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**The Inference of Phylogenetic Relationships
among the Main Lineages of Terrestrial Plants,
With Emphasis on the Positions of the Major Bryophyte Groups,
Using Small-Subunit Ribosomal RNA Gene Sequences**

Volume I

A Thesis

Submitted to the Faculty of Graduate Studies and Research

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for the Degree of

Master of Science

in Biology

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by

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ABSTRACT

The precise phylogenetic relationships among the major lineages of terrestrial plants remain uncertain. Small-subunit ribosomal RNA gene sequences were used to infer a molecular phylogeny of all major groups of land plants and contribute to the understanding of the evolutionary events underlying the transition from algal ancestors to land flora. Emphasis was placed on the phylogenetics of lower plants, with a special focus on bryophyte phylogeny. A well-constructed set of rDNA sequences from representative green plant taxa was accomplished through filling out gaps in the available data by sequencing rDNA in five additional taxa from a range of lower plants. Only sequences which are relatively error free and at least 93% complete were included. The alignment of the nuclear-encoded rRNA sequences was achieved using secondary structure information. The aligned sequences were subjected to both character and distance based methods of phylogenetic analysis. The results of this study suggest that 18S rDNA sequence data do not contain sufficient phylogenetic signal to resolve the relationship of the bryophytes amongst themselves or to other lower plants. A well constructed set of representative taxa from most major plant groups coupled with complete 18S rDNA sequence, the most appropriate alignment, and exhaustive phylogenetic analyses fail still to provide clusters with reasonable confidence as measured by bootstrap or decay analysis. Evidence for the lack of phylogenetic signal to resolve deep phylogenetic branches among lower plants is demonstrated by low confidence values in the phylogenetic trees regardless of the method employed. Further support for this contention is provided with analysis using random tree-length distribution and by phylogenetic analyses with different categories of sites classified according to substitution rate calculations. The graphical display of relative substitution rate variabilities using the lower plant 18S rRNA variability model displays the extent of conservation of the 18S rRNA gene sequence among the lower plants.

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LIST OF ABBREVIATIONS

A	adenine
bp	base pairs
C	cytosine
CTAB	cetyltrimethylammonium bromide
dATP	2'-deoxyadenosine 5'-triphosphate
DCSE	Dedicated Comparative Sequence Editor
dCTP	2'-deoxycytidine 5'-triphosphate
dGTP	2'-deoxyguanosine 5'-triphosphate
dITP	2'-deoxyinosine 5'-triphosphate
DNA	deoxyribonucleic acid
dNTP	2'-deoxynucleoside 5'-triphosphate
dTTP	2'-deoxythymidine 5'-triphosphate
EDTA	ethylenediaminetetraacetic acid
G	guanine
h	hour
indels	insertions / deletions
kb	kilobase
min	minute
OTUs	operational taxonomic units
PCR	Polymerase Chain Reaction
Physcomitrella	<i>P. patens</i>
rDNA	ribosomal RNA gene
RNA	ribonucleic acid
RNase A	Ribonuclease A
rRNA	ribosomal RNA
s	second
T	thymine
TE	10 mM Tris, 1 mM EDTA, pH 8.0 buffer
Tris	tris(hydroxymethyl)aminomethane
U	uracil

1. INTRODUCTION

The reconstruction of deep phylogenetic relationships among lower land plants is necessary for elucidating important events in the evolution of green plants and especially green terrestrial plants. Plants are defined here to include a broad assemblage of photosynthetic organisms that all contain chlorophylls *a* and *b*, store their photosynthetic products as starch inside the chloroplasts, and have cell walls made of cellulose (Raven *et al.*, 1992). This assemblage is compiled from several thousand classically-defined green algae and several hundred thousand land plants.

Two major lineages of plants have been placed in a single monophyletic group, the subkingdom Chlorobionta (Bremer, 1985). This study has focused on the evolution of one of these major lineages which includes land plants and closely related 'green algae'. This lineage of green plants has been called the Streptophytes (Bremer, 1985), and consists of some organisms conventionally considered green algae plus the more familiar green plants found mostly on land. Traditionally, the green algae in this lineage are included in the Class Charophyceae, but some members of this class are related more closely to higher plants than to other members of the class (Mattox and Stewart, 1984; Mishler, 1985; McCourt, 1995; Melkonian and Surek, 1995). Specifically, *Chara* and related algae (Order Charales) and *Coleochaete* and related algae (Order Coleochaetales) are probably the closest living 'green algal' relatives of land plants.

Ultrastructural and morphological studies were the first to support the relationship of these orders of green algae to land plants (embryophytes) (Pickett-Heaps, 1975; Mishler, 1985; Graham *et al.*, 1991). These orders of green algae have been placed in the Class Charophyceae (Mattox and Stewart, 1984), and recent analyses suggests that the Charophyceae is a paraphyletic group with the orders circumscribed within it being placed at the base of the Streptophyta lineage (Bremer, 1985; McCourt, 1995; Surek *et al.*, 1995). Some of the more recent molecular studies corroborate earlier cladistic analyses based on ultrastructural and biochemical data (Mishler and Churchill, 1985) which propose that the 'charophycean' algae, specifically Coleochaetales, are closest to the embryophyte clade (Kranz *et al.*, 1995).

Although there is considerable evidence supporting the evolution of land plants from these green algal ancestors (Chapman and Buchheim, 1991; Surek *et al.*, 1993; Bhattacharya *et al.*, 1994; Mishler *et al.*, 1994; Kranz *et al.*, 1995), the precise relationships amongst major land plant groups are still unclear. The evolutionary transition from algal ancestors to land flora will be understood best with elucidation of the phylogenetics of lower plants. Bryophyte genealogy is believed to be of paramount importance in understanding this evolutionary transition (Hedderson *et al.*, 1996). Studying bryophyte phylogeny should provide insight into the evolutionary events underlying the colonization of the terrestrial habitat and, furthermore, will help with the differentiation of major lower plant lineages.

Bryophytes consist of approximately 18,000 species which are traditionally defined to three major groups; namely hornworts (Anthoceropsida), liverworts

(Hepaticopsida), and mosses (Byropsida). Recent analyses, using diverse data sets, suggest that the phylogenetic position of bryophytes is between the green algal ancestors of the land plants and the tracheophytes or vascular plants (Mishler *et al.*, 1984; Bremer, 1985; Bremer *et al.*, 1987; Raubeson *et al.*, 1992; Waters *et al.*, 1992; Garbary *et al.*, 1993; Mishler *et al.*, 1994). Over the past two decades, several phylogenetic studies have focused on the relationships among the three main groups of bryophytes and have utilized morphological, ultrastructural and molecular data (Renzaglia, 1978; Crandall-Stotler, 1980, 1986; Mishler and Churchill, 1984, 1985; Schuster, 1984; Schofield, 1985; Duckett and Renzaglia, 1988; Mishler *et al.*, 1992; Waters *et al.*, 1992; Garbary *et al.*, 1993; Mishler *et al.*, 1994; Kranz *et al.*, 1995; Bopp and Capesius, 1995, 1996; Hedderson *et al.*, 1996).

Despite these efforts, there is no clear consensus as to the relationships among the three main groups of the bryophytes. Cladistic analyses of combined morphological, ultrastructural, and biochemical data have implied that the bryophyte clade is a paraphyletic assemblage with the Byropsida a sister group to the tracheophytes, the Anthoceroopsida a sister group to the Byropsida-tracheophyte lineage, and the Hepaticopsida as basal group among the land plants (Mishler and Churchill, 1984, 1985). Investigations using characters derived from male gametogenesis suggest different relationships, in which bryophytes are a monophyletic group with Anthoceroopsida a sister group to a Byropsida-Hepaticopsida lineage (Garbary *et al.*, 1993). The bryophyte clade is proposed to share a common ancestor with the Lycopodopsida group, and this

monophyletic bryophyte-Lycopodopsida clade is a sister group to the remaining land plants.

In recent years, the sequences of the small-subunit ribosomal RNA genes have been used increasingly to investigate plant evolution (Hamby and Zimmer, 1992). Investigation of these nuclear genes has provided evidence for their potential phylogenetic usefulness at the deeper hierarchical level (Zimmer *et al.*, 1989; Kantz *et al.*, 1990; Zechman *et al.*, 1990; Hillis and Dixon, 1991; Lewis *et al.*, 1992; Waters *et al.*, 1992; Wilcox *et al.*, 1992; Ashton *et al.*, 1994). Analysis of these gene sequences has been applied to the study of bryophyte phylogeny, and preliminary results place an Anthoceropsida-Byropsida clade as a sister to the tracheophytes with the Hepaticopsida remaining basal to the other land plants (Waters *et al.*, 1992). However, the topology of Mishler and Churchill (1984, 1985) can be imposed upon the rRNA tree of Waters *et al.* (1992) by increasing tree length by a single step. Cladistic analyses of the green plants that combine and compare available morphological and molecular data sets have resulted in poorly resolved phylogenies (Mishler *et al.*, 1994). A more comprehensive analysis to determine the origin of land plants using complete small-subunit ribosomal RNA gene sequences found conflicting resolutions depending on phylogenetic methods employed (Kranz *et al.*, 1995). This study found maximum parsimony and maximum likelihood analyses implied a successive evolution from 'charophycean' algae to bryophytes, lycopods and seed plants whereas, distance matrix methods implied a separate evolution of the lycopods and seed plants by grouping the bryophytes together with the 'charophycean' algae. Conversely, more recent work of Hedderson *et al.* (1996) resolved

hornworts as the basal group among the land plants with the liverworts and mosses forming a monophyletic clade sister to the tracheophytes.

The pattern of diversification of the bryophytes is not the only aspect of the evolution of land plants that remain imperfectly known. The relationships of the morphologically and biologically diverse lineages that constitute the vascular cryptograms also remain controversial (Bremer *et al.*, 1987; Niklas and Banks, 1990; Garbary *et al.*, 1993; Stewart and Rothwell, 1993; Hasebe *et al.*, 1995; Kranz *et al.*, 1995; Manhart, 1995; Pryer *et al.*, 1995; Kenrick and Crane, 1997; Kranz and Huss, 1997). For example, there is debate as to whether the Lycopodiophyta are polyphyletic (Garbary *et al.*, 1993) or monophyletic (Niklas and Banks, 1990; Kranz and Huss, 1997). In addition, some studies suggest that the Psilotophyta form the basal lineage of vascular plant evolution (Bremer *et al.*, 1987) whereas, others suggest Lycopodiophyta fill this role (Stewart and Rothwell, 1993; Kranz and Huss, 1997). Similarly, the relationships among and placement of the Filicophyta have also been the subject of considerable controversy (Stewart and Rothwell, 1993; Hedderson *et al.*, 1996). Clearly, precise relationships among basal lineages of land plants including the bryophytes remain unclear.

Other studies using nuclear encoded rRNA gene sequences have placed in doubt previously unquestioned bryophyte genealogy. The monophyly of the Hepaticopsida is brought into dispute by suggestions that only the complex thalloid liverworts (Class Marchantiopsida) are a sister group to the tracheophytes (Chapman and Buchheim, 1991, 1992; Mishler *et al.*, 1994). The remaining hepatics comprised of simple thalloid and leafy liverworts (Class Jungermanniopsida) are proposed to form a monophyletic group

with Anthoceropsida and Byropsida or Anthoceropsida alone. More recent studies, again using 18S rRNA genes, but including only strictly bryophyte taxa, have attempted to resolve the questionable monophyly of the liverworts (Capesius, 1995; Bopp and Capesius, 1995). These studies support placing Marchantiidae at the base of the Byropsida-Jungermanniidae clade, but disagree on the monophyly of the Bryopsida lineage. Phylogenies constructed with nucleotide sequence data of chloroplast gene *rbcL* corroborate the separation between complex thalloid and leafy/simple thalloid liverworts but could not reject the monophyly of the liverworts (Lewis *et al.*, 1997).

Some aspects of the origin and evolution of land plants do appear to be approaching resolution. It is now accepted widely that the Charophyceae are a sister group to the embryophytes (land plants including bryophytes and tracheophytes) which together form a monophyletic lineage. The bryophytes occupy a phylogenetic position between the Charophyceae and the tracheophytes. The seed plants are a monophyletic group containing the monophyletic angiosperms and the gymnosperms, although the gymnosperm monophyly is still under investigation (Chaw *et al.*, 1997). Nonetheless, several issues on the origin and evolution of land plants remain unresolved. This study set out to investigate some of the issues regarding the resolution of basal lineages of land plants with a specific focus on bryophyte genealogy.

Recent advances have provided some insight into the aforementioned discrepancies among the several studies using nuclear-encoded rRNA sequence data. The quantity and quality of sequence data, the assessment of homology provided by sequence alignment, taxon sampling and methods of phylogenetic analysis all have profound effects

on the outcome of phylogenetic investigation using molecular data (Charleston *et al.*, 1994; Lecointre *et al.*, 1993; Yang, 1997). There are five basic steps in the phylogenetic analysis of DNA sequences, although all previous investigators of bryophyte phylogeny have either excluded or at least de-emphasized some of the steps. The steps involved are discussed in the following five sections and are: 1) choice of DNA sequence, 2) sampling of appropriate taxa, 3) sequence alignment 4) assessment of phylogenetic signal and 5) choosing a method of phylogenetic inference.

1.1 Nuclear-encoded rRNA sequence data

The partial or complete DNA sequences of single genes comprise the most common type of sequence sample used in molecular phylogenetic studies. Although some effort has been made to study the phylogenetic utility of particular genes (Friedlander *et al.*, 1992; Graybeal, 1994), the location and length of sequences are chosen generally on the basis of factors other than their ability to accurately represent the genome. These factors include interest in the functional characteristics of a region, its historical use in systematic studies, and several technical considerations which are often unrelated to its ability to reconstruct whole-genome relationships. Indeed, how well a gene represents the entire genome is difficult to evaluate *a priori*.

The potential usefulness of rRNA for phylogenetic inference has been established and merits for its use include universality, functional constancy, ease of identification and isolation, and apparent lack of horizontal gene transfer (Olsen, 1988). The development of 'universal' rDNA-specific primers has revolutionized the process of rRNA gene

sequence determination (Sogin, 1990; White *et al.*, 1990). In addition, DNA sequencing technologies have progressed to a point where complete or nearly complete nucleotide sequence data for rRNA genes can be obtained readily. One of the greatest assets of rRNA-based phylogenetic analysis is the availability of pre-determined ribosomal RNA gene sequences from several public databases (Benson *et al.*, 1997; Madaik *et al.*, 1997; Van de Peer *et al.*, 1997).

The rRNAs can be identified by their typical eukaryotic sedimentation values: 5S, 5.8S, 18S, and 28S. In eukaryotes, two internal transcribed spacers designated ITS-1 and ITS-2 separate the 18S, 5.8S and 28S genes. The 18S rRNA gene was chosen for use in this study based largely on the availability of 'universal' primers, and the large body of previously determined plant 18S rDNA sequence data. Previous studies into the elucidation of bryophyte phylogeny were able to utilize only partial sequence data of genes for 18S rRNA (Waters *et al.*, 1992; Mishler *et al.*, 1992; Mishler *et al.*, 1994). Only recently has complete or nearly complete sequence data for bryophytes been available for use in phylogenetic analyses (Bopp and Capesius, 1995, 1996; Capesius, 1995; Kranz *et al.*, 1995; Hedderson *et al.*, 1996). This investigation included an 18S rRNA gene in the data set only if a minimum of 93% of its sequence had been determined.

1.2 Taxon sampling

The previous section addressed the assumption that the sampled nucleotide sequences are representative of the genomes from which they are drawn. Similarly, inferred phylogenetic relationships are presumed to represent those of the organisms

involved. Although, in general, sampling properties of DNA sequence data are assessed quite well, species sampling has received little to no attention. When a species is sampled, it is assumed implicitly that its sequence is representative of the sequences in the supposed monophyletic group of which the chosen species is a member. In practice, however, it has been shown that radically different phylogenies can be obtained if the same presumed monophyletic groups are sampled differently (Lecointre *et al.*, 1993). Several studies have shown that significant variations in robustness and topology of trees obtained depend on species representation of the taxa (Patterson, 1989; Hendriks *et al.*, 1990; Adoutte and Philippe, 1993; Lecointre *et al.*, 1993; Charleston *et al.*, 1994). The impact of species sampling can be minimized with sampling several species per presumed monophyletic group (Lecointre *et al.*, 1993).

Incomplete taxon sampling is likely to explain some of the discrepancies among the different studies of bryophyte phylogeny. Some studies into the bryophyte phylogeny have restricted the taxa to strictly bryophyte species (Capesius, 1995; Bopp and Capesius, 1995, 1996). All other studies looking at deeper phylogenetic relationships have had the foresight to include other major lineages but the inclusion of representative taxa is generally weak (Renzaglia, 1978; Crandall-Stotler, 1980, 1986; Mishler and Churchill, 1984, 1985; Schuster, 1984; Schofield, 1985; Duckett and Renzaglia, 1988; Mishler *et al.*, 1992; Waters *et al.*, 1992; Garbary *et al.*, 1993; Mishler *et al.*, 1994; Kranz *et al.*, 1995; Hedderson *et al.*, 1996). For example, the most complete taxa set used in phylogenetic investigation of bryophytes included 18 bryophyte taxa with only two hornworts and six taxa from fern and seed plants representing all tracheophytes

(Hedderson *et al.*, 1996). Kranz *et al.* (1995) propose a bryophyte phylogeny with only three representatives from the group and none from the hornworts. However, this study does include representative taxa from all major lineages of plants and is comprised of 54 species spanning seven phyla and fifteen families.

1.3 Sequence alignment and secondary structure information

One of the essential steps in the reconstruction of phylogenetic history is the determination of character homology among the taxa being studied. One of the basic principles of systematics is that only homologous characters can provide meaningful markers of genealogical descent. Consequently, the accuracy of a phylogeny from molecular data is critically dependent on the accuracy of sequence alignment. Hence, mistaken hypotheses of character-state homology are a primary source of error in evolutionary studies (Morrison and Ellis, 1997). Molecular sequence data are exceptionally susceptible to these errors since assessment of homology involves the alignment of nucleotides or amino acids with one hypothesis of homology for each position. Positional homology in orthologous sequences can fall into one of three categories; 1) identical character states, 2) substitutions representing point mutations or 3) insertions/deletions (indels). The positioning of indels is the most problematic aspect of sequence alignment which becomes more acute with divergent taxa. The purpose of sequence alignment is to introduce gaps into the sequences only when they truly represent insertion or deletion events during the course of evolution.

Insertions and deletions are common in genes for rRNA (Kjer, 1995) and require the introduction of gaps into sequences for the purpose of alignment. Until recently, studies of bryophyte phylogeny relied on computer programs to align rRNA gene sequences, often followed by manual adjustments and exclusion of regions containing ambiguous alignment (Chapman and Buchheim, 1992; Mishler *et al.*, 1992; Ashton *et al.*, 1994; Hedderson *et al.*, 1996). Mishler *et al.* (1994) stated simply that their alignments of molecular data were done 'by eye'. Some authors of studies in bryophyte phylogeny even fail to mention this very important step, perhaps providing insight into their assessment of this critical first step of systematics (Capesius, 1995; Bopp and Capesius, 1995, 1996). The central assumption for both the manual and computer-assisted alignments referred to above is that they maximize nucleotide identity. The computer programs for sequence alignment function by invoking gap penalties. These gap penalties are assigned arbitrarily by the investigator since neither the causes nor consequences of these gaps are understood (Kjer, 1995). Arguably, the assignment of gap penalties may be appropriate for polypeptide encoding sequences but is not so for rRNA genes. Some regions of rRNA genes are extremely intolerant to changes in length while other regions vary without any apparent constraints, suggesting that each region and possibly each position within the rRNA molecule would be better served with its own individual 'gap penalty' (Kjer, 1995). In addition, neither these manual nor computer-assisted alignments optimize the alignment of homologous positions within structurally conservative stems that lack significant nucleotide identity. For example, a point mutation has at least 50% likelihood that its new nucleotide state is identical to one of its nonhomologous neighbors.

If this mutation occurs near a gap, it could be systematically misaligned when alignments are based strictly on maximizing nucleotide identity.

Small-subunit rRNA is a molecule for which there is an *a priori* biological model of secondary structure (Gutell *et al.*, 1994). In addition, primary and secondary structures are well conserved even among very divergent taxa (Hillis and Dixon, 1991). Although the secondary structure of rRNA is not identical across all taxa, it is still more highly conserved than the nucleotide sequences (Kjer, 1995). Recent advances in homology assessment suggest that secondary structure information can be used to enforce sequence alignment (Kjer, 1995; Hickson *et al.*, 1996). Since the sequences of the rRNA genes (rDNA) are constrained by the secondary structures of their products, this allows knowledge of the secondary structure to be used for the alignment of the rDNA sequences. Furthermore, the helix (or stem) and single-stranded (or loop) regions can be treated as separate divisions of nucleotide site change provided the paired bases of the helices result from compensatory mutations (Vawter and Brown, 1993; Muse, 1995).

There are two main methods for predicting secondary structure of ribosomal RNA molecules: phylogeny and energy minimization studies (Jaeger *et al.*, 1990). Energy minimization utilizes thermodynamic parameters and computer algorithms which determine minimum and near minimum free energy folding of RNA. Predictions based on phylogeny relies on primary structure alignments with subsequent folding of several sequences into similar structures for functionally analogous RNA. Certainly, structural and evolutionary studies are complementary since the derivation of evolutionary trees requires a dependable sequence alignment as a starting point, and the establishment of

such an alignment is enhanced by the knowledge of secondary structure landmarks.

Secondary structure landmarks include boundaries of helices and loops and the existence of compensating substitutions in complementary strands.

Comparative sequence analysis of RNA structure is based on the simple principle that homologous RNA molecules will adopt the same secondary and tertiary structures with different primary sequences. Practically, comparative studies identify secondary structure base pairings by finding compensatory base changes or covariations in alignments of homologous sequences. This approach has been successfully applied to several classes of RNA molecules (Woese *et al.*, 1980; Glotz *et al.*, 1981; Zwieb *et al.*, 1981; Noller *et al.*, 1981; Maly and Brimacombe, 1983; Woese *et al.*, 1983; Gutell, 1996). Analysis of 16S and 23S rRNA sequences have revealed that many positions are restricted to certain types of pairing sequences, either subsets of the four Watson-Crick pairings or non-canonical base-pairs, such as A:C and G:U or A:G and G:A (Gutell, 1996). The majority of restricted variations in rRNA base pair sequences are of the purine:pyrimidine type (Gautheret and Gutell, 1997).

It has become evident to investigators of phylogenetics using rRNA gene sequence data that secondary structure information is indispensable (Morrison and Ellis, 1997). Certainly, erroneous homology statements due to alignment inaccuracies are as problematic as other well-known problems associated with sequence length and tree-inference methods (Russo *et al.*, 1996). Previous studies on the phylogeny of bryophytes have failed to consider structural aspects of rRNA when aligning their sequences and discrepancies amongst their trees are probably attributable largely to the

alignments employed (Chapman and Buchheim, 1992; Mishler *et al.*, 1992; Ashton *et al.*, 1994; Capesius, 1995; Bopp and Capesius, 1995, 1996; Hedderson *et al.*, 1996). This study utilizes both primary and secondary structure information for homology assessment.

1.4 Assessment of phylogenetic signal and variability

Once sequences have been aligned, some assessment of the presence of phylogenetic signal is necessary. Phylogenetic signal is the information used to infer phylogenetic history. There is obviously no purpose in additional analysis if all sequences are identical; likewise, extremely divergent sequences may be randomized with respect to phylogenetic history giving a phylogenetic hypothesis that might as well have been selected at random. The reasonable alignment of DNA sequences is not sufficient to justify the use of the sequences in phylogenetic analysis. Analysis of most aligned sequences may produce an optimal tree with a given method, but without some assessment of phylogenetic signal, there is insufficient reason to expect that the optimal tree is a good estimate of phylogeny (Hillis *et al.*, 1993).

DNA sequences compared among organisms may contain phylogenetic signal, or they may be randomized with respect to phylogenetic history. To order to proceed with an appropriate analysis of a particular data set, phylogenetic signal needs to be distinguished from random noise. This is especially important when considering the fact that random data frequently produce a single most-parsimonious tree often considerably shorter than the second-best alternative (Hillis and Huelsenbeck, 1992). Mistakenly,

several investigators cite confidence in their molecular data by quoting high decay indices on their single most-parsimonious tree without first establishing phylogenetic signal (Mishler *et al.*, 1994).

In addition, the usefulness of small-subunit rRNA gene sequence data for phylogenetic inference has been determined largely on groups of taxa other than lower land plants. Until recently, the greatest justification for use of rRNA had been successful analyses of bacterial evolution (Woese, 1987). The usefulness of this gene for phylogenetic inference of lower land plant relationships has not been fully justified but rather implied from other studies with bacterial (Woese, 1987; Turner *et al.*, 1989), animal (Hillis and Dixon, 1989; Larson and Wilson, 1989; Sogin *et al.*, 1989) and some higher plant (Gouy and Li, 1989; Hamby and Zimmer, 1992; Wolfe *et al.*, 1989) taxa. Plant DNA sequences shows extensive homoplasy in comparison to those derived from animals (Syvanen *et al.*, 1989) and, furthermore, it has been established that rates of synonymous substitution vary among evolutionary lineages of plants (Bousquet *et al.*, 1992; Gaut *et al.*, 1996; Eyre-Walker and Gaut, 1997). These facts justify the need to determine independently the utility of 18S rDNA for phylogenetic inference of relationships among lower land plants and specifically of those involving bryophytes.

One method of detecting the presence of phylogenetic signal in a given data set is to examine the shape of the tree-length distribution for a random subset of all trees resulting from a parsimony analysis (Hillis and Huelsenbeck, 1992; Hillis *et al.*, 1993). Only a strongly left-skewed tree-length distribution indicates good phylogenetic signal. The skewness test statistic, g_1 , can provide a rapid and efficient test for significant

structure in data matrices for phylogenetic analysis (Hillis and Huelsenbeck, 1992). Data sets with no correlated characters produce distributions that are close to symmetrical and suggest phylogenetic signal is not sufficient to warrant further phylogenetic analysis of the data.

The determination of the presence or absence of phylogenetic signal can be complemented by characterizing the extent and location of nucleotide variability. Several studies have shown that most constrained molecules, such as rRNAs, exhibit a considerable variation in substitution rate (Uzzell and Corbin, 1971; Golding, 1983; Holmquist *et al.*, 1983; Van de Peer *et al.*, 1993; Wakeley, 1993; Sullivan *et al.*, 1995; Yang, 1995). Consequently, the probability of these molecules sustaining a substitution is different for different sites, possibly because selective and functional constraints are not identical in different parts of the molecule. Several studies have revealed that rate variation among sites is a very important issue in the framework of phylogenetic inference. These studies have shown that ignorance of this variation leads to underestimation of transition/transversion rates (Wakeley, 1994), evolutionary distances and branch lengths (Jin and Nei, 1990; Ota and Nei, 1994; Tateno *et al.*, 1994). The underestimation of branch lengths can affect profoundly tree reconstruction by distance matrix methods (Olsen, 1987; Jin and Nei, 1990; Tateno *et al.*, 1994; Van de Peer *et al.*, 1996). The ignorance of site variability has especially drastic effects on the construction of universal trees (Yang and Roberts, 1995). To many systematists interested in phylogeny, the identification and acknowledgment of partitions in the sequence data provided by assessment of site variability may seem irrelevant at best. However, ignoring

details of evolutionary processes presupposes falsely that phylogeny reconstruction methods are robust to such influences (Bull *et al.*, 1993).

A recently developed method for estimating the variability of nucleotide sites in a sequence alignment was used in this study (Van de Peer *et al.*, 1996c). This method defines the variability of each nucleotide site as its evolutionary rate relative to the average evolutionary rate of all the nucleotide sites of the molecule. This variability measurement was used in a distance-based phylogenetic method (Van de Peer *et al.*, 1996c) which provided a precise estimation of evolutionary distances. The site variability information was also used to divide the alignment positions into a number of sets of increasing relative substitution rate. This provided a means to determine which nucleotides generate phylogenetic signal for phylogeny reconstruction and exactly where the signal allows resolution in the tree. In addition, site variabilities were calculated from representative taxa of all major lower plant lineages and were used to construct a quantitative map of nucleotide substitution rates in lower plant nuclear small-subunit rRNA. The variability map for lower plant 18S rRNA was constructed by superimposing on to the secondary structure model of *Physcomitrella patens* 18S rRNA the calculated site variabilities where the variability of each nucleotide site is indicated by means of a colored dot (Van de Peer *et al.*, 1996a). This colored map provides a detailed and quantitative description of positional variability with respect to secondary structure of the molecule.

1.5 Phylogenetic analysis

It is not the intention or purpose of this thesis to educate the reader on the different methods of phylogenetic inference. However, a brief synopsis of the phylogenetic methods employed in this study is justified. A phylogenetic tree is a form of a graph composed of nodes and branches. The nodes in the trees presented in this thesis represent the 18S rRNA gene and the branches represent the topological relationships between the nodes. Nodes can be divided into internal and external types where the latter are also called operational taxonomic units (OTUs). All trees in this thesis are rooted by identifying a special node, called the root, using the outgroup method. The outgroup method determines of the relationship of n sequences by adding one or more sequences that are known to be outside the clade of n sequences or basal in phylogenetic lineage. The number of possible topologies for an unrooted strictly bifurcating tree with the 67 OTUs examined in this thesis is approximately $2.1253 * 10^{109}$. Clearly, the application of computers is warranted and several software packages that are available for phylogenetic investigation have been employed in this study. The number of trees is significant since it dictates the use of heuristic methods and often the choice of phylogenetic method and the number of OTUs is limited by computational considerations (CPU time for some of the methods engaged in this study required several weeks for calculation).

Several methods have been proposed for building a phylogenetic tree from observed data. Tree-building methods can be divided into two groups in terms of the

type of data they use: distance-based methods and character-based methods. Distance-based methods use distance matrices which consist of all possible pairwise distances, whereas character-based methods use an array of character states. The distance-based analyses used in this study incorporate the neighbor-joining method (Saitou and Nei, 1987) which is applied to distance matrices calculated by using various parametric models (Jin and Nei, 1990; Jukes and Cantor, 1969; Kimura, 1980; Tajima and Nei, 1984; Van de Peer *et al.*, 1996c). The character-based methods used in this study consist of the maximum parsimony method (Fitch, 1971) and the maximum likelihood method (Felsenstein, 1981).

In comparative analysis of homologous DNA sequences, nucleotide substitution is commonly assumed to follow a homogeneous Markov process (Yang, 1994). The Markov process is specified by a rate matrix, Q , whose elements represent instantaneous substitution rates among the four nucleotides. For mathematical simplicity and ease of computation, normally extra restrictions have been placed on the structure of Q , leading to various parametric models. These parametric models are assumed to correct for superimposed mutations when calculating distance values between sequences. The parametric models or distance methods fit a tree to a matrix of pairwise distances between the sequences - the rate matrix, Q . Each distance is based on the fraction of positions in which two sequences differ and is defined as dissimilarity, S . This dissimilarity is actually an underestimation of the true evolutionary distance since some of the sequence positions likely represent multiple events. Furthermore, since mutations are fixed in DNA sequences, there is an increasing chance with time that mutations occur at the same

nucleotide position, creating a superimposed mutation. Therefore, different parametric models each try to estimate the number of unseen mutations with distinct assumptions about the nature of evolutionary changes. Such models can be used to construct estimates of evolutionary distances in pairwise sequence comparisons and are used also in maximum likelihood joint comparisons of all sequences. The evolutionary distance between two sequences is then assumed to be proportional to the time that separates them.

The neighbor-joining method of phylogenetic tree construction (Saitou and Nei, 1987) takes the distance matrix built by one the parametric models and modifies it. This modified distance matrix is constructed so that the separation between each pair of nodes is adjusted on the basis of their average divergence from all other nodes. This has the effect of normalizing the divergence of each OTU for its average clock rate (Swofford and Olsen, 1990). The neighbor-joining algorithm then joins the least distant pair of OTUs as defined by the modified distance matrix into a new OTU. This new OTU is added to the tree while the replaced OTUs and their respective branches are removed from the tree. This process converts the newly added OTU into a terminal node on a tree of reduced size. At each stage of the process, two terminal OTUs are replaced by a new one. The process is complete when only two OTU remain, separated by a single branch. The neighbor-joining approach does not depend on ultrametric data (Swofford and Olsen, 1990) which is appropriate for 18S rDNA and its unequal rates of evolution across the broad selection of taxa used in this study.

Character-based methods treat each substitution separately rather than reducing all of the variation to a single divergence value. The character-based method of maximum

parsimony (Fitch, 1971) partitions similarities between DNA sequences on a character-by-character basis. Alternative trees are evaluated, one character at a time, to determine how many evolutionary events they each require. By the criterion of parsimony, the 'best' or most parsimonious tree is the one requiring the fewest total events. Parsimony is the principle of logic that the simplest explanation should be preferred over the more complex. In the context of phylogenetic investigation, the most parsimonious tree is the one requiring the fewest evolutionary events to explain the differences between DNA sequence data. Maximum likelihood (Felsenstein, 1981) is the other character-based method used in this study. Maximum likelihood is a statistical model which considers each character separately for the likelihood of observed change given a particular topology and model of molecular evolution.

Once the topology is constructed, it is necessary to evaluate the reliability of the tree by assessing the confidence in the robustness of the phylogeny provided by the sequence data. Bootstrap analysis can be used to place confidence intervals on phylogenies and is a type of statistical analysis to test the reliability of certain branches in the evolutionary tree (Felsenstein, 1985; Swofford and Olsen, 1990). Bootstrapping involves resampling the data with replacement to create a series of bootstrap samples of the same size as the original data. Nucleic acid sequences provide resampled data in the form of nucleotides of a sequence, while the statistical significance of a cluster is given by the fraction of trees, based on the resampled data, containing that cluster. Bootstrapping is the most popular method used to evaluate tree robustness and has been extensively tested and assessed as an estimator of robustness (Hillis and Bull, 1993; Felsenstein and

Kishino, 1993). Problems with bootstrapping stem from several assumptions that the bootstrap places on the data. Two of these assumptions have been identified as most troublesome when dealing with molecular data (Hillis and Bull, 1993). These are: 1) that characters are independent and 2) that characters are a representative sample from an underlying universe of character data. DNA sequence character data are neither independent (Gatesy *et al.*, 1993) especially ribosomal RNA genes (Wheeler and Honeycutt, 1988) nor a representative sample of the universe of character data (DeSalle *et al.*, 1994). Nevertheless, bootstrapping can be applied, provided the investigator understands these caveats when interpreting bootstrap values which underestimate consistently the accuracy of a phylogenetic hypothesis.

To ensure the confidence in clusters based on bootstrap values was not being misinterpreted, another method was employed for testing the statistical significance of a particular branching pattern of a phylogenetic tree. Decay analysis is an alternative method for evaluating support for clades and has been well studied for its merits (Bremer, 1988; Donoghue *et al.*, 1992; Graham *et al.*, 1991; Källersjö *et al.*, 1992; Mishler *et al.*, 1991). Decay analysis works by obtaining the strict consensus of trees that are one step longer than the most parsimonious tree(s), two steps longer, and so on until all resolution is lost. The 'decay index' provides the confidence estimate and is defined as the number of steps maximum parsimony must be relaxed to cause a particular clade to lose its support (Mishler *et al.*, 1991).

It is now evident that different methods used in phylogenetic analysis may not give the same branching pattern in phylogenetic trees (Barnabas *et al.*, 1995; Kranz *et al.*,

1995; Yang, 1997). Hence, it is necessary to construct trees by more than one type of method and make cooperative use of information to arrive at a possible species phylogeny (Kim, 1993). It is this author's contention that none of the distance-based or character-based methods, when appropriately employed in phylogenetic inference, warrant complete condemnation or approval. Instead, a comparative approach, as taken in this study, will avoid making any unduly algorithm-weighted or philosophy-laden phylogenetic judgement.

There must be certain *a priori* knowledge to make justifiable claims for one particular treatment to be unequivocally the best in any particular situation (Avice *et al.*, 1994). Most investigators of phylogeny using a specific set of sequence data do not have precise knowledge of DNA evolutionary rates, extent of variability, or processes underlying character transformation. Consequently, determination of which phylogenetic method is the most apt for the data is not possible since the conformity of the data to a particular set of assumptions underlying each phylogenetic method is uncertain. Therefore, it is reasonable to compare results of several analyses and subsequently to place greater emphasis and confidence in phylogenetic outcomes derived from multiple approaches than in those which, idiosyncratically, are generated by only one or a few methods (Avice and Nelson, 1995). As stated by Avice *et al.* (1995); "...outcomes robust to alternative analyses and databases should be less controversial than those which are strongly analysis- or data-dependent", which is in agreement with Felsenstein's (1982) sentiment; "It is essential that we not adopt a single method [of phylogenetic inference] as a universal panacea." This study employed two distinct character-based phylogenetic

approaches, maximum parsimony and maximum likelihood, and several distance-based phylogenetic algorithms, against nine data sets for a total of 40 treatments overall.

1.6 Purpose and strategy of investigation

Previous investigations into the evolution and origin of lower land plants have failed to address thoroughly one or more of the aforementioned considerations. This study will attempt to resolve issues of bryophyte phylogenetics with all the elements of phylogenetic analyses addressed properly. The phylogenetic approach taken in this thesis, which in most contexts is appropriately conservative, was to apply a variety of conceptually and operationally distinct phylogenetic algorithms to the well-aligned, nuclear-encoded, small-subunit rRNA gene sequence data-set, coupled with a complete cross section of representative taxa from all major lineages of land plants.

This study expects that the putative secondary-structure model employed here is likely to have produced a multiple-sequence alignment that is closer to the true alignment by evidently aligning homologous nucleotides. The usefulness of the 18S rDNA gene sequence for deeper-level phylogenetic inference of lower plant evolution was addressed by examining the data set for the amount, location and level of phylogenetic signal. Furthermore, the higher-order structures, coupled with variability information presented here using substitution-rate calculations, provide a powerful way to identify phylogenetically important elements in the 18S rRNA molecular structure.

2. MATERIALS AND METHODS

2.1 Taxa and sequence data

The sequence data used in this study are derived from 18S rRNA genes in which a minimum of 93% of the nucleotides has been determined. Furthermore, only sequences which can be assumed on the basis of reasonable criteria to be error-free, have been utilized. The data comprise 67 unique sequences from 57 species of plants spanning seven phyla and fifteen families, seven species of green algae spanning two phyla and five families and three species of fungi belonging to three families. The list of taxa with classification and GenBank accession numbers are presented in Table 2 (see vol. II, p. 32).

The list of taxa contains five original sequences which include two moss species:

Physcomitrella patens, *Sphagnum palustre*; two hornworts: *Megaceros aenigmaticus*, *Notophylas breutelii*; and a fern: *Pteris vittata*.

The taxa set designations are outlined in Table 3 (vol. II, p. 35). Taxa set A is the original collection of 41 18S rDNA sequences. Taxa set B contains the 18S rDNA sequences of 67 taxa which includes all the sequences of taxa set A and 27 additional taxa. Taxa set C is a subset of taxa set B and consists of 43 18S rRNA gene sequences representing strictly lower plant taxa.

2.2 DNA isolation

Physcomitrella total genomic DNA was extracted from approximately 4 g (fresh weight) of gametophytic tissue grown on solid medium covered with cellophane overlays

(Ashton *et al.*, 1985) by the CTAB method of Doyle and Doyle (1990) after removal of excess water from the tissue by gentle vacuum filtration and grinding to a powder in liquid nitrogen. Using this procedure, about 25 µg of DNA were isolated per gram (wet weight) of moss. RNA was digested using RNase A (10 µg/ml) followed by precipitation of the DNA using a standard procedure, outlined by Doyle and Doyle (1990), and redissolving in TE. DNA concentrations and purity of the solutions were estimated from the absorbances at 260 nm and 280 nm of diluted samples and also from ethidium bromide fluorescence of electrophoresed samples.

Pteris vittata spores were collected in Barbados by Dr. N. Ashton and gametophytic tissue was provided by Ryan McDonald. Whole genomic DNA was extracted in a similar fashion to the procedure outlined above.

Total genomic DNAs from *Sphagnum palustre*, *Megaceros aenigmaticus*, and *Notophylas breutelii*, which had been purified by CsCl centrifugation and quantified, were gifts from Dr. K.S. Renzaglia (Southern Illinois University).

2.3 DNA amplification

2.3.1 Oligonucleotide synthesis

Oligonucleotides for use as primers for amplification and cycle sequencing of 18S rDNA fragments were synthesized using an Applied Biosystems 391 DNA Synthesizer PCR-MATE (Perkin-Elmer) in the 'Trityl-off' mode. In this mode, the dimethoxytrityl group at the 5' terminal of the oligonucleotide is removed at the end of the synthesis. The

oligonucleotides were cleaved from the support by incubation at room temperature for 2 h with fresh concentrated ammonium hydroxide, followed by deprotection with ammonium hydroxide at 55°C for 20–40 h. Each cleaved and deprotected oligonucleotide was dispersed into small aliquots, evaporated to dryness and stored dry or re-dissolved in a small volume of sterile distilled water and stored at -20°C. Primer concentrations in aqueous stock solutions were calculated from absorbances of diluted samples measured at 260 nm. Unpurified primers were used for both PCR and sequencing reactions.

The primers used for DNA amplification and cycle sequencing of three regions, collectively comprising approximately 94% of the complete 18S rRNA gene, are described in Table 1 (vol. II, p. 2). The location of the primer binding sites relative to the 18S rRNA gene is graphically depicted in Fig. 1 (vol. II, p. 3).

2.3.2 Reaction mixtures

Sequencing templates were generated using nested and semi-nested PCR. Long PCR constituted the first round of DNA amplification to generate an approximately 1800 bp PCR amplicon under high fidelity PCR conditions. This full length fragment was generated using primers NS1 and NS8 and 500 ng of whole genomic DNA with water substituted for the template in the negative controls. Long PCR reaction mixtures using hot-start with a total volume of 100 µl consisted of 1.25 units of *Taq* DNA polymerase (Gibco), 0.28 µg of TaqStart™ antibody (Clontech), 20 mM Tris-HCl (pH 8.4), 50 mM KCl, 2.5 mM MgCl₂, 125 µM of each dATP, dCTP, dGTP and dTTP, 0.5 µM (50 pmol) of each primer (NS1 and NS8), and approximately 500 ng of genomic DNA.

Nested and semi-nested PCR reactions constituted the second round of amplification using first round products as template, with negative controls also using first round negative reaction mixtures in place of template. Second round PCR reactions using hot-start with a total volume of 100 μ l were done in triplicate, each reaction mixture comprising 2.5 units of *Taq* DNA polymerase (Gibco), 0.55 μ g of TaqStart™ antibody (Clontech), 20 mM Tris-HCl (pH 8.4), 50 mM KCl, 2.5 mM MgCl₂, 200 μ M of each dATP, dCTP, dGTP and dTTP, 50 pmol of each forward and reverse primer, and 2 μ l of first round amplification products. The primer pairs used for semi-nested PCR were NS1 & NS2, NS5 & NS8, and NS7 & NS8. The primer pair, NS3 & NS4, was used for nested PCR. These primer pairs with their respective predicted PCR amplicon sizes are described in Table 1 (vol. II, p. 2).

2.3.3 Thermal cycling parameters and PCR product detection

All DNA amplifications were carried out on a PE 9600 thermal cycler (Perkin-Elmer). All thermal cycling regimes were preceded by a 2 min, 95°C incubation period, and succeeded by an additional extension time of 10 min at 72°C immediately followed by 4°C incubation until placed in storage at -20°C. Long PCR reactions were cycled 25 times through the following temperature regimes: 94°C for 30 s, 55°C for 45 s, 72°C for 120 s. Nested and semi-nested PCR reactions were cycled for a total of 40 cycles using a 3 stage touch-down PCR thermal cycling regime. The first stage consisted of 5 cycles through the following temperature regimes: 94°C for 10 s, 64°C for 20 s, 72°C for 30 s.

The second stage consisted of 5 cycles through the following thermal cycling regimes: 94°C for 10 s, 62°C for 15 s, 72°C for 30 s. The final stage consisted of 30 cycles through the following thermal cycling regime: 92°C for 10 s, 60°C for 10 s, 72°C for 30 s.

Thermal cycled PCR mixtures (10 µl) were resolved electrophoretically on 1.6% agarose gels and visualized by ethidium bromide fluorescence. Sizes of amplified fragments were estimated by comparison with a 100 bp DNA ladder (Gibco).

2.4 DNA sequencing

2.4.1 Template preparation

Three 90 µl post-PCR reactions mixtures in which agarose gel electrophoresis had revealed a single product of expected size, were purified using the Wizard™ PCR Prep DNA purification system (Promega), producing 50 µl of solution containing the amplified rDNA in distilled water. The purified sequencing template was quantified using the DNA DipStick™ Kit (Invitrogen).

2.4.2 Cycle sequencing

The purified and quantified double-stranded PCR products were used in fluorescence-based dideoxy cycle sequencing reactions using the PRISM™ Ready Reaction DyeDeoxy™ Terminator Cycle Sequencing Kit (Applied Biosystems). The cycle sequencing reaction mixtures of 20 µl final volume contained the following ingredients: 100-200 ng of template DNA, 9.5 µl of terminator premix [1.6 µM A-DyeDeoxy, 94.7

μM T-DyeDeoxy, $0.4 \mu\text{M}$ G-DyeDeoxy, $47.4 \mu\text{M}$ C-DyeDeoxy, $79.0 \mu\text{M}$ dITP, $15.8 \mu\text{M}$ dATP, $15.8 \mu\text{M}$ dCTP, $15.8 \mu\text{M}$ dTTP, 168.4 mM Tris-HCl (pH 9.0), 4.2 mM $(\text{NH}_4)_2\text{SO}_4$, 42.1 mM MgCl_2 , $0.42 \text{ units}/\mu\text{l}$ AmpliTaq[®] DNA polymerase], and 3.2 pmol of primer. The sequencing primers were the same as those used in the DNA amplification reactions namely; NS1, NS2, NS3, NS4, NS5, NS7 and NS8. Individual primers with appropriate rDNA PCR products were combined in seven different sequencing reactions which facilitated the acquisition of double-stranded sequence data for the majority of the 18S rRNA gene.

Cycle sequencing reactions were carried out on a PE 2400 (Perkin-Elmer) thermal cycler. The sequencing reactions were set up on ice and placed in the thermal cycler which was preheated to $96 \text{ }^\circ\text{C}$. The thermal cycle sequencing regime consisted of 25 cycles and was as follows: $96 \text{ }^\circ\text{C}$ for 10 s, $50 \text{ }^\circ\text{C}$ for 5 s, $60 \text{ }^\circ\text{C}$ for 4 min. The reactions were held at $4 \text{ }^\circ\text{C}$ until purification which was typically performed within 1 h of the final cycle.

2.4.3 Purification of sequencing reaction products

Subsequent to cycle sequencing, the reactions were passed through Sephadex G-50 spin columns (Centri-Sep[™] columns, Princeton Separations) to remove unincorporated dye terminators. The purified reaction mixtures were spun in a vacuum centrifuge (Centrivap[®], Labconco) at room temperature until just dry (typically 5 min). The

reaction pellets were resuspended in 3 μ l of loading buffer [5 parts deionized formamide to one part 50 mg blue dextran / 25 mM EDTA (pH 8.0)].

2.4.4 DNA sequence determination

Automated sequencing was employed using the ABI PRISM™ 377 DNA Sequencer (Applied Biosystems). The resuspended reaction pellets were immediately heat shocked at 94 °C for 5 min and then transferred to a Labtop™ cooler (Nalgene) to snap cool them at -20 °C. Samples were electrophoresed at 100 nucleotides/h with 1680 volts, 150 watts (floating), and 50 mAmp (floating) through denaturing polyacrylamide gels (4% acrylamide, 5% cross-linked, 6 M urea) at a run temperature of 51 °C with a gel thickness of 0.2 mm and a well-to-read distance of 36 cm. The fluorescent images were captured with virtual filter set A and 2400 scans/h using ABI PRISM™ Collection Software v1.1.0 (Applied Biosystems). The raw sequencing data were processed using multicomponent analysis, baseline subtraction and scaling with ABI PRISM™ DNA Sequencing Analysis Software v2.1.1 (Applied Biosystems) using the SemiAdaptive BaseCalling module. All base calls were confirmed manually by visual inspection of the electropherograms.

The double-stranded DNA sequence information was assembled from consensus sequence data using the simple homology matching algorithm and contig manager of MacDNASIS® Pro Sequence Analysis Software v3.6 (Hitachi Software). Sequence data corresponding to primer binding sites were removed and replaced with the ambiguity code 'N'.

2.5 Sequence alignment

The manual sequence alignment procedure used was that described by Van de Peer *et al.* (1996c). Manual alignment was facilitated with DCSE v2.54 (De Rijk and De Wachter, 1993). The alignment defines the complete secondary structure of the 18S rRNA molecule. The alignment process was iterative, beginning with the juxtaposition of regions of extensive primary structure similarity. The alignment was then refined in regions of greater variability through the use of secondary structure information available from the SSU rRNA Database (Van de Peer *et al.*, 1997). The secondary structures were inferred by comparative analysis relying on the search for compensatory base substitutions or positional covariance (Gutell, 1996).

The DCSE sequence alignment file was imported into MacClade v3.05 (Maddison and Maddison, 1992) to facilitate data manipulation, character annotation and file conversion for various phylogenetic analysis software packages.

2.6 Tree length distribution skewness

The g_1 statistic was used to test for possible presence of significant phylogenetic signal (Huelsenbeck, 1991; Hillis *et al.*, 1993). The g_1 statistic based on the length distribution of 50,000 randomly sampled trees from a set of all possible trees was generated by the RANDOM TREES command of PAUP 3.1.1 (Swofford, 1993). The g_1 statistic was determined for both taxa set B and C.

2.7 Phylogenetic analyses

All phylogenetic analyses were applied to the entire 18S rRNA sequence excluding ambiguous alignment positions. Phylogenetic analyses were performed on two taxa sets, designated A and B as indicated in Table 3 (vol. II, p. 35). Taxa set A consisted of a subset of taxa set B comprising 41 representative taxa from all major plant groups. *Chara foetida* was used as the outgroup for all phylogenetic analyses with taxa set A. Taxa set B consisted of the entire species collection comprising 57 representative taxa from all major plant groups, seven species of green algae and three species of fungi. *Saccharomyces cerevisiae* was used as the outgroup for phylogenetic analyses with taxa set B except as otherwise indicated for parsimony and neighbor-joining analyses. The outgroup selection for taxa set B was chosen after sampling successive outgroups to ensure tree topology was not dramatically affected as described by Lecointre *et al.* (1993).

2.7.1 Character-based phylogenetic analyses

2.7.1.1 Parsimony

Phylogenetic searches under the principle of maximum parsimony were conducted using PAUP 3.1.1 (Swofford, 1993). Heuristic search strategies were employed with the MULPARS option in effect, random taxon addition with 100 replicates, and tree-bisection-reconnection (TBR) branch swapping was performed. Gaps were treated as missing data and uninformative positions were excluded from analysis. The outgroup for taxa set B was defined with 2 species of fungi: *Saccharomyces cerevisiae* and *Schizosaccharomyces pombe*. Whenever more than one most parsimonious tree was

found, the output was displayed as a 50% majority-rule consensus tree with the frequency of clustering indicated at each node as a percentage.

Confidence estimates for clustering groups were obtained by bootstrap analysis using identical heuristic search parameters. Results of the bootstrap analyses are displayed on 50% majority-rule bootstrap trees with nodes showing less than 50% confidence collapsed to yield polytomies. Computational limitations dictated the maximum number of replicates performed with each data set. Taxa set B was analyzed with 50 bootstrap replicates each with 5 random taxon additions. Taxa set A was analyzed with 100 bootstrap replicates each with 10 random taxon additions.

Additional confidence estimates provided by decay analysis (Bremer, 1988) were obtained from the taxa set A parsimony analysis using TreeRot software (Sorenson, 1996). A strict consensus tree was generated from the most parsimonious trees found during the heuristic search with taxa set A. TreeRot was used to generate a PAUP command file to examine each node on the strict consensus tree using heuristic search parameters (MULPARS option in effect, 10 random taxon additions with TBR performed and only keeping trees not compatible with constraint-tree) and a constraint statement. Decay indices were generated for all nodes of the strict consensus tree using this constraint statement approach and applied to the tree at each respective node.

2.7.1.2 Maximum likelihood

Maximum likelihood analysis was performed only on taxa set A using fastDNAML v1.1.1a (Olsen *et al.*, 1994). The analysis parameters were set to randomize the sequence

addition order. The expected transition/transversion ratio was given the value 1.71 (estimated from data set B, S.E. 0.07). The distances were calculated based on the Felsenstein's (1981) model which allows the four nucleotides to have unequal frequencies. Confidence in clusters was estimated from 10 bootstrap replicates using the same analysis parameters, with corresponding bootstrap values shown in percentage displayed on a 50% majority-rule bootstrap tree.

2.7.1.3 Quartet puzzling

Maximum likelihood analysis was also applied to both taxa sets using a fast tree search algorithm called quartet puzzling (Strimmer and Haeseler, 1996). Quartet puzzling consists of three steps; maximum likelihood, puzzling and consensus. Quartet puzzling applies maximum likelihood tree reconstruction to all possible quartets from n sequences. This reconstructs all possible quartet maximum likelihood trees. The second step called puzzling involves repeatedly combining the quartet trees to an overall tree. The final step computes the majority-rule consensus of all intermediate trees giving the quartet puzzling tree. The analysis was performed by using PUZZLE v3.0 from The PUZZLE Software web page (<http://www.zi.biologie.uni-muenchen.de/~strimmer/>) by Strimmer and Haeseler (Zoologisches Institut, Universität München, Germany). Tree reconstruction used 1000 puzzling steps with the HKY (Hasegawa *et al.*, 1985) model of substitution. The model of Hasegawa *et al.* (1985) allows both different rates for transitions and transversions and different nucleotide frequencies. All parameters including rate heterogeneity were estimated from the data sets by maximum likelihood. The estimations of support for

internal branches is calculated automatically and assigned to nodes on the tree and shown in percent.

2.7.2 Distance-based phylogenetic analyses

Distance matrices were calculated from data sets A and B, and superimposed mutations were accounted for by using several distance algorithms. The distance-based phylogenetic analysis and the construction and drawing of evolutionary trees based on the calculated distance matrices were accomplished with TREECON for Windows v1.2 (Van de Peer and De Wachter, 1994).

The transformation of sequence data into evolutionary distances is performed by the program MATRIXW of TREECON for Windows v1.2 (Van de Peer and De Wachter, 1994) which produces a matrix of dissimilarities based on $N(N-1)/2$ pair-wise comparisons between N sequences. The various distance algorithms which were implemented using the software package are outlined in subsequent sections (see 2.7.2.1 - 2.7.2.8).

The neighbor-joining method (Saitou and Nei, 1987) was implemented for tree construction using the program TREE of TREECON for Windows v1.2 (Van de Peer and De Wachter, 1994). The program ROOT of TREECON for Windows v1.2 (Van de Peer and De Wachter, 1994) was used to place a root on the initially unrooted tree topologies produced by the neighbor-joining method. Taxa set B topologies were rooted with *Saccharomyces cerevisiae*. Taxa set A topologies were rooted with *Chara foetida*.

Confidence estimates in topologies were provided by bootstrap analysis based on 100 resampled data sets and conducted using the program MATRIXW of TREECON for

Windows v1.2 (Van de Peer and De Wachter, 1994). A 50% majority-rule bootstrap tree was generated for each distance calculation algorithm. The 50% majority-rule bootstrap trees constructed with TREECON are not consensus trees from a number of bootstrap trees, as constructed in the PHYLIP package (Felsenstein, 1995). The generation of bootstrap values were achieved by selecting one sequence, from each data set, as the chosen outgroup for the bootstrap analysis. The program rooted every bootstrap tree and the original tree with this particular outgroup sequence. *Saccharomyces cerevisiae* was used to root trees for taxa set B, while *Chara foetida* was used as the outgroup sequence for taxa set A. All 100 bootstrap trees were based on the 100 resampled data sets and compared with the tree based on the original sequence alignment. The fraction of bootstrap trees containing specific clusters, as defined by the original tree, was computed. This calculation was performed for all clusters found in the original tree and is displayed as a percentage adjacent to the corresponding node.

The DRAW program of TREECON for Windows v1.2 (Van de Peer and De Wachter, 1994) was used for graphical manipulation and annotation of the trees. Taxa set B topologies were rerooted at the internode between the fungi (*Saccharomyces cerevisiae*, *Schizosaccharomyces pombe* and *Dictyostelium discoideum*) and the remainder of the taxa. The trees generated from each analysis were subjected to manual branch rotations at appropriate internodes. The branch rotations maintained similar tree order thereby facilitating tree-to-tree comparisons without altering tree topology.

The series of distance-based phylogenetic analyses on taxa set A and B using various distance calculation algorithms are summarized in Tables 8 and 9 (vol. II, p. 68-

69). The results from each analysis are depicted graphically with a set of four trees. Each set consisting of a pair of trees with distance scales and a pair of trees with bootstrap values. The first pair of trees contains branch length information with the accompanying distance scale. The second pair of trees contains topology confidence measurements depicted on a 50% majority-rule bootstrap tree. The bootstrap trees contain only the clusters maintaining 50% majority-rule such that clusters with less than 50% support are drawn as polytomies. The second tree of each pair facilitates visual inspection by using triangles which represent the most consistent monophyletic clusters (clusters of taxa that share a basal node and are retained in the majority of distance calculations).

2.7.2.1 Jukes & Cantor (1969) distance calculation

The correction algorithm of Jukes and Cantor (1969) is the simplest method for nucleotide substitution estimation. The one parameter model maintains the following assumptions; 1) that all substitutions at a given sequence position are independent, 2) that all sequence positions are equally subject to change, 3) that substitutions occur randomly among the four types of nucleotides such that there is no bias in the direction of change, and 4) that no insertions or deletions have occurred. The Jukes and Cantor (1969) model estimates evolutionary distance from observed dissimilarity based on these assumptions. The number of nucleotide substitutions per site that actually have occurred since divergence of sequences A and B (evolutionary distance D_{AB}) is estimated by:

$$D_{AB} = -\frac{3}{4} \ln\left(1 - \frac{4}{3}S\right)$$

where S (dissimilarity) is the fraction of different nucleotides between two sequences.

2.7.2.2 Tajima & Nei (1984) distance calculation

The Tajima and Nei (1984) algorithm estimates the number of substitutions based on a four parameter model, assuming a so-called 'equal input' model. In this model, the rate of nucleotide substitution is assumed to be the same. Consequently, this algorithm does not take into account unequal rates of substitution among different nucleotide pairs. The TREECON program uses an average base composition computed from all sequences analyzed (Van de Peer and De Wachter, 1994). In the correction of Tajima and Nei, the evolutionary distance (D_{AB}) is estimated by:

$$D_{AB} = -b \ln \left(1 - \frac{1}{b} S \right)$$

where

$$b = 1 - \sum_{i \in N} f_i^2$$

where S (dissimilarity) is the fraction of different nucleotides between two sequences and f_i is the frequency of the i -th type of nucleotide belonging to the set of possible nucleotide types N (=A,G,C,orT) in the sequences being compared.

2.7.2.3 Kimura (1980) distance calculation

The Kimura algorithm is a two parameter model which provides a method for inferring evolutionary distance in which transitions and transversions are treated separately (Kimura, 1980). In the correction of Kimura, the evolutionary distance (D_{AB}) is estimated by:

$$D_{AB} = -\frac{1}{2} \ln[(1 - 2P - Q)\sqrt{1 - 2Q}]$$

where P is the fraction of sequence positions differing by a transition and Q is the fraction of sequence positions differing by a transversion.

2.7.2.4 Jin & Nei (1990) distance calculation

Several studies suggest that the rate of nucleotide substitution varies approximately according to the gamma distribution (Jin and Nei, 1990; Nei, 1991). This gamma distribution is specified by a parameter 'a' which is the square of the inverse of the coefficient of variation of substitution rate (Nei, 1991). The Jin and Nei (1990) model does not assume that the rate of nucleotide substitution is the same for all sites as this assumption rarely holds for real sequences (Van de Peer *et al.*, 1993). For the Jukes and Cantor's one parameter model, the evolutionary distance (D_{AB}) is calculated as follows (Jin and Nei, 1990):

for $a=1$:

$$D_{AB} = \frac{S}{1 - \frac{4}{3}S}$$

where the parameter 'a' was set to 1 for both data sets (Kumar and Rzhetsky, 1996) and where S (dissimilarity) is the fraction of different nucleotides between two sequences.

For the Kimura's two parameter evolution model, the evolutionary distance (D_{AB}) is calculated as follows (Jin and Nei, 1990):

for $a=1$

$$D_{AB} = \frac{2P + Q}{2(1 - 2P - Q)} + \frac{Q}{2(1 - 2Q)}$$

where the parameter 'a' was set to 1 for both data sets and where P is the fraction of sequence positions differing by a transition and Q is the fraction of sequence positions differing by a transversion.

2.7.2.5 Transversions only distance calculation

Distance calculation with transversions only is suggested to be more informative by removing transition 'noise' which could block phylogenetic signal (Woese *et al.*, 1991). The evolutionary distance (D_{AB}) is then estimated by (Tajima and Nei, 1984; Swofford and Olsen, 1990):

$$D_{AB} = -b \ln\left(1 - \frac{1}{b}Q\right)$$

where Q is the fraction of transversions and

$$b = 1 - \left[(f_A + f_G)^2 + (f_C + f_T)^2 \right]$$

and $f_A + f_G$ being the fraction of purines, and $f_C + f_T$ being the fraction of pyrimidines, computed over the complete alignment.

2.7.2.6 No correction distance calculation

The sequences were analyzed also using distance calculation where no algorithms were used to correct for superimposed mutations such that dissimilarity is equal to distance. The evolutionary distance (D_{AB}) is estimated by:

$$D_{AB} = S$$

where S (dissimilarity) is the fraction of different nucleotides between two sequences.

2.7.2.7 Van de Peer & De Wachter (1994) distance calculation

The Van de Peer and De Wachter algorithm computes evolutionary distance between sequences taking into account the relative substitution rate of the individual positions in an alignment (Van de Peer *et al.*, 1996c). The estimation of relative nucleotide substitution rates and computation of the evolutionary distances taking into account rate calibration were done as follows:

For an alignment of N sequences, $N(N-1)/2$ pairwise evolutionary distances d are computed according to the equation of Jukes and Cantor (1969):

$$d = -\frac{3}{4} \ln\left(1 - \frac{4}{3} f\right) \quad (1)$$

where f is the dissimilarity or fraction of observed substitutions between two sequences.

After all pairwise distances were computed, they were classified into a number of distance intervals, and the fraction of sequence pairs which fell within a certain distance interval and were characterized by a nucleotide change was calculated. This was done for every nucleotide position and the fraction was plotted against the mean distance interval (Van de Peer *et al.*, 1996c).

A curve obeying equation:

$$p_i = \frac{3}{4} \left[1 - \exp\left(-\frac{4}{3} v_i d\right) \right] \quad (2)$$

was then fitted to these points by nonlinear regression. Equation 2 expresses the probability p_i that an alignment of position i contains a different nucleotide in two

sequences, as a function of the evolutionary distance d separating these sequences. The slope of the curve in the origin yields the specific nucleotide substitution rate v_i for the position under consideration.

At this point, the estimated nucleotide substitution rates were derived on the basis of a distance matrix computed by means of equation 1 and, therefore, are not yet optimal. This equation provides only an initial approximation of the relation between dissimilarity and distance, since it starts from the unrealistic assumption that all nucleotides have the same substitution rate (Jukes and Cantor, 1969). Therefore, after estimation of all v_i values, alignment positions were grouped into sets of similar variability producing a spectrum of relative nucleotide substitutions rates. Once the shape of the spectrum was known, the following equation was used to determine dissimilarity, f , as a function of the evolutionary distance d (Van de Peer *et al.*, 1996c):

$$f = \frac{3}{4} \left\{ 1 - \exp \left[-\frac{4}{3} p \ln \left(1 + \frac{d}{p} \right) \right] \right\} \quad (3)$$

The value of the parameter p depends on the shape of the substitution rate spectrum.

The inverse of equation 3 gives:

$$d = p \left[\left(1 - \frac{4}{3} f \right)^{-\frac{3}{4p}} - 1 \right] \quad (4)$$

The equation 4 calculates a more accurate conversion of dissimilarity into distance than equation 1. Equation 4 was used to convert pairwise sequence dissimilarities into evolutionary distances. Then relative substitution rate of each alignment position was estimated again on the basis of these new evolutionary distances, as described above, and

a new spectrum of evolutionary rates is derived. This iterative process was repeated several times until the nucleotide substitution rates, v_i , stabilized. Three iterations were found empirically to be sufficient for both taxa set A and B.

2.7.2.8 No correction distance calculation with compartmentalization

The variable nucleotide positions in taxa set B were partitioned equally into five categories (1 through 5) based on their relative nucleotide variabilities (v_i). The categories were developed by determining first v_i for each nucleotide position as previously described by Van de Peer *et al.* (1996c). The v_i values were calculated using the program MATRIXW of TREECON for Windows v1.2 (Van de Peer and De Wachter, 1994). The relative nucleotide substitution rates were estimated for each of the alignment positions that were not conserved absolutely and that contained a nucleotide in at least 25% of the aligned sequences. The output file generated by MATRIXW was imported into a computer spreadsheet program (Excel v5, Microsoft) for manipulation and sorting.

Absolutely conserved alignment positions were assigned to category '6'. All alignment positions which contained nucleotides in less than 25% of the aligned sequences were placed in the category '0'. All nucleotide positions which contain alignment ambiguities were categorized in '-1'.

The distance calculation made without compensation for superimposed mutations was described above. The distance calculation was performed on a series of subsets of the original data set B. Each subset or compartment included only nucleotide positions which fitted within defined categories. Three compartments were formed as follows: A)

categories 0,1,2,3,4 B) categories 0,1,2 and C) categories 2,3,4,5,6. This compartmentalization provided subsets of alignment positions representing data of sorted variability from the taxa set B alignment. These subsets were subjected to phylogenetic analysis using distance-based calculations with no algorithmic correction for superimposed mutations.

2.7.2.9 Accounting for insertions & deletions

All distance calculations were performed also with insertions and deletions taken into account. This calculation was performed as an option using the program MATRIXW of TREECON for Windows v1.2 (Van de Peer and De Wachter, 1994). The MATRIXW program provides output that was used as an input file for the neighbor-joining tree construction method. For example, the evolutionary distance according to the Jukes and Cantor (1969) model is then calculated as (Van de Peer *et al.*, 1996c):

$$D_{AB} = -\frac{3}{4} \ln \left[1 - \frac{4}{3} \left(\frac{S_U}{I + S_U} \right) \right] \left[1 - \frac{G}{T} \right] + \frac{G}{T}$$

where I is the number of identical nucleotides, S_U is the number of positions showing a substitution and G is the number of gaps in one sequence with respect to the other. T is the sum of I, S and G. The first term of the equation accounts for substitutions and comprises the Jukes and Cantor correction factor for multiple mutations per site. The second term accounts for deletions and insertions. A row of adjacent gaps is treated as one gap, regardless of its length. Ambiguities (N) are not taken into account.

2.7.3 Overall strict consensus tree

The overall strict consensus tree represents absolute corroboration of clusters in the bootstrap trees which were generated from all tree construction methods.

Dendrograms for taxa set A and B representing a strict overall consensus tree were constructed by visual inspection of all 50% majority-rule bootstrap trees. Only clusters which are maintained strictly in the bootstrap trees from all phylogenetic analyses are depicted. Any variation of taxa clustering among the phylogenetic approaches resulted in those affected taxa being displayed as polytomies.

2.8 Secondary structure

The secondary structure model for *Physcomitrella* 18S rRNA was drawn clockwise in the 5' → 3' direction. Helix numbering is according to Van de Peer *et al.* (1997). The primary structure data for *Physcomitrella* was collected from GenBank accession X80986. The secondary structure information along with helix numbering was extracted from the DCSE alignment file described above. The model was constructed manually using a computer graphics program (FreeHand v7.0, Macromedia) with some sections preassembled using a RNA secondary structure computer program (loopDloop, Gilbert, 1992).

2.9 Variability map

A secondary structure model for lower plant small-subunit rRNA was generated by superimposing relative substitution rate variabilities on the 18S rRNA secondary structure model of *Physcomitrella* (vol. II, p. 57). The relative substitution rates (v_i) were calculated using 43 taxa representing all major groups of lower land plants, designated taxa set C and are listed in Table 3 (vol. II, p. 35). The positions belonging to five sets of average variability based on a quantitative estimate of relative substitution rates are indicated by different colors. These colors in order of most variable to least are red, orange, yellow, green and blue. The five categories represented by these colors were assigned by equally partitioning the alignment positions for which quantitative estimates of relative substitution rates could be calculated. The absolutely conserved positions ($v_i = 0$) are indicated in purple. Sites colored white belong to areas that are highly variable but are deleted in at least 75% of the sequences thereby not allowing accurate measurement of their relative evolutionary rate. Gray sites indicate alignment positions in which primary and secondary structure information did not provide enough information to allow unequivocal alignment.

3. RESULTS AND DISCUSSION

3.1 Nuclear-encoded rRNA sequence data acquisition

3.1.1 Species sampling

Green plant 18S rDNA sequence data from the several public databases provided ample representatives from most major monophyletic lineages, although, at the outset of this project, there was significantly poorer representation in a few groups. In the Bryophyta, several 18S rDNA sequences from the Hepaticopsida and Bryopsida were available, but sequence data for only one species from Sphagnopsida and two species from the Anthocerotopsida were listed. In addition, Filicophyta representation was relatively sparse.

To reduce the impact of species sampling artifacts upon inferred phylogenies, it was imperative that these putative monophyletic groups were represented properly. The Filicophyta group was strengthened by addition of another fern, *Pteris vittata*. The Anthocerotopsida class was augmented with two additional species of hornworts, *Megaceros aenigmaticus* and *Notophylas breutelii*. The Sphagnopsida class was enhanced by the addition of *Sphagnum palustre*. *P. patens* was added to the Bryopsida class, principally to determine the placement of this moss species which is used extensively in developmental studies and is becoming a quite important plant model system (Ashton *et al.*, 1993).

3.1.2 DNA amplification

The results of the first round of PCR, which generated NS1-NS8 PCR products used subsequently as templates for the semi-nested and nested PCRs, is shown in Fig. 2 (vol. II, p. 5). The resulting PCR amplicons produced the expected fragment size of approximately 1800 bp without any nonspecific amplification products. Fig. 3 (vol. II, p. 7) shows the results of the semi-nested and nested PCRs using the NS1-NS8 amplicon as template. Each reaction produced an amplicon of expected sized with the corresponding 'no DNA reamplification' controls showing no bands indicative of clean amplification reactions.

3.1.3 DNA sequencing

A typical electropherogram generated by analyzing gel images captured on the DNA sequencer is shown in Fig. 4 (vol. II, p. 9). Each electropherogram represents the results of a single DNA sequencing reaction and the assemblage of the cycle sequencing results for *P. patens*, *M. aenigmaticus*, *N. breutelii*, *P. vittata* and *S. palustre* are shown in Fig. 5, 6, 7, 8, and 9 respectively (vol. II, p. 12-26).

After initiation of the project, the sequence of the *Physcomitrella* 18S rRNA gene was determined independently by Capesius, 1995 (GenBank accession # X80986), and the sequence data obtained in this project was compared *via* a homology alignment in Fig. 10 (vol. II, p. 27). The locations of the primers used in this study relative to the complete sequence of the *Physcomitrella* 18S rRNA gene are identified in Fig. 1 (vol. II, p. 3) and correspond to the sequence presented in the lower line of the homology alignment in Fig. 10. The upper line of the homology alignment represents

Physcomitrella 18S rDNA sequence data generated in this project with 'Ns' representing unresolved nucleotides. The location of the primers used for the PCR and sequencing reactions dictated that only approximately 94% of the whole Physcomitrella 18S rRNA gene sequence could be determined. Almost 100% of the obtainable sequence data was determined for the Physcomitrella 18S rRNA gene with the exception of the four nucleotides at positions 1123, 1131, 1139 and 1171 as indicated with 'Ns' in Fig. 10.

The homology assessment between the two independently determined Physcomitrella 18S rDNA sequences revealed over 99.8% agreement. Only two differences were found. The nucleotide at position 1519 is recorded as a 'G' in GenBank # 80986 but was identified as an 'A' by the sequencing work of this project. The other disagreement between the two sequences stems from the placement of a 'C' either at position 1526 or 1530.

3.2 Sequence alignment and secondary structure information

The results of the sequence alignment of the 18S rRNA genes for 67 taxa designated as taxa set B (Table 3, vol. II, p. 35) are presented in Appendix G (vol. II, p. 220-265). The aligned sequence length is 2,198 nucleotides of which 1,165 positions (53%) are invariant across all taxa. A position with a single nucleotide aligned against a gap or missing data in the other sequences is treated as invariant, as are positions with identical nucleotides in all sequences. The remaining 1,033 nucleotide positions represent either point mutations (substitutions in one or more orthologous sequences) or indels in one or more of the sequences.

The DCSE alignment presented in Appendix G contains the postulated 18S rRNA secondary structure pattern for all 67 taxa in encoded form. The symbols used to indicate the secondary structure elements are described in Appendix G (vol. II, p. 219) and are used in concert with the helix numbering (Row 63) proposed by Van de Peer *et al.* (1997). The 18S rRNA sequences are written in the 5' to 3' orientation with the secondary structure symbols intercalated between the primary sequence characters.

The most problematic aspect of the sequence alignment was the positioning of the indels through the introduction of gaps into the sequences and ensuring that these gaps truly represented insertion and deletion events. Secondary structure information was used to assist in the alignment of these areas. A typical example illustrating the usefulness of the secondary structure information in the sequence alignment procedure is located on p. 235 (vol. II). At position 766, 767 and 768 there are three 'Cs' in the sequence of *Physcomitrella* (Row 67). These three nucleotides represent the 5' strand of helix E23-1 immediately before the terminal loop of the helix containing the sequence 'CUUGU'. The alignment of the most 3' 'C' of these three nucleotides and the nucleotides in the loop is quite arbitrary based on primary sequence structure and several alignment rearrangements would satisfy the maximized nucleotide identity principle. Typically, these nucleotides would be removed from the analysis and labeled as arbitrary alignment positions or worse they would be retained for phylogenetic analysis and perhaps aligned incorrectly with a correspondingly erroneous homology statement. However, secondary structure information about the 3' region of the E23-1

helix immediately following the terminal loop shows that three 'Gs' would pair up with the three 'Cs' on the 5' region of the E23-1 helix. Consequently, the alignment of the most 3' 'C' and likewise the most 5' 'G' are no longer arbitrary and can be used correctly in phylogenetic estimation. The alignment of 'CUUGU' remains arbitrary but is recognized as such since it is identified to lie in a highly variable terminal loop region of helix E23-1.

The alignment of 67 orthologous 18S rDNA sequences produced 66 positions for which homology assessment remained arbitrary. These positions are indicated with '-1' in the column headings of the MacClade matrix (Appendix H, vol. II, p. 268-344) and were removed for all phylogenetic estimations. This exclusion reduced the data set in a procedure termed 'culling' which has been generally accepted as an appropriate method for the objective treatment of 'alignment ambiguous' sites (Gatesy *et al.*, 1993; Wheeler *et al.*, 1994).

The proposed secondary structure can also provide corroborating evidence for sequence accuracy. The accuracy of the sequence data used in this project and their error status are indicated in Table 3 (vol. II, p. 35). The proposed secondary structure of helix 46 for 18S rRNA provided additional support for the *Physcomitrella* sequence data presented in this project. The *Physcomitrella* 18S rRNA sequence given in GenBank # 80986 requires the integrity of helix 46 to be maintained with only five conventional base pairings which are split with three unconventional base pairings containing an internal bulge with one unpaired base and forcing an unstable terminal loop of three unpaired nucleotides (Fig. 14, vol. II, p. 52). The corrected *Physcomitrella* 18S rRNA secondary

structure, deduced from primary sequence data generated in this project, proposes a much more stable helix 46 consisting of six conventional base pairings interrupted with only two unconventional pairings and with a more stable terminal loop of four unpaired bases (Fig. 15, vol. II, p. 54).

3.3 Assessment of phylogenetic signal and variability

Waters *et al.* (1992) estimated the phylogenetic signal in plant, including bryophyte, 18S rDNA using homoplasy excess ratio (HER) (Archie, 1989). Farris (1991) describes both the computational drawbacks and statistical inconsistencies of HER and associated statistics as derived by Archie (1989). Because of these statistical inconsistencies, the practical use of HER is diminished (DeSalle *et al.*, 1994). Furthermore, Ashton *et al.* (1992) pointed out that the signal was assessed on a data set containing sequencing errors and ambiguities, poorly aligned sequences and where some sequences were less than 50% complete.

Estimation of phylogenetic signal for the 18S rRNA gene sequence data used in this study are found in Appendix C (vol. II, p. 36). The shape of the tree-length distribution that results from a parsimony analysis of a random subset of all trees for taxa set B is given in the upper graph of Fig. 11 (vol. II, p. 37). The frequency distribution of lengths of random trees for taxa set B is visually close to symmetrical and the skewness, as measured by the g_1 statistic (-0.6235), indicates that a limited amount of phylogenetic signal exists in the data. The results of this test provide some indication that the data in taxa set B is more structured than random sequences and that further phylogenetic

analysis is warranted. However, the signal is weak as evidenced by the symmetry of the distribution and the g_1 statistic of -0.6235. A study by Huelsenbeck (1991) examined the relationship between length of the correct tree and skewness of the tree-length distribution in simulated phylogenies. Using simulated phylogenies, in which the true tree was known, the study found that the optimal or most parsimonious tree is likely to be the correct tree only in analyses of data sets that produced tree-length distributions which are significantly more skewed than expected from random data. The 99% confidence limit for the skewness statistic (g_1) for random DNA sequence data was for values greater than -0.46. Examples of data sets, that gave correct tree topologies, had produced also significantly skewed tree-length distributions with g_1 statistics lower than -2.0. The g_1 statistic of -0.5330 for taxa set C and the close to symmetrical frequency distribution of random tree-lengths (lower graph of Fig. 11, vol. II, p. 37) suggests that very little phylogenetic signal exists in the lower plant 18S rDNA sequences.

To complement the determination of phylogenetic signal, Appendix D (vol. II, p. 39) contains data which characterize the extent and location of nucleotide variability. Fig. 12 (vol. II, p. 40) displays graphically the distribution of relative substitution rates estimated from the alignment of 18S rDNA of taxa set B. For taxa set B, 649 positions were partitioned into 59 sets of similar variability and then further categorized into five sets of equal size as indicated by color. These five sets do not include 1152 invariant positions or 65 'alignment ambiguous' positions, nor do they include 319 positions that contain a gap in more than 75% of the aligned sequences. The graph demonstrates that the amount of variability, as measured by v_i (relative substitution rate), is not distributed

evenly with respect to the number of positions. In fact, some degrees of variability are represented over 40 times more frequently than others. Also, the graph illustrates that after disregarding absolutely conserved residues, the most variable sites have a substitution rate ~800 times higher than the least variable ones.

Certainly, this variability data invites examination of the utility of parsimony for the phylogenetic estimation using 18S rDNA sequence data. Parsimony involves very stringent assumptions concerning the process of sequence evolution which includes the constancy of substitution rates across nucleotide sites (Stewart, 1993). Clearly, this assumption is not met with 18S rRNA gene sequence data and caution should be exercised with the utility of parsimony by complementing the analyses with other methods.

A theoretical underpinning of phylogenetic analyses using 18S rDNA molecular data is the assumption that the 18S rRNA gene mutates at a rate which is slow enough to avoid spurious homoplasy. However, there are eight regions of the eukaryotic 18S gene that have been identified to be locations of increased variability (Neefs *et al.*, 1993). It is important to ensure that the regions of increased variability do not contain too much 'noise' through multiple substitutions. The inclusion of this highly variable data may be required for local analysis of closely related taxa but may translate into 'noisy data' and obscure phylogenetic signal for global analyses of the diverse plant taxa used in this study.

Certainly, some parts of the gene are invariant and are easily recognized by visual inspection of the data matrix (e.g. positions 461-480, vol. II, p. 229). Conversely, visual inspection of the data matrix for highly variable positions is very subjective and error-

prone. In addition, the variable regions for 18S rRNA genes identified by Neefs *et al.* (1993) are not consistent with the taxa used in this study. Fig. 13 (vol. II, p. 42) displays the distribution of relative substitution rates with respect to the MacClade matrix (Appendix H) estimated from the alignment of 18S rDNA sequence data of taxa set B. This figure depicts graphically that the distribution of variability is not confined to particular regions nor are there clearly delimited sections of extensive hypervariability.

Closer inspection of the alignment positions and their associated variabilities are presented in Table 4 (vol. II, p. 44). The site number refers to the alignment position in the data matrix (MacClade matrix, Appendix H) where nonvariable alignment positions are excluded from the table. All other positions that contain a nucleotide in at least 25% of the aligned sequences have a variability score which was generated with 3 iterations of the substitution rate calculation (Van de Peer *et al.*, 1996c). The category numbers and their respective color codes were derived from the defined categories displayed in Fig. 12 (vol. II, p. 40). The categories divided the variable positions equally into five subsets from highest variability (category 1) to lowest variability (category 5). The relative rate limits of the subsets and corresponding category numbers are as follows: $< 10^{-0.498}$ (category 5); $10^{-0.498} - 10^{0.051}$ (category 4); $10^{0.051} - 10^{0.502}$ (category 3); $10^{0.502} - 10^{0.850}$ (category 2); $10^{0.850} - 10^{1.625}$ (category 1). Absolutely conserved positions ($v_i=0$) are assigned to category 6. Alignment ambiguous positions are assigned to category -1. Category 0 sites correspond to alignment positions that are extremely variable, but that are deleted in too many sequences to allow a sufficiently accurate measurement of their

relative evolutionary rate. The MacClade data matrix (Appendix H) contains the category assignment for each alignment position in the individual column headings.

The most variable alignment position for which homology assessment was possible is position # 819 (refer to Table 4, vol. II, p. 62 and Appendix H, vol. II, p. 295). Visual inspection of the nucleotide variability with reference to closely related taxa suggests a small degree of spurious homoplasy. For example, the character state 'G' is found from yeast to corn, although, the character state is variable within the same genus of liverwort (*Sphaerocarpus*) and lycopod (*Selaginella*). This position would be useful for local analyses but could contribute 'noise' in global analyses of all green plant taxa. If extensive homoplasy exists in the data set, the resulting 'noise' may be obscuring phylogenetic signal.

Each position was assessed for its contribution to this 'noise' factor and analyses indicated that spurious homoplasy is not an issue with the data set used in this study. The extent of variability decreased sharply from the most variable position # 819 with a variability score of 42 to only 17 for position # 2126 which is the 15th most variable site. Consequently, the number of sites exhibiting significant variability is very low and this corroborates the initial assessment of phylogenetic signal. For example, position # 2126 maintains only two character states over all lower and higher land plants (refer to Appendix H, vol. II, p. 338).

The location and extent of nucleotide variability were used in phylogenetic tree construction with a distance-based method (Van de Peer *et al.*, 1996c) to ensure branch lengths were not underestimated. This novel method for measuring the substitution rates

of individual nucleotides provides a more realistic relationship between sequence dissimilarity and evolutionary distance. Van de Peer *et al.* (1996b) have confirmed the successful use of this method for phylogenetic inference using small subunit rRNA of green algae.

The assigned categories were used also to compartmentalize the data into subsets based on variability. This gave a means to determine which nucleotides render phylogenetic signal for phylogeny reconstruction and to show exactly where in the tree the alignment positions with their associated signal provide resolution.

3.3.1 Variability map for 18S rRNA of lower plants

Figure 16 (vol. II, p. 56) shows a secondary structure model for lower plant 18S rRNA, where the positions belonging to eight sets of site variability data are indicated by colored dots. Five of these colors indicate different average variability and are; (in order of most to least variable) red, orange, yellow, green and blue. These five sets of variability were calculated, as described previously, using taxa set C (Table 3, vol. II, p. 35) representing only lower plant taxa. Figure 17 (vol. II, p. 58) shows the distribution of relative substitution rates estimated from the alignment of lower plant taxa 18S rDNA. The colored areas in the spectra define sets of nucleotides indicated by colored dots on the variability map in Fig. 16 (vol. II, p. 56). In Fig. 16, the variability of the nucleotide sites of lower plant 18S rRNA is mapped in the shape of the secondary structure model by superimposing the colors in Table 5 (vol. II, p. 60) on to the 18S rRNA secondary structure model of *Physcomitrella* shown in Fig. 15 (vol. II, p. 54).

The alignment positions and their associated variabilities are presented in Table 5 (vol. II, p. 61-64). The site number refers to the alignment position in the data matrix (MacClade matrix, Appendix H, vol. II, p. 268-344) where nonvariable alignment positions are excluded from the table. All other positions that contain a nucleotide in at least 25% of the aligned sequences have a variability score which was generated with 3 iterations of substitution rate calculation (Van de Peer *et al.*, 1996c). The category numbers and their respective color codes were derived from the defined categories displayed in Fig. 17 (vol. II, p. 58). The categories divided equally the variable positions into five subsets from highest variability (category 1) to lowest variability (category 5). The relative rate limits of the subsets and corresponding categories and colors are as follows: $< 10^{-0.123}$ (5, blue); $10^{-0.123} - 10^{0.106}$ (4, green); $10^{0.106} - 10^{0.428}$ (3, yellow); $10^{0.428} - 10^{0.909}$ (2, orange); $10^{0.909} - 10^{1.570}$ (1, red). Absolutely conserved positions ($v_i=0$) are assigned to the category 6 and given the color purple. Alignment ambiguous positions are assigned to category -1 and the 'color' grey. Category 0 and the 'color' white correspond to alignment positions that are highly variable, but that are deleted in too many sequences to allow a sufficiently accurate measurement of their relative evolutionary rate.

The debate over differential weighting of stems and loops is influenced largely by the assumption that stem regions are more conservative than the single stranded loop regions (Wheeler and Honeycutt, 1988; Vawter and Brown, 1993). Intuitively, it seems logical that base paired regions should require compensatory mutations and consequently are less prone to point mutations. However, helix 10 (Fig. 16, vol. II, p. 56), with a green and orange dot both facing a purple dot, illustrates that this is not an absolute truth.

However, in general, the two nucleotides of a base pair have the same or a neighboring color indicating they are about equally variable.

The variability map presented in Fig. 16 illustrates also that even highly conserved stems can have highly variable positions. For example, helix 49, which is found in all small-subunit ribosomal RNA, is highly variable as evidenced by red, orange and yellow colored dots. Interestingly, helix 49 in bacterial rRNA shows extensive variability as well (Van de Peer *et al.*, 1996a). Likewise, several single stranded regions are highly conserved as indicated with purple dots (e.g. the internal bulge joining helices 33, 34, 45, and 47). These examples demonstrate the need for visualization and description of structural features as done in this study.

In addition, the possible stabilizing effect of non-Watson-Crick base pairs providing transitional structures that could serve as intermediates in the evolution of standard base pairs places the practice of differential weighting for transitions and transversions in question, especially when considering repetitive genes such as 18S rDNA. This statement can be substantiated easily with a rapid examination of the variability map in concert with the secondary structure model for 18S rRNA of *Physcomitrella* (Fig. 15 and 16) which verifies within several helices that variable positions contain non-canonical sequences (e.g. helix E23-1).

Other notable features of the variability map (Fig. 16) include the curious phenomenon of single highly variable nucleotides amongst several absolutely conserved nucleotides (e.g. the single red dot in the internal bulge adjacent to helix 26). The variability map also illustrates that the exclusion of hypervariable regions, such as E23-1,

would discard some of the most informative characters. The variability map graphically depicts the extent of nucleotide conservation in lower plant 18S rRNA as shown by the high number of purple dots.

3.4 Phylogenetic analyses

In an attempt to facilitate practical discussion and concentrate on the deeper phylogenetic relationships of lower plants, all tree descriptions will focus on significant differences from previous analyses which are characterized by confidence measurements indicating greater than 50% support. Minor changes in topology could be discussed at length, but would conflict with the overall philosophy of the approach taken in this thesis which is to place emphasis and confidence in phylogenetic outcomes shared by multiple methods.

3.4.1 Character-based phylogenetic analyses

All phylogenetic trees pertaining to character-based phylogenetic analyses using parsimony can be cross-referenced with Table 6 (vol. II, p. 66), and maximum likelihood can be similarly indexed with Table 7 (vol. II, p. 67).

3.4.1.1 Parsimony

Character-based analysis under the parsimony criterion produced four equally most parsimonious trees (length=1589; CI=0.402; RI=0.619) for phylogenetic analysis with taxa set A. These trees were based on 369 parsimony-informative characters from a data matrix containing 587 variant positions. These optimal trees were found with replicate number 5 and occupy 3 tree islands. These optimal trees were found 85

additional times in 100 random-addition replicates with tree-bisection-reconnection (TBR) branch-swapping performed (total number of rearrangements tried = 13,730,714).

The 50% majority-rule consensus tree of these four most parsimonious trees is displayed in Fig. A (vol. II, p. 72). These trees differed mostly in positioning of *Sphagnum sp.* and Filicophyta taxa. One of these trees differed considerably with respect to the placement of Bryophyta species and basal branching of all lower plant taxa. The consensus tree suggests a polyphyletic phylogeny of Byropsida and Hepaticopsida clades. The tree also proposes higher plants are a monophyletic assemblage with all lower plants including ferns evolving separately with the exception of *Selaginella sp.* which have been placed basal to all land plants. This last relationship is dramatically different from those found frequently in the literature and is most likely an artifact of the analyses due to branch attraction effects of *Selaginella sp.* to the outgroup. Van de Peer *et al.* (1993) have demonstrated that sequences of OTUs with higher evolutionary rates tend to cluster closer to the base of the tree. *Selaginella sp.* 18S rRNA gene sequences demonstrate high evolutionary rates which is evidenced by their long branches. These long branches contain over eighty steps in each of the four most parsimonious trees (data not shown) and is also seen in phylograms of all distance-based analyses. Consequently, this branch attraction effect was witnessed in all subsequent phylogenetic estimations with taxa set A and this artifact will be appropriately removed from any further discussion.

All relationships of lower plant taxa and basal lineages are dissolved completely in the 50% majority-rule bootstrap tree presented in Fig. B (vol. II, p. 73). This indicates

that the data have provided little support for these relationships, and consequently, we cannot assume any confidence in the relationships of the Byropsida or other lower plant taxa. Hillis and Bull (1993) have pointed out that bootstrap analyses resulting in 50% majority-rule bootstrap trees can result in overestimates of accuracy for correctly inferring the true phylogeny. They demonstrated that sequences which exhibit unequal rates of evolutionary change are most susceptible to this phenomenon. In an effort to ensure the bootstrap values are not being misinterpreted, decay analyses were performed and the results are presented in Fig. C (vol. II, p. 74). The decay indices concur with the bootstrap values by providing correspondingly weak support for the relationships suggesting bryophyte genealogy and other relationships among the lower plant taxa. Specifically, decay indices as low as 2 are found supporting clusters of bryophyte classes which correlate with bootstrap values of <50%. Similarly, a decay index of 30 is found supporting the relationship between two *Sphagnum* species and this corresponds to a bootstrap value of 100%. Clearly, both assessments of confidence suggest that the data set supports the resolution of closely related species but offers little confidence with respect to the revelation of deeper phylogenies.

The phylogenetic analysis of taxa set A using maximum parsimony produced dendrograms which appeared susceptible to taxon sampling and outgroup effects. The acquisition of a faster computer later in the study permitted a more comprehensive analysis with taxa set B. The addition of 26 sequences to taxa set A allowed better representation of the presumed monophyletic groups and allowed the addition of green algae and fungi to the analysis. The addition of the green algae provided a 'buffer-zone'

between the fungal outgroup and the taxa of phylogenetic investigation. Character-based analysis of taxa set B, under the parsimony criterion, produced 88 equally most parsimonious trees (length=3156; CI=0.357; RI=0.595). These trees were based on 586 parsimony-informative characters from a data matrix containing 993 variant positions. These optimal trees were found with replicate number 17 and occupy 4 tree islands. These optimal trees were found 46 additional times in 100 random-addition replicates with tree-bisection-reconnection (TBR) branch-swapping performed (total number of rearrangements tried = 159,744,909). The 50% majority-rule consensus tree of these 88 most parsimonious trees is displayed in Fig. D (vol. II, p. 75).

In all optimal trees found, the following groups were monophyletic:

Anthocerotopsida, Magnoliophyta, Pinophyta, Isoetopsida, Lycopodiella, Charaphyta, and Chlorophyta. These monophyletic clades agree with the established phylogeny for these taxa, with the exception of Charaphyta since most phylogenies suggest this group is paraphyletic with Coleochaetales closest to the embryophyte clade (Kranz *et al.*, 1995). However, several relationships of the clades presented in Fig. D concur with recent analyses in the literature. The Charophyceae and the orders circumscribed within it are placed basal to the Streptophyta lineage (Bremer, 1985; McCourt, 1995; Surek *et al.*, 1995). The angiosperms are a monophyletic group and are a sister group to the gymnosperms, which is in agreement with the widely accepted phylogeny of these groups (Chaw *et al.*, 1997).

The placement of the green algae at the base of Streptophyte lineage and higher plants at the top (e.g. Magnoliophyta and Pinophyta taxa) coincides with the

demonstrated phylogeny of this lineage. Accordingly, these taxa were included mainly to provide 'boundaries' or 'guideposts' for the focus of this investigation which is the placement of the taxa between these two groups. The placement of the lower plants has already been established to lie somewhere in the middle of these two groups and the dendrogram presented in Fig. D does not disagree with this contention. This 'framing' of the lower plant taxa provided a buffer-zone which diminishes the artifactual effects of taxon sampling and outgroup selection which was suspected to be problematic in taxa set A (the original data set).

The tracheophyte lineage begins with the monophyletic Lycopodiophyta. Recent work of Stewart and Rothwell (1993) and Kranz and Huss (1997) support the placement of Lycopodiophyta at the base of the tracheophyte lineage. The monophyly of Lycopods is in agreement with the work by Niklas and Banks (1990) and Kranz and Huss (1997) but disagrees with polyphyletic phylogeny presented by Garbary *et al.* (1993). The tracheophyte lineage continues with the paraphyletic phylogeny of the ferns, within which the horsetails and whiskferns branch, and the lineage ends with the higher plants as expected.

Debate concerning bryophyte phylogeny has been whether the group is paraphyletic (Mishler and Churchill, 1984, 1985; Waters *et al.*, 1992; Hedderson *et al.*, 1996) or monophyletic (Garbary *et al.*, 1993). The bryophyte phylogeny determined with parsimony analysis using taxa set B supports the paraphyletic assemblage with most of the Bryopsida a sister group to the tracheophytes. Two moss species, *Atrichum undulatum* and *Polytrichum formosum* formed a sister group along with some liverworts

to the monophyletic hornwort clade. This hornwort-moss-liverwort group was found to be a sister clade to the tracheophyte lineage. Interestingly, two moss species are found more closely related to higher plants than all other lower plants including ferns. The two *Sphagnum* species occurred in this position in 84 of the 88 most parsimonious trees.

Some Hepaticopsida emerge as a basal group among the land plants which is supported by earlier work (Mishler and Churchill, 1984, 1985; Mishler *et al.*, 1992; Waters *et al.*, 1992) but disagrees with the more recent work of Hedderson *et al.* (1996) which places hornworts as the basal group among land plants. Furthermore, the Hepaticopsida group is not monophyletic since some liverwort taxa group with the Anthocerotopsida clade which is contrary to work of Garbary *et al.* (1993). These liverworts which group with the hornworts comprise the subclass of leafy liverworts called Jungermannopsida. Interestingly, the liverwort taxa which form the basal group to all land plants consist of only thalloid liverworts which comprises the Subclass Marchantiopsida. There appears to be a distinctly separate evolution of these two subclasses of liverworts and this separation has been reported elsewhere (Capesius, 1995; Bopp and Capesius, 1996; Capesius and Bopp, 1997).

Clearly, the phylogeny of lower plants illustrated in Fig. D (vol. II, p. 75) presents some relationships which could find support with one or a few previous publications. However, the resolution of the debate over the phylogeny of lower plants should not be conducted by strengthening the support for particular relationships or finding weakness in others by quoting the number of published trees conforming or disagreeing with a phylogeny. This would be especially inappropriate given the concerns

identified in the introduction of this thesis which pertain to the questionable phylogenetic estimation of most if not all the published trees. Therefore, this thesis will claim support for phylogenetic relationships only by measuring confidence with bootstrap analyses.

The 50% majority-rule bootstrap tree found using maximum parsimony with taxa set B is presented in Fig. E (vol. II, p. 76). The tree is highly polytomous for deeper levels of lower plant phylogeny. The strongly supported clusters of lower plants presented in Fig. E is found primarily at the tips of the branches. Again, the data set allows the resolution of closely related species but offers little confidence in deeper phylogenetic separations. The only deep phylogenetic separation of the lower plants which is supported by bootstrap analyses is the suggested monophyly of the Lycopodiophyta group but with only 55% confidence.

3.4.1.2 Quartet puzzling

The maximum likelihood method for reconstructing phylogenetic relationships is another character-based method and works by trying to evaluate the probability of observing a particular set of data. Maximum-likelihood techniques are championed by many researchers from the population genetics tradition as the preferred phylogenetic estimation approach (Mishler, 1994). However, maximum likelihood's computational complexity prevents phylogenetic estimation for large numbers of taxa. Quartet puzzling can be applied to data sets with a much greater number of taxa than can other search algorithms such as stepwise addition (e.g. fastDNAm1, Olsen *et al.*, 1994). However, it has been recognized that quartet puzzling's ability to reconstruct the true tree is less than that of fastDNAm1 (Strimmer and Von Haeseler, 1996). Nevertheless, the large number

of taxa in this study and popular acceptance of the validity of the maximum likelihood approach justify amply the employment of quartet puzzling.

The results of quartet puzzling on taxa set A is presented in Fig. F (vol. II, p. 77) and the results using taxa set B is presented in Fig. G (vol. II, p. 78). The deeper phylogenetic relationships remain unresolved as was the case using parsimony analysis and taxa set B. However, taxa set A (Fig. F) proposes a separate evolution of the higher plant taxa (e.g. Magnoliophyta and Pinophyta) from all other tracheophytes and lower land plants. This is a controversial finding and conflicts dramatically with accepted phylogeny of the streptophyte lineage. Such a result may depend upon a taxon sampling effect confounded by the maximum likelihood principle as a similar result was obtained with fastDNAm1 (see below).

3.4.1.3 Maximum likelihood

The traditional maximum likelihood calculation with stepwise addition generated long run times. Consequently, analyses was performed only on the reduced taxa set, A. The results of the calculation are shown in Fig. H (vol. II, p. 79). The maximum likelihood analysis was conducted on 681 distinct data patterns found in taxa set A, and examined 3493 alternative trees producing the final tree with Ln likelihood of -13491.72. The results of the bootstrap analysis are given in Fig. I (vol. II, p. 80).

The phylogeny in Fig. I coincides with results obtained using quartet puzzling since it too suggests a separate evolution of the higher plant taxa. However, such a phylogeny requires confirmation derived from other analytical methods (see above). The computational requirements of this method permitted only 10 bootstraps of the data and

therefore extra caution is needed for interpretation of the confidence measurements. In fact, the bootstrap tree produces a different phylogeny from that of the original analysis by clustering the ferns and some fern allies with the higher plants. This reinforces the conviction that trees displaying a completely separate evolution of higher land plant taxa are spurious. The remainder of the tree fails also to determine deeper green plant phylogenies. However, branch tips are resolved as with previous data sets and methods.

3.4.2 Distance-based phylogenetic analyses

All phylogenetic trees pertaining to distance-based phylogenetic analyses can be cross-referenced with Table 8 for taxa set A (vol. II, p. 68) and Table 9 for taxa set B (vol. II, p. 69). Each phylogenetic estimation using distance-based approaches resulted in the production of four trees. The first pair of trees are phylograms depicting branch length and have the accompanying distance scale located at the top of the page. The second pair of trees display the results of bootstrap analyses and have the accompanying bootstrap values shown as percent. All branches yielding less than 50% support are collapsed to produce a 50% majority-rule bootstrap tree. It is important to note, however, that these bootstrap trees are not constructed with the consensus tree approach as are the 50% majority-rule bootstrap trees used in the character-based analyses. Rather, the results of the bootstrap analyses for distance-based analyses are displayed on the original phylogram with branches showing less than 50% support collapsed to produce polytomies.

Table 10 (vol. II, p. 70) summarizes consistent monophyletic clusters found in all distance-based analyses. These monophyletic clusters constitute the triangles shown in

the second tree of each pair of trees. The trees with the monophyletic clusters displayed as triangles have the same figure label as the original tree on the preceding page. The second tree with the monophyletic clusters shown as triangles is used to illustrate the relationships among the major groups more clearly.

3.4.2.1 Neighbor-joining analysis with Jukes & Cantor (1969) distance calculation

Figure A1B (vol. II, p. 147-148) shows the results of the neighbor-joining analysis on taxa set B using Jukes and Cantor (1969) distance calculation with branches showing less than 50% support by bootstrap analysis dissolved to form polytomies. The significance difference seen with this phylogenetic estimation is the placement of *Klebsormidium flaccidum* and *Coleochaete scutata*. These results place Coleochaetales as the closest green algae to the embryophyte clade. Resolution of the relationships of the bryophytes and other lower land plants remained unsupported which is indicative of the poor phylogenetic signal available for deeper phylogenetic analyses. Again, the phylogeny of closely related taxa and other overall relationships identified in previous descriptions for taxa set B remain unchanged both with and without correction for insertions and deletions.

Jukes & Cantor distance calculation on taxa set A with no correction for indels displays a separate evolution of the higher plants (i.e. Pinophyta and Magnoliophyta) from the rest of the Streptophyta lineage (Fig. A1S, vol. II, p. 81-82). This arrangement concurs with parsimony analyses of taxa set A (refer to Fig. A, vol. II, p. 72). Analysis of taxa set A with correction for insertions and deletions (Fig. A2S, vol. II, p. 85-86) produces no significant changes in tree topology but does produce reduced bootstrap

support for some deeper clustering of the lower plant taxa. However, bootstrap analyses provides 75% support with correction for indels and 73% confidence without indel correction for a separate evolution of the higher plants (Fig. A2SB, vol. II, p. 87-88). Although this arrangement has been seen in previous character-based analyses with taxa set A, this is the first time bootstrap analyses supports this separate evolution of higher plants. As discussed previously, this arrangement is highly controversial and suspect due to outgroup and other species sampling concerns with taxa set A. Notably, taxon arrangements at the tips of the branches maintain both identical clustering and high bootstrap support with all previous analyses including taxa set B (e.g. the topology of very closely related species is maintained).

Further evidence which substantiates bootstrap values is branch length. All distance-based phylogenetic trees are displayed as phylograms thereby depicting genetic distance between nodes. The branch length is congruent with the amount of phylogenetic signal supporting each cluster in the tree with short branches indicative of low phylogenetic signal supporting the corresponding cluster. All phylograms produce repeatedly short branches for deep phylogenetic relationships of the lower land plants. Conversely, closely related taxa and their consistent clustering which are demonstrating high bootstrap support have also long branch lengths. The consistency of this observation is significant but for the sake of brevity will not be reiterated with each distance calculation analysis.

3.4.2.2 Neighbor-joining analysis with Tajima & Nei (1984) distance calculation

All phylogenetic trees produced by neighbor-joining with Tajima & Nei (1984) distance calculation and with correction for indels for both taxa set A and B produced trees identical to those obtained from the previous distance calculation. Similar shuffling of branches was noted with taxa set A, although, no significant alternations to tree topology were observed. The results of neighbor-joining analyses with Tajima & Nei distance calculation can be found on p. 89-97 (vol. II) for taxa set A (Fig. B1S, B1SB, B2S, and B2SB) and p. 153-160 (vol. II) for taxa set B (Fig. B1, B1B, B2, and B2B).

3.4.2.3 Neighbor-joining analysis with Kimura (1980) distance calculation

Neighbor-joining analysis with Kimura (1980) distance calculation produced a tree topology for taxa set A (Fig. C1S, vol. II, p. 97, 98 & Fig. C2S; vol. II, p. 101, 102) which was fully restored to the topology observed with Jukes & Cantor distance calculation (Fig. A1S & A2S, vol. II, p. 81-85) regardless of treatment of indels. Taxa set B (Fig. C1, vol. II, p. 161 & Fig. C2, vol. II, p. 165) retains similar topology with all previous distance calculation analyses.

3.4.2.4 Neighbor-joining analysis with Jin & Nei (1990) distance calculation

The results of Jin & Nei distance calculation with both Jukes & Cantor and Kimura 2-p evolution models can be found on p. 105-120 (vol. II) for taxa set A (Fig. D1.1S, D1.1SB, D1.2S, D1.2SB, D2.1S, D2.1SB, D2.2S, and D2.2SB) and p. 169-184 (vol. II) for taxa set B (Fig. D1.1, D1.1B, D1.2, D1.2B, D2.1, D2.1B, D2.2, and D2.2B). Again, no significant changes are observed for either taxa set from the topologies previously described.

3.4.2.5 Neighbor-joining analysis with transversions only distance calculation

Dramatic topology changes are observed with transversion only distance calculation for both taxa sets. The results of these analyses can be found on p. 121-128 (vol. II) for taxa set A (Fig. E1S, E1SB, E2S, and E2SB) and p. 185-192 (vol. II) for taxa set B (Fig. E1, E1B, E2, and E2B). Taxa set B produced several rearrangements and changes in taxa affiliation, most of which disappeared when bootstrap confidences of $\geq 50\%$ were applied. However, taxa set A topology associated ferns and fern allies with higher plant taxa making them a sister group to the Pinophyta and Magnoliophyta lineage. These relationships were seen also in trees derived from character-based analyses using maximum likelihood methods (refer to Fig. H, vol. II, p. 79).

3.4.2.6 Neighbor-joining analysis with no correction distance calculation

Phylogenetic analysis using distance calculation with no correction for superimposed mutations was performed strictly with the intention of obtaining a reference point for the level of correction the other parametric models imposed on the raw data. Consequently, the results of these analyses were not factored into the construction of the overall consensus tree. The topologies produced with this method fit well with other methods described previously. The results of these analyses indicate that the level of influence each distance method imposes is not as great as imagined initially. The results of these analyses can be found on p. 129-136 (vol. II) for taxa set A (Fig. F1S, F1SB, F2S, and F2SB) and p. 193-200 (vol. II) for taxa set B (Fig. F1, F1B, F2, and F2B).

3.4.2.7 Neighbor-joining analysis with Van de Peer & De Wachter (1994) distance calculation

Unlike previous distance calculation models, the novelty and appropriateness of this distance calculation method merits its description here. This distance-based method is the most appropriate parametric model for phylogenetic estimation using 18S rRNA gene sequence data. This method accounts for the local sequence variability in the computation of evolutionary distance. The calculation utilizes a parameter in an equation which expresses the probability that the alignment position contains a different nucleotide in orthologous sequences. This probability is expressed as a function of the distances separating these sequences (i.e. the number of nucleotide substitutions that occurred since the divergence of two sequences). The distances computed for 18S rDNA using this method are reputed to be more precise than those obtained with other distance-based methods (Van de Peer *et al.*, 1993).

The results of neighbor-joining phylogenetic analysis using the Van de Peer and De Wachter (1994) distance calculation method for taxa set A are given in Fig. G1S & G2S (vol. II, p. 137-144) and Fig. G1 & G2 (vol. II, p. 201-208) for taxa set B. The most notable difference in topology is seen with the clustering of fern and fern allies as a sister group to the higher plants. It would appear that this proposed phylogeny is beginning to carry some weight despite the low bootstrap value of 53%. This relationship is seen with taxa set A and B and with both distance-based and character-based methods. However, this method could not produce any revelations concerning the relationships of other lower land plants or resolve any of the debates over the genealogy of the bryophytes. It is

evident now that there is simply insufficient phylogenetic signal in 18S rRNA gene sequences to resolve unequivocally and in detail the evolutionary transition from algal ancestors to land flora.

3.4.2.8 Neighbor-joining analysis with no correction distance calculation and compartmentalization

The background work required for this analysis was described in section 3.3 (Assessment of Phylogenetic Signal and Variability, I; p. 53). The categories determined by the data presented in Table 4 (vol. II, p. 44) were used to compartmentalize the data into subsets based on variability. This provided a means to determine which nucleotides render phylogenetic signal for phylogeny reconstruction and to show exactly where in the tree the alignment positions with their associated signal provide resolution. If the data matrix contains homoplasy due to excessive variability, then the more variable positions (e.g. categories 0, 1, 2) should be most appropriate for closely related taxa but only contribute 'noise' for global analyses with widely divergent taxa. Likewise, the more conserved alignment positions (e.g. categories 3, 4, 5) should contribute phylogenetic signal appropriate for deeper phylogenetic relationships. However, if phylogenetic signal is sufficiently weak due to low variability then all the variability and hence signal will be useful only for phylogenetic study of closely related taxa and resolution of deeper phylogenetic relationships would not be possible.

The first subset utilized characters assigned to categories 0, 1, 2, 3, and 4 which essentially removed only the most invariant alignment positions. The results of these analyses are presented in Fig. H1B (vol. II, p. 209, 210). The topology of this analysis is

compared most appropriately to distance-based analysis with no correction and these results are presented in Fig. F1B (vol. II, p. 195, 196). Comparison between these two trees reveal only minor differences which is somewhat contrary to what should be expected with the removal of the most conserved nucleotides if in fact these positions contribute to deeper relationships.

The next subset retains the most variable nucleotides corresponding to categories 0, 1, and 2. The results of these analyses are presented in Fig. H2B (vol. II, p. 211-212). The topology remains unchanged and indicates the vast majority of the phylogenetic signal providing the resultant tree resides in the most variable sites. Deeper phylogenetic relationships remain unresolved as in all previous analyses suggesting even the positions in categories 3, 4, and 5 are too conserved to provide phylogenetic signal.

The final subset includes all the variable nucleotides by retaining categories 2, 3, 4, and 5, except the 'hypervariable' ones within categories 0 and 1. The resulting tree is presented in Fig. H3B (vol. II, p. 213-214). The monophyly of three clades loses support with the removal of these positions but deeper relationships are not 'unmasked'. This indicates clearly that the data set and all phylogenetic analyses are not suffering from extensive homoplasy but that conversely the most variable nucleotides contain data suitable only for resolving clades of closely related taxa. Consequently, it seems that the 18S rDNA gene is too similar within lower plant taxa to allow informative deeper level phylogenetic analyses. This degree of similarity among 18S rRNA gene sequences could indicate that extant species of lower plants are evolutionary younger than other terrestrial plants.

3.4.3 Overall strict consensus tree

In an effort to summarize the many trees generated by this study, an overall strict consensus tree was developed for both taxa sets. These consensus trees based on all analyses represent a very conservative view of the clades and phylogenetic relationships for which this project provides complete support. Only clusters which were maintained strictly in the 50% majority-rule bootstrap trees from all phylogenetic analyses are depicted. Any variation of taxa clustering emanating from the various phylogenetic approaches resulted in those affected taxa being displayed as polytomies. These trees are presented in Fig. 18 (vol. II, p. 215) pertaining to taxa set A and Fig. 19 (vol. II, p. 217) pertaining to taxa set B.

These figures delineate graphically the overall conclusion of this project. The phylogenetic relationships of the bryophytes and other lower plant taxa cannot be resolved with 18S rDNA sequence data. The data provides good resolution for closely related taxa, the phylogeny of higher plants and the placement of the bryophytes between the green algae and the tracheophytes. Any other claims would be unwarranted.

3.5 Conclusion

The final inference is that 18S rDNA sequence data do not have sufficient resolving power to contribute significantly to our understanding of phylogenetic relationships of the bryophytes and other lower plant taxonomic groups. This study strove to resolve unequivocally the evolutionary transition from putative algal ancestors to the extant land flora by elucidating bryophyte phylogeny and differentiating major lower plant lineages thereby providing insight into the sequence of evolutionary events which accompanied colonization of the terrestrial habitat. In doing so, it became evident that there is insufficient phylogenetic signal in 18S rRNA to elucidate satisfactorily lower plant phylogenies.

The notion that 18S rRNA gene sequence data is not sufficient to solve the problem at hand is not a novel revelation. Capesius and Van de Peer (1997) share this contention and state "that the 18S rRNA is not very well suited to resolve all branching patterns within the different orders and families of bryophytes". A recent study by Tourasse and Gouy (1997) suggests that the large-subunit rRNAs appear to be more suitable than small-subunit rRNAs for reconstructing universal phylogenies. In addition, several studies suggest that sequence data from several independent loci must be used to obtain the correct 'species tree' (Saitou and Nei, 1986; Bull *et al.*, 1993; Kim, 1993; Cao *et al.*, 1994; Avise and Nelson, 1995).

Methods for inferring phylogeny from DNA sequences have proliferated in the last few years. Unfortunately, decisions concerning which of many described methods will be used in a given study are rarely made by weighing the advantages and

disadvantages of each approach. Similarly, the choice of DNA sequence has not been examined closely. Instead, issues of availability or historical inertia often dictate such choices. Future investigations examining the phylogeny of lower plants cannot afford to continue in this vein. This study illustrates that obtaining sufficient data and the complete application of appropriate methods of phylogenetic inference are essential for the reconstruction of deep phylogenetic relationships of lower land plants. Although this study satisfied the latter criterion, it showed clearly that plant 18S rRNA gene sequence data are insufficient for the purpose proposed at the outset of this project.

Consequently, future investigations should be scrutinized carefully when proposing bryophyte or lower plant phylogeny based strictly on this gene.

3.5.1 Future work

As the underlying theme of this thesis maintains, it is preferable that when a putative clade fails to appear consistently across several alternative phylogenetic procedures, as applied here to 18S rDNA, the fruitful approach will be to gather and analyze additional data, including those from unlinked regions, rather than to proclaim what may be a fragile truth from one preferred method of phylogeny estimation.

There are several possible strategies for improving the power of phylogenetic inference using DNA sequence data. Certainly, increasing the number of nucleotides will increase the chance of obtaining a whole-genome tree, whereby whole-genome trees are closer to species trees than any specific gene tree. Focusing on a single gene increases the chance that phylogenetic inference is affected by location-dependent processes in sequence evolution (Cummings *et al.*, 1995) in turn resulting in the generation of trees

characterized by spurious associations. This can be avoided by sampling sites from different genes or even genomes (i.e. chloroplast, nuclear).

The 18S rRNA gene sequence data should not be discarded but rather combined with more data. Furthermore, this and previous studies have not utilized all phylogenetic information in rRNA gene sequences associated with features such as base-paired versus non base-paired nucleotides, compensatory mutations, and types of secondary structure. The merits for doing so are obvious. Structural aspects of rRNA offer a number of opportunities for informed examination of the data for phylogenetic information. Different positions of the molecule may have very rigid evolutionary constraints connected with function either by sequence or structure.

The culling method used in this study for accommodating 'alignment ambiguous' sites is not the only appropriate method for dealing with the multiple putative homology statements. Certainly to date, the process of incorporating several possible alignment solutions into phylogenetic analysis has not been well delineated. Another possible solution involves stringing the multiple alignments together into a single grand alignment (Wheeler *et al.*, 1995). This procedure, termed 'elision', ensures that positions with alignment of unambiguous sites will occur more often than alignment-ambiguous sites and will thus carry greater weight. This procedure allows phylogenetic analysis to be performed on the entire data set whereby a less extreme form of weighting is implemented relative to culling.

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**The Inference of Phylogenetic Relationships
among the Main Lineages of Terrestrial Plants,
With Emphasis on the Positions of the Major Bryophyte Groups,
Using Small-Subunit Ribosomal RNA Gene Sequences**

Volume II

A Thesis

Submitted to the Faculty of Graduate Studies and Research

in Partial Fulfillment of the Requirements

for the Degree of

Master of Science

in Biology

University of Regina

by

Nick A. Antonishyn

Regina, Saskatchewan

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APPENDIX A: Raw Data

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Table 1. Primers

Name	Sequence ^a	Primer Ln	Size (bp) ^b	T _m (°C) ^c
NS1	GTA GTT ATA TGC TTG TCT C	19	521	45.4
NS2	GGC TGC TGG CAC CAG ACT TGC	21		74.7
NS3	GCA AGT CTG GTG CCA GCA GCC	21	566	74.7
NS4	CTT CCG TGA ATT CTT TTA AG	20		63.6
NS5	AAC TTA AAG GAA TTG ACG GAA	22	628	62.8
NS7	GCA ATA ACA GGT CTG TGA TGC	21	337	62.7
NS8	TCC GCA GGT TCA CCT ACG GA	20		70.7

^a Sequences are written 5'-3'. All odd-numbered primers are 5' primers; even numbers indicate 3' primers.

^b Product sizes are approximate based on the 18S rDNA gene of *Physcomitrella patens*. the size of the region amplified is the product size minus the primers.

^c T_m's were calculated by the neighbor-joining method.

Figure 1.

Locations of primers on the 18S rDNA gene.

The arrowheads represent the 3' end of each primer.

<u>Primer</u>	<u>Location*</u>
NS1	20-38
NS2	560-580
NS3	560-580
NS4	1146-1165
NS5	1143-1164
NS7	1437-1456
NS8	1794-1814

* location is identified with respect to *Physcomitrella patens* (X80986)

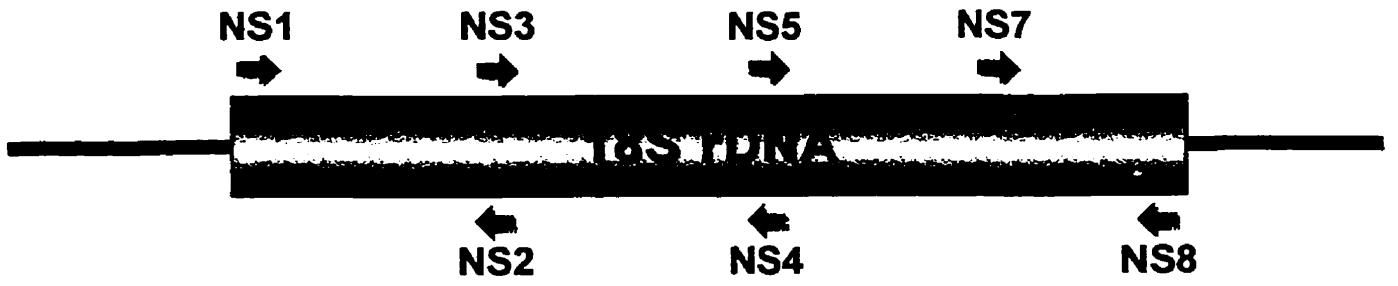


Figure 2.

Photograph of NS1-NS8 PCR products subsequently used as templates for semi-nested and nested PCRs.

NS1-NS8 PCR products used as template for subsequent PCRs were sized using electrophoresis on an 1% agarose gel; stained with ethidium bromide and visualized with UV (312nm) light. The image was captured on Polaroid 667 film.

The expected amplicon size ~1800 bp.

<u>Lane</u>	<u>Content</u>
M	100 bp DNA Ladder
1	<i>Pteris vittata</i>
2	<i>Physcomitrella patens</i>
3	<i>Notophylas breutelii</i>
4	<i>Megaceros aenigmaticus</i>
5	<i>Sphagnum palustre</i>

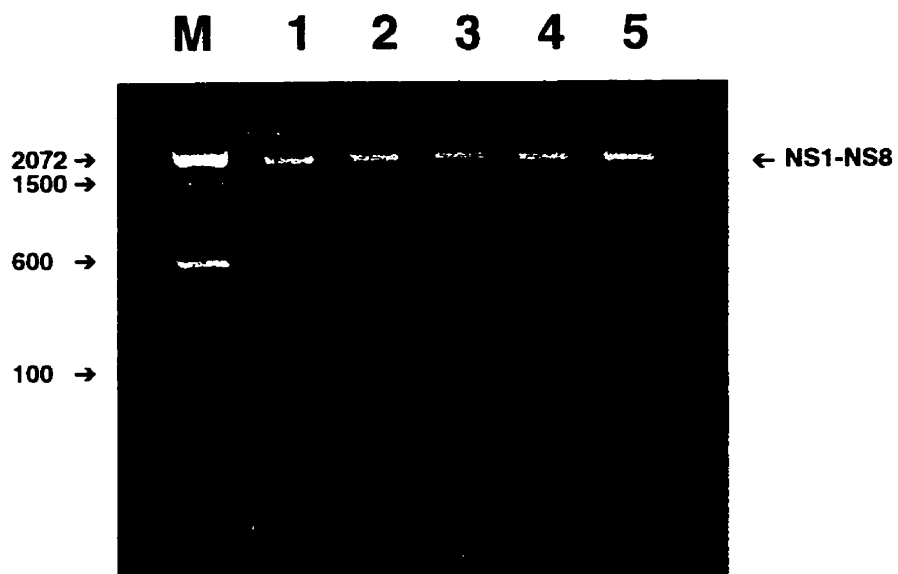


Figure 3.

Photographs of semi-nested and nested PCR products subsequently used as sequencing templates.

Semi-nested and nested PCR products using NS1-NS8 amplicon as template were sized using electrophoresis on 1.6% agarose gels; stained with ethidium bromide and visualized with UV (312nm) light. Images were captured on Polaroid 667 film.

<u>Photo</u>	<u>Taxa</u>
a	<i>Physcomitrella patens</i>
b	<i>Sphagnum palustre</i>
c	<i>Megaceros aenigmaticus</i>
d	<i>Notophylas breutelu</i>
e	<i>Pteris vittata</i>

<u>Lane</u>	<u>Content</u>
M	100 bp DNA Ladder
1	NS1-NS2 PCR product (expected size ~ 561 bp)
2	NS1-NS8; no DNA reamplification control
3	NS3-NS4 PCR product (expected size ~ 607 bp)
4	NS1-NS8; no DNA reamplification control
5	NS5-NS8 PCR product (expected size ~ 670 bp)
6	NS1-NS8; no DNA reamplification control
7	NS7-NS8 PCR product (expected size ~ 378 bp)
8	NS1-NS8; no DNA reamplification control

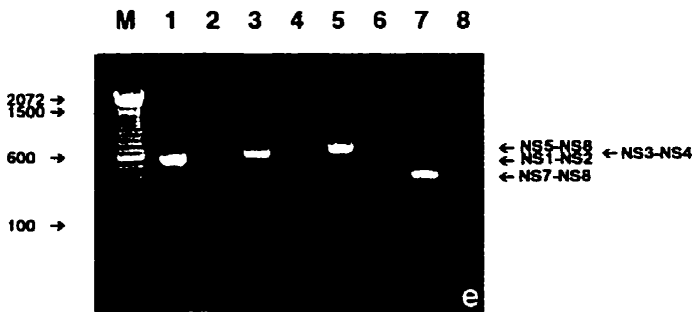
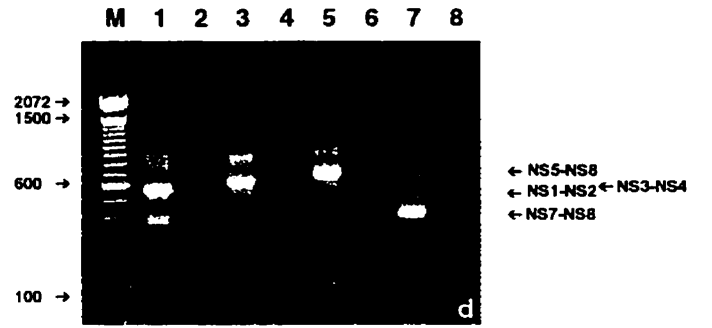
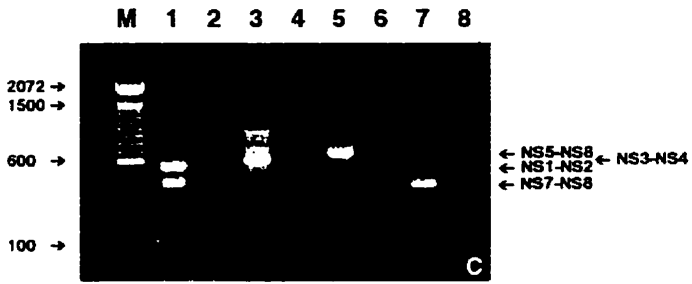
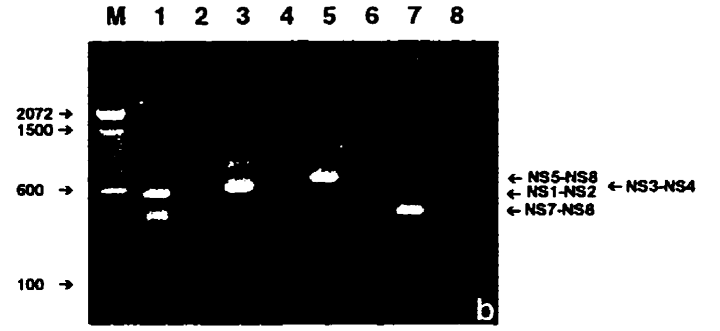
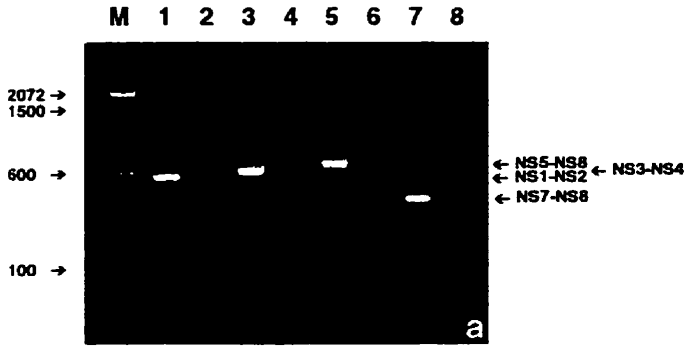


Figure 4.

Typical electropherogram.

Generated by analyzing gel images captured on the ABI 377 DNA Sequencer. The electropherograms were analyzed using ABI Prism DNA sequencing analysis software (version 2.1.1). The electropherograms represent 520 bp of double-stranded sequence data of the 5' portion of the 18S rDNA gene for *Physcomitrella patens*.

02. NS1.1 is the forward strand, Page 10

04. NS2.1 is the reverse strand, Page 11

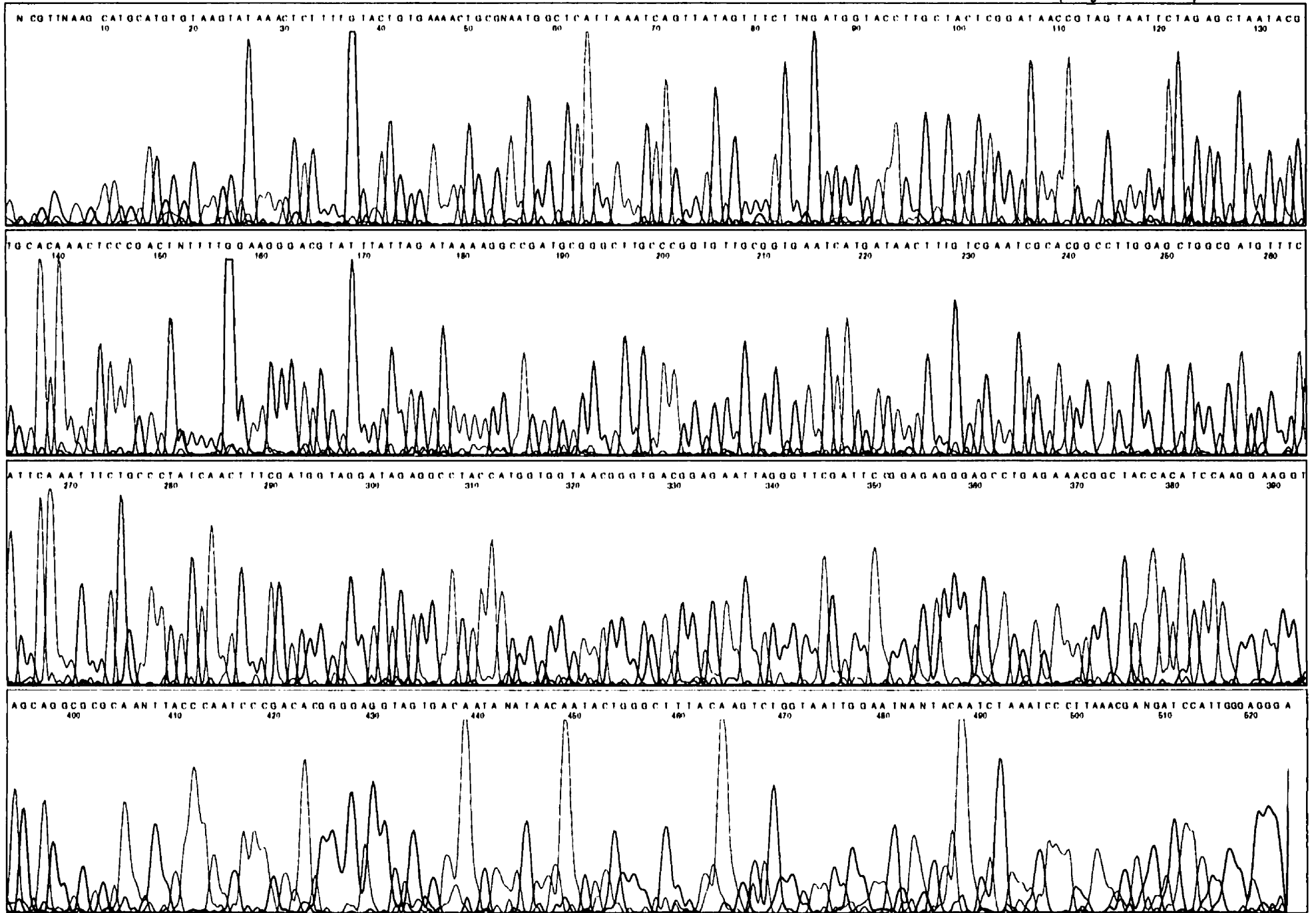


Model 377
Version 2.1.1

02•NS1.1
PATENS 1-2/1.5 1 in 5 dlutio
NS1
Lane 2

Signal G:307 A:206 T:138 C:61
DT4%Ac(A Set-AnyPrimer)
377-95020199
Points 950 to 5919 Base 1: 950

Page 1 of 1
Sun, Apr 13, 1997 2:10 PM
Mon, Oct 2, 1995 1:02 AM
Spacing: 10.39 SemiAdaptive



10



Model 377
Version 2.1.1

04•NS2.1
PATENS 1-2/1.5 1 in 5 dilutio
NS2
Lane 4

Signal G:291 A:294 T:273 C:154
DT4%Ac(A Set-AnyPrimer)
377-95020199
Points 962 to 5907 Base 1: 962

Page 1 of 1
Mon, Oct 2, 1995 10:19 AM
Mon, Oct 2, 1995 1:02 AM
Spacing: 10.40 SemiAdaptive

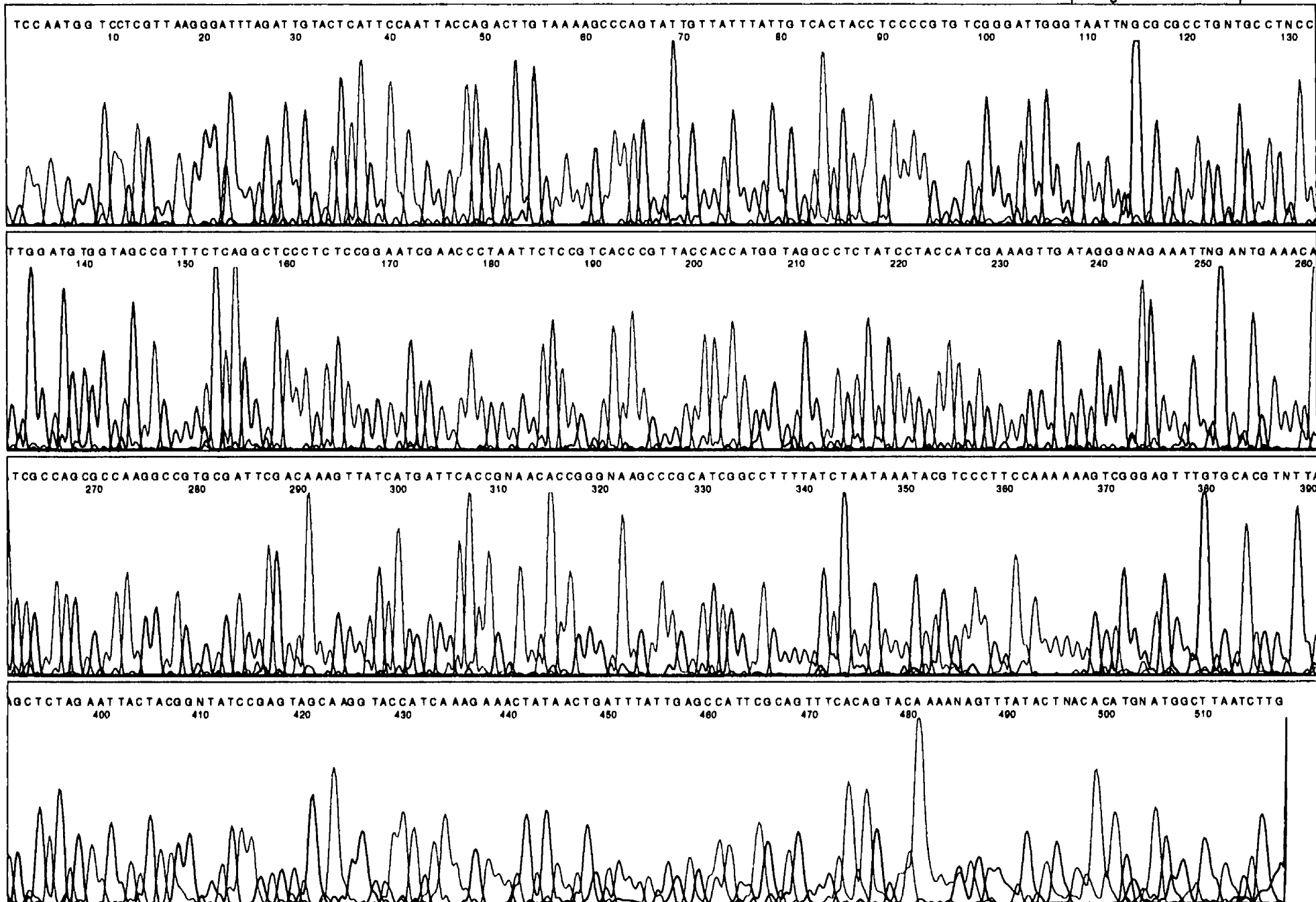


Figure 5.

18S rDNA sequence data for *Physcomitrella patens*.

The entire gene is displayed with "N" showing unsequenced regions.
The sequence was determined by double-stranded cycle sequencing using the primers listed in Table 1 (Page 2).

Base Usage

<u>Code</u>	<u>Count</u>	<u>%</u>
A	429	24
C	347	19
G	466	26
T	471	26
N	104	5
TOTAL	1818	

10	20	30	40	50	60
NNNNNNNNNN	NNNNNNNNNN	NNNNNNNNNN	NNNNNNNNAA	AGATTAAGCC	ATGCATGTGT
NNNNNNNNNN	NNNNNNNNNN	NNNNNNNNNN	NNNNNNNNTT	TCTAATTCGG	TACGTACACA
70	80	90	100	110	120
AAGTATAAAC	TCTTTTGTAC	TGTGAAACTG	CGAATGGCTC	ATTAATCAG	TTATAGTTTC
TTCATATTTG	AGAAAACATG	ACACTTTGAC	GCTTACCGAG	TAATTTAGTC	AATATCAAAG
130	140	150	160	170	180
TTTGATGGTA	CCTTGCTACT	CGGATAACCG	TAGTAATTCT	AGAGCTAATA	CGTGCACAAA
AAACTACCAT	GGAACGATGA	GCCTATTGGC	ATCATTAAAG	TCTCGATTAT	GCACGTGTTT
190	200	210	220	230	240
CTCCCGACTT	TTTTGGAAGG	GACGTATTTA	TTAGATAAAA	GGCCGATGCG	GGCTTGCCCG
GAGGGCTGAA	AAAACCTTCC	CTGCATAAAT	AATCTATTTT	CCGGCTACGC	CCGAACGGGC
250	260	270	280	290	300
GTGTTGCGGT	GAATCATGAT	AACTTTGTCG	AATCGCACGG	CCTTGGCGCT	GGCGATGTTT
CACAACGCCA	CTTAGTACTA	TTGAAACAGC	TTAGCGTGCC	GGAACCGCGA	CCGCTACAAA
310	320	330	340	350	360
CATTCAAATT	TCTGCCCTAT	CAACTTTTCGA	TGGTAGGATA	GAGGCCTACC	ATGGTGGTAA
GTAAGTTTAA	AGACGGGATA	GTTGAAAGCT	ACCATCCTAT	CTCCGGATGG	TACCACCATT
370	380	390	400	410	420
CGGGTGACGG	AGAATTAGGG	TTCGATTCCG	GAGAGGGAGC	CTGAGAAACG	GCTACCACAT
GCCCACTGCC	TCTTAATCCC	AAGCTAAGGC	CTCTCCCTCG	GACTCTTTGC	CGATGGTGTA
430	440	450	460	470	480
CCAAGGAAGG	CAGCAGGCGC	GCAAATTACC	CAATCCCGAC	ACGGGGAGGT	AGTGACAATA
GGTTCCTTCC	GTCGTCCGCG	CGTTAATGG	GTTAGGGCTG	TGCCCTCCA	TCACTGTTAT
490	500	510	520	530	540
AATAACAATA	CTGGGCTTTT	ACAAGTCTGG	TAATTGGAAT	GAGTACAATC	TAAATCCCTT
TTATTGTTAT	GACCCGAAAA	TGTTCAAGCC	ATTAACCTTA	CTCATGTTAG	ATTTAGGGAA
550	560	570	580	590	600
AACGAGGATC	CATTGGAGGN	NNNNNNNNNN	NNNNNNNNNN	GCGGTAATTC	CAGCTCCAAT
TTGCTCCTAG	GTAACCTCCN	NNNNNNNNNN	NNNNNNNNNN	CGCCATTAAG	GTCGAGGTTA
610	620	630	640	650	660
AGCGTATATT	TAAGTTGTTG	CAGTTAAAAA	GCTCGTAGTT	GGATCTTGGG	TCGGGGGGAG
TCGCATATAA	ATTCAACAAC	GTCAATTTTT	CGAGCATCAA	CCTAGAACCC	AGCCCCCTC
670	680	690	700	710	720
CGGTCCGCCC	CTTGTGGGTG	TGCACTGGTC	CACTCGGCCT	TCTTGCCGGG	GACGCGCTCC
GCCAGGCGGG	GAACACCCAC	ACGTGACCAG	GTGAGCCGGA	AGAACGGCCC	CTGCGCGAGG
730	740	750	760	770	780
TGGTTTTAAT	TAATCGGGAC	GCGGAGTCGG	CGATGTTACT	TTGAAAAAAT	TAGAGTGCTC
ACCAAAATTA	ATTAGCCCTG	CGCCTCAGCC	GCTACAATGA	AACTTTTTTA	ATCTCACGAG
790	800	810	820	830	840
AAAGCAAGCC	TATGCTCTGA	ATACATTAGC	ATGGAATAAC	GTGATAGGAC	TCTGGTCTCG
TTTCGTTCCG	ATACGAGACT	TATGTAATCG	TACCTTATTG	CACTATCCTG	AGACCAGGAC
850	860	870	880	890	900
TTCGTGTTGG	TCTTCGGGAC	CGGAGTAATG	ATTAATAGGG	ACGGTTGGGG	GCATTCGTAT
AAGCACAACC	AGAAGCCCTG	GCCTCATTAC	TAATTATCCC	TGCCAACCCC	CGTAAGCATA

910 TTCATTGTCA AAGTAACAGT	920 GAGGTGAAAT CTCCACTTTA	930 TCTTGGATTT AGAACCTAAA	940 ATGAAAGACG TACTTTCTGC	950 AACTTCTGCG TTGAAGACGC	960 AAAGCATTTG TTTCGTA AAC
970 CCAAGGATGT GGTTCCTACA	980 TTTCATTAAT AAAGTAATTA	990 CAAGAACGAA GTTCTTGCTT	1000 AGTTGGGGGC TCAACCCCCG	1010 TCGAAGACGA AGCTTCTGCT	1020 TCAGATACCG AGTCTATGGC
1030 TCCTAGTCTC AGGATCAGAG	1040 AACCATAAAC TTGGTATTTG	1050 GATGCCGACT CTACGGCTGA	1060 AGGGATTGGC TCCCTAACCG	1070 GGATGTACT CCTACAATGA	1080 TTGATGACTC AACTACTGAG
1090 CGCCAGCACC GCGGTCGTGG	1100 TTATGAGAAA AATACTCTTT	1110 TCAAAGTTTT AGTTTCAAAA	1120 TNGGTTCCGG ANCCAAGGCC	1130 GGNGAGTATG CCNCTCATA C	1140 NTCGCAAGNC NAGCGTTCNG
1150 TGAANNNNNN ACTTNNNNNN	1160 NNNNNNNNNN NNNNNNNNNN	1170 NNNNNNGCAC NNNNNNCGTG	1180 NACCAGGAGT NTGGTCCTCA	1190 GGAGCTGCGG CCTCGACGCC	1200 CTTAATTTGA GAATTA AACT
1210 CTCAACACGG GAGTTGTGCC	1220 GGAAACTTAC CCTTTGAATG	1230 CAGGTCCAGA GTCCAGGTCT	1240 CATAGTAAGG GTATCATTCC	1250 ATTGACAGAT TAACTGTCTA	1260 TGAGAGCTCT ACTCTCGAGA
1270 TTCTTGATTC AAGAACTAAG	1280 TATGGGTGGT ATACCCACCA	1290 GGTGCATGGC CCACGTACCG	1300 CGTTCCTAGT GCAAGAATCA	1310 TGGTGGAGTG ACCACCTCAC	1320 ATTTGTCTGG TAAACAGACC
1330 TTAATTCCGT AATTAAGGCA	1340 TAACGAACGA ATTGCTTGCT	1350 GACCTCAGCC CTGGAGTCGG	1360 TGCTAACTAG ACGATTGATC	1370 TTACGCGAAG AATGCGCTTC	1380 GATTTTTTCC CTAAAAAAGG
1390 TTTGCGGCCA AAACGCCGGT	1400 ACTTCTTAGA TGAAGAATCT	1410 GGGACTATCG CCCTGATAGC	1420 GCGTCTAGCC CGCAGATCGG	1430 GATGGAAGTT CTACCTTCAA	1440 TGAGGCAATA ACTCCGTTAT
1450 ACAGGTCTGT TGTCAGACA	1460 GATGCCCTTA CTACGGGAAT	1470 GATGTTCTGG CTACAAGACC	1480 GCCGCACGCG CGGCGTGCGC	1490 CGCTACACTG GCGATGTGAC	1500 ATGAATTCAA TACTTAAGTT
1510 CGAGTTTATA GCTCAAATAT	1520 ACCTGGACCG TGGACCTGGC	1530 ATAGGTCTGG TATCCAGACC	1540 GTAATCTTTT CATTAGAAAA	1550 GAAATTTTCA CTTTAAAGTA	1560 CGTGATGGGG GCACTACCCC
1570 ATAGATCATT TATCTAGTAA	1580 GCAATTATTG CGTTAATAAC	1590 ATCTTCAACG TAGAAGTTGC	1600 AGGAATTCCT TCCTTAAGGA	1610 AGTAAGCGCG TCATTGCGCG	1620 AGTCATCAGC TCAGTAGTCG
1630 TCGCGTTGAC AGCGCAACTG	1640 TACGTCCTG ATGCAGGGAC	1650 CCCTTTGTAC GGGAAACATG	1660 ACACCGCCCC TGTGGCGGGC	1670 TCGCTCCTAC AGCGAGGATG	1680 CGATTGAATG GCTAACTTAC
1690 GTCCGGTGAA CAGGCCACTT	1700 GTTTTCGGAT CAAAA GCCTA	1710 TGCGGCGATG ACGCCGCTAC	1720 CCGGCGGTTT GGCCGCCAAG	1730 GCCGCCGGTG CGGCCGCCAC	1740 ACGTTGGGAG TGCAACCCTC
1750 AAGTTCATTA TTCAAGTAAT	1760 AACCTTATCA TTGGAATAGT	1770 TTTAGAGGAA AAATCTCCTT	1780 GGAGAAGTCG CCTCTTCAGC	1790 TAACAAGGTT ATTGTTCAA	1800 TCCGTAGGTN AGGCATCCAN
1810 NNNNNNNNNN NNNNNNNNNN	1820 NNNNNNN... NNNNNNN...	1830	1840	1850	1860

Figure 6.

18S rDNA sequence data for *Megaceros aenigmaticus*.

The entire gene is displayed with "N" showing unsequenced regions.
The sequence was determined by double-stranded cycle sequencing using the primers listed in Table 1 (Page 2).

Base Usage

<u>Code</u>	<u>Count</u>	<u>%</u>
A	448	25
C	344	19
G	457	25
T	464	25
N	111	6
TOTAL	1824	

10	20	30	40	50	60
NNNNNNNNNN	NNNNNNNNNN	NNNNNNNNNN	NNNNNNNNNA	AGATNNA GCC	ATGCATGTGT
NNNNNNNNNN	NNNNNNNNNN	NNNNNNNNNN	NNNNNNNNNT	TCTANNTCGG	TACGTACACA
70	80	90	100	110	120
AAGTATAAAC	ACTTTTGTAC	TGTGAAACTG	CGAATGGCTC	ATTAAATCAG	TTATAGTTTC
TTCATATTTG	TGAAAACATG	ACACTTTGAC	GCTTACCGAG	TAATTTAGTC	AATATCAAAG
130	140	150	160	170	180
TTTGATGGTA	CCTTGCTACT	CGGATAACCG	TAGTAATTCT	AGAGCTAATA	CGTGCAACAA
AAACTACCAT	GGAACGATGA	GCCTATTGGC	ATCATTAAAG	TCTCGATTAT	GCACGTTGTT
190	200	210	220	230	240
CTCCCCGACTT	CTGGGANGGG	GCGTATTTAT	TAGATATAAG	GCCGATATGG	GTCTTGATCC
GAGGGCTGAA	GACCCTNCCC	CGCATAAATA	ATCTATATTC	CGGCTATACC	CAGAACTAGG
250	260	270	280	290	300
CGGTATTACG	GTGAATCATG	ATAACTCCTC	GAATGCTCAA	CGGCCTCTGG	CGCTGGCGAT
GCCATAATGC	CACTTAGTAC	TATTGAGGAG	CTTACGAGTT	GCCGGAGACC	GCGACCGCTA
310	320	330	340	350	360
ATTTCAATTC	AATTTCTGCC	CTATATAACT	TTCGATGGTA	GGATAGAGGC	CTACCATGGT
TAAAGTAAGT	TAAAGACGGG	GATATATTGA	AAGCTACCAT	CCTATCTCCG	GATGGTACCA
370	380	390	400	410	420
GGTAACGGGT	GACGGAGAAT	TAGGGTTCGA	TTCCGGAGAG	GGAGCCTGAG	AAACGGCTAC
CCATTGCCCA	CTGCCTCTTA	ATCCCAAGCT	AAGGCCTCTC	CCTCGGACTC	TTTGCCGATG
430	440	450	460	470	480
CACATCCAAG	GAAGGCAGCA	GGCGCGCAAA	TTACCCAATC	CCGACACGGG	GAGGTAGTGA
GTGTAGGTTT	CTTCCGTCGT	CCGCGCGTTT	AATGGGTTAG	GGCTGTGCC	CTCCATCACT
490	500	510	520	530	540
CAATAAATAA	CAATACTGGG	CTTTTACAAG	TCTGGTAATT	GGAATGAGTA	CAATCTAAAT
GTTATTTATT	GTTATGACCC	GAAAAATGTT	AGACCATTAA	CCTTACTCAT	GTTAGATTTA
550	560	570	580	590	600
CCCTNAACGA	GGATCCATTG	GAGGNNNNNN	NNNNNNNNNN	NNNNNGCGGT	AATTCCAGCT
GGGANTTGCT	CCTAGGTAAC	CTCCNNNNNN	NNNNNNNNNN	NNNNNCGCCA	TTAAGGTCSA
610	620	630	640	650	660
CCNATAGCGN	NTAATAAAGT	TGTTGCAGTT	AAAAAGCTCG	TAGTTGGATC	TCGGGGTGGG
GGNTATCGCN	NATTATTTCA	ACAACGTCAA	TTTTTCGAGC	ATCAACCTAG	AGCCCCACCC
670	680	690	700	710	720
GTGACTGGTC	CGCCTCTCAA	GGTGTGGTAC	TGGGTGACTC	CANTCCTTCT	GGTCGGGGAC
CACTGACCAG	GCGGAGAGTT	CCACACCATG	ACCCACTGAG	GTNAGGAAGA	CCAGCCCCTG
730	740	750	760	770	780
GATGCTCCTG	GACTAAACTG	CCTGGGACGC	GGAGTCGACG	ATGTTACTTT	GAAAAAATTA
CTACGAGGAC	CTGATTTGAC	GGACCCCTGC	CCTCAGCTGC	TACAATGAAA	CTTTTTTAAT
790	800	810	820	830	840
GAGTGCTCAA	AGCAAGCCTC	ATGCTCTGAA	TACATTAGCA	TGGAATAACG	TGATAGGACT
CTCAGGAGTT	TCGTTCCGGAG	TACGAGACTT	ATGTAATCGT	ACCTTATTGC	ACTATCCTGA
850	860	870	880	890	900
CCTGGTCCTA	TTGTGTTGNT	CTACGGGATC	GGAGTAATGA	TTAATAGGGA	CAGTTGGGGG
GGACCAGGAT	AACACAACNA	GATGCCCTAG	CCTCATTACT	AATTATCCCT	GTCAACCCCC

910 CATTTCGATT GTAAGCATAA	920 TCATTGTCAG AGTAACAGTC	930 AGGTGAAATT TCCACTTTAA	940 CTTGGATTTA GAACCTAAAT	950 TGAAAGACGA ACTTTCTGCT	960 ACTTCTGCCA TGAAGACGCT
970 AAGCATTTCG TTCGTAAACG	980 CAAGGCATGT GTTCCGTACA	990 TTTCATTAAT AAAGTAATTA	1000 CAAGAACGAA GTTCTTGCTT	1010 AGTTGGGGGC TCAACCCCGG	1020 TCGAAGACGA AGCTTCTGCT
1030 TCAGATACCG AGTCTATGGC	1040 TCCTAGTCTC AGGATCAGAG	1050 AACCATAAAC TTGGTATTTG	1060 GATGCCGACT CTACGGCTGA	1070 AGGGATTGGC TCCCTAACCG	1080 GGATGTTAAT CCTACAATTA
1090 TAGATGACTC ATCTACTGAG	1100 CGCCAGCACC GCGGTCGTGG	1110 TTATGAGAAA AATACTCTTT	1120 TCAAAGTTTT AGTTTCAAAA	1130 TGGNTTCCGG ACCNAAAGCC	1140 GGGNAGTATG CCCNTCATAC
1150 NTCGCAAGNC NAGCGTTCNG	1160 TGAANTNNNN ACTTNANNNN	1170 NNNNNNNNNN NNNNNNNNNN	1180 NNNNNNGNAN NNNNNNCNTN	1190 CACNAGGAGT GTGNTCCTCA	1200 GGAGCTGCCG CCTCGACGCC
1210 CTTAATTTGA GAATTAAGCT	1220 CTCAACACNG GAGTTGTGNC	1230 GGAAACTTAC CCTTTGAATG	1240 CAGGTCCAGA GTCCAGGTCT	1250 CATAGTAAGG GTATCATTCC	1260 ATTGACAGAT TAACTGTCTA
1270 TGAGAGCTCT ACTCTCGAGA	1280 TTCTTGATTC AAGAACTAAG	1290 TATGGGTGGT ATACCCACCA	1300 GGTGCATGGC CCACGTACCG	1310 CGTTCCTAGT GCAAGAATCA	1320 TGGTGGAGTG ACCACCTCAC
1330 ATTTGTCTGG TAAACAGACC	1340 TTAATTCCTG AATTAAGGCA	1350 TAACGAACGA ATTGCTTGCT	1360 GACCTCAGCC CTGGAGTCGG	1370 TGCTAACTAG ACGATTGATC	1380 TTACACGAAG AATGTGCTTC
1390 AATCTTCTTC TTAGAAAGAAG	1400 GTGGCCAACT CACCGTTGA	1410 TCTTAGAGGG AGAATCTCCC	1420 ACTATTTGCG TGATAAACGC	1430 TCTAGCGAAT AGATCGCTTA	1440 GGAAGTTTGA CCTTCAAACCT
1450 GGCAATAACA CCGTTATTGT	1460 GGTCTGTGAT CCAGACACTA	1470 GCCCTTAGAT CGGGAATCTA	1480 GTTCTGGGCC CAAGACCCGG	1490 GCACGCGCGC CGTGCGCGCG	1500 TACACTGATG ATGTGACTAC
1510 AATTCAACGA TTAAGTTGCT	1520 GTTTATAACC CAAATATTGG	1530 TGGGCCGAGA ACCCGGCTCT	1540 GGTTTGGGTA CCAAACCCAT	1550 ATCTTTTGAA TAGAAAACCT	1560 ATTTCATCGT TAAAGTAGCA
1570 GATGGGGATA CTACCCCTAT	1580 GATCATTGCA CTAGTAACGT	1590 ATTATTGATC TAATAACTAG	1600 TTCAACGAGG AAGTTGCTCC	1610 AATTCCTAGT TTAAGGATCA	1620 AAGCGCGAGT TTCGCGCTCA
1630 CATCAGCTCG GTAGTCGAGC	1640 CGTTGACTAC GCAACTGATG	1650 GTCCCTGCCC CAGGGACGGG	1660 TTTGTAACA AAACATGTGT	1670 CCGCCCGTCG GGCGGGCAGC	1680 CTCCTACCGA GAGGATGGCT
1690 TTGAATGGTC AACTTACCAG	1700 CGGTGAAGTT GCCACTTCAA	1710 TTCGGATTGC AAGCCTAACG	1720 GGCGACACCG CCGCTGTGGC	1730 GCGGGTAACT CGCCATTGA	1740 GCCGGCGACG CGGCCGCTGC
1750 TTGTGAGAAG AACACTCTTC	1760 TTCATTA AAC AAGTAATTTG	1770 CTTATCATT GAATAGTAAA	1780 AGAGGAAGGA TCTCCTTCT	1790 GAAGTCGTAA CTTCAGCATT	1800 CAAGGTTTCC GTTCCAAAGG
1810 GTAGGTGANN CATCCACTNN	1820 CTGNNNNNNN GACNNNNNNN	1830 NNNN..... NNNN.....	1840	1850	1860

Figure 7.

18S rDNA sequence data for *Notophylas breutelu*.

The entire gene is displayed with "N" showing unsequenced regions.
The sequence was determined by double-stranded cycle sequencing using the primers listed in Table 1 (Page 2).

Base Usage

<u>Code</u>	<u>Count</u>	<u>%</u>
A	455	25
C	359	20
G	473	26
T	444	24
N	102	5
TOTAL	1833	

10 NNNNNNNNNN NNNNNNNNNN	20 NNNNNNNNNN NNNNNNNNNN	30 NNNNNNNNNN NNNNNNNNNN	40 NNNNNNNNAA NNNNNNNNTT	50 AGATNAAGCC TCTANTTCGG	60 ATGCATGTGT TACGTACACA
70 AAGTATAAAC TTCATATTTG	80 TCTTATGTAC AGAATACATG	90 TGAGAAACTG ACTCTTTGAC	100 AGAATGGCTC TCTTACCGAG	110 ATTAAATCAG TAATTTAGTC	120 TGATAGTATC ACTATCATAG
130 TGAGAGGGCA ACTCTCCCGT	140 CCTCGCTACG GGAGCGATGC	150 GCGGATAACC CGCCTATTGG	160 GGAGATAATT CCTCTATTAA	170 CTAGAGCTAA GATCTCGATT	180 TACGTGCAAA ATGCACGTTT
190 AACTCCCGAC TTGAGGGCTG	200 TCTTGGGGGG AGAACCCCCC	210 GAAGGGTTTT CTTCCCAAAA	220 TTTTAGATAA AAAATCTATT	230 AGAGGACGAT TCTCCTGCTA	240 GCGGGCTTGA CGCCCCAACT
250 CCCGGTATTC GGGCCATAAG	260 ACGGTGAATC TGCCACTTAG	270 ATGGATAACT TACCTATTGA	280 CCTCGAATCG GGAGCTTAGC	290 CACGGCCCTG GTGCCGGGAC	300 GCGCTGGCGA CGCGACCGCT
310 TGTTTCATTC ACAAAGTAAG	320 AAATTCCTGC TTTAAGGACG	330 CCTATCAACT GGATAGTTGA	340 TTCGATGGTA AAGCTACCAT	350 GGATAGAGGC CCTATCTCCG	360 CTACCATGGT GATGGTACCA
370 GGTAACGGGT CCATTGCCCA	380 GACGGAGAAT CTGCCTCTTA	390 TAGGGTTCTGA ATCCCAAGCT	400 TTCCGGAGAG AAGGCCTCTC	410 GGAACCTGAG CCTTGGACTC	420 AAACGGCTAC TTTGCCGATG
430 CACATCCAAG GTGTAGGTTT	440 GAAGGCAGCA CTTCCGTCGT	450 GGCGCCCAAA CCGCGGGTTT	460 TTACCCAATC AATGGGTTAG	470 CCGACACGGG GGCTGTGCC	480 GAGGTAGTGA CTCCATCACT
490 CAATAAATAA GTTATTTATT	500 CAATACTGGG GTTATGACCC	510 CCTTTACAAG GGAAATGTTC	520 TCCTGGTAAT AGGACCATTA	530 TGGAATGAGT ACCTTACTCA	540 ACAATCTAAA TGTTAGATTT
550 TCCCTTAACG AGGGAATTGC	560 AGGATCCATT TCCTAGGTAA	570 GGAGGNNNNN CCTCCNNNNN	580 NNNNNNNNNN NNNNNNNNNN	590 NNNNNNNGCG NNNNNNNCGCC	600 TAATTCCAGC ATTAAGGTCC
610 TCCAATAGCG AGGTTATCGC	620 TACAATAAAG ATGTTATTTT	630 TTGTTGCACT AACAAAGTCA	640 TAAAAAGGTC ATTTTTCCAG	650 GTAGTTGGAT CATCAACCTA	660 CTCGGGGCGG GAGCCCCGCC
670 GGTGATTGGT CCACTAACCA	680 CCGCCTCCNG GGCGGAGGNC	690 AGGGGTCTAC TCCCCAGATG	700 TAGGGACCTC ATCCCTGGAG	710 ACCCCTCTCC TGGGGAGAGG	720 TCGGGAGCAT AGCCCTCGTA
730 GCTCCTGTTT CGAGGACAAG	740 TACCTTAGCT ATGGAATCGA	750 TGGTACGGGG ACCATGCCCC	760 AGTGCACGCA TCACGCTGCT	770 TGTTTCTTTG ACAAAGAAAC	780 GAAAAAATTA CTTTTTTAAT
790 GGAGTTCTCA CCTCAAGAGT	800 AAGGAAGCCA TTCCTTCGGT	810 TCATGCTCTA AGTACGAGAT	820 ATTACATAAG TAATGTATTC	830 AATGGATTAA TTACCTAATT	840 CGATTATAGG GCTAATATCC
850 ACTCTGGTCC TGAGACCAGG	860 TATTGGGTTG ATAACCCAAC	870 GTCTCCGGGC CAGAGGCCCG	880 ATCGGAGTAA TAGCCTCATT	890 TGATTAATAG ACTAATTATC	900 GGACAGTTGG CCTGTCAACC

910 GGGCATTCGT CCCCGTAAGCA	920 ATTTTCGATTG TAAAGCTAAC	930 TCAGAGGTGA AGTCTCCACT	940 AATTCTTGGGA TTAAGAACCT	950 TTTATGAAAG AAATACTTTC	960 ACGAACTTCT TGCTTGAAGA
970 GCGAAAGCAT CGCTTTCGTA	980 TTGCCAAGGA AACGGTTCCT	990 TGTTTTTCATT ACAAAAGTAA	1000 AATCAAGAAC TTAGTTCCTG	1010 GAAAGTTGGG CTTCAACCC	1020 GGCTCGAAGA CCGAGCTTCT
1030 CGATCAGATA GCTAGTCTAT	1040 CCGTCCTAGT GGCAGGATCA	1050 CTCAACCATA GAGTTGGTAT	1060 AACGATGCCG TTGCTACGGC	1070 ACTAGGGATT TGATCCCTAA	1080 GGCGGGTGTT CCGCCACAA
1090 AATAAGATTG TTATTCTAAC	1100 ACCCCCCGG TGGGGGGGCC	1110 CCACCCTTAT GGTGGGAATA	1120 TGAAGAAACC ACTTCTTTGG	1130 AAAGTTTTTG TTTCAAAAAC	1140 GGTTCGGGG CCAAGGCCCC
1150 GATATGTTGC CTATACAACG	1160 AAGCTGANNN TTCGACTNNN	1170 NNNNNNNNNN NNNNNNNNNN	1180 NNNNNNNNNN NNNNNNNNNN	1190 GNANCA CNAG CNTNGTGNTC	1200 GAGTGGAGCT CTCACCTCGA
1210 GCGGCTTAAT CGCCGAATTA	1220 TTGACTCAAC AACTGAGTTG	1230 ACGGGGAAAC TGCCCTTTG	1240 TCACCAGGTC AGTGGTCCAG	1250 CAGACATAGT GTCTGTATCA	1260 AAGGATTGAC TTCCTAACTG
1270 AGATTGAGAG TCTAACTCTC	1280 CTCTTCTTG GAGAAAGAAC	1290 ATTCTATGGG TAAGATACCC	1300 TGGTGGTGCA ACCACCACGT	1310 TGGCCGTTCT ACCGGCAAGA	1320 TAGTTGGTGG ATCAACCACC
1330 AGTGATTTGT TCACTAAACA	1340 CTGGTTAATT GACCAATTA	1350 CCGTAAACGA GGCAATTGCT	1360 ACGAGACCTC TGCTCTGGAG	1370 AGCCTGCTAA TCGGACGATT	1380 CTAGTTACAC GATCAATGTG
1390 GAAGAACCTT CTTCTTGAA	1400 CTGCGGGGTC GACGCCCCAG	1410 AACTTCTTAG TTGAAGAATC	1420 AGGGACTATT TCCCTGATAA	1430 TGCCTAGT ACGCAGATCA	1440 GAATAGAAGT CTTATCTTCA
1450 TTGAGGAAAT AACTCCTTTA	1460 AACAGGTCTG TTGTCCAGAC	1470 TGATGCCCTT ACTACGGGAA	1480 AGATGTTCTG TCTACAAGAC	1490 GGCCGCACGC CCGGCGTGCG	1500 GCGCTACACT CGCGATGTGA
1510 GATGAATTCA CTACTTAAGT	1520 ACGAGTTTAT TGCTCAAATA	1530 AACCTGGGCC TTGGACCCGG	1540 GAGAGGTCTG CTCTCCAGAC	1550 GGTAATCTGC CCATTAGACG	1560 TGAAATTTCA ACTTTAAAGT
1570 TCGTGATGGG AGCACTACCC	1580 GATAGATCAT CTATCTAGTA	1590 TGAAATTAAT ACTTTAATAA	1600 GCTCTCCAAC CGAGAGGTTG	1610 GAGGAATTC CTCCTTAAGG	1620 TAGTAAGCGC ATCATTCCGG
1630 GAATCATCAG CTTAGTAGTC	1640 CTCGCGTTGA GAGCGCAACT	1650 CTACGTCCCT GATGCAGGGA	1660 GCCCTTTGTA CGGGAAACAT	1670 ACACACCGCC TGTGTGGCGG	1680 CGTCGTTCT GCAGCAAGGA
1690 ACCGATTGAT TGGCTAACTA	1700 TGGTCCGGGG ACCAGGCCCC	1710 AAGTTTTTCGG TTCAAAGCC	1720 ATTGGTGGCG TAACCACCGC	1730 ACACCGGGGG TGTGGCCCCC	1740 GTCACCGCCG CAGTGGCGGG
1750 GCGACGTTGT CGCTGCAACA	1760 GAGAAGTTCA CTCTTCAAGT	1770 TCAACCCTTA AGTTGGGAAT	1780 TCATTTAGAG AGTAAATCTC	1790 GAAGGAGAAG CTTCTCTTTC	1800 TCGTCGTAAC AGCAGCATTG
1810 AAGGTTTCCG TTCCAAAGGC	1820 TAGGTGANCN ATCCACTNGN	1830 NNNNNNNNNN NNNNNNNNNN	1840 NNN. NNN.	1850	1860

Figure 8.

18S rDNA sequence data for *Pteris vittata*.

The entire gene is displayed with "N" showing unsequenced regions.
The sequence was determined by double-stranded cycle sequencing using the primers listed in Table 1 (Page 2).

Base Usage

<u>Code</u>	<u>Count</u>	<u>%</u>
A	424	23
C	363	20
G	459	25
T	446	25
N	118	7
TOTAL	1810	

10 NNNNNNNNNN NNNNNNNNNN	20 NNNNNNNNNN NNNNNNNNNN	30 NNNNNNNNNN NNNNNNNNNN	40 NNNNNNNNAA NNNNNNNNTT	50 AGNTTAAGCC TCNAATTCGG	60 ATGCATGTGT TACGTACACA
70 AAGTATAAAC TTCATATTTG	80 TCTTTTGTAC AGAAAAATG	90 TGTGAAACTG ACACTTTGAC	100 CGAATGGCTC GCTTACCGAG	110 ATTAAATCAG TAATTTAGTC	120 TTATAGTTTC AATATCAAAG
130 TTTGATGGTA AAACTACCAT	140 CCTNGCTACT GGANCGATGA	150 CGGATAACCG GCCTATTGGC	160 TAGTAATTCT ATCATTAAAG	170 AGAGCTAATA TCTCGATTAT	180 CGTGCACCAA GCACGTGGTT
190 ATCCCGACTT TAGGGCTGAA	200 CTGGAAGGGA GACCTTCCCT	210 CGCATTTATT GCGTAAATAA	220 AGATAAAAGG TCTATTTTCC	230 CCGATGCGGG GGCTACGCC	240 CTCGCCCGGT GAGCGGGCCA
250 AATGCGGTGA TTACGCCACT	260 ATCATGATAA TAGTACTATT	270 CTTCCCGAAT GAAGGGCTTA	280 CGCACGGCCT GCGTGCCGGA	290 TGGCGCCGGC ACCGCGGCCG	300 GATGCTTCAT CTACGAAGTA
310 TCAAATTTCT AGTTTAAAGA	320 GCCCTATCAA CGGGATAGTT	330 CTTTCGATGG GAAAGCTACC	340 TAGGATAGAG ATCCTATCTC	350 GCCTACCATG CGGATGGTAC	360 GTGGTGACGG CACCACTGCC
370 GTGACGGAGA CACTGCCTCT	380 ATTAGGGTTC TAATCCCAAG	390 GATTCCGGAG CTAAGGCCTC	400 AGGGAGCCTG TCCCTCGGAC	410 AGAAACGGCT TCTTTGCCGA	420 ACCACATCCA TGGTGTAGGT
430 AGGAAGGCAG TCCTTCCGTC	440 CAGGCGCGCA GTCCGCGCGT	450 AATTACCCAA TTAATGGGTT	460 TCCCGACACG AGGGCTGTGC	470 GGGAGGTAGT CCCTCCATCA	480 GACAATAAAT CTGTTATTTA
490 AACAACTAGT TTGTTATGAC	500 GGCTTTTTCA CCGAAAAAGT	510 AGTCTGGTAA TCAGACCATT	520 TTGGAATGAG AACCTTACTC	530 TACAATCTAA ATGTTAGATT	540 ATCCCTTAAC TAGGGAATTG
550 GAGGATCCAT CTCCTAGGTA	560 TGGAGGGCAA ACCTCCCGTT	570 NNNNNNNNNN NNNNNNNNNN	580 NNNNNNNNNG NNNNNNNNNC	590 GTAATTCAG CATTAAGGTC	600 CTCCAATAGC GAGGTTATCG
610 GTATATTTAA CATATAAATT	620 GTTGTTGCAG CAACAACGTC	630 TTAAAAAGCT AATTTTTCGA	640 CGTAGTTGGA GCATCAACCT	650 TCTCGGGGCG AGAGCCCCGC	660 GGGCGAGCCG CCCGCTCGCC
670 TCCGCCTCCT AGGCGGAGGA	680 TTGGTGTGCA AACCACACGT	690 CTGGTCGCTC GACCAGCGAG	700 CGCCCTTCT GCGGGAAAGA	710 GTCGGGGACG CAGCCCCCTGC	720 CGCTCCTGGC GCGAGGACCG
730 CTTAATTGGC GAATTAACCG	740 TGGGACGCGG ACCTGCGCC	750 ATTCGGCGAT TAAGCCGCTA	760 GTTACTTTGA CAATGAAACT	770 AAAAATTAGA TTTTTAATCT	780 GTGCTCAAAG CACGAGTTTC
790 CAAGCCTATG GTTCCGATAC	800 CTCTGAATAC GAGACTTATG	810 ATTAGCATGG TAATCGTACC	820 AATAACGCGA TTATTGCGCT	830 TAGGACTCTG ATCCTGAGAC	840 GTCCTATTGT CAGGATAACA
850 GTTGGTCTTC CAACCAGAAG	860 GGGACCGGAG CCCTGGCCTC	870 TAATGATTAA ATTACTAATT	880 TAGGGACGGT ATCCCTGCCA	890 TGGGGGCATT ACCCCCGTAA	900 CGTATTTTCAT GCATAAAGTA

910 TGTCAGAGGT ACAGTCTCCA	920 GAAATTCTTG CTTTAAGAAC	930 GATTTATGAA CTAAATACTT	940 AGACGAACTA TCTGCTTGAT	950 CTGCGAAAAGC GACGCTTTCG	960 ATTTGCCAAG TAAACGGTTC
970 GATGTTTTCA CTACAAAAGT	980 TTAATCAAGA AATTAGTTCT	990 ACGAAAGTTG TGCTTTCAAC	1000 GGGGCTCGAA CCCCGAGCTT	1010 GACGATCAGA CTGCTAGTCT	1020 TACCGTCTTA ATGGCAGGAT
1030 GTCTCAACCA CAGAGTTGGT	1040 TAAACGATGC ATTTGCTACG	1050 CGACTAGGGA GCTGATCCCT	1060 TTGGCGGATG AACCGCCTAC	1070 TTACTTTGAT AATGAAACTA	1080 GACTCCGCCA CTGAGGCGGT
1090 GCACCTTATG CGTGGAATAC	1100 AGAAATCAAA TCTTTAGTTT	1110 GTTTTTGTTT CAAAAACCAA	1120 TCCGGGGGGA AGGCCCCCTT	1130 GTATGGTCGC CATACCAGCG	1140 AAGGCTGAAN TTCCGACTTN
1150 NNNNNNNNNN NNNNNNNNNN	1160 NNNNNNNNNN NNNNNNNNNN	1170 NNNNNCACCA NNNNNGTGTT	1180 GGAGTGGAGC CCTCACCTCG	1190 CTGCGGCTTA GACGCCGAAT	1200 ATTTGACTCA TAAACTGAGT
1210 ACACGGGGAA TGTGCCCTT	1220 ACTTACCAGG TGAATGGTCC	1230 TCCAGACATA AGGTCTGTAT	1240 GTAAGGATTG CATTCTAAC	1250 ACAGATTGAG TGTCTAACTC	1260 AGCTCTTTCT TCGAGAAAAG
1270 TGATTCTATG ACTAAGATAC	1280 GGTGGTGGTG CCACCACCAC	1290 CATGGCCGTT GTACCGGCAA	1300 CTTAGTTGGT GAATCAACCA	1310 GGAGTGATTT CCTCACTAAA	1320 GTCTGGTTAA CAGACCAATT
1330 TTCCGTTAAC AAGGCAATTG	1340 GAACGAGACC CTTGCTCTGG	1350 TCAGCCTGCT AGTCGGACGA	1360 AACTAGTTAC TTGATCAATG	1370 ACGAAGGATC TGCTTCCTAG	1380 CTCTTCGTGG GAGAAGCACC
1390 CCAACTTCTT GGTTGAAGAA	1400 AGAGGGACTA TCTCCCTGAT	1410 TGGCCGTCTA ACCGGCAGAT	1420 GGCCATGGAA CCGGTACCTT	1430 GTTTGAGGCA CAAACCTCCG	1440 ATAACAGGTC TATTGTCCAG
1450 TGTGATGCCC ACACTACGGG	1460 TTAGATGTTC AATCTACAAG	1470 TGGGCCGCAC ACCCGGCGTG	1480 GCGCGCTACA CGCGCGATGT	1490 CTGATGAATT GACTACTTAA	1500 CAACGAGTTT GTTGCTCAAA
1510 ACCACCTGGG TGTTGGACCC	1520 CCGACAGGCC GGCTGTCCGG	1530 CGGGTAATCT GCCATTAGA	1540 TTTGAAATTT AAACTTTAAA	1550 CATCGTGATG GTAGCACTAC	1560 GGGATAGATC CCCTATCTAG
1570 ATTGCAATTA TAACGTTAAT	1580 TTGATCTTCA AACTAGAAGT	1590 ACGAGGAATT TGCTCCTTAA	1600 CCTAGTAAGC GGATCATTTC	1610 GCGAGTCATC CGCTCAGTAG	1620 AGCTCGCGTT TCGAGCGCAA
1630 GACACGGTCC CTGTGCCAGG	1640 CTGCCCTTTG GACGGGAAAC	1650 TACACACCGC ATGTGTGGCG	1660 CCGTCGCTCC GGCAGCGAGG	1670 TACCGATTGA ATGGCTAACT	1680 ATGGTCCGGT TACCAGGCCA
1690 GAAGTTTTCG CTTCAAAAAGC	1700 GATCGCGACG CTAGCGCTGC	1710 ACGCTGGCGG TGCGACCGCC	1720 NTTCTCCGCC NAAGAGGCGG	1730 GGTGACGTTG CCTGCAAC	1740 TGAGAAGTTC ACTCTTCAAG
1750 ATTAACCTTA TAATTGGAAT	1760 TCATTTAGAG AGTAAATCTC	1770 GAAGGAGAAG CTTCCTCTTC	1780 TCGTANCAGN AGCATNGTCN	1790 NNNNNNNNNN NNNNNNNNNN	1800 NNNNNNNNNN NNNNNNNNNN
1810 NNNNNNNNNN NNNNNNNNNN	1820	1830	1840	1850	1860

Figure 9.

18S rDNA sequence data for *Sphagnum palustre*.

The entire gene is displayed with "N" showing unsequenced regions.
The sequence was determined by double-stranded cycle sequencing using the primers listed in Table 1 (Page 2).

Base Usage

<u>Code</u>	<u>Count</u>	<u>%</u>
A	418	23
C	380	21
G	487	27
T	432	24
N	103	5
TOTAL	1820	

10	20	30	40	50	60
NNNNNNNNNN	NNNNNNNNNN	NNNNNNNNNN	NNNNNNNNA	AGNNTNAGCC	ATGCATGTGT
NNNNNNNNNN	NNNNNNNNNN	NNNNNNNNNN	NNNNNNNNNTT	TCNNANTCGG	TACGTACACA
70	80	90	100	110	120
AAGTATAAAC	TCTTTTGTAC	CGTGAAACTG	CGAATGGCTC	ATTAAATCAG	TTATAGTTTC
TTCATATTTG	AGAAAACATG	GCACTTTGAC	GCTTACCAG	TAATTTAGTC	AATATCAAAG
130	140	150	160	170	180
TTTGATGGTA	CCTTGCTACT	CGGATAACCG	TAGTAATTCT	AGAGCTAATA	CGTGCACCGA
AAACTACCAT	GGAACGATGA	GCCTATTGGC	ATCATTAAGA	TCTCGATTAT	GCACGTGGCT
190	200	210	220	230	240
CTCCCACCC	CGCAAGGGGA	AGGGACGTAT	TTATTAGATA	AAAGGCCGAT	GCGGGCTCGC
GAGGGCTGGG	GCGTTCCCCT	TCCCTGCATA	AATAATCTAT	TTTCCGGCTA	CGCCCCGAGC
250	260	270	280	290	300
CCGGTATTGC	GGTGAATCAT	GATAACTCGT	CGAATCGCAC	GGCCCTGGCG	CCGGCGATGT
GGCCATAACG	CCACTTAGTA	CTATTGAGCA	GCTTAGCGTG	CCGGGACCGC	GGCCGCTACA
310	320	330	340	350	360
TTCAATCAAA	TTTCTGCCCT	ATCAACTTTC	GACGGTAGGA	TAGAGGCCTA	CCGTGGTGGT
AAGTAAAGTT	AAAGACGGGA	TAGTTGAAAG	CTGCCATCCT	ATCTCCGGAT	GGCACCACCA
370	380	390	400	410	420
GACGGGTGAC	GGAGAATTAG	GGTTCGATTC	CGGAGAGGGA	GCCTGAGAAA	CGGCTACCAC
CTGCCCACTG	CCTCTTAATC	CCAAGCTAAG	GCCTCTCCCT	CGGACTCTTT	GCCGATGGTG
430	440	450	460	470	480
ATCCAAGGAA	GGCAGCAGGC	GCGCAAATTA	CCCAATCCCG	ACACGGGGAG	GTAGTGACAA
TAGGTTCCCT	CCGTCGTCCG	CGCGTTAAT	GGGTTAGGGC	TGTGCCCTC	CATCACTGTT
490	500	510	520	530	540
TAAATAACGA	TACCGGGCTT	TTACAAGTCT	GGTAATCGGA	ATGAGTACAA	TCTAAATCCC
ATTTATTGCT	ATGGCCCGAA	AATGTTCCGA	CCATTAGCCT	TACTCATGTT	AGATTTAGGG
550	560	570	580	590	600
TTAACGAGGA	TCCATTGGAG	GNNNNNNNNN	NNNNNNNNNN	NNGCGGTAAT	TCCAGCTCCA
AATTGCTCCT	AGGTAACCTC	CNNNNNNNNN	NNNNNNNNNN	NNGCCATTA	AGGTGAGGT
610	620	630	640	650	660
ATAGCGTATA	TTTAAGTTGT	TGCAGTAAA	AAGCTCGTAG	TTGGATCTCG	GGGCGGGGCG
TATCGCATAT	AAATTCAACA	ACGTCAATTT	TTCGAGCATC	AACCTAGAGC	CCCCCCCCG
670	680	690	700	710	720
GGCGGTCCGC	CCTCTTTGGG	GGTGTGCACC	GGCCTCCCCG	TCCTTAGTGC	CGGGGACGCG
CCGCCAGGGC	GGAGAAACCC	CCACACGTGG	CCGGAGGGGC	AGGAATCACG	GCCCCTGCGC
730	740	750	760	770	780
CTCCTGGCCT	TGACTGGTCG	GGACGCGGAT	TCGGCGACGT	TACTTTGAAA	AAATTAGAGT
GAGGACCGGA	ACTGACCAGC	CCTGCGCCTA	AGCCGCTGCA	ATGAAACTTT	TTAATCTCA
790	800	810	820	830	840
AATCAAAGCA	AGCCTACGCT	CTGAATACAT	TAGCATGGGA	TAGCGGATA	GGACTCTGGT
TTAGTTTCGT	TCGGATGCGA	GACTTATGTA	ATCGTACCCT	ATCGCGCTAT	CCTGAGACCA
850	860	870	880	890	900
CCTGTTGTGT	TGGTCTTCGG	GACCGGAGTA	ATGATTAATA	GGGACGGTTG	GGGGCATTCC
GGACAAACACA	ACCAGAAGCC	CTGGCCTCAT	TACTAATTAT	CCCTGCCAAC	CCCCGTAAGC

910 TATTTATTG ATAAAGTAAC	920 TCAGAGGTGA AGTCTCCACT	930 AATTCTTGG TTAAGAACCT	940 TTTACGAAAG AAATGCTTTC	950 ACGAACTTCT TGCTTGAAGA	960 GCGAAAGCAT CGCTTTCGTA
970 TTGCCAAGGA AACGGTTCCT	980 TGTTTTTATT ACAAAAGTAA	990 AATCAAGAAC TTAGTTCTTG	1000 GAAAGTTGGG CTTTCAACCC	1010 GGCTCGAAGA CCGAGCTTCT	1020 CGATCAGATA GCTAGTCTAT
1030 CCGTCCTAGT GGCAGGATCA	1040 CTCAACCGTA GAGTTGGCAT	1050 AACGATGCCG TTGCTACGGC	1060 ACTAGGGATC TGATCCCTAG	1070 GGCGGATGTT CCGCCTACAA	1080 GAATCGATGA CTTAGCTACT
1090 CTCCGCCGGC GAGGCGGCCG	1100 ACCTTATGAG TGAATACTC	1110 AAATCAAAGT TTTAGTTTCA	1120 TTTTGGGTTT AAAACCCAAG	1130 CGGGGGGAGT GCCCCCTCA	1140 ATGGTCGCAA TACCAGCGTT
1150 GGCTGANNNN CCGACTNNNN	1160 NNNNNNNNNN NNNNNNNNNN	1170 NNNNNNNNNN NNNNNNNNNN	1180 NNNCACCAGG NNNGTGGTCC	1190 AGTGGAGCCT TCACCTCGGA	1200 GCGGCTTAAT CGCCGAATTA
1210 TTGACTCAAC AACTGAGTTG	1220 ACGGGGAAAC TGCCCTTTG	1230 TCACCAGGTC AGTGGTCCAG	1240 CAGACATAGT GTCTGTATCA	1250 GAGGATTGAC CTCCTAACTG	1260 AGATTGAGAG TCTAACTCTC
1270 CTCTTCTTGG GAGAAAGAAC	1280 ATTCTATGGG TAAGATACCC	1290 TGGTGGTGCA ACCACCACGT	1300 TGGCCGTTCT ACCGGCAAGA	1310 TAGTTGGTGG ATCAACCACC	1320 AGTGATTTGT TCACTAAACA
1330 CTGGTTAATT GACCAATTA	1340 CCGTTAACGA GGCAATTGCT	1350 ACGAGACCTC TGCTCTGGAG	1360 AGCCTGTAA TCGGACAATT	1370 CTAGTACGC GATCAGTGCG	1380 GAAGGGTTC CTTCCCAAGG
1390 CCTTGCGGGG GGAACCGCCC	1400 CGACTTCTTA GCTGAAGAAT	1410 GAGGGACTAT CTCCCTGATA	1420 TGGCGTCTAG ACCGCAGATC	1430 CCAATGGAAG GGTTACCTTC	1440 TTTGAGGCAA AAACTCCGTT
1450 TAACAGGTCT ATTGTCCAGA	1460 GTGATGCCCT CACTACGGGA	1470 TAGAATGTTC ATCTTACAAG	1480 TGGGCCGCAC ACCCGGCGTG	1490 GCGCGCTACA CGCGCGATGT	1500 CTGATGAATT GACTACTTAA
1510 CAACGAGTTT GTTGCTCAAA	1520 ATAACCTGGG TATTGGACCC	1530 CCGACAGGCC GGCTGTCCGG	1540 CGGGTAATCT GCCATTAGA	1550 TTTGAAATTT AAACTTTAAA	1560 CATCGTGATG GTAGCACTAC
1570 GGGATAGATC CCCTATCTAG	1580 ATTGCAATTA TAACGTTAAT	1590 TTGATCTCCA AACTAGAGGT	1600 ACGAGGAATT TGCTCCTTAA	1610 CCTAGTAAGC GGATCATTCC	1620 GCGAGTCATC CGCTCAGTAG
1630 AGCTCGCGTT TCGAGCGCAA	1640 GACTACGTCC CTGATGCAGG	1650 CTGCCCTTTG GACGGGAAAC	1660 TACACACCGC ATGTGTGGCG	1670 CCGTCGCTCC GGCAGCGAGG	1680 TACCGATTGA ATGGCTAACT
1690 ATGGTCCGGT TACCAGGCCA	1700 GAAGTTTTTCG CTTCAAAAGC	1710 GATCGCGGCG CTAGCGCCGC	1720 ACGCCTGCGG TGCGGACGCC	1730 TTCGCCCGCG AAGCGCGGCG	1740 GCGACGTTGT CGCTGCAACA
1750 GAGAAGTTCA CTCTTCAAGT	1760 TCAAACTTA AGTTTGAAT	1770 TCATTTAGAG AGTAAATCTC	1780 GAAGGAGAAG CTTCCTCTTC	1790 TCGTAACAAG AGCATTGTTC	1800 GTTTCCGTAG CAAAGGCATC
1810 GTGAACNNNN CACTTGNNNN	1820 NNNNNNNNNN NNNNNNNNNN	1830	1840	1850	1860

Figure 10.

18S rDNA sequence data comparisons of *Physcomitrella patens*.

Homology alignment conducted with MacDNASIS sequence analysis software (Pro v3.6).

Physcomitrella patens 18S rDNA sequence data generated in this project is represented on the top line [labeled "Physcomitrella paten"]. *Physcomitrella patens* 18S rDNA sequence reported in GenBank accession # X80986 [labeled "P patens (Capesius)"]

Physcomitrel		10	20	30	40	50	
P patens (Ca	1	NNNNNNNNNN	NNNNNNNNNN	NNNNNNNNNN	NNNNNNNNAA	AGATTAAGCC	50
	1	AACCTGGTTG	ATCCTGCCAG	TAGTCATATG	CTTGTCTCAA	AGATTAAGCC	50
Physcomitrel		60	70	80	90	100	
P patens (Ca	51	ATGCATGTGT	AAGTATAAAC	TCTTTTGTAC	TGTGAAACTG	CGAATGGCTC	100
	51	ATGCATGTGT	AAGTATAAAC	TCTTTTGTAC	TGTGAAACTG	CGAATGGCTC	100
Physcomitrel		110	120	130	140	150	
P patens (Ca	101	ATTAAATCAG	TTATAGTTTC	TTTGATGGTA	CCTTGCTACT	CGGATAACCG	150
	101	ATTAAATCAG	TTATAGTTTC	TTTGATGGTA	CCTTGCTACT	CGGATAACCG	150
Physcomitrel		160	170	180	190	200	
P patens (Ca	151	TAGTAATTCT	AGAGCTAATA	CGTGCACAAA	CTCCCRACTT	TTTTGGAAGG	200
	151	TAGTAATTCT	AGAGCTAATA	CGTGCACAAA	CTCCCRACTT	TTTTGGAAGG	200
Physcomitrel		210	220	230	240	250	
P patens (Ca	201	GACGTATTTA	TTAGATAAAA	GGCCGATGCG	GGCTTGCCCG	GTGTTGCGGT	250
	201	GACGTATTTA	TTAGATAAAA	GGCCGATGCG	GGCTTGCCCG	GTGTTGCGGT	250
Physcomitrel		260	270	280	290	300	
P patens (Ca	251	GAATCATGAT	AACTTTGTCT	AATCGCACGG	CCTTGCCGCT	GGCGATGTTT	300
	251	GAATCATGAT	AACTTTGTCT	AATCGCACGG	CCTTGCCGCT	GGCGATGTTT	300
Physcomitrel		310	320	330	340	350	
P patens (Ca	301	CATTCAAATT	TCTGCCCTAT	CAACTTTCGA	TGGTAGGATA	GAGGCCTACC	350
	301	CATTCAAATT	TCTGCCCTAT	CAACTTTCGA	TGGTAGGATA	GAGGCCTACC	350
Physcomitrel		360	370	380	390	400	
P patens (Ca	351	ATGGTGGTAA	CGGGTGACGG	AGAATTAGGG	TTCGATTCCG	GAGAGGGAGC	400
	351	ATGGTGGTAA	CGGGTGACGG	AGAATTAGGG	TTCGATTCCG	GAGAGGGAGC	400
Physcomitrel		410	420	430	440	450	
P patens (Ca	401	CTGAGAAACG	GCTACCACAT	CCAAGGAAGG	CAGCAGGCGC	GCAAATTACC	450
	401	CTGAGAAACG	GCTACCACAT	CCAAGGAAGG	CAGCAGGCGC	GCAAATTACC	450
Physcomitrel		460	470	480	490	500	
P patens (Ca	451	CAATCCCAGC	ACGGGGAGGT	AGTGACAATA	AATAACAATA	CTGGGCTTTT	500
	451	CAATCCCAGC	ACGGGGAGGT	AGTGACAATA	AATAACAATA	CTGGGCTTTT	500
Physcomitrel		510	520	530	540	550	
P patens (Ca	501	ACAAGTCTGG	TAATTGGAAT	GAGTACAATC	TAAATCCCTT	AACGAGGATC	550
	501	ACAAGTCTGG	TAATTGGAAT	GAGTACAATC	TAAATCCCTT	AACGAGGATC	550
Physcomitrel		560	570	580	590	600	
P patens (Ca	551	CATTGGAGGN	NNNNNNNNNN	NNNNNNNNNN	GCGGTAATTC	CAGCTCCAAT	600
	551	CATTGGAGGG	CAAGTCTGGT	GCCAGCAGCC	GCGGTAATTC	CAGCTCCAAT	600
Physcomitrel		610	620	630	640	650	
P patens (Ca	601	AGCGTATATT	TAAGTTGTTG	CAGTTAAAAA	GCTCGTAGTT	GGATCTTGGG	650
	601	AGCGTATATT	TAAGTTGTTG	CAGTTAAAAA	GCTCGTAGTT	GGATCTTGGG	650
Physcomitrel		660	670	680	690	700	
P patens (Ca	651	TCGGGGGGAG	CGGTCCGCCC	CTTGTGGGTG	TGCACTGGTC	CACTCGGCCT	700
	651	TCGGGGGGAG	CGGTCCGCCC	CTTGTGGGTG	TGCACTGGTC	CACTCGGCCT	700
Physcomitrel		710	720	730	740	750	
P patens (Ca	701	TCTTGCCGGG	GACGCGCTCC	TGGTTTTAAT	TAATCGGGAC	GCGGAGTCGG	750
	701	TCTTGCCGGG	GACGCGCTCC	TGGTTTTAAT	TAATCGGGAC	GCGGAGTCGG	750
Physcomitrel		760	770	780	790	800	
P patens (Ca	751	CGATGTTACT	TTGAAAAAAT	TAGAGTGCTC	AAAGCAAGCC	TATGCTCTGA	800
	751	CGATGTTACT	TTGAAAAAAT	TAGAGTGCTC	AAAGCAAGCC	TATGCTCTGA	800

Physcomitrel		810	820	830	840	850	
<i>P patens</i> (Ca	801	ATACATTAGC	ATGGAATAAC	GTGATAGGAC	TCTGGTCCTG	TTCGTGTTGG	850
	801	ATACATTAGC	ATGGAATAAC	GTGATAGGAC	TCTGGTCCTG	TTCGTGTTGG	850
Physcomitrel		860	870	880	890	900	
<i>P patens</i> (Ca	851	TCTTCGGGAC	CGGAGTAATG	ATTAATAGGG	ACGGTTGGGG	GCATTTCGTAT	900
	851	TCTTCGGGAC	CGGAGTAATG	ATTAATAGGG	ACGGTTGGGG	GCATTTCGTAT	900
Physcomitrel		910	920	930	940	950	
<i>P patens</i> (Ca	901	TTCATTGTCA	GAGGTGAAAT	TCTTGGATTT	ATGAAAGACG	AACTTCTGCG	950
	901	TTCATTGTCA	GAGGTGAAAT	TCTTGGATTT	ATGAAAGACG	AACTTCTGCG	950
Physcomitrel		960	970	980	990	1000	
<i>P patens</i> (Ca	951	AAAGCATTG	CCAAGGATGT	TTTCATTAAT	CAAGAACGAA	AGTTGGGGGC	1000
	951	AAAGCATTG	CCAAGGATGT	TTTCATTAAT	CAAGAACGAA	AGTTGGGGGC	1000
Physcomitrel		1010	1020	1030	1040	1050	
<i>P patens</i> (Ca	1001	TCGAAGACGA	TCAGATACCG	TCCTAGTCTC	AACCATAAAC	GATGCCGACT	1050
	1001	TCGAAGACGA	TCAGATACCG	TCCTAGTCTC	AACCATAAAC	GATGCCGACT	1050
Physcomitrel		1060	1070	1080	1090	1100	
<i>P patens</i> (Ca	1051	AGGGATTGGC	GGATGTTACT	TTGATGACTC	CGCCAGCACC	TTATGAGAAA	1100
	1051	AGGGATTGGC	GGATGTTACT	TTGATGACTC	CGCCAGCACC	TTATGAGAAA	1100
Physcomitrel		1110	1120	1130	1140	1150	
<i>P patens</i> (Ca	1101	TCAAAGTTTT	TNGGTTCCGG	GGNGAGTATG	NTCGCAAGNC	TGAANNNNNN	1150
	1101	TCAAAGTTTT	TGGGTTCCGG	GGGGAGTATG	GTCGCAAGGC	TGAAACTTAA	1150
Physcomitrel		1160	1170	1180	1190	1200	
<i>P patens</i> (Ca	1151	NNNNNNNNNN	NNNNNNGCAC	NACCAGGAGT	GGAG-CTGCG	GCTTAATTTG	1200
	1151	AGGAATTGAC	GGAAGGGCAC	CACCAGGAGT	GGAGCCTGCG	GCTTAATTTG	1200
Physcomitrel		1210	1220	1230	1240	1250	
<i>P patens</i> (Ca	1201	ACTCAACACG	GGGAAACTTA	CCAGGTCCAG	ACATAGTAAG	GATTGACAGA	1250
	1201	ACTCAACACG	GGGAAACTTA	CCAGGTCCAG	ACATAGTAAG	GATTGACAGA	1250
Physcomitrel		1260	1270	1280	1290	1300	
<i>P patens</i> (Ca	1251	TTGAGAGCTC	TTTCTTGATT	CTATGGGTGG	TGGTGATGG	CCGTTCTTAG	1300
	1251	TTGAGAGCTC	TTTCTTGATT	CTATGGGTGG	TGGTGATGG	CCGTTCTTAG	1300
Physcomitrel		1310	1320	1330	1340	1350	
<i>P patens</i> (Ca	1301	TTGGTGGAGT	GATTTGTCTG	GTAAATCCG	TTAACGAACG	AGACCTCAGC	1350
	1301	TTGGTGGAGT	GATTTGTCTG	GTAAATCCG	TTAACGAACG	AGACCTCAGC	1350
Physcomitrel		1360	1370	1380	1390	1400	
<i>P patens</i> (Ca	1351	CTGCTAACTA	GTTACGCGAA	GGATTTTTTC	CTTTGCGGCC	AACTTCTTAG	1400
	1351	CTGCTAACTA	GTTACGCGAA	GGATTTTTTC	CTTTGCGGCC	AACTTCTTAG	1400
Physcomitrel		1410	1420	1430	1440	1450	
<i>P patens</i> (Ca	1401	AGGGACTATC	GGCGTCTAGC	CGATGGAAGT	TTGAGGCAAT	AACAGGTCTG	1450
	1401	AGGGACTATC	GGCGTCTAGC	CGATGGAAGT	TTGAGGCAAT	AACAGGTCTG	1450
Physcomitrel		1460	1470	1480	1490	1500	
<i>P patens</i> (Ca	1451	TGATGCCCTT	AGATGTTCTG	GGCCGCACGC	GCGTACTACT	GATGAATTCA	1500
	1451	TGATGCCCTT	AGATGTTCTG	GGCCGCACGC	GCGTACTACT	GATGAATTCA	1500
Physcomitrel		1510	1520	1530	1540	1550	
<i>P patens</i> (Ca	1501	ACGAGTTTAT	AACCTGGACC	GATA-GGTCT	GGGTAATCTT	TTGAAATTTT	1550
	1501	ACGAGTTTAT	AACCTGGGCC	GATACGGT-T	GGGTAATCTT	TTGAAATTTT	1550
Physcomitrel		1560	1570	1580	1590	1600	
<i>P patens</i> (Ca	1551	ATCGTGATGG	GGATAGATCA	TTGCAATTAT	TGATCTTCAA	CGAGGAATTC	1600
	1551	ATCGTGATGG	GGATAGATCA	TTGCAATTAT	TGATCTTCAA	CGAGGAATTC	1600

		1610	1620	1630	1640	1650	
Physcomitrel	1601	CTAGTAAGCG	CGAGTCATCA	GCTCGCGTTG	ACTACGTCCC	TGCCCTTTGT	1650
P patens (Ca	1601	CTAGTAAGCG	CGAGTCATCA	GCTCGCGTTG	ACTACGTCCC	TGCCCTTTGT	1650
		1660	1670	1680	1690	1700	
Physcomitrel	1651	ACACACCGCC	CGTCGCTCCT	ACCGATTGAA	TGGTCCGGTG	AAGTTTTTCGG	1700
P patens (Ca	1651	ACACACCGCC	CGTCGCTCCT	ACCGATTGAA	TGGTCCGGTG	AAGTTTTTCGG	1700
		1710	1720	1730	1740	1750	
Physcomitrel	1701	ATTGCGGCGA	TGCCGGCGGT	TCGCCGCCGG	TGACGTTGGG	AGAAGTTCAT	1750
P patens (Ca	1701	ATTGCGGCGA	TGCCGGCGGT	TCGCCGCCGG	TGACGTTGGG	AGAAGTTCAT	1750
		1760	1770	1780	1790	1800	
Physcomitrel	1751	TAAACCTTAT	CATTTAGAGG	AAGGAGAAGT	CGTAACAAGG	TTTCCGTAGG	1800
P patens (Ca	1751	TAAACCTTAT	CATTTAGAGG	AAGGAGAAGT	CGTAACAAGG	TTTCCGTAGG	1800
		1810	1820	1830	1840	1850	
Physcomitrel	1801	TNNNNNNNN	NNNNNNNN.	1850
P patens (Ca	1801	TGAACCTGCA	GAAGGATCA.	1850

APPENDIX B : Taxa

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Table 2. Taxa list with classification.

Division Chlorophyta (Green Plants)

Subdivision Embryophytina (Embryophytes)

<u>Class</u>	<u>Subclass</u>	<u>Genus species</u>	<u>Accession #</u>	
Bryophyta	Anthocerotopsida	<i>Anthoceros laevis</i>	U18491	
		<i>Anthoceros agrestis</i>	X80984	
		<i>Megaceros aenigmaticus</i>		
		<i>Notophylas breutelii</i>		
	Bryopsida	<i>Atrichum undulatum</i>	X85093	
		<i>Bryum argenteum</i>	U18529	
		<i>Eurhynchium hians</i>	U18501	
		<i>Funaria hygrometrica</i>	X74114	
		<i>Leptobryum pyriforme</i>	X80980	
		<i>Mnium hornum</i>	X80985	
		<i>Physcomitrella patens</i>		
		<i>Physcomitrella patens 2</i>	X80986	
		<i>Physcomitrium pyriforme</i>	U18517	
		<i>Polytrichum formosum</i>	X80982	
		Sphagnopsida	<i>Sphagnum cuspidatum</i>	X80213
			<i>Sphagnum palustre</i>	
		Hepaticopsida	<i>Calypogeia arguta</i>	X78439
	<i>Conocephalum conicum</i>		X80987	
	<i>Fossombronia pusilla</i>		X78341	
	<i>Marchantia polymorpha</i>		X75521	
	<i>Reboulia hemisphaerica</i>		U18521	
	<i>Riccia fluitans</i>		X78441	
	<i>Scapania nemorea</i>		X80983	
	<i>Sphaerocarpus donnelli</i>		X85094	
	<i>Sphaerocarpus texanus</i>		U18522	
	<i>Riccardia pinguis</i>		X85095	
	<i>Pellia epiphylla</i>	X80210		

Table 2. Taxa list with classification (cont'd).

Division Chlorophyta (Green Plants)

Subdivision Embryophytina (Embryophytes)

<u>Class</u>	<u>Subclass</u>	<u>Genus species</u>	<u>Accession #</u>
Equisetophyta	Equisetophyta	<i>Equisetum robustum</i>	X78890
	Sphenopsida	<i>Equisetum hyemale</i>	U18500
Filicophyta	Filicopsida	<i>Pteris vittata</i>	
		<i>Adiantum raddianum</i>	X78889
		<i>Adiantum raddianum 2</i>	U18621
		<i>Osmunda cinnamomea</i>	U18516
	Ophioglossopsida	<i>Salvinia natans</i>	X90413
		<i>Ophioglossum petiolatum</i>	U18515
Lycopodiophyta	Isoetopsida	<i>Isoetes englemannii</i>	U18506
		<i>Selaginella umbrosa</i>	X75517
		<i>Selaginella vogelii</i>	X83520
	Lycopodiopsida	<i>Huperzia lucidulum</i>	U18505
		<i>Lycopodiella inundatum</i>	U18512
		<i>Lycopodium phlegmaria</i>	X81964
		<i>Lycopodium taxifolium</i>	X83522
		<i>Lycopodium tristachyum</i>	U18511
Magnoliophyta	Liliopsida	<i>Oryza sativa</i>	X00755
		<i>Zea mays</i>	K02202
	Magnoliopsida	<i>Arabidopsis thaliana</i>	X16077
		<i>Glycine max</i>	X02623
		<i>Sinapis alba</i>	X66325
Pinophyta	Cycadicae	<i>Zamia pumila</i>	M20017
		<i>Ginkgo biloba</i>	D16448
	Pinicae	<i>Nageia nagi</i>	D16447
		<i>Pinus luchuensis</i>	D38246
		<i>Pinus wallichiana</i>	X75080
		<i>Taxus mairei</i>	D16445
	Gneticae	<i>Gnetum leyboldii</i>	L24045
Psilotophyta	Psilotopsida	<i>Psilotum nudum</i>	X81963

Table 2. Taxa list with classification (cont'd).

Outgroups

Division Chlorophyta (Green Plants)
 Subdivision Chlorophycophytina (Green Algae)

<u>Class</u>	<u>Subclass</u>	<u>Genus species</u>	<u>Accession #</u>
Charophyta	Coleochaetales	<i>Coloeochaete scutata</i>	X68825
	Klebsormidiales	<i>Klebsormidium flaccidum</i>	X75520
	Charales	<i>Chara foetida</i>	X70704
		<i>Nitella flexilis</i>	U05261
		<i>Nitella sp.</i>	M95615
Chlorophyta	Chlorococcales	<i>Parietochloris pseudoalveolaris</i>	M63002
		<i>Hydrodictyon reticulatum</i>	M74497
	Volvocales	<i>Chlamydomonas reinhardtii</i>	M32703
<u>Division</u>	<u>Class</u>	<u>Genus species</u>	<u>Accession #</u>
Eumycota	Hemiascomycetes	<i>Saccharomyces cerevisiae</i>	J01353
	Archaeascomycetes	<i>Schizosaccharomyces pombe</i>	X58056
Myxomycota	Dictyosteliida	<i>Dictyostelium discoideum</i>	K02641

Table 3. Taxa set designations.

Taxa	Author	Accession	Error Status	Taxa Set A	Taxa Set B	Taxa Set C
<i>Megaceros aenigmaticus</i>	Ashton		OK	/	/	/
<i>Notophylas brütelu</i>	Ashton		OK	/	/	/
<i>Physcomitrella patens</i>	Ashton		OK	/	/	/
<i>Pteris vittata</i>	Ashton		OK	/	/	/
<i>Sphagnum palustre</i>	Ashton		OK	/	/	/
<i>Anthoceros agrestis</i>	Capesius	X80984	N/A	/	/	/
<i>Atrichum undulatum</i>	Capesius	X85093	OK	/	/	/
<i>Calypogeia arguta</i>	Capesius	X78439	OK	/	/	/
<i>Conocophalum conicum</i>	Capesius	X80987	OK	/	/	/
<i>Fossombronia pusilla</i>	Capesius	X78341	OK	/	/	/
<i>Funaria hygrometrica</i>	Capesius	X74114	OK	/	/	/
<i>Leptobryum pyriforme</i>	Capesius	X80980	OK	/	/	/
<i>Mnium hornum</i>	Capesius	X80985	OK	/	/	/
<i>Pellia epiphylla</i>	Capesius	X80210	OK	/	/	/
<i>Physcomitrella patens 2</i>	Capesius	X80986	OK	/	/	/
<i>Polytrichum formosum</i>	Capesius	X80982	OK	/	/	/
<i>Riccardia pinguis</i>	Capesius	X85095	OK	/	/	/
<i>Riccia fluitans</i>	Capesius	X78441	OK	/	/	/
<i>Scapania nemorea</i>	Capesius	X80983	OK	/	/	/
<i>Sinapis alba</i>	Capesius	X66325	OK	/	/	/
<i>Sphaerocarpos donnellii</i>	Capesius	X85094	OK	/	/	/
<i>Sphagnum cuspidatum</i>	Capesius	X80213	OK	/	/	/
<i>Anthoceros laevis</i>	Chapman	U18491	N/A	/	/	/
<i>Bryum argenteum</i>	Chapman	U18529	OK	/	/	/
<i>Equisetum hyemale</i>	Chapman	U18500	OK	/	/	/
<i>Eurhynchium hians</i>	Chapman	U18501	OK	/	/	/
<i>Huperzia lucidulum</i>	Chapman	U18505	OK	/	/	/
<i>Isoetes engelmannii</i>	Chapman	U18506	OK	/	/	/
<i>Lycopodiella inundatum</i>	Chapman	U18512	OK	/	/	/
<i>Lycopodium tristachyum</i>	Chapman	U18511	OK	/	/	/
<i>Ophioglossum petiolatum</i>	Chapman	U18515	OK	/	/	/
<i>Osmunda cinnamomea</i>	Chapman	U18516	OK	/	/	/
<i>Physcomitrium pyriforme</i>	Chapman	U18517	OK	/	/	/
<i>Reboulia hemisphaerica</i>	Chapman	U18521	OK	/	/	/
<i>Sphaerocarpus texanus</i>	Chapman	U18522	OK	/	/	/
<i>Ginkgo biloba</i>	Chaw	D16448	OK	/	/	/
<i>Nageia nagi</i>	Chaw	D16447	OK	/	/	/
<i>Pinus luchuensis</i>	Chaw	D38246	OK	/	/	/
<i>Taxus mairei</i>	Chaw	D16445	OK	/	/	/
<i>Schizosaccharomyces pombe</i>	De Wachter	X58056	N/A	/	/	/
<i>Glycine max</i>	Eckenrode	X02623	OK	/	/	/
<i>Arabidopsis thaliana</i>	Gruendler	X16077	OK	/	/	/
<i>Chlamydomonas reinhardtii</i>	Gunderson	M32703	OK	/	/	/
<i>Adiantum raddianum</i>	Huss	X78889	OK	/	/	/
<i>Chara foetida</i>	Huss	X70704	OK	/	/	/
<i>Coleochaete scutata</i>	Huss	X68825	OK	/	/	/
<i>Equisetum robustum</i>	Huss	X78890	OK	/	/	/
<i>Klebsormidium flaccidum</i>	Huss	X75520	OK	/	/	/
<i>Lycopodium phlegmaria</i>	Huss	X81964	OK	/	/	/
<i>Lycopodium taxifolium</i>	Huss	X83522	OK	/	/	/
<i>Marchantia polymorpha</i>	Huss	X75521	OK	/	/	/
<i>Psilotum nudum</i>	Huss	X81963	OK	/	/	/
<i>Salvinia natans</i>	Huss	X90413	OK	/	/	/
<i>Selaginella umbrosa</i>	Huss	X75517	OK	/	/	/
<i>Selaginella vogelii</i>	Huss	X83520	OK	/	/	/
<i>Parietochloris pseudoalveolaris</i>	Lewis	M63002	OK	/	/	/
<i>Saccharomyces cerevisiae</i>	Mankin	J01353	N/A	/	/	/
<i>Dictyostelium discoideum</i>	McCarroll	K02641	N/A	/	/	/
<i>Zea mays</i>	Messing	K02202	OK	/	/	/
<i>Zamia pumila</i>	Naire	M20017	OK	/	/	/
<i>Gnetum leyboldii</i>	Nickrent	L24045	OK	/	/	/
<i>Nitella flexilis</i>	Ragan	U05261	OK	/	/	/
<i>Pinus wallichiana</i>	Sensen	X75080	OK	/	/	/
<i>Oryza sativa</i>	Takaiwa	X00755	OK	/	/	/
<i>Hydrodictyon reticulatum</i>	Wilcox	M74497	OK	/	/	/
<i>Nitella sp.</i>	Wilcox	M95615	OK	/	/	/
<i>Adiantum raddianum 2</i>	Wolf	U18621	OK	/	/	/

OK: determined by Hedderson (personal communication)
 OK : determined by double-stranded sequencing
 N/A: Not Available

APPENDIX C : Tree Length Distribution

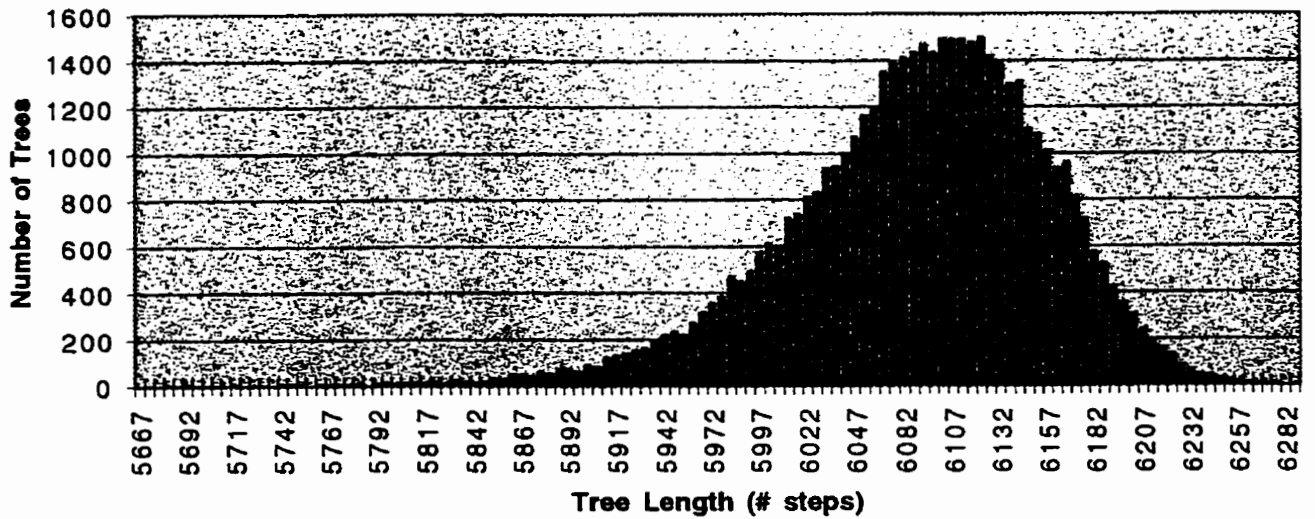
Figure 11 Frequency distribution of lengths of random trees for taxa set B and C.. 37

Figure 11.

Frequency distribution of lengths of random trees for taxa set B and C.

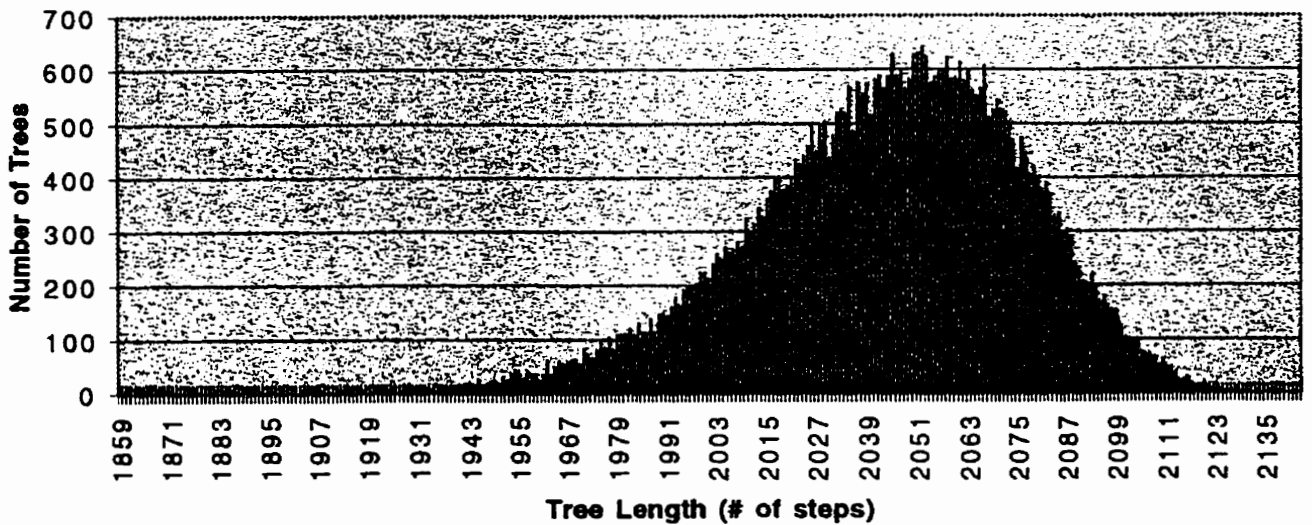
The g_1 statistic for tree length distribution skewness was determined for both taxa set B and C. The g_1 statistic based on the length distribution of 50,000 randomly sampled trees from a set of all possible trees.

Taxa set B frequency distribution of lengths of random trees



mean	sd	g_1	g_2
6086.17	70.21	-0.6235	0.6492

Taxa set C frequency distribution of lengths of random trees



mean	sd	g_1	g_2
2043.27	33.62	-0.5330	0.3880

APPENDIX D : Substitution Rate Calculation

Figure 12	Distribution of relative substitution rates estimated from alignment of 18S rDNA of taxa set B.....	40
Figure 13	Distribution of relative substitution rates with respect to MacClade matrix, estimated from alignment of 18S rDNA of taxa set B.....	42
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Figure 12.

Distribution of relative substitution rates, estimated from alignment of 18S rDNA of taxa set B.

Rates were estimated for each of the alignment positions not absolutely conserved and that contain a nucleotide in at least 25% of the aligned sequences. The colored areas in spectra define sets of nucleotides indicated in different categories used for phylogenetic analysis.

Distribution of Nucleotide Substitution Rates

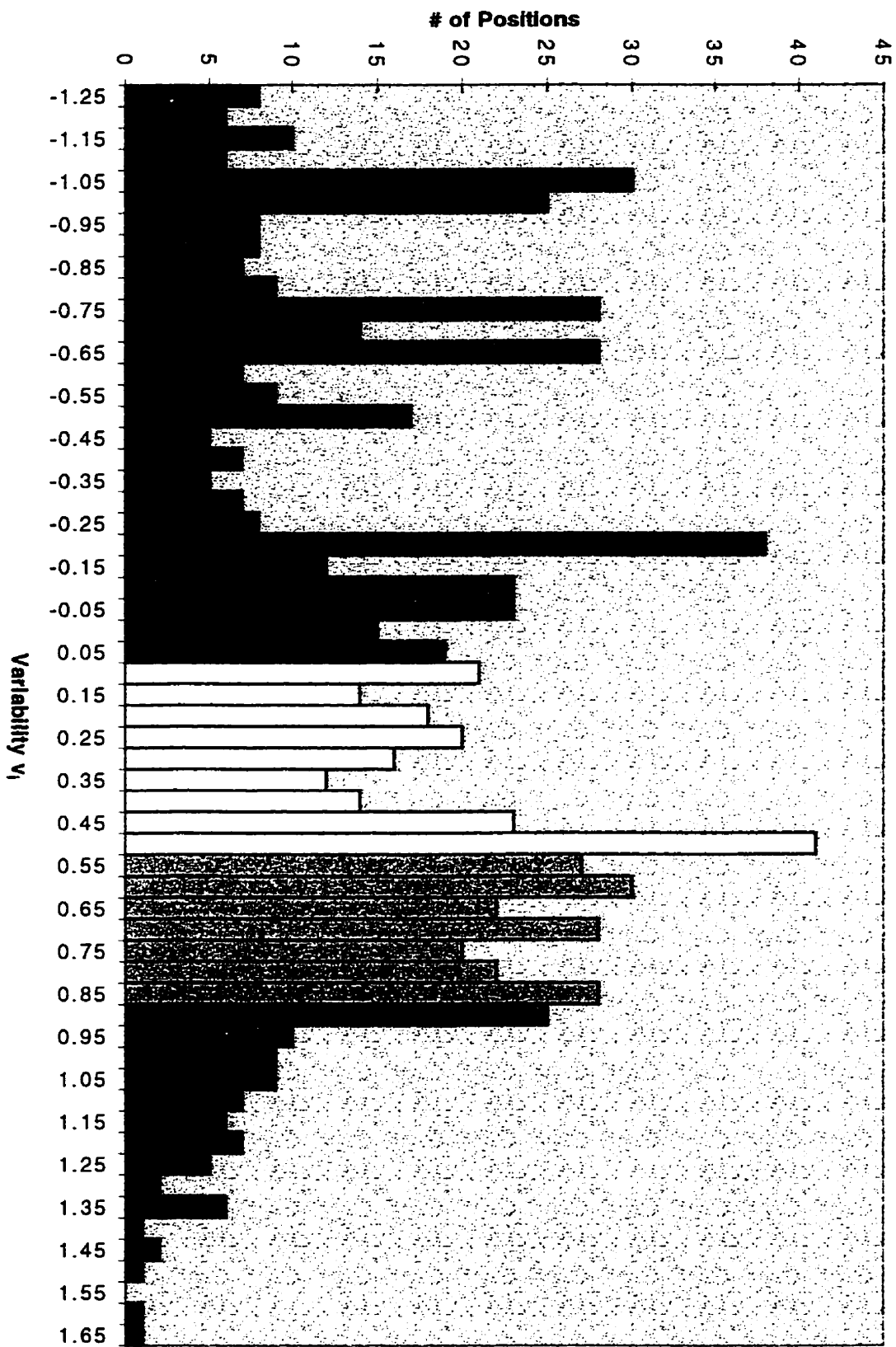
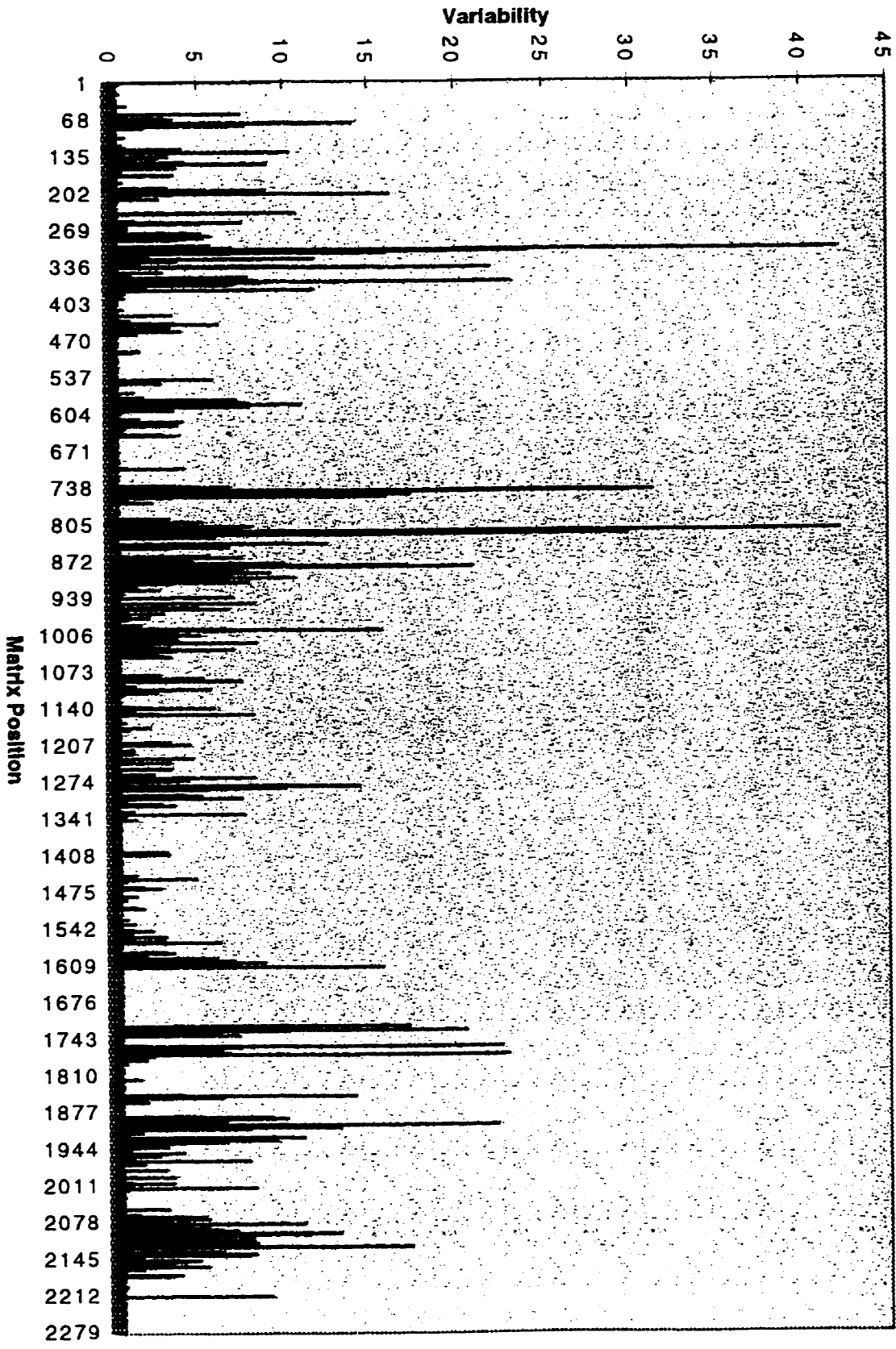


Figure 13.

Distribution of relative substitution rates with respect to MacClade matrix, estimated from alignment of 18S rDNA of taxa set B.

Variability is the log of the values in the variability table for taxa set B (Table 4).



Nucleotide Variability

Table 4.

Variability table for taxa set B.

The variabilities were calculated using taxa set B.

Column Headings


Site :refers to matrix location of MacClade Matrix

Variability :Score generated with 3 iterations of substitution rate calculation.

Cat :Variability category (shown in columns of MacClade Matrix)

Color Key

Cat #	Color
- 1	Grey
0	White
1	Red
2	Pink/Orange
3	Yellow
4	Green
5	Blue
6	Purple



Note: Nonvariable sites represented by purple are not included in table.

Table 4. Variability table for taxa set B.

Site	Variability	Cat	Site	Variability	Cat	Site	Variability	Cat	Site	Variability	Cat	Site	Variability	Cat
	-	-1		0.619	4	208	1.727746	3		0.856233	4	315	-	0
	9.269674	1		0.223341	5		0.170722	5		0.757058	4	317	-	0
	0.552329	4		0.767443	4	211	-	0		0.078532	5	318	2.160407	3
	0.529134	4		0.317833	4		0.182153	5		0.952773	4		0.651236	4
	0.148367	5		0.767447	4	214	2.739964	3		0.689048	4	320	-	0
	0.327716	4		3.282056	2		-	-1		0.929514	4		0.181397	5
	0.138438	5		1.078825	4		-	-1		0.917046	4	323	-	0
	0.90967	4		0.223339	5		-	-1		3.281521	2		11.752676	1
	0.446531	4	140	2.728873	3		-	-1		5.17309	2		3.810957	2
	7.509933	1	141	1.397547	3		-	-1	278	2.100022	3	330	-	0
	0.682403	4	142	1.496357	3		-	-1		0.687948	4	331	1.629031	3
	0.128236	5	145	-	0		-	-1	280	-	0		0.718257	4
	0.103599	5	146	-	0		-	-1	281	-	0		0.651259	4
	0.871057	4	147	-	0		-	-1		5.740174	2		0.920512	4
68	3.06303	3		4.286476	2		-	-1	283	2.145807	3		0.601508	4
	3.585126	2	149	1.890271	3		-	-1		5.2746	2		9.703156	1
72	-	0	150	-	0		-	-1		4.666056	2		14.84859	1
73	-	0		8.999743	1		-	-1		5.366636	2		21.933331	1
74	-	0		0.266337	5		-	-1		4.367602	2	340	-	0
75	3.011281	3		0.867829	4		-	-1	288	1.88005	3		0.319632	4
	14.107607	1	156	-	0		-	-1		3.845947	2	348	2.963082	3
	12.137928	1	157	1.657986	3		-	-1		-	-1		0.068799	5
78	1.663868	3		3.796199	2		-	-1		-	-1		0.419559	4
	0.344254	4		0.076108	5		-	-1		-	-1		0.476006	4
80	2.036346	3	161	-	0		-	-1		-	-1	353	1.156994	3
	7.768451	1		0.672955	4		-	-1		-	-1		0.09905	5
	6.805617	2	169	-	0		-	-1		-	-1	355	-	0
83	-	0		3.589689	2		-	-1		-	-1	356	-	0
86	1.905839	3	173	-	0		-	-1		-	-1		7.874473	1
87	-	0	178	-	0		3.289062	2		-	-1		0.323949	4
88	-	0		0.668554	4		10.726263	1		-	-1		0.777658	4
91	-	0		0.176978	5		5.156658	2		5.695101	2		0.631455	4
	0.084587	5	192	-	0	242	-	0	301	2.225938	3	361	-	0
	0.103866	5		0.068687	5	243	-	0		4.631436	2		0.596501	4
	0.343067	4		3.278543	2	244	-	0		4.675918	2		8.468694	1
	0.825009	4		0.176977	5	245	-	0		6.930992	2		3.206521	2
	0.619042	4		5.293993	2	246	-	0		4.692835	2		23.149639	1
	0.158184	5		8.955044	1		0.123009	5		42.043568	1		0.705765	4
	0.892272	4	200	-	0		0.323058	4	308	-	0		8.743754	1
	4.129072	2		4.639279	2	254	1.368369	3	309	-	0	368	-	0
	0.867849	4	202	1.137486	3	256	3.046788	3		24.016476	1		0.488794	4
	0.612641	4	203	1.58899	3		7.52297	1	311	1.855102	3		7.05522	2
127	-	0	204	-	0		5.17679	2		16.033964	1		0.943945	4
	0.650846	4		16.210125	1		0.255383	5		6.983452	2		0.076474	5
	10.379545	1		3.625143	2		0.920481	4	314	-	0	377	2.043571	3

Table 4. Variability table for taxa set B.

Site	Variability	Cat	Site	Variability	Cat	Site	Variability	Cat	Site	Variability	Cat	Site	Variability	Cat
	0.2255	5		0.380907	4	622	-	0		15.156329	1	802	-	0
379	-	0		0.103866	5	626	2.214879	3	753	-	0		5.207098	2
	11.742348	1		5.792713	2		3.791995	2		4.028542	2		0.826669	4
381	1.561765	3	544	1.734027	3	628	-	0		12.693307	1		3.440878	2
	0.317832	4	546	2.247888	3		0.788289	4		15.880737	1	807	1.243845	3
	0.085336	5	547	2.015177	3		0.651262	4		3.526304	2	808	3.050692	3
	0.646627	4	548	-	0		0.067634	5	758	-	0	809	-	0
	0.771534	4		0.515175	4		0.768031	4		0.715596	4	810	-	0
388	-	0	550	2.825728	3	634	-	0		0.825691	4		8.033676	1
	0.767469	4		0.103866	5		0.628	4	765	-	0		7.444651	1
	0.085667	5	567	1.256526	3		0.093563	5	766	1.450168	3	813	-	0
405	-	0		0.073112	5		0.160887	5	767	2.265764	3	814	-	0
	0.055434	5	575	1.805796	3	640	-	0	768	-	0	815	-	0
	0.103866	5	578	1.828991	3		0.810007	4	769	-	0	816	-	0
	0.666214	4		0.176978	5	643	2.917701	3	770	-	0	817	-	0
	0.123009	5		7.237089	1		3.864982	2	771	-	0	818	-	0
	0.089922	5		0.895812	4		0.160887	5	772	-	0		42.17226	1
	3.470867	2		0.627969	4		0.093563	5	773	-	0		16.798563	1
	1.082919	4		5.861557	2		0.108727	5	774	-	0		12.115381	1
	0.176977	5		7.878078	1		0.089327	5	775	-	0		5.566195	2
	3.5805	2		3.665234	2		0.110319	5		-	-1		10.407763	1
441	2.942682	3		0.39326	4		0.110319	5		-	-1	824	-	0
442	1.125725	3		11.017945	1		0.058113	5		-	-1		29.837286	1
	6.185495	2	590	1.367205	3		0.08572	5		-	-1		15.620416	1
	3.397484	2	591	-	0		0.223341	5		-	-1	827	3.118919	3
	0.093562	5		7.87618	1		0.874578	4		-	-1		4.957371	2
	3.97311	2		5.325319	2		0.757713	4		-	-1		5.009516	2
	0.093562	5		0.628004	4		4.149544	2		-	-1	830	2.072786	3
	1.004536	4	597	-	0		0.213441	5		-	-1		5.860284	2
460	1.485064	3	598	1.688656	3		0.104755	5		-	-1	832	-	0
	0.066862	5		3.542102	2		0.179315	5		-	-1	833	-	0
	0.223341	5		0.718146	4		0.22334	5		-	-1	834	-	0
	0.071828	5		0.476982	4		3.608802	2		-	-1	835	-	0
	0.123009	5		0.545833	4		4.396942	2		-	-1	836	-	0
	0.085336	5		0.093563	5		6.770321	2		-	-1	837	-	0
491	1.606322	3		0.103866	5		6.163895	2		-	-1	838	-	0
493	1.34938	3		0.093563	5		0.58855	4		-	-1	839	-	0
	0.084587	5		0.11128	5	743	1.77866	3	793	-	0	840	-	0
	0.176976	5		0.628063	4		31.307409	1	795	1.626606	3	841	-	0
	0.176976	5	614	1.569786	3	746	-	0		3.264218	2		12.496162	1
	0.055434	5		0.600258	4		29.58567	1		0.531552	4		5.20372	2
	0.159313	5		0.600123	4		4.048969	2	798	-	0		3.781062	2
	0.239817	5		0.651231	4		1.047038	4	799	-	0		6.75635	2
	0.251453	5		4.038343	2		3.501297	2		0.978878	4		3.219445	2
	0.223342	5		0.405938	4		17.272007	1	801	2.577457	3		0.176977	5

Table 4. Variability table for taxa set B.

Site	Variability	Cat	Site	Variability	Cat	Site	Variability	Cat	Site	Variability	Cat	Site	Variability	Cat
	6.69502	2	908	2.942903	3	984	-	0		3.279169	2		0.073112	5
850	3.159932	3	909	2.401639	3		0.22334	5		0.366882	4		3.278544	2
851	-	0		3.559604	2		0.17698	5		0.099254	5		4.380527	2
	0.939649	4		3.848646	2	990	1.578121	3	1081	2.670779	3		0.176977	5
	5.553166	2		7.812299	1		0.223341	5		0.317833	4		0.176977	5
	4.199852	2		3.441546	2		0.609693	4		0.149706	5	1213	-	0
	7.43768	1		3.278681	2		0.190065	5		0.318916	4		0.085336	5
868	-	0		0.609673	4		3.278607	2	1087	2.711495	3		0.184502	5
	3.184855	2		0.223341	5		6.50707	2		5.190529	2		0.085336	5
870	2.143478	3	919	1.400151	3	998	-	0		7.37346	1	1218	1.125726	3
871	-	0		0.609689	4		15.625661	1		0.067051	5		1.004646	4
	3.849394	2	922	-	0		7.443014	1		0.600254	4	1223	1.125726	3
	4.558035	2	925	1.556025	3	1002	1.239992	3	1101	1.125724	3	1225	1.125726	3
874	1.209866	3	926	2.695115	3		0.176979	5	1102	1.229711	3		4.509048	2
875	1.131517	3		0.600138	4	1007	2.709673	3		0.249082	5		0.176977	5
876	2.950408	3	933	-	0		3.856979	2		5.574201	2		3.278598	2
877	2.627478	3	938	-	0		5.012696	2		3.278647	2		0.190064	5
	9.923486	1		0.085131	5	1011	-	0	1110	-	0		3.278607	2
	0.433646	4		6.942472	2		3.676152	2	1111	1.490349	3	1262	2.319566	3
880	1.947697	3	942	-	0	1014	1.18634	3		0.83844	4		0.199949	5
	0.322677	4	947	-	0		0.82716	4	1113	2.452124	3		0.176979	5
	17.108433	1		0.223341	5	1016	3.03937	3		0.587425	4		8.144135	1
	7.019657	2		8.182248	1		1.035093	4		0.185878	5	1270	1.196749	3
	20.887762	1		0.223341	5	1019	-	0		0.895822	4		3.2786	2
885	3.144126	3	954	-	0		3.760838	2		0.711259	4		4.27888	2
886	-	0	955	-	0		0.142231	5	1140	1.711748	3	1274	1.778439	3
887	-	0	956	-	0	1022	2.032823	3		3.279149	2		3.766665	2
	0.308956	5	957	-	0		8.269547	1		5.804338	2		0.651223	4
	7.545074	1		4.916144	2	1024	2.510505	3		0.651224	4	1280	-	0
	6.175605	2		4.032711	2	1025	-	0	1148	1.229711	3	1281	-	0
	6.876983	2		4.33321	2		0.598423	4		0.353988	4	1282	-	0
892	1.125726	3		6.676934	2	1031	2.294927	3	1152	2.550627	3		13.920209	1
	3.280788	2	962	-	0		3.276029	2		8.073201	1		14.335297	1
	0.606773	4	963	-	0	1033	-	0		0.600253	4	1285	2.057786	3
	9.101762	1	964	-	0		6.934744	2		0.067052	5		0.840473	4
897	-	0		0.085336	5		5.591353	2		0.085667	5	1287	-	0
	3.656609	2		0.819166	4		3.487582	2	1176	-	0		9.944017	1
	4.568418	2	969	2.942651	3	1039	-	0	1177	2.073606	3		6.991466	2
	6.871687	2		0.88833	4		0.223338	5		0.870256	4	1290	1.699787	3
901	1.734086	3		0.952234	4	1042	2.550945	3		0.318915	4		5.278444	2
902	1.181077	3	973	-	0	1043	-	0		0.867922	4	1292	-	0
903	2.64297	3	976	2.073654	3	1044	-	0		0.099254	5	1295	-	0
	10.580979	1	977	-	0		1.088647	4		0.318916	4	1296	-	0
	6.437685	2		0.994332	4	1046	1.721671	3		0.085336	5	1297	-	0
906	-	0		0.767389	4	1047	2.945104	3		0.16287	5	1298	-	0

Table 4. Variability table for taxa set B.

Site	Variability	Cat	Site	Variability	Cat	Site	Variability	Cat	Site	Variability	Cat	Site	Variability	Cat
	4.137196	2		0.342442	4		0.055434	5	1621	-	0	1691	-	0
1301	2.066523	3	1399	-	0		0.10037	5	1622	-	0	1692	-	0
	5.050073	2	1400	-	0		1.004537	4	1623	-	0	1693	-	0
	3.748783	2	1404	3.044568	3	1547	1.201387	3	1624	-	0	1694	-	0
	4.030158	2	1406	-	0	1548	2.176435	3	1625	-	0	1695	-	0
	1.032752	4		0.122122	5		1.004549	4	1626	-	0	1696	-	0
	7.372039	1	1409	3.144292	3		1.004549	4	1627	-	0	1697	-	0
1308	1.208915	3		0.110174	5		1.004549	4	1628	-	0	1698	-	0
1309	-	0		0.110174	5	1559	2.942643	3	1629	-	0	1699	-	0
	0.105014	5	1413	-	0	1560	-	0	1630	-	0	1700	-	0
	0.897775	4		0.286228	5		0.777568	4	1631	-	0	1701	-	0
1316	2.788565	3		0.207483	5		1.004538	4	1632	-	0	1702	-	0
1317	-	0		0.282539	5	1567	2.942643	3	1633	-	0	1703	-	0
1318	-	0		0.197163	5		6.064565	2	1634	-	0	1704	-	0
	3.521061	2	1421	-	0		0.055434	5	1635	-	0	1705	-	0
1322	-	0		0.249173	5		0.103866	5	1636	-	0	1706	-	0
	0.223338	5		0.17698	5		0.055434	5	1637	-	0	1707	-	0
	0.085335	5		0.103867	5		0.084587	5	1638	-	0	1708	-	0
1330	-	0	1447	1.217765	3	1587	2.942936	3	1639	-	0	1709	-	0
	1.11505	4		0.194847	5		0.59067	4	1640	-	0	1710	-	0
	0.176979	5		0.282144	5		3.39748	2	1641	-	0	1711	-	0
	7.536263	1		4.684387	2	1593	1.802592	3	1642	-	0	1712	-	0
	0.107145	5		0.618602	4		0.801753	4	1643	-	0	1713	-	0
	0.317832	4	1468	1.61105	3		5.880794	2	1644	-	0	1714	-	0
	0.651223	4		0.609042	4		0.718139	4	1645	-	0	1715	-	0
	0.595197	4	1470	1.496837	3	1601	-	0	1646	-	0	1716	-	0
	0.282144	5	1471	2.657177	3		0.845176	4	1647	-	0		-	-1
	0.282812	5		0.473366	4		6.894108	2	1648	-	0		-	-1
1345	1.219445	3		0.138306	5		3.490597	2	1649	-	0		-	-1
	0.224716	5	1485	1.243285	3		3.642759	2	1650	-	0		-	-1
	0.223342	5		0.639934	4		5.469203	2	1651	-	0		-	-1
	0.223342	5		0.103866	5		8.669055	1	1652	-	0	17.088163	1	
	0.223342	5		0.640092	4	1608	-	0	1653	-	0	11.813357	1	
	0.199227	5		0.093563	5		5.1464	2	1654	-	0	10.049682	1	
	0.223341	5		1.120804	4		6.075157	2	1655	-	0	8.316597	1	
	0.093563	5	1507	1.611039	3		7.233139	1	1656	-	0	11.036735	1	
	0.282811	5	1512	-	0		6.954473	2	1657	-	0	4.970957	2	
	0.20249	5		0.076474	5		4.510309	2	1658	-	0	7.209176	1	
	0.190807	5		0.066073	5		15.581428	1	1659	-	0	3.906911	2	
	0.194068	5		0.194847	5	1615	-	0	1685	-	0	20.366158	1	
	0.089327	5		0.194847	5	1616	-	0	1686	-	0	4.22669	2	
	0.122557	5		0.715707	4	1617	-	0	1687	-	0	9.901206	1	
	0.290471	5	1532	-	0	1618	-	0	1688	-	0	4.685752	2	
1392	-	0	1535	1.125725	3	1619	-	0	1689	-	0	3.330656	2	
	0.294554	5		0.176979	5	1620	-	0	1690	-	0	1735	-	0

Table 4. Variability table for taxa set B.

Site	Variability	Cat	Site	Variability	Cat	Site	Variability	Cat	Site	Variability	Cat	Site	Variability	Cat
	5.540837	2		0.435026	4		7.689672	1		0.062285	5		4.300284	2
1737	2.858823	3		6.035917	2	1914	1.382366	3		0.594578	4	2110	-	0
	3.757946	2		6.102921	2	1917	1.464771	3		0.090382	5	2111	-	0
	7.08152	1	1858	1.524477	3		0.651237	4		0.078533	5		5.711187	2
1752	-	0		0.619124	4		0.441986	4	2039	-	0		5.251099	2
	0.103866	5		0.619124	4		10.916348	1		0.176977	5		7.885318	1
	0.103866	5	1861	1.816442	3	1927	1.984201	3		0.066072	5		6.253412	2
	0.103866	5	1862	-	0		5.979513	2		0.223341	5	2116	-	0
1756	-	0	1863	-	0		0.176994	5	2056	2.942847	3	2117	-	0
1757	1.68322	3	1864	-	0		0.317833	4	2057	1.125725	3	2118	-	0
	22.372915	1	1865	-	0		9.271133	1		0.079143	5		0.393321	4
1760	2.009835	3	1866	-	0		0.318912	4	2067	1.201386	3		8.07118	1
1761	-	0	1867	-	0		6.408242	2	2068	1.30576	3		5.272329	2
	7.231117	1	1868	-	0		3.925524	2		0.185877	5	2122	2.273361	3
1763	1.273527	3	1869	-	0	1938	-	0		3.794235	2		6.766105	2
	3.736887	2	1870	-	0	1943	2.942646	3		5.066354	2		6.587655	2
1765	2.161109	3	1871	-	0	1944	1.221869	3		3.278123	2		5.908135	2
	6.133319	2	1875	-	0		0.777568	4	2074	-	0		17.14352	1
	4.891889	2	1876	-	0		3.848276	2		5.309021	2		5.260294	2
1769	-	0	1877	-	0	1955	2.518901	3	2076	2.942675	3	2128	-	0
1770	-	0	1878	-	0	1956	1.41215	3		0.223971	5	2129	-	0
	0.140775	5	1879	-	0	1960	2.460857	3	2081	3.081217	3	2130	-	0
	6.563599	2	1883	-	0		0.882488	4		0.185878	5	2131	-	0
1773	1.802534	3	1884	-	0		0.886979	4		4.173811	2	2132	-	0
	22.762793	1	1885	-	0	1968	1.301808	3		10.869044	1	2133	-	0
	0.804438	4	1886	-	0		7.662072	1		0.159675	5		6.14751	2
1777	2.106157	3		4.897605	2		0.315614	5		3.982403	2		5.072736	2
	0.223338	5		4.530867	2	1974	1.579151	3		3.635116	2		4.278468	2
	0.089922	5		8.996841	1		0.066073	5		0.092489	5		4.554476	2
1784	1.739936	3	1890	3.117699	3	1985	2.826697	3		0.17479	5	2138	-	0
	0.22334	5		7.973341	1		0.055434	5		5.252111	2		7.962576	1
	0.609045	4		9.877855	1		0.23334	5	2093	1.839357	3		5.387676	2
	0.055434	5		6.367309	2		0.232166	5		6.056272	2		7.606258	1
	0.066073	5		0.101542	5		1.021165	4	2095	-	0		0.394096	4
	0.068799	5	1898	2.552688	3		3.281151	2		4.981063	2		0.249173	5
	0.09646	5	1899	1.201123	3		0.170739	5		6.971542	2		0.104422	5
	0.133313	5		0.140775	5		0.223339	5		6.634578	2		0.104422	5
1818	-	0		0.085336	5		0.803576	4		4.767314	2	2147	1.373574	3
1819	1.37357	3	1902	2.658007	3		3.280293	2		13.032255	1		0.210552	5
	0.098881	5		22.094467	1		0.240836	5		7.91533	1		1.063467	4
	0.133313	5	1905	-	0		0.106225	5		11.219984	1		3.437787	2
	0.09646	5		0.16286	5		0.873594	4		6.605432	2		4.73329	2
	0.09646	5		13.04104	1		0.093562	5		6.592031	2		4.137398	2
	3.844656	2	1909	-	0		7.980903	1	2106	1.247594	3		3.932129	2
	13.952195	1		3.297023	2		0.062285	5		7.838114	1		3.315608	2

Table 4. Variability table for taxa set B.

Site	Variability	Cat	Site	Variability	Cat
	0.389642	4	2251	-	0
	3.177977	2	2252	-	0
2159	-	0	2253	-	0
2160	1.27842	3	2254	-	0
	3.17761	2	2255	-	0
	5.208125	2	2256	-	0
2163	1.867603	3	2257	-	0
	0.148508	5	2258	-	0
	0.210897	5	2259	-	0
2166	1.278405	3	2260	-	0
2167	1.457551	3	2261	-	0
	0.32395	4	2262	-	0
2176	1.520867	3	2263	-	0
	3.70476	2	2264	-	0
	3.17618	2	2265	-	0
2187	-	0	2266	-	0
2188	-	0	2267	-	0
2189	-	0	2268	-	0
	0.166579	5	2269	-	0
	0.512631	4	2270	-	0
	0.230657	5	2271	-	0
	8.887003	1	2272	-	0
	0.471042	4	2273	-	0
	-	-1	2274	-	0
	-	-1	2275	-	0
	-	-1	2276	-	0
	-	-1	2277	-	0
	-	-1	2278	-	0
2234	-	0	2279	-	0
2235	-	0	2280	-	0
2236	-	0	2281	-	0
2237	-	0	2282	-	0
2238	-	0	2283	-	0
2239	-	0			
2240	-	0			
2241	-	0			
2242	-	0			
2243	-	0			
2244	-	0			
2245	-	0			
2246	-	0			
2247	-	0			
2248	-	0			
2249	-	0			
2250	-	0			

APPENDIX E : Secondary Structure

Figure 14	Secondary structure model for 18S rDNA of <i>Physcomitrella patens</i>	52
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Figure 14.

Secondary structure model for 18S rDNA of *Physcomitrella patens*.

The primary structure data was obtained from GenBank accession X80986. The sequence is written clockwise 5' → 3'. Helix numbering is according to Van de Peer *et al.* (1997).

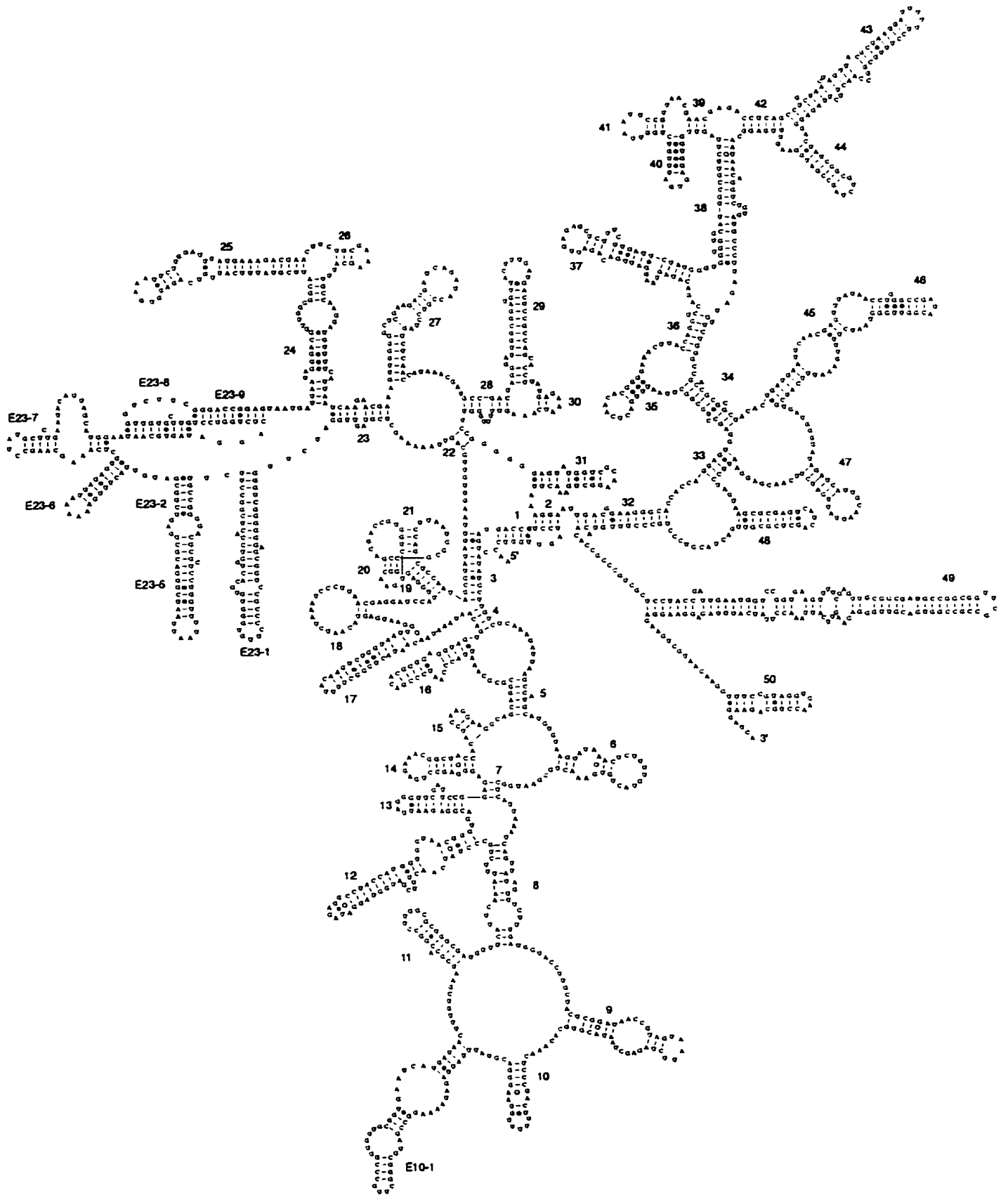


Figure 15.

Secondary structure model for 18S rDNA of *Physcomitrella patens* with primary and secondary structure corrections on helix 46.

The uncorrected primary structure data was obtained from Genbank accession X80986. The sequence is written clockwise 5' → 3'. Helix numbering is according to Van de Peer *et al.* (1997).

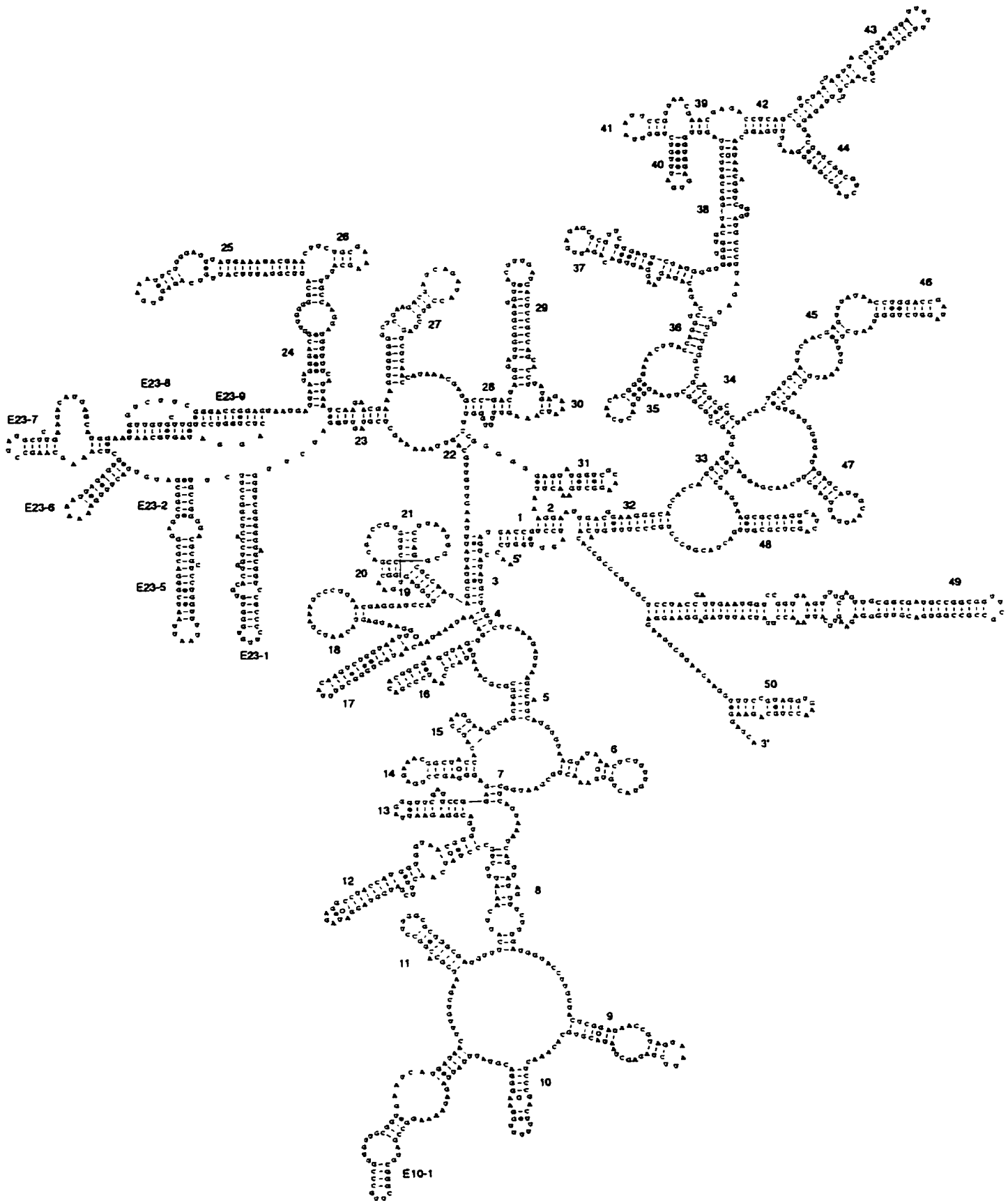
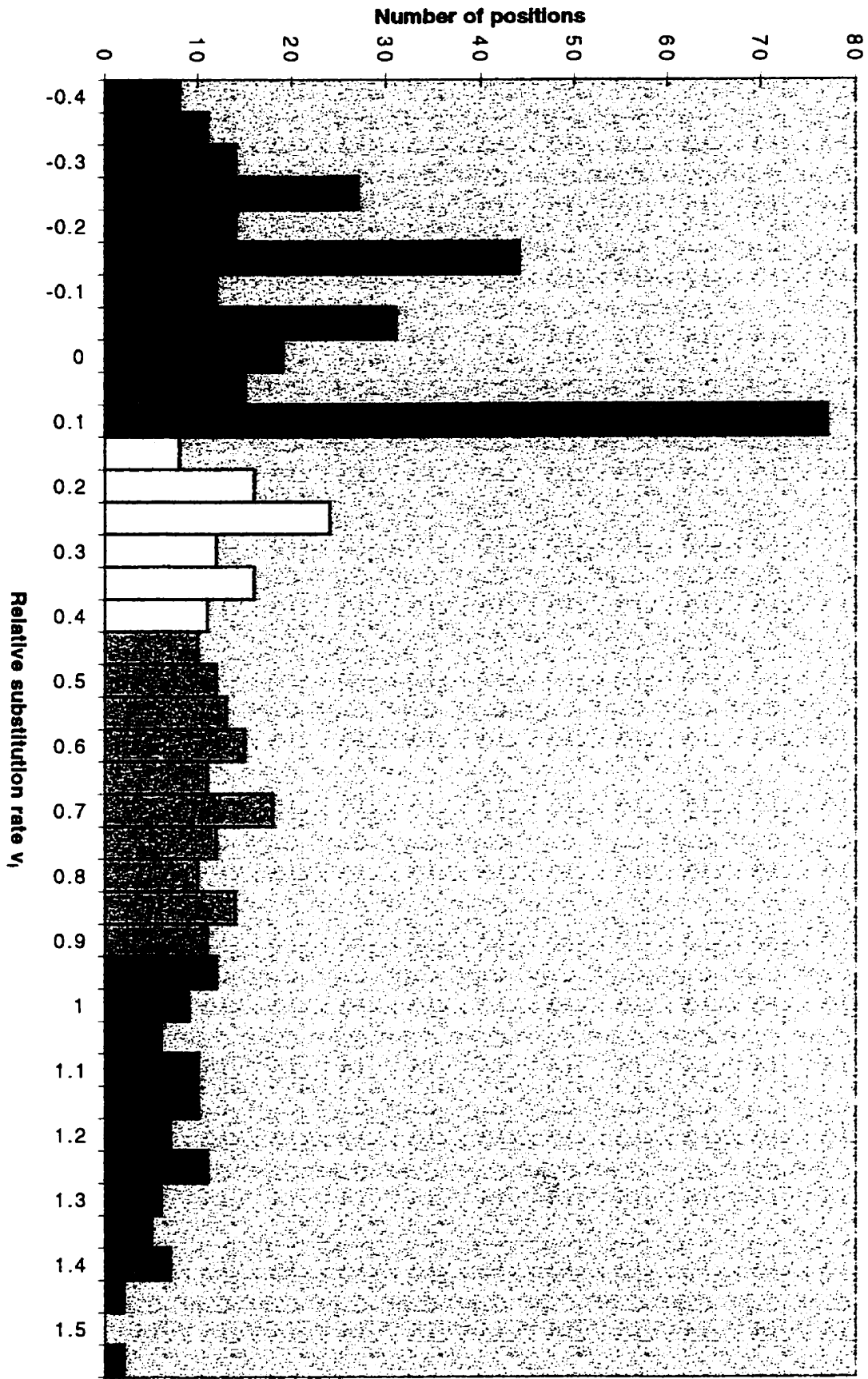


Figure 16.

Variability map of lower plants superimposed on the 18S rDNA secondary structure model of *Physcomitrella patens* shown in Figure 15.

Nucleotides are subdivided into five groups of increasing variability as described by Van de Peer *et al.* (1997). The most variable positions are in red (category 1) and, the most conserved in blue (category 5). Purple indicates the positions which are absolutely conserved, (category 6). Nucleotides present in *Physcomitrella patens* but absent in 75% of the lower plants considered are indicated in white. Grey indicates regions of undetermined variability due to ambiguous alignment.



Distribution of Relative Substitution Rates

Table 5.

Variability table for taxa set C.

The variabilities were calculated using taxa set C.

Column Headings


Site :refers to matrix location of MacClade Matrix

Variability :Score generated with 3 iterations of substitution rate calculation.

Cat :Variability category (shown in columns of MacClade Matrix)

Color Key

Cat #	Color
- 1	Grey
0	White
1	Red
2	Pink/Orange
3	Yellow
4	Green
5	Blue
6	Purple



Note: Nonvariable sites represented by purple are not included in table.

Table 5. Variability table for taxa set C.

Site	Variability	Cat	Site	Variability	Cat	Site	Variability	Cat	Site	Variability	Cat	Site	Variability	Cat	Site	Variability	Cat
	-	-1		7.313498	2		-	-1		3.631387	2	379	-	0		0.875497	4
23	2.400065	3	156	-	0		-	-1		23.843922	1		12.590858	1		2.717394	2
	1.003543	4		1.232586	4		-	-1	308	-	0		0.71236	5	590	2.505148	3
39	2.598499	3	158	1.852433	3		-	-1	309	-	0		4.848178	2	591	-	0
	0.926294	4		0.585814	5		-	-1		22.692989	1	387	1.568456	3		0.896541	4
	0.45451	5	161	-	0		11.689941	1		8.727041	1	388	-	0	597	-	0
	0.606526	5	165	1.983505	3		16.944672	1		18.401386	1		1.232585	4		0.376893	5
	1.160694	4	169	-	0		22.61495	1		8.233348	1		0.579161	5	598	2.20891	3
72	-	0	173	-	0	242	-	0	314	-	0	405	-	0		1.230248	4
73	-	0	178	-	0	243	-	0	315	-	0		0.976897	4	605	1.717242	3
74	-	0		1.230248	4	244	-	0	317	-	0		5.038437	2		0.896539	4
	4.224235	2	192	-	0	245	-	0	320	-	0	426	2.645902	3		0.976897	4
	12.306859	1		0.415194	5	246	-	0	321	1.34663	3		7.998219	2		0.896541	4
	8.566137	1		1.230248	4		6.590549	2	323	-	0		1.230248	4		0.896541	4
	7.440287	2		0.749565	4		10.867265	1		14.763712	1		12.991754	1		0.896541	4
	1.232585	4		4.523172	2		9.573917	1		0.517846	5		0.896539	4		3.302628	2
	0.79191	4	200	-	0		0.527933	5	330	-	0		3.853834	2		0.79191	4
83	-	0		9.901134	1		0.681978	5		3.460521	2		0.896539	4		1.230247	4
	2.953751	2		4.949099	2		0.385379	5		1.230251	4	460	1.325305	3		0.585813	5
87	-	0	204	-	0		0.997563	4		0.415194	5		0.385349	5	622	-	0
88	-	0		14.679193	1		5.063408	2		15.974884	1		1.232585	4		2.964889	2
91	-	0		5.13748	2		3.559758	2		16.96855	1		0.445487	5		0.744925	4
	0.897031	4		0.897031	4		9.07279	1		23.797117	1		0.71236	5	628	-	0
	0.976897	4	210	1.602081	3		1.232587	4	340	-	0		3.049086	2		0.79191	4
100	2.324674	3	211	-	0	279	1.75157	3	348	1.320645	3		0.590929	5	634	-	0
	0.71236	5	212	1.590599	3	280	-	0		0.744925	4		0.897031	4		0.896539	4
117	1.497844	3		7.277153	2	281	-	0		0.579161	5		1.230248	4		0.896539	4
121	2.452257	3		-	-1		0.944067	4	352	1.793893	3		1.230248	4	640	-	0
	0.68883	5		-	-1		3.564573	2		0.579162	5		0.590929	5		0.43188	5
127	-	0		-	-1		5.754246	2		0.579162	5	514	1.33712	3		0.896539	4
	0.85119	4		-	-1		0.579162	5	355	-	0	519	1.856973	3		0.842553	4
	3.128895	2		-	-1		1.144802	4	356	-	0	520	1.901095	3		0.655874	5
	0.71236	5		-	-1		3.371231	2		3.885621	2		1.232586	4		0.859801	4
	1.232585	4		-	-1		-	-1	361	-	0		0.590929	5		0.859801	4
	1.232585	4		-	-1		-	-1	362	1.712447	3		0.976897	4		0.625936	5
	1.232585	4		-	-1		-	-1		13.417314	1		1.094154	4		0.896683	4
	3.07286	2		-	-1		-	-1		12.757014	1	548	-	0		1.232586	4
	1.232587	4		-	-1		-	-1		24.381956	1		0.976897	4	703	1.87433	3
	0.897031	4		-	-1		-	-1	366	1.290763	3		0.517845	5		5.497921	2
	0.681978	5		-	-1		-	-1		6.684884	2	575	1.717251	3		5.252469	2
	2.868198	2		-	-1		-	-1	368	-	0		0.585814	5		0.90715	4
145	-	0		-	-1		-	-1		15.167713	1		1.230248	4		0.995272	4
146	-	0		-	-1		-	-1		0.590929	5		4.65137	2		1.239812	4
147	-	0		-	-1	300	1.558337	3		0.47586	5		0.906842	4		1.23259	4
	0.831836	4		-	-1		0.79191	4		0.744925	4		0.896539	4		1.230249	4
150	-	0		-	-1		1.132634	4	378	1.276647	3		1.232586	4		14.297762	1

Table 5. Variability table for taxa set C.

Site	Variability	Cat	Site	Variability	Cat	Site	Variability	Cat	Site	Variability	Cat	Site	Variability	Cat	Site	Variability	Cat
	0.608078	5	799	-	0		1.230248	4		0.475859	5		1.230249	4		0.71236	5
	26.794685	1		0.976897	4	849	1.844533	3	922	-	0		4.21441	2	1216	1.65957	3
746	-	0	801	2.59143	3	851	-	0	925	1.949421	3	1019	-	0		0.71236	5
	34.862289	1	802	-	0		0.79191	4		0.79191	4		18.699017	1		1.230248	4
748	1.788468	3	803	2.378535	3		1.232595	4	933	-	0	1022	2.668656	3		1.064245	4
750	1.963703	3	806	2.620925	3		3.220134	2	938	-	0		5.175593	2		0.415192	5
	13.608665	1		4.219388	2		4.82024	2		0.572653	5		8.259002	1		1.230248	4
	20.423738	1		4.644421	2	868	-	0		0.571523	5	1025	-	0		9.787221	1
753	-	0	809	-	0	869	1.426172	3	942	-	0		0.694456	5	1270	2.138923	3
	5.828836	2	810	-	0	870	1.572389	3	947	-	0	1033	-	0		1.100646	4
	16.300552	1		9.427018	1	871	-	0		1.232585	4		6.062812	2		8.953848	1
	11.987606	1		7.609166	2		5.294254	2		7.020597	2	1036	2.576682	3	1280	-	0
	4.937298	2	813	-	0	873	2.430647	3		1.232586	4	1039	-	0	1281	-	0
758	-	0	814	-	0		1.230248	4	954	-	0		1.232587	4	1282	-	0
765	-	0	815	-	0		0.976897	4	955	-	0		5.817531	2		7.964884	2
766	1.463314	3	816	-	0	877	2.59143	3	956	-	0	1043	-	0		16.169518	1
	5.38996	2	817	-	0		14.121339	1	957	-	0	1044	-	0	1285	1.583347	3
768	-	0	818	-	0		3.206238	2		8.899158	1	1045	2.301372	3	1287	-	0
769	-	0		23.75489	1	881	1.874331	3		6.581316	2		1.115244	4		21.747038	1
770	-	0		9.600008	1		20.268888	1		7.403722	2		0.385379	5		0.590929	5
771	-	0		17.612123	1		7.130877	2		7.041693	2		3.611351	2	1292	-	0
772	-	0		4.920897	2		17.795233	1	962	-	0	1085	1.522126	3	1295	-	0
773	-	0		19.445169	1	886	-	0	963	-	0	1087	1.832407	3	1296	-	0
774	-	0	824	-	0	887	-	0	964	-	0		6.514496	2	1297	-	0
775	-	0		37.154118	1	888	2.137763	3		0.71236	5		0.596708	5	1298	-	0
	-	-1		16.188332	1		10.991977	1	973	-	0		3.81175	2		12.166841	1
	-	-1		6.8196	2		11.666423	1		0.79191	4	1110	-	0	1301	1.833651	3
	-	-1		7.968797	2		13.117917	1	977	-	0		6.093278	2		1.100648	4
	-	-1		0.668578	5		1.232586	4		1.234016	4		2.764985	2		1.232587	4
	-	-1		5.035216	2		5.570765	2	984	-	0		4.214374	2	1305	1.623335	3
	-	-1		7.122333	2	897	-	0		1.230249	4		0.590928	5		8.358266	1
	-	-1	832	-	0		0.528368	5		1.064245	4		2.998769	2		0.585814	5
	-	-1	833	-	0		3.757417	2		0.475859	5	1139	1.649982	3	1309	-	0
	-	-1	834	-	0		0.681978	5		1.064239	4		5.743462	2		0.452203	5
	-	-1	835	-	0		0.791911	4	897	2.251053	3		3.596749	2		7.901381	2
	-	-1	836	-	0		8.97216	1	998	-	0		0.596708	5	1316	2.341105	3
	-	-1	837	-	0		0.681978	5		17.344212	1	1149	1.649981	3	1317	-	0
	-	-1	838	-	0	906	-	0		5.807863	2		12.240114	1	1318	-	0
	-	-1	839	-	0		9.565826	1		4.712387	2		0.579161	5		0.897033	4
	-	-1	840	-	0		0.79191	4		1.230248	4	1176	-	0	1322	-	0
	-	-1	841	-	0	911	2.072435	3		1.230248	4		0.385382	5		1.232586	4
	-	-1		19.986122	1		0.681978	5	1008	1.886223	3		0.71236	5		0.71236	5
793	-	0		19.772421	1		3.976127	2		4.285595	2		0.517843	5	1330	-	0
	1.232587	4		1.232585	4		0.475859	5	1011	-	0		1.230248	4		1.230248	4
	1.230249	4		12.967735	1		1.232585	4		10.062966	1		1.230248	4		7.170249	2
798	-	0		0.681978	5		1.094154	4		1.230249	4	1213	-	0		0.450999	5

Table 5. Variability table for taxa set C.

Site	Variability	Cat	Site	Variability	Cat	Site	Variability	Cat	Site	Variability	Cat	Site	Variability	Cat	Site	Variability	Cat	
1342	2.179529	3		0.590929	5	1638	-	0	1709	-	0		0.517846	5		3.930546	2	
1343	2.208907	3		1.094154	4	1639	-	0	1710	-	0		3.766107	2		0.451622	5	
	1.232586	4	1560	-	0	1640	-	0	1711	-	0		1.232584	4		0.71236	5	
	1.232586	4		0.590929	5	1641	-	0	1712	-	0		1.230249	4		5.529709	2	
	1.232586	4		0.976897	4	1642	-	0	1713	-	0		1.232584	4		23.22554	1	
1358	1.321306	3		0.590929	5	1643	-	0	1714	-	0		0.590929	5	1905	-	0	
	1.232584	4		0.897031	4	1644	-	0	1715	-	0		0.694456	5		0.744925	4	
	0.896539	4	1593	1.281927	3	1645	-	0	1716	-	0		0.744925	4		14.152463	1	
1365	2.208908	3	1594	1.89385	3	1646	-	0		-	-1		0.640894	5	1909	-	0	
1372	1.293607	3		1.230248	4	1647	-	0		-	-1		0.745625	4		9.584916	1	
1373	1.601586	3	1601	-	0	1648	-	0		-	-1	1818	-	0		13.889804	1	
1374	1.651814	3	1602	2.325185	3	1649	-	0		-	-1		0.640894	5		0.681978	5	
	0.736933	5		6.72905	2	1650	-	0		-	-1		0.745625	4		1.232585	4	
1379	1.286471	3		3.328416	2	1651	-	0	21.542904	1			0.640894	5		4.126468	2	
1390	1.379363	3	1605	2.325188	3	1652	-	0	10.897581	1			0.640894	5	1928	1.952729	3	
1392	-	0	1606	1.7309	3	1653	-	0	12.854319	1			8.114097	1	1830	1.424817	3	
1393	1.491091	3		19.256275	1	1654	-	0	13.896381	1			4.267783	2		8.436602	1	
1399	-	0	1608	-	0	1655	-	0	13.521935	1			0.71236	5	1938	-	0	
1400	-	0	1609	1.919692	3	1656	-	0	0.897031	4			0.71236	5		1.232587	4	
1406	-	0		3.223252	2	1657	-	0	5.125933	2	1862		-	0	1969	1.706745	3	
	1.036602	4		6.568538	2	1658	-	0	6.135058	2	1863		-	0	1973	1.783382	3	
	0.628137	5		3.374982	2	1659	-	0	26.879246	1	1864		-	0	1874	1.377244	3	
	0.628137	5		18.423962	1	1685	-	0	17.859688	1	1865		-	0		0.694456	5	
1413	-	0	1615	-	0	1686	-	0	0.639935	5	1866		-	0		0.590929	5	
	0.513402	5	1616	-	0	1687	-	0	1735	-	0	1867	-	0		1.232585	4	
	1.130359	4	1617	-	0	1688	-	0				3.564716	2	1868	-	0	0.560479	5
	1.26734	4	1618	-	0	1689	-	0				0.744925	4	1869	-	0	0.896539	4
1419	1.920922	3	1619	-	0	1690	-	0	1738	1.930106	3	1870	-	0		3.82684	2	
1421	-	0	1620	-	0	1691	-	0		8.480633	1	1871	-	0		0.436233	5	
	1.211314	4	1621	-	0	1692	-	0	1752	-	0	1875	-	0		0.436233	5	
	1.230248	4	1622	-	0	1693	-	0		0.9769	4	1876	-	0		0.436233	5	
	0.976897	4	1623	-	0	1694	-	0		0.9769	4	1877	-	0		0.508977	5	
1449	1.7309	3	1624	-	0	1695	-	0		0.9769	4	1878	-	0		0.681978	5	
1450	2.179529	3	1625	-	0	1696	-	0	1756	-	0	1879	-	0	2039	-	0	
1453	1.783381	3	1626	-	0	1697	-	0		4.327728	2	1883	-	0		1.230248	4	
	1.064241	4	1627	-	0	1698	-	0		26.128506	1	1884	-	0		0.694456	5	
	0.976897	4	1628	-	0	1699	-	0		9.143496	1	1885	-	0		1.232585	4	
	0.976897	4	1629	-	0	1700	-	0	1761	-	0	1886	-	0		0.68883	5	
	0.896541	4	1630	-	0	1701	-	0		7.401666	2		3.997552	2		1.232585	4	
1512	-	0	1631	-	0	1702	-	0		3.642895	2		4.140663	2		0.79191	4	
	0.47586	5	1632	-	0	1703	-	0		3.766142	2		2.784576	2	2074	-	0	
	0.694456	5	1633	-	0	1704	-	0	1769	-	0		6.760604	2		1.230247	4	
1520	1.7309	3	1634	-	0	1705	-	0	1770	-	0		16.033895	1		0.803386	4	
1521	1.7309	3	1635	-	0	1706	-	0		9.037307	1		6.582469	2		5.125934	2	
1532	-	0	1636	-	0	1707	-	0		7.610094	2	1896	1.809742	3		9.114459	1	
	1.230248	4	1637	-	0	1708	-	0		24.898895	1		1.094154	4		0.68883	5	

Table 5. Variability table for taxa set C.

Site	Variability	Cat	Site	Variability	Cat	Site	Variability	Cat
	10.471128	1		1.23911	4	2261	-	0
	0.465398	5		5.718729	2	2262	-	0
	3.220219	2		0.745625	4	2263	-	0
2095	-	0	2153	2.283268	3	2264	-	0
	3.912166	2	2154	1.822172	3	2265	-	0
	2.679604	2	2156	2.672781	3	2266	-	0
	11.410489	1	2159	-	0	2267	-	0
	6.03823	2		3.127488	2	2268	-	0
	5.755039	2		0.485144	5	2269	-	0
	3.362382	2	2187	-	0	2270	-	0
	3.315792	2	2188	-	0	2271	-	0
2106	2.120934	3	2189	-	0	2272	-	0
	5.2207	2		0.916255	4	2273	-	0
2108	1.862846	3	2201	1.658721	3	2274	-	0
2110	-	0		-	-1	2275	-	0
2111	-	0		-	-1	2276	-	0
	4.871907	2		-	-1	2277	-	0
	2.860922	2		-	-1	2278	-	0
	3.509642	2		-	-1	2279	-	0
	4.956572	2	2234	-	0	2280	-	0
2116	-	0	2235	-	0	2281	-	0
2117	-	0	2236	-	0	2282	-	0
2118	-	0	2237	-	0	2283	-	0
	0.656031	5	2238	-	0			
	4.686344	2	2239	-	0			
2121	1.587528	3	2240	-	0			
	7.657166	2	2241	-	0			
	4.619767	2	2242	-	0			
	0.745625	4	2243	-	0			
	15.034582	1	2244	-	0			
	0.70407	5	2245	-	0			
2128	-	0	2246	-	0			
2129	-	0	2247	-	0			
2130	-	0	2248	-	0			
2131	-	0	2249	-	0			
2132	-	0	2250	-	0			
2133	-	0	2251	-	0			
2134	1.774515	3	2252	-	0			
	0.745625	4	2253	-	0			
	0.745625	4	2254	-	0			
2138	-	0	2255	-	0			
2141	2.345036	3	2256	-	0			
2142	2.004233	3	2257	-	0			
	1.211314	4	2258	-	0			
	0.541494	5	2259	-	0			
	0.541494	5	2260	-	0			

APPENDIX F : Trees

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Table 6. Character-Based phylogenetic trees; Parsimony

Figure	Data Set	Description	Page
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Table 7. Character-based phylogenetic trees; Maximum likelihood

Figure	Data Set	Description	Page
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Table 8. Distance-Based phylogenetic trees; Taxa set A.

Figure	Distance Calculation	Insertion & Deletions taken into account	Alignment Positions	Bootstrap	Evolution Model	Page
A1S	Jukes & Cantor (1969)	No	All	No	~	81
A1BS	"	No	All	100	~	83
A2S	"	Yes	All	No	~	85
A2BS	"	Yes	All	100	~	87
B1S	Tajima & Nei (1984)	No	All	No	~	89
B1BS	"	No	All	100	~	91
B2S	"	Yes	All	No	~	93
B2BS	"	Yes	All	100	~	95
C1S	Kimura (1980)	No	All	No	~	97
C1BS	"	No	All	100	~	99
C2S	"	Yes	All	No	~	101
C2BS	"	Yes	All	100	~	103
D1.1S	Jin & Nei (1990)	No	All	No	Jukes & Cantor	105
D1.1BS	"	No	All	100	Jukes & Cantor	107
D1.2S	"	Yes	All	No	Jukes & Cantor	109
D1.2BS	"	Yes	All	100	Jukes & Cantor	111
D2.1S	"	No	All	No	Kimura 2-p	113
D2.1BS	"	No	All	100	Kimura 2-p	115
D2.2S	"	Yes	All	No	Kimura 2-p	117
D2.2BS	"	Yes	All	100	Kimura 2-p	119
E1S	Transversions Only	No	All	No	~	121
E1BS	"	No	All	100	~	123
E2S	"	Yes	All	No	~	125
E2BS	"	Yes	All	100	~	127
F1S	No Correction	No	All	No	~	129
F1BS	"	No	All	100	~	131
F2S	"	Yes	All	No	~	133
F2BS	"	Yes	All	100	~	135
G1S	Van de Peer & De Wachter (1996)	No	All	No	~	137
G1BS	"	No	All	100	~	139
G2S	"	Yes	All	No	~	141
G2BS	"	Yes	All	100	~	143

Figure Label Key:

- A First letter: Distance Calculation (i.e. A= Jukes & Cantor)
- 1 Number: indel status (e.g. 1= not taken into account, 2= taken into account)
- B If B is present = 50% majority-rule bootstrap tree
- S Indicates taxa set A

Table 9. Distance-Based phylogenetic trees; Taxa set B.

Figure	Distance Calculation	Insertion & Deletions taken into account	Alignment Positions	Bootstrap	Evolution Model	Page
A1	Jukes & Cantor (1969)	No	All	No	~	145
A1B	"	No	All	100	~	147
A2	"	Yes	All	No	~	149
A2B	"	Yes	All	100	~	151
B1	Tajima & Nei (1984)	No	All	No	~	153
B1B	"	No	All	100	~	155
B2	"	Yes	All	No	~	157
B2B	"	Yes	All	100	~	159
C1	Kimura (1980)	No	All	No	~	161
C1B	"	No	All	100	~	163
C2	"	Yes	All	No	~	165
C2B	"	Yes	All	100	~	167
D1.1	Jin & Nei (1990)	No	All	No	Jukes & Cantor	169
D1.1B	"	No	All	100	Jukes & Cantor	171
D1.2	"	Yes	All	No	Jukes & Cantor	173
D1.2B	"	Yes	All	100	Jukes & Cantor	175
D2.1	"	No	All	No	Kimura 2-p	177
D2.1B	"	No	All	100	Kimura 2-p	179
D2.2	"	Yes	All	No	Kimura 2-p	181
D2.2B	"	Yes	All	100	Kimura 2-p	183
E1	Transversions Only	No	All	No	~	185
E1B	"	No	All	100	~	187
E2	"	Yes	All	No	~	189
E2B	"	Yes	All	100	~	191
F1	No Correction	No	All	No	~	193
F1B	"	No	All	100	~	195
F2	"	Yes	All	No	~	197
F2B	"	Yes	All	100	~	199
G1	Van de Peer & De Wachter (1996)	No	All	No	~	201
G1B	"	No	All	100	~	203
G2	"	Yes	All	No	~	205
G2B	"	Yes	All	100	~	207
H1B	No Correction	No	0,1,2,3,4	100	~	209
H2B	"	No	0,1,2	100	~	211
H3B	"	No	2,3,4,5,6	100	~	213

Figure Label Key:

- A First letter: Distance Calculation (i.e. A= Jukes & Cantor)
- 1 Number: indel status (e.g. 1= not taken into account, 2= taken into account)
- B If B is present = 50% majority-rule bootstrap tree

Table 10.

Consistent monophyletic clusters in distance-based analyses.

These monophyletic clusters constitute the triangles shown in the second tree of each pair for all distance-based analysis.

Taxa set A : Monophyletic Clusters
Taxa set B : Monophyletic Clusters

Pinophyta	<i>Pinus luchuensis</i> <i>Taxus mairei</i> <i>Nageia nagi</i> <i>Gnetum leyboldii</i>
Magnoliophyta	<i>Zea mays</i> <i>Oryza sativa</i> <i>Glycine max</i> <i>Sinapis alba</i> <i>Arabidopsis thaliana</i>
Isoetopsida	<i>Selaginella vogelii</i> <i>Selaginella umbrosa</i>
Lycopodiopsida	<i>Lycopodium taxifolium</i> <i>Lycopodium phlegmaria</i>
Bryopsida	<i>Physcomitrella patens</i> <i>Funaria hygrometrica</i> <i>Mnium hornum</i> <i>Leptobryum pyriforme</i>
Anthocerotopsida	<i>Anthoceros laevis</i> <i>Notophylas breutelu</i> <i>Megaceros aenigmaticus</i>
Hepaticopsida	<i>Riccia fluitans</i> <i>Conocephalum conicum</i> <i>Marchantia polymorpha</i> <i>Sphaerocarpos donnelli</i>

Pinophyta	<i>Pinus wallichiana</i> <i>Pinus luchuensis</i> <i>Taxus mairei</i> <i>Nageia nagi</i> <i>Gnetum leyboldii</i>
Magnoliophyta	<i>Zea mays</i> <i>Oryza sativa</i> <i>Glycine max</i> <i>Sinapis alba</i> <i>Arabidopsis thaliana</i>
Isoetopsida	<i>Selaginella vogelii</i> <i>Selaginella umbrosa</i> <i>Isoetes engelmannii</i>
Lycopodiopsida	<i>Lycopodium taxifolium</i> <i>Lycopodium phlegmaria</i> <i>Lycopodium tristachyum</i> <i>Huperzia lucidulum</i>
Bryopsida	<i>Physcomitrella patens</i> <i>Physcomitrella patens 2</i> <i>Funaria hygrometrica</i> <i>Physcomitrium pyriforme</i> <i>Mnium hornum</i> <i>Leptobryum pyriforme</i> <i>Eurhynchium hians</i> <i>Bryum argenteum</i>
Anthocerotopsida	<i>Anthoceros laevis</i> <i>Notophylas breutelu</i> <i>Megaceros aenigmaticus</i>
Hepaticopsida	<i>Riccia fluitans</i> <i>Conocephalum conicum</i> <i>Marchantia polymorpha</i> <i>Reboulia hemisphaerica</i> <i>Sphaerocarpos texanus</i> <i>Sphaerocarpos donnelli</i>
Charales	<i>Chara foetida</i> <i>Nitella sp.</i> <i>Nitella flexilis</i>
Chlorophyta	<i>Parietochloris pseudoalveolaris</i> <i>Hydrodictyon reticulatum</i> <i>Chlamydomonas reinhardtii</i>

Majority rule

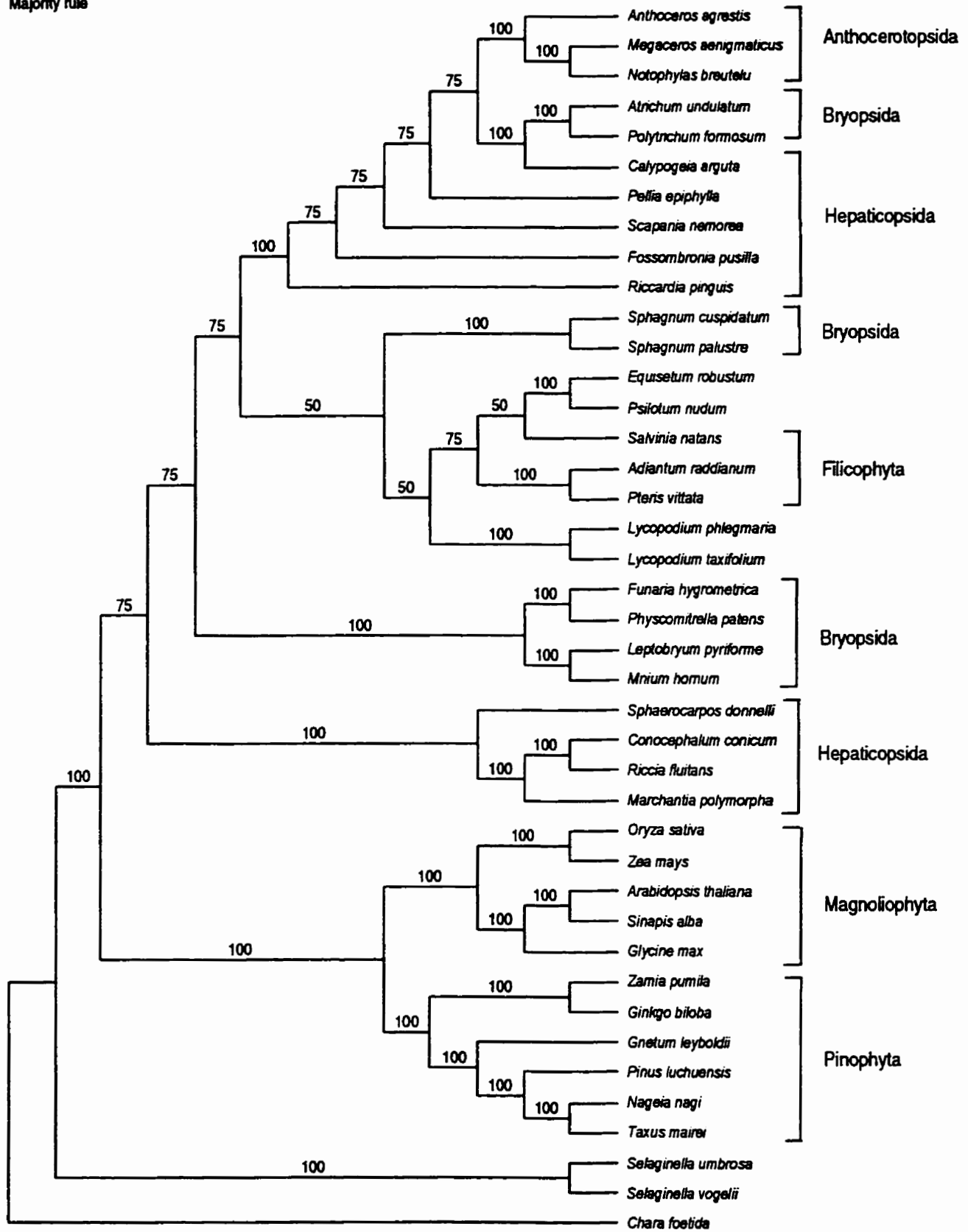


Figure A Parsimony; 50% Majority-Rule Consensus Tree of four most parsimonious trees (tree length = 1589) Taxa set A; informative sites only; Heuristic (100 replicates; random taxon addition; TBR)

50% Majority-rule Bootstrap

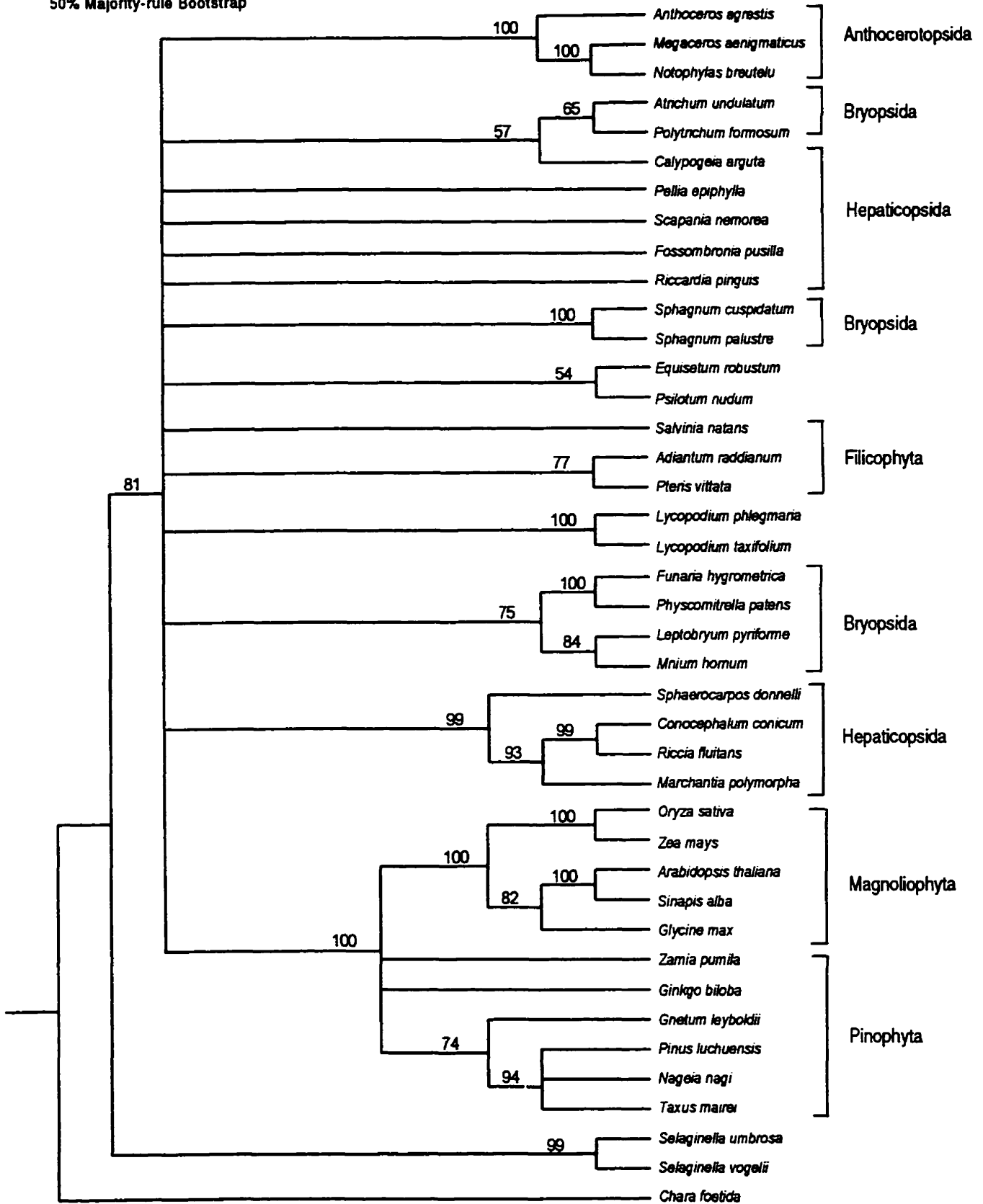


Figure B Parsimony; 50% Majority-Rule Bootstrap Tree
 Taxa set A; informative sites only;
 Heuristic (100 bootstrap replicates; each with 10 replicates of random taxon addition; TBR)

Strict

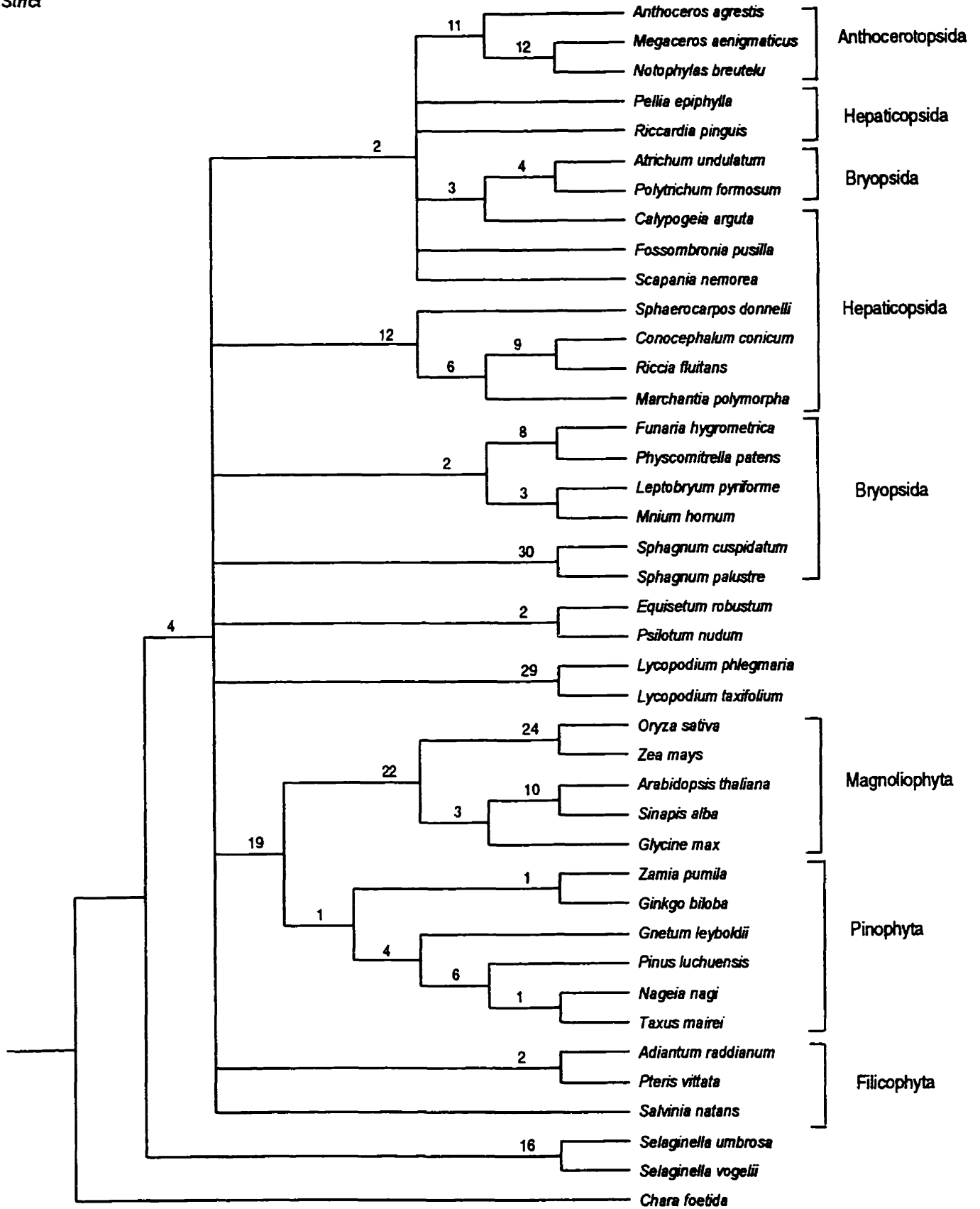


Figure C Parsimony; Strict Consensus Tree of four most parsimonious trees (tree length = 1589) Taxa set A; informative sites only; Heuristic (100 replicates; random taxon addition; TBR) Decay Indices are shown above each branch

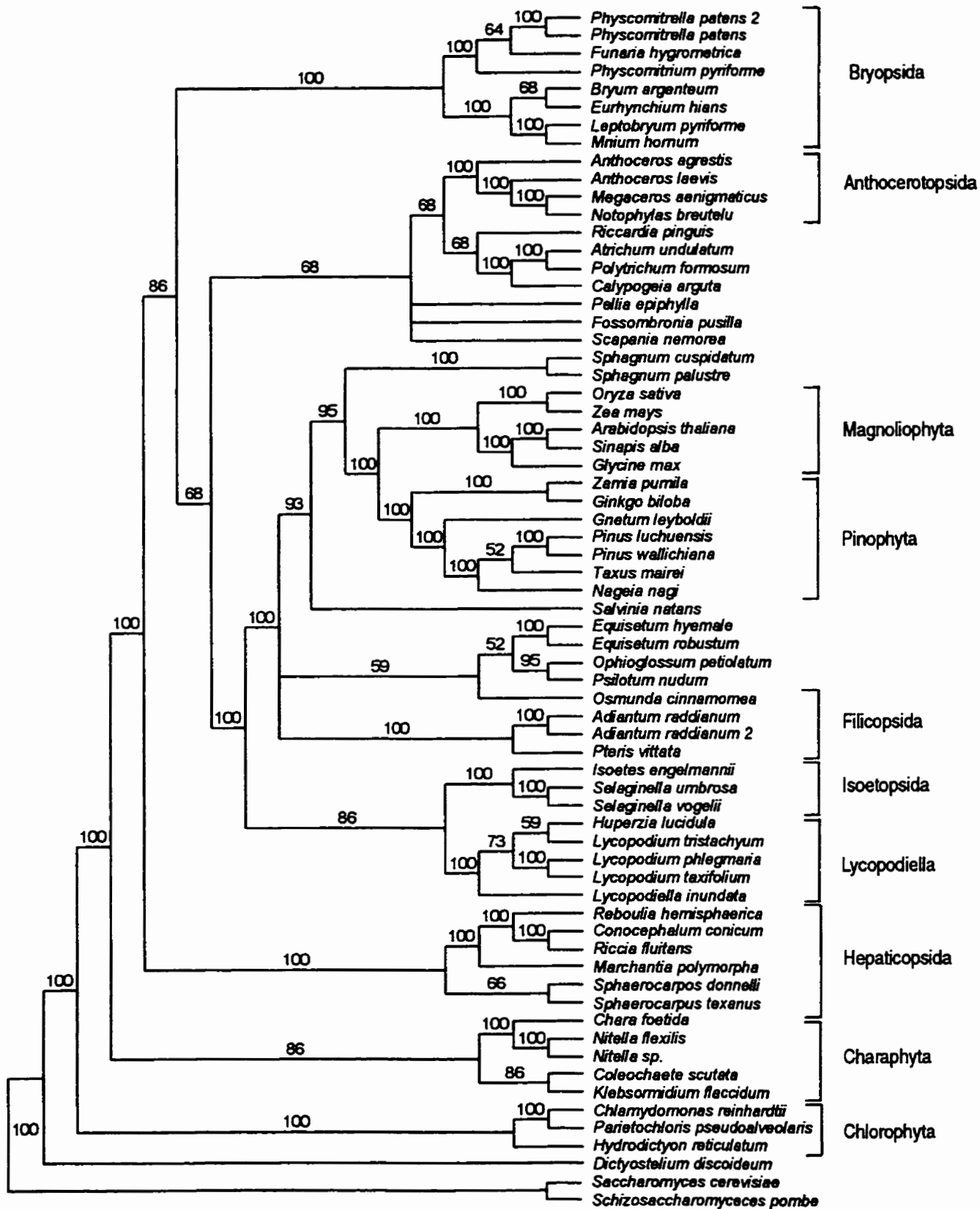


Figure D Parsimony; 50% Majority-Rule Consensus Tree of 88 most parsimonious trees (tree length = 3156) Taxa set B; informative sites only; Heuristic (100 replicates; random taxon addition; TBR)

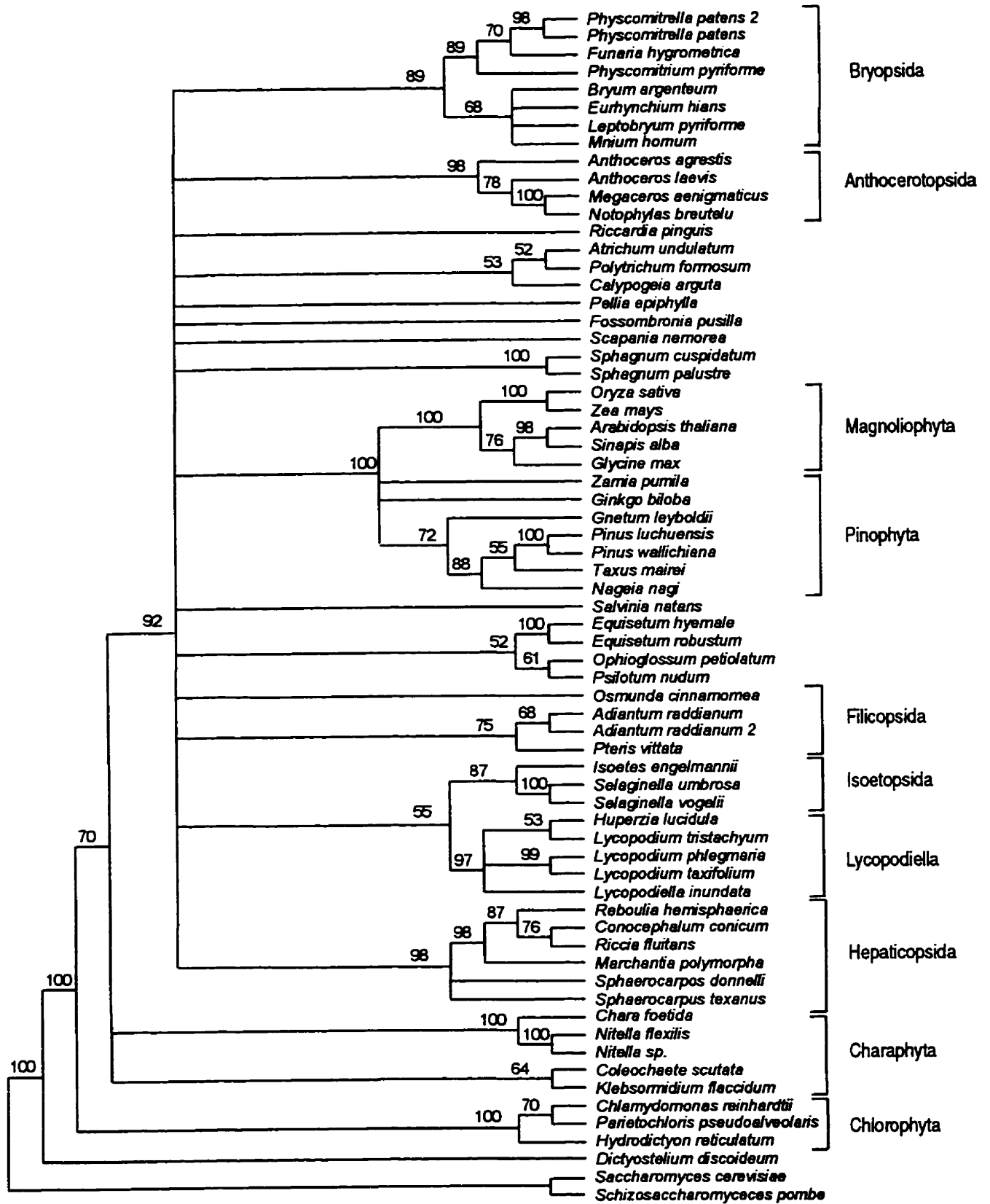


Figure E Parsimony; 50% Majority-Rule Bootstrap Tree
 Taxa set B; informative sites only;
 Heuristic (50 bootstrap replicates; each with 5 replicates of random taxon addition; TBR)

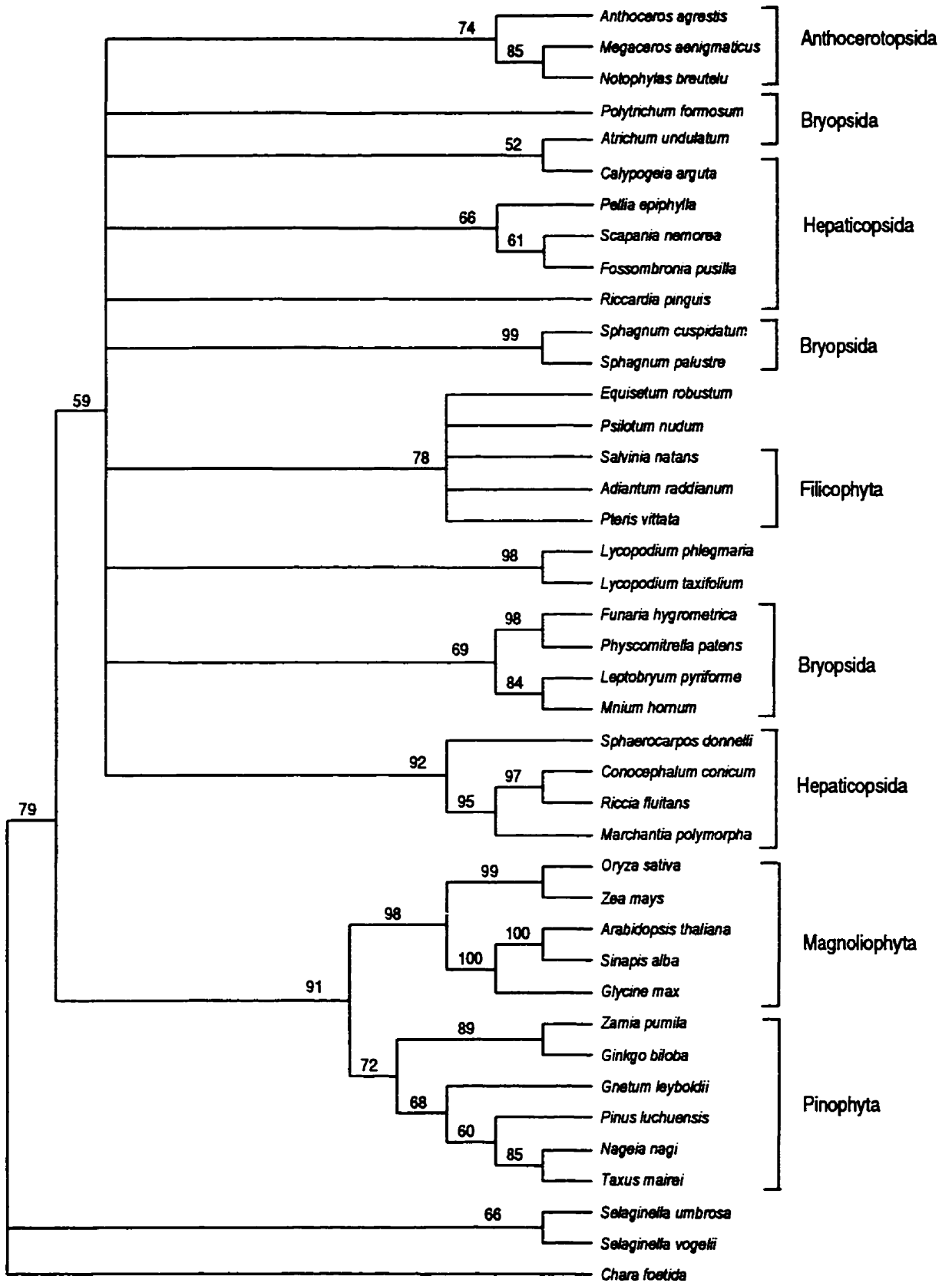


Figure F Quartet puzzling; Taxa set A

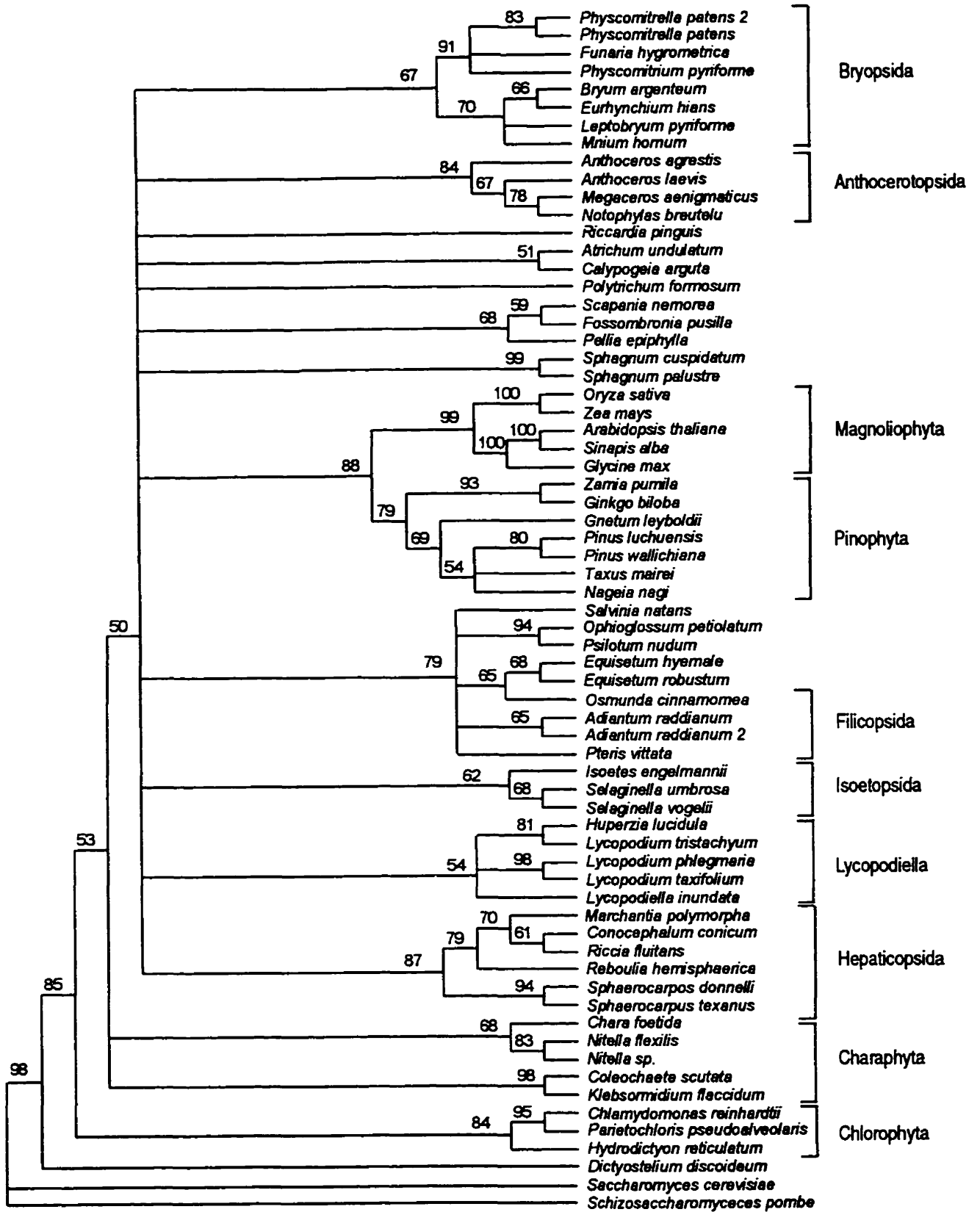


Figure G Quartet puzzling; Taxa set B

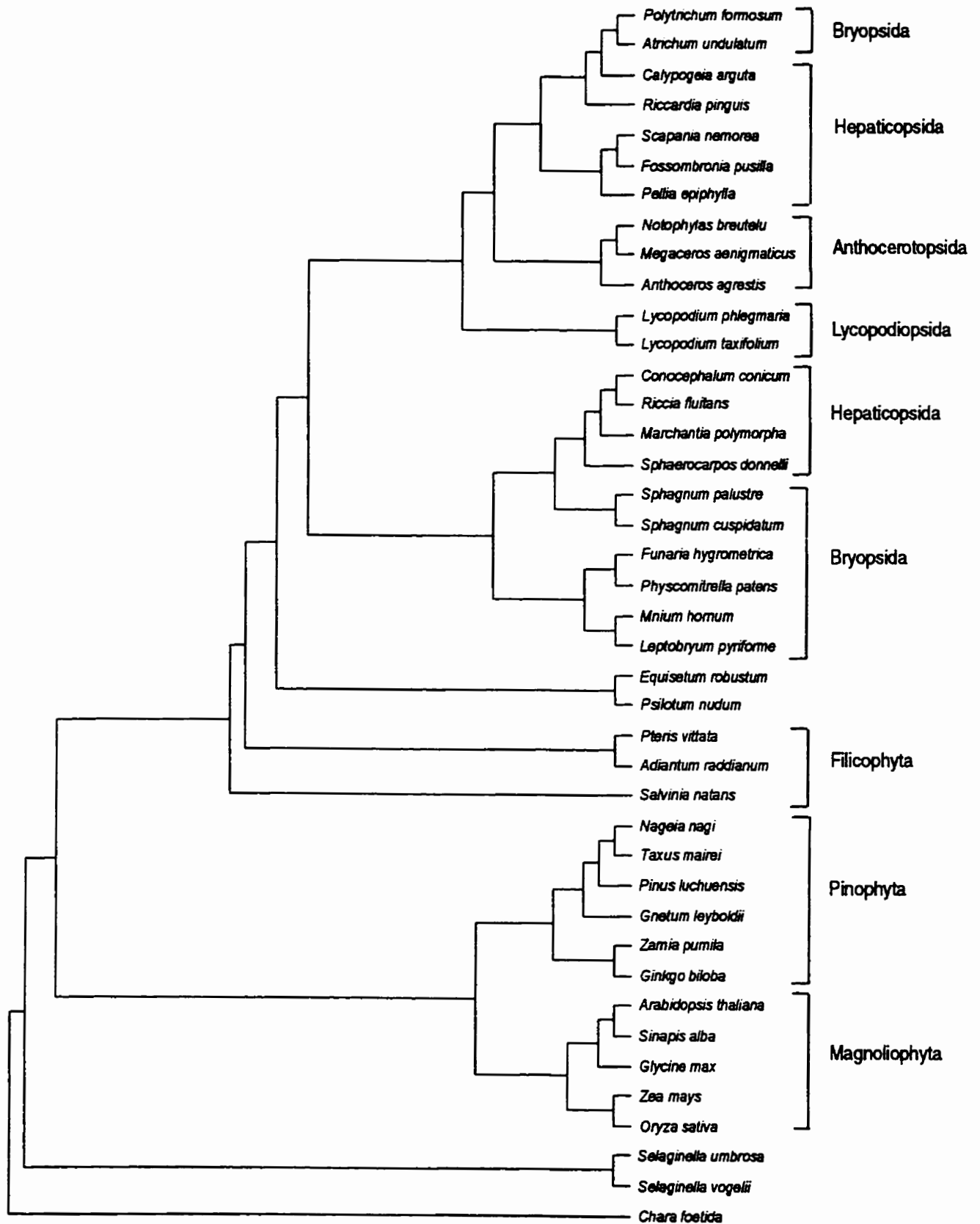


Figure H Maximum Likelihood; Taxa set A
Ln Likelihood= -13291.72141

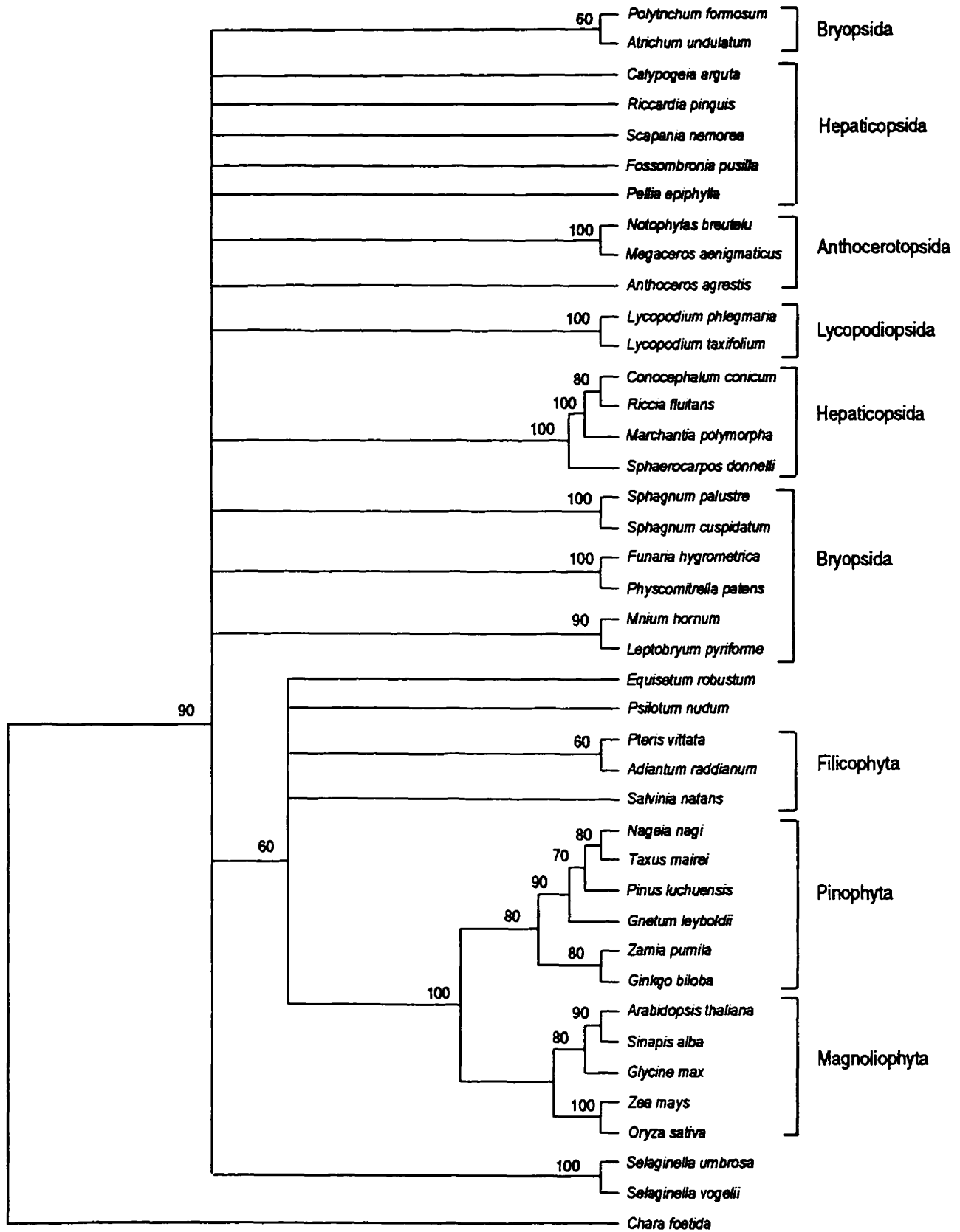


Figure 1 Maximum Likelihood; 50% Majority-Rule Bootstrap Tree; Taxa set A 10 Bootstraps

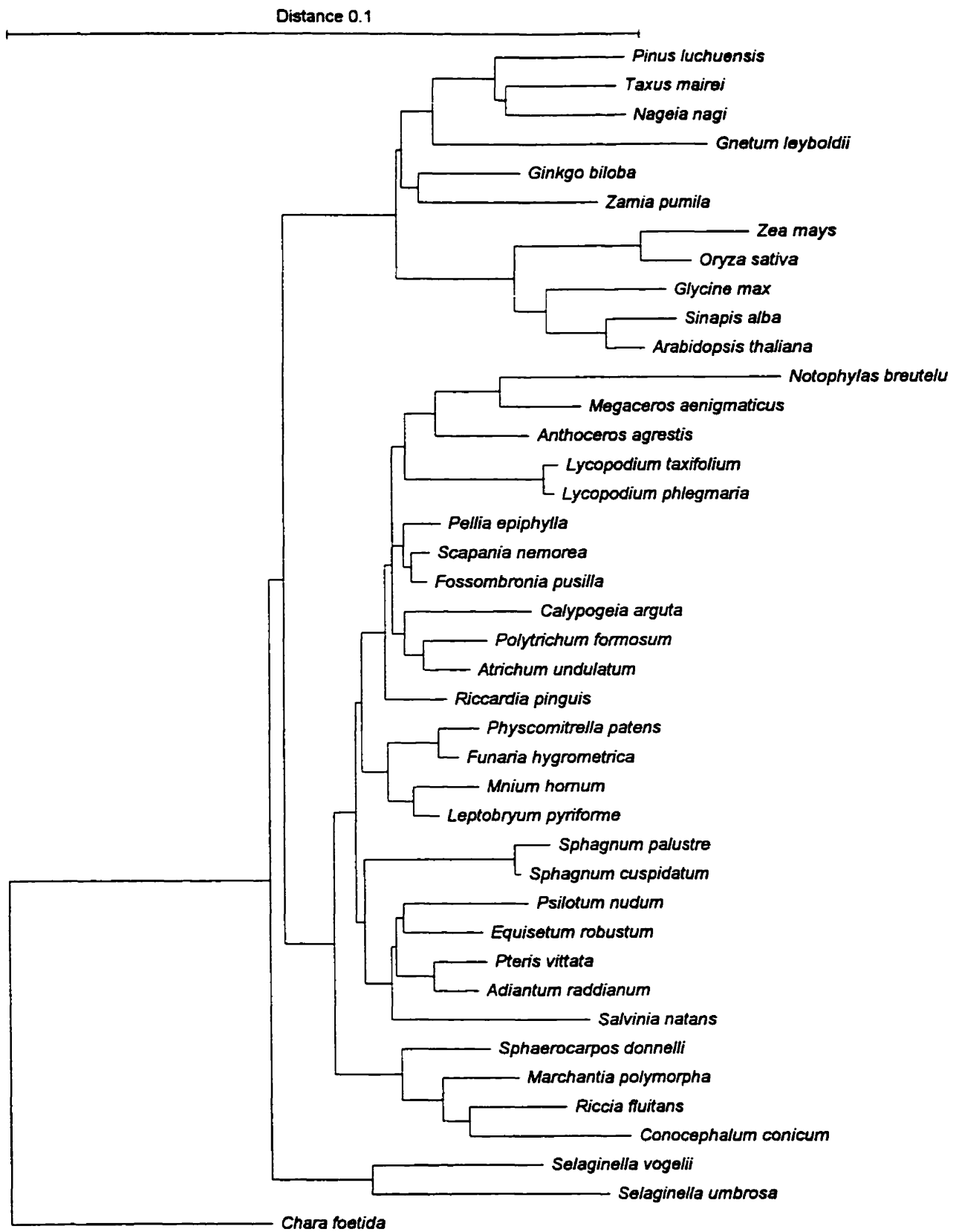


Figure A1S Jukes & Cantor, Neighbor-joining

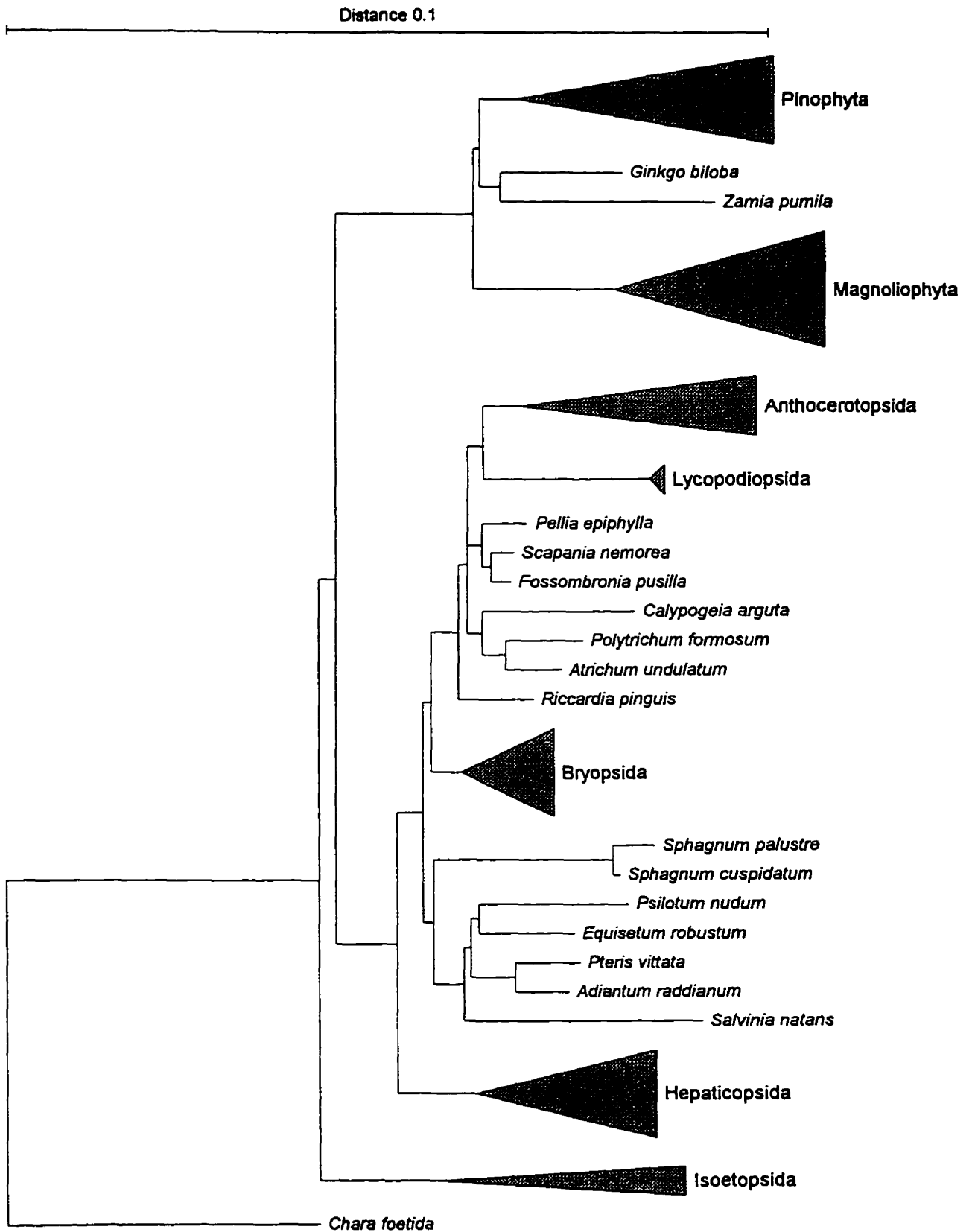


Figure A1S Jukes & Cantor, Neighbor-joining

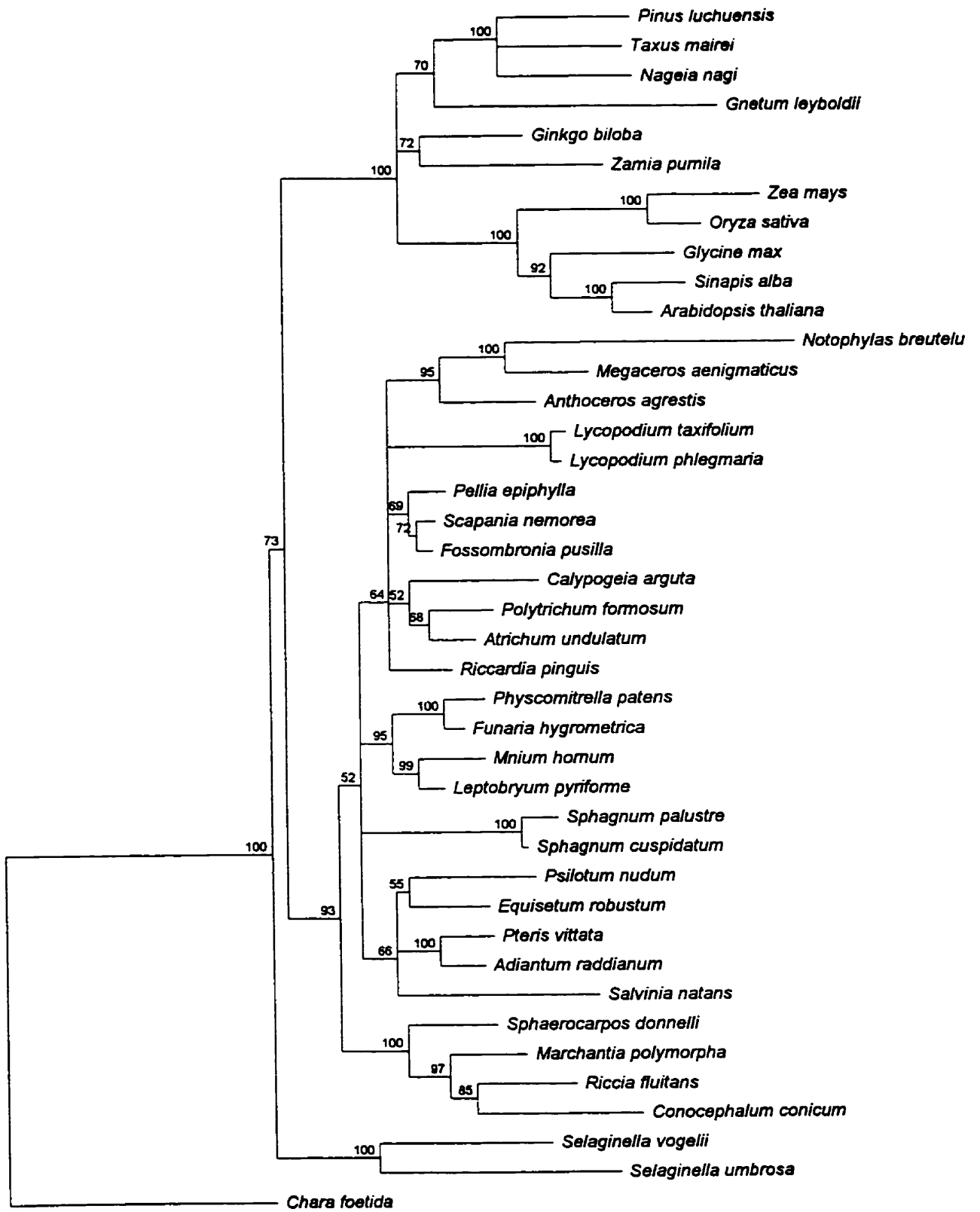


Figure A1SB Jukes & Cantor, Neighbor-joining
50% Majority-Rule Bootstrap Tree

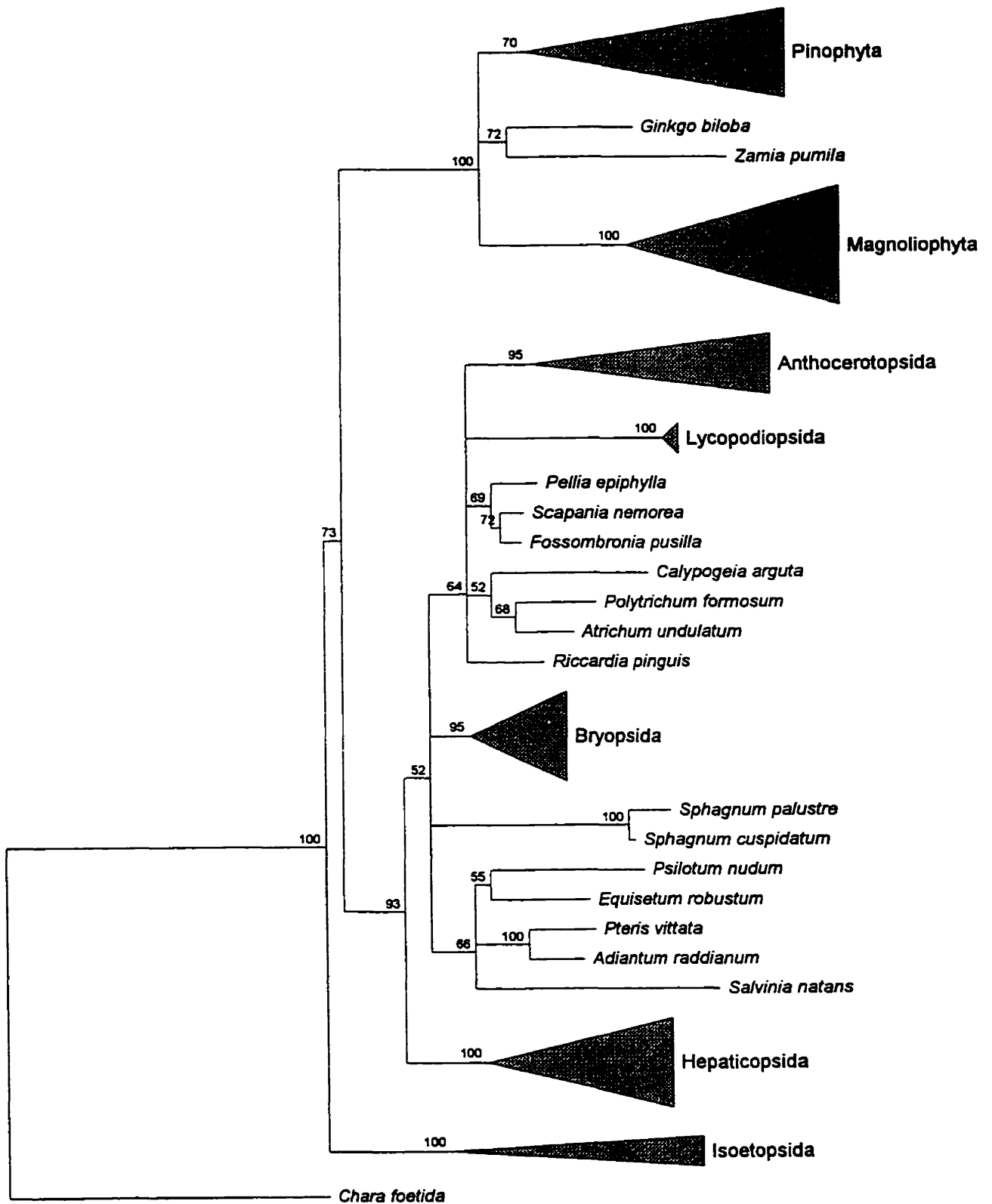


Figure A1SB Jukes & Cantor; Neighbor-joining
50% Majority-Rule Bootstrap Tree

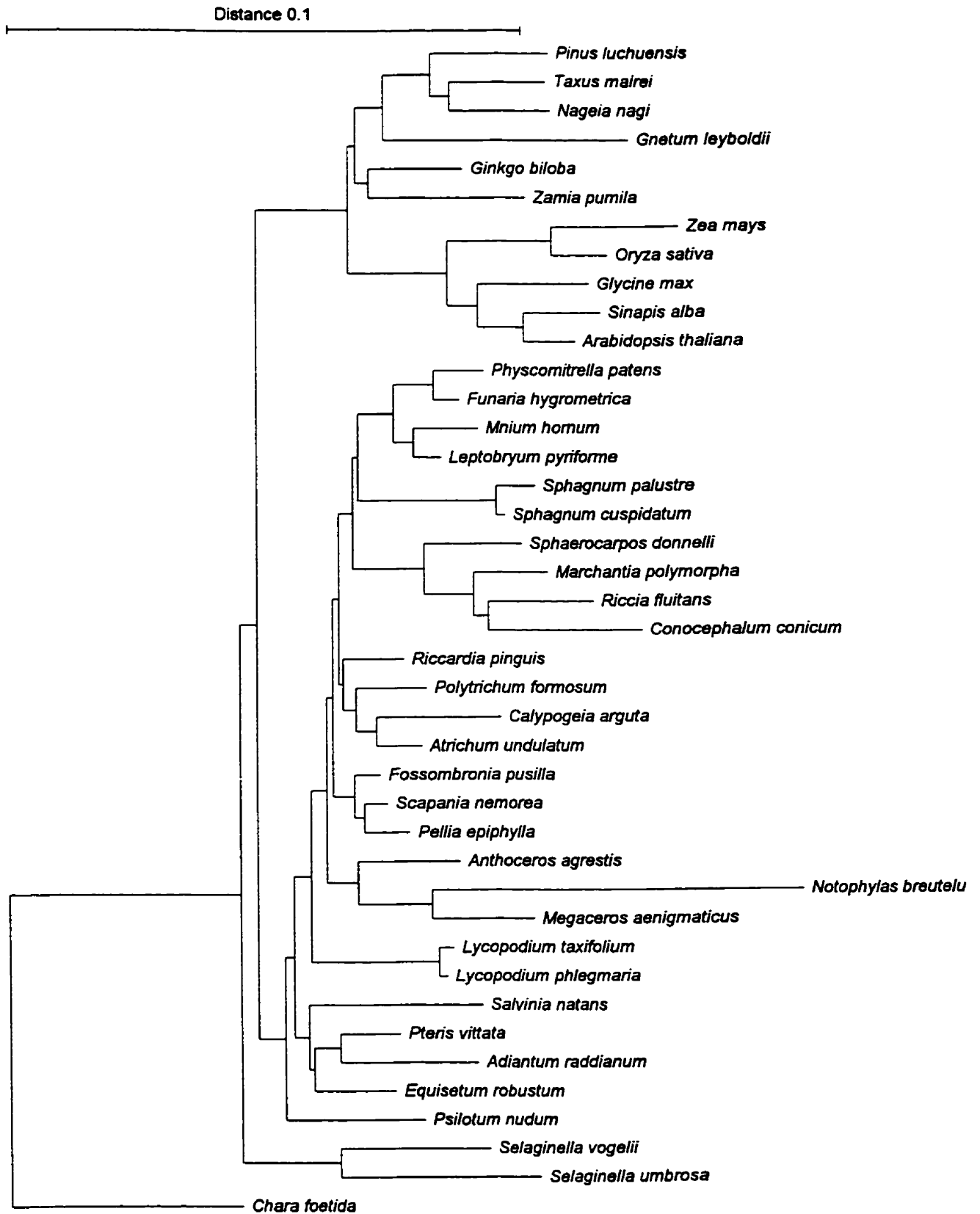


Figure A2S Jukes & Cantor
 Insertions & Deletions taken into account

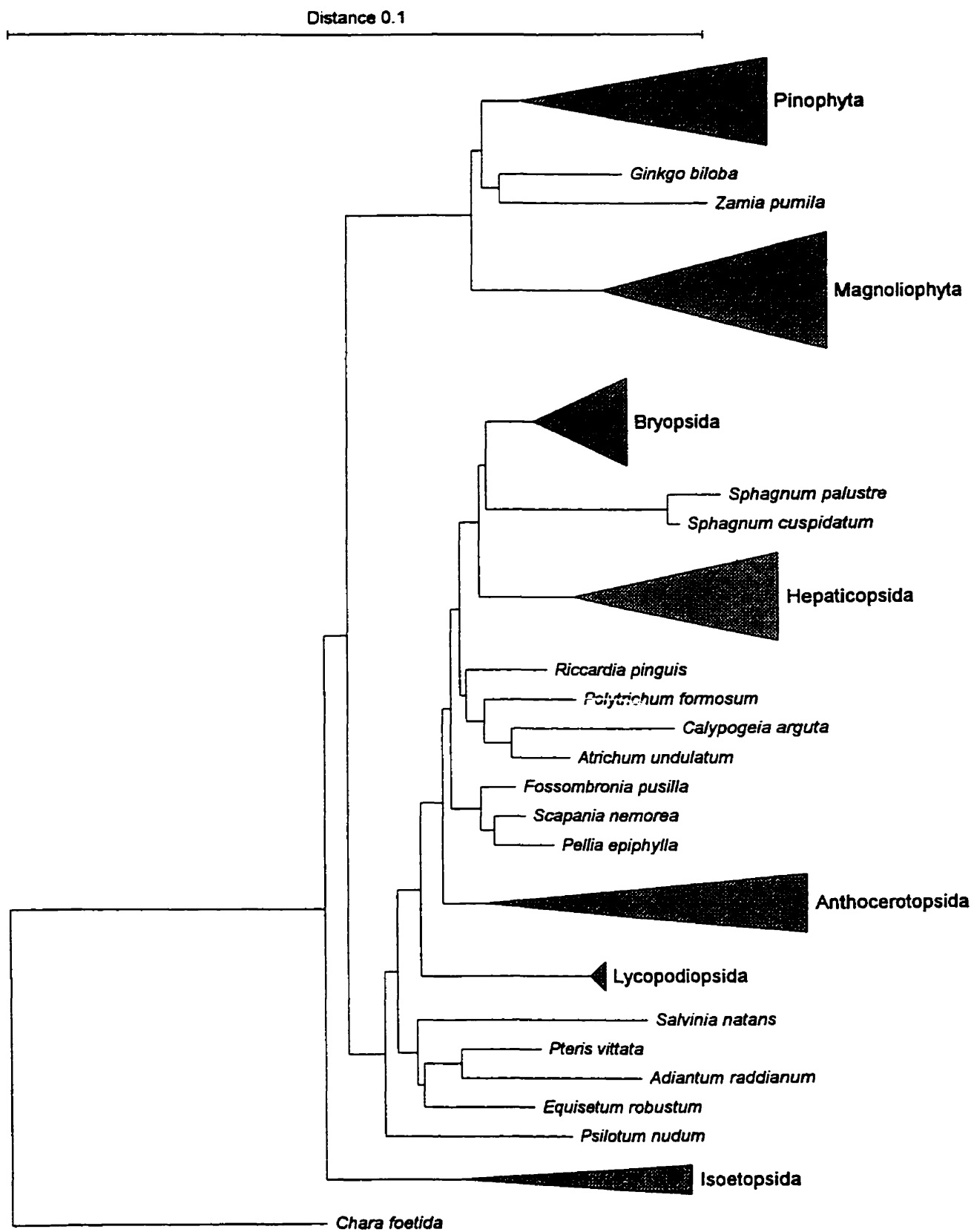


Figure A2S Jukes & Cantor
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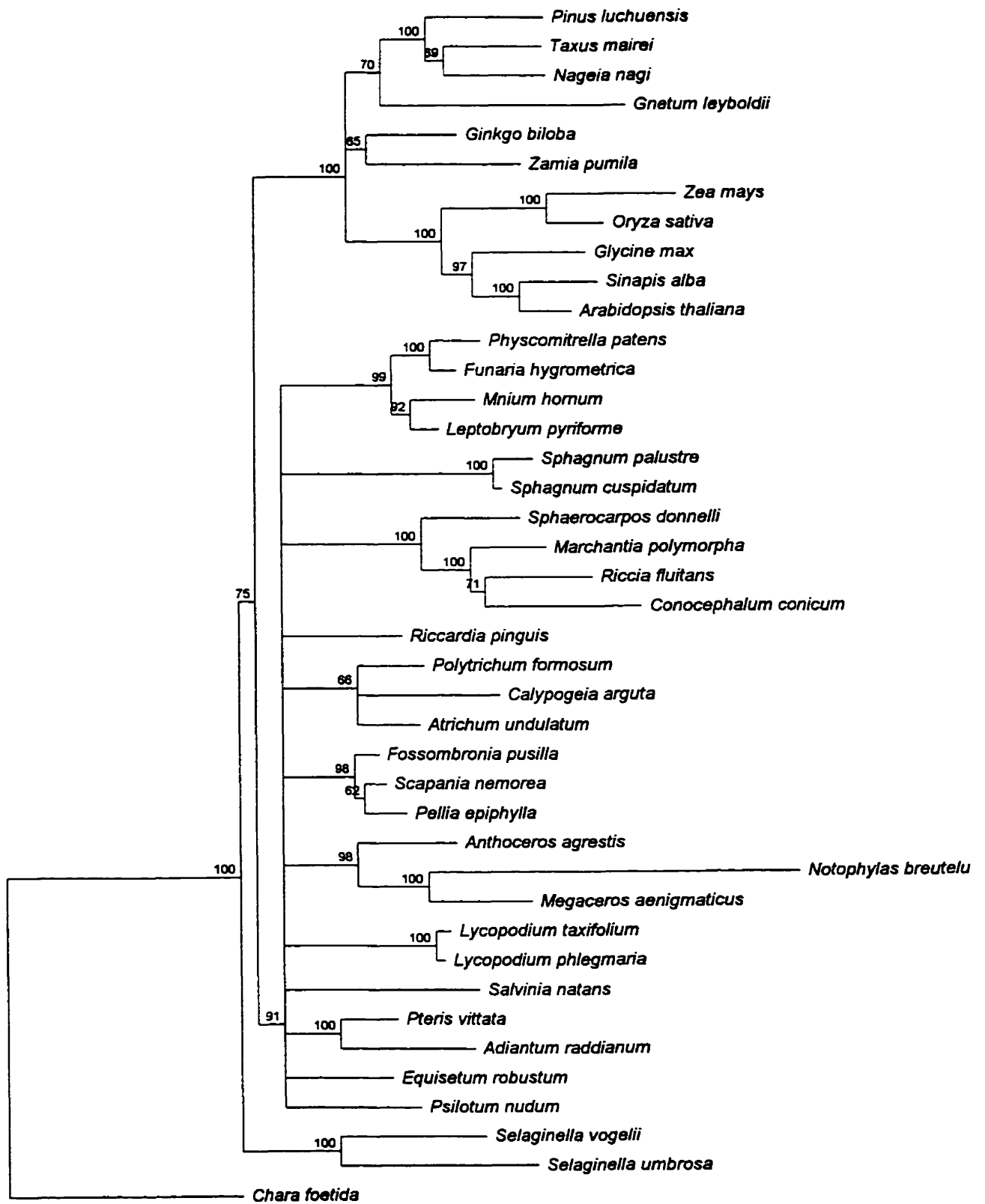


Figure A2SB Jukes & Cantor
 Insertions & Deletions taken into account
 50% Majority-Rule Bootstrap Tree

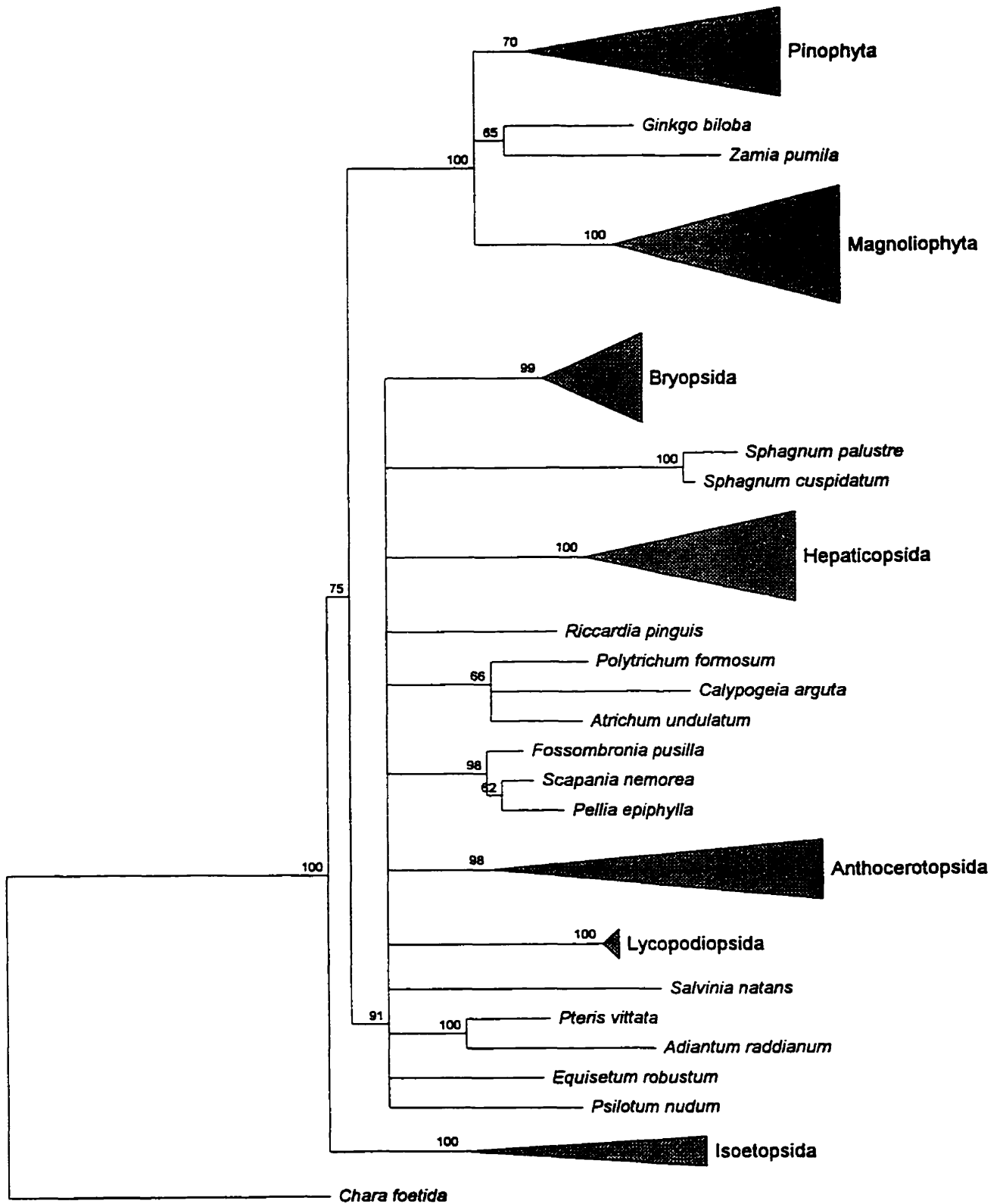


Figure A2SB Jukes & Cantor
 Insertions & Deletions taken into account
 50% Majority-Rule Bootstrap Tree

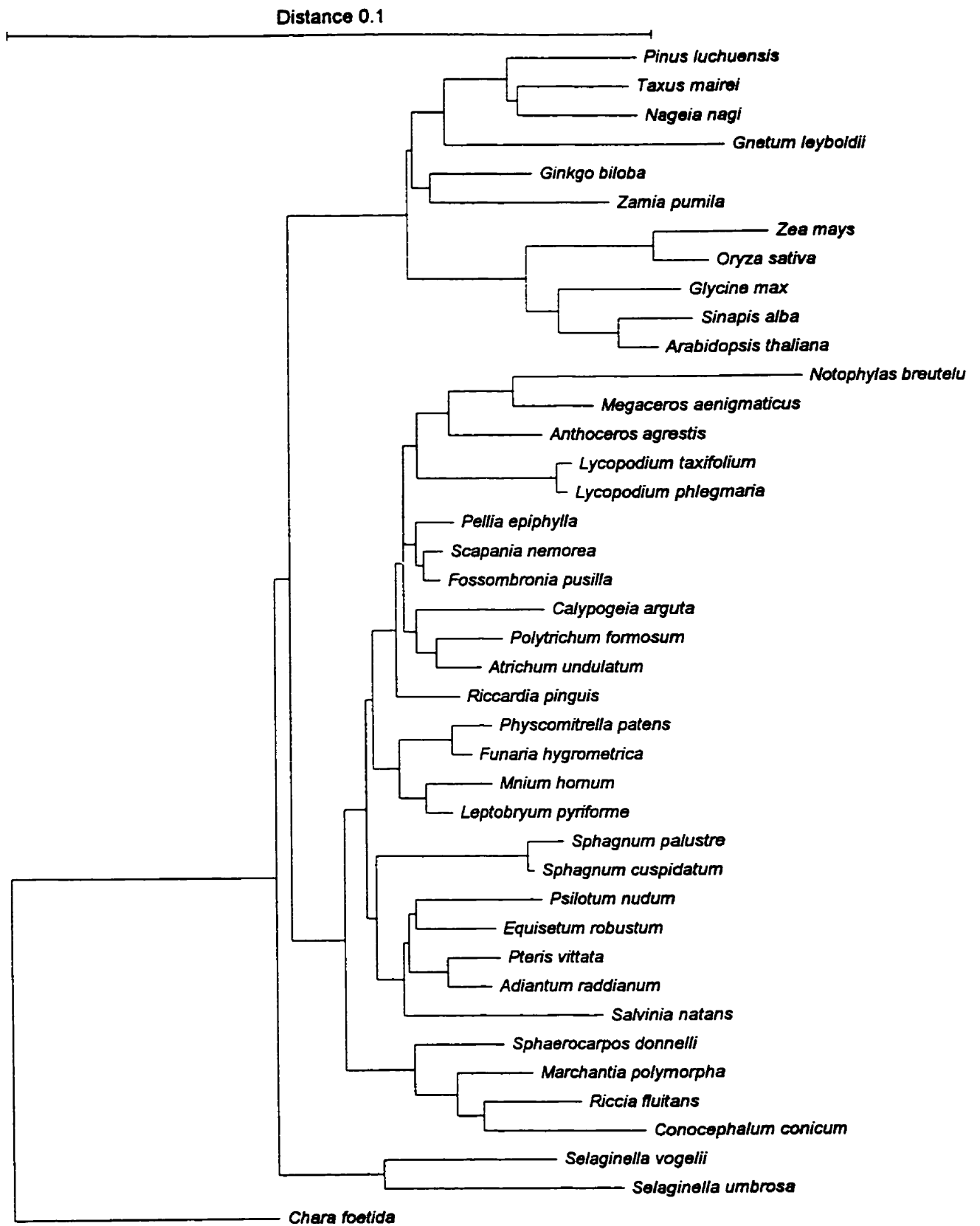


Figure B1S Tajima & Nei; Neighbor-joining

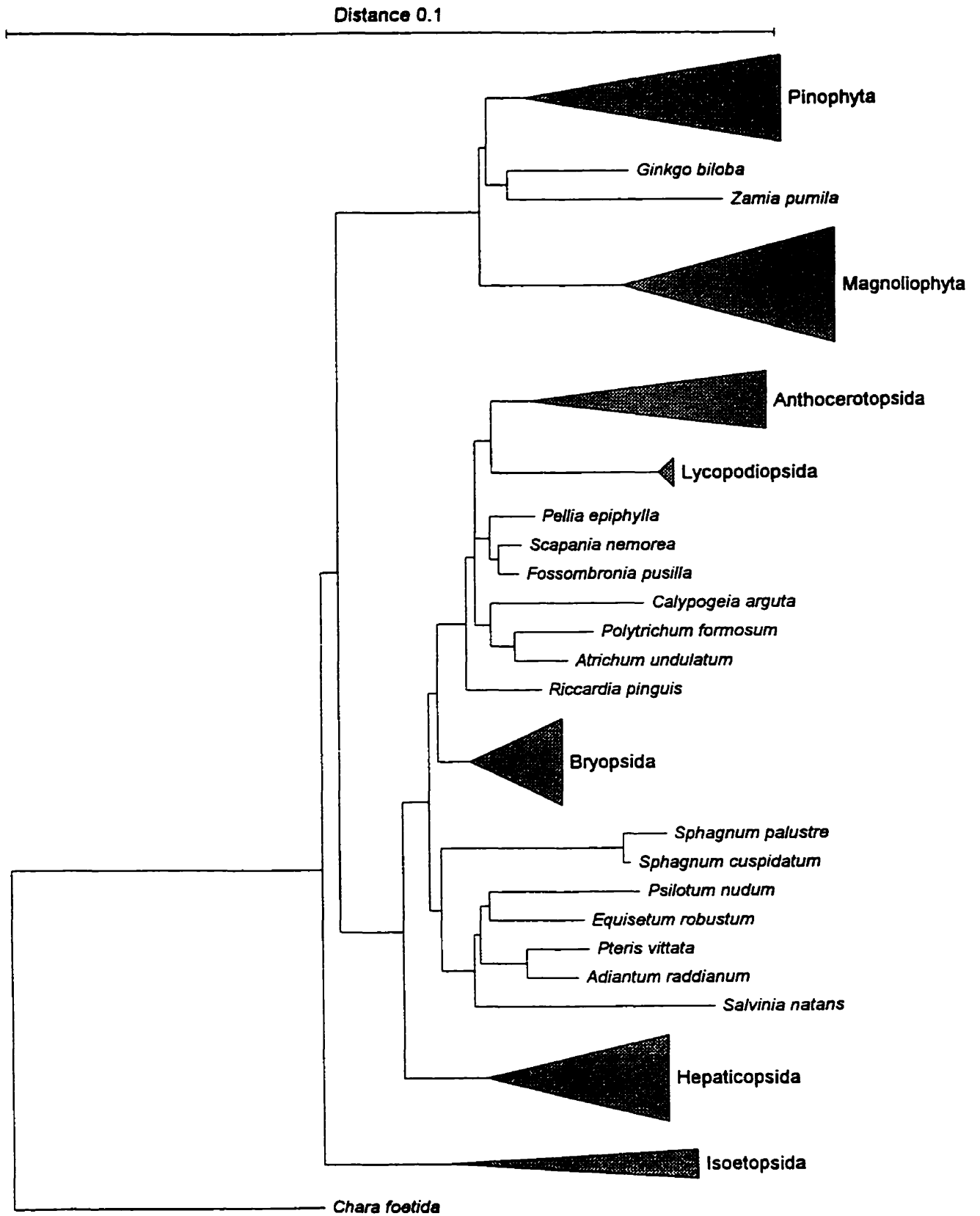


Figure B1S Tajima & Nei; Neighbor-joining

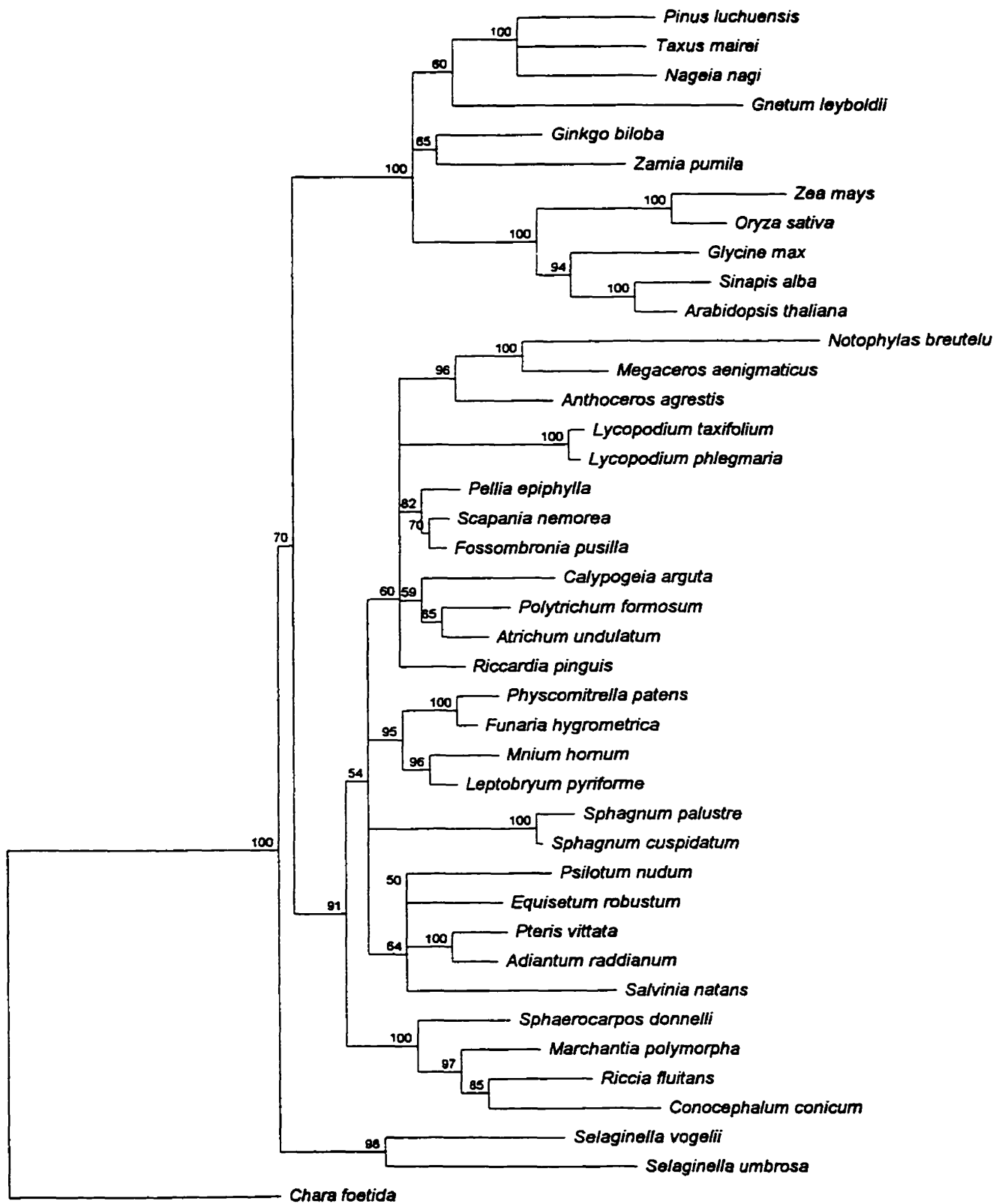


Figure B1SB Tajima & Nei; Neighbor-joining
 50% Majority-Rule Bootstrap Tree

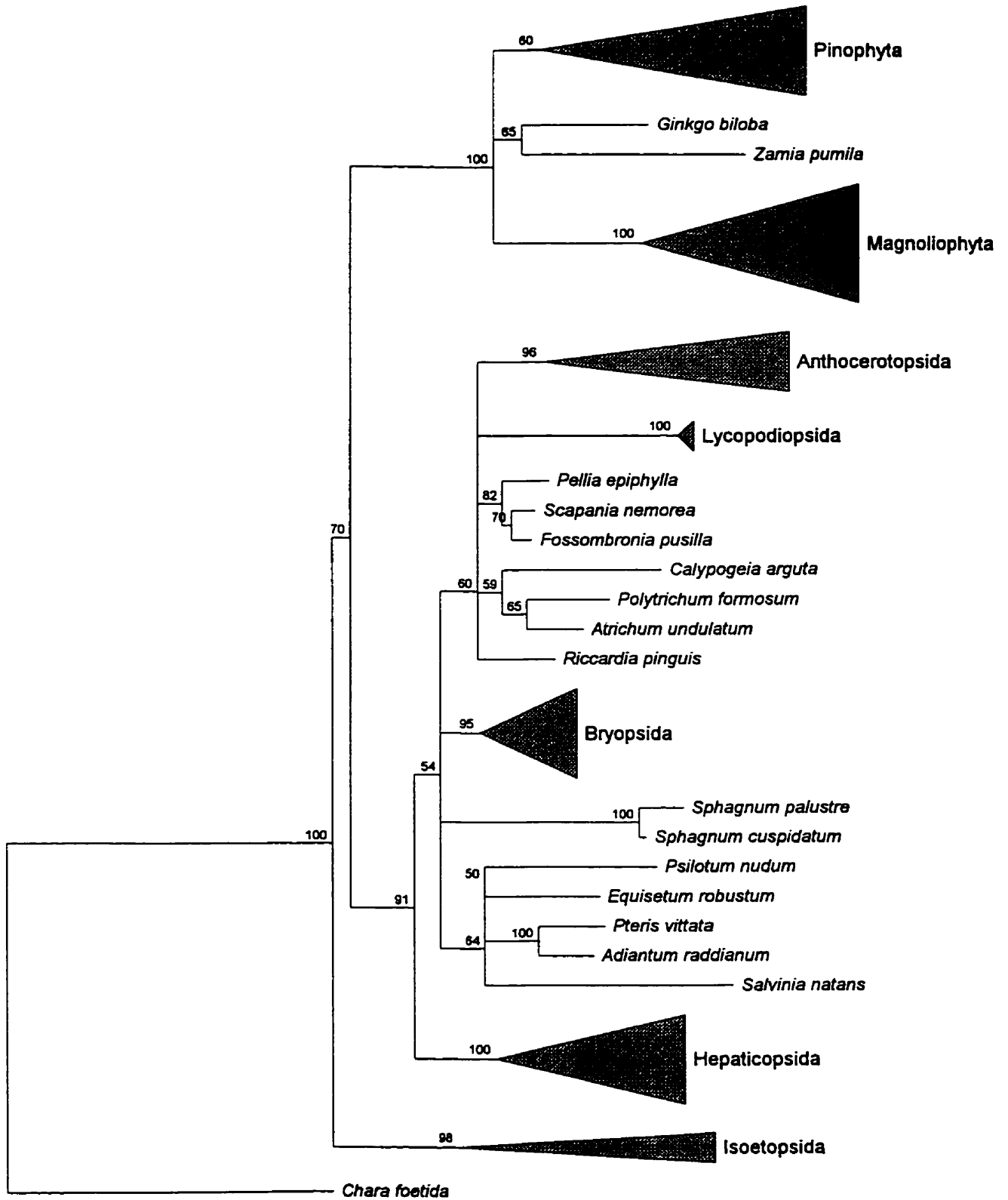


Figure B1SB Tajima & Nei; Neighbor-joining
50% Majority-Rule Bootstrap Tree

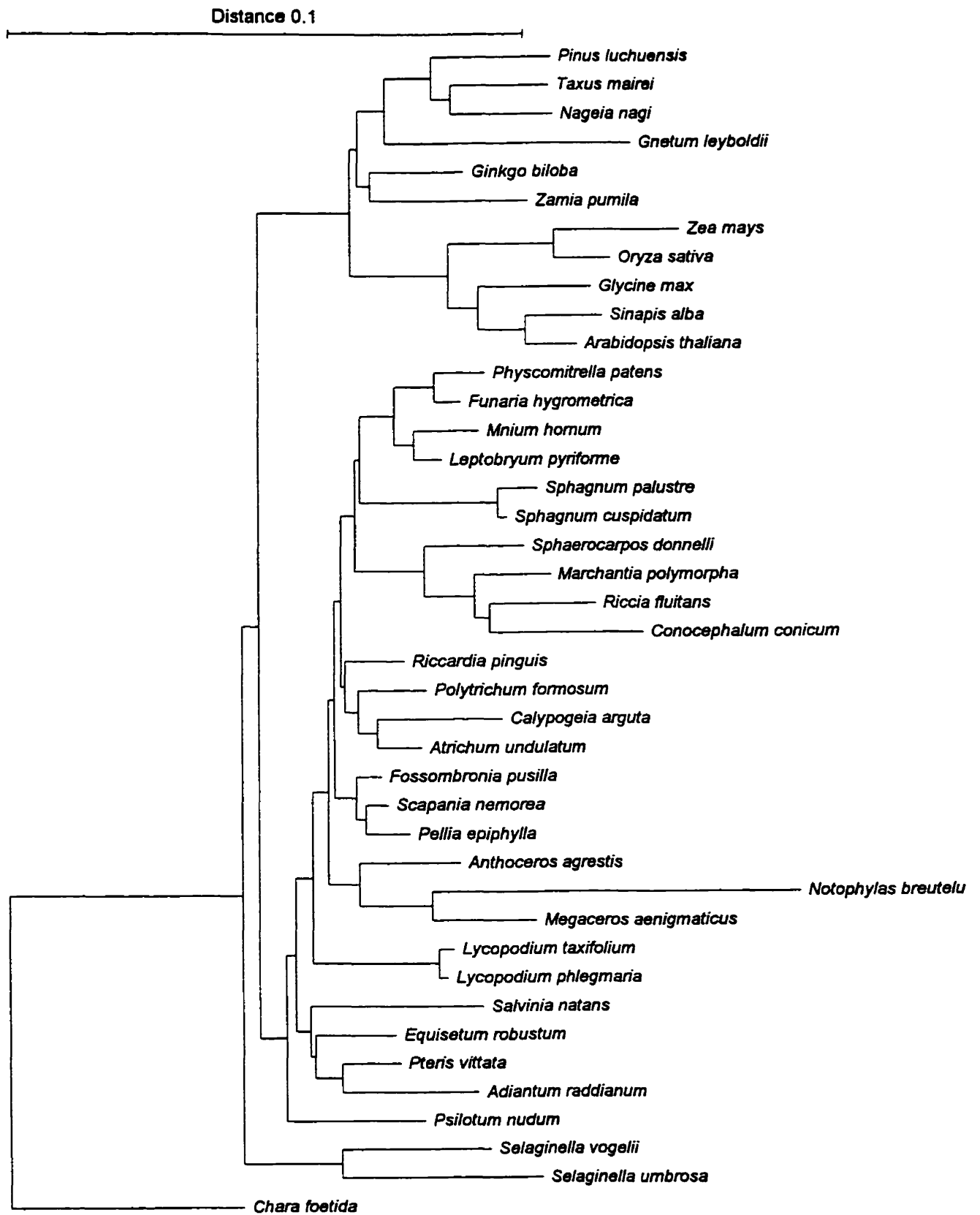


Figure B2S Tajima & Nei
Insertions & Deletions taken into account

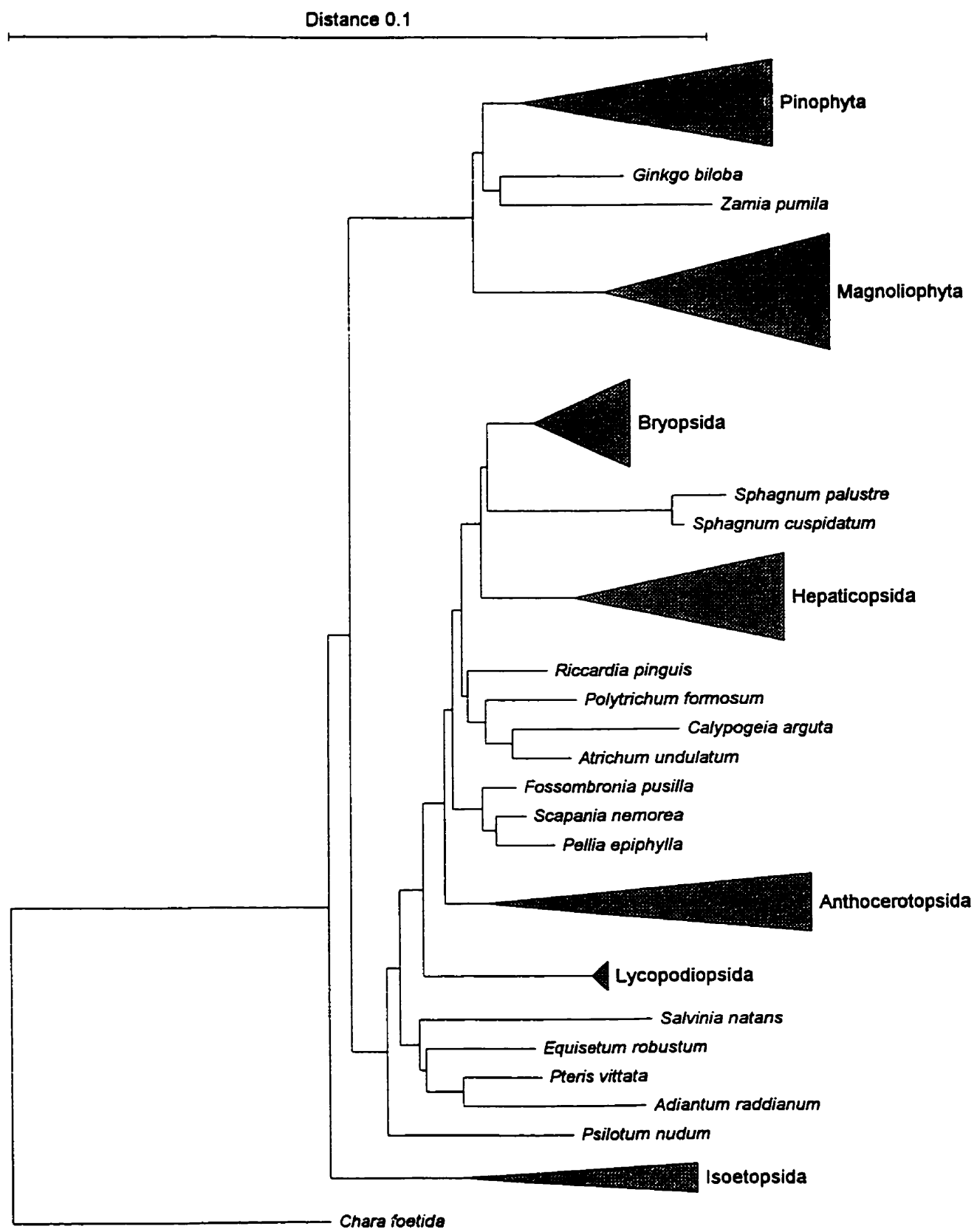


Figure B2S Tajima & Nei
Insertions & Deletions taken into account

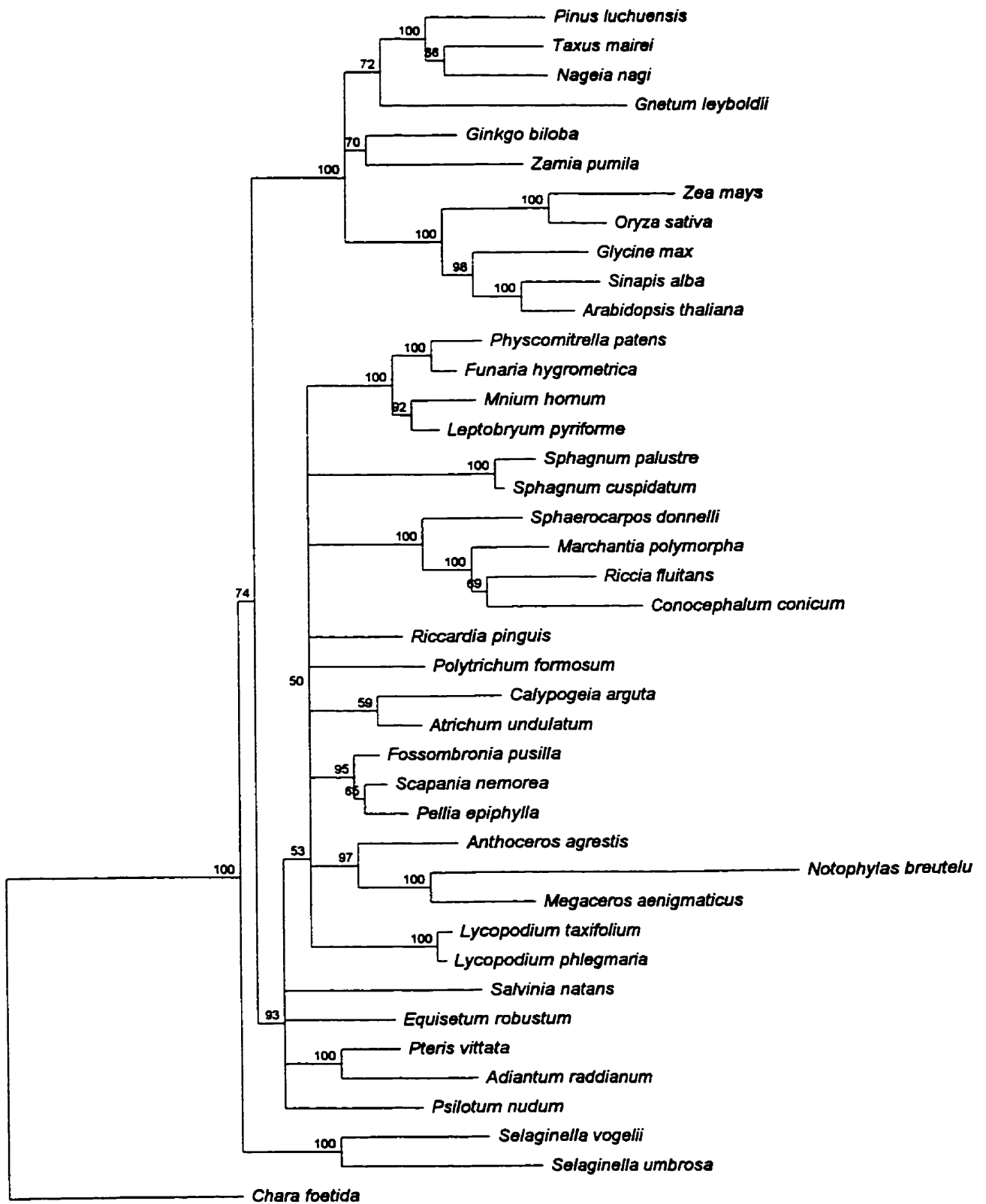


Figure B2SB Tajima & Nei
 Insertions & Deletions taken into account
 50% Majority-Rule Bootstrap Tree

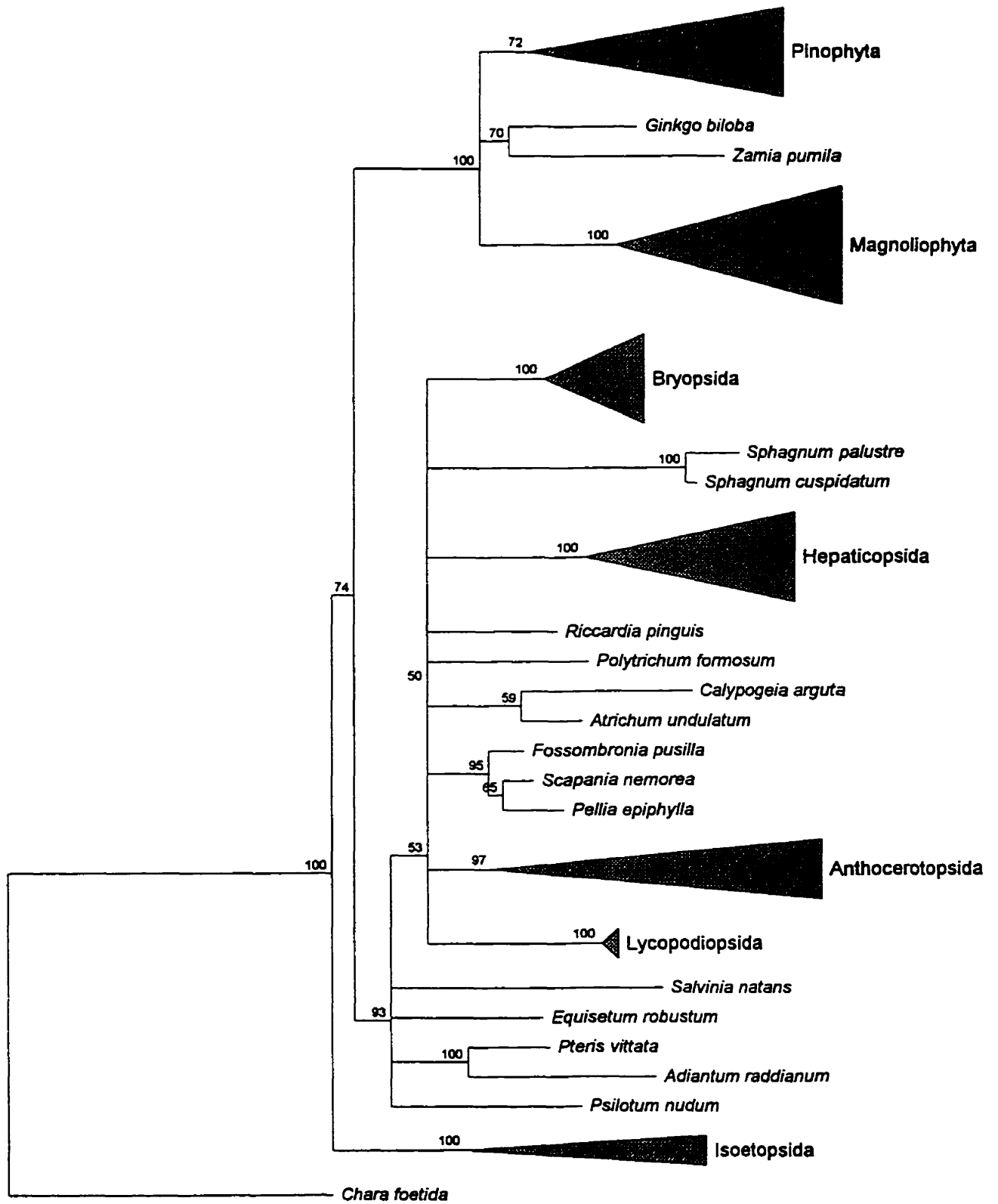


Figure B2SB Tajima & Nei
 Insertions & Deletions taken into account
 50% Majority-Rule Bootstrap Tree



Figure C1S Kimura; Neighbor-joining

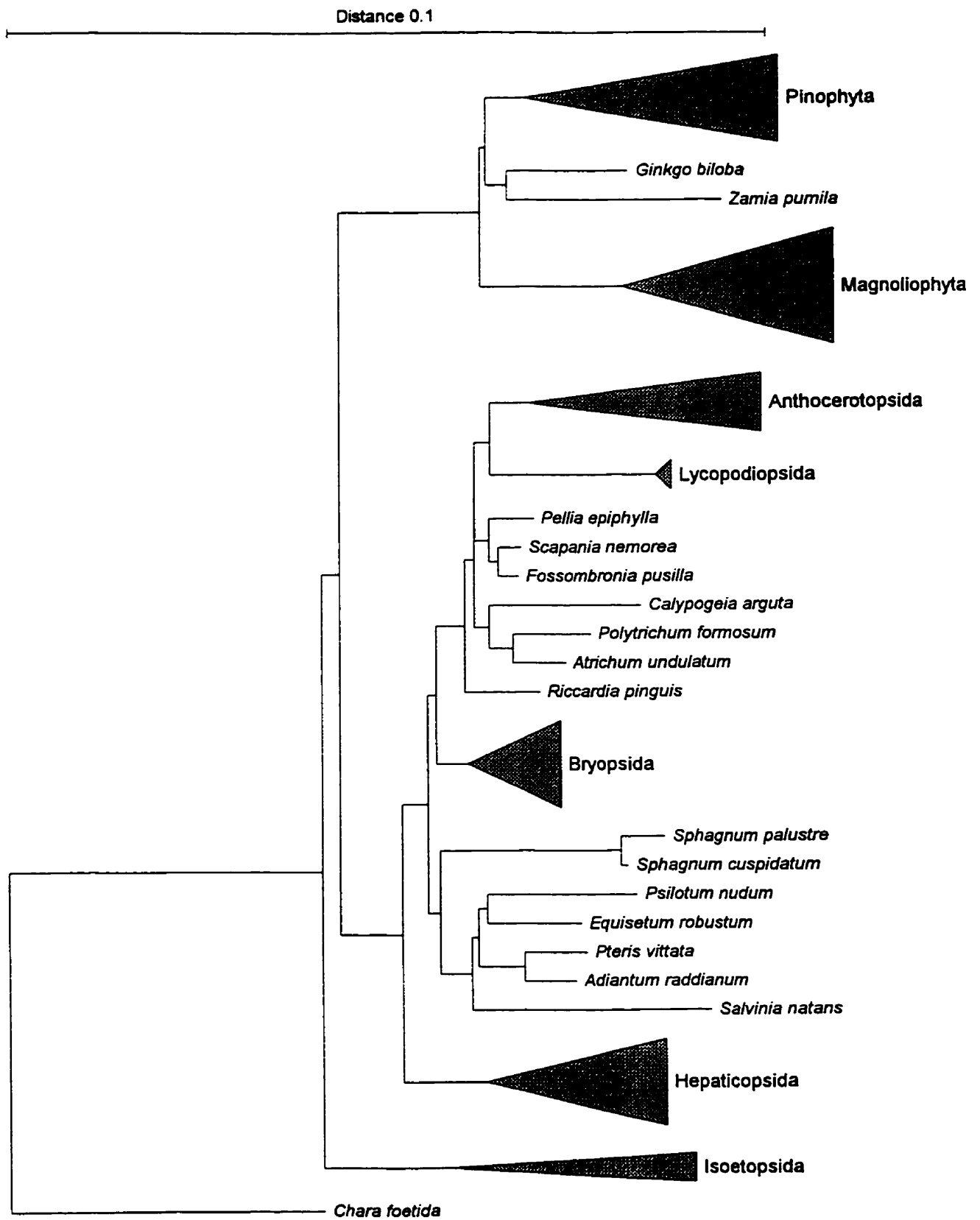


Figure C1S Kimura; Neighbor-joining

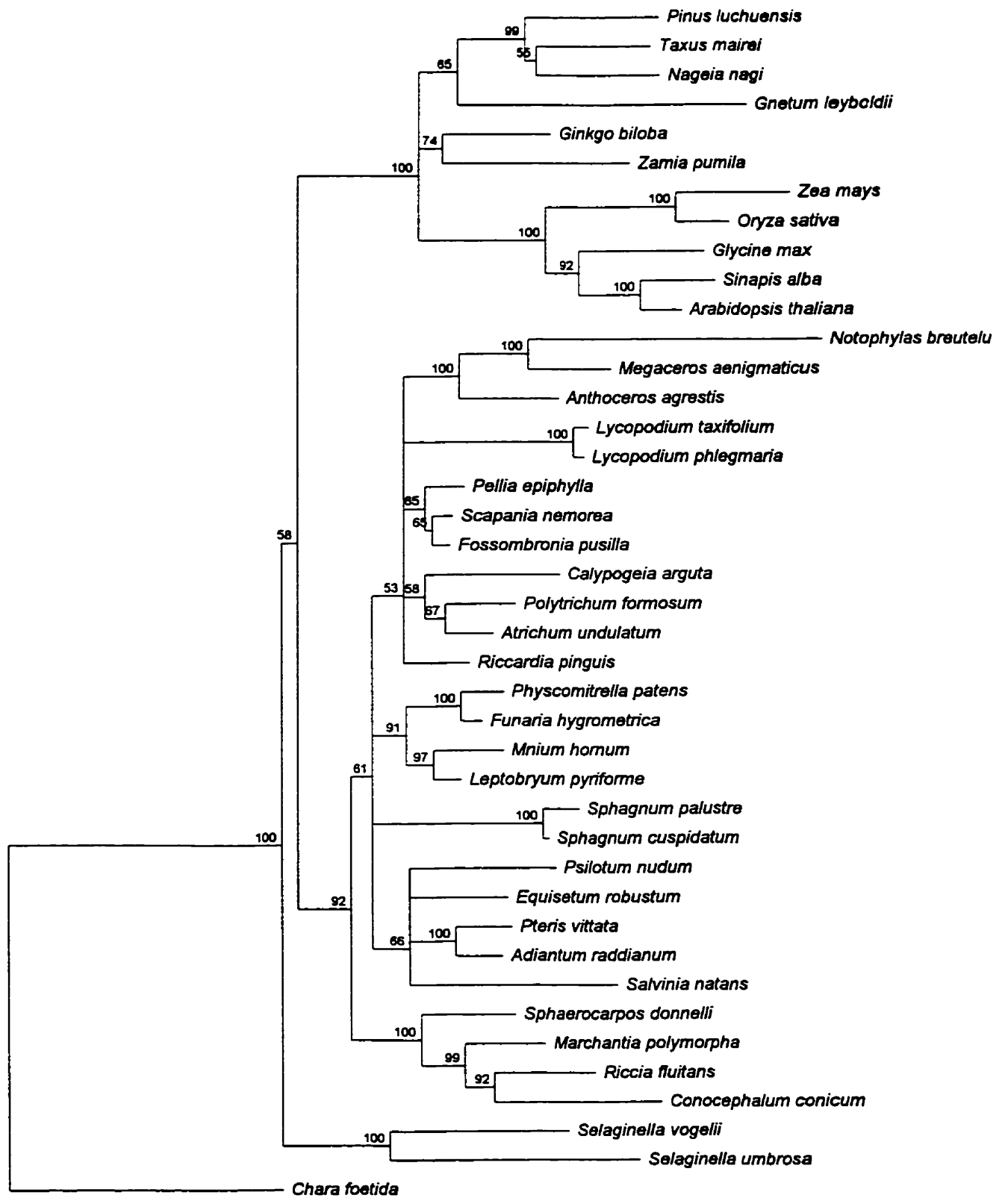


Figure C1SB Kimura; Neighbor-joining
50% Majority-Rule Bootstrap Tree

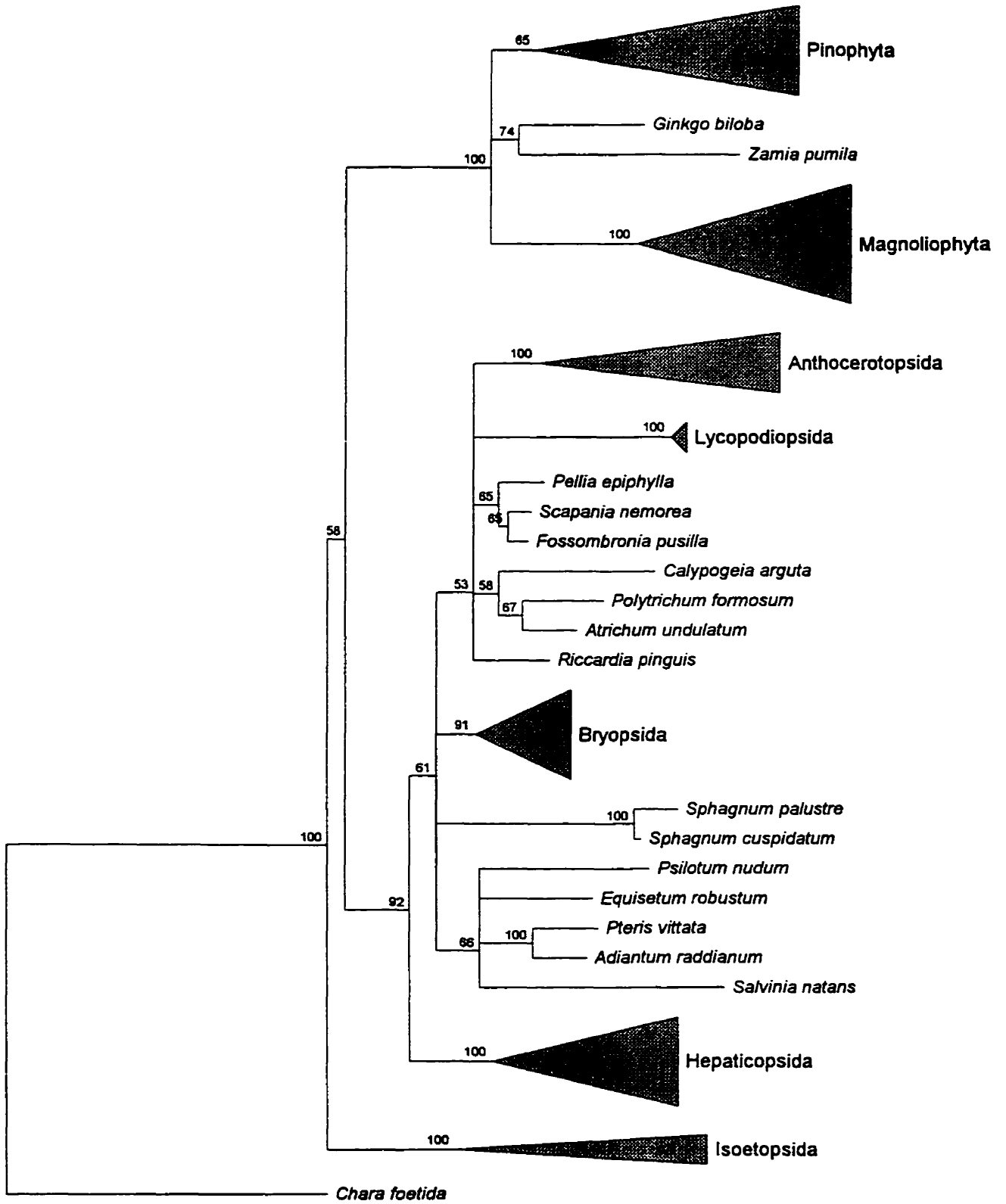


Figure C1SB Kimura; Neighbor-joining
50% Majority-Rule Bootstrap Tree

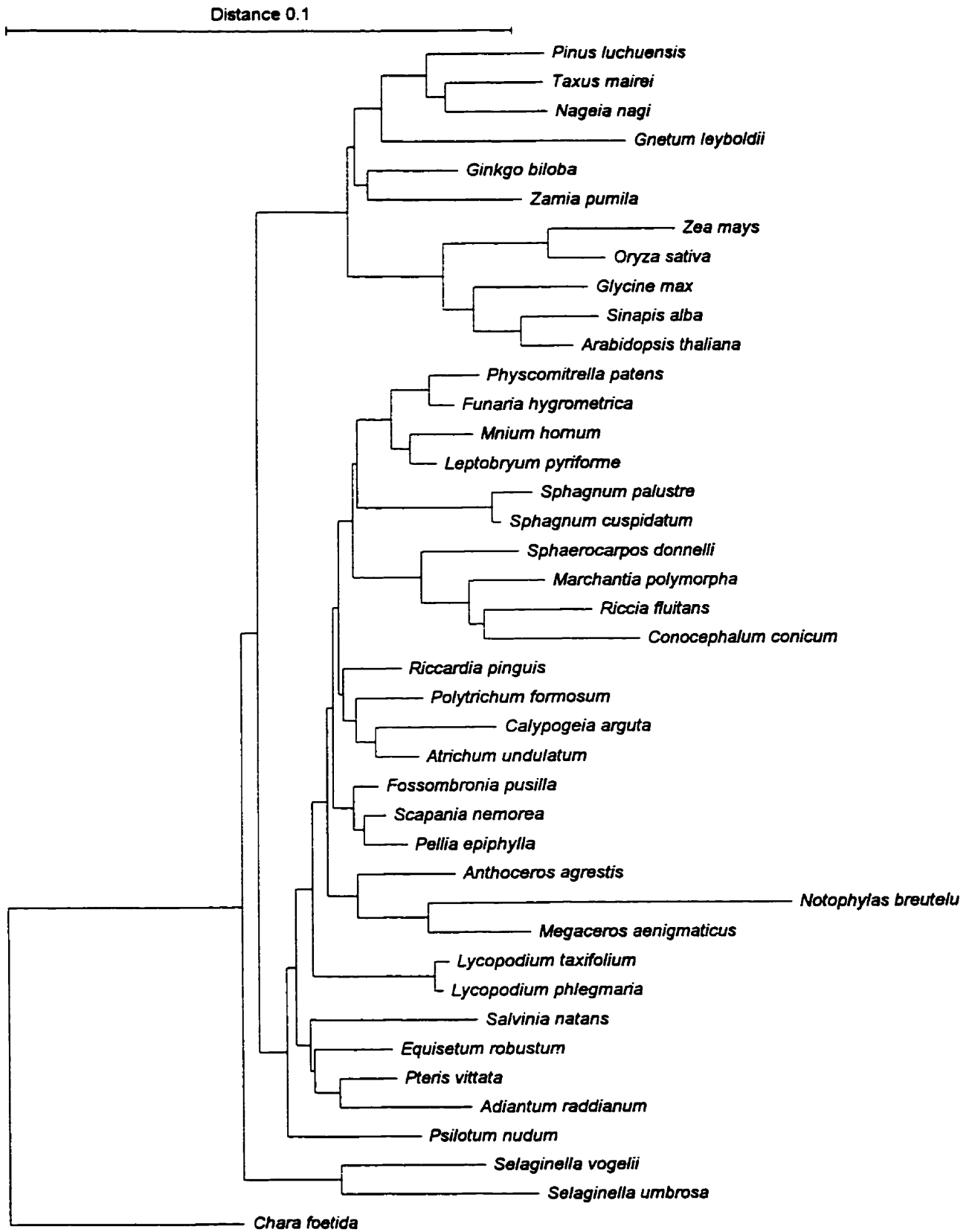


Figure C2S Kimura; (Insertions & Deletions taken into account)

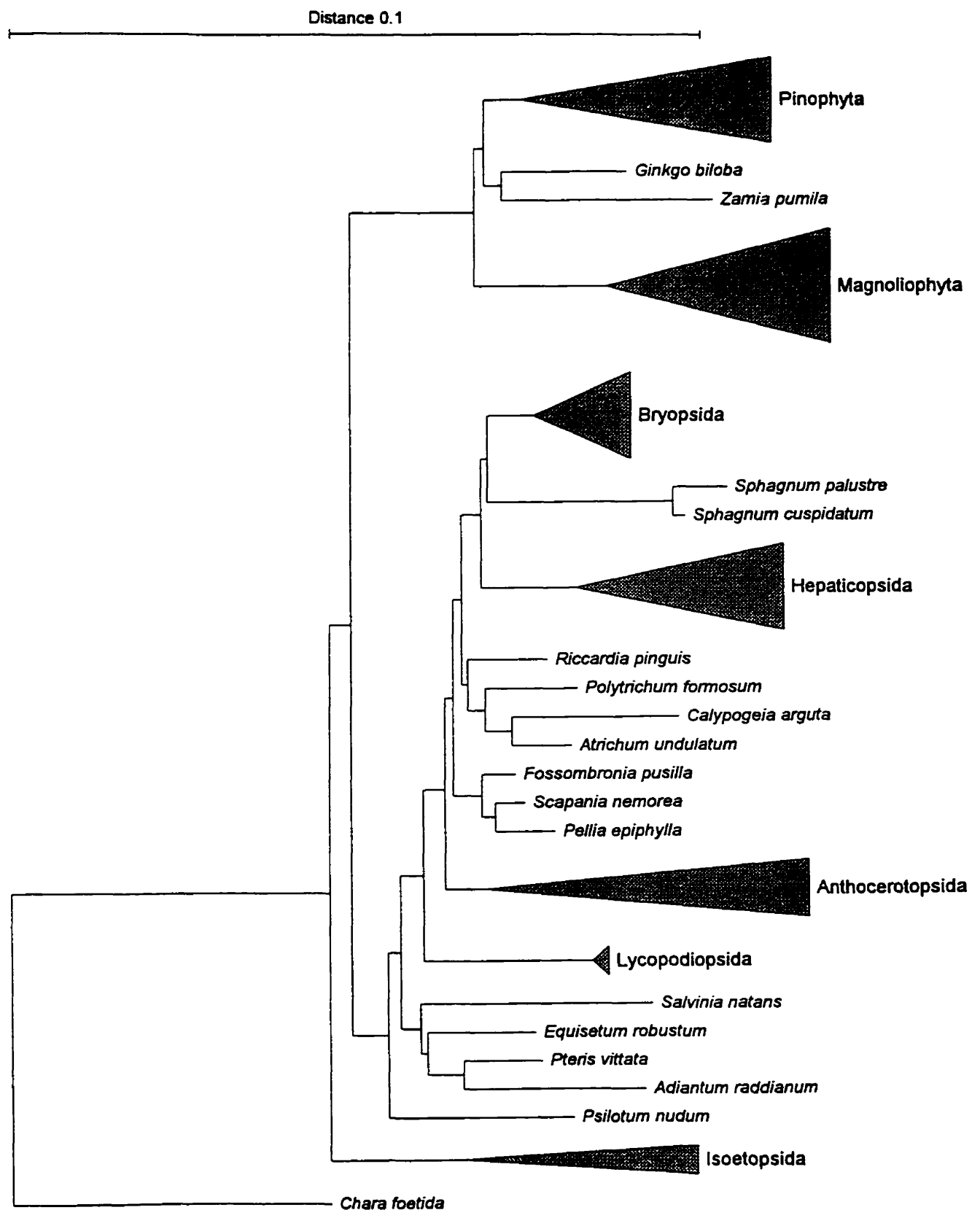


Figure C2S Kimura; (Insertions & Deletions taken into account)

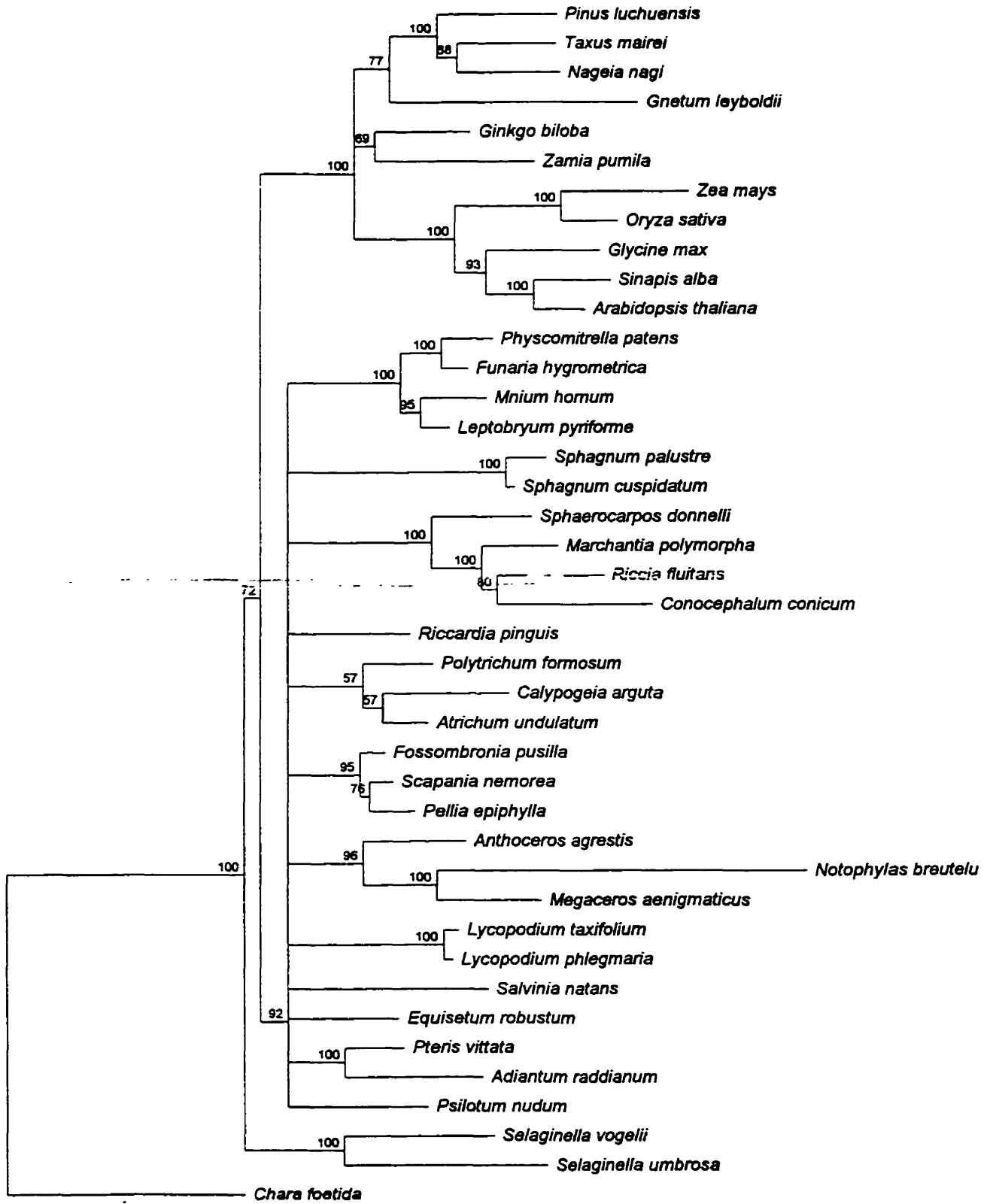


Figure C2SB Kimura; (Insertions & Deletions taken into account)
 50% Majority-Rule Bootstrap Tree

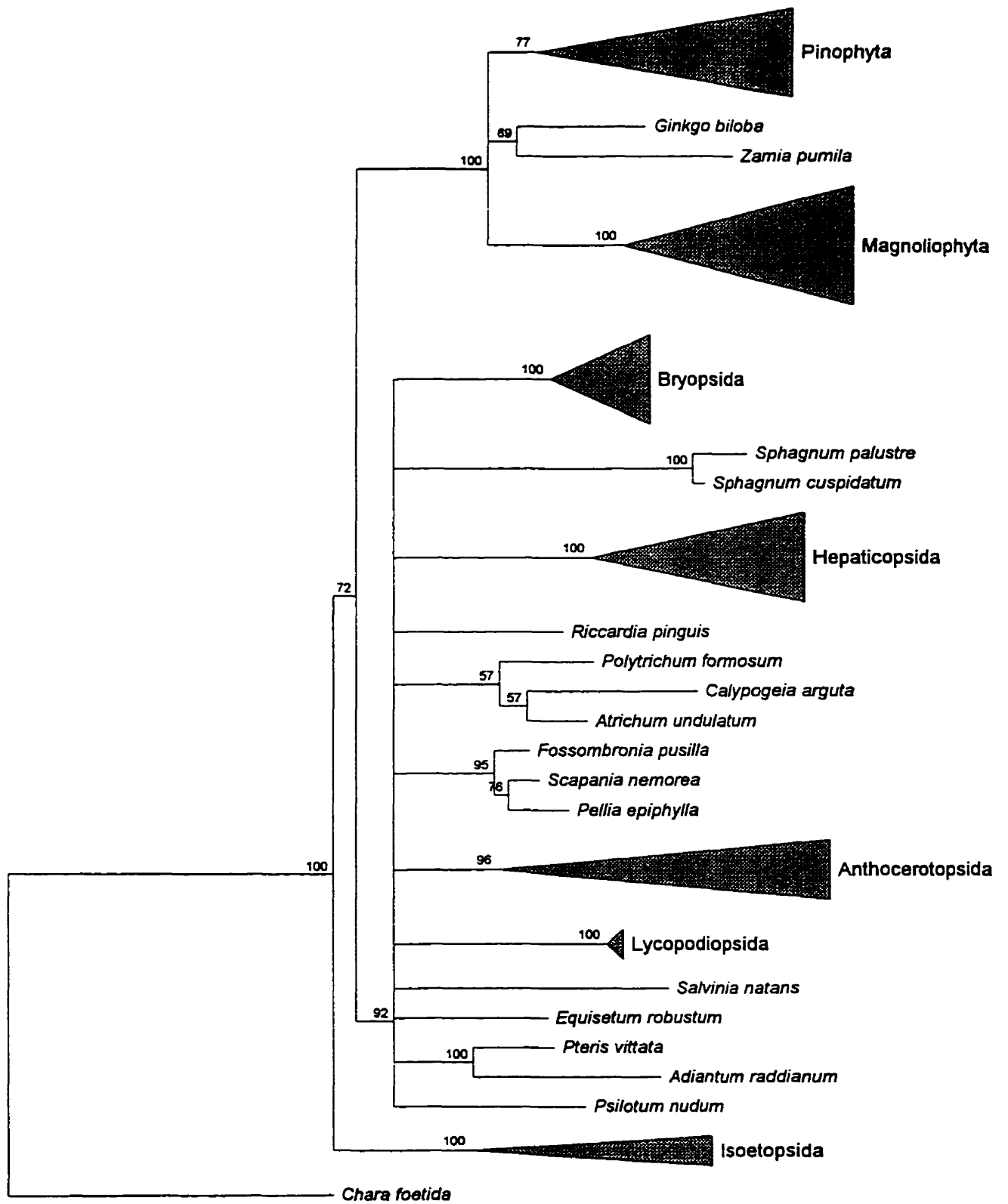


Figure C2SB Kimura; (Insertions & Deletions taken into account)
 50% Majority-Rule Bootstrap Tree

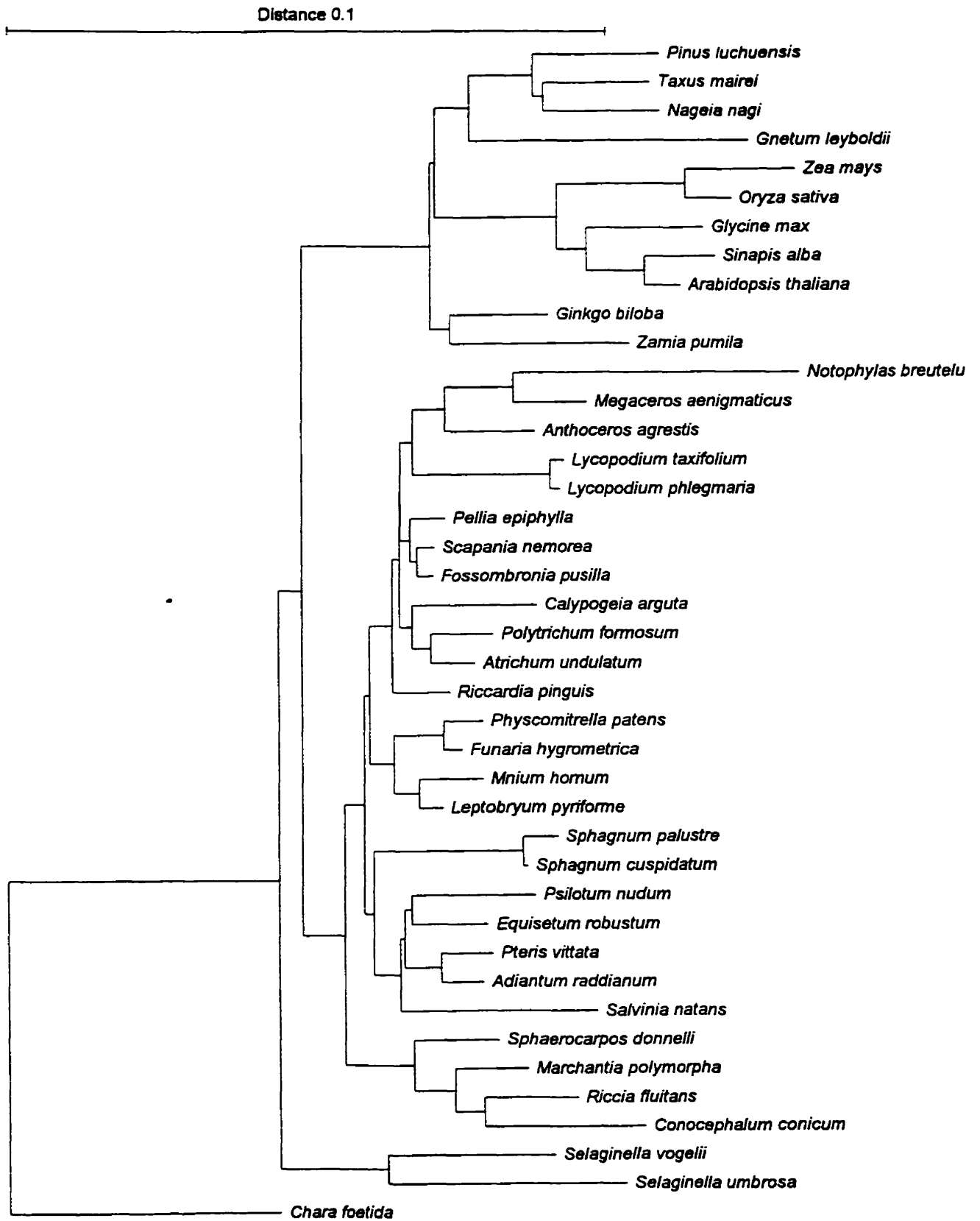


Figure D1.1S Jin & Nei (Distance Calculation); Jukes & Cantor (Evolution Model), Gamma distance = 1

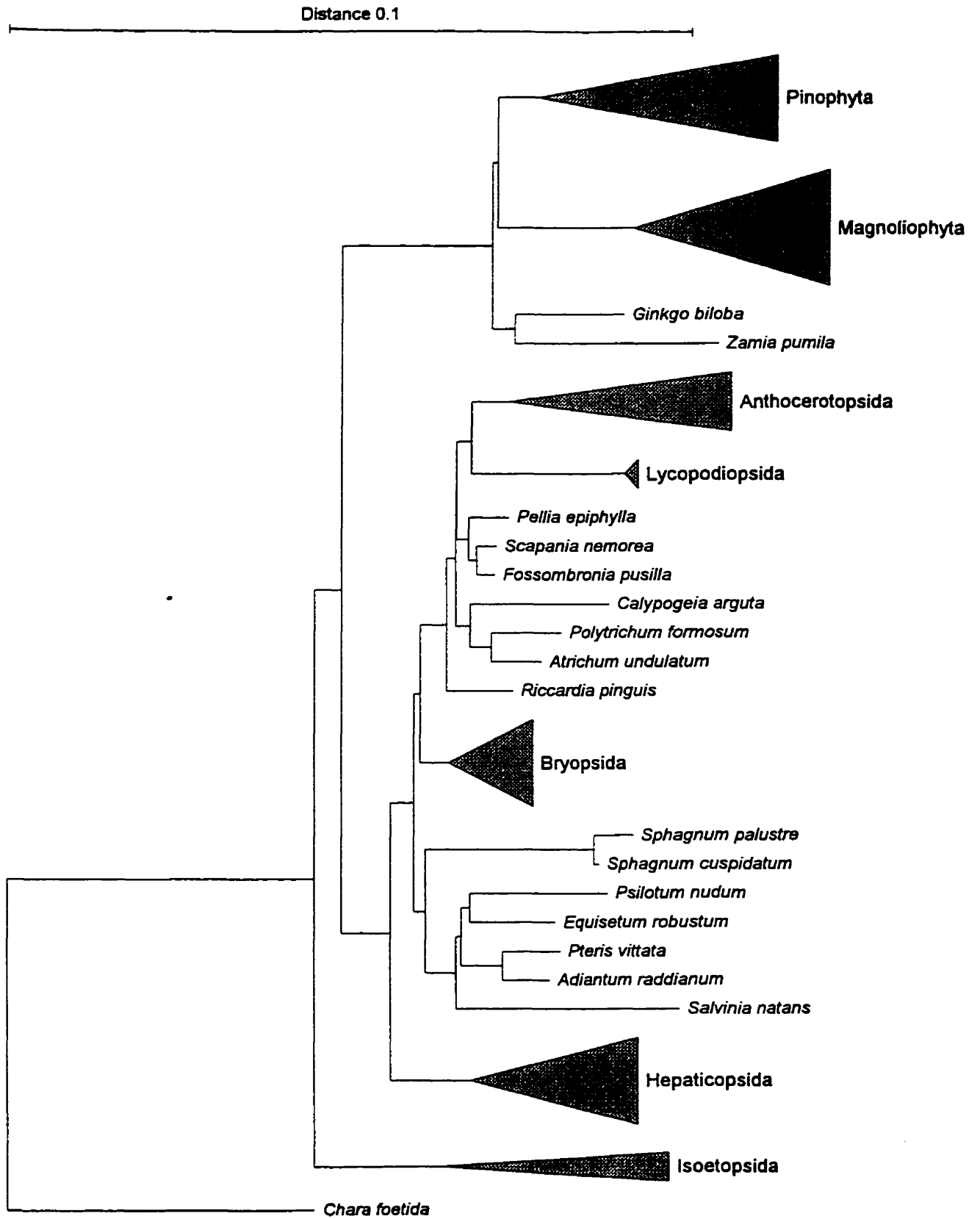


Figure D1.1S Jin & Nei (Distance Calculation); Jukes & Cantor (Evolution Model), Gamma distance = 1

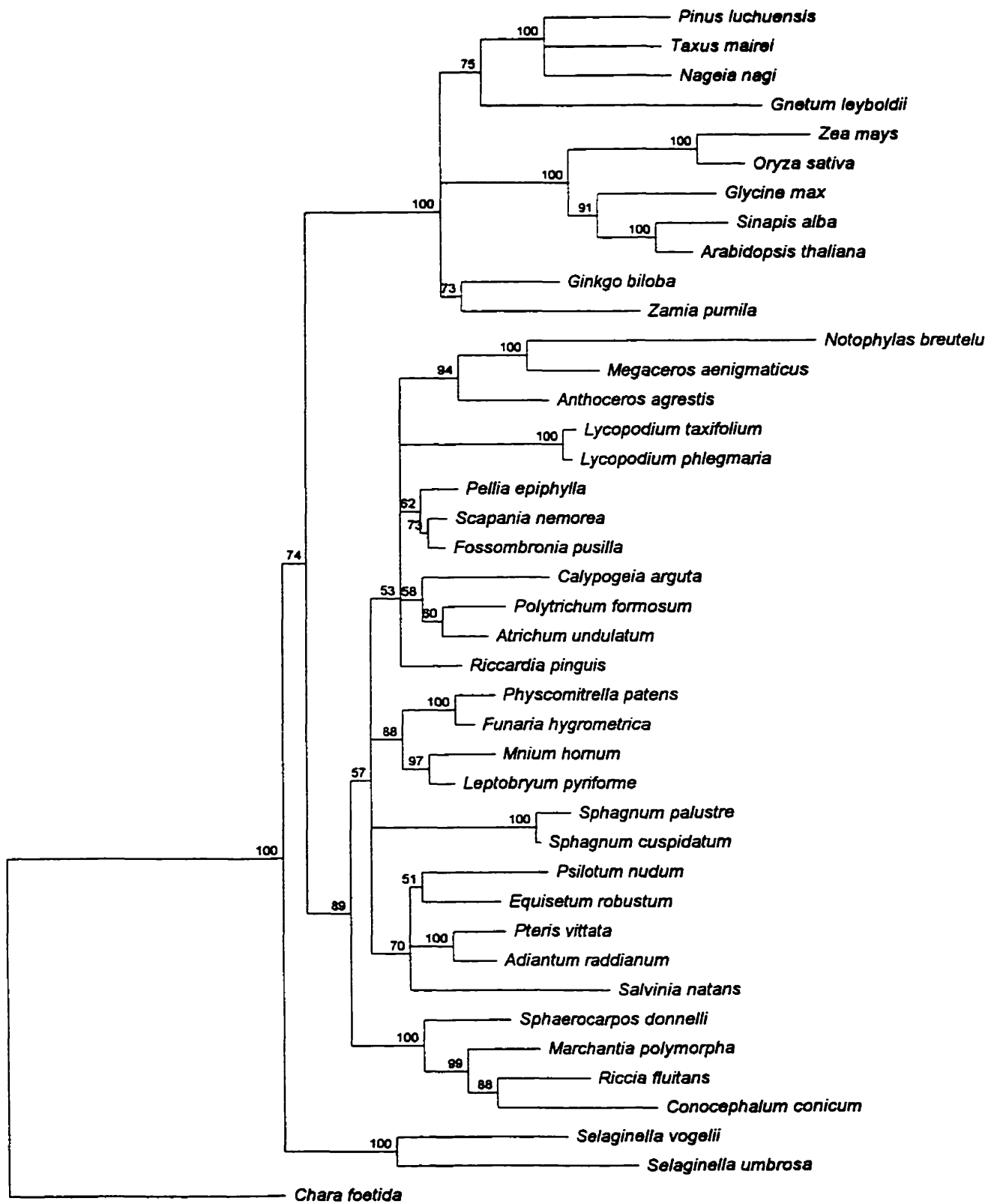


Figure D1.1SB Jin & Nei (Distance Calculation); Jukes & Cantor (Evolution Model), Gamma distance = 1
50% Majority-Rule Bootstrap Tree

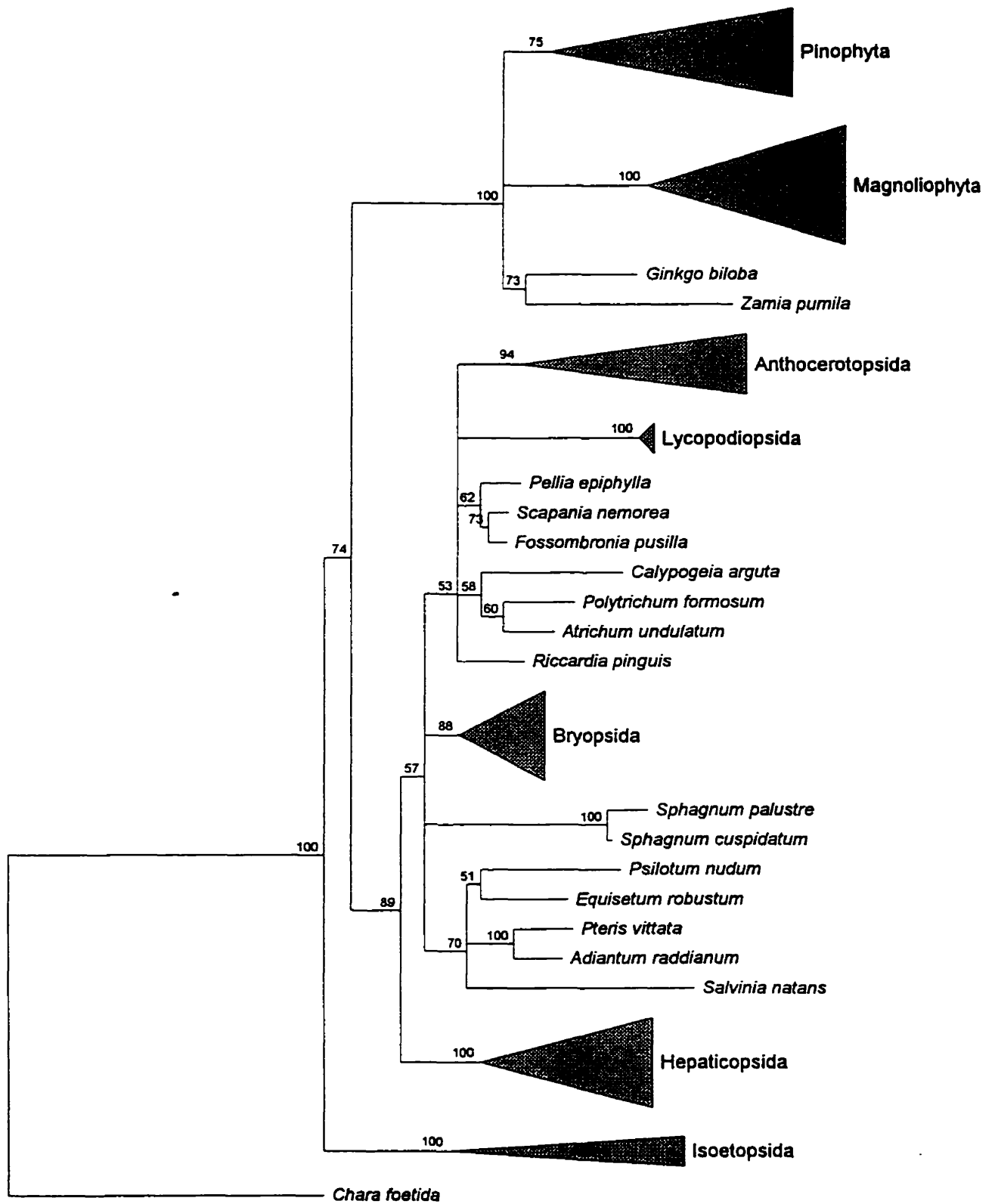


Figure D1.1SB Jin & Nei (Distance Calculation); Jukes & Cantor (Evolution Model), Gamma distance = 1
50% Majority-Rule Bootstrap Tree

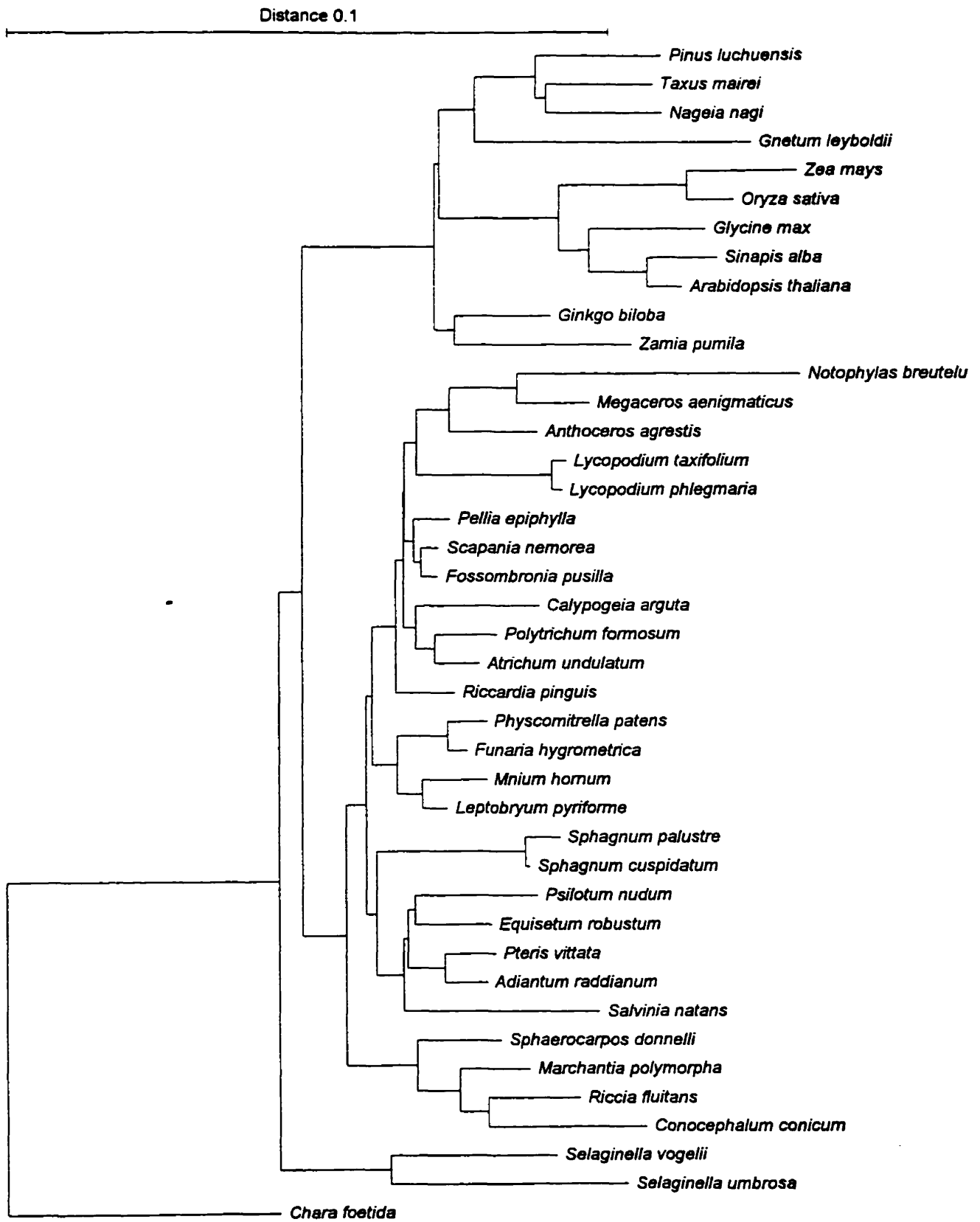


Figure D1.2S Jin & Nei (Distance Calculation); Jukes & Cantor (Evolution Model), Gamma distance = 1
Insertions & Deletions taken into account

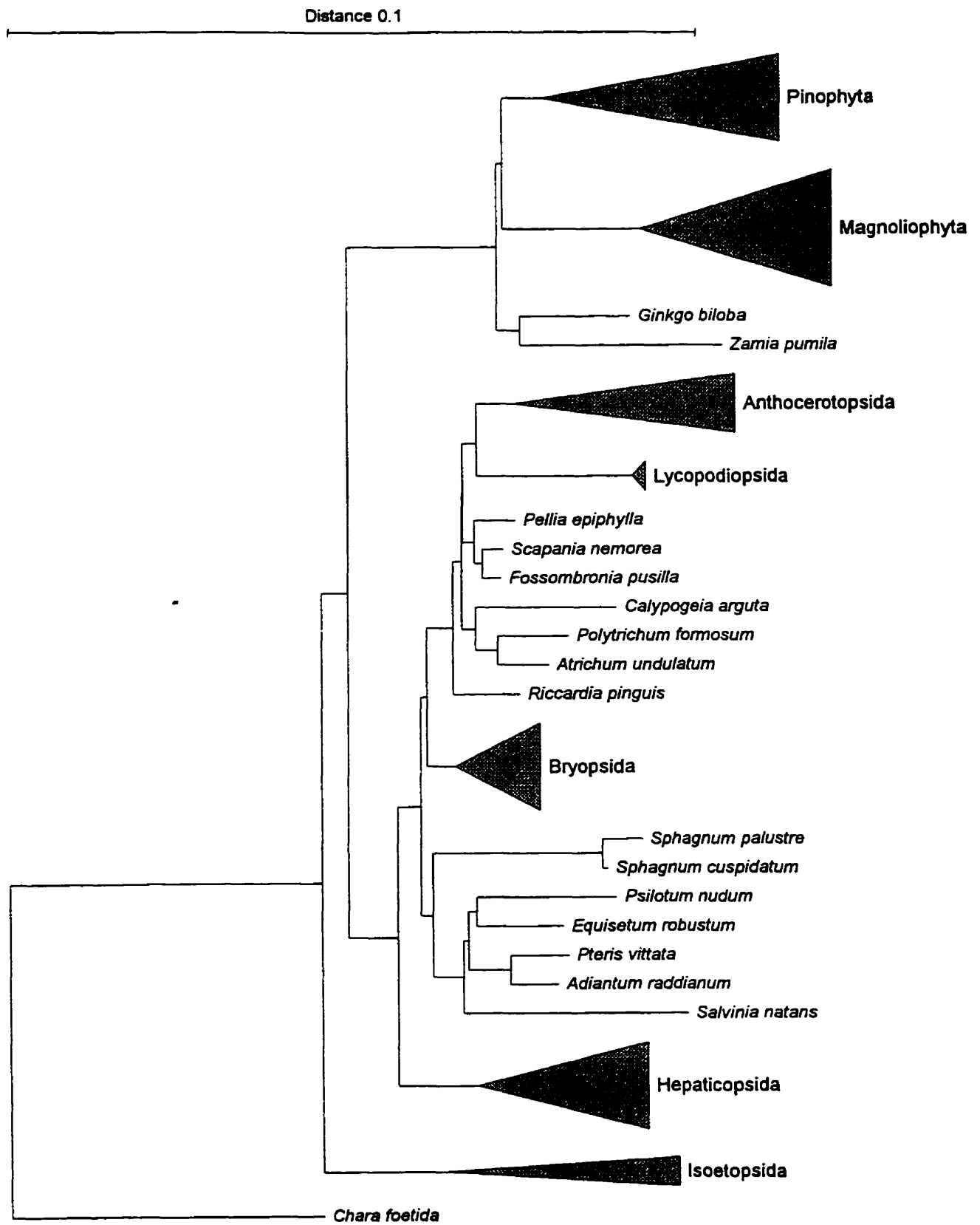


Figure D1.2S Jin & Nei (Distance Calculation); Jukes & Cantor (Evolution Model), Gamma distance = 1
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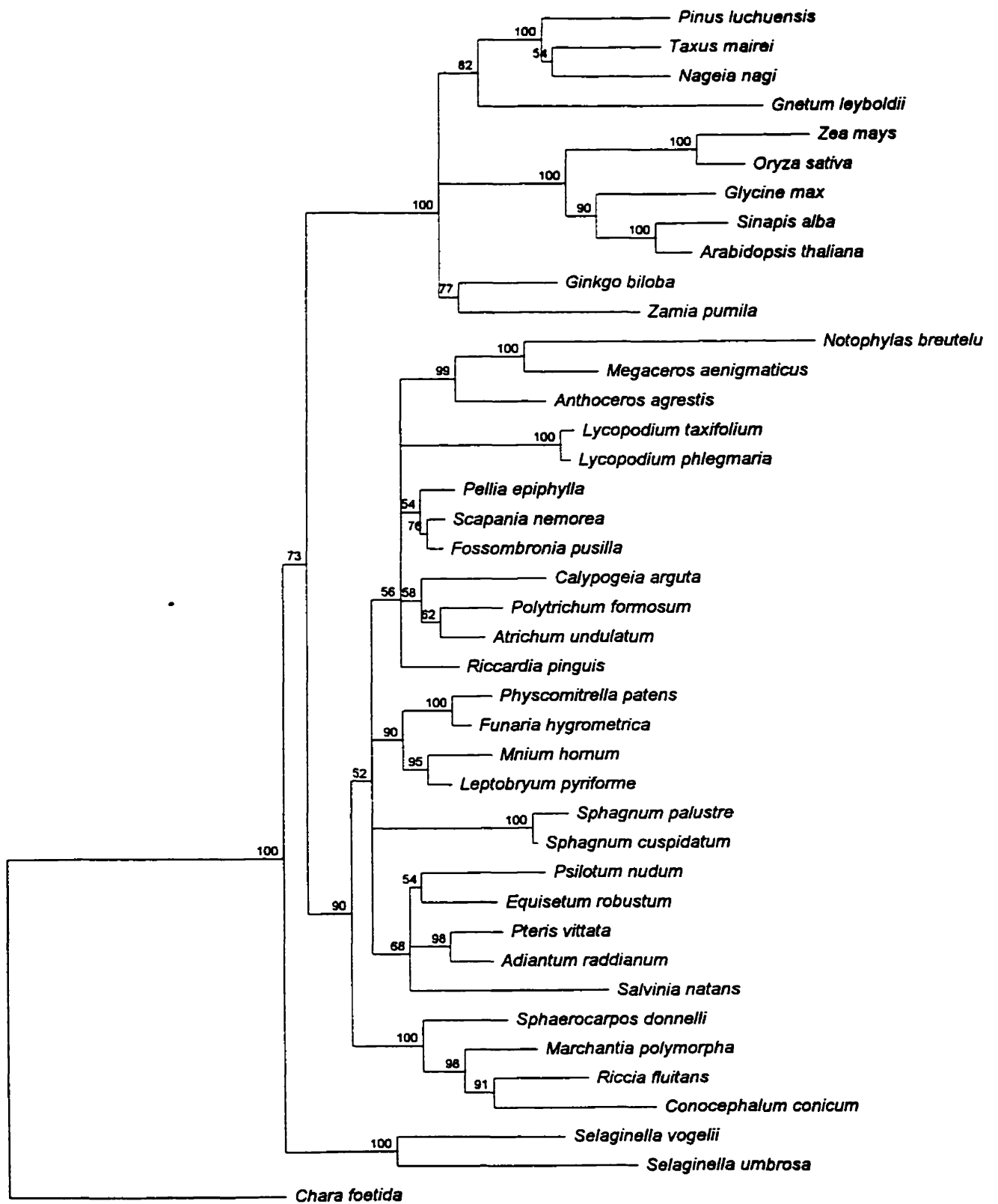


Figure D1.2SB Jin & Nei (Distance Calculation); Jukes & Cantor (Evolution Model), Gamma distance = 1
 Insertions & Deletions taken into account
 50% Majority-Rule Bootstrap Tree

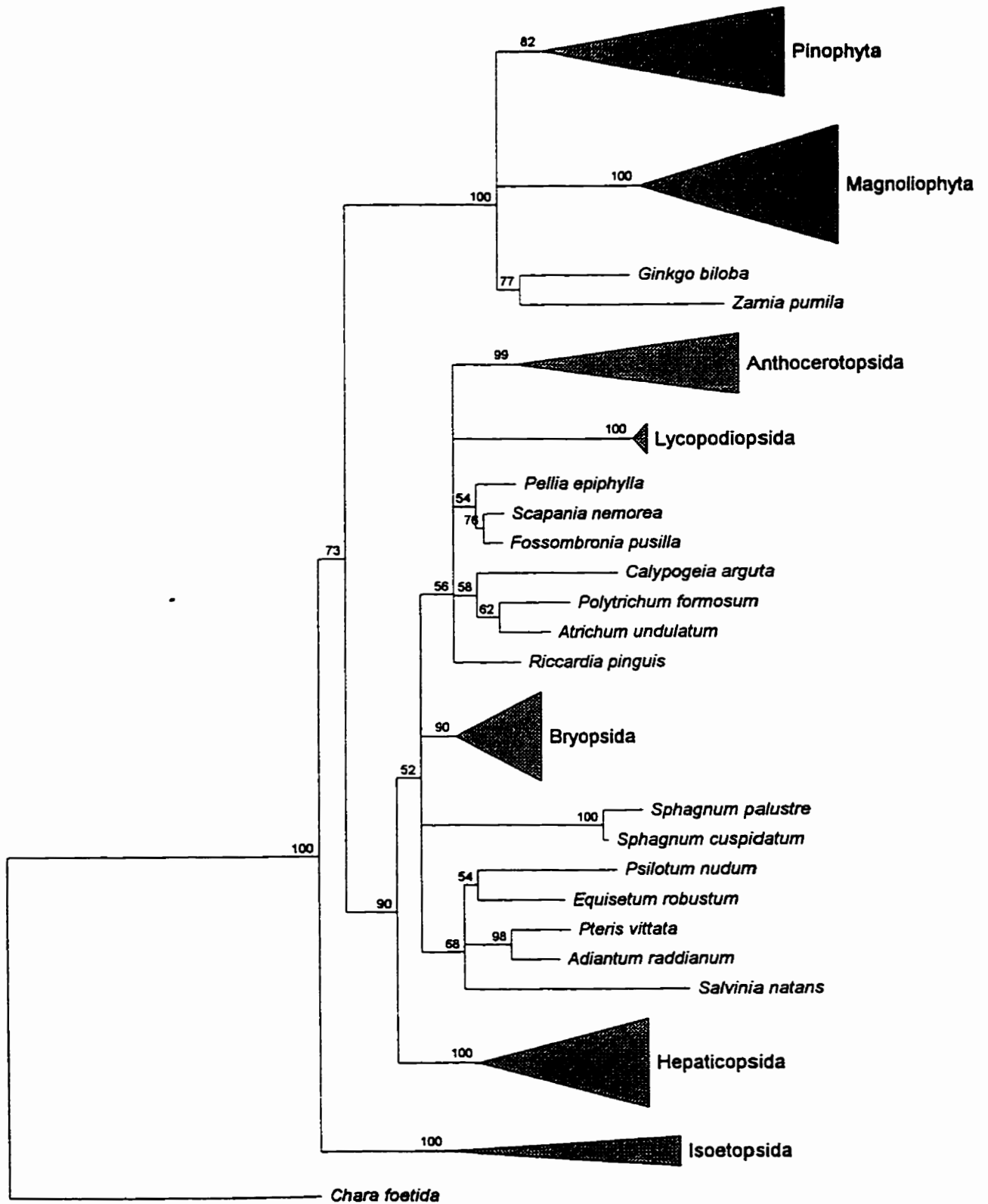


Figure D1.2SB Jin & Nei (Distance Calculation); Jukes & Cantor (Evolution Model), Gamma distance = 1
 Insertions & Deletions taken into account
 50% Majority-Rule Bootstrap Tree

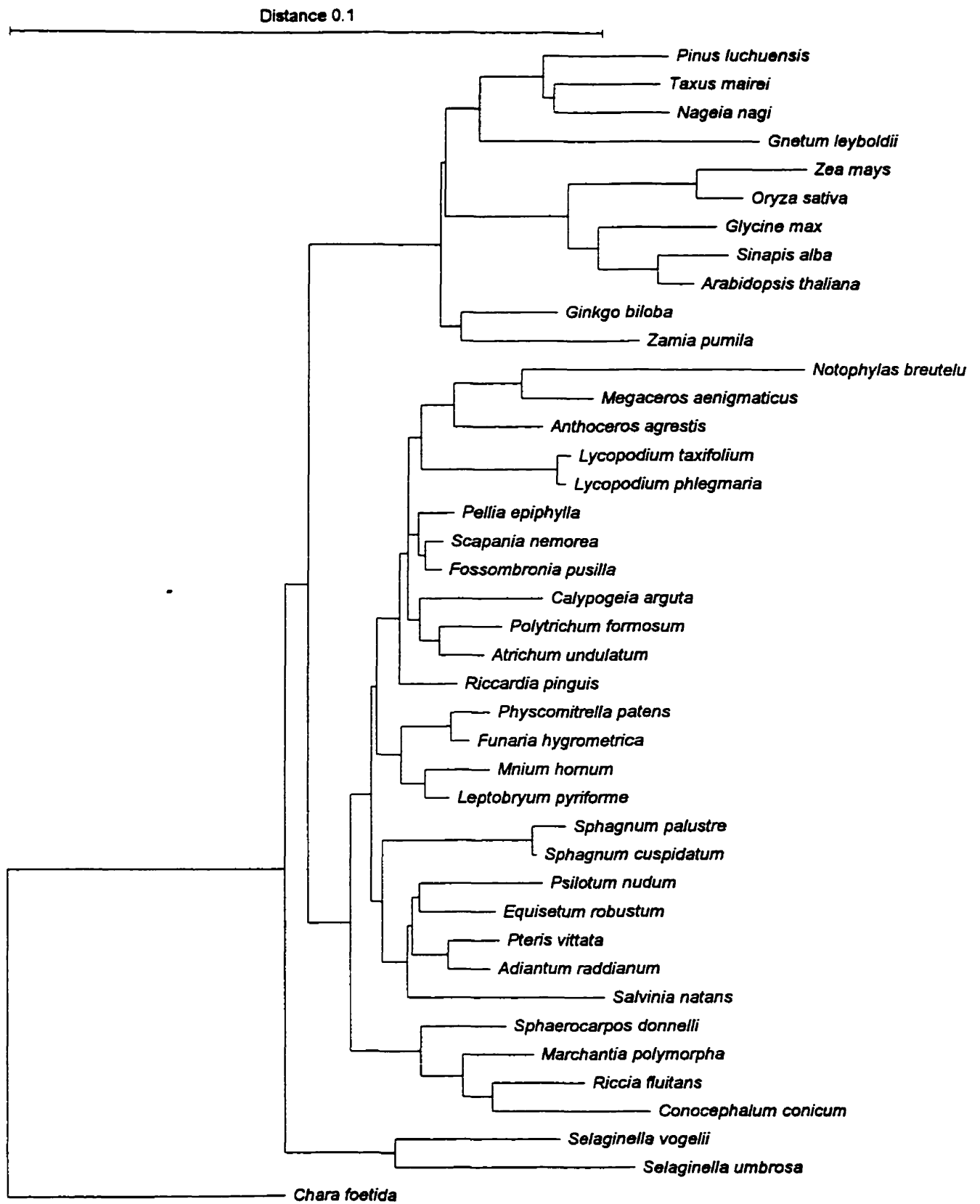


Figure D2.1S Jin & Nei (Distance Calculation); Kimura 2-p (Evolution Model), Gamma distance = 1

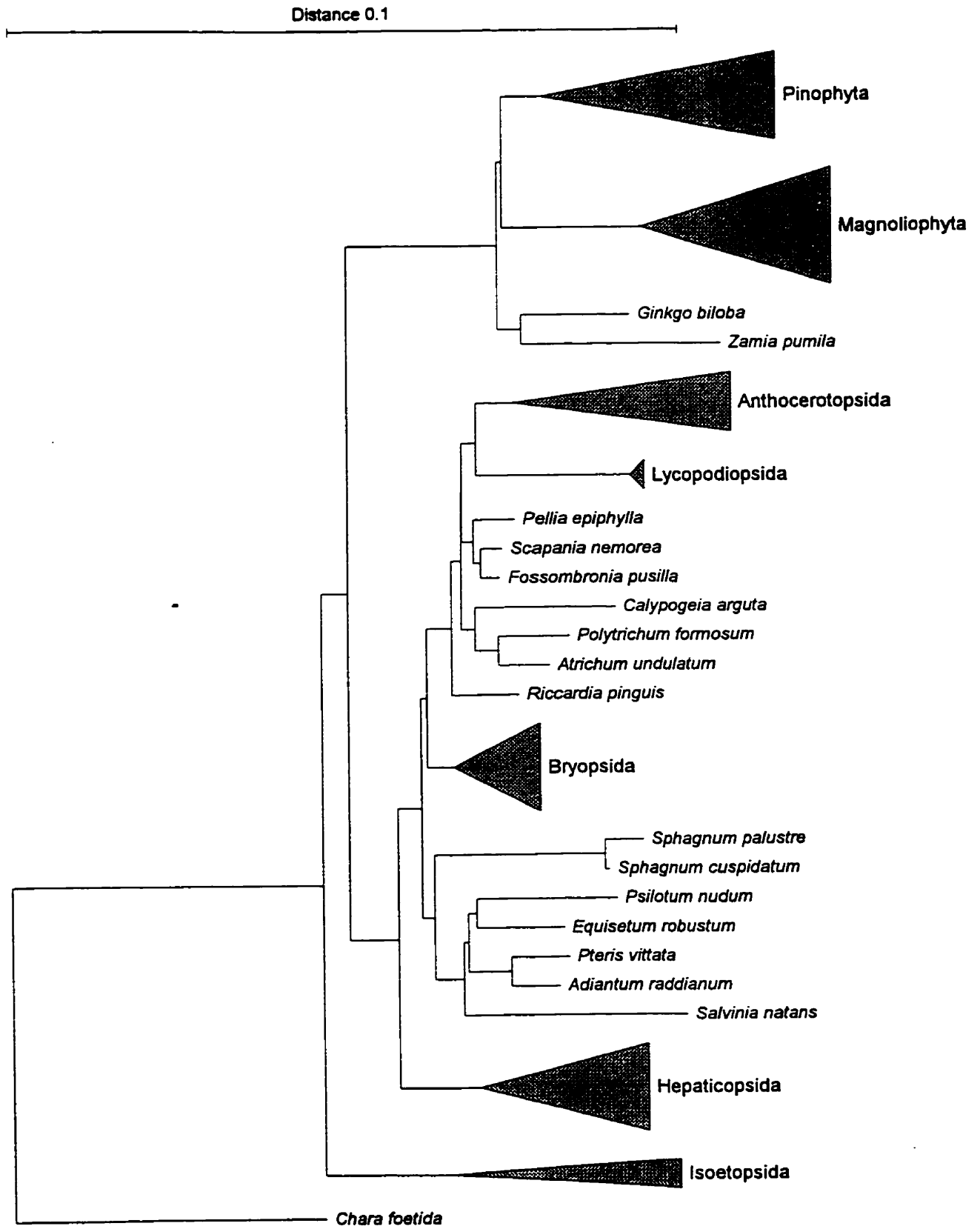


Figure D2.1S Jin & Nei (Distance Calculation); Kimura 2-p (Evolution Model), Gamma distance = 1

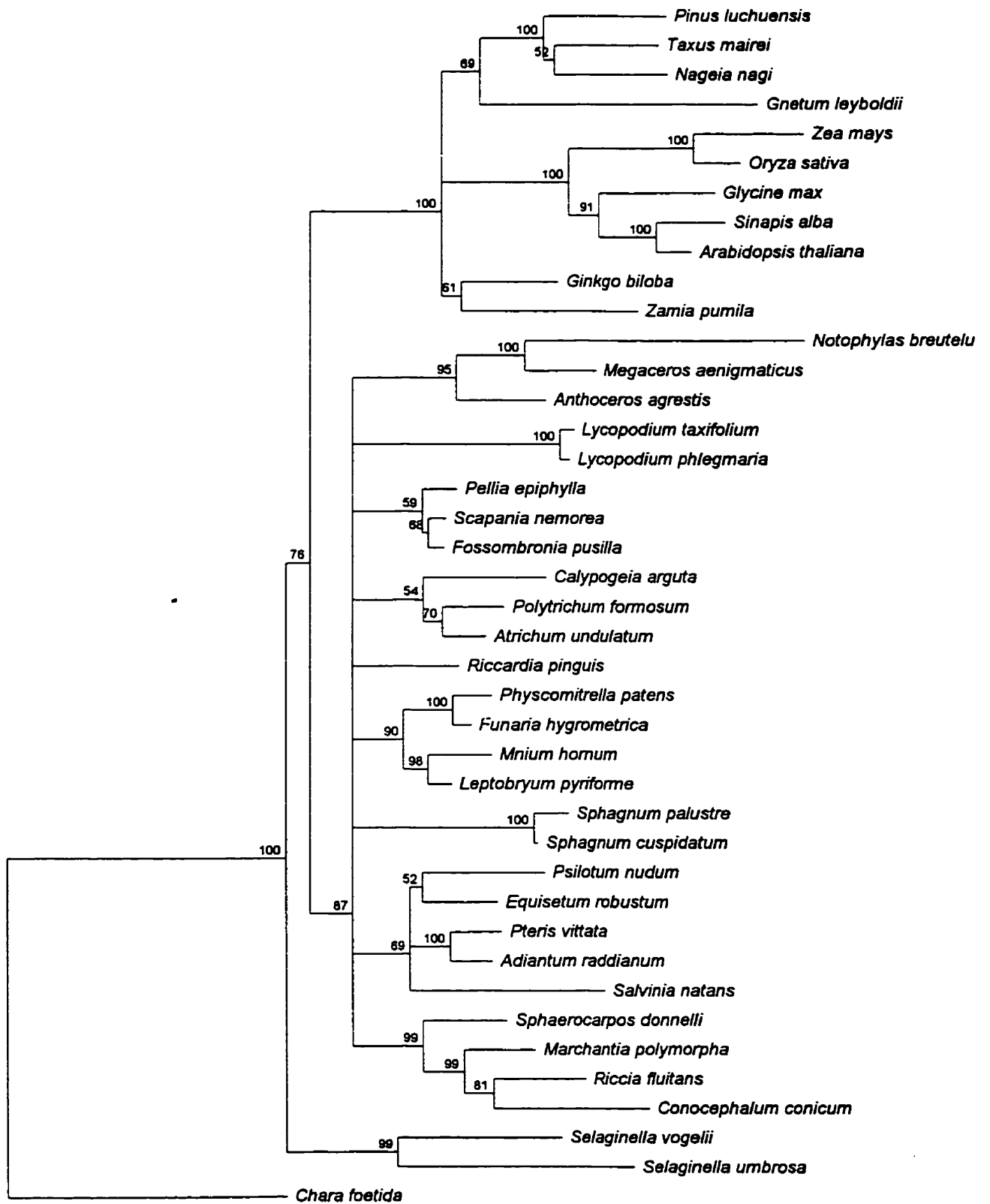


Figure D2.1SB Jin & Nei (Distance Calculation); Kimura 2-p (Evolution Model), Gamma distance = 1
50% Majority-Rule Bootstrap Tree

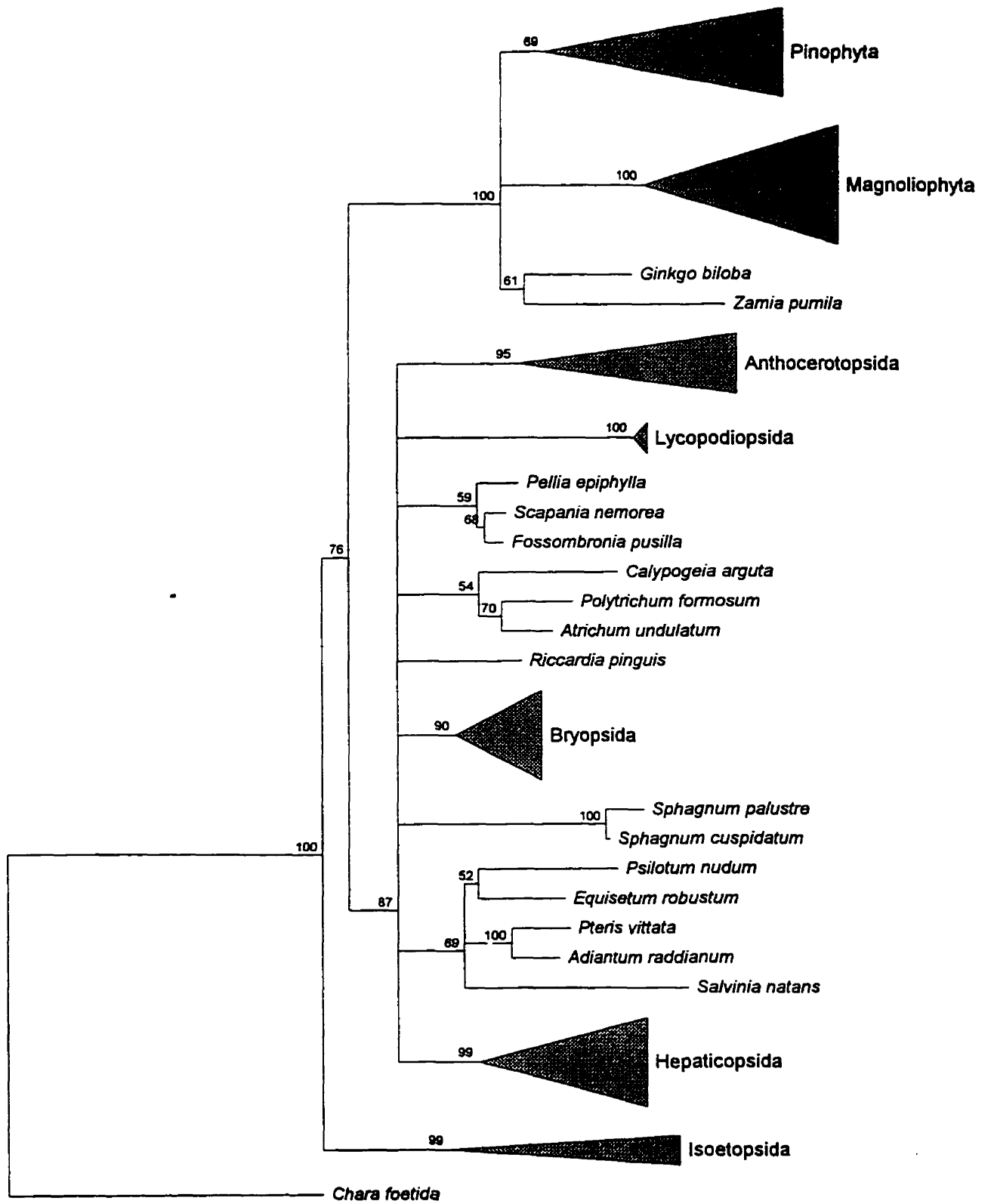


Figure D2.1SB Jin & Nei (Distance Calculation); Kimura 2-p (Evolution Model), Gamma distance = 1
50% Majority-Rule Bootstrap Tree



Figure D2.2S Jin & Nei (Distance Calculation); Kimura 2-p (Evolution Model), Gamma distance = 1
Insertions & Deletions taken into account

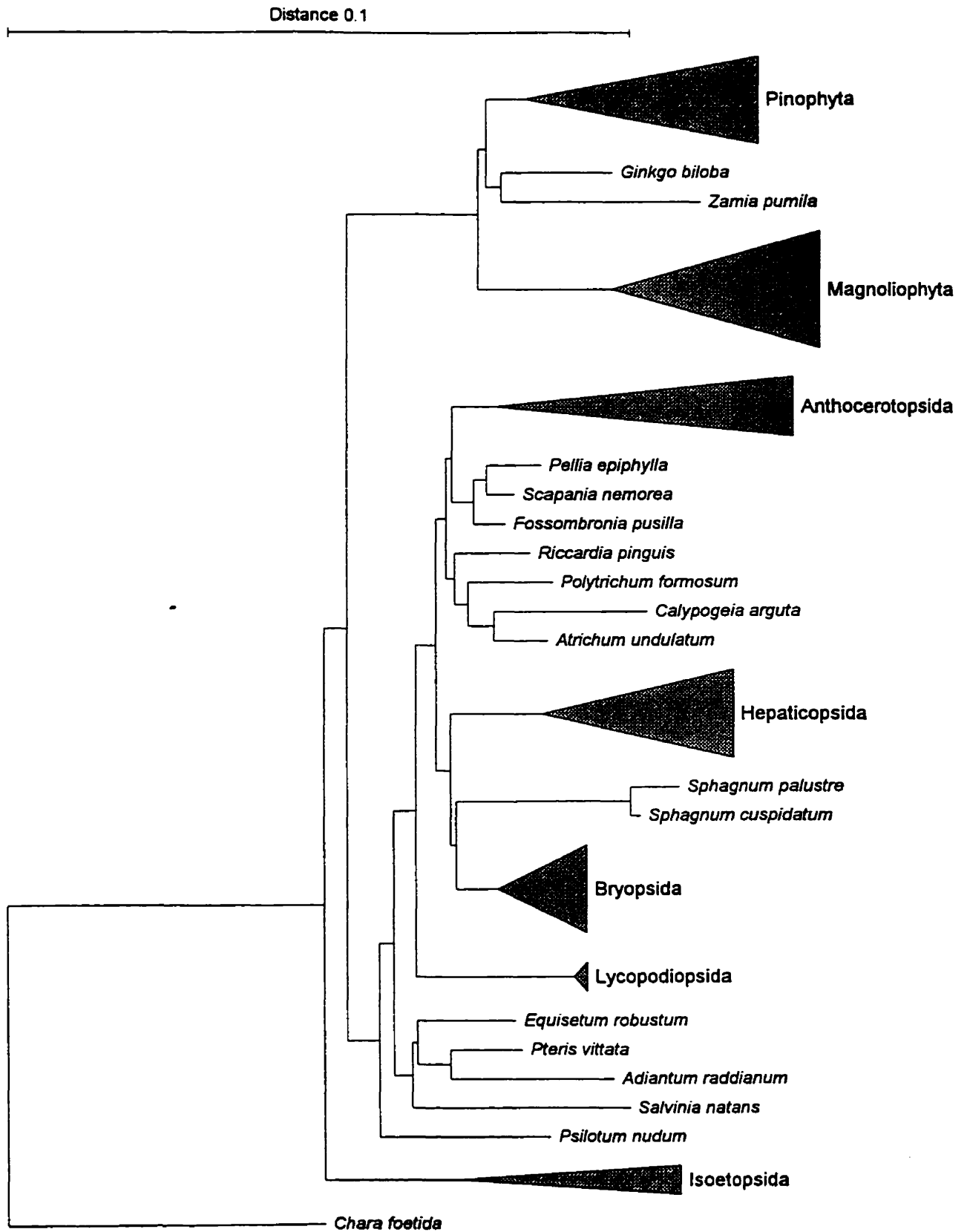


Figure D2.2S Jin & Nei (Distance Calculation); Kimura 2-p (Evolution Model), Gamma distance = 1
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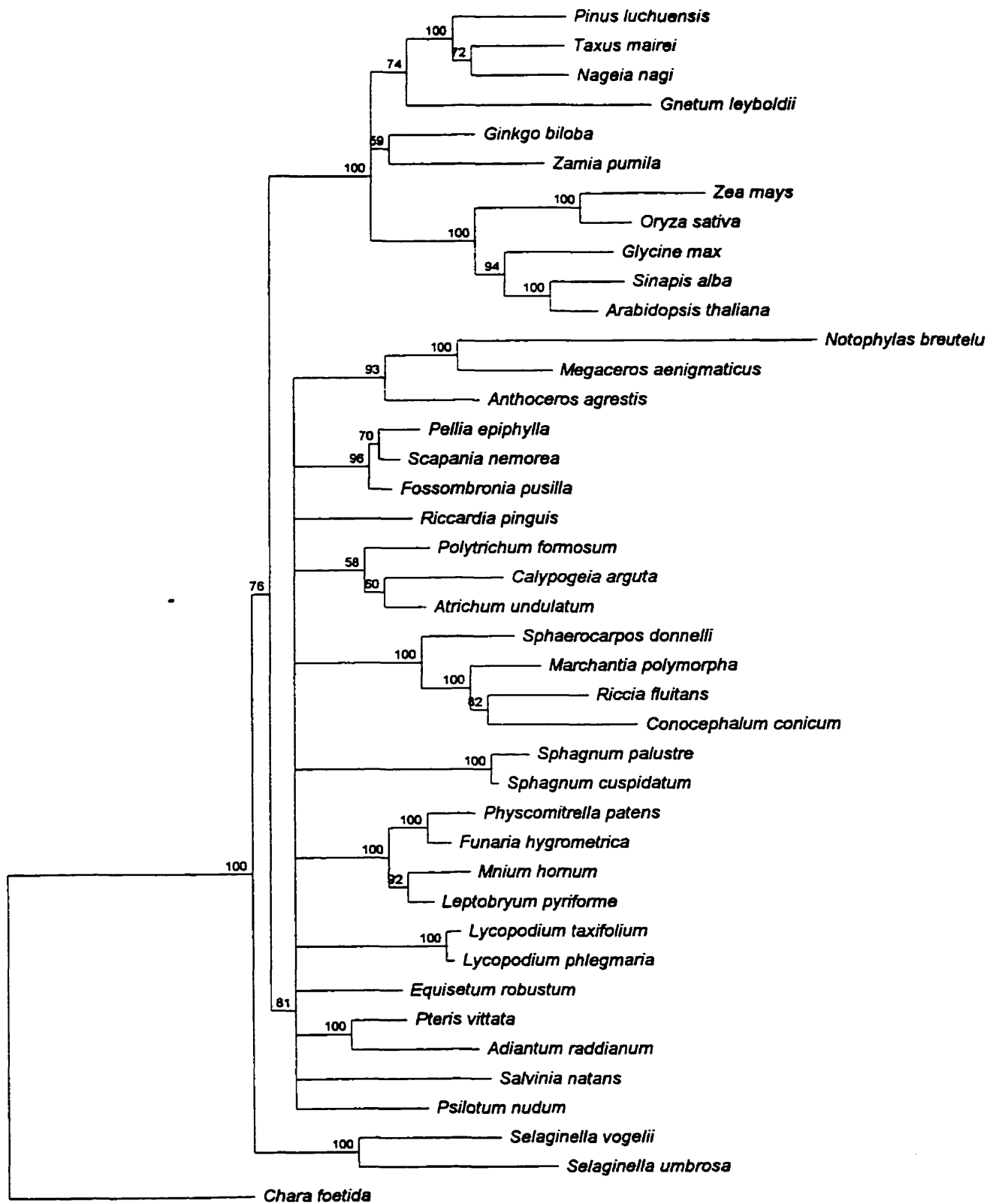


Figure D2.2SB Jin & Nei (Distance Calculation); Kimura 2-p (Evolution Model), Gamma distance = 1
 Insertions & Deletions taken into account
 50% Majority-Rule Bootstrap Tree

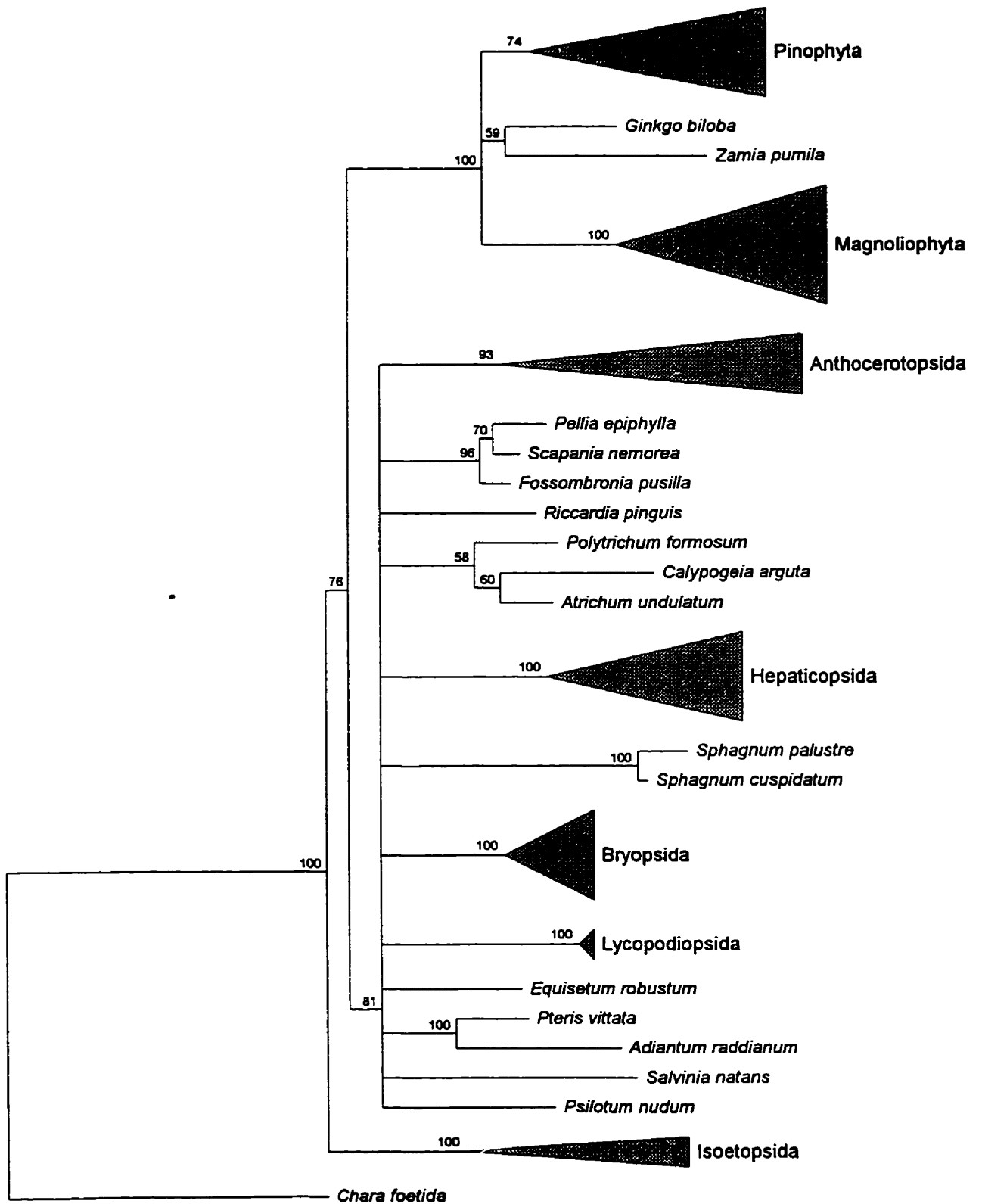


Figure D2.2SB Jin & Nei (Distance Calculation); Kimura 2-p (Evolution Model), Gamma distance = 1
 Insertions & Deletions taken into account
 50% Majority-Rule Bootstrap Tree

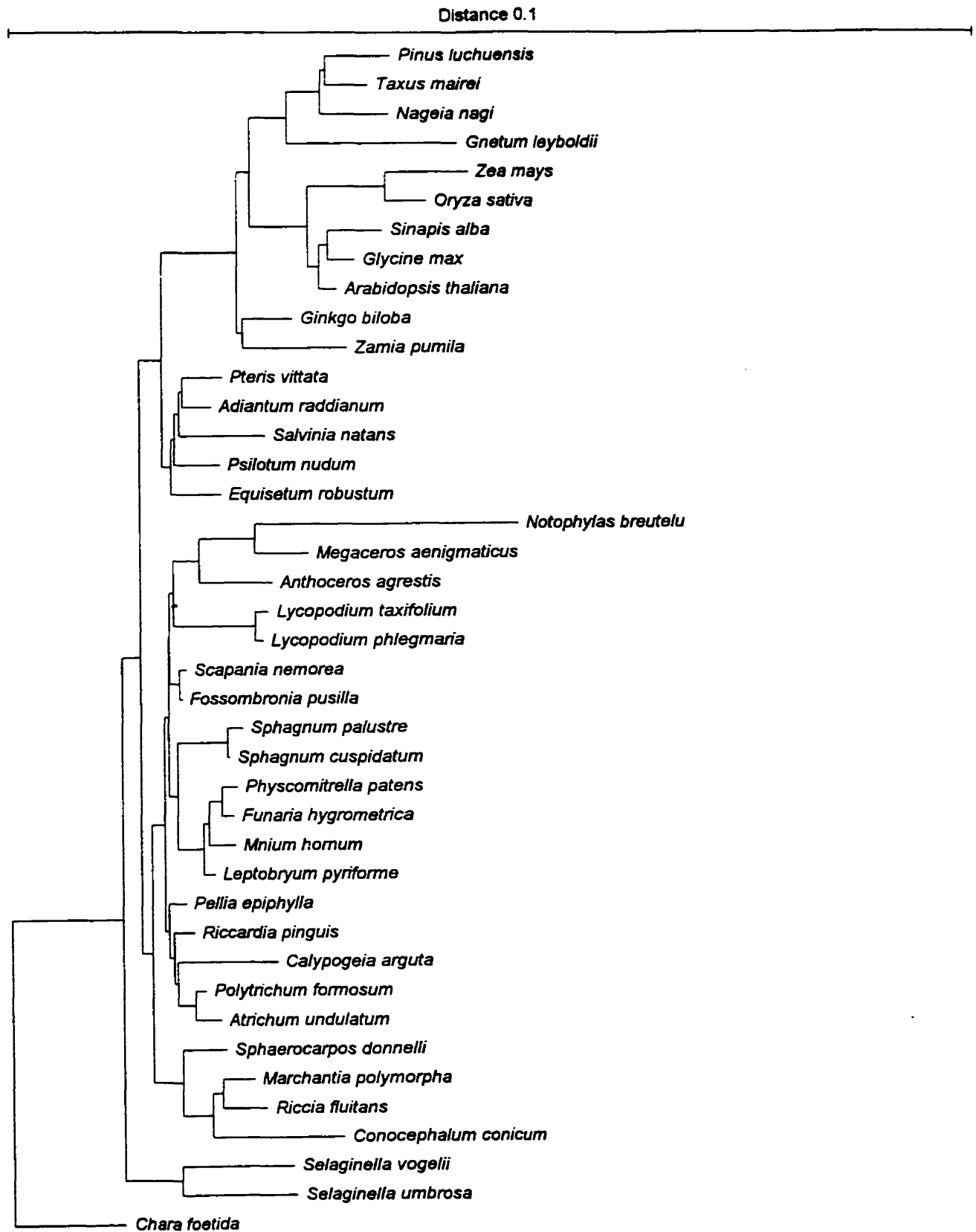


Figure E1S Transversions Only; Neighbor-joining

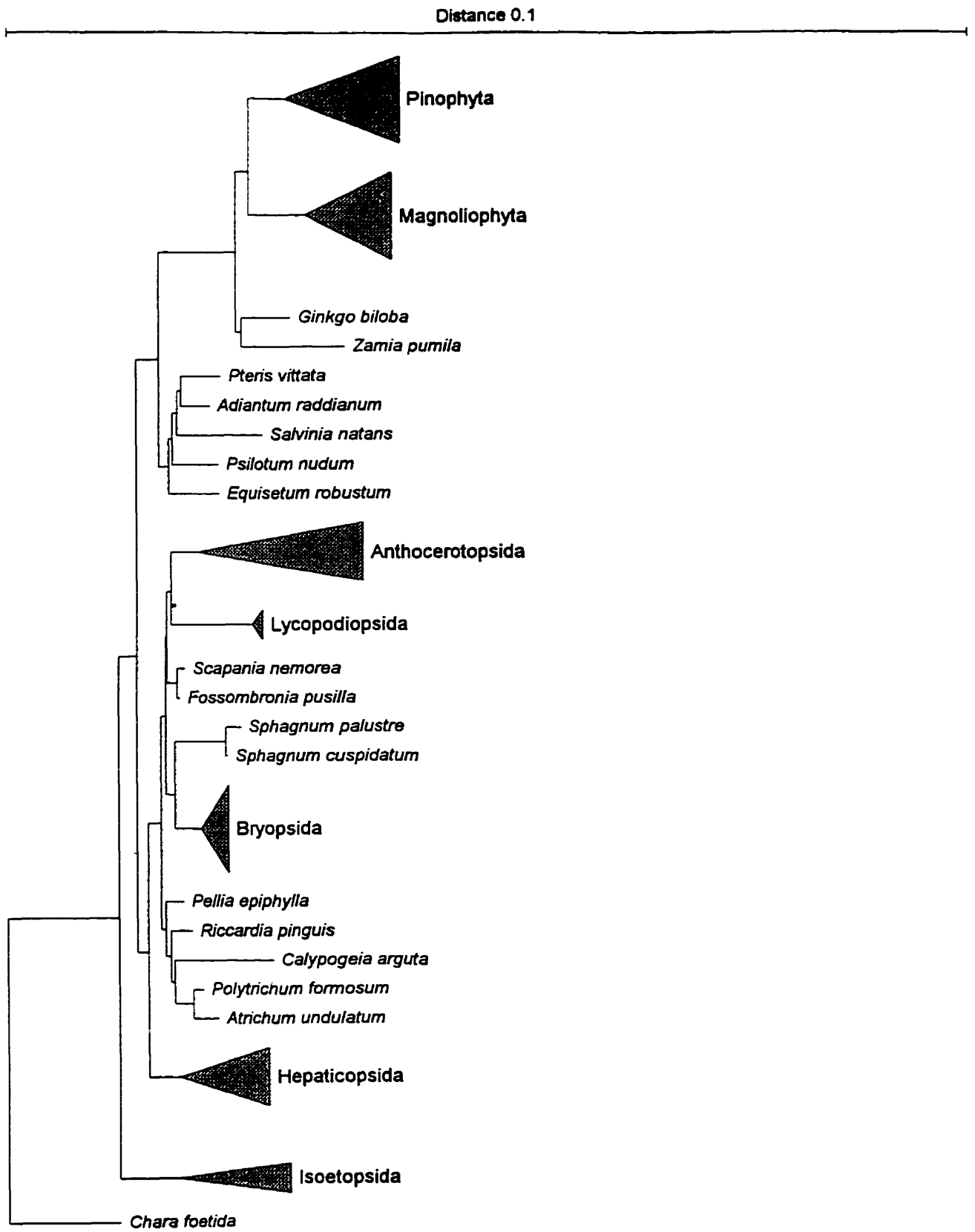


Figure E1S Transversions Only; Neighbor-joining

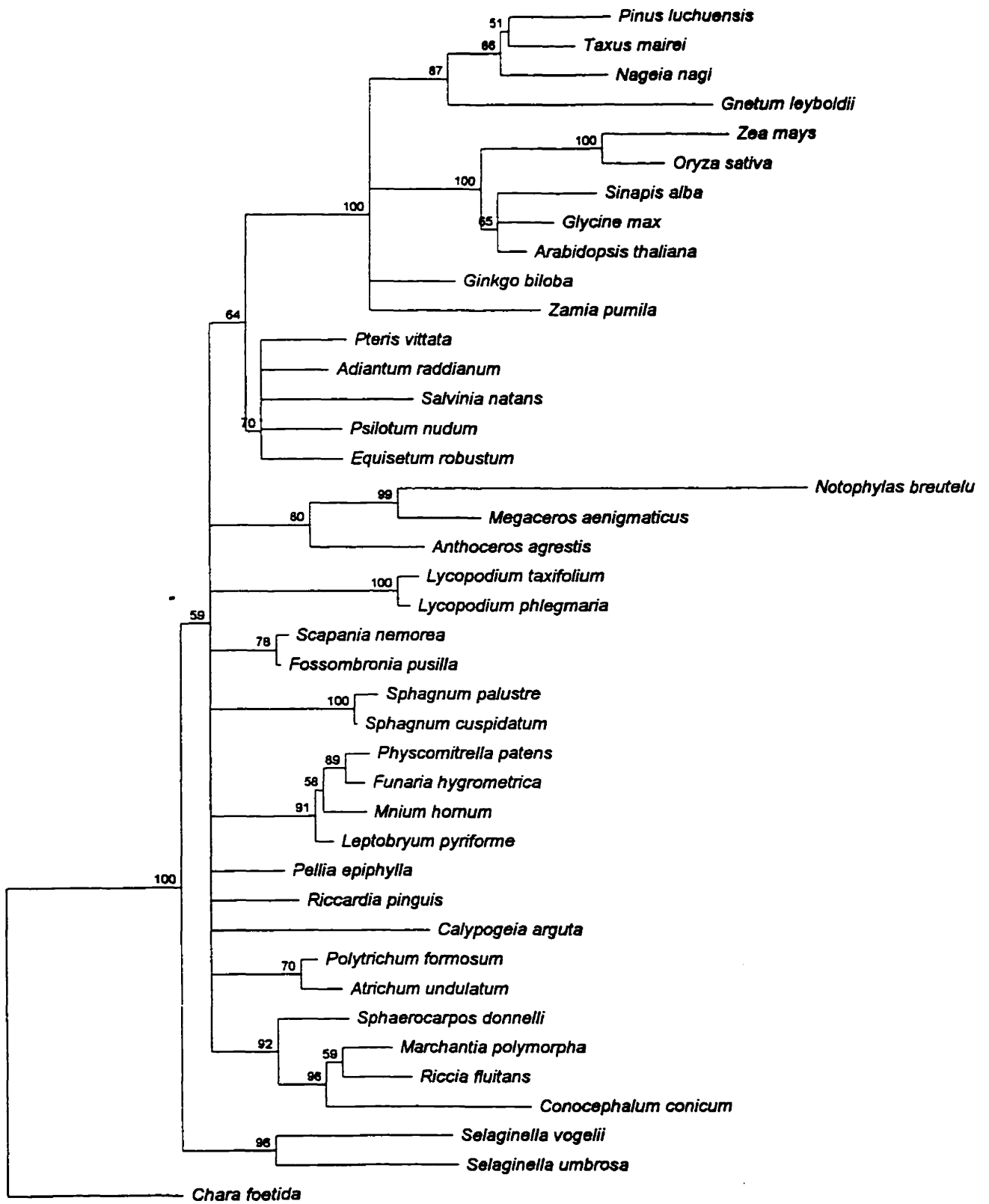


Figure E1SB Transversions Only; Neighbor-joining
50% Majority-Rule Bootstrap Tree

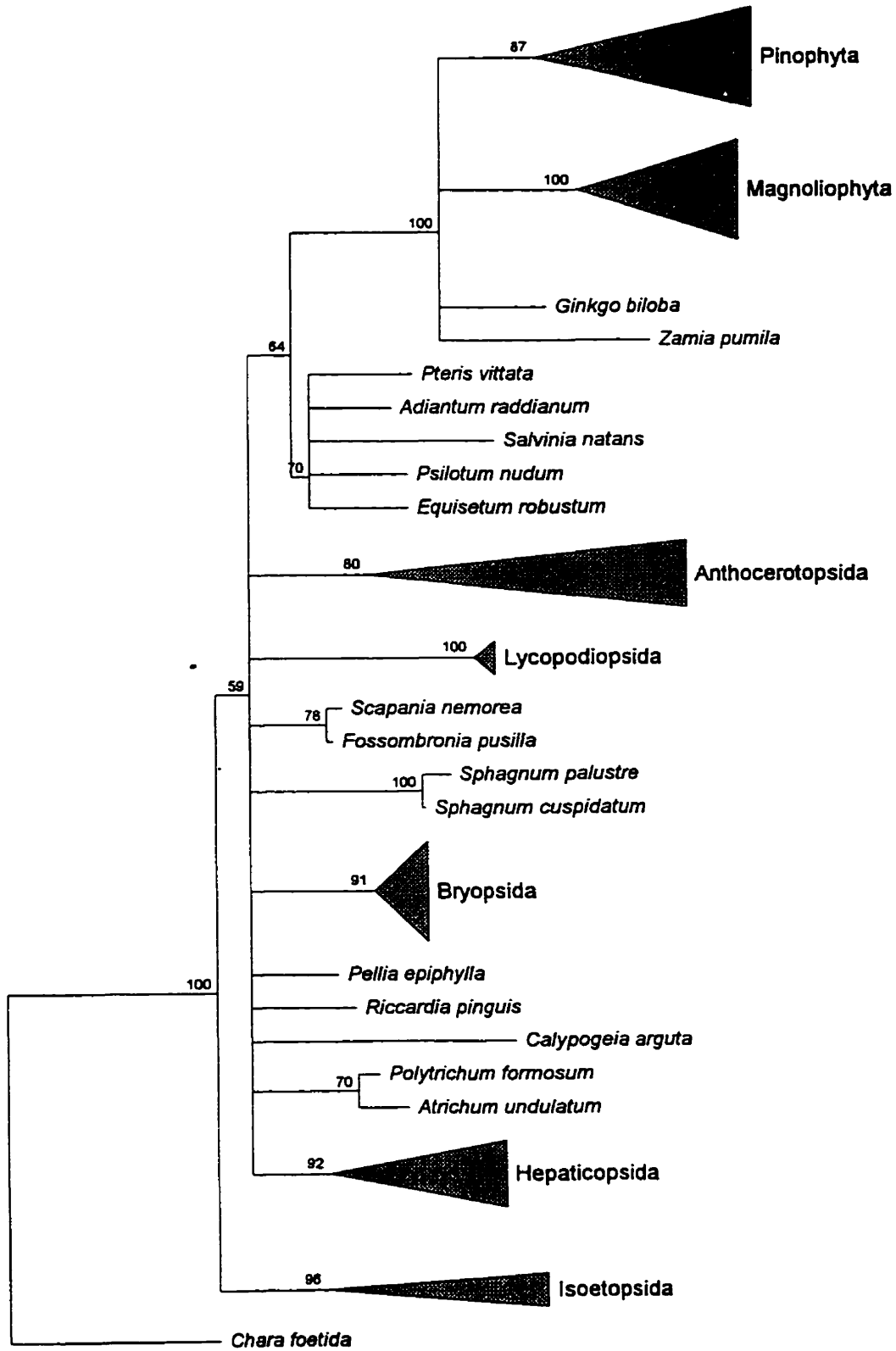


Figure E1SB Transversions Only; Neighbor-joining
50% Majority-Rule Bootstrap Tree

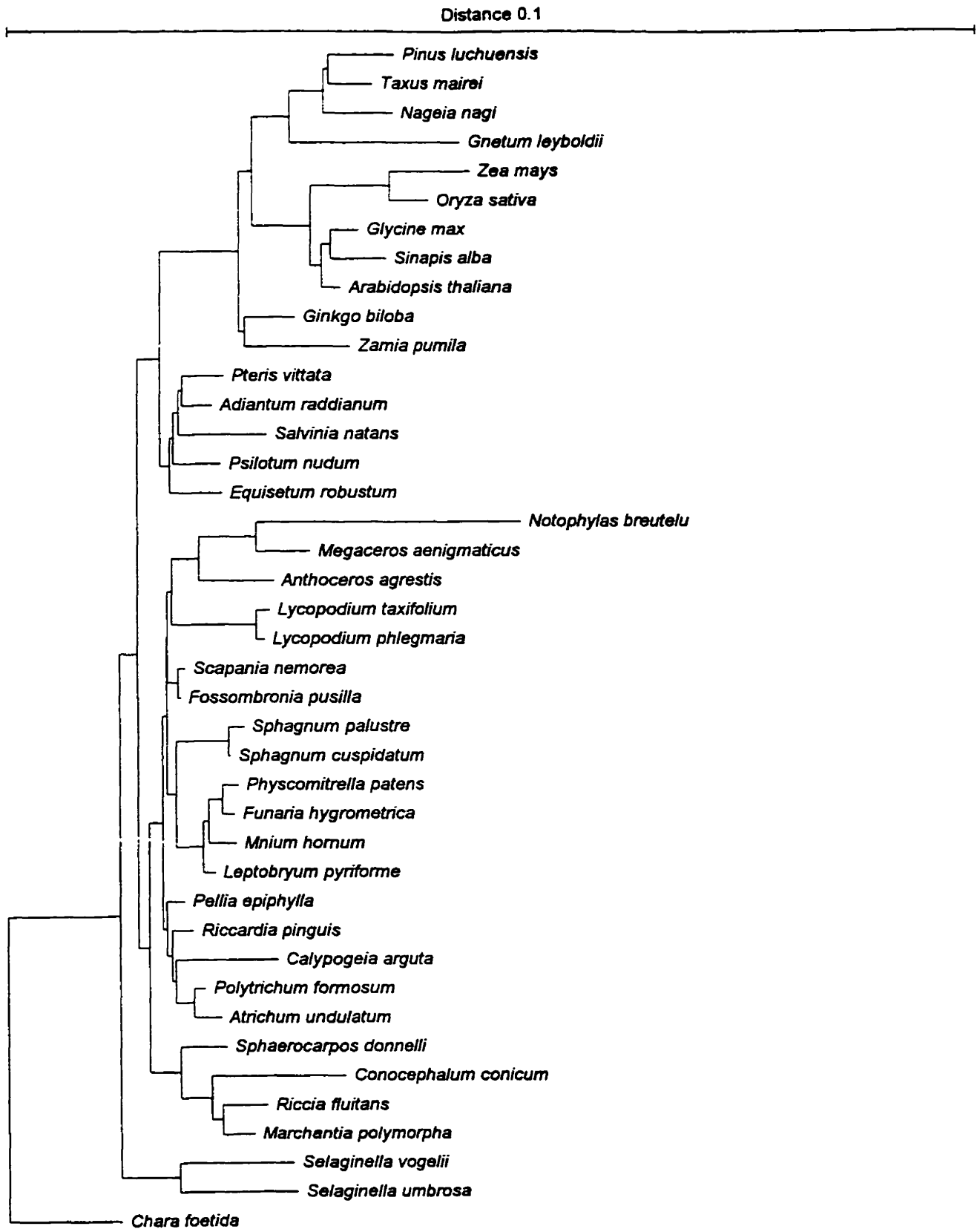


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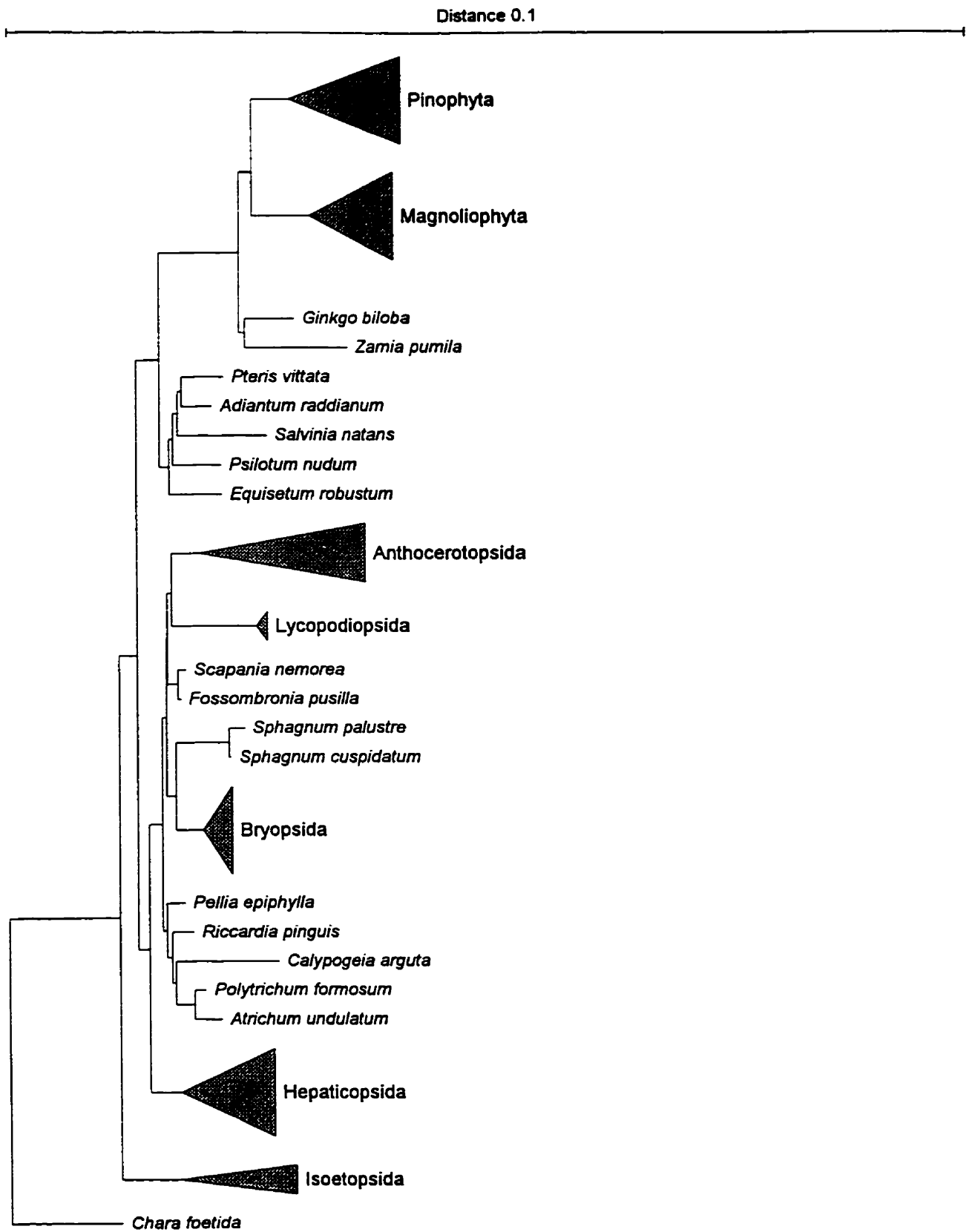


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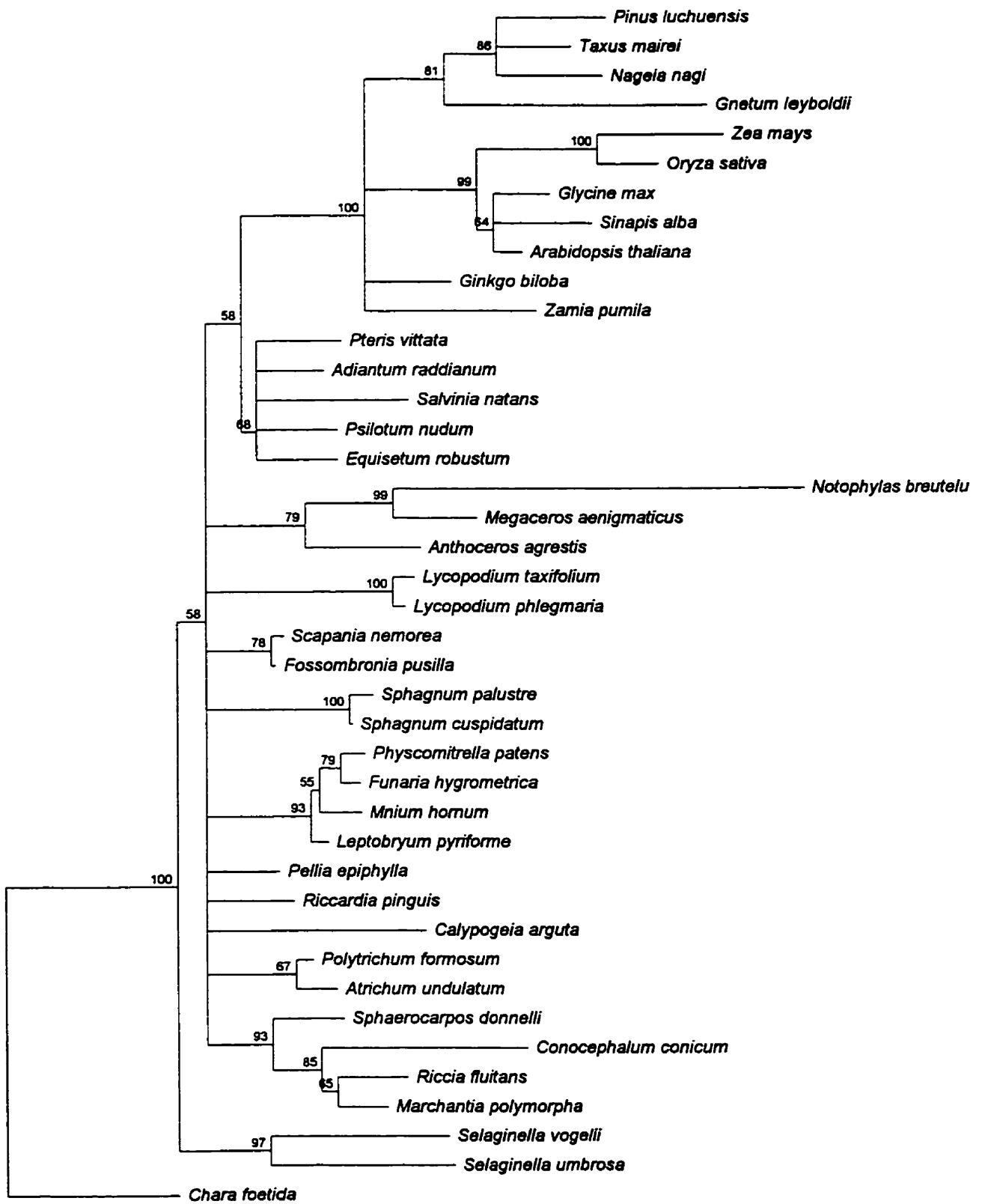


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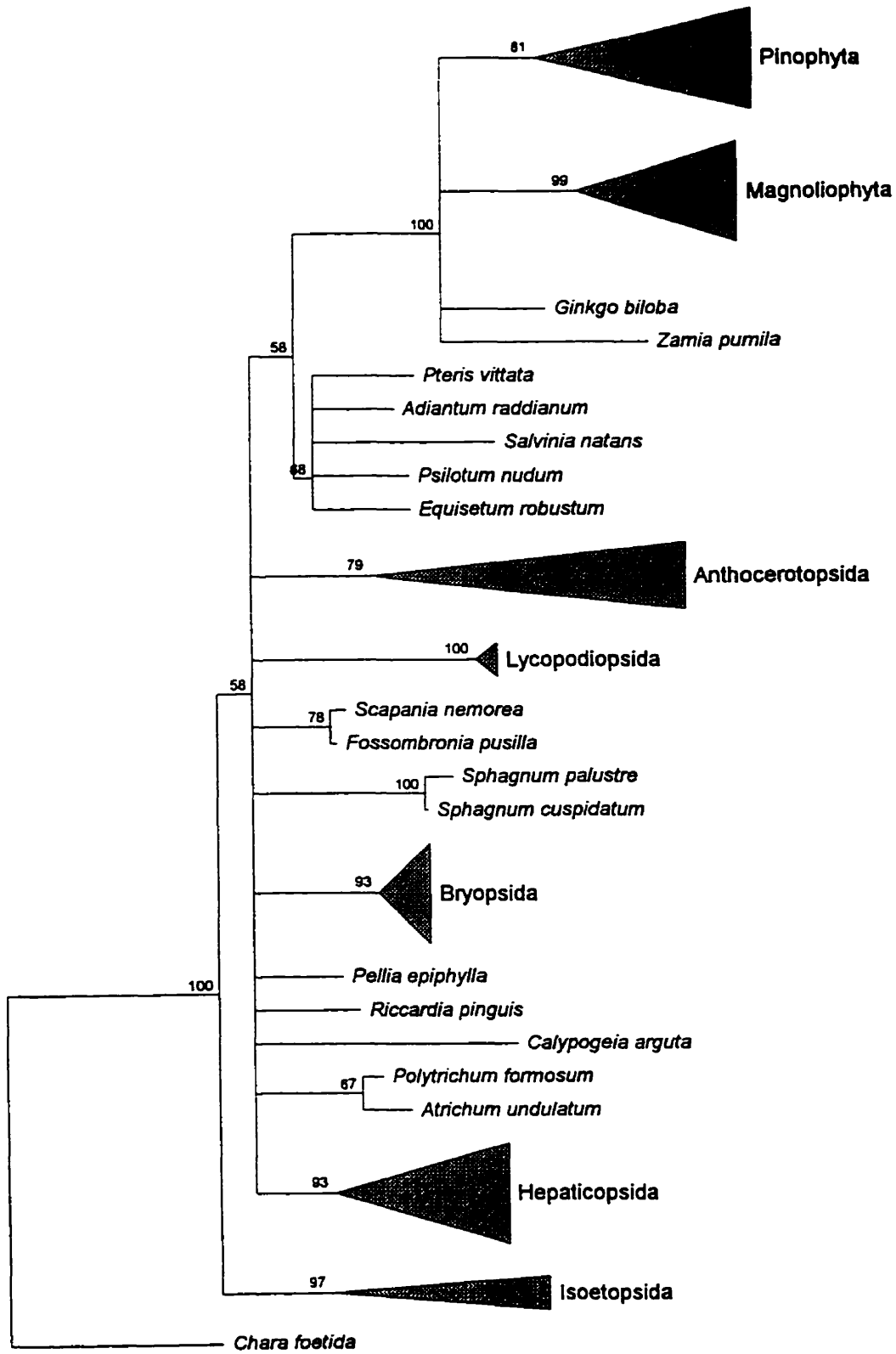


Figure E2SB Transversions Only; Insertions & Deletions taken into account
50% Majority-Rule Bootstrap Tree

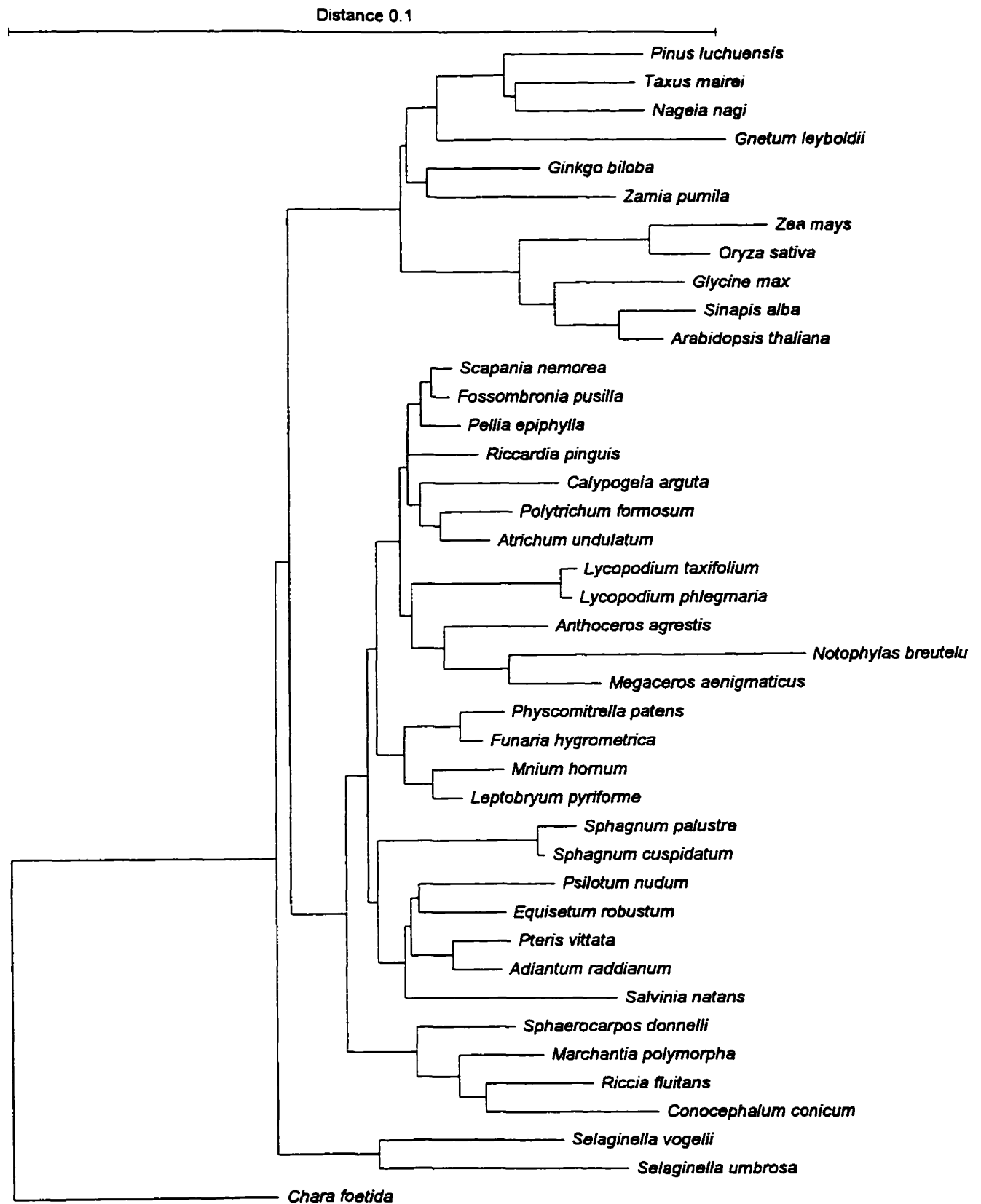


Figure F1S No Correction; Neighbor-joining

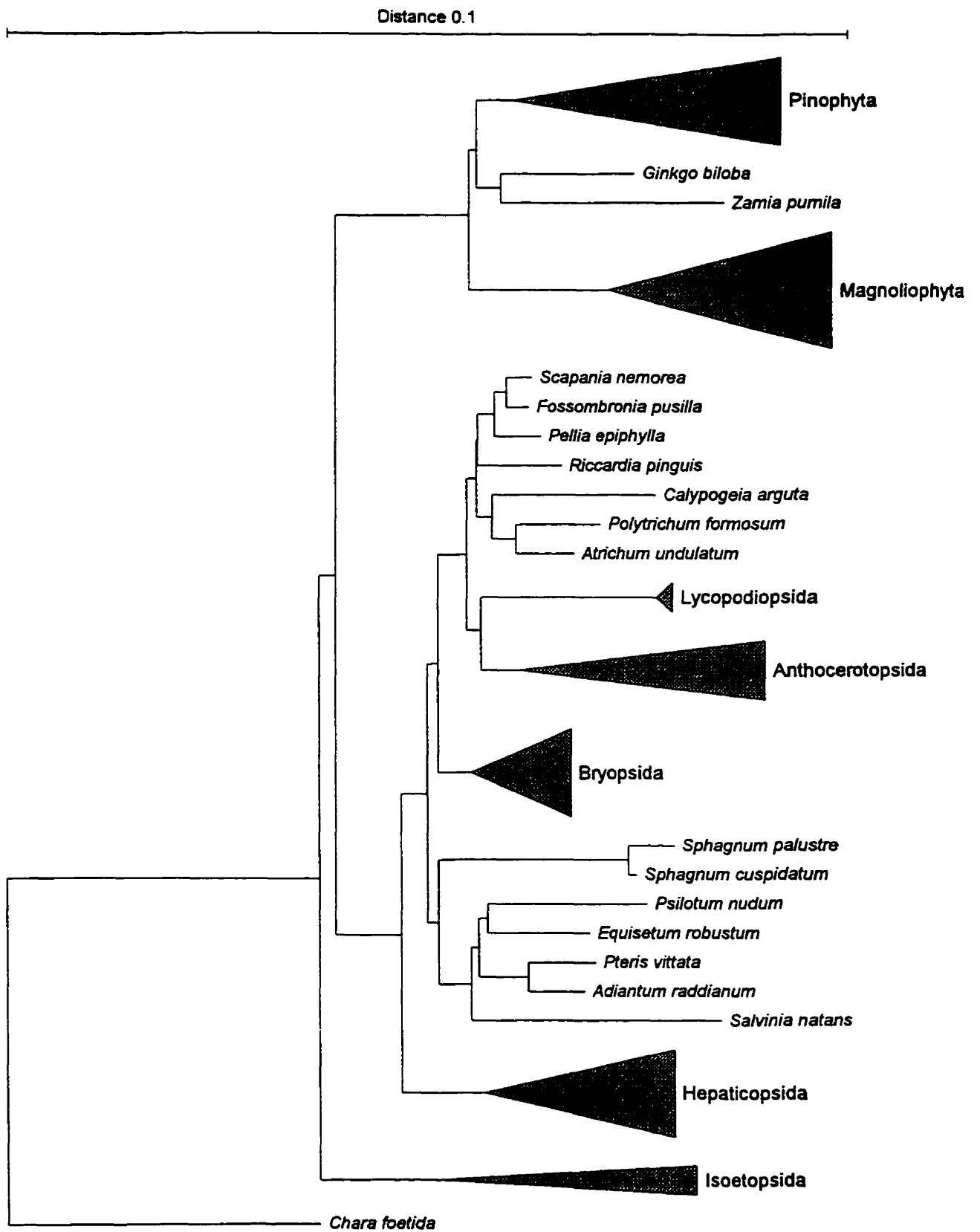


Figure F1S No Correction; Neighbor-joining

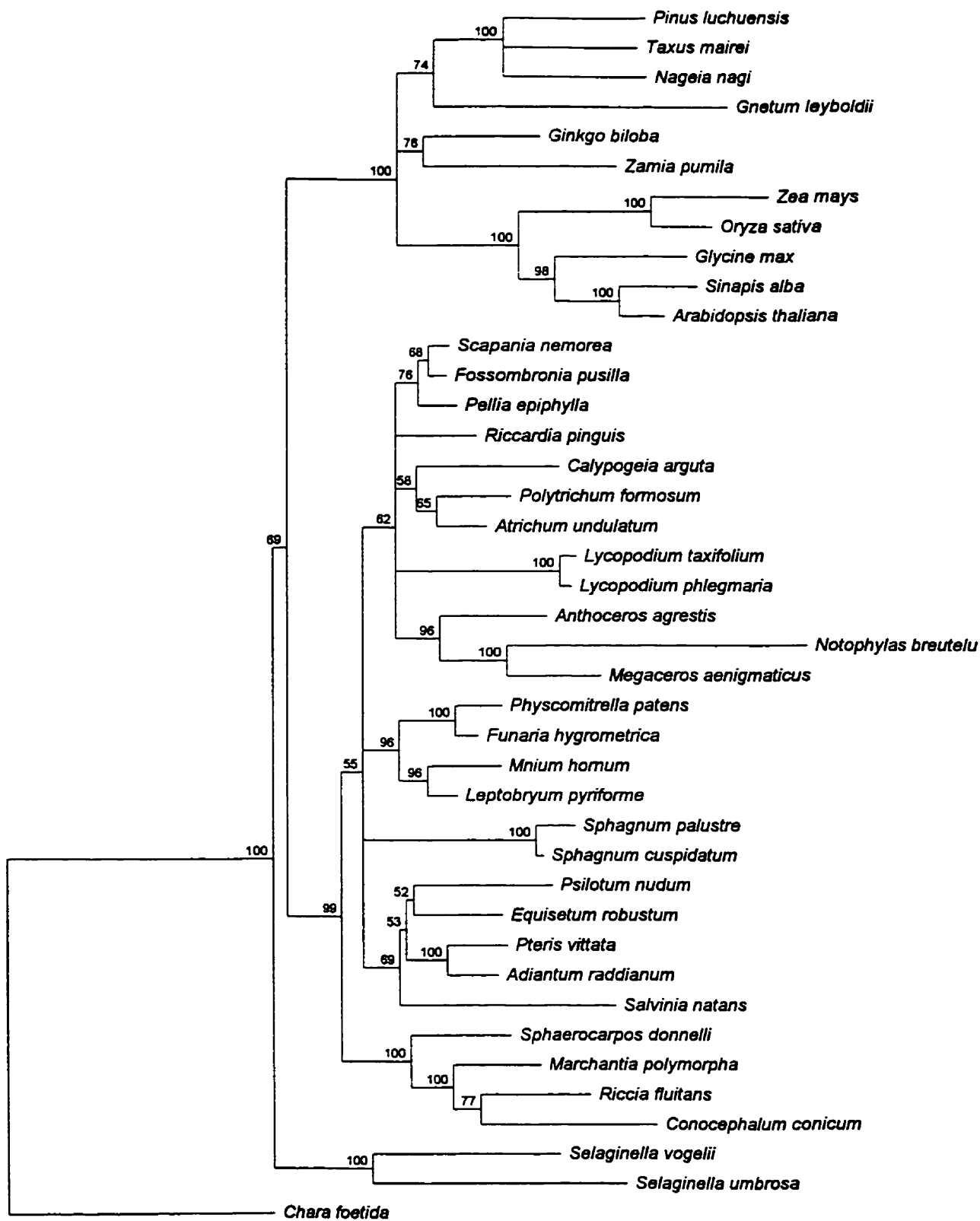


Figure F1SB No Correction; Neighbor-joining
50% Majority-Rule Bootstrap Tree

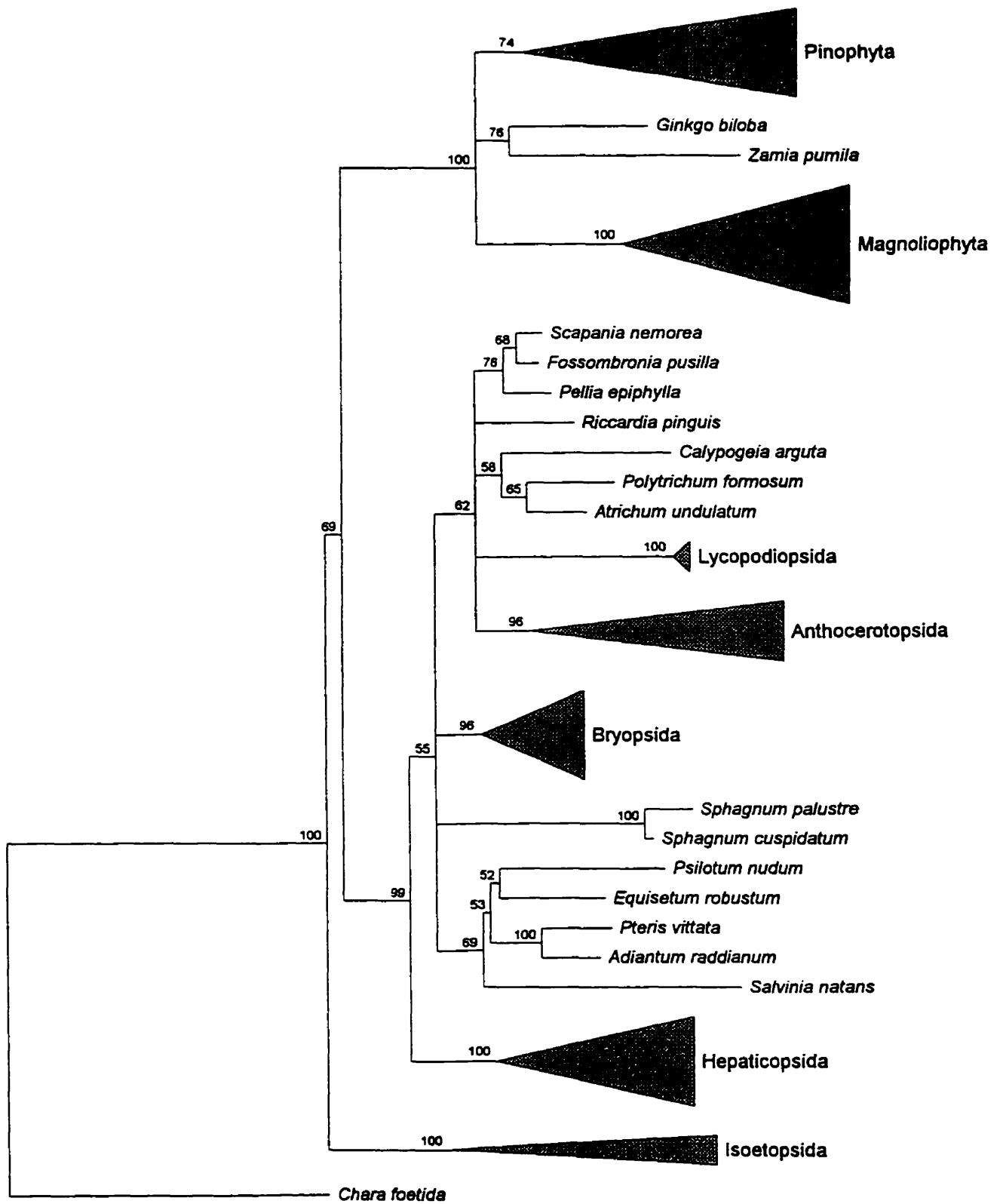


Figure F1SB No Correction; Neighbor-joining
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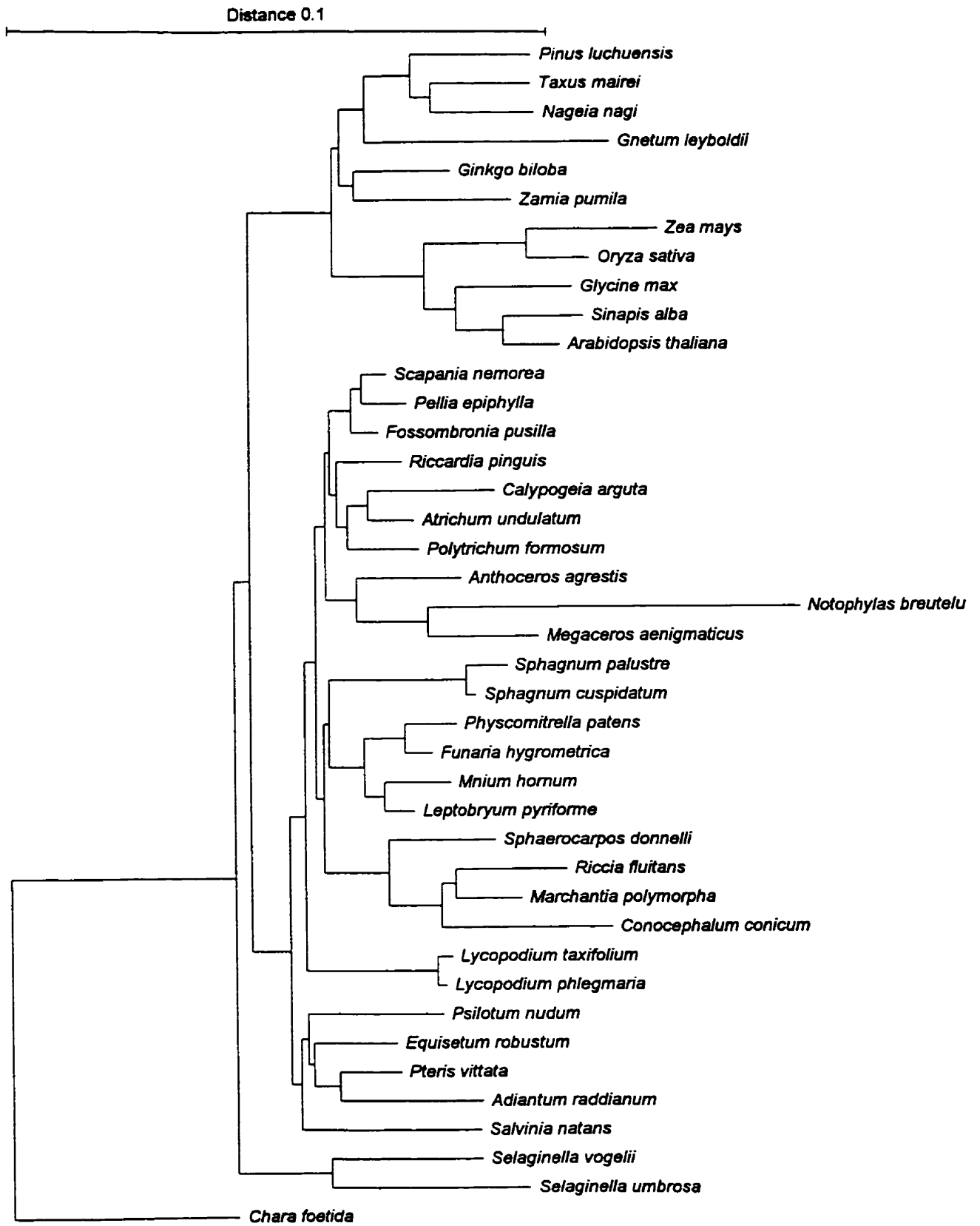
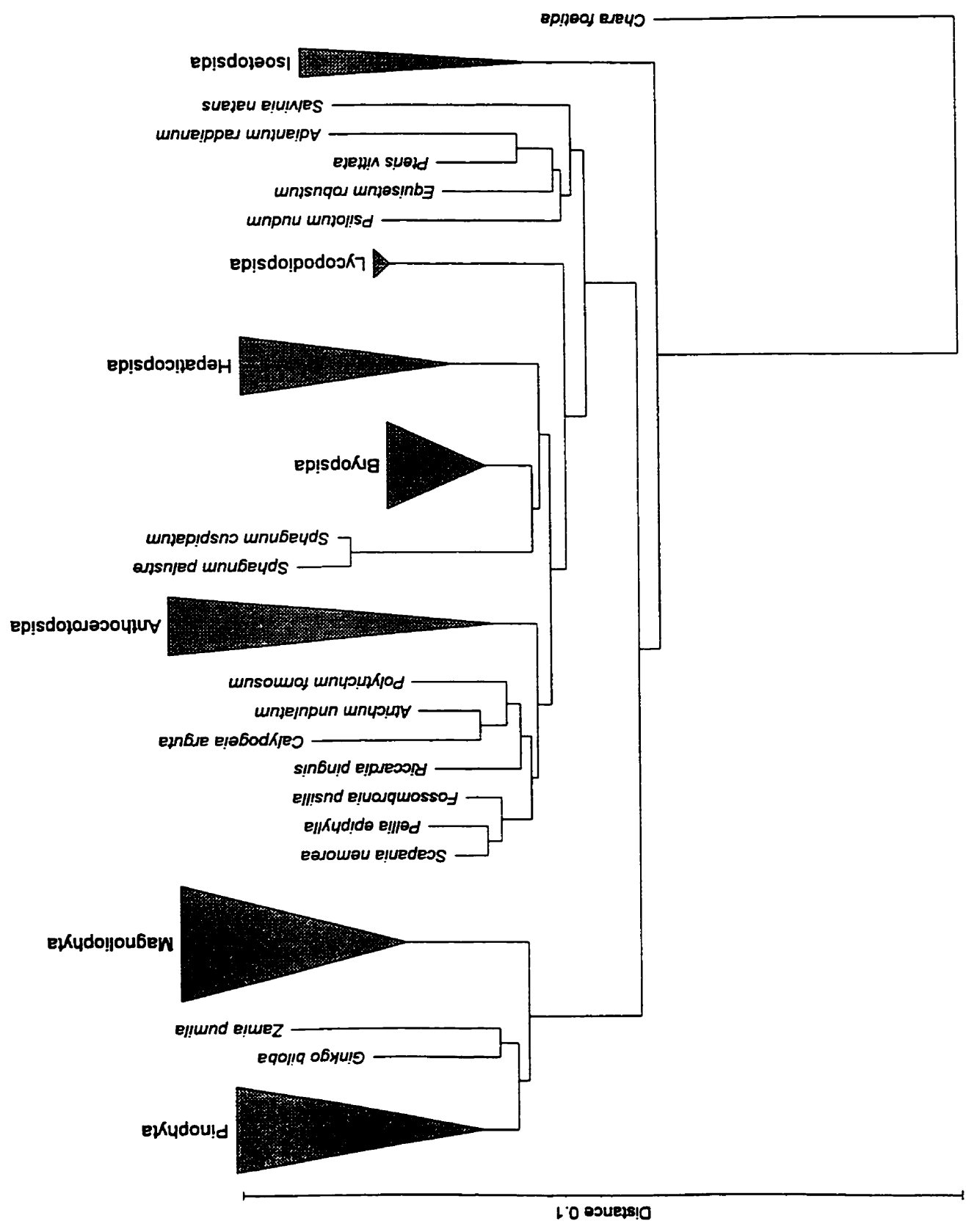


Figure F2S No Correction; Insertions & Deletions taken into account

Figure F-25 No Correction; Insertions & Deletions taken into account



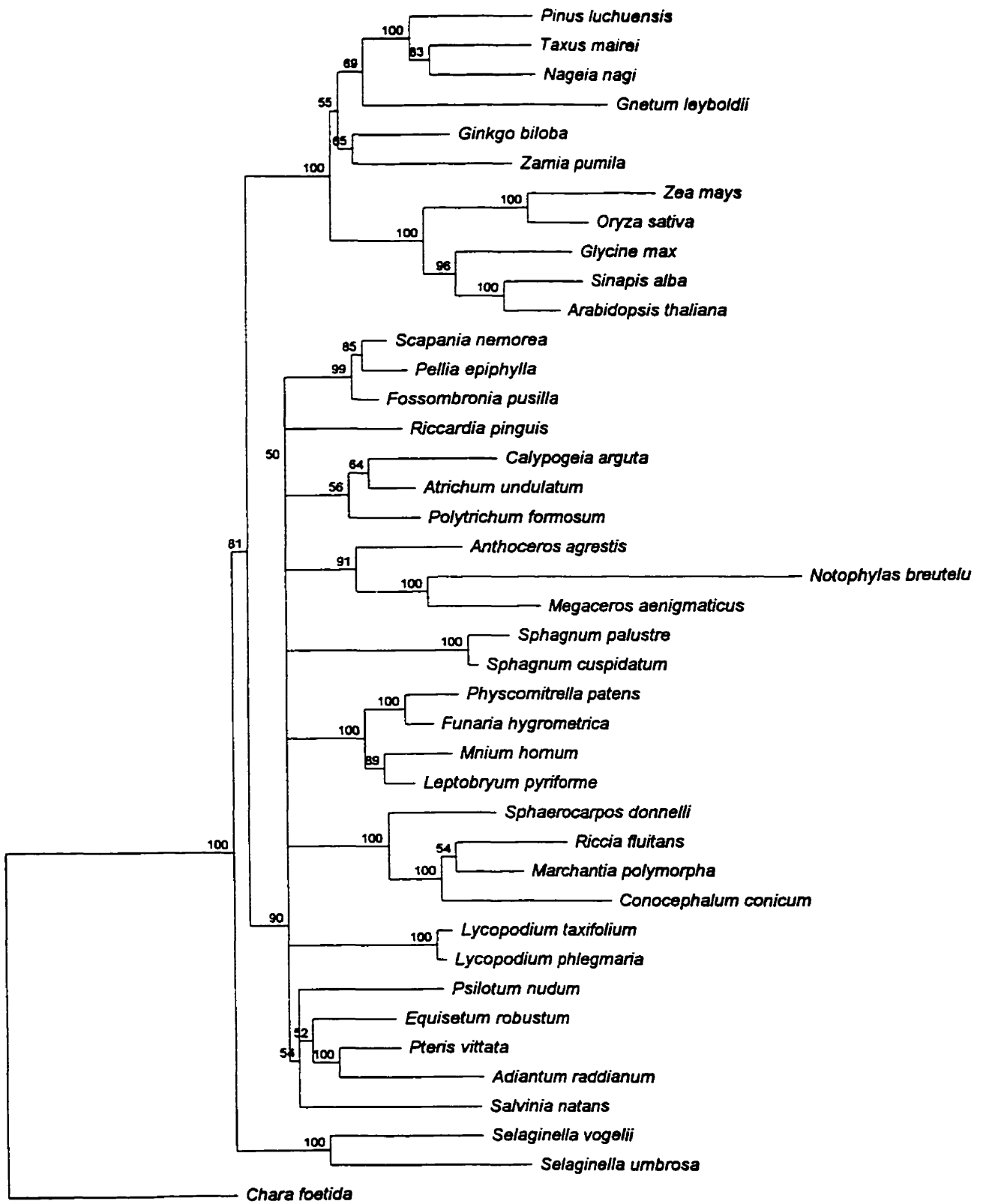


Figure F2SB No Correction; Insertions & Deletions taken into account
 50% Majority-Rule Bootstrap Tree

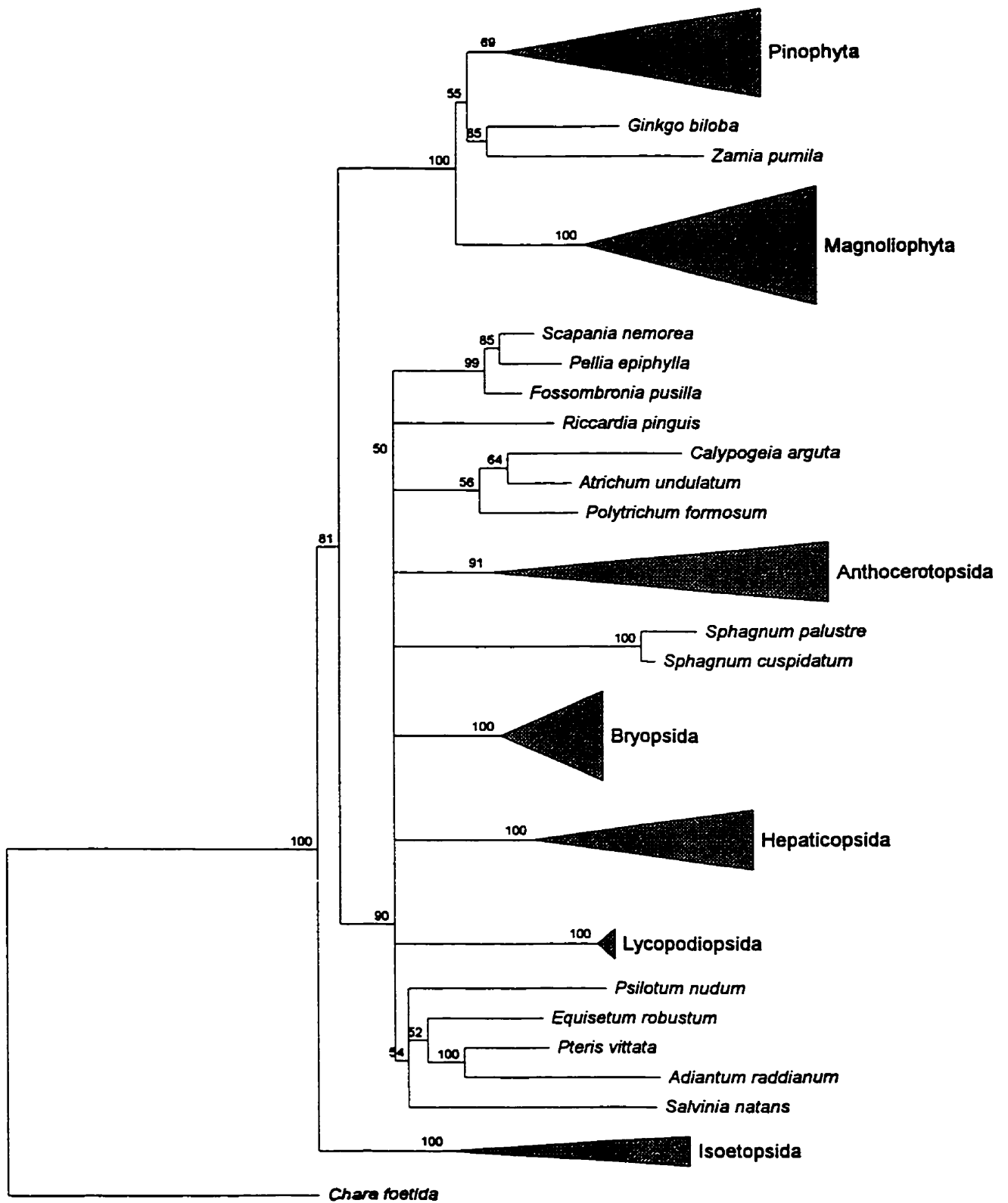


Figure F2SB No Correction; Insertions & Deletions taken into account
50% Majority-Rule Bootstrap Tree

Distance 0.1

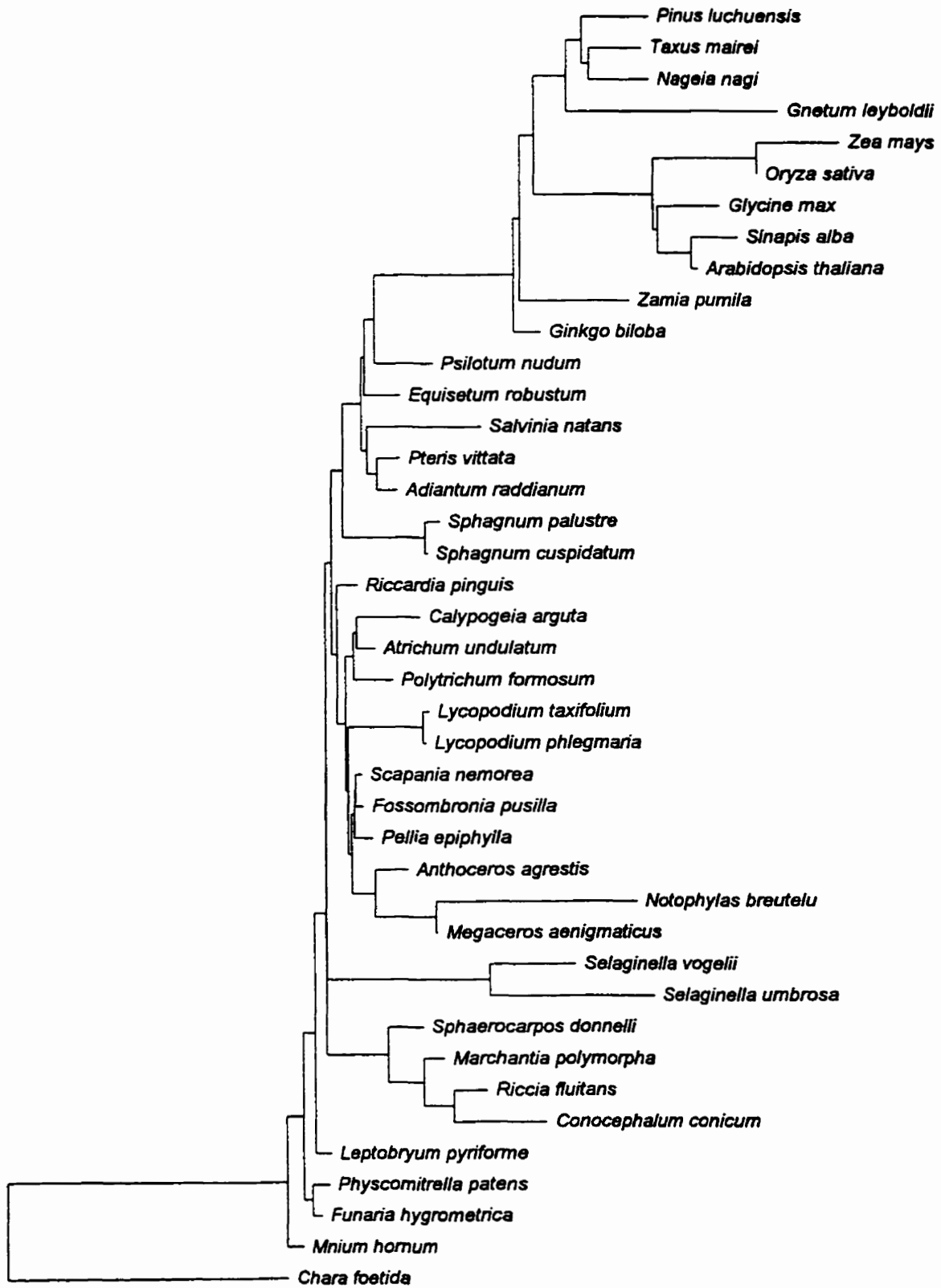


Figure G1S Van de Peer & De Wachter, Neighbor-joining

Distance 0.1

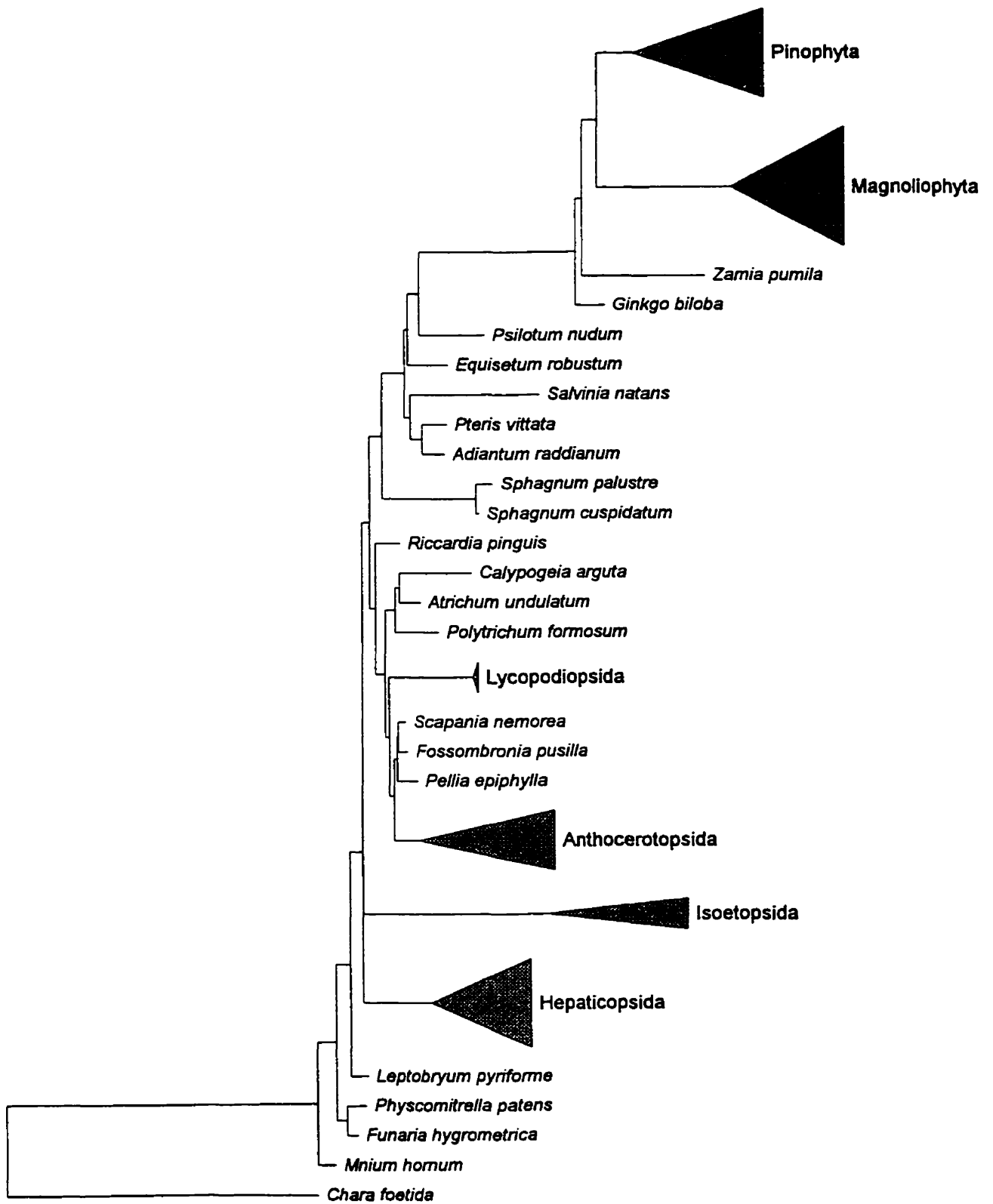


Figure G1S Van de Peer & De Wachter; Neighbor-joining

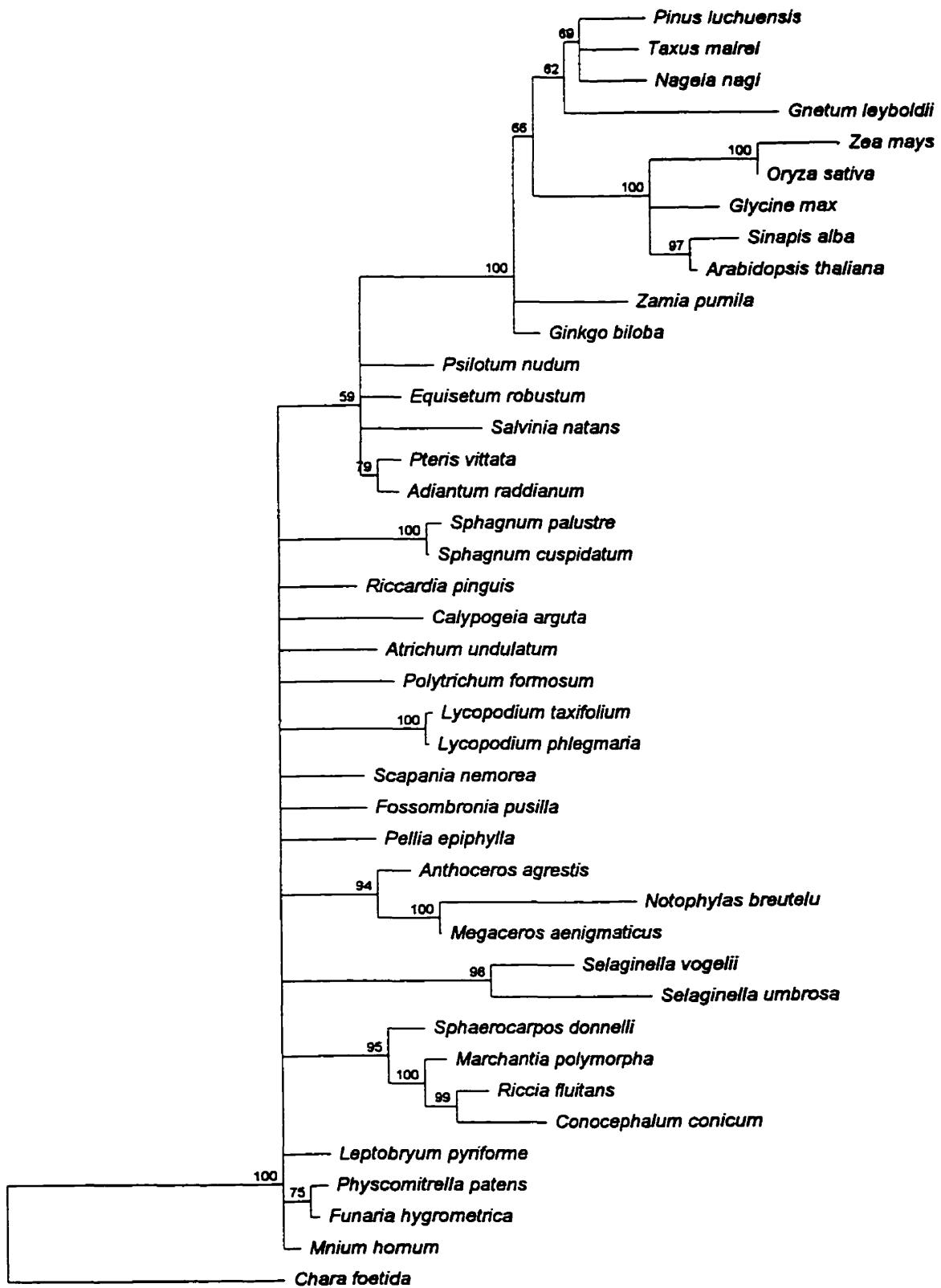


Figure G1SB Van de Peer & De Wachter, Neighbor-joining
50% Majority-Rule Bootstrap Tree

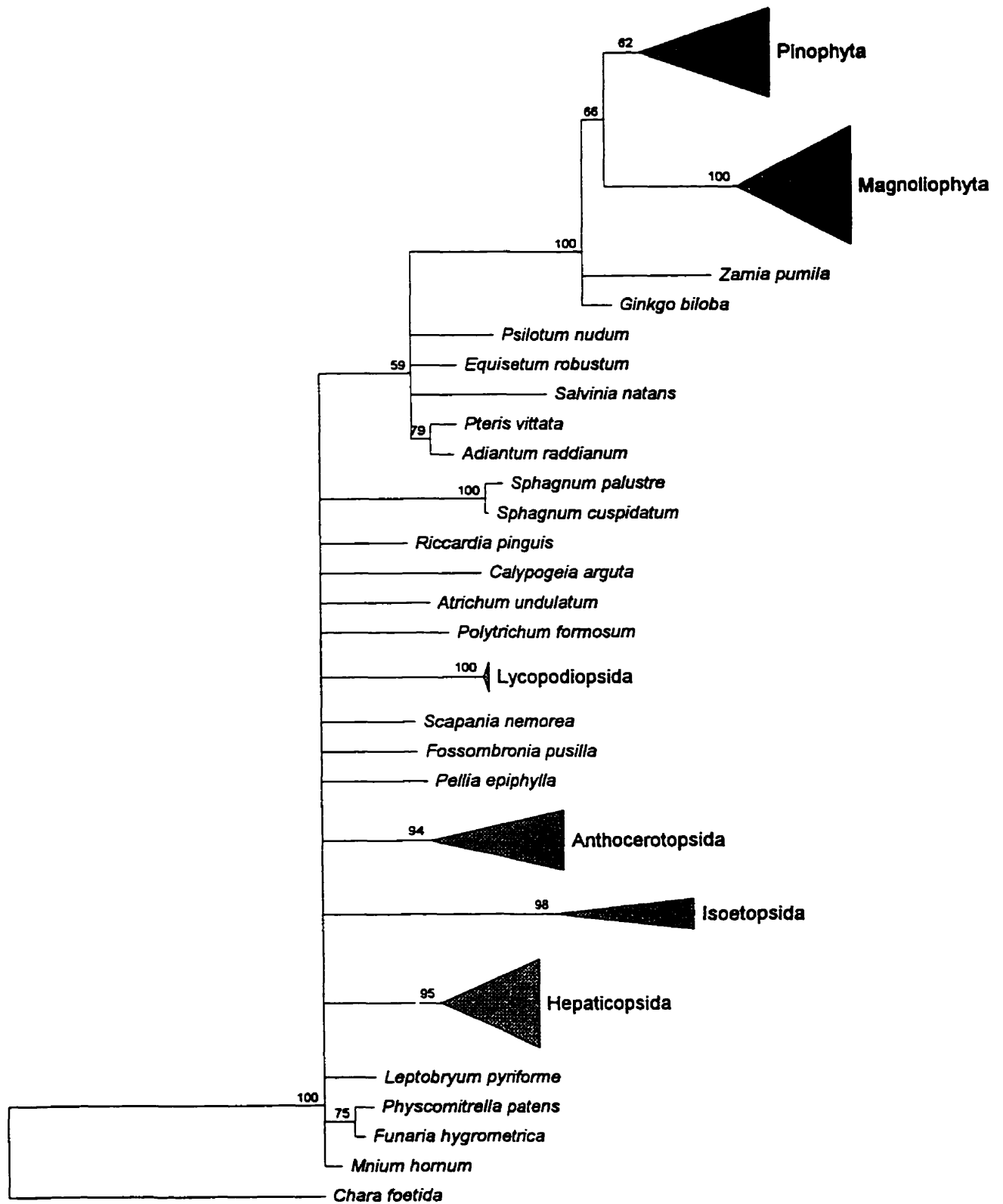


Figure G1SB Van de Peer & De Wachter, Neighbor-joining
50% Majority-Rule Bootstrap Tree

Distance 0.1

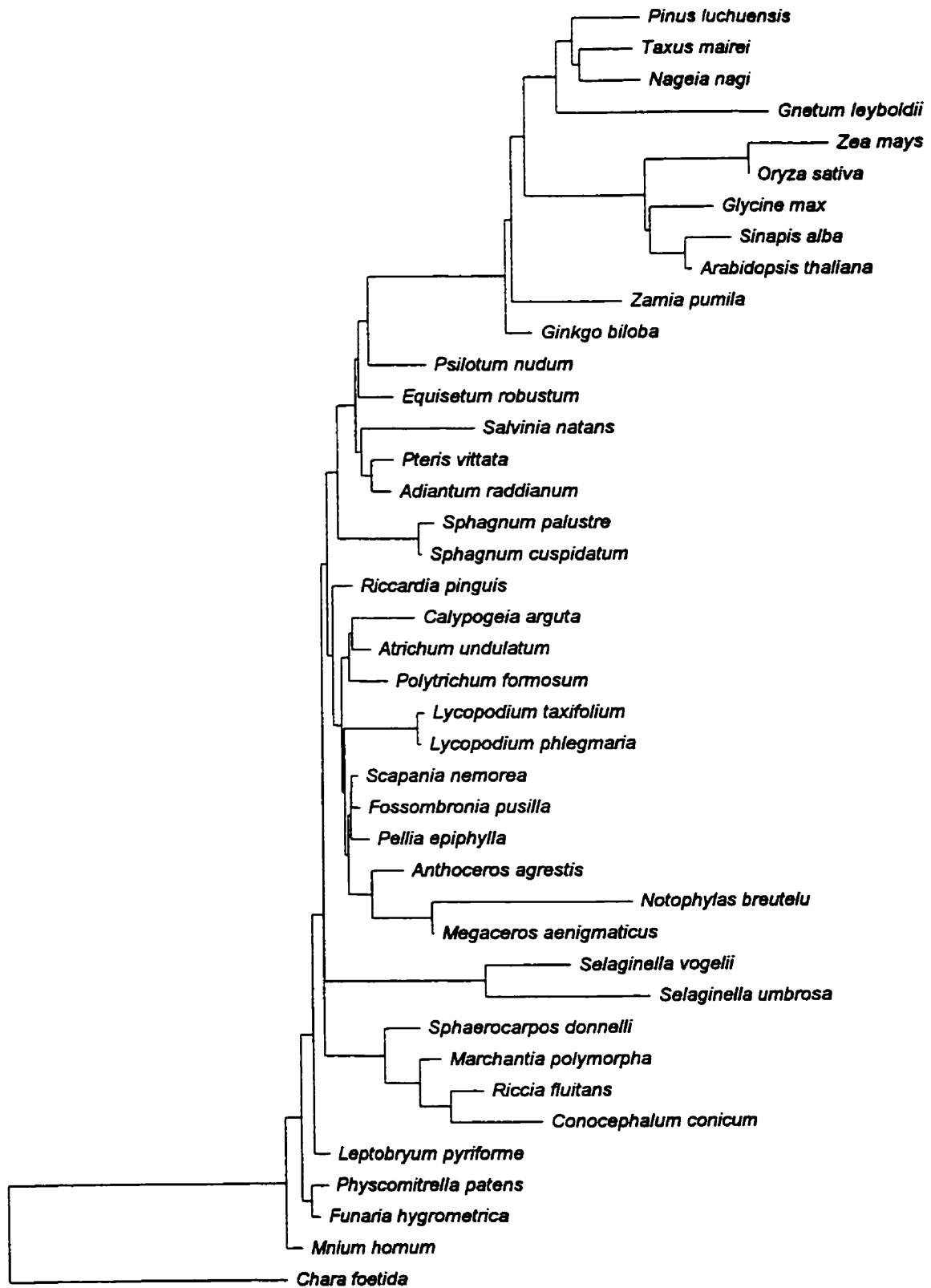


Figure G2S Van de Peer & De Wachter, Insertions & Deletions taken into account

Distance 0.1

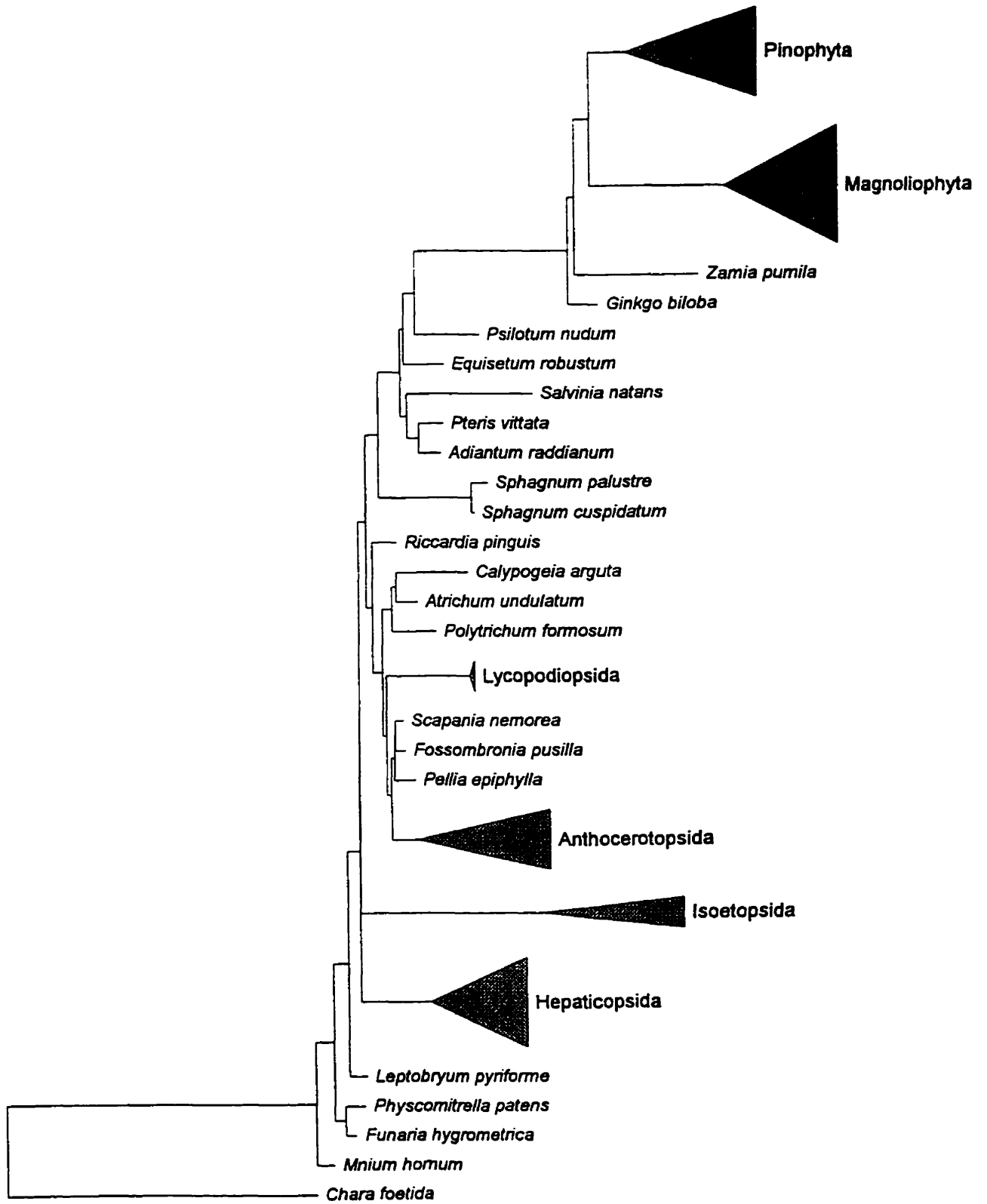


Figure G2S Van de Peer & De Wachter, Insertions & Deletions taken into account

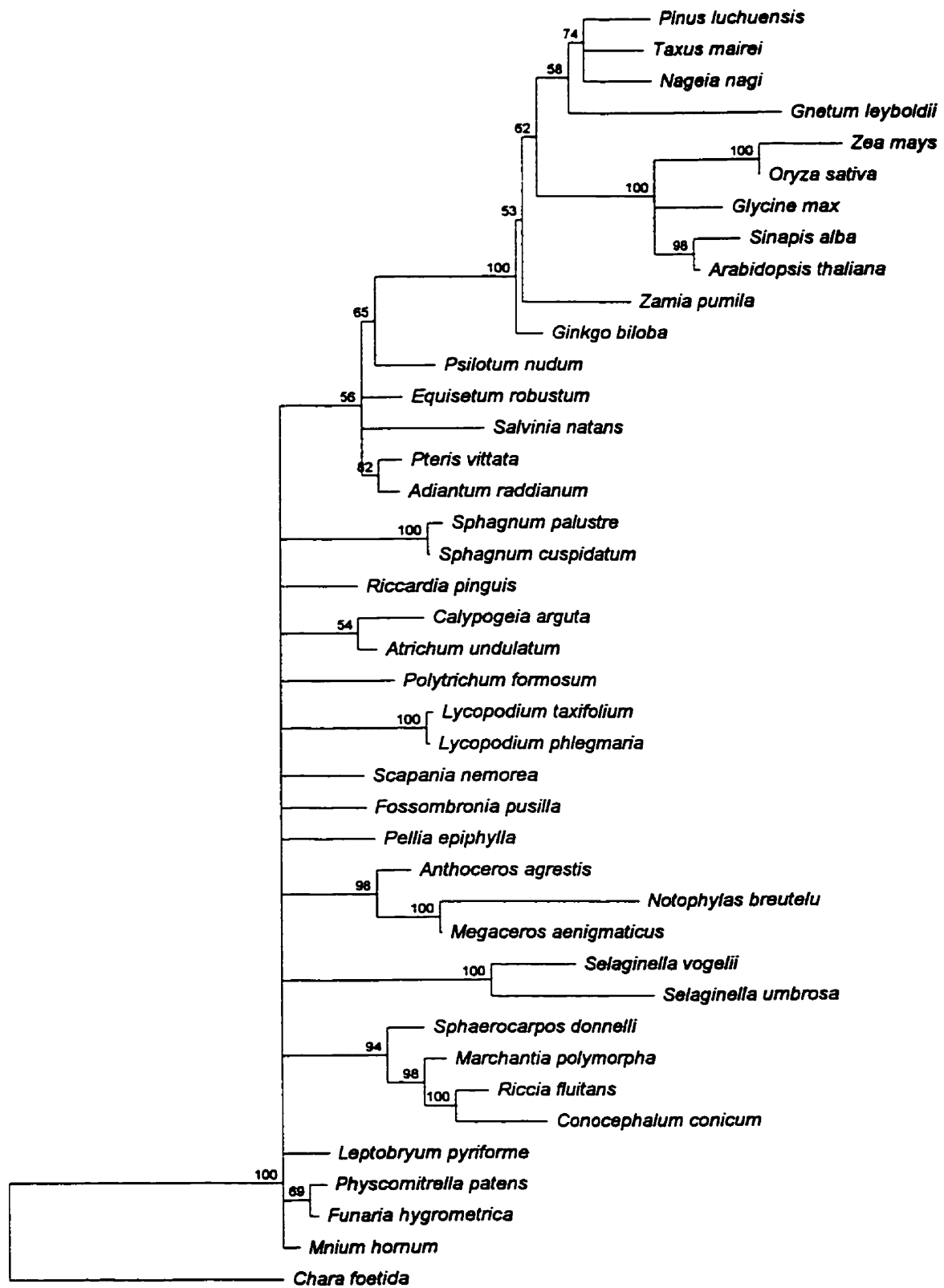


Figure G2SB Van de Peer & De Wachter, Insertions & Deletions taken into account
50% Majority-Rule Bootstrap Tree

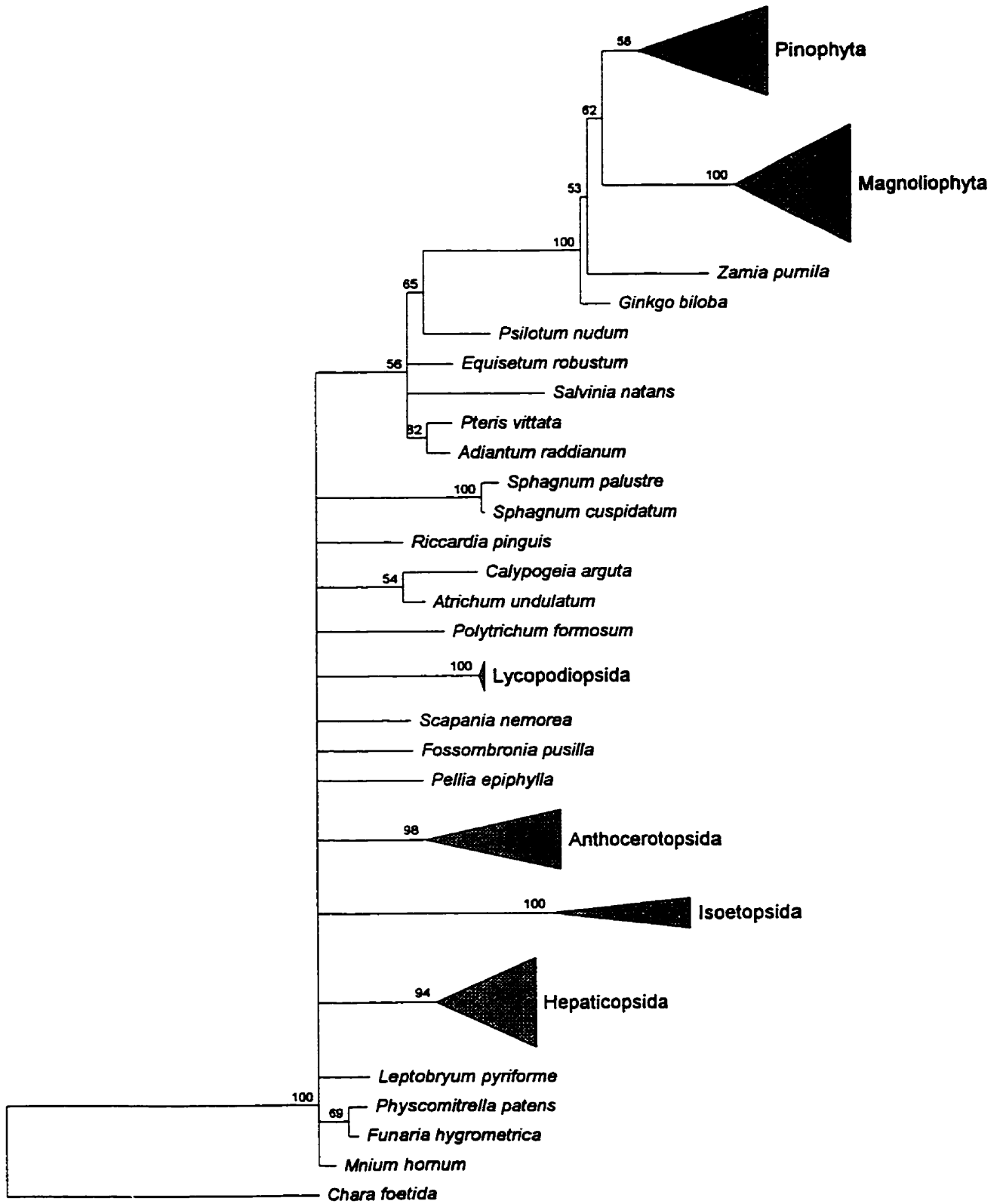


Figure G2SB Van de Peer & De Wachter, Insertions & Deletions taken into account
50% Majority-Rule Bootstrap Tree

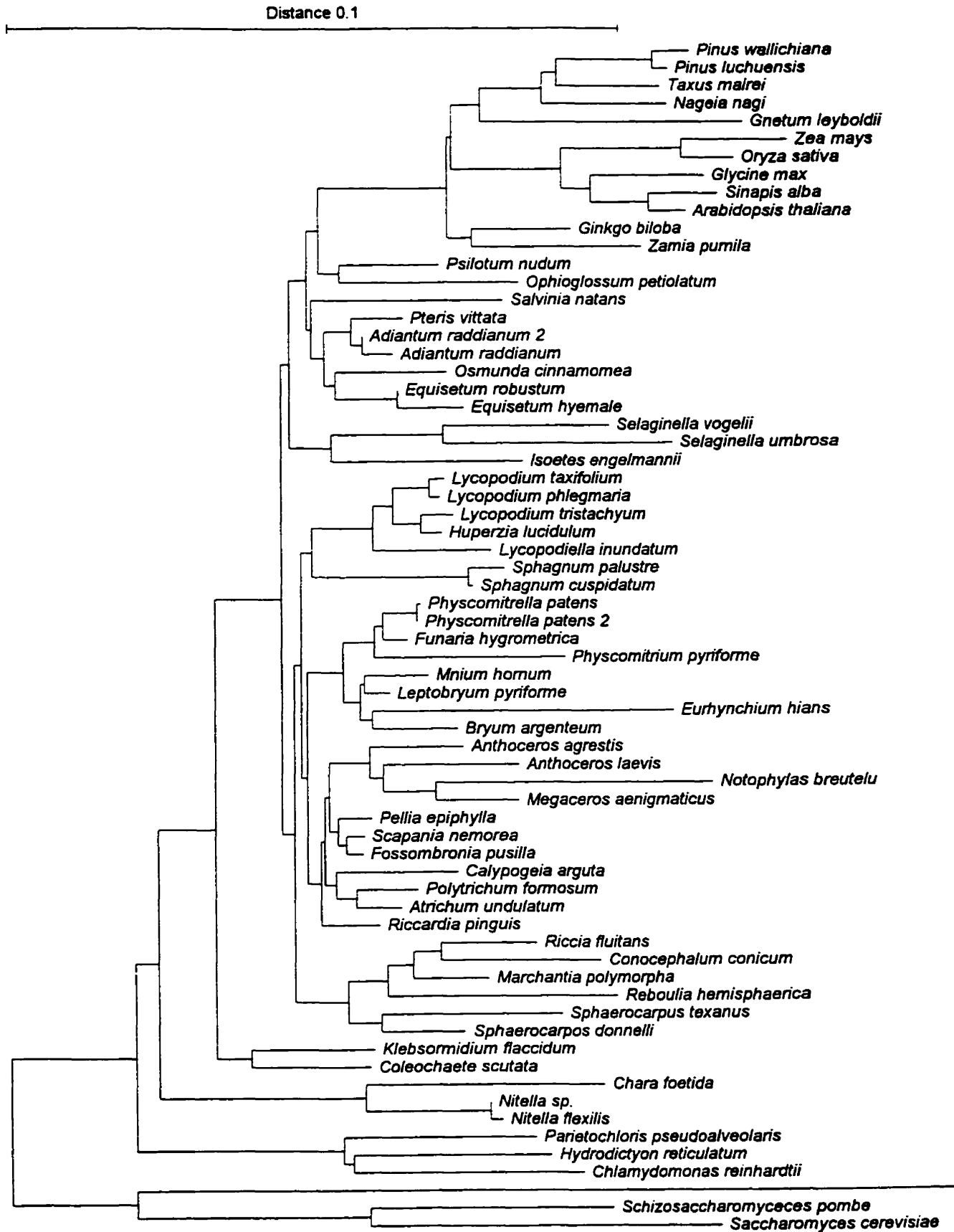


Figure A1 Jukes & Cantor, Neighbor-joining

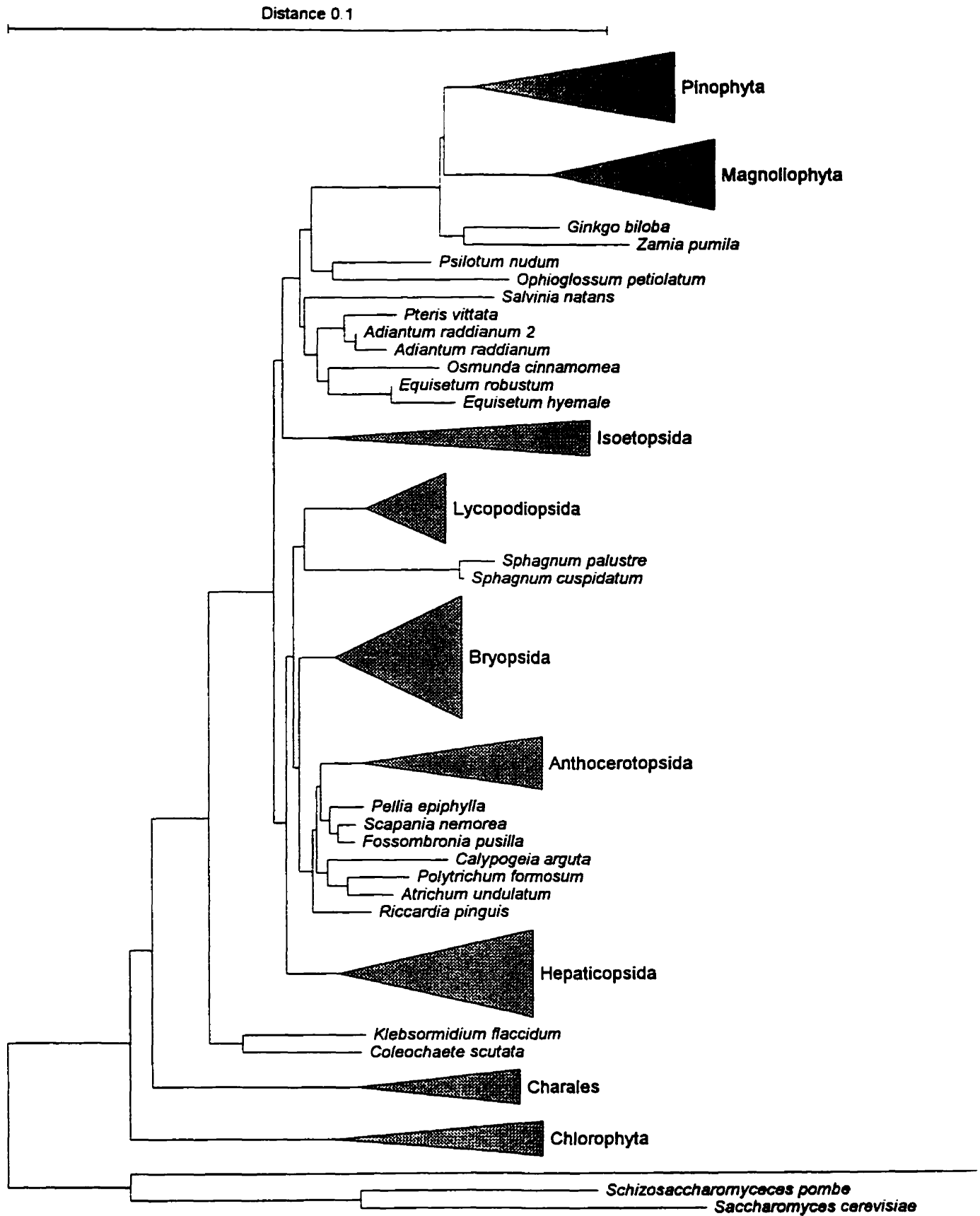


Figure A1 Jukes & Cantor, Neighbor-joining

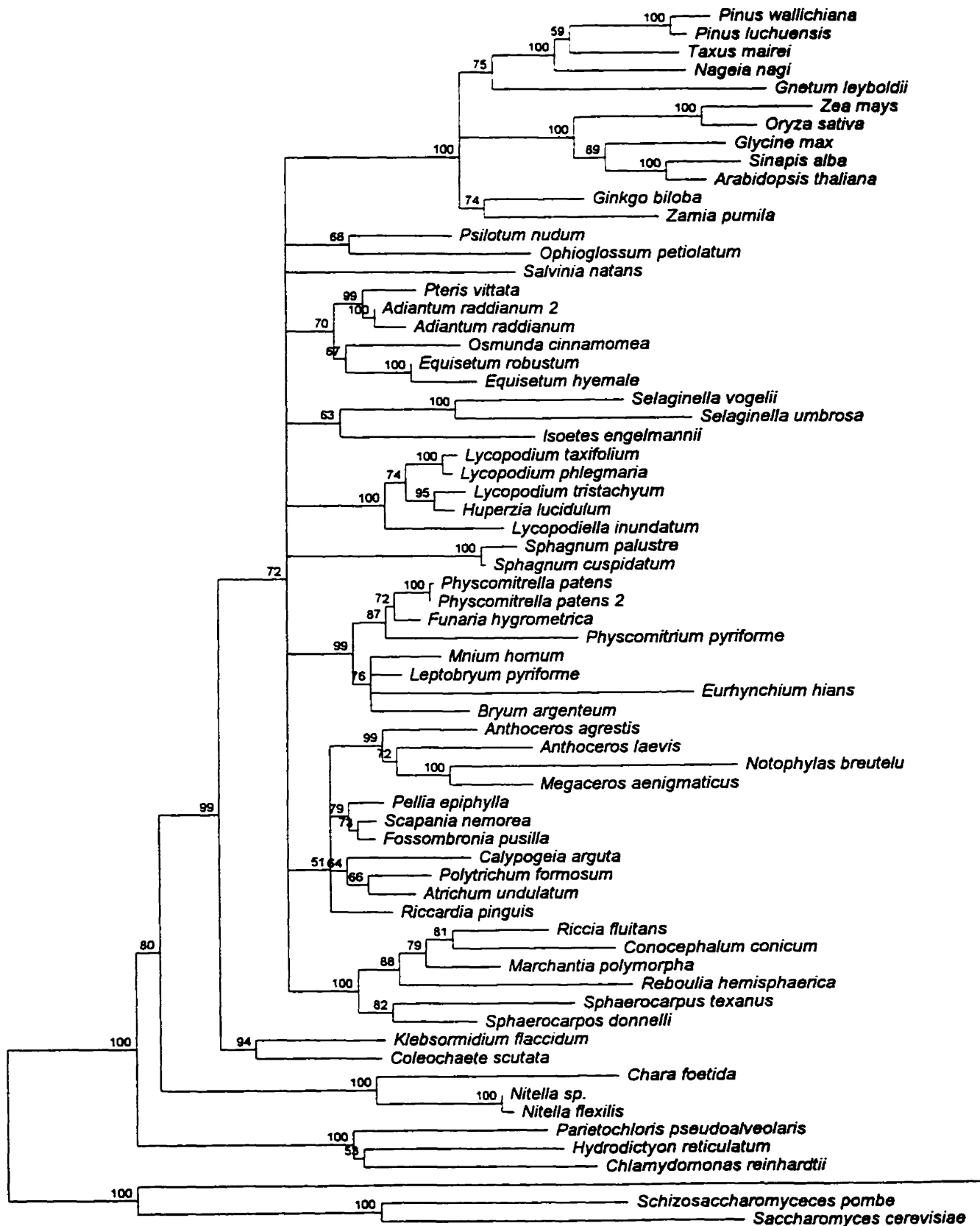


Figure A1B Jukes & Cantor, 50% Majority-Rule Bootstrap Tree

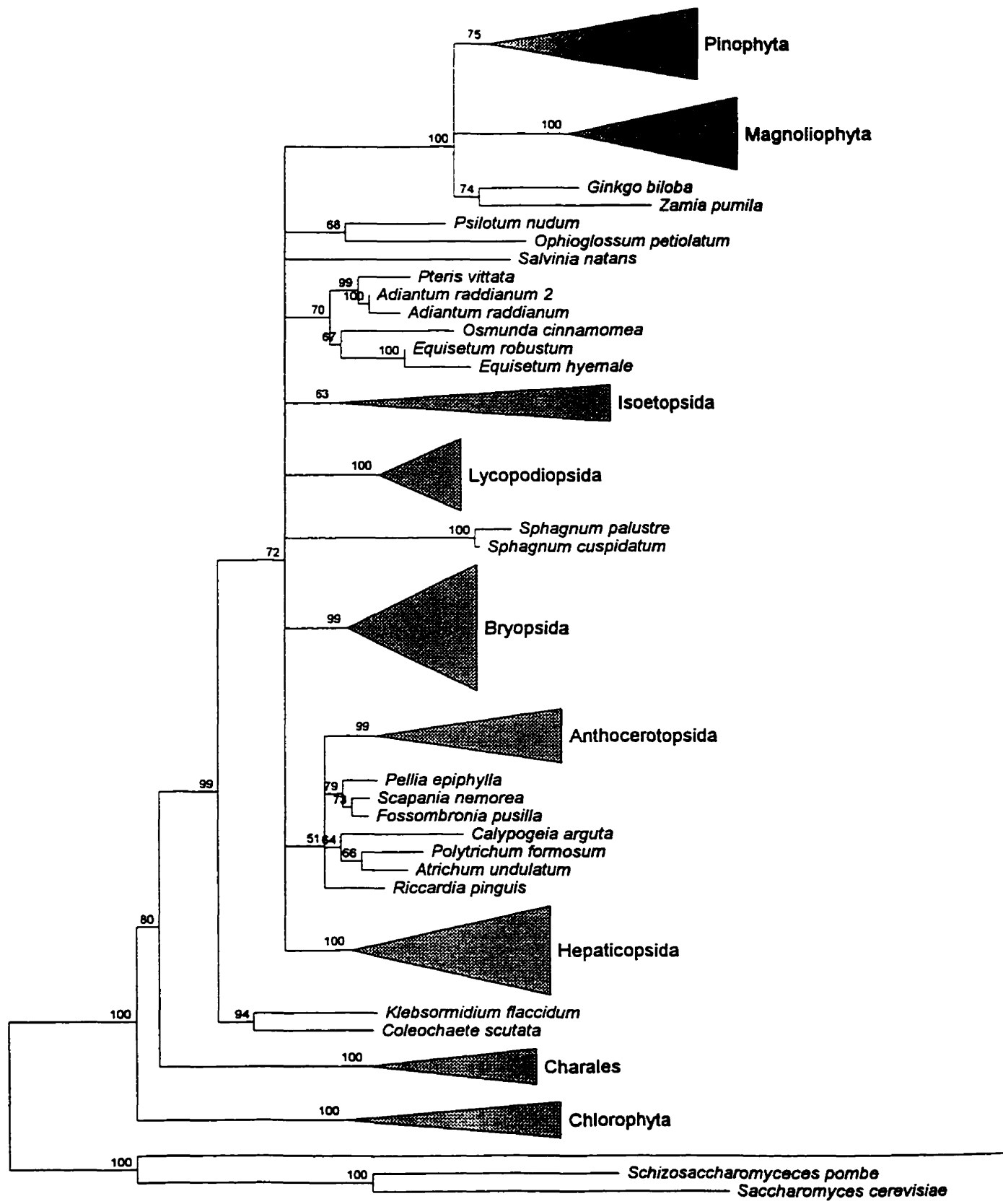


Figure A1B Jukes & Cantor, 50% Majority-Rule Bootstrap Tree

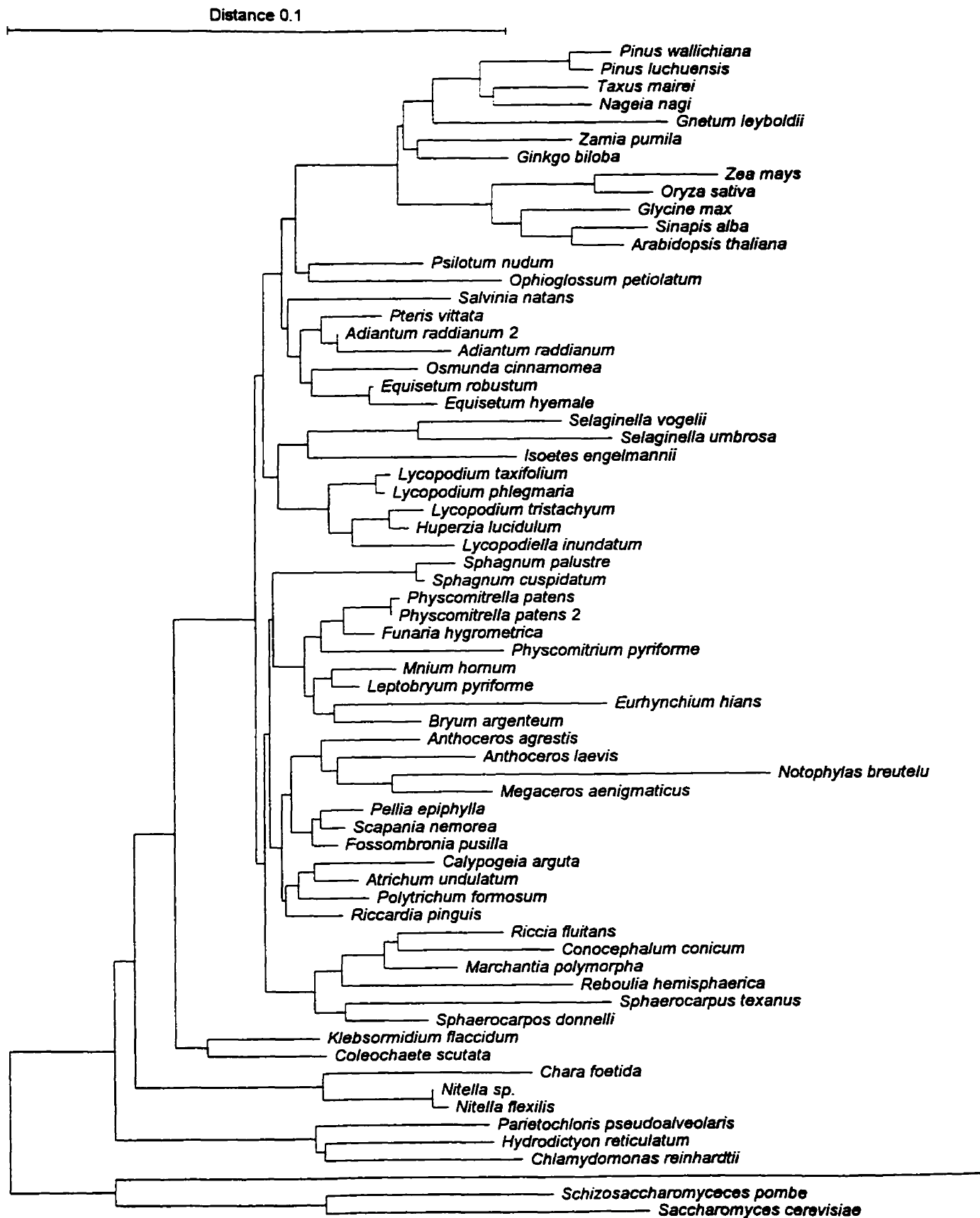


Figure A2 Jukes & Cantor, Neighbor-joining (Insertions & Deletions taken into account)

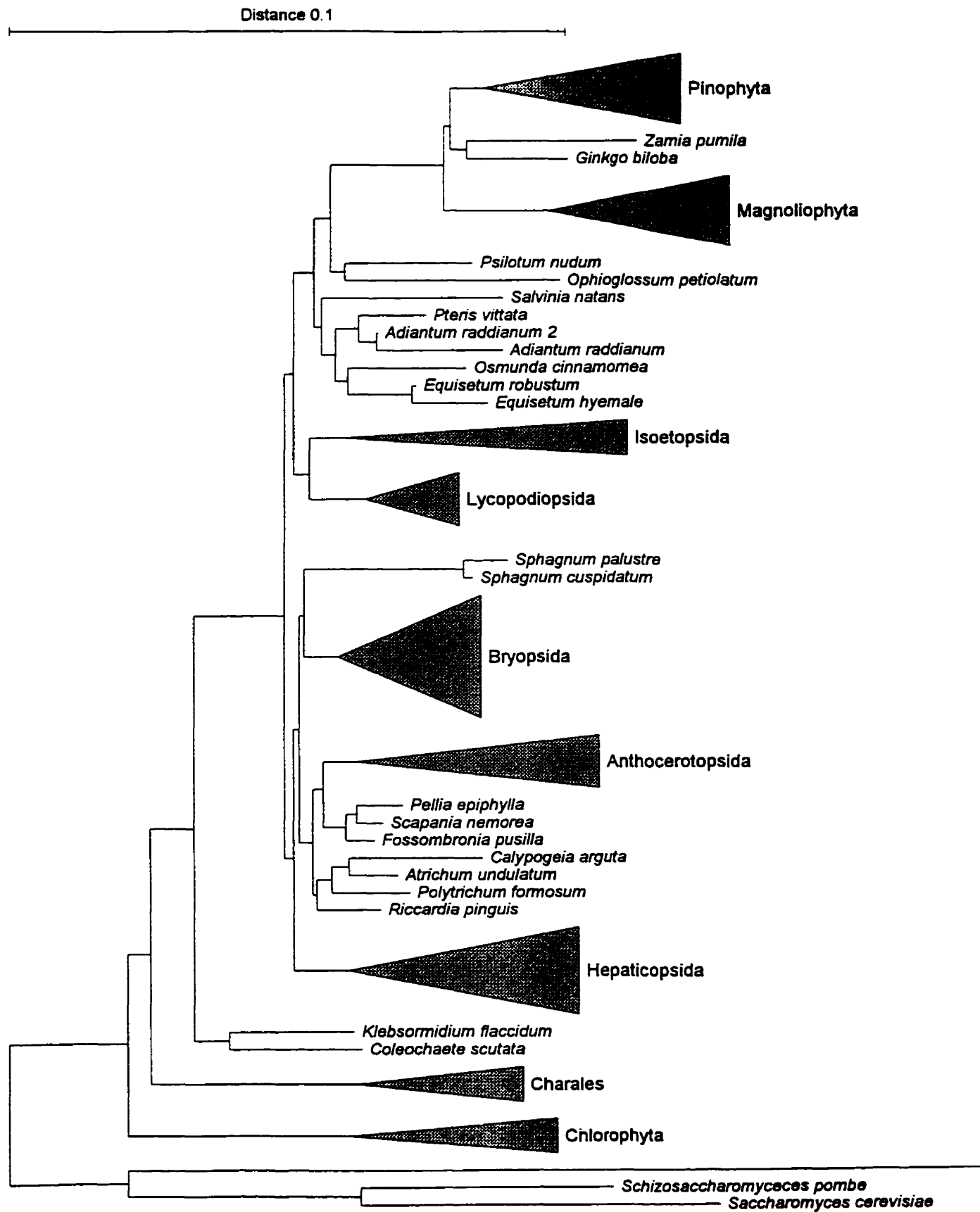


Figure A2 Jukes & Cantor, Neighbor-joining (Insertions & Deletions taken into account)

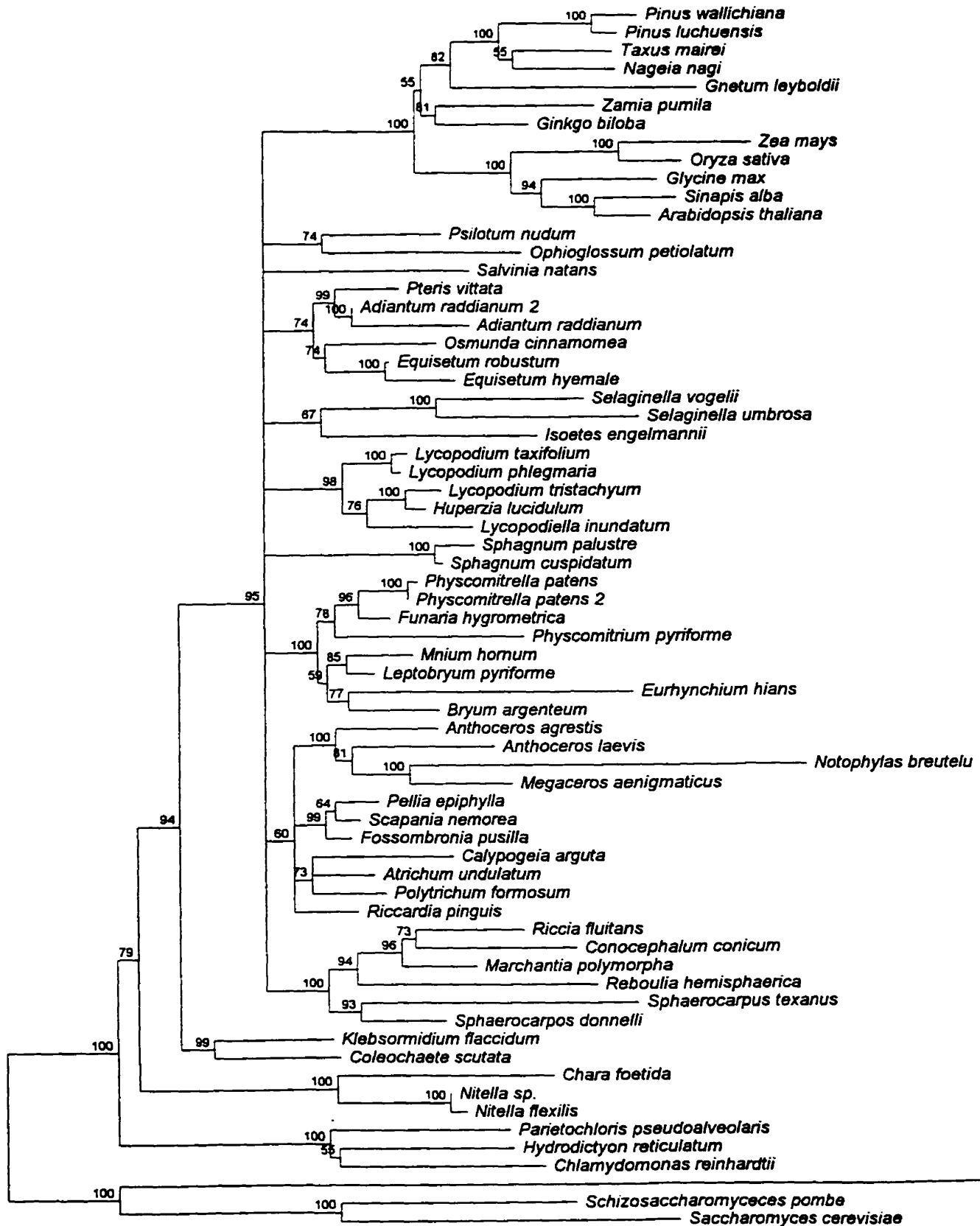
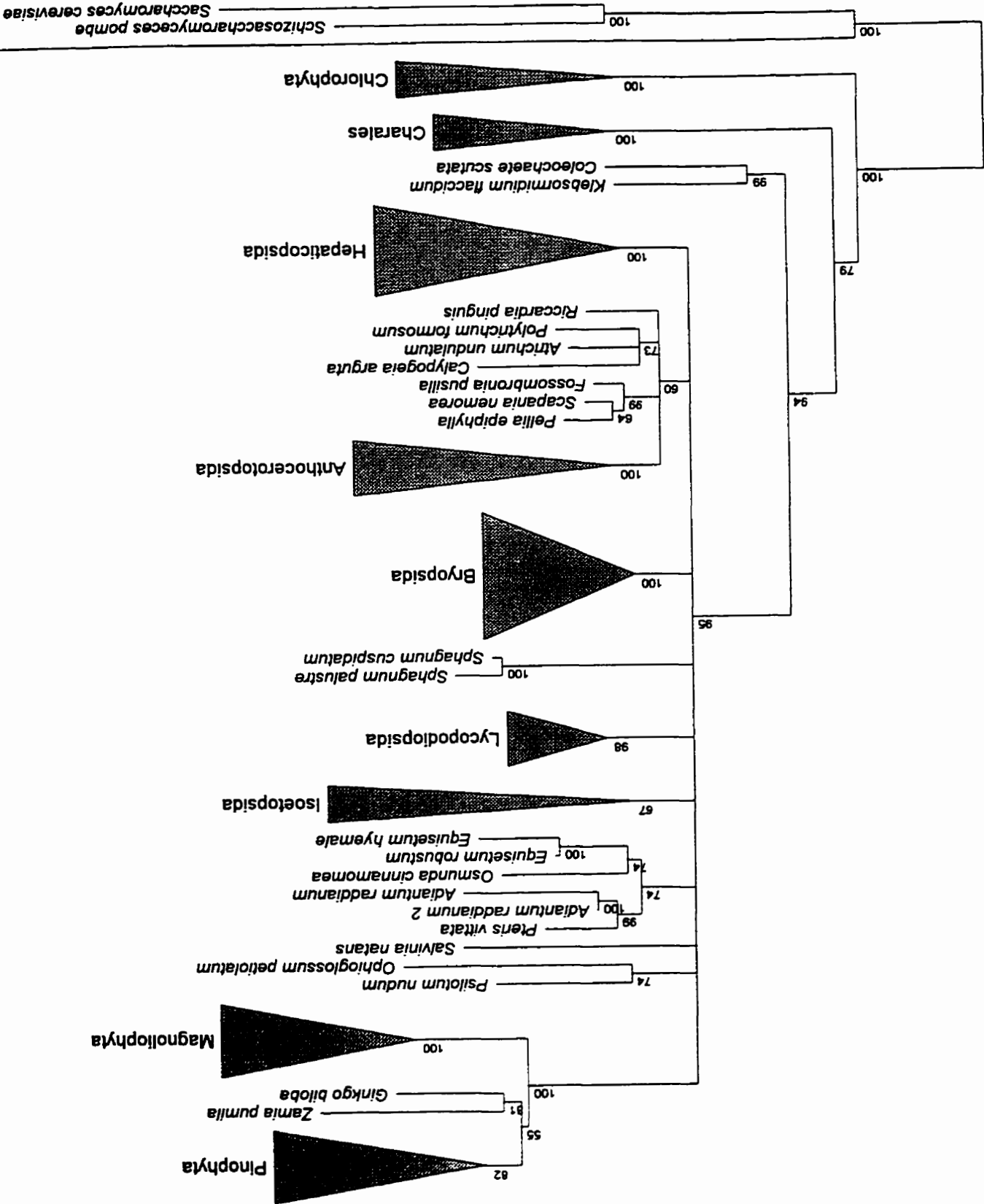


Figure A2B Jukes & Cantor, (Insertions & Deletions taken into account)
50% Majority-Rule Bootstrap Tree

Figure A2B Jukes & Cantor, (Insertions & Deletions taken into account) 50% Majority-Rule Bootstrap Tree



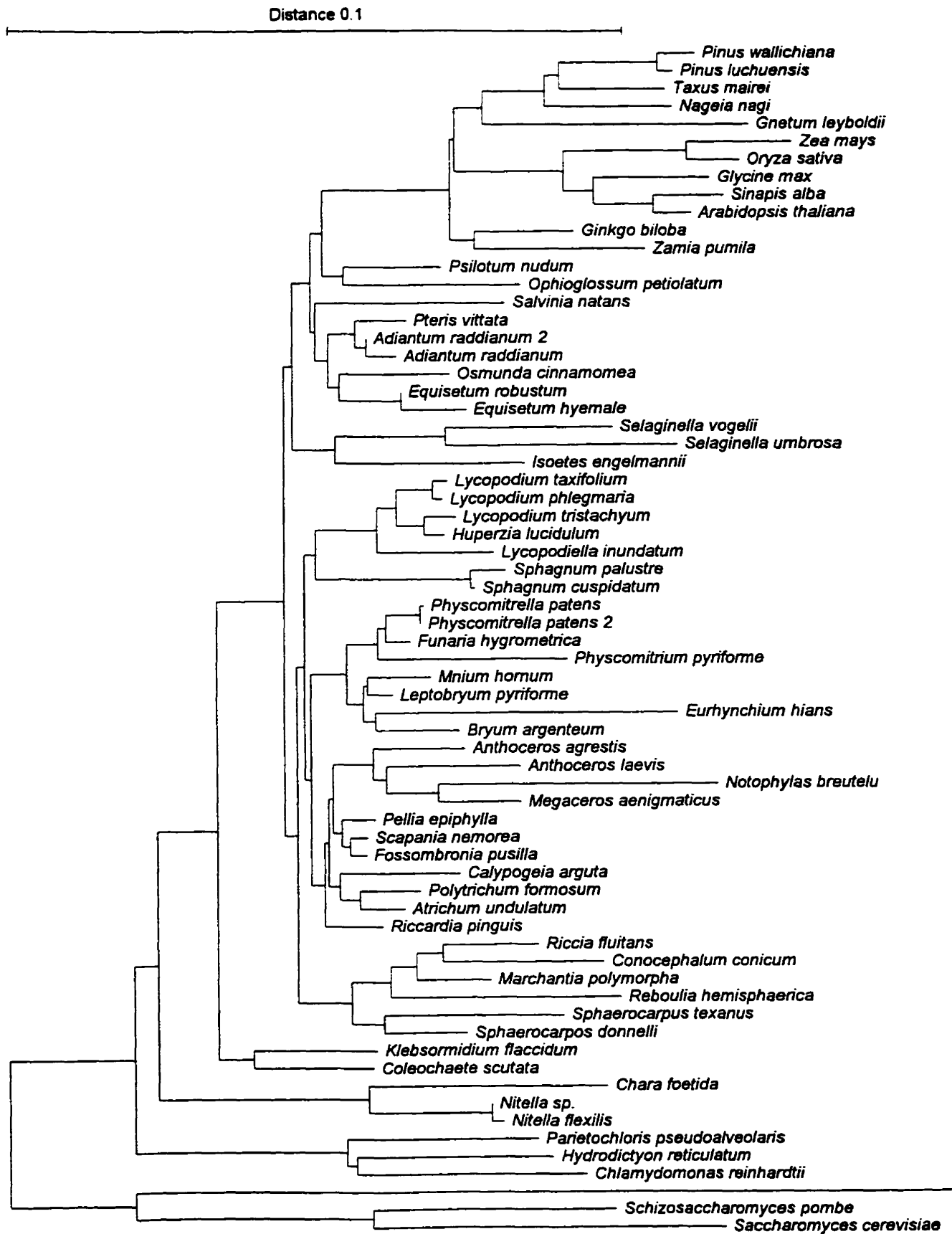


Figure B1 Tajima & Nei; Neighbor-joining

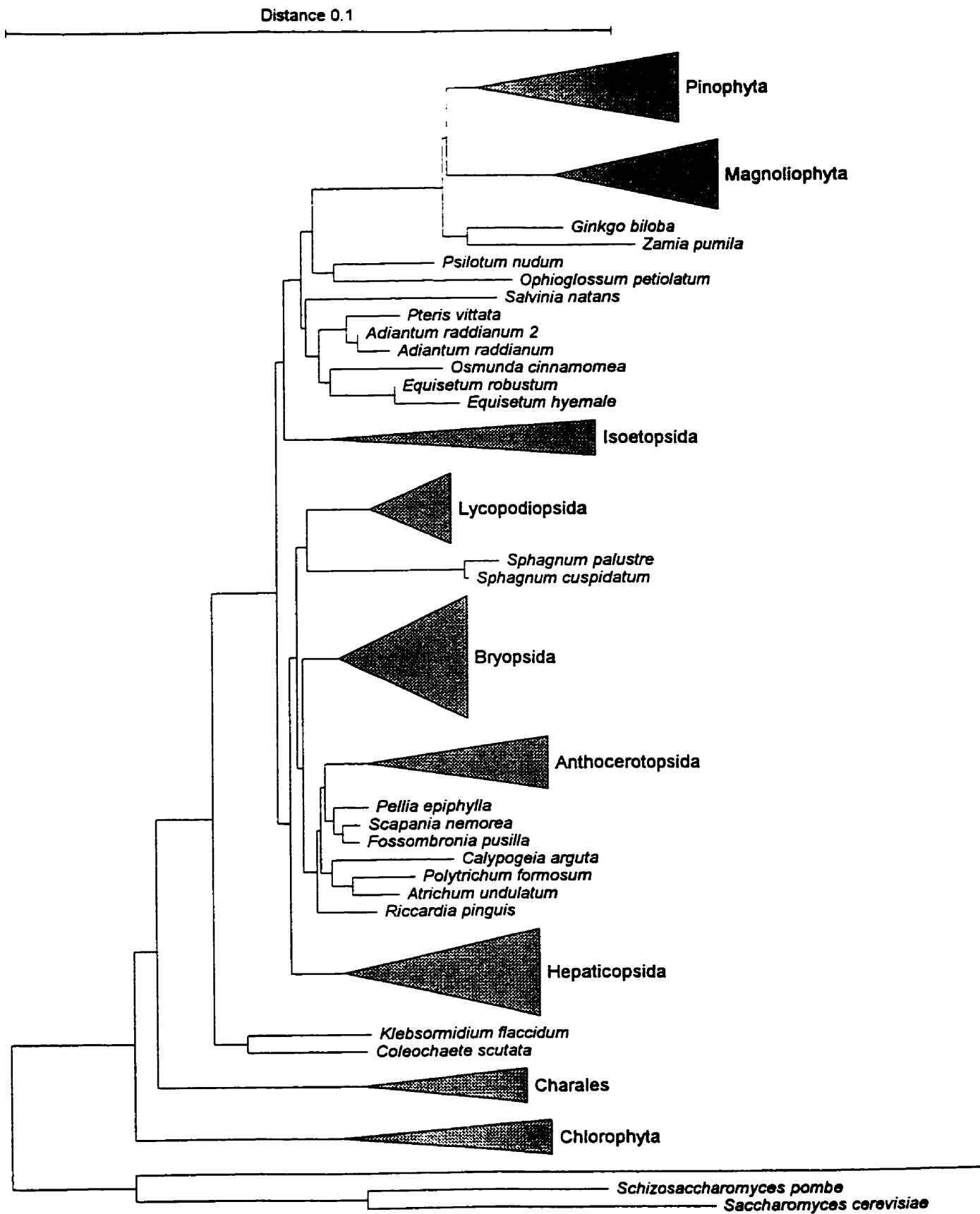


Figure B1 Tajima & Nei; Neighbor-joining

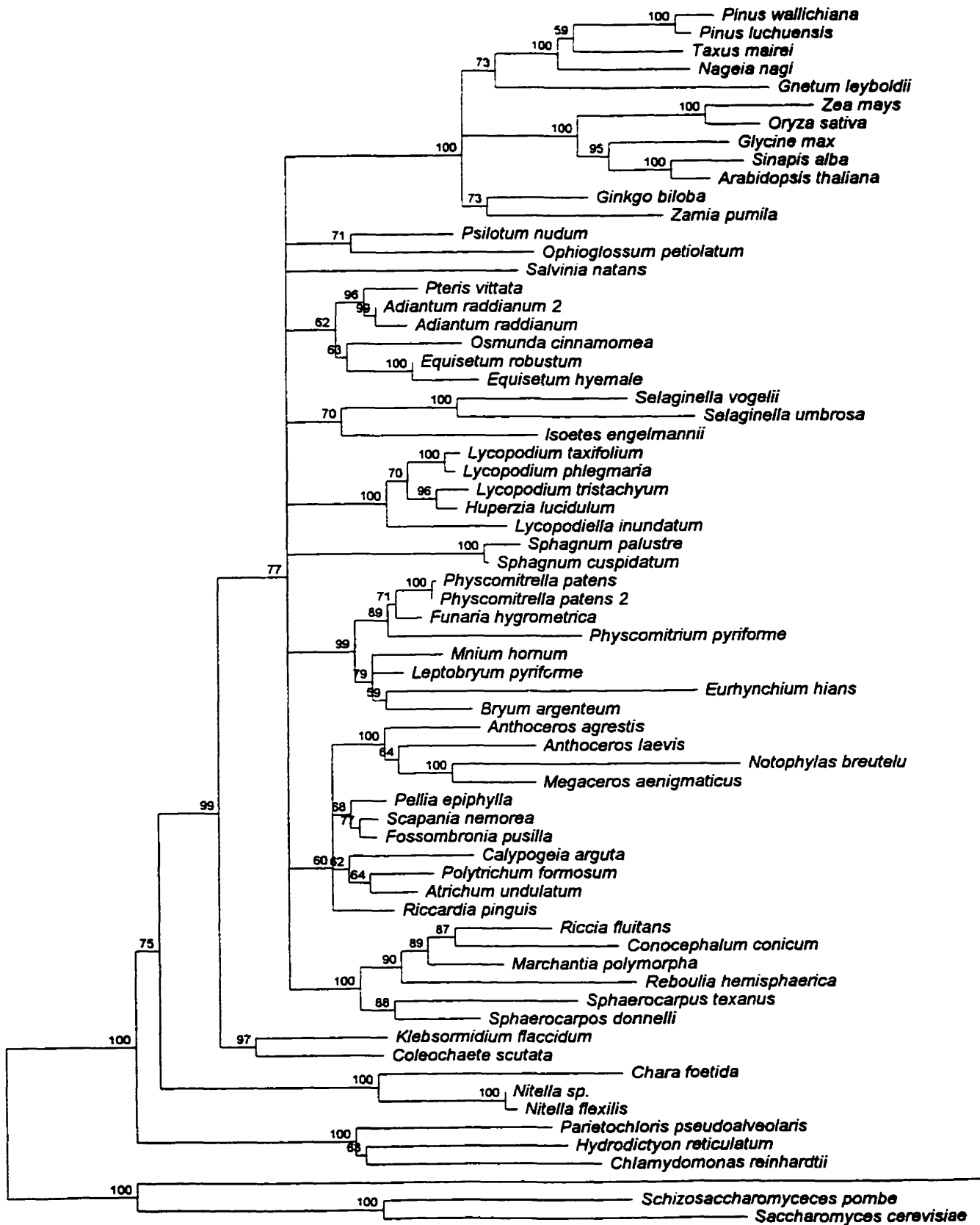


Figure B1B Tajima & Nei; 50% Majority-Rule Bootstrap Tree

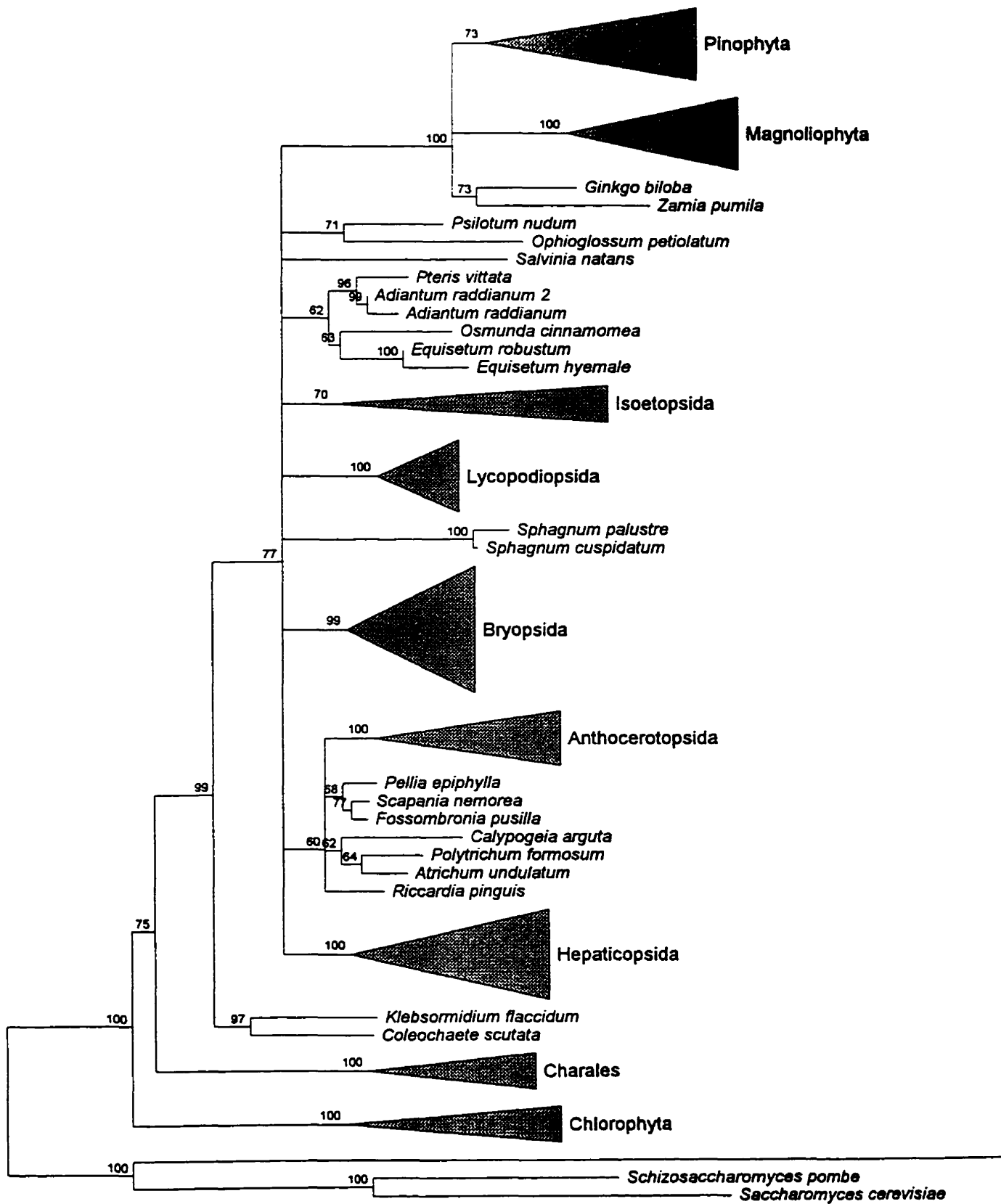


Figure B1B Tajima & Nei; 50% Majority-Rule Bootstrap Tree

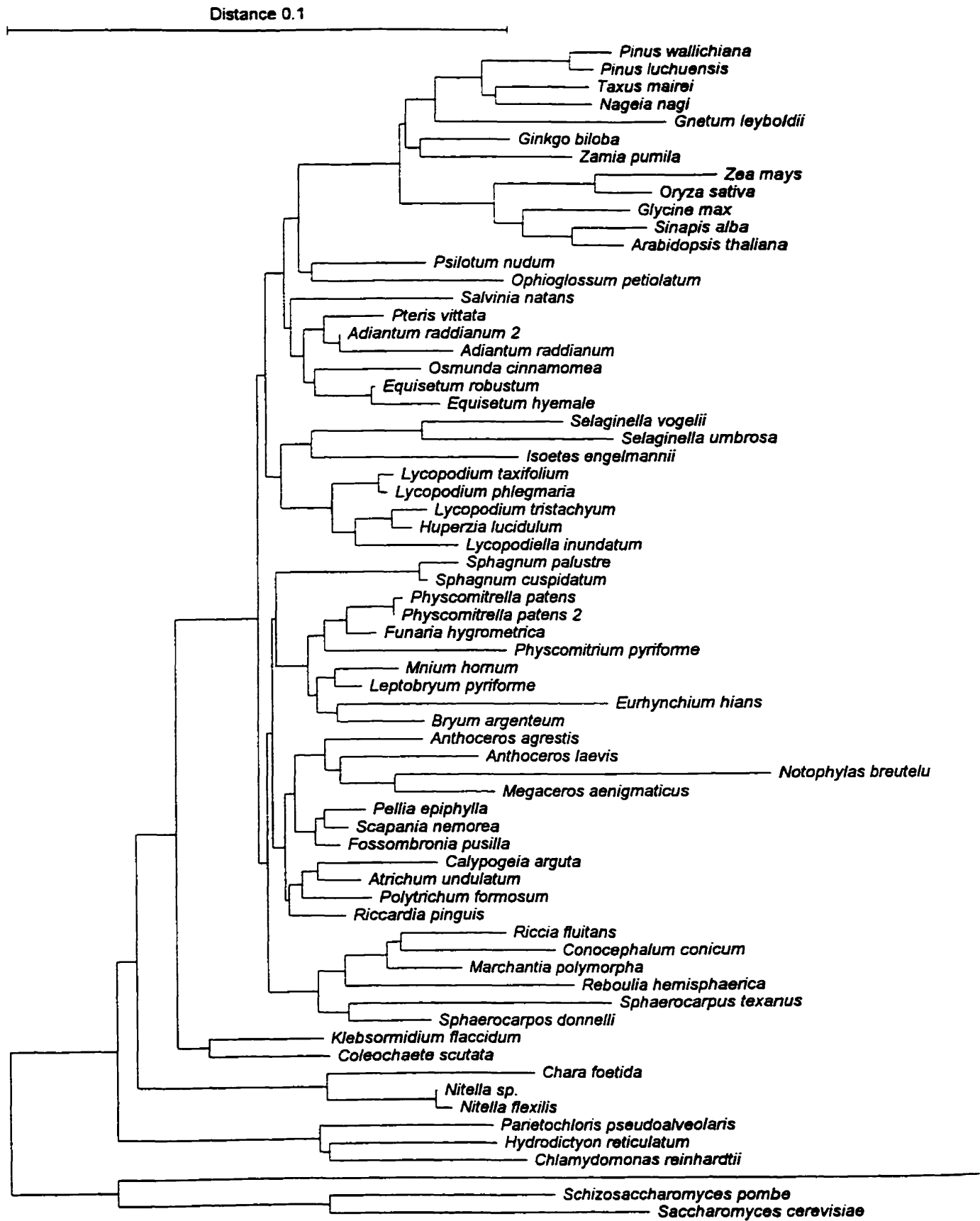


Figure B2 Tajima & Nei; Neighbor-joining (Insertions & Deletions taken into account)

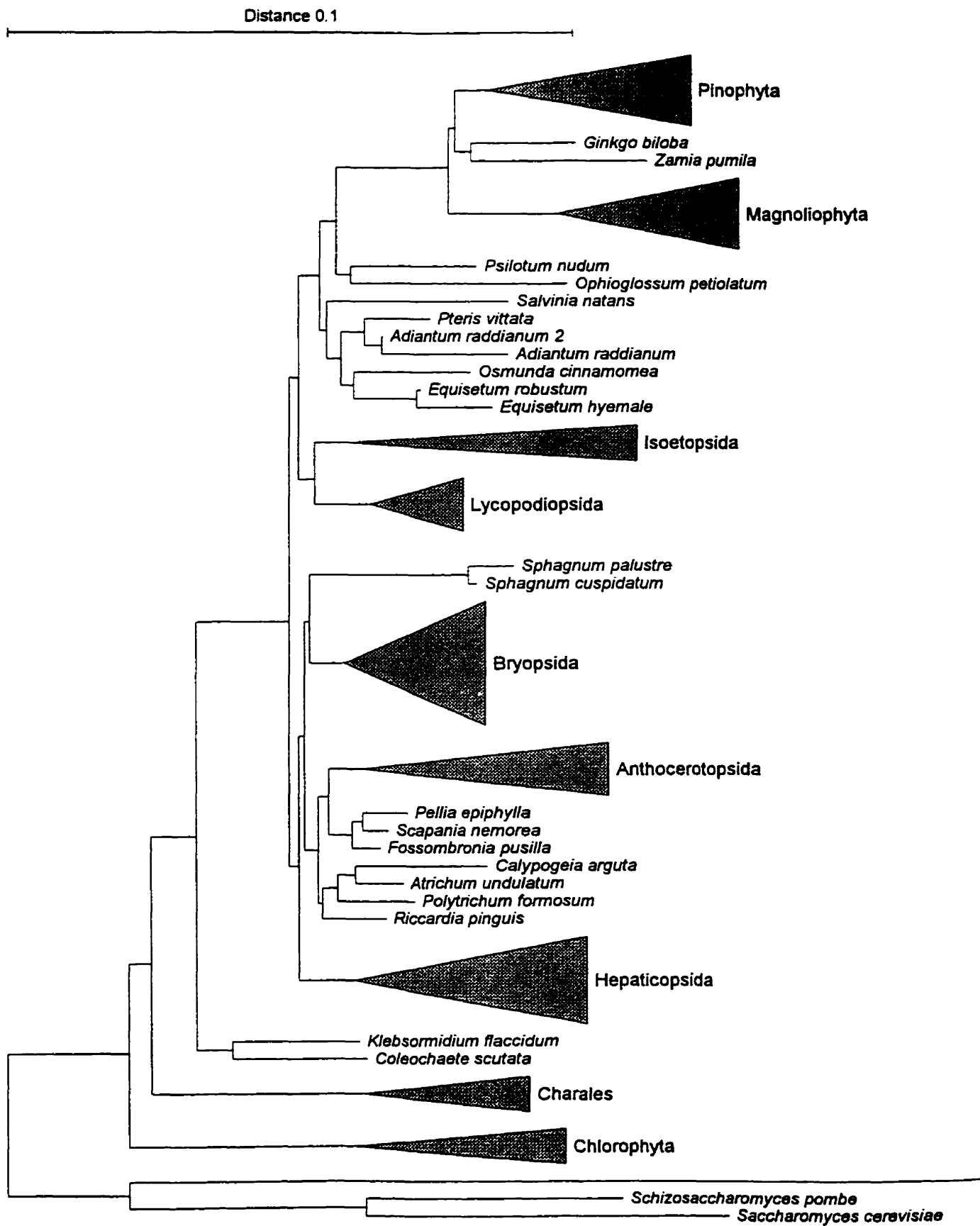


Figure B2 Tajima & Nei; Neighbor-joining (Insertions & Deletions taken into account)

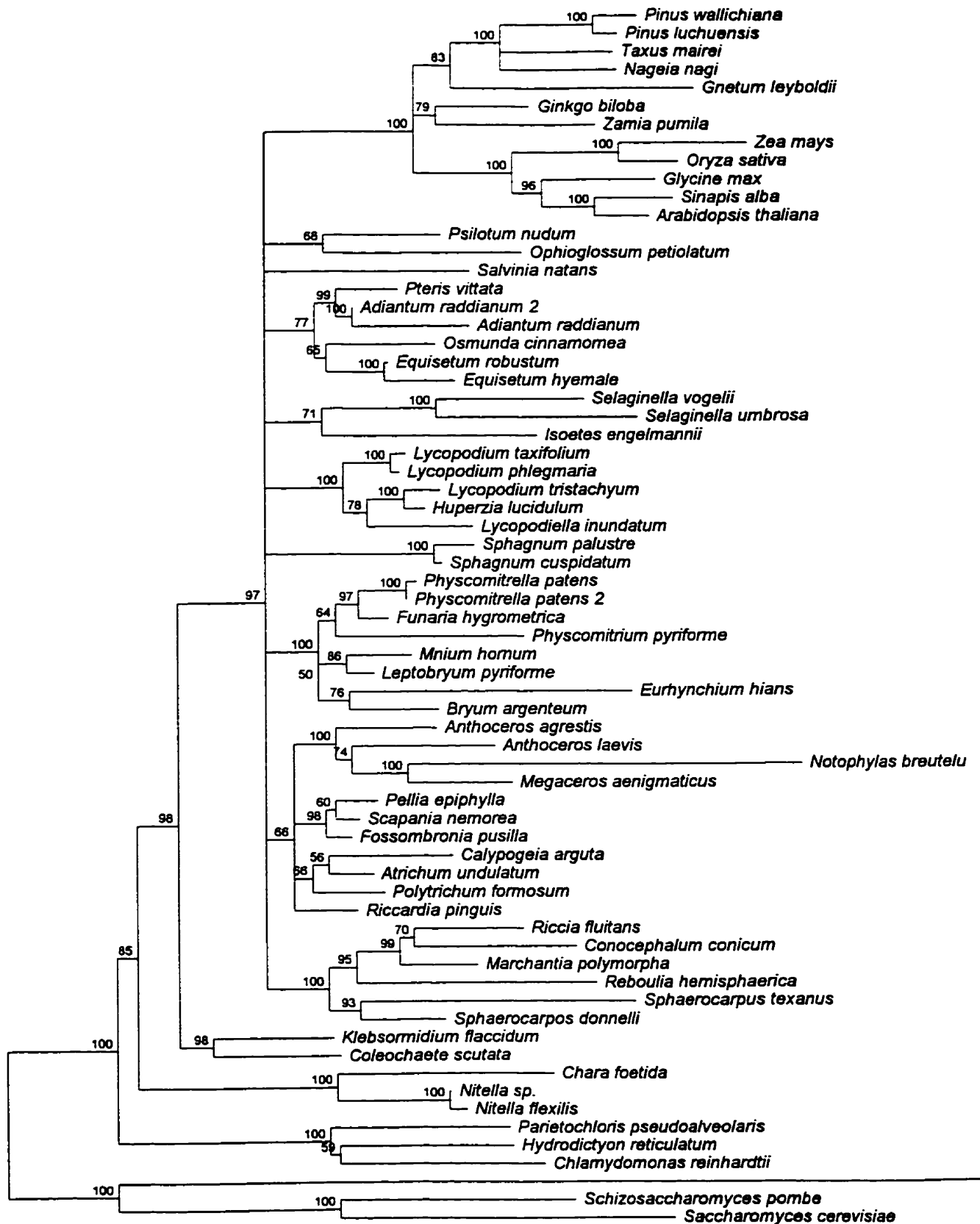


Figure B2B Tajima & Nei; (Insertions and Deletions taken into account)
50% Majority-Rule Bootstrap Tree

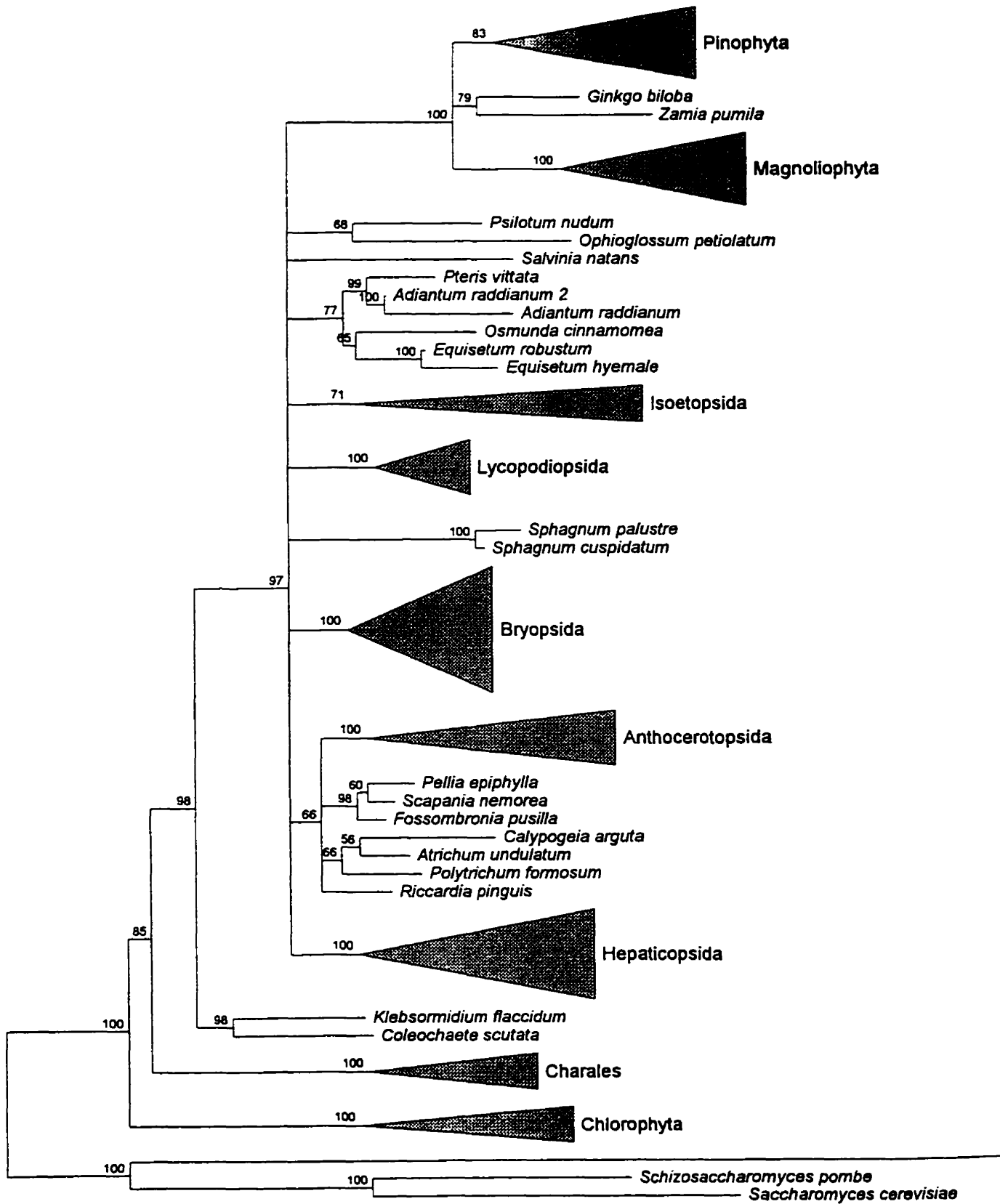


Figure B2B Tajima & Nei; (Insertions and Deletions taken into account)
50% Majority-Rule Bootstrap Tree

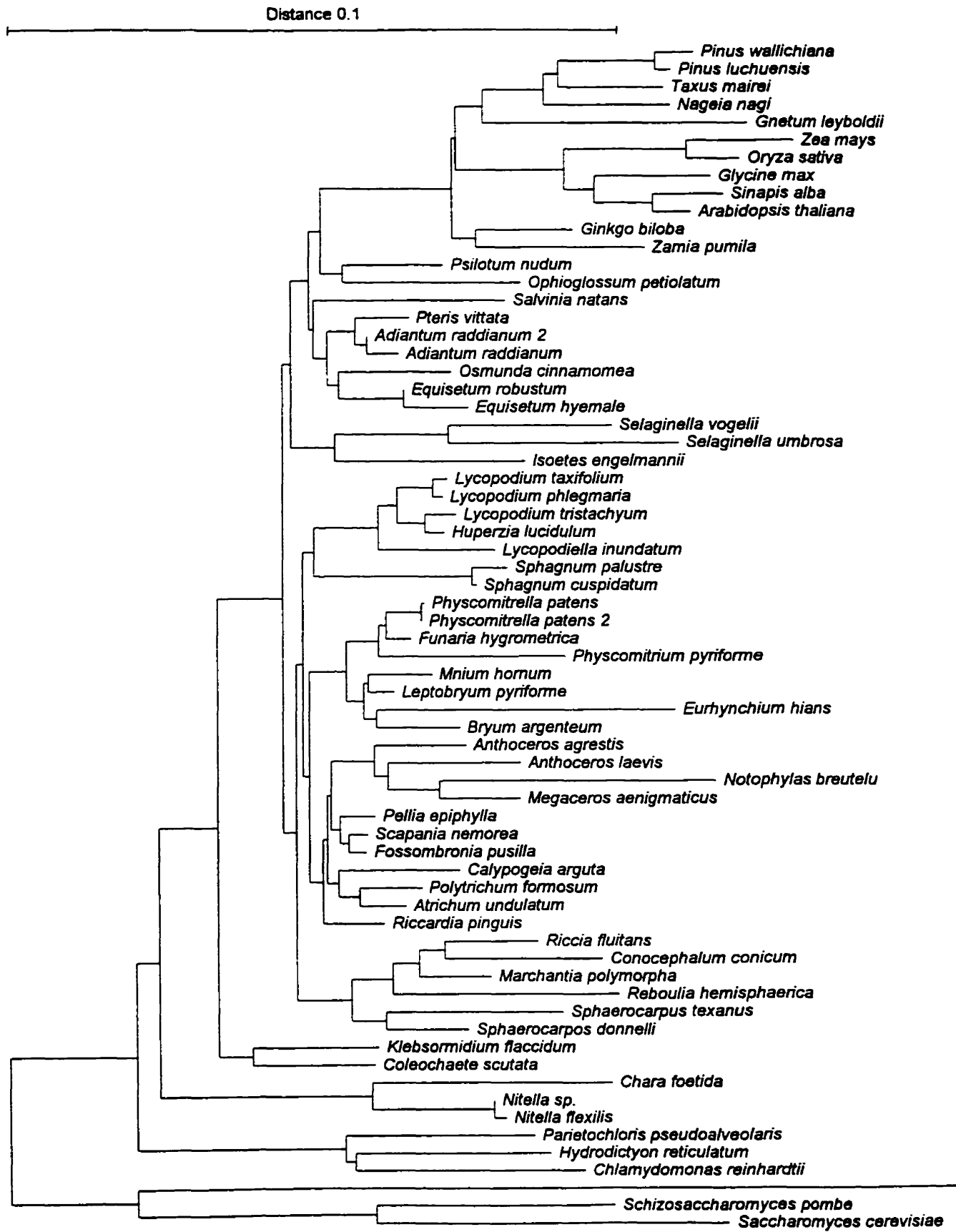


Figure C1 Kimura; Neighbor-joining

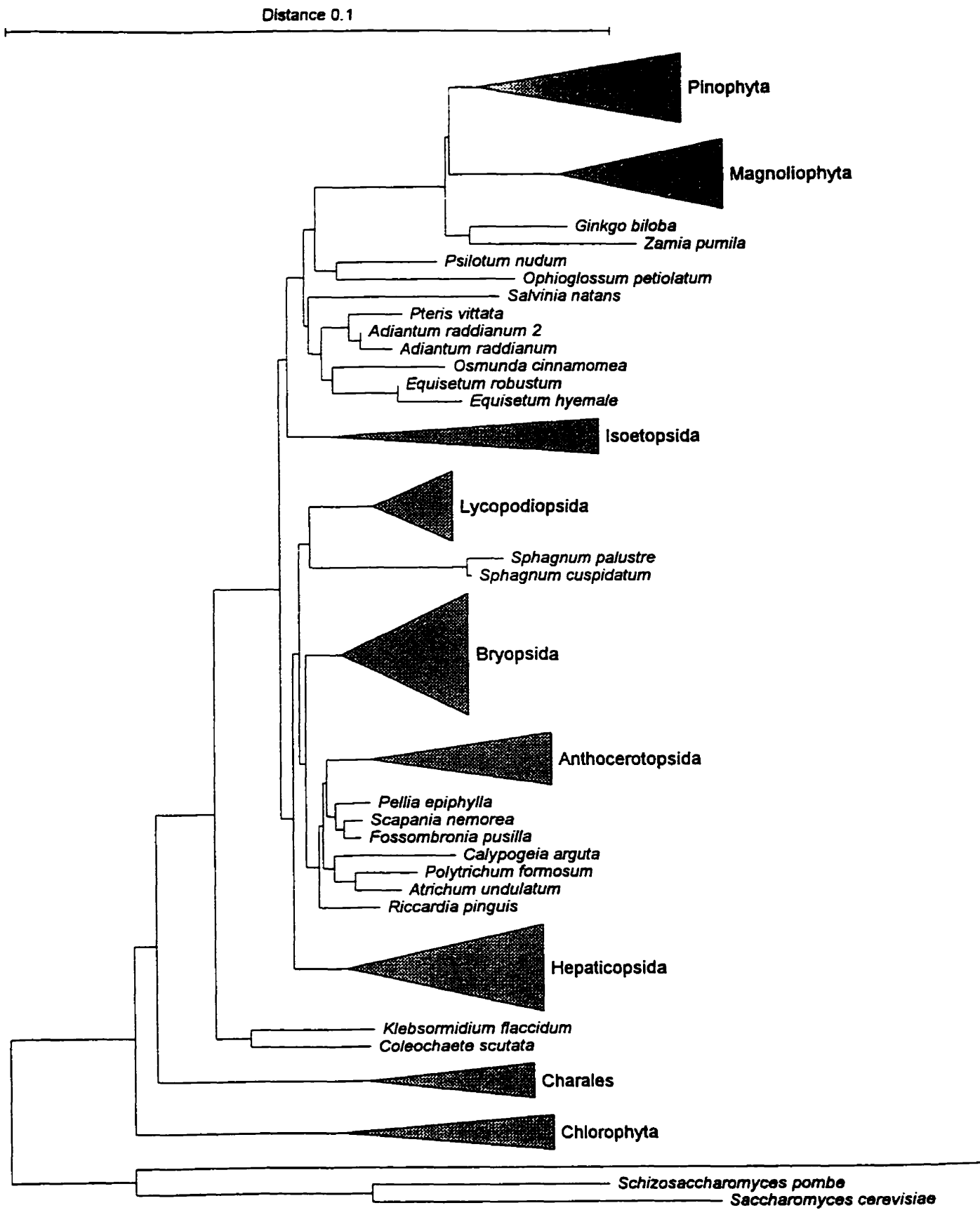


Figure C1 Kimura; Neighbor-joining

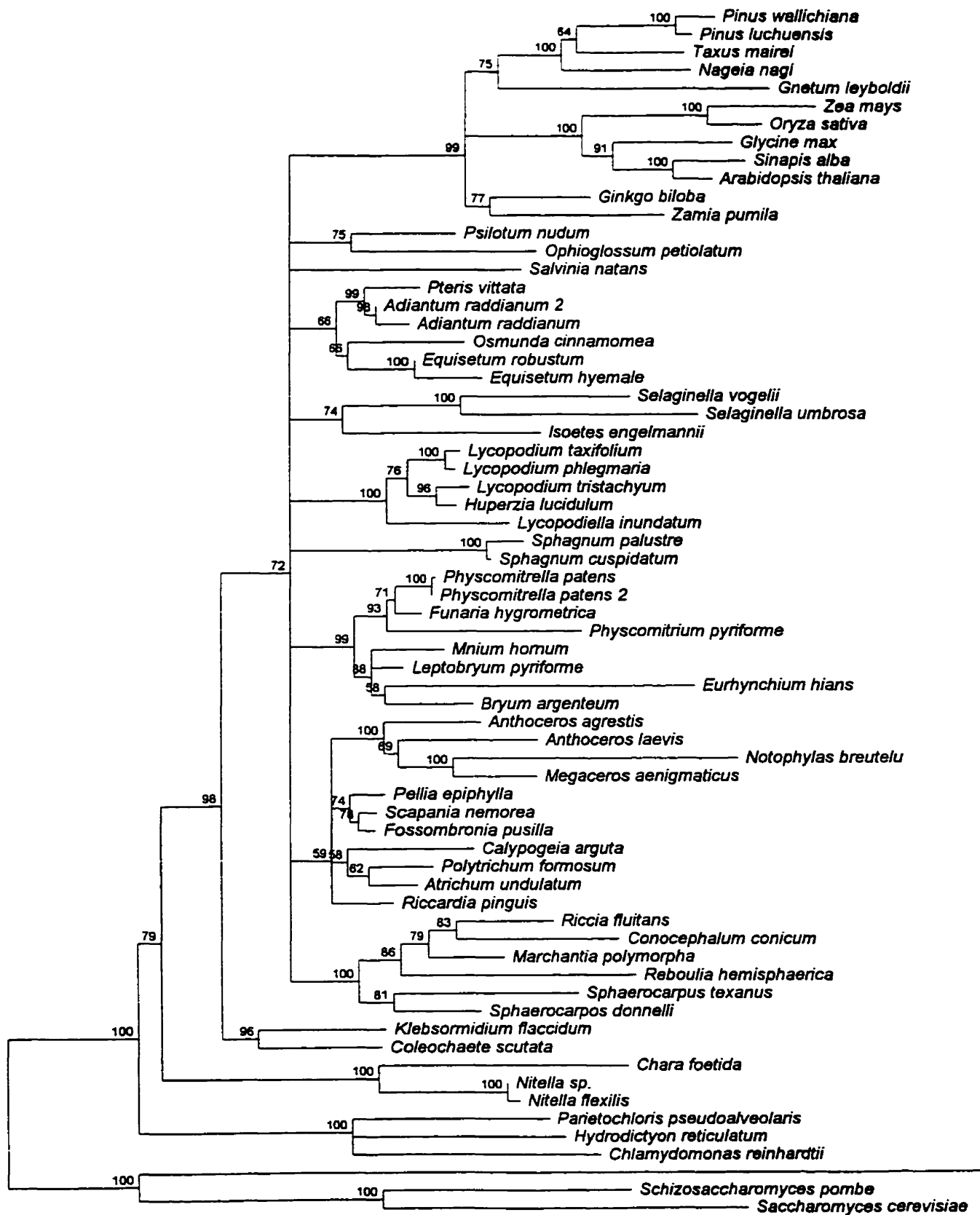


Figure C1B Kimura; 50% Majority-Rule Bootstrap Tree

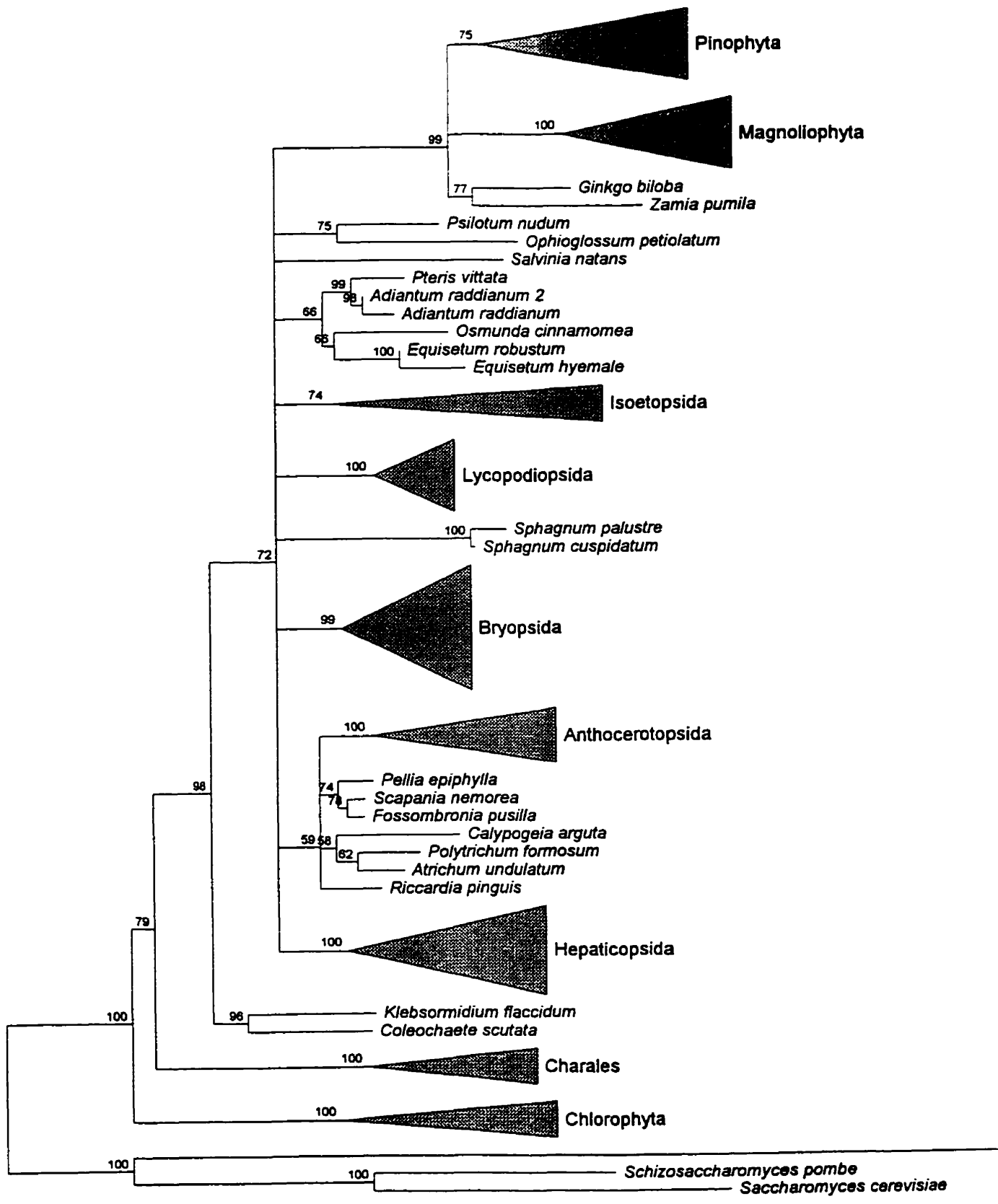


Figure C1B Kimura; 50% Majority-Rule Bootstrap Tree

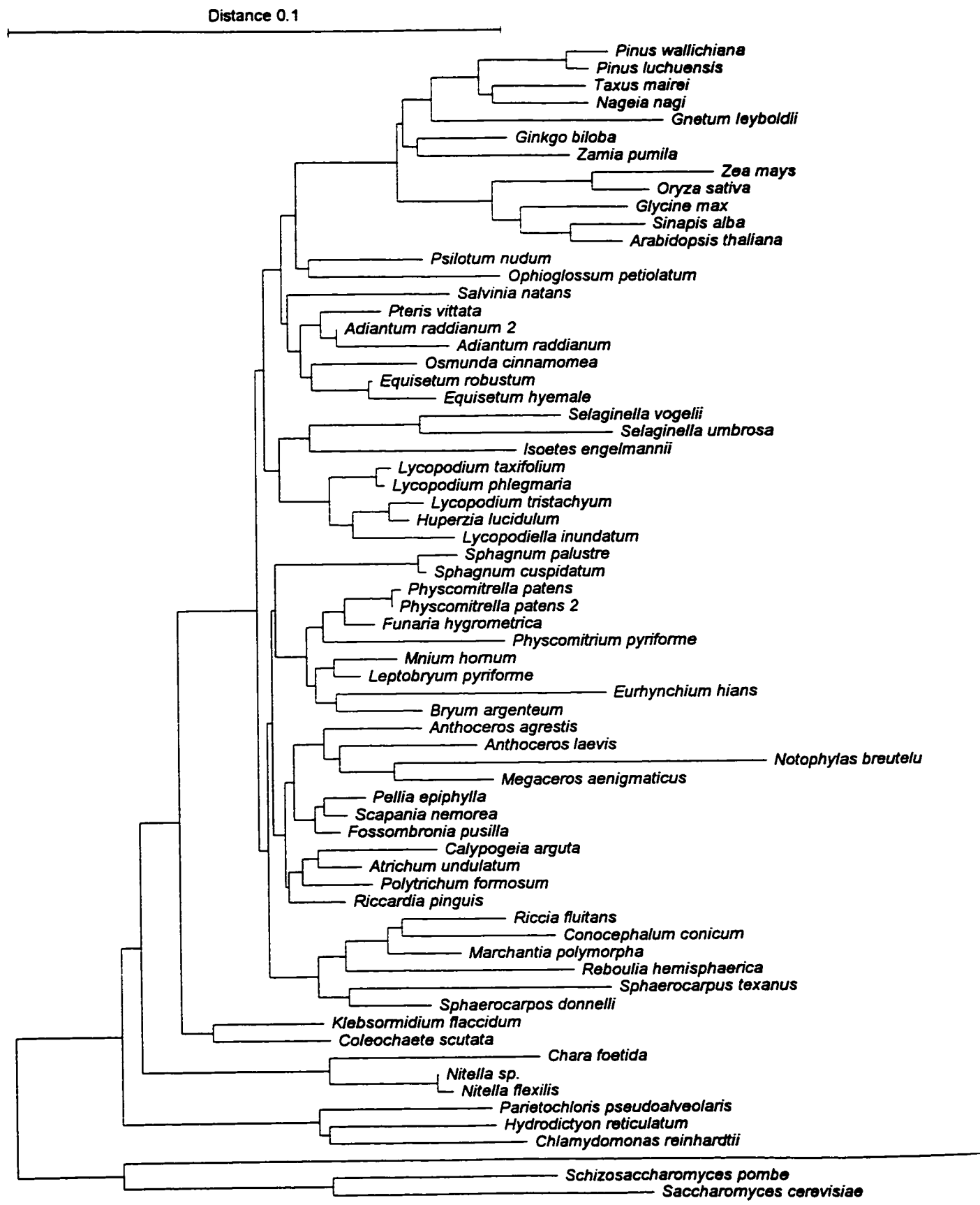


Figure C2 Kimura; Neighbor-joining (Insertions & Deletions taken into account)

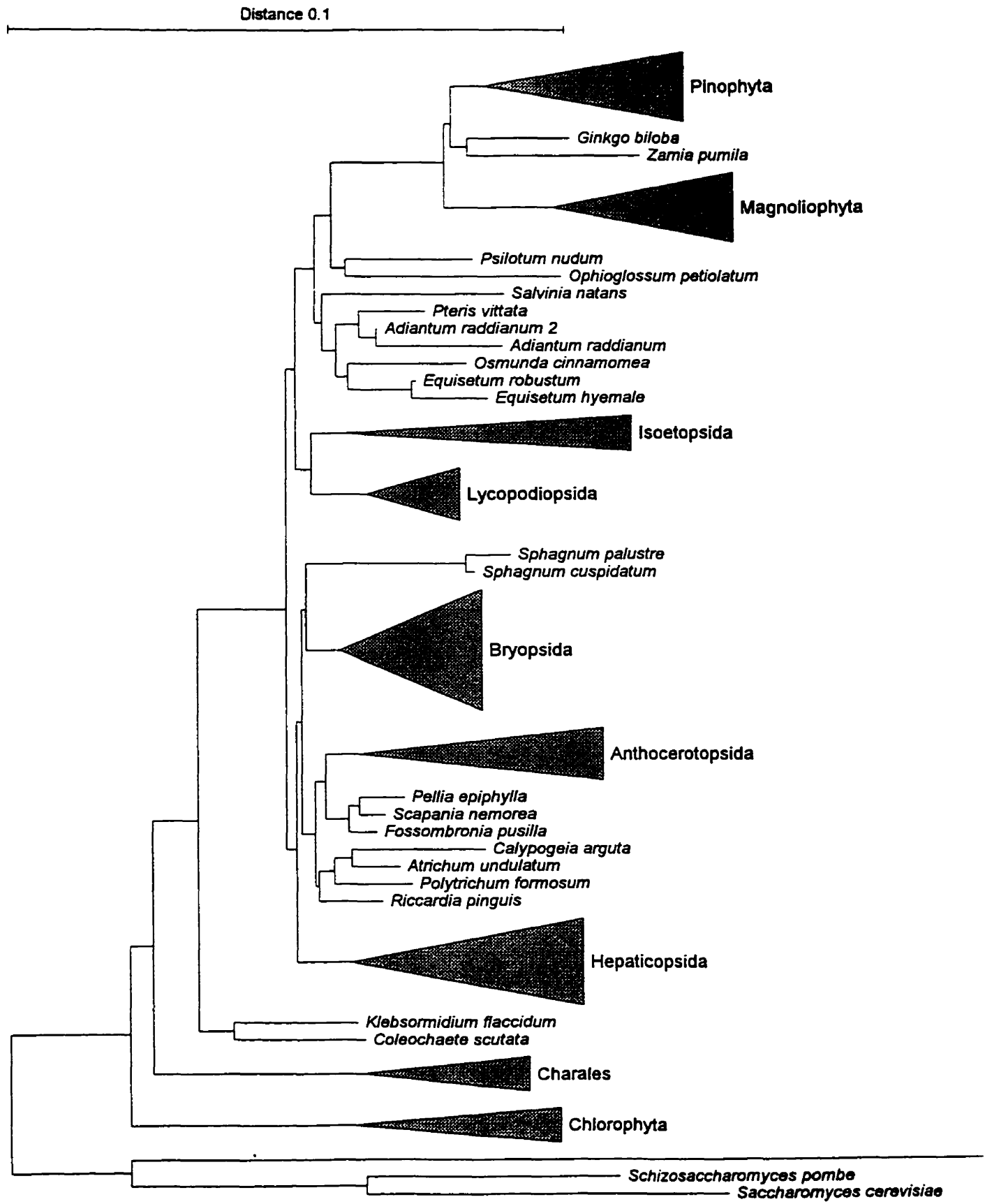


Figure C2 Kimura; Neighbor-joining (Insertions & Deletions taken into account)

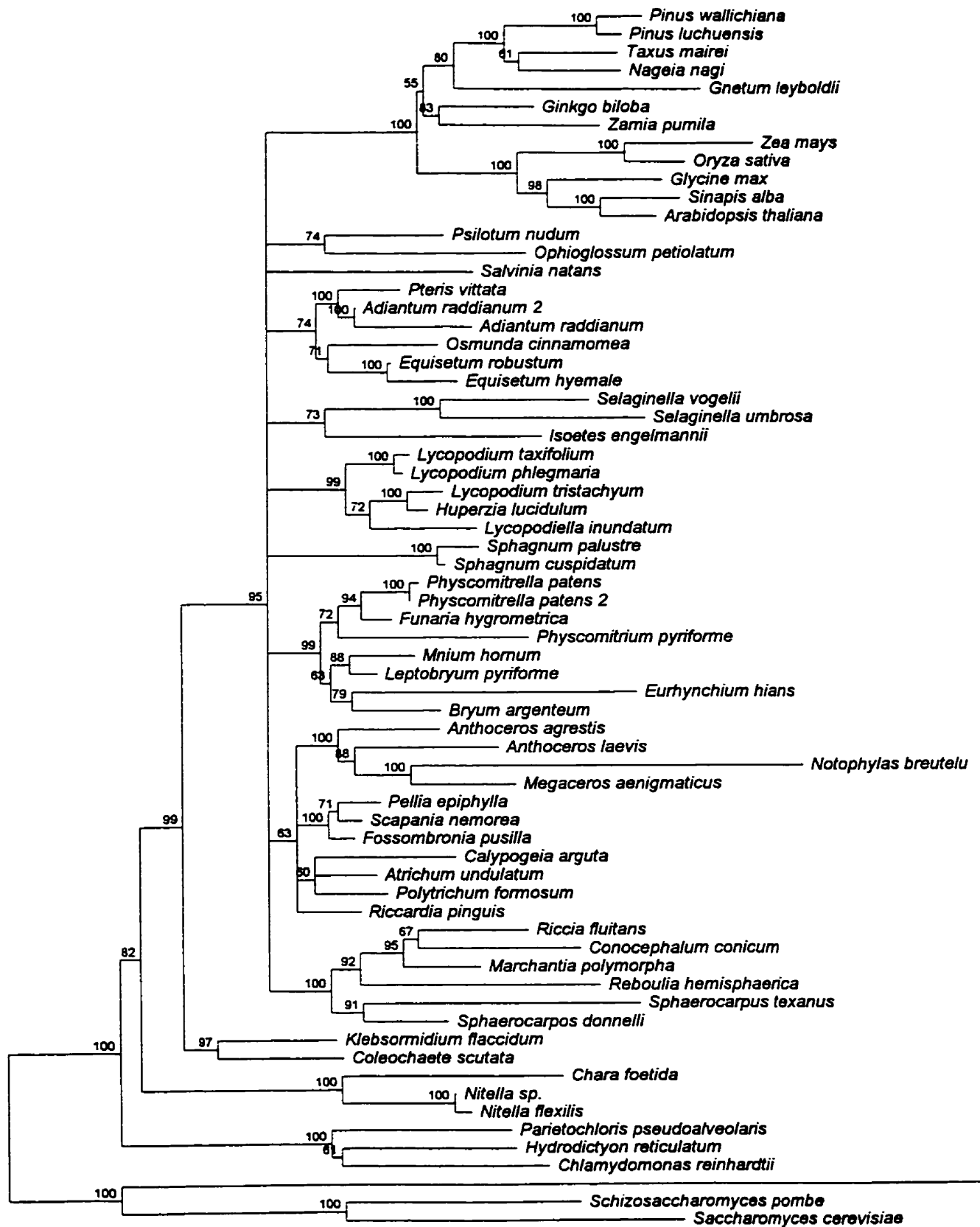


Figure C2B Kimura; (Insertions & Deletions taken into account)
50% Majority-Rule Bootstrap Tree

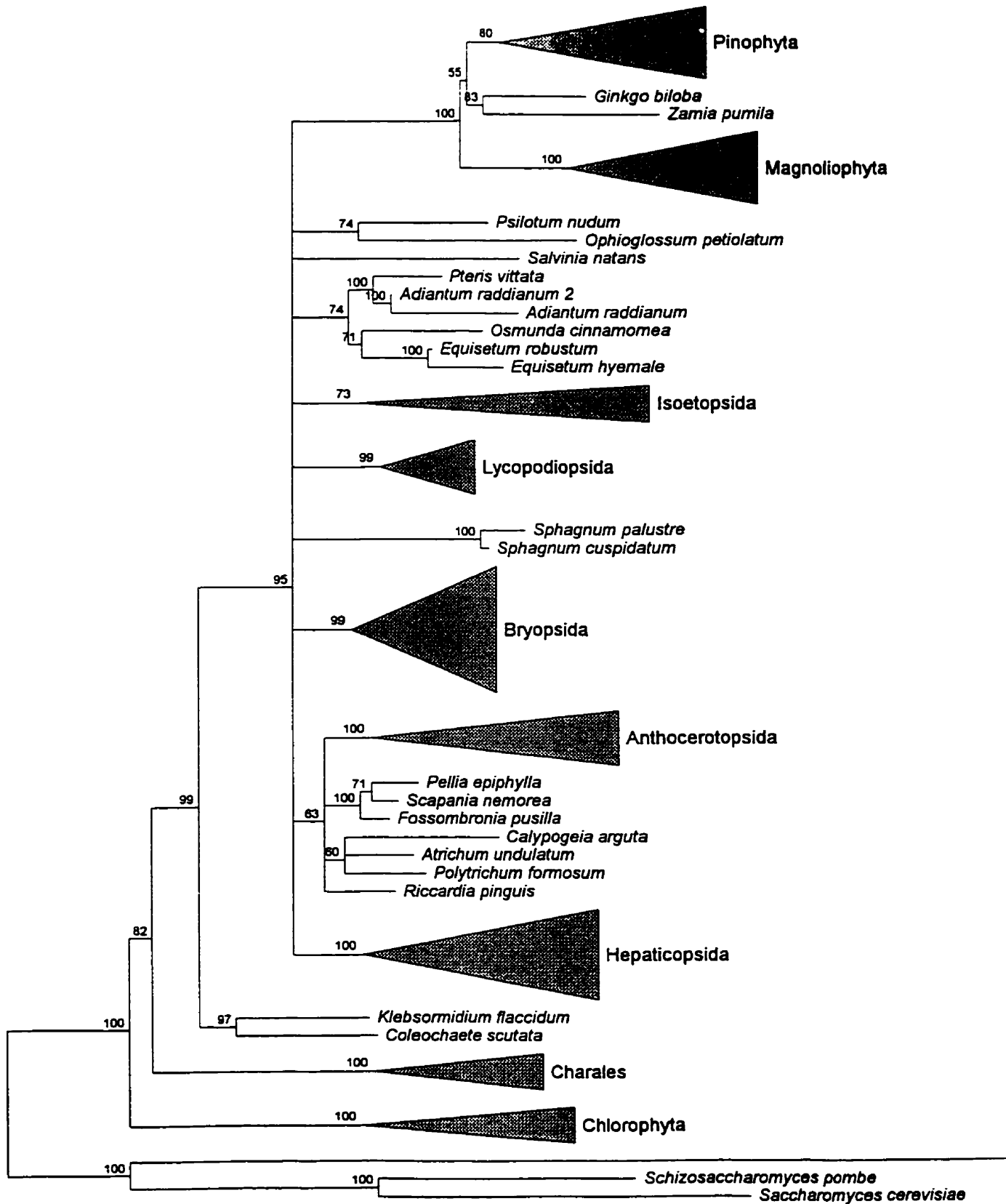


Figure C2B Kimura; (Insertions & Deletions taken into account)
 50% Majority-Rule Bootstrap Tree

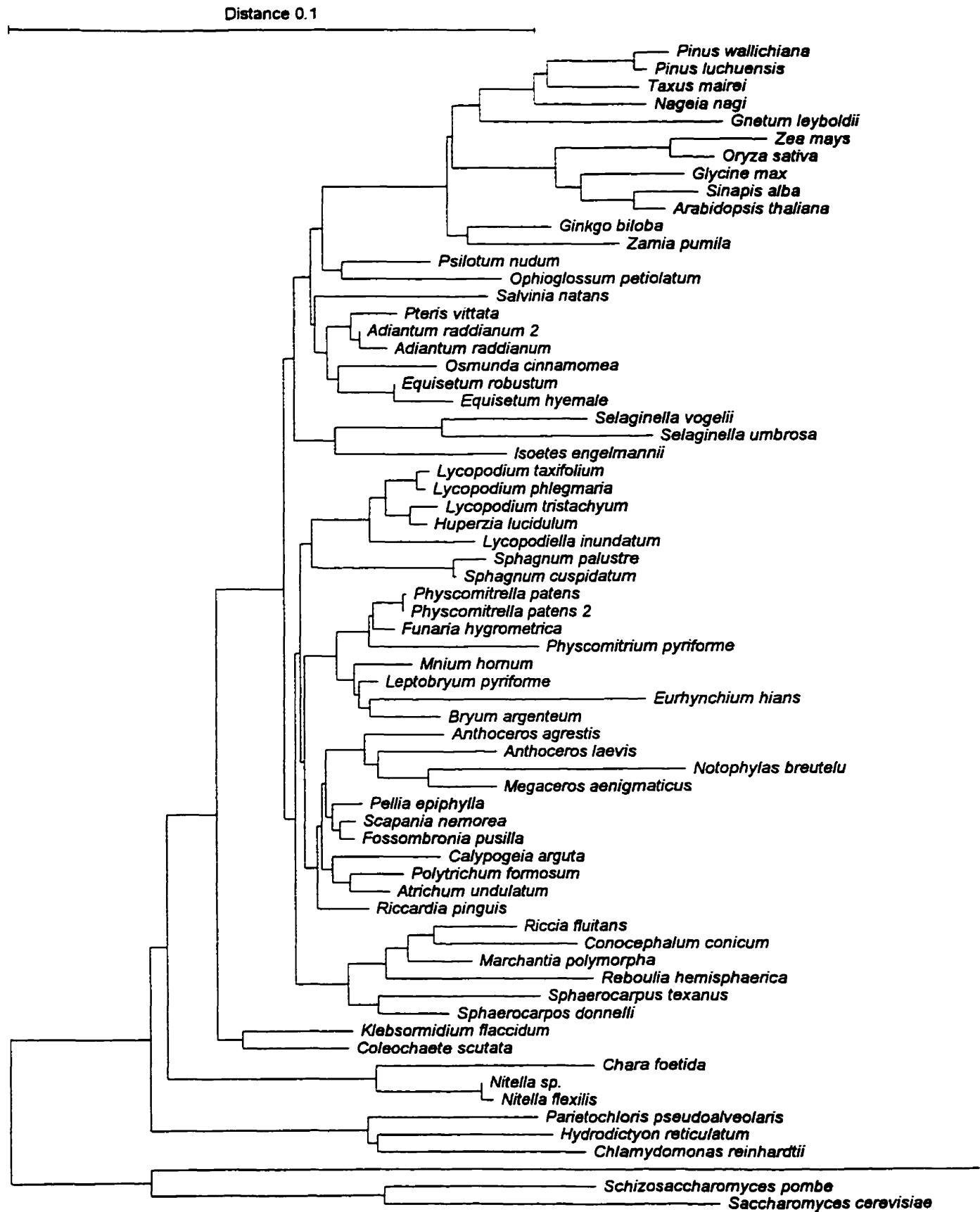


Figure D1.1 Jin & Nei (Distance Calculation); Jukes & Cantor (Evolution Model), Gamma distance = 1

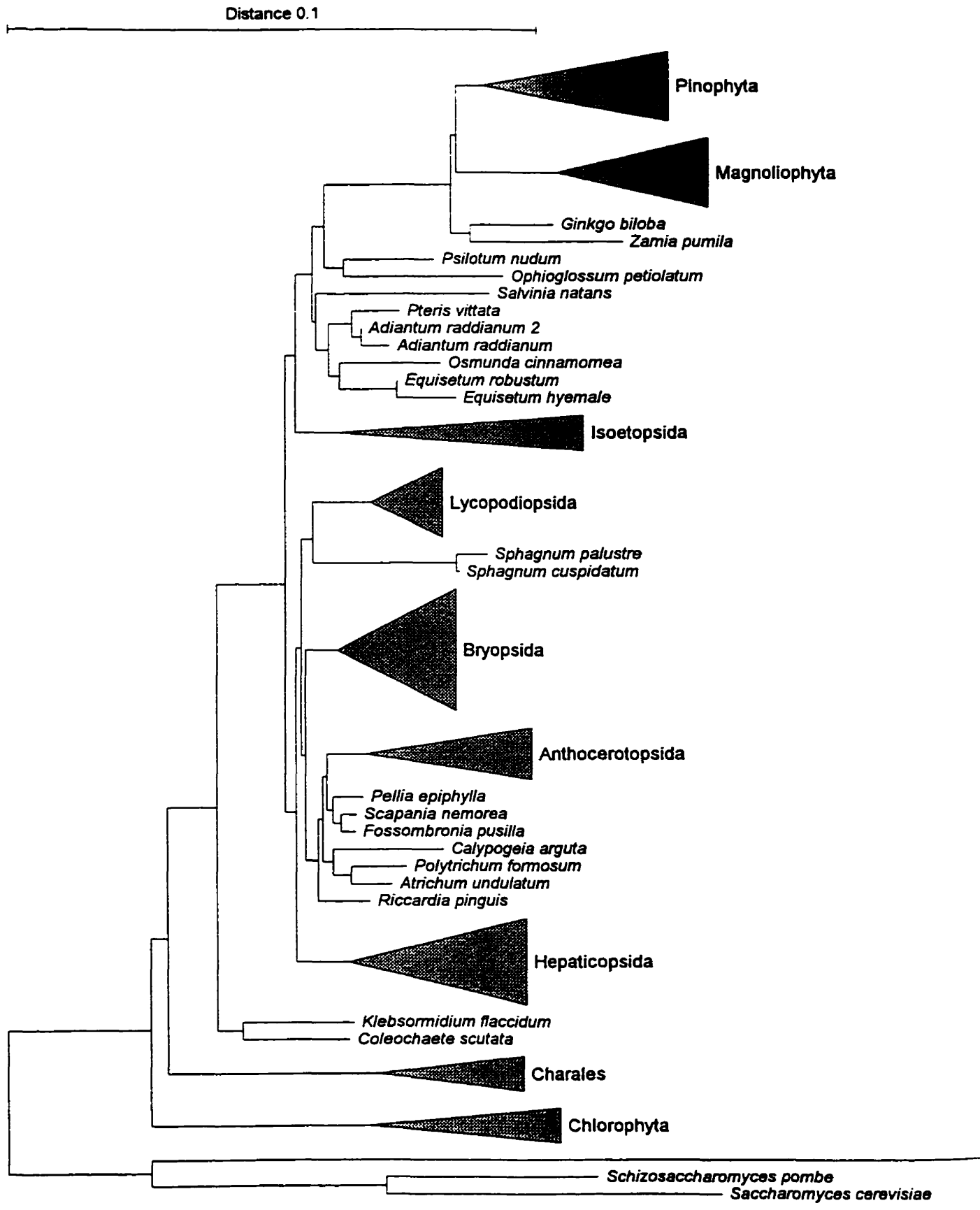


Figure D1.1 Jin & Nei (Distance Calculation); Jukes & Cantor (Evolution Model), Gamma distance = 1

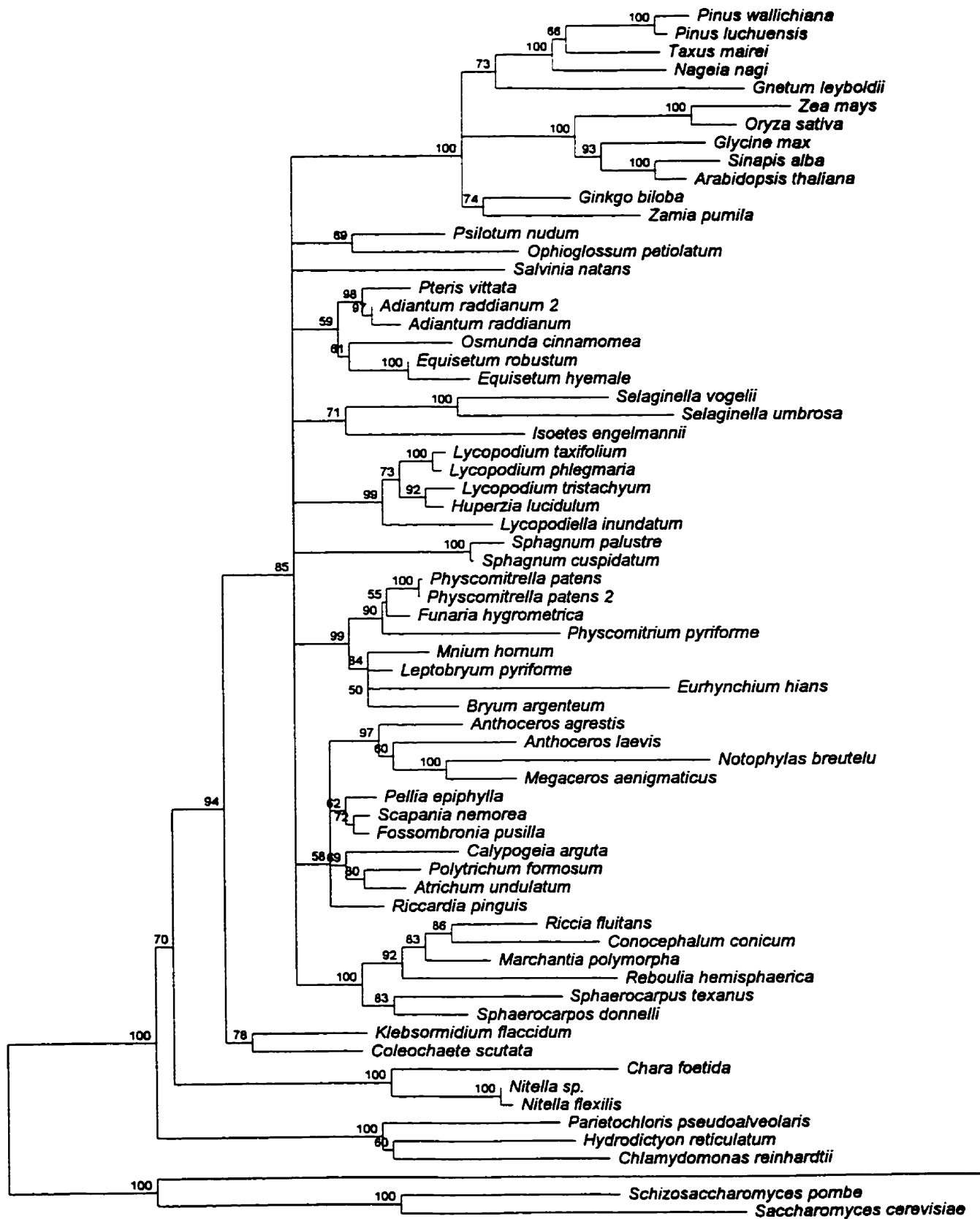


Figure D1.1B Jin & Nei (Distance Calculation); Jukes & Cantor (Evolution Model), Gamma distance = 1
50% Majority-Rule Bootstrap Tree

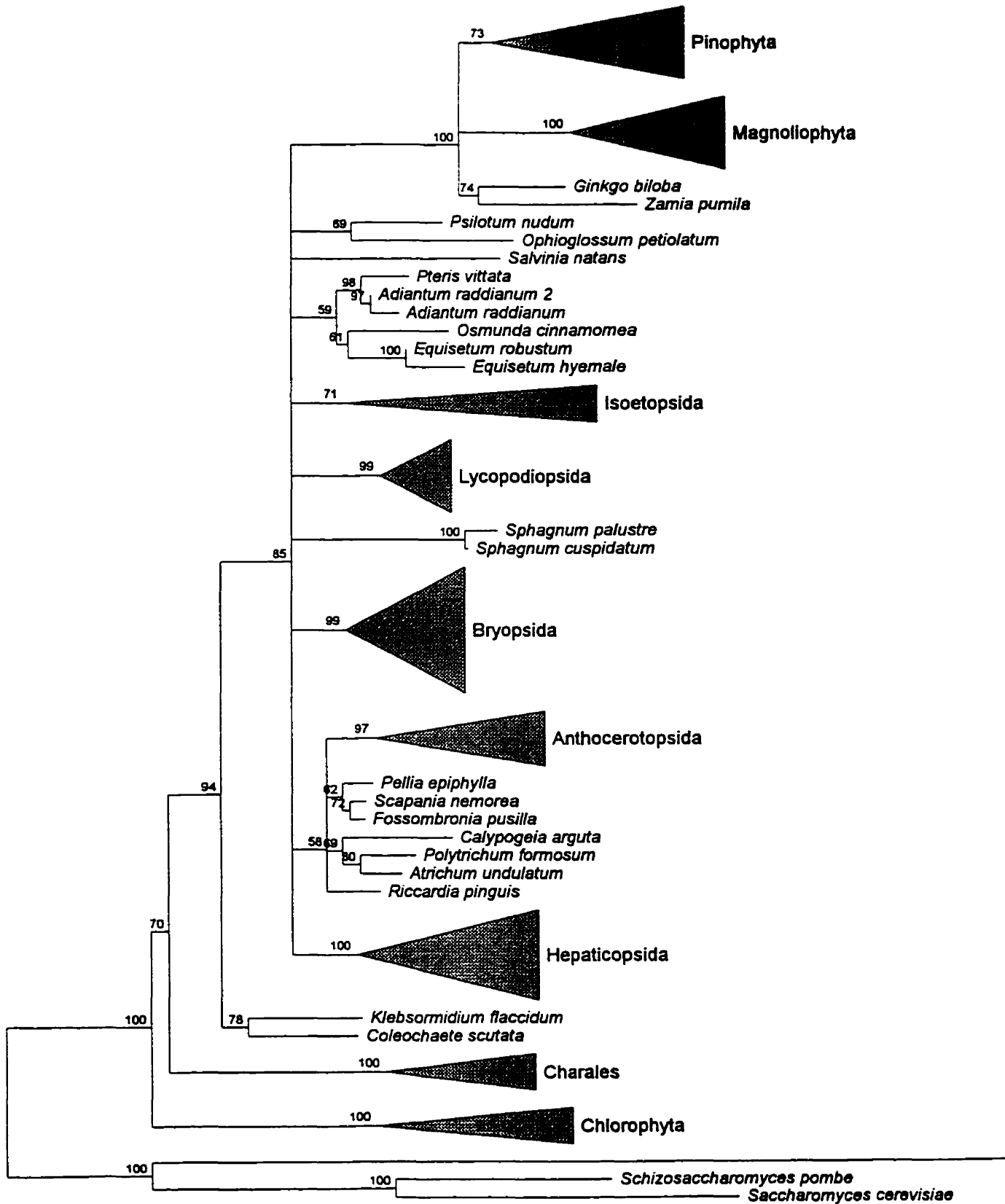


Figure D1.18 Jin & Nei (Distance Calculation); Jukes & Cantor (Evolution Model), Gamma distance = 1
50% Majority-Rule Bootstrap Tree

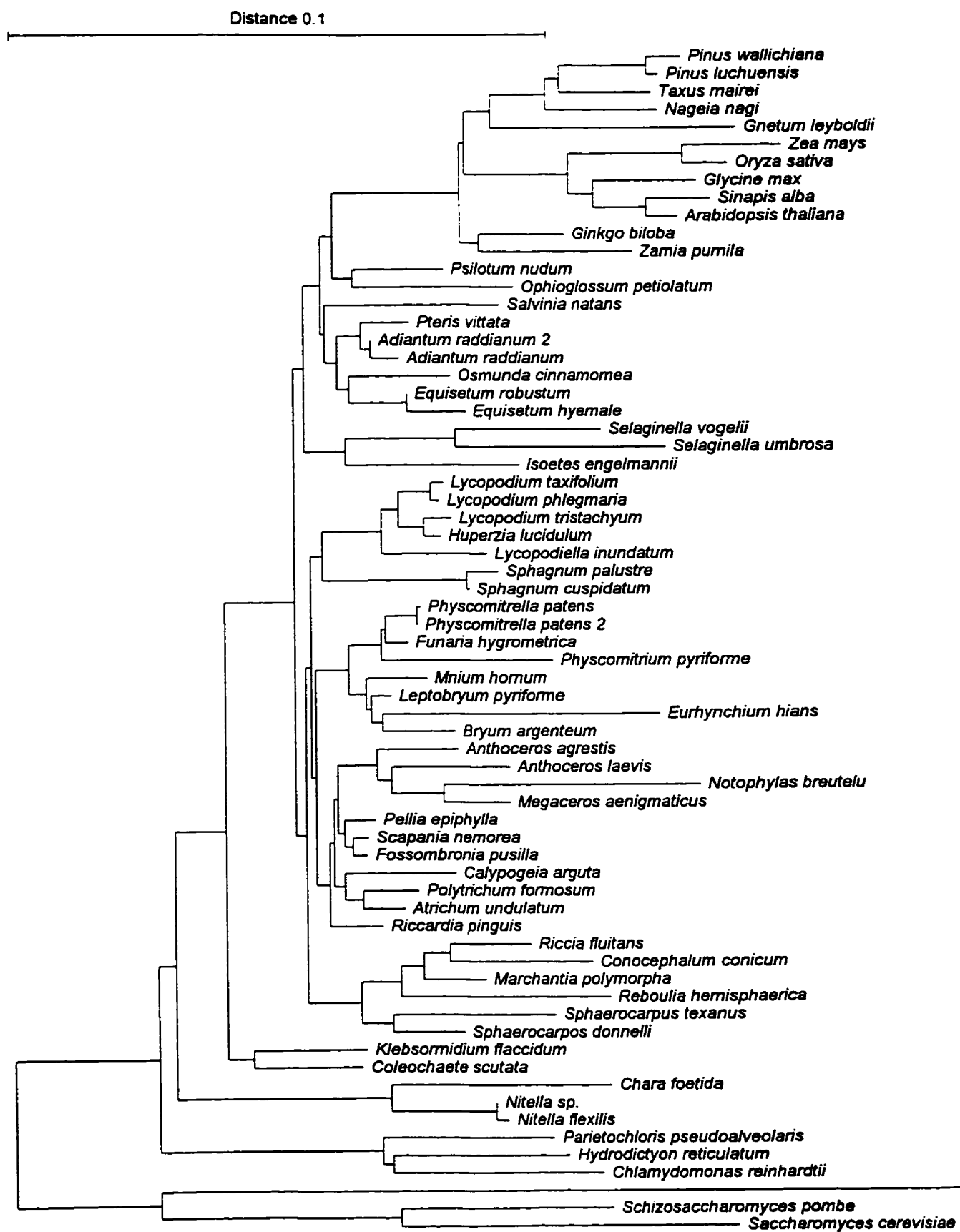


Figure D1.2 Jin & Nei (Distance Calculation); Jukes & Cantor (Evolution Model), Gamma distance = 1
Insertions & Deletions taken into account

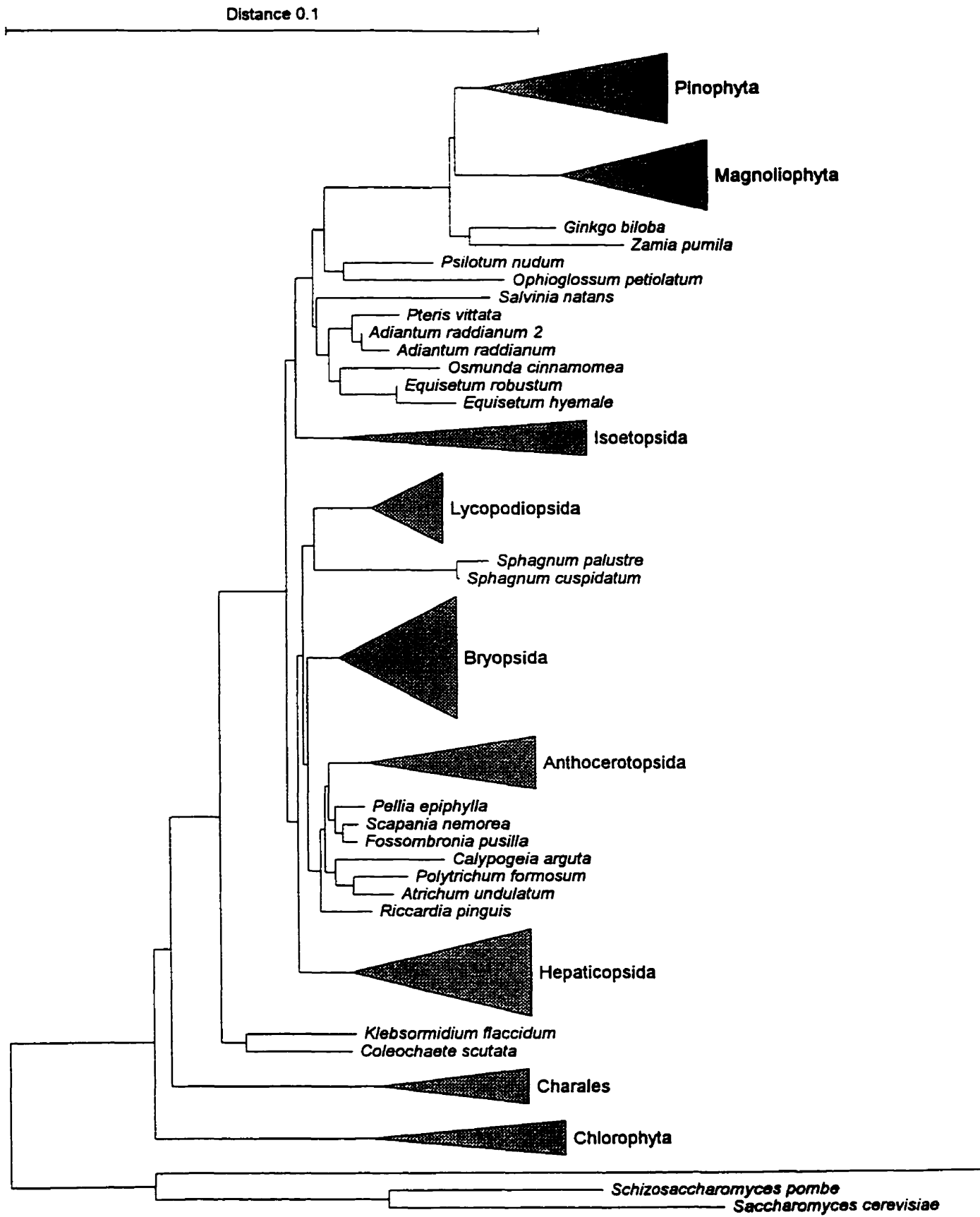


Figure D1.2 Jin & Nei (Distance Calculation); Jukes & Cantor (Evolution Model), Gamma distance = 1 insertions & Deletions taken into account

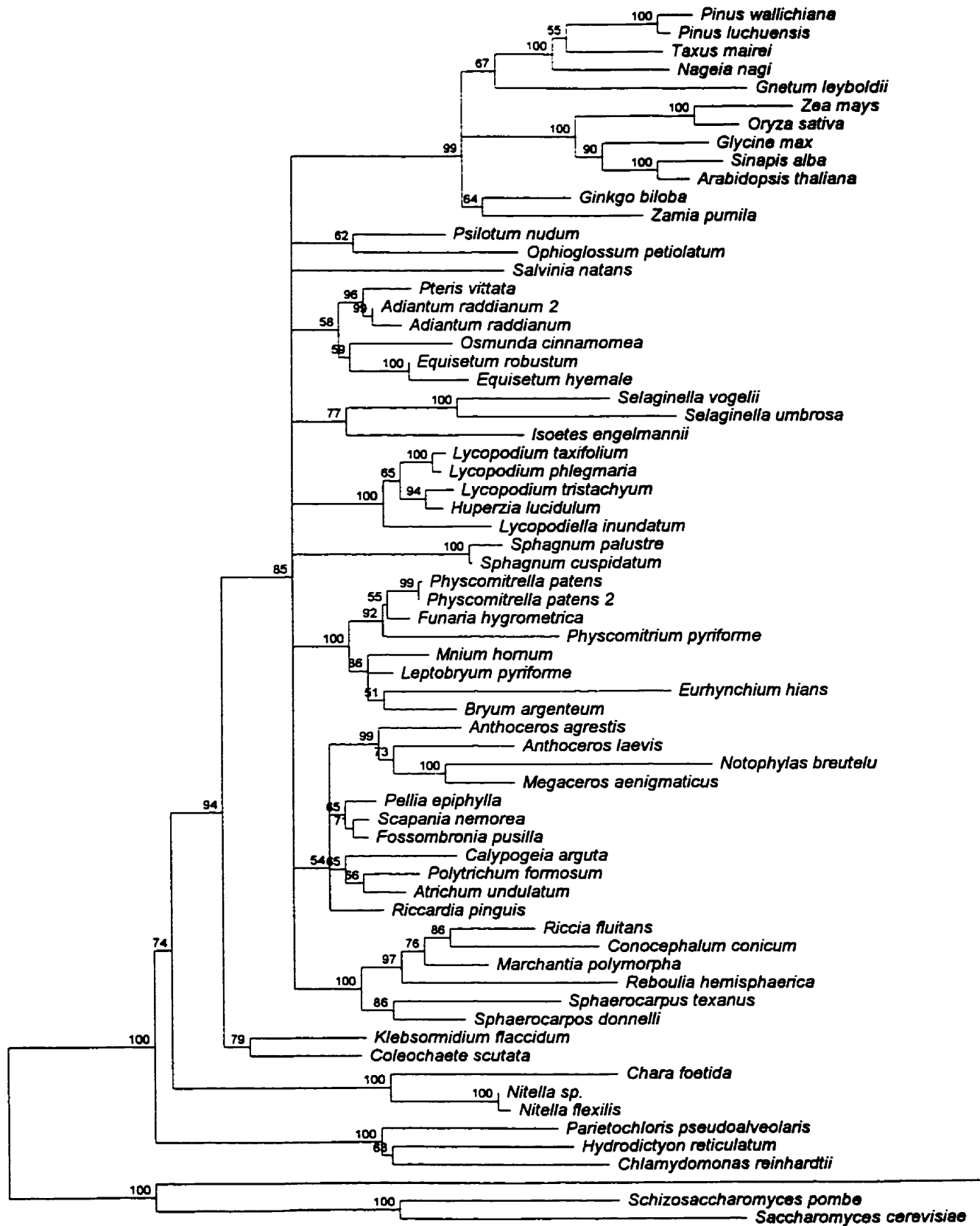


Figure D1.2B Jin & Nei (Distance Calculation); Jukes & Cantor (Evolution Model), Gamma distance = 1
 Insertions & Deletions taken into account
 50% Majority-Rule Bootstrap Tree

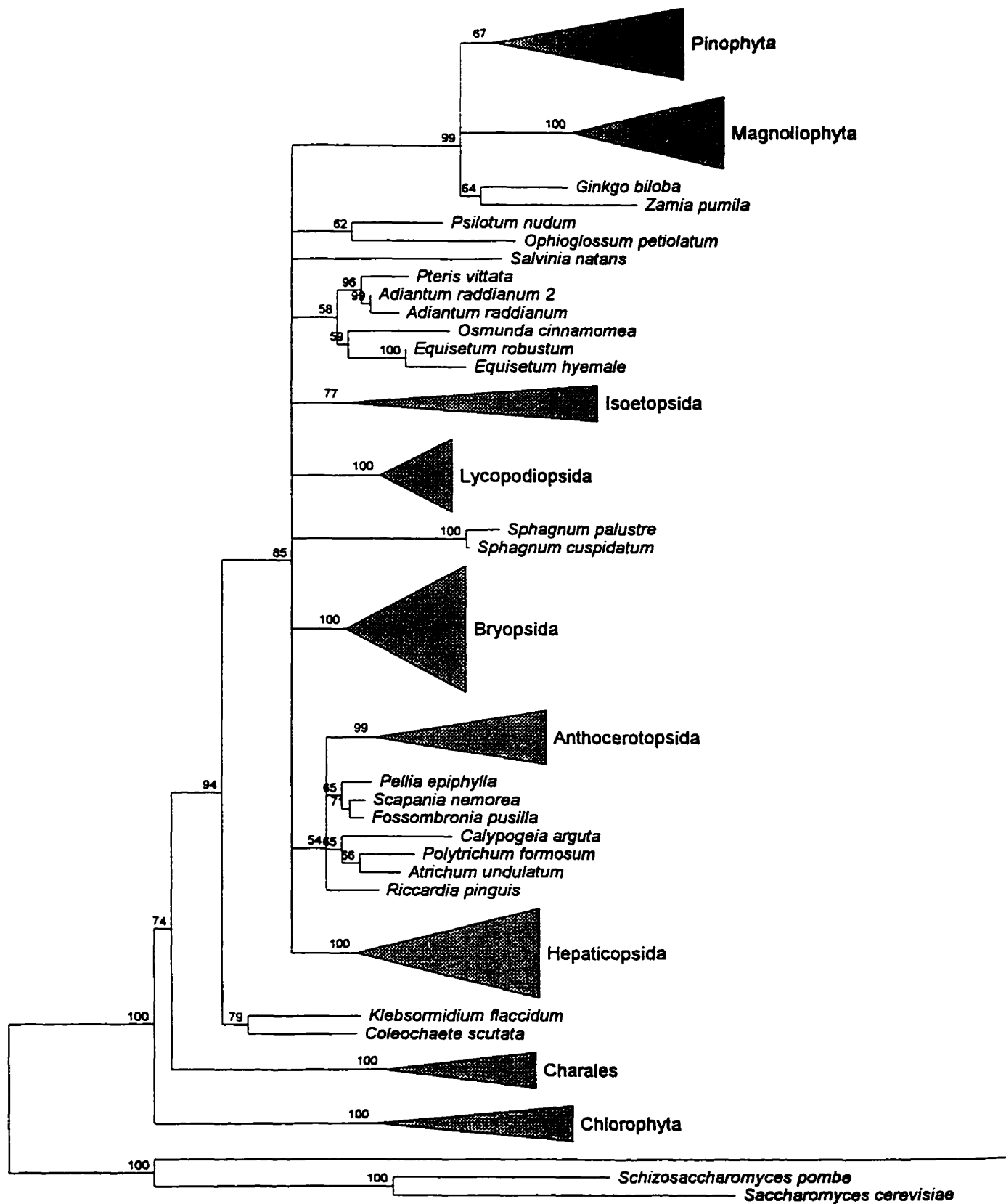


Figure D1.2B Jin & Nei (Distance Calculation); Jukes & Cantor (Evolution Model), Gamma distance = 1
 Insertions & Deletions taken into account
 50% Majority-Rule Bootstrap Tree

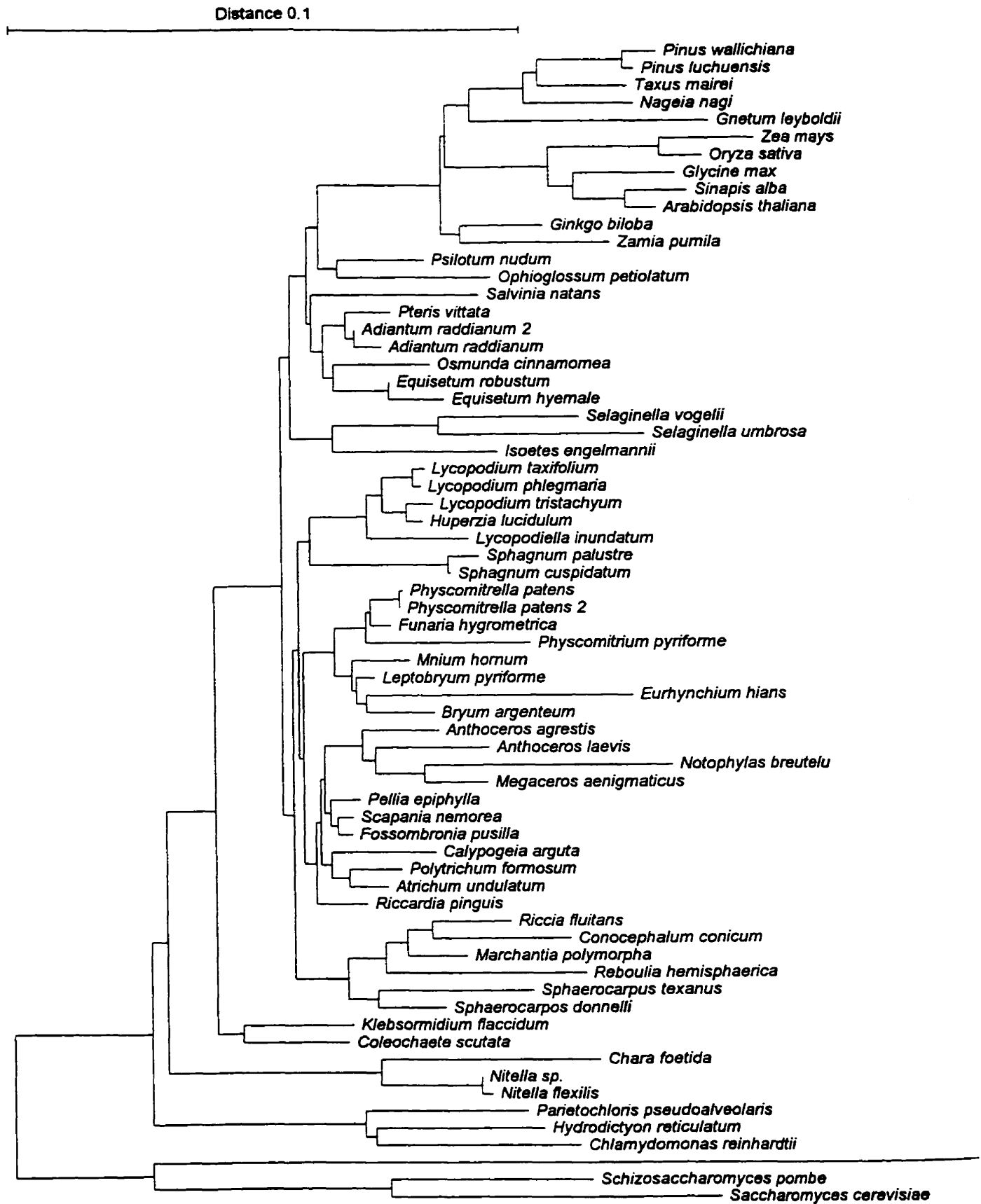


Figure D2.1 Jin & Nei (Distance Calculation); Kimura 2-p (Evolution Model), Gamma distance = 1

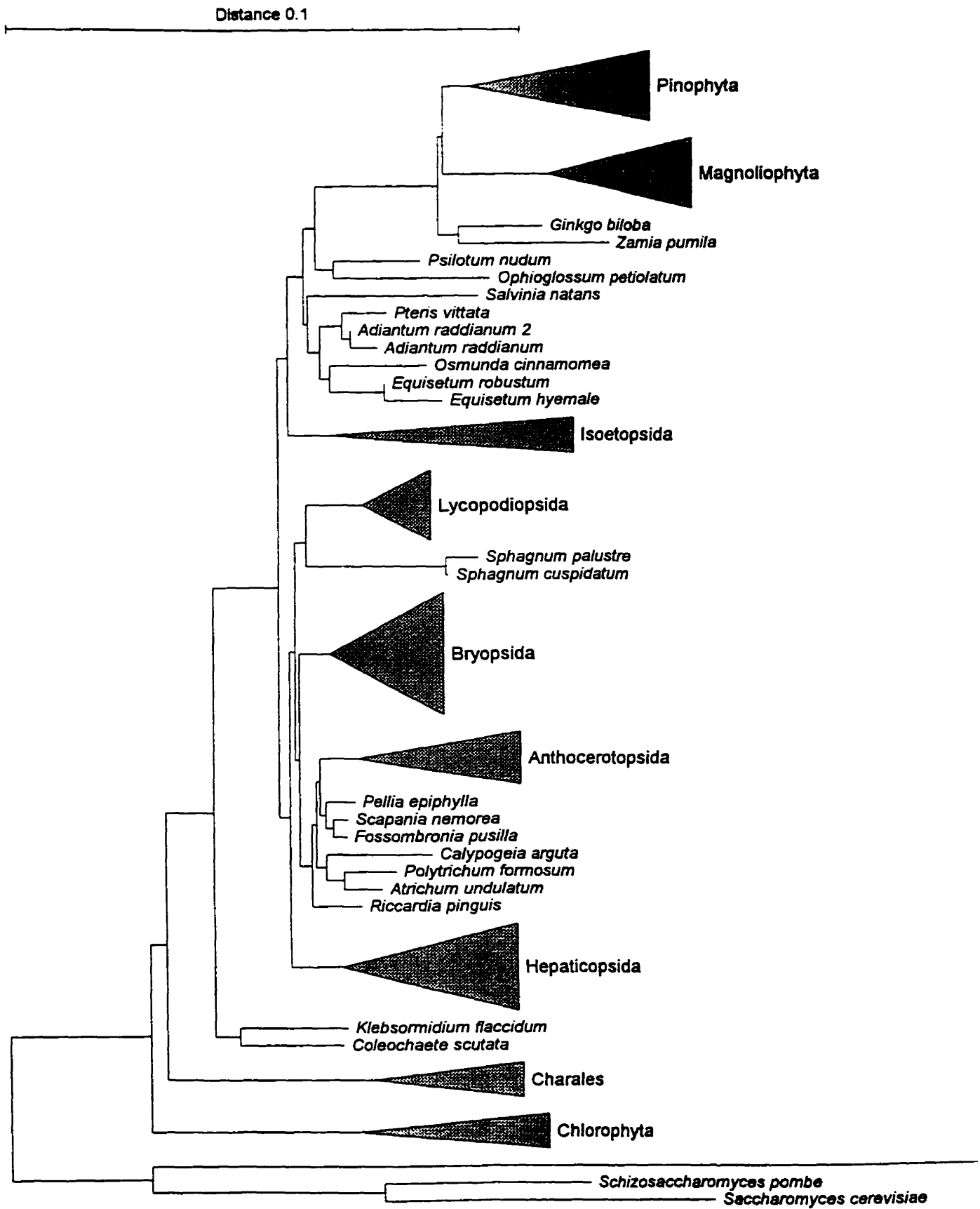


Figure D2.1 Jin & Nei (Distance Calculation); Kimura 2-p (Evolution Model), Gamma distance = 1

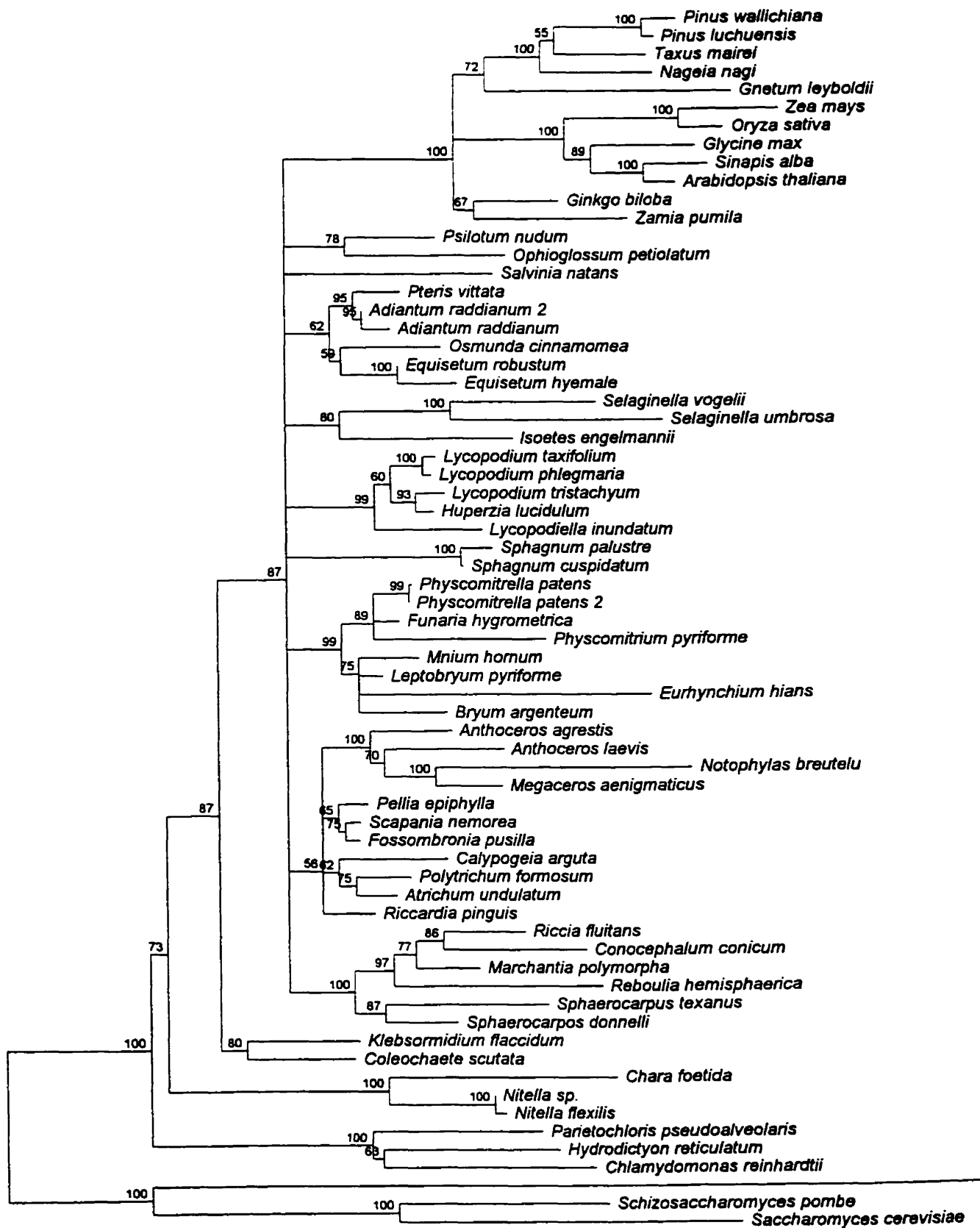


Figure D2.18 Jin & Nei (Distance Calculation); Kimura 2-p (Evolution Model), Gamma distance = 1
50% Majority-Rule Bootstrap Tree

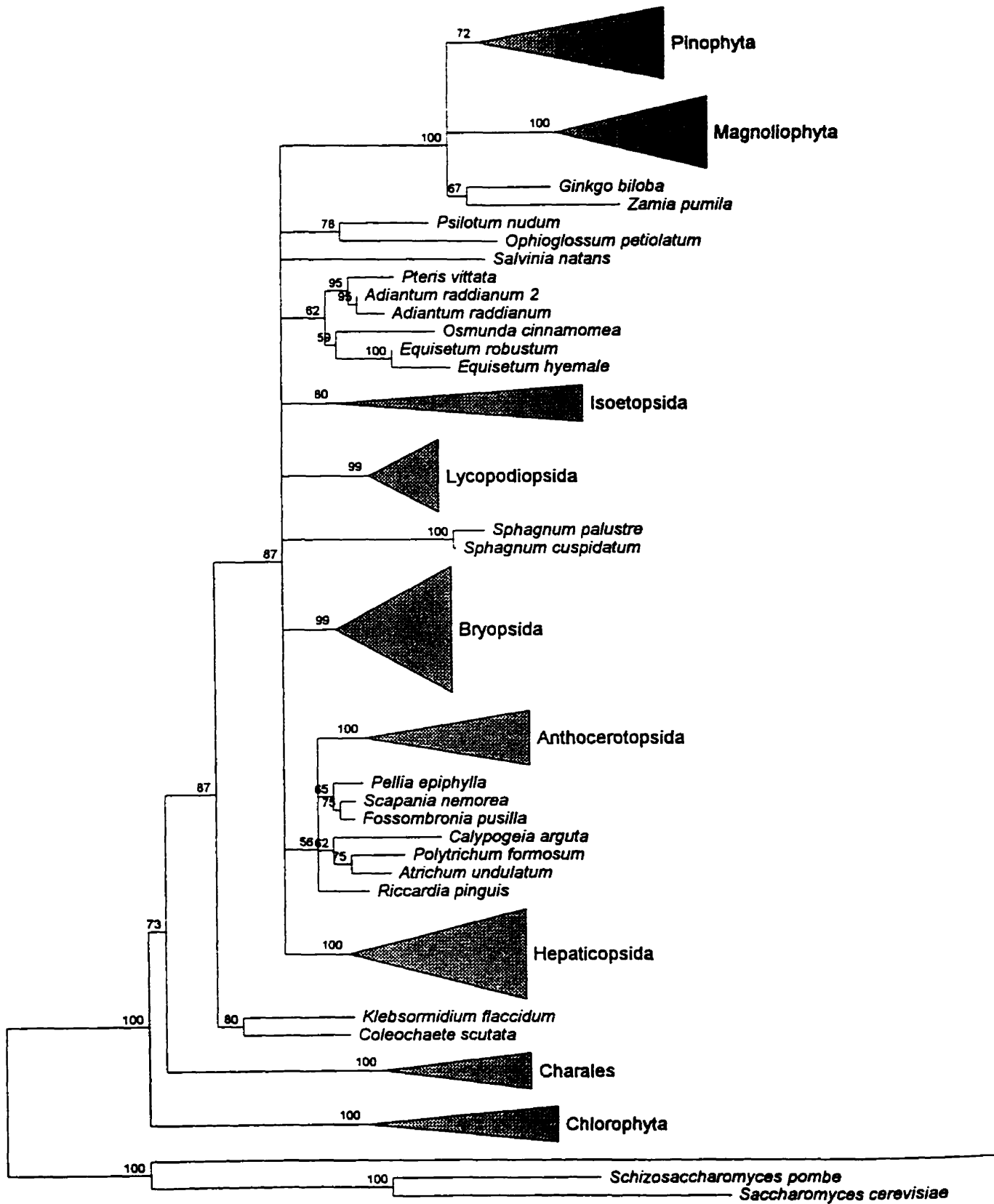


Figure D2.1B Jin & Nei (Distance Calculation); Kimura 2-p (Evolution Model), Gamma distance = 1
50% Majority-Rule Bootstrap Tree

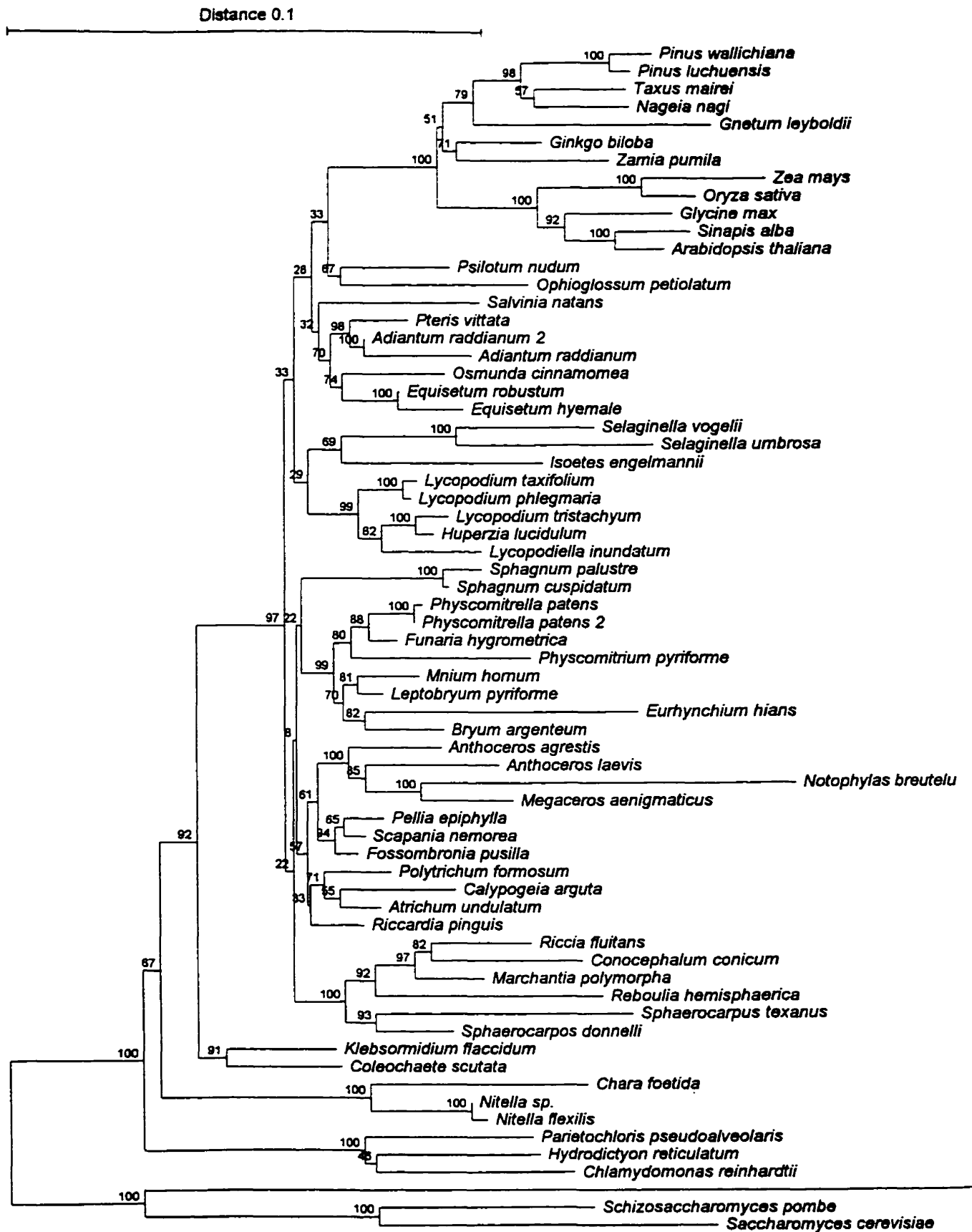


Figure D2.2 Jin & Nei (Distance Calculation); Kimura 2-p (Evolution Model), Gamma distance = 1
Insertions and Deletions taken into account

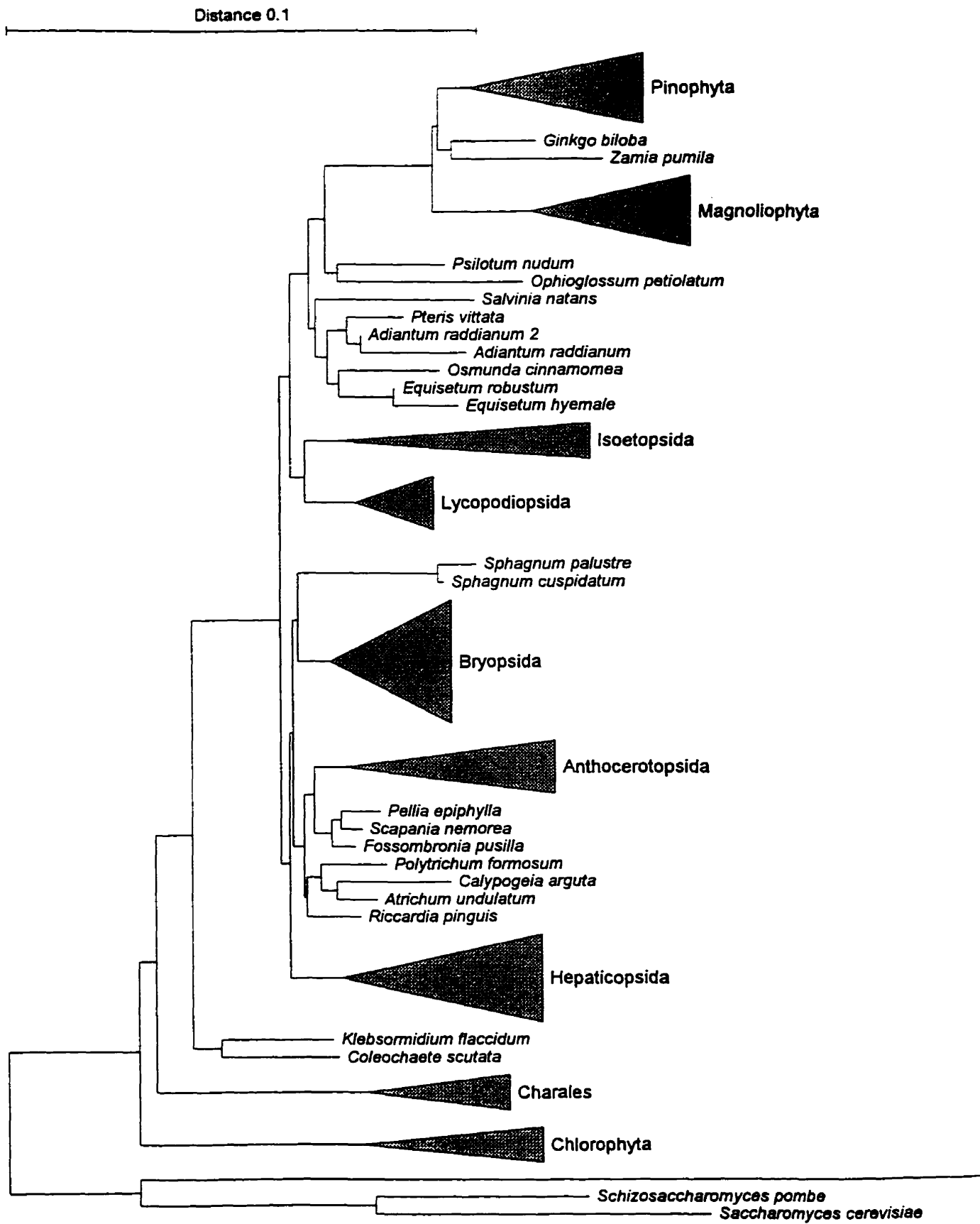


Figure D2.2 Jin & Nei (Distance Calculation); Kimura 2-p (Evolution Model). Gamma distance = 1
Insertions and Deletions taken into account

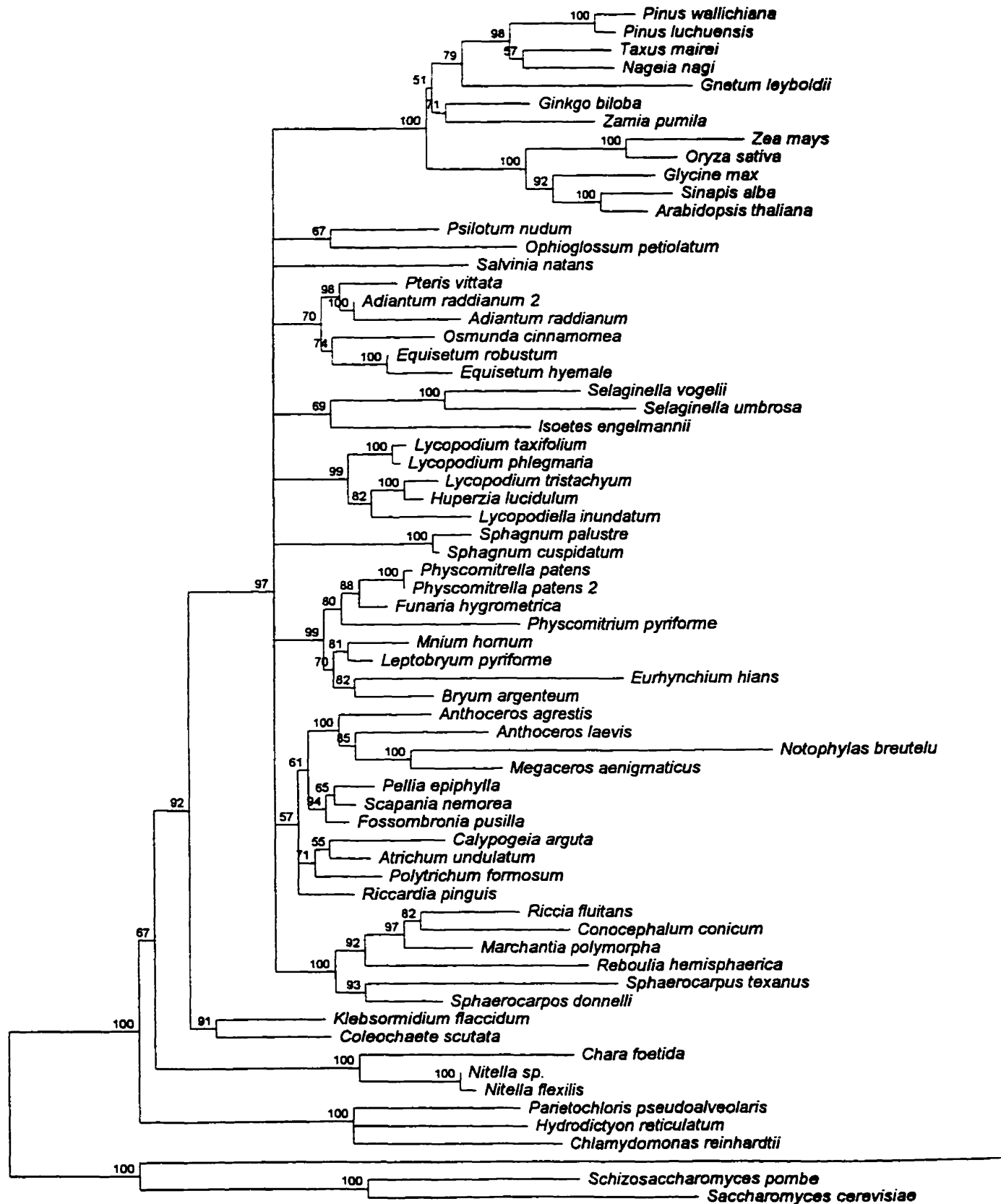


Figure D2.2B Jin & Nei (Distance Calculation); Kimura 2-p (Evolution Model), Gamma distance = 1
 Insertions & Deletions taken into account
 50% Majority-Rule Bootstrap Tree

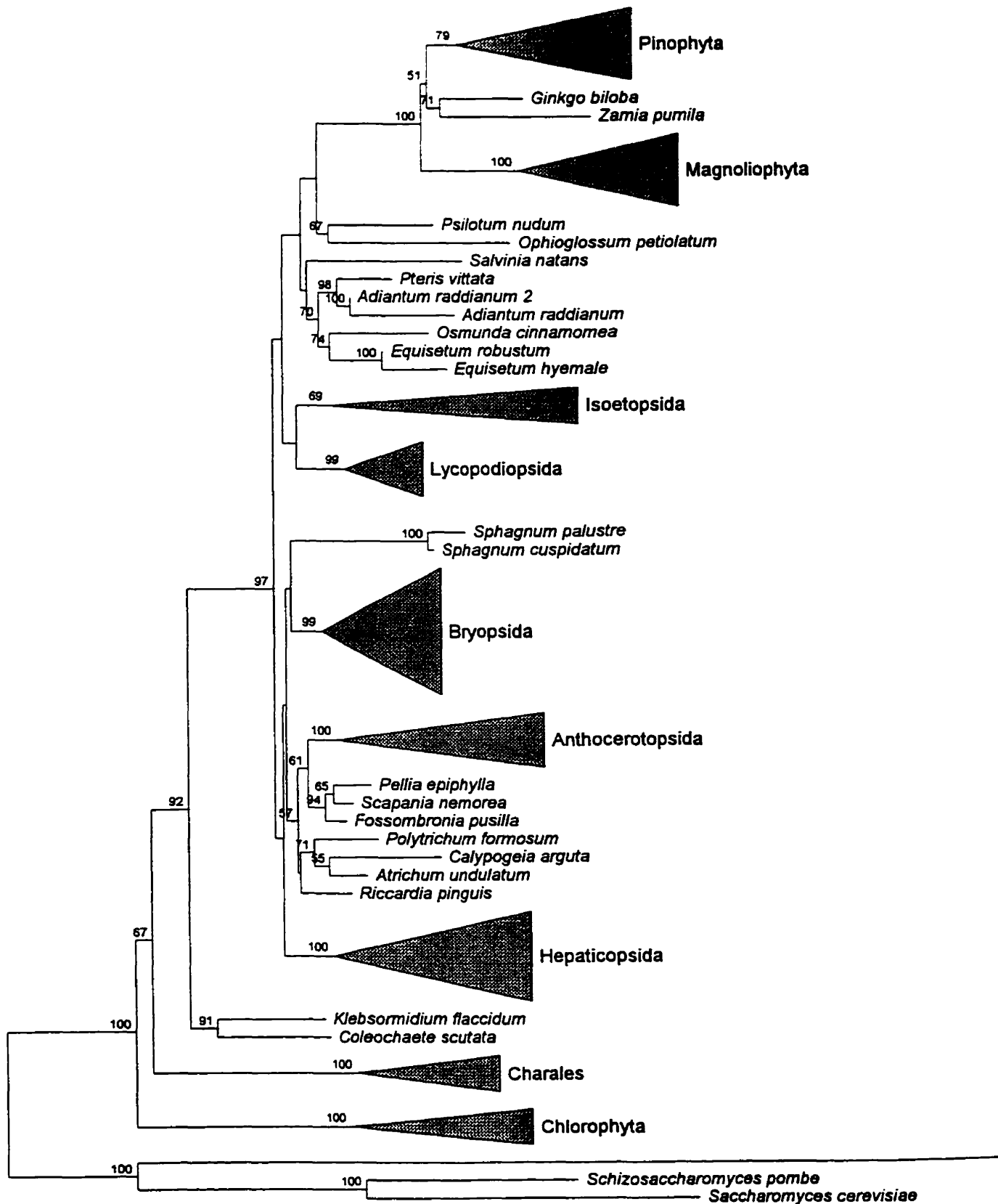


Figure D2.2B Jin & Nei (Distance Calculation); Kimura 2-p (Evolution Model), Gamma distance = 1
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 50% Majority-Rule Bootstrap Tree

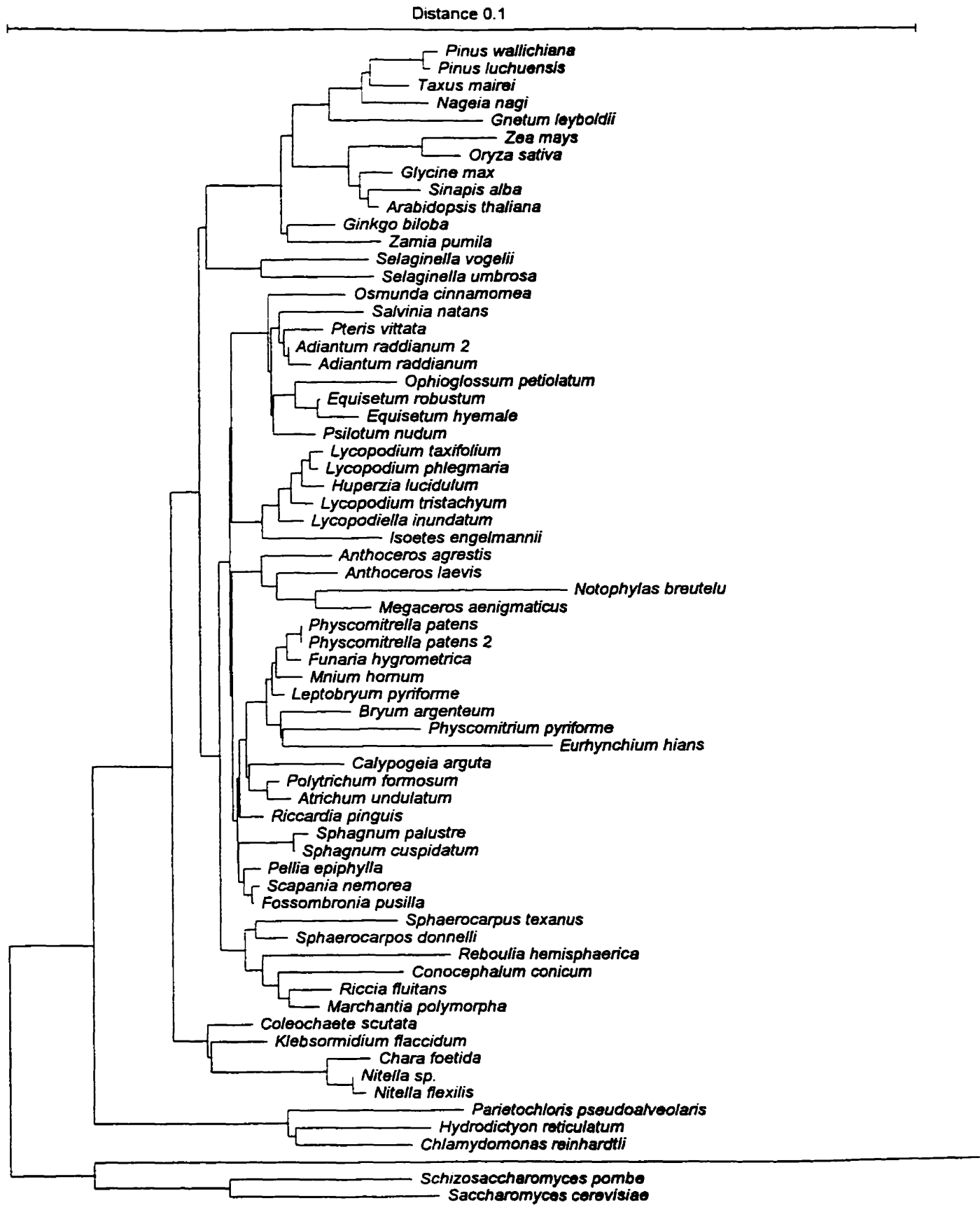


Figure E1 Transversions Only; Neighbor-joining

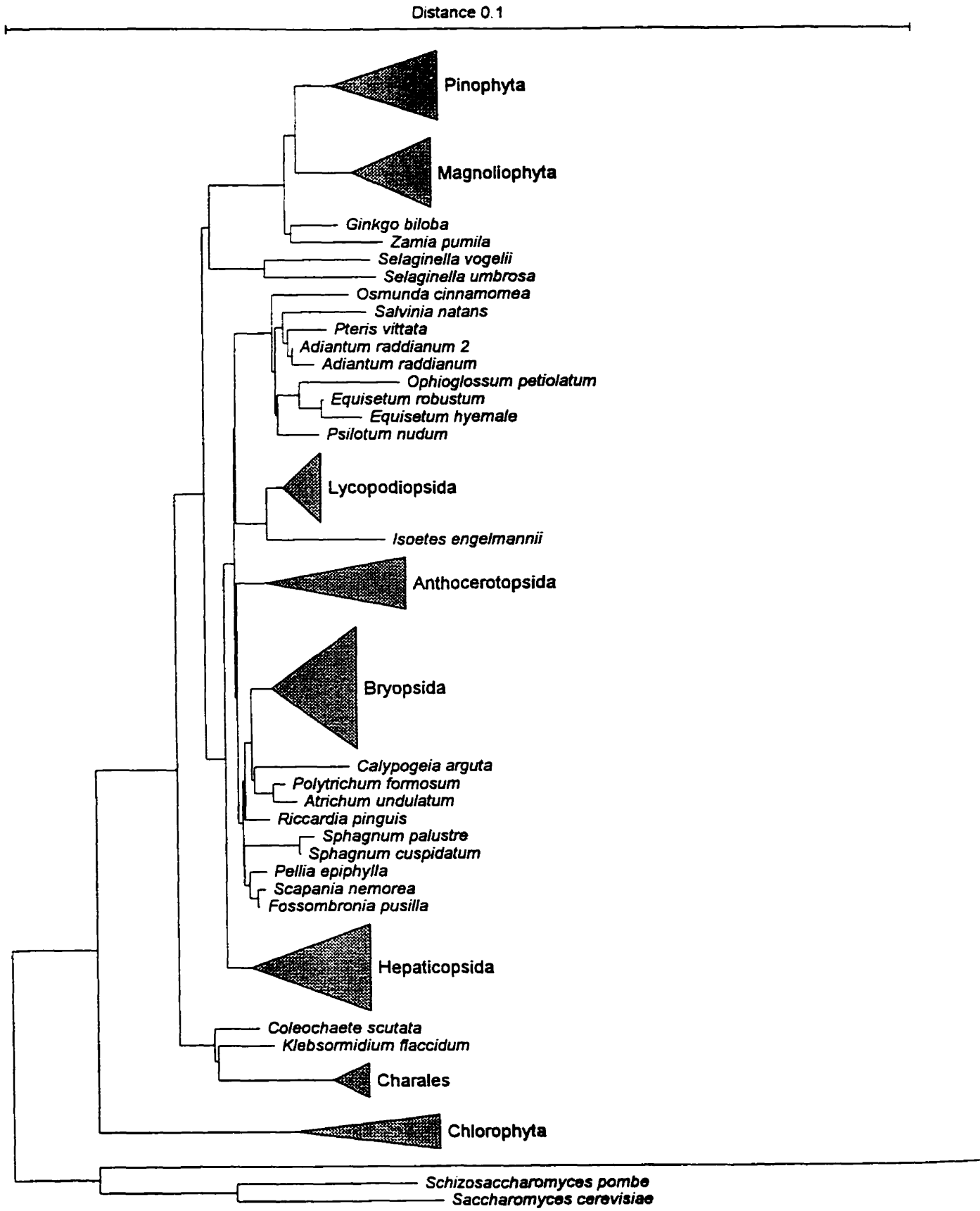


Figure E1 Transversions Only; Neighbor-joining

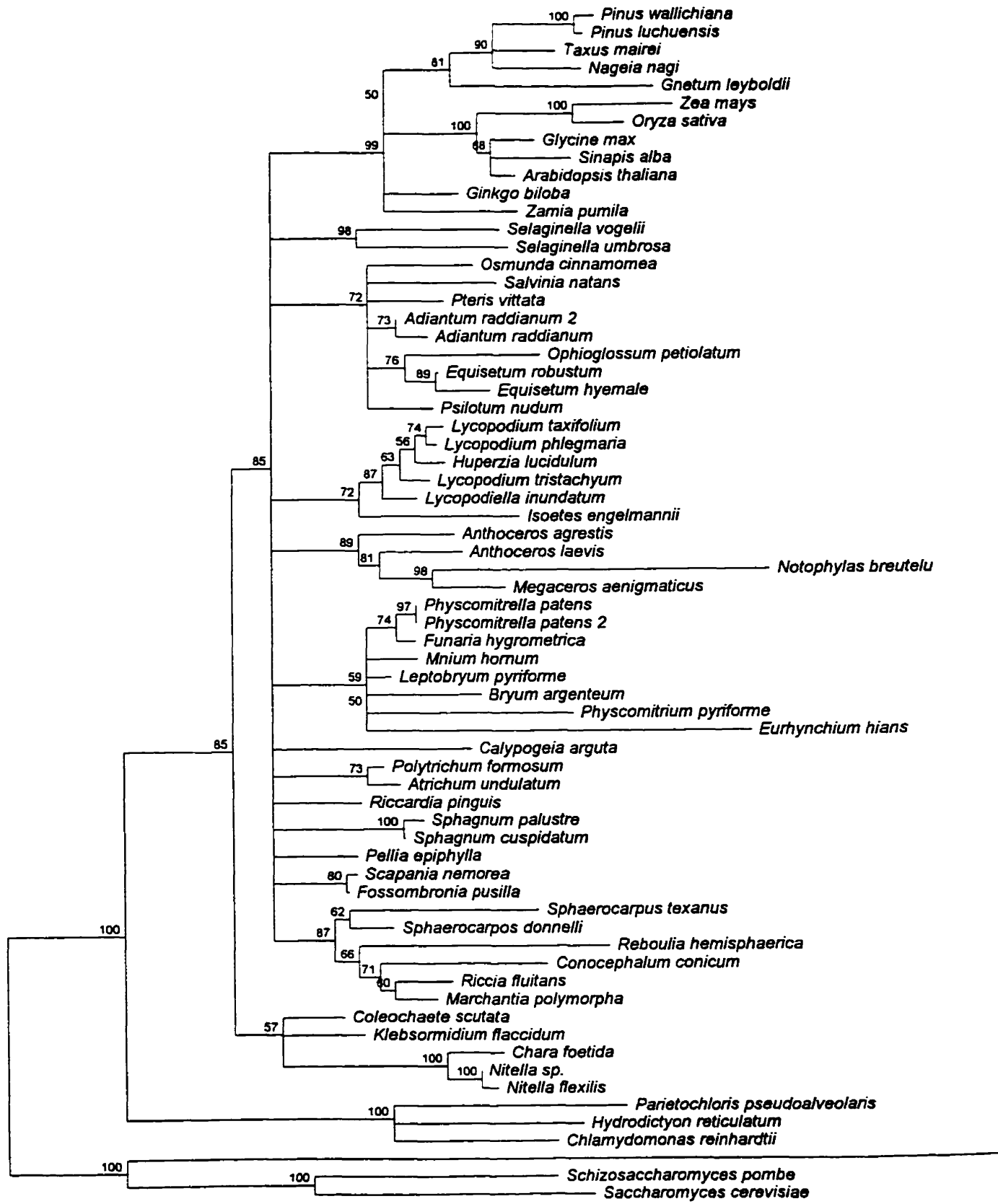


Figure E1B Transversions Only; Neighbor-joining
50% Majority-Rule Bootstrap Tree

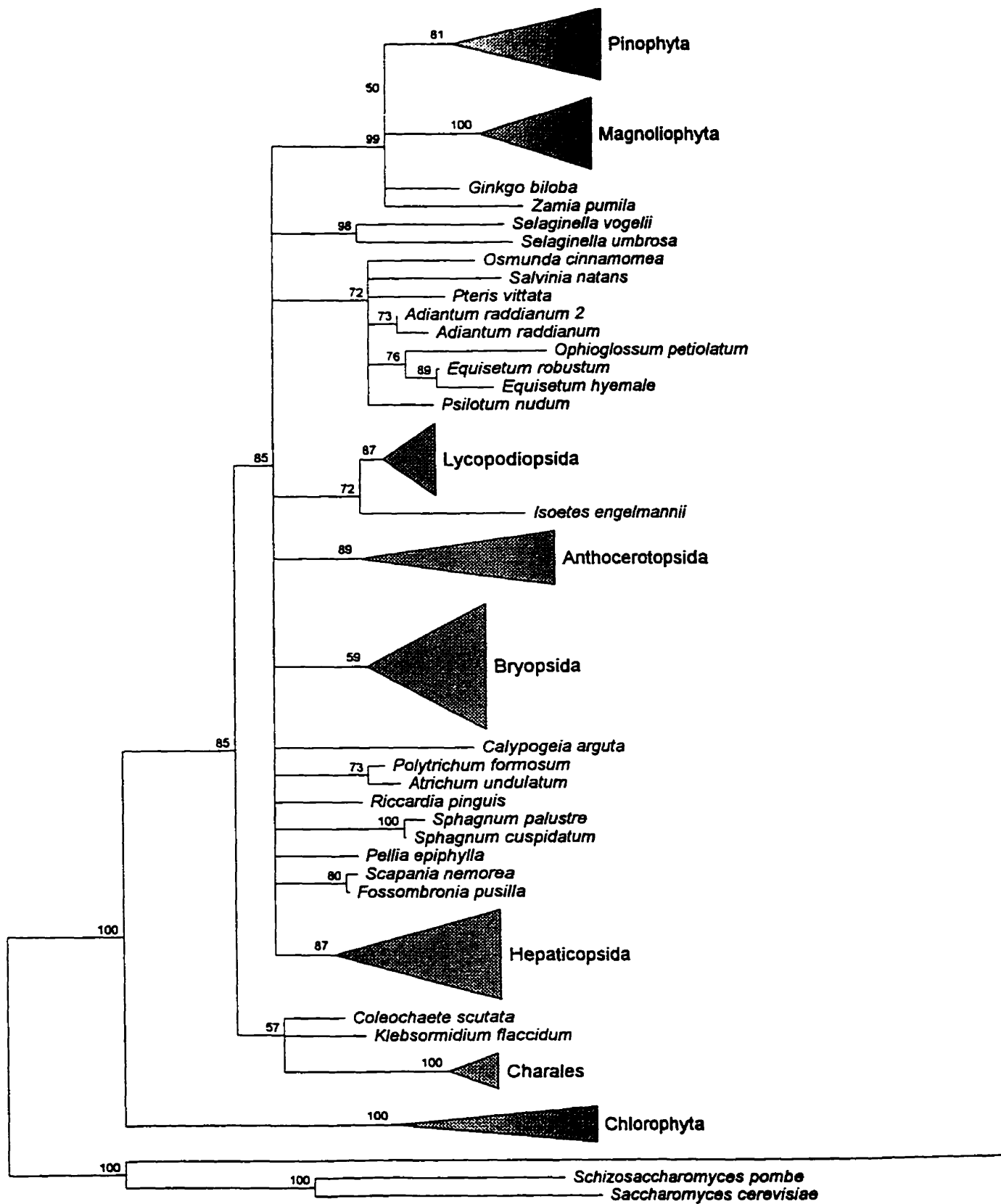


Figure E1B Transversions Only: Neighbor-joining
50% Majority-Rule Bootstrap Tree

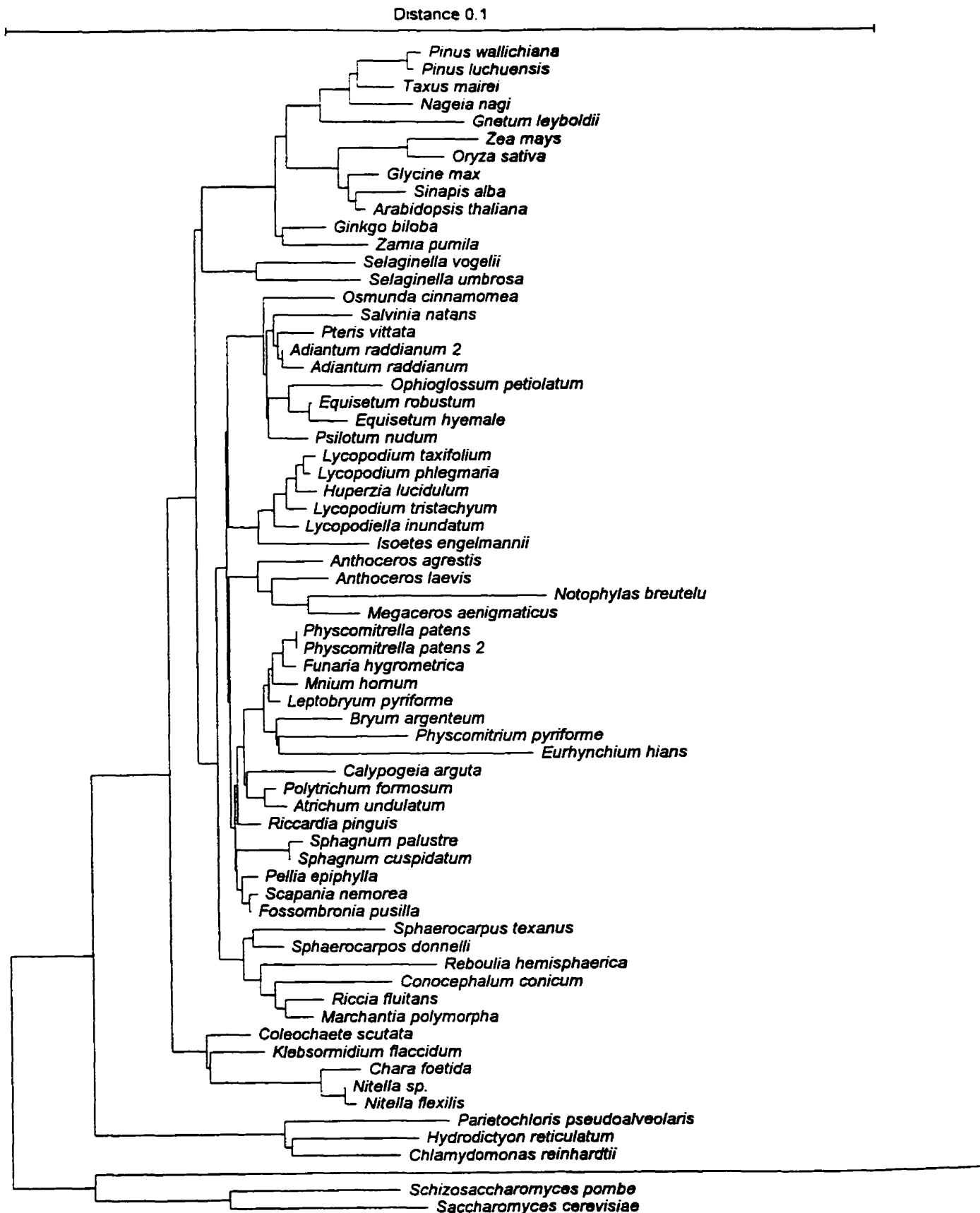


Figure E2 Transversions Only; Neighbor-joining
Insertions & Deletions taken into account

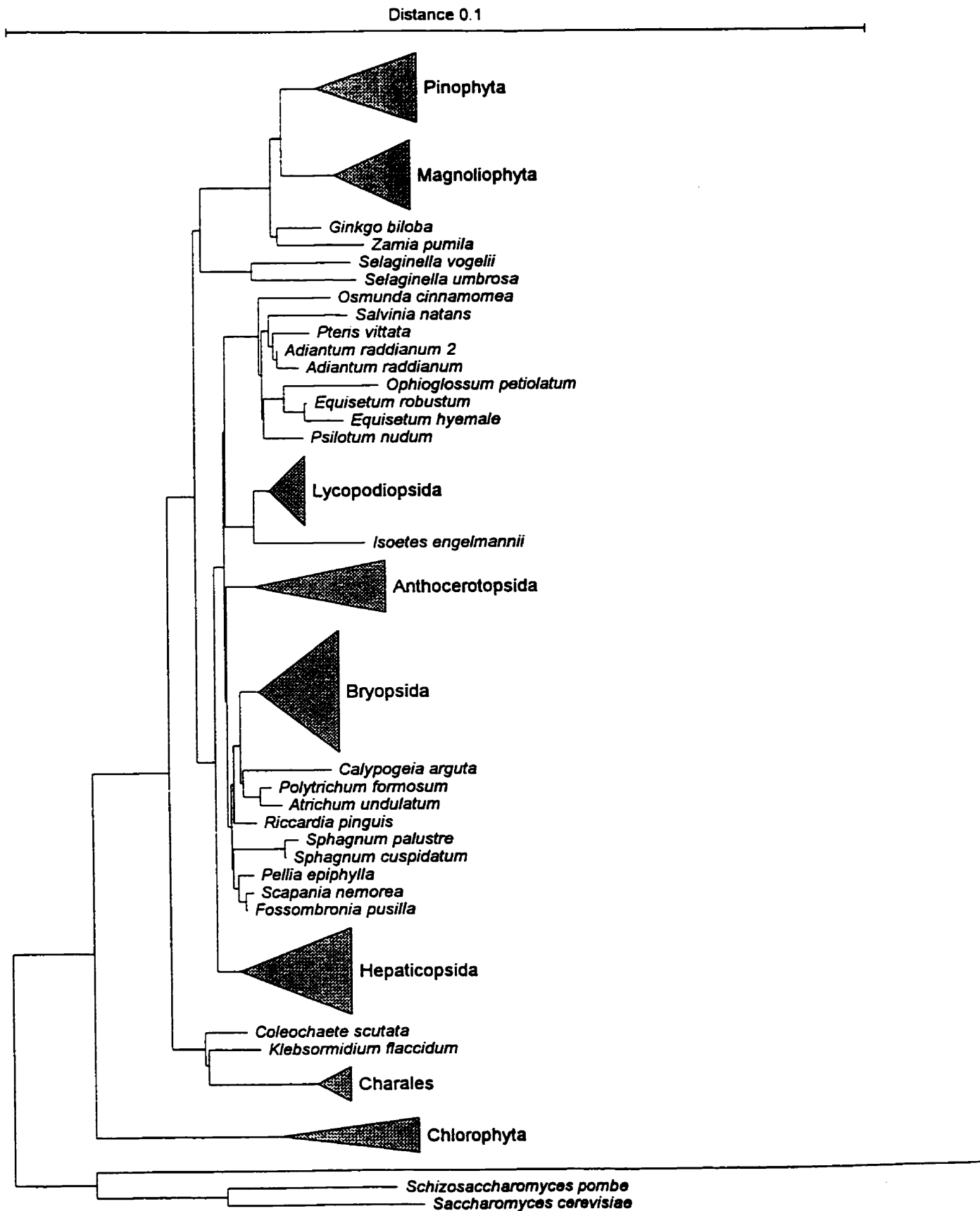


Figure E2 Transversions Only; Neighbor-joining
Insertions & Deletions taken into account

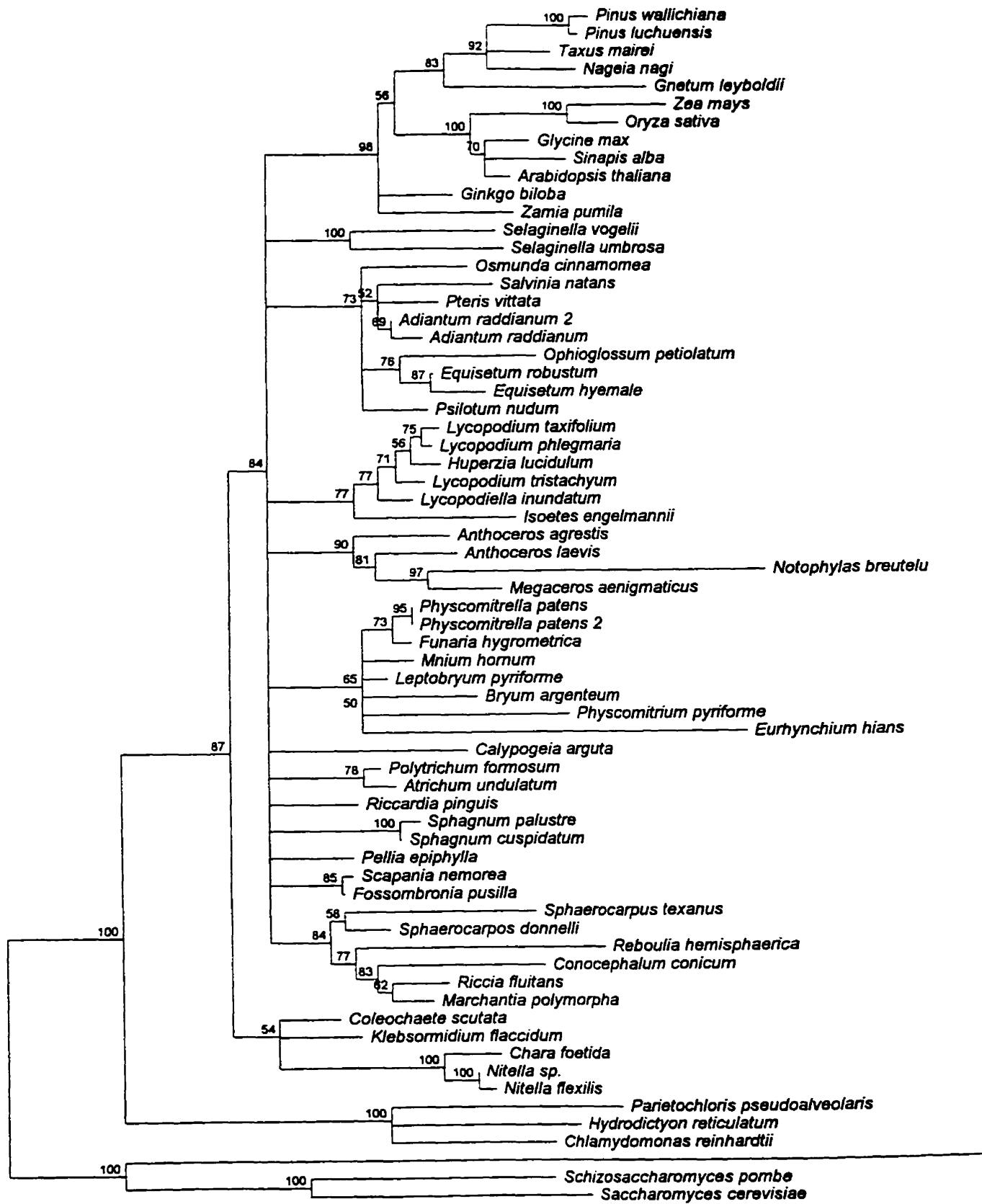


Figure E2B Transversions Only; Neighbor-joining
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 50% Majority-Rule Bootstrap Tree

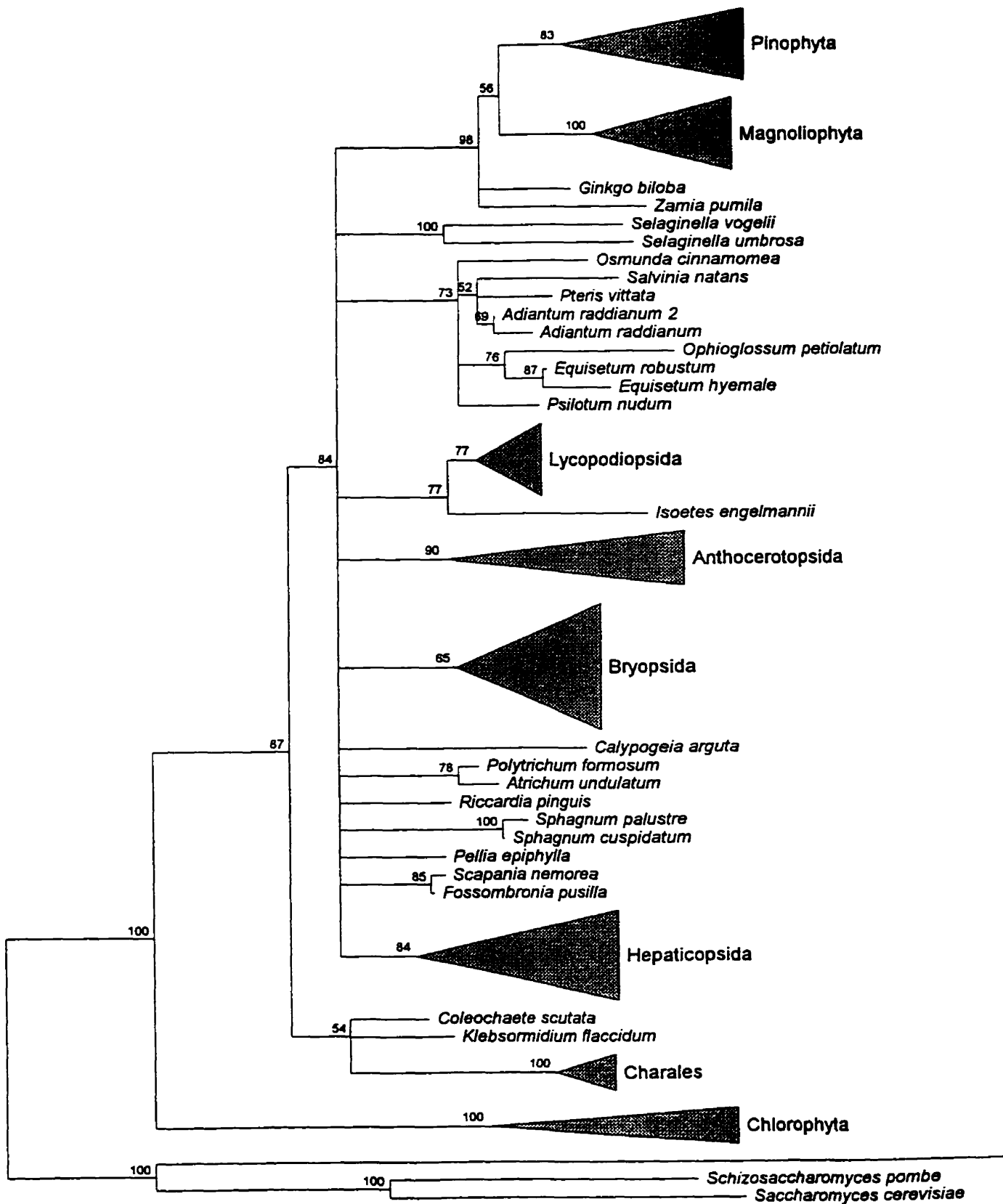


Figure E2B Transversions Only; Neighbor-joining
 Insertions & Deletions taken into account
 50% Majority-Rule Bootstrap Tree

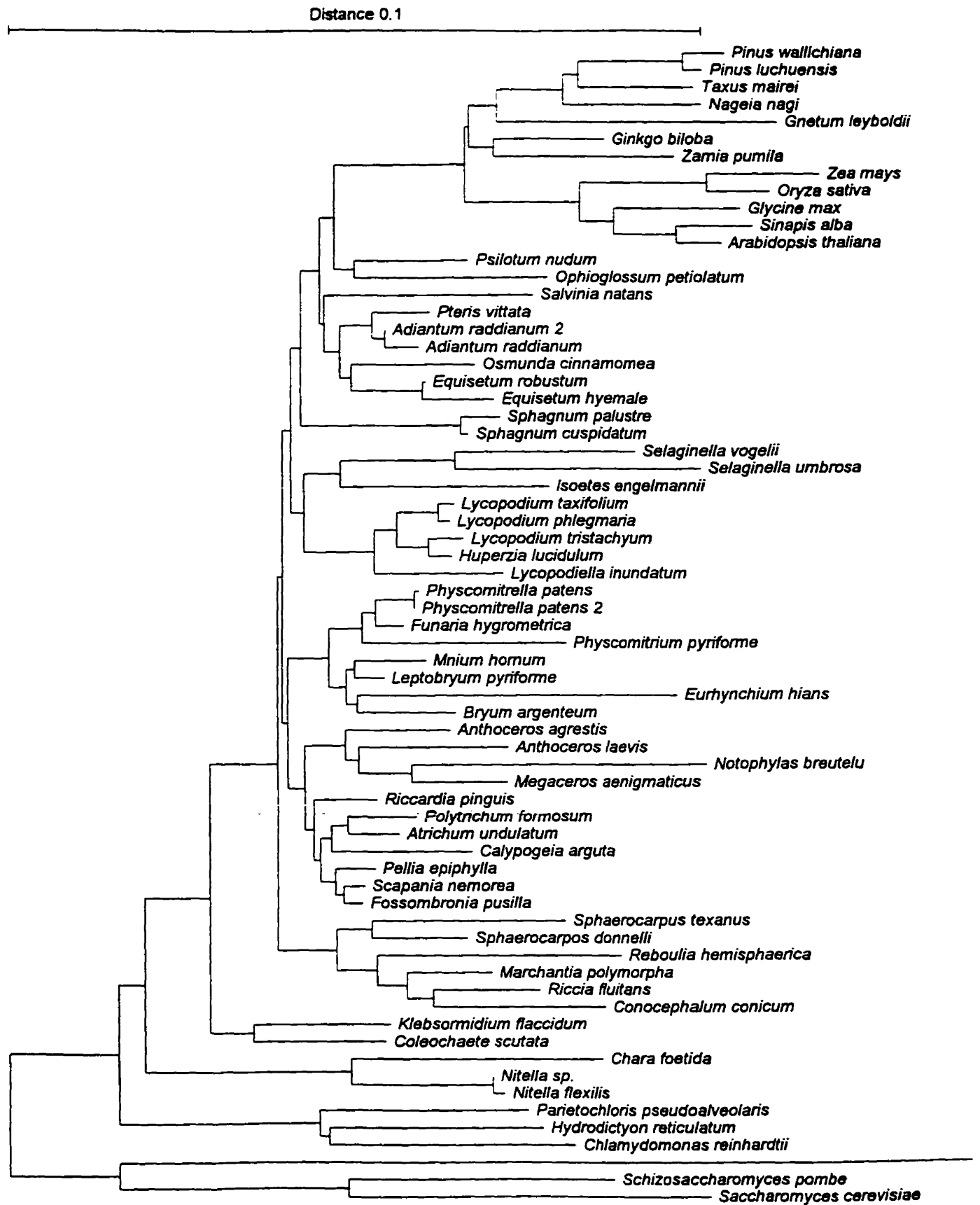


Figure F1 No Correction; Neighbor-joining

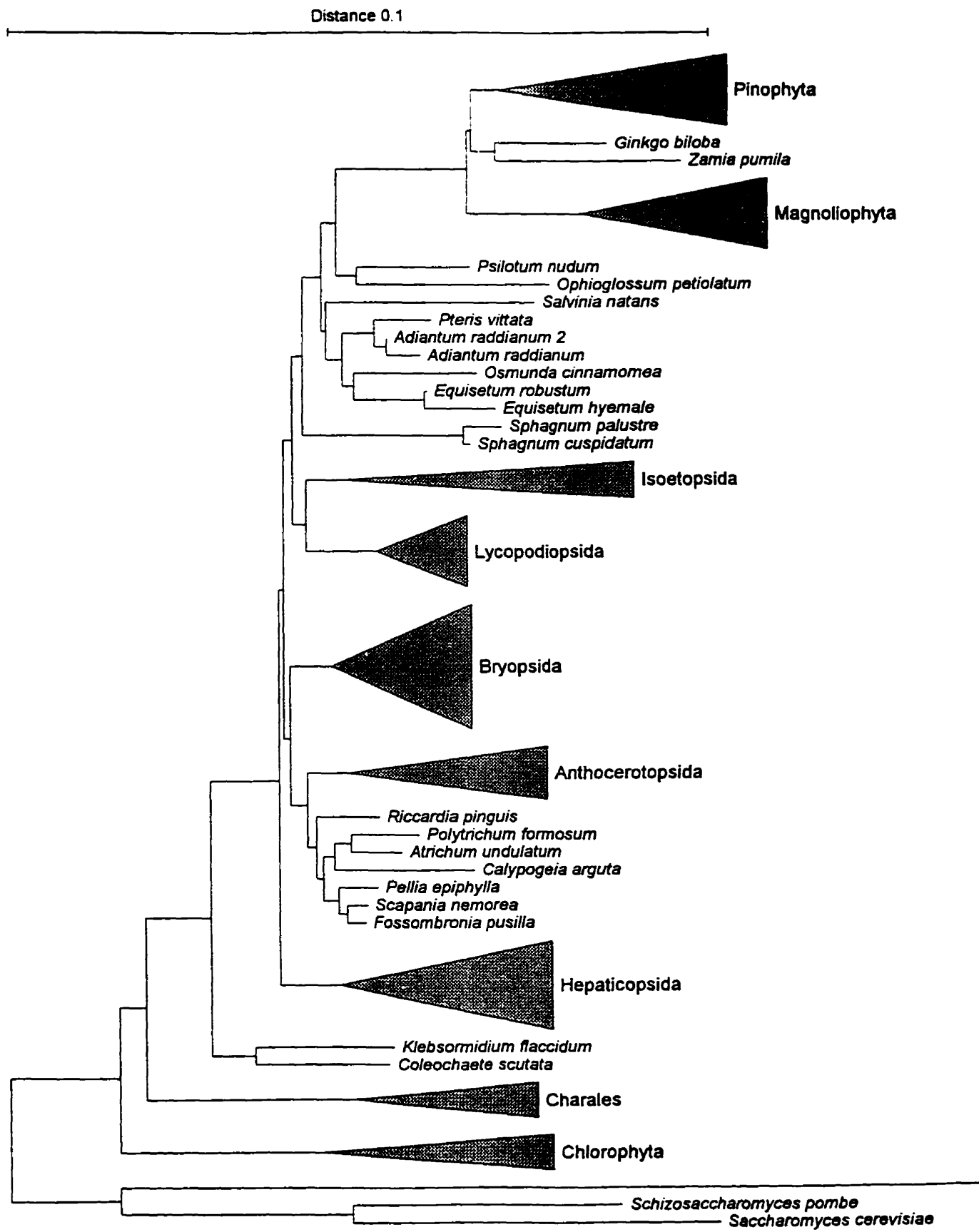


Figure F1 No Correction; Neighbor-joining

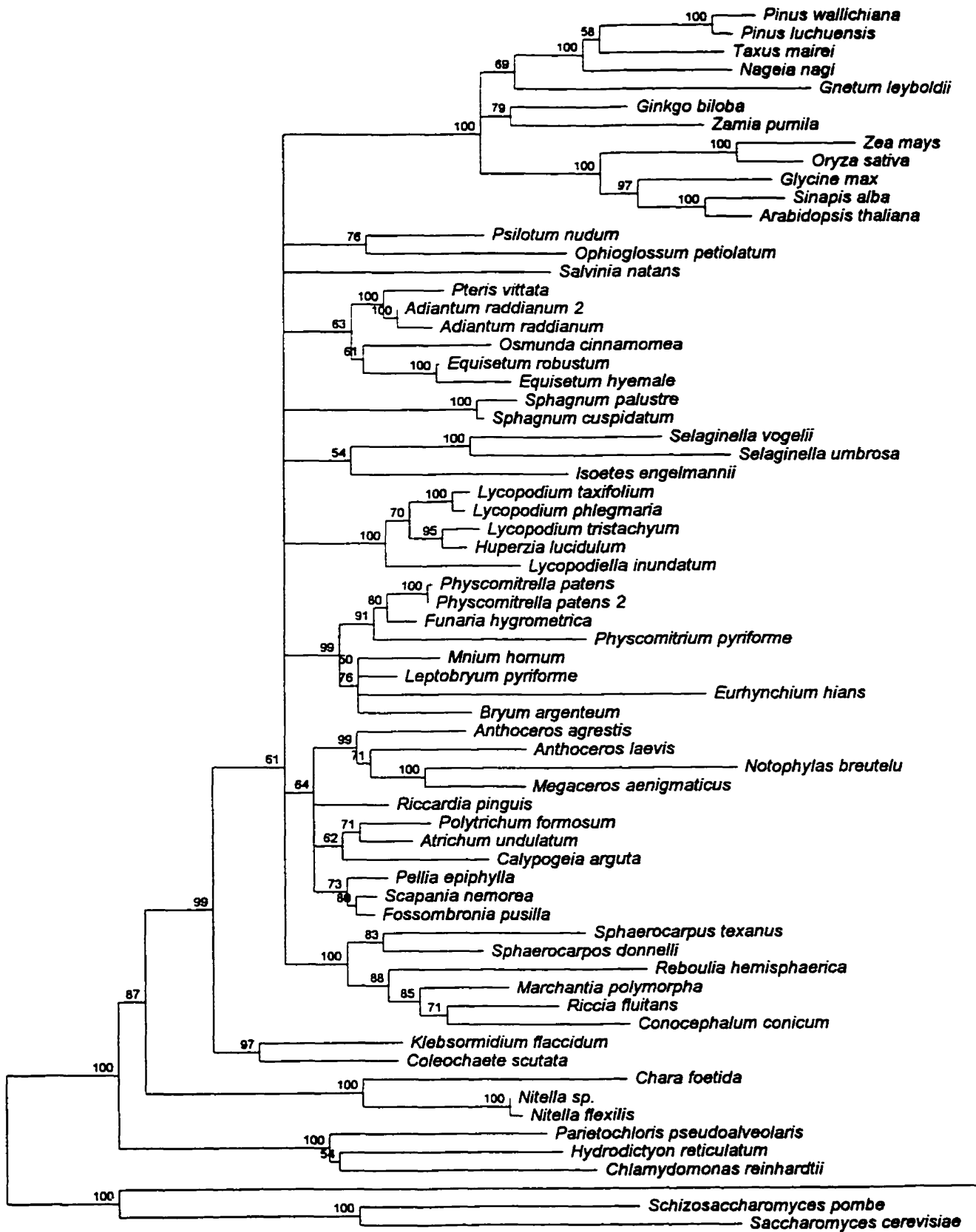


Figure F1B No Correction; Neighbor-joining
50% Majority-Rule Bootstrap Tree

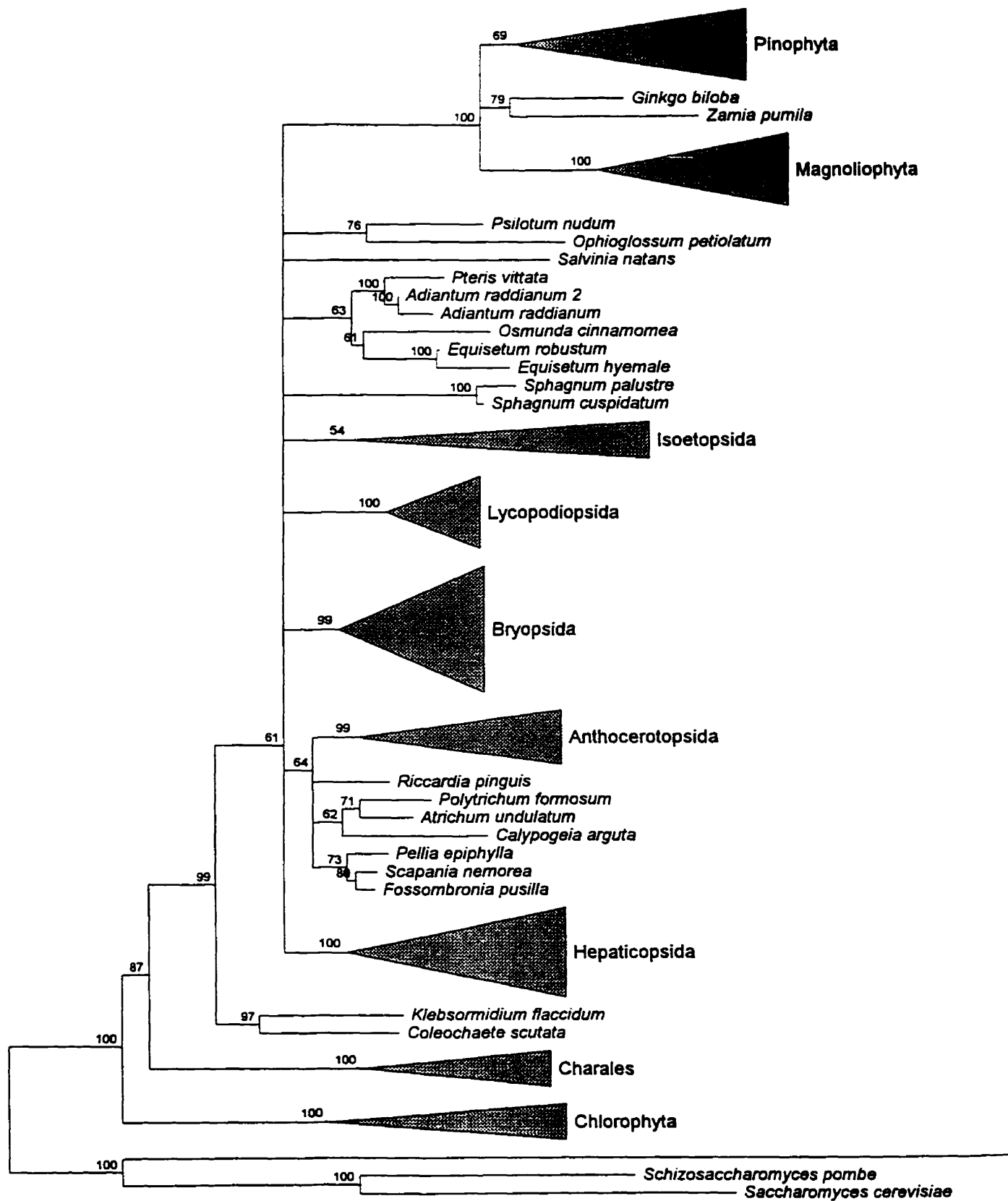


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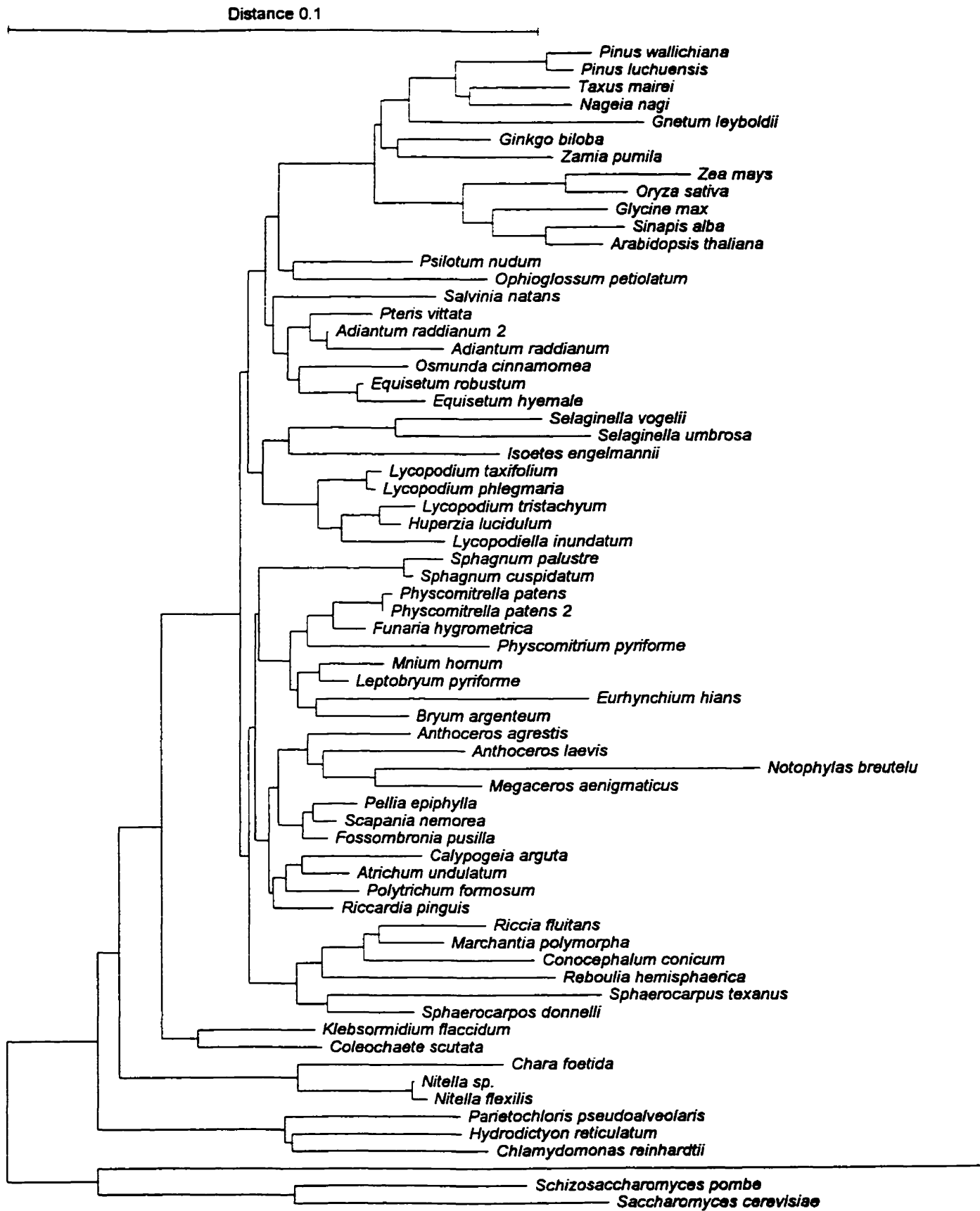


Figure F2 No Correction; Neighbor-joining
Insertions & Deletions taken into account

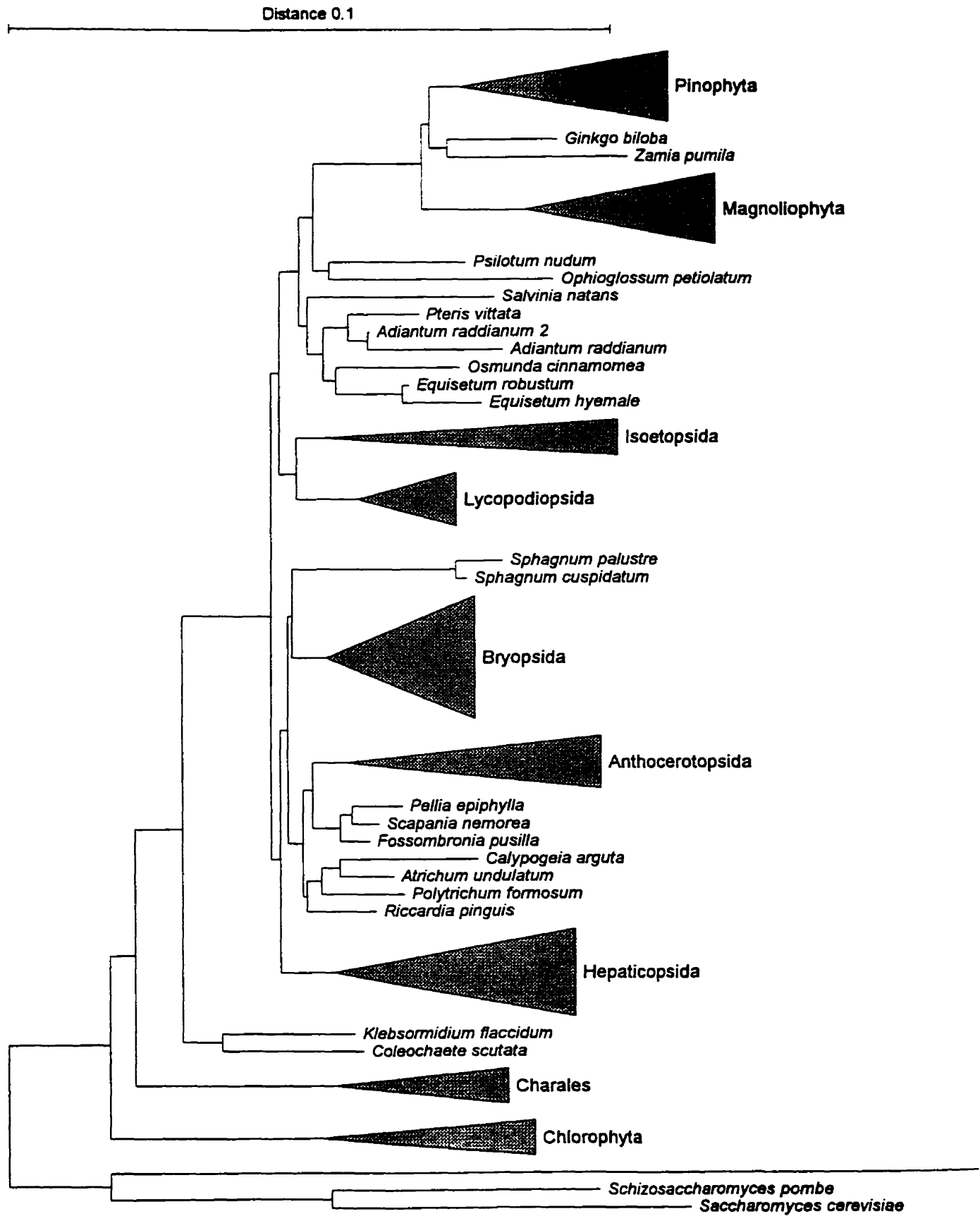


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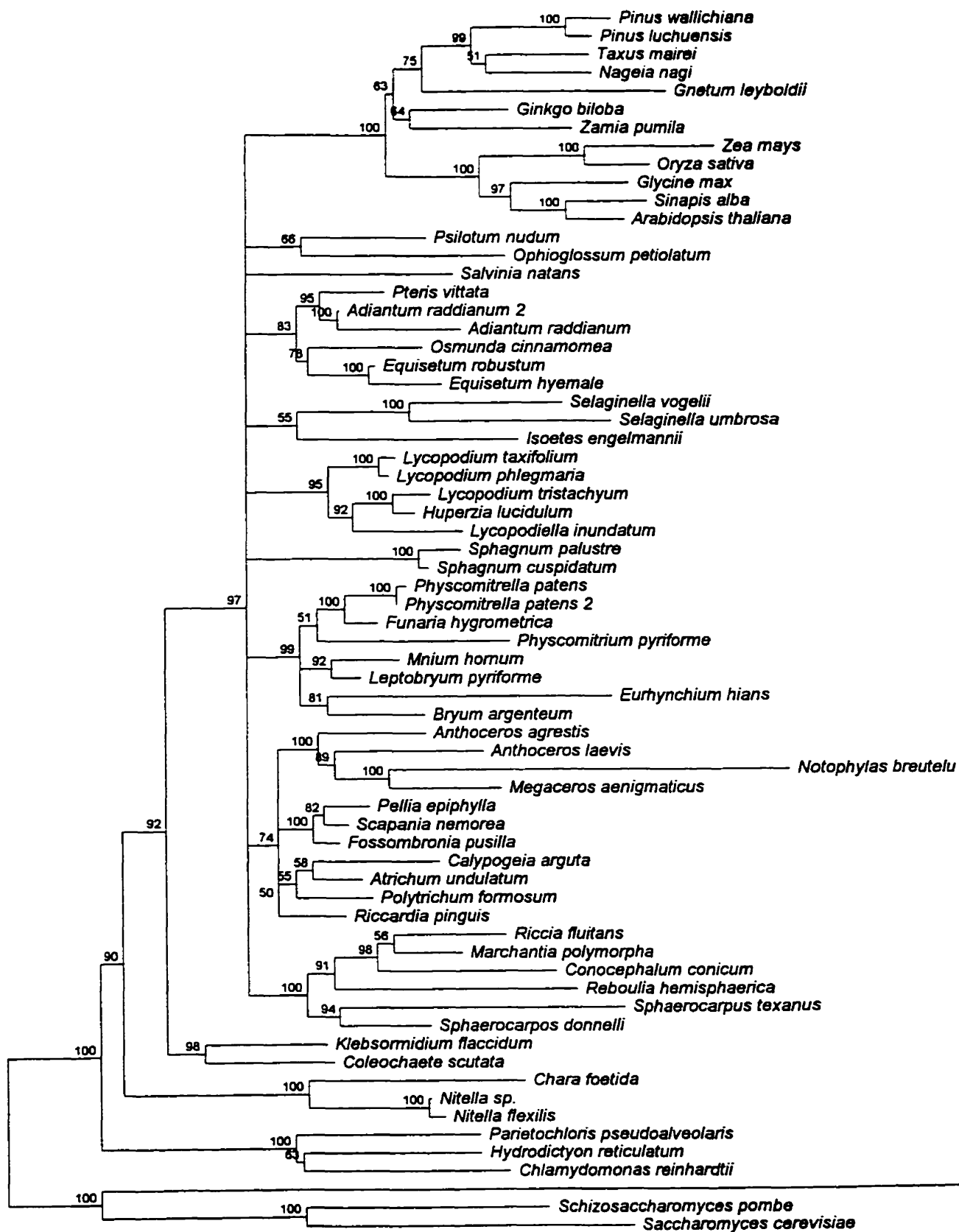


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 Insertions & Deletions taken into account
 50% Majority-Rule Bootstrap Tree

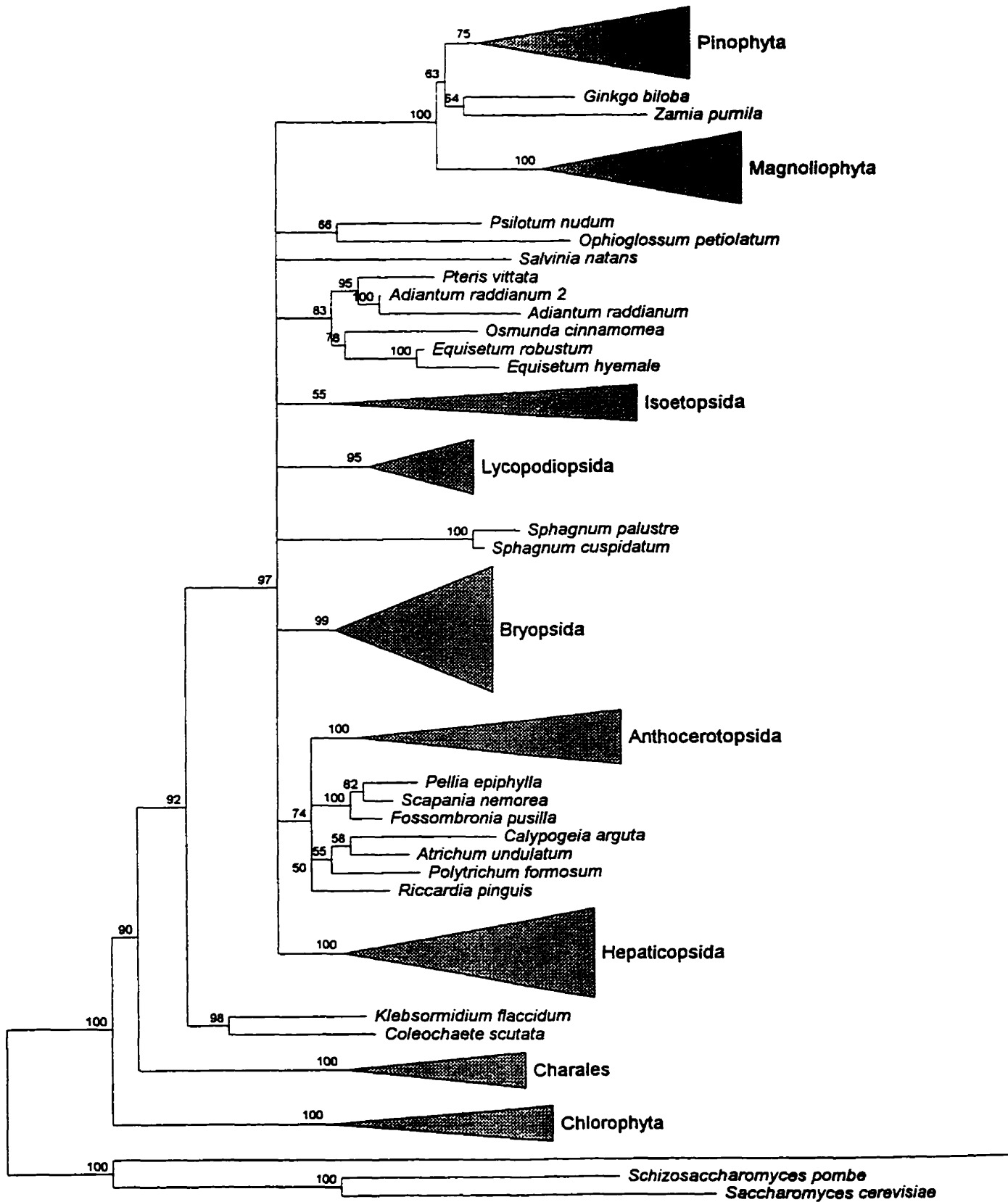


Figure F2B No Correction; Neighbor-joining
 Insertions & Deletions taken into account
 50% Majority-Rule Bootstrap Tree

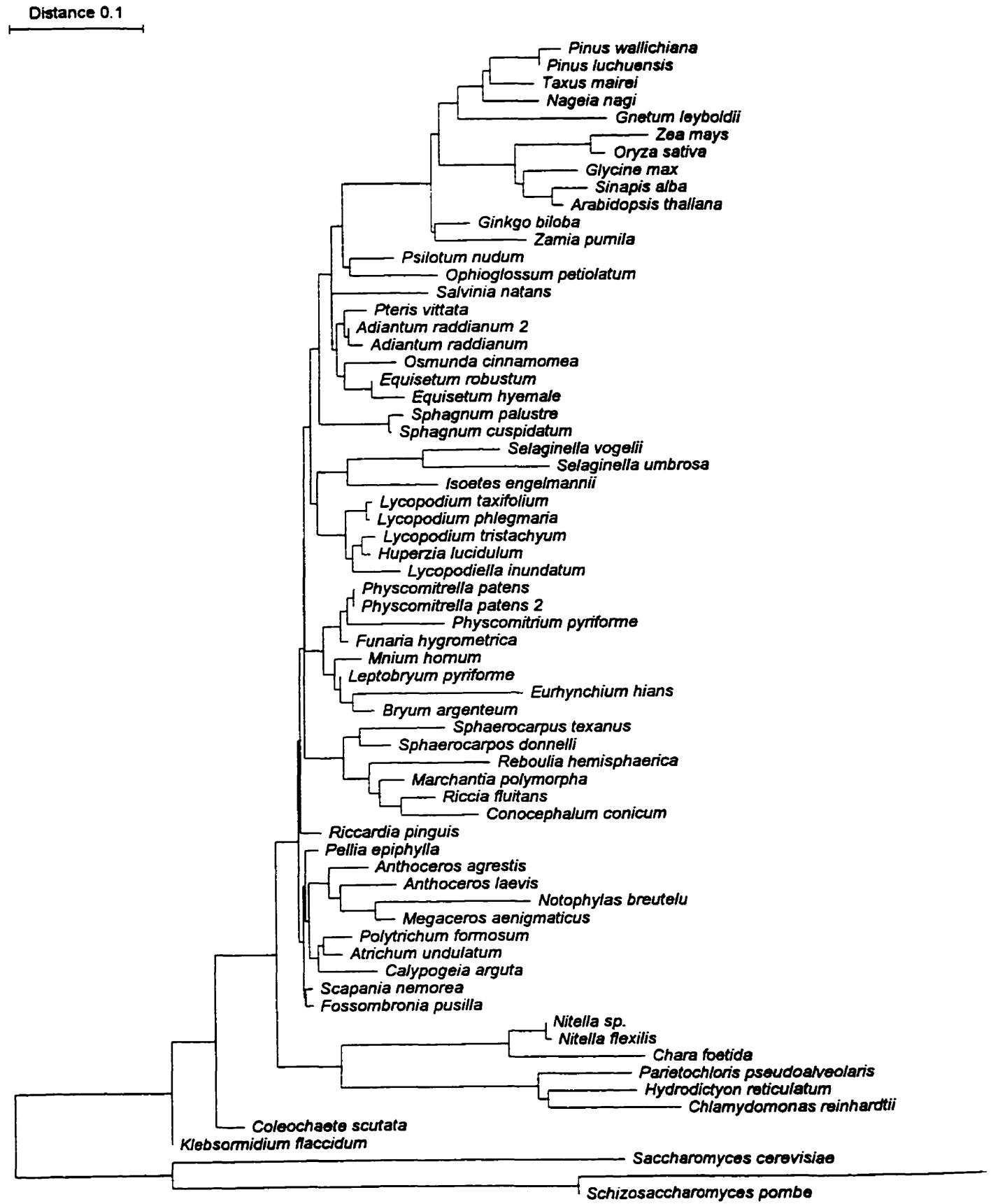


Figure G1 Van de Peer & De Wachter, Neighbor-joining

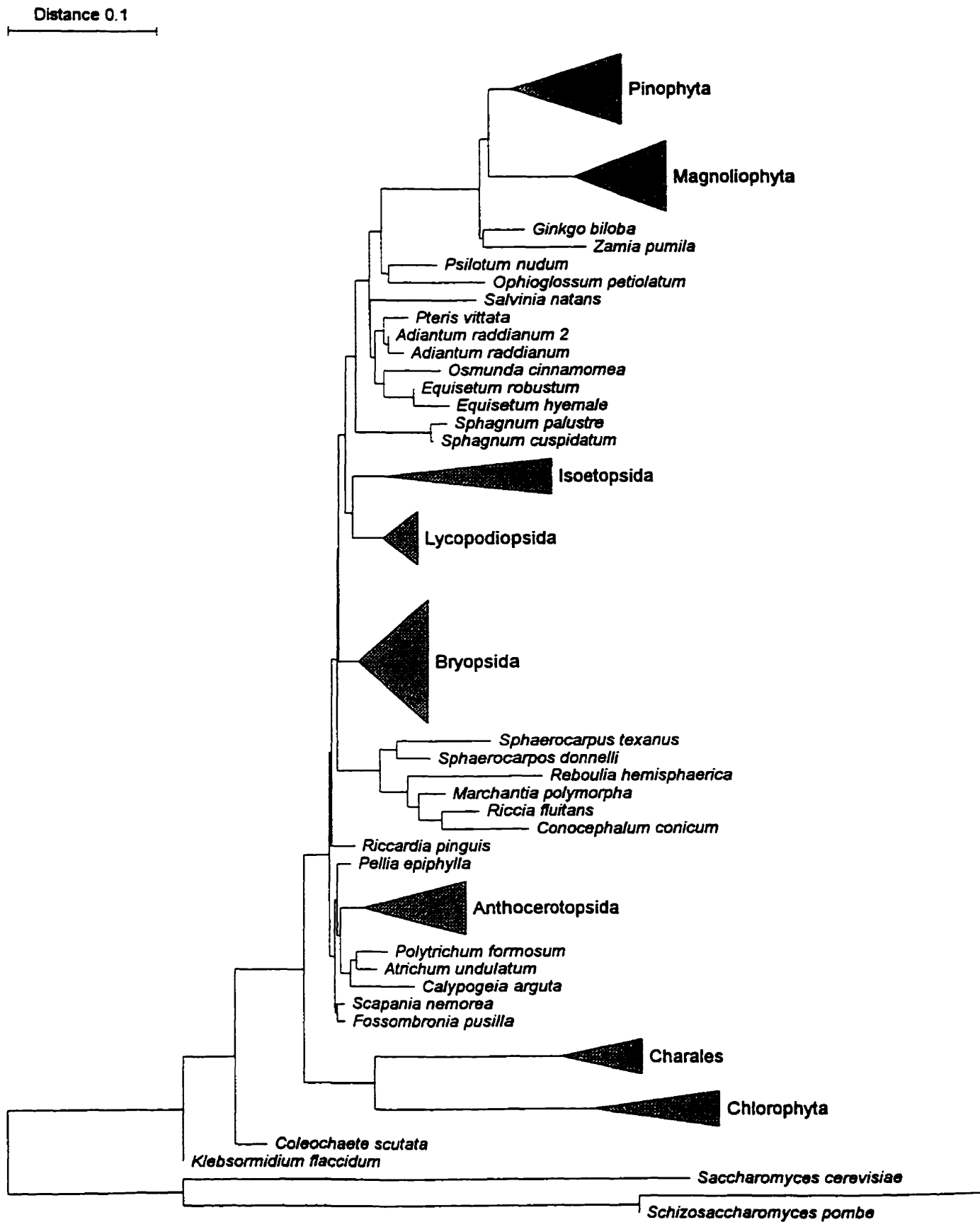


Figure G1 Van de Peer & De Wachter, Neighbor-joining

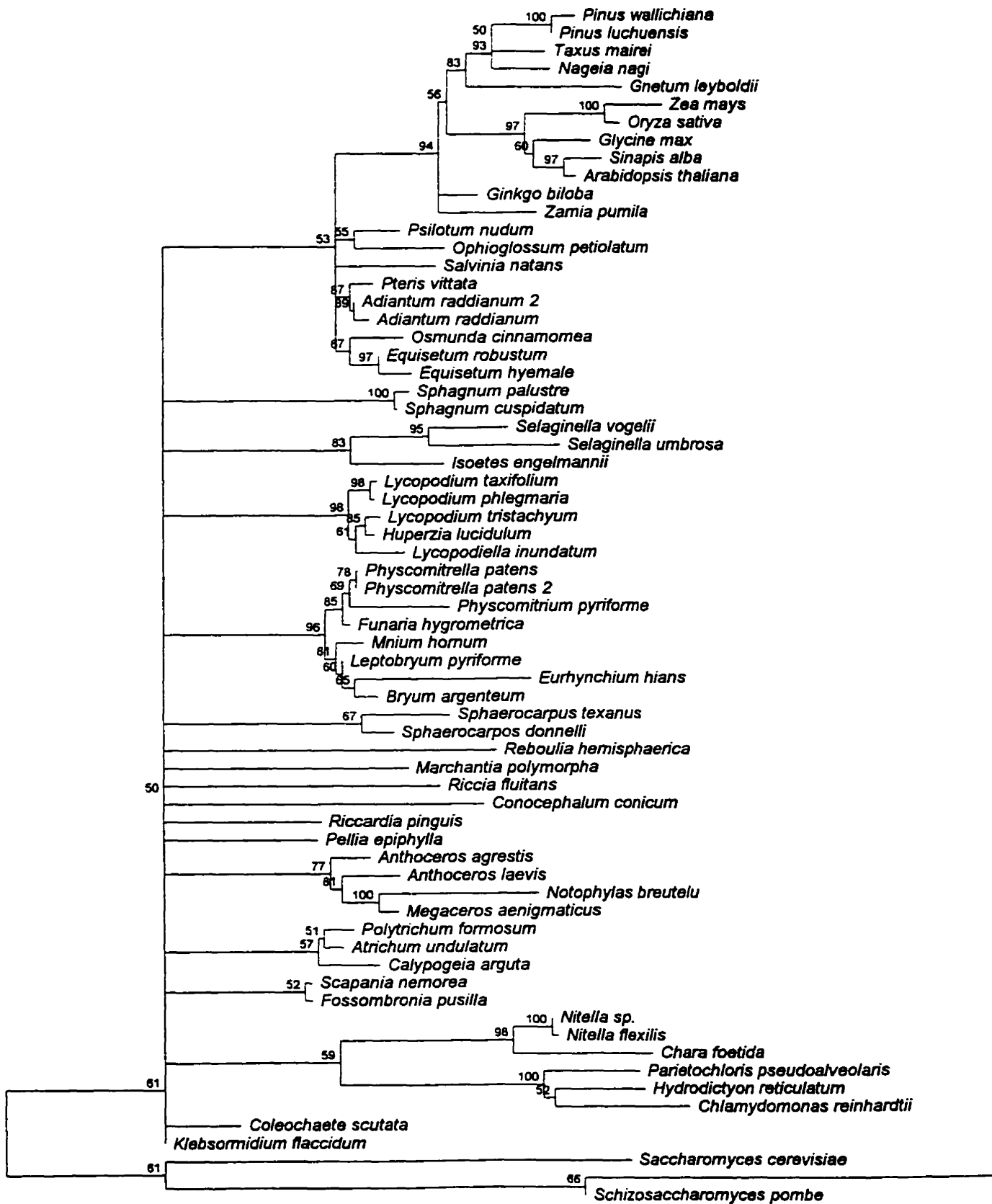


Figure G1B Van de Peer & De Wachter, Neighbor-joining
50% Majority-Rule Bootstrap Tree

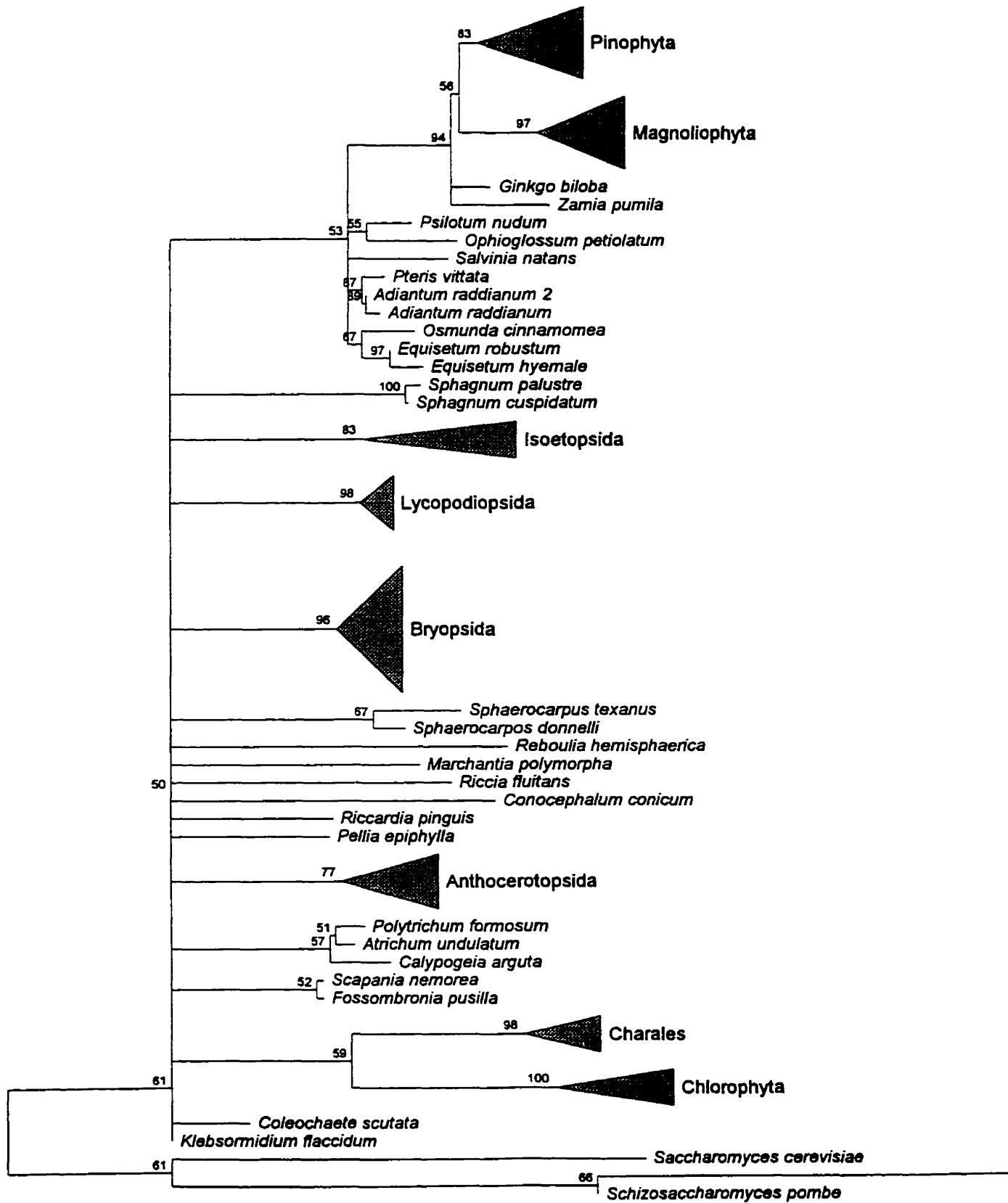


Figure G1B Van de Peer & De Wachter, Neighbor-joining
50% Majority-Rule Bootstrap Tree

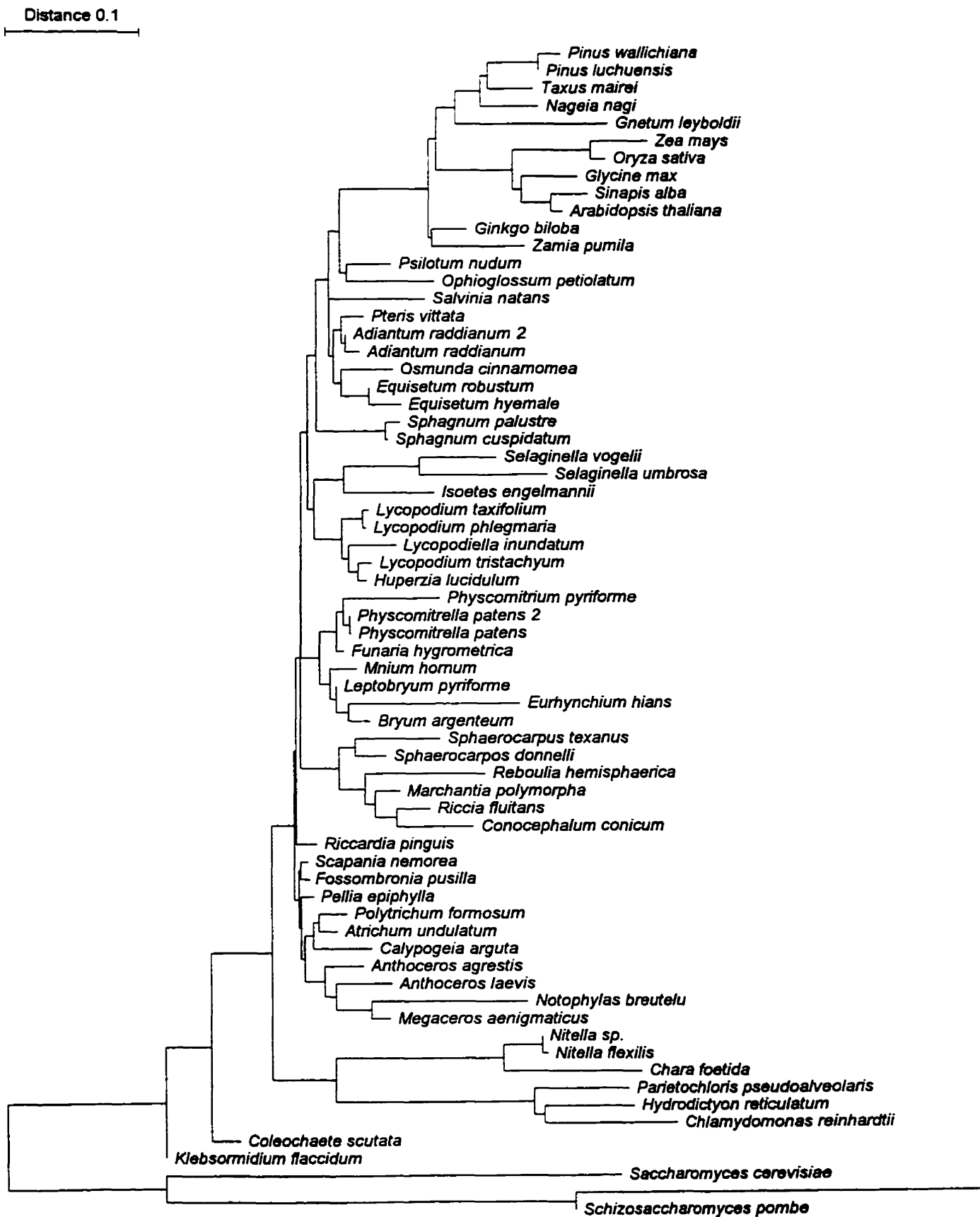


Figure G2 Van de Peer & De Wachter
Insertions & Deletions taken into account

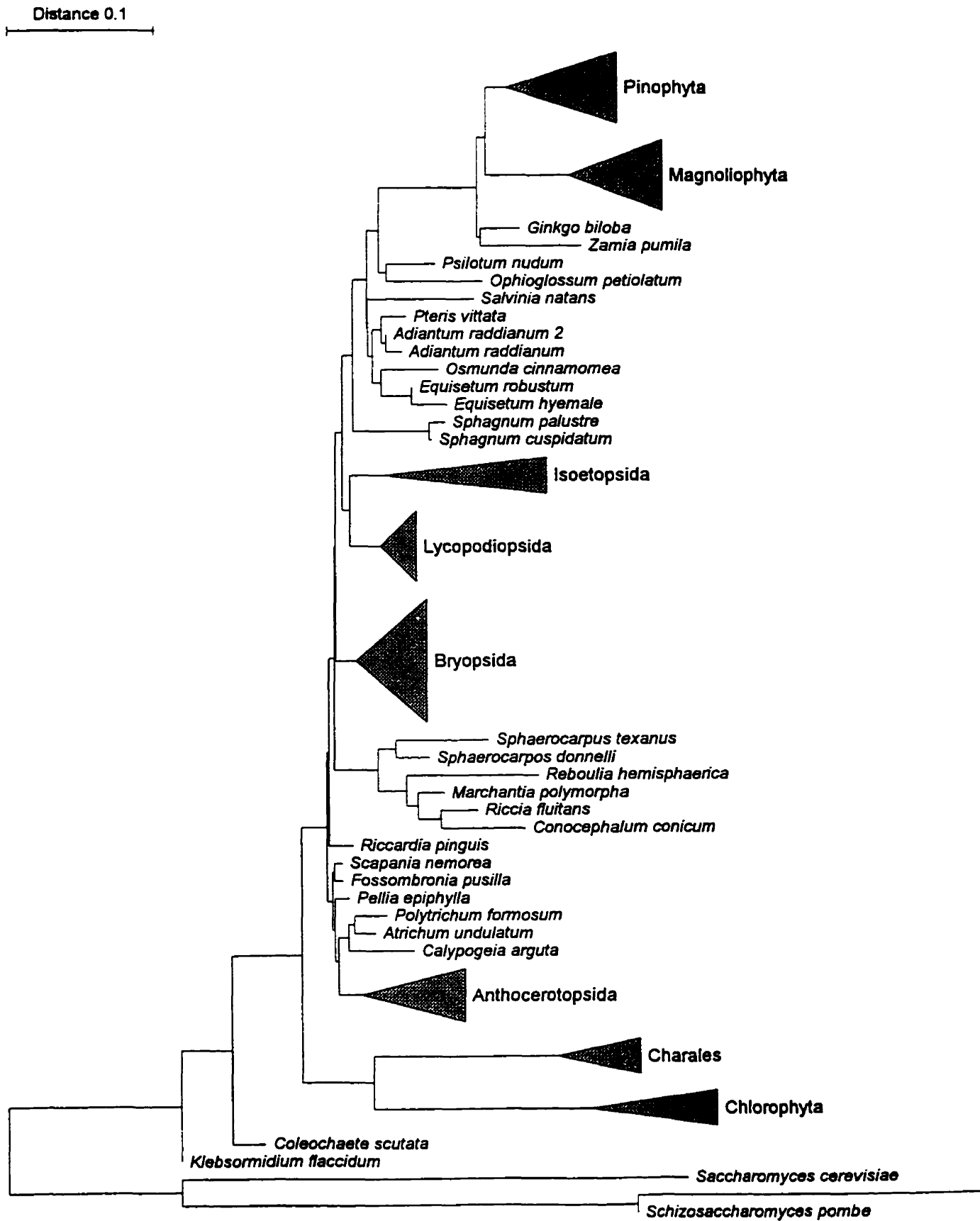


Figure G2 Van de Peer & De Wachter
Insertions & Deletions taken into account

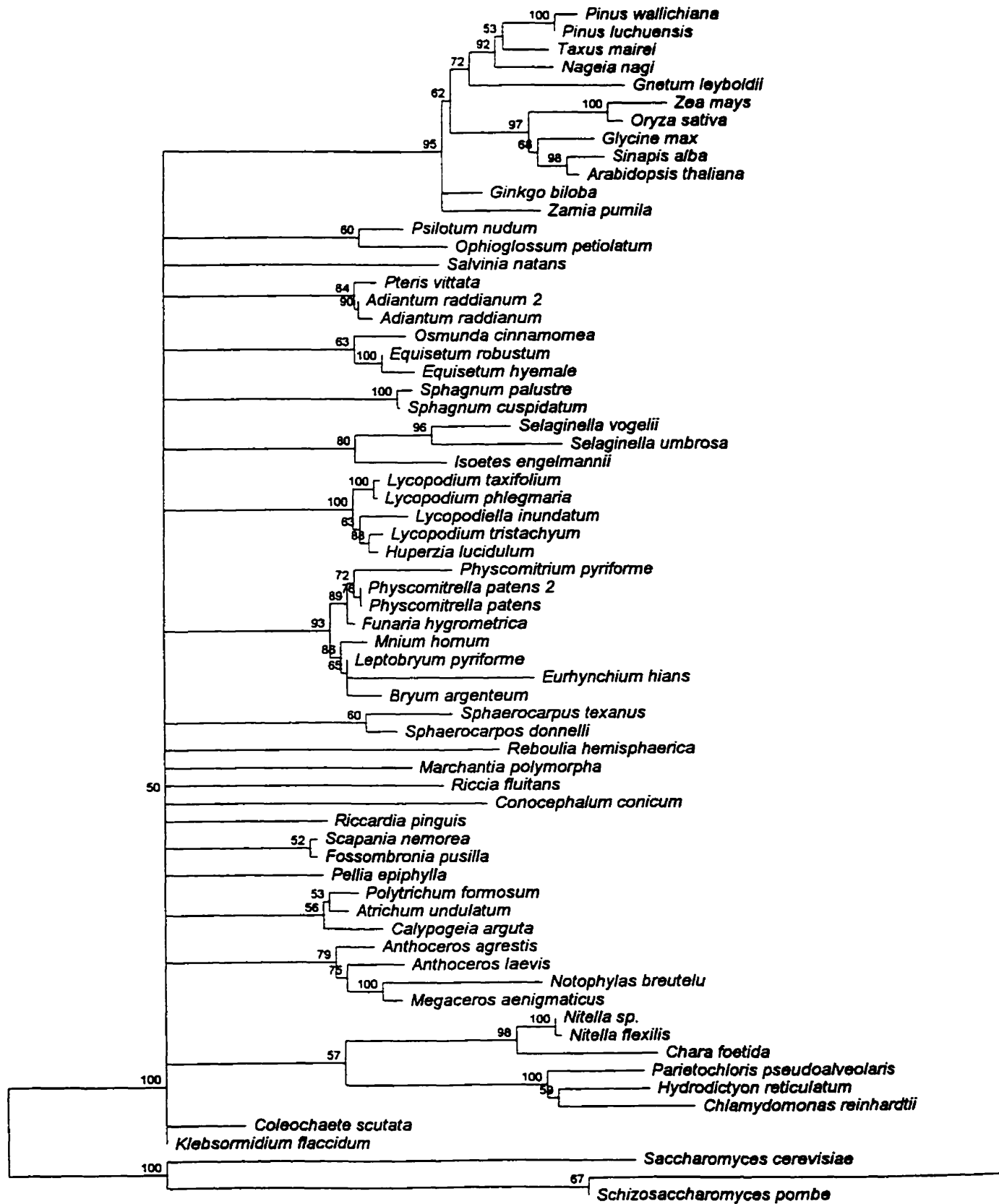


Figure G2B Van de Peer & De Wachter
 Insertions & Deletions taken into account
 50% Majority-Rule Bootstrap Tree

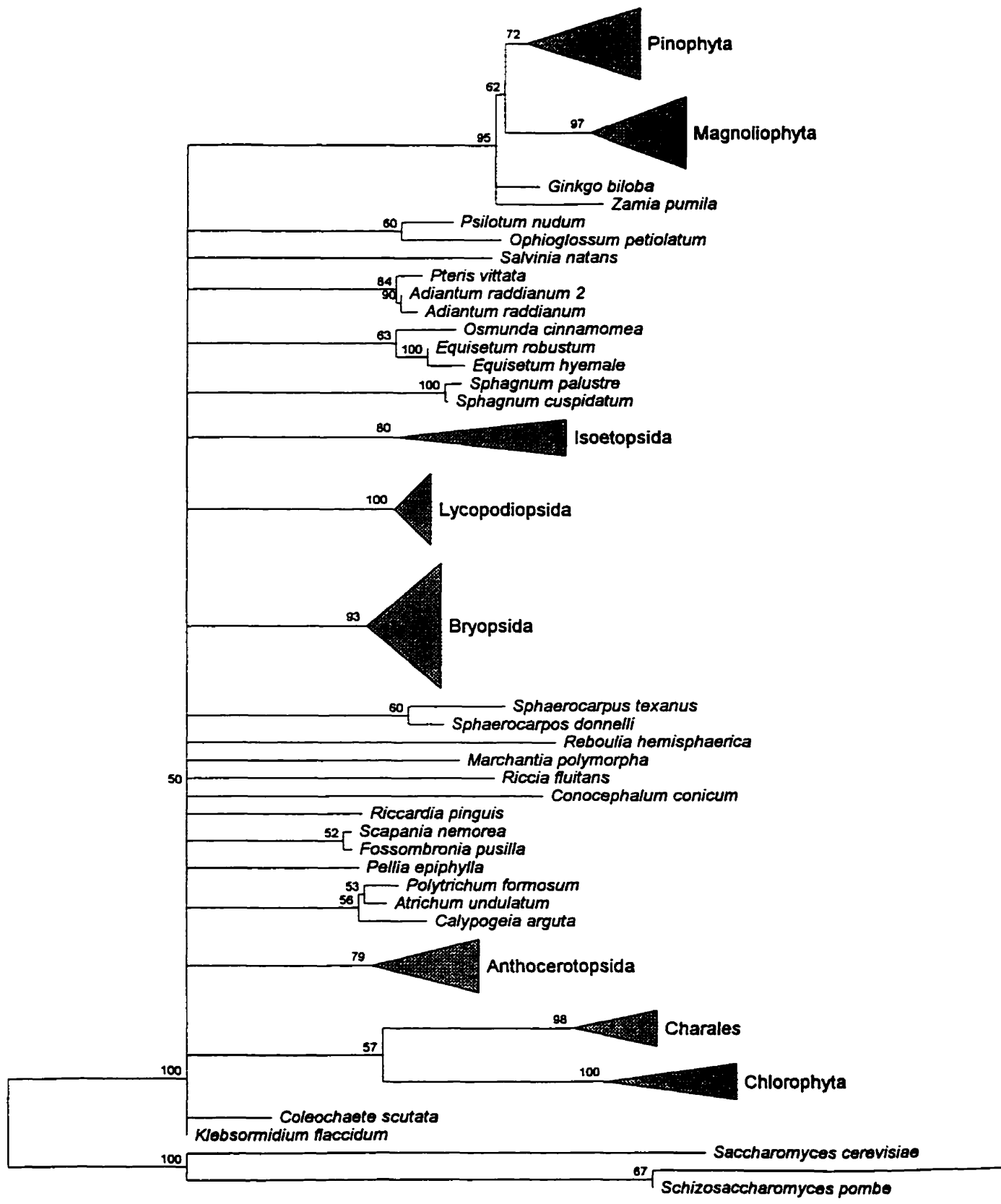


Figure G2B Van de Peer & De Wachter
 Insertions & Deletions taken into account
 50% Majority-Rule Bootstrap Tree

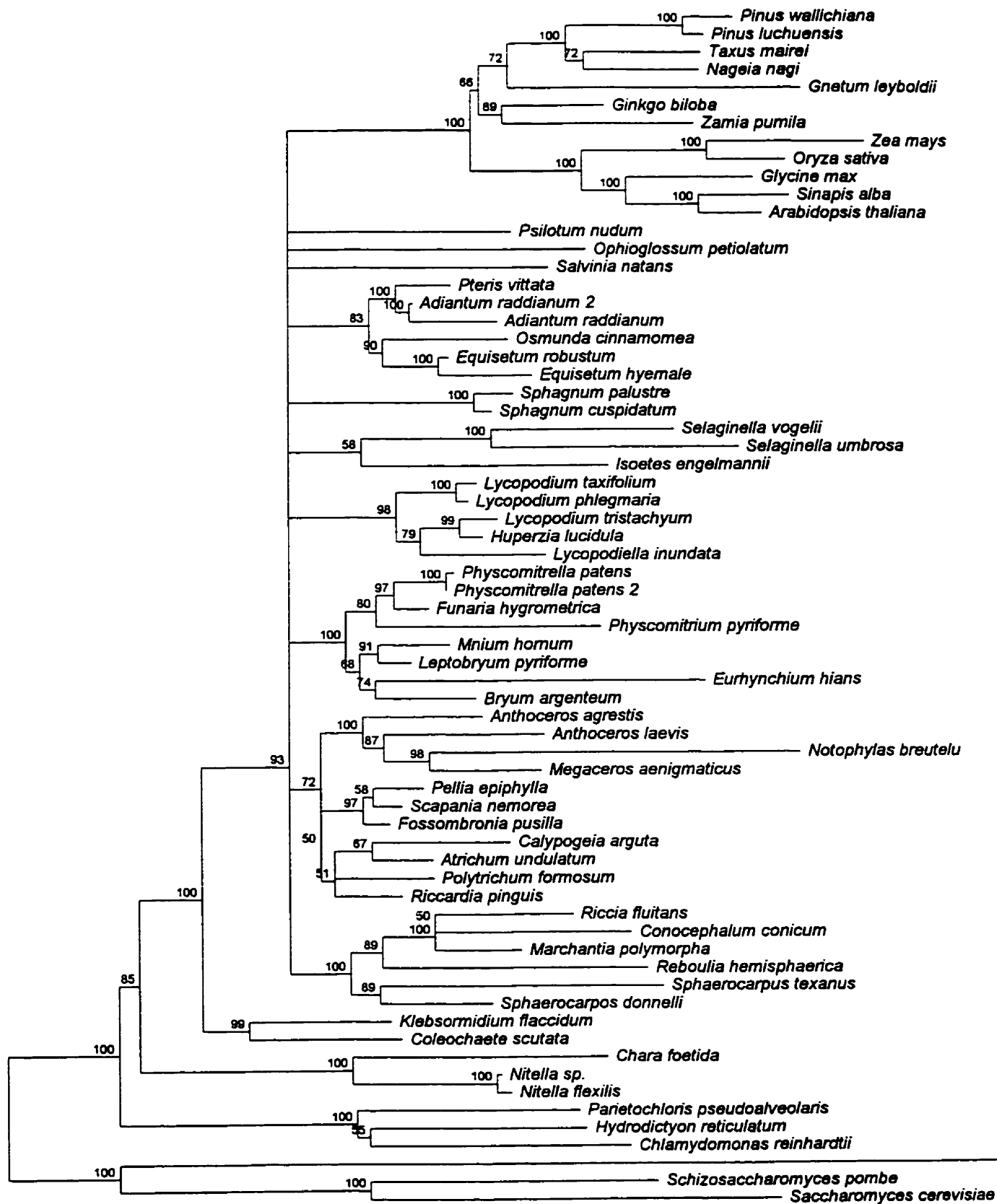


Figure H1B No Correction; Neighbor-joining
 Insertions & Deletions taken into account
 50% Majority-Rule Bootstrap Tree
 Only Categories 0, 1, 2, 3, 4

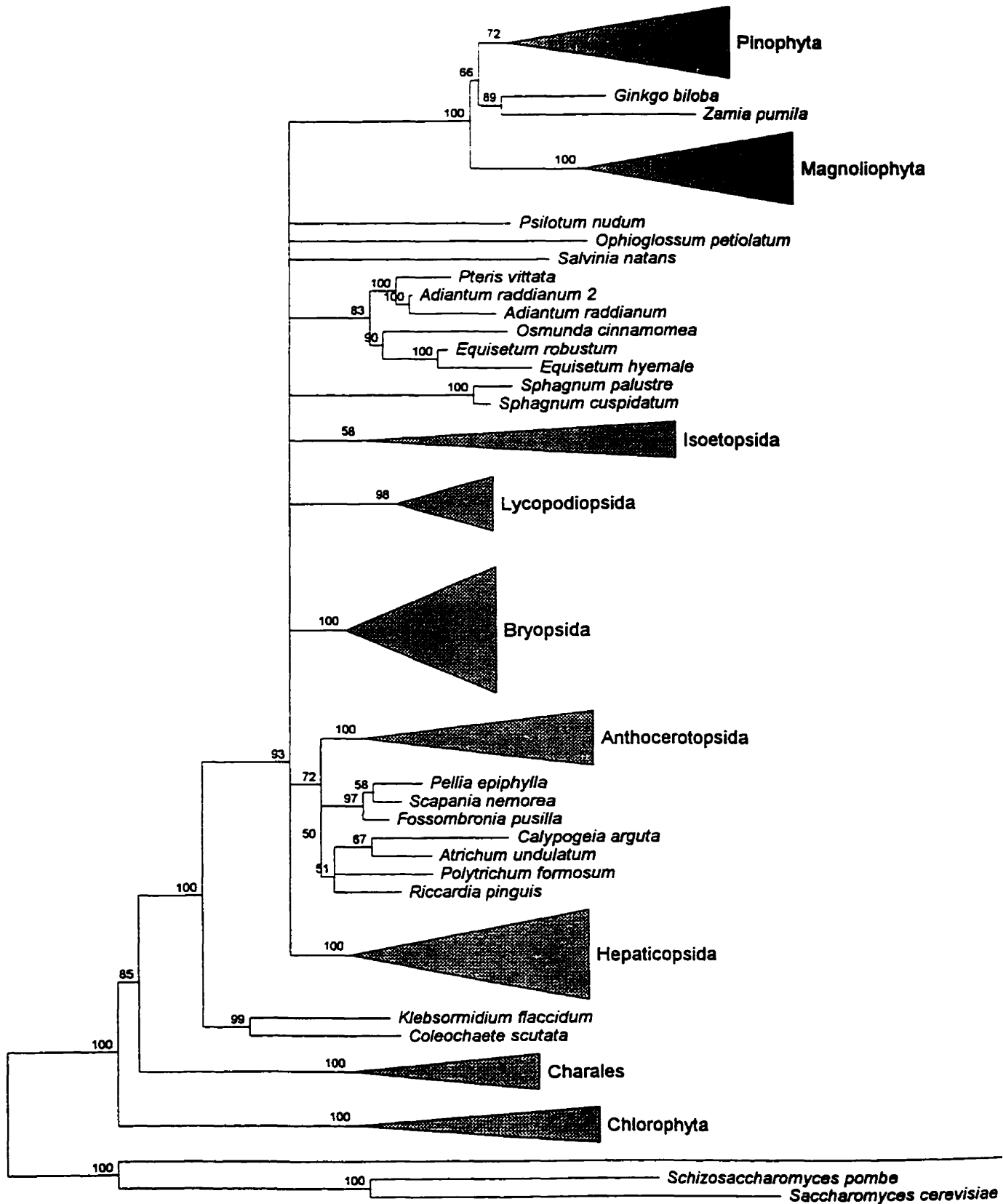


Figure H1B No Correction; Neighbor-joining
 Insertions & Deletions taken into account
 50% Majority-Rule Bootstrap Tree
 Only Categories 0, 1, 2, 3, 4

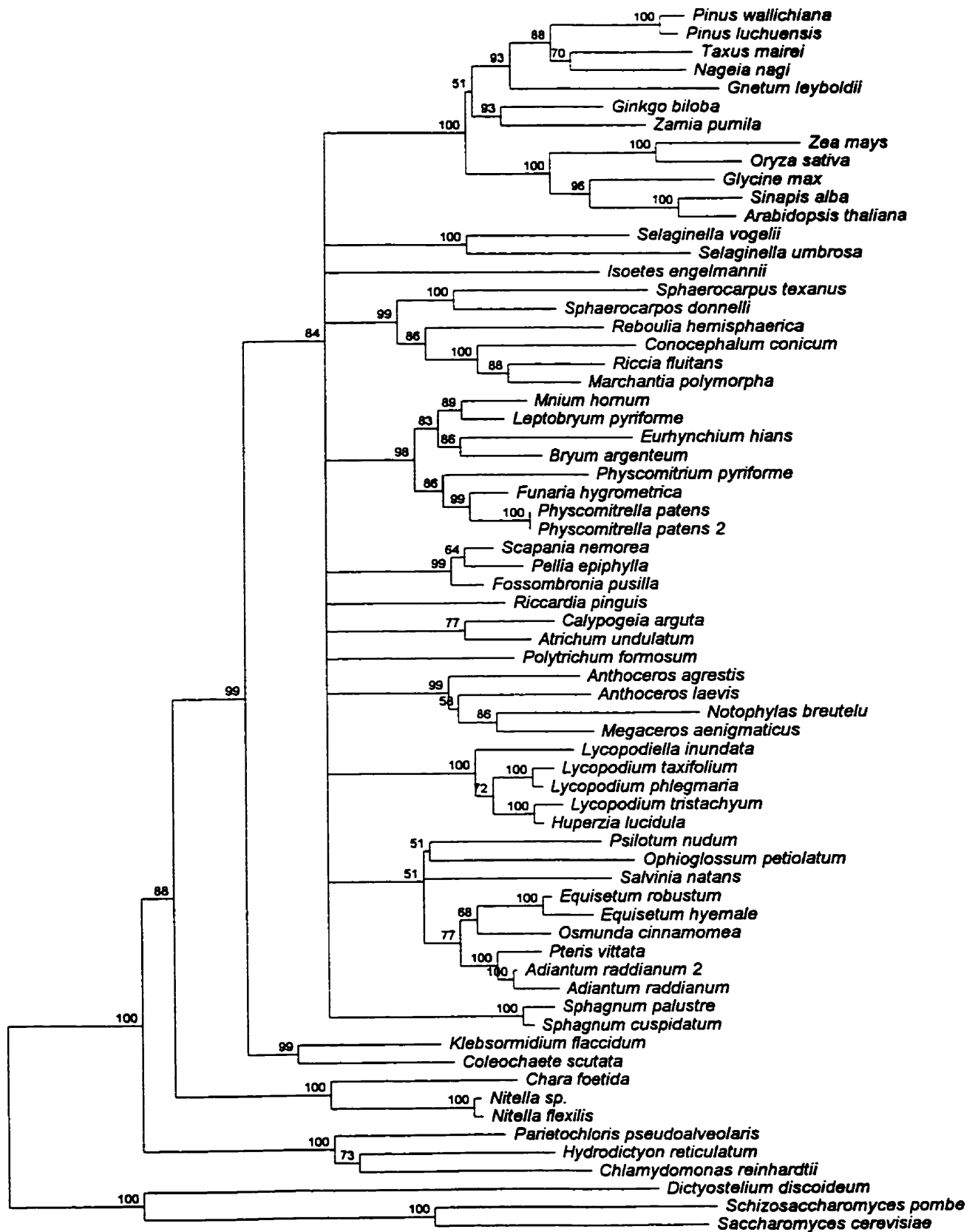


Figure H2B No Correction; Neighbor-joining
 Insertions & Deletions taken into account
 50% Majority-Rule Bootstrap Tree
 Only Categories 0, 1, 2

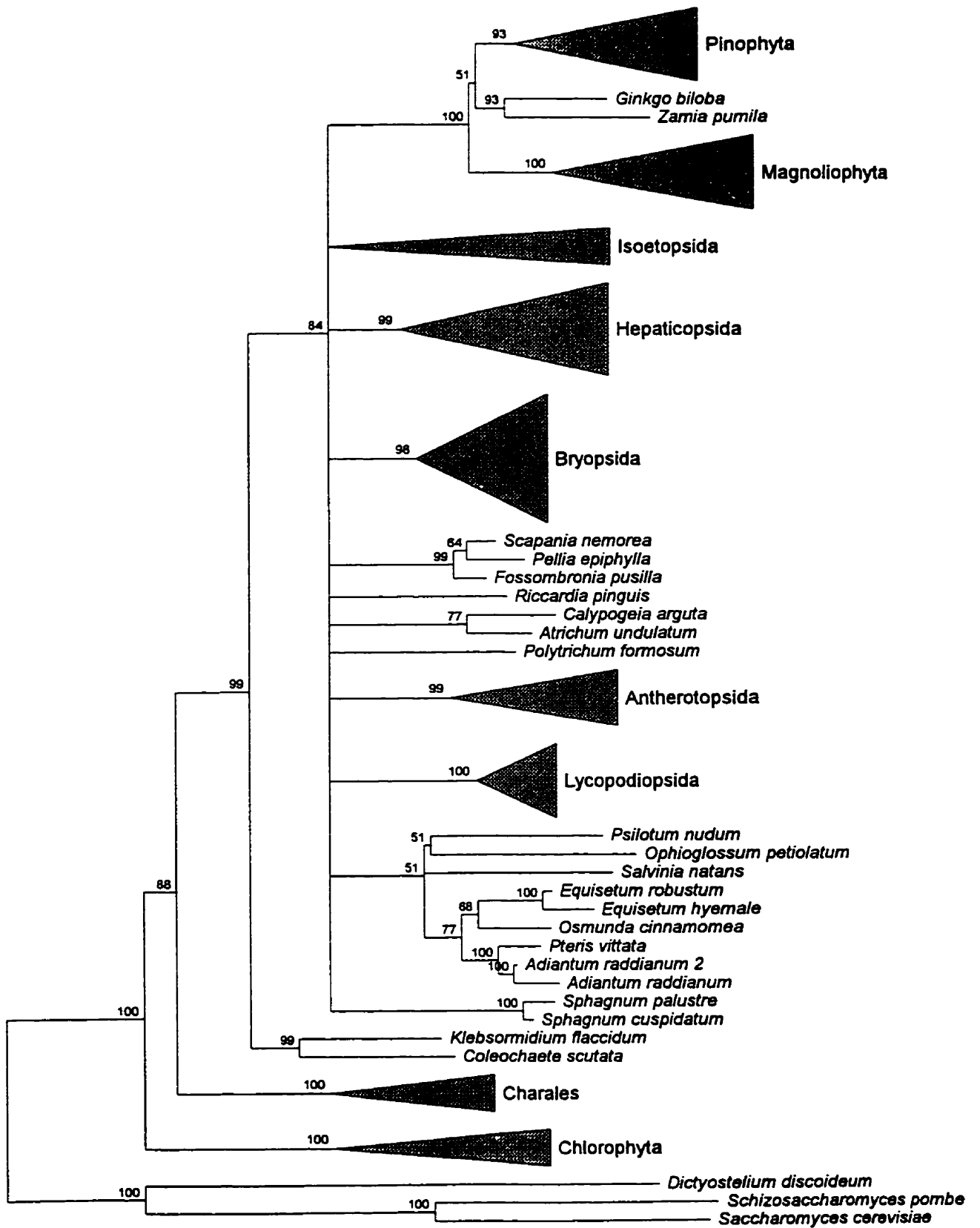


Figure H2B No Correction; Neighbor-joining
 Insertions & Deletions taken into account
 50% Majority-Rule Bootstrap Tree
 Only Categories 0, 1, 2

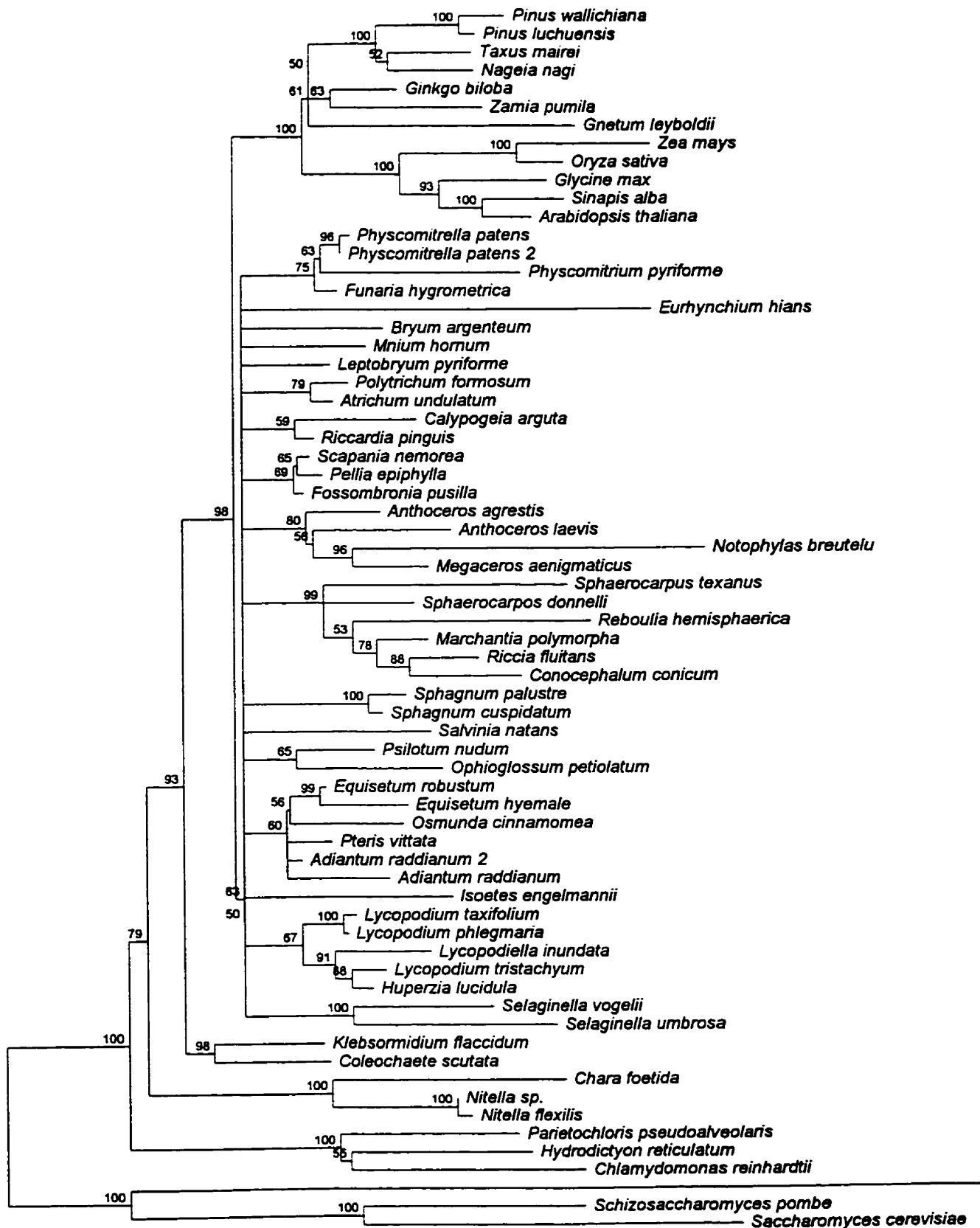


Figure H3B No Correction; Neighbor-joining
 Insertions & Deletions taken into account
 50% Majority-Rule Bootstrap Tree
 Only Categories 2, 3, 4, 5, 6

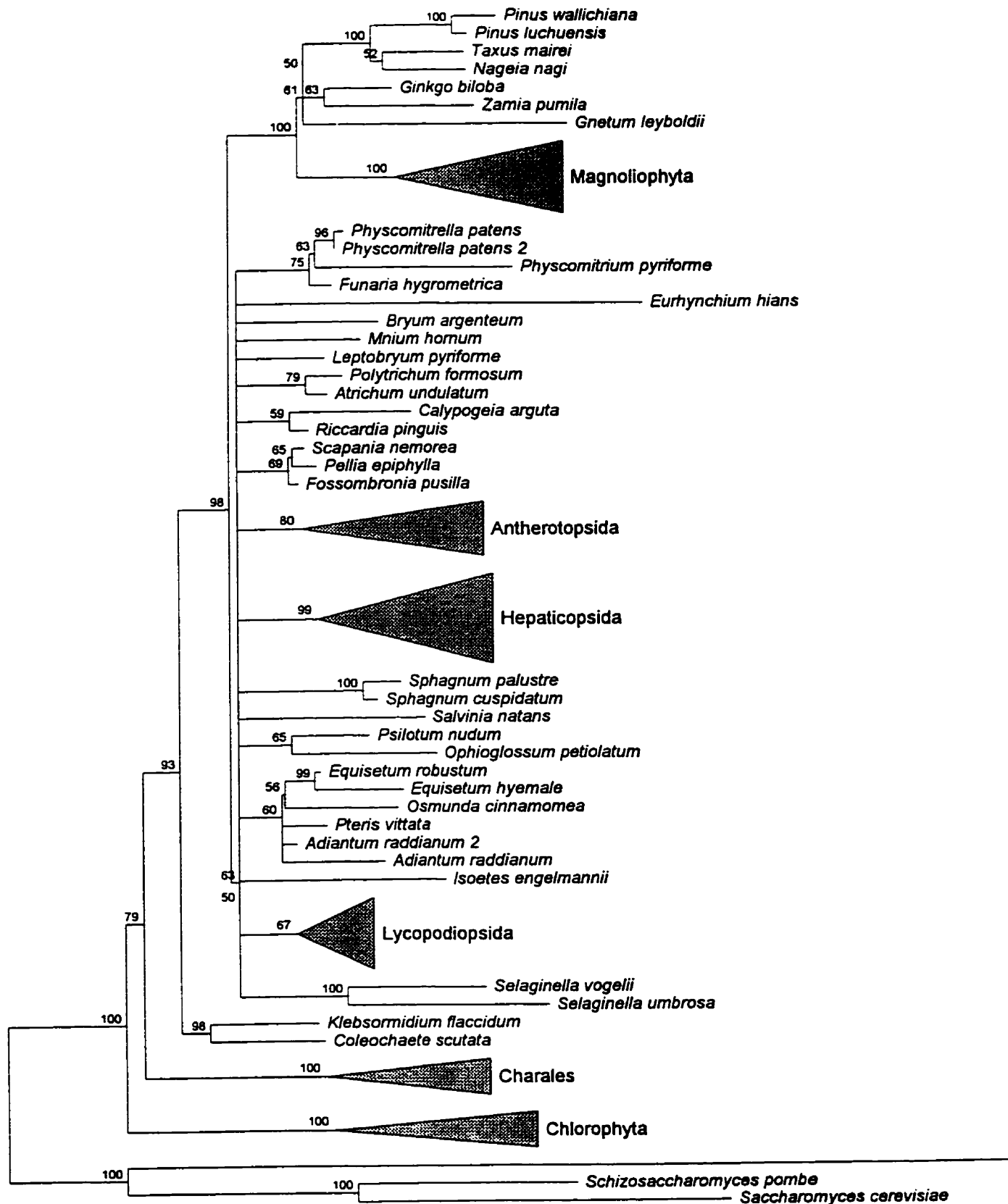


Figure H3B No Correction; Neighbor-joining
 Insertions & Deletions taken into account
 50% Majority-Rule Bootstrap Tree
 Only Categories 2, 3, 4, 5, 6

Figure 18

Overall strict consensus tree; Taxa set A

The overall strict consensus tree represents absolute corroboration of clusters in the bootstrap trees which were generated from all tree construction methods. Dendrograms for taxa set A and B representing a strict overall consensus tree were constructed by visual inspection of all 50% majority-rule bootstrap trees. Only clusters which are maintained strictly in the bootstrap trees from all phylogenetic analyses are depicted. Any variation of taxa clustering among the phylogenetic approaches resulted in those affected taxa being displayed as polytomies.

Color Key

—	Magnoliophyta
—	Pinophyta
—	Equisetophyta
—	Psilotophyta
—	Filicophyta
—	Lycopodiophyta
—	Anthocerotopsida
—	Bryopsida
—	Hepaticopsida
—	Charophyta

Overall Strict Consensus Tree

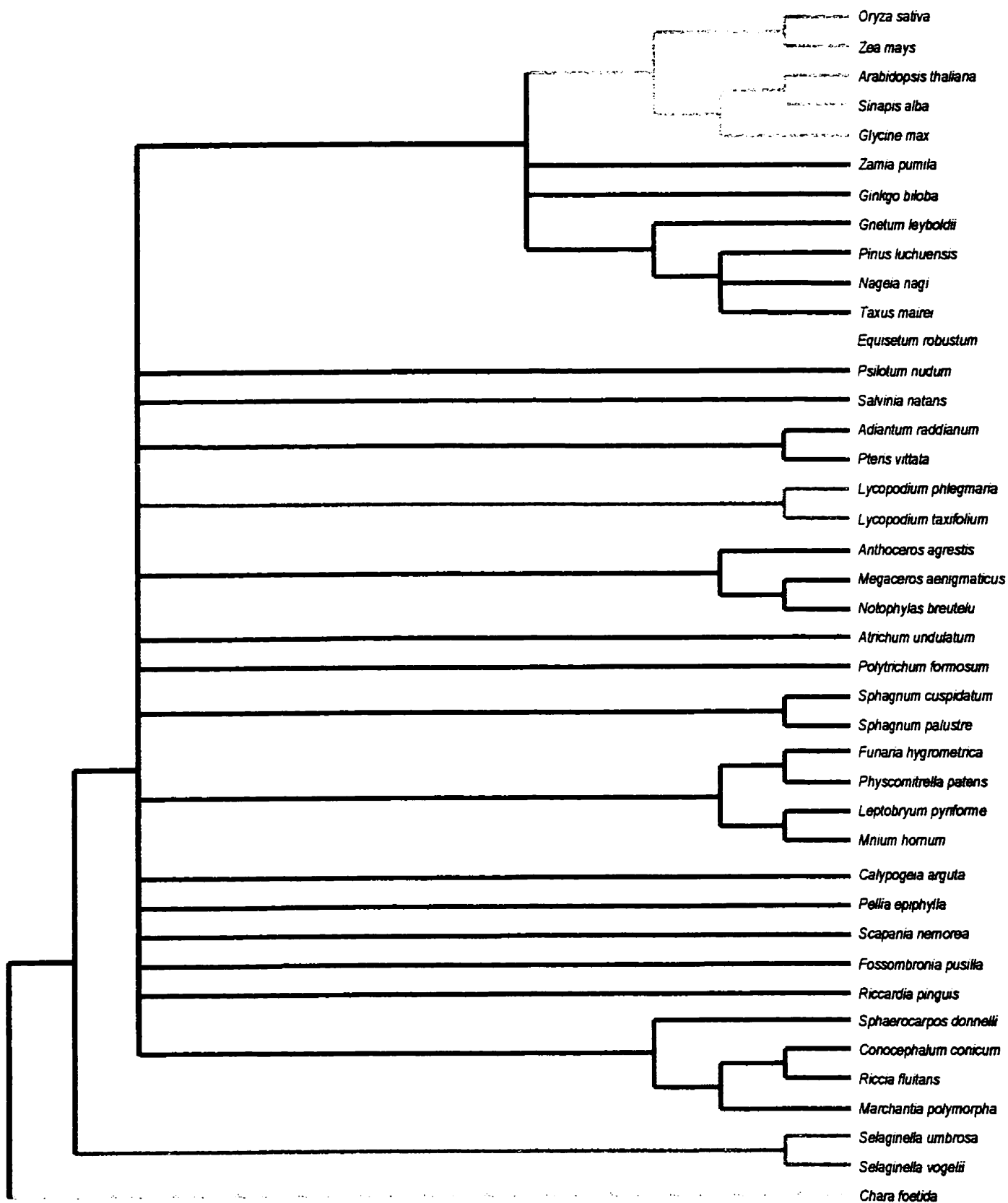


Figure 19

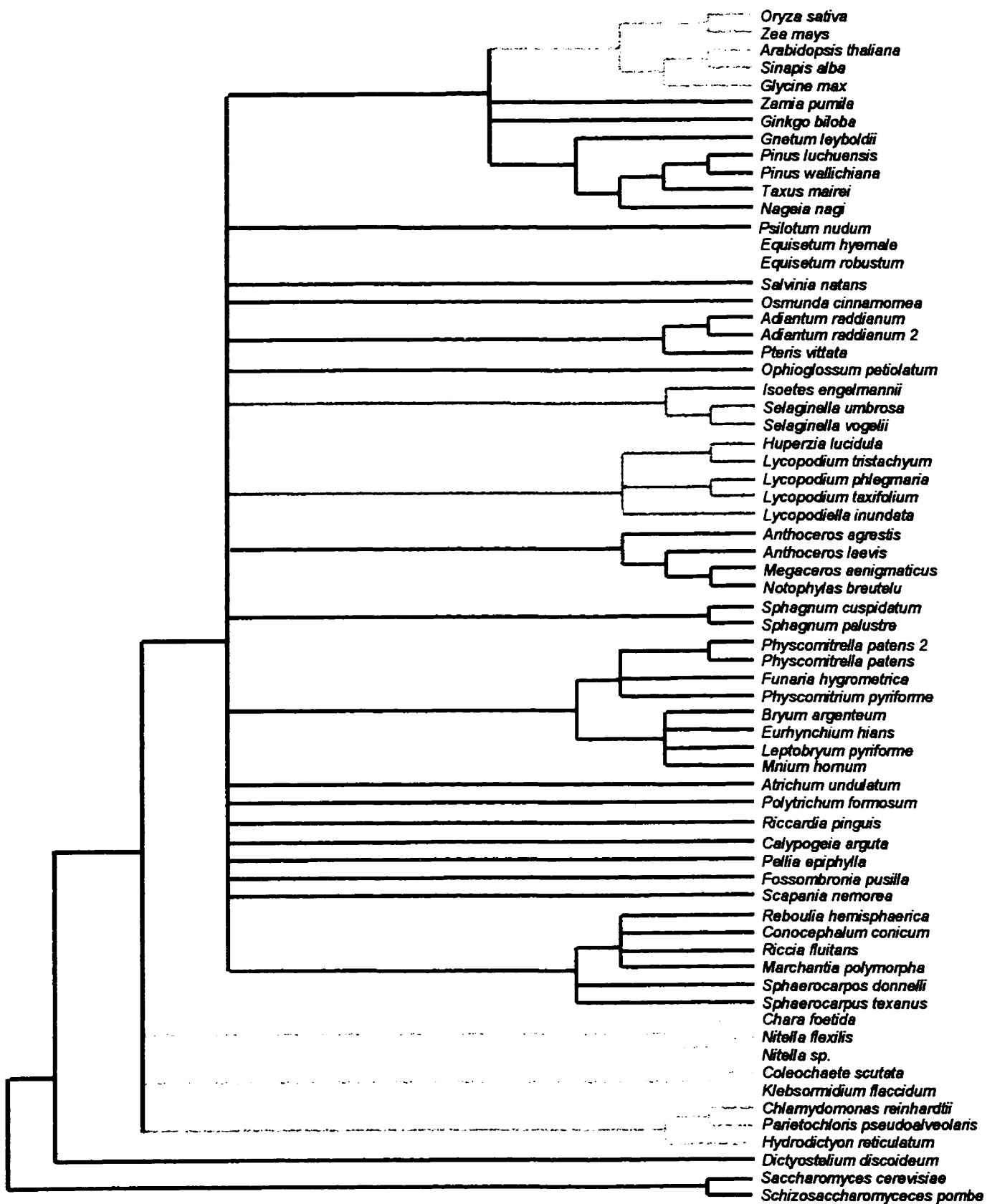
Overall strict consensus tree; Taxa set B

The overall strict consensus tree represents absolute corroboration of clusters in the bootstrap trees which were generated from all tree construction methods. Dendrograms for taxa set A and B representing a strict overall consensus tree were constructed by visual inspection of all 50% majority-rule bootstrap trees. Only clusters which are maintained strictly in the bootstrap trees from all phylogenetic analyses are depicted. Any variation of taxa clustering among the phylogenetic approaches resulted in those affected taxa being displayed as polytomies.

Color Key

———	Magnoliophyta
———	Pinophyta
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———	Lycopodiophyta
———	Anthocerotopsida
———	Bryopsida
———	Hepaticopsida
———	Charophyta
———	Chlorophyta
———	Fungi

Overall Strict Consensus Tree



APPENDIX G : DCSE Alignment

Alignment matrix using Dedicated Comparative Sequence Editor (DCSE) containing secondary structure information.

The standard ambiguity codes are used to apply partial identity to unresolved nucleotides.

<u>Code</u>	<u>Represented nucleotides</u>
Y	U or C
R	A or G
M	A or C
K	U or G
W	U or A
S	C or G
B	U, C, or G
D	U, A, or G
H	U, C, or A
V	C, A, or G
N	U, C, A, or G

Additional Codes

- N : unidentified nucleotide, length of unidentified are probably known.
- o : unidentified nucleotide, length of unidentified area unknown. In this case, the symbol "o" is intercalated for the number of nucleotides from the most closely related species.
- : denotes the presence of a gap at an alignment positions

Secondary Structure Symbols

- [and] : beginning and end of one strand of a helix.
- ^ : symbolizes][, a new helix starting immediately after a previous one.
- { and } : beginning and end of an internal loop
- (and) ; enclose a base forming part of a non-standard pair [any other pair other than G-C, A-U, or G-U].

DCSE Alignment

1 2283

	10	20	30	40	50	
- U A C U G G U U G A U C C U G C C A G U A G U (C) A U A U G C U U G U C U C A A A G A U U A A I G C						1 Saccharomyces cerevisiae
- U A C U G G U U G A U C C U G C C A G U A G U (C) A U A U G C U U G U C U C A A A G A U U A A I G C						2 Schizosaccharomyces pombe
- A A C U G G U U G A U C C U G C C A G U A G U (C) A U A U G C U U G U C U C A A A G A U U A A I G C						3 Physcomitrella patens
- A A C U G G U U G A U C C U G C C A G U A G U (C) A U A U G C U U G U C U C A A A G A U U A A I G C						4 Anthoceros agrestis
- A A C U G G U U G A U C C U G C C A G U A G U (C) A U A U G C U U G U C U C A A A G A U U A A I G C						5 Anthoceros laevis
- A A C U G G U U G A U C C U G C C A G U A G U (C) A U A U G C U U G U C U C A A A G A U U A A I G C						6 Pellia epiphylla
- A A C U G G U U G A U C C U G C C A G U A G U (C) A U A U G C U U G U C U C A A A G A U U A A I G C						7 Reboulia hemisphaerica 2
- A A C U G G U U G A U C C U G C C A G U A G U (C) A U A U G C U U G U C U C A A A G A U U A A I G C						8 Riccardia pinguis
- A A C U G G U U G A U C C U G C C A G U A G U (C) A U A U G C U U G U C U C A A A G A U U A A I G C						9 Sphaerocarpos donnellii
- A A C U G G U U G A U C C U G C C A G U A G U (C) A U A U G C U U G U C U C A A A G A U U A A I G C						10 Sphaerocarpos texanus
- A A C U G G U U G A U C C U G C C A G U A G U (C) A U A U G C U U G U C U C A A A G A U U A A I G C						11 Atrichum undulatum
- A A C U G G U U G A U C C U G C C A G U A G U (C) A U A U G C U U G U C U C A A A G A U U A A I G C						12 Bryum argenteum
- A A C U G G U U G A U C C U G C C A G U A G U (C) A U A U G C U U G U C U C A A A G A U U A A I G C						13 Eurhynchium hians
- A A C U G G U U G A U C C U G C C A G U A G U (C) A U A U G C U U G U C U C A A A G A U U A A I G C						14 Funaria hygrometrica
- A A C U G G U U G A U C C U G C C A G U A G U (C) A U A U G C U U G U C U C A A A G A U U A A I G C						15 Leptobryum pyriforme
- A A C U G G U U G A U C C U G C C A G U A G U (C) A U A U G C U U G U C U C A A A G A U U A A I G C						16 Mnium hornum
- A A C U G G U U G A U C C U G C C A G U A G U (C) A U A U G C U U G U C U C A A A G A U U A A I G C						17 Physcomitrium pyriforme
- A A C U G G U U G A U C C U G C C A G U A G U (C) A U A U G C U U G U C U C A A A G A U U A A I G C						18 Polytrichum formosum
- A A C U G G U U G A U C C U G C C A G U A G U (C) A U A U G C U U G U C U C A A A G A U U A A I G C						19 Sphagnum cuspidatum
- A A C U G G U U G A U C C U G C C A G U A G U (C) A U A U G C U U G U C U C A A A G A U U A A I G C						20 Calypogeia arguta
- A A C U G G U U G A U C C U G C C A G U A G U (C) A U A U G C U U G U C U C A A A G A U U A A I G C						21 Conocephalum conicum
- A A C U G G U U G A U C C U G C C A G U A G U (C) A U A U G C U U G U C U C A A A G A U U A A I G C						22 Fossombronina pusilla
- A A C U G G U U G A U C C U G C C A G U A G U (C) A U A U G C U U G U C U C A A A G A U U A A I G C						23 Marchantia polymorpha
- A A C U G G U U G A U C C U G C C A G U A G U (C) A U A U G C U U G U C U C A A A G A U U A A I G C						24 Riccia fluitans
- A A C U G G U U G A U C C U G C C A G U A G U (C) A U A U G C U U G U C U C A A A G A U U A A I G C						25 Scapania nemorea
- A A C U G G U U G A U C C U G C C A G U A G U (C) A U A U G C U U G U C U C A A A G A U U A A I G C						26 Equisetum hyemale
- A A C U G G U U G A U C C U G C C A G U A G U (C) A U A U G C U U G U C U C A A A G A U U A A I G C						27 Equisetum robustum
- A A C U G G U U G A U C C U G C C A G U A G U (C) A U A U G C U U G U C U C A A A G A U U A A I G C						28 Isoetes engelmannii
- A A C U G G U U G A U C C U G C C A G U A G U (C) A U A U G C U U G U C U C A A A G A U U A A I G C						29 Huperzia lucidula
- A A C U G G U U G A U C C U G C C A G U A G U (C) A U A U G C U U G U C U C A A A G A U U A A I G C						30 Lycopodiella inundata
- A A C U G G U U G A U C C U G C C A G U A G U (C) A U A U G C U U G U C U C A A A G A U U A A I G C						31 Lycopodium phlegmaria
- A A C U G G U U G A U C C U G C C A G U A G U (C) A U A U G C U U G U C U C A A A G A U U A A I G C						32 Lycopodium taxifolium
- A A C U G G U U G A U C C U G C C A G U A G U (C) A U A U G C U U G U C U C A A A G A U U A A I G C						33 Lycopodium tristachyum
- A A C U G G U U G A U C C U G C C A G U A G U (C) A U A U G C U U G U C U C A A A G A U U A A I G C						34 Selaginella galleottii
- A A C U G G U U G A U C C U G C C A G U A G U (C) A U A U G C U U G U C U C A A A G A U U A A I G C						35 Selaginella sp.
- U A C U G G U U G A U C C U G C C A G U A G U (C) A U A U G C U U G U C U C A A A G A U U A A I G C						36 Oryza sativa
- U A C U G G U U G A U C C U G C C A G U A G U (C) A U A U G C U U G U C U C A A A G A U U A A I G C						37 Zea mays
- U A C U G G U U G A U C C U G C C A G U A G U (C) A U A U G C U U G U C U C A A A G A U U A A I G C						38 Arabidopsis thaliana
- U A C U G G U U G A U C C U G C C A G U A G U (C) A U A U G C U U G U C U C A A A G A U U A A I G C						39 Glycine max
- U A C U G G U U G A U C C U G C C A G U A G U (C) A U A U G C U U G U C U C A A A G A U U A A I G C						40 Sinapis alba
- U A C U G G U U G A U C C U G C C A G U A G U (C) A U A U G C U U G U C U C A A A G A U U A A I G C						41 Zamia pumila
- U A C U G G U U G A U C C U G C C A G U A G U (C) A U A U G C U U G U C U C A A A G A U U A A I G C						42 Gnetum leyboldii
- U A C U G G U U G A U C C U G C C A G U A G U (C) A U A U G C U U G U C U C A A A G A U U A A I G C						43 Ginkgo biloba
- U A C U G G U U G A U C C U G C C A G U A G U (C) A U A U G C U U G U C U C A A A G A U U A A I G C						44 Pinus luchuensis
- U A C U G G U U G A U C C U G C C A G U A G U (C) A U A U G C U U G U C U C A A A G A U U A A I G C						45 Pinus wallichiana
- U A C U G G U U G A U C C U G C C A G U A G U (C) A U A U G C U U G U C U C A A A G A U U A A I G C						46 Podocarpus nakaii
- U A C U G G U U G A U C C U G C C A G U A G U (C) A U A U G C U U G U C U C A A A G A U U A A I G C						47 Taxus mairei
- U A C U G G U U G A U C C U G C C A G U A G U (C) A U A U G C U U G U C U C A A A G A U U A A I G C						48 Adiantum raddianum
- U A C U G G U U G A U C C U G C C A G U A G U (C) A U A U G C U U G U C U C A A A G A U U A A I G C						49 Adiantum raddianum 2
- U A C U G G U U G A U C C U G C C A G U A G U (C) A U A U G C U U G U C U C A A A G A U U A A I G C						50 Ophioglossum petiolatum
- U A C U G G U U G A U C C U G C C A G U A G U (C) A U A U G C U U G U C U C A A A G A U U A A I G C						51 Osmunda cinnamomea
- U A C U G G U U G A U C C U G C C A G U A G U (C) A U A U G C U U G U C U C A A A G A U U A A I G C						52 Salvinia natans
- U A C U G G U U G A U C C U G C C A G U A G U (C) A U A U G C U U G U C U C A A A G A U U A A I G C						53 Psilotum nudum 2
- U A C U G G U U G A U C C U G C C A G U A G U (C) A U A U G C U U G U C U C A A A G A U U A A I G C						54 Chara foetida
- U A C U G G U U G A U C C U G C C A G U A G U (C) A U A U G C U U G U C U C A A A G A U U A A I G C						55 Coleochaete scutata
- U A C U G G U U G A U C C U G C C A G U A G U (C) A U A U G C U U G U C U C A A A G A U U A A I G C						56 Klebsormidium flaccidum
- U A C U G G U U G A U C C U G C C A G U A G U (C) A U A U G C U U G U C U C A A A G A U U A A I G C						57 Nitella flexilis
- U A C U G G U U G A U C C U G C C A G U A G U (C) A U A U G C U U G U C U C A A A G A U U A A I G C						58 Nitella sp.
- U A C U G G U U G A U C C U G C C A G U A G U (C) A U A U G C U U G U C U C A A A G A U U A A I G C						59 Chlamydomonas reinhardtii
- U A C U G G U U G A U C C U G C C A G U A G U (C) A U A U G C U U G U C U C A A A G A U U A A I G C						60 Hydrodictyon reticulatum
- U A C U G G U U G A U C C U G C C A G U A G U (C) A U A U G C U U G U C U C A A A G A U U A A I G C						61 Parietochloris pseudoalveolaris
- U A C U G G U U G A U C C U G C C A G U A G U (C) A U A U G C U U G U C U C A A A G A U U A A I G C						62 Dictyostelium discoideum
- U A C U G G U U G A U C C U G C C A G U A G U (C) A U A U G C U U G U C U C A A A G A U U A A I G C						63 Helix numbering 1
- U A C U G G U U G A U C C U G C C A G U A G U (C) A U A U G C U U G U C U C A A A G A U U A A I G C						64 Pteris vittata
- U A C U G G U U G A U C C U G C C A G U A G U (C) A U A U G C U U G U C U C A A A G A U U A A I G C						65 Megaceros aenigmaticus
- U A C U G G U U G A U C C U G C C A G U A G U (C) A U A U G C U U G U C U C A A A G A U U A A I G C						66 Notophylas breutelii
- U A C U G G U U G A U C C U G C C A G U A G U (C) A U A U G C U U G U C U C A A A G A U U A A I G C						67 Physcomitrella patens
- U A C U G G U U G A U C C U G C C A G U A G U (C) A U A U G C U U G U C U C A A A G A U U A A I G C						68 Sphagnum palustre

DCSE Alignment

1 :10
 2 :20
 3 :30
 4 :40
 5 :50
 6 :60
 7 :70
 8 :80
 9 :90
 10 :100
 11 :110
 12 :120
 13 :130
 14 :140
 15 :150
 16 :160
 17 :170
 18 :180
 19 :190
 20 :200
 21 :210
 22 :220
 23 :230
 24 :240
 25 :250
 26 :260
 27 :270
 28 :280
 29 :290
 30 :300
 31 :310
 32 :320
 33 :330
 34 :340
 35 :350
 36 :360
 37 :370
 38 :380
 39 :390
 40 :400
 41 :410
 42 :420
 43 :430
 44 :440
 45 :450
 46 :460
 47 :470
 48 :480
 49 :490
 50 :500
 51 :510
 52 :520
 53 :530
 54 :540
 55 :550
 56 :560
 57 :570
 58 :580
 59 :590
 60 :600
 61 :610
 62 :620
 63 :630
 64 :640
 65 :650
 66 :660
 67 :670
 68 :680

- 1 Saccharomyces cerevisiae
- 2 Schizosaccharomyces pombe
- 3 Physcometrella patens
- 4 Anthoceros agrestis
- 5 Anthoceros laevis
- 6 Pellia epiphylla
- 7 Reboulia hemisphaerica 2
- 8 Riccardia pinguis
- 9 Sphaerocarpos donnellii
- 10 Sphaerocarpos texanus
- 11 Atrichum undulatum
- 12 Bryum argenteum
- 13 Eurlimichium hians
- 14 Funaria hygrometrica
- 15 Lepidobryum pyriforme
- 16 Mnium hornum
- 17 Physcomatritium pyriforme
- 18 Polytretichum formosum
- 19 Sphagnum cuspidatum
- 20 Calypogeia arguta
- 21 Conocephalum conicum
- 22 Fossombronia pusilla
- 23 Marchantia polymorpha
- 24 Riccia fluitans
- 25 Scapania nemorea
- 26 Equisetum hyemale
- 27 Equisetum rostratum
- 28 Isoetes engelmannii
- 29 Isoetes lucidula
- 30 Lycopodium inundata
- 31 Lycopodium phlegmaria
- 32 Lycopodium taxifolium
- 33 Lycopodium tristachyum
- 34 Selaginella galleottii
- 35 Selaginella sp.
- 36 Oryza sativa
- 37 Zea mays
- 38 Arabidopsis thaliana
- 39 Glycine max
- 40 Sinapis alba
- 41 Zama pumila
- 42 Gnetum leyboldii
- 43 Ginkgo biloba
- 44 Pinus luchuensis
- 45 Pinus wallichiana
- 46 Podocarpus nakiellii
- 47 Taxus mairei
- 48 Adiantum raddianum
- 49 Adiantum raddianum 2
- 50 Ophioglossum petiolatum
- 51 Osmunda cinnamomea
- 52 Salvinia natans
- 53 Ptilocnemum 2
- 54 Chara foetida
- 55 Coleochaete scutata
- 56 Klebsormidium flaccidum
- 57 Nitella flexilis
- 58 Nitella sp.
- 59 Chlamydomonas reinhardtii
- 60 Hydrodictyon reticulatum
- 61 Paratetochloris pseudovalvolaris
- 62 Dictyostelium discoideum
- 63 Helix numbering 1
- 64 Petasites vitreata
- 65 Heloglossum dentigulatus
- 66 Notophytas breutelei
- 67 Physcometrella patens
- 68 Sphagnum palustre

	310	320	330	340	350	
-	-	-	-	-	-	1 Saccharomyces cerevisiae
(G	A	C	I	-	-	2 Schizosaccharomyces pombe
C	G	G	C	I	-	3 Physcomitrella patens
C	C	G	G	I	-	4 Anthoceros agrestis
C	C	G	G	I	-	5 Anthoceros laevis
C	C	G	G	I	-	6 Pellia epiphylla
C	C	G	G	I	-	7 Reboulia hemisphaerica 2
C	C	G	G	I	-	8 Riccardia pinguis
C	C	G	G	I	-	9 Sphaerocarpos donnelli
C	C	G	G	I	-	10 Sphaerocarpos texanus
C	C	G	G	I	-	11 Atrichum undulatum
C	C	G	G	I	-	12 Bryum argenteum
C	C	G	G	I	-	13 Eurhynchium hians
C	C	G	G	I	-	14 Funaria hygrometrica
C	C	G	G	I	-	15 Leptobryum pyriforme
C	C	G	G	I	-	16 Mnium hornum
C	C	G	G	I	-	17 Physcomitrium pyriforme
C	C	G	G	I	-	18 Polytrichum formosum
C	C	G	G	I	-	19 Sphagnum cuspidatum
C	C	G	G	I	-	20 Calypogeia arguta
C	C	G	G	I	-	21 Conocephalum conicum
C	C	G	G	I	-	22 Fossombronina pusilla
C	C	G	G	I	-	23 Marchantia polymorpha
C	C	G	G	I	-	24 Riccia fluitans
C	C	G	G	I	-	25 Scapania nemorea
C	C	G	G	I	-	26 Equisetum hyemale
C	C	G	G	I	-	27 Equisetum robustum
(C	U	G	I	-	-	28 Isoetes engelmannii
(C	C	G	G	I	-	29 Huperzia lucidula
(C	C	G	G	I	-	30 Lycopodiella inundata
C	C	G	G	I	-	31 Lycopodium phlegmaria
C	C	G	G	I	-	32 Lycopodium taxifolium
(C	C	G	G	I	-	33 Lycopodium tristachyum
C	C	G	G	I	-	34 Selaginella galleottii
C	C	G	G	I	-	35 Selaginella sp.
C	C	G	G	I	-	36 Oryza sativa
C	C	G	G	I	-	37 Zea mays
U	U	G	C	I	-	38 Arabidopsis thaliana
C	C	G	G	I	-	39 Glycine max
(G	U	G	C	I	-	40 Sinapis alba
C	C	G	G	I	-	41 Zamia pumila
U	U	G	C	I	-	42 Gnetum leyboldii
C	C	G	G	I	-	43 Ginkgo biloba
C	C	G	G	I	-	44 Pinus luchuensis
C	C	G	G	I	-	45 Pinus wallichiana
C	C	G	G	I	-	46 Podocarpus nakaii
C	C	G	G	I	-	47 Taxus mairei
C	C	G	G	I	-	48 Adiantum raddianum
C	C	G	G	I	-	49 Adiantum raddianum 2
C	C	G	G	I	-	50 Ophioglossum petiolatum
U	U	G	C	I	-	51 Osmunda cinnamomea
C	C	G	G	I	-	52 Salvinia natans
(C	C	G	G	I	-	53 Psilotum nudum 2
(C	C	G	G	I	-	54 Chara foetida
(C	C	G	G	I	-	55 Coleochaete scutata
(C	C	G	G	I	-	56 Klebsormidium flaccidum
(C	C	G	G	I	-	57 Nitella flexilis
(C	C	G	G	I	-	58 Nitella sp.
C	C	G	G	I	-	59 Chlamydomonas reinhardtii
U	U	G	C	I	-	60 Hydrodictyon reticulatum
C	C	G	G	I	-	61 Parietochloris pseudoalveolaris
-	(G	U	I	-	-	62 Dictyostelium discoideum
-	-	-	-	-	-	63 Helix numbering 1
C	C	G	G	I	-	64 Pteris vittata
C	C	G	G	I	-	65 Megaceros senigmaticus
C	C	G	G	I	-	66 Notophylas breutelii
C	C	G	G	I	-	67 Physcomitrella patens
C	C	G	G	I	-	68 Sphagnum palustre

	360	370	380	390	400	
C G C I A	-1U G G C	- C U U G G U	-G C U G G C G A J U G	- G U U C A I U U C	-A A A U U U U G J C C	1 Saccharomyces cerevisiae
C G C I A	-1U G G G C	- C U U G G C C	-G C U G G C G A J U G	- G U U C A I U U C	-A A A U U U U G J C C	2 Schizosaccharomyces pombe
C G C I A	-1C G G G C	- C U U G G C C	-G C U G G C G A J U G	- U U U C A I U U C	-A A A U U U U G J C C	3 Physcomitrella patens
C G C I A	-1C G G G C	- C U U G G C C	-G C U G G C G A J U G	- U U U C A I U U C	-A A A U U U U G J C C	4 Anthoceros agrestis
(C) G C I A	-1C G G G C	- C U U G G C C	-G C U G G C G A J U G	- U U U C A I U U C	-A A A U U U U G J C C	5 Anthoceros laevis
C G C I A	-1C G G G C	- C U U G G C C	-G C U G G C G A J U G	- U U U C A I U U C	-A A A U U U U G J C C	6 Pellia epiphylla
C G C I A	-1C G G G C	- C U U G G C C	-G C U G G C G A J U G	- U U U C A I U U C	-A A A U U U U G J C C	7 Reboulia hemisphaerica 2
C G C I A	-1C G G G C	- C U U G G C C	-G C U G G C G A J U G	- U U U C A I U U C	-A A A U U U U G J C C	8 Riccardia pinguis
C G C I A	-1C G G G C	- C U U G G C C	-G C U G G C G A J U G	- U U U C A I U U C	-A A A U U U U G J C C	9 Sphaerocarpos donnelli
C G C I A	-1C G G G C	- C U U G G C C	-G C U G G C G A J U G	- U U U C A I U U C	-A A A U U U U G J C C	10 Sphaerocarpos texanus
C G C I A	-1C G G G C	- C U U G G C C	-G C U G G C G A J U G	- U U U C A I U U C	-A A A U U U U G J C C	11 Atrichum undulatum
C G C I A	-1C G G G C	- C U U G G C C	-G C U G G C G A J U G	- U U U C A I U U C	-A A A U U U U G J C C	12 Bryum argenteum
C(C) C I A	-1C G G G C	- C U U G G C C	-G C U G G C G A J U G	- U U U C A I U U C	-A A A U U U U G J C C	13 Eurhynchium hians
C G C I A	-1C G G G C	- C U U G G C C	-G C U G G C G A J U G	- U U U C A I U U C	-A A A U U U U G J C C	14 Funaria hygrometrica
C G C I A	-1C G G G C	- C U U G G C C	-G C U G G C G A J U G	- U U U C A I U U C	-A A A U U U U G J C C	15 Leptobryum pyriforme
C G C I A	-1C G G G C	- C U U G G C C	-G C U G G C G A J U G	- U U U C A I U U C	-A A A U U U U G J C C	16 Mnium hornum
C G C I A	-1C G G G C	- C U U G G C C	-G C U G G C G A J U G	- U U U C A I U U C	-A A A U U U U G J C C	17 Physcomitrium pyriforme
C G C I A	-1C G G G C	- C U U G G C C	-G C U G G C G A J U G	- U U U C A I U U C	-A A A U U U U G J C C	18 Polytrichum formosum
C G C I A	-1C G G G C	- C U U G G C C	-G C U G G C G A J U G	- U U U C A I U U C	-A A A U U U U G J C C	19 Sphagnum cuspidatum
C G C I A	-1C G G G C	- C U U G G C C	-G C U G G C G A J U G	- U U U C A I U U C	-A A A U U U U G J C C	20 Calypogeia arguta
C G C I A	-1C G G G C	- C U U G G C C	-G C U G G C G A J U G	- U U U C A I U U C	-A A A U U U U G J C C	21 Conocephalum conicum
C G C I A	-1C G G G C	- C U U G G C C	-G C U G G C G A J U G	- U U U C A I U U C	-A A A U U U U G J C C	22 Fossombronina pusilla
C G C I A	-1C G G G C	- C U U G G C C	-G C U G G C G A J U G	- U U U C A I U U C	-A A A U U U U G J C C	23 Marchantia polymorpha
C G C I A	-1C G G G C	- C U U G G C C	-G C U G G C G A J U G	- U U U C A I U U C	-A A A U U U U G J C C	24 Riccia fluitans
C G C I A	-1C G G G C	- C U U G G C C	-G C U G G C G A J U G	- U U U C A I U U C	-A A A U U U U G J C C	25 Scapania nemorea
C G C I A	-1C G G G C	- C U U G G C C	-G C U G G C G A J U G	- U U U C A I U U C	-A A A U U U U G J C C	26 Equisetum hyemale
C G C I A	-1C G G G C	- C U U G G C C	-G C U G G C G A J U G	- U U U C A I U U C	-A A A U U U U G J C C	27 Equisetum robustum
C G C I A	-1C G G G C	- C U U G G C C	-G C U G G C G A J U G	- U U U C A I U U C	-A A A U U U U G J C C	28 Isoetes engelmannii
C G C I A	-1C G G G C	- C U U G G C C	-G C U G G C G A J U G	- U U U C A I U U C	-A A A U U U U G J C C	29 Huperzia lucidula
C G C I A	-1C G G G C	- C U U G G C C	-G C U G G C G A J U G	- U U U C A I U U C	-A A A U U U U G J C C	30 Lycopodiella inundata
C G C I A	-1C G G G C	- C U U G G C C	-G C U G G C G A J U G	- U U U C A I U U C	-A A A U U U U G J C C	31 Lycopodium phlegmaria
C G C I A	-1C G G G C	- C U U G G C C	-G C U G G C G A J U G	- U U U C A I U U C	-A A A U U U U G J C C	32 Lycopodium taxifolium
C G C I A	-1C G G G C	- C U U G G C C	-G C U G G C G A J U G	- U U U C A I U U C	-A A A U U U U G J C C	33 Lycopodium tristachyum
C G C I A	-1C G G G C	- C U U G G C C	-G C U G G C G A J U G	- U U U C A I U U C	-A A A U U U U G J C C	34 Selaginella galleottii
C G C I A	-1C G G G C	- C U U G G C C	-G C U G G C G A J U G	- U U U C A I U U C	-A A A U U U U G J C C	35 Selaginella sp.
C G C I A	-1C G G G C	- C U U G G C C	-G C U G G C G A J U G	- U U U C A I U U C	-A A A U U U U G J C C	36 Oryza sativa
C G C I A	-1C G G G C	- C U U G G C C	-G C U G G C G A J U G	- U U U C A I U U C	-A A A U U U U G J C C	37 Zea mays
C G C I A	-1C G G G C	- C U U G G C C	-G C U G G C G A J U G	- U U U C A I U U C	-A A A U U U U G J C C	38 Arabidopsis thaliana
C G C I A	-1C G G G C	- C U U G G C C	-G C U G G C G A J U G	- U U U C A I U U C	-A A A U U U U G J C C	39 Glycine max
C G C I A	-1C G G G C	- C U U G G C C	-G C U G G C G A J U G	- U U U C A I U U C	-A A A U U U U G J C C	40 Sinapis alba
U G C I A	-1U G G G C	- C U U G G C C	-G C U G G C G A J U G	- U U U C A I U U C	-A A A U U U U G J C C	41 Zamia pumila
G C G A	-1C G G G C	- C U U G G C C	-G C U G G C G A J U G	- U U U C A I U U C	-A A A U U U U G J C C	42 Gnetum leyboldii
C G C I A	-1C G G G C	- C U U G G C C	-G C U G G C G A J U G	- U U U C A I U U C	-A A A U U U U G J C C	43 Ginkgo biloba
C G C I A	-1C A G G C	- C U U G G C C	-G C U G G C G A J U G	- U U U C A I U U C	-A A A U U U U G J C C	44 Pinus luchuensis
C G C I A	-1C G G G C	- C U U G G C C	-G C U G G C G A J U G	- U U U C A I U U C	-A A A U U U U G J C C	45 Pinus wallichiana
C G C I A	-1C G G G C	- C U U G G C C	-G C U G G C G A J U G	- U U U C A I U U C	-A A A U U U U G J C C	46 Podocarpus nakaii
C G C I A	-1C G G G C	- C U U G G C C	-G C U G G C G A J U G	- U U U C A I U U C	-A A A U U U U G J C C	47 Taxus mairei
C G C I A	-1C G G G C	- C U U G G C C	-G C U G G C G A J U G	- U U U C A I U U C	-A A A U U U U G J C C	48 Adiantum raddianum
C G C I A	-1C G G G C	- C U U G G C C	-G C U G G C G A J U G	- U U U C A I U U C	-A A A U U U U G J C C	49 Adiantum raddianum 2
(C) G C I A	-1C G G G C	- C U U G G C C	-G C U G G C G A J U G	- U U U C A I U U C	-A A A U U U U G J C C	50 Ophioglossum petiolatum
C G C I A	-1C G G G C	- C U U G G C C	-G C U G G C G A J U G	- U U U C A I U U C	-A A A U U U U G J C C	51 Osmunda cinnamomea
C G C I A	-1C G G G C	- C U U G G C C	-G C U G G C G A J U G	- U U U C A I U U C	-A A A U U U U G J C C	52 Salvinia natans
C G C I A	-1C G G G C	- C U U G G C C	-G C U G G C G A J U G	- U U U C A I U U C	-A A A U U U U G J C C	53 Psilotum nudum 2
C G C I A	-1C G G G C	- C U U G G C C	-G C U G G C G A J U G	- U U U C A I U U C	-A A A U U U U G J C C	54 Chara foetida
C G C I A	-1C G G G C	- C U U G G C C	-G C U G G C G A J U G	- U U U C A I U U C	-A A A U U U U G J C C	55 Coleochaete scutata
C G C I A	-1C G G G C	- C U U G G C C	-G C U G G C G A J U G	- U U U C A I U U C	-A A A U U U U G J C C	56 Klebsoraidium flaccidum
C G C I A	-1U G G G C	- C U U G G C C	-G C U G G C G A J U G	- U U U C A I U U C	-A A A U U U U G J C C	57 Nitella flexilis
C G U I A	-1U G G G C	- C U U G G C C	-G C U G G C G A J U G	- U U U C A I U U C	-A A A U U U U G J C C	58 Nitella sp.
C G C I A	-1C G G G C	- C U U G G C C	-G C U G G C G A J U G	- U U U C A I U U C	-A A A U U U U G J C C	59 Chlamydomonas reinhardtii
C G C I A	-1C G G G C	- C U U G G C C	-G C U G G C G A J U G	- U U U C A I U U C	-A A A U U U U G J C C	60 Hydrodictyon reticulatum
C G C I A	-1U G G G C	- C U U G G C C	-G C U G G C G A J U G	- U U U C A I U U C	-A A A U U U U G J C C	61 Parietochloris pseudoalveolaris
C G A	-1G A	- U U U A	-G C U G G C G A J U G	- U U U C A I U U C	-A A A U U U U G J C C	62 Dictyostelium discoideum
11						
(C) G C I A	-1C G G G C	- C U U G G C C	-G C U G G C G A J U G	- U U U C A I U U C	-A A A U U U U G J C C	63 Helix numbering 1
(G) C U C A A C	-1C G G G C	- C U U G G C C	-G C U G G C G A J U G	- U U U C A I U U C	-A A A U U U U G J C C	64 Pteris vittata
(C) G C I A	-1C G G G C	- C U U G G C C	-G C U G G C G A J U G	- U U U C A I U U C	-A A A U U U U G J C C	65 Megaceros aenigmaticus
C G C I A	-1C G G G C	- C U U G G C C	-G C U G G C G A J U G	- U U U C A I U U C	-A A A U U U U G J C C	66 Notophyas breutelii
C G C I A	-1C G G G C	- C U U G G C C	-G C U G G C G A J U G	- U U U C A I U U C	-A A A U U U U G J C C	67 Physcomitrella patens
C G C I A	-1C G G G C	- C U U G G C C	-G C U G G C G A J U G	- U U U C A I U U C	-A A A U U U U G J C C	68 Sphagnum palustre

	410	420	430	440	450	
U(A)U(C) - A)A C U(U)U C G)A U G G U A G G(A)U)A G U(G)(G) C U A C C A U G G U(U)U C A)A(C)G G G)U A						1 Saccharomyces cerevisiae
U(A)U(C) - A)A C U(U)U C G)A U G G U A G G(A)U)A G A(G)(G) C U A C C A U G G U(U)U U A)A(C)G G G)U A						2 Schizosaccharomyces pombe
U(A)U(C) - A)A C U(U)U C G)A U G G U A G G(A)U)A G A(G)(G) C U A C C A U G G U(U)U U A)A(C)G G G)U G						3 Physcomitrella patens
U(A)U(C) - A)A C U(U)U C G)A U G G U A G G(A)U)A G A(G)(G) C U A C C A U G G U(U)U U A)A(C)G G G)U G						4 Anthoceros agrestis
U(A)U(C) - A)A C U(U)U C G)A U G G U A G G(A)U)A G A(G)(G) C U A C C A U G G U(U)U U A)A(C)G G G)U G						5 Anthoceros laevis
U(A)U(C) - A)A C U(U)U C G)A U G G U A G G(A)U)A G A(G)(G) C U A C C A U G G U(U)U U A)A(C)G G G)U G						6 Pellia epiphylla
U(A)U(C) - A)A C U(U)U C G)A U G G U A G G(A)U)A G A(G)(G) C U A C C A U G G U(U)U U A)A(C)G G G)U G						7 Reboulia hemisphaerica 2
U(A)U(C) - A)A C U(U)U C G)A U G G U A G G(A)U)A G A(G)(G) C U A C C A U G G U(U)U U A)A(C)G G G)U G						8 Riccardia pinguis
U(A)U(C) - A)A C U(U)U C G)A U G G U A G G(A)U)A G A(G)(G) C U A C C A U G G U(U)U U A)A(C)G G G)U G						9 Sphaerocarpos donnellii
U(A)U(C) - A)A C U(U)U C G)A U G G U A G G(A)U)A G A(G)(G) C U A C C A U G G U(U)U U A)A(C)G G G)U G						10 Sphaerocarpos texanus
U(A)U(C) - A)A C U(U)U C G)A U G G U A G G(A)U)A G A(G)(G) C U A C C A U G G U(U)U U A)A(C)G G G)U G						11 Atrichum undulatum
U(A)U(C) - A)A C U(U)U C G)A U G G U A G G(A)U)A G A(G)(G) C U A C C A U G G U(U)U U A)A(C)G G G)U G						12 Bryum argenteum
U(A)U(C) - A)A C U(U)U C G)A U G G U A G G(A)U)A G A(G)(G) C U A C C A U G G U(U)U U A)A(C)G G G)U G						13 Eurhynchium hians
U(A)U(C) - A)A C U(U)U C G)A U G G U A G G(A)U)A G A(G)(G) C U A C C A U G G U(U)U U A)A(C)G G G)U G						14 Funaria hygrometrica
U(A)U(C) - A)A C U(U)U C G)A U G G U A G G(A)U)A G A(G)(G) C U A C C A U G G U(U)U U A)A(C)G G G)U G						15 Leptobryum pyriforme
U(A)U(C) - A)A C U(U)U C G)A U G G U A G G(A)U)A G A(G)(G) C U A C C A U G G U(U)U U A)A(C)G G G)U G						16 Mnium hornum
U(A)U(C) - A)A C U(U)U C G)A U G G U A G G(A)U)A G A(G)(G) C U A C C A U G G U(U)U U A)A(C)G G G)U G						17 Physcomitrium pyriforme
U(A)U(C) - A)A C U(U)U C G)A U G G U A G G(A)U)A G A(G)(G) C U A C C A U G G U(U)U U A)A(C)G G G)U G						18 Polytrichum formosum
U(A)U(C) - A)A C U(U)U C G)A U G G U A G G(A)U)A G A(G)(G) C U A C C A U G G U(U)U U A)A(C)G G G)U G						19 Sphagnum cuspidatum
U(A)U(C) - A)A C U(U)U C G)A U G G U A G G(A)U)A G A(G)(G) C U A C C A U G G U(U)U U A)A(C)G G G)U G						20 Calypogeia arguta
U(A)U(C) - A)A C U(U)U C G)A U G G U A G G(A)U)A G A(G)(G) C U A C C A U G G U(U)U U A)A(C)G G G)U G						21 Conocephalum concium
U(A)U(C) - A)A C U(U)U C G)A U G G U A G G(A)U)A G A(G)(G) C U A C C A U G G U(U)U U A)A(C)G G G)U G						22 Fossombronina pusilla
U(A)U(C) - A)A C U(U)U C G)A U G G U A G G(A)U)A G A(G)(G) C U A C C A U G G U(U)U U A)A(C)G G G)U G						23 Marchantia polymorpha
U(A)U(C) - A)A C U(U)U C G)A U G G U A G G(A)U)A G A(G)(G) C U A C C A U G G U(U)U U A)A(C)G G G)U G						24 Riccia fluitans
U(A)U(C) - A)A C U(U)U C G)A U G G U A G G(A)U)A G A(G)(G) C U A C C A U G G U(U)U U A)A(C)G G G)U G						25 Scapania nemorea
U(A)U(C) - A)A C U(U)U C G)A U G G U A G G(A)U)A G A(G)(G) C U A C C A U G G U(U)U U A)A(C)G G G)U G						26 Equisetum hyemale
U(A)U(C) - A)A C U(U)U C G)A U G G U A G G(A)U)A G A(G)(G) C U A C C A U G G U(U)U U A)A(C)G G G)U G						27 Equisetum robustum
U(A)U(C) - A)A C U(U)U C G)A U G G U A G G(A)U)A G A(G)(G) C U A C C A U G G U(U)U U A)A(C)G G G)U G						28 Isoetes engelmannii
U(A)U(C) - A)A C U(U)U C G)A U G G U A G G(A)U)A G A(G)(G) C U A C C A U G G U(U)U U A)A(C)G G G)U G						29 Huperzia lucidula
U(A)U(C) - A)A C U(U)U C G)A U G G U A G G(A)U)A G A(G)(G) C U A C C A U G G U(U)U U A)A(C)G G G)U G						30 Lycopodiella inundata
U(A)U(C) - A)A C U(U)U C G)A U G G U A G G(A)U)A G A(G)(G) C U A C C A U G G U(U)U U A)A(C)G G G)U G						31 Lycopodium phlegmaria
U(A)U(C) - A)A C U(U)U C G)A U G G U A G G(A)U)A G A(G)(G) C U A C C A U G G U(U)U U A)A(C)G G G)U G						32 Lycopodium taxifolium
U(A)U(C) - A)A C U(U)U C G)A U G G U A G G(A)U)A G A(G)(G) C U A C C A U G G U(U)U U A)A(C)G G G)U G						33 Lycopodium tristachyum
U(A)U(C) - A)A C U(U)U C G)A U G G U A G G(A)U)A G A(G)(G) C U A C C A U G G U(U)U U A)A(C)G G G)U A						34 Selaginella galleottii
U(A)U(C) - A)A C U(U)U C G)A U G G U A G G(A)U)A G A(G)(G) C U A C C A U G G U(U)U U A)A(C)G G G)U G						35 Selaginella sp.
U(A)U(C) - A)A C U(U)U C G)A U G G U A G G(A)U)A G A(G)(G) C U A C C A U G G U(U)U U A)A(C)G G G)U G						36 Orzyza sativa
U(A)U(C) - A)A C U(U)U C G)A U G G U A G G(A)U)A G A(G)(G) C U A C C A U G G U(U)U U A)A(C)G G G)U G						37 Zea mays
U(A)U(C) - A)A C U(U)U C G)A U G G U A G G(A)U)A G A(G)(G) C U A C C A U G G U(U)U U A)A(C)G G G)U G						38 Arabidopsis thaliana
U(A)U(C) - A)A C U(U)U C G)A U G G U A G G(A)U)A G A(G)(G) C U A C C A U G G U(U)U U A)A(C)G G G)U G						39 Glycine max
U(A)U(C) - A)A C U(U)U C G)A U G G U A G G(A)U)A G A(G)(G) C U A C C A U G G U(U)U U A)A(C)G G G)U G						40 Sinapis alba
U(A)U(C) - A)A C U(U)U C G)A U G G U A G G(A)U)A G A(G)(G) C U A C C A U G G U(U)U U A)A(C)G G G)U G						41 Zamia pumila
U(A)U(C) - A)A C U(U)U C G)A U G G U A G G(A)U)A G A(G)(G) C U A C C A U G G U(U)U U A)A(C)G G G)U G						42 Gnetum leyboldii
U(A)U(C) - A)A C U(U)U C G)A U G G U A G G(A)U)A G A(G)(G) C U A C C A U G G U(U)U U A)A(C)G G G)U G						43 Ginkgo biloba
U(A)U(C) - A)A C U(U)U C G)A U G G U A G G(A)U)A G A(G)(G) C U A C C A U G G U(U)U U A)A(C)G G G)U G						44 Pinus luchuensis
U(A)U(C) - A)A C U(U)U C G)A U G G U A G G(A)U)A G A(G)(G) C U A C C A U G G U(U)U U A)A(C)G G G)U G						45 Pinus walllichiana
U(A)U(C) - A)A C U(U)U C G)A U G G U A G G(A)U)A G A(G)(G) C U A C C A U G G U(U)U U A)A(C)G G G)U G						46 Podocarpus nakaai
U(A)U(C) - A)A C U(U)U C G)A U G G U A G G(A)U)A G A(G)(G) C U A C C A U G G U(U)U U A)A(C)G G G)U G						47 Taxus mairei
U(A)U(C) - A)A C U(U)U C G)A U G G U A G G(A)U)A G A(G)(G) C U A C C A U G G U(U)U U A)A(C)G G G)U G						48 Adiantum raddianum
U(A)U(C) - A)A C U(U)U C G)A U G G U A G G(A)U)A G A(G)(G) C U A C C A U G G U(U)U U A)A(C)G G G)U G						49 Adiantum raddianum 2
U(A)U(C) - A)A C U(U)U C G)A U G G U A G G(A)U)A G A(G)(G) C U A C C A U G G U(U)U U A)A(C)G G G)U G						50 Ophioglossum petiolatum
U(A)U(C) - A)A C U(U)U C G)A U G G U A G G(A)U)A G A(G)(G) C U A C C A U G G U(U)U U A)A(C)G G G)U G						51 Osmunda cinnamomea
U(A)U(C) - A)A C U(U)U C G)A U G G U A G G(A)U)A G A(G)(G) C U A C C A U G G U(U)U U A)A(C)G G G)U G						52 Salvinia natans
U(A)U(C) - A)A C U(U)U C G)A U G G U A G G(A)U)A G A(G)(G) C U A C C A U G G U(U)U U A)A(C)G G G)U G						53 Psilotum nudum 2
U(A)U(C) - A)A C U(U)U C G)A U G G U A G G(A)U)A G A(G)(G) C U A C C A U G G U(U)U U A)A(C)G G G)U G						54 Chara foetida
U(A)U(C) - A)A C U(U)U C G)A U G G U A G G(A)U)A G A(G)(G) C U A C C A U G G U(U)U U A)A(C)G G G)U G						55 Coleochaete scutata
U(A)U(C) - A)A C U(U)U C G)A U G G U A G G(A)U)A G A(G)(G) C U A C C A U G G U(U)U U A)A(C)G G G)U G						56 Klebsormidium flaccidum
U(A)U(C) - A)A C U(U)U C G)A U G G U A G G(A)U)A G A(G)(G) C U A C C A U G G U(U)U U A)A(C)G G G)U G						57 Nitella flexilis
U(A)U(C) - A)A C U(U)U C G)A U G G U A G G(A)U)A G A(G)(G) C U A C C A U G G U(U)U U A)A(C)G G G)U G						58 Nitella sp.
U(A)U(C) - A)A C U(U)U C G)A U G G U A G G(A)U)A G A(G)(G) C U A C C A U G G U(U)U U A)A(C)G G G)U G						59 Chlamydomonas reinhardtii
U(A)U(C) - A)A C U(U)U C G)A U G G U A G G(A)U)A G A(G)(G) C U A C C A U G G U(U)U U A)A(C)G G G)U G						60 Hydrodictyon reticulatum
U(A)U(C) - A)A C U(U)U C G)A U G G U A G G(A)U)A G A(G)(G) C U A C C A U G G U(U)U U A)A(C)G G G)U G						61 Parietochloris pseudoalveolaris
U(A)U(C) - A)A C U(U)U C G)A U G G U A G G(A)U)A G A(G)(G) C U A C C A U G G U(U)U U A)A(C)G G G)U A						62 Dictyostelium discoideum
U(A)U(C) - A)A C U(U)U C G)A U G G U A G G(A)U)A G A(G)(G) C U A C C A U G G U(U)U U A)A(C)G G G)U A						63 Helix numbering 1
U(A)U(C) - A)A C U(U)U C G)A U G G U A G G(A)U)A G A(G)(G) C U A C C A U G G U(U)U U A)A(C)G G G)U G						64 Pteris vittata
U(A)U(C) - A)A C U(U)U C G)A U G G U A G G(A)U)A G A(G)(G) C U A C C A U G G U(U)U U A)A(C)G G G)U G						65 Megaceros aenigmaticus
U(A)U(C) - A)A C U(U)U C G)A U G G U A G G(A)U)A G A(G)(G) C U A C C A U G G U(U)U U A)A(C)G G G)U G						66 Notophylas breutelii
U(A)U(C) - A)A C U(U)U C G)A U G G U A G G(A)U)A G A(G)(G) C U A C C A U G G U(U)U U A)A(C)G G G)U G						67 Physcomitrella patens
U(A)U(C) - A)A C U(U)U C G)A U G G U A G G(A)U)A G A(G)(G) C U A C C A U G G U(U)U U A)A(C)G G G)U G						68 Sphagnum palustre

DCSE Alignment

460	470	480	490	500
A[C G G G A A U]A A G G I G U U C I G A U]U C C G*G A G]A[G G I G]A G C C]U G A G A A A C[G G C U(A)C C]				1 Saccharomyces cerevisiae
A[C G G G A G A A U]U A G G I G U U C I G A U]U C C G*G A G]A[G G I G]A G C C]U G A G A A A C[G G C U(A)C C]				2 Schizosaccharomyces pombe
A[C G G G A G A A U]U A G G I G U U C I G A U]U C C G*G A G]A[G G I G]A G C C]U G A G A A A C[G G C U(A)C C]				3 Physcomitrella patens
A[C G G G A G A A U]U A G G I G U U C I G A U]U C C G*G A G]A[G G I G]A G C C]U G A G A A A C[G G C U(A)C C]				4 Anthoceros agrestis
A[C G G G A G A A U]U A G G I G U U C I G A U]U C C G*G A G]A[G G I G]A G C C]U G A G A A A C[G G C U(A)C C]				5 Anthoceros laevis
A[C G G G A G A A U]U A G G I G U U C I G A U]U C C G*G A G]A[G G I G]A G C C]U G A G A A A C[G G C U(A)C C]				6 Pellia epiphylla
A[C G G G A G A A U]U A G G I G U U C I G A U]U C C G*G A G]A[G G I G]A G C C]U G A G A A A C[G G C U(A)C C]				7 Reboulia hemisphaerica 2
A[C G G G A G A A U]U A G G I G U U C I G A U]U C C G*G A G]A[G G I G]A G C C]U G A G A A A C[G G C U(A)C C]				8 Riccardia pinguis
A[C G G G A G A A U]U A G G I G U U C I G A U]U C C G*G A G]A[G G I G]A G C C]U G A G A A A C[G G C U(A)C C]				9 Sphaerocarpos donnellii
A[C G G G A G A A U]U A G G I G U U C I G A U]U C C G*G A G]A[G G I G]A G C C]U G A G A A A C[G G C U(A)C C]				10 Sphaerocarpos texanus
A[C G G G A G A A U]U A G G I G U U C I G A U]U C C G*G A G]A[G G I G]A G C C]U G A G A A A C[G G C U(A)C C]				11 Atrichum undulatum
A[C G G G A G A A U]U A G G I G U U C I G A U]U C C G*G A G]A[G G I G]A G C C]U G A G A A A C[G G C U(A)C C]				12 Bryum argenteum
A[C G G G A G A A U]U A G G I G U U C I G A U]U C C G*G A G]A[G G I G]A G C C]U G A G A A A C[G G C U(A)C C]				13 Eurhynchium hians
A[C G G G A G A A U]U A G G I G U U C I G A U]U C C G*G A G]A[G G I G]A G C C]U G A G A A A C[G G C U(A)C C]				14 Funaria hygrometrica
A[C G G G A G A A U]U A G G I G U U C I G A U]U C C G*G A G]A[G G I G]A G C C]U G A G A A A C[G G C U(A)C C]				15 Leptobryum pyriformae
A[C G G G A G A A U]U A G G I G U U C I G A U]U C C G*G A G]A[G G I G]A G C C]U G A G A A A C[G G C U(A)C C]				16 Mnium hornum
A[C G G G A G A A U]U A G G I G U U C I G A U]U C C G*G A G]A[G G I G]A G C C]U G A G A A A C[G G C U(A)C C]				17 Physcomitrium pyriforme
A[C G G G A G A A U]U A G G I G U U C I G A U]U C C G*G A G]A[G G I G]A G C C]U G A G A A A C[G G C U(A)C C]				18 Polytrichum formosum
A[C G G G A G A A U]U A G G I G U U C I G A U]U C C G*G A G]A[G G I G]A G C C]U G A G A A A C[G G C U(A)C C]				19 Sphagnum cuspidatum
A[C G G G A G A A U]U A G G I G U U C I G A U]U C C G*G A G]A[G G I G]A G C C]U G A G A A A C[G G C U(A)C C]				20 Calypogeia arguta
A[C G G G A G A A U]U A G G I G U U C I G A U]U C C G*G A G]A[G G I G]A G C C]U G A G A A A C[G G C U(A)C C]				21 Conocephalum concicum
A[C G G G A G A A U]U A G G I G U U C I G A U]U C C G*G A G]A[G G I G]A G C C]U G A G A A A C[G G C U(A)C C]				22 Fossoambrosia pusilla
A[C G G G A G A A U]U A G G I G U U C I G A U]U C C G*G A G]A[G G I G]A G C C]U G A G A A A C[G G C U(A)C C]				23 Marchantia polymorpha
A[C G G G A G A A U]U A G G I G U U C I G A U]U C C G*G A G]A[G G I G]A G C C]U G A G A A A C[G G C U(A)C C]				24 Riccia fluitans
A[C G G G A G A A U]U A G G I G U U C I G A U]U C C G*G A G]A[G G I G]A G C C]U G A G A A A C[G G C U(A)C C]				25 Scapania nemorea
A[C G G G A G A A U]U A G G I G U U C I G A U]U C C G*G A G]A[G G I G]A G C C]U G A G A A A C[G G C U(A)C C]				26 Equisetum hyemale
A[C G G G A G A A U]U A G G I G U U C I G A U]U C C G*G A G]A[G G I G]A G C C]U G A G A A A C[G G C U(A)C C]				27 Equisetum robustum
A[C G G G A G A A U]U A G G I G U U C I G A U]U C C G*G A G]A[G G I G]A G C C]U G A G A A A C[G G C U(A)C C]				28 Isoetes engelmannii
A[C G G G A G A A U]U A G G I G U U C I G A U]U C C G*G A G]A[G G I G]A G C C]U G A G A A A C[G G C U(A)C C]				29 Huperzia lucidula
A[C G G G A G A A U]U A G G I G U U C I G A U]U C C G*G A G]A[G G I G]A G C C]U G A G A A A C[G G C U(A)C C]				30 Lycopodiella inundata
A[C G G G A G A A U]U A G G I G U U C I G A U]U C C G*G A G]A[G G I G]A G C C]U G A G A A A C[G G C U(A)C C]				31 Lycopodium phlegmaria
A[C G G G A G A A U]U A G G I G U U C I G A U]U C C G*G A G]A[G G I G]A G C C]U G A G A A A C[G G C U(A)C C]				32 Lycopodium taxifolium
A[C G G G A G A A U]U A G G I G U U C I G A U]U C C G*G A G]A[G G I G]A G C C]U G A G A A A C[G G C U(A)C C]				33 Lycopodium tristachyum
A[C G G G A G A A U]U A G G I G U U C I G A U]U C C G*G A G]A[G G I G]A G C C]U G A G A A A C[G G C U(A)C C]				34 Selaginella galleottii
A[C G G G A G A A U]U A G G I G U U C I G A U]U C C G*G A G]A[G G I G]A G C C]U G A G A A A C[G G C U(A)C C]				35 Selaginella sp.
A[C G G G A G A A U]U A G G I G U U C I G A U]U C C G*G A G]A[G G I G]A G C C]U G A G A A A C[G G C U(A)C C]				36 Oryza sativa
A[C G G G A G A A U]U A G G I G U U C I G A U]U C C G*G A G]A[G G I G]A G C C]U G A G A A A C[G G C U(A)C C]				37 Zea mays
A[C G G G A G A A U]U A G G I G U U C I G A U]U C C G*G A G]A[G G I G]A G C C]U G A G A A A C[G G C U(A)C C]				38 Arabidopsis thaliana
A[C G G G A G A A U]U A G G I G U U C I G A U]U C C G*G A G]A[G G I G]A G C C]U G A G A A A C[G G C U(A)C C]				39 Glycine max
A[C G G G A G A A U]U A G G I G U U C I G A U]U C C G*G A G]A[G G I G]A G C C]U G A G A A A C[G G C U(A)C C]				40 Sinapis alba
A[C G G G A G A A U]U A G G I G U U C I G A U]U C C G*G A G]A[G G I G]A G C C]U G A G A A A C[G G C U(A)C C]				41 Zamia pumila
A[C G G G A G A A U]U A G G I G U U C I G A U]U C C G*G A G]A[G G I G]A G C C]U G A G A A A C[G G C U(A)C C]				42 Gnetum leyboldii
A[C G G G A G A A U]U A G G I G U U C I G A U]U C C G*G A G]A[G G I G]A G C C]U G A G A A A C[G G C U(A)C C]				43 Ginkgo biloba
A[C G G G A G A A U]U A G G I G U U C I G A U]U C C G*G A G]A[G G I G]A G C C]U G A G A A A C[G G C U(A)C C]				44 Pinus luchuensis
A[C G G G A G A A U]U A G G I G U U C I G A U]U C C G*G A G]A[G G I G]A G C C]U G A G A A A C[G G C U(A)C C]				45 Pinus wallichiana
A[C G G G A G A A U]U A G G I G U U C I G A U]U C C G*G A G]A[G G I G]A G C C]U G A G A A A C[G G C U(A)C C]				46 Podocarpus nakaui
A[C G G G A G A A U]U A G G I G U U C I G A U]U C C G*G A G]A[G G I G]A G C C]U G A G A A A C[G G C U(A)C C]				47 Taxus mairei
A[C G G G A G A A U]U A G G I G U U C I G A U]U C C G*G A G]A[G G I G]A G C C]U G A G A A A C[G G C U(A)C C]				48 Adiantum raddianum
A[C G G G A G A A U]U A G G I G U U C I G A U]U C C G*G A G]A[G G I G]A G C C]U G A G A A A C[G G C U(A)C C]				49 Adiantum raddianum 2
A[C G G G A G A A U]U A G G I G U U C I G A U]U C C G*G A G]A[G G I G]A G C C]U G A G A A A C[G G C U(A)C C]				50 Ophioglossum petiolatum
A[C G G G A G A A U]U A G G I G U U C I G A U]U C C G*G A G]A[G G I G]A G C C]U G A G A A A C[G G C U(A)C C]				51 Osmunda cinnamomea
A[C G G G A G A A U]U A G G I G U U C I G A U]U C C G*G A G]A[G G I G]A G C C]U G A G A A A C[G G C U(A)C C]				52 Salvinia natans
A[C G G G A G A A U]U A G G I G U U C I G A U]U C C G*G A G]A[G G I G]A G C C]U G A G A A A C[G G C U(A)C C]				53 Psilotum nudum 2
A[C G G G A G A A U]U A G G I G U U C I G A U]U C C G*G A G]A[G G I G]A G C C]U G A G A A A C[G G C U(A)C C]				54 Chara foetida
A[C G G G A G A A U]U A G G I G U U C I G A U]U C C G*G A G]A[G G I G]A G C C]U G A G A A A C[G G C U(A)C C]				55 Coleochaete scutata
A[C G G G A G A A U]U A G G I G U U C I G A U]U C C G*G A G]A[G G I G]A G C C]U G A G A A A C[G G C U(A)C C]				56 Klebsormidium flaccidum
A[C G G G A G A A U]U A G G I G U U C I G A U]U C C G*G A G]A[G G I G]A G C C]U G A G A A A C[G G C U(A)C C]				57 Nitella flexilis
A[C G G G A G A A U]U A G G I G U U C I G A U]U C C G*G A G]A[G G I G]A G C C]U G A G A A A C[G G C U(A)C C]				58 Nitella sp.
A[C G G G A G A A U]U A G G I G U U C I G A U]U C C G*G A G]A[G G I G]A G C C]U G A G A A A C[G G C U(A)C C]				59 Chlamydomonas reinhardtii
A[C G G G A G A A U]U A G G I G U U C I G A U]U C C G*G A G]A[G G I G]A G C C]U G A G A A A C[G G C U(A)C C]				60 Hydrodictyon reticulatum
A[C G G G A G A A U]U A G G I G U U C I G A U]U C C G*G A G]A[G G I G]A G C C]U G A G A A A C[G G C U(A)C C]				61 Parietochloris pseudoalveolaris
A[C G G G A G A A U]U A G G I G U U C I G A U]U C C G*G A G]A[G G I G]A G C C]U G A G A A A C[G G C U(A)C C]				62 Dictyostelium discoideum
A[C G G G A G A A U]U A G G I G U U C I G A U]U C C G*G A G]A[G G I G]A G C C]U G A G A A A C[G G C U(A)C C]				63 Helix numbering 1
A[C G G G A G A A U]U A G G I G U U C I G A U]U C C G*G A G]A[G G I G]A G C C]U G A G A A A C[G G C U(A)C C]				64 Pteris vittata
A[C G G G A G A A U]U A G G I G U U C I G A U]U C C G*G A G]A[G G I G]A G C C]U G A G A A A C[G G C U(A)C C]				65 Megaceros senigmaticus
A[C G G G A G A A U]U A G G I G U U C I G A U]U C C G*G A G]A[G G I G]A G C C]U G A G A A A C[G G C U(A)C C]				66 Notophyas breutelii
A[C G G G A G A A U]U A G G I G U U C I G A U]U C C G*G A G]A[G G I G]A G C C]U G A G A A A C[G G C U(A)C C]				67 Physcomitrella patens
A[C G G G A G A A U]U A G G I G U U C I G A U]U C C G*G A G]A[G G I G]A G C C]U G A G A A A C[G G C U(A)C C]				68 Sphagnum palustre

	510	520	530	540	550
}A(C(A)U	C(C A A G I G A - (A)G I G C A I G C A - G G C I G C G G C A A A (U U A C C I C A A)U C C U)A A U U - C I A				1 Saccharomyces cerevisiae
}A(C(A)U	C(C A A G I G A - (A)G I G C A I G C A - G G C I G C G G C A A A (U U A C C I C A A)U C C C G)A C A - [C G				2 Schizosaccharomyces pombe
}A(C(A)U	C(C A A G I G A - (A)G I G C A I G C A - G G C I G C G G C A A A (U U A C C I C A A)U C C C G)A C A - [C G				3 Physcomitrella patens
}A(C(A)U	C(C A A G I G A - (A)G I G C A I G C A - G G C I G C G G C A A A (U U A C C I C A A)U C C C G)A C A - [C G				4 Anthoceros agrestis
}A(C(A)U	C(C A A G I G A - (A)G I G C A I G C A - G G C I G C G G C A A A (U U A C C I C A A)U C C C G)A C A - [C G				5 Anthoceros laevis
}A(C(A)U	C(C A A G I G A - (A)G I G C A I G C A - G G C I G C G G C A A A (U U A C C I C A A)U C C C G)A C A - [C G				6 Pellia epiphylla
}A(C(A)U	C(C A A G I G A - (A)G I G C A I G C A - G G C I G C G G C A A A (U U A C C I C A A)U C C C G)A C A - [C G				7 Reboulia hemisphaerica 2
}A(C(A)U	C(C A A G I G A - (A)G I G C A I G C A - G G C I G C G G C A A A (U U A C C I C A A)U C C C G)A C A - [C G				8 Riccardia pinquus
}A(C(A)U	C(C A A G I G A - (A)G I G C A I G C A - G G C I G C G G C A A A (U U A C C I C A A)U C C C G)A C A - [C G				9 Sphaerocarpos donnelli
}A(C(A)U	C(C A A G I G A - (A)G I G C A I G C A - G G C I G C G G C A A A (U U A C C I C A A)U C C C G)A C A - [C G				10 Sphaerocarpos texanus
}A(C(A)U	C(C A A G I G A - (A)G I G C A I G C A - G G C I G C G G C A A A (U U A C C I C A A)U C C C G)A C A - [C G				11 Atrichum undulatum
}A(C(A)U	C(C A A G I G A - (A)G I G C A I G C A - G G C I G C G G C A A A (U U A C C I C A A)U C C C G)A C A - [C G				12 Bryum argenteum
}A(C(A)U	C(C A A G I G A - (A)G I G C A I G C A - G G C I G C G G C A A A (U U A C C I C A A)U C C C G)A C A - [C G				13 Eurhynchium hians
}A(C(A)U	C(C A A G I G A - (A)G I G C A I G C A - G G C I G C G G C A A A (U U A C C I C A A)U C C C G)A C A - [C G				14 Funaria hygrometrica
}A(C(A)U	C(C A A G I G A - (A)G I G C A I G C A - G G C I G C G G C A A A (U U A C C I C A A)U C C C G)A C A - [C G				15 Leptobryum pyriforme
}A(C(A)U	C(C A A G I G A - (A)G I G C A I G C A - G G C I G C G G C A A A (U U A C C I C A A)U C C C G)A C A - [C G				16 Mnium hornum
}A C A (U	C(C A A G I G A - (A)G I G C A I G C A - G G C I G C G G C A A A (U U A C C I C A A)U C C C G)A C A - [C G				17 Physcomitrium pyriforme
}A(C(A)U	C(C A A G I G A - (A)G I G C A I G C A - G G C I G C G G C A A A (U U A C C I C A A)U C C C G)A C A - [C G				18 Polytrichum formosum
}A(C(A)U	C(C A A G I G A - (A)G I G C A I G C A - G G C I G C G G C A A A (U U A C C I C A A)U C C C G)A C A - [C G				19 Sphagnum cuspidatum
}A(C(A)U	C(C A A G I G A - (A)G I G C A I G C A - G G C I G C G G C A A A (U U A C C I C A A)U C C C G)A C A - [C G				20 Calypogeia arguta
}A(C(A)U	C(C A A G I G A - (A)G I G C A I G C A - G G C I G C G G C A A A (U U A C C I C A A)U C C C G)A C A - [C G				21 Conocephalum conicum
}A C A (U	C(C A A G I G A - (A)G I G C A I G C A - G G C I G C G G C A A A (U U A C C I C A A)U C C C G)A C A - [C G				22 Fossombronina pusilla
}A(C(A)U	C(C A A G I G A - (A)G I G C A I G C A - G G C I G C G G C A A A (U U A C C I C A A)U C C C G)A C A - [C G				23 Marchantia polymorpha
}A(C(A)U	C(C A A G I G A - (A)G I G C A I G C A - G G C I G C G G C A A A (U U A C C I C A A)U C C C G)A C A - [C G				24 Riccia fluitans
}A(C(A)U	C(C A A G I G A - (A)G I G C A I G C A - G G C I G C G G C A A A (U U A C C I C A A)U C C C G)A C A - [C G				25 Scapania nemorea
}A(C(A)U	C(C A A G I G A - (A)G I G C A I G C A - G G C I G C G G C A A A (U U A C C I C A A)U C C C G)A C A - [C G				26 Equisetum hyemale
}A(C(A)U	C(C A A G I G A - (A)G I G C A I G C A - G G C I G C G G C A A A (U U A C C I C A A)U C C C G)A C A - [C G				27 Equisetum robustum
}A(C(A)U	C(C A A G I G A - (A)G I G C A I G C A - G G C I G C G G C A A A (U U A C C I C A A)U C C C G)A C A - [C G				28 Isoetes engelmannii
}A(C(A)U	C(C A A G I G A - (A)G I G C A I G C A - G G C I G C G G C A A A (U U A C C I C A A)U C C C G)A C A - [C G				29 Huperzia lucidula
}A(C(A)U	C(C A A G I G A - (A)G I G C A I G C A - G G C I G C G G C A A A (U U A C C I C A A)U C C C G)A C A - [C G				30 Lycopodiella inundata
}A(C(A)U	C(C A A G I G A - (A)G I G C A I G C A - G G C I G C G G C A A A (U U A C C I C A A)U C C C G)A C A - [C G				31 Lycopodium phlegmaria
}A(C(A)U	C(C A A G I G A - (A)G I G C A I G C A - G G C I G C G G C A A A (U U A C C I C A A)U C C C G)A C A - [C G				32 Lycopodium taxifolium
}A(C(A)U	C(C A A G I G A - (A)G I G C A I G C A - G G C I G C G G C A A A (U U A C C I C A A)U C C C G)A C A - [C G				33 Lycopodium tristachyum
}A(C(A)U	C(C A A G I G A - (A)G I G C A I G C A - G G C I G C G G C A A A (U U A C C I C A A)U C C C G)A C A - [C G				34 Selaginella gallectii
}A(C(A)U	C(C A A G I G A - (A)G I G C A I G C A - G G C I G C G G C A A A (U U A C C I C A A)U C C C G)A C A - [C G				35 Selaginella sp.
}A(C(A)U	C(C A A G I G A - (A)G I G C A I G C A - G G C I G C G G C A A A (U U A C C I C A A)U C C C G)A C A - [C G				36 Oryza sativa
}A(C(A)U	C(C A A G I G A - (A)G I G C A I G C A - G G C I G C G G C A A A (U U A C C I C A A)U C C C G)A C A - [C G				37 Zea mays
}A(C(A)U	C(C A A G I G A - (A)G I G C A I G C A - G G C I G C G G C A A A (U U A C C I C A A)U C C C G)A C A - [C G				38 Arabidopsis thaliana
}A(C(A)U	C(C A A G I G A - (A)G I G C A I G C A - G G C I G C G G C A A A (U U A C C I C A A)U C C C G)A C A - [C G				39 Glycine max
}A(C(A)U	C(C A A G I G A - (A)G I G C A I G C A - G G C I G C G G C A A A (U U A C C I C A A)U C C C G)A C A - [C G				40 Sinapis alba
}A(C(A)U	C(C A A G I G A - (A)G I G C A I G C A - G G C I G C G G C A A A (U U A C C I C A A)U C C C G)A C A - [C G				41 Zamia pumila
}A(C(A)U	C(C A A G I G A - (A)G I G C A I G C A - G G C I G C G G C A A A (U U A C C I C A A)U C C C G)A C A - [C G				42 Gnetum leyboldii
}A(C(A)U	C(C A A G I G A - (A)G I G C A I G C A - G G C I G C G G C A A A (U U A C C I C A A)U C C C G)A C A - [C G				43 Ginkgo biloba
}A(C(A)U	C(C A A G I G A - (A)G I G C A I G C A - G G C I G C G G C A A A (U U A C C I C A A)U C C C G)A C A - [C G				44 Pinus luchuensis
}A(C(A)U	C(C A A G I G A - (A)G I G C A I G C A - G G C I G C G G C A A A (U U A C C I C A A)U C C C G)A C A - [C G				45 Pinus wallichiana
}A(C(A)U	C(C A A G I G A - (A)G I G C A I G C A - G G C I G C G G C A A A (U U A C C I C A A)U C C C G)A C A - [C G				46 Podocarpus nakaii
}A(C(A)U	C(C A A G I G A - (A)G I G C A I G C A - G G C I G C G G C A A A (U U A C C I C A A)U C C C G)A C A - [C G				47 Taxus mairei
}A(C(A)U	C(C A A G I G A - (A)G I G C A I G C A - G G C I G C G G C A A A (U U A C C I C A A)U C C C G)A C A - [C G				48 Adiantum raddianum
}A(C(A)U	C(C A A G I G A - (A)G I G C A I G C A - G G C I G C G G C A A A (U U A C C I C A A)U C C C G)A C A - [C G				49 Adiantum raddianum 2
}A(C(A)U	C(C A A G I G A - (A)G I G C A I G C A - G G C I G C G G C A A A (U U A C C I C A A)U C C C G)A C A - [C G				50 Ophioglossum petiolatum
}A(C(A)U	C(C A A G I G A - (A)G I G C A I G C A - G G C I G C G G C A A A (U U A C C I C A A)U C C C G)A C A - [C G				51 Osmunda cinnamomea
}A(C(A)U	C(C A A G I G A - (A)G I G C A I G C A - G G C I G C G G C A A A (U U A C C I C A A)U C C C G)A C A - [C G				52 Salvinia natans
}A(C(A)U	C(C A A G I G A - (A)G I G C A I G C A - G G C I G C G G C A A A (U U A C C I C A A)U C C C G)A C A - [C G				53 Psilotum nudum 2
}A(C(A)U	C(C A A G I G A - (A)G I G C A I G C A - G G C I G C G G C A A A (U U A C C I C A A)U C C C G)A C A - [C G				54 Chara foetida
}A(C(A)U	C(C A A G I G A - (A)G I G C A I G C A - G G C I G C G G C A A A (U U A C C I C A A)U C C C G)A C A - [C G				55 Coleochaete scutata
}A(C(A)U	C(C A A G I G A - (A)G I G C A I G C A - G G C I G C G G C A A A (U U A C C I C A A)U C C C G)A C A - [C G				56 Klebsormidium flaccidum
}A(C(A)U	C(C A A G I G A - (A)G I G C A I G C A - G G C I G C G G C A A A (U U A C C I C A A)U C C C G)A C A - [C G				57 Nitella flexilis
}A(C(A)U	C(C A A G I G A - (A)G I G C A I G C A - G G C I G C G G C A A A (U U A C C I C A A)U C C C G)A C A - [C G				58 Nitella sp.
}A(C(A)U	C(C A A G I G A - (A)G I G C A I G C A - G G C I G C G G C A A A (U U A C C I C A A)U C C C G)A C A - [C G				59 Chlamydomonas reinhardtii
}A(C(A)U	C(C A A G I G A - (A)G I G C A I G C A - G G C I G C G G C A A A (U U A C C I C A A)U C C C G)A C A - [C G				60 Hydrodictyon reticulatum
}A(C(A)U	C(C A A G I G A - (A)G I G C A I G C A - G G C I G C G G C A A A (U U A C C I C A A)U C C C G)A C A - [C G				61 Parietochloris pseudoalveolaris
}A(C U U	C(U A C G I G A - A G I G C A I G C A - G G C I G C G G C A A A (U U A C C I C A A)U C C - C)A A U A C I G				62 Dictyostelium discoideum
- - 15 - -	- - 15 - -	- - 5 - -	- - - -	- - 16 - -	- - - -
}A(C(A)U	C(C A A G I G A - (A)G I G C A I G C A - G G C I G C G G C A A A (U U A C C I C A A)U C C C G)A C A - [C G				63 Helix numbering 1
}A(C(A)U	C(C A A G I G A - (A)G I G C A I G C A - G G C I G C G G C A A A (U U A C C I C A A)U C C C G)A C A - [C G				64 Pteris vittata
}A(C(A)U	C(C A A G I G A - (A)G I G C A I G C A - G G C I G C G G C A A A (U U A C C I C A A)U C C C G)A C A - [C G				65 Megaceros aenigmaticus
}A(C(A)U	C(C A A G I G A - (A)G I G C A I G C A - G G C I G C G G C A A A (U U A C C I C A A)U C C C G)A C A - [C G				66 Notophylas breutelii
}A(C(A)U	C(C A A G I G A - (A)G I G C A I G C A - G G C I G C G G C A A A (U U A C C I C A A)U C C C G)A C A - [C G				67 Physcomitrella patens
}A(C(A)U	C(C A A G I G A - (A)G I G C A I G C A - G G C I G C G G C A A A (U U A C C I C A A)U C C C G)A C A - [C G				68 Sphagnum palustre

560	570	580	590	600	
G G G(A -1G G U A G)U(G A C A)A U A A A U A A(C G A A U A C C A G G G G G C C C)U U U C -[G G G U C - U U G					1 Saccharomyces cerevisiae
G G G(A -1G G U A G)U(G A C A)A U A A A U A A(C G A A U A C C A G G G G G C C C)U U U C -[G G G U C - U U G					2 Schizosaccharomyces pombe
G G G(A -1G G U A G)U(G A C A)A U A A A U A A(C G A A U A C C A G G G G G C C C)U U U C -[G G G U C - U U G					3 Physcomitrella patens
G G G(A -1G G U A G)U(G A C A)A U A A A U A A(C G A A U A C C A G G G G G C C C)U U U C -[G G G U C - U U G					4 Anthoceros agrestis
G G G(A -1G G U A G)U(G A C A)A U A A A U A A(C G A A U A C C A G G G G G C C C)U U U C -[G G G U C - U U G					5 Anthoceros laevis
G G G(A -1G G U A G)U(G A C A)A U A A A U A A(C G A A U A C C A G G G G G C C C)U U U C -[G G G U C - U U G					6 Pellia epiphylla
G G G(A -1G G U A G)U(G A C A)A U A A A U A A(C G A A U A C C A G G G G G C C C)U U U C -[G G G U C - U U G					7 Rebutia hemisphaerica 2
G G G(A -1G G U A G)U(G A C A)A U A A A U A A(C G A A U A C C A G G G G G C C C)U U U C -[G G G U C - U U G					8 Riccardia pinguis
G G G(A -1G G U A G)U(G A C A)A U A A A U A A(C G A A U A C C A G G G G G C C C)U U U C -[G G G U C - U U G					9 Sphaerocarpos donnellii
G G G(A -1G G U A G)U(G A C A)A U A A A U A A(C G A A U A C C A G G G G G C C C)U U U C -[G G G U C - U U G					10 Sphaerocarpos texanus
G G G(A -1G G U A G)U(G A C A)A U A A A U A A(C G A A U A C C A G G G G G C C C)U U U C -[G G G U C - U U G					11 Atrichum undulatum
G G G(A -1G G U A G)U(G A C A)A U A A A U A A(C G A A U A C C A G G G G G C C C)U U U C -[G G G U C - U U G					12 Bryum argenteum
G G G(A -1G G U A G)U(G A C A)A U A A A U A A(C G A A U A C C A G G G G G C C C)U U U C -[G G G U C - U U G					13 Eurhynchium hians
G G G(A -1G G U A G)U(G A C A)A U A A A U A A(C G A A U A C C A G G G G G C C C)U U U C -[G G G U C - U U G					14 Funaria hygrometrica
G G G(A -1G G U A G)U(G A C A)A U A A A U A A(C G A A U A C C A G G G G G C C C)U U U C -[G G G U C - U U G					15 Leptobryum pyriforme
G G G(A -1G G U A G)U(G A C A)A U A A A U A A(C G A A U A C C A G G G G G C C C)U U U C -[G G G U C - U U G					16 Mnium hornum
G G G(A -1G G U A G)U(G A C A)A U A A A U A A(C G A A U A C C A G G G G G C C C)U U U C -[G G G U C - U U G					17 Physcomitrium pyriforme
G G G(A -1G G U A G)U(G A C A)A U A A A U A A(C G A A U A C C A G G G G G C C C)U U U C -[G G G U C - U U G					18 Polytrichum formosum
G G G(A -1G G U A G)U(G A C A)A U A A A U A A(C G A A U A C C A G G G G G C C C)U U U C -[G G G U C - U U G					19 Sphagnum cuspidatum
G G G(A -1G G U A G)U(G A C A)A U A A A U A A(C G A A U A C C A G G G G G C C C)U U U C -[G G G U C - U U G					20 Calypogeia arguta
G G G(A -1G G U A G)U(G A C A)A U A A A U A A(C G A A U A C C A G G G G G C C C)U U U C -[G G G U C - U U G					21 Conocephalum conicum
G G G(A -1G G U A G)U(G A C A)A U A A A U A A(C G A A U A C C A G G G G G C C C)U U U C -[G G G U C - U U G					22 Fossombronia pusilla
G G G(A -1G G U A G)U(G A C A)A U A A A U A A(C G A A U A C C A G G G G G C C C)U U U C -[G G G U C - U U G					23 Marchantia polymorpha
G G G(A -1G G U A G)U(G A C A)A U A A A U A A(C G A A U A C C A G G G G G C C C)U U U C -[G G G U C - U U G					24 Riccia fluitans
G G G(A -1G G U A G)U(G A C A)A U A A A U A A(C G A A U A C C A G G G G G C C C)U U U C -[G G G U C - U U G					25 Scapania nemorea
G G G(A -1G G U A G)U(G A C A)A U A A A U A A(C G A A U A C C A G G G G G C C C)U U U C -[G G G U C - U U G					26 Equisetum hyemale
G G G(A -1G G U A G)U(G A C A)A U A A A U A A(C G A A U A C C A G G G G G C C C)U U U C -[G G G U C - U U G					27 Equisetum robustum
G G G(A -1G G U A G)U(G A C A)A U A A A U A A(C G A A U A C C A G G G G G C C C)U U U C -[G G G U C - U U G					28 Isoetes engelmannii
G G G(A -1G G U A G)U(G A C A)A U A A A U A A(C G A A U A C C A G G G G G C C C)U U U C -[G G G U C - U U G					29 Huperzia lucidula
G G G(A -1G G U A G)U(G A C A)A U A A A U A A(C G A A U A C C A G G G G G C C C)U U U C -[G G G U C - U U G					30 Lycopodiella inundata
G G G(A -1G G U A G)U(G A C A)A U A A A U A A(C G A A U A C C A G G G G G C C C)U U U C -[G G G U C - U U G					31 Lycopodium phlegmaria
G G G(A -1G G U A G)U(G A C A)A U A A A U A A(C G A A U A C C A G G G G G C C C)U U U C -[G G G U C - U U G					32 Lycopodium taxifolium
G G G(A -1G G U A G)U(G A C A)A U A A A U A A(C G A A U A C C A G G G G G C C C)U U U C -[G G G U C - U U G					33 Lycopodium tristachyum
G G G(A -1G G U A G)U(G A C A)A U A A A U A A(C G A A U A C C A G G G G G C C C)U U U C -[G G G U C - U U G					34 Selaginella galleottii
G G G(A -1G G U A G)U(G A C A)A U A A A U A A(C G A A U A C C A G G G G G C C C)U U U C -[G G G U C - U U G					35 Selaginella sp.
G G G(A -1G G U A G)U(G A C A)A U A A A U A A(C G A A U A C C A G G G G G C C C)U U U C -[G G G U C - U U G					36 Oryza sativa
G G G(A -1G G U A G)U(G A C A)A U A A A U A A(C G A A U A C C A G G G G G C C C)U U U C -[G G G U C - U U G					37 Zea mays
G G G(A -1G G U A G)U(G A C A)A U A A A U A A(C G A A U A C C A G G G G G C C C)U U U C -[G G G U C - U U G					38 Arabidopsis thaliana
G G G(A -1G G U A G)U(G A C A)A U A A A U A A(C G A A U A C C A G G G G G C C C)U U U C -[G G G U C - U U G					39 Glycine max
G G G(A -1G G U A G)U(G A C A)A U A A A U A A(C G A A U A C C A G G G G G C C C)U U U C -[G G G U C - U U G					40 Sinapis alba
G G G(A -1G G U A G)U(G A C A)A U A A A U A A(C G A A U A C C A G G G G G C C C)U U U C -[G G G U C - U U G					41 Zamia pumila
G G G(A -1G G U A G)U(G A C A)A U A A A U A A(C G A A U A C C A G G G G G C C C)U U U C -[G G G U C - U U G					42 Gnetum leyboldii
G G G(A -1G G U A G)U(G A C A)A U A A A U A A(C G A A U A C C A G G G G G C C C)U U U C -[G G G U C - U U G					43 Ginkgo biloba
G G G(A -1G G U A G)U(G A C A)A U A A A U A A(C G A A U A C C A G G G G G C C C)U U U C -[G G G U C - U U G					44 Pinus luchuensis
G G G(A -1G G U A G)U(G A C A)A U A A A U A A(C G A A U A C C A G G G G G C C C)U U U C -[G G G U C - U U G					45 Pinus wallichiana
G G G(A -1G G U A G)U(G A C A)A U A A A U A A(C G A A U A C C A G G G G G C C C)U U U C -[G G G U C - U U G					46 Podocarpus nakaii
G G G(A -1G G U A G)U(G A C A)A U A A A U A A(C G A A U A C C A G G G G G C C C)U U U C -[G G G U C - U U G					47 Taxus mairei
G G G(A -1G G U A G)U(G A C A)A U A A A U A A(C G A A U A C C A G G G G G C C C)U U U C -[G G G U C - U U G					48 Adiantum raddianum
G G G(A -1G G U A G)U(G A C A)A U A A A U A A(C G A A U A C C A G G G G G C C C)U U U C -[G G G U C - U U G					49 Adiantum raddianum 2
G G G(A -1G G U A G)U(G A C A)A U A A A U A A(C G A A U A C C A G G G G G C C C)U U U C -[G G G U C - U U G					50 Ophioglossum petiolatum
G G G(A -1G G U A G)U(G A C A)A U A A A U A A(C G A A U A C C A G G G G G C C C)U U U C -[G G G U C - U U G					51 Osmunda cinnamomea
G G G(A -1G G U A G)U(G A C A)A U A A A U A A(C G A A U A C C A G G G G G C C C)U U U C -[G G G U C - U U G					52 Salvinia natans
G G G(A -1G G U A G)U(G A C A)A U A A A U A A(C G A A U A C C A G G G G G C C C)U U U C -[G G G U C - U U G					53 Psilotum nudum 2
G G G(A -1G G U A G)U(G A C A)A U A A A U A A(C G A A U A C C A G G G G G C C C)U U U C -[G G G U C - U U G					54 Chara foetida
G G G(A -1G G U A G)U(G A C A)A U A A A U A A(C G A A U A C C A G G G G G C C C)U U U C -[G G G U C - U U G					55 Coleochaete scutata
G G G(A -1G G U A G)U(G A C A)A U A A A U A A(C G A A U A C C A G G G G G C C C)U U U C -[G G G U C - U U G					56 Klebsormidium flaccidum
G G G(A -1G G U A G)U(G A C A)A U A A A U A A(C G A A U A C C A G G G G G C C C)U U U C -[G G G U C - U U G					57 Nitella flexilis
G G G(A -1G G U A G)U(G A C A)A U A A A U A A(C G A A U A C C A G G G G G C C C)U U U C -[G G G U C - U U G					58 Nitella sp.
G G G(A -1G G U A G)U(G A C A)A U A A A U A A(C G A A U A C C A G G G G G C C C)U U U C -[G G G U C - U U G					59 Chlamydomonas reinhardtii
G G G(A -1G G U A G)U(G A C A)A U A A A U A A(C G A A U A C C A G G G G G C C C)U U U C -[G G G U C - U U G					60 Hydrodictyon reticulatum
G G G(A -1G G U A G)U(G A C A)A U A A A U A A(C G A A U A C C A G G G G G C C C)U U U C -[G G G U C - U U G					61 Parietochloris pseudoalveolaris
G G G(A -1G G U A G)U(G A C A)A U A A A U A A(C G A A U A C C A G G G G G C C C)U U U C -[G G G U C - U U G					62 Dictyostelium discoideum
G G G(A -1G G U A G)U(G A C A)A U A A A U A A(C G A A U A C C A G G G G G C C C)U U U C -[G G G U C - U U G					63 Helix numbering 1
G G G(A -1G G U A G)U(G A C A)A U A A A U A A(C G A A U A C C A G G G G G C C C)U U U C -[G G G U C - U U G					64 Pteris vittata
G G G(A -1G G U A G)U(G A C A)A U A A A U A A(C G A A U A C C A G G G G G C C C)U U U C -[G G G U C - U U G					65 Megaceros senigmaticus
G G G(A -1G G U A G)U(G A C A)A U A A A U A A(C G A A U A C C A G G G G G C C C)U U U C -[G G G U C - U U G					66 Notophylax breutelii
G G G(A -1G G U A G)U(G A C A)A U A A A U A A(C G A A U A C C A G G G G G C C C)U U U C -[G G G U C - U U G					67 Physcomitrella patens
G G G(A -1G G U A G)U(G A C A)A U A A A U A A(C G A A U A C C A G G G G G C C C)U U U C -[G G G U C - U U G					68 Sphagnum palustre

	610	620	630	640	650		
U A(A)U U G	G	A	A	U	G	A	1 Saccharomyces cerevisiae
U A(A)U U G	G	A	A	U	G	A	2 Schizosaccharomyces pombe
U A(A)U U G	G	A	A	U	G	A	3 Physcomitrella patens
U A(A)U U G	G	A	A	U	G	A	4 Anthoceros agrestis
U A(A)U U G	G	A	A	U	G	A	5 Anthoceros laevis
U A(A)U U G	G	A	A	U	G	A	6 Pellia epiphylla
U A(A)U U G	G	A	A	U	G	A	7 Reboulia hemisphaerica 2
U A(A)U U G	G	A	A	U	G	A	8 Riccardia pinguis
U A(A)U U G	G	A	A	U	G	A	9 Sphaerocarpos donnellii
U A(A)U U G	G	A	A	U	G	A	10 Sphaerocarpos texanus
U A(A)U U G	G	A	A	U	G	A	11 Atrichum undulatum
U A(A)U U G	G	A	A	U	G	A	12 Bryum argenteum
U A(A)U U G	G	A	A	U	G	A	13 Eurhynchium hians
U A(A)U U G	G	A	A	U	G	A	14 Funaria hygrometrica
U A(A)U U G	G	A	A	U	G	A	15 Leptobryum pyriforme
U A(A)U U G	G	A	A	U	G	A	16 Mnium hornum
U A(A)U U G	G	A	A	U	G	A	17 Physcomitrium pyriforme
U A(A)U U G	G	A	A	U	G	A	18 Polytrichum formosum
U A(A)U U G	G	A	A	U	G	A	19 Sphagnum cuspidatum
U A(A)U U G	G	A	A	U	G	A	20 Calypogeia arguta
U A(A)U U G	G	A	A	U	G	A	21 Conocephalum conicum
U A(A)U U G	G	A	A	U	G	A	22 Fossombronina pusilla
U A(A)U U G	G	A	A	U	G	A	23 Marchantia polymorpha
U A(A)U U G	G	A	A	U	G	A	24 Riccia frutitans
U A(A)U U G	G	A	A	U	G	A	25 Scapania nemorea
U A(A)U U G	G	A	A	U	G	A	26 Equisetum hyemale
U A(A)U U G	G	A	A	U	G	A	27 Equisetum robustum
U A(A)U U G	G	A	A	U	G	A	28 Isoetes engelmannii
U A(A)U U G	G	A	A	U	G	A	29 Huperzia lucidula
U A(A)U U G	G	A	A	U	G	A	30 Lycopodiella inundata
U A(A)U U G	G	A	A	U	G	A	31 Lycopodium phlegmaria
U A(A)U U G	G	A	A	U	G	A	32 Lycopodium taxifolium
U A(A)U U G	G	A	A	U	G	A	33 Lycopodium tristachyum
U A(A)U U G	G	A	A	U	G	A	34 Selaginella gallectonii
U A(A)U U G	G	A	A	U	G	A	35 Selaginella sp.
U A(A)U U G	G	A	A	U	G	A	36 Oryza sativa
U A(A)U U G	G	A	A	U	G	A	37 Zea mays
U A(A)U U G	G	A	A	U	G	A	38 Arabidopsis thaliana
U A(A)U U G	G	A	A	U	G	A	39 Glycine max
U A(A)U U G	G	A	A	U	G	A	40 Sinapis alba
U A(A)U U G	G	A	A	U	G	A	41 Zamia pumila
U A(A)U U G	G	A	A	U	G	A	42 Gnetum leyboldii
U A(A)U U G	G	A	A	U	G	A	43 Ginkgo biloba
U A(A)U U G	G	A	A	U	G	A	44 Pinus luchuensis
U A(A)U U G	G	A	A	U	G	A	45 Pinus wallichiana
U A(A)U U G	G	A	A	U	G	A	46 Podocarpus nakaii
U A(A)U U G	G	A	A	U	G	A	47 Taxus mairei
U A(A)U U G	G	A	A	U	G	A	48 Adiantum raddianum
U A(A)U U G	G	A	A	U	G	A	49 Adiantum raddianum 2
U A(A)U U G	G	A	A	U	G	A	50 Ophioglossum petiolatum
U A(A)U U G	G	A	A	U	G	A	51 Osmunda cinnamomea
U A(A)U U G	G	A	A	U	G	A	52 Salvinia natans
U A(A)U U G	G	A	A	U	G	A	53 Psilotum nudum 2
U A(A)U U G	G	A	A	U	G	A	54 Chara foetida
U A(A)U U G	G	A	A	U	G	A	55 Coleochaete scutata
U A(A)U U G	G	A	A	U	G	A	56 Klebsormidium flaccidum
U A(A)U U G	G	A	A	U	G	A	57 Nitella flexilis
U A(A)U U G	G	A	A	U	G	A	58 Nitella sp.
U A(A)U U G	G	A	A	U	G	A	59 Chlamydomonas reinhardtii
U A(A)U U G	G	A	A	U	G	A	60 Hydrodictyon reticulatum
U A(A)U U G	G	A	A	U	G	A	61 Parietochloris pseudoalveolaris
U A(A)U U G	G	A	A	U	G	A	62 Dictyostelium discoideum
U A(A)U U G	G	A	A	U	G	A	63 Helix numbering 1
U A(A)U U G	G	A	A	U	G	A	64 Pteris vittata
U A(A)U U G	G	A	A	U	G	A	65 Megaceros senigmaticus
U A(A)U U G	G	A	A	U	G	A	66 Notophylas breutelu
U A(A)U U G	G	A	A	U	G	A	67 Physcomitrella patens
U A(A)U U G	G	A	A	U	G	A	68 Sphagnum palustre

660	670	680	690	700	
A G G C A G U U G G U G C C A G C A G C C G G U A A U U C A G C U C C A A U A G C G U A					1 Saccharomyces cerevisiae
A G G G C A A G U U G G U G C C C A G C C A G C C G G U A A U U C C A G C U C C C A A U A G C G U A					2 Schizosaccharomyces pombe
A G G G C A A G U U G G U G C C C C A G C C A G C C G G U A A U U C C A G C U C C C A A U A G C G U A					3 Physcomitrella patens
A G G G C A A G U U G G U G C C C C A G C C A G C C G G U A A U U C C A G C U C C C A A U A G C G U A					4 Anthoceros agrestis
A G G G C A A G U U G G U G C C C C A G C C A G C C G G U A A U U C C A G C U C C C A A U A G C G U A					5 Anthoceros laevis
A G G G C A A G U U G G U G C C C C A G C C A G C C G G U A A U U C C A G C U C C C A A U A G C G U A					6 Pellia epiphylla
A G G G C A A G U U G G U G C C C C A G C C A G C C G G U A A U U C C A G C U C C C A A U A G C G U A					7 Reboulia hemisphaerica 2
A G G G C A A G U U G G U G C C C C A G C C A G C C G G U A A U U C C A G C U C C C A A U A G C G U A					8 Riccardia pinquius
A G G G C A A G U U G G U G C C C C A G C C A G C C G G U A A U U C C A G C U C C C A A U A G C G U A					9 Sphaerocarpos donnelli
A G G G C A A G U U G G U G C C C C A G C C A G C C G G U A A U U C C A G C U C C C A A U A G C G U A					10 Sphaerocarpos texanus
A G G G C A A G U U G G U G C C C C A G C C A G C C G G U A A U U C C A G C U C C C A A U A G C G U A					11 Atrichum undulatum
A G G G C A A G U U G G U G C C C C A G C C A G C C G G U A A U U C C A G C U C C C A A U A G C G U A					12 Bryum argenteum
A G G G C A A G U U G G U G C C C C A G C C A G C C G G U A A U U C C A G C U C C C A A U A G C G U A					13 Eurhynchium hians
A G G G C A A G U U G G U G C C C C A G C C A G C C G G U A A U U C C A G C U C C C A A U A G C G U A					14 Funaria hygrometrica
A G G G C A A G U U G G U G C C C C A G C C A G C C G G U A A U U C C A G C U C C C A A U A G C G U A					15 Leptobryum pyriforme
A G G G C A A G U U G G U G C C C C A G C C A G C C G G U A A U U C C A G C U C C C A A U A G C G U A					16 Mnium hornum
A G G G C A A G U U G G U G C C C C A G C C A G C C G G U A A U U C C A G C U C C C A A U A G C G U A					17 Physcomitrium pyriforme
A G G G C A A G U U G G U G C C C C A G C C A G C C G G U A A U U C C A G C U C C C A A U A G C G U A					18 Polytrichum formosum
A G G G C A A G U U G G U G C C C C A G C C A G C C G G U A A U U C C A G C U C C C A A U A G C G U A					19 Sphagnum cuspidatum
A G G G C A A G U U G G U G C C C C A G C C A G C C G G U A A U U C C A G C U C C C A A U A G C G U A					20 Calypogeia arguta
A G G G C A A G U U G G U G C C C C A G C C A G C C G G U A A U U C C A G C U C C C A A U A G C G U A					21 Conocephalum conicum
A G G G C A A G U U G G U G C C C C A G C C A G C C G G U A A U U C C A G C U C C C A A U A G C G U A					22 Fossombronia pusilla
A G G G C A A G U U G G U G C C C C A G C C A G C C G G U A A U U C C A G C U C C C A A U A G C G U A					23 Marchantia polymorpha
A G G G C A A G U U G G U G C C C C A G C C A G C C G G U A A U U C C A G C U C C C A A U A G C G U A					24 Riccia fluitans
A G G G C A A G U U G G U G C C C C A G C C A G C C G G U A A U U C C A G C U C C C A A U A G C G U A					25 Scapania nemorea
A G G G C A A G U U G G U G C C C C A G C C A G C C G G U A A U U C C A G C U C C C A A U A G C G U A					26 Equisetum hyemale
A G G G C A A G U U G G U G C C C C A G C C A G C C G G U A A U U C C A G C U C C C A A U A G C G U A					27 Equisetum robustum
A G G G C A A G U U G G U G C C C C A G C C A G C C G G U A A U U C C A G C U C C C A A U A G C G U A					28 Isoetes engelmannii
A G G G C A A G U U G G U G C C C C A G C C A G C C G G U A A U U C C A G C U C C C A A U A G C G U A					29 Huperzia lucidula
A G G G C A A G U U G G U G C C C C A G C C A G C C G G U A A U U C C A G C U C C C A A U A G C G U A					30 Lycopodiella inundata
A G G G C A A G U U G G U G C C C C A G C C A G C C G G U A A U U C C A G C U C C C A A U A G C G U A					31 Lycopodium phlegmaria
A G G G C A A G U U G G U G C C C C A G C C A G C C G G U A A U U C C A G C U C C C A A U A G C G U A					32 Lycopodium taxifolium
A G G G C A A G U U G G U G C C C C A G C C A G C C G G U A A U U C C A G C U C C C A A U A G C G U A					33 Lycopodium tristachyum
A G G G C A A G U U G G U G C C C C A G C C A G C C G G U A A U U C C A G C U C C C A A U A G C G U A					34 Selaginella galieottii
A G G G C A A G U U G G U G C C C C A G C C A G C C G G U A A U U C C A G C U C C C A A U A G C G U A					35 Selaginella sp.
A G G G C A A G U U G G U G C C C C A G C C A G C C G G U A A U U C C A G C U C C C A A U A G C G U A					36 Oryza sativa
A G G G C A A G U U G G U G C C C C A G C C A G C C G G U A A U U C C A G C U C C C A A U A G C G U A					37 Zea mays
A G G G C A A G U U G G U G C C C C A G C C A G C C G G U A A U U C C A G C U C C C A A U A G C G U A					38 Arabidopsis thaliana
A G G G C A A G U U G G U G C C C C A G C C A G C C G G U A A U U C C A G C U C C C A A U A G C G U A					39 Glycine max
A G G G C A A G U U G G U G C C C C A G C C A G C C G G U A A U U C C A G C U C C C A A U A G C G U A					40 Sinapis alba
A G G G C A A G U U G G U G C C C C A G C C A G C C G G U A A U U C C A G C U C C C A A U A G C G U A					41 Zamia pumila
A G G G C A A G U U G G U G C C C C A G C C A G C C G G U A A U U C C A G C U C C C A A U A G C G U A					42 Gnetum leyboldii
A G G G C A A G U U G G U G C C C C A G C C A G C C G G U A A U U C C A G C U C C C A A U A G C G U A					43 Ginkgo biloba
A G G G C A A G U U G G U G C C C C A G C C A G C C G G U A A U U C C A G C U C C C A A U A G C G U A					44 Pinus luchuensis
A G G G C A A G U U G G U G C C C C A G C C A G C C G G U A A U U C C A G C U C C C A A U A G C G U A					45 Pinus wallichiana
A G G G C A A G U U G G U G C C C C A G C C A G C C G G U A A U U C C A G C U C C C A A U A G C G U A					46 Podocarpus nakaii
A G G G C A A G U U G G U G C C C C A G C C A G C C G G U A A U U C C A G C U C C C A A U A G C G U A					47 Taxus mairei
A G G G C A A G U U G G U G C C C C A G C C A G C C G G U A A U U C C A G C U C C C A A U A G C G U A					48 Adiantum raddianum
A G G G C A A G U U G G U G C C C C A G C C A G C C G G U A A U U C C A G C U C C C A A U A G C G U A					49 Adiantum raddianum 2
A G G G C A A G U U G G U G C C C C A G C C A G C C G G U A A U U C C A G C U C C C A A U A G C G U A					50 Ophioglossum petiolatum
A G G G C A A G U U G G U G C C C C A G C C A G C C G G U A A U U C C A G C U C C C A A U A G C G U A					51 Osmunda cinnamomea
A G G G C A A G U U G G U G C C C C A G C C A G C C G G U A A U U C C A G C U C C C A A U A G C G U A					52 Salvinia natans
A G G G C A A G U U G G U G C C C C A G C C A G C C G G U A A U U C C A G C U C C C A A U A G C G U A					53 Psilotum nudum 2
A G G G C A A G U U G G U G C C C C A G C C A G C C G G U A A U U C C A G C U C C C A A U A G C G U A					54 Chara foetida
A G G G C A A G U U G G U G C C C C A G C C A G C C G G U A A U U C C A G C U C C C A A U A G C G U A					55 Coleochaete scutata
A G G G C A A G U U G G U G C C C C A G C C A G C C G G U A A U U C C A G C U C C C A A U A G C G U A					56 Klebsormidium flaccidum
A G G G C A A G U U G G U G C C C C A G C C A G C C G G U A A U U C C A G C U C C C A A U A G C G U A					57 Nitella flexilis
A G G G C A A G U U G G U G C C C C A G C C A G C C G G U A A U U C C A G C U C C C A A U A G C G U A					58 Nitella sp.
A G G G C A A G U U G G U G C C C C A G C C A G C C G G U A A U U C C A G C U C C C A A U A G C G U A					59 Chlamydomonas reinhardtii
A G G G C A A G U U G G U G C C C C A G C C A G C C G G U A A U U C C A G C U C C C A A U A G C G U A					60 Hydrodictyon reticulatum
A G G G C A A G U U G G U G C C C C A G C C A G C C G G U A A U U C C A G C U C C C A A U A G C G U A					61 Parietochloris pseudoalveolaris
A G G G C A A G U U G G U G C C C C A G C C A G C C G G U A A U U C C A G C U C C C A A U A G C G U A					62 Dictyostelium discoideum
- - - 20 - - - - 21 - - - - - 20' - - - - - 21' - - - - 19' - - - - - 3'					63 Helix numbering 1
A G G G C A A G U U G G U G C C C C A G C C A G C C G G U A A U U C C A G C U C C C A A U A G C G U A					64 Pteris vittata
A G G G C A A G U U G G U G C C C C A G C C A G C C G G U A A U U C C A G C U C C C A A U A G C G U A					65 Megaceros aenigmaticus
A G G G C A A G U U G G U G C C C C A G C C A G C C G G U A A U U C C A G C U C C C A A U A G C G U A					66 Nectophylax breutelii
A G G G C A A G U U G G U G C C C C A G C C A G C C G G U A A U U C C A G C U C C C A A U A G C G U A					67 Physcomitrella patens
A G G G C A A G U U G G U G C C C C A G C C A G C C G G U A A U U C C A G C U C C C A A U A G C G U A					68 Sphagnum palustre

DCSE Alignment

	710	720	730	740	750	
U A U U A J A A G U U G U U G C A G U U A A A A A G C U C G U I A G U U G A J A C U U U G G G C C C I G I G						1 Saccharomyces cerevisiae
U A U U U A J A A G U U G U U G C A G U U A A A A A G C U C G U I A G U U G A J A C U U U G G A I G I C C U G G G						2 Schizosaccharomyces pombe
U A U U U U A A A G U U G U U G C A G U U A A A A A G C U C G U I A G U U G G J A C U U U G G G G U C I G I G G						3 Physcomitrella patens
U A U U U U A A A G U U G U U G C A G U U A A A A A G C U C G U I A G U U G G J A C U U U G G G G U C I G I G G						4 Anthoceros agrestis
U A U U U U A A A G U U G U U G C A G U U A A A A A G C U C G U I A G U U G G J A C U U U G G G G U C I G I G G						5 Anthoceros laevis
U A U U U U A A A G U U G U U G C A G U U A A A A A G C U C G U I A G U U G G J A C U U U G G G A U G G G G G						6 Peelia epiphylla
U A U U U U A A A G U U G U U G C A G U U A A A A A G C U C G U I A G U U G G J A C U U U G G G G A C G G G G						7 Reboulia hemisphaerica 2
U A U U U U A A A G U U G U U G C A G U U A A A A A G C U C G U I A G U U G G J A C U U U G G G G A C G G G G						8 Riccardia pinquus
U A U U U U A A A G U U G U U G C A G U U A A A A A G C U C G U I A G U U G G J A C U U U G G G G A C G G G G						9 Sphaerocarpos donnelli
U A U U U U A A A G U U G U U G C A G U U A A A A A G C U C G U I A G U U G G J A C U U U G G G G A C G G G G						10 Sphaerocarpos texanus
U A U U U U A A A G U U G U U G C A G U U A A A A A G C U C G U I A G U U G G J A C U U U G G G G A C G G G G						11 Atrichum undulatum
U A U U U U A A A G U U G U U G C A G U U A A A A A G C U C G U I A G U U G G J A C U U U G G G G A C G G G G						12 Bryum argenteum
U A U U U U A A A G U U G U U G C A G U U A A A A A G C U C G U I A G U U G G J A C U U U G G G G A C G G G G						13 Eurhynchium hians
U A U U U U A A A G U U G U U G C A G U U A A A A A G C U C G U I A G U U G G J A C U U U G G G G A C G G G G						14 Funaria hygrometrica
U A U U U U A A A G U U G U U G C A G U U A A A A A G C U C G U I A G U U G G J A C U U U G G G G A C G G G G						15 Leptobryum pyriforme
U A U U U U A A A G U U G U U G C A G U U A A A A A G C U C G U I A G U U G G J A C U U U G G G G A C G G G G						16 Mnium hornum
U A U U U U A A A G U U G U U G C A G U U A A A A A G C U C G U I A G U U G G J A C U U U G G G G A C G G G G						17 Physcomitrium pyriforme
U A U U U U A A A G U U G U U G C A G U U A A A A A G C U C G U I A G U U G G J A C U U U G G G G A C G G G G						18 Polytichum formosum
U A U U U U A A A G U U G U U G C A G U U A A A A A G C U C G U I A G U U G G J A C U U U G G G G A C G G G G						19 Sphagnum cuspidatum
U A U U U U A A A G U U G U U G C A G U U A A A A A G C U C G U I A G U U G G J A C U U U G G G G A C G G G G						20 Calypogeia arguta
U A U U U U A A A G U U G U U G C A G U U A A A A A G C U C G U I A G U U G G J A C U U U G G G G A C G G G G						21 Conocephalum conicum
U A U U U U A A A G U U G U U G C A G U U A A A A A G C U C G U I A G U U G G J A C U U U G G G G A C G G G G						22 Fossombronina pusilla
U A U U U U A A A G U U G U U G C A G U U A A A A A G C U C G U I A G U U G G J A C U U U G G G G A C G G G G						23 Marchantia polymorpha
U A U U U U A A A G U U G U U G C A G U U A A A A A G C U C G U I A G U U G G J A C U U U G G G G A C G G G G						24 Riccia fluitans
U A U U U U A A A G U U G U U G C A G U U A A A A A G C U C G U I A G U U G G J A C U U U G G G G A C G G G G						25 Scapania nemorea
U A U U U U A A A G U U G U U G C A G U U A A A A A G C U C G U I A G U U G G J A C U U U G G G G A C G G G G						26 Equisetum hyemale
U A U U U U A A A G U U G U U G C A G U U A A A A A G C U C G U I A G U U G G J A C U U U G G G G A C G G G G						27 Equisetum robustum
U A U U U U A A A G U U G U U G C A G U U A A A A A G C U C G U I A G U U G G J A C U U U G G G G A C G G G G						28 Isoetes engelmannii
U A U U U U A A A G U U G U U G C A G U U A A A A A G C U C G U I A G U U G G J A C U U U G G G G A C G G G G						29 Huperzia lucidula
U A U U U U A A A G U U G U U G C A G U U A A A A A G C U C G U I A G U U G G J A C U U U G G G G A C G G G G						30 Lycopodiella inundata
U A U U U U A A A G U U G U U G C A G U U A A A A A G C U C G U I A G U U G G J A C U U U G G G G A C G G G G						31 Lycopodium phlegmaria
U A U U U U A A A G U U G U U G C A G U U A A A A A G C U C G U I A G U U G G J A C U U U G G G G A C G G G G						32 Lycopodium taxifolium
U A U U U U A A A G U U G U U G C A G U U A A A A A G C U C G U I A G U U G G J A C U U U G G G G A C G G G G						33 Lycopodium tristachyum
U A U U U U A A A G U U G U U G C A G U U A A A A A G C U C G U I A G U U G G J A C U U U G G G G A C G G G G						34 Selaginella galieortzii
U A U U U U A A A G U U G U U G C A G U U A A A A A G C U C G U I A G U U G G J A C U U U G G G G A C G G G G						35 Selaginella sp.
U A U U U U A A A G U U G U U G C A G U U A A A A A G C U C G U I A G U U G G J A C U U U G G G G A C G G G G						36 Oryza sativa
U A U U U U A A A G U U G U U G C A G U U A A A A A G C U C G U I A G U U G G J A C U U U G G G G A C G G G G						37 Zea mays
U A U U U U A A A G U U G U U G C A G U U A A A A A G C U C G U I A G U U G G J A C U U U G G G G A C G G G G						38 Arabidopsis thaliana
U A U U U U A A A G U U G U U G C A G U U A A A A A G C U C G U I A G U U G G J A C U U U G G G G A C G G G G						39 Glycine max
U A U U U U A A A G U U G U U G C A G U U A A A A A G C U C G U I A G U U G G J A C U U U G G G G A C G G G G						40 Sinapis alba
U A U U U U A A A G U U G U U G C A G U U A A A A A G C U C G U I A G U U G G J A C U U U G G G G A C G G G G						41 Zamia pumila
U A U U U U A A A G U U G U U G C A G U U A A A A A G C U C G U I A G U U G G J A C U U U G G G G A C G G G G						42 Gnetum leyboldii
U A U U U U A A A G U U G U U G C A G U U A A A A A G C U C G U I A G U U G G J A C U U U G G G G A C G G G G						43 Ginkgo biloba
U A U U U U A A A G U U G U U G C A G U U A A A A A G C U C G U I A G U U G G J A C U U U G G G G A C G G G G						44 Pinus luchuensis
U A U U U U A A A G U U G U U G C A G U U A A A A A G C U C G U I A G U U G G J A C U U U G G G G A C G G G G						45 Pinus wallichiana
U A U U U U A A A G U U G U U G C A G U U A A A A A G C U C G U I A G U U G G J A C U U U G G G G A C G G G G						46 Podocarpus nakaii
U A U U U U A A A G U U G U U G C A G U U A A A A A G C U C G U I A G U U G G J A C U U U G G G G A C G G G G						47 Taxus mairei
U A U U U U A A A G U U G U U G C A G U U A A A A A G C U C G U I A G U U G G J A C U U U G G G G A C G G G G						48 Adiantum raddianum
U A U U U U A A A G U U G U U G C A G U U A A A A A G C U C G U I A G U U G G J A C U U U G G G G A C G G G G						49 Adiantum raddianum 2
U A U U U U A A A G U U G U U G C A G U U A A A A A G C U C G U I A G U U G G J A C U U U G G G G A C G G G G						50 Ophioglossum petiolatum
U A U U U U A A A G U U G U U G C A G U U A A A A A G C U C G U I A G U U G G J A C U U U G G G G A C G G G G						51 Osmunda cinnamomea
U A U U U U A A A G U U G U U G C A G U U A A A A A G C U C G U I A G U U G G J A C U U U G G G G A C G G G G						52 Salvinia natans
U A U U U U A A A G U U G U U G C A G U U A A A A A G C U C G U I A G U U G G J A C U U U G G G G A C G G G G						53 Psilotum nudum 2
U A U U U U A A A G U U G U U G C A G U U A A A A A G C U C G U I A G U U G G J A C U U U G G G G A C G G G G						54 Chara foetida
U A U U U U A A A G U U G U U G C A G U U A A A A A G C U C G U I A G U U G G J A C U U U G G G G A C G G G G						55 Coleochaete scutata
U A U U U U A A A G U U G U U G C A G U U A A A A A G C U C G U I A G U U G G J A C U U U G G G G A C G G G G						56 Klebsormidium flaccidum
U A U U U U A A A G U U G U U G C A G U U A A A A A G C U C G U I A G U U G G J A C U U U G G G G A C G G G G						57 Nitella flexilis
U A U U U U A A A G U U G U U G C A G U U A A A A A G C U C G U I A G U U G G J A C U U U G G G G A C G G G G						58 Nitella sp.
U A U U U U A A A G U U G U U G C A G U U A A A A A G C U C G U I A G U U G G J A C U U U G G G G A C G G G G						59 Chlamydomonas reinhardtii
U A U U U U A A A G U U G U U G C A G U U A A A A A G C U C G U I A G U U G G J A C U U U G G G G A C G G G G						60 Hydrodictyon reticulatum
U A U U U U A A A G U U G U U G C A G U U A A A A A G C U C G U I A G U U G G J A C U U U G G G G A C G G G G						61 Parietochloris pseudoalveolaris
U A U U U U A A A G U U G U U G C A G U U A A A A A G C U C G U I A G U U G G J A C U U U G G G G A C G G G G						62 Dictyostelium discoideum
U A U U U U A A A G U U G U U G C A G U U A A A A A G C U C G U I A G U U G G J A C U U U G G G G A C G G G G						63 Helix numbering 1
U A U U U U A A A G U U G U U G C A G U U A A A A A G C U C G U I A G U U G G J A C U U U G G G G A C G G G G						64 Pteris vittata
U A U U U U A A A G U U G U U G C A G U U A A A A A G C U C G U I A G U U G G J A C U U U G G G G A C G G G G						65 Megaceros senigmaticus
U A U U U U A A A G U U G U U G C A G U U A A A A A G C U C G U I A G U U G G J A C U U U G G G G A C G G G G						66 Notophylas breutelii
U A U U U U A A A G U U G U U G C A G U U A A A A A G C U C G U I A G U U G G J A C U U U G G G G A C G G G G						67 Physcomitrella patens
U A U U U U A A A G U U G U U G C A G U U A A A A A G C U C G U I A G U U G G J A C U U U G G G G A C G G G G						68 Sphagnum palustre

DCSE Alignment

	760	770	780	790	800	
U U G(- G)C C - G G U(C)C G - A) - - - - - U U U U U - - - - - (U C G) - U						1 Saccharomyces cerevisiae
U C G(- A)C U - G G U(C)C G - C(C)G C] - [G C G U G] - U						2 Schizosaccharomyces pombe
G G G(A)G - C - G G U(C)C G - C C C] - [G G G U G] - U						3 Physcomitrella patens
G U(G) - G C U - G G U(C)C G - C C U] - [G G G U G] - U						4 Anthoceros agrestis
G(U G - G A)C - G G U(C)C G - C U] - [G G U G] - U						5 Anthoceros laevis
(C) - - G A(G)C - G G U(C)C G - C C] - [G G U G] - U						6 Pellia epiphylla
U C - C G G)C - G G U(C)C G - (G)C] - [G(G) U G] - C						7 Reboulia hemisphaerica 2
G(C - G A)G C - G G U(C)C G - C C] - [G G U G] - U						8 Riccardia pinguis
U C - G G(G)C - G G U(C)C G - C U] - [G G U G] - U						9 Sphaerocarpos donnelli
U(C - G G G - -G G U(C)C G - C C] - [G G U G] - U						10 Sphaerocarpos texanus
G(C - G A)G C - G G U(C)C G - C C] - [G G U G] - U						11 Atrichum undulatum
(G)G - G A(G)C - G G U(C)C G - C C] - [G G U G] - U						12 Bryum argenteum
U G - G A(G U C)G G U(C)C G - C U] - [G G U(C) - U						13 Eurhynchium hians
(G)G - G A(G)C - G G U(C)C G - C C C] - [G G G U G] - U						14 Funaria hygrometrica
(G)G - G A(G)C - G G U(C)C G - C C C] - [G G G U G] - U						15 Leptobryum pyriforme
(G)G - G A(G)C - G G U(C)C G - C C C] - [G G G U G] - U						16 Mnium hornum
I G G - I G A(G)C - G G U(C)C G - C C] - [G G G U G] - U						17 Physcomitrium pyriforme
G A - G A(G)C - G G U(C)C G - C C] - [G G U G] - U						18 Polytrichum formosum
G(C - I G G(G)C - G G U(C)C G - C U] - [G G U G] - U						19 Sphagnum cuspidatum
G(C - I G(A)G C - G G U(C)C G - C C] - [G G U G] - U						20 Calypogeia arguta
(C) U - U(G)G C - G G U(C)C G - C U] - [G G U G] - U						21 Conocephalum conicum
(C) - - G A(G)C - G G U(C)C G - C C] - [G G U G] - U						22 Fossombronia pusilla
C - - C G(G)C - G G U(C)C G - C U] - [G G U G] - U						23 Marchantia polymorpha
(C) - - G A(G)C - G G U(C)C G - C C] - [G G U G] - U						24 Riccia fluitans
I G C - G U G)C - G G U(C)C G - C C] - [G G U G] - U						25 Scapania nemorea
I G C - G U G)C - G G U(C)C G - C C] - [G G U G] - U						26 Equisetum hyemale
A U - G C(A)C - G G U(C)C G - C C C] - [G G U G] - U						27 Equisetum robustum
G C - G A(A)C - I G G U(C)C G - C C] - [G G U G] - U						28 Isoetes engelmannii
G C - G G(A)C - G G U(C)C G - C U U] - [G G U G] - U						29 Huperzia lucidula
G C - G A(A)C - I G G U(C)C G - C C] U - [G G U G] - U						30 Lycopodiella inundata
G C - G A(A)C - I G G U(C)C G - C C] U - [G G U G] - U						31 Lycopodium phlegmaria
G C - G A(A)C - I G G U(C)C G - C C] U - [G G U G] - U						32 Lycopodium taxifolium
G C - G A(A)C - G G U(C)C G - C C] - [G G U G] - U						33 Lycopodium tristachyum
G - - G(A)C U - G G U(C)C G - C C C] - [G G U G] - U						34 Selaginella galleottii
C - - G A(A)C - G G U(C)C G - C C C] - [G G U G] - U						35 Selaginella sp.
(C G G G G)C - G G U(C)C G - C C] - [G G C]A - U						36 Oryza sativa
(C - - - - -)C - G G(G - I)U G - C C] - [G G C A] - -						37 Zea mays
U C - G G C C - G G U(C)C G - C C] - [G G U G] - U						38 Arabidopsis thaliana
U C - G G A U C - G G U(C)C G - C C] - [G G U G] - U						39 Glycine max
U C - G(G)C C - G G U(C)C G - C] - [G U G] - A						40 Sinapis alba
C)C - G G C C - G G U(C)C G - C U] - [G G U G] - U						41 Zamia pumila
G U - G G C C - G G U(C)C G - C C] - [G G U G] - U						42 Gnetum leyboldii
(U)C - G G C C - G G U(C)C G - C C] - [G G U G] - U						43 Ginkgo biloba
A(C) - G G C C - G G U(C)C G - C C] - [G G U G] - U						44 Pinus luchuensis
A(A) - G G U(C) - G G U(C)C G - C C] - [G G U G] - U						45 Pinus wallichiana
(U C - I)U G(U)C - G G U(C)C G - C C] - [G G U G] - U						46 Podocarpus nakaii
A(C) - G G U U - G G U(C)U G - C C] - [G G U G] - U						47 Taxus mairei
G C - G A(G)C(- G)G U(C)C G - C C] - [G G U G] - U						48 Adiantum raddianum
G C - G A(G)C(- G)G U(C)C G - C C] - [G G U G] - U						49 Adiantum raddianum 2
G C - G(G)G C - G G U(C)C G - C C] - [G G G U G] - U						50 Ophioglossum petiolatum
G(C - I G A)G C - G G U(C)C G - C U] - [G G U G] - U						51 Osmunda cinnamomea
G C - G G G C - G G U(C)C G - C C] - [G G U G] - U						52 Salvinia natans
G C - G A(G)C - G G U(C)C G - C C] - [G G U G] - U						53 Psilotum nudum 2
G G G - G U C - G G U(C)C G - C U C] - [G G G C] - U						54 Chara foetida
G(C)G - A C C - G G U(C)C G - C C] - [G G U G] - U						55 Coleochaete scutata
(G)C G - G C C - G G U(C)C G - C C] - [G G U G] - U						56 Klebsormidium flaccidum
A G G - (A)C U - G G U(C)C G - C C] - [G G C] - U						57 Nitella flexilis
A G G - (A)C U - G G U(C)C G - C C] U] - [G G C] - U						58 Nitella sp.
U G G - (U)G C - G G U(C)C G - C C] - [G G U G] - U						59 Chlamydomonas reinhardtii
U(C)U - A G C - G G U(C)C G - C C] - [G G U G] - A						60 Hydrodictyon reticulatum
C G C - G C C - G G U(C)C G - C C G] - [C G U G] - U						61 Parietochloris pseudoalveolaris
C C G - G G U - U A(- U G U C A U U)A C C A C] - [G U G U] - -						62 Dictyostelium discoideum
1 -						63 Helix numbering 1
G C - G(A)G C - G G U(C)C G - C C] - [G G U G] - U						64 Pteris vittata
G(U G - A C)U - G G U(C)C G - C C] - [G G U G] - U						65 Megaceros aenigmaticus
G(U - G A)U] - G G U(C)C G - C C] - [G G G] - U						66 Notophylas breutelii
G G G(A)G - C - G G U(C)C G - C C C] - [G G G U G] - U						67 Physcomitrella patens
G(C - I G G(G)C - G G U(C)C G - C C] - [G G U G] - U						68 Sphagnum palustre

860	870	880	890	900	
A A C U	A A C U	A A C U	A A C U	A A C U	1 Saccharomyces cerevisiae
A A C U	A A C U	A A C U	A A C U	A A C U	2 Schizosaccharomyces pombe
A A C U	A A C U	A A C U	A A C U	A A C U	3 Physcomitrella patens
A A C U	A A C U	A A C U	A A C U	A A C U	4 Anthoceros agrestis
A A C U	A A C U	A A C U	A A C U	A A C U	5 Anthoceros laevis
A A C U	A A C U	A A C U	A A C U	A A C U	6 Pellia epiphylla
A A C U	A A C U	A A C U	A A C U	A A C U	7 Reboulia hemisphaerica 2
A A C U	A A C U	A A C U	A A C U	A A C U	8 Riccardia pinguis
A A C U	A A C U	A A C U	A A C U	A A C U	9 Sphaerocarpos donnellii
A A C U	A A C U	A A C U	A A C U	A A C U	10 Sphaerocarpos texanus
A A C U	A A C U	A A C U	A A C U	A A C U	11 Atrichum undulatum
A A C U	A A C U	A A C U	A A C U	A A C U	12 Bryum argenteum
A A C U	A A C U	A A C U	A A C U	A A C U	13 Eurhynchium hians
A A C U	A A C U	A A C U	A A C U	A A C U	14 Funaria hygrometrica
A A C U	A A C U	A A C U	A A C U	A A C U	15 Leptobryum pyriforme
A A C U	A A C U	A A C U	A A C U	A A C U	16 Mnium hornum
A A C U	A A C U	A A C U	A A C U	A A C U	17 Physcomitrium pyriforme
A A C U	A A C U	A A C U	A A C U	A A C U	18 Polytrichum formosum
A A C U	A A C U	A A C U	A A C U	A A C U	19 Sphagnum cuspidatum
A A C U	A A C U	A A C U	A A C U	A A C U	20 Calypogeia arguta
A A C U	A A C U	A A C U	A A C U	A A C U	21 Conocephalum conicum
A A C U	A A C U	A A C U	A A C U	A A C U	22 Fossombronina pusilla
A A C U	A A C U	A A C U	A A C U	A A C U	23 Marchantia polymorpha
A A C U	A A C U	A A C U	A A C U	A A C U	24 Riccia fluitans
A A C U	A A C U	A A C U	A A C U	A A C U	25 Scapania nemorea
A A C U	A A C U	A A C U	A A C U	A A C U	26 Equisetum hyemale
A A C U	A A C U	A A C U	A A C U	A A C U	27 Equisetum robustum
A A C U	A A C U	A A C U	A A C U	A A C U	28 Isoetes engelmannii
A A C U	A A C U	A A C U	A A C U	A A C U	29 Huperzia lucidula
A A C U	A A C U	A A C U	A A C U	A A C U	30 Lycopodiella inundata
A A C U	A A C U	A A C U	A A C U	A A C U	31 Lycopodium phlegmaria
A A C U	A A C U	A A C U	A A C U	A A C U	32 Lycopodium taxifolium
A A C U	A A C U	A A C U	A A C U	A A C U	33 Lycopodium tristachyum
A A C U	A A C U	A A C U	A A C U	A A C U	34 Selaginella galleottii
A A C U	A A C U	A A C U	A A C U	A A C U	35 Selaginella sp.
A A C U	A A C U	A A C U	A A C U	A A C U	36 Oryza sativa
A A C U	A A C U	A A C U	A A C U	A A C U	37 Zea mays
A A C U	A A C U	A A C U	A A C U	A A C U	38 Arabidopsis thaliana
A A C U	A A C U	A A C U	A A C U	A A C U	39 Glycine max
A A C U	A A C U	A A C U	A A C U	A A C U	40 Sinapis alba
A A C U	A A C U	A A C U	A A C U	A A C U	41 Zamia pumila
A A C U	A A C U	A A C U	A A C U	A A C U	42 Gnetum leyboldii
A A C U	A A C U	A A C U	A A C U	A A C U	43 Ginkgo biloba
A A C U	A A C U	A A C U	A A C U	A A C U	44 Pinus luchuensis
A A C U	A A C U	A A C U	A A C U	A A C U	45 Pinus wallichiana
A A C U	A A C U	A A C U	A A C U	A A C U	46 Podocarpus nakaai
A A C U	A A C U	A A C U	A A C U	A A C U	47 Taxus mairei
A A C U	A A C U	A A C U	A A C U	A A C U	48 Adiantum raddianum
A A C U	A A C U	A A C U	A A C U	A A C U	49 Adiantum raddianum 2
A A C U	A A C U	A A C U	A A C U	A A C U	50 Ophioglossum petiolatum
A A C U	A A C U	A A C U	A A C U	A A C U	51 Osmunda cinnamomea
A A C U	A A C U	A A C U	A A C U	A A C U	52 Salvinia natans
A A C U	A A C U	A A C U	A A C U	A A C U	53 Psilotum nudum 2
A A C U	A A C U	A A C U	A A C U	A A C U	54 Chara foetida
A A C U	A A C U	A A C U	A A C U	A A C U	55 Coleochaete scutata
A A C U	A A C U	A A C U	A A C U	A A C U	56 Klebsormidium flaccidum
A A C U	A A C U	A A C U	A A C U	A A C U	57 Nitella flexilis
A A C U	A A C U	A A C U	A A C U	A A C U	58 Nitella sp.
A A C U	A A C U	A A C U	A A C U	A A C U	59 Chlamydomonas reinhardtii
A A C U	A A C U	A A C U	A A C U	A A C U	60 Hydrodictyon reticulatum
A A C U	A A C U	A A C U	A A C U	A A C U	61 Parietochloris pseudoalveolaris
A A C U	A A C U	A A C U	A A C U	A A C U	62 Dictyostelium discoideum
A A C U	A A C U	A A C U	A A C U	A A C U	63 Helix numbering 1
A A C U	A A C U	A A C U	A A C U	A A C U	64 Pteris vittata
A A C U	A A C U	A A C U	A A C U	A A C U	65 Megaceros aenigmaticus
A A C U	A A C U	A A C U	A A C U	A A C U	66 Notophylas breutelii
A A C U	A A C U	A A C U	A A C U	A A C U	67 Physcomitrella patens
A A C U	A A C U	A A C U	A A C U	A A C U	68 Sphagnum palustre

	910	920	930	940	950																																																	
J	C	G	A	-	A	I	C	-	C	A	G	G	A	I	C	U	U	U	A	C	U	U	-	G	A	A	A	A	U	U	A	-	G	A	G	U	-	G	U	U	C	A	A	A	-	G	I	C	A		1	<i>Saccharomyces cerevisiae</i>		
J	G	G	A	-	A	I	C	-	C	A	G	G	A	I	C	U	U	U	A	C	U	U	-	G	A	A	A	A	U	U	A	-	G	A	G	U	-	G	U	U	C	A	A	A	-	G	I	C	A		2	<i>Schizosaccharomyces pombe</i>		
G	G	A	-	G	I	U	-	C	G	G	C	G	I	A	U	G	U	U	A	C	U	U	-	G	A	A	A	A	U	U	A	-	G	A	G	U	-	G	I	C	U	C	A	A	A	-	G	I	C	A		3	<i>Physcomitrella patens</i>	
G	G	A	-	G	I	U	-	C	G	G	C	G	I	A	U	G	U	U	A	C	U	U	-	G	A	A	A	A	U	U	A	-	G	A	G	U	-	G	I	C	U	C	A	A	A	-	G	I	C	A		4	<i>Anthoceros agrestis</i>	
G	G	A	-	G	I	U	-	C	G	G	C	G	I	A	U	G	U	U	A	C	U	U	-	G	A	A	A	A	U	U	A	-	G	A	G	U	-	G	I	C	U	C	A	A	A	-	G	I	C	A		5	<i>Anthoceros laevis</i>	
G	G	A	-	G	I	U	-	C	G	G	C	G	I	A	U	G	U	U	A	C	U	U	-	G	A	A	A	A	U	U	A	-	G	A	G	U	-	G	I	C	U	C	A	A	A	-	G	I	C	A		6	<i>Pellia epiphylla</i>	
G	G	A	-	G	I	U	-	C	G	G	C	G	I	A	U	G	U	U	A	C	U	U	-	G	A	A	A	A	U	U	A	-	G	A	G	U	-	G	I	C	U	C	A	A	A	-	G	I	C	A		7	<i>Reboulia hemisphaerica 2</i>	
G	G	A	-	G	I	U	-	C	G	G	C	G	I	A	U	G	U	U	A	C	U	U	-	G	A	A	A	A	U	U	A	-	G	A	G	U	-	G	I	C	U	C	A	A	A	-	G	I	C	A		8	<i>Riccardia pinguis</i>	
G	G	A	-	G	I	U	-	C	G	G	C	G	I	A	U	G	U	U	A	C	U	U	-	G	A	A	A	A	U	U	A	-	G	A	G	U	-	G	I	C	U	C	A	A	A	-	G	I	C	A		9	<i>Sphaerocarpos donnellii</i>	
G	G	A	-	G	I	U	-	C	G	G	C	G	I	A	U	G	U	U	A	C	U	U	-	G	A	A	A	A	U	U	A	-	G	A	G	U	-	G	I	C	U	C	A	A	A	-	G	I	C	A		10	<i>Sphaerocarpos texanus</i>	
G	G	A	-	G	I	U	-	C	G	G	C	G	I	A	U	G	U	U	A	C	U	U	-	G	A	A	A	A	U	U	A	-	G	A	G	U	-	G	I	C	U	C	A	A	A	-	G	I	C	A		11	<i>Atrichum undulatum</i>	
G	G	A	-	G	I	U	-	C	G	G	C	G	I	A	U	G	U	U	A	C	U	U	-	G	A	A	A	A	U	U	A	-	G	A	G	U	-	G	I	C	U	C	A	A	A	-	G	I	C	A		12	<i>Bryum argenteum</i>	
G	G	A	-	G	I	U	-	C	G	G	C	G	I	A	U	G	U	U	A	C	U	U	-	G	A	A	A	A	U	U	A	-	G	A	G	U	-	G	I	C	U	C	A	A	A	-	G	I	C	A		13	<i>Eurhynchium hians</i>	
G	G	A	-	G	I	U	-	C	G	G	C	G	I	A	U	G	U	U	A	C	U	U	-	G	A	A	A	A	U	U	A	-	G	A	G	U	-	G	I	C	U	C	A	A	A	-	G	I	C	A		14	<i>Punaria hygrometrica</i>	
G	G	A	-	G	I	U	-	C	G	G	C	G	I	A	U	G	U	U	A	C	U	U	-	G	A	A	A	A	U	U	A	-	G	A	G	U	-	G	I	C	U	C	A	A	A	-	G	I	C	A		15	<i>Leptobryum pyriforme</i>	
G	G	A	-	G	I	U	-	C	G	G	C	G	I	A	U	G	U	U	A	C	U	U	-	G	A	A	A	A	U	U	A	-	G	A	G	U	-	G	I	C	U	C	A	A	A	-	G	I	C	A		16	<i>Mnium hornum</i>	
G	G	A	-	G	I	U	-	C	G	G	C	G	I	A	U	G	U	U	A	C	U	U	-	G	A	A	A	A	U	U	A	-	G	A	G	U	-	G	I	C	U	C	A	A	A	-	G	I	C	A		17	<i>Physcomitrium pyriforme</i>	
G	G	A	-	G	I	U	-	C	G	G	C	G	I	A	U	G	U	U	A	C	U	U	-	G	A	A	A	A	U	U	A	-	G	A	G	U	-	G	I	C	U	C	A	A	A	-	G	I	C	A		18	<i>Polytrichum formosum</i>	
G	G	A	-	G	I	U	-	C	G	G	C	G	I	A	U	G	U	U	A	C	U	U	-	G	A	A	A	A	U	U	A	-	G	A	G	U	-	G	I	C	U	C	A	A	A	-	G	I	C	A		19	<i>Sphagnum cuspidatum</i>	
G	G	A	-	G	I	U	-	C	G	G	C	G	I	A	U	G	U	U	A	C	U	U	-	G	A	A	A	A	U	U	A	-	G	A	G	U	-	G	I	C	U	C	A	A	A	-	G	I	C	A		20	<i>Calypogeia arguta</i>	
G	G	A	-	G	I	U	-	C	G	G	C	G	I	A	U	G	U	U	A	C	U	U	-	G	A	A	A	A	U	U	A	-	G	A	G	U	-	G	I	C	U	C	A	A	A	-	G	I	C	A		21	<i>Conocephalum conicum</i>	
G	G	A	-	G	I	U	-	C	G	G	C	G	I	A	U	G	U	U	A	C	U	U	-	G	A	A	A	A	U	U	A	-	G	A	G	U	-	G	I	C	U	C	A	A	A	-	G	I	C	A		22	<i>Fossombronia pusilla</i>	
G	G	A	-	G	I	U	-	C	G	G	C	G	I	A	U	G	U	U	A	C	U	U	-	G	A	A	A	A	U	U	A	-	G	A	G	U	-	G	I	C	U	C	A	A	A	-	G	I	C	A		23	<i>Marchantia polymorpha</i>	
G	G	A	-	G	I	U	-	C	G	G	C	G	I	A	U	G	U	U	A	C	U	U	-	G	A	A	A	A	U	U	A	-	G	A	G	U	-	G	I	C	U	C	A	A	A	-	G	I	C	A		24	<i>Riccia fluitans</i>	
G	G	A	-	G	I	U	-	C	G	G	C	G	I	A	U	G	U	U	A	C	U	U	-	G	A	A	A	A	U	U	A	-	G	A	G	U	-	G	I	C	U	C	A	A	A	-	G	I	C	A		25	<i>Scapania nemorea</i>	
G	G	A	-	G	I	U	-	C	G	G	C	G	I	A	U	G	U	U	A	C	U	U	-	G	A	A	A	A	U	U	A	-	G	A	G	U	-	G	I	C	U	C	A	A	A	-	G	I	C	A		26	<i>Equisetum hyemale</i>	
G	G	A	-	G	I	U	-	C	G	G	C	G	I	A	U	G	U	U	A	C	U	U	-	G	A	A	A	A	U	U	A	-	G	A	G	U	-	G	I	C	U	C	A	A	A	-	G	I	C	A		27	<i>Equisetum robustum</i>	
J	C	G	A	-	G	I	C	-	C	G	G	C	G	I	A	U	G	U	U	A	C	U	U	-	G	A	A	A	A	U	U	A	-	G	A	G	U	-	G	I	C	U	C	A	A	A	-	G	I	C	A		28	<i>Isoetes engelmannii</i>
G	G	A	-	G	I	U	-	C	G	G	C	G	I	A	U	G	U	U	A	C	U	U	-	G	A	A	A	A	U	U	A	-	G	A	G	U	-	G	I	C	U	C	A	A	A	-	G	I	C	A		29	<i>Huperzia lucidula</i>	
G	G	A	-	G	I	U	-	C	G	G	C	G	I	A	U	G	U	U	A	C	U	U	-	G	A	A	A	A	U	U	A	-	G	A	G	U	-	G	I	C	U	C	A	A	A	-	G	I	C	A		30	<i>Lycopodiella inundata</i>	
G	G	A	-	G	I	U	-	C	G	G	C	G	I	A	U	G	U	U	A	C	U	U	-	G	A	A	A	A	U	U	A	-	G	A	G	U	-	G	I	C	U	C	A	A	A	-	G	I	C	A		31	<i>Lycopodium phlegmaria</i>	
G	G	A	-	G	I	U	-	C	G	G	C	G	I	A	U	G	U	U	A	C	U	U	-	G	A	A	A	A	U	U	A	-	G	A	G	U	-	G	I	C	U	C	A	A	A	-	G	I	C	A		32	<i>Lycopodium taxifolium</i>	
G	G	A	-	G	I	U	-	C	G	G	C	G	I	A	U	G	U	U	A	C	U	U	-	G	A	A	A	A	U	U	A	-	G	A	G	U	-	G	I	C	U	C	A	A	A	-	G	I	C	A		33	<i>Lycopodium tristachyum</i>	
G	G	A	-	G	I	U	-	C	G	G	C	G	I	A	U	G	U	U	A	C	U	U	-	G	A	A	A	A	U	U	A	-	G	A	G	U	-	G	I	C	U	C	A	A	A	-	G	I	C	A		34	<i>Selaginella galleottii</i>	
G	G	A	-	G	I	U	-	C	G	G	C	G	I	A	U	G	U	U	A	C	U	U	-	G	A	A	A	A	U	U	A	-	G	A	G	U	-	G	I	C	U	C	A	A	A	-	G	I	C	A		35	<i>Selaginella sp.</i>	
G	G	A	-	G	I	U	-	C	G	G	C	G	I	A	U	G	U	U	A	C	U	U	-	G	A	A	A	A	U	U	A	-	G	A	G	U	-	G	I	C	U	C	A	A	A	-	G	I	C	A		36	<i>Oryza sativa</i>	
G	G	A	-	G	I	U	-	C	G	G	C	G	I	A	U	G	U	U	A	C	U	U	-	G	A	A	A	A	U	U	A	-	G	A	G	U	-	G	I	C	U	C	A	A	A	-	G	I	C	A		37	<i>Zea mays</i>	
G	G	A	-	G	I	U	-	C	G	G	C	G	I	A	U	G	U	U	A	C	U	U	-	G	A	A	A	A	U	U	A	-	G	A	G	U	-	G	I	C	U	C	A	A	A	-	G	I	C	A		38	<i>Arabidopsis thaliana</i>	
G	G	A	-	G	I	U	-	C	G	G	C	G	I	A	U	G	U	U	A	C	U	U	-	G	A	A	A	A	U	U	A	-	G	A	G	U	-	G	I	C	U	C	A	A	A	-	G	I	C	A		39	<i>Glycine max</i>	
G	G	A	-	G	I	U	-	C	G	G	C	G	I	A	U	G	U	U	A	C	U	U	-	G	A	A	A	A	U	U	A	-	G	A	G	U	-	G	I	C	U	C	A	A	A	-	G	I	C	A		40	<i>Sinapis alba</i>	
G	G	A	-	G	I	U	-	C	G	G	C	G	I	A	U	G	U	U	A	C	U	U	-	G	A	A	A	A	U	U	A	-	G	A	G	U	-	G	I	C	U	C	A	A	A	-	G	I	C	A		41	<i>Zamia pumila</i>	

DCSE Alignment

	960	970	980	990	1000	
I G G C G	- - - U A U	- - - U G C U	- - - A U U	- - - A G - C A	- - - U A A U	1 Saccharomyces cerevisiae
I G G C A A	- - - G U U	- - - U G C U	- - - A U U	- - - A G - C A	- - - U A A U	2 Schizosaccharomyces pombe
A G C	- - - C U A U	- - - G C U(C)U	- - - A U U	- - - A G - C A	- - - U A A C	3 Physcomitrella patens
(U)G C	- - - C U A U	- - - G C U(C)U	- - - A U U	- - - A G - C A	- - - U A A C	4 Anthoceros agrestis
A G C	- - - C U A U	- - - G C U(C)U	- - - A U U	- - - A G - C A	- - - U A A C	5 Anthoceros laevis
A G C	- - - C U A U	- - - G C U(C)U	- - - A U U	- - - A G - C A	- - - U A A C	6 Pellia epiphylla
(A)G C	- - - C A U A C	- - - G C U(C)U	- - - A U U	- - - A G - C A	- - - U A A C	7 Reboulia hemisphaerica 2
A G C	- - - C U A C	- - - G C U(C)U	- - - A U U	- - - A G - C A	- - - U A A C	8 Riccardia pinguis
A G C	- - - C U A C	- - - G C U(C)U	- - - A U U	- - - A G - C A	- - - U A A C	9 Sphaerocarpos donnellii
(A)G C	- - - C U A C	- - - G C U(C)U	- - - A U U	- - - A G - C A	- - - U A A C	10 Sphaerocarpos texanus
A G C	- - - C U A U	- - - G C U(C)U	- - - A U U	- - - A G - C A	- - - U A A C	11 Atrichum undulatum
A G C	- - - C U A U	- - - G C U(C)U	- - - A U U	- - - A G - C A	- - - U A A C	12 Bryum argenteum
A G C	- - - C U A U	- - - G C U(C)U	- - - A U U	- - - A G - C A	- - - U A A C	13 Eurhynchium hians
A G C	- - - C U A U	- - - G C U(C)U	- - - A U U	- - - A G - C A	- - - U A A C	14 Funaria hygrometrica
A G C	- - - C U A U	- - - G C U(C)U	- - - A U U	- - - A G - C A	- - - U A A C	15 Leptobryum pyriforme
A G C	- - - C U A U	- - - G C U(C)U	- - - A U U	- - - A G - C A	- - - U A A C	16 Mnium hornum
A G C	- - - C U A U	- - - G C U(C)U	- - - A U U	- - - A G - C A	- - - U A A C	17 Physcomitrium pyriforme
A G C	- - - C U A U	- - - G C U(C)U	- - - A U U	- - - A G - C A	- - - U A A C	18 Polytrichum formosum
A G C	- - - C U A C	- - - G C U(C)U	- - - A U U	- - - A G - C A	- - - U A A C	19 Sphagnum cuspidatum
A G C	- - - C U A U	- - - G C U(C)U	- - - A U U	- - - A G - C A	- - - U A A C	20 Calypogeia arguta
A G C	- - - C A U A U	- - - G C U(C)U	- - - A U U	- - - A G - C A	- - - U A A C	21 Conocephalum conicum
A G C	- - - C U A U	- - - G C U(C)U	- - - A U U	- - - A G - C A	- - - U A A C	22 Fossombronina pusilla
A G C	- - - C A U A C	- - - G C U(C)U	- - - A U U	- - - A G - C A	- - - U A A C	23 Marchantia polymorpha
A G C	- - - C G U A C	- - - G C U(C)U	- - - A U U	- - - A G - C A	- - - U A A C	24 Riccia fluitans
A G C	- - - C G U A U	- - - G C U(C)U	- - - A U U	- - - A G - C A	- - - U A A C	25 Scapania nemorea
A G C	- - - C U G U	- - - G C U(C)U	- - - A U U	- - - A G - C A	- - - U A A C	26 Equisetum hyemale
A G C	- - - C U A U	- - - G C U(C)U	- - - A U U	- - - A G - C A	- - - U A A C	27 Equisetum robustum
G G C	- - - C U A U	- - - G C U(C)U	- - - A U U	- - - A G - C A	- - - U A A C	28 Isoetes engelmannii
G G C	- - - C U A U	- - - G C U(C)U	- - - A U U	- - - A G - C A	- - - U A A C	29 Huperzia lucidula
G G C	- - - C U A U	- - - G C U(C)U	- - - A U U	- - - A G - C A	- - - U A A C	30 Lycopodiella inundata
G G C	- - - C U A U	- - - G C U(C)U	- - - A U U	- - - A G - C A	- - - U A A C	31 Lycopodium phlegmaria
G G C	- - - C U A U	- - - G C U(C)U	- - - A U U	- - - A G - C A	- - - U A A C	32 Lycopodium taxifolium
A G C	- - - C U A U	- - - G C U(C)U	- - - A U U	- - - A G - C A	- - - U A A C	33 Lycopodium tristachyum
A G C	- - - A A U U	- - - G C U(C)U	- - - A U U	- - - A G - C A	- - - U A A C	34 Selaginella galleottii
G G C	- - - U C C U	- - - G C U(C)U	- - - A U U	- - - A G - C A	- - - U A A C	35 Selaginella sp.
A G C	- - - C A U C	- - - G C U(C)U	- - - A U U	- - - A G - C A	- - - U A A C	36 Oryza sativa
A G C	- - - C A U C	- - - G C U(C)U	- - - A U U	- - - A G - C A	- - - U A A C	37 Zea mays
A G C	- - - C U A C	- - - G C U(C)U	- - - A U U	- - - A G - C A	- - - U A A C	38 Arabidopsis thaliana
A G C	- - - C U A C	- - - G C U(C)U	- - - A U U	- - - A G - C A	- - - U A A C	39 Glycine max
A G C	- - - C U A C	- - - G C U(C)U	- - - A U U	- - - A G - C A	- - - U A A C	40 Sinapis alba
A G C	- - - U U A U	- - - G C U(C)U	- - - A U U	- - - A G - C A	- - - U A A C	41 Zamia pumila
A G C	- - - C U A U	- - - G C U(C)U	- - - A U U	- - - A G - C A	- - - U A A C	42 Gnetum leyboldii
A G C	- - - C U A C	- - - G C U(C)U	- - - A U U	- - - A G - C A	- - - U A A C	43 Ginkgo biloba
A G C	- - - C U A C	- - - G C U(C)U	- - - A U U	- - - A G - C A	- - - U A A C	44 Pinus luchuensis
A G C	- - - U U A U	- - - G C U(C)U	- - - A U U	- - - A G - C A	- - - U A A C	45 Pinus wallichiana
A G C	- - - C U A C	- - - G C U(C)U	- - - A U U	- - - A G - C A	- - - U A A C	46 Podocarpus nakaii
A G C	- - - C U A C	- - - G C U(C)U	- - - A U U	- - - A G - C A	- - - U A A C	47 Taxus mairei
A G C	- - - C U A U	- - - G C U(C)U	- - - A U U	- - - A G - C A	- - - U A A C	48 Adiantum raddianum
A G C	- - - C U A U	- - - G C U(C)U	- - - A U U	- - - A G - C A	- - - U A A C	49 Adiantum raddianum 2
A G C	- - - C U A U	- - - G C U(C)U	- - - A U U	- - - A G - C A	- - - U A A C	50 Ophioglossum petiolatum
A G C	- - - C U A U	- - - G C U(C)U	- - - A U U	- - - A G - C A	- - - U A A C	51 Osmunda cinnamomea
A G C	- - - C U G U	- - - G C U(C)U	- - - A U U	- - - A G - C A	- - - U A A C	52 Salvinia natans
A G C	- - - C U G U	- - - G C U(C)U	- - - A U U	- - - A G - C A	- - - U A A C	53 Psilotum nudum 2
G G C	- - - C U G U	- - - G C U(C)U	- - - A U U	- - - A G - C A	- - - U A A C	54 Chara foetida
G G C	- - - A U A G	- - - G C U(C)U	- - - A U U	- - - A G - C A	- - - U A A C	55 Coleochaete scutata
G G C	- - - C U A C	- - - G C U(C)U	- - - A U U	- - - A G - C A	- - - U A A C	56 Klebsormidium flaccidum
G G C	- - - C C A C	- - - G C U(C)U	- - - A U U	- - - A G - C A	- - - U A A C	57 Nitella flexilis
G G C	- - - C C A C	- - - G C U(C)U	- - - A U U	- - - A G - C A	- - - U A A C	58 Nitella sp.
G G C	- - - C U A C	- - - G C U(C)U	- - - A U U	- - - A G - C A	- - - U A A C	59 Chlamydomonas reinhardtii
G G C	- - - A U A C	- - - G C U(C)U	- - - A U U	- - - A G - C A	- - - U A A C	60 Hydrodictyon reticulatum
G G C	- - - C U A C	- - - G C U(C)U	- - - A U U	- - - A G - C A	- - - U A A C	61 Parietochloris pseudoalveolaris
G G C	- - - U C U	- - - G C U(C)U	- - - A U U	- - - A G - C A	- - - U A A C	62 Dictyostelium discoideum
- - - - -	- - - - -	- - - - -	- - - - -	- - - - -	- - - - -	63 Helix numbering 1
A G C	- - - C U A U	- - - G C U(C)U	- - - A U U	- - - A G - C A	- - - U A A C	64 Pteris vittata
A G C	- - - C U A U	- - - G C U(C)U	- - - A U U	- - - A G - C A	- - - U A A C	65 Megaceros aenigmaticus
G C	- - - C A U A U	- - - G C U(C)U	- - - A U U	- - - A G - C A	- - - U A A C	66 Notophylas breuteli
A G C	- - - C U A U	- - - G C U(C)U	- - - A U U	- - - A G - C A	- - - U A A C	67 Physcomitrella patens
A G C	- - - C U A C	- - - G C U(C)U	- - - A U U	- - - A G - C A	- - - U A A C	68 Sphagnum palustre

DCSE Alignment

	1010	1020	1030	1040	1050	
A U)A G G A I C G U	- U)U G G U U C U	- (A U U U U U	- G U U G I G U U	- U U C U)A G -	G A C - - C A(U)C G U	1 Saccharomyces cerevisiae
A U)A G G A I C G U	- U)U G G U U C U	- (A U U U U U	- G U U G I G U U	- U U C U)A G -	G A C - - C G C C G U	2 Schizosaccharomyces pombe
A U)A G G A I C U C	- U)U G G U U C U	- (A U U U U U	- G U U G I G U U	- U U C U)A G -	G A C - - C G G A G U	3 Physcomitrella patens
A U)A G G A I C U C	- U)U G G U U C U	- (A U U U U U	- G U U G I G U U	- U U C U)A G -	G A C - - U G G A G U	4 Anthoceros agrestis
A U)A G G A I C U C	- U)U G G U U C U	- (A U U U U U	- G U U G I G U U	- U U C U)A G -	G A C - - C G G A G U	5 Anthoceros laevis
A U)A G G A I C U C	- U)U G G U U C U	- (A U U U U U	- G U U G I G U U	- U U C U)A G -	G A C - - C G G A G U	6 Pellia epiphylla
A U)A G G A I C U C	- U)U G G U U C U	- (A U U U U U	- G U U G I G U U	- U U C U)A G -	G A C - - C G G A G U	7 Reboulia hemisphaerica 2
A U)A G G A I C U C	- U)U G G U U C U	- (A U U U U U	- G U U G I G U U	- U U C U)A G -	G A C - - C G G A G U	8 Riccardia pinguis
A U)A G G A I C U C	- U)U G G U U C U	- (A U U U U U	- G U U G I G U U	- U U C U)A G -	G A C - - C G G A G U	9 Sphaerocarpos donnelli
A U)A G G A I C U C	- U)U G G U U C U	- (A U U U U U	- G U U G I G U U	- U U C U)A G -	G A C - - C G G A G U	10 Sphaerocarpos texanus
A U)A G G A I C U C	- U)U G G U U C U	- (A U U U U U	- G U U G I G U U	- U U C U)A G -	G A C - - C G G A G U	11 Atrichum undulatum
A U)A G G A I C U C	- U)U G G U U C U	- (A U U U U U	- G U U G I G U U	- U U C U)A G -	G A C - - C G G A G U	12 Bryum argenteum
A U)A G G A I C U C	- U)U G G U U C U	- (A U U U U U	- G U U G I G U U	- U U C U)A G -	G A C - - C G G A G U	13 Eurhynchium nians
A U)A G G A I C U C	- U)U G G U U C U	- (A U U U U U	- G U U G I G U U	- U U C U)A G -	G A C - - C G G A G U	14 Funaria hygrometrica
A U)A G G A I C U C	- U)U G G U U C U	- (A U U U U U	- G U U G I G U U	- U U C U)A G -	G A C - - C G G A G U	15 Leptobryum pyriforme
A U)A G G A I C U C	- U)U G G U U C U	- (A U U U U U	- G U U G I G U U	- U U C U)A G -	G A C - - C G G A G U	16 Mnium hornum
A U)A G G A I C U C	- U)U G G U U C U	- (A U U U U U	- G U U G I G U U	- U U C U)A G -	G A C - - C G G A G U	17 Physcomitrium pyriforme
A U)A G G A I C U C	- U)U G G U U C U	- (A U U U U U	- G U U G I G U U	- U U C U)A G -	G A C - - C G G A G U	18 Polytrichum formosum
A U)A G G A I C U C	- U)U G G U U C U	- (A U U U U U	- G U U G I G U U	- U U C U)A G -	G A C - - C G G A G U	19 Sphagnum cuspidatum
A U)A G G A I C U C	- U)U G G U U C U	- (A U U U U U	- G U U G I G U U	- U U C U)A G -	G A C - - C G G A G U	20 Calypogeia arguta
A U)A G G A I C U C	- U)U G G U U C U	- (A U U U U U	- G U U G I G U U	- U U C U)A G -	G A C - - C G G A G U	21 Conocephalum concum
A U)A G G A I C U C	- U)U G G U U C U	- (A U U U U U	- G U U G I G U U	- U U C U)A G -	G A C - - C G G A G U	22 Fossombronina pustilla
A U)A G G A I C U C	- U)U G G U U C U	- (A U U U U U	- G U U G I G U U	- U U C U)A G -	G A C - - C G G A G U	23 Marchantia polymorpha
A U)A G G A I C U C	- U)U G G U U C U	- (A U U U U U	- G U U G I G U U	- U U C U)A G -	G A C - - C G G A G U	24 Riccia fluitans
A U)A G G A I C U C	- U)U G G U U C U	- (A U U U U U	- G U U G I G U U	- U U C U)A G -	G A C - - C G G A G U	25 Scapania nemorea
A U)A G G A I C U C	- U)U G G U U C U	- (A U U U U U	- G U U G I G U U	- U U C U)A G -	G A C - - C G G A G U	26 Equisetum hyemale
A U)A G G A I C U C	- U)U G G U U C U	- (A U U U U U	- G U U G I G U U	- U U C U)A G -	G A C - - C G G A G U	27 Equisetum robustum
A U)A G G A I C U C	- U)U G G U U C U	- (A U U U U U	- G U U G I G U U	- U U C U)A G -	G A C - - C G G A G U	28 Isoetes engelmannii
A U)A G G A I C U C	- U)U G G U U C U	- (A U U U U U	- G U U G I G U U	- U U C U)A G -	G A C - - C G G A G U	29 Huperzia lucidula
A U)A G G A I C U C	- U)U G G U U C U	- (A U U U U U	- G U U G I G U U	- U U C U)A G -	G A C - - C G G A G U	30 Lycopodiella inundata
A U)A G G A I C U C	- U)U G G U U C U	- (A U U U U U	- G U U G I G U U	- U U C U)A G -	G A C - - C G G A G U	31 Lycopodium phlegmaria
A U)A G G A I C U C	- U)U G G U U C U	- (A U U U U U	- G U U G I G U U	- U U C U)A G -	G A C - - C G G A G U	32 Lycopodium taxifolium
A U)A G G A I C U C	- U)U G G U U C U	- (A U U U U U	- G U U G I G U U	- U U C U)A G -	G A C - - C G G A G U	33 Lycopodium tristachyum
A U)A G G A I C U C	- U)U G G U U C U	- (A U U U U U	- G U U G I G U U	- U U C U)A G -	G A C - - C G G A G U	34 Selaginella galleottii
A U)A G G A I C U C	- U)U G G U U C U	- (A U U U U U	- G U U G I G U U	- U U C U)A G -	G A C - - C G G A G U	35 Selaginella sp.
A U)A G G A I C U C	- U)U G G U U C U	- (A U U U U U	- G U U G I G U U	- U U C U)A G -	G A C - - C G G A G U	36 Oryza sativa
A U)A G G A I C U C	- U)U G G U U C U	- (A U U U U U	- G U U G I G U U	- U U C U)A G -	G A C - - C G G A G U	37 Zea mays
A U)A G G A I C U C	- U)U G G U U C U	- (A U U U U U	- G U U G I G U U	- U U C U)A G -	G A C - - C G G A G U	38 Arabidopsis thaliana
A U)A G G A I C U C	- U)U G G U U C U	- (A U U U U U	- G U U G I G U U	- U U C U)A G -	G A C - - C G G A G U	39 Glycine max
A U)A G G A I C U C	- U)U G G U U C U	- (A U U U U U	- G U U G I G U U	- U U C U)A G -	G A C - - C G G A G U	40 Sinapis alba
A U)A G G A I C U C	- U)U G G U U C U	- (A U U U U U	- G U U G I G U U	- U U C U)A G -	G A C - - C G G A G U	41 Zamia pumila
A U)A G G A I C U C	- U)U G G U U C U	- (A U U U U U	- G U U G I G U U	- U U C U)A G -	G A C - - C G G A G U	42 Gnetum leyboldii
A U)A G G A I C U C	- U)U G G U U C U	- (A U U U U U	- G U U G I G U U	- U U C U)A G -	G A C - - C G G A G U	43 Ginkgo biloba
A U)A G G A I C U C	- U)U G G U U C U	- (A U U U U U	- G U U G I G U U	- U U C U)A G -	G A C - - C G G A G U	44 Pinus luchuensis
A U)A G G A I C U C	- U)U G G U U C U	- (A U U U U U	- G U U G I G U U	- U U C U)A G -	G A C - - C G G A G U	45 Pinus walllichiana
A U)A G G A I C U C	- U)U G G U U C U	- (A U U U U U	- G U U G I G U U	- U U C U)A G -	G A C - - C G G A G U	46 Podocarpus nakaii
A U)A G G A I C U C	- U)U G G U U C U	- (A U U U U U	- G U U G I G U U	- U U C U)A G -	G A C - - C G G A G U	47 Taxus mairei
A U)A G G A I C U C	- U)U G G U U C U	- (A U U U U U	- G U U G I G U U	- U U C U)A G -	G A C - - C G G A G U	48 Adiantum raddianum
A U)A G G A I C U C	- U)U G G U U C U	- (A U U U U U	- G U U G I G U U	- U U C U)A G -	G A C - - C G G A G U	49 Adiantum raddianum 2
A U)A G G A I C U C	- U)U G G U U C U	- (A U U U U U	- G U U G I G U U	- U U C U)A G -	G A C - - C G G A G U	50 Ophioglossum petiolatum
A U)A G G A I C U C	- U)U G G U U C U	- (A U U U U U	- G U U G I G U U	- U U C U)A G -	G A C - - C G G A G U	51 Osmunda cinnamomea
A U)A G G A I C U C	- U)U G G U U C U	- (A U U U U U	- G U U G I G U U	- U U C U)A G -	G A C - - C G G A G U	52 Salvinia natans
A U)A G G A I C U C	- U)U G G U U C U	- (A U U U U U	- G U U G I G U U	- U U C U)A G -	G A C - - C G G A G U	53 Psilotum nudum 2
A U)A G G A I C U C	- U)U G G U U C U	- (A U U U U U	- G U U G I G U U	- U U C U)A G -	G A C - - C G G A G U	54 Chara foetida
A U)A G G A I C U C	- U)U G G U U C U	- (A U U U U U	- G U U G I G U U	- U U C U)A G -	G A C - - C G G A G U	55 Coleochaete scutata
A U)A G G A I C U C	- U)U G G U U C U	- (A U U U U U	- G U U G I G U U	- U U C U)A G -	G A C - - C G G A G U	56 Klebsormidium flaccidum
A U)A G G A I C U C	- U)U G G U U C U	- (A U U U U U	- G U U G I G U U	- U U C U)A G -	G A C - - C G G A G U	57 Nitella flexilis
A U)A G G A I C U C	- U)U G G U U C U	- (A U U U U U	- G U U G I G U U	- U U C U)A G -	G A C - - C G G A G U	58 Nitella sp.
A U)A G G A I C U C	- U)U G G U U C U	- (A U U U U U	- G U U G I G U U	- U U C U)A G -	G A C - - C G G A G U	59 Chlamydomonas reinhardtii
A U)A G G A I C U C	- U)U G G U U C U	- (A U U U U U	- G U U G I G U U	- U U C U)A G -	G A C - - C G G A G U	60 Hydrodictyon reticulatum
A U)A G G A I C U C	- U)U G G U U C U	- (A U U U U U	- G U U G I G U U	- U U C U)A G -	G A C - - C G G A G U	61 Parietochloris pseudovalveolaris
A U)A G G A I C U C	- U)U G G U U C U	- (A U U U U U	- G U U G I G U U	- U U C U)A G -	G A C - - C G G A G U	62 Dictyostelium discoideum
A U)A G G A I C U C	- U)U G G U U C U	- (A U U U U U	- G U U G I G U U	- U U C U)A G -	G A C - - C G G A G U	63 Helix numbering 1
A U)A G G A I C U C	- U)U G G U U C U	- (A U U U U U	- G U U G I G U U	- U U C U)A G -	G A C - - C G G A G U	64 Pteris vittata
A U)A G G A I C U C	- U)U G G U U C U	- (A U U U U U	- G U U G I G U U	- U U C U)A G -	G A C - - C G G A G U	65 Megaceros aenigmaticus
A U)A G G A I C U C	- U)U G G U U C U	- (A U U U U U	- G U U G I G U U	- U U C U)A G -	G A C - - C G G A G U	66 Notophylas breutelii
A U)A G G A I C U C	- U)U G G U U C U	- (A U U U U U	- G U U G I G U U	- U U C U)A G -	G A C - - C G G A G U	67 Physcomitrella patens
A U)A G G A I C U C	- U)U G G U U C U	- (A U U U U U	- G U U G I G U U	- U U C U)A G -	G A C - - C G G A G U	68 Sphagnum palustre

	1060	1070	1080	1090	1100		
A			A U G A(U	U A A(U	A)G G	G A C G I G U - C G G	1 Saccharomyces cerevisiae
A			A U G A(U	U A A(U	A)G G	G A U A I G U - C G G	2 Schizosaccharomyces pombe
A			A U G A(U	U A A(U	A)G G	G A C G I G U - U G G	3 Physcomitrella patens
A			A U G A(U	U A A(U	A)G G	G A C A I G U - U G G	4 Anthoceros agrestis
A			A U G A(U	U A A(U	A)G G	G A C A I G U - U G G	5 Anthoceros laevis
A			A U G A(U	U A A(U	A)G G	G A C A I G U - U G G	6 Pellia epiphylla
A			A U G A(U	U A A(U	A)G G	G A C A I G U - U G G	7 Reboulia hemisphaerica 2
A			A U G A(U	U A A(U	A)G G	G A C A I G U - U G G	8 Riccardia pinguis
A			A U G A(U	U A A(U	A)G G	G A C A I G U - U G G	9 Sphaerocarpos donnelli
A			A U G A(U	U A A(U	A)G G	G A C A I G U - U G G	10 Sphaerocarpos texanus
A			A U G A(U	U A A(U	A)G G	G A C A I G U - U G G	11 Atrichum undulatum
A			A U G A(U	U A A(U	A)G G	G A C A I G U - U G G	12 Bryum argenteum
A			A U G A(U	U A A(U	A)G G	G A U A I G U - U G G	13 Eurhynchium hians
A			A U G A(U	U A A(U	A)G G	G A C G I G U - U G G	14 Funaria hygrometrica
A			A U G A(U	U A A(U	A)G G	G A C A I G U - U G G	15 Leptobryum pyriforme
A			A U G A(U	U A A(U	A)G G	G A C A I G U - U G G	16 Mnium hornum
A			A U G A(U	U A A(U	A)G G	G A C G I G U - U G G	17 Physcomitrium pyriforme
A			A U G A(U	U A A(U	A)G G	G A C G I G U - U G G	18 Polytrichum formosum
A			A U G A(U	U A A(U	A)G G	G A C G I G U - U G G	19 Sphagnum cuspidatum
A			A U G A(U	U A A(U	A)G G	G A C A I G U - U G G	20 Calypogeia arguta
A			A U G A(U	U A A(U	A)G G	G A C A I G U - U G G	21 Conocephalum concium
A			A U G A(U	U A A(U	A)G G	G A C A I G U - U G G	22 Fossombronia pusilla
A			A U G A(U	U A A(U	A)G G	G A C A I G U - U G G	23 Marchantia polymorpha
A			A U G A(U	U A A(U	A)G G	G A C A I G U - U G G	24 Riccia fluitans
A			A U G A(U	U A A(U	A)G G	G A C A I G U - U G G	25 Scapania nemorea
A			A U G A(U	U A A(U	A)G G	G A C A I G U - U G G	26 Equisetum hyemale
A			A U G A(U	U A A(U	A)G G	G A C A I G U - U G G	27 Equisetum robustum
A			A U G A(U	U A A(U	A)G G	(C)A C A I G U - U G G	28 Isoetes engelmannii
A			A U G A(U	U A A(U	A)G G	G A C A I G U - U G G	29 Huperzia lucidula
A			A U G A(U	U A A(U	A)G G	G A C A I G U - U G G	30 Lycopodiella inundata
A			A U G A(U	U A A(U	A)G G	G A C A I G U - U G G	31 Lycopodium phlegmaria
A			A U G A(U	U A A(U	A)G G	G A C A I G U - U G G	32 Lycopodium taxifolium
A			A U G A(U	U A A(U	A)G G	G A C G I G U - U G G	33 Lycopodium tristachyum
A			A U G A(U	U A A(U	A)G G	G A C A I G U - U G G	34 Selaginella galleottii
A			A U G A(U	U A A(U	A)G G	G A C A I G U - U G G	35 Selaginella sp.
A			A U G A(U	U A A(U	A)G G	G A C A I G U - C G G	36 Oryza sativa
A			A U G A(U	U A A(U	A)G G	G A C A I G U - C G G	37 Zea mays
A			A U G A(U	U A A(U	A)G G	G A C A I G U - C G G	38 Arabidopsis thaliana
A			A U G A(U	U A A(U	A)G G	G A C A I G U - C G G	39 Glycine max
A			A U G A(U	U A A(U	A)G G	G A C A I G U - C G G	40 Sinapis alba
A			A U G A(U	U A A(U	A)G G	G A C G I G U - C G G	41 Zamia pumila
A			A U G A(U	U A A(U	A)G G	G A C I U G U - C G G	42 Gnetum leyboldii
A			A U G A(U	U A A(U	A)G G	G A C G I G U - C G G	43 Ginkgo biloba
A			A U G A(U	U A A(U	A)G G	G A C I U G U - C G G	44 Pinus luchuensis
A			A U G A(U	U A A(U	A)G G	G A C I U G U - C G G	45 Pinus wallichiana
A			A U G A(U	U A A(U	A)G G	G A C I U G U - C G G	46 Podocarpus nakaii
A			A U G A(U	U A A(U	A)G G	G A C I U G U - C G G	47 Taxus mairei
A			A U G A(U	U A A(U	A)G G	G A C A I G U - U G G	48 Adiantum raddianum
A			A U G A(U	U A A(U	A)G G	G A C A I G U - U G G	49 Adiantum raddianum 2
A			A U G A(U	U A A(U	A)G G	G A C A I G U - U G G	50 Ophioglossum petiolatum
A			A U G A(U	U A A(U	A)G G	G A C A I G U - U G G	51 Osmunda cinnamomea
A			A U G A(U	U A A(U	A)G G	G A C A I G U - U G G	52 Salvinia natans
A			A U G A(U	U A A(U	A)G G	G A C A I G U - U G G	53 Psilotum nudum 2
A			A U G A(U	U A A(U	A)G G	G A C G I G U - U G G	54 Chara foetida
A			A U G A(U	U A A(U	A)G G	G A C A I G U - U G G	55 Coleochaete scutata
A			A U G A(U	U A A(U	A)G G	G A C A I G U - U G G	56 Klebsormidium flaccidum
A			A U G A(U	U A A(U	A)G G	G A C A I G U - U G G	57 Nitella flexilis
A			A U G A(U	U A A(U	A)G G	(C)A A U A I G G	58 Nitella sp.
A			A U G A(U	U A A(U	A)G G	G U A I G U - C G G	59 Chlamydomonas reinhardtii
A			A U G A(U	U A A(U	A)G G	G A C A I G U - C G G	60 Hydrodictyon reticulatum
A			A U G A(U	U A A(U	A)G G	G A C A I G U - C G G	61 Parietochloris pseudoalveolaris
A			A U G A(U	U A A(U	A)G G	G A C A I G U - U G G	62 Dictyostelium discoideum
A			A U G A(U	U A A(U	A)G G	G A C A I G U - U G G	63 Helix numbering 1
A			A U G A(U	U A A(U	A)G G	G A C G I G U - U G G	64 Pteris vittata
A			A U G A(U	U A A(U	A)G G	G A C A I G U - U G G	65 Megaceros aenigmaticus
A			A U G A(U	U A A(U	A)G G	G A C A I G U - U G G	66 Notophylas breutelu
A			A U G A(U	U A A(U	A)G G	G A C G I G U - U G G	67 Physcomitrella patens
A			A U G A(U	U A A(U	A)G G	G A C G I G U - U G G	68 Sphagnum palustre

	1110	1120	1130	1140	1150	
(C)G	-	-	-	-	-	1 Saccharomyces cerevisiae
G	-	-	-	-	-	2 Schizosaccharomyces pombe
U(A)U	-	-	-	-	-	3 Physcomitrella patens
U	-	-	-	-	-	4 Anthoceros agrestis
C	-	-	-	-	-	5 Anthoceros laevis
A	-	-	-	-	-	6 Pellia epiphylla
-	-	-	-	-	-	7 Reboulia hemisphaerica 2
-	-	-	-	-	-	8 Riccardia pinguis
-	-	-	-	-	-	9 Sphaerocarpos donnelli
-	-	-	-	-	-	10 Sphaerocarpos texanus
-	-	-	-	-	-	11 Atrichum undulatum
-	-	-	-	-	-	12 Bryum argenteum
-	-	-	-	-	-	13 Euzhynchium hiens
-	-	-	-	-	-	14 Funaria hygrometrica
-	-	-	-	-	-	15 Leptobryum pyriforme
-	-	-	-	-	-	16 Mnium hornum
-	-	-	-	-	-	17 Physcomitrium pyriforme
-	-	-	-	-	-	18 Polytrichum formosum
-	-	-	-	-	-	19 Sphagnum cuspidatum
-	-	-	-	-	-	20 Calypogeia arguta
-	-	-	-	-	-	21 Conocephalum conicum
-	-	-	-	-	-	22 Fossombronina pusilla
-	-	-	-	-	-	23 Marchantia polymorpha
-	-	-	-	-	-	24 Riccia fluitans
-	-	-	-	-	-	25 Scapania nemorea
-	-	-	-	-	-	26 Equisetum hyemale
-	-	-	-	-	-	27 Equisetum robustum
-	-	-	-	-	-	28 Isoetes engelmannii
-	-	-	-	-	-	29 Huperzia lucidula
-	-	-	-	-	-	30 Lycopodiella inundata
-	-	-	-	-	-	31 Lycopodium phlegmaria
-	-	-	-	-	-	32 Lycopodium taxifolium
-	-	-	-	-	-	33 Lycopodium tristachyum
-	-	-	-	-	-	34 Selaginella galleottii
-	-	-	-	-	-	35 Selaginella sp.
-	-	-	-	-	-	36 Oryza sativa
-	-	-	-	-	-	37 Zea mays
-	-	-	-	-	-	38 Arabidopsis thaliana
-	-	-	-	-	-	39 Glycine max
-	-	-	-	-	-	40 Sinapis alba
-	-	-	-	-	-	41 Zamia pumila
-	-	-	-	-	-	42 Gnetum leyboldii
-	-	-	-	-	-	43 Ginkgo biloba
-	-	-	-	-	-	44 Pinus luchuensis
-	-	-	-	-	-	45 Pinus wallichiana
-	-	-	-	-	-	46 Podocarpus nakaii
-	-	-	-	-	-	47 Taxus mairei
-	-	-	-	-	-	48 Adiantum raddianum
-	-	-	-	-	-	49 Adiantum raddianum 2
-	-	-	-	-	-	50 Ophioglossum petiolatum
-	-	-	-	-	-	51 Osmunda cinnamomea
-	-	-	-	-	-	52 Salvinia natans
-	-	-	-	-	-	53 Psilotum nudum 2
-	-	-	-	-	-	54 Chara foetida
-	-	-	-	-	-	55 Coleochaete scutata
-	-	-	-	-	-	56 Klebsormidium flaccidum
-	-	-	-	-	-	57 Nitella flexilis
-	-	-	-	-	-	58 Nitella sp.
-	-	-	-	-	-	59 Chlamydomonas reinhardtii
-	-	-	-	-	-	60 Hydrodictyon reticulatum
-	-	-	-	-	-	61 Parietochloris pseudoalveolaris
-	-	-	-	-	-	62 Dictyostelium discoideum
-	-	-	-	-	-	63 Helix numbering 1
-	-	-	-	-	-	64 Pteris vittata
-	-	-	-	-	-	65 Megaceros aenigmaticus
-	-	-	-	-	-	66 Notophylas breuteli
-	-	-	-	-	-	67 Physcomitrella patens
-	-	-	-	-	-	68 Sphagnum palustre

	1160	1170	1180	1190	1200	
C U A C U G C G A A A G C A U U U G C C A A G G A - U G U U U U(C A)U U A A^U C A A G A A C G G A A A						1 Saccharomyces cerevisiae
C U A C U G C G A A A G C A U U U G C C A A G G A - U G U U U U(C A)U U A A^U C A A G A A C G G A A A						2 Schizosaccharomyces pombe
C U U C U G C G A A A G C A U U U G C C A A G G A - U G U U U U(C A)U U A A^U C A A G A A C G G A A A						3 Physcomitrella patens
C U U C U G C G A A A G C A U U U G C C A A G G A - U G U U U U(C A)U U A A^U C A A G A A C G G A A A						4 Anthoceros agrestis
C U U C U G C G A A A G C A U U U G C C A A G G A - U G U U U U(C A)U U A A^U C A A G A A C G G A A A						5 Anthoceros laevis
C U U C U G C G A A A G C A U U U G C C A A G G A - U G U U U U(C A)U U A A^U C A A G A A C G G A A A						6 Pellia epiphylla
C U U C U G C G A A A G C A U U U G C C A A G G A - U G U U U U(C A)U U A A^U C A A G A A C G G A A A						7 Reboulia hemisphaerica 2
C U U C U G C G A A A G C A U U U G C C A A G G A - U G U U U U(C A)U U A A^U C A A G A A C G G A A A						8 Riccardia pinquius
C U U C U G C G A A A G C A U U U G C C A A G G A - U G U U U U(C A)U U A A^U C A A G A A C G G A A A						9 Sphaerocarpos donnelli
C U U C U G C G A A A G C A U U U G C C A A G G A - U G U U U U(C A)U U A A^U C A A G A A C G G A A A						10 Sphaerocarpos texanus
C U U C U G C G A A A G C A U U U G C C A A G G A - U G U U U U(C A)U U A A^U C A A G A A C G G A A A						11 Atrichum undulatum
C U U C U G C G A A A G C A U U U G C C A A G G A - U G U U U U(C A)U U A A^U C A A G A A C G G A A A						12 Bryum argenteum
C U U C U G C G A A A G C A U U U G C C A A G G A - U G U U U U(C A)U U A A^U C A A G A A C G G A A A						13 Eurhynchium hians
C U U C U G C G A A A G C A U U U G C C A A G G A - U G U U U U(C A)U U A A^U C A A G A A C G G A A A						14 Funaria hygrometrica
C U U C U G C G A A A G C A U U U G C C A A G G A - U G U U U U(C A)U U A A^U C A A G A A C G G A A A						15 Leptobryum pyriforme
C U U C U G C G A A A G C A U U U G C C A A G G A - U G U U U U(C A)U U A A^U C A A G A A C G G A A A						16 Mnium hornum
C U U C U G C G A A A G C A U U U G C C A A G G A - U G U U U U(C A)U U A A^U C A A G A A C G G A A A						17 Physcomitrium pyriforme
C U U C U G C G A A A G C A U U U G C C A A G G A - U G U U U U(C A)U U A A^U C A A G A A C G G A A A						18 Polytrichum formosum
C U U C U G C G A A A G C A U U U G C C A A G G A - U G U U U U(C A)U U A A^U C A A G A A C G G A A A						19 Sphagnum cuspidatum
C U U C U G C G A A A G C A U U U G C C A A G G A - U G U U U U(C A)U U A A^U C A A G A A C G G A A A						20 Calypogeia arguta
C U U C U G C G A A A G C A U U U G C C A A G G A - U G U U U U(C A)U U A A^U C A A G A A C G G A A A						21 Conocephalum concicum
C U U C U G C G A A A G C A U U U G C C A A G G A - U G U U U U(C A)U U A A^U C A A G A A C G G A A A						22 Fossombronia pusilla
C U U C U G C G A A A G C A U U U G C C A A G G A - U G U U U U(C A)U U A A^U C A A G A A C G G A A A						23 Marchantia polymorpha
C U U C U G C G A A A G C A U U U G C C A A G G A - U G U U U U(C A)U U A A^U C A A G A A C G G A A A						24 Riccia fluitans
C U U C U G C G A A A G C A U U U G C C A A G G A - U G U U U U(C A)U U A A^U C A A G A A C G G A A A						25 Scapania nemorea
C U A C U G C G A A A G C A U U U G C C A A G G A - U G U U U U(C A)U U A A^U C A A G A A C G G A A A						26 Equisetum hyemale
C U A C U G C G A A A G C A U U U G C C A A G G A - U G U U U U(C A)U U A A^U C A A G A A C G G A A A						27 Equisetum robustum
C U A C U G C G A A A G C A U U U G C C A A G G A - U G U U U U(C A)U U A A^U C A A G A A C G G A A A						28 Isoetes engelmannii
C U A C U G C G A A A G C A U U U G C C A A G G A - U G U U U U(C A)U U A A^U C A A G A A C G G A A A						29 Huperzia lucidula
C U A C U G C G A A A G C A U U U G C C A A G G A - U G U U U U(C A)U U A A^U C A A G A A C G G A A A						30 Lycopodiella inundata
C U A C U G C G A A A G C A U U U G C C A A G G A - U G U U U U(C A)U U A A^U C A A G A A C G G A A A						31 Lycopodium phlegmaria
C U A C U G C G A A A G C A U U U G C C A A G G A - U G U U U U(C A)U U A A^U C A A G A A C G G A A A						32 Lycopodium taxifolium
C U A C U G C G A A A G C A U U U G C C A A G G A - U G U U U U(C A)U U A A^U C A A G A A C G G A A A						33 Lycopodium tristachyum
C U A C U G C G A A A G C A U U U G C C A A G G A - U G U U U U(C A)U U A A^U C A A G A A C G G A A A						34 Selaginella galileottii
C U A C U G C G A A A G C A U U U G C C A A G G A - U G U U U U(C A)U U A A^U C A A G A A C G G A A A						35 Selaginella sp.
C A A C U G C G A A A G C A U U U G C C A A G G A - U G U U U U(C A)U U A A^U C A A G A A C G G A A A						36 Oryza sativa
C A A C U G C G A A A G C A U U U G C C A A G G A - U G U U U U(C A)U U A A^U C A A G A A C G G A A A						37 Zea mays
C A A C U G C G A A A G C A U U U G C C A A G G A - U G U U U U(C A)U U A A^U C A A G A A C G G A A A						38 Arabidopsis thaliana
C A A C U G C G A A A G C A U U U G C C A A G G A - U G U U U U(C A)U U A A^U C A A G A A C G G A A A						39 Glycine max
C A A C U G C G A A A G C A U U U G C C A A G G A - U G U U U U(C A)U U A A^U C A A G A A C G G A A A						40 Sinapis alba
C C A C U G C G A A A G C A U U U G C C A A G G A - U G U U U U(C A)U U A A^U C A A G A A C G G A A A						41 Zamia pumila
C C A C U G C G A A A G C A U U U G C C A A G G A - U G U U U U(C A)U U A A^U C A A G A A C G G A A A						42 Gnetum leyboldii
C C A C U G C G A A A G C A U U U G C C A A G G A - U G U U U U(C A)U U A A^U C A A G A A C G G A A A						43 Ginkgo biloba
C C A C U G C G A A A G C A U U U G C C A A G G A - U G U U U U(C A)U U A A^U C A A G A A C G G A A A						44 Pinus luchuensis
C C A C U G C G A A A G C A U U U G C C A A G G A - U G U U U U(C A)U U A A^U C A A G A A C G G A A A						45 Pinus wallichiana
C C A C U G C G A A A G C A U U U G C C A A G G A - U G U U U U(C A)U U A A^U C A A G A A C G G A A A						46 Podocarpus nakaii
C C A C U G C G A A A G C A U U U G C C A A G G A - U G U U U U(C A)U U A A^U C A A G A A C G G A A A						47 Taxus mairei
C U A C U G C G A A A G C A U U U G C C A A G G A - U G U U U U(C A)U U A A^U C A A G A A C G G A A A						48 Adiantum raddianum
C U A C U G C G A A A G C A U U U G C C A A G G A - U G U U U U(C A)U U A A^U C A A G A A C G G A A A						49 Adiantum raddianum 2
C U A C U G C G A A A G C A U U U G C C A A G G A - U G U U U U(C A)U U A A^U C A A G A A C G G A A A						50 Ophioglossum petiolatum
C U A C U G C G A A A G C A U U U G C C A A G G A - U G U U U U(C A)U U A A^U C A A G A A C G G A A A						51 Osmunda cinnamomea
C U A C U G C G A A A G C A U U U G C C A A G G A - U G U U U U(C A)U U A A^U C A A G A A C G G A A A						52 Salvinia natans
C U U C U G C G A A A G C A U U U G C C A A G G A - U G U U U U(C A)U U A A^U C A A G A A C G G A A A						53 Psilotum nudum 2
C U U C U G C G A A A G C A U U U G C C A A G G A - U G U U U U(C A)U U A A^U C A A G A A C G G A A A						54 Chara foetida
C U U C U G C G A A A G C A U U U G C C A A G G A - U G U U U U(C A)U U A A^U C A A G A A C G G A A A						55 Coleochaete scutata
C U U C U G C G A A A G C A U U U G C C A A G G A - U G U U U U(C A)U U A A^U C A A G A A C G G A A A						56 Klebsormidium flaccidum
C U U C U G C G A A A G C A U U U G C C A A G G A - U G U U U U(C A)U U A A^U C A A G A A C G G A A A						57 Nitella flexilis
C U U C U G C G A A A G C A U U U G C C A A G G A - U G U U U U(C A)U U A A^U C A A G A A C G G A A A						58 Nitella sp.
C U A C U G C G A A A G C A U U U G C C A A G G A - U G U U U U(C A)U U A A^U C A A G A A C G G A A A						59 Chlamydomonas reinhardtii
C U A C U G C G A A A G C A U U U G C C A A G G A - U G U U U U(C A)U U A A^U C A A G A A C G G A A A						60 Hydrodictyon reticulatum
C U A C U G C G A A A G C A U U U G C C A A G G A - U G U U U U(C A)U U A A^U C A A G A A C G G A A A						61 Parietochloris pseudoalveolaris
C U U C U G C G A A A G C A U U U G C C A A G G A - U G U U U U(C A)U U A A^U C A A G A A C G G A A A						62 Dictyostelium discoideum
- - - - - 26' - - - - - 26' - - - - - 24' - - - - - 23' - - - - -						63 Helix numbering 1
C U A C U G C G A A A G C A U U U G C C A A G G A - U G U U U U(C A)U U A A^U C A A G A A C G G A A A						64 Pteris vittata
C U U C U G C G A A A G C A U U U G C C A A G G A - U G U U U U(C A)U U A A^U C A A G A A C G G A A A						65 Megaceros aenigmaticus
C U U C U G C G A A A G C A U U U G C C A A G G A - U G U U U U(C A)U U A A^U C A A G A A C G G A A A						66 Notophylas breutelii
C U U C U G C G A A A G C A U U U G C C A A G G A - U G U U U U(C A)U U A A^U C A A G A A C G G A A A						67 Physcomitrella patens
C U U C U G C G A A A G C A U U U G C C A A G G A - U G U U U U(C A)U U A A^U C A A G A A C G G A A A						68 Sphagnum palustre

1210	1220	1230	1240	1250
[G U U A G G G A U I C I G - A A) G A U G) A U C U G I U A C I C G U C I G U A) G U - C U U A A C) C A U A A A A				1 Saccharomyces cerevisiae
[G U U A G G G A U I C I G - A A) G A C G) A U C A G A U A C I C G U C I G U A) G U - C U U A A C) C A U A A A A				2 Schizosaccharomyces pombe
[G U U G G G G G I G C U I C I G - A A) G A C G) A U C A G A U A C I C G U C I G U A) G U - C U U A A C) C A U A A A A				3 Physcomitrella patens
[G U U G G G G G I G C U I C I G - A A) G A C G) A U C A G A U A C I C G U C I G U A) G U - C U U A A C) C A U A A A A				4 Anthoceros agrestis
[G U U G G G G G I G C U I C I G - A A) G A C G) A U C A G A U A C I C G U C I G U A) G U - C U U A A C) C A U A A A A				5 Anthoceros laevis
[G U U G G G G G I G C U I C I G - A A) G A C G) A U C A G A U A C I C G U C I G U A) G U - C U U A A C) C A U A A A A				6 Pellia epiphylla
[G U U G G G G G I G C U I C I G - A A) G A C G) A U C A G A U A C I C G U C I G U A) G U - C U U A A C) C A U A A A A				7 Reboulia hemisphaerica 2
[G U U G G G G G I G C U I C I G - A A) G A C G) A U C A G A U A C I C G U C I G U A) G U - C U U A A C) C A U A A A A				8 Riccardia pinguis
[G U U G G G G G I G C U I C I G - A A) G A C G) A U C A G A U A C I C G U C I G U A) G U - C U U A A C) C A U A A A A				9 Sphaerocarpos donnelli
[G U U G G G G G I G C U I C I G - A A) G A C G) A U C A G A U A C I C G U C I G U A) G U - C U U A A C) C A U A A A A				10 Sphaerocarpos texanus
[G U U G G G G G I G C U I C I G - A A) G A C G) A U C A G A U A C I C G U C I G U A) G U - C U U A A C) C A U A A A A				11 Atrichum undulatum
[G U U G G G G G I G C U I C I G - A A) G A C G) A U C A G A U A C I C G U C I G U A) G U - C U U A A C) C A U A A A A				12 Bryum argenteum
[G U U G G G G G I G C U I C I G - A A) G A C G) A U C A G A U A C I C G U C I G U A) G U - C U U A A C) C A U A A A A				13 Eurhynchium hians
[G U U G G G G G I G C U I C I G - A A) G A C G) A U C A G A U A C I C G U C I G U A) G U - C U U A A C) C A U A A A A				14 Funaria hygrometrica
[G U U G G G G G I G C U I C I G - A A) G A C G) A U C A G A U A C I C G U C I G U A) G U - C U U A A C) C A U A A A A				15 Leptobryum pyriforme
[G U U G G G G G I G C U I C I G - A A) G A C G) A U C A G A U A C I C G U C I G U A) G U - C U U A A C) C A U A A A A				16 Mnium hornum
[G U U G G G G G I G C U I C I G - A A) G A C G) A U C A G A U A C I C G U C I G U A) G U - C U U A A C) C A U A A A A				17 Physcomitrium pyriforme
[G U U G G G G G I G C U I C I G - A A) G A C G) A U C A G A U A C I C G U C I G U A) G U - C U U A A C) C A U A A A A				18 Polytrichum formosum
[G U U G G G G G I G C U I C I G - A A) G A C G) A U C A G A U A C I C G U C I G U A) G U - C U U A A C) C A U A A A A				19 Sphagnum cuspidatum
[G U U G G G G G I G C U I C I G - A A) G A C G) A U C A G A U A C I C G U C I G U A) G U - C U U A A C) C A U A A A A				20 Calypogeia arguta
[G U U G G G G G I G C U I C I G - A A) G A C G) A U C A G A U A C I C G U C I G U A) G U - C U U A A C) C A U A A A A				21 Conocephalum conicum
[G U U G G G G G I G C U I C I G - A A) G A C G) A U C A G A U A C I C G U C I G U A) G U - C U U A A C) C A U A A A A				22 Fossombronina pusilla
[G U U G G G G G I G C U I C I G - A A) G A C G) A U C A G A U A C I C G U C I G U A) G U - C U U A A C) C A U A A A A				23 Marchantia polymorpha
[G U U G G G G G I G C U I C I G - A A) G A C G) A U C A G A U A C I C G U C I G U A) G U - C U U A A C) C A U A A A A				24 Riccia fluitans
[G U U G G G G G I G C U I C I G - A A) G A C G) A U C A G A U A C I C G U C I G U A) G U - C U U A A C) C A U A A A A				25 Scapania nemorea
[G U U G G G G G I G C U I C I G - A A) G A C G) A U C A G A U A C I C G U C I G U A) G U - C U U A A C) C A U A A A A				26 Equisetum hyemale
[G U U G G G G G I G C U I C I G - A A) G A C G) A U C A G A U A C I C G U C I G U A) G U - C U U A A C) C A U A A A A				27 Equisetum robustum
[G U U G G G G G I G C U I C I G - A A) G A C G) A U C A G A U A C I C G U C I G U A) G U - C U U A A C) C A U A A A A				28 Isoetes engelmannii
[G U U G G G G G I G C U I C I G - A A) G A C G) A U C A G A U A C I C G U C I G U A) G U - C U U A A C) C A U A A A A				29 Huperzia lucidula
[G U U G G G G G I G C U I C I G - A A) G A C G) A U C A G A U A C I C G U C I G U A) G U - C U U A A C) C A U A A A A				30 Lycopodiella inundata
[G U U G G G G G I G C U I C I G - A A) G A C G) A U C A G A U A C I C G U C I G U A) G U - C U U A A C) C A U A A A A				31 Lycopodium phlegmaria
[G U U G G G G G I G C U I C I G - A A) G A C G) A U C A G A U A C I C G U C I G U A) G U - C U U A A C) C A U A A A A				32 Lycopodium taxifolium
[G U U G G G G G I G C U I C I G - A A) G A C G) A U C A G A U A C I C G U C I G U A) G U - C U U A A C) C A U A A A A				33 Lycopodium tristachyum
[G U U G G G G G I G C U I C I G - A A) G A C G) A U C A G A U A C I C G U C I G U A) G U - C U U A A C) C A U A A A A				34 Selaginella galleottii
[G U U G G G G G I G C U I C I G - A A) G A C G) A U C A G A U A C I C G U C I G U A) G U - C U U A A C) C A U A A A A				35 Selaginella sp.
[G U U G G G G G I G C U I C I G - A A) G A C G) A U C A G A U A C I C G U C I G U A) G U - C U U A A C) C A U A A A A				36 Oryza sativa
[G U U G G G G G I G C U I C I G - A A) G A C G) A U C A G A U A C I C G U C I G U A) G U - C U U A A C) C A U A A A A				37 Zea mays
[G U U G G G G G I G C U I C I G - A A) G A C G) A U C A G A U A C I C G U C I G U A) G U - C U U A A C) C A U A A A A				38 Arabidopsis thaliana
[G U U G G G G G I G C U I C I G - A A) G A C G) A U C A G A U A C I C G U C I G U A) G U - C U U A A C) C A U A A A A				39 Glycine max
[G U U G G G G G I G C U I C I G - A A) G A C G) A U C A G A U A C I C G U C I G U A) G U - C U U A A C) C A U A A A A				40 Sinapis alba
[G U U G G G G G I G C U I C I G - A A) G A C G) A U C A G A U A C I C G U C I G U A) G U - C U U A A C) C A U A A A A				41 Zamia pumila
[G U U G G G G G I G C U I C I G - A A) G A C G) A U C A G A U A C I C G U C I G U A) G U - C U U A A C) C A U A A A A				42 Gnetum leyboldii
[G U U G G G G G I G C U I C I G - A A) G A C G) A U C A G A U A C I C G U C I G U A) G U - C U U A A C) C A U A A A A				43 Ginkgo biloba
[G U U G G G G G I G C U I C I G - A A) G A C G) A U C A G A U A C I C G U C I G U A) G U - C U U A A C) C A U A A A A				44 Pinus luchuensis
[G U U G G G G G I G C U I C I G - A A) G A C G) A U C A G A U A C I C G U C I G U A) G U - C U U A A C) C A U A A A A				45 Pinus wallichiana
[G U U G G G G G I G C U I C I G - A A) G A C G) A U C A G A U A C I C G U C I G U A) G U - C U U A A C) C A U A A A A				46 Podocarpus nakaii
[G U U G G G G G I G C U I C I G - A A) G A C G) A U C A G A U A C I C G U C I G U A) G U - C U U A A C) C A U A A A A				47 Taxus mairei
[G U U G G G G G I G C U I C I G - A A) G A C G) A U C A G A U A C I C G U C I G U A) G U - C U U A A C) C A U A A A A				48 Adiantum raddianum
[G U U G G G G G I G C U I C I G - A A) G A C G) A U C A G A U A C I C G U C I G U A) G U - C U U A A C) C A U A A A A				49 Adiantum raddianum 2
[G U U G G G G G I G C U I C I G - A A) G A C G) A U C A G A U A C I C G U C I G U A) G U - C U U A A C) C A U A A A A				50 Ophioglossum petiolatum
[G U U G G G G G I G C U I C I G - A A) G A C G) A U C A G A U A C I C G U C I G U A) G U - C U U A A C) C A U A A A A				51 Osmunda cinnamomea
[G U U G G G G G I G C U I C I G - A A) G A C G) A U C A G A U A C I C G U C I G U A) G U - C U U A A C) C A U A A A A				52 Salvinia natans
[G U U G G G G G I G C U I C I G - A A) G A C G) A U C A G A U A C I C G U C I G U A) G U - C U U A A C) C A U A A A A				53 Psilotum nudum 2
[G U U G G G G G I G C U I C I G - A A) G A C G) A U C A G A U A C I C G U C I G U A) G U - C U U A A C) C A U A A A A				54 Chara foetida
[G U U G G G G G I G C U I C I G - A A) G A C G) A U C A G A U A C I C G U C I G U A) G U - C U U A A C) C A U A A A A				55 Coleochaete scutata
[G U U G G G G G I G C U I C I G - A A) G A C G) A U C A G A U A C I C G U C I G U A) G U - C U U A A C) C A U A A A A				56 Klebsormidium flaccidum
[G U U G G G G G I G C U I C I G - A A) G A C G) A U C A G A U A C I C G U C I G U A) G U - C U U A A C) C A U A A A A				57 Nitella flexilis
[G U U G G G G G I G C U I C I G - A A) G A C G) A U C A G A U A C I C G U C I G U A) G U - C U U A A C) C A U A A A A				58 Nitella sp.
[G U U G G G G G I G C U I C I G - A A) G A C G) A U C A G A U A C I C G U C I G U A) G U - C U U A A C) C A U A A A A				59 Chlamydomonas reinhardtii
[G U U G G G G G I G C U I C I G - A A) G A C G) A U C A G A U A C I C G U C I G U A) G U - C U U A A C) C A U A A A A				60 Hydrodictyon reticulatum
[G U U G G G G G I G C U I C I G - A A) G A C G) A U C A G A U A C I C G U C I G U A) G U - C U U A A C) C A U A A A A				61 Parietochloris pseudoalveolaris
[G U U G G G G G I G C U I C I G - A A) G A C G) A U C A G A U A C I C G U C I G U A) G U - C U U A A C) C A U A A A A				62 Dictyostelium discoideum
[G U U G G G G G I G C U I C I G - A A) G A C G) A U C A G A U A C I C G U C I G U A) G U - C U U A A C) C A U A A A A				63 Helix numbering 1
[G U U G G G G G I G C U I C I G - A A) G A C G) A U C A G A U A C I C G U C I G U A) G U - C U U A A C) C A U A A A A				64 Pteris vittata
[G U U G G G G G I G C U I C I G - A A) G A C G) A U C A G A U A C I C G U C I G U A) G U - C U U A A C) C A U A A A A				65 Megaceros aenigmaticus
[G U U G G G G G I G C U I C I G - A A) G A C G) A U C A G A U A C I C G U C I G U A) G U - C U U A A C) C A U A A A A				66 Notophyas breutelii
[G U U G G G G G I G C U I C I G - A A) G A C G) A U C A G A U A C I C G U C I G U A) G U - C U U A A C) C A U A A A A				67 Physcomitrella patens
[G U U G G G G G I G C U I C I G - A A) G A C G) A U C A G A U A C I C G U C I G U A) G U - C U U A A C) C A U A A A A				68 Sphagnum palustre

	1260	1270	1280	1290	1300	
C U A U I G C C C - G A C U I A G I - A U C G G G U G G I U I G U U - U U U U - U A A U - (G A - - - C C						1 Saccharomyces cerevisiae
C U A U I G C C C - G A C U I A G G I G A U U C G G G C G G A I U I G U U - - U C A U U U U A U C - (G A - - - C U						2 Schizosaccharomyces pombe
C G A U I G C C C - G A C U I A G G I G A U U G G G C G G G A I U I G U U - - A A U U - U G A U - (G A - - - C U						3 Physcomitrella patens
C G A U I G C C C - G A C U I A G G I G A U U G G G C G G G A I U I G U U - - A A U U - U G A U - (G A - - - C (G						4 Anthoceros agrestis
C G A U I G C C C - G A C U I A G G I G A U U G G G C G G G A I U I G U U - - A A U U - U G A U - (G A - - - C U						5 Anthoceros laevis
C G A U I G C C C - G A C U I A G G I G A U U G G G C G G G A I U I G U U - - A A U U - U G A U - (G A - - - C U						6 Pellia epiphylla
C G A U I G C C C - G A C U I A G G I G A U U G G G C G G G A I U I G U U - - G A U U - A G A U - (G A - - - C U						7 Rebolia hemisphaerica 2
C G A U I G C C C - G A C U I A G G I G A U U G G G C G G G A I U I G U U - - A A U U - U G A U - (G A - - - C U						8 Riccardia pinguis
C G A U I G C C C - G A C U I A G G I G A U U G G G C G G G A I U I G U U - - A A U U - U G A U - (G A - - - C U						9 Sphaerocarpos donnellii
C G A U I G C C C - G A C U I A G G I G A U U G G G C G G G A I U I G U U - - A A U U - U G A U - (G A - - - C U						10 Sphaerocarpos texanus
C G A U I G C C C - G A C U I A G G I G A U U G G G C G G G A I U I G U U - - A A U U - U G A U - (G A - - - C C						11 Atrichum undulatum
C G A U I G C C C - G A C U I A G G I G A U U G G G C G G G A I U I G U U - - G C U U - U G A U - (G A - - - C C						12 Bryum argenteum
C G A U I G C C C - G A C U I A G G I G A U U G G G C G G G A I U I G U U - - C A U U - U G A U - (G A - - - C (C						13 Eurhynchium hians
C G A U I G C C C - G A C U I A G G I G A U U G G G C G G G A I U I G U U - - A C U U - U G A U - (G A - - - C U						14 Funaria hygrometrica
C G A U I G C C C - G A C U I A G G I G A U U G G G C G G G A I U I G U U - - G A U U - U G A U - (G A - - - C C						15 Leptobryum pyriforme
C G A U I G C C C - G A C U I A G G I G A U U G G G C G G G A I U I G U U - - G A U U - C G A U - (G A - - - C C						16 Mnium hornum
C G A U I G C C C - G A C U I A G G I G A U U G G G C G G G A I U I G U U - - A C U U - U G A U - (C A - - - C U						17 Physcomitrium pyriforme
C G A U I G C C C - G A C U I A G G I G A U U C G G C G G G A I U I G U U - - A A U U - U G A U - (G A - - - C C						18 Polytrichum formosum
C G A U I G C C C - G A C U I A G G I G A U U C G G C G G G A I U I G U U - - G A A U - C G A U - (G A - - - C U						19 Sphagnum cuspidatum
C G A U I G C C C - G A C U I A G G I G A U U C G G C G G G A I U I G U U - - A A U U - U G A U - (G A - - - C U						20 Calypogeia arguta
C G A U I G C C C - G A C U I A G G I G A U U C G G C G G G A I U I G U U - - U C U U U U G A U - (G A - - - C U						21 Conocephalum conicum
C G A U I G C C C - G A C U I A G G I G A U U C G G C G G G A I U I G U U - - A A U U - U G A U - (G A - - - C U						22 Fossombronina pusilla
C G A U I G C C C - G A C U I A G G I G A U U C G G C G G G A I U I G U U - - A A U U - U G A U - (G A - - - C U						23 Marchantia polymorpha
C G A U I G C C C - G A C U I A G G I G A U U C G G C G G G A I U I G U U - - U G A U - A G A U - (G A - - - C U						24 Riccia fluitans
C G A U I G C C C - G A C U I A G G I G A U U C G G C G G G A I U I G U U - - A A U U - U G A U - (G A - - - C U						25 Scapania nemorea
C G A U I G C C C - G A C U I A G G I G A U U C G G C G G G A I U I G U U - - A C U U - C G A U - (G A - - - C U						26 Equisetum hyemale
C G A U I G C C C - G A C U I A G G I G A U U C G G C G G G A I U I G U U - - A C U U - C G A U - (G A - - - C U						27 Equisetum robustum
C G A U I G C C C - G A C U I A G G I G A U U C G G C G G G A I U I G U U - - A C U U - C G A U - (G A - - - C U						28 Isoetes engelmannii
C G A U I G C C C - G A C U I A G G I G A U U C G G C G G G A I U I G U U - - A A U U - C G A U - (G A - - - C C						29 Huperzia lucidula
C G A U I G C C C - G A C U I A G G I G A U U C G G C G G G A I U I G U U - - N A U U - C G A U - (G A - - - C C						30 Lycopodiella inundata
C G A U I G C C C - G A C U I A G G I G A U U C G G C G G G A I U I G U U - - A A U U - C G A U - (G A - - - C C						31 Lycopodium phlegmaria
C G A U I G C C C - G A C U I A G G I G A U U C G G C G G G A I U I G U U - - A A U U - C G A U - (G A - - - C C						32 Lycopodium taxifolium
C G A U I G C C C - G A C U I A G G I G A U U C G G C G G G A I U I G U U - - A A U U - C G A U - (G A - - - C C						33 Lycopodium tristachyum
C G A U I G C C C - G A C U I A G G I G A U U C G G C G G G A I U I G U U - - A U U U - G A U - (G A - - - C U						34 Selaginella galleottii
C G A U I G C C C - G A C U I A G G I G A U U C G G C G G G A U I G U U - - A U U U - G G A U - (G A - - - C (A						35 Selaginella sp.
C G A U I G C C C - G A C U I A G G I G A U U C G G C G G G A I U I G U U - - G C U U - U A A G - (G A - - - C U						36 Oryza sativa
C G A U I G C C C - G A C U I A G G I G A U U C G G C G G G A I U I G U U - - A C U A - A U A G - (G A - - - C (C						37 Zea mays
C G A U I G C C C - G A C U I A G G I G A U U C G G C G G G A I U I G U U - - G C U U - U A A G - (G A - - - C U						38 Arabidopsis thaliana
C G A U I G C C C - G A C U I A G G I G A U U C G G C G G G A I U I G U U - - G C U U - U A A G - (G A - - - C U						39 Glycine max
C G A U I G C C C - G A C U I A G G I G A U U C G G C G G G A I U I G U U - - G C U U - U A A G - (G A - - - C U						40 Sinapis alba
C G A U I G C C C - G A C U I A G G I G A U U C G G C G G G A I U I G U U - - G C U U - U A A G - (G A - - - C U						41 Zamia pumila
C G A U I G C C C - G A C U I A G G I G A U U C G G C G G G A I U I G U U - - G C U U - U A A G - (G A - - - C U						42 Gnetum leyboldii
C G A U I G C C C - G A C U I A G G I G A U U C G G C G G G A I U I G U U - - G C U U - U A A G - (G A - - - C U						43 Ginkgo biloba
C G A U I G C C C - G A C U I A G G I G A U U C G G C G G G A I U I G U U - - G C U U - U A A G - (G A - - - C U						44 Pinus luchuensis
C G A U I G C C C - G A C U I A G G I G A U U C G G C G G G A I U I G U U - - G C U U - U A A G - (G A - - - C U						45 Pinus wallichiana
C G A U I G C C C - G A C U I A G G I G A U U C G G C G G G A I U I G U U - - G C U U - U A A G - (G A - - - C U						46 Podocarpus nakaii
C G A U I G C C C - G A C U I A G G I G A U U C G G C G G G A I U I G U U - - G C U U - U A A G - (G A - - - C U						47 Taxus mairei
C G A U I G C C C - G A C U I A G G I G A U U C G G C G G G A I U I G U U - - A C U U - U G A U - (G A - - - C U						48 Adiantum raddianum
C G A U I G C C C - G A C U I A G G I G A U U C G G C G G G A I U I G U U - - A C U U - U G A U - (G A - - - C U						49 Adiantum raddianum 2
C G A U I G C C C - G A C U I A G G I G A U U C G G C G G G A I U I G U U - - A C U U - C G A U - (G A - - - C U						50 Ophioglossum petiolatum
C G A U I G C C C - G A C U I A G G I G A U U C G G C G G G A I U I G U U - - A C U U - C A A U - (G A - - - C U						51 Osmunda cinnamomea
C G A U I G C C C - G A C U I A G G I G A U U C G G C G G G A I U I G U U - - A C U U - C G A U - (G A - - - C U						52 Salvinia natans
C G A U I G C C C - G A C U I A G G I G A U U C G G C G G G A I U I G U U - - A C U U - U G A U - (G A - - - C U						53 Psilotum nudum 2
C G A U I G C C C - G A C U I A G G I G A U U C G G C G G G A I U I G U U - - U A U U - G G A U - (G A - - - C U						54 Chara foetida
C G A U I G C C C - G A C U I A G G I G A U U C G G C G G G A I U I G U U - - A A U U - U G A U - (G A - - - C U						55 Coleochaete scutata
C G A U I G C C C - G A C U I A G G I G A U U C G G C G G G A I U I G U U - - A A U U - U G A U - (G A - - - C U						56 Klebsormidium flaccidum
C G A U I G C C C - G A C U I A G G I G A U U C G G C G G G A I U I G U U - - U A U U - U G A U - (G A - - - C U						57 Nitella flexilis
C G A U I G C C C - G A C U I A G G I G A U U C G G C G G G A I U I G U U - - U A U U - U G A U - (G A - - - C U						58 Nitella sp.
C G A U I G C C C - G A C U I A G G I G A U U C G G C G G G A I U I G U U - - C U U U - U G A U - (G A - - - C U						59 Chlamydomonas reinhardtii
C G A U I G C C C - G A C U I A G G I G A U U C G G C G G G A I U I G U U - - U U U U - U G A U - (G A - - - C U						60 Hydrodictyon reticulatum
C G A U I G C C C - G A C U I A G G I G A U U C G G C G G G A I U I G U U - - U A U U - C G A U - (G A - - - C U						61 Parietochloris pseudoalveolaris
C U A U I G U C C - G A C U I A G G I G A U U C G G U U A A I A U U U U - - U U C - - - - [A A A U U - 29						62 Dictyostelium discoideum
C G A U I G C C C - 28 - G A C U I A G G I G A U U C G G C G G G A I U I G U U - - A C U U - U G A U - (G A - - - C U						63 Helix numbering 1
C G A U I G C C C - G A C U I A G G I G A U U C G G C G G G A I U I G U U - - A A U U - A G A U - (G A - - - C U						64 Pteris vittata
C G A U I G C C C - G A C U I A G G I G A U U C G G C G G G A I U I G U U - - A A U - A G A U U (G A - - - C C						65 Megaceros aenigmaticus
C G A U I G C C C - G A C U I A G G I G A U U C G G C G G G A I U I G U U - - A C U U - U G A U - (G A - - - C U						66 Notophylas breutelu
C G A U I G C C C - G A C U I A G G I G A U U C G G C G G G A I U I G U U - - G A A U - C G A U - (G A - - - C U						67 Physcomitrella patens
C G A U I G C C C - G A C U I A G G I G A U U C G G C G G G A I U I G U U - - G A A U - C G A U - (G A - - - C U						68 Sphagnum palustre

	1310	1320	1330	1340	1350		
..	C A C U C G G U	- A C -	C U U A	- C I G A I -	G A A A U C J A	- A A I G U C U U U G I G G U U U C U G G G G	1 Saccharomyces cerevisiae
U G C C U C C G G I C	- A I C -	C U U A	- C I G A I -	G A A A U C J A	- A A I G U C U U U G I G G U U U C U G G G G	2 Schizosaccharomyces pombe	
C C G C C A G I C	- A I C -	C U U A	- U I G A I -	G A A A U C J A	- A A I G U U U U U G I G G U U U C U G G G G	3 Physcomitrella patens	
U C C G C C A G I C	- A I C -	C U U A	- U I G A I -	G A A A U C J A	- A A I G U U U U U G I G G U U U C U G G G G	4 Anthoceros agrestis	
C C G C C A G I C	- A I C -	C U U A	- U I G A I -	G A A A U C J A	- A A I G U U U U U G I G G U U U C U G G G G	5 Anthoceros laevis	
C C G C C A G I C	- A I C -	C U U A	- U I G A I -	G A A A U C J A	- A A I G U U U U U G I G G U U U C U G G G G	6 Pellia epiphylla	
C C G C C A G I C	- A I C -	C U U A	- U I G A I -	G A A A U C J A	- A A I G U U U U U G I G G U U U C U G G G G	7 Reboulia hemisphaerica 2	
C C G C C A G I C	- A I C -	C U U A	- U I G A I -	G A A A U C J A	- A A I G U U U U U G I G G U U U C U G G G G	8 Riccardia pinguis	
C C G C C A G I C	- A I C -	C U U A	- U I G A I -	G A A A U C J A	- A A I G U U U U U G I G G U U U C U G G G G	9 Sphaerocarpos donnelli	
C C G C C A G I C	- A I C -	C U U A	- U I G A I -	G A A A U C J A	- A A I G U U U U U G I G G U U U C U G G G G	10 Sphaerocarpos texanus	
C C G C C A G I C	- A I C -	C U U A	- U I G A I -	G A A A U C J A	- A A I G U U U U U G I G G U U U C U G G G G	11 Atrichum undulatum	
C C G C C A G I C	- A I C -	C U U A	- U I G A I -	G A A A U C J A	- A A I G U U U U U G I G G U U U C U G G G G	12 Bryum argenteum	
U C C G C C A G I C	- A I C -	C U U A	- U I G A I -	G A A A U C J A	- A A C U U U U U U G I G G C U U U C U G G G G	13 Eurhynchium hians	
C C G C C A G I C	- A I C -	C U U A	- U I G A I -	G A A A U C J A	- A A I G U U U U U G I G G U U U C U G G G G	14 Funaria hygrometrica	
C C G C C A G I C	- A I C -	C U U A	- U I G A I -	G A A A U C J A	- A A I G U U U U U G I G G U U U C U G G G G	15 Leptobryum pyriforme	
C C G C C A G I C	- A I C -	C U U A	- U I G A I -	G A A A U C J A	- A A I G U U U U U G I G G U U U C U G G G G	16 Mnium hornum	
C C G C C A G I C	- A I C -	C U U A	- U I G A I -	G A A A U C J A	- A A I G U U U U U G I G G C U U C U G G G G	17 Physcomitrium pyriforme	
C C G C C A G I C	- A I C -	C U U A	- U I G A I -	G A A A U C J A	- A A I G U U U U U G I G G U U U C U G G G G	18 Polytichum formosum	
C C G C C A G I C	- A I C -	C U U A	- U I G A I -	G A A A U C J A	- A A I G U U U U U G I G G U U U C U G G G G	19 Sphagnum cuspidatum	
C C G C C A G I C	- A I C -	C U U A	- U I G A I -	G A A A U C J A	- A A I G U U U U U G I G G U U U C U G G G G	20 Calypogeia arguta	
C C G C C A G I C	- A I C -	C U U C G	- A I G A I -	G A A A U C J A	- A A I G U U U U U G I G G U U U C U G G G G	21 Conocephalum conicum	
C C G C C A G I C	- A I C -	C U U A	- U I G A I -	G A A A U C J A	- A A I G U U U U U G I G G U U U C U G G G G	22 Fossombronina pusilla	
C C G C C A G I C	- A I C -	C U U C A	- U I G A I -	G A A A U C J A	- A A I G U U U U U G I G G U U U C U G G G G	23 Marchantia polymorpha	
C C G C C A G I C	- A I C -	C U U C A	- U I G A I -	G A A A U C J A	- A A I G U U U U U G I G G U U U C U G G G G	24 Riccia fluitans	
C C G C C A G I C	- A I C -	C U U A	- U I G A I -	G A A A U C J A	- A A I G U U U U U G I G G U U U C U G G G G	25 Scapania nemorea	
C U G C C A G I C	- A I C -	C U U A	- U I G A I -	G A A A U C J A	- A A I G U C U U U G I G G U U U C U G G G G	26 Equisetum hyemale	
C C G C C A G I C	- A I C -	C U U A	- U I G A I -	G A A A U C J A	- A A I G U C U U U G I G G U U U C U G G G G	27 Equisetum robustum	
C C G C C A G I C	- A I C -	C U U C	- U I G A I -	G A A A U C J A	- A A I G U U U U U G I G G U U U C U G G G G	28 Isoetes engelmannii	
C C G C C A G I C	- A I C -	C U U A	- U I G A I -	G A A A U C J A	- A A I G U U U U U G I G G U U U C U G G G G	29 Huperzia lucidula	
C C G C C A G I C	- A I C -	C U U A	- U I G A I -	G A A A U C J A	- A A I G U U U U U G I G G U U U C U G G G G	30 Lycopodiella inundata	
C C G C C A G I C	- A I C -	C U U A	- U I G A I -	G A A A U C J A	- A A I G U U U U U G I G G U U U C U G G G G	31 Lycopodium phlegmaria	
C C G C C A G I C	- A I C -	C U U A	- U I G A I -	G A A A U C J A	- A A I G U U U U U G I G G U U U C U G G G G	32 Lycopodium taxifolium	
C C G C C A G I C	- A I C -	C U U A	- U I G A I -	G A A A U C J A	- A A I G U U U U U G I G G U U U C U G G G G	33 Lycopodium tristachyum	
C C G C C A G I C	- A I C -	C U U C	- U I G A I -	G A A A U C J A	- A A I G U C U U U G I G G U U U C U G G G G	34 Selaginella galleortii	
U C C G C C A G I C	- A I C -	C U U A	- U I G A I -	G A A A U C J A	- A A I G U C U U U G I G G U U U C U G G G G	35 Selaginella sp.	
C C G C C A G I C	- A I C -	C U U A	- U I G A I -	G A A A U C J A	- A A I G U C U U U G I G G U U U C U G G G G	36 Oryza sativa	
U C C G C C U G G I C	- A I C -	C U U A	- U I G A I -	G A A A U C J A	- A A I G U C U U U G I G G U U U C U G G G G	37 Zea mays	
C C G C C U G G I C	- A I C -	C U U A	- U I G A I -	G A A A U C J A	- A A I G U U U U U G I G G U U U C U G G G G	38 Arabidopsis thaliana	
C C G C C U G G I C	- A I C -	C U U A	- U I G A I -	G A A A U C J A	- A A I G U C U U U G I G G U U U C U G G G G	39 Glycine max	
C C G C C U G G I C	- A I C -	C U U A	- U I G A I -	G A A A U C J A	- A A I G U U U U U G I G G U U U C U G G G G	40 Sinapis alba	
C C G C C U G G I C	- A I C -	C U U A	- U I G A I -	G A A A U C J A	- A A I G U U U U U G I G G U U U C U G G G G	41 Zamia pumila	
C C G C C U G G I C	- A I C -	C U U C	- U I G A I -	G A A A U C J A	- G A I G U U U U U G I G G U U U C U G G G G	42 Gnetum leyboldii	
C C G C C U G G I C	- A I C -	C U U C	- U I G A I -	G A A A U C J A	- G A I G U U U U U G I G G U U U C U G G G G	43 Ginkgo biloba	
C C G C C A G I C	- A I C -	C U U C	- U I G A I -	G A A A U C J A	- G A I G U U U U U G I G G U U U C U G G G G	44 Pinus luchuensis	
C C G C C A G I C	- A I C -	C U U C	- U I G A I -	G A A A U C J A	- G A I G U U U U U G I G G U U U C U G G G G	45 Pinus wallichiana	
C C G C C A G I C	- A I C -	C U U C	- U I G A I -	G A A A U C J A	- G A I G U U U U U G I G G U U U C U G G G G	46 Podocarpus nakaii	
C C G C C A G I C	- A I C -	C U U A	- U I G A I -	G A A A U C J A	- G A I G U U U U U G I G G U U U C U G G G G	47 Taxus mairei	
C C G C C A G I C	- A I C -	C U U A	- U I G A I -	G A A A U C J A	- A A I G U U U U U G I G G U U U C U G G G G	48 Adiantum raddianum	
C C G C C A G I C	- A I C -	C U U A	- U I G A I -	G A A A U C J A	- A A I G U C U U U G I G G U U U C U G G G G	49 Adiantum raddianum 2	
C C G C C A G I C	- A I C -	C U U A	- U I G A I -	G A A A U C J A	- A A I G U U U U U G I G G U U U C U G G G G	50 Ophioglossum petiolatum	
C C G C C A G I C	- A I C -	C U U A	- U I G A I -	G A A A U C J A	- A A I G U U U U U G I G G U U U C U G G G G	51 Osmunda cinnamomea	
U C C G C C A G I C	- A I C -	C U U A	- U I G A I -	G A A A U C J A	- A A I G U U U U U G I G G U U U C U G G G G	52 Salvinia natans	
C C G C C A G I C	- A I C -	C U U G	- U I G A I -	G A A A U C J A	- A A I G U C U U U G I G G U U U C U G G G G	53 Psilotum nudum 2	
C C G C C A G I C	- A I C -	C U U A	- U I G A I -	G A A A U C J A	- A A I G U U U U U G I G G U U U C U G G G G	54 Chara foetida	
C C G C C A G I C	- A I C -	C U U A	- U I G A I -	G A A A U C J A	- A A I G U U U U U G I G G U U U C U G G G G	55 Coleochaete scutata	
C C G C C A G I C	- A I C -	C U U G	- U I G A I -	G A A A U C J A	- A A I G U U U U U G I G G U U U C U G G G G	56 Klebsormidium flaccidum	
C C G C C A G I C	- A I C -	C U U G	- U I G A I -	G A A A U C J A	- A A I G U U U U U G I G G U U U C U G G G G	57 Nitella flexilis	
C C G C C A G I C	- A I C -	C U U G	- U I G A I -	G A A A U C J A	- A A I G U U U U U G I G G U U U C U G G G G	58 Nitella sp.	
C U G C C A G I C	- A I C -	C U U A	- U I G A I -	G A A A U C J A	- A A I G U U U U U G I G G U U U C U G G G G	59 Chlamydomonas reinhardtii	
C C G C C A G I C	- A I C -	C U U A	- U I G A I -	G A A A U C J A	- A A I G U U U U U G I G G U U U C U G G G G	60 Hydrodictyon reticulatum	
C C G C C A G I C	- A I C -	C U U A	- U I G A I -	G A A A U C J A	- A A I G U U U U U G I G G U U U C U G G G G	61 Parietochloris pseudoalveolaris	
U A A U C G G I C	- A I C -	C U U G	- U I G A I -	G A A A U C J A	- U A I G U U U U U A I G A U U U C U G G G G	62 Dictyostelium discoideum	
9' - - - - -	- - - - -	- - - - -	- - - - -	- - - - -	- - - - -	- - - - -	63 Helix numbering 1
C C G C C A G I C	- A I C -	C U U A	- U I G A I -	G A A A U C J A	- A A I G U U U U U G I G G U U U C U G G G G	64 Pteris vittata	
C C G C C A G I C	- A I C -	C U U A	- U I G A I -	G A A A U C J A	- A A I G U U U U U G I G G U U U C U G G G G	65 Megaceros aenigmaticus	
C C G C C A G I C	- A I C -	C U U A	- U I G A I -	G A A A U C J A	- A A I G U U U U U G I G G U U U C U G G G G	66 Notophylus breutelii	
C C G C C A G I C	- A I C -	C U U A	- U I G A I -	G A A A U C J A	- A A I G U U U U U G I G G U U U C U G G G G	67 Physcomitrella patens	
C C G C C A G I C	- A I C -	C U U A	- U I G A I -	G A A A U C J A	- A A I G U U U U U G I G G U U U C U G G G G	68 Sphagnum palustre	

DCSE Alignment

	1460	1470	1480	1490	1500	
C	A	A	A	A	A	1 Saccharomyces cerevisiae
U	G	A	G	G	G	2 Schizosaccharomyces pombe
C	C	A	G	G	G	3 Physcomitrella patens
C	A	G	G	G	G	4 Anthoceros agrestis
U	C	A	G	G	G	5 Anthoceros laevis
C	A	G	G	G	G	6 Pellia epiphylla
U	C	A	G	G	G	7 Reboulia hemisphaerica 2
C	A	G	G	G	G	8 Riccardia pinguis
U	C	A	G	G	G	9 Sphaerocarpos donnellii
C	A	G	G	G	G	10 Sphaerocarpos texanus
U	C	A	G	G	G	11 Atrichum undulatum
C	A	G	G	G	G	12 Bryum argenteum
U	C	A	G	G	G	13 Eurhynchium hians
C	A	G	G	G	G	14 Funaria hygrometrica
U	C	A	G	G	G	15 Leptobryum pyriforme
C	A	G	G	G	G	16 Mnium hornum
U	C	A	G	G	G	17 Physcomitrium pyriforme
C	A	G	G	G	G	18 Polytrichum formosum
U	C	A	G	G	G	19 Sphagnum cuspidatum
C	A	G	G	G	G	20 Calypogeia arguta
U	C	A	G	G	G	21 Conocephalum conicum
C	A	G	G	G	G	22 Fossombronina pusilla
U	C	A	G	G	G	23 Marchantia polymorpha
C	A	G	G	G	G	24 Riccia fluitans
U	C	A	G	G	G	25 Scapania nemorea
C	A	G	G	G	G	26 Equisetum hyemale
U	C	A	G	G	G	27 Equisetum robustum
C	A	G	G	G	G	28 Isoetes engelmannii
U	C	A	G	G	G	29 Huperzia lucidula
C	A	G	G	G	G	30 Lycopodiella inundata
U	C	A	G	G	G	31 Lycopodium phlegmaria
C	A	G	G	G	G	32 Lycopodium taxifolium
U	C	A	G	G	G	33 Lycopodium tristachyum
C	A	G	G	G	G	34 Selaginella galleottii
U	C	A	G	G	G	35 Selaginella sp.
C	A	G	G	G	G	36 Oryza sativa
U	C	A	G	G	G	37 Zea mays
C	A	G	G	G	G	38 Arabidopsis thaliana
U	C	A	G	G	G	39 Glycine max
C	A	G	G	G	G	40 Sinapis alba
U	C	A	G	G	G	41 Zania pumila
C	A	G	G	G	G	42 Gnetum leyboldii
U	C	A	G	G	G	43 Ginkgo biloba
C	A	G	G	G	G	44 Pinus luchuensis
U	C	A	G	G	G	45 Pinus wallichiana
C	A	G	G	G	G	46 Podocarpus naki
U	C	A	G	G	G	47 Taxus mairei
C	A	G	G	G	G	48 Adiantum raddianum
U	C	A	G	G	G	49 Adiantum raddianum 2
C	A	G	G	G	G	50 Ophioglossum petiolatum
U	C	A	G	G	G	51 Osmunda cinnamomea
C	A	G	G	G	G	52 Salvinia natans
U	C	A	G	G	G	53 Psilotum nudum 2
C	A	G	G	G	G	54 Chara foetida
U	C	A	G	G	G	55 Coleochaete scutata
C	A	G	G	G	G	56 Klebsormidium flaccidum
U	C	A	G	G	G	57 Nitella flexilis
C	A	G	G	G	G	58 Nitella sp.
U	C	A	G	G	G	59 Chlamydomonas reinhardtii
C	A	G	G	G	G	60 Hydrodictyon reticulatum
U	C	A	G	G	G	61 Parietochloris pseudoalveolaris
C	A	G	G	G	G	62 Dictyostellium discoideum
U	C	A	G	G	G	63 Helix numbering 1
C	A	G	G	G	G	64 Pteris vittata
U	C	A	G	G	G	65 Megaceros aenigmaticus
C	A	G	G	G	G	66 Notophylas breutelu
U	C	A	G	G	G	67 Physcomitrella patens
C	A	G	G	G	G	68 Sphagnum palustre

	1510	1520	1530	1540	1550	
G A U U U U G U G J G G	- U I G G (U) G G I U G I C A U G	- G C C G U U (C) U J C A (G U - U*G	G U G G A J G U G A			1 Saccharomyces cerevisiae
G A U U U C U A U G J G G	- U I G G (U) G G I U G I C A U G	- G C C G U U (C) U J A (G U - U*G	G U G G A J G U G A			2 Schizosaccharomyces pombe
G A U U U C U A U G J G G	- U I G G (U) G G I U G I C A U G	- G C C G U U (C) U J A (G U - U*G	G U G G A J G U G A			3 Physcomitrella patens
G A U U U C U A U G J G G	- U I G G (U) G G I U G I C A U G	- G C C G U U (C) U J A (G U - U*G	G U G G A J G U G A			4 Anthoceros agrestis
G A U U U C U A U G J G G	- U I G G (U) G G I U G I C A U G	- G C C G U U (C) U J A (G U - U*G	G U G G A J G U G A			5 Anthoceros laevis
G A U U U C U A U G J G G	- U I G G (U) G G I U G I C A U G	- G C C G U U (C) U J A (G U - U*G	G U G G A J G U G A			6 Pellia epiphylla
G A U U U C U A U G J G G	- U I G G (U) G G I U G I C A U G	- G C C G U U (C) U J A (G U - U*G	G U G G A J G U G A			7 Reboulia hemisphaerica 2
G A U U U C U A U G J G G	- U I G G (U) G G I U G I C A U G	- G C C G U U (C) U J A (G U - U*G	G U G G A J G U G A			8 Riccardia pinguis
G A U U U C U A U G J G G	- U I G G (U) G G I U G I C A U G	- G C C G U U (C) U J A (G U - U*G	G U G G A J G U G A			9 Sphaerocarpos donnelli
G A U U U C U A U G J G G	- U I G G (U) G G I U G I C A U G	- G C C G U U (C) U J A (G U - U*G	G U G G A J G U G A			10 Sphaerocarpos texanus
G A U U U C U A U G J G G	- U I G G (U) G G I U G I C A U G	- G C C G U U (C) U J A (G U - U*G	G U G G A J G U G A			11 Atrichum undulatum
G A U U U C U A U G J G G	- U I G G (U) G G I U G I C A U G	- G C C G U U (C) U J A (G U - U*G	G U G G A J G U G A			12 Bryum argenteum
G A U U U C U A U G J G G	- U I G G (U) G G I U G I C A U G	- G C C G U U (C) U J A (G U - U*G	G U G G A J G U G A			13 Eurhynchium hians
G A U U U C U A U G J G G	- U I G G (U) G G I U G I C A U G	- G C C G U U (C) U J A (G U - U*G	G U G G A J G U G A			14 Funaria hygrometrica
G A U U U C U A U G J G G	- U I G G (U) G G I U G I C A U G	- G C C G U U (C) U J A (G U - U*G	G U G G A J G U G A			15 Leptobryum pyriforme
G A U U U C U A U G J G G	- U I G G (U) G G I U G I C A U G	- G C C G U U (C) U J A (G U - U*G	G U G G A J G U G A			16 Mnium hornum
G A U U U C U A U G J G G	- U I G G (U) G G I U G I C A U G	- G C C G U U (C) U J A (G U - U*G	G U G G A J G U G A			17 Physcomitrium pyriforme
G A U U U C U A U G J G G	- U I G G (U) G G I U G I C A U G	- G C C G U U (C) U J A (G U - U*G	G U G G A J G U G A			18 Polytrichum formosum
G A U U U C U A U G J G G	- U I G G (U) G G I U G I C A U G	- G C C G U U (C) U J A (G U - U*G	G U G G A J G U G A			19 Sphagnum cuspidatum
G A U U U C U A U G J G G	- U I G G (U) G G I U G I C A U G	- G C C G U U (C) U J A (G U - U*G	G U G G A J G U G A			20 Calypogeia arguta
G A U U U C U A U G J G G	- U I G G (U) G G I U G I C A U G	- G C C G U U (C) U J A (G U - U*G	G U G G A J G U G A			21 Conocephalum conicum
G A U U U C U A U G J G G	- U I G G (U) G G I U G I C A U G	- G C C G U U (C) U J A (G U - U*G	G U G G A J G U G A			22 Fossombronia pusilla
G A U U U C U A U G J G G	- U I G G (U) G G I U G I C A U G	- G C C G U U (C) U J A (G U - U*G	G U G G A J G U G A			23 Marchantia polymorpha
G A U U U C U A U G J G G	- U I G G (U) G G I U G I C A U G	- G C C G U U (C) U J A (G U - U*G	G U G G A J G U G A			24 Riccia fluitans
G A U U U C U A U G J G G	- U I G G (U) G G I U G I C A U G	- G C C G U U (C) U J A (G U - U*G	G U G G A J G U G A			25 Scapania nemorea
G A U U U C U A U G J G G	- U I G G (U) G G I U G I C A U G	- G C C G U U (C) U J A (G U - U*G	G U G G A J G U G A			26 Equisetum hyemale
G A U U U C U A U G J G G	- U I G G (U) G G I U G I C A U G	- G C C G U U (C) U J A (G U - U*G	G U G G A J G U G A			27 Equisetum robustum
G A U U U C U A U G J G G	- U I G G (U) G G I U G I C A U G	- G C C G U U (C) U J A (G U - U*G	G U G G A J G U G A			28 Isoetes engelmannii
G A U U U C U A U G J G G	- U I G G (U) G G I U G I C A U G	- G C C G U U (C) U J A (G U - U*G	G U G G A J G U G A			29 Huperzia lucidula
G A U U U C U A U G J G G	- U I G G (U) G G I U G I C A U G	- G C C G U U (C) U J A (G U - U*G	G U G G A J G U G A			30 Lycopodiella inundata
G A U U U C U A U G J G G	- U I G G (U) G G I U G I C A U G	- G C C G U U (C) U J A (G U - U*G	G U G G A J G U G A			31 Lycopodium phlegmaria
G A U U U C U A U G J G G	- U I G G (U) G G I U G I C A U G	- G C C G U U (C) U J A (G U - U*G	G U G G A J G U G A			32 Lycopodium taxifolium
G A U U U C U A U G J G G	- U I G G (U) G G I U G I C A U G	- G C C G U U (C) U J A (G U - U*G	G U G G A J G U G A			33 Lycopodium tristachyum
G A U U U C U A U G J G G	- U I G G (U) G G I U G I C A U G	- G C C G U U (C) U J A (G U - U*G	G U G G A J G U G A			34 Selaginella galleottii
G A U U U C U A U G J G G	- U I G G (U) G G I U G I C A U G	- G C C G U U (C) U J A (G U - U*G	G U G G A J G U G A			35 Selaginella sp.
G A U U U C U A U G J G G	- U I G G (U) G G I U G I C A U G	- G C C G U U (C) U J A (G U - U*G	G U G G A J G U G A			36 Oryza sativa
G A U U U C U A U G J G G	- U I G G (U) G G I U G I C A U G	- G C C G U U (C) U J A (G U - U*G	G U G G A J G U G A			37 Zea mays
G A U U U C U A U G J G G	- U I G G (U) G G I U G I C A U G	- G C C G U U (C) U J A (G U - U*G	G U G G A J G U G A			38 Arabidopsis thaliana
G A U U U C U A U G J G G	- U I G G (U) G G I U G I C A U G	- G C C G U U (C) U J A (G U - U*G	G U G G A J G U G A			39 Glycine max
G A U U U C U A U G J G G	- U I G G (U) G G I U G I C A U G	- G C C G U U (C) U J A (G U - U*G	G U G G A J G U G A			40 Sinapis alba
G A U U U C U A U G J G G	- U I G G (U) G G I U G I C A U G	- G C C G U U (C) U J A (G U - U*G	G U G G A J G U G A			41 Zamia pumila
G A U U U C U A U G J G G	- U I G G (U) G G I U G I C A U G	- G C C G U U (C) U J A (G U - U*G	G U G G A J G U G A			42 Gnetum leyboldii
G A U U U C U A U G J G G	- U I G G (U) G G I U G I C A U G	- G C C G U U (C) U J A (G U - U*G	G U G G A J G U G A			43 Ginkgo biloba
G A U U U C U A U G J G G	- U I G G (U) G G I U G I C A U G	- G C C G U U (C) U J A (G U - U*G	G U G G A J G U G A			44 Pinus luchuensis
G A U U U C U A U G J G G	- U I G G (U) G G I U G I C A U G	- G C C G U U (C) U J A (G U - U*G	G U G G A J G U G A			45 Pinus wallichiana
G A U U U C U A U G J G G	- U I G G (U) G G I U G I C A U G	- G C C G U U (C) U J A (G U - U*G	G U G G A J G U G A			46 Podocarpus nakai
G A U U U C U A U G J G G	- U I G G (U) G G I U G I C A U G	- G C C G U U (C) U J A (G U - U*G	G U G G A J G U G A			47 Taxus mairei
G A U U U C U A U G J G G	- U I G G (U) G G I U G I C A U G	- G C C G U U (C) U J A (G U - U*G	G U G G A J G U G A			48 Adiantum raddianum
G A U U U C U A U G J G G	- U I G G (U) G G I U G I C A U G	- G C C G U U (C) U J A (G U - U*G	G U G G A J G U G A			49 Adiantum raddianum 2
G A U U U C U A U G J G G	- U I G G (U) G G I U G I C A U G	- G C C G U U (C) U J A (G U - U*G	G U G G A J G U G A			50 Ophioglossum petiolatum
G A U U U C U A U G J G G	- U I G G (U) G G I U G I C A U G	- G C C G U U (C) U J A (G U - U*G	G U G G A J G U G A			51 Osmunda cinnamomea
G A U U U C U A U G J G G	- U I G G (U) G G I U G I C A U G	- G C C G U U (C) U J A (G U - U*G	G U G G A J G U G A			52 Salvinia natans
G A U U U C U A U G J G G	- U I G G (U) G G I U G I C A U G	- G C C G U U (C) U J A (G U - U*G	G U G G A J G U G A			53 Psilotum nudum 2
G A U U U C U A U G J G G	- U I G G (U) G G I U G I C A U G	- G C C G U U (C) U J A (G U - U*G	G U G G A J G U G A			54 Chara foetida
G A U U U C U A U G J G G	- U I G G (U) G G I U G I C A U G	- G C C G U U (C) U J A (G U - U*G	G U G G A J G U G A			55 Coleochaete scutata
G A U U U C U A U G J G G	- U I G G (U) G G I U G I C A U G	- G C C G U U (C) U J A (G U - U*G	G U G G A J G U G A			56 Klebsormidium flaccidum
G A U U U C U A U G J G G	- U I G G (U) G G I U G I C A U G	- G C C G U U (C) U J A (G U - U*G	G U G G A J G U G A			57 Nitella flexilis
G A U U U C U A U G J G G	- U I G G (U) G G I U G I C A U G	- G C C G U U (C) U J A (G U - U*G	G U G G A J G U G A			58 Nitella sp.
G A U U U C U A U G J G G	- U I G G (U) G G I U G I C A U G	- G C C G U U (C) U J A (G U - U*G	G U G G A J G U G A			59 Chlamydomonas reinhardtii
G A U U U C U A U G J G G	- U I G G (U) G G I U G I C A U G	- G C C G U U (C) U J A (G U - U*G	G U G G A J G U G A			60 Hydrodictyon reticulatum
G A U U U C U A U G J G G	- U I G G (U) G G I U G I C A U G	- G C C G U U (C) U J A (G U - U*G	G U G G A J G U G A			61 Parietochloris pseudoalveolaris
G A U U U C U A U G J G G	- U I G G (U) G G I U G I C A U G	- G C C G U U (C) U J A (G U - U*G	G U G G A J G U G A			62 Dictyostelium discoideum
G A U U U C U A U G J G G	- U I G G (U) G G I U G I C A U G	- G C C G U U (C) U J A (G U - U*G	G U G G A J G U G A			63 Helix numbering 1
G A U U U C U A U G J G G	- U I G G (U) G G I U G I C A U G	- G C C G U U (C) U J A (G U - U*G	G U G G A J G U G A			64 Pteris vittata
G A U U U C U A U G J G G	- U I G G (U) G G I U G I C A U G	- G C C G U U (C) U J A (G U - U*G	G U G G A J G U G A			65 Megaceros aenigmaticus
G A U U U C U A U G J G G	- U I G G (U) G G I U G I C A U G	- G C C G U U (C) U J A (G U - U*G	G U G G A J G U G A			66 Notophylas breutei
G A U U U C U A U G J G G	- U I G G (U) G G I U G I C A U G	- G C C G U U (C) U J A (G U - U*G	G U G G A J G U G A			67 Physcomitrella patens
G A U U U C U A U G J G G	- U I G G (U) G G I U G I C A U G	- G C C G U U (C) U J A (G U - U*G	G U G G A J G U G A			68 Sphagnum palustre

1560	1570	1580	1590	1600	
[U U U G U C U G C]	- U U A A U U (G C G) A U A A C G I A A	- C I G A G A I C C U U A A A C C U (A) I C U A A (A) U A			1 Saccharomyces cerevisiae
[U U U G U C U G G]	- U U A A U U (C C G) U U A A C G I A A	- C I G A G A I C C U U A A A C C U (I) G I C U A A (A) U A			2 Schizosaccharomyces pombe
[U U U G U C U G G]	- U U A A U U (C C G) U U A A C G I A A	- C I G A G A I C C U U A A A C C U (A) G C C U (I) G I C U A A (A) U A			3 Physcomitrella patens
[U U U G U C U G G]	- U U A A U U (C C G) U U A A C G I A A	- C I G A G A I C C U U A A A C C U (A) G C C U (I) G I C U A A (A) U A			4 Anthoceros agrestis
[U U U G U C U G G]	- U U A A U U (C C G) U U A A C G I A A	- C I G A G A I C C U U A A A C C U (A) G C C U (I) G I C U A A (A) U A			5 Anthoceros laevis
[U U U G U C U G G]	- U U A A U U (C C G) U U A A C G I A A	- C I G A G A I C C U U A A A C C U (A) G C C U (I) G I C U A A (A) U A			6 Pellia epiphylla
[U U U G U C U G G]	- U U A A U U (C C G) U U A A C G I A A	- C I G A G A I C C U U A A A C C U (A) G C C U (I) G I C U A A (A) U A			7 Reboulia hemisphaerica 2
[U U U G U C U G G]	- U U A A U U (C C G) U U A A C G I A A	- C I G A G A I C C U U A A A C C U (A) G C C U (I) G I C U A A (A) U A			8 Riccardia pinguis
[U U U G U C U G G]	- U U A A U U (C C G) U U A A C G I A A	- C I G A G A I C C U U A A A C C U (A) G C C U (I) G I C U A A (A) U A			9 Sphaerocarpos donnellii
[U U U G U C U G G]	- U U A A U U (C C G) U U A A C G I A A	- C I G A G A I C C U U A A A C C U (A) G C C U (I) G I C U A A (A) U A			10 Sphaerocarpos texanus
[U U U G U C U G G]	- U U A A U U (C C G) U U A A C G I A A	- C I G A G A I C C U U A A A C C U (A) G C C U (I) G I C U A A (A) U A			11 Atrichum undulatum
[U U U G U C U G G]	- U U A A U U (C C G) U U A A C G I A A	- C I G A G A I C C U U A A A C C U (A) G C C U (I) G I C U A A (A) U A			12 Bryum argenteum
[U U U G U C U G G]	- U U A A U U (C C G) U U A A C G I A A	- C I G A G A I C C U U A A A C C U (A) G C C U (I) G I C U A A (A) U A			13 Eurhynchium hians
[U U U G U C U G G]	- U U A A U U (C C G) U U A A C G I A A	- C I G A G A I C C U U A A A C C U (A) G C C U (I) G I C U A A (A) U A			14 Funaria hygrometrica
[U U U G U C U G G]	- U U A A U U (C C G) U U A A C G I A A	- C I G A G A I C C U U A A A C C U (A) G C C U (I) G I C U A A (A) U A			15 Leptobryum pyriforme
[U U U G U C U G G]	- U U A A U U (C C G) U U A A C G I A A	- C I G A G A I C C U U A A A C C U (A) G C C U (I) G I C U A A (A) U A			16 Mnium hornum
[U U U G U C U G G]	- U U A A U U (C C G) U U A A C G I A A	- C I G A G A I C C U U A A A C C U (A) G C C U (I) G I C U A A (A) U A			17 Physcomitrium pyriforme
[U U U G U C U G G]	- U U A A U U (C C G) U U A A C G I A A	- C I G A G A I C C U U A A A C C U (A) G C C U (I) G I C U A A (A) U A			18 Polytrichum formosum
[U U U G U C U G G]	- U U A A U U (C C G) U U A A C G I A A	- C I G A G A I C C U U A A A C C U (A) G C C U (I) G I C U A A (A) U A			19 Sphagnum cuspidatum
[U U U G U C U G G]	- U U A A U U (C C G) U U A A C G I A A	- C I G A G A I C C U U A A A C C U (A) G C C U (I) G I C U A A (A) U A			20 Calypogeia arguta
[U U U G U C U G G]	- U U A A U U (C C G) U U A A C G I A A	- C I G A G A I C C U U A A A C C U (A) G C C U (I) G I C U A A (A) U A			21 Conocephalum conicum
[U U U G U C U G G]	- U U A A U U (C C G) U U A A C G I A A	- C I G A G A I C C U U A A A C C U (A) G C C U (I) G I C U A A (A) U A			22 Fossombronia pusilla
[U U U G U C U G G]	- U U A A U U (C C G) U U A A C G I A A	- C I G A G A I C C U U A A A C C U (A) G C C U (I) G I C U A A (A) U A			23 Marchantia polymorpha
[U U U G U C U G G]	- U U A A U U (C C G) U U A A C G I A A	- C I G A G A I C C U U A A A C C U (A) G C C U (I) G I C U A A (A) U A			24 Riccia fluitans
[U U U G U C U G G]	- U U A A U U (C C G) U U A A C G I A A	- C I G A G A I C C U U A A A C C U (A) G C C U (I) G I C U A A (A) U A			25 Scapania nemorea
[U U U G U C U G G]	- U U A A U U (C C G) U U A A C G I A A	- C I G A G A I C C U U A A A C C U (A) G C C U (I) G I C U A A (A) U A			26 Equisetum hyemale
[U U U G U C U G G]	- U U A A U U (C C G) U U A A C G I A A	- C I G A G A I C C U U A A A C C U (A) G C C U (I) G I C U A A (A) U A			27 Equisetum robustum
[U U U G U C U G G]	- U U A A U U (C C G) U U A A C G I A A	- C I G A G A I C C U U A A A C C U (A) G C C U (I) G I C U A A (A) U A			28 Isoetes engelmannii
[U U U G U C U G G]	- U U A A U U (C C G) U U A A C G I A A	- C I G A G A I C C U U A A A C C U (A) G C C U (I) G I C U A A (A) U A			29 Huperzia lucidula
[U U U G U C U G G]	- U U A A U U (C C G) U U A A C G I A A	- C I G A G A I C C U U A A A C C U (A) G C C U (I) G I C U A A (A) U A			30 Lycopodiella inundata
[U U U G U C U G G]	- U U A A U U (C C G) U U A A C G I A A	- C I G A G A I C C U U A A A C C U (A) G C C U (I) G I C U A A (A) U A			31 Lycopodium phlegmaria
[U U U G U C U G G]	- U U A A U U (C C G) U U A A C G I A A	- C I G A G A I C C U U A A A C C U (A) G C C U (I) G I C U A A (A) U A			32 Lycopodium tristachyum
[U U U G U C U G G]	- U U A A U U (C C G) U U A A C G I A A	- C I G A G A I C C U U A A A C C U (A) G C C U (I) G I C U A A (A) U A			33 Lycopodium tristachyum
[U U U G U C U G G]	- U U A A U U (C C G) U U A A C G I A A	- C I G A G A I C C U U A A A C C U (A) G C C U (I) G I C U A A (A) U A			34 Selaginella galieottii
[U U U G U C U G G]	- U U A A U U (C C G) U U A A C G I A A	- C I G A G A I C C U U A A A C C U (A) G C C U (I) G I C U A A (A) U A			35 Selaginella sp.
[U U U G U C U G G]	- U U A A U U (C C G) U U A A C G I A A	- C I G A G A I C C U U A A A C C U (A) G C C U (I) G I C U A A (A) U A			36 Oryza sativa
[U U U G U C U G G]	- U U A A U U (C C G) U U A A C G I A A	- C I G A G A I C C U U A A A C C U (A) G C C U (I) G I C U A A (A) U A			37 Zea mays
[U U U G U C U G G]	- U U A A U U (C C G) U U A A C G I A A	- C I G A G A I C C U U A A A C C U (A) G C C U (I) G I C U A A (A) U A			38 Arabidopsis thaliana
[U U U G U C U G G]	- U U A A U U (C C G) U U A A C G I A A	- C I G A G A I C C U U A A A C C U (A) G C C U (I) G I C U A A (A) U A			39 Glycine max
[U U U G U C U G G]	- U U A A U U (C C G) U U A A C G I A A	- C I G A G A I C C U U A A A C C U (A) G C C U (I) G I C U A A (A) U A			40 Sinapis alba
[U U U G U C U G G]	- U U A A U U (C C G) U U A A C G I A A	- C I G A G A I C C U U A A A C C U (A) G C C U (I) G I C U A A (A) U A			41 Zamia pumila
[U U U G U C U G G]	- U U A A U U (C C G) U U A A C G I A A	- C I G A G A I C C U U A A A C C U (A) G C C U (I) G I C U A A (A) U A			42 Gnetum leyboldii
[U U U G U C U G G]	- U U A A U U (C C G) U U A A C G I A A	- C I G A G A I C C U U A A A C C U (A) G C C U (I) G I C U A A (A) U A			43 Ginkgo biloba
[U U U G U C U G G]	- U U A A U U (C C G) U U A A C G I A A	- C I G A G A I C C U U A A A C C U (A) G C C U (I) G I C U A A (A) U A			44 Pinus luchuensis
[U U U G U C U G G]	- U U A A U U (C C G) U U A A C G I A A	- C I G A G A I C C U U A A A C C U (A) G C C U (I) G I C U A A (A) U A			45 Pinus wallichiana
[U U U G U C U G G]	- U U A A U U (C C G) U U A A C G I A A	- C I G A G A I C C U U A A A C C U (A) G C C U (I) G I C U A A (A) U A			46 Podocarpus nakaii
[U U U G U C U G G]	- U U A A U U (C C G) U U A A C G I A A	- C I G A G A I C C U U A A A C C U (A) G C C U (I) G I C U A A (A) U A			47 Taxus mairei
[U U U G U C U G G]	- U U A A U U (C C G) U U A A C G I A A	- C I G A G A I C C U U A A A C C U (A) G C C U (I) G I C U A A (A) U A			48 Adiantum raddianum
[U U U G U C U G G]	- U U A A U U (C C G) U U A A C G I A A	- C I G A G A I C C U U A A A C C U (A) G C C U (I) G I C U A A (A) U A			49 Adiantum raddianum 2
[U U U G U C U G G]	- U U A A U U (C C G) U U A A C G I A A	- C I G A G A I C C U U A A A C C U (A) G C C U (I) G I C U A A (A) U A			50 Ophioglossum petiolatum
[U U U G U C U G G]	- U U A A U U (C C G) U U A A C G I A A	- C I G A G A I C C U U A A A C C U (A) G C C U (I) G I C U A A (A) U A			51 Osmunda cinnamomea
[U U U G U C U G G]	- U U A A U U (C C G) U U A A C G I A A	- C I G A G A I C C U U A A A C C U (A) G C C U (I) G I C U A A (A) U A			52 Salvinia natans
[U U U G U C U G G]	- U U A A U U (C C G) U U A A C G I A A	- C I G A G A I C C U U A A A C C U (A) G C C U (I) G I C U A A (A) U A			53 Psilotum nudum 2
[U U U G U C U G G]	- U U A A U U (C C G) U U A A C G I A A	- C I G A G A I C C U U A A A C C U (A) G C C U (I) G I C U A A (A) U A			54 Chara foetida
[U U U G U C U G G]	- U U A A U U (C C G) U U A A C G I A A	- C I G A G A I C C U U A A A C C U (A) G C C U (I) G I C U A A (A) U A			55 Coleochaete scutata
[U U U G U C U G G]	- U U A A U U (C C G) U U A A C G I A A	- C I G A G A I C C U U A A A C C U (A) G C C U (I) G I C U A A (A) U A			56 Klebsormidium flaccidum
[U U U G U C U G G]	- U U A A U U (C C G) U U A A C G I A A	- C I G A G A I C C U U A A A C C U (A) G C C U (I) G I C U A A (A) U A			57 Nitella flexilis
[U U U G U C U G G]	- U U A A U U (C C G) U U A A C G I A A	- C I G A G A I C C U U A A A C C U (A) G C C U (I) G I C U A A (A) U A			58 Nitella sp.
[C U U G U C] A I G G]	- U U G A U U (C C) G U U A A C G I A A	- C I G A G A I C C U U A A A C C U (A) G C C U (I) G I C U A A (A) U A			59 Chlamydomonas reinhardtii
[C U U G U C] A I G G]	- U U G A U U (C C) G U U A A C G I A A	- C I G A G A I C C U U A A A C C U (A) G C C U (I) G I C U A A (A) U A			60 Hydrodictyon reticulatum
[C U U G U C] A I G G]	- U U G A U U (C C) G U U A A C G I A A	- C I G A G A I C C U U A A A C C U (A) G C C U (I) G I C U A A (A) U A			61 Parietochloris pseudoalveolaris
[U U U G U C U G G]	- U U A A U U (C C G) U U A A C G I A A	- C I G A G A I C C U U A A A C C U (A) G C C U (I) G I C U A A (A) U A			62 Dictyostelium discoideum
- 40' - - - - 41' - - - - 41' - - - - 39' - - - - 42' - - - - - - - - - -					63 Helix numbering 1
[U U U G U C U G G]	- U U A A U U (C C G) U U A A C G I A A	- C I G A G A I C C U U A A A C C U (A) G C C U (I) G I C U A A (A) U A			64 Pteris vittata
[U U U G U C U G G]	- U U A A U U (C C G) U U A A C G I A A	- C I G A G A I C C U U A A A C C U (A) G C C U (I) G I C U A A (A) U A			65 Megaceros enigmaticus
[U U U G U C U G G]	- U U A A U U (C C G) U U A A C G I A A	- C I G A G A I C C U U A A A C C U (A) G C C U (I) G I C U A A (A) U A			66 Notophyllus breutelii
[U U U G U C U G G]	- U U A A U U (C C G) U U A A C G I A A	- C I G A G A I C C U U A A A C C U (A) G C C U (I) G I C U A A (A) U A			67 Physcomitrella patens
[U U U G U C U G G]	- U U A A U U (C C G) U U A A C G I A A	- C I G A G A I C C U U A A A C C U (A) G C C U (I) G I C U A A (A) U A			68 Sphagnum palustre

DCSE Alignment

	1610	1620	1630	1640	1650	
- G U G G I U G	- C U A G C A					1 Saccharomyces cerevisiae
- G C U G G A	- U C A G C C A U U					2 Schizosaccharomyces pombe
- G U U A I C G	- C G A A G G A					3 Physcomitrella patens
- G U U A I C A	- C G A A G A I					4 Anthoceros agrestis
- G U U A I C A	- C G A A G A A					5 Anthoceros laevis
- G U U A I C A	- C G A A G A I					6 Pellia epiphylla
- C U A I C G C	- G G G G G A I					7 Reboulia hemisphaerica 2
- G U U A I C G	- C G A A G A A					8 Riccardia pinguis
- G C U A I C G	- C G A - G G A I					9 Sphaerocarpos donnellii
- G C U A I C G	- C G A G G C A					10 Sphaerocarpos texanus
- G U U A I C A	- C G A A G A A					11 Atrichum undulatum
- G U U A I C G	- C G A A G G A I					12 Bryum argenteum
- (C) U U G C G	- C I C A A G G A I					13 Eurhynchium hians
- G U U A I C G	- C G A A G G A I					14 Funaria hygrometrica
- G U U A I C G	- C G A A G G A I					15 Leptobryum pyriforme
- G U U A I C G	- C G A A G G A I					16 Mnium hornum
- G U U A I C A	- U G A A G G A I					17 Physcomitrium pyriforme
- G U U A I C G	- C G A A G A A					18 Polytrichum formosum
- G U C I A I C G	- C G A A G G G I					19 Sphagnum cuspidatum
- G U U A I C A	- C G A A G A A					20 Calypogeia arguta
(C) G C U A I C G	- C G G A G U C I					21 Conocephalum conicum
- G U U A I C A	- C G A A G A A					22 Fossombronina pusilla
- G C U A I C G	- C G G G G A I					23 Marchantia polymorpha
- G C U A I C G	- C G G G G G I					24 Riccia fluitans
- G U U A I C A	- C G A A G A I					25 Scapania nemorea
- G U U A I C G	- C G A A G A I					26 Equisetum hyemale
- G U U A I C G	- C G A A G A I					27 Equisetum robustum
- G U (- A I C G A I C G A A G G A I						28 Isoetes engelmannii
- G U U A I C A	- C I G A A G A I					29 Huperzia lucidula
- G U U A I C A	- C I G A A G A I					30 Lycopodiella inundata
- G U U A I C A	- C I G A A G A I					31 Lycopodium phlegmaria
- G U U A I C A	- C I G A A G A I					32 Lycopodium taxifolium
- G U U A I C A	- C I G A A G A I					33 Lycopodium cristatum
- G C U A I C G	- C A G A G G I					34 Selaginella galleottii
- G C U A I C U	- C G G A G G I					35 Selaginella sp.
- G C U A U G	- C G G A G I C					36 Oryza sativa
- G C U A I C G	- U G G A G G I					37 Zea mays
- G C U A I C G	- U G G A G G I					38 Arabidopsis thaliana
- G C U A I C G	- U G G A G G I					39 Glycine max
- G C U A I C G	- U G G A G G I					40 Sinapis alba
- G C U A I C G	- C G G A G G I					41 Zamia pumila
- G C U A I C G	- U G G A G G I					42 Gnetum leyboldii
- G C U A I C G	- C G G A G G I					43 Ginkgo biloba
- G C U A I C G	- C G G A G G I					44 Pinus luchuensis
- G C U A I C G	- C G G A G G I					45 Pinus wallichiana
- G C U A I C A	- C G G A G G I					46 Podocarpus nakaai
- G C U A I C G	- C G G A G G I					47 Taxus mairei
- G U U A I C A	- C I G A A G I G					48 Adiantum raddianum
- G U U A I C A	- C I G A A G I G					49 Adiantum raddianum 2
- G U U A I C G	- C G A A G I C					50 Ophioglossum petiolatum
- G U U A C I A	- C I G A A G I A					51 Osmunda cinnamomea
- G U U A I C G	- C G A A G G I					52 Salvinia natans
- G U U A I C G	- C G A A G G I					53 Psilotum nudum 2
- G C U A I C G	- C G G G G A I					54 Chara foetida
- G U U A I C A	- C A A A G G I					55 Coleochaete scutata
- G U U A I C A	- C G A A G A I					56 Klebsormidium flaccidum
- G C U A I C G	- C G A G G A I					57 Nitella flexilis
- G C U A I C G	- C G A G G A I					58 Nitella sp.
- G U C I A I C G	- A U C G C I					59 Chlamydomonas reinhardtii
- G U C I A I C G	- G U C G C I					60 Hydrodictyon reticulatum
- G U C I A I C G	- G U U G G I					61 Parietochloris pseudoalveolaris
- A G U A I G U	- A U U A U U A G U C I G A U A U I A G A C I G A U A G C U U U U C I U G G G G U U U G G A					62 Dictyostelium discoideum
- 43-						63 Helix numbering 1
- G U U A I C A	- C G A A G I G					64 Pteris vittata
- G U U A I C A	- C G A A G I A A					65 Megaceros enigmaticus
- G U U A I C A	- C G A A G I A A					66 Notophylas breutelii
- G U U A I C G	- C G A A G G A I					67 Physcomitrella patens
- G U C I A I C G	- C I G A A G G I					68 Sphagnum palustre

1660	1670	1680	1690	1700	
.	1 Saccharomyces cerevisiae
.	2 Schizosaccharomyces pombe
.	3 Physcomitrella patens
.	4 Anthoceros agrestis
.	5 Anthoceros laevis
.	6 Pellia epiphylla
.	7 Reboulia hemisphaerica 2
.	8 Riccardia pinguis
.	9 Sphaerocarpos donnellii
.	10 Sphaerocarpos texanus
.	11 Atrichum undulatum
.	12 Bryum argenteum
.	13 Eurhynchium hians
.	14 Funaria hygrometrica
.	15 Leptobryum pyriforme
.	16 Mnium hornum
.	17 Physcomitrium pyriforme
.	18 Polytrichum formosum
.	19 Sphagnum cuspidatum
.	20 Calypogeia arguta
.	21 Conocephalum conicum
.	22 Fossombronina pusilla
.	23 Marchantia polymorpha
.	24 Riccia fluitans
.	25 Scapania nemorea
.	26 Equisetum hyemale
.	27 Equisetum robustum
.	28 Isoetes engelmannii
.	29 Huperzia lucidula
.	30 Lycopodiella inundata
.	31 Lycopodium phlegmaria
.	32 Lycopodium taxifolium
.	33 Lycopodium tristachyum
.	34 Selaginella galleottii
.	35 Selaginella sp.
.	36 Oryza sativa
.	37 Zea mays
.	38 Arabidopsis thaliana
.	39 Glycine max
.	40 Sinapis alba
.	41 Zamia pumila
.	42 Gnetum leyboldii
.	43 Ginkgo biloba
.	44 Pinus luchuensis
.	45 Pinus wallichiana
.	46 Podocarpus nakaii
.	47 Taxus mairei
.	48 Adiantum raddianum
.	49 Adiantum raddianum 2
.	50 Ophioglossum petiolatum
.	51 Osmunda cinnamomea
.	52 Salvinia natans
.	53 Psilotum nudum 2
.	54 Chara foetida
.	55 Coleochaete scutata
.	56 Klebsormidium flaccidum
.	57 Nitella flexilis
.	58 Nitella sp.
.	59 Chlamydomonas reinhardtii
.	60 Hydrodictyon reticulatum
.	61 Parietochloris pseudoalveolaris
.	62 Dictyostelium discoideum
A U G A U U U C G					63 Helix numbering 1
-E43_1'	-E43_2	E43_3	E43_4	E43_5	64 Pteris vittata
					65 Megaceros enigmaticus
					66 Notophylas breutelii
					67 Physcomitrella patens
					68 Sphagnum palustre

1710	1720	1730	1740	1750
				1 Saccharomyces cerevisiae
				2 Schizosaccharomyces pombe
				3 Physcomitrella patens
				4 Anthoceros agrestis
				5 Anthoceros laevis
				6 Pellia epiphylla
				7 Reboulia hemisphaerica 2
				8 Riccardia pinguis
				9 Sphaerocarpos donnelli
				10 Sphaerocarpos texanus
				11 Atrichum undulatum
				12 Bryum argenteum
				13 Eurhynchium hians
				14 Funaria hygrometrica
				15 Leptobryum pyriforme
				16 Mnium hornum
				17 Physcomitrium pyriforme
				18 Polytrichum formosum
				19 Sphagnum cuspidatum
				20 Calypogeia arguta
				21 Conocephalum conicum
				22 Fossombronia pusilla
				23 Marchantia polymorpha
				24 Riccia fluitans
				25 Scapania nemorea
				26 Equisetum hyemale
				27 Equisetum robustum
				28 Isoetes engelmannii
				29 Huperzia lucidula
				30 Lycopodiella inundata
				31 Lycopodium phlegmaria
				32 Lycopodium taxifolium
				33 Lycopodium tristachyum
				34 Selaginella galleottii
				35 Selaginella sp.
				36 Oryza sativa
				37 Zea mays
				38 Arabidopsis thaliana
				39 Glycine max
				40 Sinapis alba
				41 Zamia pumila
				42 Gnetum leyboldii
				43 Ginkgo biloba
				44 Pinus luchuensis
				45 Pinus wallichiana
				46 Podocarpus nakaii
				47 Taxus mairei
				48 Adiantum raddianum
				49 Adiantum raddianum 2
				50 Ophioglossum petiolatum
				51 Osmunda cinnamomea
				52 Salvinia natans
				53 Psilotum nudum 2
				54 Chara foetida
				55 Coleochaete scutata
				56 Klebsormidium flaccidum
				57 Nitella flexilis
				58 Nitella sp.
				59 Chlamydomonas reinhardtii
				60 Hydrodictyon reticulatum
				61 Parietochloris pseudoalveolaris
				62 Dictyostelium discoideum
				63 Helix numbering 1
				64 Pteris vittata
				65 Megaceros aenigmaticus
				66 Notophylas breutelii
				67 Physcomitrella patens
				68 Sphagnum palustre

	1760	1770	1780	1790	1800	
1G	- A(C U - A U C G - G U)U U C A A A -	- [G C C G - A U G G]A A G U(U U G A G	G J C A(A)U)A A C(A)G G U			1 Saccharomyces cerevisiae
1G	- A(C U - A U U G - G C)A U C A A A -	- [G C C G - A U G G]A A G U(U U G A G	G J C A A(U)A A C(A)G G U			2 Schizosaccharomyces pombe
1G	- A(C U - A U C G - G C)G U C U A A -	- [G C C G - A U G G]A A G U(U U G A G	G J C A A(U)A A C(A)G G U			3 Physcomitrella patens
1G	- A(C U - A U(C)G - G C)G U C U A A -	- [G C C (A) - A U G G]A A G U(U U G A G	G J C A A(U)A A C(A)G G U			4 Anthoceros agrestis
1G	- A(C U - A U U G - G C)G U C U A A -	- [G C C G A - A U G G]A A G U(U U G A G	G J C A A(U)A A C(A)G G U			5 Anthoceros laevis
1G	- A(C U - A U U G - G C)G U C U A A -	- [G C C C A - A U G G]A A G U(U U G A G	G J C A A(U)A A C(A)G G U			6 Pellia epiphylla
1G	A C A(C U G U C - - G C)G U C U A A -	- [G C(C)G - A C G G]A A G U(U U G A G	- G J C A A(U)A A C(A)G G U			7 Reboulia hemisphaerica 2
1G	- A(C U - A U U G - G C)G U C U A A -	- [G C C A - A U G G]A A G U(U U G A G	G J C A A(U)A A C(A)G G U			8 Riccardia pinquus
1G	- A(C U - A U C G - G C)G U C U A A -	- [G C C G - A U G G]A A G U(U U G A G	G J C A A(U)A A C(A)G G U			9 Sphaerocarpos donnellii
1G	- A(C U - A U C G - G C)G U C U A A -	- [G C C G - A U G G]A A G U(U U G A G	G J C A A(U)A A C(A)G G -			10 Sphaerocarpos texanus
1G	- A(C U - A U U G - G C)G U C U A A -	- [G C C A - A U G G]A A G U(U U G A G	G J C A A(U)A A C(A)G G U			11 Atrichum undulatum
1G	- A(C U - A U C G - G C)G U C U A A -	- [G C C G - A U G G]A A G U(U U G A G	G J C A A(U)A A C(A)G G U			12 Bryum argenteum
1G	- A(C U - A U C G - G C)G U C U A A -	- [G C C G - A U G G]A A G U(U U G A G	G J C A A(U)A A C(A)G G U			13 Euzhynchium hians
1G	- A(C U - A U C G - G C)G U C U A A -	- [G C C G - A U G G]A A G U(U U G A G	G J C A A(U)A A C(A)G G U			14 Funaria hygrometrica
1G	- A(C U - A U C G - G C)G U C U A A -	- [G C C G - A U G G]A A G U(U U G A G	G J C A A(U)A A C(A)G G U			15 Leptobryum pyriforme
1G	- A(C U - A U C G - G C)G U C U A A -	- [G C C G - A U G G]A A G U(U U G A G	G J C A A(U)A A C(A)G G U			16 Mnium hornum
1G	- A(C U - A U(C)G - G C)G U C U A A -	- [G C C (A) - A U G G]A A G U(U U G A G	G J C A A(U)A A C(A)G G U			17 Physcomitrium pyriforme
1G	- A(C U - A U C G - G C)G U C U A A -	- [G C C G - A U G G]A A G U(U U G A G	G J C A A(U)A A C(A)G G U			18 Polytrichum formosum
1G	- A(C U - A U U G - G C)G U C U A A -	- [G C C G - A U G G]A A G U(U U G A G	G J C A A(U)A A C(A)G G U			19 Sphagnum cuspidatum
1G	- A(C U - A U U G - G C)G U C U A A -	- [G C C A - A U G G]A A G U(U U G A G	G J C A A(U)A A C(A)G G U			20 Calypogeia arguta
1G	- A(C U - G U C G - G C)G U C U A A -	- [G C C G - A C G G]A A G U(U U G A G	G J C A A(U)A A C(A)G G U			21 Conocephalum concium
1G	- A(C U - A U U G - G C)G U C U A A -	- [G C C A - A U G G]A A G U(U U G A G	G J C A A(U)A A C(A)G G U			22 Fossombronina pusilla
1G	- A(C U - A U C G - G C)G U C U A A -	- [G C C G - A U G G]A A G U(U U G A G	G J C A A(U)A A C(A)G G U			23 Marchantia polymorpha
1G	- A(C U - G U C G - G C)G U C U A A -	- [G C C G - A C G G]A A G U(U U G A G	G J C A A(U)A A C(A)G G U			24 Riccia fluitans
1G	- A(C U - A U U G - G C)G U C U A A -	- [G C C A - A U G G]A A G U(U U G A G	G J C A A(U)A A C(A)G G U			25 Scapania nemorea
1G	- A(C U - A U G G - C C)G U C U A A -	- [G G C C - A U G G]A A G U(U U G A G	G J C A A(U)A A C(A)G G U			26 Equisetum hyemale
1G	- A(C U - A U U G - C C)G U C U A A -	- [G G C C - A U G G]A A G U(U U G A G	G J C A A(U)A A C(A)G G U			27 Equisetum robustum
1G	- A(C U - A U U C - G C)G U C U A A -	- [G C G G - A U G G]A A G U(U U G A G	G J C A A(U)A A C(A)G G U			28 Isoetes engelmannii
1G	- A(C U - A U U C - G C)G U C C A A -	- [G C G G - A U G G]A A G U(U U G A G	G J C A A(U)A A C(A)G G U			29 Huperzia lucidula
1G	- A(C U - A U U C - G C)G U C C A A -	- [G C G G - A U G G]A A G U(U U G A G	G J C A A(U)A A C(A)G G U			30 Lycopodiella inundata
1G	- A(C U - A U U C - G C)G U C C A A -	- [G C G G - A U G G]A A G U(U U G A G	G J C A A(U)A A C(A)G G U			31 Lycopodium phlegmaria
1G	- A(C U - A U U C - G C)G U C C A A -	- [G C G G - A U G G]A A G U(U U G A G	G J C A A(U)A A C(A)G G U			32 Lycopodium taxifolium
1G	- A(C U - A U U C - G C)G U C C A A -	- [G C G G - A U G G]A A G U(U U G A G	G J C A A(U)A A C(A)G G U			33 Lycopodium tristachyum
1G	- A(C U - A U C G - G C)G U C C A A -	- [G C G G - A U G G]A A G U(U U G A G	G J C A A(U)A A C(A)G G U			34 Selaginella galleottii
1G	- A(C U - A U U C - G C)G U C C A A -	- [G C G G - A U G G]A A G U(U U G A G	G J C A A(U)A A C(A)G G U			35 Selaginella sp.
1G	- A(C U - (A)U G G - C C)G U U U A A -	- [G G C C - A(C)G]A A G U(U U G A G	G J C A A(U)A A C(A)G G U			36 Oryza sativa
1G	- A(C U - (A)U G G - C C)G U U U A A -	- [G G C C - G(C)G] - A A G U(U U G A G	G J C A A(U)A A C(A)G G U			37 Zea mays
1G	- A(C U - (A)U G G - C C)G U U U A A -	- [G G C C - A(A)G]A A G U(U U G A G	G J C A A(U)A A C(A)G G U			38 Arabidopsis thaliana
1G	- A(C U - (A)U G G - C C)G U U U A A -	- [G G C C - A(C)G]A A G U(U U G A G	G J C A A(U)A A C(A)G G U			39 Glycine max
1G	- A(C U - (A)U G G - C C)G U U U A A -	- [G G C C - A(A)G]A A G U(U U G A G	G J C A A(U)A A C(A)G G U			40 Sinapis alba
1G	- A(C U - A U G G - C C)G U U U A A -	- [G G C C - A U G G]A A G U(U U G A G	G J C A A(U)A A C(A)G G U			41 Zamia pumila
1G	- A(C U - A U G G - C C)G U U U A A -	- [G G C C - A U G G]A A G U(U U G A G	G J C A A(U)A A C(A)G G U			42 Gnetum leyboldii
1G	- A(C U - A U G G - C C)G U U U A A -	- [G G C C - A U G G]A A G U(U U G A G	G J C A A(U)A A C(A)G G U			43 Ginkgo biloba
1G	- A(C U - A U G G - C C)G U U U A A -	- [G G C C - A U G G]A A G U(U U G A G	G J C A A(U)A A C(A)G G U			44 Pinus luchuensis
1G	- A(C U - A U G G - C C)G U U U A A -	- [G G C C - A U G G]A A G U(U U G A G	G J C A A(U)A A C(A)G G U			45 Pinus wallichiana
1G	- A(C U - A U G G - C C)G U U U A A -	- [G G C C - A U G G]A A G U(U U G A G	G J C A A(U)A A C(A)G G U			46 Podocarpus nakaai
1G	- A(C U - A U G G - C C)G U U U A A -	- [G G C C - A U G G]A A G U(U U G A G	G J C A A(U)A A C(A)G G U			47 Taxus mairei
1G	- A(C U - A U G G - C C)G U U U A A -	- [G G C C - A U G G]A A G U(U U G A G	G J C A A(U)A A C(A)G G U			48 Adiantum raddianum
1G	- A(C U - A U G G - C C)G U U U A A -	- [G G C C - A U G G]A A G U(U U G A G	G J C A A(U)A A C(A)G G U			49 Adiantum raddianum 2
1G	- A(C U - G U G G - C C)G U U U A A -	- [G G C C - A C G G]A A G U(U U G A G	G J C A A(U)A A C(A)G G U			50 Ophioglossum petiolatum
1G	- A(C U - A U G G - C C)G U U U A A -	- [G G C C - A U G G]A A G U(U U G A G	G J C A A(U)A A C(A)G G U			51 Osmunda cinnamomea
1G	- A(C U - G U G G - C C)R U C U A A -	- [G G C C - A C G G]A A G U(U U G A G	G J C A A(U)A A C(A)G G U			52 Salvinia natans
1G	- A(C U - A U G G - G(A)C)G A C U A A -	- [G G C C - A U G G]A A G U(U U G A G	G J C A A(U)A A C(A)G G U			53 Psilotum nudum 2
1G	- A(C U - G U U G G(A)C)G A C U A A -	- [G C C A - A C G G]A A G U(U U G A G	G J C A A(U)A A C(A)G G U			54 Chara foetida
1G	- A(C U - A U C G - G C)U U C U A A -	- [G C C G - A U G G]A A G U(U U G A G	G J C A A(U)A A C(A)G G U			55 Coleochaete scutata
1G	- A(C U - A U U(U)G C)G U C U A A C A I G C A -	- [G C C A - A U G G]A A G U(U U G A G	G J C A A(U)A A C(A)G G U			56 Klebsormidium flaccidum
1G	- A(C U - G U U G G(A)C)G A C U A A -	- [G C C A - A U G G]A A G U(U U G A G	G J C A A(U)A A C(A)G G U			57 Nitella flexilis
1G	- A(C U - G U U G G(A)C)G A C U A A -	- [G C C A - A U G G]A A G U(U U G A G	G J C A A(U)A A C(A)G G U			58 Nitella sp.
1G	- A(C U - A U U G - G C)G U U U A A -	- [G C C A - A U G G]A A G U(U U G A G	G J C A A(U)A A C(A)G G U			59 Chlamydomonas reinhardtii
1G	- A(C U - A U U G - G C)G U U U A A -	- [G U G A - A U G G]A A G U(U U G A G	G J C A A(U)A A C(A)G G U			60 Hydrodictyon reticulatum
1G	- A(C U - (C)U C G - G C)G A C U A A -	- [G C C G - A(U)G]A A G U(U U G A G	G J C A A(U)A A C(A)G G U			61 Parietochloris pseudoalveolaris
1G	- A(C U - (A)C C U - G C)C U C A A A -	- [G C A G - G(C)G]A A G U C(C G A G	G J C A A(U)A A C(A)G G U			62 Dictyostelium discoideum
1G	- A(C U - A U G G - C C)G U U U A A -	- [G G C C - A U G G]A A G U(U U G A G	G J C A A(U)A A C(A)G G U			63 Helix numbering 1
1G	- A(C U - A U U U - G C)G U U U A A -	- [G C G A - A U G G]A A G U(U U G A G	G J C A A(U)A A C(A)G G U			64 Pteris vittata
1G	- A(C U - A U U U - G C)G U U U A A -	- [G C G A - A U G G]A A G U(U U G A G	G J C A A(U)A A C(A)G G U			65 Megaceros aenigmaticus
1G	- A(C U - A U U U - G C)G U U U A A -	- [G U G A - A U G G]A A G U(U U G A G	G J C A A(U)A A C(A)G G U			66 Notophylas breuteli
1G	- A(C U - A U C G - G C)G U U U A A -	- [G C C G - A U G G]A A G U(U U G A G	G J C A A(U)A A C(A)G G U			67 Physcomitrella patens
1G	- A(C U - A U U G - G C)G U U U A A -	- [G C C A - A U G G]A A G U(U U G A G	G J C A A(U)A A C(A)G G U			68 Sphaqnum palustre

1810	1820	1830	1840	1850
Ci-U G U G I A U G G C C C U U A G A - U (G I U) U C U G G I G C C I G C I A C G C G (C) G C U A C A C U G A C G G G				1 Saccharomyces cerevisiae
Ci-U G U G I A U G G C C C U U A G A - U (G I U) U C U G G I G C C I G C I A C G C G (C) G C U A C A C U G A C G G G				2 Schizosaccharomyces pombe
Ci-U G U G I A U G G C C C U U A G A - U (G I U) U C U G G I G C C I G C I A C G C G (C) G C U A C A C U G A C G G G				3 Physcomitrella patens
Ci-U G U G I A U G G C C C U U A G A - U (G I U) U C U G G I G C C I G C I A C G C G (C) G C U A C A C U G A C G G G				4 Anthoceros agrestis
Ci-U G U G I A U G G C C C U U A G A - U (G I U) U C U G G I G C C I G C I A C G C G (C) G C U A C A C U G A C G G G				5 Anthoceros laevis
Ci-U G U G I A U G G C C C U U A G A - U (G I U) U C U G G I G C C I G C I A C G C G (C) G C U A C A C U G A C G G G				6 Pellia epiphylla
Ci-U G U G I A U G G C C C U U A G A - U (G I U) U C U G G I G C C I G C I A C G C G (C) G C U A C A C U G A C G G G				7 Reboulia hemisphaerica 2
Ci-U G U G I A U G G C C C U U A G A - U (G I U) U C U G G I G C C I G C I A C G C G (C) G C U A C A C U G A C G G G				8 Riccardia pinguis
Ci-U G U G I A U G G C C C U U A G A - U (G I U) U C U G G I G C C I G C I A C G C G (C) G C U A C A C U G A C G G G				9 Sphaerocarpos donnellii
Ci-U G U G I A U G G C C C U U A G A - U (G I U) U C U G G I G C C I G C I A C G C G (C) G C U A C A C U G A C G G G				10 Sphaerocarpos texanus
Ci-U G U G I A U G G C C C U U A G A - U (G I U) U C U G G I G C C I G C I A C G C G (C) G C U A C A C U G A C G G G				11 Atrichum undulatum
Ci-U G U G I A U G G C C C U U A G A - U (G I U) U C U G G I G C C I G C I A C G C G (C) G C U A C A C U G A C G G G				12 Bryum argenteum
Ci-U G U G I A U G G C C C U U A G A - U (G I U) U C U G G I G C C I G C I A C G C G (C) G C U A C A C U G A C G G G				13 Eurhynchium hians
Ci-U G U G I A U G G C C C U U A G A - U (G I U) U C U G G I G C C I G C I A C G C G (C) G C U A C A C U G A C G G G				14 Funaria hygrometrica
Ci-U G U G I A U G G C C C U U A G A - U (G I U) U C U G G I G C C I G C I A C G C G (C) G C U A C A C U G A C G G G				15 Leptobryum pyriforme
Ci-U G U G I A U G G C C C U U A G A - U (G I U) U C U G G I G C C I G C I A C G C G (C) G C U A C A C U G A C G G G				16 Mnium hornum
Ci-U G U G I A U G G C C C U U A G A - U (G I U) U C U G G I G C C I G C I A C G C G (C) G C U A C A C U G A C G G G				17 Physcomitrium pyriforme
Ci-U G U G I A U G G C C C U U A G A - U (G I U) U C U G G I G C C I G C I A C G C G (C) G C U A C A C U G A C G G G				18 Polytrichum formosum
Ci-U G U G I A U G G C C C U U A G A - U (G I U) U C U G G I G C C I G C I A C G C G (C) G C U A C A C U G A C G G G				19 Sphagnum cuspidatum
Ci-U G U G I A U G G C C C U U A G A - U (G I U) U C U G G I G C C I G C I A C G C G (C) G C U A C A C U G A C G G G				20 Calypogeia arguta
Ci-U G U G I A U G G C C C U U A G A - U (G I U) U C U G G I G C C I G C I A C G C G (C) G C U A C A C U G A C G G G				21 Conocephalum concium
Ci-U G U G I A U G G C C C U U A G A - U (G I U) U C U G G I G C C I G C I A C G C G (C) G C U A C A C U G A C G G G				22 Fossombronina pusilla
Ci-U G U G I A U G G C C C U U A G A - U (G I U) U C U G G I G C C I G C I A C G C G (C) G C U A C A C U G A C G G G				23 Marchantia polymorpha
Ci-U G U G I A U G G C C C U U A G A - U (G I U) U C U G G I G C C I G C I A C G C G (C) G C U A C A C U G A C G G G				24 Riccia fluitans
Ci-U G U G I A U G G C C C U U A G A - U (G I U) U C U G G I G C C I G C I A C G C G (C) G C U A C A C U G A C G G G				25 Scapania nemorea
Ci-U G U G I A U G G C C C U U A G A - U (G I U) U C U G G I G C C I G C I A C G C G (C) G C U A C A C U G A C G G G				26 Equisetum hyemale
Ci-U G U G I A U G G C C C U U A G A - U (G I U) U C U G G I G C C I G C I A C G C G (C) G C U A C A C U G A C G G G				27 Equisetum robustum
Ci-U G U G I A U G G C C C U U A G A - U (G I U) U C U G G I G C C I G C I A C G C G (C) G C U A C A C U G A C G G G				28 Isoetes engelmannii
Ci-U G U G I A U G G C C C U U A G A - U (G I U) U C U G G I G C C I G C I A C G C G (C) G C U A C A C U G A C G G G				29 Huperzia lucidula
Ci-U G U G I A U G G C C C U U A G A - U (G I U) U C U G G I G C C I G C I A C G C G (C) G C U A C A C U G A C G G G				30 Lycopodiella inundata
Ci-U G U G I A U G G C C C U U A G A - U (G I U) U C U G G I G C C I G C I A C G C G (C) G C U A C A C U G A C G G G				31 Lycopodium phlegmaria
Ci-U G U G I A U G G C C C U U A G A - U (G I U) U C U G G I G C C I G C I A C G C G (C) G C U A C A C U G A C G G G				32 Lycopodium taxifolium
Ci-U G U G I A U G G C C C U U A G A - U (G I U) U C U G G I G C C I G C I A C G C G (C) G C U A C A C U G A C G G G				33 Lycopodium tristachyum
Ci-U G U G I A U G G C C C U U A G A - U (G I U) U C U G G I G C C I G C I A C G C G (C) G C U A C A C U G A C G G G				34 Selaginella galleottii
Ci-U G U G I A U G G C C C U U A G A - U (G I U) U C U G G I G C C I G C I A C G C G (C) G C U A C A C U G A C G G G				35 Selaginella sp.
Ci-U G U G I A U G G C C C U U A G A - U (G I U) U C U G G I G C C I G C I A C G C G (C) G C U A C A C U G A C G G G				36 Oryza sativa
Ci-U G U G I A U G G C C C U U A G A - U (G I U) U C U G G I G C C I G C I A C G C G (C) G C U A C A C U G A C G G G				37 Zea mays
Ci-U G U G I A U G G C C C U U A G A - U (G I U) U C U G G I G C C I G C I A C G C G (C) G C U A C A C U G A C G G G				38 Arabidopsis thaliana
Ci-U G U G I A U G G C C C U U A G A - U (G I U) U C U G G I G C C I G C I A C G C G (C) G C U A C A C U G A C G G G				39 Glycine max
Ci-U G U G I A U G G C C C U U A G A - U (G I U) U C U G G I G C C I G C I A C G C G (C) G C U A C A C U G A C G G G				40 Sinapis alba
Ci-U G U G I A U G G C C C U U A G A - U (G I U) U C U G G I G C C I G C I A C G C G (C) G C U A C A C U G A C G G G				41 Zamia pumila
Ci-U G U G I A U G G C C C U U A G A - U (G I U) U C U G G I G C C I G C I A C G C G (C) G C U A C A C U G A C G G G				42 Gnetum leyboldii
Ci-U G U G I A U G G C C C U U A G A - U (G I U) U C U G G I G C C I G C I A C G C G (C) G C U A C A C U G A C G G G				43 Ginkgo biloba
Ci-U G U G I A U G G C C C U U A G A - U (G I U) U C U G G I G C C I G C I A C G C G (C) G C U A C A C U G A C G G G				44 Pinus luchuensis
Ci-U G U G I A U G G C C C U U A G A - U (G I U) U C U G G I G C C I G C I A C G C G (C) G C U A C A C U G A C G G G				45 Pinus wallichiana
Ci-U G U G I A U G G C C C U U A G A - U (G I U) U C U G G I G C C I G C I A C G C G (C) G C U A C A C U G A C G G G				46 Podocarpus nakaii
Ci-U G U G I A U G G C C C U U A G A - U (G I U) U C U G G I G C C I G C I A C G C G (C) G C U A C A C U G A C G G G				47 Taxus mairei
Ci-U G U G I A U G G C C C U U A G A - U (G I U) U C U G G I G C C I G C I A C G C G (C) G C U A C A C U G A C G G G				48 Adiantum raddianum
Ci-U G U G I A U G G C C C U U A G A - U (G I U) U C U G G I G C C I G C I A C G C G (C) G C U A C A C U G A C G G G				49 Adiantum raddianum 2
Ci-U G U G I A U G G C C C U U A G A - U (G I U) U C U G G I G C C I G C I A C G C G (C) G C U A C A C U G A C G G G				50 Ophioglossum peticiatum
Ci-U G U G I A U G G C C C U U A G A - U (G I U) U C U G G I G C C I G C I A C G C G (C) G C U A C A C U G A C G G G				51 Osmunda cinnamomea
Ci-U G U G I A U G G C C C U U A G A - U (G I U) U C U G G I G C C I G C I A C G C G (C) G C U A C A C U G A C G G G				52 Salvinia natans
Ci-U G U G I A U G G C C C U U A G A - U (G I U) U C U G G I G C C I G C I A C G C G (C) G C U A C A C U G A C G G G				53 Psilotum nudum 2
Ci-U G U G I A U G G C C C U U A G A - U (G I U) U C U G G I G C C I G C I A C G C G (C) G C U A C A C U G A C G G G				54 Chara foetida
Ci-U G U G I A U G G C C C U U A G A - U (G I U) U C U G G I G C C I G C I A C G C G (C) G C U A C A C U G A C G G G				55 Coleochaete scutata
Ci-U G U G I A U G G C C C U U A G A - U (G I U) U C U G G I G C C I G C I A C G C G (C) G C U A C A C U G A C G G G				56 Klebsormidium fiacccidum
Ci-U G U G I A U G G C C C U U A G A - U (G I U) U C U G G I G C C I G C I A C G C G (C) G C U A C A C U G A C G G G				57 Nitella flexilis
Ci-U G U G I A U G G C C C U U A G A - U (G I U) U C U G G I G C C I G C I A C G C G (C) G C U A C A C U G A C G G G				58 Nitella sp.
Ci-U G U G I A U G G C C C U U A G A - U (G I U) U C U G G I G C C I G C I A C G C G (C) G C U A C A C U G A C G G G				59 Chlamydomonas reinhardtii
Ci-U G U G I A U G G C C C U U A G A - U (G I U) U C U G G I G C C I G C I A C G C G (C) G C U A C A C U G A C G G G				60 Hydrodictyon reticulatum
Ci-U G U G I A U G G C C C U U A G A - U (G I U) U C U G G I G C C I G C I A C G C G (C) G C U A C A C U G A C G G G				61 Parietochloris pseudoalveolaris
Ci-U G U G I A U G G C C C U U A G A - U (G I U) U C U G G I G C C I G C I A C G C G (C) G C U A C A C U G A C G G G				62 Dictyostelium discoideum
38' - - - - - 36' - - - - - 34' - - - - - 45-				63 Helix numbering 1
Ci-U G U G I A U G G C C C U U A G A - U (G I U) U C U G G I G C C I G C I A C G C G (C) G C U A C A C U G A C G G G				64 Pteris vittata
Ci-U G U G I A U G G C C C U U A G A - U (G I U) U C U G G I G C C I G C I A C G C G (C) G C U A C A C U G A C G G G				65 Megaceros aenigmaticus
Ci-U G U G I A U G G C C C U U A G A - U (G I U) U C U G G I G C C I G C I A C G C G (C) G C U A C A C U G A C G G G				66 Notophylas breutelii
Ci-U G U G I A U G G C C C U U A G A - U (G I U) U C U G G I G C C I G C I A C G C G (C) G C U A C A C U G A C G G G				67 Physcomitrella patens
Ci-U G U G I A U G G C C C U U A G A - U (G I U) U C U G G I G C C I G C I A C G C G (C) G C U A C A C U G A C G G G				68 Sphagnum palustre

DCSE Alignment

	1660	1670	1880	1890	1900	
A G I C A G C I G A G I U						1 Saccharomyces cerevisiae
A G I C C A A C I G A G I U						2 Schizosaccharomyces pombe
A I U U C C A A C I G A G I U						3 Physcomitrella patens
A I U U C C A A C I G A G I U						4 Anthoceros agrestis
A I U U C C A A C I G A G I U						5 Anthoceros laevis
A I U U C C A A C I G A G I U						6 Pellia epiphylla
A I C U C C A A C I G A G I U						7 Reboulia hemisphaerica 2
A I U U C C A A C I G A G I U						8 Riccardia pinguis
A I C U C C A A C I G A G I U						9 Sphaerocarpos donnelli
A I U U C C A A C I G A G I U						10 Sphaerocarpos texanus
A I U U C C A A C I G A G I U						11 Atrichum undulatum
A I U U C C A A C I G A G I U						12 Bryum argenteum
A I U U C C A A C I G A G I U						13 Eurhynchium hians
A I U U C C A A C I G A G I U						14 Funaria hygrometrica
A I U U C C A A C I G A G I U						15 Leptobryum pyriforme
A I U U C C A A C I G A G I U						16 Mnium hornum
A I U U C C A A C I G A G I U						17 Physcomitrium pyriforme
A I U U C C A A C I G A G I U						18 Polytrichum formosum
A I U U C C A A C I G A G I U						19 Sphagnum cuspidatum
A I U U C C A A C I G A G I U						20 Calypogeia arguta
A I G U C C A A C I G A G I U						21 Conocephalum conicum
A I U U C C A A C I G A G I U						22 Fossombronina pusilla
A I C U C C A A C I G A G I U						23 Marchantia polymorpha
A I C U C C A A C I G A G I U						24 Riccia fluitans
A I U U C C A A C I G A G I U						25 Scapania nemorea
A I U U C C A A C I G A G I U						26 Equisetum hyemale
A I U U C C A A C I G A G I U						27 Equisetum robustum
A I U U C C A A C I G A G I U						28 Isoetes engelmannii
A I U U C C A A C I G A G I U						29 Huperzia lucidula
A I U U C C A A C I G A G I U						30 Lycopodiella inundata
A I U U C C A A C I G A G I U						31 Lycopodium phlegmaria
A I U U C C A A C I G A G I U						32 Lycopodium taxifolium
A I U U C C A A C I G A G I U						33 Lycopodium tristachyum
A I U U C C A A C I G A G I U						34 Selaginella galleottii
A I U U C C A A C I G A G I U						35 Selaginella sp.
A I U U C C A A C I G A G I U						36 Oryza sativa
A I U U C C A A C I G A G I U						37 Zea mays
A I U U C C A A C I G A G I U						38 Arabidopsis thaliana
A I U U C C A A C I G A G I U						39 Glycine max
A I U U C C A A C I G A G I U						40 Sinapis alba
A I U U C C A A C I G A G I U						41 Zamia pumila
A I U U C C A A C I G A G I U						42 Gnetum leyboldii
A I U U C C A A C I G A G I U						43 Ginkgo biloba
A I A C C A A C I G A G I U						44 Pinus luchuensis
A I G C C A A C I G A G I U						45 Pinus wallichiana
A I A U C C A A C I G A G I C						46 Podocarpus nakaii
A I A C C A A C I G A G I U						47 Taxus mairei
A I U U C C A A C I G A G I U						48 Adiantum raddianum
A I U U C C A A C I G A G I U						49 Adiantum raddianum 2
A I U U C C A A C I G A G I U						50 Ophioglossum petriclatum
A I U U C C A A C I G A G I U						51 Osmunda cinnamomea
A I U U C C A A C I G A G I U						52 Salvinia natans
A I U U C C A A C I G A G I U						53 Psilotum nudum 2
A I U U C C A A C I G A G I U						54 Chara foetida
A I U U C C A A C I G A G I U						55 Coleochaete scutata
A I U U C C A A C I G A G I U						56 Klebsormidium flaccidum
A I U U C C A A C I G A G I U						57 Nitella flexilis
A I U U C C A A C I G A G I U						58 Nitella sp.
A I U U C C A A C I G A G I C						59 Chlamydomonas reinhardtii
A I U U C C A A C I G A G I C						60 Hydrodictyon reticulatum
A I U U C C A A C I G A G I C						61 Parietochloris pseudoalveolaris
A A A C A A A A A A G G						62 Dictyostelium discoideum
- - - - -						63 Helix numbering 1
A I U U C C A A C I G A G I U						64 Pteris vittata
A I U U C C A A C I G A G I U						65 Megaceros aenigmaticus
A I U U C C A A C I G A G I U						66 Notophylas breutelii
A I U U C C A A C I G A G I U						67 Physcomitrella patens
A I U U C C A A C I G A G I U						68 Sphagnum palustre

	1910	1920	1930	1940	1950	
GA	GA	GA	GA	GA	GA	1 Saccharomyces cerevisiae
GA	GA	GA	GA	GA	GA	2 Schizosaccharomyces pombe
GA	GA	GA	GA	GA	GA	3 Physcomitrella patens
GA	GA	GA	GA	GA	GA	4 Anthoceros agrestis
GA	GA	GA	GA	GA	GA	5 Anthoceros laevis
GA	GA	GA	GA	GA	GA	6 Pellia epiphylla
GA	GA	GA	GA	GA	GA	7 Reboulia hemisphaerica 2
GA	GA	GA	GA	GA	GA	8 Riccardia pinguis
GA	GA	GA	GA	GA	GA	9 Sphaerocarpos donnelli
GA	GA	GA	GA	GA	GA	10 Sphaerocarpos texanus
GA	GA	GA	GA	GA	GA	11 Atrichum undulatum
GA	GA	GA	GA	GA	GA	12 Bryum argenteum
GA	GA	GA	GA	GA	GA	13 Eurhynchium hians
GA	GA	GA	GA	GA	GA	14 Funaria hygrometrica
GA	GA	GA	GA	GA	GA	15 Leptobryum pyriforme
GA	GA	GA	GA	GA	GA	16 Mnium hornum
GA	GA	GA	GA	GA	GA	17 Physcomitrium pyriforme
GA	GA	GA	GA	GA	GA	18 Polytrichum formosum
GA	GA	GA	GA	GA	GA	19 Sphagnum cuspidatum
GA	GA	GA	GA	GA	GA	20 Calypogeia arguta
GA	GA	GA	GA	GA	GA	21 Conocephalum concium
GA	GA	GA	GA	GA	GA	22 Fossombronia pusilla
GA	GA	GA	GA	GA	GA	23 Marchantia polymorpha
GA	GA	GA	GA	GA	GA	24 Riccia fluitans
GA	GA	GA	GA	GA	GA	25 Scapania nemorea
GA	GA	GA	GA	GA	GA	26 Equisetum hyemale
GA	GA	GA	GA	GA	GA	27 Equisetum robustum
GA	GA	GA	GA	GA	GA	28 Isoetes engelmannii
GA	GA	GA	GA	GA	GA	29 Huperzia lucidula
GA	GA	GA	GA	GA	GA	30 Lycopodiella inundata
GA	GA	GA	GA	GA	GA	31 Lycopodium phlegmaria
GA	GA	GA	GA	GA	GA	32 Lycopodium taxifolium
GA	GA	GA	GA	GA	GA	33 Lycopodium tristachyum
GA	GA	GA	GA	GA	GA	34 Selaginella galleottii
GA	GA	GA	GA	GA	GA	35 Selaginella sp.
GA	GA	GA	GA	GA	GA	36 Oryza sativa
GA	GA	GA	GA	GA	GA	37 Zea mays
GA	GA	GA	GA	GA	GA	38 Arabidopsis thaliana
GA	GA	GA	GA	GA	GA	39 Glycine max
GA	GA	GA	GA	GA	GA	40 Sinapis alba
GA	GA	GA	GA	GA	GA	41 Zamia pumila
GA	GA	GA	GA	GA	GA	42 Gnetum leyboldii
GA	GA	GA	GA	GA	GA	43 Ginkgo biloba
GA	GA	GA	GA	GA	GA	44 Pinus luchuensis
GA	GA	GA	GA	GA	GA	45 Pinus wallichiana
GA	GA	GA	GA	GA	GA	46 Podocarpus nakaii
GA	GA	GA	GA	GA	GA	47 Taxus mairei
GA	GA	GA	GA	GA	GA	48 Adiantum raddianum
GA	GA	GA	GA	GA	GA	49 Adiantum raddianum 2
GA	GA	GA	GA	GA	GA	50 Ophioglossum petiolatum
GA	GA	GA	GA	GA	GA	51 Osmunda cinnamomea
GA	GA	GA	GA	GA	GA	52 Salvinia natans
GA	GA	GA	GA	GA	GA	53 Psilotum nudum 2
GA	GA	GA	GA	GA	GA	54 Chara foetida
GA	GA	GA	GA	GA	GA	55 Coleochaete scutata
GA	GA	GA	GA	GA	GA	56 Klebsormidium flaccidum
GA	GA	GA	GA	GA	GA	57 Nitella flexilis
GA	GA	GA	GA	GA	GA	58 Nitella sp.
GA	GA	GA	GA	GA	GA	59 Chlamydomonas reinhardtii
GA	GA	GA	GA	GA	GA	60 Hydrodictyon reticulatum
GA	GA	GA	GA	GA	GA	61 Parietochloris pseudoalveolaris
GA	GA	GA	GA	GA	GA	62 Dictyostelium discoideum
GA	GA	GA	GA	GA	GA	63 Helix numbering 1
GA	GA	GA	GA	GA	GA	64 Pteris vittata
GA	GA	GA	GA	GA	GA	65 Megaceros aenigmaticus
GA	GA	GA	GA	GA	GA	66 Notophylas breutelii
GA	GA	GA	GA	GA	GA	67 Physcomitrella patens
GA	GA	GA	GA	GA	GA	68 Sphagnum palustre

DCSE Alignment

1960	1970	1980	1990	2000
[A G A G C A] U U G U A A U U A U U G C U C U U C A A C G A G G A A U U C C U A G U A [A G C G C A A G				1 Saccharomyces cerevisiae
[A G A G C A] U U G C C A A U U A U U U G C U C U U C A A C G A G G A A U U C C U A G U A [A G C G C A A G				2 Schizosaccharomyces pombe
[A G A U C A] U U G C C A A U U A U U U G A U C U U C A A C G A G G A A U U C C U A G U A [A G C G C G A G				3 Physcomitrella patens
[A G A U C A] U U G C C A A U U A U U U G A U C U U C A A C G A G G A A U U C C U A G U A [A G C G C G A G				4 Anthoceros agrestis
[A G A U C A] U U G C C A A U U A U U U G A U C U U C A A C G A G G A A U U C C U A G U A [A G C G C G A G				5 Anthoceros laevis
[A G A U C A] U U G C C A A U U A U U U G A U C U U C A A C G A G G A A U U C C U A G U A [A G C G C G A G				6 Pellia epiphylla
[A G A U C A] U U G C C A A U U A U U U G A U C U U C A A C G A G G A A U U C C U A G U A [A G C G C G A G				7 Reboulia hemisphaerica 2
[A G A U C A] U U G C C A A U U A U U U G A U C U U C A A C G A G G A A U U C C U A G U A [A G C G C G A G				8 Riccardia pinguis
[A G A - C A] U U G C C A A U U A U U U G (A) U C U U C A A C G A G G A A U U C C U A G U A [A (G) C G C G A G				9 Sphaerocarpos donnelli
[A G A U C A] U U G C C A A U U A U U U G A U C U U C A A C G A G G A A U U C C U A G U A [A G C G C G A G				10 Sphaerocarpos texanus
[A G A U C A] U U G C C A A U U A U U U G A U C U U C A A C G A G G A A U U C C U A G U A [A G C G C G A G				11 Atrichum undulatum
[A G A U C A] U U G C C A A U U A U U U G A U C U U C A A C G A G G A A U U C C U A G U A [A G C G C G A G				12 Bryum argenteum
[A G A U C A] U U G C C A A U U A U U U G A U C U U C A A C G A G G A A U U C C U A G U A [A G C G C G A G				13 Eurhynchium hians
[A G A U C A] U U G C C A A U U A U U U G A U C U U C A A C G A G G A A U U C C U A G U A [A G C G C G A G				14 Funaria hygrometrica
[A G A U C A] U U G C C A A U U A U U U G A U C U U C A A C G A G G A A U U C C U A G U A [A G C G C G A G				15 Leptobryum pyriforme
[A G A U C A] U U G C C A A U U A U U U G A U C U U C A A C G A G G A A U U C C U A G U A [A G C G C G A G				16 Mnium hornum
[A G A U C A] U U G C C A A U U A U U U G A U C U U C A A C G A G G A A U U C C U A G U A [A G C G C G A G				17 Physcomitrium pyriforme
[A G A U C A] U U G C C A A U U A U U U G A U C U U C A A C G A G G A A U U C C U A G U A [A G C G C G A G				18 Polytrichum formosum
[A G A U C A] U U G C C A A U U A U U U G A U C U U C A A C G A G G A A U U C C U A G U A [A G C G C G A G				19 Sphagnum cuspidatum
[A G A U C A] U U G C C A A U U A U U U G A U C U U C A A C G A G G A A U U C C U A G U A [A G C G C G A G				20 Calypogeia arguta
[A G A U C A] U U G C C A A U U A U U U G A U C U U C A A C G A G G A A U U C C U A G U A [A G C G C G A G				21 Conocephalum concinnum
[A G A U C A] U U G C C A A U U A U U U G A U C U U C A A C G A G G A A U U C C U A G U A [A G C G C G A G				22 Fossombronina pusilla
[A G A U C A] U U G C C A A U U A U U U G A U C U U C A A C G A G G A A U U C C U A G U A [A G C G C G A G				23 Marchantia polymorpha
[A G A U C A] U U G C C A A U U A U U U G A U C U U C A A C G A G G A A U U C C U A G U A [A G C G C G A G				24 Riccia fluitans
[A G A U C A] U U G C C A A U U A U U U G A U C U U C A A C G A G G A A U U C C U A G U A [A G C G C G A G				25 Scapania nemorea
[A G A U C A] U U G C C A A U U A U U U G A U C U U C A A C G A G G A A U U C C U A G U A [A G C G C G A G				26 Equisetum hyemale
[A G A U C A] U U G C C A A U U A U U U G A U C U U C A A C G A G G A A U U C C U A G U A [A G C G C G A G				27 Equisetum robustum
[A G A U C A] U U G C C A A U U A U U U G A U C U U C A A C G A G G A A U U C C U A G U A [A G C G C G A G				28 Isoetes engelmannii
[A G A U C A] U U G C C A A U U A U U U G A U C U U C A A C G A G G A A U U C C U A G U A [A G C G C G A G				29 Huperzia lucidula
[A G A U C A] U U G C C A A U U A U U U G A U C U U C A A C G A G G A A U U C C U A G U A [A G C G C G A G				30 Lycopodiella inundata
[A G A U C A] U U G C C A A U U A U U U G A U C U U C A A C G A G G A A U U C C U A G U A [A G C G C G A G				31 Lycopodium phlegmaria
[A G A U C A] U U G C C A A U U A U U U G A U C U U C A A C G A G G A A U U C C U A G U A [A G C G C G A G				32 Lycopodium taxifolium
[A G A U C A] U U G C C A A U U A U U U G A U C U U C A A C G A G G A A U U C C U A G U A [A G C G C G A G				33 Lycopodium tristachyum
[A G A U C A] U U G C C A A U U A U U U G A U C U U C A A C G A G G A A U U C C U A G U A [A G C G C G A G				34 Selaginella galleottii
[A G A U C A] U U G C C A A U U A U U U G A U C U U C A A C G A G G A A U U C C U A G U A [A G C G C G A G				35 Selaginella sp.
[A G A U C A] U U G C C A A U U A U U U G G U C U U C A A C G A G G A A U U C C U A G U A [A G C G C G A G				36 Oryza sativa
[A G A U C A] U U G C C A A U U A U U U G G U C U U C A A C G A G G A A U U C C U A G U A [A G C G C G A G				37 Zea mays
[A G A U C A] U U G C C A A U U A U U U G G U C U U C A A C G A G G A A U U C C U A G U A [A G C G C G A G				38 Arabidopsis thaliana
[A G A U C A] U U G C C A A U U A U U U G G U C U U C A A C G A G G A A U U C C U A G U A [A G C G C G A G				39 Glycine max
[A G A U C A] U U G C C A A U U A U U U G G U C U U C A A C G A G G A A U U C C U A G U A [A G C G C G A G				40 Sinapis alba
[A G A U C A] U U G C C A A U U A U U U G G U C U U C A A C G A G G A A U U C C U A G U A [A G C G C G A G				41 Zamia pumila
[A G A U C] U U U G C C A A U U A U U C [G G U C U U C A A C G A G G A A U U C C U A G U A [A G C G C G A G				42 Gnetum boldii
[A G A U C A] U U G C C A A U U A U U U G A U C U U C A A C G A G G A A U U C C U A G U A [A G C G C G A G				43 Ginkgo biloba
[A G A (C) A] U U G C C A A U U A U U U G (A) U C U U C A A C G A G G A A U U C C U A G U A [A G C G C G A G				44 Pinus luchuensis
[A G A C C A] U U G C C A A U U A U U U G G U C U U C A A C G A G G A A U U C C U A G U A [A G C G C G (U) G				45 Pinus wallichiana
[A G A (C) A] U U G C C A A U U A U U U G (A) U C U U C A A C G A G G A A U U C C U A G U A [A G C G C G A G				46 Podocarpus naki
[A G A (C) A] U U G C C A A U U A U U U G (A) U C U U C A A C G A G G A A U U C C U A G U A [A G C G C G A G				47 Taxus mairei
[G A U C A] U U G C C A A U U A U U U G A U C U U C A A C G A G G A A U U C C U A G U A [A G C G C (G) A G				48 Adiantum raddianum
[A G A U C A] U U G C C A A U U A U U U G A U C U U C A A C G A G G A A U U C C U A G U A [A G C G C G A G				49 Adiantum raddianum 2
[A G A U C A] U U G C C A A U U A U U U G A U C U U C A A C G A G G A A U U C C U A G U A [A G C G C G A G				50 Ophioglossum petiolatum
[A G A U C A] U U G C C A A U U A U U U G A U C U U C A A C G A G G A A U U C C U A G U A [A G C G C G A G				51 Osmunda cinnamomea
[A G A U C A] U U G C C A A U U A U U U G A U C U U C A A C G A G G A A U U C C U A G U A [A G C G C G A G				52 Salvinia natans
[A G A U U A] U U G C C A A U U A U U U G A U C U U C A A C G A G G A A U U C C U A G U A [A G C G C G A G				53 Psilotum nudum 2
[A G A U U A] U U G C C A A U U A U U U G A U C U U C A A C G A G G A A U U C C U A G U A [A G C G C G A G				54 Chara foetida
[A G A U U A] U U G C C A A U U A U U U G A U C U U C A A C G A G G A A U U C C U A G U A [A G C G C G A G				55 Coleochaete scutata
[A G A U U A] U U G C C A A U U A U U U G A U C U U C A A C G A G G A A U U C C U A G U A [A G C G C G A G				56 Klebsormidium flaccidum
[A G A U U A] U U G C C A A U U A U U U G A U C U U C A A C G A G G A A U U C C U A G U A [A G C G C G A G				57 Nitella flexilis
[A G A U U A] U U G C C A A U U A U U U G A U C U U C A A C G A G G A A U U C C U A G U A [A G C G C G A G				58 Nitella sp.
[A G A U U A] U U G C C A A U U A U U U G A U C U U C A A C G A G G A A U U C C U A G U A [A G C G C G A G				59 Chlamydomonas reinhardtii
[A G A U U A] U U G C C A A U U A U U U G A U C U U C A A C G A G G A A U U C C U A G U A [A G C G C G A G				60 Hydrodictyon reticulatum
[A G A U U A] U U G C C A A U U A U U U G A U C U U C A A C G A G G A A U U C C U A G U A [A G C G C G A G				61 Parietochloris pseudoalveolaris
[U G A U C] U U U G C C A A U U A U U U G A U C U U C A A C G A G G A A U U C C U A G U A [A G C G C U A A G]				62 Dictyostelium discoideum
- 47 - - - - - 47 - - - - - 33 - - - - - 48 - - -				63 Helix numbering 1
[A G A U C A] U U G C C A A U U A U U U G A U C U U C A A C G A G G A A U U C C U A G U A [A G C G C G A G				64 Pteris vittata
[A G A U C A] U U G C C A A U U A U U U G A U C U U C A A C G A G G A A U U C C U A G U A [A G C G C G A G				65 Megaceros aenigmaticus
[A G A U C A] U U G C C A A U U A U U U G A U C U U C A A C G A G G A A U U C C U A G U A [A G C G C G A G				66 Notophylas breutelii
[A G A U C A] U U G C C A A U U A U U U G A U C U U C A A C G A G G A A U U C C U A G U A [A G C G C G A G				67 Physcomitrella patens
[A G A U C A] U U G C C A A U U A U U U G A U C U U C A A C G A G G A A U U C C U A G U A [A G C G C G A G				68 Sphagnum palustre

DCSE Alignment

	2010	2020	2030	2040	2050																									
U C	A	U	C	A	I	G	C	U	U	G	C	G	U	U	G	A	-	C	A	I	C	A	C	C	G	C	C	G	1	<i>Saccharomyces cerevisiae</i>
U C	A	U	C	A	I	G	C	U	U	G	C	G	U	U	G	A	-	C	A	I	C	A	C	C	G	C	C	G	2	<i>Schizosaccharomyces pombe</i>
U C	A	U	C	A	I	G	C	U	U	G	C	G	U	U	G	A	-	C	A	I	C	A	C	C	G	C	C	G	3	<i>Physcomitrella patens</i>
U C	A	U	C	A	I	G	C	U	U	G	C	G	U	U	G	A	-	C	A	I	C	A	C	C	G	C	C	G	4	<i>Anthoceros agrestis</i>
U C	A	U	C	A	I	G	C	U	U	G	C	G	U	U	G	A	-	C	A	I	C	A	C	C	G	C	C	G	5	<i>Anthoceros laevis</i>
U C	A	U	C	A	I	G	C	U	U	G	C	G	U	U	G	A	-	C	A	I	C	A	C	C	G	C	C	G	6	<i>Pellia epiphylla</i>
U C	A	U	C	A	I	G	C	U	U	G	C	G	U	U	G	A	-	C	A	I	C	A	C	C	G	C	C	G	7	<i>Reboulia hemisphaerica 2</i>
U C	A	U	C	A	I	G	C	U	U	G	C	G	U	U	G	A	-	C	A	I	C	A	C	C	G	C	C	G	8	<i>Riccardia pinquius</i>
U C	A	U	C	A	I	G	C	U	U	G	C	G	U	U	G	A	-	C	A	I	C	A	C	C	G	C	C	G	9	<i>Sphaerocarpos donnellii</i>
U C	A	U	C	A	I	G	C	U	U	G	C	G	U	U	G	A	-	C	A	I	C	A	C	C	G	C	C	G	10	<i>Sphaerocarpos texanus</i>
U C	A	U	C	A	I	G	C	U	U	G	C	G	U	U	G	A	-	C	A	I	C	A	C	C	G	C	C	G	11	<i>Atrichum undulatum</i>
U C	A	U	C	A	I	G	C	U	U	G	C	G	U	U	G	A	-	C	A	I	C	A	C	C	G	C	C	G	12	<i>Bryum argenteum</i>
U C	A	U	C	A	I	G	C	U	U	G	C	G	U	U	G	A	-	C	A	I	C	A	C	C	G	C	C	G	13	<i>Eurhynchium hians</i>
U C	A	U	C	A	I	G	C	U	U	G	C	G	U	U	G	A	-	C	A	I	C	A	C	C	G	C	C	G	14	<i>Funaria hygrometrica</i>
U C	A	U	C	A	I	G	C	U	U	G	C	G	U	U	G	A	-	C	A	I	C	A	C	C	G	C	C	G	15	<i>Leptobryum pyriforme</i>
U C	A	U	C	A	I	G	C	U	U	G	C	G	U	U	G	A	-	C	A	I	C	A	C	C	G	C	C	G	16	<i>Mnium hornum</i>
U C	A	U	C	A	I	G	C	U	U	G	C	G	U	U	G	A	-	C	A	I	C	A	C	C	G	C	C	G	17	<i>Physcomitrium pyriforme</i>
U C	A	U	C	A	I	G	C	U	U	G	C	G	U	U	G	A	-	C	A	I	C	A	C	C	G	C	C	G	18	<i>Polytrichum formosum</i>
U C	A	U	C	A	I	G	C	U	U	G	C	G	U	U	G	A	-	C	A	I	C	A	C	C	G	C	C	G	19	<i>Sphagnum cuspidatum</i>
U C	A	U	C	A	I	G	C	U	U	G	C	G	U	U	G	A	-	C	A	I	C	A	C	C	G	C	C	G	20	<i>Calyptogelea arguta</i>
U C	A	U	C	A	I	G	C	U	U	G	C	G	U	U	G	A	-	C	A	I	C	A	C	C	G	C	C	G	21	<i>Conocephalum conicum</i>
U C	A	U	C	A	I	G	C	U	U	G	C	G	U	U	G	A	-	C	A	I	C	A	C	C	G	C	C	G	22	<i>Fossombronina pusilla</i>
U C	A	U	C	A	I	G	C	U	U	G	C	G	U	U	G	A	-	C	A	I	C	A	C	C	G	C	C	G	23	<i>Marchantia polymorpha</i>
U C	A	U	C	A	I	G	C	U	U	G	C	G	U	U	G	A	-	C	A	I	C	A	C	C	G	C	C	G	24	<i>Riccia fluitans</i>
U C	A	U	C	A	I	G	C	U	U	G	C	G	U	U	G	A	-	C	A	I	C	A	C	C	G	C	C	G	25	<i>Scapania nemorea</i>
U C	A	U	C	A	I	G	C	U	U	G	C	G	U	U	G	A	-	C	A	I	C	A	C	C	G	C	C	G	26	<i>Equisetum hyemale</i>
U C	A	U	C	A	I	G	C	U	U	G	C	G	U	U	G	A	-	C	A	I	C	A	C	C	G	C	C	G	27	<i>Equisetum robustum</i>
U C	A	U	C	A	I	G	C	U	U	G	C	G	U	U	G	A	-	C	A	I	C	A	C	C	G	C	C	G	28	<i>Isoetes engelmannii</i>
U C	A	U	C	A	I	G	C	U	U	G	C	G	U	U	G	A	-	C	A	I	C	A	C	C	G	C	C	G	29	<i>Huperzia lucidula</i>
U C	A	U	C	A	I	G	C	U	U	G	C	G	U	U	G	A	-	C	A	I	C	A	C	C	G	C	C	G	30	<i>Lycopodiella inundata</i>
U C	A	U	C	A	I	G	C	U	U	G	C	G	U	U	G	A	-	C	A	I	C	A	C	C	G	C	C	G	31	<i>Lycopodium phlegmaria</i>
U C	A	U	C	A	I	G	C	U	U	G	C	G	U	U	G	A	-	C	A	I	C	A	C	C	G	C	C	G	32	<i>Lycopodium taxifolium</i>
U C	A	U	C	A	I	G	C	U	U	G	C	G	U	U	G	A	-	C	A	I	C	A	C	C	G	C	C	G	33	<i>Lycopodium tristachyum</i>
U C	A	U	C	A	I	G	C	U	U	G	C	G	U	U	G	A	-	C	A	I	C	A	C	C	G	C	C	G	34	<i>Selaginella galleottii</i>
U C	A	U	C	A	I	G	C	U	U	G	C	G	U	U	G	A	-	C	A	I	C	A	C	C	G	C	C	G	35	<i>Selaginella sp.</i>
U C	A	U	C	A	I	G	C	U	U	G	C	G	U	U	G	A	-	C	A	I	C	A	C	C	G	C	C	G	36	<i>Oryza sativa</i>
U C	A	U	C	A	I	G	C	U	U	G	C	G	U	U	G	A	-	C	A	I	C	A	C	C	G	C	C	G	37	<i>Zea mays</i>
U C	A	U	C	A	I	G	C	U	U	G	C	G	U	U	G	A	-	C	A	I	C	A	C	C	G	C	C	G	38	<i>Arabidopsis thaliana</i>
U C	A	U	C	A	I	G	C	U	U	G	C	G	U	U	G	A	-	C	A	I	C	A	C	C	G	C	C	G	39	<i>Glycine max</i>
U C	A	U	C	A	I	G	C	U	U	G	C	G	U	U	G	A	-	C	A	I	C	A	C	C	G	C	C	G	40	<i>Sinapis alba</i>
U C	A	U	C	A	I	G	C	U	U	G	C	G	U	U	G	A	-	C	A	I	C	A	C	C	G	C	C	G	41	<i>Zamia pumila</i>
U C	A	U	C	A	I	G	C	U	U	G	C	G	U	U	G	A	-	C	A	I	C	A	C	C	G	C	C	G	42	<i>Gnetum leyboldii</i>
U C	A	U	C	A	I	G	C	U	U	G	C	G	U	U	G	A	-	C	A	I	C	A	C	C	G	C	C	G	43	<i>Ginkgo biloba</i>
U C	A	U	C	A	I	G	C	U	U	G	C	G	U	U	G	A	-	C	A	I	C	A	C	C	G	C	C	G	44	<i>Pinus luchuensis</i>
U C	A	U	C	A	I	G	C	U	U	G	C	G	U	U	G	A	-	C	A	I	C	A	C	C	G	C	C	G	45	<i>Pinus wallichiana</i>
U C	A	U	C	A	I	G	C	U	U	G	C	G	U	U	G	A	-	C	A	I	C	A	C	C	G	C	C	G	46	<i>Podocarpus nakaii</i>
U C	A	U	C	A	I	G	C	U	U	G	C	G	U	U	G	A	-	C	A	I	C	A	C	C	G	C	C	G	47	<i>Taxus mairei</i>
U C	A	U	C	A	I	G	C	U	U	G	C	G	U	U	G	A	-	C	A	I	C	A	C	C	G	C	C	G	48	<i>Adiantum raddianum</i>
U C	A	U	C	A	I	G	C	U	U	G	C	G	U	U	G	A	-	C	A	I	C	A	C	C	G	C	C	G	49	<i>Adiantum raddianum 2</i>
U C	A	U	C	A	I	G	C	U	U	G	C	G	U	U	G	A	-	C	A	I	C	A	C	C	G	C	C	G	50	<i>Ophioglossum petiolatum</i>
U C	A	U	C	A	I	G	C	U	U	G	C	G	U	U	G	A	-	C	A	I	C	A	C	C	G	C	C	G	51	<i>Osmunda cinnamomea</i>
U C	A	U	C	A	I	G	C	U	U	G	C	G	U	U	G	A	-	C	A	I	C	A	C	C	G	C	C	G	52	<i>Salvinia natans</i>
U C	A	U	C	A	I	G	C	U	U	G	C	G	U	U	G	A	-	C	A	I	C	A	C	C	G	C	C	G	53	<i>Psilotum nudum 2</i>
U C	A	U	C	A	I	G	C	U	U	G	C	G	U	U	G	A	-	C	A	I	C	A	C	C	G	C	C	G	54	<i>Chara foetida</i>
U C	A	U	C	A	I	G	C	U	U	G	C	G	U	U	G	A	-	C	A	I	C	A	C	C	G	C	C	G	55	<i>Coleochaete scutata</i>
U C	A	U	C	A	I	G	C	U	U	G	C	G	U	U	G	A	-	C	A	I	C	A	C	C	G	C	C	G	56	<i>Klebsormidium flaccidum</i>
U C	A	U	C	A	I	G	C	U	U	G	C	G	U	U	G	A	-	C	A	I	C	A	C	C	G	C	C	G	57	<i>Nitella flexilis</i>
U C	A	U	C	A	I	G	C	U	U	G	C	G	U	U	G	A	-	C	A	I	C	A	C	C	G	C	C	G	58	<i>Nitella sp.</i>
U C	A	U	C	A	I	G	C	U	U	G	C	G	U	U	G	A	-	C	A	I	C	A	C	C	G	C	C	G	59	<i>Chlamydomonas reinhardtii</i>
U C	A	U	C	A	I	G	C	U	U	G	C	G	U	U	G	A	-	C	A	I	C	A	C	C	G	C	C	G	60	<i>Hydrodictyon reticulatum</i>
U C	A	U	C	A	I	G	C	U	U	G	C	G	U	U	G	A	-	C	A	I	C	A	C	C	G	C	C	G	61	<i>Parietochloris pseudoaerolaris</i>
U C	A	U	C	A	I	G	C	U	U	G	C	G	U	U	G	A	-	C	A	I	C	A	C	C	G	C	C	G	62	<i>Dictyostelium discoideum</i>
U C	A	U	C	A	I	G	C	U	U	G	C	G	U	U	G	A	-	C	A	I	C	A	C	C	G	C	C	G	63	<i>Helix numbering 1</i>
U C	A	U	C	A	I	G	C	U	U	G	C	G	U	U	G	A	-	C	A	I	C	A	C	C	G	C	C	G	64	<i>Pteris vittata</i>
U C	A	U	C	A	I	G	C	U	U	G	C	G	U	U	G	A	-	C	A	I	C	A	C	C	G	C	C	G	65	<i>Megaceros aenigmaticus</i>
U C	A	U	C	A	I	G	C	U	U	G	C	G	U	U	G	A	-	C	A	I	C	A	C	C	G	C	C	G	66	<i>Notophylas breutelu</i>
U C	A	U	C	A																										

DCSE Alignment

2060		2070		2080		2090		2100			
U	C	G	C	U	A	G	U	A	C	1	<i>Saccharomyces cerevisiae</i>
U	C	G	C	U	A	C	U	A	C	2	<i>Schizosaccharomyces pombe</i>
U	C	G	C	U	A	C	U	A	C	3	<i>Physcomitrella patens</i>
U	C	G	C	U	A	C	U	A	C	4	<i>Anthoceros agrestis</i>
U	C	G	C	U	A	C	U	A	C	5	<i>Anthoceros laevis</i>
U	C	G	C	U	A	C	U	A	C	6	<i>Pellia epiphylla</i>
U	C	G	C	U	A	C	U	A	C	7	<i>Reboulia hemisphaerica</i> 2
U	C	G	C	U	A	C	U	A	C	8	<i>Riccardia pinguis</i>
U	C	G	C	U	A	C	U	A	C	9	<i>Sphaerocarpos donnelli</i>
U	C	G	C	U	A	C	U	A	C	10	<i>Sphaerocarpos texanus</i>
U	C	G	C	U	A	C	U	A	C	11	<i>Atrichum undulatum</i>
U	C	G	C	U	A	C	U	A	C	12	<i>Bryum argenteum</i>
U	C	G	C	U	A	C	U	A	C	13	<i>Eurhynchium hians</i>
U	C	G	C	U	A	C	U	A	C	14	<i>Funaria hygrometrica</i>
U	C	G	C	U	A	C	U	A	C	15	<i>Leptobryum pyriforme</i>
U	C	G	C	U	A	C	U	A	C	16	<i>Mnium hornum</i>
U	C	G	C	U	A	C	U	A	C	17	<i>Physcomitrium pyriforme</i>
U	C	G	C	U	A	C	U	A	C	18	<i>Polytrichum formosum</i>
U	C	G	C	U	A	C	U	A	C	19	<i>Sphagnum cuspidatum</i>
U	C	G	C	U	A	C	U	A	C	20	<i>Calypogeia arguta</i>
U	C	G	C	U	A	C	U	A	C	21	<i>Conocephalum concicum</i>
U	C	G	C	U	A	C	U	A	C	22	<i>Fossombronia pusilla</i>
U	C	G	C	U	A	C	U	A	C	23	<i>Marchantia polymorpha</i>
U	C	G	C	U	A	C	U	A	C	24	<i>Riccia fluitans</i>
U	C	G	C	U	A	C	U	A	C	25	<i>Scapania nemorea</i>
U	C	G	C	U	A	C	U	A	C	26	<i>Equisetum hyemale</i>
U	C	G	C	U	A	C	U	A	C	27	<i>Equisetum robustum</i>
U	C	G	C	U	A	C	U	A	C	28	<i>Isoetes engelmannii</i>
U	C	G	C	U	A	C	U	A	C	29	<i>Huperzia lucidula</i>
U	C	G	C	U	A	C	U	A	C	30	<i>Lycopodiella inundata</i>
U	C	G	C	U	A	C	U	A	C	31	<i>Lycopodium phlegmaria</i>
U	C	G	C	U	A	C	U	A	C	32	<i>Lycopodium taxifolium</i>
U	C	G	C	U	A	C	U	A	C	33	<i>Lycopodium tristachyum</i>
U	C	G	C	U	A	C	U	A	C	34	<i>Selaginella galleottii</i>
U	C	G	C	U	A	C	U	A	C	35	<i>Selaginella</i> sp.
U	C	G	C	U	A	C	U	A	C	36	<i>Oryza sativa</i>
U	C	G	C	U	A	C	U	A	C	37	<i>Zea mays</i>
U	C	G	C	U	A	C	U	A	C	38	<i>Arabidopsis thaliana</i>
U	C	G	C	U	A	C	U	A	C	39	<i>Glycine max</i>
U	C	G	C	U	A	C	U	A	C	40	<i>Sinapis alba</i>
U	C	G	C	U	A	C	U	A	C	41	<i>Zamia pumila</i>
U	C	G	C	U	A	C	U	A	C	42	<i>Gnetum leyboldii</i>
U	C	G	C	U	A	C	U	A	C	43	<i>Ginkgo biloba</i>
U	C	G	C	U	A	C	U	A	C	44	<i>Pinus luchuensis</i>
U	C	G	C	U	A	C	U	A	C	45	<i>Pinus wallichiana</i>
U	C	G	C	U	A	C	U	A	C	46	<i>Podocarpus nakaii</i>
U	C	G	C	U	A	C	U	A	C	47	<i>Taxus mairei</i>
U	C	G	C	U	A	C	U	A	C	48	<i>Adiantum raddianum</i>
U	C	G	C	U	A	C	U	A	C	49	<i>Adiantum raddianum</i> 2
U	C	G	C	U	A	C	U	A	C	50	<i>Ophioglossum petiolatum</i>
U	C	G	C	U	A	C	U	A	C	51	<i>Osmunda cinnamomea</i>
U	C	G	C	U	A	C	U	A	C	52	<i>Salvinia natans</i>
U	C	G	C	U	A	C	U	A	C	53	<i>Psilotum nudum</i> 2
U	C	G	C	U	A	C	U	A	C	54	<i>Chara foetida</i>
U	C	G	C	U	A	C	U	A	C	55	<i>Coleochaete scutata</i>
U	C	G	C	U	A	C	U	A	C	56	<i>Klebsormidium flaccidum</i>
U	C	G	C	U	A	C	U	A	C	57	<i>Nitella flexilis</i>
U	C	G	C	U	A	C	U	A	C	58	<i>Nitella</i> sp.
U	C	G	C	U	A	C	U	A	C	59	<i>Chlamydomonas reinhardtii</i>
U	C	G	C	U	A	C	U	A	C	60	<i>Hydrodictyon reticulatum</i>
U	C	G	C	U	A	C	U	A	C	61	<i>Parietochloris pseudoalveolaris</i>
U	C	G	C	U	A	C	U	A	C	62	<i>Dictyostelium discoideum</i>
U	C	G	C	U	A	C	U	A	C	63	<i>Helix numbering 1</i>
U	C	G	C	U	A	C	U	A	C	64	<i>Pteris vittata</i>
U	C	G	C	U	A	C	U	A	C	65	<i>Megaceros aenigmaticus</i>
U	C	G	C	U	A	C	U	A	C	66	<i>Notophylax breutelii</i>
U	C	G	C	U	A	C	U	A	C	67	<i>Physcomitrella patens</i>
U	C	G	C	U	A	C	U	A	C	68	<i>Sphagnum palustre</i>

2160	2170	2180	2190	2200
G(A)C(A)A(A)C(U)U(G)G(U)C(A)U(U)U(A)G(A)G(A)A(C)U(A)A(A)G(U)C(G)U(-)A(A)C(A)A(G)G(U)U(U)C(-)				1 Saccharomyces cerevisiae
G(A)C(A)A(A)C(U)U(G)G(U)C(A)U(U)U(A)G(A)G(A)A(G)U(A)A(A)G(U)C(G)U(-)A(A)C(A)A(G)G(U)U(U)C(-)				2 Schizosaccharomyces pombe
JA(U)U(A)A(A)C(C)U(U)U(A)U(C)A(U)U(U)U(A)G(A)G(A)A(G)G(A)G(A)A(G)U(C)G(U)U(U)C(-)A(A)C(A)A(G)G(U)U(U)C(-)				3 Physcomitrella patens
JA(U)U(A)A(A)C(C)U(U)U(A)U(C)A(U)U(U)U(A)G(A)G(A)A(G)G(A)G(A)A(G)U(C)G(U)U(U)C(-)A(A)C(A)A(G)G(U)U(U)C(-)				4 Anthoceros agrestis
JA(U)U(A)A(A)C(C)U(U)U(A)U(C)A(U)U(U)U(A)G(A)G(A)A(G)G(A)G(A)A(G)U(C)G(U)U(U)C(-)A(A)C(A)A(G)G(U)U(U)C(-)				5 Anthoceros laevis
JA(U)U(A)A(A)C(C)U(U)U(A)U(C)A(U)U(U)U(A)G(A)G(A)A(G)G(A)G(A)A(G)U(C)G(U)U(U)C(-)A(A)C(A)A(G)G(U)U(U)C(-)				6 Pellia epiphylla
U(U)U(A)A(A)C(C)U(U)U(A)U(C)A(U)U(U)U(A)G(A)G(A)A(G)G(A)G(A)A(G)U(C)G(U)U(U)C(-)A(A)C(A)A(G)G(U)U(U)C(-)				7 Reboulia hemisphaerica 2
JA(U)U(A)A(A)C(C)U(U)U(A)U(C)A(U)U(U)U(A)G(A)G(A)A(G)G(A)G(A)A(G)U(C)G(U)U(U)C(-)A(A)C(A)A(G)G(U)U(U)C(-)				8 Riccardia pinguis
U(U)U(A)A(A)C(C)U(U)U(A)U(C)A(U)U(U)U(A)G(A)G(A)A(G)G(A)G(A)A(G)U(C)G(U)U(U)C(-)A(A)C(A)A(G)G(U)U(U)C(-)				9 Sphaerocarpos donnellii
C(U)U(A)A(A)C(C)U(U)U(A)U(C)A(U)U(U)U(A)G(A)G(A)A(G)G(A)G(A)A(G)U(C)G(U)U(U)C(-)A(A)C(A)A(G)G(U)U(U)C(-)				10 Sphaerocarpos texanus
JA(U)U(A)A(A)C(C)U(U)U(A)U(C)A(U)U(U)U(A)G(A)G(A)A(G)G(A)G(A)A(G)U(C)G(U)U(U)C(-)A(A)C(A)A(G)G(U)U(U)C(-)				11 Atrichum undulatum
JA(U)U(A)A(A)C(C)U(U)U(A)U(C)A(U)U(U)U(A)G(A)G(A)A(G)G(A)G(A)A(G)U(C)G(U)U(U)C(-)A(A)C(A)A(G)G(U)U(U)C(-)				12 Bryum argenteum
JA(U)U(A)A(A)C(C)U(U)U(A)U(C)A(U)U(U)U(A)G(A)G(A)A(G)G(A)G(A)A(G)U(C)G(U)U(U)C(-)A(A)C(A)A(G)G(U)U(U)C(-)				13 Eurhynchium hians
JA(U)U(A)A(A)C(C)U(U)U(A)U(C)A(U)U(U)U(A)G(A)G(A)A(G)G(A)G(A)A(G)U(C)G(U)U(U)C(-)A(A)C(A)A(G)G(U)U(U)C(-)				14 Funaria hygrometrica
JA(U)U(A)A(A)C(C)U(U)U(A)U(C)A(U)U(U)U(A)G(A)G(A)A(G)G(A)G(A)A(G)U(C)G(U)U(U)C(-)A(A)C(A)A(G)G(U)U(U)C(-)				15 Leptobryum pyriforme
JA(U)U(A)A(A)C(C)U(U)U(A)U(C)A(U)U(U)U(A)G(A)G(A)A(G)G(A)G(A)A(G)U(C)G(U)U(U)C(-)A(A)C(A)A(G)G(U)U(U)C(-)				16 Mnium hornum
JA(C)C(A)A(A)C(C)U(U)U(A)U(C)A(U)U(U)U(A)G(A)G(A)A(G)G(A)G(A)A(G)U(C)G(U)U(U)C(-)A(A)C(A)A(G)G(U)U(U)C(-)				17 Physcomitrium pyriforme
JA(U)U(A)A(A)C(C)U(U)U(A)U(C)A(U)U(U)U(A)G(A)G(A)A(G)G(A)G(A)A(G)U(C)G(U)U(U)C(-)A(A)C(A)A(G)G(U)U(U)C(-)				18 Polytrichum formosum
JA(U)U(A)A(A)C(C)U(U)U(A)U(C)A(U)U(U)U(A)G(A)G(A)A(G)G(A)G(A)A(G)U(C)G(U)U(U)C(-)A(A)C(A)A(G)G(U)U(U)C(-)				19 Sphagnum cuspidatum
JA(U)U(A)A(A)C(C)U(U)U(A)U(C)A(U)U(U)U(A)G(A)G(A)A(G)G(A)G(A)A(G)U(C)G(U)U(U)C(-)A(A)C(A)A(G)G(U)U(U)C(-)				20 Calypogeia arguta
JU(U)U(A)A(A)C(C)U(U)U(A)U(C)A(U)U(U)U(A)G(A)G(A)A(G)G(A)G(A)A(G)U(C)G(U)U(U)C(-)A(A)C(A)A(G)G(U)U(U)C(-)				21 Conocephalum conicum
JA(U)U(A)A(A)C(C)U(U)U(A)U(C)A(U)U(U)U(A)G(A)G(A)A(G)G(A)G(A)A(G)U(C)G(U)U(U)C(-)A(A)C(A)A(G)G(U)U(U)C(-)				22 Fossombronia pusilla
JU(U)U(A)A(A)C(C)U(U)U(A)U(C)A(U)U(U)U(A)G(A)G(A)A(G)G(A)G(A)A(G)U(C)G(U)U(U)C(-)A(A)C(A)A(G)G(U)U(U)C(-)				23 Marchantia polymorpha
JU(U)U(A)A(A)C(C)U(U)U(A)U(C)A(U)U(U)U(A)G(A)G(A)A(G)G(A)G(A)A(G)U(C)G(U)U(U)C(-)A(A)C(A)A(G)G(U)U(U)C(-)				24 Riccia fluitans
JA(U)U(A)A(A)C(C)U(U)U(A)U(C)A(U)U(U)U(A)G(A)G(A)A(G)G(A)G(A)A(G)U(C)G(U)U(U)C(-)A(A)C(A)A(G)G(U)U(U)C(-)				25 Scapania nemorea
JA(U)U(A)A(A)C(C)U(U)U(A)U(C)A(U)U(U)U(A)G(A)G(A)A(G)G(A)G(A)A(G)U(C)G(U)U(U)C(-)A(A)C(A)A(G)G(U)U(U)C(-)				26 Equisetum hyemale
JA(U)U(A)A(A)C(C)U(U)U(A)U(C)A(U)U(U)U(A)G(A)G(A)A(G)G(A)G(A)A(G)U(C)G(U)U(U)C(-)A(A)C(A)A(G)G(U)U(U)C(-)				27 Equisetum robustum
A(U)U(A)A(A)C(C)U(U)U(A)U(C)A(U)U(U)U(A)G(A)G(A)A(G)G(A)G(A)A(G)U(C)G(U)U(U)C(-)A(A)C(A)A(G)G(U)U(U)C(-)				28 Isoetes engelmannii
JA(U)U(A)A(A)C(C)U(U)U(A)U(C)A(U)U(U)U(A)G(A)G(A)A(G)G(A)G(A)A(G)U(C)G(U)U(U)C(-)A(A)C(A)A(G)G(U)U(U)C(-)				29 Huperzia lucidula
JA(U)U(A)A(A)C(C)U(U)U(A)U(C)A(U)U(U)U(A)G(A)G(A)A(G)G(A)G(A)A(G)U(C)G(U)U(U)C(-)A(A)C(A)A(G)G(U)U(U)C(-)				30 Lycopodiella inundata
JA(U)U(A)A(A)C(C)U(U)U(A)U(C)A(U)U(U)U(A)G(A)G(A)A(G)G(A)G(A)A(G)U(C)G(U)U(U)C(-)A(A)C(A)A(G)G(U)U(U)C(-)				31 Lycopodium phlegmaria
JA(U)U(A)A(A)C(C)U(U)U(A)U(C)A(U)U(U)U(A)G(A)G(A)A(G)G(A)G(A)A(G)U(C)G(U)U(U)C(-)A(A)C(A)A(G)G(U)U(U)C(-)				32 Lycopodium taxifolium
JA(U)U(A)A(A)C(C)U(U)U(A)U(C)A(U)U(U)U(A)G(A)G(A)A(G)G(A)G(A)A(G)U(C)G(U)U(U)C(-)A(A)C(A)A(G)G(U)U(U)C(-)				33 Lycopodium tristachyum
JA(U)U(A)A(A)C(C)U(U)U(A)U(C)A(U)U(U)U(A)G(A)G(A)A(G)G(A)G(A)A(G)U(C)G(U)U(U)C(-)A(A)C(A)A(G)G(U)U(U)C(-)				34 Selaginella galleottii
JA(U)U(A)A(A)C(C)U(U)U(A)U(C)A(U)U(U)U(A)G(A)G(A)A(G)G(A)G(A)A(G)U(C)G(U)U(U)C(-)A(A)C(A)A(G)G(U)U(U)C(-)				35 Selaginella sp.
A(U)U(A)A(A)C(C)U(U)U(A)U(C)A(U)U(U)U(A)G(A)G(A)A(G)G(A)G(A)A(G)U(C)G(U)U(U)C(-)A(A)C(A)A(G)G(U)U(U)C(-)				36 Oryza sativa
A(U)U(A)A(A)C(C)U(U)U(A)U(C)A(U)U(U)U(A)G(A)G(A)A(G)G(A)G(A)A(G)U(C)G(U)U(U)C(-)A(A)C(A)A(G)G(U)U(U)C(-)				37 Zea mays
A(C)U(A)A(A)C(C)U(U)U(A)U(C)A(U)U(U)U(A)G(A)G(A)A(G)G(A)G(A)A(G)U(C)G(U)U(U)C(-)A(A)C(A)A(G)G(U)U(U)C(-)				38 Arabidopsis thaliana
A(C)U(A)A(A)C(C)U(U)U(A)U(C)A(U)U(U)U(A)G(A)G(A)A(G)G(A)G(A)A(G)U(C)G(U)U(U)C(-)A(A)C(A)A(G)G(U)U(U)C(-)				39 Glycine max
(U)C(U)A(A)A(C)C(U)U(U)U(A)U(C)A(U)U(U)U(A)G(A)G(A)A(G)G(A)G(A)A(G)U(C)G(U)U(U)C(-)A(A)C(A)A(G)G(U)U(U)C(-)				40 Sinapis alba
A(U)U(A)A(A)C(C)U(U)U(A)U(C)A(U)U(U)U(A)G(A)G(A)A(G)G(A)G(A)A(G)U(C)G(U)U(U)C(-)A(A)C(A)A(G)G(U)U(U)C(-)				41 Zamia pumila
A(U)U(A)A(A)C(C)U(U)U(A)U(C)A(U)U(U)U(A)G(A)G(A)A(G)G(A)G(A)A(G)U(C)G(U)U(U)C(-)A(A)C(A)A(G)G(U)U(U)C(-)				42 Gnetum leyboldii
A(U)U(A)A(A)C(C)U(U)U(A)U(C)A(U)U(U)U(A)G(A)G(A)A(G)G(A)G(A)A(G)U(C)G(U)U(U)C(-)A(A)C(A)A(G)G(U)U(U)C(-)				43 Ginkgo biloba
A(U)U(A)A(A)C(C)U(U)U(A)U(C)A(U)U(U)U(A)G(A)G(A)A(G)G(A)G(A)A(G)U(C)G(U)U(U)C(-)A(A)C(A)A(G)G(U)U(U)C(-)				44 Pinus luchuensis
A(U)U(A)A(A)C(C)U(U)U(A)U(C)A(U)U(U)U(A)G(A)G(A)A(G)G(A)G(A)A(G)U(C)G(U)U(U)C(-)A(A)C(A)A(G)G(U)U(U)C(-)				45 Pinus wallichiana
A(U)U(A)A(A)C(C)U(U)U(A)U(C)A(U)U(U)U(A)G(A)G(A)A(G)G(A)G(A)A(G)U(C)G(U)U(U)C(-)A(A)C(A)A(G)G(U)U(U)C(-)				46 Podocarpus nakaii
A(U)U(A)A(A)C(C)U(U)U(A)U(C)A(U)U(U)U(A)G(A)G(A)A(G)G(A)G(A)A(G)U(C)G(U)U(U)C(-)A(A)C(A)A(G)G(U)U(U)C(-)				47 Taxus mairei
JA(U)U(A)A(A)G(C)C(U)U(U)U(A)U(C)A(U)U(U)U(A)G(A)G(A)A(G)G(A)G(A)A(G)U(C)G(U)U(U)C(-)A(A)C(A)A(G)G(U)U(U)C(-)				48 Adiantum raddianum
JA(U)U(A)A(A)G(C)C(U)U(U)U(A)U(C)A(U)U(U)U(A)G(A)G(A)A(G)G(A)G(A)A(G)U(C)G(U)U(U)C(-)A(A)C(A)A(G)G(U)U(U)C(-)				49 Adiantum raddianum 2
JA(U)U(A)A(A)G(C)C(U)U(U)U(A)U(C)A(U)U(U)U(A)G(A)G(A)A(G)G(A)G(A)A(G)U(C)G(U)U(U)C(-)A(A)C(A)A(G)G(U)U(U)C(-)				50 Ophioglossum petiolatum
JA(U)U(A)A(A)G(C)C(U)U(U)U(A)U(C)A(U)U(U)U(A)G(A)G(A)A(G)G(A)G(A)A(G)U(C)G(U)U(U)C(-)A(A)C(A)A(G)G(U)U(U)C(-)				51 Osmunda cinnamomea
JA(U)U(A)A(A)G(C)C(U)U(U)U(A)U(C)A(U)U(U)U(A)G(A)G(A)A(G)G(A)G(A)A(G)U(C)G(U)U(U)C(-)A(A)C(A)A(G)G(U)U(U)C(-)				52 Salvinia natans
JA(U)U(A)A(A)C(C)U(U)U(A)U(C)A(U)U(U)U(A)G(A)G(A)A(G)G(A)G(A)A(G)U(C)G(U)U(U)C(-)A(A)C(A)A(G)G(U)U(U)C(-)				53 Psilotum nudum 2
A(U)U(A)A(A)C(C)U(U)U(A)U(C)A(U)U(U)U(A)G(A)G(A)A(G)G(A)G(A)A(G)U(C)G(U)U(U)C(-)A(A)C(A)A(G)G(U)U(U)C(-)				54 Chara foetida
JA(U)U(A)A(A)C(C)U(U)U(A)U(C)A(U)U(U)U(A)G(A)G(A)A(G)G(A)G(A)A(G)U(C)G(U)U(U)C(-)A(A)C(A)A(G)G(U)U(U)C(-)				55 Coleochaete scutata
JA(U)U(A)A(A)C(C)U(U)U(A)U(C)A(U)U(U)U(A)G(A)G(A)A(G)G(A)G(A)A(G)U(C)G(U)U(U)C(-)A(A)C(A)A(G)G(U)U(U)C(-)				56 Klebsormidium flaccidum
A(U)U(A)A(A)C(C)U(U)U(A)U(C)A(U)U(U)U(A)G(A)G(A)A(G)G(A)G(A)A(G)U(C)G(U)U(U)C(-)A(A)C(A)A(G)G(U)U(U)C(-)				57 Nitella flexilis
A(U)U(A)A(A)C(C)U(U)U(A)U(C)A(U)U(U)U(A)G(A)G(A)A(G)G(A)G(A)A(G)U(C)G(U)U(U)C(-)A(A)C(A)A(G)G(U)U(U)C(-)				58 Nitella sp.
A(U)U(A)A(A)C(C)U(U)U(A)U(C)A(U)U(U)U(A)G(A)G(A)A(G)G(A)G(A)A(G)U(C)G(U)U(U)C(-)A(A)C(A)A(G)G(U)U(U)C(-)				59 Chlamydomonas reinhardtii
A(U)U(A)A(A)C(C)U(U)U(A)U(C)A(U)U(U)U(A)G(A)G(A)A(G)G(A)G(A)A(G)U(C)G(U)U(U)C(-)A(A)C(A)A(G)G(U)U(U)C(-)				60 Hydrodictyon reticulatum
A(U)U(A)A(A)C(C)U(U)U(A)U(C)A(U)U(U)U(A)G(A)G(A)A(G)G(A)G(A)A(G)U(C)G(U)U(U)C(-)A(A)C(A)A(G)G(U)U(U)C(-)				61 Parietochloris pseudoalveolaris
A(U)U(A)A(A)C(C)U(U)U(A)U(C)A(U)U(U)U(A)G(A)G(A)A(G)G(A)G(A)A(G)U(C)G(U)U(U)C(-)A(A)C(A)A(G)G(U)U(U)C(-)				62 Dictyostelium discoideum
(U)U(U)A(A)A(U)C(U)U(C)A(U)U(U)U(A)G(A)G(A)A(G)G(A)G(A)A(G)U(C)G(U)U(U)C(-)A(A)C(A)A(G)G(U)U(U)C(-)				63 Helix numbering 1
- - - - - 49' - - - - -				
JA(U)U(A)A(A)C(C)U(U)U(A)U(C)A(U)U(U)U(A)G(A)G(A)A(G)G(A)G(A)A(G)U(C)G(U)U(U)C(-)A(A)C(A)A(G)G(U)U(U)C(-)				64 Pteris vittata
JA(U)U(A)A(A)C(C)U(U)U(A)U(C)A(U)U(U)U(A)G(A)G(A)A(G)G(A)G(A)A(G)U(C)G(U)U(U)C(-)A(A)C(A)A(G)G(U)U(U)C(-)				65 Megaceros aenigmaticus
JA(U)U(A)A(A)C(C)U(U)U(A)U(C)A(U)U(U)U(A)G(A)G(A)A(G)G(A)G(A)A(G)U(C)G(U)U(U)C(-)A(A)C(A)A(G)G(U)U(U)C(-)				66 Notophylas breutelii
JA(U)U(A)A(A)C(C)U(U)U(A)U(C)A(U)U(U)U(A)G(A)G(A)A(G)G(A)G(A)A(G)U(C)G(U)U(U)C(-)A(A)C(A)A(G)G(U)U(U)C(-)				67 Physcomitrella patens
JA(U)U(A)A(A)C(C)U(U)U(A)U(C)A(U)U(U)U(A)G(A)G(A)A(G)G(A)G(A)A(G)U(C)G(U)U(U)C(-)A(A)C(A)A(G)G(U)U(U)C(-)				68 Sphagnum palustre

2210	2220	2230	2240	2250	
C G U A G	G J U G A	A I C C U G C	G G A A G J G A U C	A U U A	1 Saccharomyces cerevisiae
(C)G U A G	G J U G A	A I C C U G C	(A)G A A G J G A U C	A U U A	2 Schizosaccharomyces pombe
(C)G U A G	G J U G A	A I C C U G C	(A)G A A G J G A U C	A o o o	3 Physcomitrella patens
(C)G U A G	G J U G A	A I C C U G C	(A)G A A G J G A U C	A o o o	4 Anthoceros agrestis
o o o o	o j o o	o o o o	o j o o	o o o o	5 Anthoceros laevis
(C)G U A G	G J U G A	A I C C U G C	(A)G A A G J G A U C	A o o o	6 Pellia epiphylla
(o) o o o	o j o o	o o o o	o j o o	o o o o	7 Reboulia hemisphaerica 2
(C)G U A G	G J U G A	A I C C U G C	(A)G A A G J G A U C	A o o o	8 Riccardia pinguis
(C)G U A G	G J U G A	A I C C U G C	(A)G A A G J G A U C	A o o o	9 Sphaerocarpos donnelli
(o) o o o	o j o o	o o o o	o j o o	o o o o	10 Sphaerocarpos texanus
(C)G U A G	G J U G A	A I C C U G C	(A)G A A G J G A U C	A o o o	11 Atrichum undulatum
(o) o o o	o j o o	o o o o	o j o o	o o o o	12 Bryum argenteum
(o) o o o	o j o o	o o o o	o j o o	o o o o	13 Eurhynchium hians
C G U A G	G J U G A	A I C C U G C	G G A A G J G A U C	A U U G C A C A C A C A C A A A A A A A A C U G	14 Funaria hygrometrica
(C)G U A G	G J U G A	A I C C U G C	(A)G A A G J G A U C	A C U G G G C C G	15 Leptobryum pyriforme
(C)G U A G	G J U G A	A I C C U G C	(A)G A A G J G A U C	A o o o	16 Mnium hornum
(o) o o o	o j o o	o o o o	o j o o	o o o o	17 Physcomitrium pyriforme
(C)G U A G	G J U G A	A I C C U G C	(A)G A A G J G A U C	A o o o	18 Polytrichum formosum
(C)G U A G	G J U G A	A I C C U G C	(A)G A A G J G A U C	A o o o	19 Sphagnum cuspidatum
(C)G U A G	G J U G A	A I C C U G C	(A)G A A G J G A U C	A o o o	20 Calypogeia arguta
U G U A G	G J U G A	A I C C U G C	A G A A G J G A U C	A o o o	21 Conocephalum conicum
(C)G U A G	G J U G A	A I C C U G C	(A)G A A G J G A U C	A o o o	22 Fossombronina pusilla
(C) o o o	o j o o	o o o o	o j o o	o o o o	23 Marchantia polymorpha
(C)G U A G	G J U G A	A I C C U G C	(A)G A A G J G A U C	A o o o	24 Riccia fluitans
(C)G U A G	G J U G A	A I C C U G C	(A)G A A G J G A U C	A o o o	25 Scapania nemorea
(o) o o o	o j o o	o o o o	o j o o	o o o o	26 Equisetum hyemale
(C) o o o	o j o o	o o o o	o j o o	o o o o	27 Equisetum robustum
o o o o	o j o o	o o o o	o j o o	o o o o	28 Isoetes engelmannii
o o o o	o j o o	o o o o	o j o o	o o o o	29 Huperzia lucidula
(o) o o o	o j o o	o o o o	o j o o	o o o o	30 Lycopodiella inundata
(C) o o o	o j o o	o o o o	o j o o	o o o o	31 Lycopodium phlegmaria
(C) o o o	o j o o	o o o o	o j o o	o o o o	32 Lycopodium taxifolium
(o) o o o	o j o o	o o o o	o j o o	o o o o	33 Lycopodium cristachyum
(C) o o o	o j o o	o o o o	o j o o	o o o o	34 Selaginella galleottii
(C) o o o	o j o o	o o o o	o j o o	o o o o	35 Selaginella sp.
C G U A G	G J U G A	A I C C U G C	G G A A G J G A U C	A U U G	36 Oryza sativa
C G U A G	G J U G A	A I C C U G C	G G A A G J G A U C	A U U G	37 Zea mays
C G U A G	G J U G A	A I C C U G C	G G A A G J G A U C	A U U G	38 Arabidopsis thaliana
C G U A G	G J U G A	A I C C U G C	G G A A G J G A U C	A U U G	39 Glycine max
C G U A G	G J U G A	A I C C U G C	G G A A G J G A U C	A U U G	40 Sinapis alba
C G U A G	G J U G A	A I C C U G C	G G A A G J G A U C	A U U G	41 Zamia pumila
C G U A G	G J U G A	A I C C U G C	G G A A G J G A U C	A U U G	42 Gnetum leyboldii
o o o o	o j o o	o o o o	o j o o	o o o o	43 Ginkgo biloba
o o o o	o j o o	o o o o	o j o o	o o o o	44 Pinus luchuensis
o o o o	o j o o	o o o o	o j o o	o o o o	45 Pinus wallichiana
o o o o	o j o o	o o o o	o j o o	o o o o	46 Podocarpus nakaai
o o o o	o j o o	o o o o	o j o o	o o o o	47 Taxus mairei
(o) o o o	o j o o	o o o o	o j o o	o o o o	48 Adiantum raddianum
(C) o o o	o j o o	o o o o	o j o o	o o o o	49 Adiantum raddianum 2
o o o o	o j o o	o o o o	o j o o	o o o o	50 Ophioglossum petiolatum
o o o o	o j o o	o o o o	o j o o	o o o o	51 Osmunda cinnamomea
C o o o o	o j o o	o o o o	o j o o	o o o o	52 Salvinia natans
(C) o o o	o j o o	o o o o	o j o o	o o o o	53 Psilotum nudum 2
(o) o o o	o j o o	o o o o	o j o o	o o o o	54 Chara foetida
(C) o o o	o j o o	o o o o	o j o o	o o o o	55 Coleochaete scutata
(C)G U A G	G J U G A	A I C C U G C	(A)G A A G J G A U C	A o o o	56 Klebsormidium flaccidum
(C) o o o	o j o o	o o o o	o j o o	o o o o	57 Nitella flexilis
(C)G U A G	G J U G A	A I C C U G C	(A)G A A G J G A U C	A o o o	58 Nitella sp.
C G U A G	G J U G A	A I C C U G C	G G A A G J G A U C	A U U G	59 Chlamydomonas reinhardtii
C G U A G	G J U G A	A I C C U G C	G G A A G J G A U C	A U U G	60 Hydrodictyon reticulatum
C G U A G	G J U G A	A I C C U G C	G G A A G J G A U C	A U U G	61 Parietochloris pseudoalveolaris
C G U A G	G J U G A	A I C C U G C	G G A A G J G A U C	A U U U	62 Dictyostelium discoideum
- 50 -	- 50 -	- 50 -	- 50 -	- 50 -	63 Helix numbering 1
o o o o	o j o o	o o o o	o j o o	o o o o	64 Pteris vittata
C G U A G	G J U G A	A I C C U G C	G G A A G J G A U C	A o o o	65 Megaceros aenigmaticus
C G U A G	G J U G A	A I C C U G C	G G A A G J G A U C	A o o o	66 Notophylas breutelii
(C)G U A G	G J U G A	A I C C U G C	(A)G A A G J G A U C	A o o o	67 Physcomitrella patens
(C)G U A G	G J U G A	A I C C U G C	(A)G A A G J G A U C	A o o o	68 Sphagnum palustre

NOTE TO USERS

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UMI

APPENDIX H : MacClade Matrix

Variability categories are assigned to each alignment position, or column. The categories were transcribed to this matrix from Table 4 (Page 44).

Taxa with Variability Info

	1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20	21	22	23	24	25	26	27	28	29	30
1 Saccharomyces cerevisiae	.	T	A	T	C	T	G	G	T	T	G	A	T	C	C	T	G	C	C	A	G	T	A	G	T	C	A	T	A	T
2 Schizosaccharomyces pombe	.	T	A	C	C	T	G	G	T	T	G	A	T	C	C	T	G	C	C	A	G	T	A	G	T	C	A	T	A	T
3 Physcomitrella patens 2	.	A	A	C	C	T	G	G	T	T	G	A	T	C	C	T	G	C	C	A	G	T	A	G	T	C	A	T	A	T
4 Anthoceros agrestis	.	A	A	C	C	T	G	G	T	T	G	A	T	C	C	T	G	C	C	A	G	T	A	G	T	C	A	T	A	T
5 Anthoceros laevis	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?
6 Pedia epiphyta	.	A	A	C	C	T	G	G	T	T	G	A	T	C	C	T	G	C	C	A	G	T	A	G	T	C	A	T	A	T
7 Reboulia hemisphaerica	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?
8 Riccardia pinguis	.	A	A	C	C	T	G	G	T	T	G	A	T	C	C	T	G	C	C	A	G	T	A	G	T	C	A	T	A	T
9 Sphaerocarpos donnellii	.	A	A	C	C	T	G	G	T	T	G	A	T	C	C	T	G	C	C	A	G	T	A	G	T	C	A	T	A	T
10 Sphaerocarpos lezaricus	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?
11 Atrichum undulatum	.	A	A	C	C	T	G	G	T	T	G	A	T	C	C	T	G	C	C	A	G	T	A	G	T	C	A	T	A	T
12 Bryum argenteum	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?
13 Eurymochium hiare	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?
14 Funaria hygrometrica	.	T	A	C	C	T	G	G	T	T	G	A	T	C	C	T	G	C	C	A	G	T	A	G	T	C	A	T	A	T
15 Lophoceros pyriforme	.	A	A	C	C	T	G	G	T	T	G	A	T	C	C	T	G	C	C	A	G	T	A	G	T	C	A	T	A	T
16 Mnium haerum	.	A	A	C	C	T	G	G	T	T	G	A	T	C	C	T	G	C	C	A	G	T	A	G	T	C	A	T	A	T
17 Physcomitrium pyriforme	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?
18 Polytrichum formosum	.	A	A	C	C	T	G	G	T	T	G	A	T	C	C	T	G	C	C	A	G	T	A	G	T	C	A	T	A	T
19 Sphagnum cuspidatum	.	A	A	C	C	T	G	G	T	T	G	A	T	C	C	T	G	C	C	A	G	T	A	G	T	C	A	T	A	T
20 Calypogeia arguta	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?
21 Cerocephalum carinatum	.	A	A	C	C	T	G	G	T	T	G	A	T	C	C	T	G	C	C	A	G	T	A	G	T	C	A	T	A	T
22 Fossombronia pusilla	.	A	A	C	C	T	G	G	T	T	G	A	T	C	C	T	G	C	C	A	G	T	A	G	T	C	A	T	A	T
23 Marchantia polymorpha	.	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?
24 Riccia fluitans	.	A	A	C	C	T	G	G	T	T	G	A	T	C	C	T	G	C	C	A	G	T	A	G	T	C	A	T	A	T
25 Scapania nemorea	.	A	A	C	C	T	G	G	T	T	G	A	T	C	C	T	G	C	C	A	G	T	A	G	T	C	A	T	A	T
26 Equisetum hyemale	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?
27 Equisetum robustum	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?
28 Isoetes engelmannii	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?
29 Huperzia lucidulum	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?
30 Lycopodium truncatum	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?
31 Lycopodium phlegmaria	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?
32 Lycopodium lucidulum	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?
33 Lycopodium tristichyum	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?
34 Selaginella selaginoides	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?
35 Selaginella selaginoides	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?
36 Oryza sativa	.	T	A	C	C	T	G	G	T	T	G	A	T	C	C	T	G	C	C	A	G	T	A	G	T	C	A	T	A	T
37 Zea mays	.	T	A	C	C	T	G	G	T	T	G	A	T	C	C	T	G	C	C	A	G	T	A	G	T	C	A	T	A	T
38 Arabidopsis thaliana	.	T	A	C	C	T	G	G	T	T	G	A	T	C	C	T	G	C	C	A	G	T	A	G	T	C	A	T	A	T
39 Glycine max	.	T	A	C	C	T	G	G	T	T	G	A	T	C	C	T	G	C	C	A	G	T	A	G	T	C	A	T	A	T
40 Sinapis alba	.	T	A	C	C	T	G	G	T	T	G	A	T	C	C	T	G	C	C	A	G	T	A	G	T	C	A	T	A	T
41 Zinnia punctata	.	T	A	C	C	T	G	G	T	T	G	A	T	C	C	T	G	C	C	A	G	T	A	G	T	C	A	T	A	T
42 Onoclea sensibilis	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?
43 Girardinia bicolor	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?
44 Pinus luchuanensis	.	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?
45 Pinus massoniana	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?
46 Nagaiia nagii	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?
47 Taxus malayana	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?
48 Adiantum raddianum	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?
49 Adiantum raddianum 2	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?
50 Ophioglossum petiolatum	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?
51 Osmunda cinnamomea	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?
52 Selaginella selaginoides	.	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?
53 Polypodium nudum	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?
54 Chara foetida	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?
55 Coleochaete scutata	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?
56 Klebsormidium flaccidum	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?
57 Nitella flexilis	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?
58 Nitzschia sp.	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?
59 Chlamydomonas reinhardtii	.	T	A	C	C	T	G	G	T	T	G	A	T	C	C	T	G	C	C	A	G	T	A	G	T	C	A	T	A	T
60 Hydrodictyon reticulatum	.	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?
61 Parletochloris pseudocylindrica	.	T	A	C	C	T	G	G	T	T	G	A	T	C	C	T	G	C	C	A	G	T	A	G	T	C	A	T	A	T
62 Dictyosphaerium diosporum	.	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?
63 Plectonicon balticum	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?
64 Megaceros aeruginosus	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?
65 Netophyes brevifolium	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?
66 Physcomitrella patens	.	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?
67 Sphagnum palustre	.	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?

Taxa with Variability Info 5

	121	122	123	124	125	126	127	128	129	130	131	132	133	134	135	136	137	138	139	140	141	142	143	144	145	146	147	148	149	150
	4	6	8	2	4	4	0	4	0	1	4	5	4	0	4	4	2	4	5	3	3	3			0	0	0	2	3	0
1 Saccharomyces cerevisiae	T	A	T	C	G	T	.	T	T	A	T	T	T	G	A	T	A	G	T	T	C	C	T	T	T	.
2 Schizosaccharomyces pombe	T	A	T	C	G	T	.	T	T	A	T	T	T	G	A	T	A	G	T	A	C	C	T	C	A	.
3 Physcomitrella patens 2	T	A	T	A	G	T	.	T	T	C	T	T	T	G	A	T	G	G	T	A	C	C	T	C	A	.
4 Anthoceros agrestis	T	A	T	A	G	T	.	T	T	C	T	T	T	G	A	T	G	G	T	A	C	C	T	A	.	.
5 Anthoceros laevis	T	A	T	A	G	T	.	T	T	A	T	T	T	G	A	T	G	G	T	A	C	C	T	T	.	.
6 Pellia epiphylla	T	A	T	A	G	T	.	T	T	C	T	T	T	G	A	T	G	G	T	A	C	C	T	T	.	.
7 Reboulia hemisphaerica	T	A	T	A	G	T	.	T	T	C	T	T	T	G	A	T	G	G	T	A	C	C	T	T	.	.
8 Riccardia purpurea	T	A	T	A	G	T	.	T	T	C	T	T	T	G	A	T	G	G	T	A	C	C	T	T	.	.
9 Sphaerocarpos donnellii	T	A	T	A	G	T	.	T	T	C	T	T	T	G	A	T	G	G	T	A	C	C	T	T	.	.
10 Sphaerocarpos lasarus	T	A	T	A	G	T	.	C	T	.	C	T	T	G	A	T	G	G	T	A	C	C	T	T	.	.
11 Atrichum undulatum	T	A	T	A	G	T	.	T	T	C	T	T	T	G	A	T	G	G	T	A	C	C	T	T	.	.
12 Bryum argenteum	T	A	T	A	G	T	.	T	T	C	T	T	T	G	A	T	G	G	T	A	C	C	T	T	.	.
13 Eurytrichum hiemale	T	A	T	A	G	T	.	T	T	C	T	T	T	G	A	T	G	G	T	A	C	C	T	T	.	.
14 Funaria hygrometrica	T	A	T	A	G	T	.	T	T	C	T	T	T	G	A	T	G	G	T	A	C	C	T	T	.	.
15 Leptobryum pyriforme	T	A	T	A	G	T	.	T	T	C	T	T	T	G	A	T	G	G	T	A	C	C	T	T	.	.
16 Mnium hornum	T	A	T	A	G	T	.	T	T	C	T	T	T	G	A	T	G	G	T	A	C	C	T	T	.	.
17 Physcomitrium pyriforme	T	A	T	A	G	T	.	T	T	C	T	T	T	G	A	T	G	G	T	A	C	C	T	T	.	.
18 Ptilothecium formosum	T	A	T	A	G	T	.	T	T	C	T	T	T	G	A	T	G	G	T	A	C	C	T	T	.	.
19 Sphagnum cuspidatum	T	A	T	A	G	T	.	T	T	C	T	T	T	G	A	T	G	G	T	A	C	C	T	T	.	.
20 Calypogeia rigida	T	A	T	A	G	T	.	T	T	C	T	T	T	G	A	T	G	G	T	A	C	C	T	T	.	.
21 Conocephalum conicum	T	A	T	A	G	T	.	T	T	C	T	T	T	G	A	T	G	G	T	A	C	C	T	T	.	.
22 Fossombronia pusilla	T	A	T	A	G	T	.	T	T	C	T	T	T	G	A	T	G	G	T	A	C	C	T	T	.	.
23 Marchantia polymorpha	T	A	T	A	G	T	.	T	T	C	T	T	T	G	A	T	G	G	T	A	C	C	T	T	.	.
24 Riccia fluitans	T	A	T	A	G	C	.	T	T	C	T	T	T	G	A	T	G	G	T	A	C	C	T	A	.	.
25 Isoetes macrospora	T	A	T	A	G	T	.	T	T	C	T	T	T	G	A	T	G	G	T	A	C	C	T	A	.	.
26 Equisetum hyemale	T	A	T	A	G	T	.	T	T	C	T	T	T	G	A	T	G	G	T	A	C	C	T	.	.	.
27 Equisetum robustum	T	A	T	A	G	T	.	T	T	C	T	T	T	G	A	T	G	G	T	A	C	C	T	.	.	.
28 Isoetes engelmannii	T	A	T	A	G	T	.	T	T	C	T	T	T	G	A	T	G	G	T	A	C	C	T	.	.	.
29 Huperzia lucidulum	T	A	T	A	G	T	.	T	T	C	T	T	T	G	A	T	G	G	T	A	C	C	T	.	.	.
30 Lycopodium lucidulum	T	A	T	A	G	T	.	T	T	C	T	T	T	G	A	T	G	G	T	A	C	C	T	.	.	.
31 Lycopodium pliegmaria	T	A	T	A	G	T	.	T	T	C	T	T	T	G	A	T	G	G	T	A	C	C	T	.	.	.
32 Lycopodium lucidulum	T	A	T	A	G	T	.	T	T	C	T	T	T	G	A	T	G	G	T	A	C	C	T	.	.	.
33 Lycopodium tristichyum	T	A	T	A	G	T	.	T	T	C	T	T	T	G	A	T	G	G	T	A	C	C	T	.	.	.
34 Selaginella umbrosa	T	A	T	A	G	T	.	T	T	A	T	T	T	G	A	T	G	G	T	A	C	C	T	.	.	.
35 Selaginella vogelii	A	A	T	A	G	T	.	T	T	A	T	T	T	G	A	T	G	T	T	A	C	C	T	.	.	.
36 Oryza sativa	T	A	T	A	G	T	.	T	T	G	T	T	T	G	A	T	G	G	T	A	C	G	T	.	.	.
37 Zea mays	T	A	T	A	G	T	.	T	T	G	T	T	T	G	A	T	G	G	T	A	C	G	T	.	.	.
38 Arabidopsis thaliana	T	A	T	A	G	T	.	T	T	G	T	T	T	G	A	T	G	G	T	A	A	C	T	.	.	.
39 Glycine max	T	A	T	A	G	T	.	T	T	G	T	T	T	G	A	T	G	G	T	A	T	C	T	.	.	.
40 Strapsis alba	T	A	T	A	G	T	.	T	T	G	T	T	T	G	A	T	G	G	T	A	A	C	T	.	.	.
41 Zizia aurea	T	A	T	A	G	T	.	T	T	C	T	T	T	G	A	T	G	G	T	A	C	T	C	T	.	.
42 Quercus laevis	T	A	T	A	G	T	.	T	T	C	T	T	T	G	A	T	G	G	T	A	C	A	T	T	.	.
43 Quercus bicolor	T	A	T	A	G	T	.	T	T	C	T	T	T	G	A	T	G	G	T	A	C	C	T	T	.	.
44 Pinus taeda	T	A	T	A	G	T	.	T	T	T	T	T	T	G	A	T	G	G	T	A	C	C	A	T	.	.
45 Pinus strobus	T	A	T	A	G	T	.	T	T	T	T	T	T	G	A	T	G	G	T	A	C	C	T	T	.	.
46 Haploxyloium nigrum	T	A	T	A	G	T	.	T	T	A	T	T	T	G	A	T	G	G	T	A	C	C	T	T	.	.
47 Taxus canadensis	T	A	T	A	G	T	.	T	T	C	T	T	T	G	A	T	G	G	T	A	C	C	T	T	.	.
48 Adiantum radicans	T	A	.	A	G	T	.	T	T	C	T	T	T	G	A	T	G	G	T	A	C	C	T	.	.	.
49 Adiantum radicans 2	T	A	T	A	G	T	.	T	T	C	T	T	T	G	A	T	G	G	T	A	C	C	T	.	.	.
50 Ophioglossum petiolatum	T	A	T	A	G	T	.	T	T	C	T	T	T	G	A	T	G	G	T	A	C	C	T	.	.	.
51 Osmunda cinnamomea	T	A	T	A	G	T	.	T	T	C	T	T	T	G	A	T	G	G	T	A	C	C	T	.	.	.
52 Salvinia natans	T	A	T	A	G	T	.	T	T	C	T	T	T	G	A	T	G	G	T	A	C	C	T	.	.	.
53 Pilea pumila	T	A	T	A	G	T	.	T	T	C	T	T	T	G	A	T	G	G	T	A	C	C	C	T	T	.
54 Chara foeniculacea	T	A	T	C	A	T	.	T	T	A	T	T	T	G	A	T	G	G	T	A	C	C	T	T	.	.
55 Coleochaete scutata	T	A	T	A	G	T	.	T	T	A	T	T	T	G	A	T	G	G	T	A	C	C	T	T	.	.
56 Habrocallia flaccida	T	A	T	A	G	T	.	T	T	A	T	T	T	G	A	T	G	G	T	A	C	C	T	T	.	.
57 Nitella flexilis	T	A	T	C	G	T	.	T	T	A	T	T	T	G	A	T	G	T	T	C	C	C	C	C	A	.
58 Nitella sp	T	A	T	C	G	T	.	T	T	A	T	T	T	G	A	T	G	T	T	C	C	C	C	C	A	.
59 Chlamydomonas reinhardtii	T	A	T	A	G	T	.	T	T	A	T	T	T	G	A	T	G	G	T	A	C	C	T	.	.	.
60 Hydrodictyon reticulatum	T	A	T	A	G	T	.	T	T	A	T	T	T	G	A	T	G	G	T	A	C	C	T	.	.	.
61 Parileichthys pseudocostata	T	A	T	A	G	T	.	T	T	A	T	T	T	G	A	T	G	G	T	A	C	C	T	.	.	.
62 Dictyosphaerium discoideum	G	A	T	A	A	A	.	C	T	A	A	T	A	G	A	C	T	T	T	C	G	G	.	.	G	T	T	T	.	.
63 Pteris vittata	T	A	T	A	G	T	.	T	T	C	T	T	T	G	A	T	G	G	T	A	C	C	T	T	.	.
64 Marsippospermum angustatum	T	A	T	A	G	T	.	T	T	C	T	T	T	G	A	T	G	G	T	A	C	C	T	T	.	.
65 Notophyes brutus	G	A	T	A	G	T	.	T	C	T	G	A	G	A	G	G	G	G	C	A	C	C	T	T	.	C
66 Physcomitrella patens	T	A	T	A	G	T	.	T	T	C	T	T	T	G	A	T	G	G	T	A	C	C	T	T	.	.
67 Sphagnum palustre	T	A	T	A	G	T	.	T	T	C	T	T	T	G	A	T	G	G	T	A	C	C	T	T	.	.

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	181	182	183	184	185	186	187	188	189	190	191	192	193	194	195	196	197	198	199	200	201	202	203	204	205	206	207	208	209	210
	0	0	0	4	0	0	0	0	0	5	0	5	2	0	0	5	2	1	0	2	3	3	0	1	2	0	3	0	5	
1 Saccharomyces cerevisiae	T	A	G	A	G	C	T	A	A	T	A	-	C	A	T	G	C	T	-	-	-	A	A	A	A	T	C	T	C	G
2 Schizosaccharomyces pombe	T	A	G	A	G	C	T	A	A	T	A	-	C	A	T	G	C	T	A	-	-	A	A	A	A	T	C	C	C	G
3 Physcomitrella patens 2	T	A	G	A	G	C	T	A	A	T	A	-	C	G	T	G	C	A	C	-	A	A	A	-	C	T	C	C	G	
4 Anthoceros agrestis	T	A	G	A	G	C	T	A	A	T	A	-	C	G	T	G	C	A	A	-	C	A	A	-	C	T	C	C	G	
5 Anthoceros laevis	T	A	G	A	G	C	T	A	A	T	A	-	C	G	T	G	C	A	A	-	C	A	A	-	C	T	C	C	G	
6 Peltia epiphylla	T	A	G	A	G	C	T	A	A	T	A	-	C	G	T	G	C	A	C	-	C	A	A	-	C	T	C	C	G	
7 Reboulia hemisphaerica	T	A	G	A	G	C	T	A	A	T	A	-	C	G	T	G	C	A	C	-	C	A	A	-	C	T	C	C	G	
8 Riccardia pinguis	T	A	G	A	G	C	T	A	A	T	A	-	C	G	T	G	C	A	C	-	C	A	A	-	C	T	C	C	G	
9 Sphaerocarpos cornelli	T	A	G	A	G	C	T	A	A	T	A	-	C	G	T	G	C	A	C	-	C	A	A	-	C	T	C	C	G	
10 Sphaerocarpos lecanus	T	A	G	A	G	C	T	A	A	T	A	-	C	G	T	G	C	A	C	-	C	A	A	-	C	T	C	C	G	
11 Anichium undulatum	T	A	G	A	G	C	T	A	A	T	A	-	C	G	T	G	C	A	C	-	A	A	A	-	C	T	C	C	G	
12 Bryum argenteum	T	A	G	A	G	C	T	A	A	T	A	-	C	G	T	G	C	A	C	-	A	A	A	-	C	T	C	C	G	
13 Eurynchium hiemale	T	A	G	A	G	C	T	A	A	T	A	-	C	G	T	G	C	A	C	-	A	A	A	-	C	T	C	C	G	
14 Funaria hygrometrica	T	A	G	A	G	C	T	A	A	T	A	-	C	G	T	G	C	A	C	-	A	A	A	-	C	T	C	C	G	
15 Leptobryum pyriforme	T	A	G	A	G	C	T	A	A	T	A	-	C	G	T	G	C	A	C	-	A	A	A	-	C	T	C	C	G	
16 Mnium hornum	T	A	G	A	G	C	T	A	A	T	A	-	C	G	T	G	C	A	C	-	A	A	A	-	C	T	C	C	G	
17 Physcomitrium pyriforme	T	A	G	A	G	C	T	A	A	T	A	-	C	G	T	G	C	A	C	-	A	A	A	-	C	T	C	C	G	
18 Polytichum formosum	T	A	G	A	G	C	T	A	A	T	A	-	C	G	T	G	C	A	C	-	A	A	A	-	C	T	C	C	G	
19 Sphagnum cuspidatum	T	A	G	A	G	C	T	A	A	T	A	-	C	G	T	G	C	A	C	-	A	A	A	-	C	T	C	C	G	
20 Calypogeia arguta	T	A	G	A	G	C	T	A	A	T	A	-	C	G	T	G	C	A	C	-	A	A	A	-	C	T	C	C	G	
21 Conocephalum coriolum	T	A	G	A	G	C	T	A	A	T	A	-	C	G	T	G	C	A	C	-	C	A	A	-	C	T	C	C	G	
22 Fossombronia pusilla	T	A	G	A	G	C	T	A	A	T	A	-	C	G	T	G	C	A	C	-	C	A	A	-	C	T	C	C	G	
23 Marchantia polymorpha	T	A	G	A	G	C	T	A	A	T	A	-	C	G	T	G	C	A	C	-	C	A	A	-	C	T	C	C	G	
24 Riccia fluitans	T	A	G	A	G	C	T	A	A	T	A	-	C	G	T	G	C	A	C	-	C	A	A	-	C	T	C	C	G	
25 Scapania nemorea	T	A	G	A	G	C	T	A	A	T	A	-	C	G	T	G	C	A	C	-	C	A	A	-	C	T	C	C	G	
26 Equisetum hyemale	T	A	G	A	G	C	T	A	A	T	A	-	C	G	T	G	C	A	C	-	C	A	A	-	C	T	C	C	G	
27 Equisetum robustum	T	A	G	A	G	C	T	A	A	T	A	-	C	G	T	G	C	A	C	-	C	A	A	-	C	T	C	C	G	
28 Isoetes engelmannii	T	A	G	A	G	C	T	A	A	T	A	-	C	G	T	G	C	A	C	-	C	A	A	-	C	T	C	C	G	
29 Huperzia lucidulum	T	A	G	A	G	C	T	A	A	T	A	-	C	G	T	G	C	A	C	-	C	A	A	-	C	T	C	C	G	
30 Lycopodium inundatum	T	A	G	A	G	C	T	A	A	T	A	-	C	G	T	G	C	A	C	-	C	A	A	-	C	T	C	C	G	
31 Lycopodium phlegmaria	T	A	G	A	G	C	T	A	A	T	A	-	C	G	T	G	C	A	C	-	C	A	A	-	C	T	C	C	G	
32 Lycopodium latifolium	T	A	G	A	G	C	T	A	A	T	A	-	C	G	T	G	C	A	C	-	C	A	A	-	C	T	C	C	G	
33 Lycopodium tristachyum	T	A	G	A	G	C	T	A	A	T	A	-	C	G	T	G	C	A	C	-	C	A	A	-	C	T	C	C	G	
34 Selaginella umbrosa	T	A	G	A	G	C	T	A	A	T	A	-	C	G	T	G	C	A	C	-	A	A	A	-	C	T	C	C	G	
35 Selaginella vogelii	T	A	G	A	G	C	T	A	A	T	A	-	C	G	T	G	C	A	C	-	C	A	A	-	C	T	C	C	G	
36 Oryza sativa	T	A	G	A	G	C	T	A	A	T	A	-	C	G	T	G	C	A	A	-	C	A	A	-	C	T	C	C	G	
37 Zea mays	T	A	G	A	G	C	T	A	A	T	A	-	C	G	T	G	C	A	A	-	C	A	A	-	C	T	C	C	G	
38 Arabidopsis thaliana	T	A	G	A	G	C	T	A	A	T	A	-	C	G	T	G	C	A	A	-	C	A	A	-	C	T	C	C	G	
39 Glycine max	T	A	G	A	G	C	T	A	A	T	A	-	C	G	T	G	C	A	A	-	C	A	A	-	C	T	C	C	G	
40 Strapsia alba	T	A	G	A	G	C	T	A	A	T	A	-	C	G	T	G	C	A	A	-	C	A	A	-	C	T	C	C	G	
41 Zinnia purpurea	T	A	G	A	G	C	T	A	A	T	A	-	C	G	T	G	C	A	A	-	C	A	A	-	C	T	C	C	G	
42 Onoclea sensibilis	T	A	G	A	G	C	T	A	A	T	A	-	C	G	T	G	C	A	A	-	C	A	A	-	C	T	C	C	G	
43 Citrigo lobata	T	A	G	A	G	C	T	A	A	T	A	-	C	G	T	G	C	A	A	-	C	A	A	-	C	T	C	C	G	
44 Pinus luchuanensis	T	A	G	A	G	C	T	A	A	T	A	-	C	G	T	G	C	A	A	-	C	A	A	-	C	T	C	C	G	
45 Pinus wallichiana	T	A	G	A	G	C	T	A	A	T	A	-	C	G	T	G	C	A	A	-	C	A	A	-	C	T	C	C	G	
46 Nigella arvensis	T	A	G	A	G	C	T	A	A	T	A	-	C	G	T	G	C	A	A	-	C	A	A	-	C	T	C	C	G	
47 Triticum aestivum	T	A	G	A	G	C	T	A	A	T	A	-	C	G	T	G	C	A	A	-	C	A	A	-	C	T	C	C	G	
48 Adiantum eschlerianum	T	A	G	A	G	C	T	A	A	T	A	-	C	G	T	G	C	A	A	-	C	A	A	-	C	T	C	C	G	
49 Adiantum eschlerianum 2	T	A	G	A	G	C	T	A	A	T	A	-	C	G	T	G	C	A	A	-	C	A	A	-	C	T	C	C	G	
50 Ophioglossum petiolatum	T	A	G	A	G	C	T	A	A	T	A	-	C	G	T	G	C	A	A	-	C	A	A	-	C	T	C	C	G	
51 Osmunda cinnamomea	T	A	G	A	G	C	T	A	A	T	A	-	C	G	T	G	C	A	A	-	C	A	A	-	C	T	C	C	G	
52 Salvinia natans	T	A	G	A	G	C	T	A	A	T	A	-	C	G	T	G	C	A	A	-	C	A	A	-	C	T	C	C	G	
53 Psilotum nudum	T	A	G	A	G	C	T	A	A	T	A	-	C	G	T	G	C	A	A	-	C	A	A	-	C	T	C	C	G	
54 Chara foetida	T	A	G	A	G	C	T	A	A	T	A	-	C	G	T	G	C	A	A	-	C	A	A	-	C	T	C	C	G	
55 Coleochaete scutata	T	A	G	A	G	C	T	A	A	T	A	-	C	G	T	G	C	A	A	-	C	A	A	-	C	T	C	C	G	
56 Nostoc commune	T	A	G	A	G	C	T	A	A	T	A	-	C	G	T	G	C	A	A	-	C	A	A	-	C	T	C	C	G	
57 Nostoc sp.	T	A	G	A	G	C	T	A	A	T	A	-	C	G	T	G	C	A	A	-	C	A	A	-	C	T	C	C	G	
58 Chlamydomonas reinhardtii	T	A	G	A	G	C	T	A	A	T	A	-	C	G	T	G	C	A	A	-	C	A	A	-	C	T	C	C	G	
59 Hydrodictyon reticulatum	T	A	G	A	G	C	T	A	A	T	A	-	C	G	T	G	C	A	A	-	C	A	A	-	C	T	C	C	G	
60 Paritochloris pseudosphaerula	T	A	G	A	G	C	T	A	A	T	A	-	C	G	T	G	C	A	A	-	C	A	A	-	C	T	C	C	G	
61 Dictyostelium discoideum	-	G	G	G	G	C	T	A	A	T	A	-	C	A	T	A	C	A	A	G	C	G	A	-	T	G	G	T	G	
62 Pharis vittata	T	A	G	A	G	C	T	A	A	T	A	-	C	G	T	G	C	A	A	-	C	A	A	-	C	T	C	C	G	
63 Megascopus asio	T	A	G	A	G	C	T	A	A	T	A	-	C	G	T	G	C	A	A	-	C	A	A	-	C	T	C	C	G	
64 Metaphysa brucei	T	A	G	A	G	C	T	A	A	T	A	-	C	G	T	G	C	A	A	-	C	A	A	-	C	T	C	C	G	
65 Physcomitrella patens	T	A	G	A	G	C	T	A	A	T	A	-	C	G	T	G	C	A	A	-	C	A	A	-	C	T	C	C	G	
66 Sphagnum paludosum	T	A	G	A	G	C	T	A	A	T	A	-	C	G	T	G	C	A	A	-	C	A	A	-	C	T	C	C	G	

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	301	302	303	304	305	306	307	308	309	310	311	312	313	314	315	316	317	318	319	320	321	322	323	324	325	326	327	328	329	330	331	332	333	334	335	336	337	338	339	340				
1 Saccharomyces cerevisiae	.	.	C	T	T	Q	T	.	Q	C	T	Q	Q	C	Q	A	T	Q	.	Q	T	T	C	A	T	T	C	A	T	T	C	.	A	A	.	A	A	.	A	A	.	A	A	
2 Schizosaccharomyces pombe	.	.	C	T	T	Q	C	.	Q	C	C	Q	Q	C	Q	A	T	Q	.	Q	T	T	C	A	T	T	C	A	T	T	C	.	A	A	.	A	A	.	A	A	.	A	A	
3 Physcomitrella patens 2	.	C	T	T	Q	Q	C	.	Q	C	T	Q	Q	C	Q	A	T	Q	.	T	T	T	C	A	T	T	C	A	T	T	C	.	A	A	.	A	A	.	A	A	.	A	A	
4 Arabidopsis thaliana	.	C	C	T	Q	Q	C	.	Q	C	T	Q	Q	C	Q	A	T	Q	.	T	T	T	C	A	T	T	C	A	T	T	C	.	A	A	.	A	A	.	A	A	.	A	A	
5 Arabidopsis lyrata	.	C	C	C	T	Q	C	.	Q	C	C	Q	Q	C	C	A	T	Q	.	T	T	T	C	A	T	T	C	A	T	T	C	.	A	A	.	A	A	.	A	A	.	A	A	
6 Pellaea epiphylla	.	C	T	T	Q	Q	C	.	Q	C	T	Q	Q	C	Q	A	T	Q	.	T	T	T	C	A	T	T	C	A	T	T	C	.	A	A	.	A	A	.	A	A	.	A	A	
7 Reboulia hemisphaerica	.	C	T	T	A	Q	C	.	Q	C	T	Q	Q	C	Q	A	T	Q	.	T	T	T	C	A	T	T	C	A	T	T	C	.	A	A	.	A	A	.	A	A	.	A	A	
8 Riccardia pinguis	.	C	C	C	Q	Q	C	.	Q	C	C	Q	Q	C	Q	A	T	Q	.	T	T	T	C	A	T	T	C	A	T	T	C	.	A	A	.	A	A	.	A	A	.	A	A	
9 Sphaerocarpos donnellii	C	T	A	T	T	Q	C	.	Q	C	T	Q	Q	C	Q	A	T	Q	.	T	T	T	C	A	T	T	C	A	T	T	C	.	A	A	.	A	A	.	A	A	.	A	A	
10 Sphaerocarpos lasarum	.	C	T	T	T	Q	C	.	Q	C	T	Q	Q	C	Q	A	T	Q	.	T	T	T	C	A	T	T	C	A	T	T	C	.	A	A	.	A	A	.	A	A	.	A	A	
11 Atrichum undulatum	.	C	T	T	A	Q	C	.	Q	C	T	Q	Q	C	Q	A	T	Q	.	T	T	T	C	A	T	T	C	A	T	T	C	.	A	A	.	A	A	.	A	A	.	A	A	
12 Bryum argenteum	.	C	T	T	A	Q	C	.	Q	C	T	Q	Q	C	Q	A	T	Q	.	T	T	T	C	A	T	T	C	A	T	T	C	.	A	A	.	A	A	.	A	A	.	A	A	
13 Eurynchium hians	.	C	C	C	A	Q	C	.	Q	C	T	Q	Q	C	Q	A	T	Q	.	T	T	T	C	A	T	T	C	A	T	T	C	.	A	A	.	A	A	.	A	A	.	A	A	
14 Funaria hygrometrica	.	C	C	C	A	Q	C	.	Q	C	T	Q	Q	C	Q	A	T	Q	.	T	T	T	C	A	T	T	C	A	T	T	C	.	A	A	.	A	A	.	A	A	.	A	A	
15 Leptobryum pyriforme	.	C	C	C	A	Q	C	.	Q	C	T	Q	Q	C	Q	A	T	Q	.	T	T	T	C	A	T	T	C	A	T	T	C	.	A	A	.	A	A	.	A	A	.	A	A	
16 Mnium hornum	.	C	T	T	Q	Q	C	.	Q	C	C	Q	Q	C	Q	A	T	Q	.	T	T	T	C	A	T	T	C	A	T	T	C	.	A	A	.	A	A	.	A	A	.	A	A	
17 Physcomitrium pyriforme	.	C	T	T	Q	Q	C	.	Q	C	T	Q	Q	C	Q	A	T	Q	.	T	T	T	C	A	T	T	C	A	T	T	C	.	A	A	.	A	A	.	A	A	.	A	A	
18 Psittidium tomesum	.	C	T	C	A	Q	C	.	Q	C	T	Q	Q	C	Q	A	T	Q	.	T	T	T	C	A	T	T	C	A	T	T	C	.	A	A	.	A	A	.	A	A	.	A	A	
19 Sphagnum cuspidatum	.	C	C	T	Q	Q	C	.	Q	C	C	Q	Q	C	Q	A	T	Q	.	T	T	T	C	A	T	T	C	A	T	T	C	.	A	A	.	A	A	.	A	A	.	A	A	
20 Calypogeia angusta	.	.	C	C	C	A	Q	C	.	Q	C	T	Q	Q	C	Q	A	T	Q	.	T	T	T	C	A	T	T	C	A	T	T	C	.	A	A	.	A	A	.	A	A	.	A	A
21 Conocephalum conicum	C	T	T	T	T	Q	C	.	Q	C	C	Q	Q	C	Q	A	T	Q	.	T	T	T	C	A	T	T	C	A	T	T	C	.	A	A	.	A	A	.	A	A	.	A	A	
22 Foscomitrium pusilla	.	C	G	T	A	Q	C	.	Q	C	T	Q	Q	C	Q	A	T	Q	.	T	T	T	C	A	T	T	C	A	T	T	C	.	A	A	.	A	A	.	A	A	.	A	A	
23 Marchantia polymorpha	.	C	T	T	A	Q	C	.	Q	C	T	Q	Q	C	Q	A	T	Q	.	T	T	T	C	A	T	T	C	A	T	T	C	.	A	A	.	A	A	.	A	A	.	A	A	
24 Riccia fluitans	.	C	T	C	C	Q	C	.	Q	C	C	Q	Q	C	Q	A	T	Q	.	T	T	T	C	A	T	T	C	A	T	T	C	.	A	A	.	A	A	.	A	A	.	A	A	
25 Scapania nemorea	.	C	T	T	T	Q	C	.	Q	C	T	Q	Q	C	Q	A	T	Q	.	T	T	T	C	A	T	T	C	A	T	T	C	.	A	A	.	A	A	.	A	A	.	A	A	
26 Equisetum hyemale	.	C	T	T	Q	Q	C	.	Q	C	C	Q	Q	C	Q	A	T	Q	.	T	T	T	C	A	T	T	C	A	T	T	C	.	A	A	.	A	A	.	A	A	.	A	A	
27 Equisetum robustum	.	C	T	T	Q	.	C	.	Q	C	C	Q	Q	C	Q	A	T	Q	.	T	T	T	C	A	T	T	C	A	T	T	C	.	A	A	.	A	A	.	A	A	.	A	A	
28 Isoetes engelmannii	.	C	T	T	A	Q	T	.	Q	C	C	Q	Q	C	Q	A	T	A	.	C	T	T	C	A	T	T	C	A	T	T	C	.	A	A	.	A	A	.	A	A	.	A	A	
29 Huperzia lucidulum	.	C	T	T	T	Q	C	.	Q	C	C	Q	Q	C	Q	A	T	Q	.	T	T	T	C	A	T	T	C	A	T	T	C	.	A	A	.	A	A	.	A	A	.	A	A	
30 Lycopodium truncatum	.	C	T	C	A	Q	C	.	Q	C	C	Q	Q	C	Q	A	T	Q	.	T	T	T	C	A	T	T	C	A	T	T	C	.	A	A	.	A	A	.	A	A	.	A	A	
31 Lycopodium phlegmaria	.	C	C	T	C	Q	C	.	Q	C	C	Q	Q	C	Q	A	T	Q	.	T	T	T	C	A	T	T	C	A	T	T	C	.	A	A	.	A	A	.	A	A	.	A	A	
32 Lycopodium lucidulum	.	C	C	T	C	Q	C	.	Q	C	C	Q	Q	C	Q	A	T	Q	.	C	T	T	C	A	T	T	C	A	T	T	C	.	A	A	.	A	A	.	A	A	.	A	A	
33 Lycopodium tristachyum	.	C	T	C	A	Q	T	.	Q	C	C	Q	Q	C	Q	A	T	Q	.	C	T	T	C	A	T	T	C	A	T	T	C	.	A	A	.	A	A	.	A	A	.	A	A	
34 Selaginella umbrosa	.	C	T	C	T	Q	T	.	Q	C	T	Q	Q	C	Q	A	T	Q	.	C	T	T	C	A	T	T	C	A	T	T	C	.	A	A	.	A	A	.	A	A	.	A	A	
35 Selaginella vogelii	.	.	C	C	A	Q	A	.	Q	C	C	Q	Q	C	Q	A	T	Q	.	C	A	T	C	A	T	T	C	A	T	T	C	.	A	A	.	A	A	.	A	A	.	A	A	
36 Oryza sativa	.	C	C	T	C	Q	T	.	Q	C	C	Q	Q	C	Q	A	C	Q	.	C	A	T	C	A	T	T	C	A	T	T	C	.	A	A	.	A	A	.	A	A	.	A	A	
37 Zea mays	.	C	T	T	C	Q	T	.	Q	C	C	Q	Q	C	Q	A	C	Q	.	C	A	T	C	A	T	T	C	A	T	T	C	.	A	A	.	A	A	.	A	A	.	A	A	
38 Arabidopsis thaliana	.	C	T	C	T	Q	T	.	Q	C	T	Q	Q	C	Q	A	C	Q	.	C	A	T	C	A	T	T	C	A	T	T	C	.	A	A	.	A	A	.	A	A	.	A	A	
39 Glycine max	.	C	T	T	T	Q	T	.	Q	C	C	Q	Q	C	Q	A	C	Q	.	C	A	T	C	A	T	T	C	A	T	T	C	.	A	A	.	A	A	.	A	A	.	A	A	
40 Brasica alba	.	C	T	T	T	Q	T	.	Q	C	T	Q	Q	C	Q	A	C	Q	.	C	A	T	C	A	T	T	C	A	T	T	C	.	A	A	.	A	A	.	A	A	.	A	A	
41 Zania punctata	.	C	C	T	C	Q	A	.	Q	C	C	Q	Q	C	Q	A	C	Q	.	C	T	T	C	A	T	T	C	A	T	T	C	.	A	A	.	A	A	.	A	A	.	A	A	
42 Ononis laevis	.	C	T	T	T	Q	T	.	Q	C	C	Q	Q	C	Q	A	C	Q	.	C	T	T	C	A	T	T	C	A	T	T	C	.	A	A	.	A	A	.	A	A	.	A	A	
43 Citrus bibe	.	C	C	T	Q	Q	T	.	Q	C	T	Q	Q	C	Q	A	T	Q	.	C	T	T	C	A	T	T	C	A	T	T	C	.	A	A	.	A	A	.	A	A	.	A	A	
44 Pinus luchuensis	.	C	C	T	T	Q	T	.	Q	C	T	Q	Q	C	Q	A	T	Q	.	C	T	T	C	A	T	T	C	A	T	T	C	.	A	A	.	A	A	.	A	A	.	A	A	
45 Pinus wallichiana	.	C	C	T	T	Q	T	.	Q	C	T	Q	Q	C	Q	A	T	Q	.	C	T	T	C	A	T	T	C	A	T	T	C	.	A	A	.	A	A	.	A	A	.	A	A	
46 Nepenthes rafflesia	.	C	C	T	Q	Q	T	.	Q	C	C	Q	Q	C	Q	A	C	Q	.	C	T	T	C	A	T	T	C	A	T	T	C	.	A	A	.	A	A	.	A	A	.	A	A	
47 Taxus mairei	.	C	T	T	T	Q	T	.	Q	C	T	Q	Q	C	Q	A	C	Q	.	C	T	T	C	A	T	T	C	A	T	T	C	.	A	A	.	A	A	.	A	A	.	A	A	
48 Adiantum redonianum	.	C	T	T	Q	Q	C	.	Q	C	C	Q	Q	C	Q	A	T	Q	.	C	T	T	C	A	T	T	C	A	T	T	C	.	A	A	.	A	A	.	A	A	.	A	A	
49 Adiantum redonianum 2	.	C	T	T	Q	Q	C	.	Q	C	C	Q	Q	C	Q	A	T	Q	.	C	T	T	C	A	T	T	C	A	T	T	C	.	A	A	.	A	A	.	A	A	.	A	A	
50 Ophioglossum petiolatum	.	C	T	T	.	Q	C	.	Q	C	C	Q	Q	C	Q	A	C	Q	.	C	T	T	C	A	T	T	C	A	T	T	C	.	A	A	.	A	A	.	A	A	.	A	A	
51 Osmunda cinnamomea	.	C	T	T	Q	Q	C	.	Q	C	C	Q	Q	C	Q	A	T	Q	.	C	T	T	C	A	T	T	C	A	T	T	C	.	A	A	.	A	A	.	A	A	.	A	A	
52 Salvinia natans	.	C	C	T	T	Q	C	.	Q	C	C	Q	Q	C	Q	A																												

1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20	21	22	23	24	25	26	27	28	29	30	31	32	33	34	35	36	37	38	39	40	41	42	43	44	45	46	47	48	49	50	51	52	53	54	55	56	57	58	59	60	61	62	63	64	65	66	67	68	69	70	71	72	73	74	75	76	77	78	79	80	81	82	83	84	85	86	87	88	89	90	91	92	93	94	95	96	97	98	99	100
1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20	21	22	23	24	25	26	27	28	29	30	31	32	33	34	35	36	37	38	39	40	41	42	43	44	45	46	47	48	49	50	51	52	53	54	55	56	57	58	59	60	61	62	63	64	65	66	67	68	69	70	71	72	73	74	75	76	77	78	79	80	81	82	83	84	85	86	87	88	89	90	91	92	93	94	95	96	97	98	99	100

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1 Saccharomyces cerevisiae	A	.	A	G	G	C	A	G	C	A	.	G	G	C	G	C	G	C	A	A	A	T	T	A	C	C	C	A	A	T
2 Schizosaccharomyces pombe	A	.	A	G	G	C	A	G	C	A	.	G	G	C	G	C	G	C	A	A	A	T	T	A	C	C	C	A	A	T
3 Physcomitrella patens 2	A	.	A	G	G	C	A	G	C	A	.	G	G	C	G	C	G	C	A	A	A	T	T	A	C	C	C	A	A	T
4 Anthraceros agreste	A	.	A	G	G	C	A	G	C	A	.	G	G	C	G	C	G	C	A	A	A	T	T	A	C	C	C	A	A	T
5 Anthraceros laevis	A	.	A	G	G	C	A	G	C	A	.	G	G	C	G	G	G	C	A	A	A	T	T	A	C	C	C	A	A	T
6 Pellia epiphylla	A	.	A	G	G	C	A	G	C	A	.	G	G	C	G	C	G	C	A	A	A	T	T	A	C	C	C	A	A	T
7 Reboulia hemisphaerica	A	.	A	G	G	C	A	G	C	A	.	G	G	C	G	C	G	C	A	A	A	T	T	A	C	C	C	A	A	T
8 Riccardia pinguis	A	.	A	G	G	C	A	G	C	A	.	G	G	C	G	C	G	C	A	A	A	T	T	A	C	C	C	A	A	T
9 Sphaerocarpos donnellii	A	.	A	G	G	C	A	G	C	A	.	G	G	C	G	C	G	C	A	A	A	T	T	A	C	C	C	A	A	T
10 Sphaerocarpos lasarus	A	.	A	G	G	C	A	G	C	A	.	G	G	C	G	C	G	C	A	A	A	T	T	A	C	C	C	A	A	T
11 Atrichum undulatum	A	.	A	G	G	C	A	G	C	A	.	G	G	C	G	C	G	C	A	A	A	T	T	A	C	C	C	A	A	T
12 Bryum argenteum	A	.	A	G	G	C	A	G	C	A	.	G	G	C	G	C	G	C	A	A	A	T	T	A	C	C	C	A	A	T
13 Eurhynchium hians	A	.	A	G	G	C	A	G	A	T	.	G	G	C	G	C	G	C	A	A	A	T	T	A	C	C	C	A	A	T
14 Funaria hygrometrica	A	.	A	G	G	C	A	G	C	A	.	G	G	C	G	C	G	C	A	A	A	T	T	A	C	C	C	A	A	T
15 Leptobryum pyriforme	A	.	A	G	G	C	A	G	C	A	.	G	G	C	G	C	G	C	A	A	A	T	T	A	C	C	C	A	A	T
16 Mnium hornum	A	.	A	G	G	C	A	G	C	A	.	G	G	C	G	C	G	C	A	A	A	T	T	A	C	C	C	A	A	T
17 Physcomitrium pyriforme	A	.	A	G	G	C	A	G	C	A	.	G	G	C	G	C	G	C	A	A	A	T	T	A	C	C	C	A	A	T
18 Polytrichum formosum	A	.	A	G	G	C	A	G	C	A	.	G	G	C	G	C	G	C	A	A	A	T	T	A	C	C	C	A	A	T
19 Sphagnum cuspidatum	A	.	A	G	G	C	A	G	C	A	.	G	G	C	G	C	G	C	A	A	A	T	T	A	C	C	C	A	A	T
20 Catepogon argutus	A	.	A	G	G	C	A	G	C	A	.	G	G	C	G	C	G	C	A	A	A	T	T	A	C	C	C	A	A	T
21 Conocephalum conicum	A	.	A	G	G	C	A	G	C	A	.	G	G	C	G	C	G	C	A	A	A	T	T	A	C	C	C	A	A	T
22 Foscombronia pusilla	A	.	A	G	G	C	A	G	C	A	.	G	G	C	G	C	G	C	A	A	A	T	T	A	C	C	C	A	A	T
23 Marchantia polymorpha	A	.	A	G	G	C	A	G	C	A	.	G	G	C	G	C	G	C	A	A	A	T	T	A	C	C	C	A	A	T
24 Riccia fluitans	A	.	A	G	G	C	A	G	C	A	.	G	G	C	G	C	G	C	A	A	A	T	T	A	C	C	C	A	A	T
25 Scapania nemorea	A	.	A	G	G	C	A	G	C	A	.	G	G	C	G	C	G	C	A	A	A	T	T	A	C	C	C	A	A	T
26 Equisetum hyemale	A	.	A	G	G	C	A	G	C	G	.	G	G	C	G	C	G	C	A	A	A	T	T	A	C	C	C	A	A	T
27 Equisetum robustum	A	.	A	G	G	C	A	G	C	A	.	G	G	C	G	C	G	C	A	A	A	T	T	A	C	C	C	A	A	T
28 Isetes engelmannii	A	.	A	G	G	C	A	G	C	A	.	G	G	C	G	C	G	C	A	A	A	T	T	A	C	C	C	A	A	T
29 Huperzia lucidulum	A	.	A	G	G	C	A	G	C	A	.	G	G	C	G	C	G	C	A	A	A	T	T	A	C	C	C	A	A	T
30 Lycopodium inundatum	A	.	A	G	G	C	A	G	C	A	.	G	G	C	G	G	G	C	A	A	A	T	T	A	C	C	C	A	A	T
31 Lycopodium phlegmaria	A	.	A	G	G	C	A	G	C	A	.	G	G	C	G	C	G	C	A	A	A	T	T	A	C	C	C	A	A	T
32 Lycopodium taxifolium	A	.	A	G	G	C	A	G	C	A	.	G	G	C	G	C	G	C	A	A	A	T	T	A	C	C	C	A	A	T
33 Lycopodium tristachyum	A	.	A	G	G	C	A	G	C	A	.	G	G	C	G	C	G	C	A	A	A	T	T	A	C	C	C	A	A	T
34 Selaginella umbrosa	A	.	A	G	G	C	A	G	C	A	.	G	G	C	G	C	G	C	A	A	A	T	T	A	C	C	C	A	A	T
35 Selaginella vogelii	A	.	A	G	G	C	A	G	C	A	.	G	G	C	G	C	G	C	A	A	A	T	T	A	C	C	C	A	A	T
36 Oryza sativa	A	.	A	G	G	C	A	G	C	A	.	G	G	C	G	C	G	C	A	A	A	T	T	A	C	C	C	A	A	T
37 Zea mays	A	.	A	G	G	C	A	G	C	A	.	G	G	C	G	C	G	C	A	A	A	T	T	A	C	C	C	A	A	T
38 Arabidopsis thaliana	A	.	A	G	G	C	A	G	C	A	.	G	G	C	G	C	G	C	A	A	A	T	T	A	C	C	C	A	A	T
39 Glycine max	A	.	A	G	G	C	A	G	C	A	.	G	G	C	G	C	G	C	A	A	A	T	T	A	C	C	C	A	A	T
40 Brassica oleracea	A	.	A	G	G	C	A	G	C	A	.	G	G	C	G	C	G	C	A	A	A	T	T	A	C	C	C	A	A	T
41 Zinnia punctata	A	.	A	G	G	C	A	G	C	A	.	G	G	C	G	G	G	C	A	A	A	T	T	A	C	C	C	A	A	T
42 Arabidopsis lyrata	A	.	A	G	G	C	A	G	C	A	.	G	G	C	G	C	G	C	A	A	A	T	T	A	C	C	C	A	A	T
43 Arabidopsis thaliana	A	.	A	G	G	C	A	G	C	A	.	G	G	C	G	C	G	C	A	A	A	T	T	A	C	C	C	A	A	T
44 Pinus taeda	A	.	A	G	G	C	A	G	C	A	.	G	G	C	G	C	G	C	A	A	A	T	T	A	C	C	C	A	A	T
45 Pinus strobus	A	.	A	G	G	C	A	G	C	A	.	G	G	C	G	C	G	C	A	A	A	T	T	A	C	C	C	A	A	T
46 Magnolia grandiflora	A	.	A	G	G	C	A	G	C	A	.	G	G	C	G	C	G	C	A	A	A	T	T	A	C	C	C	A	A	T
47 Taxus canadensis	A	.	A	G	G	C	A	G	C	A	.	G	G	C	G	C	G	C	A	A	A	T	T	A	C	C	C	A	A	T
48 Adiantum pedatum	A	.	A	G	G	C	A	G	C	A	.	G	G	C	G	C	G	C	A	A	A	T	T	A	C	C	C	A	A	T
49 Adiantum pedatum 2	A	.	A	G	G	C	A	G	C	A	.	G	G	C	G	C	G	C	A	A	A	T	T	A	C	C	C	A	A	T
50 Ophioglossum pedatum	A	.	A	G	G	C	A	G	C	A	.	G	G	C	G	C	G	C	A	A	A	T	T	A	C	C	C	A	A	T
51 Osmunda cinnamomea	A	.	A	G	G	C	A	G	T	A	.	G	G	C	G	C	G	T	A	A	A	T	T	A	C	C	C	A	A	T
52 Salvinia natans	A	.	A	G	G	C	A	G	C	A	.	G	G	C	G	C	G	C	A	A	A	T	T	A	C	C	C	A	A	T
53 Pteris aquilina	A	.	A	G	G	C	A	G	C	A	.	G	G	C	G	C	G	C	A	A	A	T	T	A	C	C	C	A	A	T
54 Chara foeniculacea	A	.	A	G	G	C	A	G	C	A	.	G	G	C	G	C	G	T	A	A	A	T	T	A	C	C	C	A	A	T
55 Coleochaete scutata	A	.	A	G	G	C	A	G	C	A	.	G	G	C	G	C	G	C	A	A	A	T	T	A	C	C	C	A	A	T
56 Nostoc commune	A	.	A	G	G	C	A	G	C	A	.	G	G	C	G	C	G	C	A	A	A	T	T	A	C	C	C	A	A	T
57 Nitella saxatilis	A	.	A	G	G	C	A	G	C	A	.	G	G	C	G	C	G	C	A	A	A	T	T	A	C	C	C	A	A	T
58 Nitella sp.	A	.	A	G	G	C	A	G	C	A	.	G	G	C	G	C	G	C	A	A	A	T	T	A	C	C	C	A	A	T
59 Chara domingensis	A	.	A	G	G	C	A	G	C	A	.	G	G	C	G	C	G	C	A	A	A	T	T	A	C	C	C	A	A	T
60 Hydrodictyon reticulatum	A	.	A	G	G	C	A	G	C	A	.	G	G	C	G	C	G	C	A	A	A	T	T	A	C	C	C	A	A	T
61 Parietochloris pseudohyalinaria	A	.	A	G	G	C	A	G	C	A	.	G	G	C	G	C	G	C	A	A	A	T	T	A	C	C	C	A	A	T
62 Dictyosphaerium diocoides	A	.	A	G	G	C	A	G	C	A	.	G	G	C	G	C	G	C	A	A	A	T	T	A	C	T	C	A	A	T
63 Pteris vittata	A	.	A	G	G	C	A	G	C	A	.	G	G	C	G	C	G	C	A	A	A	T	T	A	C	C	C	A	A	T
64 Megaceros serrigulosus	A	.	A	G	G	C	A	G	C	A	.	G	G	C	G	C	G	C	A	A	A	T	T	A	C	C	C	A	A	T
65 Nostoc commune	A	.	A	G	G	C	A	G	C	A	.	G	G	C	G	C	G	C	A	A	A	T	T	A	C	C	C	A	A	T
66 Physcomitrella patens	A	.	A	G	G	C	A	G	C	A	.	G	G	C	G	C	G	C	A	A	A	T	T	A	C	C	C	A	A	T
67 Sphagnum palustre	A	.	A	G	G	C	A	G	C	A	.	G	G	C	G	C	G	C	A	A	A	T	T	A	C	C	C	A	A	T

	011	012	013	014	015	016	017	018	019	020	021	022	023	024	025	026	027	028	029	030	031	032	033	034	035	036	037	038	039	040
	1	1	0	0	0	0	0	0	1	1	1	2	1	0	1	1	3	2	2	3	2	0	0	0	0	0	0	0	0	0
1 Saccharomyces cerevisiae	T	T	T	C	C	A	A	C	G	.	G	G	C	C	T	T	T	G	C
2 Schizosaccharomyces pombe	.	T	C	A	T	G	A	C	C	G	.	G	G	T	C	G	T	T	A	A	C
3 Physcomitrella patens 2	T	C	C	A	C	T	C	Q	.	Q	C	C	T	C	T
4 Anisoceros agrestis	T	G	A	C	C	T	C	A	.	T	C	C	C	T	C	T
5 Anisoceros laevis	T	G	A	C	C	T	C	A	.	T	C	C	T	A	C
6 Pellia epiphylla	T	C	C	T	T	C	C	.	T	C	T	C	T	T
7 Reboulia hemisphaerica	T	C	C	T	T	C	T	C	T	.	C	C	T	T
8 Riccia phlegula	T	C	T	C	C	C	C	Q	T	C	C	T	T	T
9 Sphaerocarpos donnellii	.	C	C	Q	A	C	T	C	.	T	C	T	C	T	T	T	T
10 Sphaerocarpos laxus	.	C	.	.	.	C	Q	G	C	G	C	C	T	.	C	T	C	C	T	T	T
11 Atrichum undulatum	T	C	T	C	T	C	Q	A	.	T	C	C	T	T	T
12 Bryum argenteum	T	C	G	A	C	C	G	Q	.	A	C	C	T	T	Q	T	T	T
13 Euhynchium hians	T	C	C	A	C	C	C	Q	.	A	.	C	C	T	T	T
14 Funaria hygrometrica	T	C	C	A	C	T	.	C	.	Q	G	C	C	T	T	C
15 Leptobryum pyriforme	T	C	C	A	C	C	.	C	.	Q	A	C	C	T	T	T
16 Mnium hornum	T	C	G	A	C	C	.	C	.	Q	A	C	C	T	T	G	T	T	T
17 Physcomitrium pyriforme	T	C	C	A	C	T	C	Q	.	Q	C	C	T
18 Ptilotidium tomentosum	T	C	T	C	T	C	C	.	.	Q	T	C	C	T	T	T
19 Sphagnum cuspidatum	C	C	T	C	C	G	C	.	.	Q	T	C	C	T	T	A
20 Calypogeia arguta	T	C	T	C	T	C	C	A	.	T	C	C	T	T	T
21 Conocephalum conicum	Q	Q	Q	T	C	T	T	.	C	T	T	T	T	T	Q	T	A	T
22 Fossombronia pusilla	T	C	T	C	T	C	C	.	A	T	C	C	T	T	T
23 Marchantia polymorpha	C	Q	Q	C	C	C	C	.	T	T	T	C	T	T	T
24 Riccia fluitans	C	Q	Q	Q	C	C	T	.	T	C	T	C	T	T	Q	T	T
25 Scapania nemorea	T	C	T	C	T	C	C	.	A	C	C	C	T	T
26 Equisetum hyemale	T	C	Q	C	T	C	G	.	Q	T	C	C	T	T	T
27 Equisetum robustum	T	C	Q	C	T	C	G	.	Q	T	C	C	T	T	T
28 Isoetes engelmannii	Q	C	A	T	T	T	C	.	A	C	C	G	T	T
29 Huperzia lucidulum	T	C	Q	C	T	C	C	.	A	T	T	C	T	T	T
30 Lycopodium inundatum	C	T	Q	C	T	C	C	.	A	C	T	C	T	T	T
31 Lycopodium phlegmaria	T	C	Q	C	T	C	C	.	A	T	C	C	T	T	T
32 Lycopodium laetifolium	T	C	Q	C	T	C	C	.	A	T	C	C	T	T	T
33 Lycopodium tristichyum	T	C	Q	C	T	C	C	.	A	T	Y	C	T	T	T
34 Selaginella umbrosa	C	C	C	C	Q	T	C	.	A	G	C	C	T	T	T
35 Selaginella vogelii	T	C	Q	C	T	C	C	.	A	T	C	C	T	T	T
36 Oryza sativa	C	C	T	Q	.	C	T	C	Q	A	C	C	C	T
37 Zea mays	C	C	Q	Q	C	T	C	Q	A	C	C	C	T	T
38 Arabidopsis thaliana	T	C	Q	Q	C	T	T	.	Q	T	C	C	C	T	T
39 Glycine max	T	C	Q	Q	C	T	C	.	Q	T	C	C	C	T	T
40 Brassica oleracea	T	C	Q	Q	C	T	T	.	Q	T	C	C	C	T	T
41 Zizia aurea	C	C	Q	T	T	T	C	.	Q	T	C	C	C	T	T
42 Oenothera lutea	C	C	Q	C	T	C	C	.	Q	T	T	C	C	T	T
43 Oenothera biennis	C	C	Q	C	T	C	C	.	Q	T	C	C	C	T	T
44 Pinus taeda	C	C	C	T	T	O	C	.	Q	T	C	C	C	T	T
45 Pinus strobus	C	C	C	T	T	C	C	.	Q	T	C	C	C	T	T
46 Pinus resinosa	C	C	C	T	T	C	C	.	Q	T	C	C	C	T	T
47 Taxus canadensis	C	A	C	T	C	Q	C	.	Q	T	C	C	C	T	T
48 Taxus virginiana	C	A	C	T	C	A	C	.	Q	T	C	C	G	T	T
49 Adiantum raddianum	T	C	Q	C	T	C	C	.	Q	C	C	C	T	T	T
50 Adiantum raddianum 2	T	C	Q	C	T	C	C	.	Q	C	C	C	T	T	T
51 Ophioglossum petiolatum	Q	C	Q	C	T	C	C	.	Q	C	C	C	T	T	T
52 Osmunda cinnamomea	T	C	C	T	O	C	.	A	T	C	C	T	T	T
53 Salvinia natans	T	T	Q	C	C	T	C	.	Q	G	C	C	T	T	T
54 Polypodium vulgatum	T	C	Q	C	T	C	.	C	.	Q	T	C	C	T	T	C
55 Chara foeniculacea	C	C	C	T	T	T	C	.	Q	C	C	T	T	T	C
56 Coleochaete scutata	T	T	T	C	T	C	C	.	A	T	C	C	T	T	C
57 Nitzschia communis	C	C	Q	A	C	C	C	.	A	T	C	C	T	T	C
58 Nitzschia sp.	C	C	C	T	T	C	C	.	Q	C	C	T	C	T	C
59 Chlamydomonas reinhardtii	T	C	T	Q	C	T	C	.	C	A	C	C	T	T	C
60 Hydrodictyon reticulatum	T	A	T	Q	Q	C	C	.	C	T	C	C	T	T	T
61 Parachlorella chlorella	C	Q	T	Q	Q	C	C	.	C	A	C	C	T	T	Q
62 Dictyosphaerula diatomum	.	C	Q	A	C	.	A	C	C	Q	Q	T	A	.	T	C	T	C	T	T	Y	C	T	T	A	A	.	.	.	
63 Plectonon balticum	T	C	Q	C	T	Q	C	.	Q	C	C	C	T	T	T
64 Mopane serripalpus	T	Q	A	C	T	C	C	A	.	?	T	C	C	T	T
65 Notophytes brucei	.	G	A	C	C	T	C	A	.	C	C	C	C	T
66 Physcomitrella patens	T	C	C	A	C	T	C	Q	.	Q	C	C	T	T	C	T
67 Sphagnum palustre	C	C	T	C	C	.	C	.	Q	T	C	C	T	T	A

	001	002	003	004	005	006	007	008	009	010	011	012	013	014	015	016	017	018	019	020	021	022	023	024	025	026	027	028	029	030	031	032	033	034	035	036	037	038	039	040
1 Saccharomyces cerevisiae	T	.	.	T	G	C	T	.	C	G	A	A	.	T	A	T	.	A	T	T	.	A	G	.	C	A	T	G	G	A										
2 Schizosaccharomyces pombe	T	.	T	T	G	C	T	.	C	G	A	A	.	T	A	C	.	A	T	T	.	A	G	.	C	A	T	G	G	A										
3 Physcomitrella patens 2	T	.	.	.	G	C	T	C	T	G	A	A	.	T	A	C	.	A	T	T	.	A	G	.	C	A	T	G	G	A										
4 Anthoceros agrestis	T	.	.	.	G	C	T	C	T	G	A	A	.	T	A	C	.	A	T	T	.	A	G	.	C	A	T	G	G	A										
5 Anthoceros laevis	T	.	.	.	G	C	T	C	T	G	A	A	.	T	A	C	.	A	T	T	.	A	G	.	C	A	T	G	G	A										
6 Pellia epiphylla	T	.	.	.	G	C	T	C	T	G	A	A	.	T	A	C	.	A	T	T	.	A	G	.	C	A	T	G	G	A										
7 Reboulia hemisphaerica	A	C	.	.	G	C	.	C	T	G	A	A	.	T	A	C	.	A	T	T	.	A	G	.	C	A	T	G	G	A										
8 Riccardia phyloides	C	.	.	.	G	C	T	C	T	G	A	A	.	T	A	C	.	A	T	T	.	A	G	.	C	A	T	G	G	A										
9 Sphaerocarpos dornalii	C	.	.	.	G	C	T	C	T	G	A	A	.	T	A	C	.	A	T	T	.	A	G	.	C	A	T	G	G	A										
10 Sphaerocarpos lezaricus	C	.	.	.	G	T	.	C	T	G	A	A	.	T	A	C	.	A	T	T	.	A	G	.	C	A	T	G	G	A										
11 Atrichum undulatum	T	.	.	.	G	C	T	C	T	G	A	A	.	T	A	C	.	A	T	T	.	A	G	.	C	A	T	G	G	A										
12 Bryum argenteum	T	.	.	.	G	C	T	C	T	G	A	A	.	T	A	C	.	A	T	T	.	A	G	.	C	A	T	G	G	A										
13 Eurythidum flans	T	.	.	.	G	C	T	C	T	G	A	A	.	T	A	C	.	A	T	T	.	A	G	.	C	A	T	G	G	A										
14 Funaria hygrometrica	T	.	.	.	G	C	T	C	T	G	A	A	.	T	A	C	.	A	T	T	.	A	G	.	C	A	T	G	G	A										
15 Lepadobryum pyriforme	T	.	.	.	G	C	T	C	T	G	A	A	.	T	A	C	.	A	T	T	.	A	G	.	C	A	T	G	G	A										
16 Mnium hornum	T	.	.	.	G	C	T	C	T	G	A	A	.	T	A	C	.	A	T	T	.	A	G	.	C	A	T	G	G	A										
17 Physcomitrium pyriforme	T	.	.	.	G	C	T	C	T	G	A	A	.	T	A	C	.	A	T	T	.	A	G	.	C	A	T	G	G	A										
18 Ptilothidium tomentosum	T	.	.	.	G	C	T	C	T	G	A	A	.	T	A	C	.	A	T	T	.	A	G	.	C	A	T	G	G	A										
19 Sphagnum cuspidatum	C	.	.	.	G	C	T	C	T	G	A	A	.	T	A	C	.	A	T	T	.	A	G	.	C	A	T	G	G	A										
20 Calypogeia angusta	T	.	.	.	G	C	T	C	T	G	A	A	.	T	A	C	.	A	T	T	.	A	G	.	C	A	T	G	G	A										
21 Conoclophium caritum	T	.	.	.	G	C	T	C	T	G	A	A	.	T	A	C	.	A	T	T	.	A	G	.	C	A	T	G	G	A										
22 Fescomitronia pusilla	T	.	.	.	G	C	T	C	T	G	A	A	.	T	A	C	.	A	T	T	.	A	G	.	C	A	T	G	G	A										
23 Marchantia polymorpha	C	.	.	.	G	C	T	C	T	G	A	A	.	T	A	C	.	A	T	T	.	A	G	.	C	A	T	G	G	A										
24 Riccia fluitans	C	.	.	.	G	C	T	C	T	G	A	A	.	T	A	C	.	A	T	T	.	A	G	.	C	A	T	G	G	A										
25 Scapania nemoralis	T	.	.	.	G	C	T	C	T	G	A	A	.	T	A	C	.	A	T	T	.	A	G	.	C	A	T	G	G	A										
26 Equisetum hyemale	T	.	.	.	G	C	T	C	T	G	A	A	.	T	A	C	.	A	T	T	.	A	G	.	C	A	T	G	G	A										
27 Equisetum robustum	T	.	.	.	G	C	T	C	T	G	A	A	.	T	A	C	.	A	T	T	.	A	G	.	C	A	T	G	G	A										
28 Isoetes engelmannii	T	.	.	.	G	C	T	C	T	G	A	A	.	T	A	C	.	A	T	T	.	A	G	.	C	A	T	G	G	A										
29 Huperzia lucidulum	T	.	.	.	G	C	T	C	T	G	A	A	.	T	A	C	.	A	T	T	.	A	G	.	C	A	T	G	G	A										
30 Lycopodium inundatum	T	.	.	.	G	C	T	C	T	G	A	A	.	T	A	C	.	A	T	T	.	A	G	.	C	A	T	G	G	A										
31 Lycopodium phlegmaria	T	.	.	.	G	C	T	C	T	G	A	A	.	T	A	C	.	A	T	T	.	A	G	.	C	A	T	G	G	A										
32 Lycopodium lucidulum	T	.	.	.	G	C	T	C	T	G	A	A	.	T	A	C	.	A	T	T	.	A	G	.	C	A	T	G	G	A										
33 Lycopodium tristachyum	T	.	.	.	G	C	T	C	T	G	A	A	.	T	A	C	.	A	T	T	.	A	G	.	C	A	T	G	G	A										
34 Selaginella umbrosa	T	.	.	.	G	C	T	C	T	G	A	A	.	T	A	C	.	A	T	T	.	A	G	.	C	A	T	G	G	A										
35 Selaginella vogelii	T	.	.	.	G	C	T	C	T	G	A	A	.	T	A	C	.	A	T	T	.	A	G	.	C	A	T	G	G	A										
36 Oryza sativa	C	.	.	.	G	C	T	C	T	G	G	A	.	T	A	C	.	A	T	T	.	A	G	.	C	A	T	G	G	A										
37 Zea mays	C	.	.	.	G	C	T	C	T	G	G	A	.	T	A	C	.	A	T	T	.	A	G	.	C	A	T	G	G	A										
38 Arabidopsis thaliana	C	.	.	.	G	C	T	C	T	G	G	A	.	T	A	C	.	A	T	T	.	A	G	.	C	A	T	G	G	A										
39 Glycine max	C	.	.	.	G	C	T	C	T	G	T	A	.	T	A	C	.	A	T	T	.	A	G	.	C	A	T	G	G	A										
40 Brachypodium distachyon	C	.	.	.	G	C	T	C	T	G	T	A	.	T	A	C	.	A	T	T	.	A	G	.	C	A	T	G	G	A										
41 Zinnia elegans	T	.	.	.	G	C	T	C	T	G	A	A	.	T	A	C	.	A	T	T	.	A	G	.	C	A	T	G	G	A										
42 Oenothera lutea	T	.	.	.	G	C	T	C	T	G	A	A	.	T	A	C	.	A	T	T	.	A	G	.	C	A	T	G	G	A										
43 Oenothera biennis	C	.	.	.	G	C	T	C	T	G	A	A	.	T	A	C	.	A	T	T	.	A	G	.	C	A	T	G	G	A										
44 Pinus luchuanensis	T	.	.	.	G	C	T	C	T	G	A	A	.	T	A	C	.	A	T	T	.	A	G	.	C	A	T	G	G	A										
45 Pinus wallichiana	T	.	.	.	G	C	T	C	T	G	A	A	.	T	A	C	.	A	T	T	.	A	G	.	C	A	T	G	G	A										
46 Nagebia nagebia	C	.	.	.	G	C	T	C	T	G	A	A	.	T	A	C	.	A	T	T	.	A	G	.	C	A	T	G	G	A										
47 Taxus mairei	C	.	.	.	G	C	T	C	T	G	A	A	.	T	A	C	.	A	T	T	.	A	G	.	C	A	T	G	G	A										
48 Adiantum radicans	T	.	.	.	G	C	T	C	T	G	A	A	.	T	A	C	.	A	T	T	.	A	G	.	C	A	T	G	G	A										
49 Adiantum radicans 2	T	.	.	.	G	C	T	C	T	G	A	A	.	T	A	C	.	A	T	T	.	A	G	.	C	A	T	G	G	A										
50 Ophioglossum petiolatum	C	.	.	.	G	C	T	?	T	G	A	A	.	T	A	C	.	G	T	T	.	A	G	.	C	A	T	G	G	A										
51 Osmunda cinnamomea	T	.	.	.	G	C	T	C	T	G	A	A	.	T	A	C	.	A	T	T	.	A	G	.	C	A	T	G	G	A										
52 Selaginella selaginoides	T	.	.	.	G	C	T	C	T	G	A	A	.	T	A	C	.	A	T	T	.	A	G	.	C	A	T	G	G	A										
53 Pellaea rotundifolia	T	.	.	.	G	C	T	C	T	G	A	A	.	T	A	C	.	A	T	T	.	A	G	.	C	A	T	G	G	A										
54 Chara foetida	T	.	.	.	G	C	T	C	T	G	A	A	.	T	A	C	.	A	T	T	.	A	G	.	C	A	T	G	G	A										
55 Ceteochorda scutellata	G	.	.	.	G	C	T	C	T	G	A	A	.	T	A	C	.	A	T	T	.	A	G	.	C	A	T	G	G	A										
56 Heterosporium radicans	C	.	.	.	G	C	T	C	T	G	A	A	.	T	A	C	.	A	T	T	.	A	G	.	C	A	T	G	G	A										
57 Nitella flexilis	C	.	.	.	G	C	T	C	T	G	A	A	.	T	A	C	.	A	T	T	.	A	G	.	C	A	T	G	G	A										
58 Nitella sp.	C	.	.	.	G	C	T	C	T	G	A	A	.	T	A	C	.	A	T	T	.	A	G	.	C	A	T	G	G	A										
59 Chara dimorpha rethardii	C	.	.	.	G	C	T	C	T	G	A	A	.	T	A	C	.	A	T	T	.	A	G	.	C	A	T	G	G	A										
60 Hydrocotyle radicans	C	.	.	.	G	C	T	C	T	G	A	A	.	T	A	C	.	A	T	T	.	A	G	.	C	A	T	G	G	A										
61 Peristichia pseudohyalocarpa	C	.	.	.	G	C	T	C	T	G	A	A	.	T	A	C	.	A	T	T	.	A	G	.	C	A	T	G	G	A										
62 Oxytropis discoides	T	C	.	.	G	C	C	.	T	G	A	T	.	C	T	T	T	T	G	C	.	A	G	.	C	A	T	G	G	A										
63 Pteris vittata	T	.	.	.	G	C	T	C	T	G	A	A	.	T	A	C	.	A	T	T	.	A	G	.	C	A	T	G	G	A										
64 Megaceros serriguttatus	T	.	.	.	G	C	T	C	T	G	A	A	.	T	A	C	.	A	T	T	.	A	G	.	C	A	T	G	G	A										
65 Notophysis bracteata	T	.	.	.	G	C	T	C	T	G	A	A	.	T	A	C	.	A	T	T	.	A	G	.	C	A	T	G	G	A										
66 Physcomitrella patens	T	.	.	.	G	C	T	C	T	G	A	A	.	T	A	C	.	A	T	T	.	A	G	.	C	A	T	G	G	A										
67 Sphagnum palustre	C	.	.	.	G	C	T	C	T	G	A	A	.	T	A	C	.	A	T	T	.	A	G	.	C	A	T	G	G	A										

	1171	1172	1173	1174	1175	1176	1177	1178	1179	1180	1181	1182	1183	1184	1185	1186	1187	1188	1189	1190	1191	1192	1193	1194	1195	1196	1197	1198	1199	1200
	0	0	0	0	5	0	3	4	4	0	4	0	0	5	0	0	4	0	5	5	6	0	0	5	0	5	0	0	0	0
1 Saccharomyces cerevisiae	A	A	G	G	A	-	C	Q	T	T	T	T	C	A	T	T	A	A	T	C	A	A	G	A	A	C	Q	A	A	A
2 Schizosaccharomyces pombe	A	A	G	G	A	-	T	Q	T	T	T	T	C	A	T	T	A	A	T	C	A	A	G	A	A	C	Q	A	A	A
3 Physcomitrella patens 2	A	A	G	G	A	-	T	Q	T	T	T	T	C	A	T	T	A	A	T	C	A	A	G	A	A	C	Q	A	A	A
4 Arabidopsis thaliana	A	A	G	G	A	-	T	Q	T	T	T	T	C	A	T	T	A	A	T	C	A	A	G	A	A	C	Q	A	A	A
5 Arabidopsis lyrata	A	A	G	G	A	-	T	Q	T	T	T	T	C	A	T	T	A	A	T	C	A	A	G	A	A	C	Q	A	A	A
6 Phlox sp.	A	A	G	G	A	-	T	Q	T	T	T	T	C	A	T	T	A	A	T	C	A	A	G	A	A	C	Q	A	A	A
7 Reboulia hemisphaerica	A	A	G	G	A	-	T	Q	T	T	T	T	C	A	T	T	A	A	T	C	A	A	G	A	A	C	Q	A	A	A
8 Riccia pinguis	A	A	G	G	A	-	T	Q	T	T	T	T	C	A	T	T	A	A	T	C	A	A	G	A	A	C	Q	A	A	A
9 Sphaerocarpos cornelli	A	A	G	G	A	-	T	Q	T	T	T	T	C	A	T	T	A	A	T	C	A	A	G	A	A	C	Q	A	A	A
10 Sphaerocarpos lasarus	A	A	G	G	A	-	T	Q	T	T	T	T	C	A	T	T	A	A	T	C	A	A	G	A	A	C	Q	A	A	A
11 Albidum undulatum	A	A	G	G	A	-	T	Q	T	T	T	T	C	A	T	T	A	A	T	C	A	A	G	A	A	C	Q	A	A	A
12 Bryum argenteum	A	A	G	G	A	-	T	Q	T	T	T	T	C	A	T	T	A	A	T	C	A	A	G	A	A	C	Q	A	A	A
13 Eurhynchium hiemale	A	A	G	G	A	-	T	Q	T	T	T	T	C	A	T	T	A	A	T	C	A	A	G	A	A	C	Q	A	A	A
14 Funaria hygrometrica	A	A	G	G	A	-	T	Q	T	T	T	T	C	A	T	T	A	A	T	C	A	A	G	A	A	C	Q	A	A	A
15 Leptobryum pyriforme	A	A	G	G	A	-	T	Q	T	T	T	T	C	A	T	T	A	A	T	C	A	A	G	A	A	C	Q	A	A	A
16 Mnium hornum	A	A	G	G	A	-	T	Q	T	T	T	T	C	A	T	T	A	A	T	C	A	A	G	A	A	C	Q	A	A	A
17 Physcomitrium pyriforme	A	A	G	G	A	-	T	Q	T	T	T	T	C	A	T	T	A	A	T	C	A	A	G	A	A	C	Q	A	A	A
18 Polytrichum formosum	A	A	G	G	A	-	T	Q	T	T	T	T	C	A	T	T	A	A	T	C	A	A	G	A	A	C	Q	A	A	A
19 Sphagnum cuspidatum	A	A	G	G	A	-	T	Q	T	T	T	T	C	A	T	T	A	A	T	C	A	A	G	A	A	C	Q	A	A	A
20 Calypogeia arguta	A	A	G	G	A	-	T	Q	T	T	T	T	C	A	T	T	A	A	T	C	A	A	G	A	A	C	Q	A	A	A
21 Conocephalum conicum	A	A	G	G	A	-	T	Q	T	T	T	T	C	A	T	T	A	A	T	C	A	A	G	A	A	C	Q	A	A	A
22 Foscaribria pusilla	A	A	G	G	A	-	T	Q	T	T	T	T	C	A	T	T	A	A	T	C	A	A	G	A	A	C	Q	A	A	A
23 Marchantia polymorpha	A	A	G	G	A	-	T	Q	T	T	T	T	C	A	T	T	A	A	T	C	A	A	G	A	A	C	Q	A	A	A
24 Riccia fluitans	A	A	G	G	A	-	T	Q	T	T	T	T	C	A	T	T	A	A	T	C	A	A	G	A	A	C	Q	A	A	A
25 Scleroparia nemoralis	A	A	G	G	A	-	T	Q	T	T	T	T	C	A	T	T	A	A	T	C	A	A	G	A	A	C	Q	A	A	A
26 Equisetum hyemale	A	A	G	G	A	-	T	Q	T	T	T	T	C	A	T	T	A	A	T	C	A	A	G	A	A	C	Q	A	A	A
27 Equisetum robustum	A	A	G	G	A	-	T	Q	T	T	T	T	C	A	T	T	A	A	T	C	A	A	G	A	A	C	Q	A	A	A
28 Isoetes engelmannii	A	A	G	G	A	-	T	Q	T	T	T	T	C	A	T	T	A	A	T	C	A	A	G	A	A	C	Q	A	A	A
29 Hyperbaena lucidulum	A	A	G	G	A	-	T	Q	T	T	T	T	C	A	T	T	A	A	T	C	A	A	G	A	A	C	Q	A	A	A
30 Lycopodium inundatum	A	A	G	G	A	-	T	Q	T	T	T	T	C	A	T	T	A	A	T	C	A	A	G	A	A	C	Q	A	A	A
31 Lycopodium phlegmaria	A	A	G	G	A	-	T	Q	T	T	T	T	C	A	T	T	A	A	T	C	A	A	G	A	A	C	Q	A	A	A
32 Lycopodium lucidulum	A	A	G	G	A	-	T	Q	T	T	T	T	C	A	T	T	A	A	T	C	A	A	G	A	A	C	Q	A	A	A
33 Lycopodium tristichium	A	A	G	G	A	-	T	Q	T	T	T	T	C	A	T	T	A	A	T	C	A	A	G	A	A	C	Q	A	A	A
34 Selaginella selaginoides	A	A	G	G	A	-	T	Q	T	T	T	T	C	A	T	T	A	A	T	C	A	A	G	A	A	C	Q	A	A	A
35 Selaginella selaginoides	A	A	G	G	A	-	T	Q	T	T	T	T	C	A	T	T	A	A	T	C	A	A	G	A	A	C	Q	A	A	A
36 Selaginella selaginoides	A	A	G	G	A	-	T	Q	T	T	T	T	C	A	T	T	A	A	T	C	A	A	G	A	A	C	Q	A	A	A
37 Oryza sativa	A	A	G	G	A	-	T	Q	T	T	T	T	C	A	T	T	A	A	T	C	A	A	G	A	A	C	Q	A	A	A
38 Zea mays	A	A	G	G	A	-	T	Q	T	T	T	T	C	A	T	T	A	A	T	C	A	A	G	A	A	C	Q	A	A	A
39 Arabidopsis thaliana	A	A	G	G	A	-	T	Q	T	T	T	T	C	A	T	T	A	A	T	C	A	A	G	A	A	C	Q	A	A	A
40 Oryza sativa	A	A	G	G	A	-	T	Q	T	T	T	T	C	A	T	T	A	A	T	C	A	A	G	A	A	C	Q	A	A	A
41 Sinapis alba	A	A	G	G	A	-	T	Q	T	T	T	T	C	A	T	T	A	A	T	C	A	A	G	A	A	C	Q	A	A	A
42 Zinnia elegans	A	A	G	G	A	-	T	Q	T	T	T	T	C	A	T	T	A	A	T	C	A	A	G	A	A	C	Q	A	A	A
43 Arabidopsis thaliana	A	A	G	G	A	-	T	Q	T	T	T	T	C	A	T	T	A	A	T	C	A	A	G	A	A	C	Q	A	A	A
44 Arabidopsis thaliana	A	A	G	G	A	-	T	Q	T	T	T	T	C	A	T	T	A	A	T	C	A	A	G	A	A	C	Q	A	A	A
45 Arabidopsis thaliana	A	A	G	G	A	-	T	Q	T	T	T	T	C	A	T	T	A	A	T	C	A	A	G	A	A	C	Q	A	A	A
46 Arabidopsis thaliana	A	A	G	G	A	-	T	Q	T	T	T	T	C	A	T	T	A	A	T	C	A	A	G	A	A	C	Q	A	A	A
47 Arabidopsis thaliana	A	A	G	G	A	-	T	Q	T	T	T	T	C	A	T	T	A	A	T	C	A	A	G	A	A	C	Q	A	A	A
48 Arabidopsis thaliana	A	A	G	G	A	-	T	Q	T	T	T	T	C	A	T	T	A	A	T	C	A	A	G	A	A	C	Q	A	A	A
49 Arabidopsis thaliana	A	A	G	G	A	-	T	Q	T	T	T	T	C	A	T	T	A	A	T	C	A	A	G	A	A	C	Q	A	A	A
50 Arabidopsis thaliana	A	A	G	G	A	-	T	Q	T	T	T	T	C	A	T	T	A	A	T	C	A	A	G	A	A	C	Q	A	A	A
51 Arabidopsis thaliana	A	A	G	G	A	-	T	Q	T	T	T	T	C	A	T	T	A	A	T	C	A	A	G	A	A	C	Q	A	A	A
52 Arabidopsis thaliana	A	A	G	G	A	-	T	Q	T	T	T	T	C	A	T	T	A	A	T	C	A	A	G	A	A	C	Q	A	A	A
53 Arabidopsis thaliana	A	A	G	G	A	-	T	Q	T	T	T	T	C	A	T	T	A	A	T	C	A	A	G	A	A	C	Q	A	A	A
54 Arabidopsis thaliana	A	A	G	G	A	-	T	Q	T	T	T	T	C	A	T	T	A	A	T	C	A	A	G	A	A	C	Q	A	A	A
55 Arabidopsis thaliana	A	A	G	G	A	-	T	Q	T	T	T	T	C	A	T	T	A	A	T	C	A	A	G	A	A	C	Q	A	A	A
56 Arabidopsis thaliana	A	A	G	G	A	-	T	Q	T	T	T	T	C	A	T	T	A	A	T	C	A	A	G	A	A	C	Q	A	A	A
57 Arabidopsis thaliana	A	A	G	G	A	-	T	Q	T	T	T	T	C	A	T	T	A	A	T	C	A	A	G	A	A	C	Q	A	A	A
58 Arabidopsis thaliana	A	A	G	G	A	-	T	Q	T	T	T	T	C	A	T	T	A	A	T	C	A	A	G	A	A	C	Q	A	A	A
59 Arabidopsis thaliana	A	A	G	G	A	-	T	Q	T	T	T	T	C	A	T	T	A	A	T	C	A	A	G	A	A	C	Q	A	A	A
60 Arabidopsis thaliana	A	A	G	G	A	-	T	Q	T	T	T	T	C	A	T	T	A	A	T	C	A	A	G	A	A	C	Q	A	A	A
61 Arabidopsis thaliana	A	A	G	G	A	-	T	Q	T	T	T	T	C	A	T	T	A	A	T	C	A	A	G	A	A	C	Q	A	A	A
62 Arabidopsis thaliana	A	A	G	G	A	-	T	Q	T	T	T	T	C	A	T	T	A	A	T	C	A	A	G	A	A	C	Q	A	A	A
63 Arabidopsis thaliana	A	A	G	G	A	-	T	Q	T	T	T	T	C	A	T	T	A	A	T	C	A	A	G	A	A	C	Q	A	A	A
64 Arabidopsis thaliana	A	A	G	G	A	-	T	Q	T	T	T	T	C	A	T	T	A	A	T	C	A	A	G	A	A	C	Q	A	A	A
65 Arabidopsis thaliana	A	A	G	G	A	-	T	Q	T	T	T	T	C	A	T	T	A	A	T	C	A	A	G	A	A	C	Q	A	A	A
66 Arabidopsis thaliana	A	A	G	G	A	-	T	Q	T	T	T	T	C	A	T	T	A	A	T	C	A	A	G	A	A	C	Q	A	A	A
67 Arabidopsis thaliana	A	A	G	G	A	-	T	Q	T	T	T	T	C	A	T	T	A	A	T	C	A	A	G	A	A	C	Q	A	A	A

	1201	1202	1203	1204	1205	1206	1207	1208	1209	1270	1271	1272	1273	1274	1275	1276	1277	1278	1279	1280	1281	1282	1283	1284	1285	1286	1287	1288	1289	1290	1291	1292	1293	1294	1295	1296	1297	1298	1299	1300	
	0	3	0	0	0	3	0	5	1	3	0	2	2	3	2	0	0	0	4	0	0	0	1	1	3	4	0	1	2	3											
1 Saccharomyces cerevisiae	C	T	A	G	-	-	A	T	C	G	G	G	T	Q	Q	T	Q	T	T	-	-	-	T	T	T	T	-	T	A	A											
2 Schizosaccharomyces pombe	C	T	A	G	G	Q	A	T	C	G	G	G	C	A	A	T	Q	T	T	-	-	-	T	C	A	T	T	T	A	T											
3 Physcomitrella patens 2	C	T	A	G	G	G	A	T	T	G	G	C	G	Q	A	T	Q	T	T	-	-	-	A	C	T	T	-	T	Q	A											
4 Arabidopsis thaliana	C	T	A	G	G	Q	A	T	T	Q	Q	C	Q	Q	A	T	Q	T	T	-	-	-	A	A	T	T	-	T	Q	A											
5 Arabidopsis lyrata	C	T	A	G	G	Q	A	T	T	Q	Q	C	Q	Q	A	T	Q	T	T	-	-	-	A	A	T	T	-	T	Q	A											
6 Poa annua	C	T	A	G	G	Q	A	T	T	Q	Q	C	Q	Q	A	T	Q	T	T	-	-	-	A	A	T	T	-	T	Q	A											
7 Rabaia hamaphysalis	C	T	A	G	G	Q	A	T	C	Q	Q	C	Q	Q	A	T	Q	T	T	-	-	-	Q	A	T	T	-	A	Q	A											
8 Ricarda pinguis	C	T	A	G	G	Q	A	T	T	Q	Q	C	Q	Q	A	T	Q	T	T	-	-	-	A	A	T	T	-	T	Q	A											
9 Sphaerocarpos darwini	C	T	A	G	G	Q	A	T	T	Q	Q	C	Q	Q	A	T	Q	T	T	-	-	-	A	A	T	T	-	T	Q	A											
10 Sphaerocarpos lasarus	C	T	A	G	G	Q	A	T	T	Q	Q	C	Q	Q	A	T	Q	T	T	-	-	-	A	A	T	T	-	T	Q	A											
11 Abrikum undulatum	C	T	A	G	G	Q	A	T	T	T	Q	C	Q	Q	Q	T	Q	T	T	-	-	-	A	A	T	T	-	T	Q	A											
12 Bryum argenteum	C	T	A	G	G	A	A	T	T	Q	Q	C	Q	Q	Q	T	Q	T	T	-	-	-	Q	C	T	T	-	T	Q	A											
13 Euthymium hians	C	T	A	G	G	Q	A	C	T	T	Q	C	Q	Q	C	T	Q	T	T	-	-	-	C	A	T	T	-	T	Q	A											
14 Ranaria hygrometrica	C	T	A	G	G	Q	A	T	T	Q	G	C	Q	Q	A	T	Q	T	T	-	-	-	A	C	T	T	-	T	Q	A											
15 Leptobryum pyriforme	C	T	A	G	G	Q	A	T	T	Q	Q	C	Q	Q	Q	T	Q	T	T	-	-	-	Q	A	T	T	-	T	Q	A											
16 Mnium hornum	C	T	A	G	G	Q	A	T	T	Q	Q	C	Q	Q	Q	T	Q	T	T	-	-	-	Q	A	T	T	-	C	Q	A											
17 Physcomitrium pyriforme	C	T	A	G	G	Q	A	T	T	Q	Q	C	Q	Q	A	T	Q	T	T	-	-	-	A	C	T	T	-	T	Q	A											
18 Polytrichum formosum	C	T	A	G	G	Q	A	T	T	C	Q	C	Q	Q	Q	T	Q	T	T	-	-	-	A	A	T	T	-	T	Q	A											
19 Sphagnum cuspidatum	C	T	A	G	G	Q	A	T	C	Q	Q	C	Q	Q	A	T	Q	T	T	-	-	-	Q	A	A	T	-	C	Q	A											
20 Calypogeia angusta	C	T	A	G	G	Q	A	T	T	Q	Q	C	Q	Q	A	T	Q	T	T	-	-	-	A	A	T	T	-	T	Q	A											
21 Conocarpus ciliatus	C	T	A	G	G	Q	A	T	C	Q	Q	C	Q	Q	A	T	Q	T	T	-	-	-	T	C	T	T	T	T	Q	A											
22 Fossombronia pusilla	C	T	A	G	G	Q	A	T	T	Q	Q	C	Q	Q	A	T	Q	T	T	-	-	-	A	A	T	T	-	T	Q	A											
23 Marchantia polymorpha	C	T	A	G	G	Q	A	T	C	Q	Q	C	Q	Q	A	T	Q	T	T	-	-	-	A	A	T	T	-	T	Q	A											
24 Riccia fluitans	C	T	A	G	G	Q	A	T	C	Q	Q	C	Q	Q	A	T	Q	T	T	-	-	-	T	Q	A	T	T	-	A	Q	A										
25 Scapania nemorea	C	T	A	G	G	Q	A	T	T	Q	Q	C	Q	Q	A	T	Q	T	T	-	-	-	A	A	T	T	-	T	Q	A											
26 Equisetum hyemale	C	T	A	G	G	Q	A	T	T	Q	Q	C	A	Q	A	T	Q	T	T	-	-	-	A	C	T	T	-	C	Q	A											
27 Equisetum robustum	C	T	A	G	G	Q	A	T	T	Q	Q	C	A	Q	A	T	Q	T	T	-	-	-	A	C	T	T	-	C	Q	A											
28 Isoetes engelmannii	C	T	A	G	G	Q	A	T	C	Q	Q	C	Q	Q	A	T	Q	T	T	-	-	-	A	C	T	T	-	C	Q	A											
29 Huperzia lucidulum	C	T	A	G	G	Q	A	T	C	Q	Q	C	Q	Q	Q	T	Q	T	T	-	-	-	A	A	T	T	-	C	Q	A											
30 Lycopodium truncatum	C	T	A	G	G	Q	A	T	C	Q	Q	C	Q	Q	Q	T	Q	T	T	-	-	-	?	A	T	T	-	C	Q	A											
31 Lycopodium phlegmaria	C	T	A	G	G	Q	A	T	C	Q	Q	C	Q	Q	Q	T	Q	T	T	-	-	-	A	A	T	T	-	C	Q	A											
32 Lycopodium lasiocarpum	C	T	A	G	G	Q	A	T	C	Q	Q	C	Q	Q	Q	T	Q	T	T	-	-	-	A	A	T	T	-	C	Q	A											
33 Lycopodium tristichyum	C	T	A	G	G	Q	A	T	C	Q	Q	C	Q	Q	Q	T	Q	T	T	-	-	-	A	A	T	T	-	C	Q	A											
34 Selaginella selaginoides	C	T	A	G	G	Q	A	T	T	Q	Q	C	Q	Q	A	T	Q	T	T	-	-	-	A	T	T	T	-	Q	Q	A											
35 Selaginella selaginoides	C	T	A	G	G	Q	A	T	T	Q	Q	C	Q	Q	A	T	Q	T	T	-	-	-	A	T	T	T	-	Q	Q	A											
36 Dryas octopetala	C	C	A	Q	Q	Q	A	T	C	Q	Q	C	Q	Q	A	T	Q	T	T	-	-	-	Q	C	T	T	-	A	T	A											
37 Zizia aurea	C	C	A	Q	Q	Q	A	T	C	A	Q	C	Q	Q	T	-	Q	T	T	-	-	-	A	C	T	A	-	A	T	A											
38 Arabidopsis thaliana	C	G	A	Q	Q	Q	A	T	C	A	Q	C	Q	Q	A	T	Q	T	T	-	-	-	Q	C	T	T	-	A	T	A											
39 Glycine max	C	C	A	Q	Q	Q	A	T	C	A	Q	C	Q	Q	A	T	Q	T	T	-	-	-	Q	C	T	T	-	T	T	A											
40 Sinapis alba	C	C	A	Q	Q	Q	A	T	C	A	Q	C	Q	Q	A	T	Q	T	T	-	-	-	Q	C	T	T	-	T	T	A											
41 Zinnia elegans	C	C	A	Q	Q	Q	A	T	C	A	Q	C	Q	Q	A	T	Q	T	T	-	-	-	Q	C	T	C	-	T	A	A											
42 Ononis spinosa	C	C	A	Q	Q	Q	A	T	C	A	Q	C	Q	Q	A	T	Q	T	T	-	-	-	Q	C	T	T	-	T	T	A											
43 Girardinia spiralis	C	T	A	G	G	Q	A	T	C	Q	Q	C	Q	Q	A	T	Q	T	T	-	-	-	Q	C	T	T	-	T	A	A											
44 Pinus taeda	C	C	A	Q	Q	Q	A	T	C	A	Q	C	Q	Q	A	T	Q	T	T	-	-	-	Q	C	T	C	-	T	A	A											
45 Pinus strobus	C	C	A	Q	Q	Q	A	T	C	A	Q	C	Q	Q	A	T	Q	T	T	-	-	-	Q	C	T	C	-	T	A	A											
46 Nipah palm	C	C	A	Q	Q	Q	A	T	C	A	Q	C	Q	Q	A	T	Q	T	T	-	-	-	Q	C	T	C	-	T	A	A											
47 Taxus canadensis	C	C	A	Q	Q	Q	A	T	C	A	Q	C	Q	Q	A	T	Q	T	T	-	-	-	Q	C	T	C	-	T	A	A											
48 Adiantum redticatum	C	T	A	G	G	Q	A	T	T	Q	Q	C	Q	Q	A	T	Q	T	T	-	-	-	A	C	T	T	-	T	Q	A											
49 Adiantum redticatum 2	C	T	A	G	G	Q	A	T	T	Q	Q	C	Q	Q	A	T	Q	T	T	-	-	-	A	C	T	T	-	T	Q	A											
50 Ophioglossum petiolatum	C	T	A	G	G	Q	A	T	C	Q	Q	C	Q	Q	A	T	Q	T	T	-	-	-	A	C	T	T	-	T	Q	A											
51 Osmunda cinnamomea	C	T	A	G	G	Q	A	T	T	Q	Q	C	Q	Q	A	T	Q	T	T	-	-	-	A	C	T	T	-	C	A	A											
52 Selaginella selaginoides	C	T	A	G	G	Q	A	T	T	Q	Q	C	Q	Q	A	T	Q	T	T	-	-	-	A	C	T	T	-	C	Q	A											
53 Pteridium aquilinum	C	T	A	G	G	Q	A	T	C	Q	Q	C	Q	Q	A	T	Q	T	T	-	-	-	A	C	T	T	-	T	Q	A											
54 Chara corallina	C	T	A	G	G	Q	A	T	T	Q	Q	C	Q	Q	A	T	Q	T	C	-	-	-	T	A	T	T	-	G	Q	A											
55 Coloclelea sulcata	C	T	A	G	G	Q	A	T	C	Q	Q	C	Q	Q	A	T	Q	T	T	-	-	-	A	A	T	T	-	T	Q	A											
56 Hebesaridium laevis	C	T	A	G	G	Q	A	T	T	Q	Q	C	Q	Q	A	T	Q	T	T	-	-	-	A	A	T	T	-	T	Q	A											
57 Nitella flexilis	C	T	A	G	G	Q	A	T	T	Q	Q	C	Q	Q	A	T	Q	T	C	-	-	-	T	A	T	T	-	T	Q	A											
58 Nitella sp	C	T	A	G	G	Q	A	T	T	Q	Q	C	Q	Q	A	T	Q	T	C	-	-	-	T	A	T	T	-	T	Q	A											
59 Chara domingensis rethardii	C	T	A	G	G	Q	A	T	T	Q	Q	C	A	Q	A	T	Q	T	T	-	-	-	C	T	T	T	-	T	Q	A											
60 Hydrodictyon reticulatum	C	T	A	G	G	Q	A	T	T	Q	Q	C	Q	Q	A	T	Q	T	T	-	-	-	T	T	T	T	-	T	Q	A											
61 Paratetraspora pseudocubensis	C	T	A	G	G	Q	A	T	T	Q	Q	C	Q	Q	A	T	Q	T	T	-	-	-	T	T	A	T	-	C	Q	A											
62 Dictyosphaerula discoides	C	C	A	Q	Q	Q	A	T	C	Q	Q	T	T	A	A	A	A	T	T	-	-	-	-	-	T	T	-	C	-	-											
63 Pteris vittata	C	T	A	G	G	Q	A	T	T	Q	Q	C	Q	Q	A	T	Q	T	T	-	-	-	A	C	T	T	-	T	Q	A											
64 Megaceros senigallensis	C	T	A	G	G	Q	A	T	T	Q	Q	C	Q	Q	A	T	Q	T	T	-	-	-	A	A	T	T	-	A	Q	A											
65 Notophylax brevis	C	T	A	G	G	Q	A	T	T	Q	Q	C	Q	Q	Q	T	Q	T	T	-	-	-	A	A	T	-	A	A	Q	A											
66 Physcomitrella patens	C	T	A	G	G	Q	A	T	T	Q	Q	C	Q	Q	A	T	Q	T	T	-	-	-	A	C	T	T	-	T	Q	A											
67 Sphagnum palustre	C	T	A	G	G	Q	A	T	C</																																

	1291	1292	1293	1294	1295	1296	1297	1298	1299	1300	1301	1302	1303	1304	1305	1306	1307	1308	1309	1310	1311	1312	1313	1314	1315	1316	1317	1318	1319	1320
	2	0	6	6	0	0	0	6	2	3	2	2	2	4	1	6	3	0	5	6	6	6	6	4	3	0	0	2	6	
1 Saccharomyces cerevisiae	T	.	G	A	C	C	C	A	C	T	C	G	G	T	.	A	C	.	C	T	T	A	.	.	C	G
2 Schizosaccharomyces pombe	C	.	G	A	C	T	T	G	C	T	C	G	G	C	.	A	C	.	C	T	T	A	.	.	C	G
3 Physcomitrella patens 2	T	.	G	A	C	T	C	G	G	C	C	A	G	C	.	A	C	.	C	T	T	A	.	.	T	G
4 Arthrocosme agrestis	T	.	G	A	C	G	T	C	G	C	C	A	G	C	.	A	C	.	C	T	T	A	.	.	T	G
5 Arthrocosme lasio	T	.	G	A	C	T	C	C	G	C	C	A	G	C	.	A	C	.	C	T	T	A	.	.	T	G
6 Pellaea epiphylla	T	.	G	A	C	T	C	C	G	C	C	A	G	C	.	A	C	.	C	T	T	A	.	.	T	G
7 Reboulia hemisphaerica	T	.	G	A	C	T	C	C	G	C	C	A	G	C	.	A	C	.	C	T	T	A	.	.	T	G
8 Riccardia pinguis	T	.	G	A	C	T	C	C	G	C	C	A	G	C	.	A	C	.	C	T	T	A	.	.	T	G
9 Sphaerocarpos dornalii	T	.	G	A	C	T	C	C	G	C	C	A	G	C	.	A	C	.	C	T	T	A	.	.	T	G
10 Sphaerocarpos lasarus	T	.	G	A	C	T	C	C	G	.	C	A	G	C	.	A	C	.	C	T	T	A	.	.	T	G
11 Atrichum undulatum	T	.	G	A	C	C	C	C	G	C	A	A	G	C	.	A	C	.	C	T	T	A	.	.	T	G
12 Bryum argenteum	T	.	G	A	C	C	C	C	G	C	C	A	G	C	.	A	C	.	C	T	T	A	.	.	T	G
13 Eurythyzium hiemale	T	.	G	A	C	C	C	C	G	C	C	A	G	C	.	A	C	.	C	T	T	A	.	.	T	G
14 Funaria hygrometrica	T	.	G	A	C	T	C	C	G	C	C	A	G	C	.	A	C	.	C	T	T	A	.	.	T	G
15 Lophozium pyriforme	T	.	G	A	C	C	C	C	G	C	C	A	G	C	.	A	C	.	C	T	T	A	.	.	T	G
16 Lophozium pyriforme	T	.	G	A	C	C	C	C	G	C	C	A	G	C	.	A	C	.	C	T	T	A	.	.	T	G
17 Physcomitrium pyriforme	T	.	G	A	C	T	C	C	G	C	C	A	G	C	.	A	C	.	C	T	T	A	.	.	T	G
18 Polytrichum commune	T	.	G	A	C	C	C	C	G	C	G	A	G	C	.	A	C	.	C	T	T	A	.	.	T	G
19 Sphagnum cuspidatum	T	.	G	A	C	C	C	C	G	C	G	A	G	C	.	A	C	.	C	T	T	A	.	.	T	G
20 Calypogeia angusta	T	.	G	A	C	T	T	C	G	C	C	A	G	C	.	A	C	.	C	T	T	A	.	.	T	G
21 Conocephalum conicum	T	.	G	A	C	T	G	C	G	C	C	A	G	C	.	A	C	.	C	T	T	A	.	.	T	G
22 Fossombronia pusilla	T	.	G	A	C	T	C	C	G	C	C	A	G	C	.	A	C	.	C	T	T	A	.	.	T	G
23 Marchantia polymorpha	T	.	G	A	C	T	C	C	G	C	C	A	G	C	.	A	C	.	C	T	T	A	.	.	T	G
24 Riccia fluitans	T	.	G	A	C	T	C	C	G	C	C	A	G	C	.	A	C	.	C	T	T	A	.	.	T	G
25 Sphaerocarpos dornalii	T	.	G	A	C	T	C	C	G	C	C	A	G	C	.	A	C	.	C	T	T	A	.	.	T	G
26 Equisetum hyemale	T	.	G	A	C	T	C	T	G	C	C	A	G	C	.	A	C	.	C	T	T	A	.	.	T	G
27 Equisetum robustum	T	.	G	A	C	T	C	T	G	C	C	A	G	C	.	A	C	.	C	T	T	A	.	.	T	G
28 Isoetes engelmannii	T	.	G	A	C	T	C	C	G	C	C	A	G	C	.	A	C	.	C	T	T	A	.	.	T	G
29 Hypoxis lucidulum	T	.	G	A	C	C	C	C	G	C	C	A	G	C	.	A	C	.	C	T	T	A	.	.	T	G
30 Lycopodium lucidulum	T	.	G	A	C	C	C	C	G	C	C	A	G	C	.	A	C	.	C	T	T	A	.	.	T	G
31 Lycopodium ptiligera	T	.	G	A	C	C	C	C	G	C	C	A	G	C	.	A	C	.	C	T	T	A	.	.	T	G
32 Lycopodium lucidulum	T	.	G	A	C	C	C	C	G	C	C	A	G	C	.	A	C	.	C	T	T	A	.	.	T	G
33 Lycopodium trilete	T	.	G	A	C	C	C	C	G	C	C	A	G	C	.	A	C	.	C	T	T	A	.	.	T	G
34 Selaginella selaginoides	T	.	G	A	C	T	C	C	G	C	C	A	G	C	.	A	C	.	C	T	T	A	.	.	T	G
35 Selaginella selaginoides	T	.	G	A	C	A	C	C	G	C	C	A	G	C	.	A	C	.	C	T	T	A	.	.	T	G
36 Oryza sativa	G	.	G	A	C	T	C	C	G	C	C	A	G	C	.	A	C	.	C	T	T	A	.	.	T	G
37 Zea mays	G	.	G	A	C	C	C	C	G	C	T	G	G	C	.	A	C	.	C	T	T	A	.	.	T	G
38 Arabidopsis thaliana	G	.	G	A	C	T	C	C	G	C	T	G	G	C	.	A	C	.	C	T	T	A	.	.	T	G
39 Glycine max	G	.	G	A	C	T	C	C	G	C	T	G	G	C	.	A	C	.	C	T	T	A	.	.	T	G
40 Binaria alba	G	.	G	A	C	T	C	C	G	C	T	G	G	C	.	A	C	.	C	T	T	A	.	.	T	G
41 Zinnia purpurea	G	.	G	A	C	T	C	C	G	C	C	A	G	C	.	A	C	.	C	T	T	A	.	.	T	G
42 Arabidopsis thaliana	G	.	G	A	C	T	C	C	G	C	T	G	G	C	.	A	C	.	C	T	T	A	.	.	T	G
43 Arabidopsis thaliana	G	.	G	A	C	T	C	C	G	C	T	G	G	C	.	A	C	.	C	T	T	A	.	.	T	G
44 Pinus taeda	G	.	G	A	C	T	C	C	G	C	C	A	G	C	.	A	C	.	C	T	T	A	.	.	T	G
45 Pinus taeda	G	.	G	A	C	T	C	C	G	C	C	A	G	C	.	A	C	.	C	T	T	A	.	.	T	G
46 Pinus taeda	G	.	G	A	C	T	C	C	G	C	C	A	G	C	.	A	C	.	C	T	T	A	.	.	T	G
47 Pinus taeda	G	.	G	A	C	T	C	C	G	C	C	A	G	C	.	A	C	.	C	T	T	A	.	.	T	G
48 Pinus taeda	G	.	G	A	C	T	C	C	G	C	C	A	G	C	.	A	C	.	C	T	T	A	.	.	T	G
49 Pinus taeda	G	.	G	A	C	T	C	C	G	C	C	A	G	C	.	A	C	.	C	T	T	A	.	.	T	G
50 Pinus taeda	G	.	G	A	C	T	C	C	G	C	C	A	G	C	.	A	C	.	C	T	T	A	.	.	T	G
51 Pinus taeda	G	.	G	A	C	T	C	C	G	C	C	A	G	C	.	A	C	.	C	T	T	A	.	.	T	G
52 Pinus taeda	G	.	G	A	C	T	C	C	G	C	C	A	G	C	.	A	C	.	C	T	T	A	.	.	T	G
53 Pinus taeda	G	.	G	A	C	T	C	C	G	C	C	A	G	C	.	A	C	.	C	T	T	A	.	.	T	G
54 Pinus taeda	G	.	G	A	C	T	C	C	G	C	C	A	G	C	.	A	C	.	C	T	T	A	.	.	T	G
55 Pinus taeda	G	.	G	A	C	T	C	C	G	C	C	A	G	C	.	A	C	.	C	T	T	A	.	.	T	G
56 Pinus taeda	G	.	G	A	C	T	C	C	G	C	C	A	G	C	.	A	C	.	C	T	T	A	.	.	T	G
57 Pinus taeda	G	.	G	A	C	T	C	C	G	C	C	A	G	C	.	A	C	.	C	T	T	A	.	.	T	G
58 Pinus taeda	G	.	G	A	C	T	C	C	G	C	C	A	G	C	.	A	C	.	C	T	T	A	.	.	T	G
59 Pinus taeda	G	.	G	A	C	T	C	C	G	C	C	A	G	C	.	A	C	.	C	T	T	A	.	.	T	G
60 Pinus taeda	G	.	G	A	C	T	C	C	G	C	C	A	G	C	.	A	C	.	C	T	T	A	.	.	T	G
61 Pinus taeda	G	.	G	A	C	T	C	C	G	C	C	A	G	C	.	A	C	.	C	T	T	A	.	.	T	G
62 Pinus taeda	G	.	G	A	C	T	C	C	G	C	C	A	G	C	.	A	C	.	C	T	T	A	.	.	T	G
63 Pinus taeda	G	.	G	A	C	T	C	C	G	C	C	A	G	C	.	A	C	.	C	T	T	A	.	.	T	G
64 Pinus taeda	G	.	G	A	C	T	C	C	G	C	C	A	G	C	.	A	C	.	C	T	T	A	.	.	T	G
65 Pinus taeda	G	.	G	A	C	T	C	C	G	C	C	A	G	C	.	A	C	.	C	T	T	A	.	.	T	G
66 Pinus taeda	G	.	G	A	C	T	C	C	G	C	C	A	G	C	.	A	C	.	C	T	T	A	.	.	T	G
67 Pinus taeda	G	.	G	A	C	T	C	C	G	C	C	A	G	C	.	A	C	.	C	T	T	A	.	.	T	G

	1441	1442	1443	1444	1448	1449	1447	1450	1449	1450	1491	1452	1453	1494	1455	1457	1456	1459	1460	1461	1462	1463	1464	1468	1466	1467	1468	1469	1470		
	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
1	Saccharomyces cerevisiae	C	A	C	G	G	G	G	A	A	A	C	T	C	A	C	C	A	G	G	T	C	C	A	G	A	C	A	C	A	
2	Schizosaccharomyces pombe	C	A	C	G	G	G	G	A	A	A	C	T	C	A	C	C	A	G	G	T	C	C	A	G	A	C	A	T	A	G
3	Physcomitrella patens 2	C	A	C	G	G	G	G	A	A	A	C	T	T	A	C	C	A	G	G	T	C	C	A	G	A	C	A	T	A	G
4	Anthoceros agrestis	C	A	C	G	G	G	G	A	A	A	C	T	T	A	C	C	A	G	G	T	C	C	A	G	A	C	A	T	A	G
5	Anthoceros laevis	C	A	C	G	G	G	G	A	A	A	C	T	T	A	C	C	A	G	G	T	C	C	A	G	A	C	A	T	A	G
6	Pellia epiphylla	C	A	C	G	G	G	G	A	A	A	C	T	T	A	C	C	A	G	G	T	C	C	A	G	A	C	A	T	A	G
7	Reboulia hemisphaerica	C	A	C	G	G	G	G	A	A	A	C	T	T	A	C	C	A	G	G	T	C	C	A	G	A	C	A	T	A	G
8	Riocardia pinguis	C	A	C	G	G	G	G	A	A	A	C	T	T	A	C	C	A	G	G	T	C	C	A	G	A	C	A	T	A	G
9	Sphaerocarpos donnellii	C	A	C	G	G	G	G	A	A	A	C	T	T	A	C	C	A	G	G	T	C	C	A	G	A	C	A	T	A	G
10	Sphaerocarpos letaricus	C	A	C	G	G	G	G	A	A	A	C	T	T	A	C	C	A	G	G	T	C	C	A	G	A	C	A	T	A	G
11	Atrichum undulatum	C	A	C	G	G	G	G	A	A	A	C	T	T	A	C	C	A	G	G	T	C	C	A	G	A	C	A	T	A	G
12	Bryum argenteum	C	A	C	G	G	G	G	A	A	A	C	T	T	A	C	C	A	G	G	T	C	C	A	G	A	C	A	T	A	G
13	Eurymedusa hians	C	A	C	G	G	G	G	A	A	A	C	T	T	A	C	C	A	G	G	T	C	C	A	G	A	C	A	T	A	G
14	Puraria hygrometrica	C	A	C	G	G	G	G	A	A	A	C	T	T	A	C	C	A	G	G	T	C	C	A	G	A	C	A	T	A	G
15	Lepidobryum pyriforme	C	A	C	G	G	G	G	A	A	A	C	T	T	A	C	C	A	G	G	T	C	C	A	G	A	C	A	T	A	G
16	Isidium haerum	C	A	C	G	G	G	G	A	A	A	C	T	T	A	C	C	A	G	G	T	C	C	A	G	A	C	A	T	A	G
17	Physcomitrium pyriforme	C	A	C	G	G	G	G	A	A	A	C	T	T	A	C	C	A	G	G	T	C	C	A	G	A	C	A	T	A	G
18	Polytrichum formosum	C	A	C	G	G	G	G	A	A	A	C	T	T	A	C	C	A	G	G	T	C	C	A	G	A	C	A	T	A	G
19	Sphagnum cuspidatum	C	A	C	G	G	G	G	A	A	A	C	T	T	A	C	C	A	G	G	T	C	C	A	G	A	C	A	T	A	G
20	Calypogeia arguta	C	A	C	G	G	G	G	A	A	A	C	T	T	A	C	C	A	G	G	T	C	C	A	G	A	C	A	T	A	G
21	Conocephalum conicum	C	A	C	G	G	G	G	A	A	A	C	T	T	A	C	C	A	G	G	T	C	C	A	G	A	C	A	T	A	G
22	Fesemburgia pusilla	C	A	C	G	G	G	G	A	A	A	C	T	T	A	C	C	A	G	G	T	C	C	A	G	A	C	A	T	A	G
23	Marsipposiphonia polymorpha	C	A	C	G	G	G	G	A	A	A	C	T	T	A	C	C	A	G	G	T	C	C	A	G	A	C	A	T	A	G
24	Riccia fluitans	C	A	C	G	G	G	G	A	A	A	C	T	T	A	C	C	A	G	G	T	C	C	A	G	A	C	A	T	A	G
25	Soparia nemoralis	C	A	C	G	G	G	G	A	A	A	C	T	T	A	C	C	A	G	G	T	C	C	A	G	A	C	A	T	A	G
26	Equisetum byemale	C	A	C	G	G	G	G	A	A	A	C	T	T	A	C	C	A	G	G	T	C	C	A	G	A	C	A	T	A	G
27	Equisetum robustum	C	A	C	G	G	G	G	A	A	A	C	T	T	A	C	C	A	G	G	T	C	C	A	G	A	C	A	T	A	G
28	Isetes engelmannii	C	A	C	G	G	G	G	A	A	A	C	T	T	A	C	C	A	G	G	T	C	C	A	G	A	C	A	T	A	G
29	Huperzia lucidulum	C	A	C	G	G	G	G	A	A	A	C	T	T	A	C	C	A	G	G	T	C	C	A	G	A	C	A	T	A	G
30	Lycopodium inundatum	C	A	C	G	G	G	G	A	A	A	C	T	T	A	C	C	A	G	G	T	C	C	A	G	A	C	A	T	A	G
31	Lycopodium phlegmaria	C	A	C	G	G	G	G	A	A	A	C	T	T	A	C	C	A	G	G	T	C	C	A	G	A	C	A	T	A	G
32	Lycopodium lasiolobum	C	A	C	G	G	G	G	A	A	A	C	T	T	A	C	C	A	G	G	T	C	C	A	G	A	C	A	T	A	G
33	Lycopodium tristichyum	C	A	C	G	G	G	G	A	A	A	C	T	T	A	C	C	A	G	G	T	C	C	A	G	A	C	A	T	A	G
34	Salvinella umbrosa	C	A	C	G	G	G	G	A	A	A	C	T	T	A	C	C	A	G	G	T	C	C	A	G	A	C	A	T	A	G
35	Salvinella vogalii	C	A	C	G	G	G	G	A	A	A	C	T	T	A	C	C	A	G	G	T	C	C	A	G	A	C	A	T	A	G
36	Oryza sativa	C	A	C	G	G	G	G	A	A	A	C	T	T	A	C	C	A	G	G	T	C	C	A	G	A	C	A	T	A	G
37	Zea mays	C	A	C	G	G	G	G	A	A	A	C	T	T	A	C	C	A	G	G	T	C	C	A	G	A	C	A	T	A	G
38	Arabidopsis thaliana	C	A	C	G	G	G	G	A	A	A	C	T	T	A	C	C	A	G	G	T	C	C	A	G	A	C	A	T	A	G
39	Oryza rufipolis	C	A	C	G	G	G	G	A	A	A	C	T	T	A	C	C	A	G	G	T	C	C	A	G	A	C	A	T	A	G
40	Striga alba	C	A	C	G	G	G	G	A	A	A	C	T	T	A	C	C	A	G	G	T	C	C	A	G	A	C	A	T	A	G
41	Zinnia purpurea	C	A	C	G	G	G	G	A	A	A	C	T	T	A	C	C	A	G	G	T	C	C	A	G	A	C	A	T	A	G
42	Onotum lyboides	C	A	C	G	G	G	G	A	A	A	C	T	T	A	C	C	A	G	G	T	C	C	A	G	A	C	A	T	A	G
43	Ginkgo biloba	C	A	C	G	G	G	G	A	A	A	C	T	T	A	C	C	A	G	G	T	C	C	A	G	A	C	A	T	A	G
44	Pinus luchuanensis	C	A	C	G	G	G	G	A	A	A	C	T	T	A	C	C	A	G	G	T	C	C	A	G	A	C	A	T	A	G
45	Pinus wallichiana	C	A	C	G	G	G	G	A	A	A	C	T	T	A	C	C	A	G	G	T	C	C	A	G	A	C	A	T	A	G
46	Megala rugifolia	C	A	C	G	G	G	G	A	A	A	C	T	T	A	C	C	A	G	G	T	C	C	A	G	A	C	A	T	A	G
47	Taxus malayana	C	A	C	G	G	G	G	A	A	A	C	T	T	A	C	C	A	G	G	T	C	C	A	G	A	C	A	T	A	G
48	Adiantum redonianum	C	A	C	G	G	G	G	A	A	A	C	T	T	A	C	C	A	G	G	T	C	C	A	G	A	C	A	T	A	G
49	Adiantum redonianum 2	C	A	C	G	G	G	G	A	A	A	C	T	T	A	C	C	A	G	G	T	C	C	A	G	A	C	A	T	A	G
50	Ophiodon elongatum	C	A	C	G	G	G	G	A	A	A	C	T	T	A	C	C	A	G	G	T	C	C	A	G	A	C	A	T	A	G
51	Osmunda cinnamomea	C	A	C	G	G	G	G	A	A	A	C	T	T	A	C	C	A	G	G	T	C	C	A	G	A	C	A	T	A	G
52	Salvinia natans	C	A	C	G	G	G	G	A	A	A	C	T	T	A	C	C	A	G	G	T	C	C	A	G	A	C	A	T	A	G
53	Polypodium nudum	C	A	C	G	G	G	G	A	A	A	C	T	T	A	C	C	A	G	G	T	C	C	A	G	A	C	A	T	A	G
54	Chamaecytisus	C	A	C	G	G	G	G	A	A	A	C	T	T	A	C	C	A	G	G	T	C	C	A	G	A	C	A	T	A	G
55	Coleochaete scutata	C	A	C	G	G	G	G	A	A	A	C	T	T	A	C	C	A	G	G	T	C	C	A	G	A	C	A	T	A	G
56	Hebeomidium floccidum	C	A	C	G	G	G	G	A	A	A	C	T	T	A	C	C	A	G	G	T	C	C	A	G	A	C	A	T	A	G
57	Nitzschia baileyi	C	A	C	G	G	G	G	A	A	A	C	T	T	A	C	C	A	G	G	T	C	C	A	G	A	C	A	T	A	G
58	Nitzschia sp.	C	A	C	G	G	G	G	A	A	A	C	T	T	A	C	C	A	G	G	T	C	C	A	G	A	C	A	T	A	G
59	Chlamydomonas reinhardtii	C	A	C	G	G	G	G	A	A	A	C	T	T	A	C	C	A	G	G	T	C	C	A	G	A	C	A	T	A	G
60	Hydrodictyon reticulatum	C	A	C	G	G	G	G	A	A	A	C	T	T	A	C	C	A	G	G	T	C	C	A	G	A	C	A	T	A	G
61	Paratetodon pseudohyalinae	C	A	C	G	G	G	G	A	A	A	C	T	T	A	C	C	A	G	G	T	C	C	A	G	A	C	A	T	A	G
62	Dicystotellum discoideum	C	T	C	G	G	G	G	A	A	A	C	T	T	A	C	C	A	G	G	T	C	C	A	G	A	C	A	T	A	G
63	Pteris vittata	C	A	C	G	G	G	G	A	A	A	C	T	T	A	C	C	A	G	G	T	C	C	A	G	A	C	A	T	A	G
64	Megaceros eornigatus	C	A	C	G	G	G	G	A	A	A	C	T	T	A	C	C	A	G	G	T	C	C	A	G	A	C	A	T	A	G
65	Metaphysis brucei	C	A	C	G	G	G	G	A	A	A	C	T	T	A	C	C	A	G	G	T	C	C	A	G	A	C	A	T	A	G
66	Physcomitrella patens	C	A	C	G	G	G	G	A	A	A	C	T	T	A	C	C	A	G	G	T	C	C	A	G	A	C	A	T	A	G
67	Sphagnum palustre	C	A	C	G	G	G	G	A	A	A	C	T	T	A	C	C	A	G	G	T	C	C	A	G	A	C	A	T	A	G

	1881	1882	1883	1884	1885	1886	1887	1888	1889	1890	1891	1892	1893	1894	1895	1896	1897	1898	1899	1900	1901	1902	1903	1904	1905	1906	1907	1908	1909	1910
	0	4	4	0	0	0	3	0	0	2	0	5	0	0	0	0	0	5	0	0	5	0	5	0	0	0	3	4	2	0
1 Saccharomyces cerevisiae	T	T	A	A	T	T	Q	C	Q	A	T	A	A	C	Q	A	A	-	C	Q	A	Q	A	C	C	T	T	A	A	C
2 Schizosaccharomyces pombe	T	T	A	A	T	T	Q	C	Q	A	T	A	A	C	Q	A	A	-	C	Q	A	Q	A	C	C	T	T	A	A	C
3 Physcomitrella patens 2	T	T	A	A	T	T	C	C	Q	T	T	A	A	C	Q	A	A	-	C	Q	A	Q	A	C	C	T	C	A	Q	C
4 Anthoceros agrestis	T	T	A	A	T	T	C	C	Q	T	T	A	A	C	Q	A	A	-	C	Q	A	Q	A	C	C	T	C	A	Q	C
5 Anthoceros laevis	T	T	A	A	T	T	C	C	Q	T	T	A	A	C	Q	A	A	-	C	Q	A	Q	A	C	C	T	C	A	Q	C
6 Pellia epiphylla	T	T	A	A	T	T	C	C	Q	T	T	A	A	C	Q	A	A	-	C	Q	A	Q	A	C	C	T	C	A	Q	C
7 Reboulia hemisphaerica	T	T	A	A	T	T	C	C	Q	T	T	A	A	C	Q	A	A	-	C	Q	A	Q	A	C	C	T	C	A	Q	C
8 Riccardia pinguis	T	T	A	A	T	T	C	C	Q	T	T	A	A	C	Q	A	A	-	C	Q	A	Q	A	C	C	T	C	A	Q	C
9 Sphaerocarpos donnellii	T	T	A	A	T	T	C	C	Q	T	T	A	A	C	Q	A	A	-	C	Q	A	Q	A	C	C	T	C	A	Q	C
10 Sphaerocarpos lasianus	T	T	A	A	T	T	C	C	Q	T	T	A	A	C	Q	A	A	-	C	Q	A	Q	A	C	C	T	C	A	Q	C
11 Atrichum undulatum	T	T	A	A	T	T	C	C	Q	T	T	A	A	C	Q	A	A	-	C	Q	A	Q	A	C	C	T	C	A	Q	C
12 Bryum argenteum	T	T	A	A	T	T	C	C	Q	T	T	A	A	C	Q	A	A	-	C	Q	A	Q	A	C	C	T	C	A	Q	C
13 Eurymochium hians	T	T	A	A	T	T	C	C	Q	T	T	A	A	C	Q	A	A	-	C	Q	A	Q	A	C	C	T	C	A	Q	C
14 Funaria hygrometrica	T	T	A	A	T	T	C	C	Q	T	T	A	A	C	Q	A	A	-	C	Q	A	Q	A	C	C	T	C	A	Q	C
15 Leptobryum pyriforme	T	T	A	A	T	T	C	C	Q	T	T	A	A	C	Q	A	A	-	C	Q	A	Q	A	C	C	T	C	A	Q	C
16 Atrichum horum	T	T	A	A	T	T	C	C	Q	T	T	A	A	C	Q	A	A	-	C	Q	A	Q	A	C	C	T	C	A	Q	C
17 Physcomitrium pyriforme	T	T	A	A	T	T	C	C	Q	T	T	A	A	C	Q	A	A	-	C	Q	A	Q	A	C	C	T	C	A	Q	C
18 Polytichum formosum	T	T	A	A	T	T	C	C	Q	T	T	A	A	C	Q	A	A	-	C	Q	A	Q	A	C	C	T	C	A	Q	C
19 Sphagnum cuspidatum	T	T	A	A	T	T	C	C	Q	T	T	A	A	C	Q	A	A	-	C	Q	A	Q	A	C	C	T	C	A	Q	C
20 Calypogeia arguta	T	T	A	A	T	T	C	C	Q	T	T	A	A	C	Q	A	A	-	C	Q	A	Q	A	C	C	T	C	A	Q	C
21 Conocephalum conicum	T	T	A	A	T	T	C	C	Q	T	T	A	A	C	Q	A	A	-	C	Q	A	Q	A	C	C	T	C	A	Q	C
22 Fossombronia pusilla	T	T	A	A	T	T	C	C	Q	T	T	A	A	C	Q	A	A	-	C	Q	A	Q	A	C	C	T	C	A	Q	C
23 Marchantia polymorpha	T	T	A	A	T	T	C	C	Q	T	T	A	A	C	Q	A	A	-	C	Q	A	Q	A	C	C	T	C	A	Q	C
24 Riccia fluitans	T	T	A	A	T	T	C	C	Q	T	T	A	A	C	Q	A	A	-	C	Q	A	Q	A	C	C	T	C	A	Q	C
25 Scoparia nemorea	T	T	A	A	T	T	C	C	Q	T	T	A	A	C	Q	A	A	-	C	Q	A	Q	A	C	C	T	C	A	Q	C
26 Equisetum hyemale	T	T	A	A	T	T	C	C	Q	T	T	A	A	C	Q	A	A	-	C	Q	A	Q	A	C	C	T	C	A	Q	C
27 Equisetum robustum	T	T	A	A	T	T	C	C	Q	T	T	A	A	C	Q	A	A	-	C	Q	A	Q	A	C	C	T	C	A	Q	C
28 Isoetes engelmannii	T	T	A	A	T	T	C	C	Q	T	T	A	A	C	Q	A	A	-	C	Q	A	Q	A	C	C	T	C	A	Q	C
29 Huperzia lucidulum	T	T	A	A	T	T	C	C	Q	T	T	A	A	C	Q	A	A	-	C	Q	A	Q	A	C	C	T	C	A	Q	C
30 Lycopodium trundatum	T	T	A	A	T	T	C	C	Q	T	T	A	A	C	Q	A	A	-	C	Q	A	Q	A	C	C	T	C	A	Q	C
31 Lycopodium platyneurum	T	T	A	A	T	T	C	C	Q	T	T	A	A	C	Q	A	A	-	C	Q	A	Q	A	C	C	T	C	A	Q	C
32 Lycopodium laetivolum	T	T	A	A	T	T	C	C	Q	T	T	A	A	C	Q	A	A	-	C	Q	A	Q	A	C	C	T	C	A	Q	C
33 Lycopodium tristichyum	T	T	A	A	T	T	C	C	Q	T	T	A	A	C	Q	A	A	-	C	Q	A	Q	A	C	C	T	C	A	Q	C
34 Selaginella umbrosa	T	T	A	A	T	T	C	C	Q	T	T	A	A	C	Q	A	A	-	C	Q	A	Q	A	C	C	T	C	A	Q	C
35 Selaginella vogelii	T	T	A	A	T	T	C	C	Q	T	T	A	A	C	Q	A	A	-	C	Q	A	Q	A	C	C	T	C	A	Q	C
36 Oryza sativa	T	T	A	A	T	T	C	C	Q	T	T	A	A	C	Q	A	A	-	C	Q	A	Q	A	C	C	T	C	A	Q	C
37 Zea mays	T	T	A	A	T	T	C	C	Q	T	T	A	A	C	Q	A	A	-	C	Q	A	Q	A	C	C	T	C	A	Q	C
38 Arabidopsis thaliana	T	T	A	A	T	T	C	C	Q	T	T	A	A	C	Q	A	A	-	C	Q	A	Q	A	C	C	T	C	A	Q	C
39 Glycine max	T	T	A	A	T	T	C	C	Q	T	T	A	A	C	Q	A	A	-	C	Q	A	Q	A	C	C	T	C	A	Q	C
40 Brassica oleracea	T	T	A	A	T	T	C	C	Q	T	T	A	A	C	Q	A	A	-	C	Q	A	Q	A	C	C	T	C	A	Q	C
41 Zizia aurea	T	T	A	A	T	T	C	C	Q	T	T	A	A	C	Q	A	A	-	C	Q	A	Q	A	C	C	T	C	A	Q	C
42 Quercus laevis	T	T	A	A	T	T	C	C	Q	T	T	A	A	C	Q	A	A	-	C	Q	A	Q	A	C	C	T	C	A	Q	C
43 Girardinia bicolor	T	T	A	A	T	T	C	C	Q	T	T	A	A	C	Q	A	A	-	C	Q	A	Q	A	C	C	T	C	A	Q	C
44 Pinus taeda	T	T	A	A	T	T	C	C	Q	T	T	A	A	C	Q	A	A	-	C	Q	A	Q	A	C	C	T	C	A	Q	C
45 Pinus strobus	T	T	A	A	T	T	C	C	Q	T	T	A	A	C	Q	A	A	-	C	Q	A	Q	A	C	C	T	C	A	Q	C
46 Populus nigra	T	T	A	A	T	T	C	C	Q	T	T	A	A	C	Q	A	A	-	C	Q	A	Q	A	C	C	T	C	A	Q	C
47 Taxus canadensis	T	T	A	A	T	T	C	C	Q	T	T	A	A	C	Q	A	A	-	C	Q	A	Q	A	C	C	T	C	A	Q	C
48 Adiantum raddianum	T	T	A	A	T	T	C	C	Q	T	T	A	A	C	Q	A	A	-	C	Q	A	Q	A	C	C	T	C	A	Q	C
49 Adiantum raddianum 2	T	T	A	A	T	T	C	C	Q	T	T	A	A	C	Q	A	A	-	C	Q	A	Q	A	C	C	T	C	A	Q	C
50 Ophioglossum petiolatum	T	T	A	A	T	T	C	C	Q	T	T	A	A	C	Q	A	A	-	C	Q	A	Q	A	C	C	T	C	A	Q	C
51 Camarisia dimorpha	T	T	A	A	T	T	C	C	Q	T	T	A	A	C	Q	A	A	-	C	Q	A	Q	A	C	C	T	C	A	Q	C
52 Selaginella selaginoides	T	T	A	A	T	T	C	C	Q	T	T	A	A	C	Q	A	A	-	C	Q	A	Q	A	C	C	T	C	A	Q	C
53 Pteridium aquilinum	T	T	A	A	T	T	C	C	Q	T	T	A	A	C	Q	A	A	-	C	Q	A	Q	A	C	C	T	C	A	Q	C
54 Chara foeniculacea	T	T	A	A	T	T	C	C	Q	T	T	A	A	C	Q	A	A	-	C	Q	A	Q	A	C	C	T	C	A	Q	C
55 Coleochaete scutellata	T	T	A	A	T	T	C	C	Q	T	T	A	A	C	Q	A	A	-	C	Q	A	Q	A	C	C	T	C	A	Q	C
56 Klebsormidium flaccidum	T	T	A	A	T	T	C	C	Q	T	T	A	A	C	Q	A	A	-	C	Q	A	Q	A	C	C	T	C	A	Q	C
57 Nitella flexilis	T	T	A	A	T	T	C	C	Q	T	T	A	A	C	Q	A	A	-	C	Q	A	Q	A	C	C	T	C	A	Q	C
58 Nitella sp.	T	T	A	A	T	T	C	C	Q	T	T	A	A	C	Q	A	A	-	C	Q	A	Q	A	C	C	T	C	A	Q	C
59 Chlamydomonas reinhardtii	T	T	Q	A	T	T	C	C	Q	Q	T	A	A	C	Q	A	A	-	C	Q	A	Q	A	C	C	T	C	A	Q	C
60 Hydrodictyon reticulatum	T	T	Q	A	T	T	C	C	Q	Q	T	A	A	C	Q	A	A	-	C	Q	A	Q	A	C	C	T	C	A	Q	C
61 Paratetodon pseudobrevicollis	T	C	G	A	T	T	C	C	Q	Q	T	A	A	C	Q	A	A	-	C	Q	A	Q	A	C	C	T	C	A	Q	C
62 Dictyostelium discoideum	T	C	A	A	T	T	C	C	Q	Q	T	A	A	C	Q	A	A	-	C	Q	A	Q	A	C	C	T	C	A	Q	C
63 Pteris vittata	T	T	A	A	T	T	C	C	Q	T	T	A	A	C	Q	A	A	-	C	Q	A	Q	A	C	C	T	C	A	Q	C
64 Megaceros acuminatus	T	T	A	A	T	T	C	C	Q	T	T	A	A	C	Q	A	A	-	C	Q	A	Q	A	C	C	T	C	A	Q	C
65 Notophytis bracteata	T	T	A	A	T	T	C	C	Q	T	T	A	A	C	Q	A	A	-	C	Q	A	Q	A	C	C	T	C	A	Q	C
66 Physcomitrella patens	T	T	A	A	T	T	C	C	Q	T	T	A	A	C	Q	A	A	-	C	Q	A	Q	A	C	C	T	C	A	Q	C
67 Sphagnum palustre	T	T	A	A	T	T	C	C	Q	T	T	A	A	C	Q	A	A	-	C	Q	A	Q	A	C	C	T	C	A	Q	C

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	1771	1772	1773	1774	1775	1776	1777	1778	1779	1780	1781	1782	1783	1784	1785	1786	1787	1788	1789	1790	1791	1792	1793	1794	1795	1796	1797	1798	1799	1800
1 Saccharomyces cerevisiae	0	C	C	G	.	A	T	G	G	A	A	G	T	T	T	G	A	G	G	C	A	A	T	A	A	C	A	G	G	T
2 Schizosaccharomyces pombe	0	C	C	A	.	A	T	G	G	A	A	G	T	T	T	G	A	G	G	C	A	A	T	A	A	C	A	G	G	T
3 Physcomitrella patens 2	0	C	C	G	.	A	T	G	G	A	A	G	T	T	T	G	A	G	G	C	A	A	T	A	A	C	A	G	G	T
4 Arabidopsis thaliana	0	C	C	A	.	A	T	G	G	A	A	G	T	T	T	G	A	G	G	C	A	A	T	A	A	C	A	G	G	T
5 Arabidopsis lyrata	0	C	G	A	.	A	T	G	G	A	A	G	T	T	T	G	A	G	G	C	A	A	T	A	A	C	A	G	G	T
6 Petta epiphyta	0	C	C	A	.	A	T	G	G	A	A	G	T	T	T	G	A	G	G	C	A	A	T	A	A	C	A	G	G	T
7 Rhabdula hemisphaerica	0	C	C	G	.	A	C	G	G	A	A	G	T	T	T	G	A	G	G	C	A	A	T	A	A	C	A	G	G	T
8 Riccia pinguis	0	C	C	A	.	A	T	G	G	A	A	G	T	T	T	G	A	G	G	C	A	A	T	A	A	C	A	G	G	T
9 Sphaerocarpos donnellii	0	C	C	G	.	A	T	G	G	A	A	G	T	T	T	G	A	G	G	C	A	A	T	A	A	C	A	G	G	T
10 Sphaerocarpos lasarus	0	C	C	G	.	A	T	G	G	A	A	G	T	T	T	G	A	G	G	C	A	A	T	A	A	C	A	G	G	T
11 Atrichum undulatum	0	C	C	A	.	A	T	G	G	A	A	G	T	T	T	G	A	G	G	C	A	A	T	A	A	C	A	G	G	T
12 Bryum argenteum	0	C	C	G	.	A	T	G	G	A	A	G	T	T	T	G	A	G	G	C	A	A	T	A	A	C	A	G	G	T
13 Eurythochium hians	0	C	C	G	.	A	T	G	G	A	A	G	T	A	T	G	A	G	G	C	A	A	T	A	A	C	A	G	G	T
14 Funaria hygrometrica	0	C	C	G	.	A	T	G	G	A	A	G	T	T	T	G	A	G	G	C	A	A	T	A	A	C	A	G	G	T
15 Lophocolea boryana	0	C	C	G	.	A	T	G	G	A	A	G	T	T	T	G	A	G	G	C	A	A	T	A	A	C	A	G	G	T
16 Mnium hornum	0	C	C	G	.	A	T	G	G	A	A	G	T	T	T	G	A	G	G	C	A	A	T	A	A	C	A	G	G	T
17 Physcomitrium pyriforme	0	C	C	A	.	A	T	G	G	A	A	G	T	T	T	G	A	G	G	C	A	A	T	A	A	C	A	G	G	T
18 Polytrichum formosum	0	C	C	G	.	A	T	G	G	A	A	G	T	T	T	G	A	G	G	C	A	A	T	A	A	C	A	G	G	T
19 Sphagnum cuspidatum	0	C	C	A	.	A	T	G	G	A	A	G	T	T	T	G	A	G	G	C	A	A	T	A	A	C	A	G	G	T
20 Calypogeia arguta	0	C	C	A	.	T	T	G	G	A	A	G	T	T	T	G	A	G	G	C	A	A	T	A	A	C	A	G	G	T
21 Conocephalum conicum	0	C	C	G	.	A	C	G	G	A	A	G	T	T	T	G	A	G	G	C	A	A	T	A	A	C	A	G	G	T
22 Fossombronia pusilla	0	C	C	A	.	A	T	G	G	A	A	G	T	T	T	G	A	G	G	C	A	A	T	A	A	C	A	G	G	T
23 Marchantia polymorpha	0	C	C	G	.	A	T	G	G	A	A	G	T	T	T	G	A	G	G	C	A	A	T	A	A	C	A	G	G	T
24 Riccia fluitans	0	C	C	G	.	A	C	G	G	A	A	G	T	T	T	G	A	G	G	C	A	A	T	A	A	C	A	G	G	T
25 Soperia nemorea	0	C	C	A	.	A	T	G	G	A	A	G	T	T	T	G	A	G	G	C	A	A	T	A	A	C	A	G	G	T
26 Equisetum hyemale	0	C	C	C	.	A	T	G	G	A	A	G	T	T	T	G	A	G	G	C	A	A	T	A	A	C	A	G	G	T
27 Equisetum robustum	0	C	C	C	.	A	T	G	G	A	A	G	T	T	T	G	A	G	G	C	A	A	T	A	A	C	A	G	G	T
28 Isoetes engelmannii	0	C	G	G	.	A	T	G	G	A	A	G	T	T	T	G	A	G	G	C	A	A	T	A	A	C	A	G	G	T
29 Huperzia lucidulum	0	C	G	G	.	A	T	G	G	A	A	G	T	T	T	G	A	G	G	C	A	A	T	A	A	C	A	G	G	T
30 Lycopodium truncatum	0	C	G	G	.	A	T	G	G	A	A	G	T	T	T	G	A	G	G	C	A	A	T	A	A	C	A	G	G	T
31 Lycopodium phlegmaria	0	C	G	G	.	A	T	G	G	A	A	G	T	T	T	G	A	G	G	C	A	A	T	A	A	C	A	G	G	T
32 Lycopodium laxifolium	0	C	G	G	.	A	T	G	G	A	A	G	T	T	T	G	A	G	G	C	A	A	T	A	A	C	A	G	G	T
33 Lycopodium tristichyum	0	C	G	G	.	A	T	G	G	A	A	G	T	T	T	G	A	G	G	C	A	A	T	A	A	C	A	G	G	T
34 Selaginella selaginoides	A	G	C	G	.	A	T	G	G	A	A	G	T	T	T	G	A	G	G	C	A	A	T	A	A	C	A	G	G	T
35 Selaginella selaginoides	0	C	G	G	.	A	T	G	G	A	A	G	T	T	T	G	A	G	G	C	A	A	T	A	A	C	A	G	G	T
36 Oryza sativa	0	C	C	C	.	A	C	G	G	A	A	G	T	T	T	G	A	G	G	C	A	A	T	A	A	C	A	G	G	T
37 Zea mays	0	C	C	C	.	G	C	G	.	A	A	G	T	T	T	G	A	G	G	C	A	A	T	A	A	C	A	G	G	T
38 Arabidopsis thaliana	0	C	C	C	.	A	A	G	G	A	A	G	T	T	T	G	A	G	G	C	A	A	T	A	A	C	A	G	G	T
39 Glycine max	0	C	C	C	.	A	C	G	G	A	A	G	T	T	T	G	A	G	G	C	A	A	T	A	A	C	A	G	G	T
40 Brinjal	0	C	C	C	.	A	A	G	G	A	A	G	T	T	T	G	A	G	G	C	A	A	T	A	A	C	A	G	G	T
41 Zinnia	0	C	C	C	.	A	T	G	G	A	A	G	T	T	T	G	A	G	G	C	A	A	T	A	A	C	A	G	G	T
42 Arabidopsis thaliana	0	C	C	C	.	A	T	G	G	A	A	G	T	T	T	G	A	G	G	C	A	A	T	A	A	C	A	G	G	T
43 Arabidopsis thaliana	0	C	C	C	.	A	T	G	G	A	A	G	T	T	T	G	A	G	G	C	A	A	T	A	A	C	A	G	G	T
44 Pinus taeda	0	C	C	C	.	A	T	G	G	A	A	G	T	T	T	G	A	G	G	C	A	A	T	A	A	C	A	G	G	T
45 Pinus taeda	0	C	C	C	.	A	T	G	G	A	A	G	T	T	T	G	A	G	G	C	A	A	T	A	A	C	A	G	G	T
46 Pinus taeda	0	C	C	C	.	A	T	G	G	A	A	G	T	T	T	G	A	G	G	C	A	A	T	A	A	C	A	G	G	T
47 Taxus canadensis	0	C	C	C	.	A	T	G	G	A	A	G	T	T	T	G	A	G	G	C	A	A	T	A	A	C	A	G	G	T
48 Adiantum species	0	C	C	C	.	A	T	G	G	A	A	G	T	T	T	G	A	G	G	C	A	A	T	A	A	C	A	G	G	T
49 Adiantum species	0	C	C	C	.	A	T	G	G	A	A	G	T	T	T	G	A	G	G	C	A	A	T	A	A	C	A	G	G	T
50 Ophioglossum species	0	C	C	C	.	A	T	G	G	A	A	G	T	T	T	G	A	G	G	C	A	A	T	A	A	C	A	G	G	T
51 Osmunda species	0	C	C	C	.	A	T	G	G	A	A	G	T	T	T	G	A	G	G	C	A	A	T	A	A	C	A	G	G	T
52 Salvinia species	0	C	C	C	.	A	T	G	G	A	A	G	T	T	T	G	A	G	G	C	A	A	T	A	A	C	A	G	G	T
53 Peltandra species	0	C	C	C	.	A	T	G	G	A	A	G	T	T	T	G	A	G	G	C	A	A	T	A	A	C	A	G	G	T
54 Chara species	0	C	C	A	.	A	C	G	G	A	A	G	T	T	T	G	A	G	G	C	A	A	T	A	A	C	A	G	G	T
55 Coleochaete species	0	C	C	G	.	A	T	G	G	A	A	G	T	T	T	G	A	G	G	C	A	A	T	A	A	C	A	G	G	T
56 Klebsormidium species	0	C	C	A	.	A	T	G	G	A	A	G	T	T	T	G	A	G	G	C	A	A	T	A	A	C	A	G	G	T
57 Nitella species	0	C	C	A	.	A	T	G	G	A	A	G	T	T	T	G	A	G	G	C	A	A	T	A	A	C	A	G	G	T
58 Nitella sp.	0	C	C	A	.	A	T	G	G	A	A	G	T	T	T	G	A	G	G	C	A	A	T	A	A	C	A	G	G	T
59 Chara species	0	C	C	A	.	A	T	G	G	A	A	G	T	T	T	G	A	G	G	C	A	A	T	A	A	C	A	G	G	T
60 Hydrodictyon species	0	T	C	A	.	A	T	G	G	A	A	G	T	A	T	G	A	G	G	C	A	A	T	A	A	C	A	G	G	T
61 Parietaria species	0	C	C	G	.	A	T	G	G	A	A	G	T	G	T	G	A	G	G	C	A	A	T	A	A	C	A	G	G	T
62 Dictyosphaerium species	0	C	A	G	.	G	C	G	G	A	A	G	T	C	C	G	A	G	G	C	A	A	T	A	A	C	A	G	G	T
63 Plectamblypus species	0	C	C	C	.	A	T	G	G	A	A	G	T	T	T	G	A	G	G	C	A	A	T	A	A	C	A	G	G	T
64 Megaceros species	0	C	G	A	.	A	T	G	G	A	A	G	T	T	T	G	A	G	G	C	A	A	T	A	A	C	A	G	G	T
65 Notiphya species	0	T	G	A	.	A	T	A	G	A	A	G	T	T	T	G	A	G	G	A	A	A	T	A	A	C	A	G	G	T
66 Physcomitrella patens	0	C	C	G	.	A	T	G	G	A	A	G	T	T	T	G	A	G	G	C	A	A	T	A	A	C	A	G	G	T
67 Sphagnum species	0	C	C	A	.	A	T	G	G	A	A	G	T	T	T	G	A	G	G	C	A	A	T	A	A	C	A	G	G	T

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		1601	1602	1603	1604	1605	1606	1607	1608	1609	1610	1611	1612	1613	1614	1615	1616	1617	1618	1619	1620	1621	1622	1623	1624	1625	1626	1627	1628	1629	1630	
		5	6	7	8	9	0	1	2	3	4	5	6	7	8	9	0	1	2	3	4	5	6	7	8	9	0	1	2	3	4	
1	<i>Saccharomyces cerevisiae</i>	C	.	T	G	T	G	A	T	G	C	C	C	T	T	A	G	A	A	C	G	T	T	C	T	G	Q	Q	Q	C	C	Q
2	<i>Schizosaccharomyces pombe</i>	C	.	T	G	T	G	A	T	G	C	C	C	T	T	A	G	A	A	T	Q	T	T	C	T	G	Q	Q	Q	C	C	Q
3	<i>Physcomitrella patens 2</i>	C	.	T	G	T	G	A	T	G	C	C	C	T	T	A	G	A	A	T	Q	T	T	C	T	G	Q	Q	Q	C	C	Q
4	<i>Anthoceros agrestis</i>	C	.	T	G	T	G	A	T	G	C	C	C	T	T	A	G	A	A	T	Q	T	T	C	T	G	Q	Q	Q	C	C	Q
5	<i>Anthoceros laevis</i>	C	.	T	G	T	G	A	T	G	C	C	C	T	T	A	G	A	A	T	Q	T	T	C	T	G	Q	Q	Q	C	C	Q
6	<i>Pellia epiphylla</i>	C	.	T	G	T	G	A	T	G	C	C	C	T	T	A	G	A	A	T	Q	T	T	C	T	G	Q	Q	Q	C	C	Q
7	<i>Reboulia hemisphaerica</i>	C	.	T	G	T	G	A	T	G	C	C	C	T	T	A	G	A	A	T	Q	T	T	C	T	G	Q	Q	Q	C	C	Q
8	<i>Riccardia pinguis</i>	C	.	T	G	T	G	A	T	G	C	C	C	T	T	A	G	A	A	T	Q	T	T	C	T	G	Q	Q	Q	C	C	Q
9	<i>Sphaerocarpus donnellii</i>	C	.	T	G	T	G	A	T	G	C	C	C	T	T	A	G	A	A	T	Q	T	T	C	T	G	Q	Q	Q	C	C	Q
10	<i>Sphaerocarpus lazarus</i>	C	.	T	G	T	G	A	T	G	C	C	C	T	T	A	G	A	A	T	Q	T	T	C	T	G	Q	Q	Q	C	C	Q
11	<i>Asterum unidentatum</i>	C	.	T	G	T	G	A	T	G	C	C	C	T	T	A	G	A	A	T	Q	T	T	C	T	G	Q	Q	Q	C	C	Q
12	<i>Bryum argenteum</i>	C	.	T	G	T	G	A	T	G	C	C	C	T	T	A	G	A	A	T	Q	T	T	C	T	G	Q	Q	Q	C	C	Q
13	<i>Butyrium hians</i>	C	.	T	G	T	G	A	T	G	C	C	C	T	T	A	G	A	A	T	Q	T	T	C	T	G	Q	Q	Q	C	C	Q
14	<i>Purpura hyemalis</i>	C	.	T	G	T	G	A	T	G	C	C	C	T	T	A	G	A	A	T	Q	T	T	C	T	G	Q	Q	Q	C	C	Q
15	<i>Lepidobryum pyriforme</i>	C	.	T	G	T	G	A	T	G	C	C	C	T	T	A	G	A	A	T	Q	T	T	C	T	G	Q	Q	Q	C	C	Q
16	<i>Mitrium horvathi</i>	C	.	T	G	T	G	A	T	G	C	C	C	T	T	A	G	A	A	T	Q	T	T	C	T	G	Q	Q	Q	C	C	Q
17	<i>Physcomitrium pyriforme</i>	C	.	T	G	T	G	A	T	G	C	C	C	T	T	A	G	A	A	T	Q	T	T	C	T	G	Q	Q	Q	C	C	Q
18	<i>Polypodium formosum</i>	C	.	T	G	T	G	A	T	G	C	C	C	T	T	A	G	A	A	T	Q	T	T	C	T	G	Q	Q	Q	C	C	Q
19	<i>Sphagnum cuspidatum</i>	C	.	T	G	T	G	A	T	G	C	C	C	T	T	A	G	A	A	T	Q	T	T	C	T	G	Q	Q	Q	C	C	Q
20	<i>Calyptopogon argutus</i>	C	.	T	G	T	G	A	T	G	C	C	C	T	T	A	G	A	A	T	Q	T	T	C	T	G	Q	Q	Q	C	C	Q
21	<i>Cerocephalum conicum</i>	C	.	T	G	T	G	A	T	G	C	C	C	T	T	A	G	A	A	T	Q	T	T	C	T	G	Q	Q	Q	C	C	Q
22	<i>Fissidens punctatus</i>	C	.	T	G	T	G	A	T	G	C	C	C	T	T	A	G	A	A	T	Q	T	T	C	T	G	Q	Q	Q	C	C	Q
23	<i>Marchantia polymorpha</i>	C	.	T	G	T	G	A	T	G	C	C	C	T	T	A	G	A	A	T	Q	T	T	C	T	G	Q	Q	Q	C	C	Q
24	<i>Riccia fluitans</i>	C	.	T	G	T	G	A	T	G	C	C	C	T	T	A	G	A	A	T	Q	T	T	C	T	G	Q	Q	Q	C	C	Q
25	<i>Scoparia nemorea</i>	C	.	T	G	T	G	A	T	G	C	C	C	T	T	A	G	A	A	T	Q	T	T	C	T	G	Q	Q	Q	C	C	Q
26	<i>Equisetum hyemale</i>	C	.	T	G	T	G	A	T	G	C	C	C	T	T	A	G	A	A	T	Q	T	T	C	T	G	Q	Q	Q	C	C	Q
27	<i>Equisetum robustum</i>	C	.	T	G	T	G	A	T	G	C	C	C	T	T	A	G	A	A	T	Q	T	T	C	T	G	Q	Q	Q	C	C	Q
28	<i>Isetes engelmannii</i>	C	.	T	G	T	G	A	T	G	C	C	C	T	T	A	G	A	A	T	Q	T	T	C	T	G	Q	Q	Q	C	C	Q
29	<i>Huperzia lucidulum</i>	C	.	T	G	T	G	A	T	G	C	C	C	T	T	A	G	A	A	T	Q	T	T	C	T	G	Q	Q	Q	C	C	Q
30	<i>Lycopecton inundatum</i>	C	.	T	G	T	G	A	T	G	C	C	C	T	T	A	G	A	A	T	Q	T	T	C	T	G	Q	Q	Q	C	C	Q
31	<i>Lycopecton phlegmaria</i>	C	.	T	G	T	G	A	T	G	C	C	C	T	T	A	G	A	A	T	Q	T	T	C	T	G	Q	Q	Q	C	C	Q
32	<i>Lycopecton latifolium</i>	C	.	T	G	T	G	A	T	G	C	C	C	T	T	A	G	A	A	T	Q	T	T	C	T	G	Q	Q	Q	C	C	Q
33	<i>Lycopecton trielidatum</i>	C	.	T	G	T	G	A	T	G	C	C	C	T	T	A	G	A	A	T	Q	T	T	C	T	G	Q	Q	Q	C	C	Q
34	<i>Botrychium umbrosum</i>	C	.	T	G	T	G	A	T	G	C	C	C	T	T	A	G	A	A	T	Q	T	T	C	T	G	Q	Q	Q	C	C	Q
35	<i>Botrychium vogelii</i>	C	.	T	G	T	G	A	T	G	C	C	C	T	T	A	G	A	A	T	Q	T	T	C	T	G	Q	Q	Q	C	C	Q
36	<i>Oryza sativa</i>	C	.	T	G	T	G	A	T	G	C	C	C	T	T	A	G	A	A	T	Q	T	T	C	T	G	Q	Q	Q	C	C	Q
37	<i>Zea mays</i>	C	.	T	G	T	G	A	T	G	C	C	C	T	T	A	G	A	A	T	Q	T	T	C	T	G	Q	Q	Q	C	C	Q
38	<i>Arabis thaliana</i>	C	.	T	G	T	G	A	T	G	C	C	C	T	T	A	G	A	A	T	Q	T	T	C	T	G	Q	Q	Q	C	C	Q
39	<i>Oryza sativa</i>	C	.	T	G	T	G	A	T	G	C	C	C	T	T	A	G	A	A	T	Q	T	T	C	T	G	Q	Q	Q	C	C	Q
40	<i>Strapiz alba</i>	C	.	T	G	T	G	A	T	G	C	C	C	T	T	A	G	A	A	T	Q	T	T	C	T	G	Q	Q	Q	C	C	Q
41	<i>Zinnia purpurea</i>	C	.	T	G	T	G	A	T	G	C	C	C	T	T	A	G	A	A	T	Q	T	T	C	T	G	Q	Q	Q	C	C	Q
42	<i>Oenothera lutea</i>	C	.	T	G	T	G	A	T	G	C	C	C	T	T	A	G	A	A	T	Q	T	T	C	T	G	Q	Q	Q	C	C	Q
43	<i>Oenothera biennis</i>	C	.	T	G	T	G	A	T	G	C	C	C	T	T	A	G	A	A	T	Q	T	T	C	T	G	Q	Q	Q	C	C	Q
44	<i>Prunella vulgaris</i>	C	.	T	G	T	G	A	T	G	C	C	C	T	T	A	G	A	A	T	Q	T	T	C	T	G	Q	Q	Q	C	C	Q
45	<i>Prunella domestica</i>	C	.	T	G	T	G	A	T	G	C	C	C	T	T	A	G	A	A	T	Q	T	T	C	T	G	Q	Q	Q	C	C	Q
46	<i>Hesperis matronalis</i>	C	.	T	G	T	G	A	T	G	C	C	C	T	T	A	G	A	A	T	Q	T	T	C	T	G	Q	Q	Q	C	C	Q
47	<i>Taxus canadensis</i>	C	.	T	G	T	G	A	T	G	C	C	C	T	T	A	G	A	A	T	Q	T	T	C	T	G	Q	Q	Q	C	C	Q
48	<i>Adiantum redolens</i>	C	.	T	G	T	G	A	T	G	C	C	C	T	T	A	G	A	A	T	Q	T	T	C	T	G	Q	Q	Q	C	C	Q
49	<i>Adiantum redolens 2</i>	C	.	T	G	T	G	A	T	G	C	C	C	T	T	A	G	A	A	T	Q	T	T	C	T	G	Q	Q	Q	C	C	Q
50	<i>Ophioglossum pedicellatum</i>	C	.	T	G	T	G	A	T	G	C	C	C	T	T	A	G	A	A	T	Q	T	T	C	T	G	Q	Q	Q	C	C	Q
51	<i>Cornus amomum</i>	C	.	T	G	T	G	A	T	G	C	C	C	T	T	A	G	A	A	T	Q	T	T	C	T	G	Q	Q	Q	C	C	Q
52	<i>Salvia nemorosa</i>	C	.	T	G	T	G	A	T	G	C	C	C	T	T	A	G	A	A	T	Q	T	T	C	T	G	Q	Q	Q	C	C	Q
53	<i>Phlox paniculata</i>	C	.	T	G	T	G	A	T	G	C	C	C	T	T	A	G	A	A	T	Q	T	T	C	T	G	Q	Q	Q	C	C	Q
54	<i>Chrysanthemum leucanthemum</i>	C	.	T	G	T	G	A	T	G	C	C	C	T	T	A	G	A	A	T	Q	T	T	C	T	G	Q	Q	Q	C	C	Q
55	<i>Colobosolenia caudata</i>	C	.	T	G	T	G	A	T	G	C	C	C	T	T	A	G	A	A	T	Q	T	T	C	T	G	Q	Q	Q	C	C	Q
56	<i>Nabeosolenia caudata</i>	C	.	T	G	T	G	A	T	G	C	C	C	T	T	A	G	A	A	T	Q	T	T	C	T	G	Q	Q	Q	C	C	Q
57	<i>Mitella sp.</i>	C	.	T	G	T	G	A	T	G	C	C	C	T	T	A	G	A	A	T	Q	T	T	C	T	G	Q	Q	Q	C	C	Q
58	<i>Chlamydomonas reinhardtii</i>	C	.	T	G	T	G	A	T	G	C	C	C	T	T	A	G	A	A	T	Q	T	T	C	T	G	Q	Q	Q	C	C	Q
59	<i>Hydrodictyon reticulatum</i>	C	.	T	G	T	G	A	T	G	C	C	C	T	T	A	G	A	A	T	Q	T	T	C	T	G	Q	Q	Q	C	C	Q
60	<i>Parthenocarpus pseudocarpus</i>	C	.	T	G	T	G	A	T	G	C	C	C	T	T	A	G	A	A	T	Q	T	T	C	T	G	Q	Q	Q	C	C	Q
61	<i>Dicranella diandra</i>	C	.	T	G	T	G	A	T	G	C	C	C	T	T	A	G	A	A	T	Q	T	T	C	T	G	Q	Q	Q	C	C	Q
62	<i>Perla vittata</i>	C	.	T	G	T	G	A	T	G	C	C	C	T	T	A	G	A	A	T	Q	T	T	C	T	G	Q	Q	Q	C	C	Q
63	<i>Megaceros aeruginosus</i>	C	.	T	G	T	G	A	T	G	C	C	C	T	T	A	G	A	A	T	Q	T	T	C	T	G	Q	Q	Q	C	C	Q
64	<i>Notophylax brucei</i>	C	.	T	G	T	G	A	T	G	C																					

Taxa with Variability Info	09	1921	1922	1923	1924	1925	1926	1927	1928	1929	1930	1931	1932	1933	1934	1935	1936	1937	1938	1939	1940	1941	1942	1943	1944	1945	1946	1947	1948	1949	1950
	0	0	4			1	3	2	0	0	4	1	4	2	0	2	0	0	0	0	0	0	3	3	0	0	0	0	4	0	
1 Saccharomyces cerevisiae	C	T	T	.	.	G	T	Q	A	A	A	C	T	C	C	G	T	.	C	G	T	Q	C	T	Q	Q	Q	Q	A	T	
2 Schizosaccharomyces pombe	C	T	T	.	.	Q	T	T	A	A	A	C	T	C	C	G	T	.	C	G	T	Q	C	T	Q	Q	Q	Q	A	T	
3 Physcomitrella patens 2	C	T	T	.	.	T	T	Q	A	A	A	T	T	T	C	A	T	.	C	Q	T	Q	A	T	Q	Q	Q	Q	A	T	
4 Arabisopsis agrestis	C	T	T	.	.	T	T	Q	A	A	A	T	T	T	C	A	T	.	C	Q	T	Q	A	T	Q	Q	Q	Q	A	T	
5 Arabisopsis laevis	C	T	T	.	.	T	T	Q	A	A	A	T	T	T	C	A	T	.	C	Q	T	Q	A	T	Q	Q	Q	Q	A	T	
6 Pellia epiphylla	C	T	T	.	.	T	T	Q	A	A	A	T	T	T	C	A	T	.	C	Q	T	Q	A	T	Q	Q	Q	Q	A	T	
7 Reboulia hemisphaerica	C	T	T	.	.	T	T	Q	A	A	A	T	T	T	C	A	T	.	C	Q	T	Q	A	T	Q	Q	Q	Q	A	T	
8 Riccardia pinguis	C	T	T	.	.	T	T	Q	A	A	A	T	T	T	C	A	T	.	C	Q	T	Q	A	T	Q	Q	Q	Q	A	T	
9 Sphaerocarpos dorvilli	C	T	T	.	.	T	T	C	A	A	A	C	T	T	C	A	T	.	C	Q	T	Q	A	T	Q	Q	Q	Q	A	T	
10 Sphaerocarpos texanus	C	T	T	.	.	T	T	C	A	A	A	C	T	T	C	A	T	.	C	Q	T	Q	A	T	Q	Q	Q	Q	A	T	
11 Atrichum undulatum	C	T	T	.	.	T	T	Q	A	A	A	T	T	T	C	A	T	.	C	Q	T	Q	A	T	Q	Q	Q	Q	A	T	
12 Bryum argenteum	C	T	T	.	.	T	T	Q	A	A	A	T	T	T	C	A	T	.	C	Q	T	Q	A	T	Q	Q	Q	Q	A	T	
13 Euthymidium hians	C	T	T	.	.	T	T	Q	A	A	A	T	T	T	C	A	T	.	C	Q	T	Q	A	T	Q	Q	Q	Q	A	T	
14 Funaria hygrometrica	C	T	T	.	.	T	T	Q	A	A	A	T	T	T	C	A	T	.	C	Q	T	Q	A	T	Q	Q	Q	Q	A	T	
15 Leptobryum pyriforme	C	T	T	.	.	T	T	Q	A	A	A	T	T	T	C	A	T	.	C	Q	T	Q	A	T	Q	Q	Q	Q	A	T	
16 Mnium hornum	C	T	T	.	.	C	T	Q	A	A	A	T	T	T	C	A	T	.	C	Q	T	Q	A	T	Q	Q	Q	Q	A	T	
17 Physcomitrium pyriforme	C	T	T	.	.	C	T	Q	A	A	A	T	T	T	C	A	T	.	C	Q	T	Q	A	T	Q	Q	Q	Q	A	T	
18 Polytichum lomosum	C	T	T	.	.	T	T	Q	A	A	A	T	T	T	C	A	T	.	C	Q	T	Q	A	T	Q	Q	Q	Q	A	T	
19 Sphagnum cuspidatum	C	T	T	.	.	T	T	Q	A	A	A	T	T	T	C	A	T	.	C	Q	T	Q	A	T	Q	Q	Q	Q	A	T	
20 Calypogeia anguta	C	T	T	.	.	T	T	Q	A	A	A	T	T	T	C	A	T	.	C	Q	T	Q	A	T	Q	Q	Q	Q	A	T	
21 Conocephalum conicum	C	T	T	.	.	T	T	Q	A	A	A	T	T	T	C	A	T	.	C	Q	T	Q	A	T	Q	Q	Q	Q	A	T	
22 Foscomitris pusilla	C	T	T	.	.	T	T	Q	A	A	A	T	T	T	C	A	T	.	C	Q	T	Q	A	T	Q	Q	Q	Q	A	T	
23 Marchantia polymorpha	C	T	T	.	.	T	T	Q	A	A	A	T	T	T	C	A	T	.	C	Q	T	Q	A	T	Q	Q	Q	Q	A	T	
24 Riccia fluitans	C	T	T	.	.	T	T	Q	A	A	A	T	T	T	C	A	T	.	C	Q	T	Q	A	T	Q	Q	Q	Q	A	T	
25 Scapania nemorea	C	T	T	.	.	T	T	Q	A	A	A	T	T	T	C	A	T	.	C	Q	T	Q	A	T	Q	Q	Q	Q	A	T	
26 Equisetum hyemale	C	T	T	.	.	T	T	Q	A	A	A	T	T	T	C	A	T	.	C	Q	T	Q	A	T	Q	Q	Q	Q	A	T	
27 Equisetum robustum	C	T	T	.	.	T	T	Q	A	A	A	T	T	T	C	A	T	.	C	Q	T	Q	A	T	Q	Q	Q	Q	A	T	
28 Isoetes engelmannii	C	T	T	.	.	T	T	Q	A	A	A	T	T	T	C	A	T	.	C	Q	T	Q	A	T	Q	Q	Q	Q	A	T	
29 Huperzia lucidulum	C	T	T	.	.	T	T	Q	A	A	A	T	T	T	C	A	T	.	C	Q	T	Q	A	T	Q	Q	Q	Q	A	T	
30 Lycopodium truncatum	C	T	T	.	.	T	T	Q	A	A	A	T	T	T	C	A	T	.	C	Q	T	Q	A	T	Q	Q	Q	Q	A	T	
31 Lycopodium phlegmaria	C	T	T	.	.	T	T	Q	A	A	A	T	T	T	C	A	T	.	C	Q	T	Q	A	T	Q	Q	Q	Q	A	T	
32 Lycopodium lucidulum	C	T	T	.	.	T	T	Q	A	A	A	T	T	T	C	A	T	.	C	Q	T	Q	A	T	Q	Q	Q	Q	A	T	
33 Lycopodium tristachyum	C	T	T	.	.	T	T	Q	A	A	A	T	T	T	C	A	T	.	C	Q	T	Q	A	T	Q	Q	Q	Q	A	T	
34 Selaginella umbrosa	C	T	T	.	.	T	T	Q	A	A	A	C	T	T	C	A	T	.	C	Q	T	Q	A	T	Q	Q	Q	Q	A	T	
35 Selaginella vogelii	C	T	T	.	.	C	T	Q	A	A	A	T	T	T	C	A	T	.	C	Q	T	Q	A	T	Q	Q	Q	Q	A	T	
36 Oryza sativa	C	T	T	.	.	G	G	Q	A	A	A	T	T	T	C	A	T	.	C	Q	T	Q	A	T	Q	Q	Q	Q	A	T	
37 Zea mays	C	T	T	.	.	Q	Q	Q	A	A	A	T	T	T	C	A	T	.	C	Q	T	Q	A	T	Q	Q	Q	Q	A	T	
38 Arabidopsis thaliana	C	T	T	.	.	T	.	Q	A	A	A	T	T	T	C	A	T	.	C	Q	T	Q	A	T	Q	Q	Q	Q	A	T	
39 Glycine max	C	T	T	.	.	T	.	Q	A	A	A	T	T	T	C	A	T	.	C	Q	T	Q	A	T	Q	Q	Q	Q	A	T	
40 Brassica oleracea	C	T	T	.	.	T	.	Q	A	A	A	T	T	T	C	A	T	.	C	Q	T	Q	A	T	Q	Q	Q	Q	A	T	
41 Zania pusilla	C	T	G	.	.	G	C	Q	A	A	A	T	T	T	C	A	T	.	C	Q	T	Q	A	T	Q	Q	Q	Q	A	T	
42 Onoclea sensibilis	C	T	T	.	.	C	T	Q	A	A	A	T	T	T	C	A	T	.	C	Q	T	Q	A	T	Q	Q	Q	Q	A	T	
43 Oenothera biennis	C	T	G	.	.	C	C	Q	A	A	A	T	T	T	C	A	T	.	C	Q	T	Q	A	T	Q	Q	Q	Q	A	T	
44 Pinus taeda	C	T	T	.	.	C	T	C	A	A	A	T	T	T	C	A	T	.	C	Q	T	Q	A	T	Q	Q	Q	Q	A	T	
45 Pinus strobus	C	T	T	.	.	C	T	C	A	A	A	T	T	T	C	A	T	.	C	Q	T	Q	A	T	Q	Q	Q	Q	A	T	
46 Pinus resinosa	C	T	T	.	.	C	T	C	A	A	A	T	T	T	C	A	T	.	C	Q	T	Q	A	T	Q	Q	Q	Q	A	T	
47 Nagia nag	C	T	T	.	.	G	C	C	A	A	A	T	T	Q	C	A	T	.	C	Q	T	Q	A	T	Q	Q	Q	Q	A	T	
48 Taxus canadensis	C	T	T	.	.	G	C	C	A	A	A	T	T	Q	C	A	T	.	C	Q	T	Q	A	T	Q	Q	Q	Q	A	T	
49 Adiantum raddeanum	C	T	T	.	.	T	T	Q	A	A	A	T	T	T	C	A	T	.	C	Q	T	Q	A	T	Q	Q	Q	Q	A	T	
50 Adiantum raddeanum 2	C	T	T	.	.	T	T	Q	A	A	A	T	T	T	C	A	T	.	C	Q	T	Q	A	T	Q	Q	Q	Q	A	T	
51 Ophioglossum petiolatum	C	T	T	.	.	T	T	Q	A	A	A	T	T	T	C	A	T	.	C	Q	T	Q	A	T	Q	Q	Q	Q	A	T	
52 Osmunda cinnamomea	C	T	T	.	.	T	T	T	A	A	A	T	T	T	C	A	T	.	C	Q	T	Q	A	T	Q	Q	Q	Q	A	T	
53 Salvinia natans	C	T	T	.	.	T	T	Q	A	A	A	T	T	T	C	A	T	.	C	Q	T	Q	A	T	Q	Q	Q	Q	A	T	
54 Psilotum nudum	C	T	T	.	.	Q	T	Q	A	A	A	T	T	T	C	A	T	.	C	Q	T	Q	A	T	Q	Q	Q	Q	A	T	
55 Chara foeniculacea	C	T	T	.	.	C	T	C	A	A	G	T	T	T	C	A	T	.	C	Q	T	Q	A	T	Q	Q	Q	Q	A	T	
56 Colocletoleto aculeata	C	T	T	.	.	T	T	Q	A	A	A	T	T	T	C	A	T	.	C	Q	T	Q	A	T	Q	Q	Q	Q	A	T	
57 Riccomidium raddeanum	C	T	T	.	.	Q	T	Q	A	A	A	T	T	T	C	A	T	.	C	Q	T	Q	A	T	Q	Q	Q	Q	A	T	
58 Nitella flexilis	C	T	T	.	.	C	T	C	A	A	A	C	T	T	C	A	T	.	C	Q	T	Q	A	T	Q	Q	Q	Q	A	T	
59 Nitella sp	C	T	T	.	.	C	T	C	A	A	A	C	T	C/T	C	A	T	.	C	Q	T	Q	A	T	Q	Q	Q	Q	A	T	
60 Chlamydomonas reinhardtii	C	T	T	.	.	Q	T	A	A	A	A	C	C	Q	C	Q	T	.	C	Q	T	Q	A	T	Q	Q	Q	Q	A	T	
61 Hydrodictyon reticulatum	C	T	T	.	.	T	Q	A	A	A	A	C	T	G	C	A	T	.	C	Q	T	Q	A	T	Q	Q	Q	Q	C	T	
62 Paratetodonia pseudocylindrica	C	T	T	.	.	Q	G	A	A	A	A	C	T	Q	C	A	T	.	C	Q	T	Q	A	T	Q	Q	Q	Q	C	T	
63 Dictyosphaerium discoideum	C	A	T	.	.	T	T	Q	A	A	A	T	T	T	C	A	T	.	C	G	T	A	C	T	Q	Q	Q	Q	C	T	
64 Pleurothallis vittata	C	T	T	.	.	T	T	Q	A	A	A	T	T	T	C	A	T	.	C	Q	T	Q	A	T	Q	Q	Q	Q	A	T	
65 Megascopus beringianus	C	T	T	.	.	T	T	Q	A	A	A	T	T	T	C	A	T	.	C	Q	T	Q	A	T	Q	Q	Q	Q	A	T	
66 Notophthalma bruesii	C	T	G	.	.	C	T	Q	A	A	A	T	T	T	C	A	T	.	C	Q	T	Q	A	T	Q	Q	Q	Q	A	T	
67 Physcomitrella patens	C	T	T	.	.	T	T	Q	A	A	A	T	T	T	C	A	T	.	C	Q	T	Q	A	T	Q	Q	Q	Q	A	T	
68 Sphagnum palustre	C	T	T	.	.	T	T	Q	A	A	A	T	T	T	C	A	T	.	C	Q	T	Q	A	T	Q	Q	Q	Q	A	T	

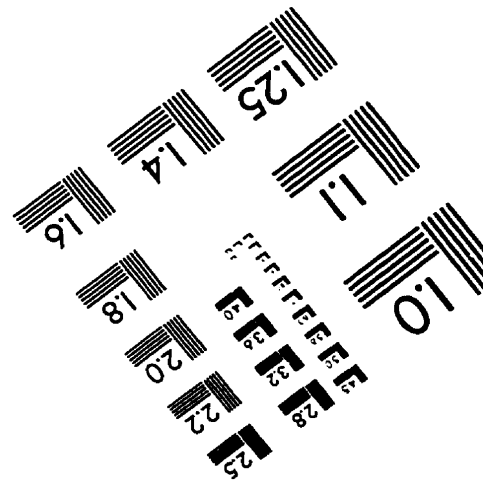
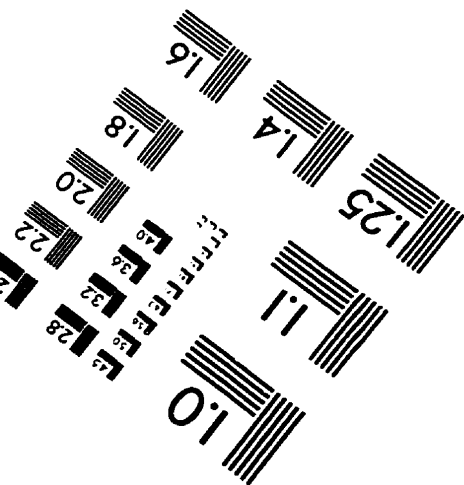
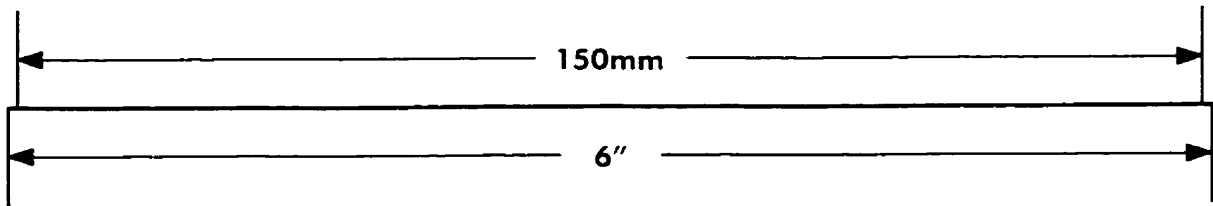
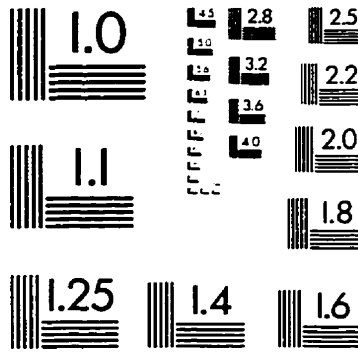
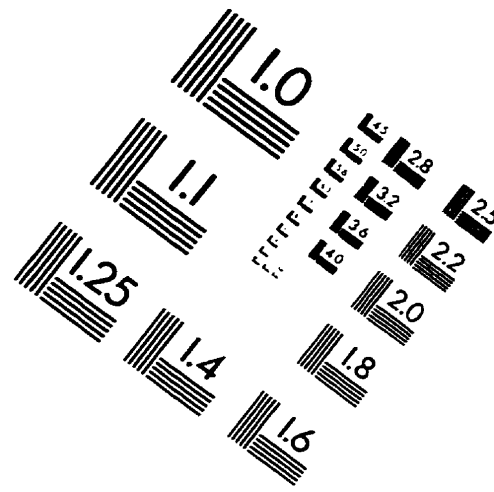
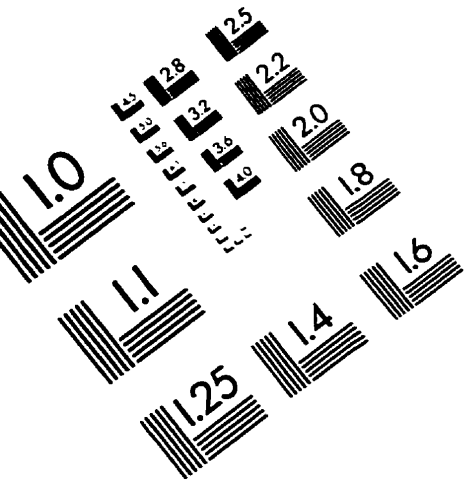
Taxa with Variability Info		00	2011	2012	2013	2014	2015	2016	2017	2018	2019	2020	2021	2022	2023	2024	2025	2026	2027	2028	2029	2030	2031	2032	2033	2034	2035	2036	2037	2038	2039	2040
		0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
1	<i>Saccharomyces cerevisiae</i>	Q	C	Q	T	T	Q	A	T	T	A	C	Q	T	C	C	C	T	Q	C	C	C	C	T	T	-	T	Q	T	A	-	C
2	<i>Schizosaccharomyces pombe</i>	Q	C	Q	T	T	Q	A	A	T	A	C	Q	T	C	C	C	T	Q	C	C	C	T	T	-	T	Q	T	A	-	C	
3	<i>Physcomitrella patens 2</i>	Q	C	Q	T	T	Q	A	C	T	A	C	Q	T	C	C	C	T	Q	C	C	C	T	T	-	T	Q	T	A	-	C	
4	<i>Arabidopsis thaliana</i>	Q	C	Q	T	T	Q	A	C	T	A	C	Q	T	C	C	C	T	Q	C	C	C	T	T	-	T	Q	T	A	-	C	
5	<i>Arabidopsis lyrata</i>	Q	?	Q	T	T	Q	A	C	T	A	C	Q	T	C	C	C	T	Q	C	C	C	T	T	-	T	Q	T	A	-	C	
6	<i>Pellaea epiphylla</i>	Q	C	Q	T	T	Q	A	C	T	A	C	Q	T	C	C	C	T	Q	C	C	C	T	T	-	T	Q	T	A	-	C	
7	<i>Rubus idaeus</i>	Q	C	Q	T	T	Q	A	C	T	A	C	Q	T	C	C	C	T	Q	C	C	C	T	T	-	T	Q	T	A	-	C	
8	<i>Rubus fruticosus</i>	Q	C	Q	T	T	Q	A	C	T	A	C	Q	T	C	C	C	T	Q	C	C	C	T	T	-	T	Q	T	A	-	C	
9	<i>Sphaerocarpos dornali</i>	Q	C	Q	T	T	Q	A	C	T	A	C	Q	T	C	C	C	T	Q	C	C	C	T	T	-	T	Q	T	A	-	C	
10	<i>Sphaerocarpos lasianus</i>	Q	C	Q	T	T	Q	A	C	T	A	C	Q	T	C	C	C	T	Q	C	C	C	T	T	-	T	Q	T	A	-	C	
11	<i>Abrotanum urabatum</i>	Q	C	Q	T	T	Q	A	T	T	A	C	Q	T	C	C	C	T	Q	C	C	C	T	T	-	T	Q	T	A	-	C	
12	<i>Bryum argenteum</i>	Q	C	Q	T	T	Q	A	C	T	A	C	Q	T	C	C	C	T	Q	C	C	C	T	T	-	T	Q	T	A	-	C	
13	<i>Eurhynchium hiemale</i>	Q	C	Q	T	T	Q	A	C	T	A	C	Q	T	C	C	C	T	Q	C	C	C	T	T	-	T	Q	T	A	-	C	
14	<i>Punaria hygrometrica</i>	Q	C	Q	T	T	Q	A	C	T	A	C	Q	T	C	C	C	T	Q	C	C	C	T	T	-	T	Q	T	A	-	C	
15	<i>Lepidobryum pyriforme</i>	Q	C	Q	T	T	Q	A	C	T	A	C	Q	T	C	C	C	T	Q	C	C	C	T	T	-	T	Q	T	A	-	C	
16	<i>Mnium hornum</i>	Q	C	Q	T	T	Q	A	C	T	A	C	Q	T	C	C	C	T	Q	C	C	C	T	T	-	T	Q	T	A	-	C	
17	<i>Physcomitrium pyriforme</i>	Q	C	Q	T	T	Q	A	C	T	A	C	Q	T	C	C	C	T	Q	C	C	C	T	T	-	T	Q	T	A	-	C	
18	<i>Polytrichum commune</i>	Q	C	Q	T	T	Q	A	T	T	A	C	Q	T	C	C	C	T	Q	C	C	C	T	T	-	T	Q	T	A	-	C	
19	<i>Sphagnum cuspidatum</i>	Q	C	Q	T	T	Q	A	C	T	A	C	Q	T	C	C	C	T	Q	C	C	C	T	T	-	T	Q	T	A	-	C	
20	<i>Calyptella arguta</i>	Q	C	A	T	T	Q	A	C	T	A	C	Q	T	C	C	C	T	Q	C	C	C	T	T	-	T	Q	T	A	-	C	
21	<i>Conocarpus elatior</i>	Q	C	Q	T	T	Q	A	T	T	A	C	Q	T	C	C	C	T	Q	C	C	C	T	T	-	T	Q	T	A	-	C	
22	<i>Festuca rubra</i>	Q	C	Q	T	T	Q	A	C	T	A	C	Q	T	C	C	C	T	Q	C	C	C	T	T	-	T	Q	T	A	-	C	
23	<i>Mercurialis polymorpha</i>	Q	C	Q	T	T	Q	A	C	T	A	C	Q	T	C	C	C	T	Q	C	C	C	T	T	-	T	Q	T	A	-	C	
24	<i>Riccia fluitans</i>	Q	C	Q	T	T	Q	A	C	T	A	C	Q	T	C	C	C	T	Q	C	C	C	T	T	-	T	Q	T	A	-	C	
25	<i>Saxifraga hypnoides</i>	Q	C	Q	T	T	Q	A	C	T	A	C	Q	T	C	C	C	T	Q	C	C	C	T	T	-	T	Q	T	A	-	C	
26	<i>Equisetum hyemale</i>	Q	C	Q	T	T	Q	A	C	T	A	C	Q	T	C	C	C	T	Q	C	C	C	T	T	-	T	Q	T	A	-	C	
27	<i>Equisetum robustum</i>	Q	C	Q	T	T	Q	A	C	T	A	C	Q	T	C	C	C	T	Q	C	C	C	T	T	-	T	Q	T	A	-	C	
28	<i>Isetes engelmannii</i>	Q	C	Q	T	T	Q	A	C	T	A	C	Q	T	C	C	C	T	Q	C	C	C	T	T	-	T	Q	T	A	-	C	
29	<i>Hypozia lucidulum</i>	?	?	?	T	T	Q	A	C	T	A	C	Q	T	C	C	C	T	Q	C	C	C	T	T	-	T	Q	T	A	-	C	
30	<i>Lycopodium inundatum</i>	Q	C	Q	T	T	Q	A	C	T	A	C	Q	T	C	C	C	T	Q	C	C	C	T	T	-	T	Q	T	A	-	C	
31	<i>Lycopodium phlegmaria</i>	Q	C	Q	T	T	Q	A	C	T	A	C	Q	T	C	C	C	T	Q	C	C	C	T	T	-	T	Q	T	A	-	C	
32	<i>Lycopodium lasiolobum</i>	Q	C	Q	T	T	Q	A	C	T	A	C	Q	T	C	C	C	T	Q	C	C	C	T	T	-	T	Q	T	A	-	C	
33	<i>Lycopodium tristachyum</i>	Q	C	Q	T	T	Q	A	C	T	A	C	Q	T	C	C	C	T	Q	C	C	C	T	T	-	T	Q	T	A	-	C	
34	<i>Salvinia selaginoides</i>	Q	C	Q	T	T	Q	A	T	T	A	C	Q	T	C	C	C	T	Q	C	C	C	T	T	-	T	Q	T	A	-	C	
35	<i>Salvinia selaginoides</i>	Q	C	Q	T	T	Q	A	C	T	A	C	Q	T	C	C	C	T	Q	C	C	C	T	T	-	T	Q	T	A	-	C	
36	<i>Oryza sativa</i>	Q	C	Q	T	T	Q	A	C	T	A	C	Q	T	C	C	C	T	Q	C	C	C	T	T	-	T	Q	T	A	-	C	
37	<i>Zea mays</i>	Q	C	Q	T	T	Q	A	C	T	A	C	Q	T	C	C	C	T	Q	C	C	C	T	T	-	T	Q	T	A	-	C	
38	<i>Arabidopsis thaliana</i>	Q	C	Q	T	T	Q	A	C	T	A	C	Q	T	C	C	C	T	Q	C	C	C	T	T	-	T	Q	T	A	-	C	
39	<i>Glycine max</i>	Q	C	Q	T	T	Q	A	C	T	A	C	Q	T	C	C	C	T	Q	C	C	C	T	T	-	T	Q	T	A	-	C	
40	<i>Syntherisma</i>	Q	C	Q	T	T	Q	A	C	T	A	C	Q	T	C	C	C	T	Q	C	C	C	T	T	-	T	Q	T	A	-	C	
41	<i>Zinnia purpurea</i>	Q	C	Q	T	T	Q	A	C	T	A	C	Q	T	C	C	C	T	Q	C	C	C	T	T	-	T	Q	T	A	-	C	
42	<i>Gnaphalium leucolobum</i>	Q	C	Q	T	T	Q	A	C	T	A	C	Q	T	C	C	C	T	Q	C	C	C	T	T	-	T	Q	T	A	-	C	
43	<i>Girardinia</i>	Q	C	Q	T	T	Q	A	C	T	A	C	Q	T	C	C	C	T	Q	C	C	C	T	T	-	T	Q	T	A	-	C	
44	<i>Pinus luchuanensis</i>	Q	T	Q	T	T	Q	A	C	T	A	C	Q	T	C	C	C	T	Q	C	C	C	T	T	-	T	Q	T	A	-	C	
45	<i>Pinus wallichiana</i>	Q	T	Q	T	T	Q	A	C	T	A	C	Q	T	C	C	C	T	Q	C	C	C	T	T	-	T	Q	T	A	-	C	
46	<i>Nageia nageia</i>	Q	C	Q	T	T	Q	A	C	T	A	C	Q	T	C	C	C	T	Q	C	C	C	T	T	-	T	Q	T	A	-	C	
47	<i>Taxus mairei</i>	Q	C	Q	T	T	Q	A	C	T	A	C	Q	T	C	C	C	T	Q	C	C	C	T	T	-	T	Q	T	A	-	C	
48	<i>Adiantum radicans</i>	Q	C	Q	T	T	Q	A	C	T	A	C	Q	T	C	C	C	T	Q	C	C	C	T	T	-	T	Q	T	A	-	C	
49	<i>Adiantum radicans 2</i>	Q	C	Q	T	T	Q	A	C	T	A	C	Q	T	C	C	C	T	Q	C	C	C	T	T	-	T	Q	T	A	-	C	
50	<i>Ophioglossum petiolatum</i>	Q	C	Q	T	T	Q	A	C	T	A	C	Q	T	C	C	C	T	Q	C	C	C	T	T	-	T	Q	T	A	-	C	
51	<i>Camunda chinensis</i>	Q	C	Q	T	T	Q	A	C	T	A	C	Q	T	C	C	C	T	Q	C	C	C	T	T	-	T	Q	T	A	-	C	
52	<i>Salvinia natans</i>	Q	C	Q	T	T	Q	A	C	T	A	C	Q	T	C	C	C	T	Q	C	C	C	T	T	-	T	Q	T	A	-	C	
53	<i>Pellaea nudum</i>	Q	C	Q	T	T	Q	A	C	T	A	C	Q	T	C	C	C	T	Q	C	C	C	T	T	-	T	Q	T	A	-	C	
54	<i>Chara foeniculacea</i>	Q	C	Q	T	T	Q	A	C	T	A	C	Q	T	C	C	C	T	Q	C	C	C	T	T	-	T	Q	T	A	-	C	
55	<i>Coleochaete ecitata</i>	Q	C	Q	T	T	Q	A	T	T	A	C	Q	T	C	C	C	T	Q	C	C	C	T	T	-	T	Q	T	A	-	C	
56	<i>Kribiastrum flaccidum</i>	Q	C	Q	T	T	Q	A	T	T	A	C	Q	T	C	C	C	T	Q	C	C	C	T	T	-	T	Q	T	A	-	C	
57	<i>Nitzschia flexilis</i>	Q	C	Q	T	T	Q	A	T	T	A	C	Q	T	C	C	C	T	Q	C	C	C	T	T	-	T	Q	T	A	-	C	
58	<i>Nitzschia sp.</i>	Q	C	Q	T	T	Q	A	T	T	A	C	Q	T	C	C	C	T	Q	C	C	C	T	T	-	T	Q	T	A	-	C	
59	<i>Chlamydomonas reinhardtii</i>	Q	C	Q	T	T	Q	A	T	T	A	C	Q	T	C	C	C	T	Q	C	C	C	T	T	-	T	Q	T	A	-	C	
60	<i>Hydrodictyon reticulatum</i>	Q	C	Q	T	T	Q	A	T	T	A	C	Q	T	C	C	C	T	Q	C	C	C	T	T	-	T	Q	T	A	-	C	
61	<i>Parietaria pseudocochlearifera</i>	Q	C	Q	T	T	Q	A	T	T	A	C	Q	T	C	C	C	T	Q	C	C	C	T	T	-	T	Q	T	A	-	C	
62	<i>Dicystotium decoides</i>	A	T	Q	C	T	Q	A	C	T	A	C	Q	T	C	C	C	T	Q	C	C	C	T	T	-	T	Q	T	A	-	C	
63	<i>Pteris vittata</i>	Q	C	Q	T	T	Q	A	C	T	A	C	Q	T	C	C	C	T	Q	C	C	C	T	T	-	T	Q	T	A	-	C	
64	<i>Megaceros sibiricus</i>	Q	C	Q	T	T	Q	A	C	T	A	C	Q	T	C	C	C	T	Q	C	C	C	T	T	-	T	Q	T	A	-	C	
65	<i>Neophytia brevis</i>	Q	C	Q	T	T	Q	A	C	T	A	C	Q	T																		

	2131	2132	2133	2134	2136	2138	2137	2139	2140	2141	2142	2143	2144	2145	2146	2147	2148	2149	2150	2151	2152	2153	2154	2155	2156	2157	2158	2159				
	0	0	0	2	2	2	0	1	2	1	4	5	6	5	0	3	5	4	2	2	2	2	2	6	4	9	2	0	3			
1	<i>Saccharomyces cerevisiae</i>	.	.	.	A	.	G	A	.	G	C	Q	G	A	Q	T	T	T	Q	G	A	C	A	A	A	C	T	.	T			
2	<i>Schizosaccharomyces pombe</i>	.	.	.	C	.	A	T	T	Q	C	C	Q	A	Q	A	A	Q	T	T	Q	A	C	A	A	A	C	T	.	T		
3	<i>Physcomitrella patens 2</i>	.	.	.	A	C	Q	T	.	T	Q	Q	Q	A	Q	A	A	Q	T	T	C	A	T	T	A	A	A	C	C	.	T	
4	<i>Anthoceros agrestis</i>	.	.	.	A	C	Q	T	.	T	Q	T	G	A	Q	A	A	Q	T	T	C	A	T	T	A	A	A	C	C	.	T	
5	<i>Anthoceros laevis</i>	.	.	.	A	C	Q	T	.	T	Q	T	G	A	Q	A	A	Q	T	T	C	A	T	T	A	A	A	C	C	.	T	
6	<i>Pellia epiphylla</i>	.	.	.	A	C	Q	T	.	T	Q	T	G	A	Q	A	A	Q	T	T	C	A	T	T	A	A	A	C	C	.	T	
7	<i>Reboulia hemisphaerica</i>	.	.	.	A	C	Q	T	.	T	Q	T	G	A	Q	A	A	Q	T	T	C	T	T	T	A	A	A	C	C	.	T	
8	<i>Riccardia pinguis</i>	.	.	.	A	C	Q	T	.	T	Q	T	G	A	Q	A	A	Q	T	T	C	A	T	T	A	A	A	C	C	.	T	
9	<i>Sphaerocarpos donnellii</i>	.	.	.	A	C	Q	T	.	T	Q	T	Q	A	Q	A	A	Q	T	C	C	T	T	T	A	A	A	C	C	.	T	
10	<i>Sphaerocarpos lazarus</i>	.	.	.	A	C	Q	T	.	T	Q	T	G	A	Q	A	A	Q	.	T	C	C	T	T	A	A	A	C	C	.	T	
11	<i>Atrichum undulatum</i>	.	.	.	A	C	Q	T	.	T	Q	T	Q	A	Q	A	A	Q	T	T	C	A	T	T	A	A	A	C	C	.	T	
12	<i>Bryum argenteum</i>	.	.	.	A	C	Q	T	.	T	Q	T	Q	A	Q	A	A	Q	T	T	C	A	T	T	A	A	A	C	C	.	T	
13	<i>Eurymochium hiemale</i>	.	.	.	T	C	Q	T	.	T	Q	T	A	Q	Q	A	A	Q	T	T	C	A	T	T	A	A	A	C	C	.	T	
14	<i>Funaria hygrometrica</i>	.	.	.	A	C	Q	T	.	T	Q	Q	Q	A	Q	A	A	Q	T	T	C	A	T	T	A	A	A	C	G	.	T	
15	<i>Leptobryum pyriforme</i>	.	.	.	A	C	Q	T	.	T	Q	T	Q	A	Q	A	A	Q	T	T	C	A	T	T	A	A	A	C	C	.	T	
16	<i>Mnium hornum</i>	.	.	.	A	C	Q	T	.	T	Q	T	G	A	Q	A	A	Q	T	T	C	A	T	T	A	A	A	C	C	.	T	
17	<i>Physcomitrium pyriforme</i>	.	.	.	A	Q	C	T	.	T	Q	Q	T	A	Q	A	A	Q	T	T	C	A	C	C	A	A	A	C	C	.	T	
18	<i>Polytrichum lomesum</i>	.	.	.	A	C	Q	T	.	T	Q	T	Q	A	Q	A	A	Q	T	T	C	A	T	T	A	A	A	C	C	.	T	
19	<i>Sphagnum cuspidatum</i>	.	.	.	A	C	Q	T	.	T	Q	T	Q	A	Q	A	A	Q	T	T	C	A	T	T	A	A	A	C	C	.	T	
20	<i>Calypogeia arguta</i>	.	.	.	T	C	Q	T	.	T	Q	T	Q	A	A	Q	A	Q	T	T	C	A	T	T	A	A	A	C	G	.	T	
21	<i>Oreosiphium auriculatum</i>	.	.	.	A	C	Q	T	.	T	Q	T	Q	A	Q	A	A	Q	T	T	C	T	T	T	A	A	A	C	C	.	T	
22	<i>Fossombronia pusilla</i>	.	.	.	A	C	Q	T	.	T	Q	T	Q	A	Q	A	A	Q	T	T	C	A	T	T	A	A	A	C	C	.	T	
23	<i>Marchantia polymorpha</i>	.	.	.	A	C	Q	T	.	T	Q	T	Q	A	Q	A	A	Q	T	C	C	T	T	T	A	A	A	C	C	.	T	
24	<i>Riccia fluitans</i>	.	.	.	A	C	Q	T	.	T	Q	T	G	A	Q	A	A	Q	T	T	C	T	T	T	A	A	A	C	C	.	T	
25	<i>Scapania nemorensis</i>	.	.	.	A	C	Q	T	.	T	Q	T	Q	A	Q	A	A	Q	T	T	C	A	T	T	A	A	A	C	C	.	T	
26	<i>Equisetum hyemale</i>	.	.	.	A	C	Q	T	.	T	Q	T	Q	A	Q	A	A	Q	T	T	C	A	T	T	A	A	A	C	C	.	T	
27	<i>Equisetum robustum</i>	.	.	.	A	C	Q	T	.	T	Q	T	Q	A	Q	A	A	Q	T	T	C	A	T	T	A	A	A	C	C	.	T	
28	<i>Isetes engelmannii</i>	.	.	.	A	C	Q	T	.	T	Q	T	Q	A	Q	A	A	Q	T	T	C	A	T	T	A	A	A	C	C	.	T	
29	<i>Huperzia lucidulum</i>	.	.	.	A	C	Q	T	.	T	Q	T	Q	A	Q	A	A	Q	T	T	C	A	T	T	A	A	A	C	C	.	T	
30	<i>Lycopodium truncatum</i>	.	.	.	A	C	Q	T	.	T	Q	T	Q	A	Q	A	A	Q	T	T	C	A	T	T	A	A	A	C	C	.	T	
31	<i>Lycopodium phlegmaria</i>	.	.	.	A	C	Q	T	.	T	Q	T	Q	A	Q	A	A	Q	T	T	C	A	T	T	A	A	A	C	C	.	T	
32	<i>Lycopodium lasiolobum</i>	.	.	.	A	C	Q	T	.	T	Q	T	Q	A	Q	A	A	Q	T	T	C	A	T	T	A	A	A	C	C	.	T	
33	<i>Lycopodium metacladum</i>	.	.	.	A	C	Q	T	.	T	Q	T	Q	A	Q	A	A	Q	T	T	C	A	T	T	A	A	A	C	C	.	T	
34	<i>Selaginella umbrosa</i>	.	.	.	A	C	Q	T	.	T	Q	T	Q	A	Q	A	A	Q	T	T	C	A	T	T	A	A	A	C	C	.	T	
35	<i>Selaginella vogelii</i>	.	.	.	A	C	Q	T	.	T	Q	T	Q	A	Q	A	A	Q	T	T	C	A	T	T	A	A	A	C	C	.	T	
36	<i>Cryza selva</i>	.	.	.	A	C	Q	T	.	T	C	Q	C	Q	A	Q	A	A	Q	T	C	C	A	T	T	A	A	A	C	C	.	T
37	<i>Zea mays</i>	A	C	C	Q	T	C	C	.	Q	C	C	Q	A	Q	A	A	Q	T	C	C	A	T	T	A	A	A	C	C	.	T	
38	<i>Arabidopsis thaliana</i>	.	.	.	A	C	Q	T	.	T	C	Q	C	Q	A	Q	A	A	Q	T	C	C	A	C	T	A	A	A	C	C	.	T
39	<i>Glycine max</i>	.	.	.	A	C	Q	T	.	T	Q	T	Q	A	Q	A	A	Q	T	T	C	A	C	T	A	A	A	C	C	.	T	
40	<i>Brassica alba</i>	.	.	.	A	C	Q	T	.	T	C	Q	C	Q	A	Q	A	A	Q	T	T	C	T	C	T	A	A	A	C	C	.	T
41	<i>Zinnia purpurea</i>	.	.	.	A	C	Q	T	.	T	C	Q	C	Q	A	Q	A	A	Q	T	T	C	A	T	T	A	A	A	C	C	.	T
42	<i>Onoclea lyallii</i>	.	.	.	A	C	Q	T	.	T	C	Q	T	Q	A	Q	A	A	Q	T	T	C	A	T	T	A	A	A	C	C	.	T
43	<i>Onoclea sensibilis</i>	.	.	.	A	C	Q	T	.	T	C	Q	C	Q	A	Q	A	A	Q	T	T	C	A	T	T	A	A	A	C	C	.	T
44	<i>Pinus luchuanensis</i>	.	.	.	A	C	Q	T	.	T	C	Q	T	Q	A	Q	A	A	Q	T	T	C	A	T	T	A	A	A	C	C	.	T
45	<i>Pinus wallichiana</i>	.	.	.	A	C	Q	T	.	T	C	Q	T	Q	A	Q	A	A	Q	T	T	C	A	T	T	A	A	A	C	C	.	T
46	<i>Negundo negundo</i>	.	.	.	A	C	Q	T	.	T	Q	C	Q	A	Q	A	A	Q	T	T	C	A	T	T	A	A	A	C	C	.	T	
47	<i>Taxus mairei</i>	.	.	.	A	T	Q	T	.	T	Q	C	Q	A	Q	A	A	Q	T	T	C	A	T	T	A	A	A	C	C	.	T	
48	<i>Adiantum radicans</i>	.	.	.	A	C	Q	T	.	T	Q	T	Q	A	Q	A	A	Q	T	T	C	A	T	T	A	A	A	C	C	.	T	
49	<i>Adiantum radicans 2</i>	.	.	.	A	C	Q	T	.	T	Q	T	Q	A	Q	A	A	Q	T	T	C	A	T	T	A	A	A	C	C	.	T	
50	<i>Ophioglossum petiolatum</i>	.	.	.	?	?	?	?	.	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?	?
51	<i>Osmunda cinnamomea</i>	.	.	.	A	C	Q	T	.	T	Q	T	Q	A	Q	A	A	Q	T	T	C	A	T	T	A	A	A	C	C	.	T	
52	<i>Selaginella selaginoides</i>	.	.	.	A	C	Q	T	.	T	Q	T	Q	A	Q	A	A	Q	T	T	C	A	T	T	A	A	A	C	C	.	T	
53	<i>Pellaea rotundifolia</i>	.	.	.	A	C	Q	T	.	T	Q	T	Q	A	Q	A	A	Q	T	T	C	A	T	T	A	A	A	C	C	.	T	
54	<i>Chara foeniculacea</i>	.	.	.	G	T	G	C	.	T	Q	T	Q	A	Q	A	A	Q	T	T	C	A	T	T	A	A	A	C	C	.	T	
55	<i>Coleochaete scutellata</i>	.	.	.	A	C	Q	T	.	T	Q	T	Q	A	Q	A	A	Q	T	T	C	A	T	T	A	A	A	C	C	.	T	
56	<i>Habrocallia laevigata</i>	.	.	.	A	A	Q	C	.	T	Q	T	Q	A	Q	A	A	Q	T	T	C	A	T	T	A	A	A	C	C	.	T	
57	<i>Hibella laevigata</i>	.	.	.	G	C	Q	C	.	T	Q	T	Q	A	Q	A	A	Q	T	T	C	A	T	T	A	A	A	C	C	.	T	
58	<i>Hibella sp.</i>	.	.	.	G	C	Q	C	.	T	Q	T	Q	A	Q	A	A	Q	T	T	C	A	T	T	A	A	A	C	C	.	T	
59	<i>Chlamydomonas reinhardtii</i>	.	.	.	C	T	T	G	.	C	T	T	Q	A	Q	A	A	Q	T	T	C	A	T	T	A	A	A	C	C	.	C	
60	<i>Hydrodictyon reticulatum</i>	.	.	.	T	A	C	T	.	T	C	C	C	Q	A	Q	A	A	Q	T	T	C	A	T	T	A	A	A	C	C	.	C
61	<i>Parietococcus pseudovalvularis</i>	.	.	.	T	Q	C	A	.	T	C	C	Q	A	Q	A	A	Q	T	T	C	A	T	T	A	A	A	C	C	.	C	
62	<i>Diatylopusium discoides</i>	.	.	.	T	A	C	.	.	A	A	T	T	A	A	A	A	Q	T	T	A	T	T	T	A	A	A	T	C	.	T	
63	<i>Pteris vittata</i>	.	.	.	A	C	Q	T	.	T	Q	T	Q	A	Q	A	A	Q	T	T	C	A	T	T	A	A	A	C	C	.	T	
64	<i>Megaceros senigallensis</i>	.	.	.	A	C	Q	T	.	T	Q	T	Q	A	Q	A	A	Q	T	T	C	A	T	T	A	A	A	C	C	.	T	
65	<i>Notophycis brucei</i>	.	.	.	A	C	Q	T	.	T	Q	T	Q	A	Q	A	A	Q	T	T	C	A	T	C	A	A	A	C	C	.	T	
66	<i>Physcomitrella patens</i>	.	.	.	A	C	Q	T	.	T	Q	T	Q	A	Q	A	A	Q	T	T	C	A	T	T	A	A	A	C	C	.	T	
67	<i>Sphagnum palustre</i>	.	.	.	A	C	Q	T	.	T	Q	T	Q	A	Q	A	A	Q	T													

6	<i>Anthroceros laevis</i>	T	A	T	C	A	T	T	T	A	G	A	G	G	A	A	G	G	A	G	A	A	G	T	C	G	T	-	-	-	-	A	
6	<i>Pellaea epiphylla</i>	T	A	T	C	A	T	T	T	A	G	A	G	G	A	A	G	G	A	G	A	A	G	T	C	G	T	-	-	-	-	A	
7	<i>Rabouia hemisphaerica</i>	T	A	T	C	A	T	T	T	A	G	A	G	G	A	A	G	G	A	G	A	A	G	T	C	G	T	-	-	-	-	A	
8	<i>Riccardia pinguis</i>	T	A	T	C	A	T	T	T	A	G	A	G	G	A	A	G	G	A	G	A	A	G	T	C	G	T	-	-	-	-	A	
9	<i>Sphaerocarpos donnellii</i>	T	A	T	C	A	T	T	T	A	G	A	G	G	A	A	G	G	A	G	A	A	G	T	C	G	T	-	-	-	-	A	
10	<i>Sphaerocarpos lantanus</i>	T	A	T	C	A	T	T	T	A	G	A	G	G	A	A	G	G	A	G	A	A	G	T	C	G	T	-	-	-	-	A	
11	<i>Atichium undulatum</i>	T	A	T	C	A	T	T	T	A	G	A	G	G	A	A	G	G	A	G	A	A	G	T	C	G	T	-	-	-	-	A	
12	<i>Bryum argenteum</i>	T	A	T	C	A	T	T	T	A	G	A	G	G	A	A	G	G	A	G	A	A	G	T	C	G	T	-	-	-	-	A	
13	<i>Eurynchium hiemale</i>	T	A	T	C	A	T	T	T	A	G	A	G	G	A	A	G	G	A	G	A	A	G	T	C	G	T	-	-	-	-	A	
14	<i>Punaria hygrometrica</i>	T	A	T	C	A	T	T	T	A	G	A	G	G	A	A	G	G	A	G	A	A	G	T	C	G	T	-	-	-	-	A	
15	<i>Leptobryum pyriforme</i>	T	A	T	C	A	T	T	T	A	G	A	G	G	A	A	G	G	A	G	A	A	G	T	C	G	T	-	-	-	-	A	
16	<i>Mnium hornum</i>	T	A	T	C	A	T	T	T	A	G	A	G	G	A	A	G	G	A	G	A	A	G	T	C	G	T	-	-	-	-	A	
17	<i>Physcomitrium pyriforme</i>	T	A	T	C	A	T	T	T	A	G	A	G	G	A	A	G	G	A	G	A	A	G	T	C	G	T	-	-	-	-	A	
18	<i>Polypodium complanatum</i>	T	A	T	C	A	T	T	T	A	G	A	G	G	A	A	G	G	A	G	A	A	G	T	C	G	T	-	-	-	-	A	
19	<i>Sphagnum cuspidatum</i>	T	A	T	C	A	T	T	T	A	G	A	G	G	A	A	G	G	A	G	A	A	G	T	C	G	T	-	-	-	-	A	
20	<i>Calypogeia arguta</i>	T	A	T	C	A	T	T	T	A	G	A	G	G	A	A	G	G	A	G	A	A	G	T	C	G	T	-	-	-	-	A	
21	<i>Conocophium conicum</i>	T	A	T	C	A	T	T	T	A	G	A	G	G	A	A	G	G	A	G	A	A	G	T	C	G	T	-	-	-	-	A	
22	<i>Fossombronia pusilla</i>	T	A	T	C	A	T	T	T	A	G	A	G	G	A	A	G	G	A	G	A	A	G	T	C	G	T	-	-	-	-	A	
23	<i>Manzanilla polymorpha</i>	T	A	T	C	A	T	T	T	A	G	A	G	G	A	A	G	G	A	G	A	A	G	T	C	G	T	-	-	-	-	A	
24	<i>Riccia fluitans</i>	T	A	T	C	A	T	T	T	A	G	A	G	G	A	A	G	G	A	G	A	A	G	T	C	G	T	-	-	-	-	A	
25	<i>Saxipha nemorea</i>	T	A	T	C	A	T	T	T	A	G	A	G	G	A	A	G	G	A	G	A	A	G	T	C	G	T	-	-	-	-	A	
26	<i>Equisetum hyemale</i>	T	A	C	C	A	T	T	T	A	G	A	G	G	A	A	G	G	A	G	A	A	G	T	C	G	T	-	-	-	-	A	
27	<i>Equisetum robustum</i>	T	A	C	C	A	T	T	T	A	G	A	G	G	A	A	G	G	A	G	A	A	G	T	C	G	T	-	-	-	-	A	
28	<i>Isetes engelmannii</i>	T	A	T	C	A	T	T	T	A	G	A	G	G	A	A	G	G	A	G	A	A	G	T	C	G	T	-	-	-	-	A	
29	<i>Huperzia lucidulum</i>	T	A	T	C	A	T	T	T	A	G	A	G	G	A	A	G	G	A	G	A	A	G	T	C	G	T	-	-	-	-	A	
30	<i>Lycopodium inundatum</i>	T	A	C	C	A	T	T	T	A	G	A	G	G	A	A	G	G	A	G	A	A	G	T	C	G	T	-	-	-	-	A	
31	<i>Lycopodium phlegmaria</i>	T	A	T	C	A	T	T	T	A	G	A	G	G	A	A	G	G	A	G	A	A	G	T	C	G	T	-	-	-	-	A	
32	<i>Lycopodium lasiolobum</i>	T	A	T	C	A	T	T	T	A	G	A	G	G	A	A	G	G	A	G	A	A	G	T	C	G	T	-	-	-	-	A	
33	<i>Lycopodium tristichium</i>	T	A	T	C	A	T	T	T	A	G	A	G	G	A	A	G	G	A	G	A	A	G	T	C	G	T	-	-	-	-	A	
34	<i>Selaginella umbrosa</i>	T	A	C	T	A	T	C	T	A	G	A	G	G	A	A	G	G	A	G	A	A	G	T	C	G	T	-	-	-	-	A	
35	<i>Selaginella vogelii</i>	T	A	T	C	A	T	T	T	A	G	A	G	G	A	A	G	G	A	G	A	A	G	T	C	G	T	-	-	-	-	A	
36	<i>Oryza sativa</i>	T	A	T	C	A	T	T	T	A	G	A	G	G	A	A	G	G	A	G	A	A	G	T	C	G	T	-	-	-	-	A	
37	<i>Zea mays</i>	T	A	T	C	A	T	T	T	A	G	A	G	G	A	A	G	G	A	G	A	A	G	T	C	G	T	-	-	-	-	A	
38	<i>Arabis alpina</i>	T	A	T	C	A	T	T	T	A	G	A	G	G	A	A	G	G	A	G	A	A	G	T	C	G	T	-	-	-	-	A	
39	<i>Olychnis saxifraga</i>	T	A	T	C	A	T	T	T	A	G	A	G	G	A	A	G	G	A	G	A	A	G	T	C	G	T	-	-	-	-	A	
40	<i>Strapsis alba</i>	T	A	T	C	A	T	T	T	A	G	A	G	G	A	A	G	G	A	G	A	A	G	T	C	G	T	-	-	-	-	A	
41	<i>Zinnia purpurea</i>	T	A	T	C	A	T	T	T	A	G	A	G	G	A	A	G	G	A	G	A	A	G	T	C	G	T	-	-	-	-	A	
42	<i>Gnaphalium polycephalum</i>	T	A	T	C	A	T	T	T	A	G	A	G	G	A	A	G	G	A	G	A	A	G	T	C	G	T	-	-	-	-	A	
43	<i>Olychnis saxifraga</i>	T	A	T	C	A	T	T	T	A	G	A	G	G	A	A	G	G	A	G	A	A	G	T	C	G	T	-	-	-	-	A	
44	<i>Pinus banksiana</i>	T	A	T	C	A	T	T	T	A	G	A	G	G	A	A	G	G	A	G	A	A	G	T	C	G	T	-	-	-	-	A	
45	<i>Pinus strobus</i>	T	A	T	C	A	T	T	T	A	G	A	G	G	A	A	G	G	A	G	A	A	G	T	C	G	T	-	-	-	-	A	
46	<i>Pinus taeda</i>	T	A	T	C	A	T	T	T	A	G	A	G	G	A	A	G	G	A	G	A	A	G	T	C	G	T	-	-	-	-	A	
47	<i>Taxus canadensis</i>	T	A	T	C	A	T	T	T	A	G	A	G	G	A	A	G	G	A	G	A	A	G	T	C	G	T	-	-	-	-	A	
48	<i>Adiantum pedatum</i>	T	A	T	C	A	T	T	T	A	G	A	G	G	A	A	G	G	A	G	A	A	G	T	C	G	T	-	-	-	-	A	
49	<i>Adiantum pedatum 2</i>	T	A	T	C	A	T	T	T	A	G	A	G	G	A	A	G	G	A	G	A	A	G	T	C	G	T	-	-	-	-	A	
50	<i>Ophioglossum petiolatum</i>	T	-	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	A
51	<i>Osmunda cinnamomea</i>	T	A	T	C	A	T	T	T	A	G	A	G	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	A	
52	<i>Selvinia selaginella</i>	T	A	T	C	A	T	T	T	A	G	A	G	G	A	A	G	G	A	G	A	A	G	T	C	G	T	-	-	-	-	A	
53	<i>Pellaea nudum</i>	T	A	T	C	A	T	T	T	A	G	A	G	G	A	A	G	G	A	G	A	A	G	T	C	G	T	-	-	-	-	A	
54	<i>Chara foetida</i>	T	A	T	C	A	T	T	T	A	G	A	G	G	A	A	G	G	A	G	A	A	G	T	C	G	T	-	-	-	-	A	
55	<i>Colobosolenia setacea</i>	T	A	T	C	A	T	T	T	A	G	A	G	G	A	A	G	G	A	G	A	A	G	T	C	G	T	-	-	-	-	A	
56	<i>Habenaria lucida</i>	T	A	T	C	A	T	T	T	A	G	A	G	G	A	A	G	G	A	G	A	A	G	T	C	G	T	-	-	-	-	A	
57	<i>Nitella flexilis</i>	T	A	T	C	A	T	T	T	A	G	A	G	G	A	A	G	G	A	G	A	A	G	T	C	G	T	-	-	-	-	A	
58	<i>Nitella sp.</i>	T	A	T	C	A	T	T	T	A	G	A	G	G	A	A	G	G	A	G	A	A	G	T	C	G	T	-	-	-	-	A	
59	<i>Chlamydomonas reinhardtii</i>	T	C	C	C	A	C	C	T	A	G	A	G	G	A	A	G	G	A	G	A	A	G	T	C	G	T	-	-	-	-	A	
60	<i>Hydrodictyon reticulatum</i>	T	C	C	C	A	C	C	T	A	G	A	G	G	A	A	G	G	A	G	A	A	G	T	C	G	T	-	-	-	-	A	
61	<i>Parietaria pseudoalveolata</i>	T	C	C	C	A	C	C	T	A	G	A	G	T	A	A	G	C	A	G	A	A	G	T	C	G	T	-	-	-	-	A	
62	<i>Dictyosporium decoloratum</i>	C	A	T	T	G	T	T	T	A	G	A	G	G	A	A	G	G	A	G	A	A	G	T	C	G	T	-	-	-	-	A	
63	<i>Pteris vittata</i>	T	A	T	C	A	T	T	T	A	G	A	G	G	A	A	G	G	A	G	A	A	G	T	C	G	T	-	-	-	-	A	
64	<i>Megaceros sentinae</i>	T	A	T	C	A	T	T	T	A	G	A	G	G	A	A	G	G	A	G	A	A	G	T	C	G	T	-	-	-	-	A	
65	<i>Notophras brevis</i>	T	A	T	C	A	T	T	T	A	G	A	G	G	A	A	G	G	A	G	A	A	G	T	C	G	T	-	-	-	-	A	
66	<i>Physcomitrella patens</i>	T	A	T	C	A	T	T	T	A	G	A	G	G	A	A	G	G	A	G	A	A	G	T	C	G	T	-	-	-	-	A	
67	<i>Sphagnum palustre</i>	T	A	T	C	A	T	T	T	A	G	A	G	G	A	A	G	G	A	G	A	A	G	T	C	G	T	-	-	-	-	A	

	0	0	0	0
1	<i>Saccharomyces cerevisiae</i>	.	.	.
2	<i>Schizosaccharomyces pombe</i>	.	.	.
3	<i>Physcomitrella patens 2</i>	.	.	.
4	<i>Arabidopsis lyrata</i>	.	.	.
5	<i>Arabidopsis lyrata</i>	.	.	.
6	<i>Pellaea atropurpurea</i>	.	.	.
7	<i>Poa annua</i>	.	.	.
8	<i>Physcomitrella patens</i>	.	.	.
9	<i>Physcomitrella patens</i>	.	.	.
10	<i>Physcomitrella patens</i>	.	.	.
11	<i>Arabidopsis thaliana</i>	.	.	.
12	<i>Bryum alpinum</i>	.	.	.
13	<i>Bryum alpinum</i>	.	.	.
14	<i>Paradeia lycopodioides</i>	7	A	G
15	<i>Lepidobryum pyriforme</i>	.	.	.
16	<i>Bryum hirtum</i>	.	.	.
17	<i>Physcomitrella patens</i>	.	.	.
18	<i>Physcomitrella patens</i>	.	.	.
19	<i>Physcomitrella patens</i>	.	.	.
20	<i>Calyptrella spida</i>	.	.	.
21	<i>Physcomitrella patens</i>	.	.	.
22	<i>Paradeia lycopodioides</i>	.	.	.
23	<i>Physcomitrella patens</i>	.	.	.
24	<i>Physcomitrella patens</i>	.	.	.
25	<i>Physcomitrella patens</i>	.	.	.
26	<i>Physcomitrella patens</i>	.	.	.
27	<i>Physcomitrella patens</i>	.	.	.
28	<i>Physcomitrella patens</i>	.	.	.
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30	<i>Physcomitrella patens</i>	.	.	.
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34	<i>Physcomitrella patens</i>	.	.	.
35	<i>Physcomitrella patens</i>	.	.	.
36	<i>Physcomitrella patens</i>	.	.	.
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38	<i>Physcomitrella patens</i>	.	.	.
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41	<i>Physcomitrella patens</i>	.	.	.
42	<i>Physcomitrella patens</i>	.	.	.
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44	<i>Physcomitrella patens</i>	.	.	.
45	<i>Physcomitrella patens</i>	.	.	.
46	<i>Physcomitrella patens</i>	.	.	.
47	<i>Physcomitrella patens</i>	.	.	.
48	<i>Physcomitrella patens</i>	.	.	.
49	<i>Physcomitrella patens 2</i>	.	.	.
50	<i>Physcomitrella patens</i>	.	.	.
51	<i>Physcomitrella patens</i>	.	.	.
52	<i>Physcomitrella patens</i>	.	.	.
53	<i>Physcomitrella patens</i>	.	.	.
54	<i>Physcomitrella patens</i>	.	.	.
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56	<i>Physcomitrella patens</i>	.	.	.
57	<i>Physcomitrella patens</i>	.	.	.
58	<i>Physcomitrella patens</i>	.	.	.
59	<i>Physcomitrella patens</i>	.	.	.
60	<i>Physcomitrella patens</i>	.	.	.
61	<i>Physcomitrella patens</i>	.	.	.
62	<i>Physcomitrella patens</i>	.	.	.
63	<i>Physcomitrella patens</i>	.	.	.
64	<i>Physcomitrella patens</i>	.	.	.
65	<i>Physcomitrella patens</i>	.	.	.
66	<i>Physcomitrella patens</i>	.	.	.
67	<i>Physcomitrella patens</i>	.	.	.

IMAGE EVALUATION TEST TARGET (QA-3)



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