ASSEMBLING THE CHALLENGING PUZZLE OF ALGAL BIODIVERSITY: SPECIES DELIMITATION WITHIN THE GENUS *ASTEROCHLORIS* (TREBOUXIOPHYCEAE, CHLOROPHYTA)¹

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The genus Asterochloris represents one of the most common, widespread, and diverse taxa of lichen photobionts. In this report, we describe and characterize six new species (A. echinata, A. friedlii, A. gaertneri, A. leprarii, A. lobophora, and A. woessiae) that were identified during our recent investigation of photobiont diversity. We found that the species differed genetically, morphologically, ecologically, and with respect to their mycobiont partners. revealed Statistical analyses significant morphological differentiation of all six newly described species, as well as their separation from previously described Asterochloris species. Chloroplast morphology represented the best morphological marker for species delineation. In fact, each species can be recognized by the dominance and unique assemblage of particular chloroplast types. Although genetically well recognized by rapidly evolving internal transcribed spacer rDNA and actin intron markers, all 13 investigated Asterochloris species shared identical small subunit rDNA sequences. We therefore demonstrated that morphologically and ecologically diverse species can frequently be grouped into a single taxonomic unit in whole-transcriptome sequencing studies, considerably affecting the resulting estimates of species diversity. Finally, we demonstrated the presence of isogamous sexual reproduction in Asterochloris, disputing the current symbiotic dogma of the loss of sexual reproduction in algal symbionts.

Key index words: Asterochloris; cryptic species; green algae; lichens; morphology; phylogeny; speciation; symbiosis; taxonomy

Abbreviations: BI, Bayesian inference; BIC, Bayesian information criterion; CAUP, Culture Collection of Algae of the Charles University in Prague, Czech Republic; CBCs, compensatory base changes; CM, confocal microscopy; GDA, General discriminant analysis; ITS, internal transcribed spacer; MCMC, Markov chain Monte Carlo; ML, maximum likelihood; NGS, next-generation sequencing; OTUs, operational taxonomic units; PCA, principal component analysis; rbcL, ribulose-bisphosphate carboxylase; SAG, Culture Collection of Algae at the University of Göttingen, Germany; UTEX, Culture Collection of Algae at the University of Texas at Austin, USA; wMP, weighted maximum parsimony

Species are fundamental units of biology, comparable to atoms in chemistry or theorems in mathematics. Therefore, the proper delimitation of species is essential for both biologists and the general public. Species delimitation is a fundamental requirement for our understanding of ecosystems and biodiversity, which is necessary for effective decision making about conservation efforts. In addition, taxonomy is a language used by scientists to help the public recognize the diversity, ecology, distribution, and evolutionary history of living organisms. However, evolving over time, species are not unchanging entities. We are therefore surrounded by a plethora of species that vary based on their evolutionary ages, which can make it extremely difficult to perform species identification and delimitation. In fact, the issue of species delimitation has been further complicated by the species problem, that is, the difficulty in defining the concept of species. To date, a wide range of species concepts have been proposed, many of which are associated with several definitions. Moreover, many of these concepts are incompatible in that they can lead to different conclusions concerning the boundaries and number of species (De Queiroz 2007).

In protists, the problem of species delimitation is enhanced by the near absence of morphological features that could be used to clearly distinguish one species from another. As a consequence, a high level of hidden diversity is usually present within nominal, morphologically defined species and genera. Hidden, morphologically highly similar species are frequently described in green algae (Lewis and Flechtner 2004, Fawley et al. 2011, Demchenko

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et al. 2012), chrysophytes (Škaloud et al. 2012, 2014, Jo et al. 2013), diatoms (Mann et al. 2004, Lundholm et al. 2012), ciliates (Quintela-Alonso et al. 2013), and heterotrophic flagellates (Hausmann et al. 2006, Harper et al. 2009). Particularly in green algae, authors often refrain to differentiate the hidden species morphologically, and they define these species according to the phylogenetic species concept (Bock et al. 2011, Krienitz et al. 2011, 2012, Fučíková et al. 2012).

In this study, we focused on species delimitation within the genus Asterochloris Tschermak-Woess, one of the most common lichen photobionts. This genus was described by Tschermak-Woess (1980). who differentiated it from the closely related Trebouxia Puymaly based on chloroplast morphology. Subsequently, molecular investigations revealed the paraphyly of the genus Trebouxia (Friedl and Zeltner 1994, Friedl and Rokitta 1997) and the close relationship of several Trebouxia species with the genus Asterochloris (Helms et al. 2001, Piercey-Normore and DePriest 2001, Skaloud and Peksa 2008). Splitting of the genus Trebouxia, as well as formal delineation of the genus Asterochloris, was proposed by Škaloud and Peksa (2010), who also emphasized a huge amount of hidden diversity within the latter genus. According to current knowledge, the genus Asterochloris represents one of the most common lichen symbionts, occurring in thalli of more than 20 lichen genera worldwide (Piercey-Normore and DePriest 2001, Yahr et al. 2004, 2006, Cordeiro et al. 2005, Nelsen and Gargas 2006, 2008, Beiggi and Piercey-Normore 2007, Bačkor et al. 2010, Škaloud and Peksa 2010, Peksa and Skaloud 2011).

During our recent investigation of Asterochloris photobionts, we identified a number of new lineages occurring in lichen thalli sampled across Europe (Bačkor et al. 2010, Škaloud and Peksa 2010, Peksa and Škaloud 2011). Apart from their delimitation by unique internal transcribed spacer (ITS) rDNA and actin I sequences, the lineages were significantly differentiated by their substrate and climatic preferences (Peksa and Škaloud 2011), suggesting that, in fact, they represent hidden species. However, the virtual identity of all three published Asterochloris SSU rDNA sequences (i.e., A. erici —AB080310, A. magna—Z21552, and A. phycobiontica —GU017647) could call determining these lineages as hidden species into question.

The principal aim of this study was to assess whether genetically differentiated *Asterochloris* lineages could be considered to represent distinct, well-defined species. In particular, we aimed to morphologically differentiate among the 13 lineages, selected to include cultured photobiont strains obtained either from personal or public culture collections. Apart from the seven currently accepted *Asterochloris* species, we investigated six additional lineages representing the putative new species. We performed a detailed morphological investigation of all photobiont strains and investigated whether the lineages could be morphologically delineated. Since species of *Trebouxia* and *Asterochloris* are traditionally differentiated based on their chloroplast morphology, we investigated the chloroplast structure and development of cultivated photobiont cells using modern light and confocal microscopy (CM). Finally, we took advantage of previously published diversity surveys to trace the genetic diversity, ecology, biogeography, and mycobiont specificity of each lineage.

MATERIALS AND METHODS

Origin and cultivation of investigated strains. The majority of strains used in this study were isolated by Škaloud and Peksa (2010) from the lichen thalli sampled in central and eastern Europe (Table S1 in the Supporting Information). The remaining strains were obtained from the Culture Collection of Algae at the University of Texas at Austin, USA (UTEX), the Culture Collection of Algae at the University of Göttingen, Germany—(SAG), and the Culture Collection of Algae and Protozoa at Argyll, United Kingdom—CCAP. The strains were grown on 2% agarized Bold's basal medium as modified by Bischoff and Bold (1963). All cultures were maintained at a temperature of 15° C, under constant illumination of 7–15 µmol photons · m⁻² · s⁻¹ (cooling box Helkama C5G).

Morphological observations and statistical analyses. To obtain a detailed morphological characterization of particular Asterochloris lineages, we investigated the cultivated strains by both conventional light and CM. Light microscopy observations were performed using an Olympus BX51 microscope (Olympus Corp., Tokyo, Japan) equipped with a differential interference contrast. For CM, a Leica TCS SP2 laser scanning confocal microscope (Leica Microsystems, Wetzlar, Germany) equipped with an argon-krypton laser was used. We applied a 488 nm excitation line and an AOBS filter-free system collecting emitted light between 498 and 700 nm. The autofluorescence of chlorophyll was exploited for visualization of the chloroplast structure. A series of optical sections through chloroplasts were captured and used for 3-dimensional reconstruction of their morphology. The chloroplast reconstructions were produced by the Image 1.34p program (Abramoff et al. 2004), using the "Volume viewer" plugin.

Individual strains were regularly observed during the 3month period of culturing, to well characterize the overall morphological variability. Zoospore formation was induced by transferring the cultures to a 1% glucose solution (Hildreth and Ahmadjian 1981). Pyrenoid was visualized by staining with a chloriodine solution (an aqueous solution of 5 g I2 and 10 g of 2,2,2-trichlor-1,1-ethandiol in 5 mL of distilled water). Since some lineages were represented only by a single or two cultured strains, we repeated the morphological investigation of selected strains in a half-year interval, after the inoculation of cells onto the fresh agar plates. During each investigation, the following characters were observed: (i) the average cell width (calculated from at least 45 replicates); (ii) cell shape (a portion of spherical, oval, and pyriform cells); (iii) the maximum number of pyrenoids per cell; (iv) the number of aplanospores per sporangia (16, 32, 48, or 128 spores); (v) chloroplast shape (a portion of following chloroplast types as viewed in CM: shallowly lobed, deeply lobed, crenulate, parietal, echinate, flat lobed, globular); and (vi) chloroplast lobe termination (a portion of following lobe termination types as viewed in CM: elongated, simple, flat, finger like, not formed). Statistical analyses of measured data (principal component and general discriminant analyses)

were performed using Statistica 8.0 (StatSoft, Inc., Tulsa, OK, USA). All graphs were created in R (R Core Team 2014), using the package ggplot2 (Wickham 2009).

DNA extraction, PCR, and sequencing. To well characterize the particular Asterochloris lineages, we investigated the genetic variation at four loci including the slowly evolving SSU rRNA and rbcL genes, and the rapidly evolving ITS rDNA and actin type I intron markers. Most of the analyzed ITS rDNA and actin sequences originated from our previous studies (Škaloud and Peksa 2010, Peksa and Škaloud 2011). However, to gain a better resolution of species relationships, we additionally obtained 12 ITS rDNA and 7 actin sequences from lichen thalli and cultured Asterochloris strains, respectively (Table S1).

Total genomic DNA was isolated following the standard cetyl trimethylammonium bromide protocol (Doyle and Doyle 1987). The amplification of SSU rRNA and rbcL genes was performed as described in Neustupa et al. (2013), using the primers 18S-F (5'-AAC CTG GTT GAT CCT GCC AGT-3') and 18S-R (5'-TGA TCC TTC TGC AGG TTC ACC TAC G-3'; Katana et al. 2001), and primers PRASF1 (5'-ATG GTT CCA CAA ACA GAA AC-3') and PRASR1 (5'-TTG TCA ATA GTA TCA AAT TC-3'; Sherwood et al. 2000). The amplification of ITS rDNA and actin type I locus was performed as described in Peksa and Škaloud (2011), using the primers nr-SSU-1780 (5'-CTG CGG AAG GAT CAT TGA TTC-3'; Piercey-Normore and DePriest 2001) and ITS4 (5'-TCC TCC GCT TAT TGA TAT GC-3'; White et al. 1990), and primers ActinF2 Astero (5'-AGC GCG GGT ACA GCT TCA C-3') and ActinR2 Astero (5'-CAG CAC TTC AGG GCA GCG GAA-3'; Škaloud and Peksa 2010). The PCR products were quantified on a 1% agarose gel stained with ethidium bromide and purified using the JetQuick PCR Purification kit (Genomed). The purified amplification products were sequenced using the PCR primers with an Applied Biosystems (Seoul, Korea) automated sequencer (ABI 3730XL) in Macrogen Corp. (Seoul, Korea).

Sequence analyses. The newly determined sequences were aligned to other sequences from the GenBank database. Three different alignments were constructed for the phylogenetic analyses: (i) an SSU rDNA alignment of 22/38 unique/ total sequences of Trebouxiales, (ii) an *rbcL* alignment of 34/ 39 unique/total sequences of Trebouxiales, and a (iii) concatenated ITS rDNA + actin alignment of 63/79 unique/total Asterochloris sequences selected to encompass all known lineages characterized by both loci. The sequences were aligned using MAFFT v. 6 software (Katoh et al. 2002) under the Q-INS-I strategy and checked for obvious sequencing errors. The alignment of actin sequences was improved by eliminating the ambiguously aligned regions using the program Gblocks v. 0.91b (Castresana 2000). The resulting alignments had lengths of 1772 (SSU rDNA), 1158 (rbcL), and 1142 (ITS rDNA + actin) characters, respectively. All alignments were submitted to TreeBase (http://www.treebase.org/treebaseweb/home.html) and are available under No. S16886.

For each of the alignment partitions, the most appropriate substitution model was estimated using the Bayesian information criterion (BIC) as implemented in jModelTest 2.1.4 (Darriba et al. 2012). This BIC-based model selection procedure selected the following models: (i) TrNef + I + Γ for SSU rDNA, (ii) TIM2 + Γ for the first codon position of the *rbcL* gene, (iii) JC + I for the second codon position of the *rbcL* gene, (iv) TIM3 + I for the third codon position of the *rbcL* gene, (v) TrNef + Γ for ITS1, (vi) K80 + Γ for ITS2, actin exon, and actin intron 248, (vii) JC for 5.8S rDNA, and (viii) HKY + Γ for actin intron 206.

The phylogenetic trees were inferred by Bayesian inference (BI) using MrBayes version 3.2.1 (Ronquist et al. 2012). With the exception of SSU rDNA data, the analyses were carried

out on partitioned data sets using the different substitution models selected by jModelTest 2.1.4. For those models having complicated substitution types, a mixed substitution type was selected to sample across the substitution model space in the Bayesian Markov chain Monte Carlo (MCMC) analysis itself. All parameters were unlinked among partitions. Two parallel MCMC runs were carried out for five million generations, each with one cold and three heated chains. Trees and parameters were sampled every 100 generations. Convergence of the two cold chains was assessed during the run by calculating the average standard deviation of split frequencies (SDSF). The SDSF values of SSU rDNA, *rbcL*, and concatenated ITS rDNA + actin analyses were 0.0008, 0.0035, and 0.0053, respectively. Finally, the burn-in values were determined using the "sump" command.

Bootstrap analyses were performed by maximum-likelihood (ML) and weighted maximum parsimony (wMP) criteria using GARLI, version 2.01 (Zwickl 2006), and PAUP*, version 4.0b10 (Swofford 2002), respectively. ML analyses consisted of rapid heuristic searches (100 pseudoreplicates) using automatic termination (genthreshfortopoterm command set to 100,000). The analyses were performed on partitioned data sets using the different substitution models selected by jModelTest 2.1.4. The wMP bootstrapping (1,000 pseudoreplicates) was performed using heuristic searches with 100 random sequence addition replicates, tree bisection reconnection swapping, random addition of sequences, and gap characters treated as missing data. Character weights were assigned using the rescaled consistency index on a scale of 0 to 1,000. New weights were based on the mean fit values for each character over all trees in the memory.

To show the genetic diversity within the newly characterized lineages, we constructed the haplotype networks on the basis of maximum parsimony analyses of all available sequences, using the Haplotype Viewer (G. Ewing, available at www.cibiv.at/~greg/haploviewer).

RESULTS

Analyses of molecular data. BI of the SSU rDNA and *rbcL* data yielded similar tree topologies, resolving Asterochloris, Trebouxia, and Myrmecia Printz as well-defined, distinct genera (Fig. 1). In the SSU rDNA analysis, a clade of environmental sequences from soil samples (Lesaulnier et al. 2008) was additionally inferred in the affiliation of the genus Asterochloris. Comparison with other SSU rDNA sequences showed that six investigated Asterochloris strains (SAG 26.81, UTEX 911, Bayerová 3401, Peksa 183, Peksa 236, and Peksa 999) contained IB3 group I introns at position 516 relative to the Escherichia coli coding region. The exon SSU rDNA sequences of all investigated Asterochloris strains were completely identical. Resequencing of a single genetically distinct strain (A. magna UTEX 902, accession Z21552) confirmed that all nucleotide differences correspond to sequencing errors (Fig. 1a). The Asterochloris rbcL sequences were slightly different from each other, but analysis of these sequences did not reveal any highly supported clades with the exception of a lineage comprising A. glomerata and A. irregularis strains (Fig. 1b).

Bayesian analysis of the concatenated ITS rDNA and actin data set revealed the existence of more

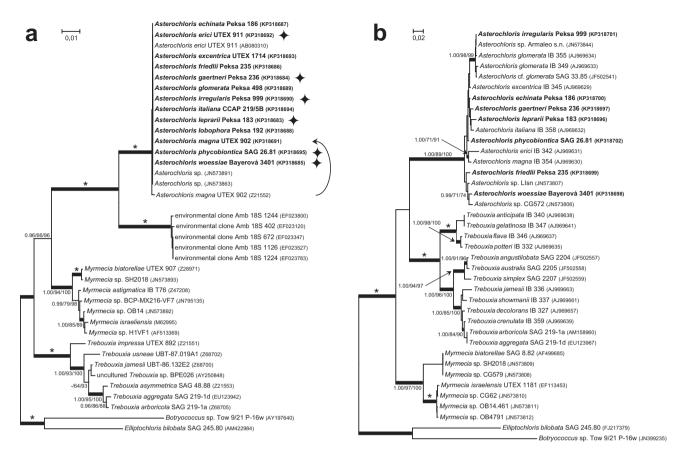


FIG. 1. Phylogeny of the Trebouxiales obtained by Bayesian inference of the SSU rDNA (a) and rbcL (b) data sets. Values at the nodes indicate statistical support estimated by three methods: MrBayes posterior-node probability (left), maximum-likelihood bootstrap (middle), and weighted maximum parsimony bootstrap (right). Full statistical support (1.00/100/100) is marked with an asterisk. Thick branches represent nodes receiving the highest posterior probability support (1.00). Newly sequenced strains are marked in bold. Those sequences containing the IB3 group I introns are marked by stars. An arrow indicated the corrected phylogenetic position of the strain UTEX 902 (*Asterochloris magna*), which SSU rDNA sequence was deposited in GenBank with putative sequencing errors. Scale bar represents the expected number of substitutions per site.

than 20 well-resolved lineages within the genus Asterochloris (Fig. 2). The relationships among the lineages correspond well with the phylogeny presented by Škaloud and Peksa (2010), including the presence of three moderately to well-supported major clades, A, B, and C. All seven formerly described species (A. erici, A. excentrica, A. glomerata, A. irregularis, A. italiana, A. magna, and A. phycobiontica) formed well-recognized, distinct lineages. The Asterochloris cultures that we isolated during our recent investigation of lichen photobionts clustered within the six distinct lineages recognized by Skaloud and Peksa (2010) as lineages 6, 7, 10, 11, 14, and 16. Lineages 6 and 7 (here referred to as A. leprarii sp. nov. and A. gaertneri sp. nov.) were inferred to belong within clade B, together with A. excentrica and three additional lineages. The remaining four lineages (here referred to as A. echinata sp. nov., A. friedlii sp. nov., A. lobophora sp. nov., and A. woessiae sp. nov.) were inferred to be members of clade C, including A. italiana, A. phycobiontica, and five additional lineages. Within clade C, the relationship among the lineages remained unresolved, with the exception of the close, significant relationship between *A. phycobiontica* and *A. lobophora*.

The comparison of all available ITS rDNA sequences pointed to wide differences in genetic variability within the six newly recognized species (Fig. 2b). A. leprarii and A. gaertneri showed almost no intraspecific genetic diversity. Of the 30 investigated A. gaertneri isolates, only one differed by a single nucleotide substitution. On the other hand, A. lobophora represented the most diverse species, containing a total of 18 different ITS rDNA genotypes. A. lobophora also stands out due to its higher occurrence in Cladonia lichens.

Morphological analyses. The diameter of vegetative cells varied, ranging from 4 to 29 μ m. The cells were generally spherical and occasionally oval or pyriform. The cell wall was thin, and occasionally, a flat, localized thickening of the wall was detected in mature cells. In old cultures, the walls of some cells were slightly thickened along their entire surface. A single nucleus with distinct nucleolus was situated parietally in the broad chloroplast infolding.

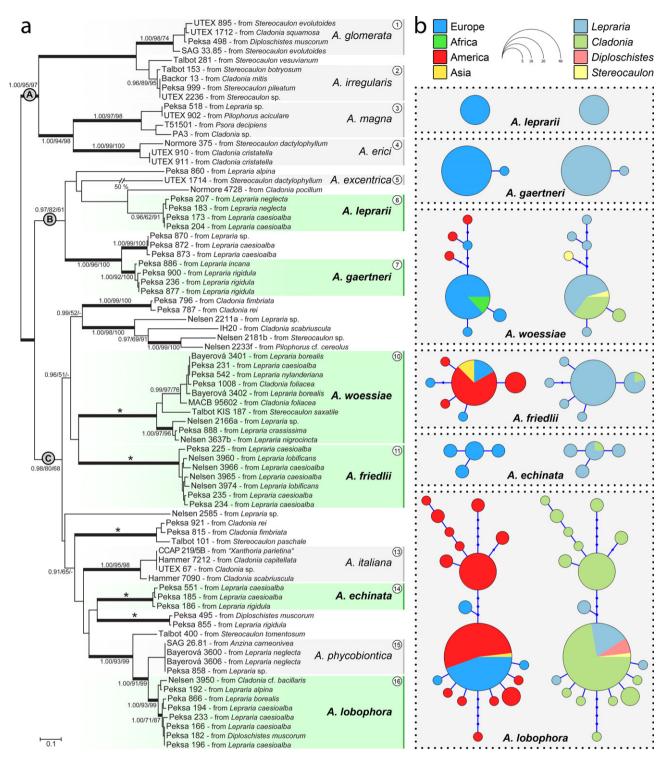


FIG. 2. Genetic diversity within the genus *Asterochloris*. (a) The Bayesian majority rule tree based on the concatenated ITS rDNA + actin alignment. See Figure 1 for the explanation of node values. Clade numbering and affiliation into the three major clades (A-C) follows Škaloud and Peksa (2010). Scale bar represents the expected number of substitutions per site. (b) Statistical parsimony haplotype networks of all available ITS rDNA sequences, showing the intraspecific diversity within the newly proposed species. Genotypes are colored according to the sampling continent (on the left) and respective mycobiont symbiotic partner (on the right). The sizes of circles representing genotypes reflect the number of sequences that share a genotype. Inferred intermediate haplotypes that were either not sampled or are extinct are represented by small noncolored circles.

The majority of the cell volume was occupied by the chloroplast. In young cells, the chloroplast was parietal or ribbon shaped. Soon, it shifted to a central position and began to develop into a massive, lobed form. Mature vegetative cells therefore contained a central axial chloroplast with variously arranged lobes reaching the cell periphery. In the late ontogenetic stages, specifically prior to zoo- or aplanosporogenesis, the chloroplast transformed into the parietal type, with smooth, never lobed margins. After a short time, it began to divide into numerous parts in preparation for asexual reproduction. Taking advantage of laser scanning CM, we recognized seven specific chloroplast types occurring in mature Asterochloris cells (Fig. 3), as follows: (i) a *deeply lobed* type, characterized by long, branched, or unbranched lobes emerging directly from the thin chloroplast layer spreading around the pyrenoid ("Tieflappig Typ" sensu, Gärtner 1985a); (ii) a shallowly lobed type, which is similar to the previous type but differs in that the chloroplast

lobes emerge from the central mass of the chloroplast layer encircling the pyrenoid ("Normaltyp" sensu, Gärtner 1985a); (iii) a *crenulate* type, distinguished by a central, massive chloroplast with a regularly nodulated surface ("Crenulater Typ" sensu, Gärtner 1985b); (iv) a *parietal* type, characterized by parietally positioned nodulated chloroplast with the margins extended into divided finger-like lobes; (v) a *flat lobed* type, representing an axial chloroplast with long lobes that appear flattened over their entire length; (vi) an *echinate* type, characterized by numerous thin radial lobes emerging uniformly from the central mass of the chloroplast layer; and (vii) a *globular* type, a simple spherical chloroplast without, or with very shallow, lobes.

In addition to these seven morphological chloroplast types, we distinguished four types of lobe terminations (Fig. 3), as follows: (i) an *elongated* type, with lobes extending longitudinally at their ends, therefore giving the chloroplast a finny appearance in surface view ("Rippenförmig Typ" sensu, Gärtner

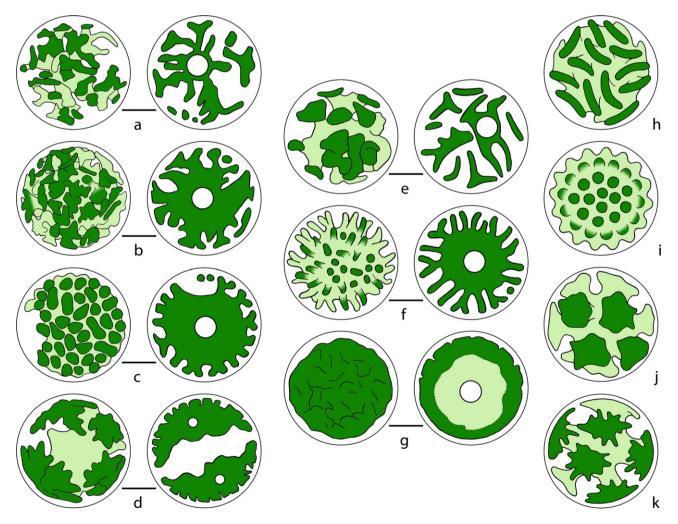


FIG. 3. Schematic drawings of particular chloroplast and lobe termination types in *Asterochloris*. (a–g) Chloroplast types (left: surface view; right: view in optical section): (a) deeply lobed, (b) shallowly lobed, (c) crenulate, (d) parietal, (e) flat lobed, (f) echinate, (g) globular. (h–k) Lobe termination types: (h) elongated, (i) simple, (j) flat, (k) finger-like.

1985a); (ii) a *simple* type, characterized by simply terminated lobes at their ends; (iii) a *flat* type of chloroplast lobe, terminated by irregular plates, perpendicularly oriented with respect to the lobe axis; and (iv) a *finger-like* type, distinguished by lobes branched into several finger-like projections. These projections were perpendicularly oriented with respect to the lobe axis, spreading below the plasma membrane.

Pyrenoids were present in all Asterochloris species except A. magna. While A. erici and A. lobophora had single pyrenoids, the cells usually contained one to several pyrenoids lying in the chloroplast center. Often, one large centrally located pyrenoid was surrounded by several smaller satellite pyrenoids, which were created by budding. The pyrenoids were generally distinct; only in A. erici did the pyrenoid gradually change over to a chloroplast matrix without a distinct pyrenoid margin. Various structures could occasionally be observed in the pyrenoid matrix by both conventional light and CM. The pyrenoids were granulated, striated, or perforated. In A. phycobiontica, the pyrenoid contained distinct rings of puzzling origin. The frequency and markedness of subpyrenoidal structures significantly increased with cell age. Pyrenoids were usually surrounded by a conspicuous starch sheet, which could be visualized by staining with chloriodine solution.

To investigate morphological differences between the 13 recognized Asterochloris species in detail, we characterized each species based on a number of features, including cell shape, dimensions, and chloroplast morphology (Table S2 in the Supporting Information). We found no significant differences in average cell dimensions among species; however, some species could be distinguished by the prevailing shape of their cells. The closely related A. glomerata and A. irregularis frequently produced oval and pyriform cells (Fig. 4a); by contrast, these cells were never observed in A. echinata. In addition, all investigated species were obviously heterogeneous in the overall morphological complexity of chloroplast types (Fig. 4b). Two to four chloroplast types were usually observed during cell ontogeny. The shallowly lobed chloroplast was the most common type, occurring in 11 of 13 investigated Asterochloris species. Four species (A. glomerata, A. irregularis, A. woessiae, and A. excentrica) were characterized by the prevailing occurrence of the deeply lobed chloroplast type, while the flat lobed type was only observed in the first three species. By contrast, deeply lobed chloroplasts were never found in A. phycobiontica or A. lobophora, which were characterized by the presence of crenulate and finger-like types. Two species, A. magna and A. echinata, could be easily recognized by their specific chloroplast morphology, the former by the presence of the simplest globular type and the latter by the combination of crenulate and echinate types. Finally, A. italiana was exceptional because it produced a single, shallowly lobed chloroplast type.

Morphological differences among the species were also observed in the shape of chloroplast lobe terminations (Fig. 4c). Elongated terminations were most commonly produced, which were observed in all species except *A. magna* and *A. echinata*. These two species could clearly be recognized by the common absence of any lobes (*A. magna*) and by the exclusive formation of single lobe terminations (*A. echinata*). Moreover, the prevalence of finger-like node terminations is characteristic of *A. phycobiontica*.

Principal component analysis (PCA) of the entire data set resulted in a relatively well-defined grouping of investigated strains belonging to particular species (Fig. 4d). For example, A. erici and A. magna were plotted in two distinct clusters in the upper left corner of the PCA plot. On the other hand, several strains belonging to different species were intermixed with each other (e.g., A. excentrica and A. woessiae). Interestingly, the closely related species pairs (A. glomerata-A. irregularis and A. phycobiontica-A. lobophora) were obviously similar based on their morphology. General discriminant analysis (GDA) yielded much better grouping of species into separate clusters (Fig. 4, e and f). A. magna formed a strong outlying cluster with a negative value on the second GDA axis (Fig. 4e). A scatter plot based on the first and third GDA axes showed the separation of all 13 Asterochloris species into distinct clusters (Fig. 4f). Discriminant analysis (DA) indicated strongly significant differentiation among all investigated species (Wilk's $\lambda < 0.00001$; P < 0.00001). Forward stepwise analysis selected the globular chloroplast, parietal chloroplast with finger-like lobes, the number of autospores, and the number of pyrenoids as the best discriminating characters. The first and third GDA axes, which well discriminated all investigated species, were highly correlated with the four factors examined. Whereas the first GDA axis was correlated with parietal and flat chloroplast types with elongated lobes (correlation coefficients 0.34 and 0.38, respectively), the third axis was correlated with deeply lobed chloroplasts with elongated lobes and globular chloroplasts (correlation coefficients 0.49 and 0.41, respectively). The globular chloroplast type was also highly correlated with the second GDA axis (correlation coefficient -0.70). The average correct discrimination of individual strains based on their morphology reached 94.3%; that is, only three investigated strains (one A. excentrica and two A. woessiae strains) were classified incorrectly by the discriminant model.

Reproduction. The life cycle and reproductive processes are schematically delineated in Figure 5. Asexual reproduction occurred by the formation of aplanospores, zoospores, and autospores. Autospore production was relatively rare; it was observed only in some species. Autospore production was initiated by slight cell enlargement and subsequent chloroplast division (Fig. 5b). In general, autospores were formed in relatively small

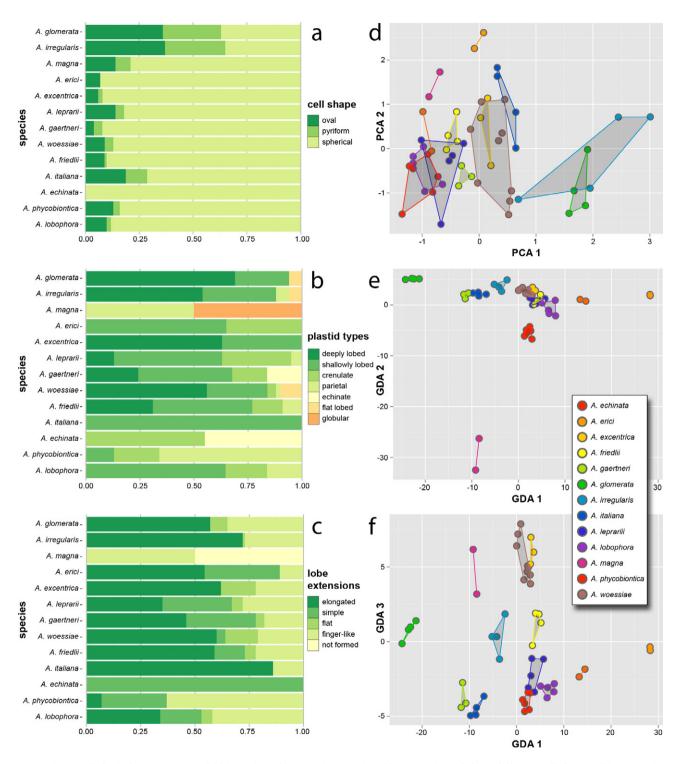


FIG. 4. Morphological comparisons of 13 investigated *Asterochloris* species. (a) Proportion of three different cell shapes. (b) Proportion of seven recognized plastid types. (c) Proportion of five distinguished lobe termination types. (d) Principal component analysis (PCA) of the entire measured morphological features data set. (e, f) General discriminant analysis (GDA) the same data set: (e) the two-dimensional plot of the 1st and 2nd GDA axes, (f) the two-dimensional plot of the 1st and 3rd GDA axes.

numbers (mostly four or eight) and were liberated by either decomposition or rupturing of the mother cell wall, without producing any special openings (Fig. 5d). The formation of aplanospores and zoospores was much more frequent. Prior to the first cleavage, the chloroplast flattened, migrated to the parietal position, and started to divide (Fig. 5f). Ultimately, a large number of daughter cells (usually 64 or 128) were present within the sporangium (Fig. 5h). Once all the daughter cells formed, a distinction between aplanosporangia and zoosporangia was observable. In the case of aplanospores, the daughter cells rounded and produced a cell wall of their own. Mature aplanospores were liberated by rupturing of the mother cell wall (Fig. 5i). Mature zoosporangia were discernible by the irregular shape of their daughter cells (Fig. 5k). The zoospores were all liberated simultaneously with the rupturing of the mother cell wall, enclosed in a gelatinous vesicle that could slightly evaginate from the mother cell wall during the liberation process. The zoospores were dorsiventrally flattened, with two equal anterior flagella, a posterior chloroplast, a medianto-posterior nucleus, and indistinct stigma. Following liberation from the sporangium, the zoospores swam in a packet, the remaining joined together by their posterior extensions (Fig. 51). Shortly thereafter, single zoospores started to detach from the packet and swam separately. Despite this, the majority of zoospores still bore posterior extensions even if observed several minutes after separation (Fig. 5n).

Sexual reproduction was very scarce and was observed only twice in *A. woessiae* (strain CAUP H1009). Biflagellate isogamous gametes were morphologically indistinguishable from zoospores. After their fusion, they gave rise to large quadriflagellate

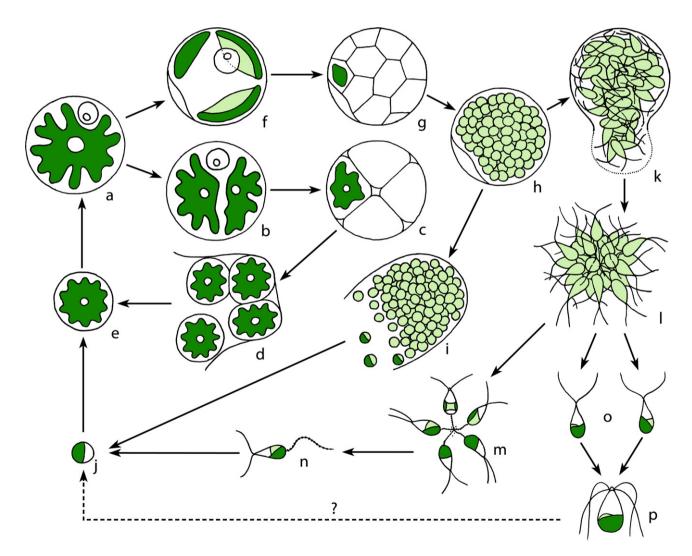


FIG. 5. Schematic representation of the life cycle and reproduction processes in *Asterochloris*. (a) Vegetative cell. (b–e). Autosporogenesis: (b) chloroplast division, (c) young autosporangium, (d) release of autospores, (e) mature autospore. (f–n). Aplano- and zoosporogenesis: (f) chloroplast flattening, note a local thickening of the cell wall, (g) young aplano/zoosporangium, (h) mature aplano/ zoosporangium, (i) release of aplanospores, (j) young aplanospore, (k) release of zoospores, note evagination of gelatinous vesicle, (l) zoospore packet, (m) releasing of zoospores from the packet, (n) single zoospore with a posterior extension. (o, p). Sexual reproduction: (o) gametes, (p) planozygote.

planozygotes (Fig. 5p). The germination of planozygotes was not observed.

TAXA DESCRIPTIONS

In this study, we revealed a clear morphological differentiation of the studied *Asterochloris* lineages. Given the genetic distinctiveness, previously published ecological diversification, and mycobiont specificity of these lineages, we are proposing that they represent new species. Descriptions and characteristics of these new taxa are provided below.

Asterochloris Tschermak-Woess; Pl. Syst. Evol. 135, pp. 291, 292 emend. Skaloud et Peksa

Type species: Asterochloris phycobiontica Tschermak-Woess 1980; Pl. Syst. Evol. 135, p. 292

Emended description: Cells spherical, occasionally oval or pyriform. Cell wall thin, occasionally with a flat local thickening. Single nucleus situated parietally in the broad chloroplast infolding. Chloroplast single, asteroid, with variously arranged lobes reaching the cell periphery. One to several pyrenoids usually lie in the chloroplast center, surrounded by a conspicuous starch sheet. Prior to aplano- and zoosporogenesis, the chloroplast flattens and assumes a parietal position. Asexual reproduction by usually 64-128 aplanospores and zoospores, occasionally by 2-8 autospores. Zoospores naked, dorsiventrally flattened, with two apical flagella, a posterior chloroplast, a median-to-posterior nucleus, and indistinct stigma. Following liberation from the sporangium, the zoospores shortly swam in a packet, joined together by their posterior extensions. Sexual reproduction scarce, by the fusion of two isogamous gametes. Photobionts of many lichens (genera Anzina, Cladia, Cladonia, Diploschistes, Lepraria, Pilophorus, Pycnothelia, Stereocaulon, etc.). Widely distributed, cosmopolitan. From the morphologically similar genus Trebouxia, it generally differs by the presence of deeply lobed chloroplast, parietal position of chloroplast prior to zoo- or aplanosporogenesis, and the production of aplanospores as a prevailing type of asexual reproduction.

Asterochloris leprarii Škaloud et Peksa sp. nov. (Fig. 6, a–k)

Vegetative cells usually spherical, occasionally oval and pyriform, (5-)7.5-24(-28) µm in diameter (Fig. 6, a–c). Cell wall thin, seldom a flat local thickening of the cell wall can be distinguished. Very rarely, the cell wall is slightly thickened along its entire surface. Chloroplast in young cells assumes the central position with several lobes spreading toward the cell's periphery. Mature cells exhibit central chloroplasts of either shallowly lobed (Fig. 6d) or crenulate form (Fig. 6e). Rarely, the deeply lobed (Fig. 6f) and parietal chloroplast (Fig. 6g) is observed as well. The chloroplast lobes can be simply terminated (Fig. 6e), elongated at their ends (Fig. 6f), or finger like (Fig. 6g). Occasionally, the lobe ends are flat (Fig. 6h). The chloroplast contains from one to many pyrenoids. Besides the typical, centrally located pyrenoid, up to seven smaller ones may be present in its vicinity (Fig. 6i). Sometimes, an indistinct granulation or striation can be visible inside pyrenoids. Starch grains are either embedded in a layer around the pyrenoid or distributed evenly throughout the chloroplast. Asexual reproduction by 64–128 aplanospores or 64 zoospores produced in spherical or ellipsoidal sporangia (Fig. 6j). Occasionally, 2–4 autospores are also produced. Zoospores dorsiventrally flattened, drop shaped, arcuate in lateral view, 6–10 μ m long and 2.8–4 μ m wide, with posterior extensions (Fig. 6k).

Holotype: Cryopreserved photobiont cells isolated from the specimen Peksa 183, deposited in the Culture Collection of Algae of the Charles University in Prague (CAUP) as the item TYPE-H 1010.

Reference strains: CAUP H 1010, SAG 2280.

Type locality: Phycobiont of *Lepraria neglecta*, collected on siliceous rock, Rybárna, Šumava Mts, Czech Republic, May 23, 2005. The lichen specimen is deposited in herbarium of O. Peksa in PL (Collection of The West Bohemian Museum in Pilsen), No. 183.

Etymology: The species is named in reference to the mycobiont genus *Lepraria* Ach.

Distribution: So far known only from Europe: Czech Republic, Norway, Slovakia (Nelsen and Gargas 2008, Škaloud and Peksa 2010, Peksa and Škaloud 2011).

Ecology: In temperate Europe, it prefers altitudes of about 800–1,000 m a.s.l; associates with ombrophilic lichens growing on acidic substrates, especially siliceous rocks (Peksa and Škaloud 2011).

Specificity: found exclusively in the thalli of lichen genus Lepraria.

Asterochloris gaertneri Škaloud et Peksa sp. nov. (Fig. 6, l-t)

Vegetative cells spherical, very rarely oval or pyriform, 5.5-26(-29.5) µm in diameter (Fig. 6, k and 1). Cell wall thin, seldom a flat local thickening of the cell wall can be observed. Very rarely, the cell wall is slightly thickened along its entire surface. Chloroplast in young cells assumes the central position with several lobes spreading toward the cell's periphery. Mature cells often possess shallowly lobed axial chloroplasts (Fig. 6, m and n). Deeply lobed (Fig. 60), crenulate (Fig. 6p), and echinate (Fig. 6, q and r) chloroplast forms observed, as well. The chloroplast lobes are simply terminated, extended longitudinally at their ends, or terminated by finger-like extensions. Flat lobes produced very rarely. The chloroplast contains from one to many distinct pyrenoids. If many, they usually jointly occupy the chloroplast's center (Fig. 6s). Sometimes, an indistinct striation can be visible inside pyrenoids. Starch grains are embedded in a layer around the pyrenoid. Asexual reproduction by 64-128-256 aplanospores or 128 zoospores produced in large spherical or ellipsoidal sporangia

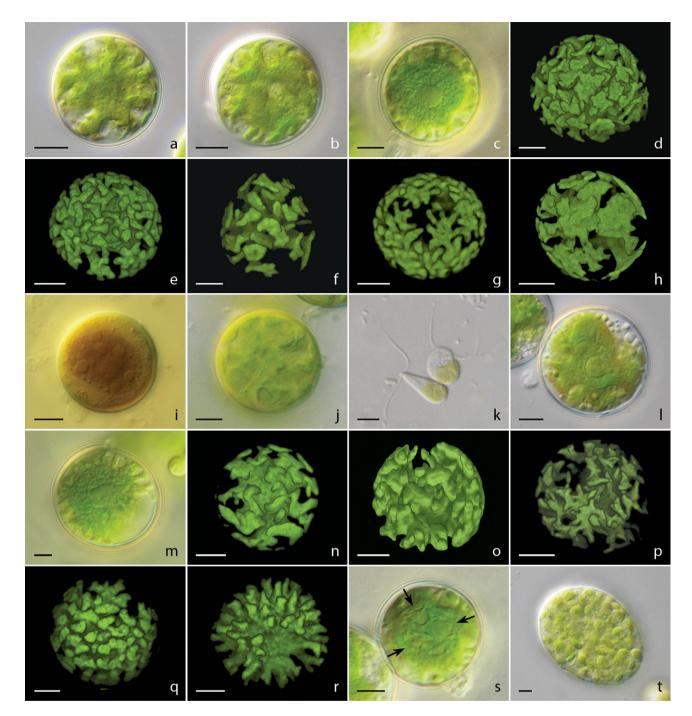


FIG. 6. Light micrographs and confocal reconstructions of chloroplast structures in *Asterochloris leprarii* and *A. gaertneri*. (a–k) *A. leprarii*. Light micrographs of mature vegetative cells possessing deeply lobed (a), shallowly lobed (b), and crenulate (c) chloroplasts. Confocal reconstructions of shallowly lobed (d), crenulate (e), deeply lobed (f), and parietal (g) chloroplast types. (h) Deeply lobed chloroplast with flat lobe ends. Several pyrenoids occur around the large central pyrenoid (i); cells stained by chloriodine solution. (j) Young aplanosporangium. (k) Zoospores. (l–t) *A. gaertneri*. Light microscopy of shallowly lobed (l) and crenulate (m) chloroplast. Confocal reconstructions of shallowly lobed (n, o), deeply lobed (p), crenulate (q), and echinate (r) chloroplasts. Several pyrenoids of equal size (arrows) occur in the chloroplast's center (s). (t) Mature aplanosporangium; scale bar—5 µm.

(Fig. 6t). Occasionally, 4–8 autospores are also produced. Zoospores dorsiventrally flattened, 6–7.5 μm long and 2.5–4 μm wide.

Holotype: Cryopreserved photobiont cells isolated from the specimen Peksa 236, deposited in the CAUP as the item TYPE-H 1013.

Reference strains: CAUP H 1013, SAG 2283.

Type locality: Phycobiont of *Lepraria rigidula*, collected on bark of *Acer pseudoplatanus*, near Stříbrnická Mt., Králický Sněžník Mts, Czech Republic, February 10, 2005. The lichen specimen is deposited in herbarium of O. Peksa in PL, No. 236.

Etymology: The species epithet is in honor of the work of Dr. Georg Gärtner, who published several reports on *Trebouxia* s.l.

Distribution: So far known only from Europe: Czech Republic, Germany, Slovakia (Škaloud and Peksa 2010).

Ecology: Associates with lichens growing on tree bark and siliceous rocks, in rain-sheltered situations (ombrophobic); in temperate Europe, it was found at altitudes of about 500–900 m a.s.l (Peksa and Škaloud 2011).

Specificity: found exclusively in the thalli of lichen genus Lepraria.

Asterochloris woessiae Škaloud et Peksa sp. nov. (Fig. 7, a-k)

Vegetative cells usually spherical, rarely oval and pyriform, (5-)6.5-19(-25.5)µm in diameter (Fig. 7a). Cell wall thin, seldom a flat local thickening of the cell wall can be made out (Fig. 7b). Very rarely, the cell wall is slightly thickened along its entire surface. Chloroplast in young cells assumes the central position with several lobes spreading toward the cell's periphery (Fig. 7c). Mature cells display a structurally complicated, central, deeply lobed chloroplast with branched lobes that emerge directly from the thin layer spreading around the pyrenoid (Fig. 7d). Mature cell chloroplasts can further exhibit several other ontogenetic stages, alternating during the cell's ontogeny. The chloroplast can be shallowly lobed (Fig. 7c), or appearing flattened over their entire length (Fig. 7e). Rarely, the crenulate (Fig. 7f) and parietal chloroplasts can be formed as well. The chloroplast lobes can be terminated by all four known types, usually by elongated, flat (Fig. 7g), or finger-like extensions. The chloroplast contains 1-3 distinctively delimited pyrenoids (Fig. 7h). Sometimes, an indistinct granulation can be visible inside the pyrenoids. Starch grains are embedded in a layer around the pyrenoid. Asexual reproduction by 128 aplanospores or zoospores produced in large spherical or ellipsoidal sporangia (Fig. 7i). Occasionally, 2-4 autospores are also produced. Zoospores dorsiventrally flattened, 4.5-7.5 µm long and 2.5-4.5 µm wide, with posterior extensions (Fig. 7j). Sexual reproduction by fusion of biflagellate isogamous gametes; planozygotes with four longitudinal flagella (Fig. 7k).

Holotype: Cryopreserved photobiont cells isolated from the specimen Bayerová 3401, deposited in the CAUP as the item TYPE-H 1009.

Reference strains: CAUP H 1009, SAG 2279.

Type locality: Phycobiont of *Lepraria borealis*, collected on sunlit slate rock, Stara planina Mts, Central Balkan National Park, Bulgaria, July 1, 2004. The lichen specimen is deposited in collection of Š. Bayerová-Slavíková in PRA (Herbarium of Institute of Botany of the ASCR, Czech Republic), No. 3401.

Etymology: The species epithet is in honor of the work of Dr. Elisabeth Tschermak-Woess, who described the genus *Asterochloris*.

Distribution: Cosmopolitan, widely distributed. Europe: Bulgaria, Czech Republic, Great Britain, Slovakia, Spain, Sweden; America: Costa Rica, USA; Africa: Canary Islands (Nelsen and Gargas 2006, Bačkor et al. 2010, Škaloud and Peksa 2010, Peksa and Škaloud 2011, Pino-Bodas et al. 2010).

Ecology: Prefers lichens growing on moderately basic substrates (shale, basalt, serpentine rocks etc.) and low altitudes of about 300–600 m a.s.l of temperate Europe (Peksa and Škaloud 2011).

Specificity: Found in a number of lichen species belonging to genera *Lepraria*, *Cladonia*, and *Stereoca-ulon*.

Asterochloris friedlii Škaloud et Peksa sp. nov. (Fig. 7, l-t)

Vegetative cells spherical or slightly oval, (4.5-)6-18(-21) µm in diameter (Fig. 7, 1 and m). Cell wall thin, seldom a flat local thickening of the cell wall can be detected. Chloroplast in young cells assumes the central position with several simple lobes spreading toward the cell's periphery (Fig. 7n). Mature cells generally display a structurally complicated, central, deeply or shallowly lobed chloroplast with branched lobes (Fig. 7, o and p). Sometimes, the crenulate (Fig. 7q) or parietal chloroplast (Fig. 7r) can be formed as well. In mature cells, the chloroplast can be slightly asymmetrically positioned. The chloroplast lobes are mostly extended longitudinally at their ends, but they can be terminated by additional three types, as well. The chloroplast generally contains a single distinct, granulated pyrenoid. Especially in older cells, the pyrenoid buds at its surface and gives rise to several smaller ones in its vicinity (Fig. 7s). Starch grains are embedded in a layer around the pyrenoid in the form of large granules. Asexual reproduction by 64-128 aplanospores or zoospores produced in large spherical, ellipsoidal, or irregular sporangia (Fig. 7t). Zoospores dorsiventrally flattened, 4.5-7 µm long and 3-3.5 µm wide, with posterior extensions.

Holotype: Cryopreserved photobiont cells isolated from the specimen Peksa 235, deposited in the CAUP as the item TYPE-H 1011.

Reference strains: CAUP H 1011, SAG 2281.

Type locality: Phycobiont of *Lepraria caesioalba*, collected on bryophytes on siliceous rock, Klenovský Vepor Mt., Slovenské Rudohorie Mts, Slovakia, July 12, 2004. The lichen thallus is deposited in herbarium of O. Peksa in PL, No. 235.

Etymology: The species epithet is in honor of the work of Dr. Thomas Friedl, who published several reports on *Trebouxia* s.l.

Distribution: Cosmopolitan, widely distributed. Europe: Czech Republic, Romania, Slovakia; America: Canada, USA; Asia: China (Nelsen and Gargas 2006, 2008, Škaloud and Peksa 2010, Peksa and Škaloud 2011).

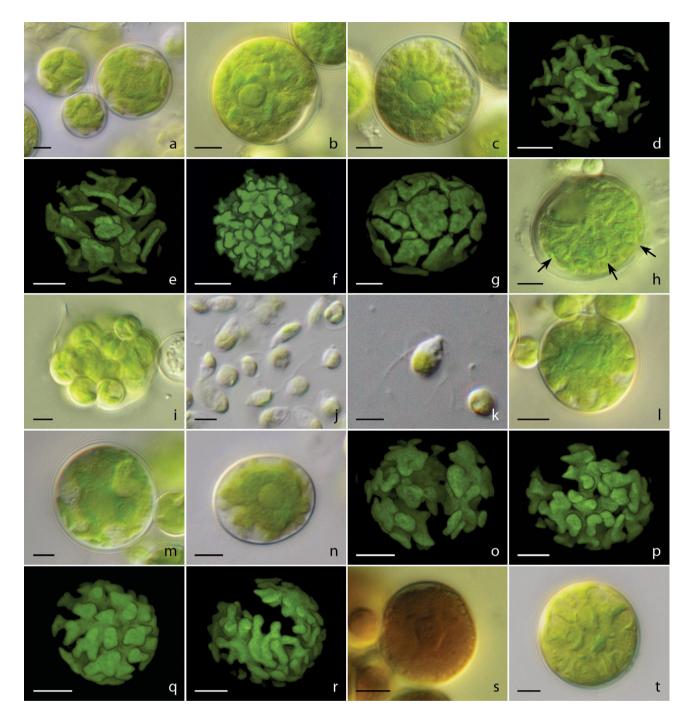


FIG. 7. Light micrographs and confocal reconstructions of chloroplast structures in *Asterochloris woessiae* and *A. friedlii*. (a–k) *A. woessiae*. Light micrographs of young (a) and mature vegetative cells (b, c). Confocal reconstructions of deeply lobed (d), flat lobed (e), and crenulate (f) chloroplasts. (g) Flat terminations of chloroplast lobes. (h) Several pyrenoids (arrows) are formed within the chloroplast. (i) Aplanosporangium. (j) Zoospores. (k) Planozygote with four flagella. (l–t) *A. woessiae*. Light microscopy of shallowly lobed (l) and deeply lobed (m) vegetative cells. (n) Young vegetative cell with simple asteroid chloroplast. Confocal reconstructions of deeply lobed (o), shallowly lobed (p), crenulate (q), and parietal (r) chloroplasts. (s) Single budding pyrenoid. (t) Young aplanosporangium. Cells in Figure (s) stained by chloriodine solution; scale bar—5 µm.

Ecology: Mainly in ombrophobic lichens growing on acidic as well as basic substrates (Peksa and Škaloud 2011).

Specificity: With a single exception (*Cladonia*) found in thalli of lichen genus *Lepraria*.

Asterochloris echinata Škaloud et Peksa sp. nov. (Fig. 8, a–i)

Vegetative cells always spherical, $(5-)7-18(-21) \mu m$ in diameter (Fig. 8a). Cell wall thin, without any local thickenings. Young and mature cells exhibit

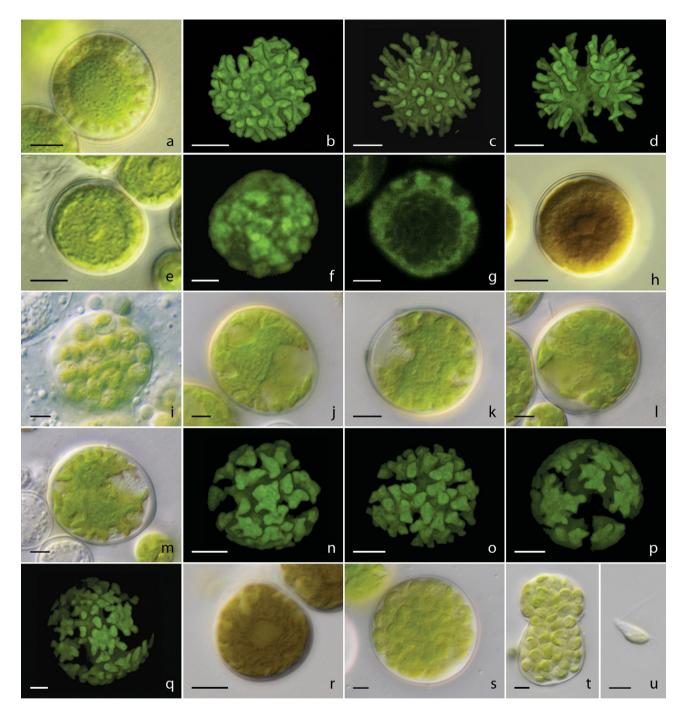


FIG. 8. Light micrographs and confocal reconstructions of chloroplast structures in *Asterochloris echinata* and *A. lobophora.* (a–i) *A. echinata*. (a) Light micrograph of mature vegetative cell. Confocal reconstructions of crenulate (b) and echinate (c, d) chloroplast. Light micrograph (e) and confocal reconstruction (f) of globular chloroplast. (g) Confocal section through the globular chloroplast. (h) Several pyrenoids occur around the large central pyrenoid. (i) Aplanosporangium. (j–u) *A. lobophora.* Light micrographs of deeply lobed (j) and shallowly lobed (k) chloroplasts. (l) Vegetative cell with a flat local thickening of the cell wall. (m) Mature vegetative cell with the thick-end cell wall. Confocal reconstructions of shallowly lobed (n), crenulate (o), and parietal (p, q) chloroplasts. A single pyrenoid is situated in the chloroplast lumen (r). (s) Mature aplanosporangium. (t) Liberated aplanospores. (u) Zoospore. Cells in Figure (h, r) stained by chloriodine solution; scale bar—5 µm.

central crenulate chloroplasts (Fig. 8b). In mature cells, the chloroplast often transform into the echinate form characterized by many thin radial lobes giving it a bristly appearance (Fig. 8, c and d). The crenulate chloroplast of old cells frequently transform into a highly specific, simple form without any lobes (Fig. 8, e and f). This form has a distinctive chloroplast ultrastructure in its central and marginal regions. In the center, the starch accumulation causes the decrease in thylakoid numbers, and subsequent modification of the chloroplast's texture (Fig. 8g). The chloroplast contains from one to many distinct or indistinct pyrenoids. In the latter case, up to eight smaller pyrenoids are present in the vicinity of the central one (Fig. 8h). Sometimes, an indistinct granulation can be visible inside pyrenoids. Starch grains are embedded evenly throughout the chloroplast. Asexual reproduction by 64-128 aplanospores produced in spherical or ellipsoidal sporangia (Fig. 8i). Zoospores very rare, dorsiventrally flattened, 6 μ m long and 4 μ m wide.

Holotype: Cryopreserved photobiont cells isolated from the specimen Peksa 186, deposited in the CAUP as the item TYPE-H 1012.

Reference strains: CAUP H 1012, SAG 2282.

Type locality: Phycobiont of *Lepraria rigidula*, collected on bryophytes on basalt rock, Klíč Mt., Lužické hory Mts, Czech Republic, September 18, 2004. The lichen specimen is deposited in herbarium of O. Peksa in PL, No. 186.

Etymology: The species epithet is named in reference to the echinate shape of chloroplast, which appeared in certain ontogenetic stages.

Distribution: So far known only from Europe: Bulgaria, Czech Republic, Portugal, Slovakia, Spain (Škaloud and Peksa 2010, Peksa and Škaloud 2011).

Ecology: Associates with ombrophilic lichens growing on acidic substrates (Peksa and Škaloud 2011).

Specificity: Found in the thalli of lichen genera Lepraria and Cladonia.

Asterochloris lobophora Škaloud et Peksa sp. nov. (Fig. 8, j–u)

Vegetative cells usually spherical, occasionally oval and pyriform, 6-23(-25.5) µm in diameter (Fig. 8, j and k). Cell wall thin, seldom a flat local thickening of the cell wall can be distinguished (Fig. 8l). Very rarely, the cell wall is slightly thickened along its entire surface (Fig. 8m). Chloroplast in young cells assumes the central position with several lobes spreading toward the cell's periphery. Mature cells exhibit shallowly lobed chloroplast (Fig. 8n), sometimes transformed into the crenulate (Fig. 8o) or parietal form (Fig. 8, p and q). Even though the chloroplast can sometimes appear to be deeply lobed, the lobes never emerge directly from the pyrenoid surroundings. The chloroplast lobes can be terminated by all four known types, usually by finger-like or elongated extensions. The chloroplast contains single distinct granulated pyrenoid (Fig. 8r). Particularly prior to cell division, the pyrenoid sometimes divides into two parts. Starch grains are embedded either in a layer around the pyrenoid or evenly throughout the chloroplast. Asexual reproduction by 64-128 aplanospores or 128-256 zoospores produced in large spherical or slightly ellipsoidal sporangia (Fig. 8, s and t). Zoospores 4–7 μm dorsiventrally flattened, long and $2.5-3.5 \mu m$ wide, with posterior extensions (Fig. 8u).

Holotype: Cryopreserved photobiont cells isolated from the specimen Peksa 166, deposited in the CAUP as the item TYPE-H 1014.

Reference strain: CAUP H 1014.

Type locality: Phycobiont of *Lepraria caesioalba*, collected on siliceous rock, Kašperk Mt., Šumava Mts, Czech Republic, March 31, 2005. The lichen specimen is deposited in herbarium of O. Peksa in PL, No. 166.

Etymology: The species epithet is named in reference to the lobed chloroplast shape.

Distribution: Cosmopolitan, widely distributed. Europe: Czech Republic, Slovakia; America: Canada, USA; Asia: India (Piercey-Normore and DePriest 2001, Yahr et al. 2004, 2006, Cordeiro et al. 2005, Nelsen and Gargas 2006, Beiggi and Piercey-Normore 2007, Bačkor et al. 2010, Kotelko and Piercey-Normore 2010, Škaloud and Peksa 2010, Peksa and Škaloud 2011, Řídká et al. 2014).

Ecology: In temperate Europe, it prefers altitudes of about 500–1,000 m a.s.l; associates with ombrophilic lichens growing mainly on acidic substrates (Peksa and Škaloud 2011).

Specificity: Found in a number of lichen species belonging to genera *Cladonia*, *Lepraria*, *Diploschistes*, and *Stereocaulon*.

DISCUSSION

Species delineation in Asterochloris. To date, a total of seven species are recognized within the genus Asterochloris. These species, established during the second half of the 20th century, were exclusively delimited based on morphological features such as cell size, cell shape, chloroplast morphology, cell wall thickness, and dissociation of aplanospores (Ahmadjian 1960, Archibald 1975, Tschermak-Woess 1980, Hildreth and Ahmadjian 1981). Each species formed a distinct, well-supported lineage in the concatenated ITS rDNA + actin phylogenetic tree (Fig. 2); however, our phylogenetic reconstruction points out the existence of several additional, wellsupported lineages within the genus. Six of these new lineages were investigated in detail in this study. In general, we can apply two alternative taxonomic approaches to assess the observed genetic diversity. First, we can consider each genetic lineage to be a separate, distinct species and based on this approach, the genus Asterochloris would encompass tens, if not hundreds, of undescribed species. Second, genetic diversity can be considered to be a manifestation of substantial infraspecific variability. Applying this "large species" concept would involve merging all described species into a single species, namely A. phycobiontica.

The existence of a single, genetically divergent species is supported by the presence of identical SSU rDNA sequences in all of the investigated strains and by the observation that the relatively low genetic variation in the ITS rDNA region is correlated with the absence of compensatory base changes (CBCs) among a number of Asterochloris lineages (Skaloud and Peksa 2010). In prokaryotes, it was proposed that species' boundaries should be defined by a fixed threshold of genetic divergence in SSU rDNA (Rosselló-Mora and Amann 2001). This concept considers that all strains showing a similarity higher than 97% belong to the same species. Applying this concept to Asterochloris would result in recognizing only a single species; however, this species concept was never applied to eukaryotic organisms, and its applicability to prokaryotes has been seriously criticized (Pedrós-Alió 2006, Stackebrandt and Ebers 2006). Similarly, the presence of CBCs in the conserved regions of the ITS2 molecule has been proposed to represent a threshold for defining species boundaries in eukaryotes (Coleman 2000); however, this concept has been subjected to mounting criticism (Caisová et al. 2011, 2013, Assunção et al. 2012). In particular, it was demonstrated that the CBCs are not diagnostic at the species level and that even genera, families, and orders of green algae can lack CBCs in such regions (Caisová et al. 2011, Škaloud and Rindi 2013). Indeed, a causal link between ITS2 secondary structure and speciation mechanisms in eukaryotes simply does not exist. Therefore, the presence and number of CBCs are most probably direct consequences of the accumulation of mutations during the evolutionary process, simply reflecting the genetic distance among organisms.

In this study and during our previous investigation of Asterochloris algae, we detected substantial genetic, morphological, and ecological differences among particular lineages. First, although the overall mean distance among the lineages was rather low in ITS rDNA (P-distance: 0.022), a considerable genetic differentiation was observed in the actin locus (P-distance: 0.168). Contrary to their broad utilization in fungal research (e.g., Grube and Kroken 2000, Daniel and Meyer 2003), actin intron sequences are still rarely used for species identification in protists. The potential of this marker for identifying and delimiting protist species has been demonstrated in photobiont genera Asterochloris (Nelsen and Gargas 2006, Skaloud and Peksa 2010) and Trebouxia (Kroken and Taylor 2000, Muggia et al. 2010) and in the heterotrophic chrysophycean genus Spumella Cienkowsky (Stoeck et al. 2008).

Second, our statistical analyses revealed a significant morphological differentiation of all investigated *Asterochloris* lineages. Although cell size was determined to be highly plastic under culture conditions, all other morphological features appear to well discriminate among species. Chloroplast morphology, in particular, could be considered to be the best morphological marker for species delineation. Škaloud and Peksa (2008) pointed to the existence of several specific chloroplast types occurring during species ontogeny. In the current study, we demonstrated that although these types are frequently shared by more than one species, particular species could be well recognized by the dominance and unique assemblage of particular types of chloroplasts (Fig. 4, b and c). For example, the parietal lobed chloroplast type prevails in A. phycobiontica, frequently occurs in A. magna, only occasionally appears in A. lobophora, and very rarely develops in A. leprarii, A. woessiae, A. friedlii, and A. irregularis. Moreover, some chloroplast types occur in a small minority of species and could be used to easily define these species. For example, the combination of crenulate and echinate chloroplast types occurs only in A. echinata, while the presence of a globular type defines A. magna. Interestingly, closely related lineages were generally similar in terms of morphology (i.e., A. gaertneri-A. irregularis and A. phycobiontica-A. lobophora).

Finally, previous molecular investigations detected clear ecological differentiation of Asterochloris lineages. Initially, Piercey-Normore and DePriest (2001) explained the photobiont switching among the symbiotic lichen associations as an analogy to human agriculture, where locally best adapted crops are selected and subsequently distributed. Later on, Piercey-Normore (2006) hypothesized that algal genotypes coexisting in the same lichen thalli may be adapted to different forest light levels. Yahr et al. (2006) then suggested the ecological specialization of photobiont lineages, depending on the local environment. Quite recently, specific ecological factors that drive the specialization of Asterochloris lineages were detected by Peksa and Skaloud (2011). Besides substrate and climatic preferences, exposure to rain and sun was the most significant environmental factor, clearly distinguishing particular lineages. The photobionts from ombrophobic and ombrophilic lichens were clustered into completely distinct clades.

In light of the above-mentioned genetic, morphological, and ecological inferences, we believe that each well-resolved *Asterochloris* lineage should be considered to be a distinct species. In fact, by applying an alternative, broader species concept, obvious differences among the lineages would be ignored. This would ultimately prevent us from understanding the true diversity, distribution, and specificity of *Asterochloris* species, as well investigating evolutionary processes that occur at the species level. Therefore, we described all lineages investigated in this study as new species.

Biogeography and specificity of new Asterochloris species. A comparison with previously published diversity surveys allows us to trace the biogeography and specificity of newly proposed Asterochloris species. Although the biogeography of microorganisms has become a highly discussed topic (Caron 2009), investigations dealing with the biogeography of symbiotic protists are very scarce. Geographic separation of particular lineages has been reported for reef-coral dinoflagellate endosymbionts (e.g., Finney et al. 2010, LaJeunesse and Thornhill 2011), as well as for endosymbiotic green algae of the ciliate *Paramecium bursaria* (Hoshina et al. 2005). By contrast, population studies on lichenized *Trebouxia* species indicated that the distribution of particular genotypes is particularly shaped by either climatic factors (Fernández-Mendoza et al. 2011) or distribution patterns of mycobiont partners (Buckley et al. 2014).

The single study dealing with the biogeography of Asterochloris photobionts indicated generally wide (eurychoric) distribution of species (Řídká et al. 2014). Nevertheless, the habitat area of common lineages seems to be more or less restricted based on climatic preferences (e.g., warm-temperate to (sub) arctic distribution of A. glomerata). Though the real diversity of Asterochloris algae is still greatly undersampled, it seems that at least some lineages exhibit restricted geographic distribution independent of climatic factors. According to the actually available genetic data, three of the six newly proposed species (A. echinata, A. gaertneri, and A. leprarii) occur only in Europe (Fig. 2b). These species have never been reported from climatically analogous regions in eastern USA, though the investigations were performed on identical or closely related lichen species (Nelsen and Gargas 2008).

In lichen associations, the term "specificity" is used to refer the range of compatible partners for a given symbiont (Yahr et al. 2006). Lichen specificity is usually perceived from the perspective of a fungal partner, that is, as the range of possible photobionts for a given fungal species. In general, both fungal specialists (having a high specificity) and generalists (having a low specificity) have been distinguished in many lichen genera (see Muggia et al. 2014). However, the specificity could be conceived from the algal perspective, as well. Using our present data, we can compare the algal specificity toward the fungal genera of the newly described Asterochloris species. All species seem to be highly specific toward the genera Cladonia and Lepraria (Fig. 2b). Two species (A. gaertneri and A. leprarii) even form the symbiotic associations exclusively with the fungal genus Lepraria. Interestingly, though a number of Stereocaulon thalli have been investigated for the diversity of their algal partners (Piercey-Normore and DePriest 2001, Nelsen and Gargas 2006, Bačkor et al. 2010, Skaloud and Peksa 2010, own unpublished data), the species herein described as new are obviously not preferred by this fungal genus. In fact, Stereocau*lon* is much widely preferred by two closely related species A. glomerata and A. irregularis (Fig. 2a).

Assessing species diversity in protists. Estimation of the total species diversity in protists remains a highly controversial topic (Caron 2009). Global protist diversity has been proposed to be extraordinarily high by some (Foissner 1999) and generally much lower and fundamentally different from the biodiversity of macroorganisms by others (Fenchel and Finlay 2003). Accordingly, while Finlay and Fenchel (1999) estimated that there are approximately 20,000 protist species, others estimate that there are several million undescribed protist species (Pawlowski et al. 2012). Such substantial differences are primarily caused by differences in methodology (e.g., morphological vs. molecular approaches), species concept (see Boenigk et al. 2012), and theoretical framework (e.g., dispersal-gene flow paradox; De Meester et al. 2002); however, the vast majority of recent investigations have provided undeniable evidence that the overall species diversity of protists is greatly underestimated (e.g., Caron et al. 2012, Pawlowski et al. 2012).

During the past decade, analysis of SSU ribosomal RNA genes has become the most commonly used approach to investigate the diversity of protists. A number of studies have revealed an extremely high proportion of SSU rDNA sequences that could not be assigned to any described species (e.g., López-García et al. 2001, Behnke et al. 2011). Exploration of SSU rDNA sequences has often revealed the existence of several novel, highly diverse lineages (Dolven et al. 2007, Howe et al. 2009). More recently, technological progress in sequencing has enabled researchers to investigate protist diversity at previously unattainable scales. Using next-generation sequencing (NGS) technologies, thousands of SSU rDNA amplicon sequences can be produced from a single sample (Edgcomb et al. 2011); however, the sequence length obtained by NGS sequencing is (at the time of writing) insufficient to characterize complete SSU rDNA genes. Therefore, only short, hypervariable SSU rDNA regions are usually targeted to assess protist diversity (Dunthorn et al. 2012).

In NGS studies, sequence data are typically converted into operational taxonomic units (OTUs) based on sequence similarity; these units are often treated as being synonymous to species (Schmidt et al. 2014). Several methods have been developed to cluster SSU rDNA sequences into OTUs; however, they often partition sequence data differently (Sun et al. 2011). In addition, the taxonomic power of OTUs generated based on sequence similarities has been questioned (Boenigk et al. 2012). In the current study, we demonstrated that morphologically and ecologically diverse species can share identical SSU rDNA sequences. As a consequence, such species would be grouped into a single OTU by NGS data processing. Considering that closely related, mostly cryptic species of protists are commonly identical in their SSU rDNA sequences, the true diversity of eukaryotes can be much greater than that estimated by NGS data. In addition, organisms sharing high SSU rDNA sequence similarity can significantly differ in their ecology and distribution. Therefore, we suggest that the rapidly evolving ITS region should be sequenced, in addition to the broadly used SSU rDNA gene, in NGSbased protist diversity investigations (e.g., Bachy et al. 2013).

Sexuality of lichen photobionts. Focusing on the main objective of this study, we performed a detailed investigation of the morphology and life cycle of many isolated photobionts, which involved hundreds of hours of microscopic observations. On two occasions, we had a brief opportunity to observe sexual reproduction in Asterochloris woessiae cultures, as evidenced by the fusing of biflagellate gametes. According to contemporary symbiotic dogma, lichen symbiosis should lead to the loss of sexual reproduction in the algal symbiont as a result of highly evolved and integrated symbiotic association (Law and Lewis 1983). The absence of sexual reproduction in lichen photobionts (except for the genus Trentepohlia) is, in fact, frequently mentioned in the literature (e.g., Ahmadjian 1987, Gärtner 1992, Friedl and Büdel 1996) and is interpreted as preventing the production of novel genotypes that would be less suited to the mycobiont (Ahmadjian 1993). However, records confirming sexual reproduction in photobionts do exist. The first exhaustive description of sexual reproduction in Trebouxia was published by Warén (1920), who observed frequent production of gametes and their subsequent fusion in photobionts of Anaptychia ciliaris, Physconia distorta, and Xanthoria parietina. Nine years later, Jaag (1929) reported sexual reproduction by iso- and anisogamy in a "Cystococcus parmeliae" photobiont isolated from Flavoparmelia caperata. Another observation was made by Ahmadjian (1959, 1960), who described frequent sexual reproduction in Trebouxia impressa isolated from Physcia stellaris. In both cases, isogamous sexual reproduction resulted in the formation of spherical, smooth-walled zygotes. Finally, indirect evidence of sexual reproduction in Trebouxia was presented by Kroken and Taylor (2000), who found a recombining population structure in photobionts of Letharia spp. by comparing ITS and actin sequences.

Although the existence of sexual reproduction has been disregarded, for example, by Gärtner (1985b), we consider that all of the abovementioned data, as well as our direct observation, clearly establish the presence of sexual reproduction in photobiont genera *Trebouxia* and *Asterochloris*. As sexual reproduction was previously reported only for *Trebouxia* s. str., gamete fusion in *A. woessiae* represents the first record of sexual reproduction in *Asterochloris*.

In conclusion, we demonstrated the obvious existence of a great number of distinct species within the genus *Asterochloris*, which differ genetically, morphologically, and ecologically. In general, the existence of an extraordinarily high number of cryptic, functionally differentiated species is probably the rule rather than the exception in protists. As a consequence, real species entities are poorly defined, providing little use in evaluating their distribution and diversity patterns or clarifying their ecological roles within ecosystems (e.g., Vyverman et al. 1996, Lilly et al. 2007). In addition, the existence of cryptic species with narrow ecological optima would significantly affect our strategies for conservation management (van Oppen and Gates 2006, Cotterill et al. 2008).

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Supporting Information

Additional Supporting Information may be found in the online version of this article at the publisher's web site:

Table S1. Asterochloris sequences used in phylogenetic reconstruction an haplotype networks, including strain and sample numbers, mycobiont species, geographic origin, and GenBank accession numbers for the ITS rDNA, actin, SSU rDNA, and rbcL loci.

Table S2. Morphological characteristics of investigated *Asterochloris* strains.

Culture	Collection	Fungal taxon	Origin		GenBank	accession		ITS		lling sensu
number	number	-	Origin	ITS	actin	SSU	rbcL	genotype	2010	Peksa & Škaloud 2011
	i leprarii Škaloud &	Peksa, sp. nov.							clade 6	clade A8
CAUP H1010	Peksa 183	Lepraria neglecta	Czech Republic, Šumava Mts.	AM906002	AM906031	KP318683	KP318696			
LEP 23	Peksa 173	Lepraria caesioalba	Czech Republic, Šumava Mts.	AM906003	AM906032	-	-			
LEP 25	Peksa 204	Lepraria caesioalba	Czech Republic, Šumava Mts.	AM906004	AM906033	-	-			
-	Peksa 932	Lepraria caesioalba	Czech Republic, Králický Sněžník Mts.	=AM906004	-	-	-			
-	Peksa 864	Lepraria caesioalba	Czech Republic, Šumava Mts.	=AM906004	-	-	-	- I.		
LEP 30	Peksa 207	Lepraria neglecta	Czech Republic, Šumava Mts.	AM906005	AM906034	-	-	۰.		
LEP 61	Peksa 515	Lepraria neglecta	Slovakia, Nízké Tatry Mts-	=AM906005	-	-	-			
-	Peksa 933	Lepraria neglecta	Czech Republic, Hanušovická vrchovina	=AM906005	-	-	-	1		
-	Peksa 857	Lepraria neglecta	Czech Republic, Slavkovský les Mts.	=AM906005	-	-	-	1		
-	Nelsen 3957	Lepraria sp.	Norway, Troms	EU008688	-	-	-	1		
Asterochloris	gaertneri Škaloud	& Peksa, sp. nov.							clade 7	clade A10
CAUP H1013	Peksa 236	Lepraria rigidula	Czech Republic, Králický Sněžník Mts.	AM905997	AM906023	KP318684	KP318697			
-	Peksa 877	Lepraria rigidula	Czech Republic, Šumava Mts.	FM955668	FM955672	-	-	1		
-	Peksa 851	Lepraria rigidula	Czech Republic, Jesenická pahorkatina	=FM955668	-	-	-	1		
-	Peksa 852	Lepraria rigidula	Czech Republic, Dolní Řasnice	=FM955668	-	-	-	1		
-	Peksa 853	Lepraria rigidula	Czech Republic, Plasská pahorkatina	=FM955668	-	-	-	1		
-	Peksa 854	Lepraria rigidula	Czech Republic, Plasská pahorkatina	=FM955668	-	-	-	1		
-	Peksa 856	Lepraria rigidula	Czech Republic, Brdy Mts.	=FM955668	-	-	-	1		
-	Peksa 875	Lepraria rigidula	Czech Republic, Svatobor Mt.	=FM955668	-	-	-	1		
-	Peksa 878	Lepraria rigidula	Czech Republic, Šumava Mts.	=FM955668	-	-	-	1		
-	Peksa 880	, Lepraria rigidula	Czech Republic, Křivoklátsko	=FM955668	-	-	-	1		
-	Peksa 881	Lepraria rigidula	Czech Republic, Křivoklátsko	=FM955668	-	-	-	1		
-	Peksa 883	Lepraria rigidula	Czech Republic, Švihovská vrchovina	=FM955668	-	-	-	1		
RIG3	Peksa 885	Lepraria rigidula	Czech Republic, Slavkovský les Mts.	=FM955668	-	-	-	1		
-	Peksa 887	Lepraria rigidula	Czech Republic, Křivoklátsko	=FM955668	-	-	-	1		
-	Peksa 901	Lepraria rigidula	Czech Republic, Šumava Mts.	=FM955668	-	-	-	I.		
-	Peksa 902	Lepraria rigidula	Czech Republic, Šumava Mts.	=FM955668	-	-	-	1		
-	Peksa 937	Lepraria rigidula	Czech Republic, Hanušovická vrchovina	=FM955668	-	-	-	1		
-	Peksa 940	Lepraria rigidula	Czech Republic, Jizerské hory Mts.	=FM955668	-	-	-	1		
-	Peksa 941	Lepraria rigidula	Czech Republic, Lužické hory Mts.	=FM955668	-	-	-	1		
-	Peksa 943	Lepraria rigidula	Germany, Lužické hory Mts.	=FM955668	-	-	-	1		
-	Peksa 944	Lepraria rigidula	Germany, Lužické hory Mts.	=FM955668	-	-	-	1		
-	Peksa 946	Lepraria rigidula	Slovakia, Nízké Tatry Mts.	=FM955668	-	-	-			
-	Peksa 948	Lepraria rigidula	Slovakia, Nízké Tatry Mts.	=FM955668	-	-	-			
-	Peksa 956	Lepraria rigidula	Czech Republic, Táborská pahorkatina	=FM955668	-	-	-	1		
-	Peksa 945	Lepraria rigidula	Czech Republic, Šumava Mts.	=FM955668	-	-	-			
-	Peksa 886	Lepraria incana	Czech Republic, Plasská pahorkatina	FN556037	KP318677	-	-			
-	Peksa 904	Lepraria membranacea	Czech Republic, Křivoklátsko	FN556038	-	-	-	1		
-	Peksa 905	Lepraria membranacea	Czech Republic, Šumava Mts.	=FN556038		-	-	1		
-	Peksa 936	Lepraria rigidula	Czech Republic, Hanušovická vrchovina	KP318664		1-	-	1		
-	Peksa 900	Lepraria rigidula	Czech Republic, Šumava Mts.	FM955669 FM955673 -		1-	-	١١.	1	
Asterochloris	woessiae Škaloud					1			clade 10	clade A5
	Bayerová 3401	Lepraria borealis	Bulgaria, Stara planina Mts.	AM900492	AM906045	KP318685	KP318698			
	Peksa 542	Lepraria nylanderiana	Czech Republic, Máslovická stráň NR	AM900493	AM906046	-	-	-		
	Peksa 231	Lepraria caesioalba	Czech Republic, Křivoklátsko	AM906014	AM906047	-	-	-		
LEP 14	Peksa 188	Lepraria caesioalba	Czech Republic, Křivoklátsko	=AM906014	-	1-	1-	1		

LEP 53	Peksa 512	Lepraria caesioalba	Czech Republic, Ralská pahorkatina	=AM906014				٦		
LEP 33	Bayerová 3402	Lepraria borealis	Bulgaria, Stara planina Mts.	AM906015	- AM906048	-	-	-		
LEP 34 LEP 37	Peksa 554	Lepraria cf. borealis	Sweden, Grebbestad	=AM906015	AIVI906048	-	-	-		
LEP 37	Peksa 534 Peksa 532	Lepraria borealis	Czech Republic, České středohoří Mts.	=AM906015	-	-	-	-		
LEP 30 LEP 39	Peksa 552 Peksa 513	Lepraria caesioalba	Czech Republic, České středohoří Mts.	=AM906015	-	-	-	-		
LEP 39 LEP 40	Peksa 513 Peksa 507	Lepraria borealis	Czech Republic, Česke stredohori Mis.	=AM906015	-	-	-	-		
LEP 40	Peksa 516	Lepraria borealis	Slovakia, Nízké Tatry Mts.	=AM906015	-			-		
LEP 45	Peksa 543	Lepraria borealis	Czech Republic, Libčice nad Vltavou	=AM906015	-			I.		
LLF JU	Peksa 865	Lepraria borealis	Czech Republic, Křivoklátsko	=AM906015	-			1.		
-	Peksa 934	Lepraria borealis	Czech Republic, Hanušovická vrchovina	=AM906015	-	-	-	-		
-	Peksa 1008	Cladonia foliacea	Czech Republic, Máslovická stráň NR	AM906016	- AM906049	-	-	-		
-	Peksa 919	Cladonia humilis		FM945348	AM906049	-	-	-		
-	Peksa 919 Peksa 925	Cladonia humilis	Czech Republic, Chvaletice Czech Republic, Chvaletice	FM945350	-	-	-	-		
-	Peksa 925 Peksa 924	Cladonia numins Cladonia subulata	Czech Republic, Chvaletice	FM945385	-	-	-	-		
-	Vancurova A66	Cladonia subulata Cladonia sp.	Canary Isles, Tenerife	KP318665	-	-	-	-		
-	Vancurova A00	Cladonia sp. Cladonia sp.	Canary Isles, Gomera	KP318666	-	-	-	-		
-	Vancurova A110 Vancurova A116		Canary Isles, Gomera	KP318667		1-	<u>[</u>			
-	Steinova CL83	Stereocaulon sp. Cladonia sp.	Czech Republic, České Švýcarsko NP	KP318668				-		
-		Cladonia sp. Cladonia coccifera	Spain, Somosierra	KP318669	-	-	-	-		
-	Steinova CL105 Talbot KIS 187	Stereocaulon saxatile	USA, Alaska	DQ229886	- DQ229897	-	-			
-					EU008710	-	-	. 		
-	Nelsen 3637b	Lepraria nigrocincta	Costa Rica, San José	EU008681 EU008687	EU008710	-	-	III. IV.		
-	Nelsen 2166a Sipman 45330	Lepraria sp. Lepraria caesioalba	Costa Rica, San José	EU008667	E0008714	-	-	1V. V.		
-	MACB 95602	Cladonia foliacea	Spain, Zamora Great Britain, Scotland	FM205731	-	-	-	۷.		
-	MACB 90622	Cladonia ronacea Cladonia convoluta	Spain, Soria	FM205726	-	-	-	VI.		
-			Czech Republic, Šumava Mts.		-	-	-	\/II		
					IKP318678					
Asterochloris	Peksa 888 s friedlii Škaloud & P	Lepraria crassissima	Czech Republic, Sumava Mts.	FN556033	KP318678	-	<u> -</u>	VII.	clade 11	clade A7
	s friedlii Škaloud & F	Peksa, sp. nov.				-	- -	VII.	clade 11	clade A7
LEP 04	s <i>friedlii</i> Škaloud & F Peksa 234	Peksa, sp. nov. Lepraria caesioalba	Slovakia, Slovenské Rudohorie Mts.	AM905994	AM906020	- - KP318686	- - KP318699		clade 11	clade A7
	s <i>friedlii</i> Škaloud & F Peksa 234 Peksa 235	Peksa, sp. nov. Lepraria caesioalba Lepraria caesioalba	Slovakia, Slovenské Rudohorie Mts. Slovakia, Slovenské Rudohorie Mts.	AM905994 AM905995	AM906020 AM906021	- - KP318686 -	- - KP318699 -		clade 11	clade A7
LEP 04	s friedlii Škaloud & F Peksa 234 Peksa 235 Nelsen 3974	Peksa, sp. nov. Lepraria caesioalba Lepraria caesioalba Lepraria lobificans	Slovakia, Slovenské Rudohorie Mts. Slovakia, Slovenské Rudohorie Mts. USA, Wisconsin	AM905994 AM905995 DQ229877	AM906020 AM906021 DQ229898	- KP318686 - -	- KP318699 - -	-	clade 11	clade A7
LEP 04	s <i>friedlii</i> Škaloud & F Peksa 234 Peksa 235 Nelsen 3974 Nelsen 2598	Peksa, sp. nov. Lepraria caesioalba Lepraria caesioalba Lepraria lobificans Lepraria atrotomentosa	Slovakia, Slovenské Rudohorie Mts. Slovakia, Slovenské Rudohorie Mts. USA, Wisconsin China, Yunnan	AM905994 AM905995 DQ229877 EU008659	AM906020 AM906021	- - KP318686 - - -	- - KP318699 - - -		clade 11	clade A7
LEP 04	s friedlii Škaloud & F Peksa 234 Peksa 235 Nelsen 3974 Nelsen 2598 Nelsen 3953	Peksa, sp. nov. Lepraria caesioalba Lepraria caesioalba Lepraria lobificans Lepraria atrotomentosa Lepraria caesiella	Slovakia, Slovenské Rudohorie Mts. Slovakia, Slovenské Rudohorie Mts. USA, Wisconsin China, Yunnan USA, Wisconsin	AM905994 AM905995 DQ229877 EU008659 EU008660	AM906020 AM906021 DQ229898	- KP318686 - - - -	- - KP318699 - - - -		clade 11	clade A7
LEP 04	s friedlii Škaloud & F Peksa 234 Peksa 235 Nelsen 3974 Nelsen 2598 Nelsen 3953 Nelsen 3962	eksa, sp. nov. Lepraria caesioalba Lepraria caesioalba Lepraria lobificans Lepraria atrotomentosa Lepraria caesiella Lepraria caesiella	Slovakia, Slovenské Rudohorie Mts. Slovakia, Slovenské Rudohorie Mts. USA, Wisconsin China, Yunnan USA, Wisconsin USA, Massachusetts	AM905994 AM905995 DQ229877 EU008659 EU008660 EU008661	AM906020 AM906021 DQ229898 EU008693 -	- KP318686 - - - - -	- - KP318699 - - - - - -		clade 11	clade A7
LEP 04	s friedlii Škaloud & F Peksa 234 Peksa 235 Nelsen 3974 Nelsen 2598 Nelsen 3953 Nelsen 3962 Nelsen 3968	eksa, sp. nov. Lepraria caesioalba Lepraria caesioalba Lepraria lobificans Lepraria atrotomentosa Lepraria caesiella Lepraria caesiella Lepraria caesiella	Slovakia, Slovenské Rudohorie Mts. Slovakia, Slovenské Rudohorie Mts. USA, Wisconsin China, Yunnan USA, Wisconsin USA, Massachusetts USA, Pennsylvania	AM905994 AM905995 DQ229877 EU008659 EU008660 EU008661 EU008662	AM906020 AM906021 DQ229898 EU008693 - EU008694 EU008695	- KP318686 - - - - - - -	- KP318699 - - - - - - - -		clade 11	clade A7
LEP 04	s friedlii Škaloud & F Peksa 234 Peksa 235 Nelsen 3974 Nelsen 2598 Nelsen 3953 Nelsen 3962 Nelsen 3968 Nelsen 3965	eksa, sp. nov. Lepraria caesioalba Lepraria caesioalba Lepraria lobificans Lepraria atrotomentosa Lepraria caesiella Lepraria caesiella	Slovakia, Slovenské Rudohorie Mts. Slovakia, Slovenské Rudohorie Mts. USA, Wisconsin China, Yunnan USA, Wisconsin USA, Massachusetts USA, Pennsylvania USA, Pennsylvania	AM905994 AM905995 DQ229877 EU008659 EU008660 EU008661 EU008662 EU008663	AM906020 AM906021 DQ229898 EU008693 - EU008694 EU008695 EU008696	- KP318686 - - - - - - - -	- KP318699 - - - - - - - - - - - - -		clade 11	clade A7
LEP 04	s friedlii Škaloud & F Peksa 234 Peksa 235 Nelsen 3974 Nelsen 2598 Nelsen 3953 Nelsen 3962 Nelsen 3968	eksa, sp. nov. Lepraria caesioalba Lepraria caesioalba Lepraria lobificans Lepraria atrotomentosa Lepraria caesiella Lepraria caesiella Lepraria caesiella Lepraria caesioalba	Slovakia, Slovenské Rudohorie Mts. Slovakia, Slovenské Rudohorie Mts. USA, Wisconsin China, Yunnan USA, Wisconsin USA, Massachusetts USA, Pennsylvania USA, Pennsylvania USA, Pennsylvania	AM905994 AM905995 DQ229877 EU008659 EU008660 EU008661 EU008662	AM906020 AM906021 DQ229898 EU008693 - EU008694 EU008695	- KP318686 - - - - - - - - - -	- KP318699 - - - - - - - - - - - - - - - -		clade 11	clade A7
LEP 04	s friedlii Škaloud & F Peksa 234 Peksa 235 Nelsen 3974 Nelsen 2598 Nelsen 3953 Nelsen 3962 Nelsen 3968 Nelsen 3965 Nelsen 3969 Nelsen 3971	eksa, sp. nov. Lepraria caesioalba Lepraria caesioalba Lepraria lobificans Lepraria atrotomentosa Lepraria caesiella Lepraria caesiella Lepraria caesiella Lepraria caesioalba Lepraria caesioalba	Slovakia, Slovenské Rudohorie Mts. Slovakia, Slovenské Rudohorie Mts. USA, Wisconsin China, Yunnan USA, Wisconsin USA, Massachusetts USA, Pennsylvania USA, Pennsylvania USA, Pennsylvania USA, Pennsylvania	AM905994 AM905995 DQ229877 EU008659 EU008660 EU008661 EU008662 EU008663 EU008666 EU008666	AM906020 AM906021 DQ229898 EU008693 - EU008694 EU008695 EU008696	- KP318686 - - - - - - - - - - - - -	- KP318699 - - - - - - - - - - - - - - - - - -		clade 11	clade A7
LEP 04	s friedlii Škaloud & F Peksa 234 Peksa 235 Nelsen 3974 Nelsen 2598 Nelsen 3953 Nelsen 3962 Nelsen 3968 Nelsen 3965 Nelsen 3969	eksa, sp. nov. Lepraria caesioalba Lepraria caesioalba Lepraria lobificans Lepraria atrotomentosa Lepraria caesiella Lepraria caesiella Lepraria caesiella Lepraria caesioalba Lepraria caesioalba Lepraria incana	Slovakia, Slovenské Rudohorie Mts. Slovakia, Slovenské Rudohorie Mts. USA, Wisconsin China, Yunnan USA, Wisconsin USA, Massachusetts USA, Pennsylvania USA, Pennsylvania USA, Pennsylvania	AM905994 AM905995 DQ229877 EU008659 EU008660 EU008661 EU008662 EU008663 EU008666 EU008668 EU008668 EU008670	AM906020 AM906021 DQ229898 EU008693 - EU008694 EU008695 EU008696 EU008699 - EU008700	- KP318686 - - - - - - - - - - - - - - -	- KP318699 - - - - - - - - - - - - - - - - - -		clade 11	clade A7
LEP 04	s friedlii Škaloud & F Peksa 234 Peksa 235 Nelsen 3974 Nelsen 2598 Nelsen 3953 Nelsen 3962 Nelsen 3965 Nelsen 3965 Nelsen 3969 Nelsen 3971 Nelsen 3950	eksa, sp. nov. Lepraria caesioalba Lepraria caesioalba Lepraria lobificans Lepraria atrotomentosa Lepraria caesiella Lepraria caesiella Lepraria caesioalba Lepraria caesioalba Lepraria incana Lepraria lobificans	Slovakia, Slovenské Rudohorie Mts. Slovakia, Slovenské Rudohorie Mts. USA, Wisconsin China, Yunnan USA, Wisconsin USA, Massachusetts USA, Pennsylvania USA, Pennsylvania USA, Pennsylvania USA, Maryland USA, Wisconsin	AM905994 AM905995 DQ229877 EU008659 EU008660 EU008661 EU008662 EU008663 EU008666 EU008666	AM906020 AM906021 DQ229898 EU008693 - EU008694 EU008695 EU008696 EU008699 -	- KP318686 - - - - - - - - - - - - - - - - - -	- KP318699 - - - - - - - - - - - - - - - - - -		clade 11	clade A7
LEP 04	s friedlii Škaloud & F Peksa 234 Peksa 235 Nelsen 3974 Nelsen 2598 Nelsen 3953 Nelsen 3962 Nelsen 3965 Nelsen 3969 Nelsen 3971 Nelsen 3950 Nelsen 3951	eksa, sp. nov.Lepraria caesioalbaLepraria caesioalbaLepraria lobificansLepraria atrotomentosaLepraria caesiellaLepraria caesiellaLepraria caesiellaLepraria caesiellaLepraria caesioalbaLepraria caesioalbaLepraria caesioalbaLepraria caesioalbaLepraria caesioalbaLepraria incanaLepraria lobificansLepraria lobificansLepraria lobificans	Slovakia, Slovenské Rudohorie Mts. Slovakia, Slovenské Rudohorie Mts. USA, Wisconsin China, Yunnan USA, Wisconsin USA, Massachusetts USA, Pennsylvania USA, Pennsylvania USA, Pennsylvania USA, Maryland USA, Wisconsin USA, Wisconsin	AM905994 AM905995 DQ229877 EU008659 EU008660 EU008661 EU008662 EU008663 EU008666 EU008668 EU008668 EU008670 EU008671 EU008672	AM906020 AM906021 DQ229898 EU008693 - EU008694 EU008695 EU008696 EU008699 - EU008700 EU008700 EU008701 -	- KP318686 - - - - - - - - - - - - - - - - - -	- KP318699		clade 11	clade A7
LEP 04	s friedlii Škaloud & F Peksa 234 Peksa 235 Nelsen 3974 Nelsen 2598 Nelsen 3953 Nelsen 3962 Nelsen 3965 Nelsen 3965 Nelsen 3971 Nelsen 3950 Nelsen 3951 Nelsen 3956	eksa, sp. nov.Lepraria caesioalbaLepraria caesioalbaLepraria lobificansLepraria atrotomentosaLepraria caesiellaLepraria caesiellaLepraria caesiellaLepraria caesioalbaLepraria caesioalbaLepraria caesioalbaLepraria caesioalbaLepraria caesioalbaLepraria incanaLepraria lobificansLepraria lobificansLepraria lobificansLepraria lobificans	Slovakia, Slovenské Rudohorie Mts. Slovakia, Slovenské Rudohorie Mts. USA, Wisconsin China, Yunnan USA, Wisconsin USA, Massachusetts USA, Pennsylvania USA, Pennsylvania USA, Pennsylvania USA, Maryland USA, Wisconsin USA, Wisconsin	AM905994 AM905995 DQ229877 EU008659 EU008660 EU008661 EU008662 EU008663 EU008663 EU008668 EU008668 EU008670 EU008671	AM906020 AM906021 DQ229898 EU008693 - EU008694 EU008695 EU008696 EU008699 - EU008700	- KP318686 - - - - - - - - - - - - - - - - - -	- KP318699		clade 11	clade A7
LEP 04	s friedlii Škaloud & F Peksa 234 Peksa 235 Nelsen 3974 Nelsen 2598 Nelsen 3953 Nelsen 3962 Nelsen 3965 Nelsen 3969 Nelsen 3971 Nelsen 3950 Nelsen 3950 Nelsen 3951 Nelsen 3956 Nelsen 3958	eksa, sp. nov.Lepraria caesioalbaLepraria caesioalbaLepraria lobificansLepraria atrotomentosaLepraria caesiellaLepraria caesiellaLepraria caesiellaLepraria caesioalbaLepraria caesioalbaLepraria caesioalbaLepraria caesioalbaLepraria caesioalbaLepraria caesioalbaLepraria incanaLepraria lobificansLepraria lobificansLepraria lobificansLepraria lobificansLepraria lobificansLepraria lobificansLepraria lobificansLepraria lobificansLepraria lobificans	Slovakia, Slovenské Rudohorie Mts. Slovakia, Slovenské Rudohorie Mts. USA, Wisconsin China, Yunnan USA, Wisconsin USA, Massachusetts USA, Pennsylvania USA, Pennsylvania USA, Pennsylvania USA, Maryland USA, Wisconsin USA, Wisconsin USA, Wisconsin	AM905994 AM905995 DQ229877 EU008659 EU008660 EU008661 EU008663 EU008663 EU008668 EU008668 EU008668 EU008670 EU008671 EU008672 EU008673	AM906020 AM906021 DQ229898 EU008693 - EU008694 EU008695 EU008696 EU008699 - EU008700 EU008700 EU008702	- KP318686 - - - - - - - - - - - - -	- KP318699		clade 11	clade A7
LEP 04	s friedlii Škaloud & F Peksa 234 Peksa 235 Nelsen 3974 Nelsen 3974 Nelsen 3974 Nelsen 3974 Nelsen 3974 Nelsen 3974 Nelsen 3962 Nelsen 3965 Nelsen 3969 Nelsen 3971 Nelsen 3950 Nelsen 3951 Nelsen 3956 Nelsen 3958 Nelsen 3957	eksa, sp. nov.Lepraria caesioalbaLepraria caesioalbaLepraria lobificansLepraria atrotomentosaLepraria caesiellaLepraria caesiellaLepraria caesiellaLepraria caesiellaLepraria caesioalbaLepraria caesioalbaLepraria caesioalbaLepraria caesioalbaLepraria caesioalbaLepraria lobificansLepraria lobificans	Slovakia, Slovenské Rudohorie Mts. Slovakia, Slovenské Rudohorie Mts. USA, Wisconsin China, Yunnan USA, Misconsin USA, Massachusetts USA, Pennsylvania USA, Pennsylvania USA, Pennsylvania USA, Maryland USA, Wisconsin USA, Maryland USA, Wisconsin USA, Wisconsin USA, Tennessee USA, Indiana USA, Massachusetts	AM905994 AM905995 DQ229877 EU008659 EU008660 EU008661 EU008662 EU008663 EU008668 EU008668 EU008670 EU008671 EU008671 EU008672 EU008673 EU008674 EU008676	AM906020 AM906021 DQ229898 EU008693 - EU008694 EU008695 EU008696 EU008699 - EU008700 EU008700 EU008701 - EU008702 EU008703 EU008705	- KP318686 - - - - - - - - - - - - -	- KP318699		clade 11	clade A7
LEP 04	s friedlii Škaloud & F Peksa 234 Peksa 235 Nelsen 3974 Nelsen 3974 Nelsen 3974 Nelsen 3974 Nelsen 3953 Nelsen 3962 Nelsen 3965 Nelsen 3969 Nelsen 3950 Nelsen 3951 Nelsen 3956 Nelsen 3957 Nelsen 3957	eksa, sp. nov.Lepraria caesioalbaLepraria caesioalbaLepraria lobificansLepraria atrotomentosaLepraria caesiellaLepraria caesiellaLepraria caesiellaLepraria caesiellaLepraria caesioalbaLepraria caesioalbaLepraria caesioalbaLepraria caesioalbaLepraria caesioalbaLepraria incanaLepraria lobificansLepraria lobificans	Slovakia, Slovenské Rudohorie Mts. Slovakia, Slovenské Rudohorie Mts. USA, Wisconsin China, Yunnan USA, Wisconsin USA, Wisconsin USA, Massachusetts USA, Pennsylvania USA, Pennsylvania USA, Pennsylvania USA, Maryland USA, Wisconsin USA, Wisconsin USA, Massachusetts USA, Maryland USA, Wisconsin USA, Tennessee USA, Indiana USA, West Virginia	AM905994 AM905995 DQ229877 EU008659 EU008660 EU008661 EU008662 EU008663 EU008668 EU008668 EU008668 EU008670 EU008671 EU008671 EU008673 EU008674	AM906020 AM906021 DQ229898 EU008693 - EU008694 EU008695 EU008696 EU008699 - EU008700 EU008701 - EU008702 EU008703	- KP318686 - - - - - - - - - - - - -	- KP318699		clade 11	clade A7
LEP 04	s friedlii Škaloud & F Peksa 234 Peksa 235 Nelsen 3974 Nelsen 3974 Nelsen 3974 Nelsen 3974 Nelsen 3953 Nelsen 3962 Nelsen 3968 Nelsen 3965 Nelsen 3969 Nelsen 3971 Nelsen 3950 Nelsen 3951 Nelsen 3956 Nelsen 3958 Nelsen 3957 Nelsen 3957 Nelsen 3970 Nelsen 3970	eksa, sp. nov.Lepraria caesioalbaLepraria caesioalbaLepraria lobificansLepraria atrotomentosaLepraria caesiellaLepraria caesiellaLepraria caesiellaLepraria caesiellaLepraria caesiellaLepraria caesiellaLepraria caesiellaLepraria caesialbaLepraria caesioalbaLepraria lobificansLepraria lobificans	Slovakia, Slovenské Rudohorie Mts. Slovakia, Slovenské Rudohorie Mts. USA, Wisconsin China, Yunnan USA, Wisconsin USA, Wisconsin USA, Massachusetts USA, Pennsylvania USA, Pennsylvania USA, Maryland USA, Wisconsin USA, Wisconsin USA, Wisconsin USA, Maryland USA, Wisconsin USA, Tennessee USA, Indiana USA, West Virginia USA, Methigan	AM905994 AM905995 DQ229877 EU008659 EU008660 EU008661 EU008663 EU008663 EU008668 EU008668 EU008670 EU008671 EU008672 EU008673 EU008674 EU008676 EU008677	AM906020 AM906021 DQ229898 EU008693 - EU008694 EU008695 EU008696 EU008699 - EU008700 EU008701 - EU008702 EU008703 EU008705 EU008706	- KP318686 - - - - - - - - - - - - -	- KP318699 KP318699		clade 11	clade A7
LEP 04	s friedlii Škaloud & F Peksa 234 Peksa 235 Nelsen 3974 Nelsen 3974 Nelsen 3974 Nelsen 3974 Nelsen 3953 Nelsen 3962 Nelsen 3968 Nelsen 3965 Nelsen 3969 Nelsen 3971 Nelsen 3950 Nelsen 3950 Nelsen 3951 Nelsen 3956 Nelsen 3957 Nelsen 3957 Nelsen 3970 Nelsen 3970	eksa, sp. nov.Lepraria caesioalbaLepraria caesioalbaLepraria lobificansLepraria atrotomentosaLepraria caesiellaLepraria caesiellaLepraria caesiellaLepraria caesiellaLepraria caesioalbaLepraria caesioalbaLepraria caesioalbaLepraria caesioalbaLepraria lobificansLepraria lobificans <td>Slovakia, Slovenské Rudohorie Mts. Slovakia, Slovenské Rudohorie Mts. USA, Wisconsin China, Yunnan USA, Wisconsin USA, Wisconsin USA, Massachusetts USA, Pennsylvania USA, Pennsylvania USA, Pennsylvania USA, Maryland USA, Wisconsin USA, Wisconsin USA, Massachusetts USA, Maryland USA, Wisconsin USA, Tennessee USA, Indiana USA, West Virginia</td> <td>AM905994 AM905995 DQ229877 EU008659 EU008660 EU008661 EU008663 EU008663 EU008668 EU008670 EU008671 EU008673 EU008674 EU008677 EU008677</td> <td>AM906020 AM906021 DQ229898 EU008693 - EU008694 EU008695 EU008696 EU008699 - EU008700 EU008701 - EU008702 EU008703 EU008705 EU008706 EU008708</td> <td>- KP318686 - - - - - - - - - - - - -</td> <td>- KP318699 KP318699</td> <td></td> <td>clade 11</td> <td>clade A7</td>	Slovakia, Slovenské Rudohorie Mts. Slovakia, Slovenské Rudohorie Mts. USA, Wisconsin China, Yunnan USA, Wisconsin USA, Wisconsin USA, Massachusetts USA, Pennsylvania USA, Pennsylvania USA, Pennsylvania USA, Maryland USA, Wisconsin USA, Wisconsin USA, Massachusetts USA, Maryland USA, Wisconsin USA, Tennessee USA, Indiana USA, West Virginia	AM905994 AM905995 DQ229877 EU008659 EU008660 EU008661 EU008663 EU008663 EU008668 EU008670 EU008671 EU008673 EU008674 EU008677 EU008677	AM906020 AM906021 DQ229898 EU008693 - EU008694 EU008695 EU008696 EU008699 - EU008700 EU008701 - EU008702 EU008703 EU008705 EU008706 EU008708	- KP318686 - - - - - - - - - - - - -	- KP318699 KP318699		clade 11	clade A7
LEP 04	s friedlii Škaloud & F Peksa 234 Peksa 235 Nelsen 3974 Nelsen 3974 Nelsen 3974 Nelsen 3974 Nelsen 3953 Nelsen 3953 Nelsen 3962 Nelsen 3965 Nelsen 3965 Nelsen 3969 Nelsen 3950 Nelsen 3951 Nelsen 3956 Nelsen 3957 Nelsen 3957 Nelsen 3957 Nelsen 3970 Nelsen 3976 Nelsen 3976 Nelsen 2590	eksa, sp. nov.Lepraria caesioalbaLepraria caesioalbaLepraria lobificansLepraria atrotomentosaLepraria caesiellaLepraria caesiellaLepraria caesiellaLepraria caesiellaLepraria caesiellaLepraria caesiellaLepraria caesiellaLepraria caesialbaLepraria lobificansLepraria lobificansLepra	Slovakia, Slovenské Rudohorie Mts. Slovakia, Slovenské Rudohorie Mts. USA, Wisconsin China, Yunnan USA, Wisconsin USA, Wisconsin USA, Massachusetts USA, Pennsylvania USA, Pennsylvania USA, Pennsylvania USA, Maryland USA, Wisconsin USA, Wisconsin USA, Wisconsin USA, Tennessee USA, Indiana USA, West Virginia USA, Michigan China, Yunnan	AM905994 AM905995 DQ229877 EU008659 EU008660 EU008661 EU008663 EU008663 EU008668 EU008670 EU008671 EU008673 EU008674 EU008677 EU008679 EU008679	AM906020 AM906021 DQ229898 EU008693 - EU008694 EU008695 EU008696 EU008699 - EU008700 EU008701 - EU008702 EU008703 EU008705 EU008706 EU008708	- KP318686 - - - - - - - - - - - - -	- KP318699 - KP318699		clade 11	clade A7
LEP 04	s friedlii Škaloud & F Peksa 234 Peksa 235 Nelsen 3974 Nelsen 3974 Nelsen 3974 Nelsen 3974 Nelsen 3953 Nelsen 3953 Nelsen 3962 Nelsen 3965 Nelsen 3965 Nelsen 3969 Nelsen 3950 Nelsen 3951 Nelsen 3955 Nelsen 3956 Nelsen 3957 Nelsen 3957 Nelsen 3970 Nelsen 3976 Nelsen 2590 Olszewski 54090	eksa, sp. nov.Lepraria caesioalbaLepraria caesioalbaLepraria lobificansLepraria atrotomentosaLepraria caesiellaLepraria caesiellaLepraria caesiellaLepraria caesiellaLepraria caesiellaLepraria caesiellaLepraria caesiellaLepraria caesialbaLepraria caesioalbaLepraria lobificansLepraria nivalisLepraria sp.	Slovakia, Slovenské Rudohorie Mts. Slovakia, Slovenské Rudohorie Mts. USA, Wisconsin China, Yunnan USA, Wisconsin USA, Wisconsin USA, Massachusetts USA, Pennsylvania USA, Pennsylvania USA, Pennsylvania USA, Pennsylvania USA, Pennsylvania USA, Maryland USA, Wisconsin USA, Wisconsin USA, Tennessee USA, Indiana USA, West Virginia USA, Michigan China, Yunnan Canada, Quebec	AM905994 AM905995 DQ229877 EU008659 EU008660 EU008661 EU008663 EU008663 EU008668 EU008670 EU008671 EU008672 EU008673 EU008676 EU008677 EU008679 EU008678	AM906020 AM906021 DQ229898 EU008693 - EU008694 EU008695 EU008696 EU008699 - EU008700 EU008701 - EU008701 - EU008703 EU008703 EU008705 EU008706 EU008708 EU008709 -	- KP318686 - - - - - - - - - - - - -	- KP318699 - KP318699		clade 11	clade A7
LEP 04	s friedlii Škaloud & F Peksa 234 Peksa 235 Nelsen 3974 Nelsen 3974 Nelsen 3974 Nelsen 3953 Nelsen 3953 Nelsen 3962 Nelsen 3968 Nelsen 3965 Nelsen 3969 Nelsen 3971 Nelsen 3950 Nelsen 3951 Nelsen 3956 Nelsen 3956 Nelsen 3957 Nelsen 3957 Nelsen 3970 Nelsen 3970 Nelsen 3976 Nelsen 2590 Olszewski 54090 Nelsen 3959	eksa, sp. nov.Lepraria caesioalbaLepraria caesioalbaLepraria lobificansLepraria atrotomentosaLepraria caesiellaLepraria caesiellaLepraria caesiellaLepraria caesiellaLepraria caesiellaLepraria caesiellaLepraria caesiellaLepraria caesialbaLepraria lobificansLepraria lobificansLepra	Slovakia, Slovenské Rudohorie Mts. Slovakia, Slovenské Rudohorie Mts. USA, Wisconsin China, Yunnan USA, Wisconsin USA, Wisconsin USA, Wisconsin USA, Massachusetts USA, Pennsylvania USA, Pennsylvania USA, Pennsylvania USA, Pennsylvania USA, Maryland USA, Wisconsin USA, Wisconsin USA, Wisconsin USA, Tennessee USA, Indiana USA, Mest Virginia USA, Michigan China, Yunnan Canada, Quebec USA, North Carolina	AM905994 AM905995 DQ229877 EU008659 EU008660 EU008661 EU008663 EU008663 EU008668 EU008670 EU008671 EU008673 EU008676 EU008677 EU008679 EU008679 EU008679	AM906020 AM906021 DQ229898 EU008693 - EU008694 EU008695 EU008696 EU008699 - EU008700 EU008701 - EU008701 - EU008703 EU008703 EU008705 EU008706 EU008708 EU008709 -	- KP318686 KP318686 KP318686	- KP318699 - KP318699		clade 11	clade A7
LEP 04	s friedlii Škaloud & F Peksa 234 Peksa 235 Peksa 235 Nelsen 3974 Nelsen 3974 Nelsen 3974 Nelsen 3953 Nelsen 3953 Nelsen 3962 Nelsen 3968 Nelsen 3965 Nelsen 3969 Nelsen 3950 Nelsen 3950 Nelsen 3951 Nelsen 3956 Nelsen 3957 Nelsen 3957 Nelsen 3957 Nelsen 3970 Nelsen 3970 Nelsen 3976 Nelsen 2590 Olszewski 54090 Nelsen 3959 Sipman 49865	eksa, sp. nov.Lepraria caesioalbaLepraria caesioalbaLepraria lobificansLepraria atrotomentosaLepraria caesiellaLepraria caesiellaLepraria caesiellaLepraria caesiellaLepraria caesiellaLepraria caesiellaLepraria caesiellaLepraria caesioalbaLepraria caesioalbaLepraria lobificansLepraria sp.Lepraria sp.	Slovakia, Slovenské Rudohorie Mts. Slovakia, Slovenské Rudohorie Mts. USA, Wisconsin China, Yunnan USA, Wisconsin USA, Wisconsin USA, Wisconsin USA, Massachusetts USA, Pennsylvania USA, Pennsylvania USA, Pennsylvania USA, Pennsylvania USA, Pennsylvania USA, Maryland USA, Wisconsin USA, Wisconsin USA, Wisconsin USA, Tennessee USA, Indiana USA, Mest Virginia USA, Mest Virginia USA, Michigan China, Yunnan Canada, Quebec USA, North Carolina China, Yunnan	AM905994 AM905995 DQ229877 EU008659 EU008660 EU008661 EU008663 EU008663 EU008668 EU008670 EU008672 EU008673 EU008674 EU008679 EU008679 EU008680 EU008680 EU008680 EU008680 EU008680 EU008680 EU008680	AM906020 AM906021 DQ229898 EU008693 - EU008694 EU008695 EU008696 EU008699 - EU008700 EU008701 - EU008701 - EU008703 EU008705 EU008706 EU008708 EU008709 - EU008713 -	- KP318686 KP318686 KP318686	- KP318699 - KP318699		clade 11	clade A7

-	Peksa 947	Lepraria rigidula	Slovakia, Slovenské Rudohorie Mts.	=FN556034	-	-	-	1	I	
-	Nelsen 3951	Cladonia caespitecia	USA, Pennsylvania	DQ229879	DQ229899	-	-			
	Nelsen 3966	Lepraria caesioalba	USA, Pennsylvania	EU008664	EU008697	-	_	-		
	Nelsen 3967	Lepraria caesioalba	USA, Pennsylvania	EU008665	EU008698	-	-	П.		
	Nelsen 3973	Lepraria lobificans	USA, Wisconsin	EU008678	EU008707	_	_			
	West & Westad s.n.		USA, Wisconsin	EU008685	EU008712	_	_	-		
EP 33	Peksa 225	Lepraria caesioalba	Romania, Paring Mts.	AM905996	AM906022	-	-	.		
	Nelsen 3972	Lepraria incana	USA, Virginia	EU008669	-	-	-	IV.		
	Nelsen 3960	Lepraria lobificans	USA, Wisconsin	EU008675	EU008704	-	-	V.		
	Peksa 1066	Lepraria elobata	Slovakia, Nízké Tatry Mts.	KP318670	-	-		VI.		
Asterochloris	echinata Škaloud &	,	Clovania, Mizice Tairy Mis.						clade 14	clade A3
CAUP H1012		Lepraria rigidula	Czech Republic, Lužické hory Mts.	AM905992	AM906017	KP318687	KP318700	1		olddo rio
	Bayerová 3324	Lepraria granulata	Bulgaria, Pirin Mts.	FN556029	-	-	-	I.		
EP 32	Peksa 185	Lepraria caesioalba	Czech Republic, Lužické hory Mts.	FM955666	- FM955670		_			
EP 55	Peksa 551	Lepraria caesioalba	Czech Republic, Lužické hory Mts.	FM955667	FM955671		_	-		
LT 55	Peksa 861	Lepraria alpina	Spain, Somiedo Mts.	FN556026	1 10/95507 1	-	-	II.		
	Steinova CL106	Cladonia diversa	Portugal, Beira Alta	KP318671				-		
	Peksa 859	Lepraria caesioalba	Spain, Somiedo Mts.	FN556028	-	-				
	Peksa 859 Peksa 930	Lepraria caesioalba Lepraria caesioalba	Czech Republic, Králický Sněžník Mts.	=FN556028	-	-		- 111.		
EP 49	Peksa 930 Peksa 520			FN556027	-	-				
LF 49	Peksa 520 Palice 11603	Lepraria caesioalba Lepraria caesioalba	Slovakia, Nízké Tatry Mts.		-	-		IV.		
Astorophlaria	I lobophora Škaloud	,	Czech Republic, Krkonoše Mts.	=FN556027	<u> </u>		<u> </u>		clade 16	clade A2
		Lepraria caesioalba	Czach Banublia, Doupovaká vroby Mto	A M006007	AM006026				claue 10	Claue Az
EP 02 CAUP H1014	Peksa 196 Peksa 166		Czech Republic, Doupovské vrchy Mts.	AM906007 AM906008	AM906036 AM906037	-	-	-		
EP 27		Lepraria caesioalba	Czech Republic, Šumava Mts. Czech Republic, Šumava Mts.		AM906037 AM906038	-	-	-		
	Peksa 194	Lepraria caesioalba		AM906009	AIVI906036	-	-	-		
EP 42	Peksa 545	Lepraria caesioalba	Czech Republic, Lužické hory Mts.	=AM906009	-	-	-	-		
EP 43	Peksa 503	Lepraria caesioalba	Czech Republic, Krkonoše Mts.	=AM906009	-	-	-	-		
EP 44	Peksa 495	Lepraria caesioalba	Czech Republic, Lužické hory Mts.	=AM906009	-	-	-	-		
EP 57	Peksa 527	Lepraria caesioalba	Czech Republic, Doupovské vrchy Mts.	=AM906009	-	-	-	-		
50.00	Peksa 867	Lepraria caesioalba	USA, California, Sierra Nevada Mts.	=AM906009	-	-	-	4		
EP 28	Peksa 192	Lepraria alpina	Czech Republic, Šumava Mts.	AM906010	AM906039	KP318688	-	4		
	Palice 11626	Lepraria alpina	Czech Republic, Krkonoše Mts.	=AM906010	-	-	-	_		
DIP 01	Peksa 182	Diploschistes muscorum	Czech Republic, Šumava Mts.	AM906011	AM906040	-	-	_		
	Peksa 576	Cladonia coniocraea	Czech Republic, Chvaletice	FM945354	-			-		
	Peksa 927	Cladonia rei	Czech Republic, Chvaletice	FM945356	-	-	-	-		
	Peksa 922	Cladonia macilenta	Czech Republic, Chvaletice	FM945366	-	-	-	-		
	Peksa 816	Cladonia ochrochlora	Slovakia, Veľká Fatra Mts.	FM945369	-	-	-	-		
	Peksa 791	Cladonia pyxidata	Slovakia, Volovské vrchy Mts.	FM945373	-	-	-	-		
	Peksa 790	Cladonia rangiformis	Slovakia, Volovské vrchy Mts.	FM945377	-	-	-	-		
	Peksa 926	Cladonia rei	Czech Republic, Chvaletice	FM945379	-	-	-	-		
	Peksa 794	Cladonia rei	Slovakia, Nízké Tatry Mts.	FM945382	-	-	-	-		
	Peksa 809	Cladonia rei	Slovakia, Slovenské Rudohorie Mts.	FM945387	-	-	-	-		
DIP 09	Peksa 928	Diploschistes muscorum	Czech Republic, Chvaletice	FM945390	-	-	-	-		
	Peksa 786	Stereocaulon tomentosum	Slovakia, Volovské vrchy Mts.	FM945391	-	-	-	-		
	Normore 289	Cladonia parasitica	Canada, Nova Scotia	AF345426	-	-	-	-		
	RY581	Cladonia pachycladodes	USA, Florida	AY712703	-	-	-	-		
	RY1002	Cladonia leporina	USA, Florida	=AY712703	-	-	-	-		
	RY572	Cladonia leporina	USA, Florida	=AY712703	-	-	-			
	RY657	Cladonia leporina	USA, Florida	=AY712703	-	-	-	Ι.		
	RY808	Cladonia leporina	USA, Florida	=AY712703	-	-	-	1.		
-	RY922	Cladonia leporina	USA, Florida	=AY712703	-	-	-			

RY979	Cladonia leporina	USA, Florida	=AY712703	-	-	_	
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	Diploschistes muscorum			-	-	-	
	Cladonia coccifera			-	-	-	
RY762	Cladonia pachycladodes	USA, Florida	AY712707	-	-	-	
RY997b	Cladonia pachycladodes	USA, Florida	=AY712707	-	-	-	
RY229	Cladonia leporina	USA, Florida	=AY712707	-	-	-	
RY649	Cladonia leporina	USA, Florida	=AY712707	-	-	-	
	Cladonia leporina	USA, Florida	=AY712707	-	-	-	
RY10	Cladonia perforata	USA, Florida	=AY712707	-	-	-	
RY576	Cladonia perforata	USA, Florida	=AY712707	-	-	-	
RY653	Cladonia perforata	USA, Florida	=AY712707	-	-	-	П.
RY802	Cladonia prostrata	USA, Florida	=AY712707	-	-	-	11.
RY924	Cladonia prostrata	USA, Florida	=AY712707	-	-	-	
RY985	Cladonia prostrata	USA, Florida	=AY712707	-	-	-	
RY230	Cladonia subtenuis	USA, Florida	=AY712707	-	-	-	
RY1186	Cladonia subtenuis	USA, North Carolina	=AY712707	-	-	-	
RY968	Cladonia pachycladodes	USA, Florida	AY712710	-	-	-	
RY954	Cladonia prostrata	USA, Florida	=AY712710	-	-	-	
-	Cladonia perforata	USA, Florida	AY842280	-	-	-	
Peksa 233	Lepraria caesioalba	Czech Republic, Vladař Mt.	AM906006	AM906035	-	-	
Peksa 509	Lepraria caesioalba	Czech Republic, Doupovské vrchy Mts.	=AM906006	-	-	-	III.
Peksa 866	Lepraria borealis			KP318679	-	-	IV.
Peksa 544	Lepraria caesioalba	Czech Republic, Chlumská hora Mt.		-	-	-	V.
				-	-	-	VI.
				-	-	-	VII.
RY765	Cladonia perforata	USA, Florida	AY712701	1-	-	-	0
					+		
RY666	Cladonia prostrata	USA, Florida	AY/12/02	-	-	-	P
RY666 RY967	Cladonia prostrata Cladonia pachycladodes	USA, Florida USA, Florida	AY712702 AY712704	-	-	-	P
	RY220 RY822 RY903 RY904 RY120 RY804 RY1006 RY763 RY854 RY926 RY1135 RY120 RY135 RY123 RY849 Peksa 496 Peksa 496 Peksa 496 Peksa 497 Steinova CL57 RY762 RY997b RY229 RY649 RY919 RY10 RY576 RY802 RY924 RY925 RY926 RY928 RY928 RY928 RY924	RY220Cladonia pachycladodesRY822Cladonia pachycladodesRY903Cladonia pachycladodesRY966Cladonia pachycladodesRY120Cladonia pachycladodesRY120Cladonia perforataRY804Cladonia perforataRY1006Cladonia prostrataRY1006Cladonia prostrataRY763Cladonia prostrataRY926Cladonia prostrataRY1135Cladonia subtenuisRY1123Cladonia subtenuisRY1123Cladonia subtenuisRY129Cladonia subtenuisRY1238Cladonia subtenuisRY1238Cladonia pachycladodesNormore 1384 (SB2Cladonia pocillumNormore 1058 (SB3Cladonia fimbriataIH17Cladonia fimbriataIH17Cladonia cocciferaRY762Cladonia pachycladodesRY997bCladonia pachycladodesRY997bCladonia pachycladodesRY997bCladonia pachycladodesRY929Cladonia pachycladodesRY929Cladonia pachycladodesRY920Cladonia pachycladodesRY921Cladonia pachycladodesRY922Cladonia pachycladodesRY933Cladonia perforataRY623Cladonia perforataRY624Cladonia perforataRY935Cladonia prostrataRY936Cladonia prostrataRY924Cladonia prostrataRY925Cladonia prostrataRY985Cladonia prostrataRY985Cladonia prostrata </td <td>RY220 Cladonia pachycladodes USA, Florida RY803 Cladonia pachycladodes USA, Florida RY966 Cladonia pachycladodes USA, Florida RY966 Cladonia perforata USA, Florida RY100 Cladonia perforata USA, Florida RY100 Cladonia prostrata USA, Florida RY100 Cladonia prostrata USA, Florida RY103 Cladonia prostrata USA, Florida RY926 Cladonia prostrata USA, Florida RY1125 Cladonia subtenuis USA, North Carolina RY1126 Cladonia subtenuis USA, North Carolina RY1123 Cladonia subtenuis USA, North Carolina RY1229 Cladonia subtenuis USA, North Carolina RY1238 Cladonia pachycladodes USA, Pennsylvania RY1239 Cladonia pachycladodes USA, Florida Normore 1058 (SB2) Cladonia pacillum Canada, Manitoba Normore 1058 Cladonia furticulosa India, Uttrakhand Kotelko 1064 Cladonia achycladodes USA, Florida RY970 Cladonia achycladodes USA, Florida</td> <td>RY220 Cladonia pachycladodes USA, Florida =AY712703 RY822 Cladonia pachycladodes USA, Florida =AY712703 RY903 Cladonia pachycladodes USA, Florida =AY712703 RY966 Cladonia pachycladodes USA, Florida =AY712703 RY120 Cladonia perforata USA, Florida =AY712703 RY106 Cladonia prostrata USA, Florida =AY712703 RY1006 Cladonia prostrata USA, Florida =AY712703 RY854 Cladonia prostrata USA, Florida =AY712703 RY125 Cladonia prostrata USA, Florida =AY712703 RY1250 Cladonia subtenuis USA, North Carolina =AY712703 RY1250 Cladonia subtenuis USA, North Carolina =AY712703 RY1220 Cladonia subtenuis USA, North Carolina =AY712703 RY1223 Cladonia subtenuis USA, Florida AY712703 RY1224 Cladonia subtenuis USA, Florida AY712703 RY1225 Cladonia subtenuis USA, Florida AY712703 RY1226 Cladonia subtenuis USA, Florida<</td> <td>RY220 Cladonia pachycladodes USA, Florida =AY712703 - RY822 Cladonia pachycladodes USA, Florida =AY712703 - RY903 Cladonia pachycladodes USA, Florida =AY712703 - RY966 Cladonia pachycladodes USA, Florida =AY712703 - RY960 Cladonia pachycladodes USA, Florida =AY712703 - RY804 Cladonia prostrata USA, Florida =AY712703 - RY763 Cladonia prostrata USA, Florida =AY712703 - RY826 Cladonia prostrata USA, Florida =AY712703 - RY1200 Cladonia subtenuis USA, North Carolina =AY712703 - RY1203 Cladonia subtenuis USA, North Carolina =AY712703 - RY1203 Cladonia subtenuis USA, Virginia =AY712703 - RY1203 Cladonia subtenuis USA, Virginia =AY712703 - RY1204 Cladonia subtenuis USA, Florida AY712703 -</td> <td>RY220 Cladonia pachycladodes USA, Florida =AYT12703 - RY822 Cladonia pachycladodes USA, Florida =AYT12703 - RY963 Cladonia pachycladodes USA, Florida =AYT12703 - RY966 Cladonia pachycladodes USA, Florida =AYT12703 - RY964 Cladonia pachycladodes USA, Florida =AYT12703 - RY964 Cladonia prostrata USA, Florida =AYT12703 - RY763 Cladonia prostrata USA, Florida =AYT12703 - RY763 Cladonia prostrata USA, Florida =AYT12703 - RY1135 Cladonia subtenuis USA, Florida =AYT12703 - RY1136 Cladonia subtenuis USA, Florida =AYT12703 - RY11270 Cladonia subtenuis USA, Florida =AYT12703 - RY11280 Cladonia subtenuis USA, Florida =AYT12703 - RY1292 Cladonia subtenuis USA, Florida =AYT12703 -</td> <td>RY220 Cladonia pachycledodes USA, Florida AY712703 - - RY822 Cladonia pachycledodes USA, Florida - - - RY803 Cladonia pachycledodes USA, Florida - - - RY806 Cladonia pachycledodes USA, Florida - - - RY804 Cladonia pachycledodes USA, Florida - - - RY804 Cladonia pachycledodes USA, Florida - - - RY106 Cladonia prostrata USA, Florida - - - - RY105 Cladonia prostrata USA, Florida - - - - RY105 Cladonia subtenuis USA, North Carolina -</td>	RY220 Cladonia pachycladodes USA, Florida RY803 Cladonia pachycladodes USA, Florida RY966 Cladonia pachycladodes USA, Florida RY966 Cladonia perforata USA, Florida RY100 Cladonia perforata USA, Florida RY100 Cladonia prostrata USA, Florida RY100 Cladonia prostrata USA, Florida RY103 Cladonia prostrata USA, Florida RY926 Cladonia prostrata USA, Florida RY1125 Cladonia subtenuis USA, North Carolina RY1126 Cladonia subtenuis USA, North Carolina RY1123 Cladonia subtenuis USA, North Carolina RY1229 Cladonia subtenuis USA, North Carolina RY1238 Cladonia pachycladodes USA, Pennsylvania RY1239 Cladonia pachycladodes USA, Florida Normore 1058 (SB2) Cladonia pacillum Canada, Manitoba Normore 1058 Cladonia furticulosa India, Uttrakhand Kotelko 1064 Cladonia achycladodes USA, Florida RY970 Cladonia achycladodes USA, Florida	RY220 Cladonia pachycladodes USA, Florida =AY712703 RY822 Cladonia pachycladodes USA, Florida =AY712703 RY903 Cladonia pachycladodes USA, Florida =AY712703 RY966 Cladonia pachycladodes USA, Florida =AY712703 RY120 Cladonia perforata USA, Florida =AY712703 RY106 Cladonia prostrata USA, Florida =AY712703 RY1006 Cladonia prostrata USA, Florida =AY712703 RY854 Cladonia prostrata USA, Florida =AY712703 RY125 Cladonia prostrata USA, Florida =AY712703 RY1250 Cladonia subtenuis USA, North Carolina =AY712703 RY1250 Cladonia subtenuis USA, North Carolina =AY712703 RY1220 Cladonia subtenuis USA, North Carolina =AY712703 RY1223 Cladonia subtenuis USA, Florida AY712703 RY1224 Cladonia subtenuis USA, Florida AY712703 RY1225 Cladonia subtenuis USA, Florida AY712703 RY1226 Cladonia subtenuis USA, Florida<	RY220 Cladonia pachycladodes USA, Florida =AY712703 - RY822 Cladonia pachycladodes USA, Florida =AY712703 - RY903 Cladonia pachycladodes USA, Florida =AY712703 - RY966 Cladonia pachycladodes USA, Florida =AY712703 - RY960 Cladonia pachycladodes USA, Florida =AY712703 - RY804 Cladonia prostrata USA, Florida =AY712703 - RY763 Cladonia prostrata USA, Florida =AY712703 - RY826 Cladonia prostrata USA, Florida =AY712703 - RY1200 Cladonia subtenuis USA, North Carolina =AY712703 - RY1203 Cladonia subtenuis USA, North Carolina =AY712703 - RY1203 Cladonia subtenuis USA, Virginia =AY712703 - RY1203 Cladonia subtenuis USA, Virginia =AY712703 - RY1204 Cladonia subtenuis USA, Florida AY712703 -	RY220 Cladonia pachycladodes USA, Florida =AYT12703 - RY822 Cladonia pachycladodes USA, Florida =AYT12703 - RY963 Cladonia pachycladodes USA, Florida =AYT12703 - RY966 Cladonia pachycladodes USA, Florida =AYT12703 - RY964 Cladonia pachycladodes USA, Florida =AYT12703 - RY964 Cladonia prostrata USA, Florida =AYT12703 - RY763 Cladonia prostrata USA, Florida =AYT12703 - RY763 Cladonia prostrata USA, Florida =AYT12703 - RY1135 Cladonia subtenuis USA, Florida =AYT12703 - RY1136 Cladonia subtenuis USA, Florida =AYT12703 - RY11270 Cladonia subtenuis USA, Florida =AYT12703 - RY11280 Cladonia subtenuis USA, Florida =AYT12703 - RY1292 Cladonia subtenuis USA, Florida =AYT12703 -	RY220 Cladonia pachycledodes USA, Florida AY712703 - - RY822 Cladonia pachycledodes USA, Florida - - - RY803 Cladonia pachycledodes USA, Florida - - - RY806 Cladonia pachycledodes USA, Florida - - - RY804 Cladonia pachycledodes USA, Florida - - - RY804 Cladonia pachycledodes USA, Florida - - - RY106 Cladonia prostrata USA, Florida - - - - RY105 Cladonia prostrata USA, Florida - - - - RY105 Cladonia subtenuis USA, North Carolina -

	RY984	Cladonia prostrata	USA, Florida	=AY712704	-	-	-	• • •	1	
	RY12	Cladonia perforata	USA, Florida	=AY712704	-	-	-	-		
	RY1011	Cladonia prostrata	USA, Florida	AY712705	-	-	-	S		
	RY242	Cladonia leporina	USA, Florida	AY712708	-	-	-			
	RY981	Cladonia leporina	USA, Florida	=AY712708	-	-	-	V		
	RY650	Cladonia prostrata	USA, Florida	AY712709	-	-	-			
	RY665	Cladonia leporina	USA, Florida	=AY712709	-	-	-	W		
	RY616	Cladonia pachycladodes	USA, Florida	AY712711	-	-	-	Y		
	Nelsen 3950	Cladonia cf. bacillaris	USA, Pennsylvania	DQ229878	DQ229892	-	-			
	RY1163	Cladonia subtenuis	USA. Missouri	DQ482671	-	-	-	КК		
	RY1242	Cladonia subtenuis	USA, Virginia	DQ482672		-	-			
	RY1161	Cladonia subtenuis	USA, Arkansas	=DQ482672		-	-	FF		
	RY1240	Cladonia subtenuis	USA, Virginia	DQ482673	_	_	_	RR		
	RY1160	Cladonia subtenuis	USA, Arkansas	DQ482674				JJ		
storochlorig	s glomerata (Warér		USA, Alkalisas	DQ402074		F	F	- 33	clade 1	clade A12
P 02	Peksa 498	Diploschistes muscorum	Czech Republic, Lužické hory Mts.	AM905998	AM906026	KP318689	1_			
- 02 \G 33.85	-	Stereocaulon evolutoides	USA, Massachusetts	FJ626732	-		- AJ969633	-		
TEX 1712		Cladonia squamosa	USA, Massachusetts	AF345406	- AM906025	<u>+</u>	AJ969634	-		
TEX 895		Stereocaulon evolutoides	USA, Massachusetts	AF345406 AF345382	AM906025 AM906024		-	-		
	s irrogularis (Hildro	th & Ahmadjian) Skaloud & Peks		AF 343362	AM900024	Г		L	clade 2	clade A13
TER 1	Peksa 999	Stereocaulon pileatum	z Czech Republic, České Švýcarsko NP	AM905999	AM906028	KP318690	KP318701	[claue 2	clade ATS
EK 1 EX 2236	reksa 999		N.A.		AM906028 AM906027	KF310090	KF310701	_		
EX 2230	- Tolbot 152	Stereocaulon sp.		AF345411	DQ229889			_		
	Talbot 153 Bačkor 13	Stereocaulon botryosum	USA, Alaska	DQ229880				_		
		Cladonia arbuscula	Slovakia, Gelnica	AM906000	AM906029			-		
- (- u bla uiu	Talbot 281	Stereocaulon vesuvianum	USA, Alaska	DQ229885	DQ229888				clade 3	
EX 902	s magna Archibald (Pilophorus aciculare	LICA Weekington	A M006012	A M006044	KP318691	A 1060620		claue 5	-
P 48	- Peksa 518		USA, Washington	AM906012 FN556046	AM906041	KF310091	AJ969630	_		
P 40	T51501	Lepraria sp.*	Slovakia, Nízké Tatry Mts.		-	-	-	_		
	Ridka PA3	Psora decipiens	Germany, Bavaria	KF907669 KP318675	-	-	-	_		
- torro oblori		Cladonia sp.	Argentina, Patagonia	KF310075	<u>I</u> -	F	<u> </u>	1	clade 4	
TEX 911	s erici (Ahmadjian)	Cladonia cristatella	LICA Magazahuzatta	AF345440	AM906018	KP318692			Claue 4	-
TEX 911	-	Cladonia cristatella	USA, Massachusetts USA, Massachusetts	AF345439	AIVI900018	KF310092	- AJ969631	-		
EX 910	-		·	AF345459 AF345442	-	-	AJ969631	-		
towo oblawie	Normore 375	Stereocaulon dactylophyllum	Canada, Nova Scotia	AF343442	<u> </u> -	<u> </u> -	-		clade 5	
	s excentrica (Archic	oald) Skaloud & Peksa		AM905993	AM906019	KD249602	A 1000000	-	claue 5	-
EX 1714	-	Stereocaulon dactylophyllum	USA, Vermont	Alvi905995	AM906019	KP318693	AJ969629		clade 13	
	s italiana (Archibald	•	lándu.	4 M000004	4 140000000	KP318694	A (000000)	-	clade 13	-
CAP 219/5B	-	Xanthoria parietina	Italy	AM906001	AM906030	KP310094	AJ969632)	-		
EX 67	-	Cladonia sp.	N.A.	AF345423	DQ229894	-	-	-		
	Hammer 7212	Cladonia capitellata	Australia	AF345421	-	-	-	4		
1	Hammer 7090	Cladonia scabriuscula	Australia	AF345424	1-	1-	<u>l</u> -	L	olodo 45	
	s phycobiontica Ts		kelu Treste	A M000 400	4 1400000 40	KD240005	KD240700		clade 15	clade A1
G 26.81	- Deversu(0000	Anzina carneonivea	Italy, Trento	AM900490	AM906042	KP318695	KP318702	4		
P 07	Bayerová 3600	Lepraria neglecta	Ukraine, the East-Carpathians	AM906013	AM906044	-	-	4		
D 00	Bayerová 3606	Lepraria neglecta	Ukraine, the East-Carpathians	AM900491	AM906043	-	-	4		
P 09		Lepraria sp.	Austria, Alps Mts.	FN556025	KP318680	-	<u> -</u>	L		
	Peksa 858	Lepiana sp.								
	s sp.			Internet		1		T	clade 17	clade A4
sterochloris	s sp. Peksa 855	Lepraria rigidula	Czech Republic, Brdy Mts.	FN556031	FN556047	-	-		clade 17	clade A4
EP 09 sterochloris IP 03 sterochloris	s sp. Peksa 855 Peksa 495			FN556031 KP318676	FN556047 KP318681	-	- -		clade 17	clade A4 clade A9

Asterochloris	s sp.							-	clade A11
-	Peksa 872	Lepraria caesioalba	USA, California	FN556041	FN556050	-	-		
	Peksa 870	Lepraria sp.*	USA, California	FN556043	FN556052	-	-		
	Peksa 873	Lepraria caesioalba	USA, California	FN556042	FN556051	-	-		
sterochloris	s sp.							clade 8	-
	Peksa 796	Cladonia fimbriata	Slovakia, Nízké Tatry Mts.	FM945358	FM955674	-	-		
	Peksa 787	Cladonia rei	Slovakia, Volovské vrchy Mts.	FM945380	FM955675	-	-		
Asterochloris	s sp.							clade 9	-
	Nelsen 2233f	Pilophorus cf. cereolus	Costa Rica, San José	DQ229883	DQ229895	-	-		
	Nelsen 2181b	Stereocaulon sp.	Costa Rica, San José	DQ229884	DQ229896	-	-		
	Nelsen 2211a	Lepraria sp.	Costa Rica, San José	EU008684	EU008711	-	-		
	Ridka IH20	Cladonia scabriuscula	India, Maharashtra	HE803038	KP318682	-	-		
sterochloris	s sp.							clade 12	-
	Talbot 101	Stereocaulon paschale	USA, Alaska	DQ229887	DQ229891	-	-		
	Peksa 815	Cladonia fimbriata	Slovakia, Veľká Fatra Mts.	FM945359	FM955676	-	-		
	Peksa 921	Cladonia rei	Czech Republic, Chvaletice	FM945378	FM955677	-	-		
Asterochloris	s sp.				-		-	clade 15	-
	Talbot 400	Stereocaulon tomentosum	USA, Alaska	DQ229882	DQ229893	-	-		
sterochloris	s sp.							-	-
	Nelsen 2585	Lepraria sp.	China, Yunnan	EU008690	EU008715	-	-		-

Morphological cha	racteristics of all in	vestigated s	trains																													
			ll shape (%) [Ce	ell size (μn	n)			Plasti	d types (%)					Lobe e	xtensions (%	.)			Combinati	on of plast	id types a	nd lobe	extension	າ ຣ (%) *ຣ∈	ee explana	tions below	M	ax.	Aplanosp	ores
Species	Culture number	spherical		, pyriform	average	min.	, max.	Shallowly	Deeply			hinate FI		lobular I	Elongated	Simple		, nger-	Not		SS SFI	SFi DE			S PN		S PFi	FE FFI	FFi num	ber of 16	T T	64 128
A glamarata	UTEX 895	41	34	25	9.3		-	lobed	lobed			lob	bed	0.0		0.0		like	formed	31			13 19	0			0 13			noids		
A. glomerata A. glomerata	UTEX 095 UTEX 1712 (1)	41 31	34 37	22 21	9,3 11.4	6,8 6.7	ອົມມາມາມາມາມາມາມາມາມາມາມາມາມາມາມາມາມາມາມ	ອັດ	56,3 73,3	0,0 0,0	0,0	0,0	12,5 0.0	0,0	68,8 60,0	0,0	12,5 0.0	18,8 40,0	0,0	31 13	0 0		0 27	0		0	0 13	0 0		more -	yes ye ves ve	es yes es ves
A. glomerata	UTEX 1712 (2)	36	42	22	11,0	6,7 6,7	í	ອີກ	75,0	0,0	0.0	0,0	0,0	0,0	66,7	0,0	0,0	33,3	0,0	25	0 0	42	0 27	0	0 0	0	0 0	0 (more -	yes ye	
A. glomerata	UTEX 1712 (3)	40	31	 29	11.1	5,3	ອີກການການການການການການການການການການການການ		69,7	0,0	0,0	0,0	12,1	0.0	33,3	0,0	18,2	48,5		 12	0 6	21 2	18 30	0	0 0	0	0 0	0 12		more -	ves ve	ากการการนี้กลึงการการการการการการการการการการการการการก
A. irregularis	UTEX 2236 (1)	34	44	22	14,7	8,7	3		50,0	0,0	16,7	0,0	0,0	0,0	66,7	0,0	0,0	33,3	0,0	33	0 0	33	0 17	0	0 0	0	17 0	0 (more yes	yes -	-
A. irregularis	UTEX 2236 (2)	38	38	25	9,8	6,3	15,4	4 29,4	64,7	0,0	0,0	0,0	5,9	0,0	88,2	0,0	0,0	11,8	0,0	29	0 0	53	0 12	0	0 0	0	0 6	0 (0 0 3 01	more yes	yes -	-
A. irregularis	UTEX 2236 (3)	26	34	40	10,5	6,5	5		56,3	0,0	6,3	0,0	18,8	0,0	81,3	0,0	0,0	18,8	0,0	19	0 0) 44	0 13	0	0 0	0	6 19	0 (more yes	yes -	-
A. irregularis	STER 1	44	32	24	11,6	6,6	5	ວຸດ	45,0	0,0	0,0	0,0	0,0	0,0	50,0	0,0	5,0	45,0	0,0	25	5 25	25	0 20	0	0 0	0	0 0	0 (0 0 3 0	more -		es yes
A. magna	UTEX 902 (1)	79	14	7	9,7	3,3	ອົງການການການການການການການການການການການການການກ	ົດ	0,0	0,0	58,8	0,0	0,0	41,2	0,0	0,0	0,0	58,8	41,2	0	0 0	0		0	41 0	0	59 0	0 (0 -		es yes
A. magna A. erici	UTEX 902 (2) UTEX 911 (1)	79 95	15	0	7,5 9,2	4,7 4.7	5	ອົດການການການການການການການການການການການການການອົງກາ	0,0 0,0	0,0 26,3	41,7 0.0	0,0 0,0	0,0 0,0	58,3 0.0	0,0 52,6	0,0 26,3	0,0 0,0	41,7 21,1	58,3 0 0	0 53	0 0			26		0	42 0	0 (0- 1 yes	- ye	es yes
A. erici	UTEX 911 (2)	93 92	5 8	0	9,2 8.3	4,7 5.0			0,0	42,9	0,0	0,0	0,0	0,0	52,0 57,1	42,9	0,0	0,0	0,0	57	0 21	0	0 0	43	0 0	0	0 0	0 (1 yes		_
A. excentrica	UTEX 1714 (1)	92	6	2	12,9	6,9	5	ແມ່ນ ແມ່ນ ແມ່ນ ແມ່ນ ແມ່ນ ແມ່ນ ແມ່ນ ແມ່ນ	76,7	0,0	0.0	0.0	0,0	0.0	53,5	0.0	9,3	37.2	0,0	16	2 5	37	7 33	0	0 0	0	0 0	0 (0 0	2 -	- ve	S -
A. excentrica	UTEX 1714 (2)	92	6	2	10,4	5,6	§	ອູດ	62,1	0,0	0,0	0,0	0,0	0,0	69,0	0,0	20,7	10,3		17	10 10	52 <i>ć</i>	10 0	0	0 0	Ũ	0 0	0 (0 0	2 -	- ye	แก้แกกกญี้แกกการการการการการการการการการการการการกา
A. excentrica	UTEX 1714 (3)	91	7	2	8,8	5,0	5		50,0	0,0	0,0	0,0	0,0	0,0	63,0	0,0	17,4	19,6	0,0	39	4 7	' 24 <i>'</i>	13 13	0	0 0	0	0 0	0 (0 0	2 -		s -
	CAUP H1010 (1)	90	6	4	18,0	7,8	28,3		25,9	11,1	0,0	0,0	0,0	0,0	40,7	11,1	11,1	37,0	0,0	33	11 19		0 19	11	0 0	0	0 0	0 (0 0 3 0	more -	- ye	es yes
	CAUP H1010 (2)	82	14	4	12,6	6,0	ā		4,3	17,4	0,0	0,0	0,0	0,0	56,5	17,4	4,3	21,7		52	4 22	2 4	0 0	17	0 0	0	0 0	0 (more -	ากกฏิ์การการการการการกฏิ์ที่การกา	es yes
	CAUP H1010 (3)	86	6	7	10,9	5,0	=	ມມ ີ ແມ່ນນັ້ນມາມັນແມ່ນມະນາມັນແມ່ນມະນາມັນແມ່ນນັ້ນມາມ⊇ິມ	21,1	21,1	10,5	0,0	0,0	0,0	39,5	21,1	5,3	34,2		32	5 11	8	0 13	21	0 0	0	11 0	0 (more -	າມມາມັນແມ່ນແມ່ນແມ່ນມາມັນນັ້ນແມ່ນ	es yes
A. leprarii	LEP 25 LEP 30	92 61	8 35	1	13,3	5,6	ອື່າການການການການການການການການການການການການການ	ອົດການການການການການການການການການການການກໍ່ມານກະອົນ	14,8	55,6	0,0 15.4	0,0	0,0	0,0 0,0	22,2	55,6 52,8	3,7 0.0	18,5	0,0	15 15	4 11 0 15			56 54		0	0 0 15 0	0 0		more -	កកម្មិតកម្មនេះការបានបានដំបូងការក	es yes
A. leprarii A. gaertneri	CAUP H1013 (1)	96	აე 2	2 2	10,8 17,7	5,4 5,9	⊜		0,0 32,1	53,8 35,7	15,4 0.0	0,0 0,0	0,0 0,0	0,0 0,0	15,4 53,6	53,8 35,7	0,0 1,8	30,8 8,9	0,0	15 32	0 15	21	2 9	34 36		0				more -	- ye yes ye	es yes es ves
	CAUP H1013 (2)	88	ے 4	2 8	17,7	5,9 5,8	3	ອັ ດແມ່ນແມ່ນແມ່ນແມ່ນແມ່ນແມ່ນແມ່ນແມ່ນ ແມ່ນນັ້ນແມ່ນ≣ີນ	20,3	13,6	0,0	47,5	0,0	0,0	30,5	61,0	1,0 1,7	6,8	0,0	19	0 0	12	2 7	14	0 0	0	0 0	0 (more -	yes ye	างการนี้เกิดการการการการการการการการการการการการการก
	CAUP H1013 (3)	94	5	2	12,3	5,5	5		21,0	0,0	0.0	0.0	0,0	0.0	54,8	0.0	8.1	37.1	0,0	44	6 29	11	2 8	0	0 0	0	0 0	0 (more -	ves ve	es ves
	CAUP H1009 (1)	78	14	8	14,2	7,2	3111111111111111111111111111111111111	ວິດ	80,0	0,0	3,3	0,0	0,0	0,0	66,7	0,0	0,0	33,3	0,0	17	0 0	50	0 30	0	0 0	0	3 0	0 (more -		yes
A. woessiae	CAUP H1009 (2)	74	20	6	14,9	8,8	23,9	9 24,4	70,7	0,0	4,9	0,0	0,0	0,0	46,3	0,0	0,0	53,7	0,0	12	0 12	. 34	0 37	0	0 0	0	50	0 (0 0 3 01	more -		yes
A. woessiae	CAUP H1009 (3)	69	19	11	15,4	9,0	ອື່າການການການການການການການການການການການການການ	ອັດການການການການການການການການການການການກໍ່ມານເອັດ	53,7	0,0	7,3	0,0	14,6	0,0	46,3	0,0	14,6	39,0	0,0	12	0 12	. 34	0 20	0	0 0	0	7 0	15 (more -		yes
A. woessiae	LEP 15 (1)	87	11	2	12,3	6,5	§		64,5	0,0	0,0	0,0	0,0	0,0	74,2	0,0	16,1	9,7		29	0 6	i 45 ´	16 3	0	0 0	0	0 0	0 (more -		yes
A. woessiae	LEP 15 (2)	87 98	11	2	10,7	5,6	í		58,6	0,0	0,0	0,0	27,6	0,0	46,6	0,0	39,7	13,8		12	0 2	34 2	12 12	0	0 0	0	0 0	28 (more -		yes
A. woessiae A. woessiae	LEP 34 (1) LEP 34 (2)	98 99	U 1	2	10,5 10,5	5,0 5,4	≣	0 50,0 1 28,6	37,5 42,9	0,0 0,0	0,0 0.0	0,0	12,5 28,6	0,0	87,5 85,7	0,0 0.0	12,5 14,3	0,0 0.0	0,0	50 29		43		0		0	0 0	13 (14 (more -		yes yes
A. woessiae	LEP 36 (1)	92	3	5	14,6	5, 4 6.8	⊜		42,5 54,8	12,9	6.5	0,0	3.2	0,0	32,3	12,9	14,3	38,7	σ,σ	23 16	0 0		0 0 13 26	13	0 0	0	6 0	3 (more -		ves
	LEP 36 (2)	96	4	0	12,9	6,2			40,0	26,7	0,0	0,0	0,0	0,0	53,3	26,7	20,0	0,0		33	0 0		20 0	27	0 0	0	0 0	0 (more -		ves
	CAUP H1011 (1)	92	6	2	12,9	6,0			35,0	0,0	10,0	0,0	0,0	0,0	70,0	0,0	5,0	25,0		50	5 0	20	0 15	0	0 0	0	10 0	0 (0 0	1 -	- ye	es yes
A. friedlii	CAUP H1011 (2)	90	10	0	10,3	4,8	16,6	6 36,7	33,3	30,0	0,0	0,0	0,0	0,0	60,0	30,0	0,0	10,0	0,0	37	0 0	23	0 10	30	0 0	0	0 0	0 (0 0	1 -	- ye	es yes
	CAUP H1011 (3)	94	6	0	11,6	6,5	and the second s		34,4	9,4	12,5	0,0	0,0	0,0	56,3	9,4	9,4	25,0	9	34	9 0	22	0 13	9	0 0	0	13 0	0 (0 0	1 -	ากกฏิสาวการการการการกู้สำนาณ	es yes
	LEP 4	87	13	1	12,7	4,4			20,8	16,7	12,5	0,0	0,0	0,0	50,0	16,7	4,2	29,2		46	4 0	4	0 17	17	0 0	0	13 0	0 (0 0	1 -	ការស្ថិតការការការការស្ថិតិការការ	es yes
	CCAP 219/5B (1)	94	5	1	8,7	4,8	.=		0,0	0,0	0,0	0,0	0,0	0,0	80,0	0,0	0,0	20,0		80	0 20			0		0	0 0	0 0	····	more -	yes -	-
	CCAP 219/5B (2) UTEX 67 (1)	93 44	8 37	10	9,1 13,1	5,6 5,7	⊜		0,0 0,0	0,0 0,0	0,0 0,0	0,0 0,0	0,0 0,0	0,0 0,0	75,0 87,5	0,0 0,0	0,0 0,0	25,0 12,5		75 88	0 25 0 13	u§		0		0		0 0		more -	yes -	- s -
B	UTEX 67 (1)	44 56	25	19 19	10,1 10,1	5,7 5,5	B	≣	0,0	0,0	0,0	0,0	0,0	0,0	87,5 100,0	0,0 0,0	0,0	۲۷,5 0,0		00 100	0 13	0	0 0	0	0 0	0	0 0	0 (more -		- ss
	CAUP H1012 (1)	100	0	0	9,6	<u>5,5</u> 4.9	5		0,0	76,2	0,0	23,8	0,0	0,0	0,0	100,0	0,0	0,0		0	0 0	0	0 0	76	0 0	0	0 0	0 (more -		s yes
	CAUP H1012 (2)	100	0	0	12,8	7,0	Buuuuuuuuuuuuuuuuuuuuuuuuuuuuuuuuuuu	ອີກ ຄວາມການການການການການການການການການການການການການກ	0,0	34,6	0,0	65,4	0,0	0,0	0,0	100,0	0,0	0,0		0	0 0	0	0 0	35	0 0	0	0 0	0 (more -	ការស្ថិតតារបានការបានស្ថិតិភាពក	es yes
A. phycobiontica	ເວັ້ນແມ່ນການການການການການການການການການການການການການ	83	4	14	11,8	5,0	20,9		0,0	44,3	55,7	0,0	0,0	0,0	0,0	52,9	0,0	47,1		0	0 0	0	0 0	44	0 0	9	47 0	0 (0 0 3 01	more -	ការស្ថិតតារបានការបានស្ថិតិតារបា	es yes
A. phycobiontica	ເຈັບບານການການການການການການການການການການການການກາ	89	8	3	11,9	6,2	2		0,0	26,8	44,6	0,0	0,0	0,0	0,0	48,2	0,0	51,8		0	0 29	0	0 0	27	0 0	21	23 0	0 (0 0 3 01	more -	านแล้วและและและและเกิดไม่และ	es yes
A. phycobiontica	())	95	4	1	12,2	5,6	2		0,0	8,8	91,2	0,0	0,0	0,0	14,7	14,7	0,0	70,6	9	0	0 0	0	0 0	9	0 15		71 0	0 (more -	កកម្មរកកម្មរកកម្មវិភាពកម្មវិភ័យការ	es yes
A. phycobiontica		91	9	0	16,1	7,0			0,0	0,0	89,6	0,0	0,0	0,0	26,9	6,0	0,0	67,2		0	0 10		0 0	0	0 27		57 0	0 (more -		es yes
A. phycobiontica	ເອັ້ນແມ່ນແມ່ນແມ່ນແມ່ນແມ່ນນັ້ນແມ່ນແມ່ນແມ່ນແມ່ນແມ່ນແມ່ນແມ່ນເມັນ	73 71	27	0	13,8	7,4 5.4	B		0,0	22,9	62,9 52.0	0,0	0,0	0,0	0,0	34,3	0,0	65,7 76 5		0	0 14			23			51 0	0 (more -	າມມີການການການການການຫຼືກັນການ	es yes
A. phycobiontica A. lobophora	LEP 7 (2) LEP 1 (1)	71 93	28 6	1	12,6 14,3	5,4 6 3	3	∋u	0,0	23,5 1/1 3	52,9 28.6	0,0	0,0	0,0 0,0	0,0 14,3	23,5 14 3	0,0	76,5 71,4	9	0 14	0 24 0 43	ມອື່ມມາມາມາມາມາມມາມອີການການການ		24 14		0	53 0 29 0	0 0		more -	ការប្រំការការការការការប្រើការការ	es yes es yes
	LEP 1 (1) LEP 1 (2)	93 87	9	ı 4	14,3	6,3 6,2			0,0 0,0	14,3 55,6	28,6 0,0	0,0 0,0	0,0 0,0	0,0	22,2	14,3 55,6	0,0 0,0	22,2		22	0 43		0 0	56	0 0	0	0 0	0 (1 -		es yes es yes
	CAUP H1014 (1)	85		- 2	13,0	0,2 9,9	3		0,0	0,0	7,1	0,0	0,0	0,0	71,4	0,0	7,1	21,4		71	7 14		0 0	0	0 0	0	7 0	0 (0 0	1 -	ນັ້ນ	s yes
	CAUP H1014 (2)	86	14	- 0	17,1	<u>8,5</u>	3		0,0	11,8	17,6	0,0	0,0	0,0	47,1	11,8	17,6	23,5		47	18 6	Ŭ 0	0 0	12	0 0	Ũ	18 O		0 0	1 -	លលក្ខណៈពេលពេលពេលក្រុកិណៈព	es yes
B	LEP 2	92	8	0	12,0	5,9	ā		0,0	14,3	28,6	0,0	0,0	0,0	14,3	14,3	0,0	71,4		·····ē	0 43	0	0 0	14	0 0	0	29 0	0 (0 0	1 -		es yes
Morphological cha	racteristics of parti	cular Asterc	ochloris spec	cies	- -										•		-															

	Species Culture number										Plast	id types ('	%)				Lobe	extensions	s (%)			Combin	ation o	ⁱ plastid	types ar	nd lobe e	extensio	ons (%) *	see exp	lanations	below		Max.	Aplanospores	
Species	Culture number	spherica	oval	pyrifo	rm a	average	min.	max.	Shallowly lobed	Deeply lobed	Crenulate	Parietal	Echinate	Flat lobed	Globular	Elongated	Simple	Flat	Finger- like	Not formed	SE	ss s	FI SF	i DE	DFI	DFi CS	6 PN	PE	PS F	PFi FE	FFI		number of pyrenoids	6 32	64 128
A. glomerata	-	37	1	36	27	10,8	5,3	22,9	9 25,3	68,6	0,0	0,0	0,0	6,2	0,0	57,2	0,0	7,7	35,1	0,0	20	0	5 3	4 8	27	0	0 (0 0	0	3	0 3	0	3 or more -	yes	yes yes
A. irregularis	-	35	j (37	28	11,7	6,3	26,3	3 34,1	54,0	0,0	5,7	0,0	6,2	0,0	71,5	0,0	1,3	27,2	0,0	27	1	6 3	9 0	15	0	0 (0 0	6	6	0 0	0	3 or more ye	s yes	
A. magna	-	79)	4	7	10,1	3,3	22,9	9,0	0,0	0,0	50,2	0,0	0,0	49,8	0,0	0,0	0,0	50,2	49,8	0	0	0	0 0	0	0 5	50 (0 0	50	0	0 0	0	0 -	-	yes yes
A. erici	-	94		7	0	8,8	4,7	15,8	8 65,4	0,0	34,6	0,0	0,0	0,0	0,0	54,9	34,6	0,0	10,5	0,0	55	0	11	0 0	0	35	0 (0 0	0	0	0 0	0	1 ye	S -	
A. excentrica	-	92	?	6	2	10,6	5,0	22,8	3 37,1	62,9	0,0	0,0	0,0	0,0	0,0	61,8	0,0	15,8	22,4	0,0	24	6	7 3	8 10	15	0	0 (0 0	0	0	0 0	0	2 -	-	yes -
A. leprarii	-	82	?	4	4	12,7	0,5	28,3	3 49,8	13,2	31,8	5,2	0,0	0,0	0,0	34,9	31,8	4,9	28,5	0,0	29	5	15	5 0	8	32	0 (0 0	5	0	0 0	0	3 or more -	-	yes yes
A. gaertneri	-	93	}	4	4	14,0	5,5	29,4	43,3	24,5	16,4	0,0	15,8	0,0	0,0	46,3	32,2	3,8	17,6	0,0	31	2	10 1	5 2	8	16	0 (0 0	0	0	0 0	16	3 or more -	yes	yes yes
A. woessiae	-	87	1	9	4	12,6	5,0	25,4	1 27,7	55,9	4,4	2,4	0,0	9,6	0,0	59,9	4,4	14,8	20,9	0,0	23	0	4 3	5 7	14	4	0 (0 0	2	2	8 0	0	3 or more -	-	- yes
A. friedlii	-	91		9	1	12,1	4,4	21,0) 46,4	30,9	14,0	8,8	0,0	0,0	0,0	59,1	14,0	4,6	22,3	0,0	42	5	0 1	7 0	14	14	0 (0 0	9	0	0 0	0	1 -	-	yes yes
A. italiana	-	72	2	19	10	10,2	4,8	22,5	5 100,0	0,0	0,0	0,0	0,0	0,0	0,0	85,6	0,0	0,0	14,4	0,0	86	0	14	0 0	0	0	0 (0 0	0	0	0 0	0	3 or more -	yes	yes -
A. echinata	-	100)	0	0	11,4	4,9	21,0	0,0	0,0	55,4	0,0	44,6	0,0	0,0	0,0	100,0	0,0	0,0	0,0	0	0	0	0 0	0	55	0 (0 0	0	0	0 0	45	3 or more -	-	yes yes
A. phycobiontica	-	84	ŀ	13	3	13,0	5,0	24,3	3 12,8	0,0	21,0	66,1	0,0	0,0	0,0	6,9	29,9	0,0	63,1	0,0	0	0	13	0 0	0	21	0	7 9	50	0	0 0	0	3 or more -	-	yes yes
A. lobophora	-	89)	0	2	13,7	5,9	25,4	1 64,4	0,0	19,2	16,4	0,0	0,0	0,0	33,9	19,2	5,0	42,0	0,0	34	5	26	0 0	0	19	0 (0 0	16	0	0 0	0	1 -	-	yes yes

* Explanation of abbreviations:

SE Shallowly lobed plastid, elongated lobe extensions
SFI Shallowly lobed plastid, flat lobe extensions
SFi Shallowly lobed plastid, finger-like lobe extensions
DE Deeply lobed plastid, elongated lobe extensions

DFI DFi

Deeply lobed plastid, flat lobe extensions Deeply lobed plastid, finger-like lobe extensions Crenulate plastid, simple lobe extensions Globular plastid, extensions not formed

CS GN

PE Parietal plastid, elongated lobe extensions
PS Parietal plastid, simple lobe extensions
PFi Parietal plastid, finger-like lobe extensions
FE Flat lobed plastid, elongated lobe extensions

FFI Flat lobed plastid, flat lobe extensionsFFi Flat lobed plastid, finger-like lobe extensionsES Echinate plastid, simple lobe extensions