

A Contribution to the Phylogeny and Taxonomy of *Phlebiopsis* (Polyporales, Basidiomycota)

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Research

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Abstract

An in-depth study of the phylogeny and taxonomy of the corticioid genus *Phlebiopsis* (Phanerochaetaceae) was conducted. Phylogenetic analyses of the ITS1-5.8S-ITS2 and nrLSU sequences demonstrated that *Phlebiopsis* is a strongly supported clade that is distinct from its sister clade that includes *Phaeophlebiopsis*, *Hapalopilus* and *Rhizochaete*. Two genera, *Australohydnum* and *Hjortstamia*, are reduced to synonyms under *Phlebiopsis* because generic type species *A. griseofuscescens* and *H. friesii*, respectively, are embedded in the *Phlebiopsis* clade. Twenty-four lineages are resolved in the ITS phylogenetic tree of *Phlebiopsis*, including six new taxa from Sri Lanka and China – *P. albescens*, *P. brunnea*, *P. cylindrospora*, *P. magnicystidiata*, *P. membranacea* and *P. sinensis*. Five new combinations are proposed – *Phaeophlebiopsis mussooriensis*, *Phlebiopsis bambusicola*, *P. dregeana*, *P. griseofuscescens* and *P. novae-granatae*. *Phlebiopsis crassa* is a morphological species complex with three distinct lineages. *Phlebiopsis lamprocystidiata* is determined to be a later synonym of *P. darjeelingensis*. The new taxa are described, illustrated, and compared and contrasted to morphologically similar species. An emended description of *Phlebiopsis* is provided along with an identification key to 27 accepted species.

Introduction

In 1978, the genus *Phlebiopsis* Jülich was erected for *Thelephora gigantea* Fr. that has effused, ceraceous basidiomata with a smooth to odontoid hymenophore, a monomitic hyphal system with hyaline, partially agglutinated, simple-septate hyphae, lamprocystidia with thick, hyaline walls, and basidiospores with hyaline, thin, smooth walls that do not react in Melzer's reagent or cotton blue (Jülich 1978; Bernicchia and Gorjón 2010). Over the next 40 years, 12 species with similar morphology were described in or transferred to the genus by Hjortstam and Ryvarden (1980), Jülich and Stalpers (1980), Dhingra (1987), Hjortstam (1987), Gilbertson & Adaskaveg (1993), Douanla-Meli and Langer (2009), Wu et al. (2010), Priyanka et al. (2011), Kaur et al. (2015) and Zhao et al. (2018). Morphologically, *Phlebiopsis* is similar to *Scopuloides* (Masse) Höhn. & Litsch. and some species of *Phanerochaete* P. Karst. It was traditionally placed in the *Phanerochaete sensu lato* group (Rattan 1977; Burdsall 1985). Although Burdsall (1985) considered *Phlebiopsis* and *Scopuloides* to be synonyms of *Phanerochaete*, most researchers at the time recognized the genera as distinct (Eriksson et al. 1978, 1981, 1984; Jülich and Stalpers 1980; Wu 1990).

The generic circumscription of *Phlebiopsis* was expanded when molecular studies showed that *Phanerochaete crassa* (Lév.) Burds. and *Phlebiopsis gigantea* (Fr.) Jülich were closely related (de Koker et al. 2003; Greslebin et al. 2004; Wu et al. 2010; Floudas and Hibbett 2015). With the inclusion of *P. crassa*, *Phlebiopsis* now also includes species with effused-reflexed, coriaceous basidiomata, a dimitic hyphal system, and lamprocystidia or skeletocystidia with light brown walls. In addition, Floudas and Hibbett (2015) created *Phaeophlebiopsis* Floudas & Hibbett to accommodate *Phlebiopsis peniophoroides* Gilb. & Adask. and similar species with ceraceous, beige-brown basidiomata and subicula, lamprocystidia with brown walls, and small basidiospores. The limits of the *Phlebiopsis* clade were extended by Miettinen et al. (2016) who transferred six species into *Phlebiopsis*. The results of their phylogenetic study showed

that the type species of *Castanoporus* Ryvar den, *Merulius castaneus* Lloyd, was nested in a clade with *P. gigantea* and, therefore, a synonym of *Phlebiopsis*. Similarly, *Dentocorticium pilatii* (Parmasto) Duehm & Michel, *Lopharia papyrina* (Mont.) Boidin, *Phanerochaete brunneocystidiata* Sheng H. Wu, and *P. laxa* Sheng H. Wu clustered in the *Phlebiopsis* clade. Based on the morphological similarity of *Thelephora friesii* Lév., the type of *Hjortstamia* Boidin & Gilles, to *L. papyrina* and *P. crassa*, they also transferred *T. friesii* to *Phlebiopsis*, thereby reducing *Hjortstamia* to a synonym of *Phlebiopsis*. Earlier, Boidin and Gilles (2003) erected the genus *Hjortstamia* for *T. friesii* and related taxa with effused-reflexed to pileate basidiomata, a dimitic hyphal system of simple-septate hyphae and thick-walled skeletal hyphae, lamprocystidia with thick walls. Ryvar den (1991) proposed *Castanoporus* for *M. castaneus*, a resupinate species with yellowish brown to purple pores, a monomitic hyphal system with simple-septate hyphae, and lamprocystidia with thick, hyaline walls. *Phlebiopsis pilatii* (Parmasto) Spirin & Miettinen is unique in the genus for it has a dimitic hyphal system of simple-septate generative and microbinding (squeletto-ligatives) hyphae and finely branched hyphidia but lacks lamprocystidia or skeletal cystidia (Larsen and Gilbertson 1977; Duhem and Michel 2009).

With *Hjortstamia* and *Castanoporus* as synonyms, *Phlebiopsis* became a morphologically heterogeneous genus with effused, effused-reflexed or pileate basidiomata with a membranous, ceraceous, corneous or coriaceous texture, hymenophore smooth to tuberculate, odontoid, or poroid, hyphal system monomitic or dimitic with a loose to compact subiculum, and typically with lamprocystidia or skeletocystidia with hyaline to brown walls. In phylogenetic analyses of Phanerochaetaceae, *Phlebiopsis* species are in a clade sister to *Rhizochaete* Gresl., Nakasone & Rajchenb., *Hapalopilus* P. Karst. and *Phaeophlebiopsis*, but distant from *Phanerochaete sensu stricto* and *Scopuloides* (Floudas and Hibbett 2015; Miettinen et al. 2016).

Another genus of interest is *Australohydnum* Jülich for it is similar to *Phlebiopsis* by its warted, irpicoid to hydroid hymenophore, a dimitic hyphal system with hyaline, encrusted skeletocystidia, and thin-walled, smooth basidiospores (Jülich 1978). The morphological similarities between *Australohydnum* and *Phanerochaete s.l.* were observed by Hjortstam and Ryvar den (1990). In a limited study of *Irpex s.s.*, sequences of *Australohydnum dregeanum* (Berk.) Hjortstam & Ryvar den and *Irpex vellereus* Berk. & Broome (a possible synonym of *A. dregeanum*) clustered together in a clade sister to two *Phanerochaete* species (Lim and Jung 2003). However, the phylogenetic relationship of *Australohydnum* within the Phanerochaetaceae remained unknown (Miettinen et al. 2016).

Among the 24 names of *Phlebiopsis* recovered in Index of Fungorum (<http://www.indexfungorum.org/>, accessed 21 June 2020), four species were transferred to *Phaeophlebiopsis*. Of the remaining 20 species, 11 were described originally from Asia (e.g., Dhingra 1987; Wu 2000, 2004; Priyanka et al. 2011; Kaur et al. 2015; Zhao et al. 2018; Xu et al. 2020). More than 150 specimens of *Phlebiopsis* were collected by the corresponding author from China and Southeast Asia in recent years. Based on these specimens and sequences obtained from GenBank, a phylogenetic analysis and taxonomic study of *Phlebiopsis* and related taxa in the Phanerochaetaceae were undertaken. This study is a contribution to the understanding of the diversity and phylogenetic relationships of crust fungi in China.

Materials And Methods

Morphological studies

Voucher specimens are deposited at the herbaria of Beijing Forestry University, Beijing, China (BJFC), Centre for Forest Mycology Research, U.S. Forest Service, Madison, Wisconsin, U.S.A. (CFMR) and National Museum of Natural Science, Taichung, Taiwan, China (TNM). Freehand sections were made from dried basidiomata and mounted in 2% (w/v) potassium hydroxide (KOH), 1% (w/v) phloxine, Melzer's reagent (IKI) or cotton blue (CB). Microscopic examinations were carried out with a Nikon Eclipse 80i microscope (Nikon Corporation, Japan) at magnifications up to 1000 ×. Drawings were made with the aid of a drawing tube. The following abbreviations are used: IKI– = neither amyloid nor dextrinoid, CB– = acyanophilous, L = mean spore length, W = mean spore width, Q = L/W ratio, n (a/b) = number of spores (a) measured from number of specimens (b). Color codes and names follow Kornerup and Wanscher (1978).

DNA extraction and sequencing

A CTAB plant genomic DNA extraction Kit DN14 (Aidlab Biotechnologies Co., Ltd, Beijing, China) was used to extract total genomic DNA from dried specimens then amplified by the polymerase chain reaction (PCR), according to the manufacturer's instructions. The ITS1-5.8S-ITS2 region was amplified with the primer pair ITS5/ITS4 (White et al. 1990) using the following protocol: initial denaturation at 95 °C for 4 min, followed by 34 cycles at 94 °C for 40 s, 58 °C for 45 s and 72 °C for 1 min, and final extension at 72 °C for 10 min. The nrLSU D1-D2 region was amplified with the primer pair LR0R/LR7 (<http://www.biology.duke.edu/fungi/mycolab/primers.htm>) employing the following procedure: initial denaturation at 94 °C for 1 min, followed by 34 cycles at 94 °C for 30 s, 50 °C for 1 min and 72 °C for 1.5 min, and final extension at 72 °C for 10 min. DNA sequencing was performed at Beijing Genomics Institute, and the sequences were deposited in GenBank (Table 1). BioEdit v.7.0.5.3 (Hall 1999) and Geneious Basic v.11.1.15 (Kearse et al. 2012) were used to review the chromatograms and for contig assembly.

Table 1

Species and sequences used in the phylogenetic analyses. New species are set in bold with type specimens indicated with an asterisk (*).

Taxa	Voucher	Locality	ITS	nrLSU	Reference
<i>Bjerkandera adusta</i>	HHB-12826-Sp	USA	KP134983	KP135198	Floudas & Hibbett 2015
<i>B. centroamericana</i>	L-13104-sp	Costa Rica	KY948791	KY948855	Justo et al. 2017
<i>Crepatura ellipsospora</i>	CLZhao 1265	China	MK343692	MK343696	Ma & Zhao 2019
<i>Donkia pulcherrima</i>	GC 1707-11	China	LC378994	LC379152	Chen et al. 2018a
<i>Geliporus exilisporus</i>	Dai 2172	China	KU598211	KU598216	Yuan et al. 2017
<i>Hapalopilus eupatorii</i>	Dammrich 10744	Germany	KX752620	KX752620	Miettinen et al. 2016
<i>H. percoctus</i>	Miettinen 2008	Botswana	KX752597	KX752597	Miettinen et al. 2016
<i>H. nidulans</i>	JV0206/2	Sweden	KX752623	KX752623	Miettinen et al. 2016
<i>Hyphodermella corrugata</i>	MA-Fungi 5527	Morocco	FN600372	JN939597	Telleria et al. 2010
<i>H. poroides</i>	Dai 10848	China	KX008368	KX011853	Zhao et al. 2017
<i>H. rosae</i>	FP-150552	USA	KP134978	KP135223	Floudas & Hibbett 2015
<i>Irpex vellereus</i>	CBS 515.92	India	AF479670	—	Lim & Jung 2003
<i>Odontoefibula orientalis</i>	GC 1703-76	China	LC379004	LC379156	Chen et al. 2018a
<i>Oxychaete cervinogilvus</i>	Schigel-5216	Australia	KX752596	KX752596	Miettinen et al. 2016
<i>Phaeophlebiopsis caribbeana</i>	HHB-6990	USA	KP135415	KP135243	Floudas & Hibbett 2015
<i>P. himalayensis</i>	He 3854	China	MT386378	MT447410	present study
<i>P. peniophoroides</i>	FP-150577	USA	KP135417	KP135273	Floudas & Hibbett 2015
<i>P. ravenelii</i>	CBS 411.50	France	MH856691	MH868208	Vu et al. 2019

Taxa	Voucher	Locality	ITS	nrLSU	Reference
<i>P. ravenelii</i>	FCUG 2216	France	—	GQ470674	Wu et al. 2010
<i>Phanerina mellea</i>	Miettinen 11393	Indonesia	KX752602	KX752602	Miettinen et al. 2016
<i>Phanerochaete arizonica</i>	RLG-10248-Sp	USA	KP135170	KP135239	Floudas & Hibbett 2015
<i>P. australis</i>	HHB-7105-Sp	USA	KP135081	KP135240	Floudas & Hibbett 2015
<i>P. bambusicola</i>	Wu 0707-2	China	MF399404	MF399395	Wu et al. 2018a
<i>P. brunnea</i>	He 1873	China	KX212220	KX212224	Liu & He 2016
<i>P. burtii</i>	HHB-4618-Sp	USA	KP135117	KP135241	Floudas & Hibbett 2015
<i>P. canobrunnea</i>	CHWC 1506-66	China	LC412095	LC412104	Wu et al. 2018b
<i>P. carnosa</i>	HHB-9195	USA	KP135129	KP135242	Floudas & Hibbett 2015
<i>P. chrysosporium</i>	HHB-6251-Sp	USA	KP135094	KP135246	Floudas & Hibbett 2015
<i>P. citrinosanguinea</i>	FP-105385-Sp	USA	KP135100	KP135234	Floudas & Hibbett 2015
<i>P. concrescens</i>	Spirin 7322	Russia	KP994380	KP994382	Volobuev et al. 2015
<i>P. cumulodentata</i>	LE 298935	Russia	KP994359	KP994386	Volobuev et al. 2015
<i>P. cystidiata</i>	Wu 1708 - 326	China	LC412097	LC412100	Wu et al. 2018b
<i>P. ericina</i>	HHB-2288	USA	KP135167	KP135247	Floudas & Hibbett 2015
<i>P. incarnata</i>	WEI 16-075	China	MF399406	MF399397	Wu et al. 2018a
<i>P. inflata</i>	Dai 10376	China	JX623929	JX644062	Jia et al. 2014
<i>P. laevis</i>	HHB-15519	USA	KP135149	KP135249	Floudas & Hibbett 2015
<i>P. livescens</i>	FD-106	USA	KP135070	KP135253	Floudas & Hibbett 2015
<i>P. magnoliae</i>	HHB-9829-Sp	USA	KP135089	KP135237	Floudas & Hibbett 2015

Taxa	Voucher	Locality	ITS	nrLSU	Reference
<i>P. porostereoides</i>	He 1902	China	KX212217	KX212221	Liu & He 2016
<i>P. pseudomagnoliae</i>	PP-25	South Africa	KP135091	KP135250	Floudas & Hibbett 2015
<i>P. pseudosanguinea</i>	FD-244	USA	KP135098	KP135251	Floudas & Hibbett 2015
<i>P. rhodella</i>	FD-18	USA	KP135187	KP135258	Floudas & Hibbett 2015
<i>P. robusta</i>	Wu 1109-69	China	MF399409	MF399400	Wu et al. 2018a
<i>P. sanguinea</i>	HHB-7524	USA	KP135101	KP135244	Floudas & Hibbett 2015
<i>P. sanguineocarnosa</i>	FD-359	USA	KP135122	KP135245	Floudas & Hibbett 2015
<i>P. sordida</i>	FD-241	USA	KP135136	KP135252	Floudas & Hibbett 2015
<i>P. stereoides</i>	He 2309	China	KX212219	KX212223	Liu & He 2016
<i>P. subceracea</i>	FP-105974-R	USA	KP135162	KP135255	Floudas & Hibbett 2015
<i>P. taiwaniana</i>	Wu 0112 - 13	China	MF399412	MF399403	Wu et al. 2018a
<i>P. velutina</i>	Kotiranta 25567	Russia	KP994354	KP994387	Volobuev et al. 2015
<i>Phlebia firma</i>	Edman K268	Sweden	EU118654	EU118654	Larsson 2007
<i>P. lilascens</i>	FCUG 2005	—	AF141622	AF141622	—
<i>Phlebiopsis albobadia</i>	He 5805 *	Sri Lanka	MT452526	—	present study
<i>P. amethystea</i>	URM 93248	Brazil	MK993644	MK993638	Xavier de Lima et al. 2020
<i>P. amethystea</i>	URM 84741	Brazil	MK993645	MK993639	Xavier de Lima et al. 2020
<i>P. brunnea</i>	He 5822 *	Sri Lanka	MT452527	MT447451	present study
<i>P. brunneocystidiata</i>	Chen 666	China	MT561707	GQ470640	Wu et al. 2010, present study
<i>P. brunneocystidiata</i>	Chen 1143	China	—	GQ470639	Wu et al. 2010
<i>P. castanea</i>	Spirin-5295	Russia	KX752610	KX752610	Miettinen et al. 2016

Taxa	Voucher	Locality	ITS	nrLSU	Reference
<i>P. castanea</i>	GC 1612-6	China	KY688208	—	Chen et al. 2018b
<i>P. castanea</i>	CLZhao 3501	China	MK269230	—	—
<i>P. castanea</i>	He 2489	China	—	MT447406	present study
<i>P. crassa</i> group A	He 5205	Vietnam	MT452523	MT447448	present study
<i>P. crassa</i> group A	He 5763	Sri Lanka	MT452524	MT447449	present study
<i>P. crassa</i> group A	He 5855	China	MT452525	MT447450	present study
<i>P. crassa</i> group A	He 6304	China	MT561714	MT598029	present study
<i>P. crassa</i> group A	Wu 0504 - 22	China	MT561715	GQ470634	Wu et al. 2010, present study
<i>P. crassa</i> group B	He 3349	China	MT561712	MT447407	present study
<i>P. crassa</i> group B	He 5866	China	MT386376	MT447408	present study
<i>P. crassa</i> group B	He 6266	China	MT561713	MT598035	present study
<i>P. crassa</i> group B	CLZhao 724	China	MG231790	—	—
<i>P. crassa</i> group B	MAFF 420737	Japan	AB809163	AB809163	—
<i>P. crassa</i> group C	KKN-86-Sp	USA	KP135394	KP135215	Floudas & Hibbett 2015
<i>P. crassa</i> group C	FP-102496-sp	USA	AY219341	—	de Koker et al. 2003
<i>P. crassa</i> group C	HHB 8834	USA	KP135393	—	Floudas & Hibbett 2015
<i>P. crassa</i> group C	ME 516	USA	KP135395	—	Floudas & Hibbett 2015
<i>P. cf. dregeana</i>	SFC 980804-4	Korea	AF479669	—	Lim & Jung 2003
<i>P. cf. dregeana</i>	UOC-DAMIA-D46	Sri Lanka	KP734203	—	—
<i>P. cf. dregeana</i>	FLAS-F-60030	USA	KY654737	—	—
<i>P. flavidoalba</i>	FD-263	USA	KP135402	KP135271	Floudas & Hibbett 2015
<i>P. flavidoalba</i>	Miettinen 17896	USA	KX752607	KX752607	Miettinen et al. 2016

Taxa	Voucher	Locality	ITS	nrLSU	Reference
<i>P. flavidoalba</i>	CFMR4167	USA	KX065957	—	—
<i>P. flavidoalba</i>	HHB-4617	USA	KP135401	—	Floudas & Hibbett 2015
<i>P. friesii</i>	He 5722	Sri Lanka	MT452528	MT447413	present study
<i>P. friesii</i>	He 5817	Sri Lanka	MT452529	MT447414	present study
<i>P. friesii</i>	He 5820	Sri Lanka	MT452530	MT447415	present study
<i>P. gigantea</i>	He 5290	China	MT386381	MT447416	present study
<i>P. gigantea</i>	Miettinen 15354	Finland	KX752605	—	Miettinen et al. 2016
<i>P. gigantea</i>	CBS 935.70	Germany	MH860011	MH871798	Vu et al. 2019
<i>P. gigantea</i>	FP-70857-Sp	USA	KP135390	KP135272	Floudas & Hibbett 2015
<i>P. griseofuscescens</i>	He 5734	Sri Lanka	MT561708	MT598032	present study
<i>P. griseofuscescens</i>	Cui 12629	China	MT561718	—	present study
<i>P. griseofuscescens</i>	CLZhao 3692	China	MT180946	MT180950	Xu et al. 2020
<i>P. griseofuscescens</i>	CLZhao 3705	China	MT180947	MT180951	Xu et al. 2020
<i>P. laxa</i>	Wu 9311-17	China	MT561710	GQ470649	Wu et al. 2010, present study
<i>P. magnicystidiata</i>	He 5648 *	China	MT386377	MT447409	present study
<i>P. magnicystidiata</i>	He 20140719_18	China	MT561719	—	present study
<i>P. magnicystidiata</i>	Wu 890805-1	China	MT561711	GQ470667	Wu et al. 2010, present study
<i>P. membranacea</i>	He 3842	China	MT386400	MT447440	present study
<i>P. membranacea</i>	He 3849 *	China	MT386401	MT447441	present study
<i>P. membranacea</i>	He 6062	China	MT386402	MT447442	present study
<i>P. pilatii</i>	He 5114	China	MT386385	MT447421	present study
<i>P. pilatii</i>	He 5165	China	MT386386	MT447422	present study
<i>P. pilatii</i>	Dai 17041	China	KY971603	KY971604	Wu et al. 2017
<i>P. pilatii</i>	Spirin 5048	Russia	KX752590	KX752590	Miettinen et al. 2016

Taxa	Voucher	Locality	ITS	nrLSU	Reference
<i>P. sinensis</i>	He 4295	China	MT386395	MT447433	present study
<i>P. sinensis</i>	He 4665	China	MT386396	MT447434	present study
<i>P. sinensis</i>	He 4673 *	China	MT386397	MT447435	present study
<i>P. sinensis</i>	He 5662	China	MT386398	MT447436	present study
<i>P. sp.</i>	FP-102937	USA	KP135391	KP135270	Floudas & Hibbett 2015
<i>P. sp.</i>	ECS1971	USA	KP135392	–	Floudas & Hibbett 2015
<i>P. sp.</i>	He 3827	China	–	MT447437	present study
<i>P. yunnanensis</i>	He 2623	China	MT386387	MT447423	present study
<i>P. yunnanensis</i>	He 3249	China	MT386375	MT447425	present study
<i>P. yunnanensis</i>	CLZhao 3958	China	MH744140	MH744142	Zhao et al. 2018
<i>P. yunnanensis</i>	CLZhao 3990	China	MH744141	MH744143	Zhao et al. 2018
<i>Pirex concentricus</i>	OSC-41587	USA	KP134984	KP135275	Floudas & Hibbett 2015
<i>Porostereum fulvum</i>	LY: 18496	France	MG649453	MG649455	–
<i>P. spadiceum</i>	CBS 474.48	France	MH856438	MH867984	Vu et al. 2019
<i>Rhizochaete americana</i>	FP-102188	USA	KP135409	KP135277	Floudas & Hibbett 2015
<i>R. belizensis</i>	FP-150712	Belize	KP135408	KP135280	Floudas & Hibbett 2015
<i>R. brunnea</i>	MR 229	Argentina	AY219389	AY219389	Greslebin et al. 2004
<i>R. violascens</i>	KHL 11169	Norway	EU118612	EU118612	Larsson 2007
<i>R. filamentosa</i>	HHB-3169-Sp	USA	KP135410	KP135278	Floudas & Hibbett 2015
<i>R. flava</i>	PR 1141	Puerto Rico	KY273030	KY273033	Nakasone et al. 2017
<i>R. fouquieriae</i>	KKN-121-sp	USA	KY948786	KY948858	Justo et al. 2017
<i>R. radicata</i>	FD-123	USA	KP135407	KP135279	Floudas & Hibbett 2015

Taxa	Voucher	Locality	ITS	nrLSU	Reference
<i>R. sulphurina</i>	HHB-5604	USA	KY273031	GU187610	Binder et al. 2010
<i>R. sulphurosa</i>	URM 87190	Brazil	KT003522	KT003519	Chikowski et al. 2015
<i>Riopa metamorphosa</i>	Spirin 2395	Russia	KX752601	KX752601	Miettinen et al. 2016
<i>R. pudens</i>	Cui 3238	China	JX623931	JX644060	Jia et al. 2014
<i>Terana caerulea</i>	FP-104073	USA	KP134980	KP135276	Floudas & Hibbett 2015
Outgroup					
<i>Ceraceomyces serpens</i>	HHB-15692-Sp	USA	KP135031	KP135200	Floudas & Hibbett 2015
<i>Phebia acerina</i>	FD-301	USA	KP135378	KP135260	Floudas & Hibbett 2015

Phylogenetic analyses

Two separate datasets, the concatenated ITS-nrLSU sequences of species in the Phanerochaetaceae and ITS only sequences of *Phlebiopsis*, were analyzed. *Ceraceomyces serpens* (Tode) Ginns and *Phlebia acerina* Peck were selected as the outgroup for the ITS-LSU dataset, whilst *Rhizochaete radicata* (Henn.) Gresl., Nakasone & Rajchenb. was used in the ITS dataset (Floudas and Hibbett 2015). For the concatenated dataset, the sequences of ITS and nrLSU were aligned separately using MAFFT v.7 (Kato et al. 2017, <http://mafft.cbrc.jp/alignment/server/>) with the G-INS-I iterative refinement algorithm, and optimized manually in BioEdit v.7.0.5.3. The separate alignments were then concatenated using Mesquite v.3.5.1 (Maddison and Maddison 2018). The datasets were deposited in TreeBase (<http://treebase.org/treebase-web/home.html>, submission ID: 26529 for Phanerochaetaceae ITS-LSU, 26530 for *Phlebiopsis* ITS).

Maximum parsimony (MP), Maximum likelihood (ML) analyses and Bayesian inference (BI) were carried out by using PAUP* v.4.0b10 (Swofford 2002), RAxML v.8.2.10 (Stamatakis 2014) and MrBayes 3.2.6 (Ronquist et al. 2012), respectively. In MP analysis, trees were generated using 100 replicates of random stepwise addition of sequence and tree-bisection reconnection (TBR) branch-swapping algorithm with all characters given equal weight. Branch supports for all parsimony analyses were estimated by performing 1000 bootstrap replicates with a heuristic search of 10 random-addition replicates for each bootstrap replicate. In ML analysis, statistical support values were obtained using rapid bootstrapping with 1000 replicates, with default settings used for other parameters. For BI, the best-fit substitution model was estimated with jModeltest v.2.17 (Darriba et al. 2012). Four Markov chains were run for 5,000,000 and 3,000,000 generations for the Phanerochaetaceae ITS-LSU and *Phlebiopsis* ITS datasets, respectively, until the split deviation frequency value was lower than 0.01. Trees were sampled every 100th generation.

The first quarter of the trees, which represented the burn-in phase of the analyses, were discarded, and the remaining trees were used to calculate posterior probabilities (BPP) in the majority rule consensus tree.

Results

Phylogenetic analyses

Forty-three ITS and 37 nrLSU sequences were generated for this study. The concatenated ITS-LSU dataset contained 101 ITS and 107 nrLSU sequences from 107 samples representing 86 Phanerochaetaceae taxa and the outgroup, while the ITS dataset contained 71 samples representing 21 *Phlebiopsis s.s.* taxa, a sample of *Irpex vellereus* and the outgroup (Table 1). The concatenated dataset had an aligned length of 2339 characters, of which 554 were parsimony-informative. MP analysis yielded one equally parsimonious tree (TL = 3603, CI = 0.360, RI = 0.695, RC = 0.250, HI = 0.640). The ITS dataset had an aligned length of 726 characters, of which 178 were parsimony-informative. MP analysis yielded 92 equally parsimonious trees (TL = 658, CI = 0.579, RI = 0.870, RC = 0.504, HI = 0.421). jModelTest suggested GTR + I + G and HKY + G were the best-fit models of nucleotide evolution for the concatenated ITS-LSU and ITS datasets, respectively. The average standard deviation of split frequencies of BI was 0.009223 and 0.007710 at the end of the run. ML and BI analyses resulted in almost identical tree topologies compared to the MP analysis. The MP trees are shown in Fig. 1 and Fig. 2 with the parsimony bootstrap values ($\geq 50\%$, first), Bayesian posterior probabilities (≥ 0.95 , second) and likelihood bootstrap values ($\geq 50\%$, third) labelled along the branches.

In the Phanerochaetaceae ITS-LSU tree (Fig. 1), *Phlebiopsis*, *Phaeophlebiopsis*, *Hapalopilus*, and *Rhizochaete* formed a strongly supported clade (98/1/100). Within this clade, the *Phlebiopsis* species clustered together with relatively strong support values (69/1/98), and species of *Phaeophlebiopsis*, *Hapalopilus* and *Rhizochaete* were in the sister subclades. In the *Phlebiopsis* ITS tree (Fig. 2), 24 lineages were resolved including 21 taxa of *Phlebiopsis* and '*Irpex vellereus*'. Samples of *P. crassa* were distributed in three distinct lineages. The six new species, *P. albescens*, *P. brunnea*, *P. cylindrospora*, *P. magnicystidiata*, *P. membranacea* and *P. sinensis*, formed distinct, strongly supported lineages.

Phlebiopsis albescens Y.N. Zhao & S.H. He, **sp. nov.** Figure 3

MycoBank: MB836023

Type – Sri Lanka, Avissawella, Salgala Forest, on fallen angiosperm twig, 3 March 2019, He 5805 (BJFC 030672, holotype).

Etymology – Refers to the white basidiomata.

Fruiting body – Basidiomata annual, resupinate, widely effused, closely adnate, inseparable from substrate, ceraceous to crustose, first as small patches, later confluent up to 15 cm long, 1 cm wide, up to 80 μm thick in section. Hymenophore smooth, white (6A1), orange white (6A2) to pale orange (6A3),

unchanged in KOH, not cracking on drying; margin indistinct, concolorous with hymenophore. Context white.

Microscopic structures – Hyphal system monomitic; generative hyphae simple-septate. Subiculum indistinct to absent. Subhymenium well developed; hyphae hyaline, thin- to slightly thick-walled, tightly agglutinated, 2.5–4 µm in diam. Lamprocystidia abundant, conical, hyaline to pale yellow, thick-walled, heavily encrusted with crystals along entire length, embedded or slightly projecting beyond hymenium, with one or two secondary septa, with a basal simple septum, 25–40 · 8–12 µm (without encrustations). Basidia clavate to cylindrical, hyaline, thin-walled, with a basal simple septum and four sterigmata, 10–16 × 3–4.5 µm; basidioles numerous, similar to basidia but slightly smaller. Basidiospores oblong ellipsoid to short cylindrical, hyaline, thin-walled, smooth, IKI–, CB–, 3.5–5 · 2–2.2 (–2.5) µm, L = 4.4 µm, W = 2.1 µm, Q = 2.1 (n = 30/1).

Distribution – Sri Lanka

Notes – *Phlebiopsis albescens* is characterized by thin, white to pale orange basidiomata, an indistinct subiculum, short lamprocystidia (< 40 µm long) and basidia (< 16 µm long), and small basidiospores (< 5 µm long). *Phlebiopsis punjabensis* G. Kaur, Avn.P. Singh & Dhingra, from India, also has thin, white basidiomata and short lamprocystidia but larger basidiospores, 5.3–8.5 × 2.5–4 µm (Kaur et al. 2015). Another species with short basidiospores, *P. yunnanensis* C.L. Zhao, from southern China, has thicker basidiomata, 100–500 µm thick, with a smooth to odontoid hymenophore, and ellipsoid basidiospores, 2.5–3.5 µm broad (Zhao et al. 2018). In the ITS phylogenetic tree (Fig. 2), *P. albescens* formed its lineage and was not closely related to any other species.

Phlebiopsis brunnea Y.N. Zhao & S.H. He, **sp. nov.** Figure 4

MycoBank: MB836024

Type – Sri Lanka, Western Province, Mitirigala Nissarana Vanaya Forest Monastery, on fallen angiosperm branch, 4 March 2019, He 5822 (BJFC 030689, holotype).

Etymology – Refers to the brown context of basidiomata.

Fruiting body – Basidiomata annual, resupinate, widely effused, closely adnate, inseparable from substrate, coriaceous, developing as small patches then confluent, up to 20 cm long, 5 cm wide, up to 350 µm thick in section. Hymenophore smooth, brownish grey (6C2–6D2), brownish orange (6C3) to greyish brown (6D3), unchanged in KOH, not cracking on drying; margin thinning out, indistinct, concolorous or darker than hymenophore. Context pale brown.

Microscopic structures – Hyphal system pseudodimitic; generative hyphae simple-septate. Subiculum well-developed, a non-agglutinated, loosely interwoven tissue; skeletocystidia (skeletal hyphae) brown, distinctly thick-walled, slightly encrusted, up to 120 µm long, 14 µm wide; hyphae hyaline to pale yellowish brown, thick-walled, smooth, moderately branched at right angles, frequently septate, 2–5 µm in

diam. Subhymenium thin; skeletocystidia as in subiculum but shorter and more heavily encrusted; generative hyphae hyaline, thin- to thick-walled, moderately branched, frequently septate, loosely interwoven, 2–4.5 µm in diam. Lamprocystidia subulate to fusiform, hyaline, thin- to thick-walled, distal end encrusted with small crystals, projecting up to 30 µm beyond hymenium, with an obtuse or acute tip, with a basal simple septum, 35–65 × 7–10 µm. Basidia clavate to subcylindrical, hyaline, thin-walled, with a basal simple septum and four sterigmata, 20–33 × 4.5–6 µm; basidioles numerous, similar to basidia but slightly smaller. Basidiospores oblong ellipsoid to subcylindrical, hyaline, thin-walled, smooth, IKI–, CB–, 6.5–7.5 (–8) × 3–3.6 (–4) µm, L = 7.3 µm, W = 3.3 µm, Q = 2.2 (n = 30/1).

Distribution – Sri Lanka.

Notes – *Phlebiopsis brunnea* is characterized by a coriaceous basidiomata with a smooth hymenophore and brown context, abundant, brown skeletocystidia in the subiculum and subhymenium, lamprocystidia, and oblong ellipsoid to subcylindrical basidiospores. *Hjortstamia bambusicola* (Berk. & Broome) Hjortstam & Ryvarden is similar with its grayish brown hymenophore and pseudodimitic hyphal system with brown skeletocystidia but with narrower basidiospores (2.5–3 µm wide) and grows on bamboo in Australia (Hjortstam and Ryvarden 2005). *Phlebiopsis brunneocystidiata* (Sheng H. Wu) Miettinen has narrower lamprocystidia (5–8 µm wide) with brown walls and a host preference for Pandanaceae in Taiwan (Wu 2004). Another similar species, *P. crassa* differs from *P. brunnea* by having effused-reflexed basidiomata with a more or less purple hymenophore and larger lamprocystidia, 50–120 × 8–20 µm (Burdson 1985; Hjortstam and Ryvarden 1990). *Phlebiopsis brunnea* formed weakly supported sister lineages to *P. brunneocystidiata* or *P. crassa* group B and C in the ITS-LSU and ITS trees, respectively (Figs. 1, 2).

Phlebiopsis cylindrospora Y.N. Zhao & S.H. He, **sp. nov.** Figure 5

MycoBank: MB836025

Type – China, Hainan Province, Lingshui County, Diaoluoshan Nature Reserve, on dead, small diameter bamboo, 2 July 2019, He 5984 (BJFC 030860, holotype).

Etymology – Refers to the cylindrical basidiospores.

Fruiting body – Basidiomata annual, resupinate, widely effused, closely adnate, inseparable from substrate, coriaceous, first as small patches, later confluent up to 20 cm long, 4 cm wide, up to 150 µm thick in section. Hymenophore smooth, orange white (6A2), orange grey (6B2) to greyish orange (6B3), turning purple in KOH, not cracking on drying; margin thinning out, indistinct, slightly fimbriate, paler than or concolorous with hymenophore. Context gray.

Microscopic structures – Hyphal system monomitic; generative hyphae simple-septate. Subiculum distinct, a somewhat agglutinated, compact tissue, arranged more or less parallel to substrate; hyphae hyaline, thick-walled, encrusted with yellow, resinous granules, infrequently branched, moderately septate, 2–4.5 µm in diam. Subhymenium indistinct; hyphae thin- to slightly thick-walled, heavily encrusted with

yellow, resinous granules, frequently septate, more or less agglutinated, 2–4 µm in diam. Lamprocystidia numerous, subfusiform, hyaline, thick-walled, apically encrusted with small crystals, embedded or slightly projecting beyond hymenium, 20–30 (–40) × 5–7 µm. Basidia clavate to subcylindrical, hyaline, thin-walled, with a basal simple septum and four sterigmata, 12–16 × 4–5 µm; basidioles numerous, similar to basidia but slightly smaller. Basidiospores cylindrical, hyaline, thin-walled, smooth, IKI–, CB–, 5.5–7 × 2–2.8 (–3) µm, L = 6.2 µm, W = 2.3 µm, Q = 2.7 (n = 30/1).

Additional specimens examined – China, Hainan Province, Qiongzong County, Limushan Nature Reserve, on fallen angiosperm twig, 8 June 2016, He 3831 (BJFC 022333); on dead, small diameter bamboo, 8 June 2016, He 3882 (BJFC 022384, CFMR); Wuzhishan County, Wuzhishan Nature Reserve, on dead, small diameter bamboo, 10 June 2016, He 3926 (BJFC 022428); 30 June 2019, He 5922 (BJFC 030797), He 5932 (BJFC 030807), He 5936 (BJFC 030811) & He 5938 (BJFC 030813); Lingshui County, Diaoluoshan Nature Reserve, on dead, small diameter bamboo, 2 July 2019, He 5981 (BJFC 030857); 5 July 2019, He 6054 (BJFC 030930), He 6061 (BJFC 030937) & He 6063 (BJFC 030939); on fallen angiosperm branch, 5 July 2019, He 6038 (BJFC 030914). Thailand, Chiang Rai, Doi Pui, on rotten bamboo, 23 July 2016, He 4080 (BJFC 023521), He 4083 (BJFC 023524) & He 4094 (BJFC 023535, CFMR).

Distribution – China and Thailand.

Notes – *Phlebiopsis cylindrospora* is characterized by pale-colored, smooth hymenophore that turns purple in KOH, a monomitic hyphal system with generative hyphae encrusted with yellow, resinous granules, small subfusiform lamprocystidia, cylindrical basidiospores, and habit on bamboo and woody angiosperms. It is similar to *P. punjabensis* that also has a pale-colored, smooth hymenophore and short lamprocystidia, but the latter species does not react with KOH and develops longer basidia (14–26 µm long), and slightly larger basidiospores (5.3–8.5 × 2.5–4 µm, Kaur et al. 2015). *Phlebiopsis albescens* differs from *P. cylindrospora* by its white hymenophore that is unchanged in KOH and distinctly smaller basidiospores (3.5–5 × 2–2.2 µm). The hymenophore in *P. friesii* (Lév.) Spirin & Miettinen turns purple in KOH also but is distinct from *P. cylindrospora* by having effused-reflexed basidiomata, a pseudodimitic hyphal system, and larger lamprocystidia, up to 80 · 20 µm (Hjortstam and Ryvarden 1990). Although the phylogenetic trees (Figs. 1, 2) show that *P. cylindrospora* and *P. pilatii* are closely related, the latter species is distinct morphologically for it lacks lamprocystidia and develops finely branched dendrohyphidia and larger basidiospores, 8–10 × 4–4.5 µm (Parmasto 1965; Larsen and Gilbertson 1977; Duhem and Michel 2009).

Phlebiopsis magnicystidiata Y.N. Zhao & S.H. He, **sp. nov.** Figure 6

MycoBank: MB836026

Type – China, Hunan Province, Guzhang County, Gaowangjie Nature Reserve, on dead angiosperm branch, 4 August 2018, He 5648 (BJFC 026710, holotype).

Etymology – Refers to the large lamprocystidia.

Fruiting body – Basidiomata annual, resupinate, widely effused, closely adnate, inseparable from substrate, ceraceous to coriaceous, up to 15 cm long, 5 cm wide, up to 400 µm thick in section. Hymenophore smooth to slightly odontoid with scattered tubercles, pruinose from projecting cystidia, greyish orange [6B(3–5)], brownish orange [6C(3–5)] to light brown [6D(4–6)], unchanged in KOH, sometimes sparsely and deeply cracked with age; margin thinning out, indistinct, concolorous with hymenophore. Context white.

Microscopic structures – Hyphal system monomitic; generative hyphae simple-septate. Subiculum indistinct to absent. Subhymenium thickening, well-developed; hyphae hyaline, thin- to slightly thick-walled, frequently septate, slightly agglutinated, vertically arranged, 2–4.5 µm in diam. Lamprocystidia numerous, fusiform to subulate, hyaline, thick-walled, heavily encrusted with crystals, embedded or projecting beyond hymenium up to 40 µm, with a basal simple septum, apex subacute, 40–80 × (7–) 9–13 (–15) µm (without encrustations). Basidia clavate, hyaline, thin-walled, with a basal simple septum and four sterigmata, 20–30 × 5–6 µm; basidioles numerous, similar to basidia but slightly smaller. Basidiospores broadly ellipsoid to subglobose, hyaline, thin-walled, smooth, IKI–, CB–, 4.5–6.5 × (3.5–) 3.8–4.8 µm, L = 5.6 µm, W = 4.4 µm, Q = 1.3 (n = 30/1).

Additional specimens examined – China, Yunnan Province, Mengla County, Wangtianshu Forest Park, on fallen angiosperm branch, 19 July 2014, He 20140719-18 (BJFC 019145); Taiwan Province, Taichung, Tunghai University, on dead branch of *Cassia siamea*, 5 August 1989, Wu 890805-1 (TNM F0022186).

Distribution – Hunan, Yunnan, and Taiwan Provinces in southern China.

Notes – *Phlebiopsis magnicystidiata* is characterized by large lamprocystidia and broadly ellipsoid to subglobose basidiospores. It is morphologically similar to and phylogenetically closely related to *P. flavidoalba* (Cooke) Hjortstam (Figs. 1, 2) that has smooth hymenophore, slightly longer ellipsoid basidiospores (6–7.5 µm long) and a distribution in North and South America (Burdshall 1985; Gilbertson and Blackwell 1985). *Phlebiopsis gigantea* and *P. magnicystidiata* have similar lamprocystidia but the former differs in its well-developed subiculum, narrowly ellipsoid basidiospores, 5–7 × 2.5–3.5 µm, and often occurs on gymnospermous wood in the North Hemisphere (Eriksson et al. 1981; Bernicchia and Gorjón 2010). Except for developing a distinct subiculum, *P. darjeelingensis* and *P. magnicystidiata* have similar sized lamprocystidia, basidia, and basidiospores (Dhingra 1987). Reports of *P. flavidoalba* from India (Rattan 1977) and Taiwan (Wu 1990) need to be confirmed for they may be *P. magnicystidiata* instead.

Phlebiopsis membranacea Y.N. Zhao & S.H. He, **sp. nov.** Figure 7

MycoBank: MB836027

Type – China, Hainan Province, Qiongzong County, Limushan Nature Reserve, on dead, small diameter bamboo, 8 June 2016, He 3849 (BJFC 022351, holotype).

Etymology – Refers to the membranaceous basidiomata.

Fruiting body – Basidiomata annual, resupinate, widely effused, adnate, separable from substrate, membranaceous, up to 20 cm long, 5 cm wide, up to 250 µm thick in section. Hymenophore smooth, orange white (6A2), orange grey (6B2), greyish orange [6B(3–5)] to brownish orange [6C(3–5)], unchanged in KOH, sometimes sparsely and finely cracked with age; margin thinning out, fimbriate, concolorous with hymenophore. Context gray.

Microscopic structures – Hyphal system pseudodimitic; generative hyphae simple-septate. Subiculum well-developed, a non-agglutinated, loosely interwoven tissue; skeletocystidia abundant, fusiform to clavate, brown, thick-walled, smooth, with an acute or obtuse apex, embedded, (30–) 40–70 × 9–13 µm; hyphae hyaline, moderately to distinctly thick-walled, smooth, rigid, frequently branched at right angles, frequently septate, 3–5 µm in diam. Subhymenium thin; hyphae hyaline, thin-walled, smooth, somewhat agglutinated, interwoven, 2–4.5 µm in diam. Hymenial cystidia scattered, similar to skeletocystidia in shape and size but with paler, thinner walls, and sparse encrustations at apex. Basidia clavate, hyaline, thin-walled, with a basal simple septum and four sterigmata, 15–22 × 4–5 µm; basidioles numerous, similar to basidia but slightly smaller. Basidiospores oblong ellipsoid to subcylindrical, hyaline, thin-walled, smooth, IKI–, CB–, 4.5–6 × 2–3 µm, L = 5.4 µm, W = 2.4 µm, Q = 2.3 (n = 30/1).

Additional specimens examined – China, Hainan Province, Qiongzong County, Limushan Nature Reserve, on dead, small diameter bamboo, 8 June 2016, He 3842 (BJFC 022344); Lingshui County, Diaoluoshan Nature Reserve, on dead, small diameter bamboo, 5 July 2019, He 6062 (BJFC 030938).

Distribution – Hainan Province, southern tropical China.

Notes – *Phlebiopsis membranacea* is characterized by membranaceous basidiomata with well-developed subicula, brown, smooth, thick-walled skeletocystidia, without lamprocystidia, and habit on bamboo in tropical China. Like *P. membranacea*, *Hjortstamia novae-gratae* (A.L. Welden) Hjortstam & Ryvarde, from Columbia, grows on bamboo but its brown, smooth skeletocystidia are tubular in shape and its basidiospores are larger, 5.5–7 × 3–4 µm (Hjortstam and Ryvarde 1990). *Phlebiopsis laxa* (Sheng H. Wu) Miettinen like *P. membranacea* has membranaceous basidiomata and loosely arranged subicular hyphae but differs in having lamprocystidia and larger basidiospores, 8–10 × 4–5 µm (Wu 2000). In the phylogenetic trees (Figs. 1, 2), *P. membranacea* is sister to *P. laxa*, though their relationship is not strongly supported.

Phlebiopsis sinensis Y.N. Zhao & S.H. He, **sp. nov.** Figure 8

MycoBank: MB836028

Type – China, Sichuan Province, Wanyuan County, Huaeshan Nature Reserve, on fallen angiosperm branch, 17 July 2013, He 4673 (BJFC 024192, holotype).

Etymology – Refers to the distribution in China.

Fruiting body – Basidiomata annual, resupinate to effused-reflexed with reflexed edges elevated and incurved with age, loosely adnate, easily detached from substrate, coriaceous, first as small patches, later confluent up to 15 cm long, 5 cm wide, up to 300 µm thick in section. Pileus projecting up to 1.5 mm; upper surface gray, slightly sulcate. Hymenophore smooth, brownish orange [6C(3–5)], greyish brown [6(D–F)3] to brown [6E(4–6)], unchanged in KOH, sometimes finely cracked with age; margin thinning out, distinct, white to gray, silky, slightly fimbriate, up to 1 mm wide. Context gray to yellowish brown.

Microscopic structures – Hyphal system pseudodimitic; generative hyphae simple-septate. Tomentum and cortex (a dark line between the tomentum and subiculum) present. Subiculum well-developed, a non-agglutinated tissue; skeletocystidia brown, thick-walled, encrusted at apex, embedded, intermediate forms between skeletocystidia and lamprocystidia observed; hyphae hyaline to pale yellow, moderately to distinctly thick-walled, smooth, rarely branched, moderately septate, easily separated, more or less parallel to substrate, 3–6 µm in diam. Subhymenium indistinct. Lamprocystidia abundant, broadly fusiform to broadly subulate, usually with a long, curved stalk and resembling skeletocystidia, hyaline to brown, thick-walled, heavily encrusted, 30–60 × 8–13 µm, projecting up to 30 µm. Basidia clavate, hyaline, thin-walled, with a basal simple septum and four sterigmata, 20–30 × 4.5–5.5 µm; basidioles numerous, similar to basidia but slightly smaller. Basidiospores oblong ellipsoid to subcylindrical, hyaline, thin-walled, smooth, IKI–, CB–, (5.5–) 6–7.5 × 2.5–3.2 (–3.5) µm, L = 6.5 µm, W = 2.8 µm, Q = 2.3 (n = 30/1).

Additional specimens examined – China, Gansu Province, Pingliang County, Kongtongshan Nature Reserve, on construction wood, 3 August 2015, He 2416 (BJFC 020870, CFMR); Hubei Province, Wufeng County, Houhe Nature Reserve, on dead angiosperm branch, 16 August 2017, He 5081 (BJFC 024599); Hunan Province, Yongshun County, Xiaoxi Nature Reserve, on dead angiosperm branch, 6 August 2018, He 5662 (BJFC 026724); Inner Mongolia, Chifeng, Aohan County, Daheishan Nature Reserve, on fallen *Quercus mongolia* branch, 3 September 2015, Tiezhi Liu et al. (CFSZ 10714), on fallen *Pinus tabuliformis* branch, 19 September 2016, Tiezhi Liu et al. (CFSZ 12436); Jiangxi Province, Ji'an County, Jinggangshan Nature Reserve, on dead *Rhododendron* branch, 11 August 2016, He 4295 (BJFC 023737, CFMR); Liaoning Province, Zhuanghe County, Xianrendong Forest Park, on dead *Quercus* branch, 5 August 2017, He 4665 (BJFC 024184); Shaanxi Province, Foping County, Foping Nature Reserve, on fallen *Betula* branch, 11 September 2013, He 1907 (BJFC 016374); Sichuan Province, Baoxing County, Fengtongzhai Nature Reserve, on fallen angiosperm trunk, 18 September 2012, He 20120918-3 (BJFC 014609).

Distribution – Gansu, Hubei, Hunan, Jiangxi, Liaoning, Shaanxi and Sichuan Provinces and Inner Mongolia Autonomous Region of China.

Notes – *Phlebiopsis sinensis* is characterized by effused to effused-reflexed, coriaceous basidiomata with well-developed subicula, brown skeletocystidia, lamprocystidia, and a temperate distribution. Submembranaceous-pellicular basidiomata, narrower cystidia (5–8 µm wide), and a tropical distribution distinguish *P. brunneocystidiata* from *P. sinensis* (Wu 2004). Both *P. crassa* and *P. sinensis* develop effused-reflexed basidiomata, but the former species has a purple-tinted hymenophore, larger lamprocystidia, 50–120 × 8–20 µm, and a tropical distribution (Hjortstam and Ryvarden 1990). Although

the ITS tree (Fig. 2) shows that *P. sinensis* and *P. friesii* are sister taxa, *P. friesii* is distinct morphologically with a hymenophore that turns purple in KOH and has a dimitic hyphal system with hyaline to yellow skeletal hyphae (Hjortstam and Ryvarde 1990).

Phaeophlebiopsis mussooriensis (Priyanka, Dhingra & N. Kaur) Nakasone & S.H. He, **comb. nov.**

MycoBank: MB836029

Phlebiopsis mussooriensis Priyanka, Dhingra & N. Kaur, Mycotaxon 115: 255, 2011.

Notes – This species is characterized by a grayish yellow hymenophore, well-developed subiculum, thin-walled generative hyphae, lamprocystidia, and ellipsoid basidiospores (Priyanka et al. 2011). As mentioned in the protologue, *P. mussooriensis* is quite similar to *P. himalayensis*, now *Phaeophlebiopsis himalayensis* (Dhingra) Zmitr., differing primarily in basidiospore size and color change of hymenophore in KOH. Based on Priyanka et al.'s (2011) description, illustration, and comments, we propose the transfer of *P. mussooriensis* into *Phaeophlebiopsis*.

Phlebiopsis bambusicola (Berk. & Broome) Nakasone & S.H. He, **comb. nov.**

MycoBank: MB836030

Corticium bambusicola Berk. & Broome, Transactions of the Linnaean Society of London 2: 64, 1882.

Peniophora bambusicola (Berk. & Broome) Sacc., Sylloge Fungorum 6: 647, 1888.

Hjortstamia bambusicola (Berk. & Broome) Hjortstam & Ryvarde, Synopsis Fungorum 20: 37, 2005.

Notes – This Australian species is known only from the type and is characterized by a grayish brown hymenophore, a dimitic hyphal system, large, brown skeletocystidia, lamprocystidia, narrowly ellipsoid to allantoid basidiospores, and a habit on bamboo (Hjortstam and Ryvarde 2005). Although similar to *P. crassa*, *P. bambusicola* has narrower basidiospores, 2.5–3 µm broad and is restricted by host preference and distribution.

Phlebiopsis dregeana (Berk.) Nakasone & S.H. He, **comb. nov.**

MycoBank: MB836031

Corticium dregeanum Berk., London Journal of Botany 5: 3, 1846.

Hymenochaete dregeana (Berk.) Masee, Botanical Journal of the Linnean Society 27: 114, 1890.

Terana dregeana (Berk.) Kuntze, Revisio generum plantarum 2: 872, 1891.

Lopharia dregeana (Berk.) P.H.B. Talbot, Bothalia 6: 57, 1951.

Irpex dregeanus (Berk.) P.H.B. Talbot, Bothalia 6: 344, 1954.

Australohydnum dregeanum (Berk.) Hjortstam & Ryvar den, Synopsis Fungorum 4: 61, 1990.

Notes – This is a poorly understood species that has been interpreted differently by various researchers. We take a narrow concept of *P. dreageana* based on studies of the type specimen and specimens restricted to Africa as described and illustrated by Masee (1891), Talbot (1951), Reid (1975), and Hjortstam and Ryvar den (1990). The ellipsoid basidiospores based on these studies are approximately $6.5\text{--}8 \times 4\text{--}5 \mu\text{m}$ in size. Note that the cylindrical basidiospores illustrated by Reid (1975) are questionable for Hjortstam (1989) noted that basidia and spores were not observed in the type. Hjortstam and Ryvar den (1990) took a broad interpretation of *A. dreageanum* when they placed *Hydnum griseofuscescens* Reichardt from Australia and *Irpex vellereus* Berk. & Broome from Sri Lanka in synonymy; see below for further discussion of these two taxa. Although *A. dreageanum* has since been reported from India (De 1998, as *Oxyporus vellereus*), South Korea (Lim 2001; Lim and Jung 2003), New Zealand (Buchanan and Ryvar den 2000), Portugal (Melo and Hjortstam 2002), Israel (Tura et al. 2011), and Italy (Saitta et al. 2014), the basidiospore size, when given, is significantly smaller than the African collections.

Sequences from authentic specimens of the species are not available at present, but ITS sequences labelled "*Australohydnum dregeanum*" in GenBank, from U.S.A., Korea and Sri Lanka, formed a strongly supported lineage within *Phlebiopsis* (Fig. 2). The identity of the taxa in this lineage needs further study.

Phlebiopsis griseofuscescens (Reichardt) Nakasone & S.H. He, **comb. nov.**

MycoBank: MB836032

Hydnum griseofuscescens Reichardt, Verhandlungen der Zoologisch-Botanischen Gesellschaft Wien 16: 374, 1866.

Irpex griseofuscescens (Reichardt) D.A. Reid, Kew Bulletin 17 (2): 273, 1963.

Australohydnum griseofuscescens (Reichardt) Jülich, Persoonia 10 (1): 138, 1978.

Irpex vellereus Berk. & Broome, Journal of the Linnean Society. Botany 14: 61, 1875.

Xylodon vellereus (Berk. & Broome) Kuntze, Revisio generum plantarum 3 (2): 541, 1898.

Hirschioporus vellereus (Berk. & Broome) Teng, Zhong Guo De Zhen Jun [Fungi of China]: 761, 1963.

Oxyporus vellereus (Berk. & Broome) A. Roy & A.B. De, J. Mycopathol. Res.: 41, 1998.

Phlebiopsis lacerata C.L. Zhao, Phytotaxa 440 (4): 274, 2020.

Hydnochaete philippinensis Lloyd (as "*philippensis*"), Mycological Writings 7 (67): 1154, 1922.

Trichaptum venustum (Berk.) G. Cunn., Bulletin of the New Zealand Department of Scientific and Industrial Research 164: 97, 1965.

Specimens examined – Sri Lanka, Western Province, Ingiriya, Dombagaskanda Forest Reserve, on fallen angiosperm branch, 27 February 2019, He 5734 (BJFC 030601). China, Sichuan Province, Miyi County, Haita Village, on fallen *Quercus* trunk, 13 September 2015, Cui 12629 (BJFC 028408) & Cui 12637 (BJFC 028416).

Notes – *Hydnum griseofuscescens* was described from Australia and is the type of *Australohydnum* (Jülich 1978). It is characterized by resupinate to effused-reflexed basidiomata with a hydroid, purplish brown hymenophore, a pseudodimitic hyphal system with simple-septate, hyaline, generative hyphae, 4–9 µm broad, encrusted hymenial cystidia with hyaline walls, and small ellipsoid basidiospores, 4–6 × 2.5–3 µm (Reid 1956 as *Irpex vellerus*, Jülich 1978). We follow Reid (1956, 1963) who determined that *H. griseofuscescens* and *I. vellerus*, described from Sri Lanka, were synonyms after studying the types of both species. Reid (1967) also reported that *T. venustum* sensu Cunningham (1965) is *H. griseofuscescens*. Based on morphological studies and sequence analyses, we determined that *P. lacerata* described from southern China (Xu et al. 2020) is conspecific with *P. griseofuscescens*.

Gilbertson and Adaskaveg (1993) described and illustrated *I. griseofuscescens* from Hawaii, but this species lacks encrusted hymenial cystidia and has small basidiospores, 4–4.5 × 2–2.5 µm. Similarly, De's (1998) description of *O. vellerus* from India appears to represent a different species with a monomitic hyphal system of hyaline to pale brown hyphae and cylindrical basidiospores, 5.2–7 × 2–3 µm. One of the specimens cited, VBMN 80451, is also at CBS, CBS 515.92, and its ITS sequence is available from GenBank (AF479670) as "*Irpex vellerus*". This sequence was included in Lim and Jung (2003) and Fig. 2, herein, where it is on a long branch, sister to *P. griseofuscescens*.

Phlebiopsis novae-gratae (A.L. Welden) Nakasone & S.H. He, **comb. nov.**

MycoBank: MB836033

Lopharia novae-gratae A.L. Welden [as '*nova-grata*'], Mycologia 67: 540, 1975.

Porostereum novae-gratum (A.L. Welden) Hjortstam & Ryvar den [as '*nova-gratum*'], Synopsis Fungorum 4: 41, 1990.

Phanerochaete novae-gratae (A.L. Welden) Sheng H. Wu [as '*nova-grata*'], Mycotaxon 88: 375, 2003.

Hjortstamia novae-gratae (A.L. Welden) Hjortstam & Ryvar den [as '*nova-grata*'], Synopsis Fungorum 25: 19, 2008.

Notes – Reported from Colombia on bamboo, this species is characterized by a pale brown hymenophore and smooth skeletocystidia but lacking lamprocystidia (Welden 1975; Hjortstam and Ryvar den 1990). Because of its morphological similarity to *P. crassa*, the transfer of *P. novae-gratae* is proposed.

Phlebiopsis crassa species complex Fig. 9

Specimens examined – *Phlebiopsis crassa* group A: Vietnam, Ho Chi Minh City, the Botanical Garden Padua, on fallen angiosperm trunk, 13 October 2017, He 5205 (BJFC 024723). Sri Lanka, Central Province, Kandy, Peradeniya Botanic Garden, on fallen angiosperm branch, 2 March 2019, He 5763 (BJFC 030630). China, Guangdong Province, Renhua County, Danxiashan Nature Reserve, on fallen angiosperm trunk, 4 June 2019, He 5855 (BJFC 030730, Fig. 10a); Yunnan Province, Qiubei County, Puzhehei Nature Reserve, 17 November 2019, He 6300 (BJFC, Fig. 10c), He 6301 (BJFC, Fig. 10d), He 6303 (BJFC, Fig. 10b) & He 6304 (BJFC); Ximeng County, Mengsuolongtan Forest Park, on fallen angiosperm branch, 15 April 2005, Wu 0504 – 22 (TNM F0018719).

Phlebiopsis crassa group B: China, Guangdong Province, Renhua County, Danxiashan Nature Reserve, on fallen angiosperm branch, 4 June 2019, He 5866 (BJFC 030741, Fig. 10f); Yunnan Province, Lushui County, Gaoligongshan Nature Reserve, on fallen angiosperm trunk, 29 November 2015, He 3349 (BJFC 021744, Fig. 10g–h); Maguan County, Gulinqing Nature Reserve, on fallen angiosperm branch, 14 November 2019, He 6266 (BJFC, Fig. 10e).

Phlebiopsis crassa group C: U.S.A., Arizona, Pima County, Santa Rita Experimental Range, on *Fouquieria splendens*, 31 July 1976, K.K. Nakasone, KKN-86-sp (CFMR); Illinois, Coles County, Fox Ridge State Park, on hardwood, 24 September 1990, A.S. Methven, FP-1024996-sp (CFMR); Mississippi, Harrison County, Harrison Experimental Forest, on *Quercus* sp., 26 March 1976, H.H. Burdsall, Jr., HHB-8834-sp (CFMR).

Notes – Our phylogenetic analyses showed that samples of *P. crassa* group A from Vietnam, Sri Lanka and southern China formed a distinct lineage and may represent *P. crassa* s.s., for the type was described from Vietnam (Figs. 1, 2). Collections from southern China and Japan, group B, and the U.S.A., group C, clustered into two lineages in the ITS tree (Fig. 2). All three lineages of *P. crassa* are morphologically similar, however. Unraveling this species complex is beyond the scope of this study, involving a number of presumed synonyms of *P. crassa*; see Lentz (1955) and Burdsall (1985).

Phlebiopsis darjeelingensis Dhingra, Nova Hedwigia 44: 222, 1987.

Phanerochaete lamprocystidiata Sheng. H. Wu, Mycotaxon 90: 426, 2004.

Phlebiopsis lamprocystidiata (Sheng H. Wu) Sheng H. Wu & Hallenb., Fungal Diversity 42: 116, 2010.

Notes – Because *P. darjeelingensis*, from India, and *P. lamprocystidiata*, from Taiwan, are nearly identical morphologically – basidiomata ceraceous when fresh then corneous when dried, well-developed subiculum of compactly packed, hyaline hyphae, and cystidia and basidiospores of similar shape and size (Dhingra 1987; Wu 2004), we consider *P. lamprocystidiata* to be a later synonym of *P. darjeelingensis*. Zmitrovich (2018) transfer *Phlebiopsis lamprocystidiata* to *Phaeophlebiopsis* based on morphology, our phylogenetic analyses show that it belongs to *Phlebiopsis* s.s., however.

Discussion

The generic limits of *Phlebiopsis* has expanded over the last 40 years since its introduction in 1978 to include significant morphological range in basidiomata habit and texture and hymenophore configuration with the aid of molecular phylogenetic methods (e.g., Floudas and Hibbett 2015; Miettinen et al. 2016; Zhao et al. 2018; Xavier de Lima et al. 2020; Xu et al. 2020). In this study, we emphasized sampling of *Phlebiopsis* taxa, and our overall results confirm those of Floudas and Hibbett (2015), Miettinen et al. (2016), and Chen et al. (2018a). In Figs. 1 and 2, *Phlebiopsis*, including the types of *Australohydnum*, *P. griseofuscescens* and *Hjortstamia*, *P. friesii*, formed a well-supported clade in the Phanerochaetaceae and is closely related to *Phaeophlebiopsis*, *Hapalopilus* and *Rhizochaete*. The genera *Phlebiopsis* and *Australohydnum* were published simultaneously (Jülich 1978) but the former is favored to avoid unnecessary name changes. Twenty-four lineages were resolved in the ITS tree of *Phlebiopsis*, among which 18 are accepted species, including the *P. crassa* species complex and six new species described herein. Further study is required to identify the taxa named *P. cf. dregeana*, *Irpex vellerus*, *Phlebiopsis* sp. FP-102937 and *Phlebiopsis* sp. ECS-1971.

Among the 24 names of *Phlebiopsis* in Index Fungorum (accessed on 21 June 2020), we accept 17 taxa in *Phlebiopsis s.s.*, including 11 that are supported by molecular data. Five taxa, *P. himalayensis* Dhingra, *P. mussooriensis*, *P. peniophoroides* Gilb. & Adask., *P. ravenelii* (Cooke) Hjortstam, and *P. roumegueri* (Bres.) Jülich & Stalpers were transferred to *Phaeophlebiopsis* based on morphology and sequence data. *Phlebiopsis lacerata* and *P. lamprocystidiata* are synonyms of *P. griseofuscescens* and *P. darjeelingensis*, respectively, as discussed above. Thus, 27 species of *Phlebiopsis* worldwide are accepted, including the six new species and four new combinations reported herein. An emended description of *Phlebiopsis* and an identification key to all species in the genus worldwide are presented below.

Phlebiopsis (Jülich) Nakasone & S.H. He, emended

Castanoporus Ryvar den, Synopsis Fungorum 5: 121, 1991.

Hjortstamia Boidin & Gilles, Bulletin de la Société Mycologique de France 118 (2): 99, 2003.

Australohydnum Jülich, Persoonia 10 (1): 138, 1978.

Description: Basidiomata annual, resupinate, effused, effused-reflexed or pileate, ceraceous, membranaceous to coriaceous. Pilei, when present, tomentose, gray to brown. Hymenophore smooth, tuberculate, odontoid, hydroid to poroid, white, gray, grayish brown, purplish brown or brown, turning purple in KOH in two species. Hyphal system monomitic or dimitic; generative hyphae simple-septate, hyaline or rarely pale brown, in dimitic species with skeletal or, in one species, micro-binding hyphae. Subiculum absent to well-developed, hyaline, brown, agglutinated or not, compact to loosely interwoven. Skeletocystidia absent or present, hyaline or brown, distinctly thick-walled, smooth or encrusted. Hymenial cystidia or lamprocystidia typically present, hyaline or light brown, thick-walled, usually encrusted. Dendrohyphidia present in one species, hyaline, thin-walled, smooth, branched. Basidia clavate or subcylindrical, with 4 sterigmata and a basal simple septum. Basidiospores cylindrical, ellipsoid, broadly ellipsoid or subglobose, hyaline, thin-walled, smooth, negative in Melzer's reagent, acyanophilous.

Type species: *Phlebiopsis gigantea* (Fr.) Jülich

Notes – The terminology relating to the cystidia observed in *Phlebiopsis* species is varied in the literature and thus confusing. There are up to three kinds of cystidia, but intermediate forms can develop to blur their distinctiveness. Lamprocystidia are found in most species of *Phlebiopsis* in the hymenium, often projecting, and may become embedded as the basidiomata thickens. They are typically conical or subfusiform with thick walls that are lightly to heavily encrusted in the upper half or apex. Skeletocystidia are found in dimitic or pseudodimitic species in which thick-walled hyphae in the subiculum curve toward the hymenium but remain embedded in the subiculum or subhymenium. The terminal ends may or may not be differentiated and usually lack encrustations. Hymenial cystidia are those structures that are similar to skeletocystidia but terminate in the hymenium and may be encrusted. In other cases, they are formed in the subhymenium and are smaller than lamprocystidia and not conical or heavily encrusted.

Key to 27 *Phlebiopsis* species

1. Hymenophore poroid, irpicoid or hydroid..... 2
1. Hymenophore smooth, tuberculate or odontoid..... 4
2. Basidiomata resupinate; hymenophore poroid to irpicoid; on gymnosperms..... *P. castanea*
2. Basidiomata effused-reflexed; hymenophore tuberculate to hydroid; on angiosperms..... 3
3. Basidiospores $6.5-8 \times 4-5 \mu\text{m}$; African species..... *P. dregeana*
3. Basidiospores $4.5-6 \times 2.5-3 \mu\text{m}$; Asian and Australasian species..... *P. griseofuscescens*
4. Dendrohyphidia present..... *P. pilatii*
4. Dendrohyphidia absent..... 5
5. Hyphal system pseudodimitic or dimitic..... 6
5. Hyphal system monomitic..... 13
6. Hymenophore turning purple in KOH..... *P. friesii*
6. Hymenophore unchanged in KOH..... 7
7. Basidiomata with well-developed pilei; skeletocystidia absent..... *P. papyrina*
7. Basidiomata resupinate to effused-reflexed; skeletocystidia present..... 8
8. Hymenophore without purple tints..... 9
8. Hymenophore more or less purple..... 12

9. Lamprocystidia none; basidiospores $\leq 6 \mu\text{m}$ long..... *P. membranacea*
9. Lamprocystidia present; basidiospores $\geq 6 \mu\text{m}$ long..... 10
10. On angiosperms and gymnosperms; from temperate China..... *P. sinensis*
10. On angiosperms or bamboo; from tropical-subtropical Asia or Australia..... 11
11. Basidiospores $6-7 \times 2.5-3 \mu\text{m}$; on bamboo; from Australia..... *P. bambusicola*
11. Basidiospores $6.5-7.5 \times 3-3.6 \mu\text{m}$; on angiospermous wood; from Sri Lanka..... *P. brunnea*
12. Lamprocystidia brown to dark brown; South American species..... *P. amethystea*
12. Lamprocystidia cystidia hyaline to pale brown; North American or Asian species... *P. crassa s.l.*
13. Lamprocystidia none; skeletocystidia or hymenial cystidia present..... 14
13. Lamprocystidia present; skeletocystidia absent..... 15
14. Basidiospores $5.5-7 \times 3-4 \mu\text{m}$; on bamboo; from Colombia..... *P. novae-granatae*
14. Basidiospores $3.7-5.5 \times 2.5-3.3 \mu\text{m}$; on hardwood; from New Zealand..... *P. afibulata*
15. Basidiospores $> 8 \mu\text{m}$ long, $> 4 \mu\text{m}$ broad..... *P. laxa*
15. Basidiospores $< 8 \mu\text{m}$ long, $< 4 \mu\text{m}$ broad..... 16
16. Lamprocystidia small, generally $< 40 \mu\text{m}$ long..... 17
16. Lamprocystidia large, generally $> 40 \mu\text{m}$ long..... 20
17. Hymenophore purple in KOH..... *P. cylindrospora*
17. Hymenophore unchanged in KOH..... 18
18. Basidiospores broadly ellipsoid, $3.5-4.5 \times 2.5-3.5 \mu\text{m}$, $Q = 1.3$ *P. yunnanensis*
18. Basidiospores narrowly ellipsoid to cylindrical..... 19
19. Basidiospores $3.5-5 \times 2-2.2 \mu\text{m}$ *P. albescens*
19. Basidiospores $5.3-8.5 \times 2.5-4 \mu\text{m}$ *P. punjabensis*
20. Lamprocystidia brown; on Pandanaceae; from Taiwan..... *P. brunneocystidiata*
20. Lamprocystidia hyaline; on other plants; from various locations..... 21

21. Subiculum indistinct to absent..... 22
- 21 .Subiculum distinct to well-developed..... 24
22. Basidia mostly with 2 sterigmata..... *P. bicornis*
22. Basidia with 4 sterigmata..... 23
23. Basidiospores $5.5-7.5 \times 3.5-4.5 \mu\text{m}$; from North and South America..... *P. flavidoalba*
23. Basidiospores $4.5-6.5 \times 3.8-4.8 \mu\text{m}$; from Asia..... *P. magnicystidiata*
24. Basidiospores narrowly ellipsoid to ellipsoid, $\leq 3 \mu\text{m}$ broad..... 25
24. Basidiospores broadly ellipsoid, $\geq 4 \mu\text{m}$ broad..... 26
25. Hymenophore smooth, pale orange to rosy; lamprocystidia $40-50 \times 6-7 \mu\text{m}$; basidiospores $< 2.5 \mu\text{m}$ wide; from Argentina..... *P. erubescens*
25. Hymenophore smooth to tuberculate, pale white to gray; lamprocystidia $60-90 \times 10-20 \mu\text{m}$; basidiospores $\geq 2.5 \mu\text{m}$ wide; from Northern Hemisphere..... *P. gigantea*
26. Lamprocystidia $< 10 \mu\text{m}$ wide; from South America..... *P. galochroa*
26. Lamprocystidia $> 10 \mu\text{m}$ wide; from Asia..... *P. darjeelingensis*

Conclusions

An in-depth study of the phylogeny and taxonomy of the genus *Phlebiopsis* was conducted based on both of morphological and molecular evidence. The results show that species of *Phlebiopsis* formed a monophyletic clade in the Phanerochaetaceae but are morphologically heterogeneous with *Australohydnum*, *Castanoporus* and *Hjortstamia* as its synonyms. The species diversity of the genus is high in subtropical and tropical areas with several new species described from East Asia. The present study also show that the DNA sequence data are very useful in exploring cryptic taxa and diversity of corticioid fungi, a large and widely distributed group poorly studied while compared with mushrooms and polypores.

List Of Abbreviations

ITS: internal transcribed spacer; nrLSU: nuclear ribosomal large subunit; BJFC: herbarium of Beijing Forestry University, Beijing, China; CFMR: Centre for Forest Mycology Research, U.S. Forest Service, Madison, Wisconsin, U.S.A.; TNM: National Museum of Natural Science, Taichung, Taiwan, China; KOH: 2% (w/v) potassium hydroxide; IKI: Melzer's reagent; CB: cotton blue; IKI-: neither amyloid nor dextrinoid; CB-: acyanophilous; L: mean spore length; W: mean spore width; Q: L/W ratio, n (a/b): number of spores

(a) measured from number of specimens (b); CTAB: cetyltrimethylammonium bromide; DNA: deoxyribonucleic acid; PCR: polymerase chain reaction; MP: maximum parsimony; ML maximum likelihood; BI: Bayesian inference; TBR: tree-bisection reconnection; BPP: Bayesian posterior probability.

Declarations

Ethics approval and consent to participate

Not applicable.

Adherence to national and international regulations

Not applicable.

Consent for publication

Not applicable.

Availability of data and materials

Details of the availability of the data and materials used in this study can be found within the text. DNA sequences were submitted to NCBI Genbank database (Table 1). Alignments were deposited at TreeBase (submission ID: 26529 for Phanerochaetaceae ITS-LSU, 26530 for *Phlebiopsis* ITS). Voucher specimens are deposited in the herbaria listed in the materials and methods section.

Competing interests

The authors declare that they have no competing interests

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Authors' contributions

Shuang-Hui He designed the research, collected most of the specimens, and wrote the text. Ya-Nan Zhao performed the phylogenetic analyses and did most of the measurement, descriptions and illustrations. Karen K. Nakasone loaned and examined type specimens of some related species, and revised language of the text. Che-Chi Chen provided with some specimens and sequences. Shi-Liang Liu and Yi-Feng Cao helped in field trips and species illustrations.

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Figures

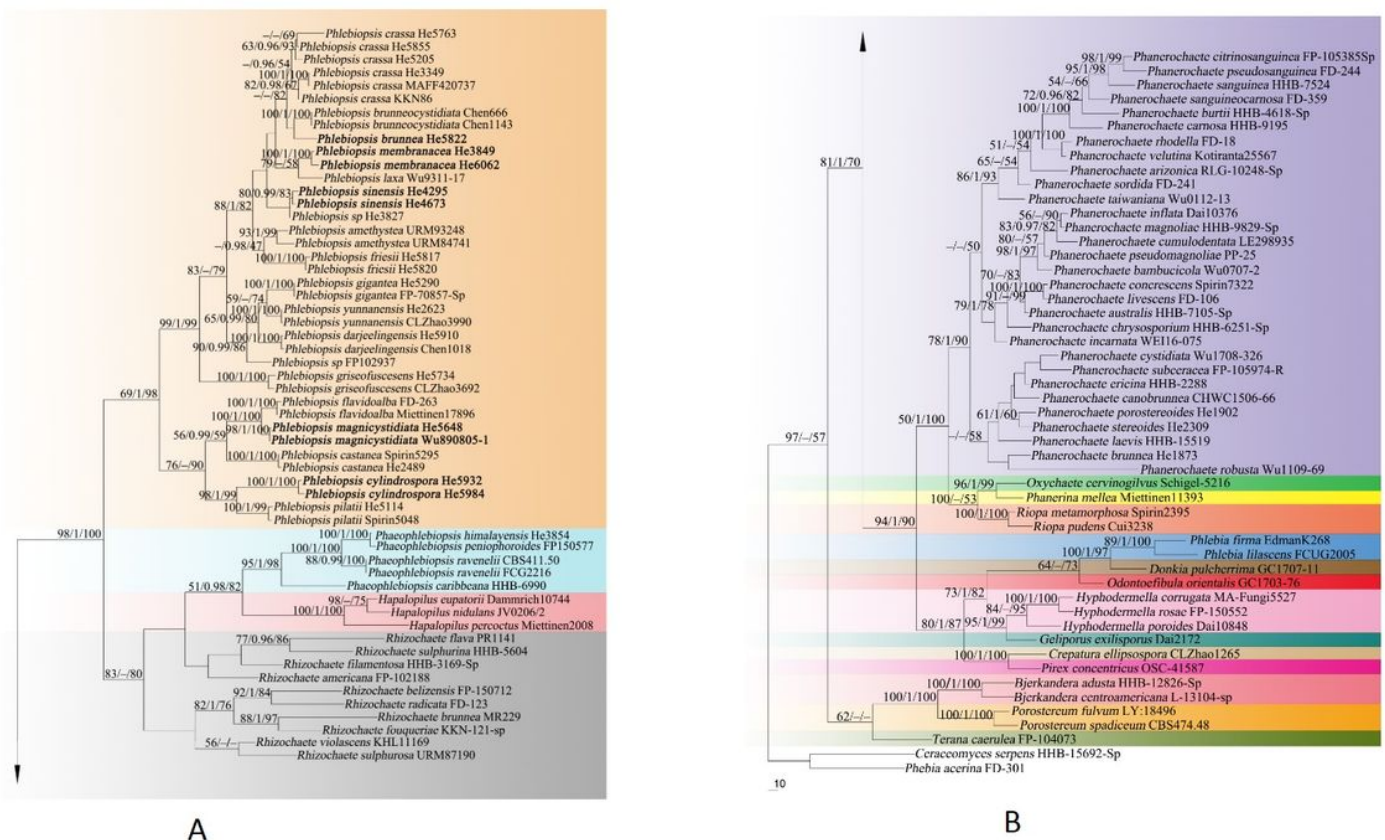


Figure 1

Phylogenetic tree from maximum parsimony analysis from the concatenated ITS and nrLSU sequences of Phanerochaetaceae taxa. Branches are labelled with parsimony bootstrap values ($\geq 50\%$, first), Bayesian posterior probabilities (≥ 0.95 , second) and likelihood bootstrap values ($\geq 50\%$, third). New species are set in bold.

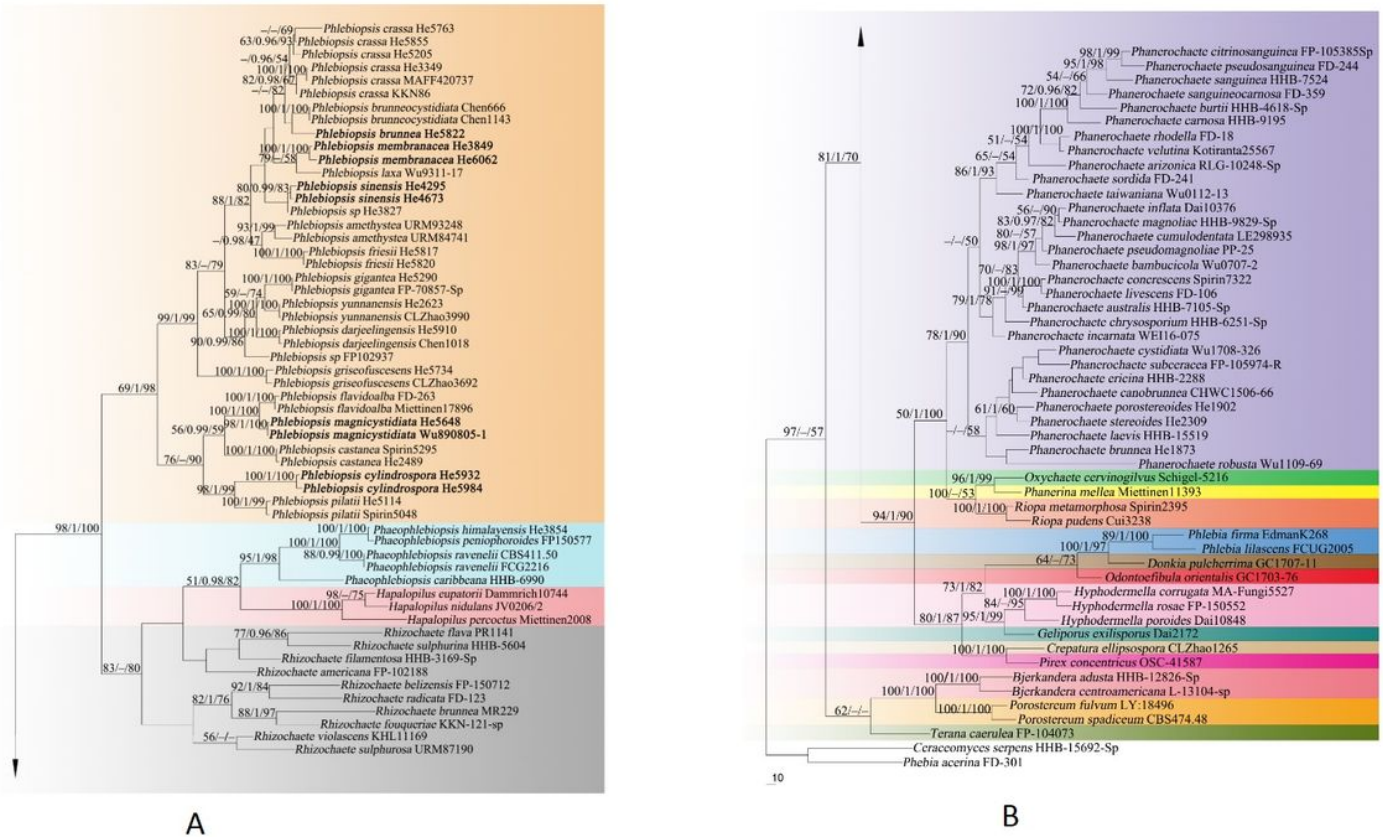


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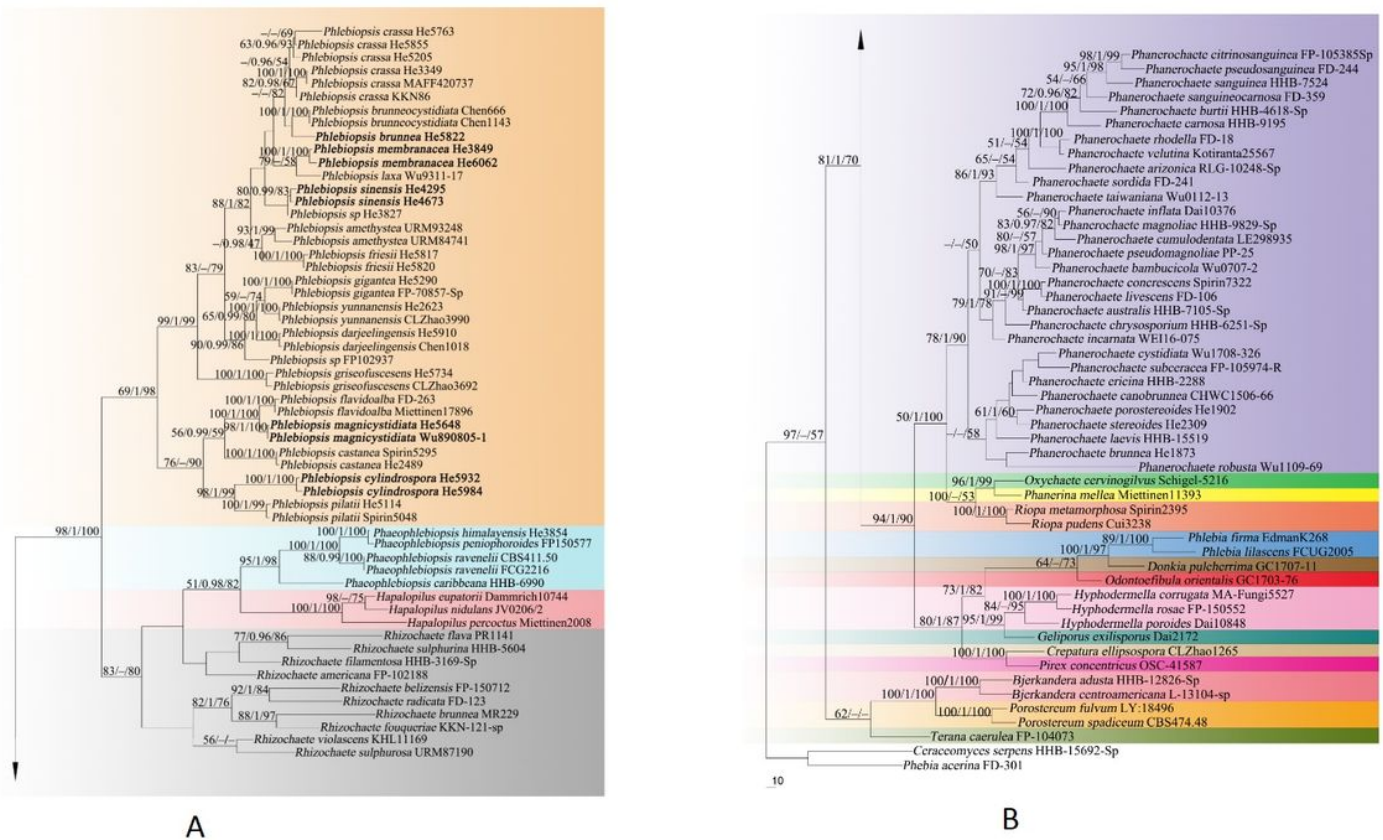
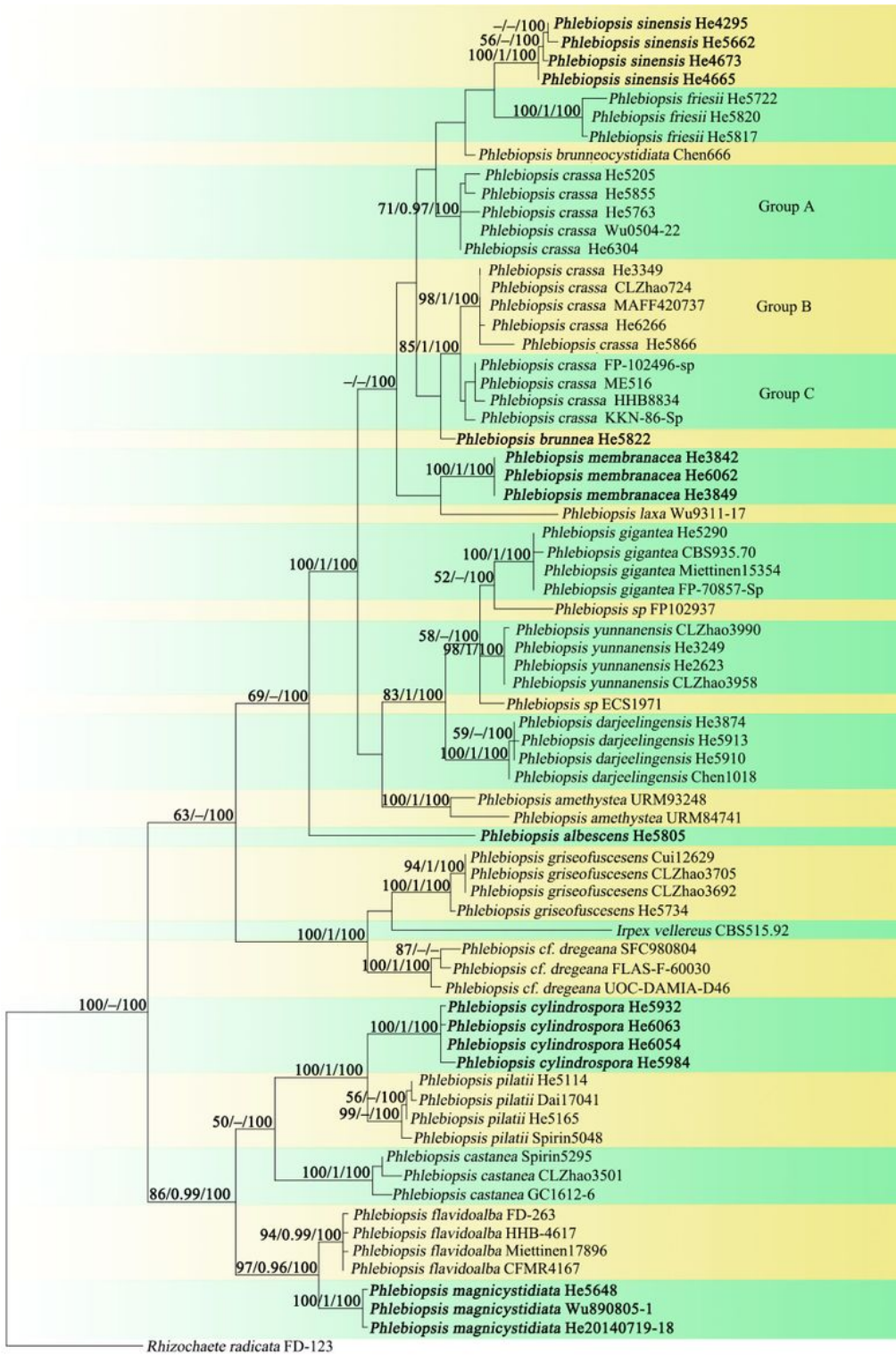


Figure 1

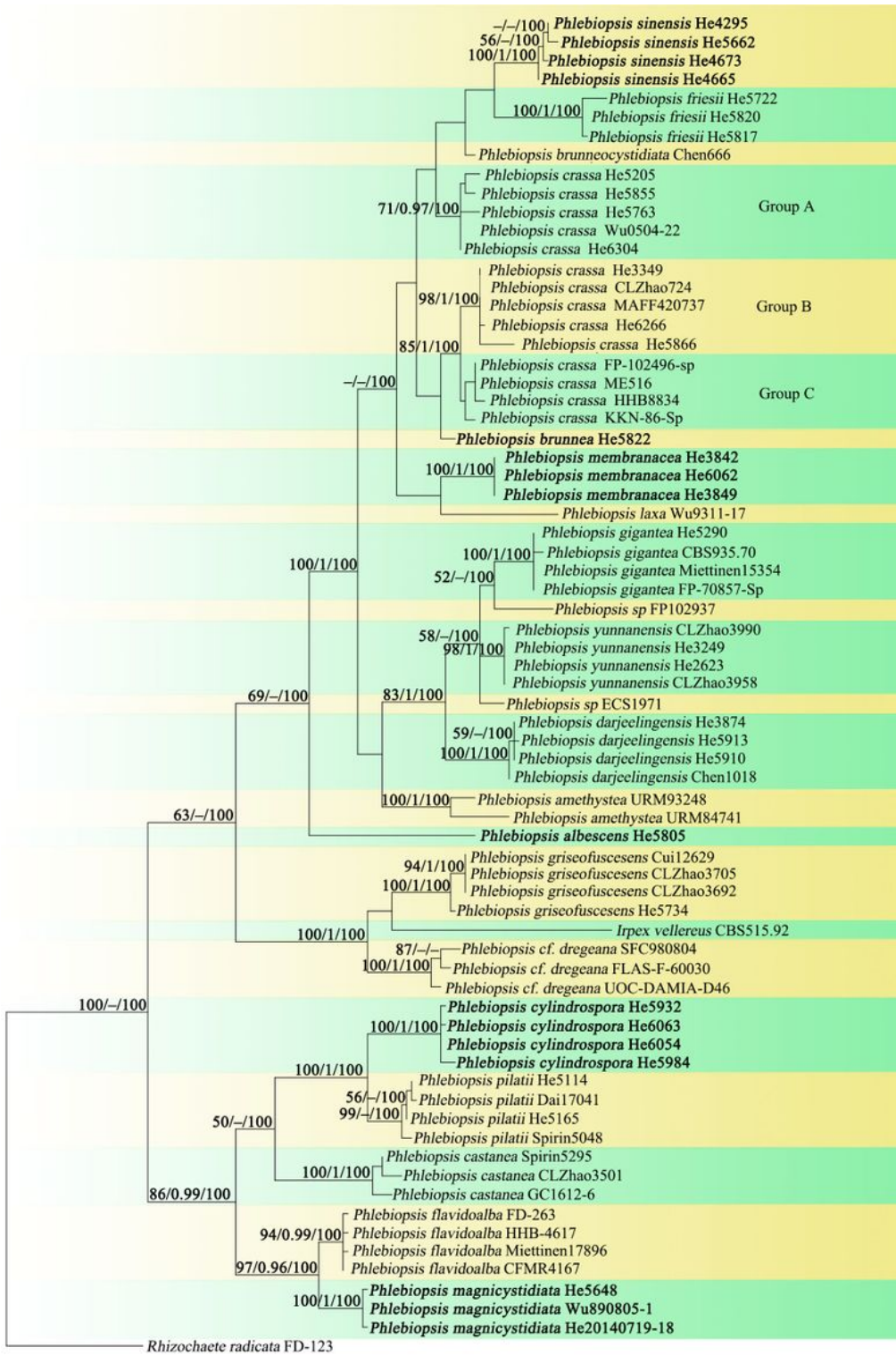
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Figure 2

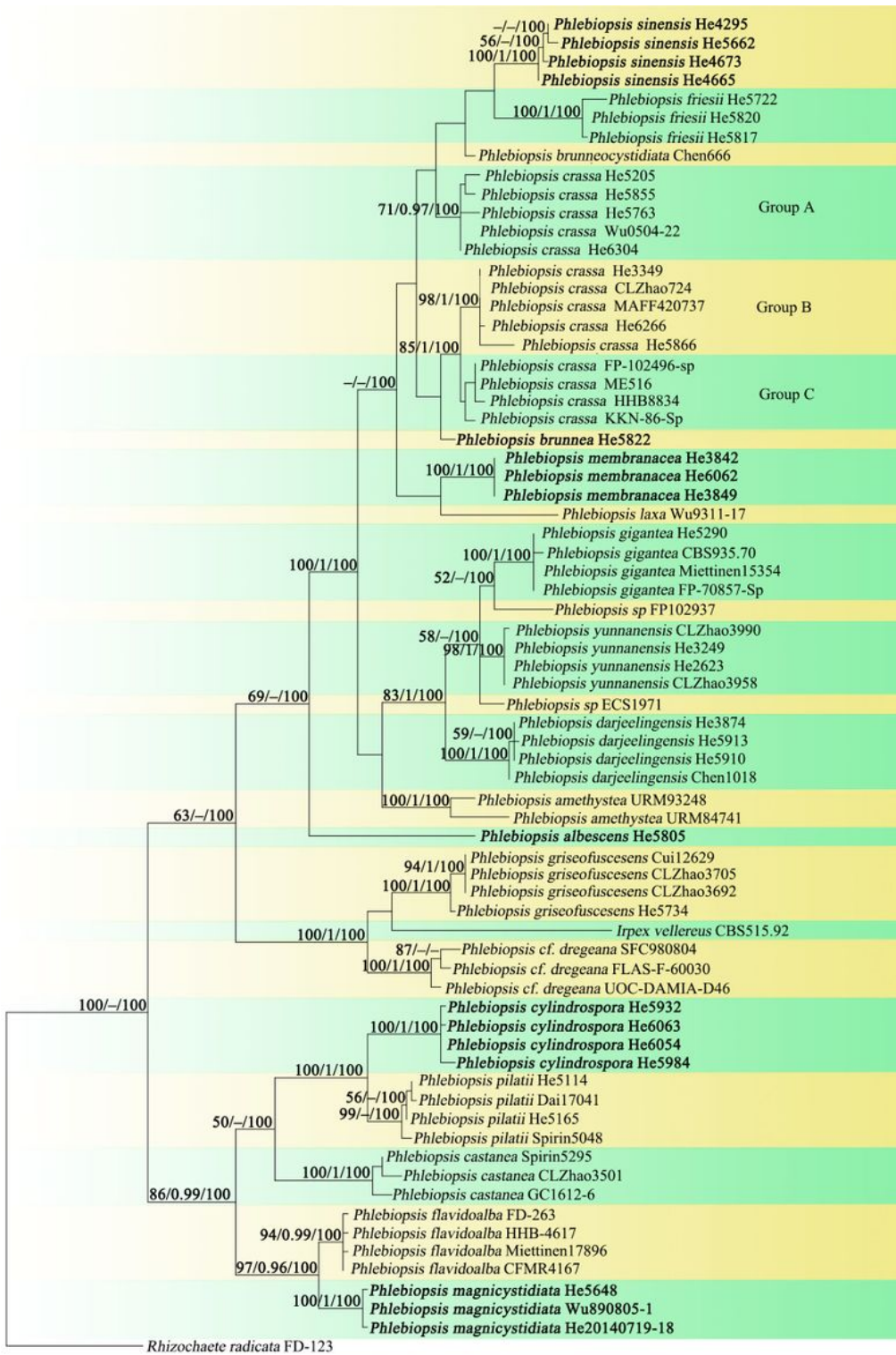
Phylogenetic tree obtained from maximum parsimony analysis of ITS sequence data of *Phlebiopsis*. Branches are labelled with parsimony bootstrap values ($\geq 50\%$, first), Bayesian posterior probabilities (≥ 0.95 , second) and likelihood bootstrap values ($\geq 50\%$, third). New species are set in bold.



10

Figure 2

Phylogenetic tree obtained from maximum parsimony analysis of ITS sequence data of *Phlebiopsis*. Branches are labelled with parsimony bootstrap values ($\geq 50\%$, first), Bayesian posterior probabilities (≥ 0.95 , second) and likelihood bootstrap values ($\geq 50\%$, third). New species are set in bold.



10

Figure 2

Phylogenetic tree obtained from maximum parsimony analysis of ITS sequence data of *Phlebiopsis*. Branches are labelled with parsimony bootstrap values ($\geq 50\%$, first), Bayesian posterior probabilities (≥ 0.95 , second) and likelihood bootstrap values ($\geq 50\%$, third). New species are set in bold.

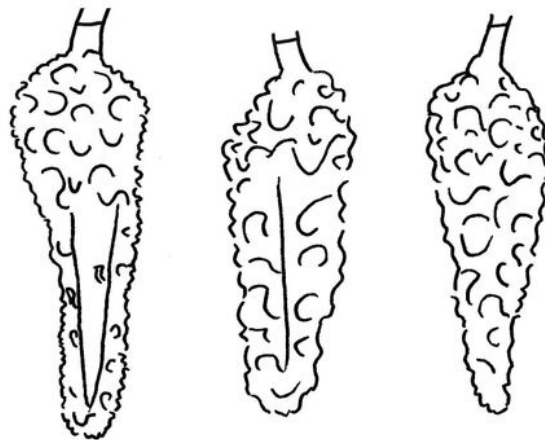
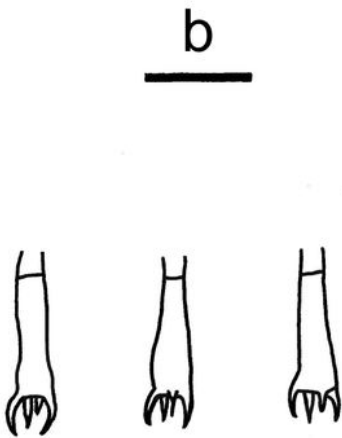
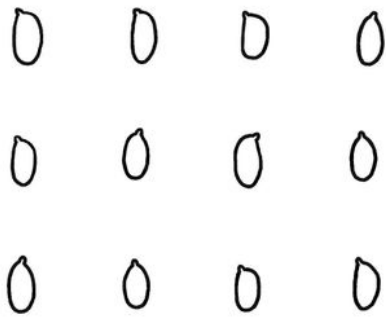
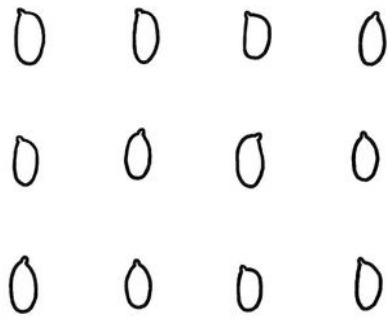


Figure 3

Phlebiopsis albescens (from the holotype He 5805; scale bars: a = 1 cm, b–e = 10 μ m). a. basidiomata; b. basidiospores; c. basidia; d. basidioles; e. lamprocystidia.



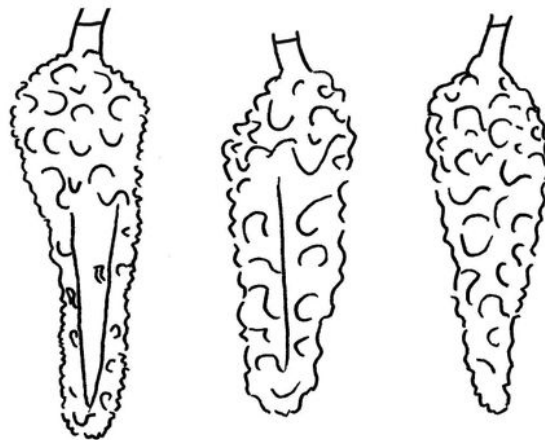
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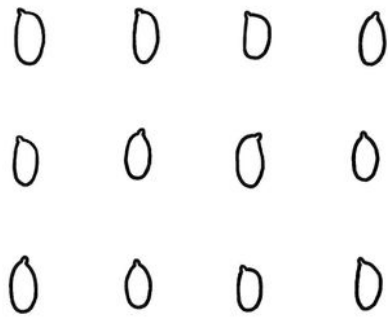
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e

Figure 3

Phlebiopsis albescens (from the holotype He 5805; scale bars: a = 1 cm, b–e = 10 μ m). a. basidiomata; b. basidiospores; c. basidia; d. basidioles; e. lamprocystidia.



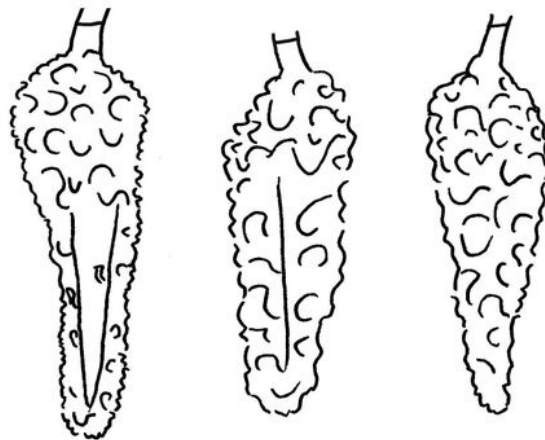
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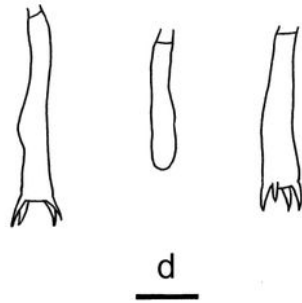
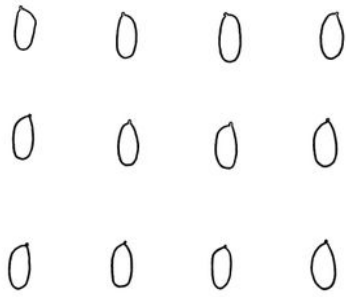
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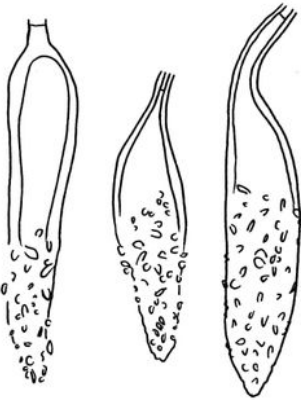
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Figure 3

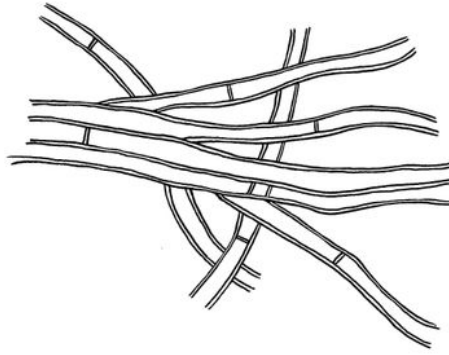
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b



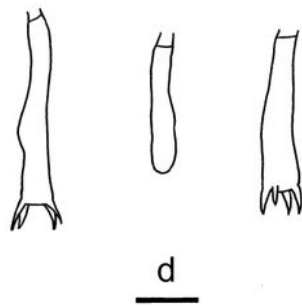
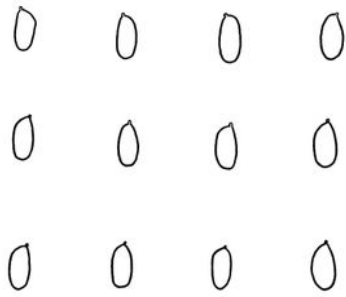
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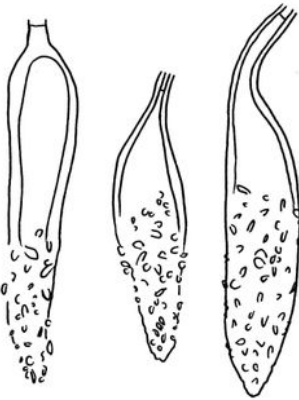
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Figure 4

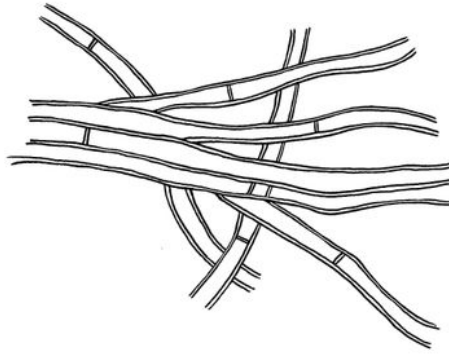
Phlebiopsis brunnea (from the holotype He 5822; scale bars: a = 1 cm, b–e = 10 μ m). a. basidiomata; b. basidiospores; c. lamprocystidia; d. basidia and basidiole; e. hyphae from subiculum.



b



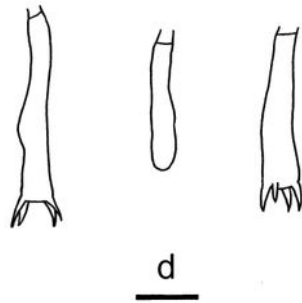
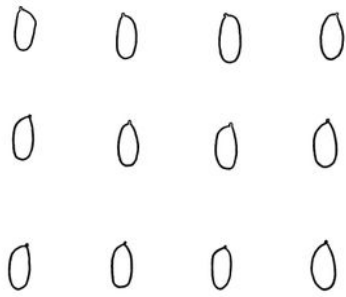
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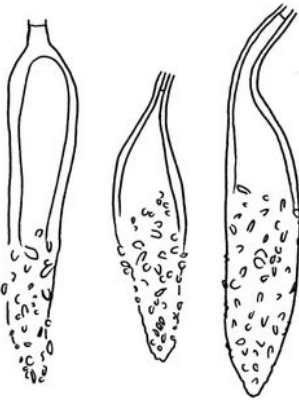
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Figure 4

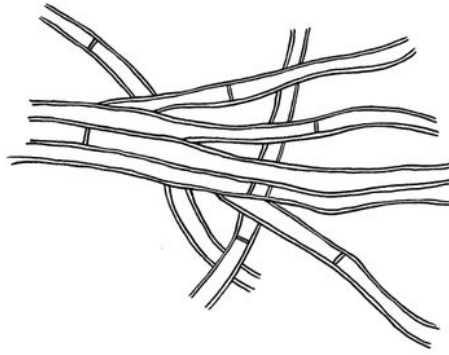
Phlebiopsis brunnea (from the holotype He 5822; scale bars: a = 1 cm, b–e = 10 μ m). a. basidiomata; b. basidiospores; c. lamprocystidia; d. basidia and basidiole; e. hyphae from subiculum.



b



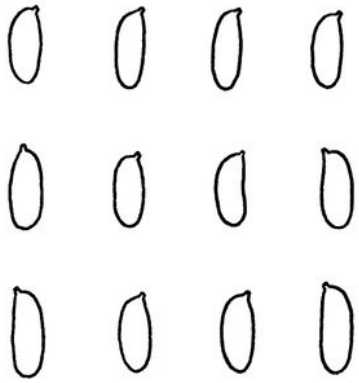
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Figure 4

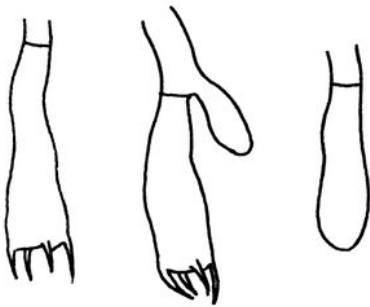
Phlebiopsis brunnea (from the holotype He 5822; scale bars: a = 1 cm, b–e = 10 μ m). a. basidiomata; b. basidiospores; c. lamprocystidia; d. basidia and basidiole; e. hyphae from subiculum.



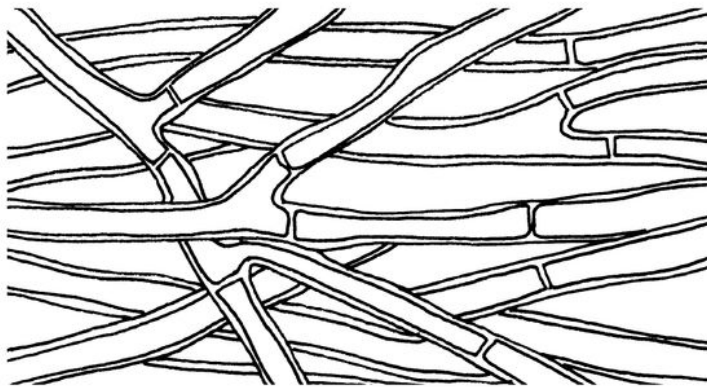
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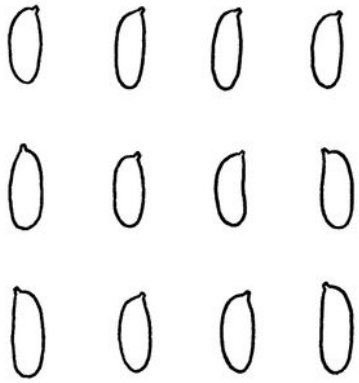


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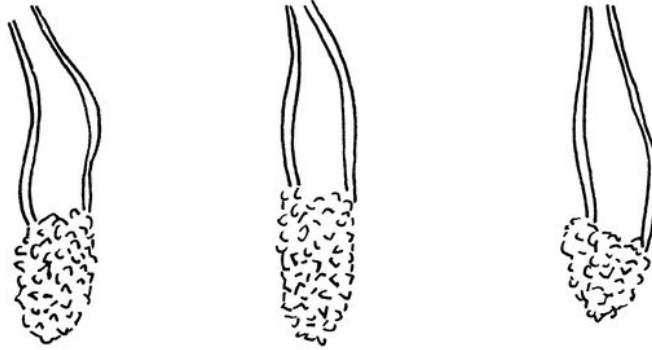


Figure 5

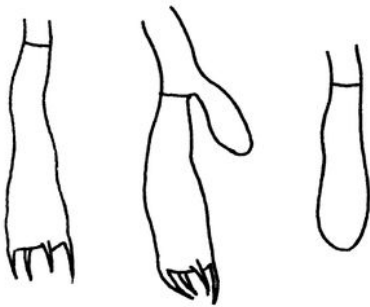
Phlebiopsis cylindrospora (from the holotype He 5984; scale bars: a = 1 cm, b–e = 10 μ m). a. basidiomata; b. basidiospores; c. basidia and basidiole; d. lamprocystidia; e. hyphae from subiculum.



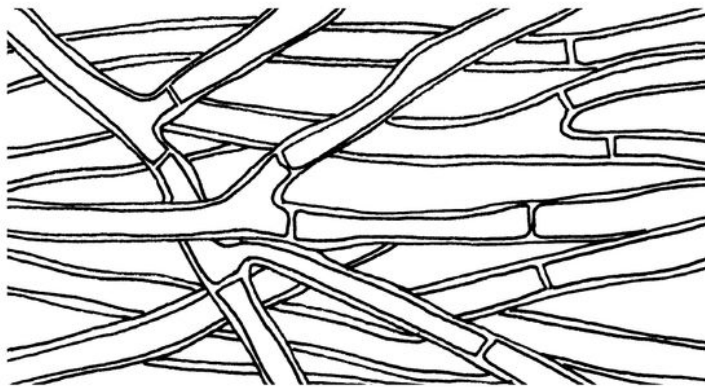
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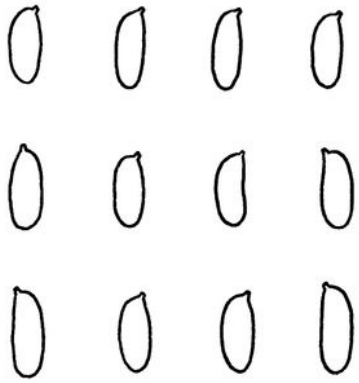


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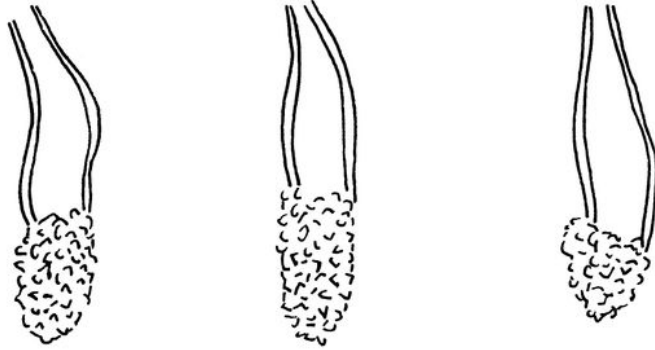


Figure 5

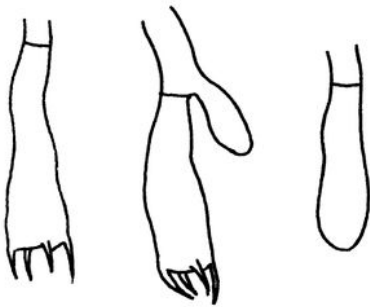
Phlebiopsis cylindrospora (from the holotype He 5984; scale bars: a = 1 cm, b–e = 10 μ m). a. basidiomata; b. basidiospores; c. basidia and basidiole; d. lamprocystidia; e. hyphae from subiculum.



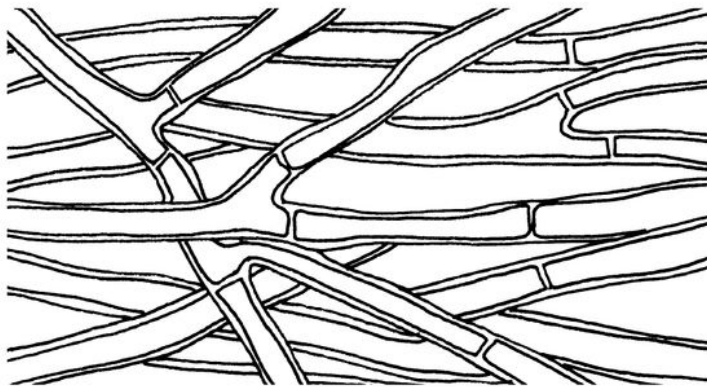
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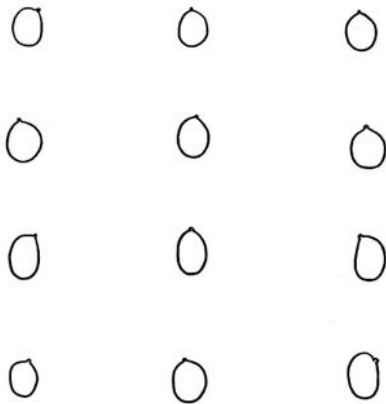


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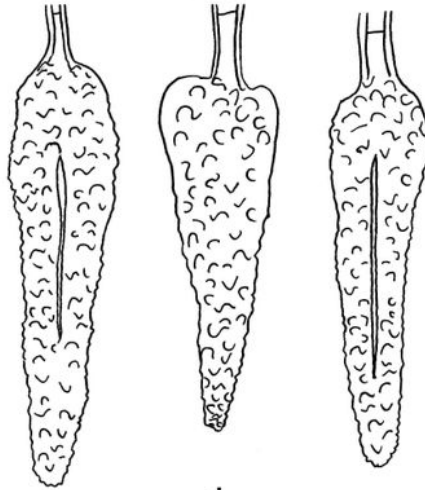


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Phlebiopsis cylindrospora (from the holotype He 5984; scale bars: a = 1 cm, b–e = 10 μ m). a. basidiomata; b. basidiospores; c. basidia and basidiole; d. lamprocystidia; e. hyphae from subiculum.



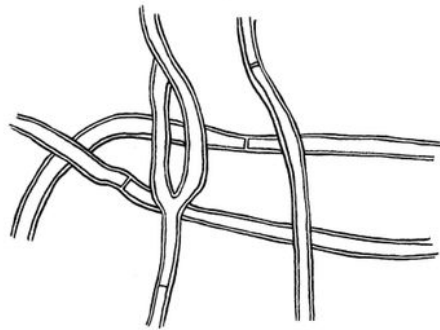
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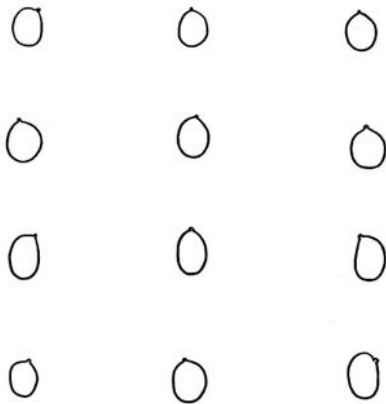
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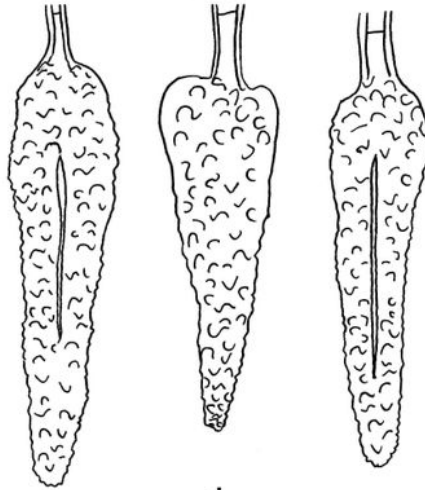
e

Figure 6

Phlebiopsis magnicystidiata (a from He 20140719-18, b–e from the holotype He 5648; scale bars: a = 1 cm, b–e = 10 μ m). a. basidiomata; b. basidiospores; c. basidia and basidiole; d. lamprocystidia; e. hyphae from subiculum.



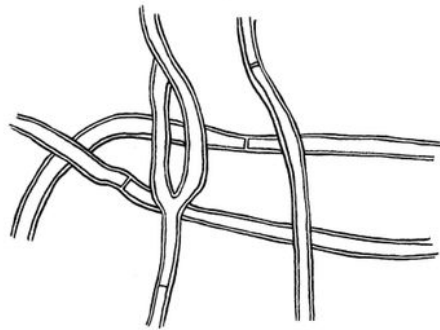
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d



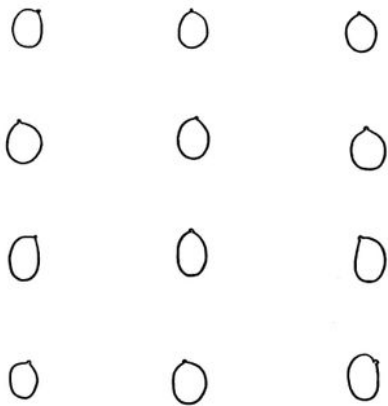
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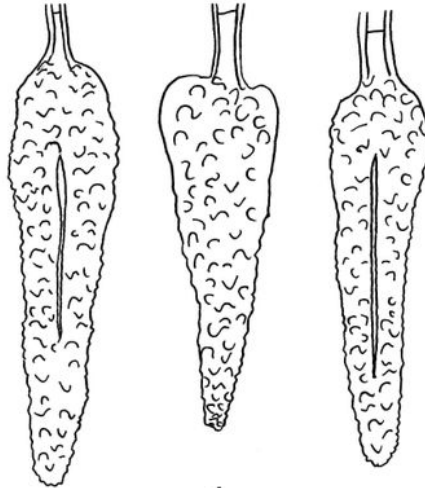
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Figure 6

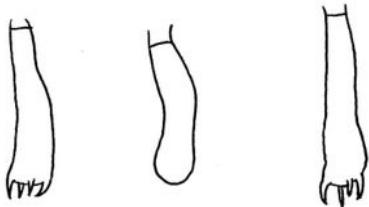
Phlebiopsis magnicystidiata (a from He 20140719-18, b–e from the holotype He 5648; scale bars: a = 1 cm, b–e = 10 μ m). a. basidiomata; b. basidiospores; c. basidia and basidiole; d. lamprocystidia; e. hyphae from subiculum.



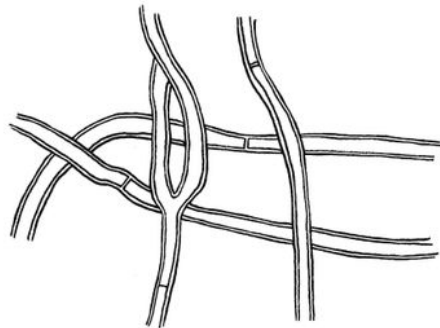
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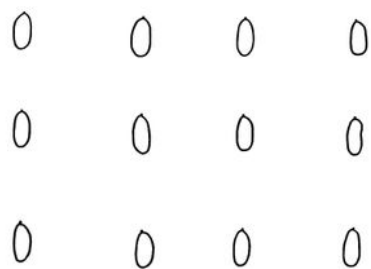
c



e

Figure 6

Phlebiopsis magnicystidiata (a from He 20140719-18, b–e from the holotype He 5648; scale bars: a = 1 cm, b–e = 10 μ m). a. basidiomata; b. basidiospores; c. basidia and basidiole; d. lamprocystidia; e. hyphae from subiculum.



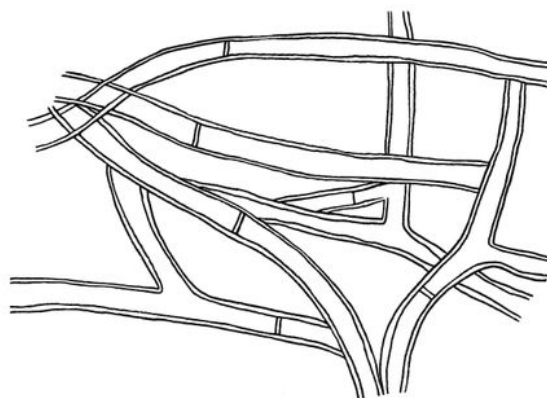
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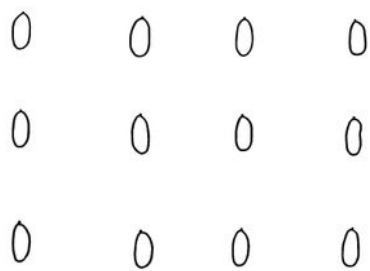
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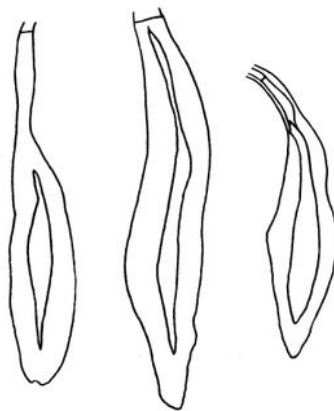
e

Figure 7

Phlebiopsis membranacea (a from He 3842, b–e from the holotype He 3849; scale bars: a = 1 cm, b–e = 10 μ m). a. basidiomata; b. basidiospores; c. basidia and basidiole; d. hymental cystidia; e. hyphae from subiculum.



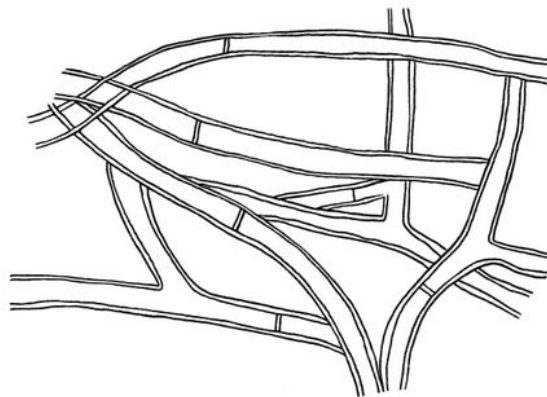
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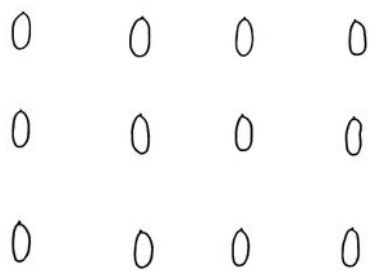
c



e

Figure 7

Phlebiopsis membranacea (a from He 3842, b–e from the holotype He 3849; scale bars: a = 1 cm, b–e = 10 μ m). a. basidiomata; b. basidiospores; c. basidia and basidiole; d. hymental cystidia; e. hyphae from subiculum.



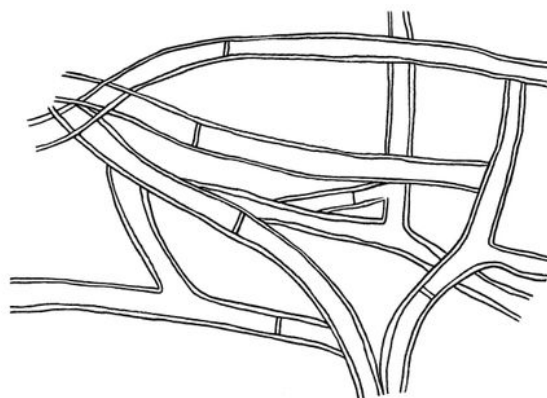
b



d



c



e

Figure 7

Phlebiopsis membranacea (a from He 3842, b–e from the holotype He 3849; scale bars: a = 1 cm, b–e = 10 μ m). a. basidiomata; b. basidiospores; c. basidia and basidiole; d. hymenial cystidia; e. hyphae from subiculum.

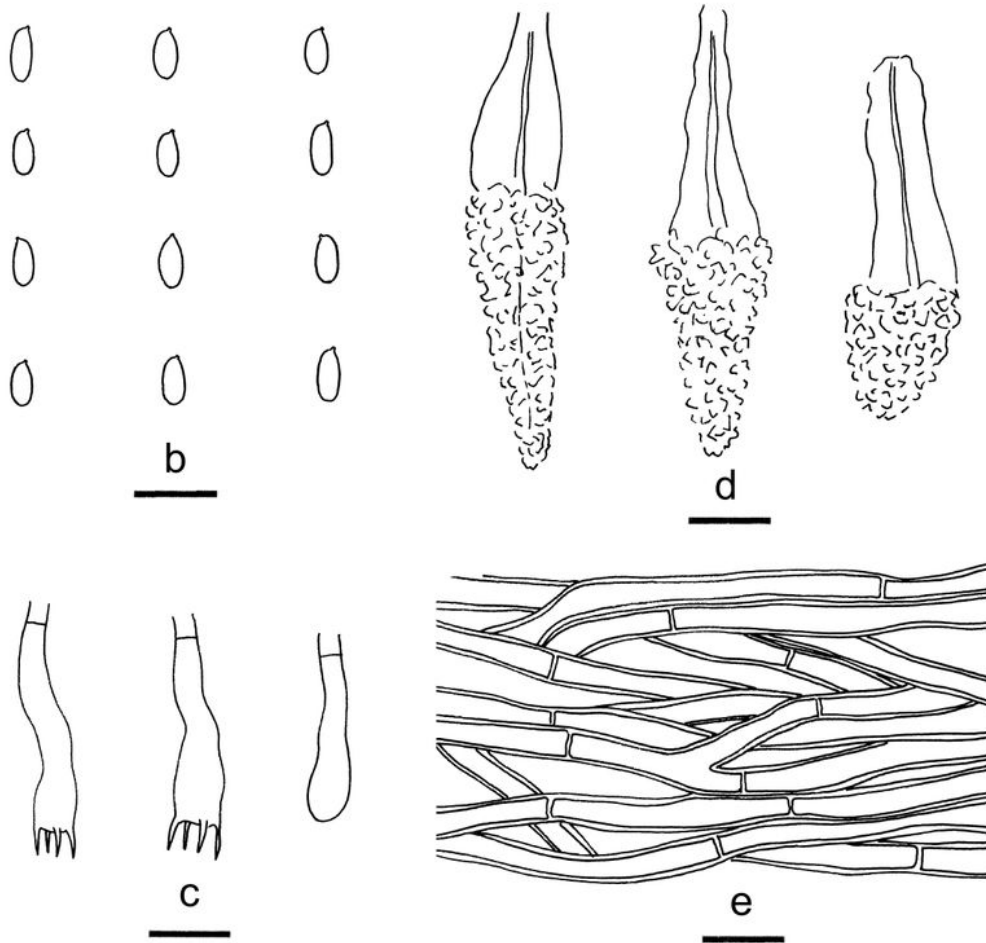


Figure 8

Phlebiopsis sinensis (from He 4673; scale bars: a = 1 cm, b–e = 10 μ m). a. basidiomata; b. basidiospores; c. basidia and basidiole; d. lamprocystidia; e. hyphae from subiculum.

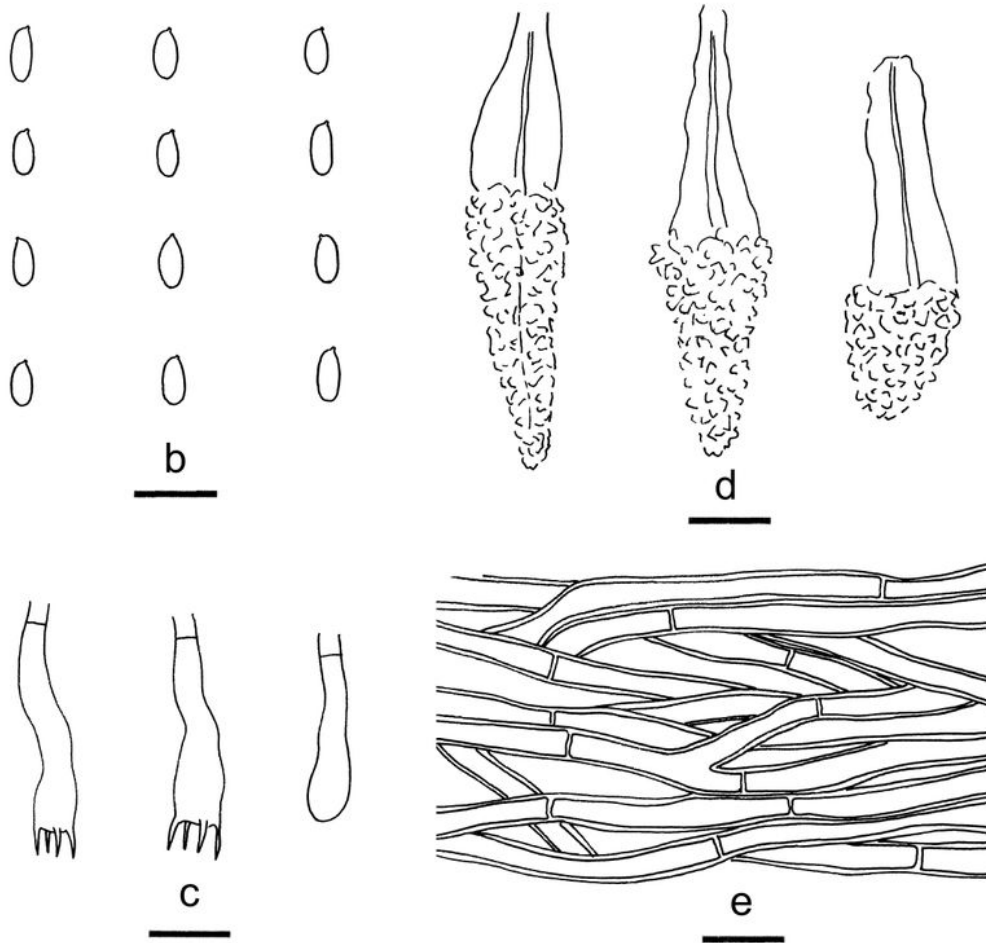


Figure 8

Phlebiopsis sinensis (from He 4673; scale bars: a = 1 cm, b–e = 10 μ m). a. basidiomata; b. basidiospores; c. basidia and basidiole; d. lamprocystidia; e. hyphae from subiculum.

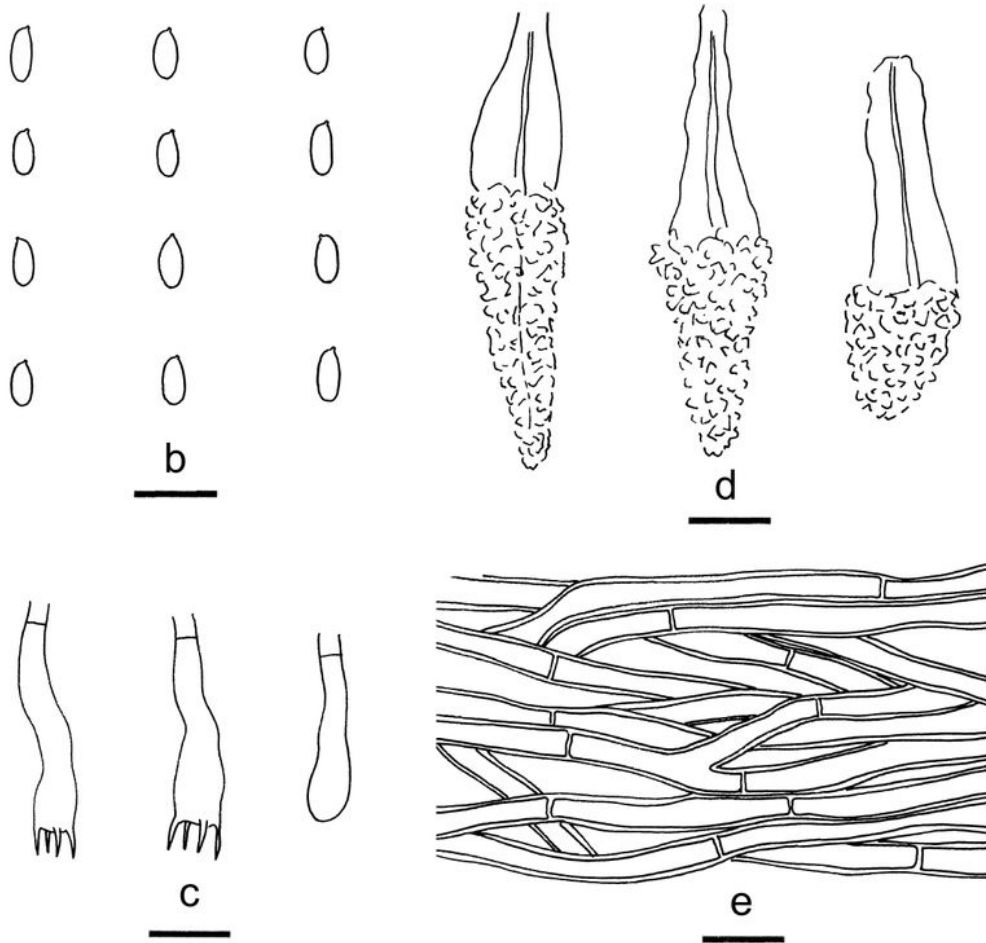


Figure 8

Phlebiopsis sinensis (from He 4673; scale bars: a = 1 cm, b–e = 10 μ m). a. basidiomata; b. basidiospores; c. basidia and basidiole; d. lamprocystidia; e. hyphae from subiculum.

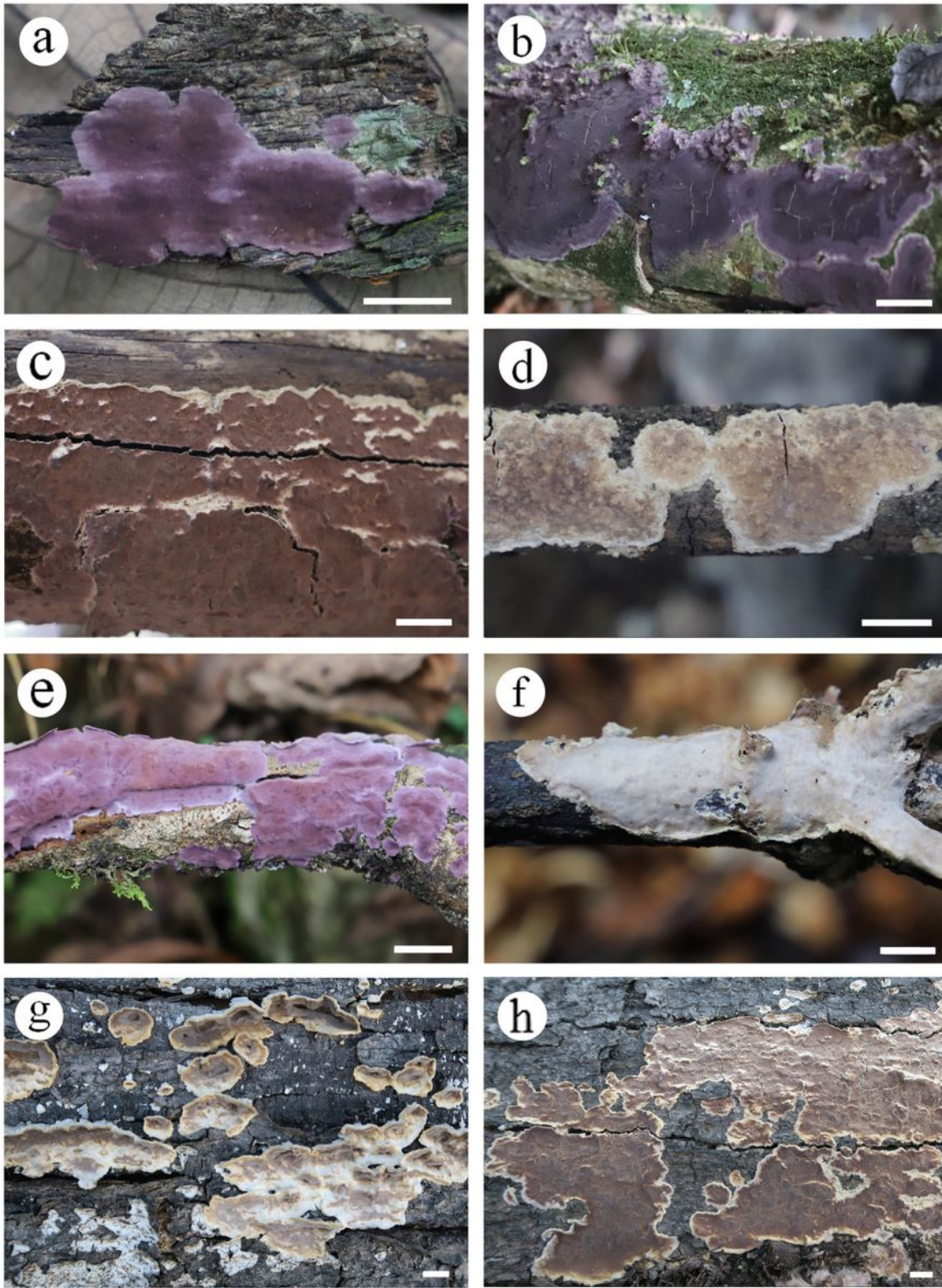


Figure 9

Basidiomata of *Phlebiopsis crassa* s.l. (a–d: *P. crassa* group A, e–h: *P. crassa* group B; scale bars: a–h = 1 cm). a. He 5855; b. He 6303; c. He 6300; d. He 6301; e. He 6266; f. He 5866; g–h. He 3349.

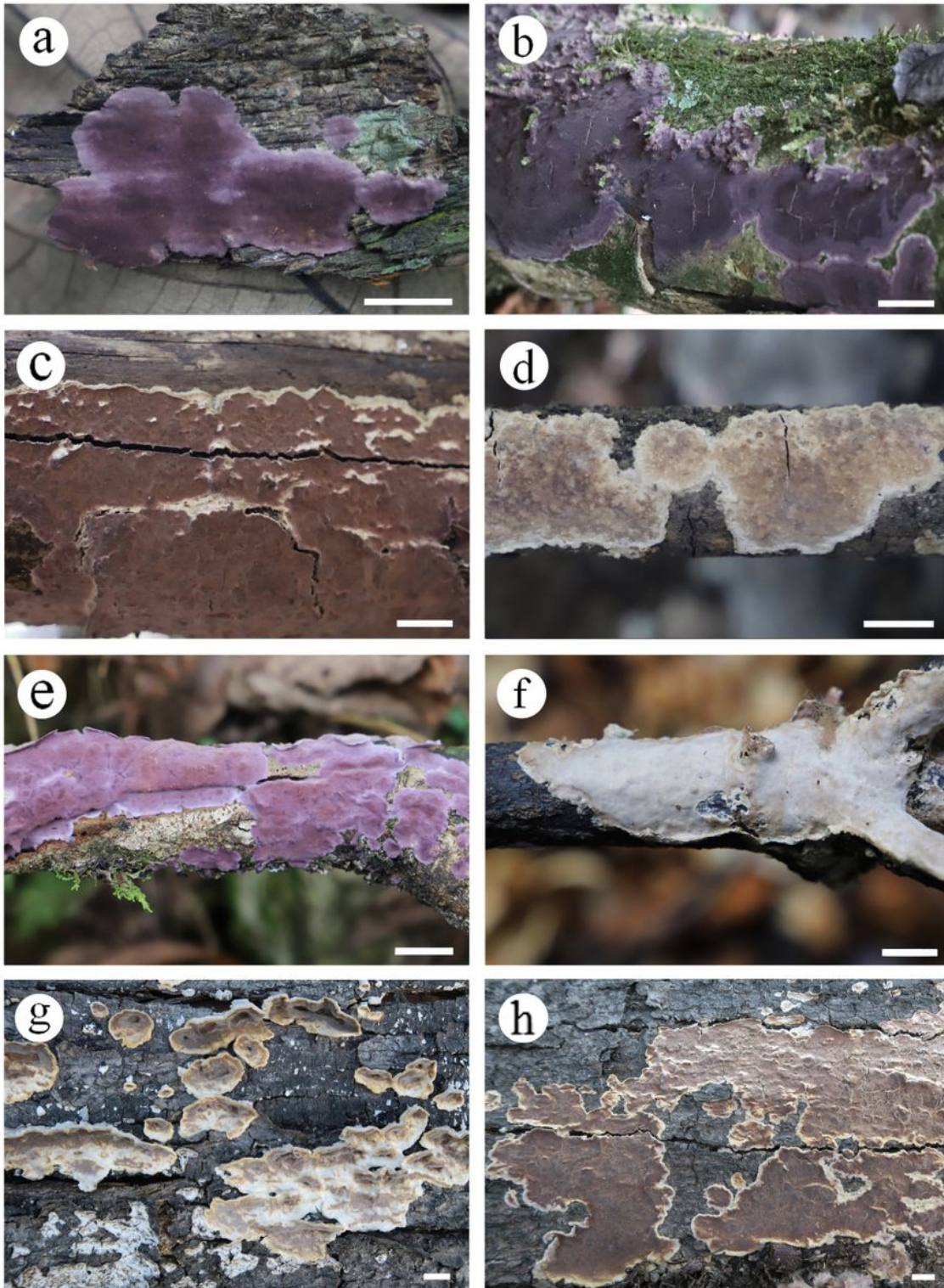


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Basidiomata of *Phlebiopsis crassa* s.l. (a–d: *P. crassa* group A, e–h: *P. crassa* group B; scale bars: a–h = 1 cm). a. He 5855; b. He 6303; c. He 6300; d. He 6301; e. He 6266; f. He 5866; g–h. He 3349.

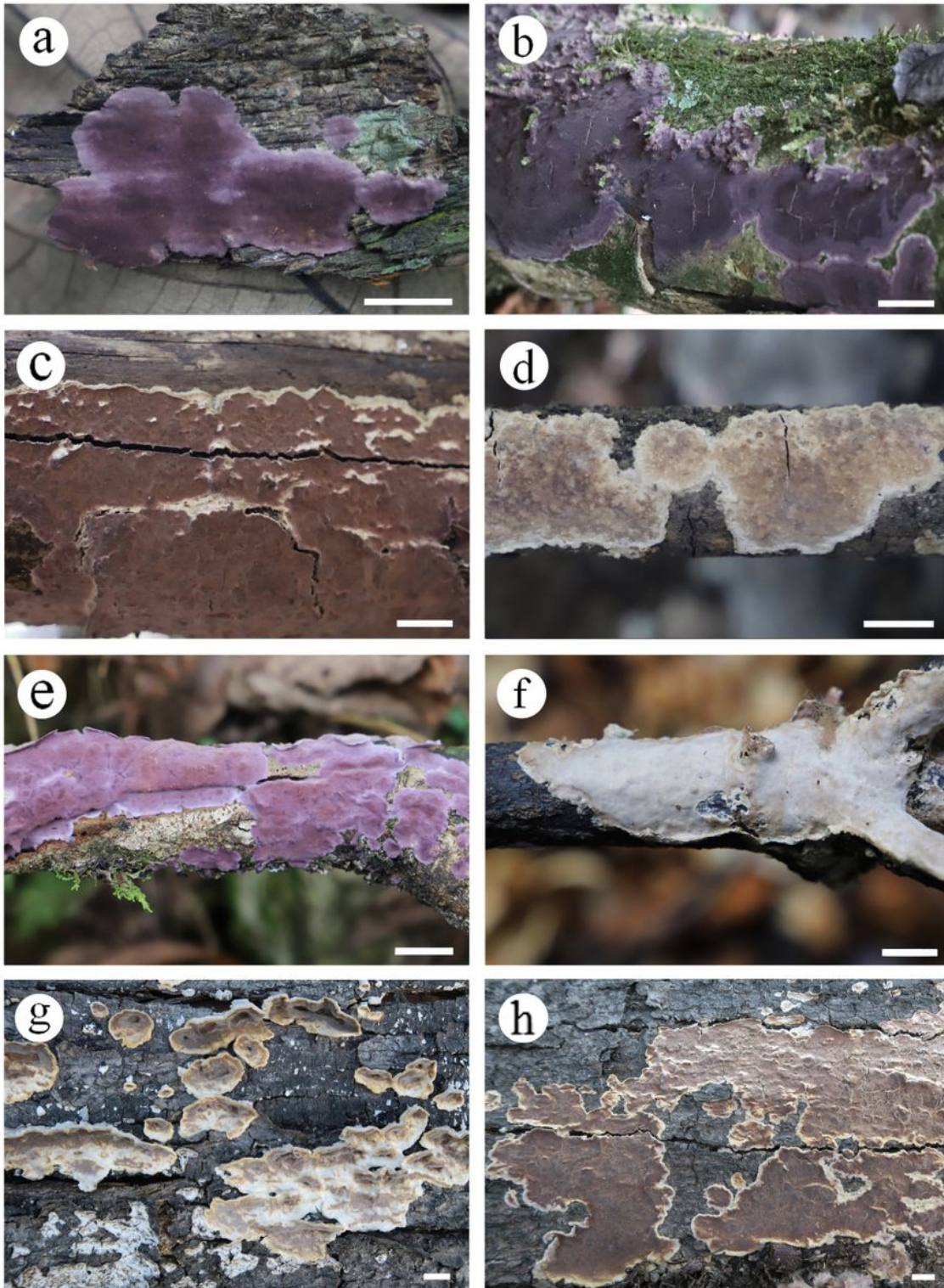


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