

Table S1. Specimens used in phylogenetic analysis and GenBank codes. Newly sequenced collections are in bold.

Species	Voucher/ Strain	GenBank Accession no.						Reference
		ITS	nrLSU	RPB2	TEF1- α	RPB1	nrSSU	
<i>Asproinocybe sinensis</i>	HMJAU59025 Holotype	OK377049 *#	OK377052 *#	OK625400	OK625330	OK625397	OK377041	Present study
<i>Asproinocybe sinensis</i>	HMJAU59026	OK377048 *#	OK377051 *#	OK625401	OK625331	OK625398	OK377040	Present study
<i>Asproinocybe sinensis</i>	M2020081289	-	OK576386 *#	-	OK625337	-	OK624826	Present study
<i>Asproinocybe daleyae</i>	PDD 106796	MN275025*#	MN275033*#	-	-	-	-	[6]
<i>Asproinocybe lyophylloides</i>	PERTH 4163559	MN275018*#	MN275027*#	-	-	-	-	[6]
<i>Asproinocybe lyophylloides</i>	MEL2432747	MN275015*#	MN275028*#	-	-	-	-	[6]
<i>Asproinocybe lyophylloides</i>	PERTH 8477001	MN275022*#	MN275030*#	-	-	-	-	[6]
<i>Asproinocybe lyophylloides</i>	PERTH E6292	MN275023*#	MN275031*#	-	-	-	-	[6]
<i>Ampulloclitocybe clavipes</i>	AFTOL-ID 542	AY789080	AY639881	AY780937	AY881022	AY788848	-	[35,36]
<i>Anupama indica</i>	AMH10033	MH989590	MH989586	MH992117	-	-	-	[37,38]
<i>Anupama indica</i>	CAL1725	MH989587	MH989583	-	-	-	-	[37,38]
<i>Asterophora lycoperdoides</i>	CBS170.86	AF357037	-	DQ367431	DQ367424	EF421021	AF357109	[32]
<i>Bonomyces afrosinopicus</i>	LIP:LYK13040015	MG696613	MG696624	MG702593	MG702590	-	MG696621	[36]
<i>Callistosporium graminicolor</i>	AFTOL978	DQ484065	AY745702	-	-	-	-	[35,36]
<i>Callistosporium imbricatum</i>	TJB9847	HM105568	HM105568	HM105567	-	-	-	[35,38]
<i>Callistosporium imbricatum</i>	SFSU:DED 8232	MF100955	MN017456	-	-	-	MN017570	[38]
<i>Callistosporium luteoolivaceum</i>	18231 (AMB)	MN017518*#&	MN017459*#&	-	-	-	-	[38]
<i>Callistosporium praemultifolium</i>	SFSU:DED 8238	MF100956	MN017464	MN018844	-	-	MN017575	[38]
<i>Callistosporium xanthophyllum</i>	IB19770276	AF325667*#&	AF261406*#&	-	-	-	-	[37]
<i>Catathelasma imperiale</i>	CORT:11CA01A	-	KR869941	KC816994	KC816900	-	KR869913	NCBI
<i>Catathelasma imperiale</i>	HKAS 84315	MK909096	MK909113	-	MK909083	-	-	NCBI
<i>Cleistocybe carneogrisea</i>	TENN:063842	-	HQ728527	-	-	-	HQ728528	[38]
<i>Cleistocybe vernalis</i>	ADP 050506 (WTU)	EF416917	EF416916	-	-	-	-	[38]
<i>Clitocybe candicans</i>	AFTOL-ID 541	DQ202268	AY645055	DQ385881	DQ408149	DQ447891	-	[32]
<i>Clitocybe dealbata</i>	IE-BSG-HC95.cp3	AF357061	AF223175	DQ825407	EF421080	DQ825414	AF357138	[32,34]
<i>Clitocybe eccentrica</i>	G0084/ PBM3851	MG663292	MK277727	-	-	-	-	NCBI

Table S1. Cont.

Species	Voucher/ Strain	GenBank Accession no.						
		ITS	nrLSU	RPB2	TEF1- α	RPB1	nrSSU	Reference
<i>Clitocybe subditopoda</i>	AFTOL-ID 533	DQ202269	AY691889	AY780942	DQ408150	DQ447892	-	[32]
<i>Clitopilus brunneiceps</i>	KUN-HKAS 80211	MN061293	MN065682	MN148121	MN166232	-	-	[22]
<i>Clitopilus brunneiceps</i>	KUN-HKAS 104510	MN061295	MN065684	MN148123	MN166234	-	-	[22]
<i>Clitopilus rugosiceps</i>	KUN-HKAS73232	MN061305	MN065695	MN148132	MN166244	-	-	[22]
<i>Clitopilus rugosiceps</i>	KUN-HKAS 107044	MT345046	MT345051	-	-	-	-	[22]
<i>Clitopilus yunnanensis</i>	KUN-HKAS104518	MN061308	MN065698	MN148136	MN166247	-	-	[22]
<i>Clitopilus yunnanensis</i>	HMJAU 24677	MN061309	MN065699	MN148116	MN166248	-	-	[22]
<i>Clitopilus sinoapalus</i>	KUN-HKAS101191	MN061322	MN065711	MN148151	MN166261	-	-	[22]
<i>Clitopilus sinoapalus</i>	KUN-HKAS 82230	MN061320	MN065712	MN148148	-	-	-	[22]
<i>Clitopilus umbilicatus</i>	KUN-HKAS80310	MN061324	MN065716	MN148153	MN166263	-	-	[22]
<i>Clitopilus umbilicatus</i>	KUN-HKAS 80945	MN061326	MN065718	MN148155	MN166265	-	-	[22]
<i>Clitopilus prunulus</i>	VHAs07/02	EF421107	EF421092	DQ825408	EF421086	DQ825416	EF421097	[32]
<i>Clitocella orientalis</i>	KUN-HKAS78763	-	MN065728	MN148165	MN166276	-	-	[22]
<i>Clitocella orientalis</i>	KUN-HKAS 77899	-	MN065725	MN148162	MN166273	-	-	[22]
<i>Clitocella mundula</i>	O:O-F71544	-	-	KC816950	KC816860	-	-	[22]
<i>Clitocella mundula</i>	HMJAU 7275	MN061331	MN065723	MN148160	MN166271	-	-	[22]
<i>Clitocella fallax</i>	O:O-F88953	-	-	KC816936	KC816845	-	-	[22]
<i>Clitocella fallax</i>	K: K(M) 116541	-	-	KC816938	KC816847	-	-	[22]
<i>Clitocella fallax</i>	CBS 605.79	AF357017	AF223165	-	-	-	AF357084	NCBI
<i>Clitopilopsis albida</i>	KUN-HKAS104519	MN061335	MN065730	MN148167	MN166278	-	-	[22]
<i>Clitopilopsis albida</i>	KUN-HKAS104520	MN061336	MN065731	MN148168	MN166279	-	-	[22]
<i>Collybia tuberosa</i>	DUKE-AOM191061	AF274376	AF261385	KP255481	KP255474	KP255479	-	[32]
<i>Dendrocollybia racemosa</i>	DUKE-DEB5575	DQ825425	AF042598	DQ825409	KP255476	DQ825417	KP255472	[32]
<i>Entocybe nitida</i>	24	KC710122	GQ289175	-	-	-	-	[6]
<i>Entocybe trachyospora</i>	414	KC710121	GQ289199	-	-	-	GQ289339	[31]
<i>Entoloma turbidum</i>	PRM915266	FJ824815	GQ244337	-	-	-	-	[6]
<i>Entoloma turbidum</i>	27	KC710060	GQ289201	GQ289269	-	-	GQ289341	[31]
<i>Entoloma hainanense</i>	GDGM 27990	NR_137783	NG_059212	-	-	-	-	NCBI

Table S1. Cont.

Species	Voucher/ Strain	GenBank Accession no.						Reference
		ITS	nrLSU	RPB2	TEF1- α	RPB1	nrSSU	
<i>Entoloma pallidocarpum</i>	GDGM 28828	JQ320106	JQ410331	JQ993080	-	-	JQ993074	NCBI
<i>Entoloma perbloomii</i>	71	KC710117	GQ289178	GQ289249	-	-	GQ289318	[31]
<i>Entoloma porphyrophaeum</i>	VHAs09/02	EF421111	-	-	EF421090	EF421053	EF421101	[32]
<i>Entoloma porphyrophaeum</i>	TB6957	-	AF261290	EF421020	-	-	-	[32]
<i>Entoloma sericeonitidum</i>	TB7144	EF421108	AF261315	EF421016	EF421087	EF421049	EF421098	[32]
<i>Entoloma sericeum</i>	VHAs03/02	DQ367430	DQ367423	DQ367435	DQ367428	DQ825424	EF421099	[32]
<i>Entoloma strictius</i>	DUKE-JM96/10	EF421109	AF042620	EF421017	EF421088	EF421050	EF421100	[32]
<i>Gerhardtia</i> sp.	HC01/025	EF421103	EF421091	EF420994	EF421060	EF421028	EF421093	[32]
<i>Guyanagarika pakaraimensis</i>	MCA4776	KX092085	KX092104	KX092139	-	-	-	[34,37]
<i>Guyanagarika anomala</i>	TH7419	KX092096	KX092110	KX092147	-	-	-	[37]
<i>Infundibulicybe geotropa</i>	ALV4344	KT122792	KT122793	-	-	-	-	[35,38]
<i>Infundibulicybe gibba</i>	AFTOL-ID 1508	DQ490635	DQ457682	DQ472727	-	DQ447913	-	[35]
<i>Lepista irina</i>	AFTOL-ID 815	DQ221109*	DQ234538*	DQ385885	DQ028591	DQ447919	-	[35]
<i>Lepista nebularis</i>	CBS362.65	AF357063	AF223217	EF421011	EF421081	DQ825415	AF357142	[35]
<i>Lepista nuda</i>	DUKE-RV84/1	AF357062*	AF042624*	EF421012	EF421082	EF421045	-	[32]
<i>Leucopaxillus subzonalis</i>	GB:0087013	KP453695	KJ417208	KJ424385	-	-	-	[33]
<i>Lyophyllum connatum</i>	DUKE-JM 90c	EF421104	AF042590	EF420995	EF421061	EF421029	-	[32]
<i>Lyophyllum decastes</i>	JM87/16	AF357059	AF042583	DQ367433	DQ367426	-	AF357136	[32]
<i>Lyophyllum decastes</i>	JM-Hon/k (T4)	AF357060	AF357078	EF421001	EF421067	F421035	-	[32]
<i>Lyophyllum favrei</i>	IE-BSG-HC96cp4	EF421102	AF223184	EF420990	EF421056	EF421024	AF357104	[32]
<i>Lyophyllum leucophaeatum</i>	IE-BSG-HAc251.97	AF357032	AF223202	DQ367434	DQ367427	DQ825419	AF357101	[32]
<i>Lyophyllum semitale</i>	IE-BSG-HC85/13	AF357049	AF042581	EF421002	EF421068	EF421036	AF357125	[32]
<i>Macrocybe sardoa</i>	29083a (MCVE)	MN017542	MN017481	-	-	-	MN017588	[38]
<i>Macrocybe titans</i>	58974 (FLAS-F)	MN017545	MN017484	-	MN026908	-	MN017589	[38]
<i>Musumecia bettlachensis</i>	TO HG2284	JF926520	JF926521	KJ681060	KJ681082	-	KJ681069	[35]
<i>Musumecia vermicularis</i>	LUG18975	-	KJ681037	KJ681061	KJ681083	-	KJ681070	[35]
<i>Neohygrophorus angelesianus</i>	AFTOL-ID 1719	DQ494678	DQ470814	-	-	-	-	[30,35]
<i>Ossicaulis lignatilis</i>	DUKE-D604(VT)	DQ825426	AF261397	DQ825410	EF421072	DQ825420	EF421094	[32]

Table S1. Cont.

Species	Voucher/ Strain	GenBank Accession no.						Reference
		ITS	nrLSU	RPB2	TEF1- α	RPB1	nrSSU	
<i>Pogonoloma macrocephalum</i>	TENN:037026	KP453700	KJ417209	-	-	-	KJ417168	[35]
<i>Pogonoloma spinulosum</i>	K(M):107286	KP453705	KJ417238	KJ424401	-	-	KU058571	[35]
<i>Porpoloma terreum</i>	CONC: F0030	KJ417306	KJ417216	-	-	-	-	[33]
<i>Porpoloma terreum</i>	REH5830	KJ417305	KJ417215	KJ424391	-	-	-	[33]
<i>Porpoloma portentosum</i>	MES531	KJ417298	KJ417210	KJ424386	-	-	-	[33]
<i>Porpoloma</i> sp.	DUKE-PR3995	EF421106	AF261395	EF421013	EF421083	EF421046	EF421095	[32]
<i>Pseudoclitocybe cyathiformis</i>	AFTOL 1998	-	EF551313	GU187815	GU187742	-	GU187659	[35]
<i>Pseudoclitopilus rhodoleucus</i>	GB:0110967	KP453696	KJ417218	KJ424393	-	-	KU058577	[35]
<i>Pseudolaccaria fellea</i>	006240 (WTU)	MN017549	MN017487	-	MN026911	-	MN017591	[38]
<i>Pseudolaccaria pachyphylla</i>	TR gmb 00672	NR_153455	NG_060151	-	-	-	-	[38]
<i>Pseudoarmillariella ectypoides</i>	AFTOL-ID 1557	DQ192175	DQ154111	DQ474127	-	DQ516076	-	[35]
<i>Pseudoomphalina kalchbrenneri</i>	T0 HG 22102011C	KR818913	KR818915	-	-	-	-	[35]
<i>Rhodocybe truncata</i>	CBS482/50	EF421110	AF223167	EF421019	KP255478	EF421052	AF357086	[32]
<i>Suillus pictus</i>	AFTOL 717	AY854069	AY684154	AY786066	AY883429	AY858965	AY662659	[35]
<i>Tephrocye boudieri</i>	IE-BSG-BSI96/84	AF357047	AF223204	DQ825411	EF421070	DQ825421	AF357122	[32]
<i>Tephrocye striaepilea</i>	FR2014091	KP192644	-	KP192524	-	-	-	NCBI
<i>Termitomyces microcarpus</i>	DUKE-PRU3900	AF357023	AF042587	EF421009	EF421077	EF421043	AF357092	[32]
<i>Tricholoma myomyces</i>	DUKE- KMS589	DQ825428	U76459	DQ367436	DQ367429	DQ842013	EF421096	[32]
<i>Tricholoma terreum</i>	C59300	EU653300	EU653304	-	-	-	JN389400	NCBI
<i>Tricholomella constricta</i>	IE-BSG-HC84/75	AF357036	AF223188	DQ825412	EF421079	DQ825422	AF357105	[32]
<i>Tricholosporum goniospermum</i>	AR122	KU559861*#&	-	KU559863	-	-	-	[20]
<i>Tricholosporum goniospermum</i>	MS41	KU559844*#&	-	-	-	-	-	[20]
<i>Tricholosporum goniospermum</i>	PeruMyc2084	MT707943*#&	-	-	-	-	-	NCBI
<i>Tricholosporum porphyrophyllum</i>	HMJAU24949	KU954553*#&	KU954556*#&	KX397356	-	-	-	NCBI
<i>Tricholosporum porphyrophyllum</i>	H6849	KU954554*#&	KU954558*#&	KX397357	-	-	-	NCBI
<i>Tricholosporum porphyrophyllum</i>	KUBOT-KRMK-2020-94	MW485792*#&	MW485793*#&	-	-	-	-	NCBI
<i>Tricholosporum guangxiense</i>	HMJAU59023	OK377045*#&	OK377053*#&	OK625399	OK625329	-	OK624824	Present study
<i>Tricholosporum guangxiense</i>	HMJAU59027	OK377046*#&	OK377055*#&	OK625402	OK625332	-	OK377042	Present study

Table S1. Cont.								
Species	Voucher/ Strain	GenBank Accession no.						
		ITS	nrLSU	RPB2	TEF1- α	RPB1	nrSSU	Reference
<i>Tricholosporum guangxiense</i>	HMJAU59028 Holotype	OK377047*#&	OK377056*#&	OK625403	OK625333	-	OK377043	Present study
<i>Tricholosporum guangxiense</i>	M2021082219 (IBK)	-	OK576387*#&	-	OK625335	-	OK624827	Present study
<i>Tricholosporum haitangshanum</i>	HMJAU59029	OK377050*#&	OK576384*#&	-	OK625334	-	-	Present study
<i>Tricholosporum haitangshanum</i>	HMJAU33972 Holotype	OK576388*#&	OK576383*#&	-	OK625338	-	OK624823	Present study
<i>Tricholosporum haitangshanum</i>	XJZ20160817	-	OK576385*#&	-	OK625336	-	OK624825	Present study
<i>Tricholosporum</i> sp.	LG218-1	MF538719*#&	-	-	-	-	-	NCBI
<i>Tricholosporum</i> sp.	LG218-9	MF538721*#&	-	-	-	-	-	NCBI
<i>Xerophorus olivascens</i>	18224 (AMB)	MN017556	MN017494	-	-	-	-	[38]
<i>Xerophorus olivascens</i>	18226 (AMB)	MN017558	MN017496	MN018856	MN026916	-	-	[38]

Note:

Sequences (GenBank Accession no.) marked with *, were used in the phylogenetic analyses of **Figure 2**.

Sequences (GenBank Accession no.) marked with #, were used in the phylogenetic analyses of **Figure S1**.

Sequences (GenBank Accession no.) marked with &, were used in the phylogenetic analyses of **Figure S2**.

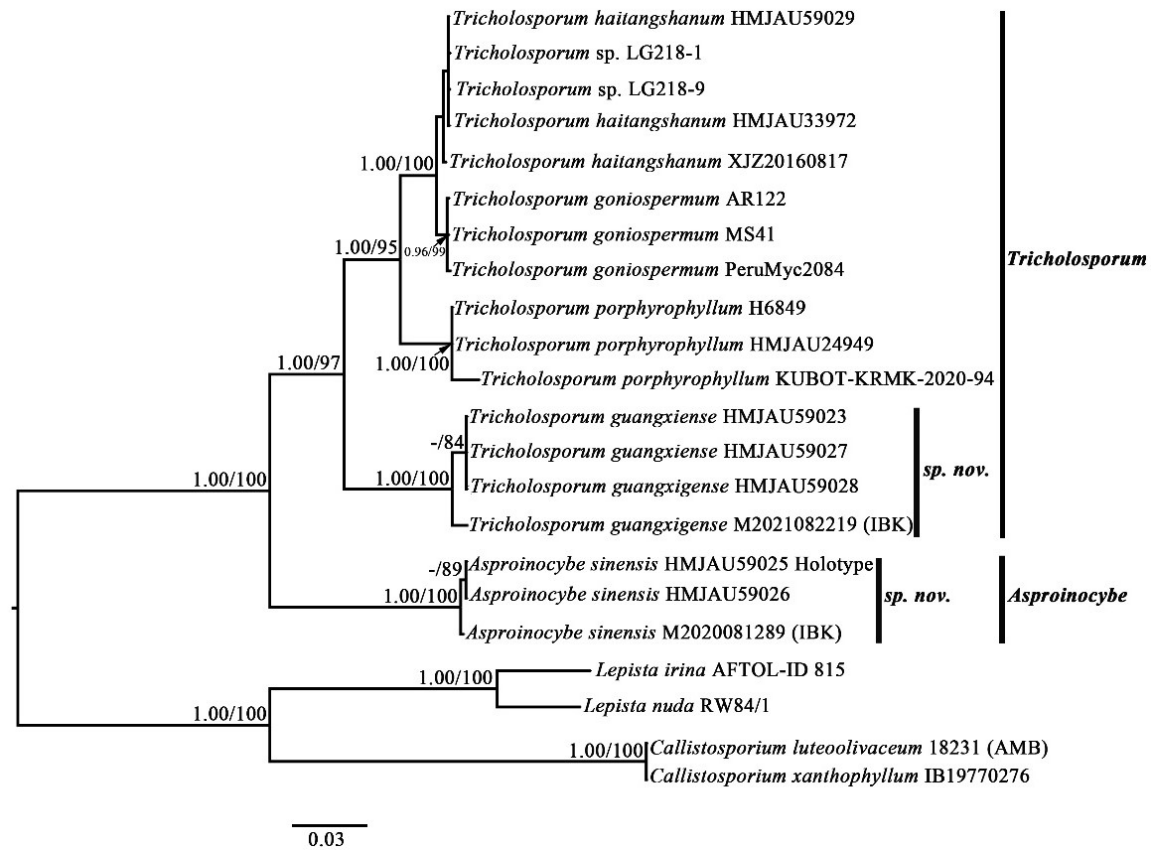


Figure S1. Phylogenetic tree inferred from partial ITS+LSU sequences showing phylogenetic relationships of *Asproinocybe* and *Tricholosporum*. Bayesian inference (BPP \geq 0.90) and maximum likelihood support values (ML \geq 70) are shown (BPP/ML).

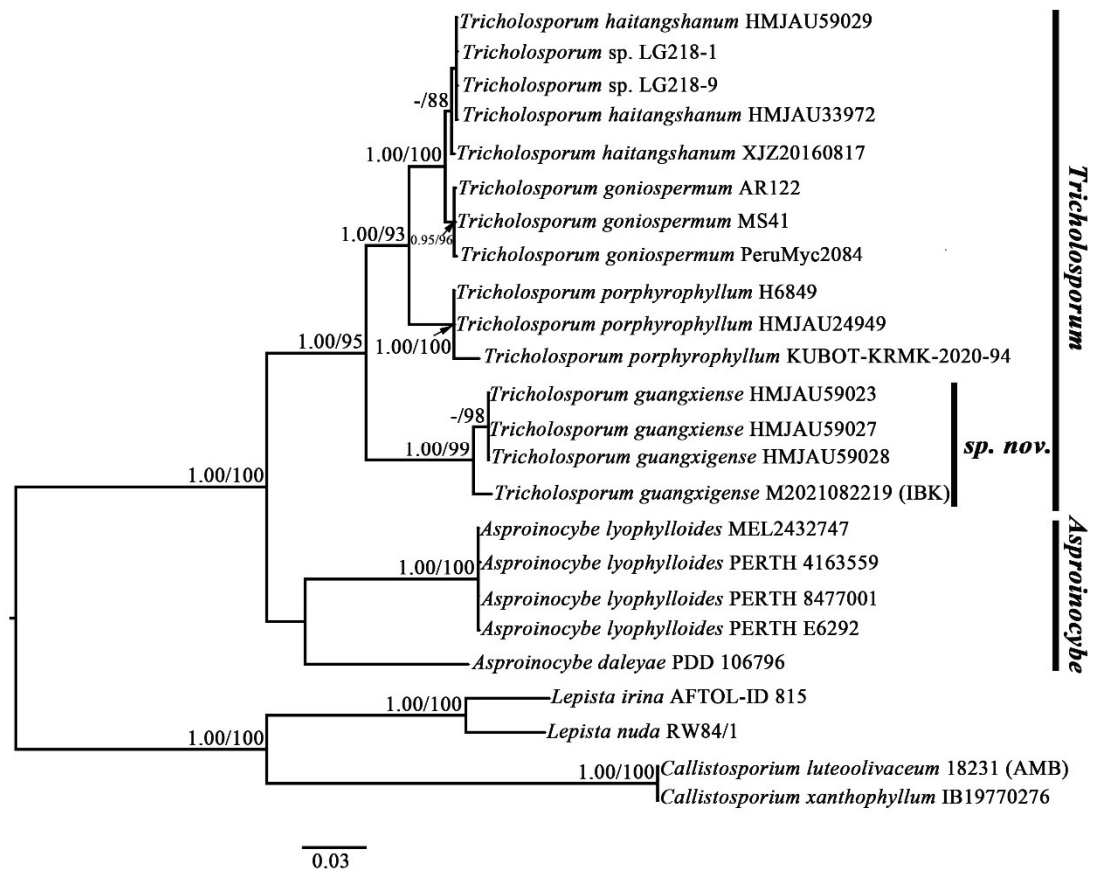


Figure S2. Phylogenetic tree inferred from partial ITS+LSU sequences showing phylogenetic relationships of *Asproinocybe* and *Tricholosporum*. Bayesian inference (BPP \geq 0.90) and maximum likelihood support values (ML \geq 70) are shown (BPP/ML)