



Kirschsteiniothelia xishuangbannaensis sp. nov. from Pará rubber (*Hevea brasiliensis*) in Yunnan, China

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Abstract

During surveys of saprobic fungi associated with Pará rubber in Yunnan Province, China, a saprobic fungus morphologically resembles *Kirschsteiniothelia* was collected from a dead branch of Pará rubber. *Kirschsteiniothelia xishuangbannaensis* is characterized by conidia having acrogenous, obclavate, rostrate, straight or curved, truncate at base, 3–8-septate, 1–2 hyaline globose/ampulliform mucilaginous sheaths around the apex. Multi-gene phylogenetic analyses of ITS, LSU and SSU showed our collection is closely affiliated to *K. thailandica* with 100% ML/1.00 BYPP statistical support. Both morphological characteristics and multi-gene phylogeny confirmed that our collection is a distinct species of *Kirschsteiniothelia*. Descriptions, micromorphological illustrations, and a phylogenetic tree showing the placement of the new taxon are provided.

Keywords – 1 new species – Hyphomycetes – *Kirschsteiniotheliaceae* – Multi-gene – Taxonomy

Introduction

Kirschsteiniotheliaceae was introduced in *Pleosporales* to accommodate a single genus *Kirschsteiniothelia* based on morphological and phylogenetic analyses (Boonmee et al. 2012). Hernandez-Restrepo et al. (2017) found that *Kirschsteiniotheliaceae* is phylogenetically distinct from other orders in *Dothideomycetes*. Thus, a new order *Kirschsteiniotheliales* was introduced to accommodate *Kirschsteiniotheliaceae*, with the type genus *Kirschsteiniothelia* (Hernandez-Restrepo et al. 2017, Hongsanan et al. 2020). The members of *Kirschsteiniotheliaceae* are commonly saprobic on dead or decaying wood (Bao et al. 2018, Rodríguez-Andrade et al. 2019, Sun et al. 2021, Verma et al. 2021).

Kirschsteiniothelia was introduced by Hawksworth (1985) and most species in this genus have been found in tropical or subtropical regions in terrestrial and freshwater habitats (Bao et al. 2018, Sun et al. 2021). The sexual morph of *Kirschsteiniothelia* has brown or black, superficial or semi-immersed, subglobose ascomata, cylindrical-clavate, bitunicate, 8-spored asci, brown to dark brown ascospores with smooth-walled and 1–2-septate (Hawksworth 1985, Boonmee et al. 2012, Hyde et al. 2013, Mehrabi et al. 2017). There are two types of asexual morphs in *Kirschsteiniothelia* as dendryphiopsis-like and sporidesmium-like (Sun et al. 2021).

Dendryphiopsis-like asexual morph was reported linking with *Kirschsteiniothelia* (*K. lignicola*, *K. emarceis*), which was confirmed by Hawksworth (1985) and Boonmee et al. (2012). Su et al. (2016) reported a sporidesmium-like asexual morph (*K. submerse*) in *Kirschsteiniothelia*. Evidence that each of these is included in the LSU-SSU-ITS matrix (Boonmee et al. 2012, Su et al. 2016).

The Pará rubber tree (*Hevea brasiliensis*) belongs to *Euphorbiaceae* and is native to the Amazon region. Rubber has great economic importance due to the latex, which is the source of natural rubber (Alvarenga & Carmo 2008). Knowledge of the fungi associated with rubber is important for pathologists, mycologists, and farmers. Senwana et al. (2021) provided a checklist of fungi and fungus-like organisms associated with rubber. The taxa in the checklist belong to 67 orders, 168 families, and 513 genera from 59 countries. Very few taxa associated with rubber have been described based on both morphology and phylogeny, while most of the previous studies on fungi associated with rubber were based only on morphology (Senwana et al. 2017, 2018, 2019, 2020, 2021, Xu et al. 2022a, b).

In this study, we introduce a new species isolated from a dead branch of *Hevea brasiliensis* in Xishuangbanna, Yunnan Province, China. The new species is confirmed based on morphological characteristics and phylogenetic analyses of combined LSU, ITS and SSU sequence data. In addition, we provide morphological characteristics, host, and location information of all known *Kirschsteiniothelia* species.

Material & methods

Sample collection and isolation

Specimens were collected from Xishuangbanna, Jinghong, Manlie, 28–30 °C, 330 m, 122°2'28"N, 100°44'23"E, Yunnan Province, China (September 2021), stored in sealable plastic bags and brought to the mycology laboratory. Morphological observations and single spore isolations were carried out following Senanayake et al. (2020). Morphological characteristics were observed using a stereomicroscope Leica S8AP0 and photographed with a Leica DM2500 camera connected to a Leica DMC4500 compound microscope. Measurements were made using Tarosoft (R) Image Frame Work software. Adobe Photoshop CS5 Extended Version 10.0 software was used to make photo-plates. Herbarium specimen of the new species was deposited at the Zhongkai University of Agriculture and Engineering (ZHKU). The living cultures were deposited at the culture collection of Zhongkai University of Agriculture and Engineering (ZHKUCC). Faces of fungi (FoF) numbers and Index Fungorum (IF) numbers were obtained as explained in Jayasiri et al. (2015) and Index Fungorum (2022).

DNA extraction, PCR amplification and sequencing

Genomic DNA was extracted directly from fresh mycelia that were scraped from a one-month-old culture using an E.Z.N.A. Forensic DNA Kit (BIO-TEK) in accordance with the manufacturer's protocol. Primer pairs ITS5 / ITS4 (White et al. 1990), LR0R / LR5 and NS1 / NS4 (Vilgalys & Hester 1990) were used to perform PCR amplifications for the large subunit ribosomal rRNA (LSU), the internal transcribed spacer (ITS) and the small subunit ribosomal rRNA (SSU), respectively. Polymerase chain reaction (PCR) amplifications were conducted using 25 µl PCR mixture containing 8.5 µl ddH₂O, 12.5 µl 2 × Master Mix (Biotek Corporation, Beijing, China), 2 µl DNA template and 1 µl each reverse and forward primer (Tibpromma et al. 2018, Xu et al. 2022b). Purification and sequencing of PCR products were carried out in Kunming Tsingke, P.R. China.

Sequence alignment and phylogenetic analyses

Sequences generated in this study were subjected to BLAST searches in GenBank to check the closely related taxa to the new taxon. Sequences were also obtained from the recently published papers (Bao et al. 2018, Hongsanan et al. 2020) (Table 1). Multiple alignments were automatically made with MAFFT v. 7 available at the web server (<http://mafft.cbrc.jp/alignment/server>), using

default settings (Kato et al. 2019). The aligned sequences were trimmed using TrimAl V 1.2 with “gappyout” option (Capella-Gutiérrez et al. 2009). Data were converted from FASTA to PHYLIP and NEXUS formats using ALTER (Alignment Transformation Environment online, <https://sing.ei.uvigo.es/ALTER>) (Glez-Peña et al. 2010). Combined multi-gene phylogenetic trees were constructed using maximum likelihood (ML) and Bayesian Inference (BYPP) analyses (Ronquist & Huelsenbeck 2003, Stamatakis 2006, Dissanayake et al. 2020). The ML analysis was done on the online platform CIPRES (<https://www.phylo.org>) using RAxML-HPC on XSEDE under the GTRGAMMA substitution model with 1,000 bootstrap replicates. Bayesian analysis was performed on the same platform CIPRES (<https://www.phylo.org>) using MrBayes. The GTR+I+G was used a substitution model of the combined ITS, LSU and SSU based on MrModeltest 2.3 (Nylander 2004). Six simultaneous Markov chains were run for 1,000,000 generations sampling every 1000th generation, halting the analyses when the average standard deviation of split frequencies was less than 0.01. Trees were visualized in FigTree v1.4.0, and the constructed tree was edited using Microsoft PowerPoint and saved as a PDF format.

Table 1 Names, strain/culture collection numbers and corresponding GenBank accession numbers of the taxa used in the phylogenetic analyses in this study. The newly generated sequences are shown in black bold while T indicates holotype or ex-type strains. Absence of GenBank accession numbers shown by “–”.

Taxa	Strains	ITS	LSU	SSU
<i>Acrospermum adeanum</i>	M133	EU940180	EU940104	EU940031
<i>Acrospermum compressum</i>	M151	EU940161	EU940084	EU940012
<i>Acrospermum gramineum</i>	M152	EU940162	EU940085	EU940013
<i>Aliquandostipite crystallinus</i>	R 76–1	–	EF175651	EF175630
<i>Aliquandostipite khaoyaiensis</i> ^T	CBS 118232	–	GU301796	–
<i>Dyfrolomyces rhizophorae</i>	JK5456A	–	GU479799	GU479766
<i>Dyfrolomyces tiomanensis</i>	NTOU3636	–	KC692156	KC692155
<i>Flavobathelium epiphyllum</i>	MPN67	–	GU327717	JN887382
<i>Halokirschsteiniothelia maritima</i>	CBS 221.60	–	AY849943	AF053726
<i>Helicomycetes roseus</i>	CBS 283.51	AY916464	AY856881	AY856928
<i>Helicomycetes roseus</i>	MFLUCC 15–0343	KY320523	KY320540	–
<i>Homortomyces combreti</i> ^T	CPC 19808	JX517281	JX517291	–
<i>Homortomyces tamaricis</i>	MFLUCC 13–0280	KU752184	KU561874	KU870905
<i>Homortomyces tamaricis</i> ^T	MFLUCC 13–0441	NR_155161	NG_059495	–
<i>Homortomyces tamaricis</i>	MFLUCC 14–0167	KU934190	KU561875	–
<i>Jahnula bipileata</i> ^T	F49–1	JN942353	EF175657	EF175635
<i>Jahnula sangamonensis</i>	A402–1B	JN942349	EF175661	EF175639
<i>Jahnula seychellensis</i>	SS 2113.2	–	EF175664	EF175643
<i>Kirschsteiniothelia aethiops</i>	CBS 109.53	–	AY016361	AY016344
<i>Kirschsteiniothelia aethiops</i>	MFLUCC 16–1104	MH182583	MH182589	MH182615
<i>Kirschsteiniothelia aethiops</i>	S–783	MH182586	MH182595	MH182617
<i>Kirschsteiniothelia aethiops</i>	MFLUCC 15–0424	KU500571	KU500578	KU500585
<i>Kirschsteiniothelia aquatica</i> ^T	MFLUCC 16–1685	MH182587	MH182594	MH182618
<i>Kirschsteiniothelia arasbaranica</i>	IRAN 2509C	KX621986	KX621987	KX621988
<i>Kirschsteiniothelia arasbaranica</i> ^T	IRAN 2508C	KX621983	KX621984	KX621985
<i>Kirschsteiniothelia atra</i> ^T	DEN	MG602687	–	–
<i>Kirschsteiniothelia cangshanensis</i> ^T	MFLUCC 16–1350	MH182584	MH182592	–
<i>Kirschsteiniothelia cangshanensis</i>	GZCC19–0515	–	MW133829	MW134609
<i>Kirschsteiniothelia ebriosa</i>	CBS H–23379	–	LT985885	–
<i>Kirschsteiniothelia emarceis</i> ^T	MFLU 10–0037	NR_138375	NG_059454	–
<i>Kirschsteiniothelia fluminicola</i> ^T	MFLUCC 16–1263	MH182582	MH182588	–
<i>Kirschsteiniothelia lignicola</i> ^T	MFLUCC 10–0036	HQ441567	HQ441568	HQ441569
<i>Kirschsteiniothelia phoenicis</i> ^T	MFLUCC 18–0216	MG859978	MG860484	MG859979
<i>Kirschsteiniothelia phoenicis</i>	MFLU 18–0153	NR_158532	NG_064508	–
<i>Kirschsteiniothelia rostrata</i>	MFLUCC15–0619	KY697280	KY697276	NG_063633

Table 1 Continued.

Taxa	Strains	ITS	LSU	SSU
<i>Kirschsteiniothelia rostrata</i>	MFLUCC 16–1124	–	MH182590	–
<i>Kirschsteiniothelia rostrata</i>	MFLU 15–1154	NR_156318	NG_059790	KY697278
<i>Kirschsteiniothelia submersa</i> ^T	MFLUCC 15–0427	KU500570	KU500577	KU500584
<i>Kirschsteiniothelia submersa</i>	S–481	–	MH182591	MH182616
<i>Kirschsteiniothelia submersa</i>	S–601	MH182585	MH182593	–
<i>Kirschsteiniothelia tectonae</i>	MFLUCC 12–0050	KU144916	KU764707	–
<i>Kirschsteiniothelia tectonae</i>	MFLUCC 13–0470	KU144924	–	–
<i>Kirschsteiniothelia thailandica</i> ^T	MFLUCC 20–0116	MT985633	MT984443	MT984280
<i>Kirschsteiniothelia thujina</i>	JF13210	KM982716	KM982718	KM982717
<i>Kirschsteiniothelia vinigena</i> ^T	CBS H–23378	–	NG_075229	–
<i>Kirschsteiniothelia xishuangbannaensis</i>^T	ZHKUCC 22–0220	OP289566	OP289564	OP303181
<i>Kirschsteiniothelia xishuangbannaensis</i>	ZHKUCC 22–0221	OP289563	OP289565	OP303182
<i>Phyllobathelium anomalum</i>	MPN 242	–	GU327722	JN887386
<i>Stemphylium vesicarium</i>	CBS 191.86	MH861935	GU238160	GU238232
<i>Stemphylium vesicarium</i>	MFLUCC 14–0920	KY659560	KY659563	KY659567
<i>Tubeufia helicomycetes</i>	CBS 271.52	AY916461	AY856887	AY856933
<i>Tubeufia javanica</i> ^T	MFLUCC 12–0545	KJ880034	KJ880036	KJ880035

Result

Phylogenetic analyses

The combined sequence alignment comprised 52 taxa, with *Stemphylium vesicarium* (CBS 191.86, MFLUCC 14–0920) as the outgroup taxa. The matrix had 2442 characters including gaps, and after trimming it had 1559 distinct alignment patterns with 28.50% of undetermined characters or gaps (ITS:554; LSU:877; SSU:1011). The RAxML analysis of the combined dataset yielded a best scoring tree (Fig. 1) with a final ML optimization likelihood value of -23356.466779. The final RAxML tree is shown in Fig. 1. Phylogenetic analyses results showed that the new strains ZHKUCC 22–0220, and ZHKUCC 22–0221 forms a sister clade with *Kirschsteiniothelia thailandica* with high statistical support (100% ML/ 1.00 BYPP, Fig. 1).

Taxonomy

Kirschsteiniothelia xishuangbannaensis R.F. Xu & Tibpromma, sp. nov. Fig. 2

Index Fungorum Number: IF 559433; Facesoffungi Number: FoF12758

Etymology – The specific epithet “xishuangbannaensis” refers to Xishuangbanna, in Yunnan Province where the holotype was collected.

Saprobic on a dead branch of *Hevea brasiliensis*. Sexual morph: Undetermined. Asexual morph: Colonies effuse on natural substrate, solitary or scattered, hairy, brown to dark brown. *Conidiophores* 35–150 × 5–15 μm (\bar{x} = 104 × 10 μm, n = 10), macronematous, septate, curved, solitary, brown to dark brown, wide and slightly swollen at base, tapering towards apex. *Conidiogenous cells* 10–50 × 5–10 μm (\bar{x} = 29 × 9 μm, n = 10), monoblastic, holoblastic, integrated, terminal, determinate, cylindrical or lageniform, smooth, brown to dark brown. *Conidia* 30–150 × 5–20 μm (\bar{x} = 93 × 16 μm, n = 20), acrogenous, solitary, obclavate, rostrate, straight or curved, truncate at base, some have guttules, 3–8-septate, yellow-brown to brown, subhyaline or pale-brown at apex, 1–2 hyaline globose/ampulliform mucilaginous sheaths around the tip (mostly 1 mucilaginous sheath).

Culture characteristics – cultures on PDA, circular, flat, velvety, smooth, entire, reddish brown from above, dark brown in the middle and the edges are reddish brown, dark-brown from below.

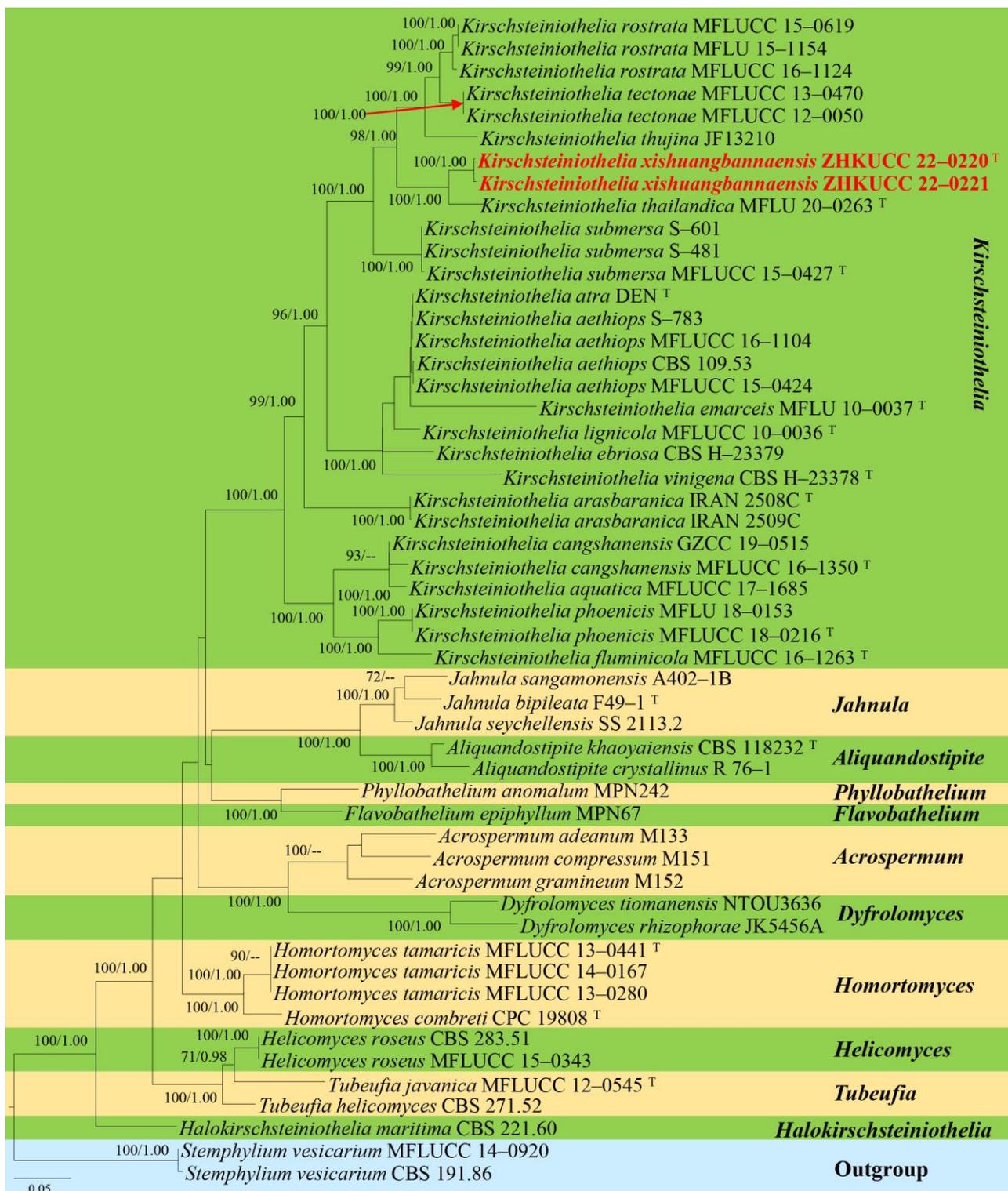


Fig. 1 – Maximum likelihood phylogenetic tree inferred from a combined dataset of ITS, LSU and SSU. The tree is rooted with *Stemphylium vesicarium* (CBS 191.86, MFLUCC 14-0920). Newly isolated strains are indicated in red bold font while T indicates holotype or ex-type strains. Bootstrap support values for maximum likelihood (ML) equal to or higher than 60% and Bayesian Probability (BYPP) equal to or higher than 0.95 are mentioned above/below the branches.

Material examined – CHINA, Yunnan Province, Xishuangbanna, Jinghong, Manlie, 28–30°C, 330 m, 122°2'28"N, 100°44'23"E, on a dead branch of *Hevea brasiliensis*, 15 September 2021, Ruifang Xu, XSBNR-44 (ZHKU 22-0126, holotype; ex-type culture ZHKUCC 22-0220 = ZHKUCC 22-0221).



Fig. 2 – *Kirschsteiniothelia xishuangbannaensis* (ZHKU 22–0126, holotype). A, B Colonies on the natural substrate. C–G Conidiophores with conidia (red arrows indicate mucilaginous sheaths). H–M Conidia (red arrows indicate mucilaginous sheaths). N Germinated conidium. O, P Colonies on PDA after one month. Scale bars: C–E, L = 50 μ m, F, G = 100 μ m, H–K, M, N = 30 μ m.

Notes – The phylogenetic analyses showed *Kirschsteiniothelia xishuangbannaensis* forms a monophyletic clade with *K. thailandica* (100% ML/1.00 BYPP, Fig. 1). The base pair comparison of *K. xishuangbannaensis* and *K. thailandica* shows 55/487 bp differences in ITS (12.11%, gaps 10 bp), 13/803 bp differences in LSU (1.62%) and 2/1,011 bp differences in SSU (0.19%). Based on the recommendation of Jeewon & Hyde (2016), a minimum of >1.5% nucleotide differences in the ITS region is indicative of a new species. Morphology comparison of the new taxon (*K. xishuangbannaensis*) and *K. thailandica* revealed that *K. xishuangbannaensis* has longer

conidiophores (35–150 vs. 55–93 μm), longer conidiogenous cells (10–50 vs. 9.5–21 μm), and longer conidia (30–150 vs. 74–110 μm). In addition, *K. thailandica* has olivaceous or brown conidia with a single hyaline mucilaginous sheath at the apex while *K. xishuangbannaensis* has yellow-brown to brown, 1–2 hyaline globose/ampulliform mucilaginous sheaths at the apex. Furthermore, the colour of culture colonies of *K. xishuangbannaensis* on PDA has dark-olivaceous whereas *K. thailandica* has reddish brown (Sun et al. 2021). Thus, *Kirschsteiniothelia xishuangbannaensis* is introduced as a new species, based on its distinct morphology, base pair differences and phylogenetic evidence.

Discussion

The survey of microfungi associated with para rubber in Yunnan Province, China revealed a new species *Kirschsteiniothelia xishuangbannaensis*. There are 29 species in *Kirschsteiniothelia* (Wijayawardene et al. 2020), including 16 sexual species (Table 2), nine asexual species (Table 3) and four species known for both morphs (Table 4), and the members of this genus are distributed in 14 countries, and most species have been found in tropical or subtropical regions (Tables 2, 3, 4). In addition, most *Kirschsteiniothelia* species have been reported from terrestrial habitat, while *K. cangshanensis*, and *K. fluminicola* found from freshwater habitat (Bao et al. 2018).

Several species were reported from various freshwater and terrestrial habitats in China viz. *K. cangshanensis*, *K. fluminicola* and *K. rostrata* (Bao et al. 2018). This shows that *Kirschsteiniothelia* species are widely distributed in Yunnan Province, China. Based on morphological investigations at different stages of rubber leaf decay (new fallen, middle stage decaying leaves, old decaying leaves), *Kirschsteiniothelia* species were discovered in Thailand (Seephueak et al. 2010). The present study expands our knowledge of *Kirschsteiniothelia* and this is the first *Kirschsteiniothelia* species associated with rubber in China that was identified based on morphology and phylogeny.

Table 2 Sexual morph morphological, host, and location information of *Kirschsteiniothelia* species. Remarks: the symbol “–” denotes no information available.

Species	Host	Country	Ascomata	Ostiole	Peridium	Pseudoparaphyses	Asci	Ascospores	References
<i>K. abietina</i>	<i>Tsuga canadensis</i>	U.S.A.	scattered, minute depressed sub-hemispherical, thin, when young and immature elevating the surface of the bark into light brown or concolorous pustules, when mature becoming sub-erumpent, black at the apex, and readily removable from the surface	minute, papilliform	100–110 \times 20 μm , oblong cylindrical	–	100–110 \times 20 μm , oblong cylindrical, 7–8 spores	23–28 \times 6–10 μm , irregularly biseriate, elliptical, rounded at the ends, slightly or not at all constricted at the septum, young hyaline, nucleosomes or granular, becoming brown	Wang et al. (2004)

Table 2 Continued.

Species	Host	Country	Ascomata	Ostiole	Peridium	Pseudoparaphyses	Asci	Ascospores	References
<i>K. acerina</i>	<i>Acer saccharum</i>	U.S.A.	185–240 × 145–200 μm, arising singly or in small groups, scattered, superficial, hemispherical, applanate at the base, black	–	20–40 μm thick, dark reddish-brown, <i>textura angularis</i>	cellular, persistent, regularly septate, branched and anastomosing, 1.5–2.0 μm thick	85–95 × 20–24 μm, subcylindrical, short-stalked, bitunicate, (4-)8-spores	22–26 × 8–11 μm, ellipsoid, rounded at the apices, 1-septate, the lower cell often somewhat smaller, slightly constricted at the septum, brown to dark brown	Hawksworth (1985)
<i>K. arasbaranica</i>	<i>Quercus petraea</i>	Iran	189–380 × 350–480 μm, slightly immersed in the substrate, subglobose to globose, scattered or loosely clustered in small groups, black, inside white	–	50–80 μm thick, comprising several layers of cells of <i>textura angularis</i> , inner layer cells hyaline, outer layer cells pale brown to dark	comprising numerous filiform 2–2.7 μm wide, hyaline, branched, embedded in a gelatinous matrix	120–180 × 30–40 μm, cylindrical to clavate, 8-spores, occasionally 4- or 6-spores, bitunicate, fissitunicate, with a short stalk up to 10 μm long, apically rounded, with an apical ocular chamber 5–8 μm wide	34–42 × 13–16 μm, arranged 2- to 4-seriate in the lower 2/3 and 1-seriate in the upper 1/3 of asci, brown to dark brown at maturity, narrowly to broadly ellipsoidal with rounded apex, thick-walled, covered with a mucilaginous sheath, 1-septate, septum deeply constricted and submedian, the upper cell distinctly larger than the lower cell, each cell containing a	Mehrabi et al. (2017)
<i>K. atkinsonii</i>	<i>Freycinetia arnottii</i>	U.S.A.	500 μm diam, subcuticular, developing under raised, black, shiny	–	–	1–1.5 μm diam, filamentous, simple or branched, septate, embedded	70–90 × 9–16 μm, 8-spores, clavate, short	distinct large guttules 14–16 × 5–6 μm, 2-seriate, clavate, light	Hyde (1997)

Table 2 Continued.

Species	Host	Country	Ascomata	Ostiole	Peridium	Pseudoparaphyses	Asci	Ascospores	References
						in mucilage	pedicellate, thick-walled, bitunicate with an ocular chamber	brown, 1-septate	
<i>K. atra</i>	Dead wood	U.S.A.	300–600 µm diam, arising singly or in small groups, scattered, colonies not delimited, superficial except at the base which can remain immersed, hemispherical to subglobose, the base slightly to strongly applanate, black	short papilliform or plane	40–60 µm thick, dark brown thick-walled, <i>textura angularis</i>	2–3.5 µm diam, persistent, abundant, regularly septate, branched and anastomosing, the centrum sometimes turning blue in iodine.	70–90 × 12–15 µm, broadly, cylindrical to cylindrical-clavate, short-stalked, fissitunicate, 4- or 8-spores	25–33 × 8.5–12 µm, ellipsoidal, rounded or somewhat constricted at the apices, 1-septate, the upper cell usually larger in size, somewhat constricted at the septum, brown, sometimes appearing almost smooth but at high magnification a delicate verruculose ornamentation is evident, not guttulate, lacking a distinct gelatinous sheath when mature	Hawksworth (1985)
<i>K. dolioloides</i>	<i>Pinus</i>	Switzerland	about 0.6 mm in diameter, singly or in small groups, superficially attached to the bare wood, conical to hemispherical, black, carbonaceous, very hard	roundish, on a short apical or lateral papilla	—	numerous, knotty, containing oil droplets	125–145 × 25–34 µm, obclavate, with a short-curved peduncle, thick walled, at the rounded apex with pore, 8-spores	84–39 × 14–15 µm, 2-celled, with a slight constriction at the septum, young olivine, old dark brown	Wang et al. (2004)

Table 2 Continued.

Species	Host	Country	Ascomata	Ostiole	Peridium	Pseudoparaphyses	Asci	Ascospores	References
<i>K. phileura</i>	<i>Tilia American</i>	New York	semi-immersed	simplicity	—	—	elliptic, uniseptate	perithecia punctiform, small	Barr (1993)
<i>K. phoenicis</i>	<i>Phoenix paludosa</i>	Thailand	135–160 × 183–235 μm, black, scattered, subglobose to globose or slightly conical in appearance, superficial with apical papilla or plane, base remaining immersed, usually flattened	—	laterally 24–34 μm wide, composed of 5–6 layers of <i>textura angularis</i> , with dark brown outer layer cells and inner layers pale brown	0.8–1.5 μm wide, branched, embedded in a gelatinous matrix	70–112 × 14–24 μm, bi- or tri-seriate in the middle of the ascus, cylindrical-clavate, straight or slightly curved, apically rounded, 8-spores, with a short pedicellate up to 5 μm long	18–27 × 5–7.5 μm, brown, ellipsoid, rounded or slightly pointed at the ends, 1-septate, septum sub-median and constricted, upper cell broader than the lower cell, each cell containing a distinct large guttulate, smooth-walled, with a mucilaginous sheath	Hyde et al. (2018)
<i>K. proteae</i>	<i>Protea cynaroides</i>	South Africa	a) up to 300 μm high, up to 390 μm wide, perithecium, intraepidermal, remain immersed with a tip of papilla reaching the surface or becoming superficial by weathering of the substrate	papillate	24–35 μm thick except for the ostiole region (43–52 μm thick), thickened and heavily pigmented towards the apex, consisting of two strata, outer stratum composed of a few layers of dark	hyaline, filamentous, septate, branched, 1–2 μm wide	54–72 × 6–8 μm, bitunicate, cylindrical to inflated cylindrical, ocular chamber	13–17 × 3–4 μm, pale brown to brown, fusoid, 1-septate, median or sub-median, smooth, at times cells become two large guttulate	Marincowitz et al. (2008)

Table 2 Continued.

Species	Host	Country	Ascomata	Ostiole	Peridium	Pseudoparaphyses	Asci	Ascospores	References
					brown, very thick-walled, compressed cells, individual cells indistinguishable, inner stratum composed of a few layers of hyaline, thin-walled, very compressed cells		0.8–1 × 0.5–2 μm		
<i>K. populi</i>	<i>Populus angustifolia</i>	Colorado	about 250–300 μm, scattered, small, globose, black, shining, not collapsing, obscurely perforate, base sunk in the wood	—	—	—	80–90 × 8 μm, cylindrical, short stipitate	12 × 6 μm, ovate, ends rounded, fuliginous, equally uniseptate, much constricted	Wang et al. (2004)
<i>K. reticulata</i>	Unknown twigs	Taiwan, China	320–450 × 180–220 μm, scattered, immersed at first, then erumpent through torn fissure of host, apex rounded, hyphae intercellular and intracellular in	—	20–40 μm thick at sides, <i>textura angularis</i> , at lateral where cells usually arranged in palisade-like rows, at base reduced, usually merged with underlying	branched, 1.5–2 μm wide	90–110 × 23–27 μm, obclavate, with a short stalk up to 15 μm long	ellipsoid, 17–23 × 7–10 μm, dark brown, 1-septate, septum constricted, median or occasionally slightly deviating from middle, with reticulate ornamentation on	Chen et al. (2006)

Table 2 Continued.

Species	Host	Country	Ascomata	Ostiole	Peridium	Pseudoparaphyses	Asci	Ascospores	References
			host, broad, 3–5 μm wide, deeply constricted where crossing host cell wall		host tissue			surface, covered with mucilaginous sheath	
<i>K. smilacis</i>	<i>Smilae</i>	Taiwan, China	450–600 \times 200–300 μm , scattered, superficial in appearance, externally covered with a thin-layer of host epidermis, apex rounded	—	35–50 μm thick, composed of opaque and melanized cells, poorly developed at base where cells usually integrate with host tissue	1–2 μm wide, frequently branched	75–100 \times 16–21 μm , obclavate, short-stalked, stalk up to 10 μm long	20–24 \times 6–8 μm , ellipsoid, pale brown, 1-septate, slightly constricted at septum, covered with mucilaginous sheath	Chen et al. (2006)
<i>K. striatispora</i>	<i>Juniperus nana</i>	Switzerland	150–250 \times 120–150 μm , arising singly, scattered, erumpent, almost entirely superficial at maturity, hemispherical, base applanate, black, lacking a distinct basal fringe	almost plane to slightly papillate	15–32 μm thick, dark reddish-brown, <i>textura angularis</i>	cellular, persistent, septate, very irregularly branched, the cells irregular in shape, 2–3 μm thick.	65–75 \times 9–11 μm , arising from the base of the ascomata cavity, vertically orientated, subcylindrical, short-stalked, bitunicate, with a distinct pimple-like internal apical beak, 4-spores	15–18 \times 5.0–6.5 μm , overlapping uniseriately arranged in the asci, ellipsoid to somewhat soleiform, 1-septate, constricted at the septum or scarcely, the cells equal in size or the lower slightly smaller, apices rounded, reddish-brown, slightly granular at first but finally with up to five longitudinal or sinuate furrows in	Hawksworth (1985)

Table 2 Continued.

Species	Host	Country	Ascomata	Ostiole	Peridium	Pseudoparaphyses	Asci	Ascospores	References
<i>K. thujina</i>	<i>Abies balsamea/ Thuja occidentalis</i>	Canada/ U.S.A.	350–500 × 300–500 μm, arising singly or in small groups, scattered, originating in the outer layers of the host but soon erumpent and superficial at maturity, hemispherical, base applanate, black	short–papilliform to conical 75 μm wide	40–60 μm thick, deep reddish–brown, <i>textura angularis</i>	evidently cellular, persistent, abundant, rather sparsely septate, branched, 2.5–3.5 μm thick	100–140 × 17–22 μm, arising from the base of the ascomata cavity, vertically orientated, subcylindrical to elongate-clavate, short-stalked, bitunicate, 8-spores	face view at high magnifications using light microscopy 36–50 × 15–17 μm, irregularly to distichously arranged in the asci, elongate–ellipsoid, slightly attenuated towards the apices, 1-septate, the cells mainly equal in size, the lower cell sometimes smaller, dark brown, apparently smooth walled, often guttulate	Hawksworth (1985)
<i>K. umbrinoidea</i>	Bark of <i>Aesculus hippocastanum</i>	Italy	superficial, subglobose	papillate	–	–	short-stalked, bitunicate	23–28 × 75–9 μm, oblong-fusiform, hyaline, two guttulate	Wang et al. (2004)
<i>K. xera</i>	bark of <i>Prunus</i>	New York	0.5 mm diam, black, carbonaceous, scattered or gregarious, covered at first, then emergent and superficial, embraced by the remains of the ruptured epidermis, smooth	inconspicuous	–	–	70–75 μm long, cylindrical, 8-spores	17–23 × 6–7 μm, uniseptate, constricted at the septum, cells somewhat unequal, guttules, with granulate contents, uniseriate or partly biseriate	Wang et al. (2004)

Table 3 Asexual morph morphological, host, and location information of *Kirschsteiniothelia* species. Remarks: the symbol “–” denotes no information available.

Species	Host	Country	Colony/Conidiomata	Conidiophore	Conidiogenous cells	Conidia	References
<i>K. cangshanensis</i>	Submerged wood	Yunnan, China	effuse, scattered, hairy, pale brown to brown	105.5–135.5 × 6–8 μm, macronematous, erect, pale brown, septate, unbranched, cylindrical, percurrent, straight or flexuous	monoblastic, integrated, terminal, determinate, pale brown, cylindrical	33–43 × 7.5–8.5 μm, solitary, dry, obclavate, septate, straight or slightly curved, pale brown to brown, with a gelatinous sheath at apex	Bao et al. (2018)
<i>K. ebriosa</i>	From sparkling wine	Spain	–	40–150 × 4 μm, macronematous, consisting of a straight or slightly sinuous, erect, dark brown, septate, thin- and smooth-walled to slightly verrucose stipe, bearing a few lateral branches and occasionally one branch at the top, branches are brown, thin- and smooth-walled, 1–5-septate, cylindrical, 30–50 × 4 μm, with rounded ends	branches, intercalary or terminal, determinate, cylindrical	8–14 × 4–5 μm, brown to dark brown, thin- and smooth-walled, 1–2(–5)-septate, sometimes solitary, mostly in branched acropetal chains of up to 5, cylindrical with rounded ends, sometimes slightly constricted at septum, produced laterally and terminally on the stipe, on the branches and on the conidia	Rodríguez-Andrade et al. (2019)
<i>K. fluminicola</i>	Submerged wood	China, Yunnan	effuse on natural substrate, hairy, dark brown	209–286 × 7–9 μm, erect, straight or flexuous, septate, smooth,	monoblastic, terminal, indeterminate,	47.5–86.5 × 8–10 μm, solitary to short-catenate,	Bao et al. (2018)

Table 3 Continued.

Species	Host	Country	Colony/Conidiomata	Conidiophore	Conidiogenous cells	Conidia	References
				cylindrical, dark brown to black, unbranched, percurrent	cylindrical, dark brown	obclavate, rostrate, truncate at base, slender and rounded at apex, aseptate when immature, multi-septate at maturity, subhyaline to dark brown, with conspicuous, spherical guttulate in almost all cells	
<i>K. rostrata</i>	Submerged wood	Thailand	effuse on natural substrate, scattered, hairy, dark brown to black	90–120 × 7.5–8.5 μm, macronematous, mononematous, solitary, wide and slightly swollen at base, tapering towards apex, straight or slightly flexuous, smooth, brown to dark brown, unbranched, septate	holoblastic, monoblastic, integrated, terminal, determinate, cylindrical or lageniform, smooth, mid to dark brown	77.5–108.5 × 17.5–20.5 μm, acrogenous, solitary, dry, olivaceous brown to brown, pale at apex, obclavate, rostrate, smooth, straight or curved, truncate at base, 6–17-septate	Bao et al. (2018)
<i>K. shimlaensis</i>	Himachal Pradesh	India	on natural substratum scattered, dark, brown to black	110–268 × 12–19 μm, macronematous, mononematous, single to loosely fasciculate, erect, straight to flexuous multi-septate, irregular or subscorpioid branched at the apex, dark brown,	18–24 × 8–10 μm, monoblastic, terminal or intercalary, cylindrical or doliiform, mostly discrete,	41–81 × 13–17.5 μm, solitary, obovoid, oblong, broad clavate or cylindrical, sometimes constricted near the middle or at	Verma et al. (2021)

Table 3 Continued.

Species	Host	Country	Colony/Conidiomata	Conidiophore	Conidiogenous cells	Conidia	References
				smooth	determinate, sometimes with enteroblastic, percurrent elongations, smooth, dark brown or brown	the supra basal cell, sub-truncate or rounded at the base, rounded at the apex, dark brown or black, smooth, guttulate, lumen aspect granulose, 2–5(–6)-septate	
<i>K. submersa</i>	decaying wood	Yunnan, China	effuse, brown or black, hairy	220–280 × 6–7 µm, macronematous, mononematous, blackish to brown, paler towards the apex, smooth, unbranched, thick-walled, multi-septate, straight to slightly flexuous, cylindrical, erect	holoblastic, integrated, terminal, determinate, percurrent, pale brown, cylindrical	37.5–51.5 × 8.5–9.5 µm, acrogenous, solitary, dry, smooth, obclavate, straight or slightly curved, truncate and darker at the base, rounded and paler at the apex, brown to pale brown, thick-walled, hyaline and thinner at the tip, 4–6-septate	Su et al. (2016)
<i>K. tectonae</i>	<i>Tectona grandis</i>	Thailand	natural substrate, superficial, hairy, dark brown, scattered, single or in groups	up to 200 µm long, 4–8 µm wide at the swollen base, superficial on host surface, macronematous, mononematous, simple, erect to slightly curved, unbranched or branched,	7.5–9.5 × 3.5–5 µm, monoblastic, integrated, terminal, cylindrical	135–150 × 16–17 µm, 9–25 or more transverse septa, cylindrical-obclavate, elongate, straight or slightly	Li et al. (2016)

Table 3 Continued.

Species	Host	Country	Colony/Conidiomata	Conidiophore	Conidiogenous cells	Conidia	References
				septate, slightly constricted at septa, pale brown to dark brown, cylindrical		curved, rounded at the apex and slightly paler, with sheath at apex, truncate at the base, dark reddish brown, thick-walled, smooth	
<i>K. thailandica</i>	<i>Ficus microcarpa</i>	Thailand	scattered, effuse, brown to dark brown, hairy on natural substrate	55–93 × 7–10 µm, macronematous, solitary, erect, cylindrical, straight or slightly flexuous, unbranched, septate, brown to dark brown	9.5–21 × 7–10 µm, monoblastic, integrated, terminal, cylindrical, brown to dark brown	74–110 × 13–20 µm, acrogenous, straight, solitary, obclavate, smooth-walled, 6–8-distoseptate, shiny, olivaceous or brown, hyaline at apex, truncate at base, tapering towards apex, with a conspicuous, gelatinous, hyaline sheath around tip	Sun et al. (2021)
<i>K. vinigena</i>	cork stopper	Spain	—	macronematous, consisting in a straight or slightly sinuous, erect, dark brown, septate, thin- and smooth-walled to coarsely verrucose (specially at the base) stipe, 100–150 × 3 µm, bearing several lateral	branches, intercalary or terminal, determinate, and cylindrical	8–80 × 4–5 µm, holoblastic, dark brown, thin- and smooth-walled to coarsely verrucose, 1–2(–7)-septate, sometimes solitary, mostly	Rodríguez-Andrade et al. (2019)

Table 3 Continued.

Species	Host	Country	Colony/Conidiomata	Conidiophore	Conidiogenous cells	Conidia	References
				branches, which also branch irregularly, branches are abundant, brown, thin- and smooth-walled to coarsely verrucose (specially at the base), non-septate to 7-septate, barrel-shaped to cylindrical, 5–80 × 4 μm, with rounded ends		disposed in branched acropetal chains of up to 4, cylindrical, with rounded ends, sometimes slightly constricted at septa, produced laterally and terminally on the stipe, on the branches, and on other conidia	
<i>K. xishuangbannaensis</i>	<i>Hevea brasiliensis</i>	Yunnan, China	effuse on natural substrate, solitary or scattered, hairy, brown to dark brown	35–150 × 5–15 μm, macronematous, septate, curved, solitary, brown to dark brown, widely and slightly swollen at base, tapering towards apex	10–50 × 5–10 μm, monoblastic, holoblastic, integrated, terminal, determinate, cylindrical or lageniform, smooth, brown to dark brown	30–150 × 5–20 μm, acrogenous, solitary, obclavate, rostrate, straight or curved, truncate at base, some have guttules, 3–8-septate, yellow–brown to brown, subhyaline or pale-brown at apex, 1–2 hyaline globose / ampulliform mucilaginous sheaths around the tip (mostly 1 mucilaginous sheath)	This study

Table 4 Sexual and asexual morph morphological, host, and location information of *Kirschsteiniothelia* species. Remarks: the symbol “–” denotes no information available.

Species	Host	Country	Sexual morph						Asexual morph				References
			Ascomata	Ostiole	Peridium	Pseudoparaphyses	Asci	Ascospores	Colony / Conidiomata	Conidiophore	Conidiogenous cells	Conidia	
<i>K. aethiops</i>	<i>Querus/ Carpinus/</i> dead wood/ submerged wood	Belgium/ Germany/ U.S.A./ China, Yunnan	300–600 μ m diameter, arising singly or in small groups, scattered, colonies not delimited, hemispherical to subglobose	20–30 μ m diameter, short papilliform	40–60 μ m thick, dark brown, thick-walled, <i>textura angularis</i>	cellular, persistent, abundant, regularly septate, branched 2.5–3.5 μ m thick	70–90 \times 12–15 μ m, arising from the base of the ascomata cavity, vertically orientated, broadly subcylindrical, short-stalked, bitunicate, 4- or 8-spores	25–33 \times 8.5–12.0 μ m, ellipsoid, rounded or somewhat constricted at the apices, 1-septate, the upper cell usually larger in size, somewhat constricted at the septum, brown, sometimes appearing almost smooth	on the substratum superficial, effuse, gregarious, hairy, velvety, dark brown	245–355 \times 8–10 μ m, mononematous, macronematous, septate, usually with one or short branches near the apex, erect, straight or flexuous, smooth, dark brown and gradually paler towards the apex, cylindrical, 5–10-septate	integrated, sometimes percurrent, terminal, becoming intercalary, pale brown or subhyaline, lageniform, new cell developing from apical or subapical part of subtending cell	54–63 \times 14–18 μ m, acrogenous, solitary, dry, cylindrical, rounded at the apex, and narrowly truncate at the base, 3–4-septate, septa thick, smooth, brown	Hawksworth (1985), Su et al. (2016)
<i>K. emarceis</i>	Dead wood	Thailand	109–280 \times 101–318 μ m, dark brown to black, clustered, solitary or scattered, superficial, subglobose to globose, slightly immersed, base flattened, with central ostiole, but unclear, with a few sparse hair-like setae	—	12–40 μ m thick, composed of 3–4 layers, comprising cells of <i>textura angularis</i> , inner layer cells pale brown, but cells of outer layer black	1–2 μ m wide, hyaline, embedded in a gelatinous matrix	88–140 \times 18–24 μ m, 8-spores, bitunicate, fissionate, cylindrical to clavate, apically rounded	25–28 \times 8–9 μ m, biserial, ellipsoidal, slightly curved, septum median to supra-median, dull green, becoming brown to dark brown at maturity, one-septate, two-celled, smooth walled, lacking a mucilaginous sheath	—	162–271 \times 7–14 μ m, erect, branched apically, brown to dark brown, smooth	—	45–56 \times 14–15 μ m, oblong to clavate, grayish, brown to dark brown, 3–4 (–5) septate, constricted at septa, smooth-walled	Boonmee et al. (2012)

Table 4 Continued.

Species	Host	Country	Sexual morph						Asexual morph				References
			Ascomata	Ostiole	Peridium	Pseudoparaphyses	Asci	Ascospores	Colony / Conidiomata	Conidiophore	Conidiogenous cells	Conidia	
<i>K. lignicola</i>	Dead wood	Thailand	152–178 × 156–214 μm, superficial, subglobose to globose, dark brown to black, membranaceous, clustered, solitary, with central papilla 24 μm wide with 24 μm high	—	18–22 mm thick, comprising 3–4 layers of cells of <i>textura angularis</i> , inner layer cells pale brown, outer layer cells dark	filiform, 1–2 mm wide, hyaline, embedded in a gelatinous matrix	107–163 × 19–28.5 μm, bitunicate, fissitunicate, cylindrical to clavate, 8-spores with along pedicel 14.5–24 μm, apically rounded	27–30 × 10–12 μm, biseriate, ellipsoidal, slightly curved, 1(–2) septate, with median septum or in lower part, some ascospores with secondary septum, dull green, brown to dark brown at maturity, thick-walled	287–406 × 11–13 μm, erect, branched apically, dark brown, smooth, conidium forming at apex	—	—	39–48 × 21–25 μm, obovoid to broadly, 1–2 transverse septa, constricted at septa, smooth-walled, dark brown, rounded at ends	Boonmee et al. 2012
<i>K. recessa</i>	<i>Pyrus</i> / dead wood	Italy/ U.S.A.	150–250 μm diameter, arising singly or more usually in small groups, scattered, superficial except at the base, hemispherical, base slightly rounded to applanate, black	short-papilliform	50–75 μm thick, dark reddish-brown, composed of thick-walled, <i>textura angularis</i> , cells mainly 4–7 μm diameter, the outer most often with the lumina becoming occluded, basal tissue similar in structure to	cellular, persistent, abundant, regularly septate, branched and anastomosing, 2–3 μm thick	90 × 10 μm, arising from the base of the ascomata cavity, vertically orientated, subcylindrical, short-stalked, bitunicate, 8-spores	15–17.5 × 5–6.5 μm, distichously arranged in the asci, elongate-ellipsoid, rounded at the apices, 1-septate, cells equal in size or the lower slightly smaller, slightly constricted at the septum, pale brown, almost smooth or with a very weak verruculose ornamentation	3.5–4.0 μm thick, erect, red-brown, septate, thick-walled	—	—	—	Hawksworth 1985

Table 4 Continued.

Species	Host	Country	Sexual morph					Asexual morph				References	
			Ascomata	Ostiole	Peridium	Pseudoparaphyses	Asci	Ascospores	Colony / Conidiomata	Conidiophore	Conidiogenous cells		Conidia
					the sides of the ascoma, well developed, mainly 8–20 µm thick, the cells tending to be vertically orientated and pallisade-like					, not guttulate, lacking a distinct gelatinous sheath			

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Accessibility of data

Provide identifiers of material and data deposited in (preferably publicly accessible) public herbaria, collections, databases and data repositories

References

- Alvarenga AA, Carmo CAF. 2008 – Seringueira. Viçosa: EPAMIG 894p.
- Bao DF, Luo ZL, Liu JK, Bhat DJ et al. 2018 – Lignicolous freshwater fungi in China III: Three new species and a new record of *Kirschsteiniothelia* from northwestern Yunnan Province. *Mycosphere* 9, 755–768. Doi 10.5943/mycosphere/9/4/4
- Barr ME. 1993 – Notes on the *Pleomassariaceae*. *Mycotaxon* 49, 129–142.
- Boonmee S, Ko Ko TW, Chukeatirote E, Hyde KD et al. 2012 – Two new *Kirschsteiniothelia* species with *Dendryphiopsis* anamorphs cluster in *Kirschsteiniotheliaceae* fam. nov. *Mycologia* 104, 698–714. Doi 10.3852/11-089

- Capella-Gutiérrez S, Silla-Martínez JM, Gabaldón T. 2009 – trimAl: a tool for automated alignment trimming in large-scale phylogenetic analyses. *Bioinformatics* 25, 1972–1973. Doi 10.1093/bioinformatics/btp348
- Chen CY, Wang CL, Huang JW. 2006 – Two new species of *Kirschsteiniothelia* from Taiwan. *Mycotaxon* 98, 153–158.
- Dissanayake AJ, Bhunjun CS, Maharachchikumbura SS, Liu JK. 2020 – Applied aspects of methods to infer phylogenetic relationships amongst fungi. *Mycosphere* 11, 2652–76. Doi 10.5943/mycosphere/11/1/18
- Glez-Peña D, Gómez-Blanco D, Reboiro-Jato M, Fdez-Riverola F et al. 2010 – ALTER: program-oriented conversion of DNA and protein alignments. *Nucleic Acids Research* 38, 14–18. Doi 10.1093/nar/gkq321
- Hawksworth DL. 1985 – *Kirschsteiniothelia*, a new genus for the *Microthelia incrustans*-group (*Dothideales*). *Botanical Journal of the Linnean Society* 91, 181–202. Doi 10.1111/j.1095-8339.1985.tb01144.x
- Hernandez-Restrepo M, Gene J, Castaneda-Ruiz RF, Mena-Portales J et al. 2017 – Phylogeny of saprobic microfungi from Southern Europe. *Studies in Mycology* 86, 53–97. Doi 10.1016/j.simyco.2017.05.002
- Hongsanan S, Hyde KD, Phookamsak R, Wanasinghe DN et al. 2020 – Refined families of *Dothideomycetes*: *Dothideomycetidae* and *Pleosporomycetidae*. *Mycosphere* 11, 1553–2107. Doi 10.5943/mycosphere/11/1/13
- Hyde KD. 1997 – Ascomycetes described on *Freycinetia*. *Sydowia* 49, 1–20.
- Hyde KD, Jones EBG, Liu JK, Ariyawansa H et al. 2013 – Families of *Dothideomycetes*. *Fungal Diversity* 63, 1–313. Doi 10.1007/s13225-013-0263-4
- Hyde KD, Chaiwan N, Norphanphoun C, Boonmee S et al. 2018 – *Mycosphere* notes 169–224. *Mycosphere* 9(2), 271–430. http://www.mycosphere.org/pdf/MYCOSPHERE_9_2_8.pdf
- Index Fungorum. 2022 – <http://www.indexfungorum.org/> (Accessed on 9/18/2022)
- Jayasiri SC, Hyde KD, Ariyawansa HA, Bhat J et al. 2015 – The Faces of Fungi database: fungal names linked with morphology, phylogeny and human impacts. *Fungal Diversity* 74, 3–18. Doi 10.1007/s13225-015-0351-8
- Jeewon R, Hyde KD. 2016 – Establishing species boundaries and new taxa among fungi: recommendations to resolve taxonomic ambiguities. *Mycosphere* 7, 1669–1677. Doi 10.5943/mycosphere/7/11/4
- Katoh K, Rozewicki J, Yamada KD. 2019 – MAFFT online service: multiple sequence alignment, interactive sequence choice and visualization. *Briefings in Bioinformatics* 20, 1160–1166. Doi 10.1093/bib/bbx108
- Li GJ, Hyde KD, Zhao RL, Hongsanan S et al. 2016 – Fungal diversity notes 253–366: taxonomic and phylogenetic contributions to fungal taxa. *Fungal diversity* 78(1), 1–237. Doi 10.1007/s13225-016-0366-9
- Marincowitz S, Crous PW, Groenewald JZ, Wingfield MJ. 2008 – Microfungi occurring on the Proteaceae in the fynbos. *CBS Biodiversity Series* 7, 1–166
- Mehrabi M, Hemmati R, Asgari B. 2017 – *Kirschsteiniothelia arasbaranica* sp. nov., and an emendation of the *Kirschsteiniotheliaceae*. *Cryptogamie, Mycologie* 38, 13–25. Doi 10.7872/crym/v38.iss1.2017.13
- Nylander JAA. 2004 – MrModeltest, version 2. Evolutionary Biology Centre, Uppsala University, Uppsala, Sweden.
- Ronquist F, Huelsenbeck JP. 2003 – MrBayes 3: Bayesian phylogenetic inference under mixed models. *Bioinformatics* 19, 1572–1574. Doi 10.1093/bioinformatics/btg180
- Rodríguez-Andrade E, Stchigel AM, Guarro J, Cano-Lira JF. 2019 – Fungal diversity of deteriorated sparkling wine and cork stoppers in Catalonia, Spain. *Microorganisms* 8(1), 12. Doi 10.3390/microorganisms8010012

- Senanayake IC, Rathnayake AR, Marasinghe DS, Calabon MS et al. 2020 – Morphological approaches in studying fungi: collection, examination, isolation, sporulation and preservation. *Mycosphere* 11, 2678–2754. Doi 10.5943/mycosphere/11/1/20
- Senwana C, Hongsanan S, Phookamsak R, Tibpromma S et al. 2019 – *Muyocopron heveae* sp. nov. and *M. dipterocarpi* appears to have host-jumped to rubber. *Mycological Progress* 18, 741–752. Doi 10.1007/s11557-019-01484-4
- Senwana C, Hongsanan S, Hyde KD, Cheewangkoon R et al. 2020 – First report of the sexual morph of *Pseudofusicoccum adansoniae* Pavlic, T.I. Burgess & M.J. Wingf. on Para rubber. *Cryptogamie, Mycologie* 41, 133–146. Doi 10.5252/cryptogamie-mycologie2020v41a7
- Senwana C, Hyde KD, Phookamsak R, Jones EBG et al. 2018 – *Coryneum heveanum* sp. nov. (*Coryneaceae, Diaporthales*) on twigs of Para rubber in Thailand. *MycoKeys* 43, 75–90. Doi 10.3897/mycokeys.43.29365
- Senwana C, Mapook A, Samarakoon MC, Karunarathna A et al. 2021 – Ascomycetes on Para rubber (*Hevea brasiliensis*). *Mycosphere* 12, 1230–1408. Doi 10.5943/mycosphere/12/1/18
- Senwana C, Phookamsak R, Doilom M, Hyde KD et al. 2017 – Novel taxa of *Diatrypaceae* from Para rubber (*Hevea brasiliensis*) in northern Thailand; introducing a novel genus *Allocryptovalsa*. *Mycosphere* 8, 1835–1855. Doi 10.5943/mycosphere/8/10/9
- Seephueak P, Petcharat V, Phongpaichit S. 2010 – Fungi associated with leaf litter of para rubber (*Hevea brasiliensis*). *Mycology* 1, 213–227. Doi 10.1080/21501203.2010.536594
- Stamatakis A. 2006 – RAxML-VI-HPC: maximum likelihood-based phylogenetic analyses with thousands of taxa and mixed models. *Bioinformatics* 22, 2688–2690. Doi 10.1093/bioinformatics/btl446
- Su HY, Hyde KD, Maharachchikumbura SSN, Ariyawansa HA et al. 2016 – The families *Distoseptisporaceae* fam. nov., *Kirschsteinioteliaceae*, *Sporormiaceae* and *Torulaceae*, with new species from freshwater in Yunnan Province, China. *Fungal Diversity* 80, 375–409. Doi 10.1007/s13225-016-0362-0
- Sun YR, Jayawardena RS, Hyde KD, Wang Y. 2021 – *Kirschsteiniotelia thailandica* sp. nov. (*Kirschsteinioteliaceae*) from Thailand. *Phytotaxa* 490, 172–182. Doi 10.11646/phytotaxa.490.2.3
- Tibpromma S, Hyde KD, McKenzie EH, Bhat DJ et al. 2018 – Fungal diversity notes 840–928: micro-fungi associated with Pandanaceae. *Fungal diversity* 93(1), 1–160. Doi 10.1007/s13225-018-0408-6
- Verma RK, Prasher IB, Sushma Gautam AK, Rajeshkumar KC, Castañeda Ruíz RF. 2021 – *Kirschsteiniotelia shimlaensis* sp. nov. from Himachal Pradesh, India. *Mycotaxon* 136(2), 401–407. Doi 10.5248/136.401
- Vilgalys R, Hester M. 1990 – Rapid genetic identification and mapping of enzymatically amplified ribosomal DNA from several *Cryptococcus* species. *Journal of Bacteriology* 172, 4238–4246. Doi 10.1128/jb.172.8.4238-4246.1990
- Wang YZ, Aptroot A, Hyde KD. 2004 – Revision of the genus *Amphisphaeria*. *Fungal Diversity Research Series* 13, 1–168.
- White TJ, Bruns T, Lee S, Taylor J. 1990 – Amplification and direct sequencing of fungal ribosomal RNA genes for phylogenetics. In: Innis MA, Gelfand DH, Sninsky JJ, White TJ, eds. *PCR protocols: a guide to methods and applications*. San Diego Academic Press, p. 315–322. Doi 10.1016/b978-0-12-372180-8.50042-1
- Wijayawardene NN, Hyde KD, Dai DQ, Sánchez-García M et al. 2020 – Outline of Fungi and fungus-like taxa. *Mycosphere* 13, 53–453. Doi 10.5943/mycosphere/13/1/2
- Xu RF, Hyde KD, Karunarathna SC, Xu JC et al. 2022b – Morphology and multi-gene phylogeny reveal a new fungal genus and species from *Hevea brasiliensis* latex in Yunnan, China. *Phytotaxa* 530, 065–076. Doi 10.11646/phytotaxa.530.1.5
- Xu RF, Thiagaraja V, Dai DQ, Karunarathna SC et al. 2022a – Additions to *Fitzroyomyces* (*Stictidaceae, Ascomycota*) from Yunnan Province, China. *Phytotaxa* 548, 253–266. Doi 10.11646/phytotaxa.548.2.8