



## Fungal Planet description sheets: 1284–1382

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### Key words

ITS nrDNA barcodes  
LSU  
new taxa  
systematics

**Abstract** Novel species of fungi described in this study include those from various countries as follows: **Antartica**, *Cladosporium australitorale* from coastal sea sand. **Australia**, *Austroboletus yourkae* on soil, *Crepidotus innuopureus* on dead wood, *Curvularia stenotaphri* from roots and leaves of *Stenotaphrum secundatum* and *Thecaphora stajscii* from capsules of *Oxalis radicata*. **Belgium**, *Paraxerochrysum coryli* (incl. *Paraxerochrysum* gen. nov.) from *Corylus avellana*. **Brazil**, *Calvatia nordestina* on soil, *Didymella tabebuicola* from leaf spots on *Tabebuia aurea*, *Fusarium subflagellisporum* from hypertrophied floral and vegetative branches of *Mangifera indica* and *Microdochium maculosum* from living leaves of *Digitaria insularis*. **Canada**, *Cuphophyllus bondii* from a grassland. **Croatia**, *Mollisia inferiseptata* from a rotten *Laurus nobilis* trunk. **Cyprus**, *Amanita exilis* on calcareous soil. **Czech Republic**, *Cytospora hippophaicola* from wood of symptomatic *Vaccinium corymbosum*. **Denmark**, *Lasiosphaeria deviata* on pieces of wood and herbaceous debris. **Dominican Republic**, *Calocybella goethei* among grass on a lawn. **France (Corsica)**, *Inocybe corsica* on wet ground. **France (French Guiana)**, *Trechispora patawaensis* on decayed branch of unknown angiosperm tree and *Trechispora subregularis* on decayed log of unknown angiosperm tree. **Germany**, *Paramicrothecium sambuci* (incl. *Paramicrothecium* gen. nov.) on dead stems of *Sambucus nigra*. **India**, *Aureobasidium microtermis* from the gut of a *Microtermes* sp. termite, *Laccaria diospyricola* on soil and *Phylloporia tamilnadensis* on branches of *Catunaregam spinosa*. **Iran**, *Pythium serotinoosporum* from soil under *Prunus dulcis*. **Italy**, *Pluteus brunneovenosus* on twigs of broadleaved trees on the ground. **Japan**, *Heterophoma rehmanniae* on leaves of *Rehmannia glutinosa* f. *hueichingensis*. **Kazakhstan**, *Murispora kazachstanica* from healthy roots of *Triticum aestivum*. **Namibia**, *Caespitomonium euphorbiae* (incl. *Caespitomonium* gen. nov.) from

## Abstract (cont.)

stems of an *Euphorbia* sp. **Netherlands**, *Alfaria junci*, *Myrmecridium junci*, *Myrmecridium juncicola*, *Myrmecridium juncigenum*, *Ophioceras junci*, *Paradinemasporium junci* (incl. *Paradinemasporium* gen. nov.), *Phialoseptomium junci*, *Sporidesmiella juncicola*, *Xenopyricularia junci* and *Zaanenomyces quadripartis* (incl. *Zaanenomyces* gen. nov.), from dead culms of *Juncus effusus*, *Cylindromonium everniae* and *Rhodoveronaea everniae* from *Evernia prunastri*, *Cyphellophora sambuci* and *Myrmecridium sambuci* from *Sambucus nigra*, *Kiflimonium junci*, *Sarocladium junci*, *Zaanenomyces moderatricis-academiae* and *Zaanenomyces versatilis* from dead culms of *Juncus inflexus*, *Microcera physciae* from *Physcia tenella*, *Myrmecridium dactylidis* from dead culms of *Dactylis glomerata*, *Neochalara spiraeae* and *Sporidesmium spiraeae* from leaves of *Spiraea japonica*, *Neofabraea salicina* from *Salix* sp., *Paradissoconium narthecii* (incl. *Paradissoconium* gen. nov.) from dead leaves of *Narthecium ossifragum*, *Polyscytalum vaccinii* from *Vaccinium myrtillus*, *Pseudosolaoacrosporiella cryptomeriae* (incl. *Pseudosolaoacrosporiella* gen. nov.) from leaves of *Cryptomeria japonica*, *Ramularia pararhabdospora* from *Plantago lanceolata*, *Sporidesmiella pini* from needles of *Pinus sylvestris* and *Xenoacrodonium juglandis* (incl. *Xenoacrodonium* gen. nov. and *Xenoacrodoniaceae* fam. nov.) from *Juglans regia*. **New Zealand**, *Cryptometrion metrosideri* from twigs of *Metrosideros* sp., *Coccomyces pycnophyllocladi* from dead leaves of *Phyllocladus alpinus*, *Hypoderma aliforme* from fallen leaves *Fuscopora solandri* and *Hypoderma subiculatum* from dead leaves *Phormium tenax*. **Norway**, *Neodevriesia kalakoutskii* from permafrost and *Variabilispora viridis* from driftwood of *Picea abies*. **Portugal**, *Entomortierella hereditatis* from a biofilm covering a deteriorated limestone wall. **Russia**, *Colpoma junipericola* from needles of *Juniperus sabina*, *Entoloma cinnamomeum* on soil in grasslands, *Entoloma verae* on soil in grasslands, *Hyphodermella pallidostraminea* on a dry dead branch of *Actinidia* sp., *Lepiota sayanensis* on litter in a mixed forest, *Papiliotrema horticola* from *Malus communis*, *Paramacroventuria ribis* (incl. *Paramacroventuria* gen. nov.) from leaves of *Ribes aureum* and *Paramyrothecium lathyri* from leaves of *Lathyrus tuberosus*. **South Africa**, *Harzia combreti* from leaf litter of *Combretum collinum* ssp. *sulvense*, *Penicillium xyleborini* from *Xyleborinus saxesenii*, *Phaeoisaria dalbergiae* from bark of *Dalbergia armata*, *Protocreopsis euphorbiae* from leaf litter of *Euphorbia ingens* and *Roi-giella syzygii* from twigs of *Syzygium chordatum*. **Spain**, *Genea zamorana* on sandy soil, *Gymnopus nigrescens* on *Scleropodium touretii*, *Hesperomyces parexochomi* on *Parexochomus quadriplagiatus*, *Paraphoma variabilis* from dung, *Phaeococcomyces kinklidomatophilus* from a blackened metal railing of an industrial warehouse and *Tuber suaveolens* in soil under *Quercus faginea*. **Svalbard and Jan Mayen**, *Inocybe nivea* associated with *Salix polaris*. **Thailand**, *Biscogniauxia whalleyi* on corticated wood. **UK**, *Parasitella quercicola* from *Quercus robur*. **USA**, *Aspergillus arizonicus* from indoor air in a hospital, *Caeliomyces tampanus* (incl. *Caeliomyces* gen. nov.) from office dust, *Cippumomyces mortalis* (incl. *Cippumomyces* gen. nov.) from a tombstone, *Cylindrium desperesense* from air in a store, *Tetracoccosporium pseudoaerium* from air sample in house, *Toxicocladosporium glendoranum* from air in a brick room, *Toxicocladosporium losalamitosense* from air in a classroom, *Valsonectria portsmouthensis* from air in men's locker room and *Varicosporellopsis americana* from sludge in a water reservoir. **Vietnam**, *Entoloma kovalenkoi* on rotten wood, *Fusarium chuoi* inside seed of *Musa itinerans*, *Micropsalliota albofelina* on soil in tropical evergreen mixed forests and *Phytophthora docyniae* from soil and roots of *Docynia indica*. Morphological and culture characteristics are supported by DNA barcodes.

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*Hyphodermella pallidostraminea*



Fungal Planet 1359 – 24 December 2021

***Hyphodermella pallidostraminea* Bukharova & Volobuev, sp. nov.**

**Etymology.** Name refers to the pale straw-yellow colour of hymenial surface.

**Classification** — *Phanerochaetaceae*, *Polyporales*, *Agaricomycetes*.

**Basidiomata** annual, resupinate, adnate, effuse, ceraceous to crustaceous when dry, at first orbicular, then confluent, up to 15 mm diam. Margin white, determinate, narrow, fibrillose, with age thinning out. **Hymenial surface** yellowish, tiger yellow (090 90 50), fresh yellow (095 90 40) or pea green (095 80 40) (all colour codes are given following the RAL Design colour chart), smooth to slightly tuberculate, cracked. Subiculum thin, indistinct, white. **Hyphal system** monomitic; generative hyphae simple-septate. Subicular hyphae 3.5–5 µm diam, hyaline, thick-walled or with slightly thickened walls, encrusted with numerous prismatic crystals, loosely interwoven, more or less parallel arranged, occasionally branched. Subhymenial hyphae 2.5–3.5 µm diam, hyaline, thin-walled, smooth, compactly interwoven, frequently ramified. **Cystidia** absent, but cylindrical to fusoid, cystidioid hyphal ends occasionally present, thin-walled, not encrusted, projecting above the hymenium up to 35 µm. **Basidia** 17–25 × 5.2–6 µm, clavate, with a simple septum at the base, 4-sterigmate. **Basidiospores** (5.0–)5.4–6.6(–6.7) × (2.9–)3.0–3.5(–3.6) µm, n = 30, L = 5.96, W = 3.23, Q = 1.82–1.87, smooth, hyaline, thin-walled, ellipsoid, inamyloid, non-dextrinoid, with tiny oil drops.

**Habitat & Distribution** — On dry, dead *Actinidia* sp. branch, in a herbaceous mixed coniferous-broadleaf forest. Hitherto only known from the type locality in the south of the Russian Far East.

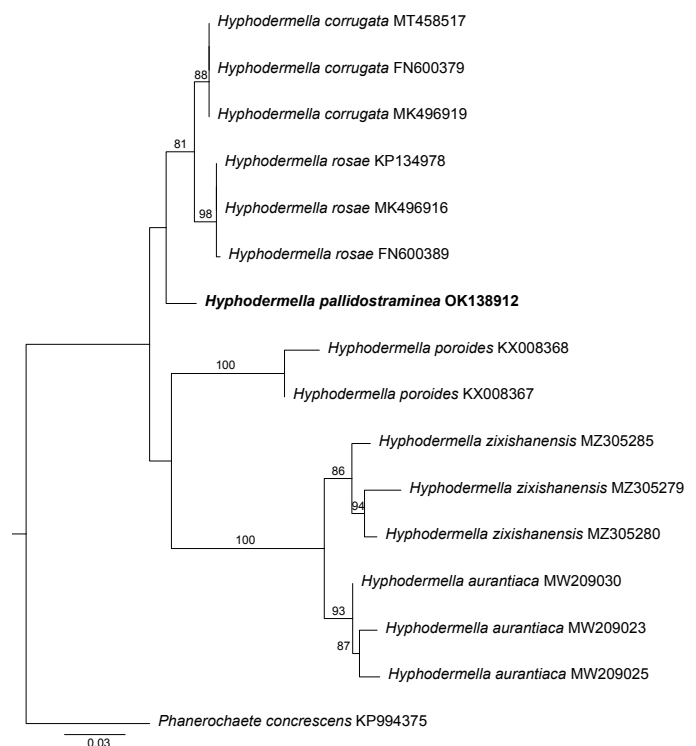
**Typus.** RUSSIA, Jewish Autonomous Oblast, Obluchensky District, Bastak State Nature Reserve, 15 km north-northeastward from the Kirga settlement, at the foot of Mount Skalistaya, N49°00'47.3" E132°53'53.6", on dry dead branch of *Actinidia* sp. (*Actinidiaceae*) in mixed coniferous-broadleaf forest, 24 Aug. 2009, N. Bukharova (*Vasilyeva*) (holotype LE 286968, isotype VLA M-22719, ITS and LSU sequences GenBank OK138912 and OK138911, MycoBank MB 841840).

**Notes** — *Hyphodermella pallidostraminea* is a minute-sized corticioid fungus characterised macroscopically by pale-yellowish smooth to tuberculate hymenophore and white pruinose margin of basidiocarps. The main distinguishing microscopic feature among other representatives of the genus *Hyphodermella* is the spore size (up to 3.5 µm in width) that is overlap-

**Colour illustrations.** Coniferous-broadleaf mixed forest in the Bastak State Nature Reserve (Jewish Autonomous Oblast, Russia). Basidiocarp; basidia and spores (all from holotype). Scale bars = 2 mm (basidiocarp), 10 µm (cystidioid hyphal ends), 5 µm (spores and basidia).

ping only with the recently described *H. zixishanensis*, but the latter species differs by the reddish to brown hymenial surface and the absence of cystidioid hyphal ends projecting above the hymenium (Wang et al. 2021). The previous attempt to find a taxonomic interpretation for the studied specimen of *H. pallidostraminea* without a molecular analysis led to the fact that this material was erroneously identified as *Sistotremastrum niveocreum* (Bukharova & Zmitrovich 2014), also being a deciduous-dwelling fungus having a monomitic hyphal system, similar spore sizes and shape, but differing by clamps on hyphae.

Based on a megablast search of NCBI's GenBank nucleotide database, the closest hits using the **ITS** sequence had highest similarity to *H. rosae* (strain FP-150552, GenBank KP134978; Identities = 625/657 (95 %), 13 gaps (1 %)), *H. corrugata* (voucher K(M) 237661, GenBank MK496919; Identities = 633/666 (95 %), 13 gaps (1 %)) and *H. rosae* (voucher CFMR DLL2011-177, GenBank KJ140672; Identities = 613/646 (95 %), 13 gaps (2 %)). Closest hits using the **LSU** sequence are *Geliporus exilisporus* (voucher TNM GC 1702-15, GenBank LC379153; Identities = 847/859 (99 %), no gaps), *H. rosae* (strain GC 1604-113, GenBank MZ637147; Identities = 849/862 (98 %), no gaps) and *H. corrugata* (voucher MA-Fungi 5527, GenBank JN939597; Identities = 849/862 (98 %), no gaps).



Phylogenetic tree derived from Maximum Likelihood analysis based on nrITS1-5.8S-ITS2 data. The analysis was performed in the IQ-TREE Web Server (Trifinopoulos et al. 2016) with 1000 ultrafast bootstrap replicates. Maximum Likelihood bootstrap support values shown above branches (BS > 80 %). The new species described in this study is in **bold** face. The scale bar represents the expected number of nucleotide changes per site.

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