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CHRYSOMYCENA PERPLEXA GEN. ET SP. NOV. (AGARICALES, POROTHELEACEAE),
A NEW ENTITY FROM THE LAZIO REGION**Abstract**

Based on morphological and molecular phylogenetic analyses, the new genus *Chrysomyca* is described for the species *C. perplexa* which shows a peculiar combination of features within the *Porothelaceae* family: a mycenoid *habitus*, a pileipellis structured as an ixocutis with some pileocystidioid elements, slightly amyloid spores and basidia, long lageniform to flagellate (flagelliform) cheilocystidia, abundant clamp-connections and sarcodimitic tissues.

Key words: *Basidiomycota*, *Agaricomycetes*, *Marasmiineae*, hydropoid clade, *Gerronema*, *Phloeomana*, *Trogia*, nrITS, nrLSU, Phylogeny, Taxonomy.

Introduction

During an investigation on the macromycetes of the Castelfusano coastline (Rome) some basidiomata of an unknown species with mycenoid *habitus* were found on a degraded stump of *Pinus pinea* L. characterized by yellowish colorations on the gills and on the stipe. The subsequent microscopic examination was able to highlight characters that did not allow a secure generic attribution on a traditional basis. The purpose of this article is therefore to provide a detailed macro- and microscopic description of the species, accompanied by colour images, and to identify, on a molecular basis, the phylogenetic position.

Materials & Methods

The annotations concerning the description of the collected samples (fresh basidiomata), of the habitat and of the plant associated with them were taken at the place of collection.

The microscopic characters were detected from fresh and dried material. In the case of dried material, rehydration was carried out with water (H₂O), Ammonia (NH₃) and Potassium hydroxide (KOH) at 5%. Glycerine buffer (L4) was also used for the observation of the preparations.

The observation of the structures and the measurements of the anatomical characters was carried out through the use of ammoniacal Red Congo (RCA) and 1% Floxin. For the observation of the parietal and intracellular pigment Blue Cotton in lactic acid (BCAL) and Toluidine Blue (BDT) was used, while Melzer's reagent (M) to test the iodine reaction of spores and other tissues.

Measurements of hyphae and spores were performed with 1000 × immersion objective lenses equipped with a calibrated micrometer. The instruments used were a Zeiss Universal R trinocular microscope with Leitz Wetzlar and PL Fluotar Lens and Zeiss Axiolab A1, with objective lenses 5 ×, 10 ×, 40 ×, 50 ×, 60 × and 100× oil immersion.

The measurement of the spores was carried out considering, from time to time, all the spores present in the visual field of the microscope, in order to satisfy the principle of randomness. In particular, the spore size was detected, excluding the apiculus, by measuring 32 basidiospores taken from several basidiomata. The values were given as minimum-average-maximum, Q = length/width with minimum and maximum values and Q_m = average quotient. The abbreviations L and l refer, respectively, to the total number of gills present in the basidioma and to the number of lamellulae between two gills. The photomicrographs were taken with Canon Powershot G10 and G11 digital cameras, while the photographs of the fresh specimens were taken *in situ* with Nikon D300 and D80 digital cameras.

The basidiomes of the examined collection have been deposited in MCVE.

Molecular analysis

DNA was extracted following the CTAB protocol according to DOYLE & DOYLE (1987). The nrITS region was amplified using the ITS1F primers (GARDES & BRUNS, 1993) and ITS4 (WHITE *ET AL.*, 1990), while the nrLSU region was amplified with the primers LR0R and LR05 (VILGALYS & HESTER, 1990). The sequences obtained were edited with Geneious v. 11.1.4 (KEARSE *ET AL.*, 2012) and then filed in GenBank (<http://www.ncbi.nlm.nih.gov/genbank>). GenBank access numbers are present in Fig. 1. The dataset used in the analyses was prepared mainly on the basis of the results present in YANG *ET AL.*, 2012, DUTTA *ET AL.*, 2017, LATHA *ET AL.*, 2018, ANTONÍN *ET AL.*, 2019, LIU *ET AL.*, 2019, and the results obtained in BLASTN. The sequences of *Gamundia leucophylla* (EU669424) and *Myxomphalina maura* (AF261378) were used as outgroups. The alignment was carried out with MAFFT v7.388 (KATOH & TOH, 2008) and the Maximum Likelihood analysis was conducted with the software RAxML 8.2.7 (STAMATAKIS, 2006) using the GTR + G model and 1000 replicas of bootstrap for statistical validation of tree nodes. Only tree support values (bootstrap-MLB) greater than or equal to 50% are shown in the tree.

Taxonomy

Chrysomyцена Vizzini, Picillo, Perrone & Dovana, nov. gen.

Mycobank MB 832697

Etiology from the ancient Greek *chrysós* (gold) + *Mycena*, referring to the golden yellow tinges on gills and stipe of this mycenoid species.

DIAGNOSIS: Habit mycenoid to omphalinoid, with yellow tinges on lamellae and stipe, lamellae adnate to subdecurrent; spore deposit whitish; spores ellipsoid, thin-walled, smooth, acyanophilous, with iodine reactions; basidia clavate, amyloid at apex; subhymenium filamentous; hymenophoral trama regular; pleurocystidia absent; cheilocystidia abundant, in clusters, lageniform to flagelliform; pileipellis as an ixocutis; caulocystidia present, similar to cheilocystidia; clamp-connections present, sometimes of the medallion-type. Context sarcodimitic. Habitat on decayed wood.

TYPE SPECIES: *Chrysomyцена perplexa*.

Chrysomyцена perplexa Picillo, Vizzini & Perrone, nov. sp. (Photo 1-31)

Mycobank MB 832698

Etiology the specific epithet is from the Latin adjective “perplexus” (unintelligible, intricate, inscrutable, ambiguous, obscure) and refers to the puzzling features of the species.

DIAGNOSIS: Pileus 7-13 mm broad, hemispheric to convex, often weakly depressed at centre, slightly undulate at margin, surface smooth, lubricous when wet, not hygrophanous, translucent-striate (pellucid) up to the centre, ochraceous-brownish at the centre, ochraceous at margin, at first covered by a dense whitish bloom which soon tends to disappear in mature specimens or due to rain. Lamellae adnate, sometimes with short decurrent tooth, spaced, up to 2 mm wide, intercalated with lamellulae, $L = 15-20$, $l = 1-2(-3)$, first whitish, then yellowish with an entire to minutely crenulate, concolorous edge. Stipe 12-25 × 2-3 mm, central, cylindrical, slightly enlarged at the base, translucent, fistulous, bright golden-yellow, surface completely decorated with a dense whitish bloom, which gives it a fluffy appearance, with evident whitish mycelial residues (rhizomorphs) at the base. Context thin, fibrous, concolorous to the external surfaces, smell and taste indistinct. Spore deposit whitish. Spores 7-8.5-9.6 × 5.2-6.1-7 μm, $Q = 1.2-1.6$, $Q_m = 1.4$, broadly ellipsoid to ellipsoid, with large central guttula or even pluriguttulate, with rather pronounced apiculus, smooth, weakly amyloid, not dextrinoid, acyanophilic. Basidia 30.4-47.8 × 6.1-9.6 μm, long clavate, pluriguttulate, tetrasporic (also bisporic) with sterigmata up to 6.1 μm long, amyloid at the apex. Subhymenium filamentous, consisting of up to 3 μm wide hyphae. Cheilocystidia 23.5-90 × 2.6-7.4 μm, with a moderately thickened wall (0.4 μm), mostly grouped in clusters (tufts), of the lageniform type but with a characteristic very thin neck of varying length (in some cases up to 70 μm long), similar to a flagellum, but also lageniform without flagellum, clavate, knotty stick shaped, bifurcate at

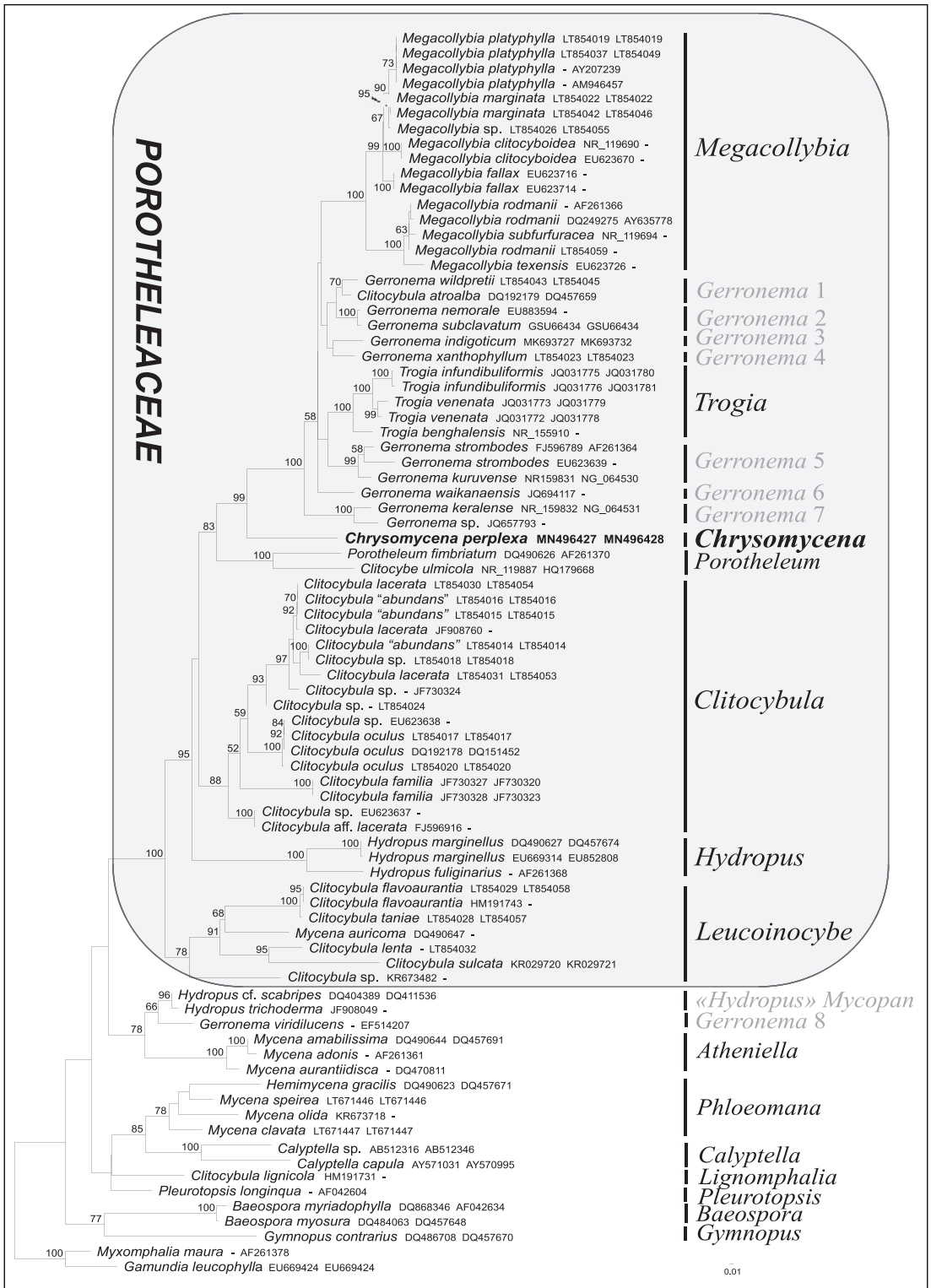


Figure 1 (tree). RAxML analysis of the combined nrITS / nrLSU (28S) dataset of *Porothelaeaceae*. Only MLB values $\geq 50\%$ are shown. Overall, an alignment of 82 sequences of the length of 1954 positions was obtained.

the apex, diverticulate. Pleurocystidia absent. Hymenophoral trama regular, sarcodimitic. Stipe structure consisting of an outer layer of thin, intertwined, knotted, diverticulate hyphae, $\times 0.9\text{-}2.6(3.2) \mu\text{m}$ and an inner layer of cylindrical hyphae, swollen in the median part, with parallel pattern, $\times 7.8\text{-}13.6(18) \mu\text{m}$, with a fairly thick wall ($0.4\text{-}0.8 \mu\text{m}$). Caulocystidia present, of shape and size comparable to that of cheilocystidia, $37.4\text{-}69.6 \times 4.3\text{-}7.8 \mu\text{m}$. Pileipellis sarcodimitic: suprapellis of the ixocutis type, gelified, with cylindrical, vesicular, utriform hyphae, with rounded apex, pointed, with a short beak or capitulum (in some cases pileocystidioid), restricted to the septa, with a fairly thick wall ($0.8 \mu\text{m}$) at the base of which no clamp-connections were observed, with terminal elements of $36.5\text{-}143 \times 14.8\text{-}34.8 \mu\text{m}$, and thinner, intertwined, knotted, diverticulate, bifurcate hyphae, with terminal elements of $61\text{-}78.3 \times 2.6\text{-}3.5 \mu\text{m}$; a brownish mixed parietal and intracellular pigment is present. Subpellis similar to the suprapellis. Oleiferous hyphae uncommon but present in all tissues of the basidioma. Clamp-connections abundant, also of the medallion type, present at the base of the cheilocystidia and caulocystidia, as well as in the thin hyphae of the pileipellis, of the hymenophoral and stipe trama; infrequent at the base of basidia. Not observed elsewhere. Habitat gregarious on a decaying *Pinus pinea* wood in the Mediterranean area, in late autumn. So far known from only one location in Lazio.

HOLOTYPE: Italy, Lazio, Pineta di Castel Fusano urban park, Castel Fusano (RM), on a *Pinus pinea* stump, 20-12-2014, leg. B. Picillo & L. Perrone (MCVE 30184).

Pileus 7-13 mm, from hemispherical to convex with slight depression to the disc, smooth, slippery in wet weather, not hygrophanous, entire margin, slightly wavy, striated for transparency up to the centre, ochre-coloured at the edge, ochre-brown to the disc; surface completely covered by a thick whitish bloom which tends to dissolve in mature specimens or by washing away.

Gills from adnate to briefly decurrent, spaced, L = 15-20, l = 1-2 (-3) wide up to 2 mm, intercalated by lamellule, at first whitish, then yellowish with the concolorous thread, from intact to slightly crenulated.

Stipe 12-25 \times 2-3 mm, central, cylindrical, slightly enlarged at the base, translucent, fistulous, golden-yellow shining, surface completely decorated with a dense whitish bloom, which gives it a fluffy appearance, there are evident whitish mycelial residues (mycelial cords/rhizomorphs) at the base.

Context fibrous, concolorous to external surfaces. No distinctive smell and taste.

Spore whitish.

Spores (Photo 2) $7\text{-}8.5\text{-}9.6 \times 5.2\text{-}6.1\text{-}7 \mu\text{m}$, Q = 1.2-1.6, Q_m = 1.4, from broadly ellipsoidal to ellipsoidal, with large central guttula or also pluriguttulate, smooth, with rather pronounced apiculus, not dextrinoid, weakly amyloid (Photo 3), acyanophilic (Photo 4).

Basidia (Photo 5) $30.4\text{-}47.8 \times 6.1\text{-}9.6 \mu\text{m}$, long clavate, pluriguttulate, tetrasporic (also bisporic) with long sterigmata up to $6.1 \mu\text{m}$, amyloid at the apex (Photo 6).

Cheilocystidia $23.5\text{-}90 \times 2.6\text{-}7.4 \mu\text{m}$, with a fairly thick wall ($0.4 \mu\text{m}$), mostly gathered in clusters, of the lageniform type but with a characteristic very thin and varied neck length (in some cases up to $70 \mu\text{m}$ long), similar to a flagellum, but also lageniforms without flagellum, clavates, claviform-nodose, bifurcated at the top, diverticulate. (Photos 7-8-9-10-11).

Subimenium filamentous, consisting of hyphae up to $3 \mu\text{m}$ wide (Photo 12).

Pleurocystidia absent.

Hymenophoral trama sarcodimitic, consisting of very variable hyphae for size and shape, with a fairly thick wall ($0.4\text{-}0.6 \mu\text{m}$). In fact, there are cylindrical, vesicular, even catenulated, utriform hyphae, with a rounded, pointed apex, restricted to the septa, and thinner, intertwined, knotted, diverticulate, bifurcated hyphae, with terminals $78\text{-}156.6 \times 3.5\text{-}63 \mu\text{m}$. A mixed brownish parietal and intracellular pigment is present. (Photos 13-14-15-16).



Photo 1. *Chrysomycena perplexa*

Photo by Bernardo Picillo

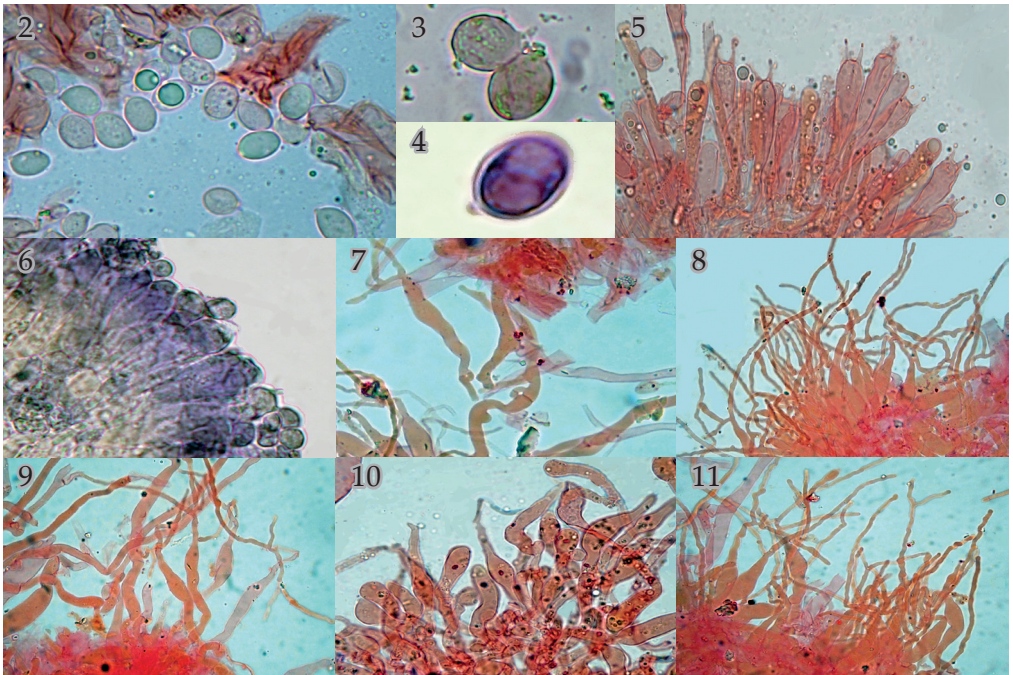


Photo 2. Spores (RCA); 3. Spores (M); 4. Spore (BCAL); 5. Basidia; 6. Hymenium (M); 7. Cheilocystidia with clamp-connections (RCA); 8-9-10-11. Cheilocystidia (RCA).

Photos by Luigi Perrone and Bernardo Picillo

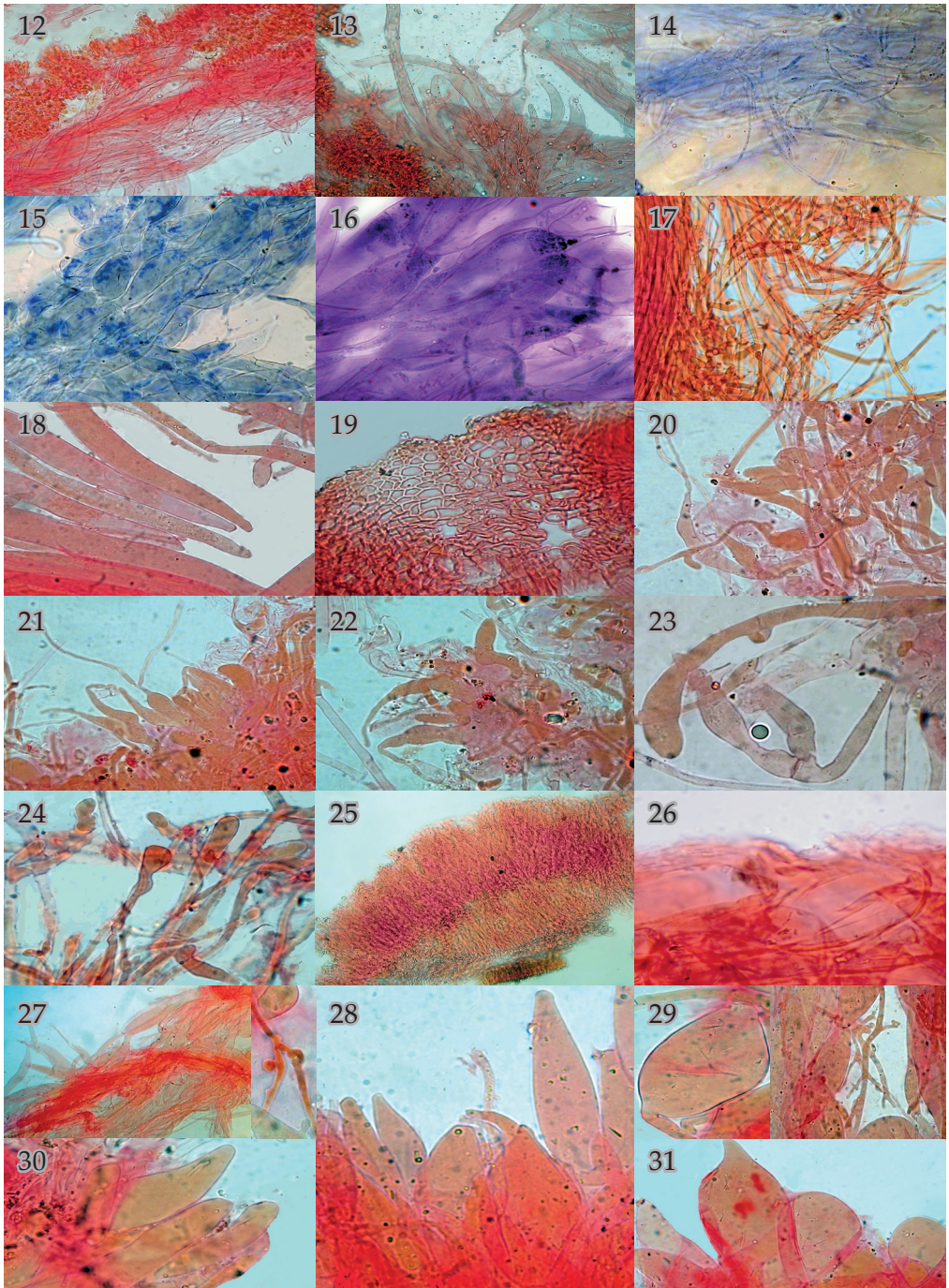


Photo 12. Hymenium and subhymenium (RCA); 13. Lamellar texture (RCA); 14-15-16. Pigmented lamellar texture (BDT and BCAL); 17. Caulopellis (RCA); 18. Stipe texture (RCA); 19. Cross section of the stipe (RCA); 20-21-22-23-24. Caulocystidia (RCA); 25. Pilepellis, complete section, without pressure on the slide (RCA); 26. Suprapellis without pressure o.t.s. (RCA); 27. Suprapellis hyphae with clamp-connections after pressure o.t.s. (RCA); 28-29-30-31. Suprapellis hypha of various morphology after pressure o.t.s. (RCA).
 Photos by Luigi Ferrone and Bernardo Picillo

Stipe structure consisting of an outer layer of thin, intertwined, knotted, diverticulated hyphae, $\times 0.9\text{-}2.6$ (3.2) μm and an inner layer of cylindrical hyphae, swollen in the median part, with parallel pattern, $\times 7.8\text{-}13.6$ (18) μm , with a fairly thick wall (0.4-0.8 μm) (Photo 17-18-19).

Caulocystidia present, of shape and size comparable to that of cheilocystidia, 37.4-69.6 \times 4.3-7.8 μm (Photos 20-21-22-23-24).

Pileipellis sarcodimitic: suprapellis ixocutis-type, gelled, with cylindrical, vesicular, utriform hyphae, with rounded apex, pointed, with short beak or capitulus (in some cases pileocystidioid), restricted to septa, with fairly thick wall (0.8 μm) at the base of which no buckle joints were observed, with terminals of 36.5-143 \times 14.8-34.8 μm , and thinner, intertwined, knotted, diverticulated, bifurcated hyphae, with terminals of 61-78, 3 \times 2.6-3.5. There is a mixed, parietal and intracellular brownish pigment (Photos 25-26-27-28-29-30-31); subpellis similar to the suprapellis.

Oleiferous hyphae non-common, but present in all tissues of the basidioma.

Clamp-connections abundant, also of the medallion-type, present at the base of the cheilocystidia and caulocystidia, as well as in the thin hyphae of the pileipellis, of the hymenophoral trama and the stipe structure. Infrequent at the base of basidia. Not observed elsewhere. (Photos 7-17-23-27).

Habitat and collections made: Italy, Lazio, Pineta di Castelfusano, urban park, gregarious on a decaying *Pinus pinea* wood, in the Mediterranean area, 20.12.2014; *legit* B. Picillo & L. Perrone (MCVE 30184). Known in that one location.

Discussion

The new genus belongs to the so-called hydropoid clade (/hydropoid). This clade was first highlighted by MONCALVO *ET AL.* (2002), in which analysis based on nrLSU sequences consisted of some species of *Hydropus* Kühner ex Singer, *Gerronema* Singer, *Megacollybia* Kotl. & Pouzar, *Clitocybula* (Singer) Singer ex Métrod and from the poroid/cifeloid genus *Porotheleum* Fr. The same configuration of the clade was present in BODENSTEINER *ET AL.* (2004). In the multigenic analysis of MATHENY (2006) the hydropoid clade (part of the larger marasmioid clade) also included *Henningsomyces candidus* (Pers.) Kuntze, some *Mycena* (Pers.) Roussel (*M. auricoma* Har. Takah., *M. amabilissima* (Peck) Sacc., *M. aurantiidisca* (Murrill) Murrill) and *Hydnopolyporus fimbriatus* (Cooke) DA Reid. *Calyptella capula* (Holmsk.) Quél. and two species of *Trogia* Fr. (*T. infundibuliformis* Berk. & Broome and *T. venenata* Zhu L. Yang) also belonged to this clade in the study of YANG *ET AL.* (2012). REDHEAD (2012, 2013, 2016a, b) included in the clade the species of the mycenoid genera *Atheniella* Redhead, Moncalvo, Vilgalys, Desjardin, B.A. Perry and *Phloeomana* Redhead. Finally ANTONÍN *ET AL.* (2019) put in the hydropoid clade the new genera *Leucoinocybe* Singer ex Antonín, Borovička, Holec & Kolařík (segregated from *Clitocybula*) and *Lignomphalia* Antonín, Borovička, Holec & Kolařík.

For this clade it is possible to use the name *Porotheleaceae* Murrill, while "*Trogiaceae*" Locq. is nom. inval. (Art. 39.1, Melbourne).

Porotheleaceae is part of the suborder *Marasmiineae* Aime, Dentinger & Gaya as circumscribed on a phylogenomic basis by DENTINGER *ET AL.* (2016) (= Marasmioid clade s. BINDER *ET AL.* 2010). Many taxa in this family have a sarcodimitic structure in the sense of Redhead (1987).

In our analysis (Fig. 1) a statistically well-supported *Porotheleaceae* family is recognized (MLB = 100%) only when it consists of the genera *Hydropus*, *Clitocybula*, *Leucoinocybe*, *Megacollybia*, *Porotheleum*, *Trogia* and some species of *Gerronema*. *Chrysomyцена* occupies an independent position in the *Porotheleaceae* and is sister (MLB = 99%) to a clade (MLB = 100%) formed by *Megacollybia*, *Trogia* and some species of *Gerronema*.

The genus *Megacollybia* is characterized by a *habitus* from collybioid to clitocyboid, a pileic fibrillary surface, abundant rhizomorphs at the base of the stipe, a sub-hymenidermic

to sub-trichodermic pileipellis (at least in the middle of the pileus), non-amyloid spores, widely clavate cheilocystidia (HUGHES ET AL., 2007; ANTONÍN ET AL. 2019), presence of clamp-connections and a sarcodimitic structure of the stipe (REDHEAD, 1987). The phylogenetically and morphologically closest genera are *Gerronema* and *Trogia*.

Gerronema is a broadly distributed genus (SINGER, 1964, 1970, 1986) with about 130 specific names (including synonyms) reported in the Index Fungorum database (<http://www.indexfungorum.org/>, last accessed 20/09/2019). The genus was originally proposed by Singer (1951) to accommodate three species (*G. melanomphax* Singer, *G. elasticum* Singer and *G. depauperatum* Singer) collected in the province of Tucumán (tropical Argentina) and characterized by a omphalinoid-to-clitocyboid growth, central or slightly eccentric long stipe, growth on rotting wood, irregular/confused lamellar texture, absence of cystidia, non-amyloid spores, intracellular pigment, present or absent clamp-connections. The definition of the genus has subsequently been met by numerous remodulations (ex. SINGER, 1964, 1970, 1975, 1986; BIGELOW, 1970; CLÉMENÇON, 1982; REDHEAD, 1986, 1987, 2002a, b; NORVELL ET AL., 1994). SINGER (1964-1986) had extended the concept of the genre by transferring some species previously included in *Omphalina* Qué. BIGELOW (1970) considered *Gerronema* synonymous posterior to *Omphalina*. Many authors (CLÉMENÇON, 1982; MOSER, 1983; KUYPER, 1986; REDHEAD, 1986, 1987, 2002a, b; NORVELL ET AL., 1994; ANTONÍN ET AL., 2008) considered *Gerronema* sensu SINGER (1964, 1975, 1986) as extremely artificial and heterogeneous. In this sense, the first molecular analyses of LUTZONI (1997), MONCALVO ET AL. (2002) and REDHEAD (2002a, b), based on the study of ribosomal genes, suggested the polyphlicity of the genus *Gerronema* as defined by Singer. NORVELL ET AL. (1994), following the studies of REDHEAD (1986, 1987) limited *Gerronema* to only lignicole species, with thin-walled spores, pileipellis in the form of cutis and with sarcodimitic structure. Thus delimited *Gerronema* would be monophyletic in the studies of MONCALVO ET AL. (2002), ANTONÍN ET AL. (2008, 2019), YANG ET AL. (2012), LIU ET AL. (2019), whose analyses, however, took into consideration only an extremely limited number of species and collections of *Gerronema* and/or of neighboring genera. Investigations in LATHA ET AL. (2018) and in the present work (Fig. 1), based on a wider taxon sampling, would indicate a polyphyletic status also for *Gerronema* s.s. in the sense of NORVELL ET AL. (1994).

Over the years, some species of *Gerronema* sensu SINGER (1951, 1964) have been included in the genera *Blasiphalia* Redhead, *Cantharellopsis* Kuyper, *Chrysomphalina* Cléménçon, *Haasiella* Kotl. & Pouzar, *Contumyces* Redhead, Moncalvo, Vilgalys & Lutzoni, *Gyroflexus* Raithehl. (= *Sphagnomphalia* Redhead, Moncalvo, Vilgalys & Lutzoni), *Lichenomphalia* Redhead, Lutzoni, Moncalvo & Vilgalys, *Loreleia* Redhead, Moncalvo, Vilgalys & Lutzoni and *Rickenella* Raithehl. (KOTLÁBA & POUZAR, 1966; CLÉMENÇON, 1982; KUYPER, 1986; REDHEAD, 2002a, b).

The genus *Trogia*, established by FRIES (1835) with the Indian species *Trogia montagnei* as type, includes generally lignicolous species with clitocyboids to omphalinoids basidioma of tenacious consistency and reviviscent *in situ* (KUMAR & MANIMOHAN, 2009, YANG ET AL., 2012). CORNER (1966), author of the first monographic work on the genus, adopted a very broad concept, recognizing 56 species united only by possessing a sarcodimitic structure. This extended concept of *Trogia* was much criticized by SINGER (1986), believing that Corner had over-emphasized the importance of the sarcodimitic structure; Singer limited the genus to only three species characterized by reviviscent basidioma, narrow and often forked gills, intricate lamellar texture and trichodermic pileipellis and distributed the others in *Gerronema*, *Hemimycena* Singer, *Hydropus* and *Mycena* (Pers.) Roussel. Redhead (1987), in a study aimed at understanding the taxonomic value of the presence of sarcodimitic tissues in the basidiomata of the *Agaricales*, emphasized the importance of this character. But instead of bringing together all the taxa with sarcodimitic tissues in a single genus (*Trogia*) as done by CORNER (1966), he recognized several genera that he included in the family *Xerulaceae* Jülich (now *Physalacriaceae* Corner partim).

In the 1991 monograph Corner reconfirmed his 1966 definition of *Trogia*, recognizing 90 species for Asia, Australasia and neotropics. In the sense of SINGER (1986) *Trogia* appears to be a good monophyletic genus (WILSON & DESJARDIN, 2005; YANG ET AL., 2012; DUTTA ET AL., 2017; ANTONÍN ET AL., 2019; Fig. 1 of this work) even though *T. montagnei*, type species of the genus, has never been sequenced (the holotypus is not traceable: CORNER, 1991; WILSON & DESJARDIN, 2005).

The genus *Hydropus* based on *Agaricus plexipes* b *fuliginaria* BATSCH (1783) contains species with mycenoid, omphalinoid to collybioid habit with amyloid or inamyloid spores and a pileipellis generally at least in a subimenniform part and/or with ventricose/swollen pileocystidia (SINGER, 1982, 1986; HAUSKNECHT ET AL., 1997; CONTU & ROBICH, 1998; BAS, 1999; ESTÈVE-RAVENTÓS ET AL., 2002). The genus is not monophyletic (see ex.: MONCALVO, 2002; ANTONÍN ET AL., 2019; and our analysis in Fig. 1) and the genus *Mycopan* Redhead, Moncalvo, Vilgalys was recently established for *Prunulus scabripes* Murrill 1916 (REDHEAD, 2013).

Among the genera similar to *Chrysomyцена* close to *Porothelaeaceae* (Fig. 1), *Phloeomana*, established by REDHEAD (2013) (type species *Agaricus speireus* Fr. 1815) and corresponding to species in *Mycena*, section *Hiemales* Konrad & Maubl. (HOLEC & KOLAŘÍK, 2017), includes species with grayish-brownish mycenoid basidiomata, growing on bark and dead wood, adnate gills, decurrent with tooth, non-amyloid spores, cheilocystidia of shape slightly different from that of basidia and hyphae of pileipellis with ramified digitations (MAAS GEESTERANUS, 1992; RONIQUIER & ARONSEN; 2007, REDHEAD, 2013; ARONSEN & LÆSSØE, 2016; ROBICH, 2016; HOLEC & KOLAŘÍK, 2017). *Atheniella* Redhead, Moncalvo, Vilgalys, Desjardin, B.A. Perry (species like *Agaricus adonis* Bull. 1793), corresponding to the adonis clade of MONCALVO ET AL., (2002) and to *Mycena*, section *Adonideae* (Fr.) Quél., is a genus consisting of mycenoid species with brightly coloured pileus of red, orange, yellow or white, ascending gills, inamyloid spores, cheilocystidia and generally fusiform pleurocystidia and elements of the pileipellis with simple diverticula or branched (MAAS GEESTERANUS, 1992; REDHEAD, 2012; ARONSEN & LÆSSØE, 2016; ROBICH, 2016).

Among the mycenoid-omphalinoid genera belonging to other families, morphologically the closest is *Chrysomphalina* (type species *Agaricus chrysophyllus* Fr. 1821), which phylogenetically belongs to the *Hygrophoraceae* (MONCALVO ET AL. 2002; MATHENY ET AL., 2006; VIZZINI ET AL., 2012; LODGE ET AL., 2014) and that stands out for the presence of carotenoid pigments, a pachipodial hymenium, a monomitic structure, the absence of clamp-connections and for poorly differentiated hymenial cystidia (CLÉMENÇON, 1982; NORVELL ET AL., 1994; LODGE ET AL., 2014).

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