1 Cortinarius sect. Riederi: taxonomy and phylogeny of the new section with European and

- 2 North American distribution
- 3

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68 Abstract

69	Cortinarius is one of the most species-rich genera of mushroom-forming fungi. Based on
70	phylogenetic and morphological evidence, a new section, Cortinarius, sect. Riederi, is
71	introduced. The taxonomy, phylogeny, ecology and distribution of mainly European but also
72	including some North American taxa of this section are treated, which includes 9 species and
73	two varieties). Of these, three taxa are described as new (C. burlinghamiae, C. pallidoriederi,
74	and C. argenteolilacinus var. dovrensis). The sect. Riederi species possess morphological
75	features similar to Phlegmacium group(s), but forms a phylogenetically isolated lineage, with
76	no supported affinity to other phlegmacioid groups. Three taxa are known from both Europe
77	and North America, two species are known only from North America, and five only from
78	Europe. Altogether eight of the ten taxa are associated with conifers or northern (boreal-
79	subalpine) deciduous trees (Betula spp.). Only two species occur in more temperate forests
80	(Fagus forests), and no species have so far been found in thermophilous Quercus forests.
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83	Key words
84	Agaricales, Basidiomycota, ecology, morphology, nrDNA ITS, phlegmacioid
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87	Introduction
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89	Cortinarius is by far the largest genus of macromycetes in the Northern Hemisphere. Many
90	groups are still poorly known, and deeper taxonomic studies (including integrative approaches
91	combining morpho-anatomical studies and molecular phylogeny) may lead to a 2-3 times
92	increase in the number of species (see e.g., Niskanen et al. 2013 on sect. Bovini; Brandrud et
93	al. 2014 on sect. Multiformes). According to Garnica et al. (2016), more than 800 species are
94	verified by distinct nrDNA ITS sequences. Niskanen et al. (2012a) estimated the number of
95	species in the Nordic countries of Europe to be approximately 900. A conservative estimate
96	for the amount of species in the Northern Hemisphere is 2000–2500 species. Already more
97	than 5000 names have been published in Cortinarius, worldwide (see Garnica et al. 2016;
98	Index Fungorum). However, this number does not reflect the real diversity in the genus, since
99	studies of type specimens have shown that many names are synonymous (see e.g.,
100	Liimatainen et al. 2014).
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102	Morphologically, Cortinarius has traditionally been subdivided into 5-8 subgenera (Moser
103	1967; Brandrud et al. 1989–2012; Knudsen and Vesterholt 2012), of which Phlegmacium,
104	Telamonia, and Dermocybe are the largest and most well-known. The subdivisions are also
105	supported by phylogenetic data to some degree, with the placement of most phlegmacioid
106	taxa into three major clades (Phlegmacium I, II and Calochroi; in the following referred
107	collectively to as "phlegmacioid species"), and most Dermocybe- + Telamonia-like species
108	forming a "superclade" (Stensrud et al. 2014; Garnica et al. 2016).
109	
110	Since the monograph of Moser (1961) on Cortinarius subgenus Phlegmacium in Europe, a
111	number of phlegmacioid groups/sections have been given an extensive taxonomic treatment:
112	sect. Phlegmacium (Brandrud 1996a, b), sect. Phlegmacioides (Brandrud 1998), sect.
113	Calochroi (Frøslev et al. 2006; 2007), (sub)sect. Elegantiores (Garnica et al. 2006), sect.
114	Claricolores (Brandrud et al. 2013), sect. Multiformes (Brandrud et al. 2014) and sect.
115	Purpurascentes (Saar et al. 2014). Furthermore, a cautiously updated taxonomy, which
116	includes most North and Central European species is provided in the descriptive key in the
117	Funga Nordica (Jeppesen et al. 2012). But comprehensive, taxonomic studies are still lacking
118	for many groups in Europe. Recent molecular studies of Phlegmacium (e.g. Garnica et al.
119	2003; 2005; 2009; 2016; Peintner et al. 2004; Frøslev et al. 2006; 2007, Liimatainen et al.
120	2014), identify a number of small, well-supported clades of phlegmacioid species with no
121	phylogenetic affinity to the three major Phlegmacium clades.
122	
123	According to phylogenetic studies (Garnica et al. 2016), Riederi is such a clade, which is also
124	morphologically well-delimited. Riederi species were formerly treated either as part of the
125	large sect. Glaucopodes (Brandrud et al. 1989–2012), or included in sections Glaucopodes
126	and Variecolores (Moser 1961) or Phlegmacium (Bidaud et al. 1999; Consiglio et al. 2006).
127	
128	The aim of this study is to present a detailed species level study of clade Riederi based on
129	molecular and morphological data, including studies of type specimens. An identification key
130	to all known European species is also provided.
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100	

133 Materials and Methods

134	
135	Molecular data
136	
137	A total of 112 specimens of clade Riederi from Europe and N America were sequenced (see
138	Collections examined under each species description), including the type specimens of C .
139	anomaloochrascens Chevassut & Rob. Henry, C. argenteolilacinus M.M. Moser, C.
140	fallaceicolor Rob. Henry (incl. "f. meridionale"), C. fulvoochrascens Rob. Henry, C.
141	fulvoochrascens var. cyanophyllus Rob. Henry, Cortinarius fulvoochrascens var.
142	subcaninicolor Rob. Henry, C. glaucocyanopus Rob. Henry, C. imbricatoides Rob. Henry, C.
143	malachioides P.D. Orton, C. marginatoochrascens Rob. Henry, C. notabilis Rob. Henry, C.
144	parksianus A.H. Smith, C. pseudoarquatus A.H. Smith, C. riederi (Weinm.) Fr., C. stilazureus
145	Rob. Henry, Phlegmacium fuscomaculatum var. smolandicum M.M. Moser, and P.
146	fuscomaculatum var. umbrinum M.M. Moser. For the majority of the specimens, the total
147	DNA was extracted from dried material with the NucleoSpin Plant II kit (Macherey-Nagel,
148	Düren, Germany). Primers ITS1F and ITS4 (White et al. 1990; Gardes and Bruns 1993) were
149	used to amplify the internal transcribed spacer region (ITS) of the ribosomal genes. The
150	protocols of the polymerase chain reaction (PCR) followed Niskanen et al. (2009). Some of
151	the specimens were amplified with the Phire® Plant Direct PCR Kit (Thermo Scientific,
152	U.S.A.) using the primers ITS1F and ITS4B (Gardes and Bruns 1993) and following the
153	method described in Papp and Dima (2017). Direct sequencing of the amplicons was
154	performed at the University of Helsinki, and LGC Genomics (Berlin, Germany) with the same
155	primers as those applied in PCR. Chromatograms were assembled and edited with Sequencher
156	4.1 (Gene Codes, Ann Arbor, MI, U.S.A.) and CodonCodeAligner 4.1. (CodonCode
157	Corporation, Centerville, MA, U.S.A.). BLAST queries of the public databases (GenBank:
158	http://www.ncbi.nlm.nih.gov/ and UNITE: http://unite.ut.ee/) were used to check for identical
159	or similar sequences.
1.00	

- 160
- 161 In order to evaluate the monophyly and phylogenetic placement of clade Riederi 108 ITS
- sequences of *Cortinarius* were sampled across the main lineages of the genus known from the
- 163 Northern Hemisphere. One sequence of each species from clade Riederi was chosen for the
- analysis, while the other *Cortinarius* lineages were represented by two to four sequences per
- 165 clade. *Hebeloma* was chosen as outgroup (Table 1). Sequences were aligned with MAFFT
- 166 (online version 7) using the E-INS-i strategy with default settings (Katoh and Toh 2008). The This is a post-peer-review, pre-copyedit version of an article published in Mycological Progress. The final authenticated version is available online at: http://dx.doi.org/<u>10.1007/s11557-018-1443-0</u>

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- 167 alignment was imported to SeaView (Gouy et al. 2010) for visual inspection. Indels were
- 168 coded as presence/absence characters with FastGap 1.2 (Borchsenius 2009) following the
- 169 simple indel coding algorithm (Simmons et al. 2001). Adding indel characters to the
- 170 nucleotide alignment of ITS sequences increases the robustness of the phylogenetic analyses
- 171 (Nagy et al. 2012; Brandrud et al. 2015; Seress et al. 2016).
- 172
- The second dataset to study the species limits within the clade Riederi was assembled using99 sequences of the target clade of which 87 were generated for this study and 12 were
- downloaded from public repositories (Table 2). Multiple sequence alignment was done with
- 176 PRANK (Löytynoja and Goldman 2005; 2008) as implemented in its graphical interface
- 177 (PRANKSTER) under default settings. The resulting alignment contained 574 positions.
- 178 Indels were coded with the same indel coding algorithm as described above. The final matrix
- 179 was composed of 609 positions (nucleotides + binary data). New sequences were submitted to
- 180 GenBank and the accession numbers are available in Table 1 and 2. The alignment was
- 181 deposited at TreeBase (TB2:S22806).
- 182
- 183 Phylogenetic analyses
- 184
- 185 Maximum Likelihood (ML) analysis was carried out for both datasets using RAxML
- 186 (Stamatakis 2014) as implemented in raxmlGUI (Silvestro and Michalak 2012). For testing
- 187 the support of the branches, rapid bootstrap analysis with 1000 replicates and ML search were
- 188 chosen. For the partitioned dataset the GTRGAMMA substitution model for the DNA and the
- 189 default set for binary (indel) characters were applied. The resulting ML phylogenetic tree (Fig.
- 190 1) was edited with MEGA6 (Tamura et al. 2013).
- 191
- 192 For the dataset 2 Bayesian inference (BI) analysis was performed with MrBayes 3.1.2
- 193 (Ronquist and Huelsenbeck 2003). The alignment was partitioned and the GTR+G model of
- 194 evolution was applied for the ITS and the 2-state Markov model for the indel characters. Four
- 195 Markov chains and two independent runs were performed for 10 000 000 generations,
- 196 sampling every 1000 steps, resulting in overall sampling of 10 001 trees. The first 4000 of the
- 197 sampled trees (40%) was excluded before constructing the 50% majority rule consensus tree
- and calculating the posterior probabilities (PP). The Markov Chain Monte Carlo (MCMC)
- convergence was checked with AWTY online (Nylander et al. 2007).
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200 Tree topologies of both BI and ML analyses were checked visually and no incongruence was

201 observed, only the ML tree is shown in Fig. 2. Branches were considered strongly supported

when BI posterior probability values (PP) exceeding at least 0.90 and ML bootstrap values

203 (BS) at least 70%. Phylogenetic trees were visualized and edited in MEGA6 (Tamura et al.

204 2013) (Figs 1 and 2).

205

206 Morphological studies

207

208 Fresh material was examined mainly from N, C, W and S Europe, and some from N America. 209 More collections than those used for DNA-sequencing were studied morphologically, but the 210 character variation described in the Taxonomy section is based on the sequenced material 211 alone, as no obvious discrepancies were observed. Collections sequenced and examined 212 morphologically are listed under each species description (and in Table 2). Most specimens 213 (except some types) were examined both macro- and micromorphologically. The 214 morphological descriptions are only based on specimens studied by the authors. The 215 measurements of macromorphological characters were based on expanded, but never old (and 216 then often aberrant) basidiocarps. The terminology of characters follows Brandrud et al. 217 (1990) and Brandrud (1996a).

218

219 Microscopical structures were observed either from fresh material mounted in H_2O , often 220 with a drop of 40% KOH added subsequently, or from dried material mounted in H₂O and 221 then in KOH. Basidiospore measurements were made at 1000× magnification with a 222 calibrated optical micrometer or on a flat screen with the program ProgRes® CapturePro from 223 Jenoptik. The measurements are based on at least 20 spores from each specimen/basidiocarp; 224 numbers in square brackets refer (in this order) to the number of collections they originate 225 from, the number of basidiocarps (sometimes more than one basidiocarp has been measured 226 per collection) and the number of spores measured, respectively. Spore measurements are 227 given as follows: length range \times width range followed by the mean values in brackets then the 228 variation of the MVs. Q values were calculated as follows: Q = length divided by width. 229 Mean values of the Q is also presented. To exclude aberrant spores the given values are based 230 only on spores within the 95% confidence interval. For comparative spore measurements see 231 also Table 3.

232

The photo micrographs of the spores (Fig. 3) are created using z-stacking (Schmidt-Stohn 233 234 2011). For this, about ten pictures taken with a Jenoptik ProgRes® C10 plus digital camera, at 235 different focus heights, were combined into the final picture with the Helicon Focus 6.5 236 program. For the correction and the final arrangement of the spores on the plates Adobe 237 Photoshop CS5 was used. The column graph of mean values of spore lenghts (Fig. 4), the scatter 238 diagram of spore mean values (Fig. 5) and the 95% confidence ellipses of spore 239 variations (Figs 6 and 7) were calculated with a special Microsoft Excel file (Documenta Geigy, 240 pers. com. of H. Clémençon). For generating figures the graphics were copied from Excel and

- 241 242
- 243
- 244 **Results and discussion**

edited with Adobe Photoshop CS5.

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246 Molecular phylogeny

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248 The ML analysis (Fig. 1) of the dataset 1 confirmed that the clade Riederi is a well-supported 249 monophyletic group within genus Cortinarius, in line with the results of Garnica et al. (2016). 250 According to multigene phylogenetic analysis using the ITS, LSU, and *rpb1* gene regions 251 performed by Garnica et al. (2016), the clade Riederi nested basally in a large well-supported group called 'Phlegmacioid clade II', together with e.g. sect. Multiformes, sect. Scauri, and 252 253 the C. lustratus and C. pinophilus groups. ITS sequences of clade Riederi species were 254 relatively dissimilar to other Cortinarius taxa when evaluating BLAST hits, with only 88% 255 similarity to the best match in GenBank (e.g. C. borgsjoeensis). Molecularly, the Riederi species also differ from typical phlegmacioid species in other lineages (e.g. Calochroi, 256 257 Infracti, Phlegmacioides) in the length of the ITS region, which in clade Riederi is more 258 similar to that of species in the Telamonia lineage (= Telamonia s. str.). Both our analyses and 259 those of Garnica et al (2016) clearly indicate that clade Riederi is an isolated lineage within 260 *Cortinarius*, and this is also supported by morphological data (see below). Therefore, we here 261 describe this well-circumscribed group formally as a new section (see Taxonomy section). For 262 a more precise infrageneric placement of this clade, multigene phylogenetic analyses 263 combined with deeper taxon sampling will need to be done in the future. 264 The analysis of the second dataset revealed 11 OTUs within the clade Riederi (Fig. 2). Nine of 265 these have a distinct 'barcoding gap', each OTU being separated by at least 1.2% (7 This is a post-peer-review, pre-copyedit version of an article published in Mycological Progress. The final authenticated version is available online at: http://dx.doi.org/10.1007/s11557-018-1443-0

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substitution and indel positions) from sister species, and with little internal variation (0-0.7%), 266 267 0-4 changes). Eight of these OTUs can also be morphologically distinguished and are here 268 treated as species (see Taxonomic part). Two OTUs in the *C. argenteolilacinus* complex 269 showed only a small 'barcoding gap' of 0.5% (3 substitutions and indel positions). This 270 percentage is smaller than normally used 1% as a minimum cut-off value for recognizing 271 species of *Cortinarius* based on ITS sequences (see e.g. Frøslev et al. 2007; Liimatainen et al. 272 2014; Garnica et al. 2016). Therefore we have decided to treat these clades as varieties although they also have morphological and ecogeographical differences (see Taxonomic part). 273 274 The one additional unsupported OTU (probably a complex of several species) from N 275 America assigned here as *Cortinarius* sp. is not further studied here due to lack of data. 276 277 Morphological and ecological characters used to circumscribe the species in sect. Riederi 278 279 Stipe shape: Already Fries (1836) and then later Moser (1961) made a major distinction 280 between phlegmacia with a marginate bulb and those with a cylindric-clavate stipe. This is 281 still largely a good diagnostic character to distinguish major phylogenetic lineages such as 282 sect. Phlegmacioides (cylindric stipe) and sect. Calochroi (flattened, marginate bulb) (see 283 Garnica et al. 2006; Frøslev et al. 2007). In sect. Riederi, however, this feature is less relevant 284 as a diagnostic character, since the species have a rather irregular stipe shape, some with a 285 mainly clavate-bulbous stipe, and some with a more marginate bulb, but usually with a fairly 286 large infraspecific variation. 287 Lamellae colour: A bright, deep violaceous colour is present in young lamellae and stipe of 288 most *Riederi* species (paler bluish in *C. malachioides*). However, this pigment vanishes with 289 age and exposure, resembling the situation with bright bluish coloured species in sect. 290 Glaucopodes. 291 Pileus colour and structure: In contrast to the above-mentioned features, pileus colour is an

important diagnostic character on the species level in sect. *Riederi*; ranging from vivid

redbrown-ochre colours in *C. fulvoochrascens* and *C. riederi* to pale (bluish-)grey in *C.*

294 argenteolilacinus. The Riederi species normally possess a finely innately fibrillose structure

of the pileus cuticle. This is due to darker pigmented, innate filaments that form a braided-

veined-structure towards the margin, and a denser, almost marbled pattern towards the centre.

297 The structure is especially distinct in specimens of *C. riederi* with a strong ochraceous-

298 redbrown colouration, but can be seen in all taxa (see also under Micromorphology). This is a post-peer-review, pre-copyedit version of an article published in Mycological Progress. The final authenticated version is available online at: http://dx.doi.org/10.1007/s11557-018-1443-0

- Although seen in some other species, such as *C. dionysae*, *C. napus* and *C.*
- 300 *aureopulverulentus*, such regular, densely braided pattern of innate fibrils is rarely
- 301 encountered in other phlegmacioid taxa.
- 302 Universal veil: Unlike most related groups, the universal veil in most taxa of sect. Riederi is
- 303 sparse, leaving almost no visible traces on the pileus or the stipe surface. Only in *C*.
- 304 *fulvoochrascens* and sometimes in *C. argenteolilacinus* can distinct veil remnants be seen,
- 305 near the bulb- and pileus margin.
- 306 *KOH-reactions*: Macrochemical reactions with KOH are always negative (slightly grey
- 307 brownish discolouring), and are not further mentioned in the Taxonomic part.
- 308 Basidiospores: The spore shape and ornamentation is fairly uniform in sect. Riederi. The
- 309 entire group is characterized by rather ellipsoid shaped spores with a more or less rounded
- 310 apex, never with a sharply pointed-elongated apex as seen in the spores of many other
- 311 phlegmacioid species (Fig. 3). Such ellipsoid-subamygdaloid spores occur also in *C*.
- 312 glaucopus agg. and in C. eucaeruleus-C. coerulescentium complex. The ornamentation also
- 313 resembles that of the spores of the latter mentioned taxa; with a pattern of rather dense,
- 314 discrete warts without interconnected ridges and crusts as seen in sect. Calochroi. A
- 315 characteristic feature of the spores of the species of sect. *Riederi* is the apex, where the
- 316 ornamentation often is more or less confluent, forming an apical "lense", sometimes also with
- 317 some prominent, subapical "teeth". This feature is especially pronounced in *C*.
- 318 argenteolilacinus. The Q value (spore length/width ratio) is very similar in all species, with
- 319 average values ranging from 1.59 to 1.69 (Table 3).
- 320 The spore size is the most important diagnostic character at the species level (Figs 4, 5, 6 and
- 321 7). When spores are measured from the stipe/cortina (excluding aberrant, unripe ones), and
- 322 mean values (MVs) from each measurement series are compared, many species show
- 323 considerable differences. For instance, among the coniferous forest species the comparatively
- 324 small-spored *C. anomaloochrascens* (spore MVs $10.9-11.9 \times 6.9-7.6 \mu m$) can be
- 325 distinguished from the sometimes co-occurring, large-spored Cortinarius riederi (spore MVs
- 326 $12.3-13.6 \times 7.9-8.7 \,\mu$ m) and *C. fulvoochrascens* (spore MVs $12.1-12.7 \times 7.6-8.2 \,\mu$ m) (Figs
- 4 and 6), and the small-spored, Betula-associated C. malachioides (spore MV 10.1–11.3×6.3–
- 328 7.1 µm) from the co-occurring, large-spored C. argenteolilacinus var. dovrensis (spore MV
- 329 $11.8-13.5 \times 7.3-8.2 \ \mu m$) (Figs 4 and 7).
- 330 *Pileus cuticle*: A major subdivision of *Phlegmacium* can be performed based on the structure
- of pileipellis (Brandrud et al. 1990). Sect. *Riederi* shares with sections such as *Glaucopodes* This is a post-peer-review, pre-copyedit version of an article published in Mycological Progress. The final authenticated version is available online at: http://dx.doi.org/10.1007/s11557-018-1443-0

and *Scauri* a duplex structure of pileipellis with a thin epicutis of narrow hyphae and a

333 subcutis/hypodermium consisting of broader, shorter, tightly cemented, "subcellular"

334 elements. Some sections such as *Calochroi*, *Cliduchi* and *Phlegmacioides*, on the other hand,

are characterized by a simplex pileipellis consisting of a thick, more or less homogeneous

336 layer of thin, long, usually loosely entangled hyphae. In sect. *Riederi* species, the transition

337 layer between the epicutis and the hypodermium consists of irregularly interconnected

bundles of rather narrow, long hyphae with more or less encrusted pigment, giving rise to the

innately fibrillose structure described under macrocharacters. The degree of encrustations of

340 these long, transition hyphae is an important diagnostic character of some species. The major

341 part of the hypodermium consists in sect. *Riederi* species of moderately inflated, 15–20 µm

342 wide elements. In C. argenteolilacinus, the hypodermium is weakly developed, and elements

343 are only little inflated.

344 Distinguishing features towards similar sections/clades: Morphologically, the species of sect.

345 *Riederi* resemble those of the *C. glaucopus* group (deep violet lamellae/stipe), the *C. talus-C.*

346 *multiformis* group (ochre yellow pileus with a similar surface structure) as well as the *C*.

347 *largus* complex (bluish-greyish pileus). The most important character for distinguishing the

348 *Riederi* species from these are the large spores $> (10.5)11 \,\mu\text{m}$ long. Furthermore, the innately

349 fibrillose surface of the pileus and the lack of a distinctly marginate bulb also clearly differ

350 from other taxa. Section *Riederi* is further characterized by the lack of distinct smells.

351 Habitat and distribution: Half of the eight European taxa are associated with coniferous trees,

the other half with deciduous ones. The widespread *C. riederi*, *C. anomaloochrascens* and *C.*

353 *fulvoochrascens* are mainly associated with *Picea abies*. *Cortinarius argenteolilacinus* var.

354 argenteolilacinus is found under Fagus sylvatica in nemoral/temperate regions of C Europe

and S Scandinavia, and under *Tilia* and *Corylus* in S Sweden and S Norway. *Cortinarius*

356 argenteolilacinus var. dovrensis and C. malachioides are recorded mainly under Betula

357 *pubescens* subsp. *tortuosa* in Scandinavian mountains.

358 For most species a preference for boreal-montane regions can be assumed. At least five of the

359 eight European taxa are associated with boreal-montane coniferous trees or with boreal

360 deciduous trees. Only C. argenteolilacinus var. argenteolilacinus and C. glaucocyanopus

361 seem to have a preference for more temperate (mixed) *Fagus* forests. Most species show an

362 overlapping distribution in the boreonemoral region of S Scandinavia, which can be regarded

363 as a region of high diversity for sect. *Riederi* in Europe. In contrast, none of the species occurs

364 in thermophilous (sub)mediterranean *Quercus* forests. The taxa of sect. *Riederi* are also This is a post-peer-review, pre-copyedit version of an article published in Mycological Progress. The final authenticated version is available online at: http://dx.doi.org/10.1007/s11557-018-1443-0

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365	characterized by less calciphilous requirements than most other phlegmacioid species.
366	Cortinarius riederi prefers oligotrophic sites, and also the other taxa rarely grow on
367	calcareous ground, and rarely occur in typical <i>Phlegmacium</i> hotspots (the exception being C.
368	fulvoochrascens).
369	Comparatively little is known about the distribution of the species of sect. Riederi outside
370	Europe. The two preferentially boreal(-subalpine) taxa C. riederi and C. argenteolilacinus var.
371	dovrensis are also found in N America, and probably have a circumpolar distribution. Two
372	western N American (non-European) species are included in sect. Riederi (C. parksianus and
373	C. burlinghamiae), as well as one unknown species complex from Canada and Alaska (Fig.
374	2), but generally, the Riederi group seems to be understudied in N America, and more taxa are
375	probably to be discovered.
376	
377	Taxonomy
378	
379	Cortinarius sect. Riederi Brandrud, Dima, Niskanen & Liimat., sect. nov.
380	MycoBank MB 827874
381	
382	Basidiocarps fleshy and phlegmacioid. Pileus glabrous and viscid-glutinous, more or less
383	innately fibrillose, vividly yellow brown to redbrown, pale greybrown to almost whitish.
384	Lamellae usually deep violaceous blue when young, soon fading to greyish brown. Stipe with
385	a clavate-bulbous base (rarely slightly marginate-bulbous), glossy, initially usually deep
386	(greyish) violet in upper part, soon fading to greyish white, often developing glossy brass
387	brown patches. Universal veil very sparse and hardly visible on most species. Context
388	fugacious violet.
389	Basidiospores large (10–14(15) \times 6–9 μ m), ellipsoid to subamygdaloid, densely vertucose
390	(with discrete warts), sometimes with an apical "lense" of confluent warts. Lamellar edge
391	more or less fertile. Pileipellis duplex, composed of thin, gelatinized epicutis and subcellular
392	subcutis/ hypodermium with brown parietal-encrusting pigment, in transition to epicutis many
393	species with bundles of zebra-striped-crustulose encrusted hyphae.
394	
395	Type species: Cortinarius riederi (Weinm.) Fr.
396	
397	Cortinarius riederi (Weinm.) Fr., Epicr. Syst. mycol.: 259 (1838).
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- 398 MycoBank: MB 155615
- 399 Figs 8a and 8b
- 400
- 401 Basionym: Agaricus riederi Weinm., Hymenom. Gasterom. Imp. Ross.: 161 (1836).
- 402 MycoBank: MB 244037. Type: Switzerland, Graubünden, Engadin, J. Favre, 26 Aug 1944
- 403 (sub nom. C. lilacinopes), GK13347 (G), neotype designated in Melot (1986). GenBank:
- 404 MH923060.
- 405
- 406 Synonyms: Cortinarius pseudoarquatus A.H. Smith, Lloydia 7(3): 181 (1944). MycoBank:
- 407 MB 285764. Holotype: U.S.A., Washington, Olympic National park, near Olympic Hot
- 408 Springs, 22 Sept 1941, A.H. Smith, AHS17141 (MICH10397).
- 409 Phlegmacium fuscomaculatum var. smolandicum M.M. Moser, Die Gatt. Phlegm.: 355
- 410 (1961). MycoBank: MB 349991. Holotype: Sweden, Småland, Femsjö, 9 Sept 1957, M.
- 411 Moser, MM 57/60 (M). Cortinarius fulvoochrascens var. smolandicus (M.M. Moser) Rob.
- 412 Henry, Bull. trimest. Soc. mycol. Fr. 105(4): 331 (1989). MycoBank: MB 126140. –
- 413 Cortinarius fuscomaculatus f. smolandicus (M.M. Moser) Nespiak, Flora Polska, Grzyby
- 414 (Mycota), 7, Basidiomycetes, Agaricales, Cortinariaceae, Cortinarius 1 (Warszawa-Krakow):
- 415 147 (1975). MycoBank: MB 347980.
- 416
- 417 Selected icones: Moser (1961), Die Gatt. Phlegm.: Pl. XVII, 95 (as C. lilacinopes neotype
- 418 for *C. riederi*), Pl. XXII, 127 (as *P. fuscomaculatum* var. *smolandicum*); Soop 2017, Pl. 5:17
- 419 as C. pseudoarquatus.
- 420
- 421
- 422 *Pileus* 5–10(12) cm, (hemi-)spherical(-campanulate), then plano-convex (to broadly
- 423 umbonate), viscid-glutinous and glossy when dry, towards margin distinctly to strongly
- 424 innately fibrillose-rivulose, centre with a fine marbled structure from darker, innate fibers,
- 425 sometimes with darker hygrophanous spots or veins towards margin, a few also with a narrow
- 426 hygrophanous ring near margin; ochraceous brown-fulvous to somewhat more orange-red
- 427 brown tinged at centre, the involute margin whitish brown. Universal veil remnants very
- 428 sparse, hardly visible.
- 429 *Lamellae* crowded (L = 80-110), 0.5–1 cm broad, initially deep violet to greyish violet, but
- soon fading to paler bluish grey and then greyish brown; edge even to crenulate.
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- 431 Stipe $6-12 \times 1-2(2.5)$ cm, with a clavate-bulbous base (up to 4.5 cm), bulb rarely slightly
- 432 (irregularly) marginate, glossy, initially deep (greyish) violet in upper part and with a pale
- 433 bluish tinge downwards, soon fading to greyish white-pale ochraceous brown, often
- 434 developing glossy brass brown patches (like *C. glaucopus*, *C. multiformis*), often with distinct
- 435 patches of watery bluish white tomentum towards base (sometimes developing strongly with
- 436 age). Universal veil very sparse and hardly visible.
- 437 *Context* initially deep violet to greyish violet in (upper part) of stipe, bluish white in pileus
- 438 and bulb, violet colour soon fading, becoming whitish with brass brown spots where
- 439 damaged.
- 440 *Macrochemical reactions* negative, KOH somewhat brownish in context.
- 441 *Smell* faint, sometimes yeast-like with age.
- 442 *Exsiccata* pale brownish with vivid brown-redbrown pileus.
- 443 *Basidiospores* [10, 15, 249], 11.6–14.2 × 7.3–9 μ m (MV = 12.9 × 8.2 μ m); variation of MVs:
- 444 $12.3-13.6 \times 7.9-8.7 \,\mu\text{m}; Q = 1.44-1.72 \,(\text{MV} = 1.58)$, ellipsoid-subamygdaloid, medium to
- 445 strongly and densely verrucose, sometimes with an apical "lense" of confluent warts;
- 446 suprahilar plague hardly visible.
- 447 *Lamellar edge* more or less fertile, sterile cells small, clavate or cylindrical.
- 448 *Pileipellis* duplex. Epicutis thin. At surface 2–5 µm wide, loosely erect-entangled, gelatinous
- 449 hyphae, hyaline or with pale, refractive pigment when collapsed. Subcutis/hypodermium
- 450 hyphae 5–20 μm wide, tightly cemented, with brown parietal-encrusting pigment, forming an
- 451 amber-like embedment; in transition to epicutis with subparallel, interconnected bundles of 3–
- 452 6(10) long and rather narrow 5–7(10) µm wide hyphae, with distinct to strong, brown, zebra-
- 453 striped-crustulose encrusted pigment. These bundles are normally easy visible when the thin
- 454 pellicle is peeled off.
- 455 *Veil at pileus surface* absent or very thin, of narrow, hyaline hyphae.
- 456 Habitat: Associated with Picea abies, probably also Pinus sylvestris (cf. Favre 1960, Moser
- 457 1961 as *C. lilacinopes*). In oligotrophic *Vaccinium myrtillus* type of forests, sometimes also in
- 458 richer/mesotrophic low-herb spruce forests, sometimes in bog rims. Frequently in young,
- 459 mossy *Picea* plantations.
- 460 *Distribution*: Mainly recorded from Fennoscandia where the species is widespread and rather
- 461 frequent in boreal regions (Moser 1961 as *Phlegmacium fuscomaculatum* var. *smolandicum*;
- 462 Soop 2011 as *C. fulvoochrascens* p.p.; Jeppesen et al. 2012). Rarely occurring also in
- 463 subalpine *Picea-Pinus* forests in the Alp-Prealp regions, so far recorded from S Tyrol, Italy This is a post-peer-review, pre-copyedit version of an article published in Mycological Progress. The final authenticated version is available online at: http://dx.doi.org/<u>10.1007/s11557-018-1443-0</u>

and Swiss national park, Engadin (neotype; see Moser 1961 as C. lilacinopes), probably also 464 in French Jura and French Alps (Bidaud et al. 1999 as C. fulvoochrascens var. umbrinus and 465 466 C. lilacinopes, material not sequenced). Recorded up to 2150 m asl. in the Swiss Alps (Favre 467 1960, as C. lilacinopes). Probably wider distributed in C Europe, but little known due to 468 confusion with the more frequent C. fulvoochrascens and C. anomaloochrascens. Recorded 469 also in western North America (Smith 1944, as C. pseudoarquatus; type conspecific with C. 470 riederi). 471 Comments: Cortinarius riederi is characterized by its (i) distinctly innately fibrillose and 472 vividly ochre brown-redbrown pileus, (ii) deeply, but fugacious violet lamellae and stipe, (iii) 473 irregularly bulbous stipe and (iv) (very) large spores. The stipe is furthermore glossy-shiny 474 often with watery blue tomentum in lower part and it soon becomes brass-brown spotted. The 475 watery blue felt is often a very characteristic feature, but is not constant, and seems often to

- 476 develop with age. The species has a mainly northern distribution in Europe, preferring
- 477 oligotrophic-acidophilous spruce forest types. It sometimes occurs in large groups in young,
- 478 mossy *Picea* plantations. In montane-subalpine sites in C Europe, acidophilous habitats such
- 479 as margins of bogs might be characterizing. *Cortinarius riederi* (= *C. fulvoochrascens* s. auct.)
- 480 has by most authors been treated collectively, including taxa with innately fibrillose and
- 481 vividly fulvous, umber to ochre brown pileus, initially deeply violet lamellae and stipe and
- 482 large, ellipsoid-subamygdaloid spores. However, according to molecular studies, the *C*.
- 483 riederi complex in its widest sense can be separated into a number of well-supported taxa. On
- 484 closer examination, these are also distinguishable morphologically and partly
- 485 ecogeographically; for instance the large-spored *C. riederi* s. str., *C. fulvoochrascens* and *C.*
- 486 *pallidoriederi* as well as *C. anomaloochrascens* with somewhat smaller spores. In
- 487 Fennoscandia, more than half of the C. riederi s. lato collections belong to the present species,
- 488 the remaining mainly to *C. anomaloochrascens*. In C Europe about fifty-fifty of the
- 489 collections belong to *C. fulvoochrascens* versus *C. anomaloochrascens*, whereas *C. riederi* s.
- 490 str. apparently is rare, and restricted to (montane-)subalpine *Picea-Pinus* forests.
- 491 Cortinarius riederi s. str. is often confused with the large-spored C. fulvoochrascens, and
- 492 there is no consistent name-use for these in the literature. *Cortinarius fulvoochrascens* can be
- 493 distinguished by the distinct veil remnants on stipe when very young, and it has furthermore
- 494 (i) a more uniform ochre-fulvous-umber brown, less glabrous, less innately fibrillose pileus,
- 495 (ii) usually a more clavate-bulbous stipe, (iii) lack of watery bluish mycelia tomentum on the
- 496 bulb, and (iv) on average slightly smaller and more strongly vertucose spores. According to This is a post-peer-review, pre-copyedit version of an article published in Mycological Progress. The final authenticated version is available online at: http://dx.doi.org/10.1007/s11557-018-1443-0

497 the spore-measurement-series performed of sequenced collections (12 specimens from 10

- 498 collections), the mean values of spore length of C. riederi in half of the measurement-series
- 499 exceed 13.0 μ m, whereas MV's >13.0 μ m are not observed from *C. fulvoochrascens* so far
- 500 (11 specimens from 7 collections; see Table 3). *Cortinarius fulvoochrascens* also apparently
- 501 has its main distribution in montane C Europe (where *C. riederi* s. str. can hardly be found).
- 502 In N European regions where both species might co-occur, *C. fulvoochrascens* occurs in
- 503 richer, often calcareous forest types, whereas C. riederi is confined to more oligotrophic/acid
- 504 sites. *Cortinarius fulvoochrascens* reported in Moser et al. (1995) from N America might
- 505 belong to *C. riederi*, based on the spore measurements given (see discussion under *C*.
- 506 *fulvoochrascens*).
- 507 According to molecular data, C. riederi is more closely related to C. anomaloochrascens than
- 508 to C. fulvoochrascens. Cortinarius anomaloochrascens sometimes occurs in the same,
- 509 acidophilous habitats as *C. riederi*, but is well distinguished by smaller and often more
- 510 amygdaloid spores. Furthermore, C. riederi normally has somewhat more vivid/warmer and
- 511 darker pileus colours, larger basidiocarps with a more clavate stipe and stronger, encrusting
- 512 pigment of pileipellis.
- 513 According to the protologue of Weinmann (1836), C. riederi might be a collective species,
- 514 including at least *C. riederi* in our sense and *C. anomaloochrascens* which is widely
- 515 distributed in boreal areas resembling the St Petersburg region where Weinmann collected.
- 516 Cortinarius fulvoochrascens is less likely to be included in the protologue, since calcareous
- 517 coniferous forests seem to be lacking or rare in the St Petersburg region. A neotype of *C*.
- 518 riederi was selected by Melot (1986), representing a collection of C. lilacinopes as described
- and depicted by J. Favre in Favre (1960) and in Moser (1961). With the wide concept in the
- 520 protologue, *C. riederi* s. str. might well be included here, so the selection of neotype can
- 521 hardly be regarded as in serious contradiction with the protologue, and should be accepted.
- 522 The type has been studied microscopically, and corresponds well with the present species, but
- 523 could also cover *C. fulvoochrascens*. However, the mean value $13.0 \times 8.2 \,\mu\text{m}$ of spores (from
- 524 stipe surface) is larger than that of all measured collections of *C. fulvoochrascens*, and fits
- 525 well with our species. The collection of Favre is from a subalpine coniferous forest of the
- 526 Engadin valley in the Alps, and this high altitude site on siliceous soils corresponds well with
- 527 the present species. Furthermore, after several sequencing attempts, the type yielded an ITS1
- 528 sequence corresponding to the present taxon (Fig. 2).
- 529

530	
531	Collections examined:
532	Finland. Savonia Borealis, Vehmersalmi, Pitkälahti, J. Vauras, JV 1475F (herb. TUR).
533	Uusimaa, Luukki, I. Kytövuori, IK93-664 (H). Kittilän Lappi, Kolari, I. Kytövuori, IK98-
534	1171 (H). Italy. S Tyrol, Anterselva, 19 Aug 2005, C. Rossi, Rossi 19-08-2005 (herb. Rossi);
535	S Tyrol, 25 Aug 2013, G. Turrini, TG25-08-2013 (herb. Turrini); 12 Aug 2012, F. Bellù, Bellu
536	12-08-2012 (herb. BOZ), 14 Aug 2012, C. Rossi (herb. Rossi). Norway. Oppland, Søndre
537	Land, Hov NW, T.E. Brandrud, B. Dima, TEB407-13/DB5109 (O); Vestre Slidre, Einang
538	Bridge S, T.E. Brandrud, TEB443-11 (O). Etnedal, Solbjør N, T.E. Brandrud & B. Dima,
539	TEB258-17 (O). Sør-Trøndelag, Trondheim, Ranheimsskogen, T.S. Jeppesen TSJ2001-004
540	(C), Nord-Trøndelag, Lierne, K. & E. Bendiksen, TE. Brandrud; KB-EB-TEB 2-10 (O).
541	Sweden. Jämtland, Bräcke, Ammersön, T.E. Brandrud, TEB141-10/SSt10-066 (O). Småland,
542	Femsjö, M. Moser, MM 57/60 (M, holotype of Phlegmacium fuscomaculatum var.
543	smolandicum). Switzerland. Graubünden, Engadin, J. Favre, 26 Aug 1944, GK13347 (G,
544	neotype). U.S.A. Washington, Olympic National Park, Olympic Hot Springs, A.H. Smith,
545	AHS17141 (MICH10397, holotype of C. pseudoarquatus).
546	
547	Cortinarius pallidoriederi Brandrud, Dima & Bellù, sp. nov.
548	MycoBank MB 827875
549	Fig. 8h
550	
551	Etymology: the epithet refers to the paler colour of the pileus than that of C. riederi
552	
553	Holotype: Italy, S Tyrol, Val d'Ega (Eggental), Novale (Rauth), Wald Lagarn, between
554	Underkor and Oberkor, E. Castellini, F. Bellù, 30 Sept 2011, Bellu 30-09-2011 (BOZ).
555	GenBank: MH923049.
556	
557	
558	Pileus 4-8 cm, (hemi-)spherical, then plano-convex, viscid-glutinous, finely innately
559	fibrillose-rivulose from darker, innate fibers; uniform pale ochraceous yellow to ochraceous
560	brown, with an olivaceous ochre tinge. Universal veil remnants sparse, rendering surface
561	shiny-micaceous-marbled when young.

- 562 *Lamellae* crowded (L = 80-90), 0.4–0.8 cm broad, initially deep violet to greyish violet, but
- soon gradually fading to greyish brown; edge even to slightly sinuate.
- 564 *Stipe* $5-8 \times 1.2-2$ cm, with a slightly marginate bulb (up to 3 cm), glossy, longitudinally
- striate, initially deep, metallic (greyish) violet in upper part and with a pale bluish tinge
- 566 downwards, soon fading to greyish white-pale ochraceous brown, often developing glossy
- brass brown-golden brown patches, especially when bruised. Universal veil very sparse and
- 568 hardly visible.
- 569 *Context* initially violet to greyish violet in upper part of stipe, mainly in cortex, otherwise
- 570 whitish, violet colour soon fading, some yellow brown spots where damaged.
- 571 *Smell* faint, indistinct.
- 572 *Exsiccata* pale brownish with almost whitish pileus.
- 573 *Basidiospores* [1, 4, 65], 11.1–13.8 × 6.3–8.5 μ m (MV = 12.4 × 7.4 μ m); variation of MVs:
- 574 $12.1-12.7 \times 6.9-7.7 \ \mu m$; Q = 1.45-1.92 (MV = 1.69), subamygdaloid-ellipsoid, strongly and
- 575 densely vertucose, sometimes with an apical "lense" of confluent warts; suprahilar plague
- 576 hardly visible.
- 577 *Pileipellis* duplex. Epicutis thin. At surface 3–4 µm wide, loosely erect-entangled, gelatinous
- 578 hyphae, hyaline or with pale, refractive pigment when collapsed, some terminal hyphae
- 579 clavate (up to 6 μm wide). Subcutis/hypodermium distinct, hyphae 5–20 μm wide, tightly
- 580 cemented, with pale yellow brown parietal-encrusting pigment, forming an amber-like
- 581 embedment, some dark brown oleifers intermixed; in transition to epicutis with subparallel,
- 582 interconnected bundles of 2–4 hyphae, rather narrow (5–8 µm wide), pale, hyaline or a few
- 583 pale zebra-striped encrusted pigment.
- 584 *Veil at pileus surface* thin, of narrow, hyaline hyphae, or collapsed with some golden yellow585 necropigment.
- 586 Habitat: Montane forest with Fagus sylvatica and some Picea abies (ca. 1300 m asl.).
- 587 Distribution: Known only from the type locality in Val d'Ega, S Tyrol, N Italy.
- 588 Comments: Cortinarius pallidoriederi is characterized by the rather pale ochraceous yellow
- 589 pileus with an olivaceous green tinge, marbled-micaceous and innately fibrillose surface,
- 590 deeply violaceous colours of lamellae and stipe (apex), glossy stipe with slightly marginate
- 591 bulb, large spores and with only few, pale encrusted hyphae in pileipellis. Phylogenetically it
- is close to *C. riederi* s. str. (distinguished by 3% dissimilarity, i.e. 16 substitution and indel
- 593 differences), but differs in its paler olive tinged pileus with less encrusted hyphae and a
- slightly marginate bulb. The spores are also slightly smaller and narrower than typical *C*. This is a post-peer-review, pre-copyedit version of an article published in Mycological Progress. The final authenticated version is available online at: http://dx.doi.org/10.1007/s11557-018-1443-0

595 *riederi* spores. The species might also be mixed with the pale *C. anomaloochrascens*, but the

- latter has smaller spores with MVs always $<12.0 \mu m$. With its pale, somewhat micaceous,
- 597 olive ochre pileus, the species may look also like *C. herpeticus* (but this is never deep violet,

598 and has much smaller spores).

599 The species is known only from the type locality, but yet its specific rank seems well-

600 documented, since it is (i) quite dissimilar from closest relatives phylogenetically (two

- 601 specimens of the type collection were sequenced in the ITS region, both were identical and
- 602 differing in 16 changes from C. riederi), and in addition (ii) possesses differential macro-
- 603 scopical characters (pale olivaceous ochre pileus) and microscopically (little encrusted
- 604 pileipellis hyphae). Thus it is described here as a new species. *Cortinarius pallidoriederi* was

found in a mixed *Fagus-Picea* forest, apparently in a forest opening or rim with some grass-

606 herb vegetation (not in deep *Fagus* litter). It remains an open question if this is a *Fagus* or

607 *Picea* associate (or with both), but according to habitat preferences of the related species C.

riederi and *C. anomaloochrascens*, it seems probable that also this could be a major *Picea* associate.

610

611 *Collections examined*: Italy. S Tyrol, Val d'Ega (Eggental), Novale (Rauth), Wald Lagarn,

between Underkor and Oberkor, leg. E. Castellini, F. Bellù, 30 Sept 2011, Bellu 30-09-2011

- 613 (BOZ, holotype; O, isotype).
- 614
- 615

616 Cortinarius fulvoochrascens Rob. Henry, Bull. trimest. Soc. myc. Fr. 59: 55 (1943).

- 617 MycoBank: MB 539413
- 618 Figs 8c and 8d
- 619

620 Holotype: France, Doubs, R. Henry 314 (PC). GenBank: MH846266.

621

622 Synonyms: Cortinarius notabilis Rob. Henry, Bull. trimest. Soc. mycol. Fr. 73(1): 53 (1957).

- 623 MycoBank: MB 295899. Holotype: France, Doubs, R. Henry 3258-a-16 (PC).
- 624 Cortinarius marginatoochrascens Rob. Henry, Doc. Mycol. 19(73): 67 (1988). MycoBank:
- 625 MB 134954. Holotype: France, Doubs, R. Henry 643 (PC0085905).
- 626 Cortinarius fallaceicolor Rob. Henry, Doc. Mycol. MycoBank: MB 111326. Holotype:

 France, Doubs, under *Picea abies*, R. Henry 80.800 (PC).
 This is a post-peer-review, pre-copyedit version of an article published in Mycological Progress. The final authenticated version is available online at: http://dx.doi.org/10.1007/s11557-018-1443-0 Brandrud et al. 2018: Taxonomy and phylogeny of Cortinarius sect. Riederi

628 Cortinarius fulvoochrascens var. cyanophyllus Rob. Henry, Doc. Mycol. 19(73): 67 (1988).

629 MycoBank: MB 134961. Type: France, Doubs, R. Henry 3258-a-29 (PC, as héterotype).

630 Cortinarius fulvoochrascens var. subcaninicolor Rob. Henry, Bull. trimest. Soc. mycol. Fr.

631 105(2): 115 (1989). MycoBank: MB 581142. Holotype: France, Doubs, under *Picea abies*, R.

632 Henry 82.77 (PC).

633 *?Cortinarius fuscomaculatus* Jul. Schäff., Ber. Bayer. Bot. Ges. 27: 211 (1947). MycoBank:

634 MB 329045.

635 Phlegmacium fuscomaculatum var. umbrinum M.M. Moser, Die Gatt. Phlegm.: 355 (1961).

636 MycoBank: MB 353720. Holotype: Austria, Tyrol, Gschnitztal, under Picea abies, Larix

637 decidua, M. Moser, MM 49/137 (M). – Cortinarius fulvoochrascens var. umbrinus (M.M.

638 Moser) Quadr., Doc. Mycol. 14(56): 29 (1984). MycoBank: MB 116631. – Cortinarius

639 fuscomaculatus f. umbrinus Nespiak, Flora Polska, Grzyby (Mycota), 7, Basidiomycetes,

640 Agaricales, Cortinariaceae, Cortinarius 1: 147 (1975). MycoBank: MB 352736.

641

642

643 *Pileus* 5–10(12) cm, (hemi-)spherical, then plano-convex, viscid-glutinous, towards margin

644 more or less distinctly innately fibrillose-rivulose, somewhat micaceous, but not glossy, often

645 slightly fibrillose and towards centre faintly tomentose due to diffuse veil remnants, when

646 young in dry weather sometimes distinctly fibrillose-tomentose(-squamulose) like a *C*.

647 *balteatus*; usually rather homogeneous ochraceous brown, fulvous brown to more umber

brown, when very young sometimes with a faint greenish ochre brown tinge towards margin;

649 young margin often white fibrillose due to abundant veil remnants, sometimes also whitish

650 from veil remnants at centre.

651 *Lamellae* crowded (L = 80-110), 0.5–1 cm broad, initially deep violet to greyish violet, but

soon fading to paler bluish grey and then greyish brown; edge even to crenulate.

653 *Stipe* $5-10 \times 1-2.5$ cm, with a more or less distinct marginate bulbous base (up to 5 cm),

654 sometimes very robust and *Boletus*-like; initially deep (greyish) violet in upper part and with a

pale bluish tinge downwards, soon fading to greyish white-pale ochraceous brown, often

turning somewhat brass brown, sometimes with diffuse patches of watery bluish white

tomentum towards base. Universal veil near the bulb margin when young distinct, often

658 forming a slight bluish white, then ochraceous yellow, partly gelatinous girdle.

- 659 *Context* initially deep violet to greyish violet in (upper part) of stipe, bluish white in pileus
- and bulb, violet colour soon fading, becoming whitish with brass brown spots where
- damaged.
- 662 *Macrochemical reactions* negative.
- 663 Smell faint, indistinct.
- 664 *Exsiccata* pale brownish with vivid brown-redbrown pileus.
- 665 *Basidiospores* [7, 11, 206], 11.1–13.8 × 7.2–8.6 μm (MV = 12.5 × 7.9 μm); variation of MVs:
- 666 $12.1-12.7 \times 7.6-8.2 \,\mu\text{m}; Q = 1.43-1.75 \,(\text{MV} = 1.59)$, ellipsoid-amygdaloid, rather strongly
- and densely verrucose, suprahilar plague hardly visible. *Lamella edge* more or less fertile,
- sterile cells small, clavate or cylindrical.
- 669 *Pileipellis* duplex. Epicutis thin. At surface 3–5 μm wide, loosely erect-entangled, gelatinous
- 670 hyphae, hyaline or with pale, refractive pigment when collapsed, terminal hyphae sometimes
- 671 clavate. Subcutis/hypodermium, hyphae 5–20 μm wide, tightly cemented, with brown
- 672 parietal-encrusting pigment, often forming an amber-like embedment; in transition to epicutis
- 673 with narrow (5–8 μm wide), long hyphae with strong, brown, parietal to zebra-striped
- 674 encrusted pigment, tightly cemented to and sometimes intermixed in hypodermium, or
- 675 forming more loose, subparallel, interconnected bundles.
- 676 *Veil at pileus surface* on young specimen often prominent, at least 2–3 layers, hyphae rather
- 677 thick (4–8(12) μm wide), some interconnected (H-shaped connections), some terminal
- 678 elements slightly inflated; many with yellow parietal pigment, some with zebra-striped
- 679 encrustrations, some filled with brown, oleiferous necropigment.
- 680 Veil at stipe surface (ochre parts) with many pale golden yellow, refractive, collapsed,
- 681 gelatinous hyphae (a few dark brown oleifers), wider hyphae (5–7(10) μm) faintly zebra-
- 682 striped-verrucose encrusted.
- 683 Habitat: Associated mainly with Picea abies, but probably also with Pinus sylvestris. In rich
- low-herb spruce-pine forests, according to Moser (1961) also on rather acid ground. In S
- 685 Scandinavia mainly in calcareous spruce forests.
- 686 *Distribution:* Mainly distributed in C Europe, especially in the Alp/Prealp region. Rare in
- 687 Fennoscandia, but exact distribution unknown due to confusion with the similar C. riederi.
- 588 Just a few collections known so far from SE and C Norway and S/C Sweden. Widespread, but
- apparently rare in montane-subalpine districts of C (-S) Europe. Reported from subalpine sites
- 690 in Tyrol, Austria (Moser 1961 as Phlegmacium fuscomaculatum incl. var. umbrinum), and
- 691 from montane areas of the Jura region of France (Henry 1943; 1966; 1988; Bidaud et al. This is a post-peer-review, pre-copyedit version of an article published in Mycological Progress. The final authenticated version is available online at: http://dx.doi.org/<u>10.1007/s11557-018-1443-0</u>

- 1999), various parts of Germany (Schäffer 1947; Moser 1961; Krieglsteiner 1991), and from
- Spain (Ballarà 1999). The species is also collected in N Italy and Romania (Transylvania) (see
 collection list). So far not confirmed from N America.
- 695 *Comments: Cortinarius fulvoochrascens* is characterized by uniformly fulvous(-umber)
- brown, somewhat fibrillose pileus when young, more or less marginate bulbous stipe with
- 697 initially distinct veil remnants and large, ellipsoid-subamygdaloid spores. The often robust
- 698 stature with veil remnants at bulb and pileus margin is well illustrated by Bidaud et al. (1999,
- 699 Pl. 217 as *C. fulvoochrascens* var. *marginato-ochrascens*). This has frequently been treated as
- part of the collective species C. riederi s. lato (see comments under C. riederi). However, this
- is well separated phylogenetically from *C. riederi* s. str., and on closer examination, these are
- normally also well-distinguished morphologically. Cortinarius riederi differs by its smooth,
- glossy pileus, with more vividly yellow-orange tinges, almost lack of veil remnants, watery
- bluish mycelial felt and a hardly marginated stipe bulb. On average, the spores are slightly
- smaller and more strongly verrucose on *C. fulvoochrascens*, but the overlap in spore features
- 706 is considerable (see comment on *C. riederi*).
- Ecologically, *C. fulvoochrascens* is characterized by occurrence mainly in C European, richer
- to calcareous spruce-pine forests, whereas C. riederi s. str. is mainly N European, in C Europe
- 709 more subalpine, and it mainly occurs in oligotrophic/acid spruce forests. *Cortinarius*
- 710 *fulvoochrascens* also resembles *C. anomaloochrascens*. The former is, however, normally
- 711 distinguished by its darker, more fibrillose pileus colours, and larger, more robust
- 512 basidiocarps with pronounced veil remnants, distinctly larger spores, and stronger encrusting
- 713 pigment of pileipellis. The species is further distinguished from *C. argenteolilacinus* and *C.*
- 714 *malachioides* by darker coloured pileus, more distinct veil remnants as well as habitat in
- 715 coniferous forests. *Cortinarius fuscomaculatus* (Schäffer 1947) is very likely synonymous
- vith our species, but no original material of Schäffer exists. However, the type of
- 717 *Phlegmacium fuscomaculatum* var. *umbrinum* described by Moser (1961) was sequenced and
- turned out to be conspecific with *C. fulvoochrascens*. The species is treated under six different
- names by R. Henry (see synonymy list), according to our sequence study of R. Henry types.
- Among the Henryian names cited as synonyms is C. fulvoochrascens var. cyanophyllus,
- which has not infrequently been regarded as a variety or a species in literature (see e.g.
- 722 Ballarà 1999). Moser et al. (1995) reported C. fulvoochrascens from W North America based
- on many collections, but the spore sizes indicated, fits better to C. riederi than to C.
- 724 *fulvoochrascens*. Future sequencing of these collections might give more information on their This is a post-peer-review, pre-copyedit version of an article published in Mycological Progress. The final authenticated version is available online at: http://dx.doi.org/<u>10.1007/s11557-018-1443-0</u>

- identity. However, only one N American collection of *C. fulvoochrascens* identified by M.
- 726 Moser was sequenced (GenBank: AF389139), but it belongs to *C. burlinghamiae*.
- 727
- 728 Collections examined:
- Austria. Tyrol, Gschnitztal, MM 49/172 (M, holotype of *Phlegmacium fuscomaculatum* var.
- *umbrinum*). **France**. Doubs, R. Henry 314 (PC, holotype); R. Henry 71.732 (PC, héterotype
- of C. fulvoochrascens); R. Henry 3258-a-16 (PC, holotype of C. notabilis); R. Henry 643 (PC,
- holotype of *C. marginatoochrascens*); R. Henry 80.800 (PC, holotype of *C. fallaceicolor*); R.
- Henry 3258-a-29 (PC, héterotype of *C. fulvoochrascens* var. *cyanophyllus*); R. Henry 82/77
- 734 (PC, holotype of *C. fulvoochrascens* var. *subcaninicolor*). Germany. Baden-Württemberg,
- 735 Schwaben, Ehingen a.d. Donau, T.E. Brandrud, TEB437-10 (O). Italy. Trentino-Alto-Adige,
- 736 Alto-Badia, La Valle Badia, 7 Sept 2008, C. Rossi, Rossi 31 (herb. Rossi); Naz-Sciaves,
- 737 Springa, 14 Oct 2000, G. Turrini, TG2000-198 (herb. Turrini). Norway. Oppland, Jevnaker,
- Rustad, T.E. Brandrud, TEB667-80 (O); Lunner, S. Oppdalen, T.E. Brandrud, TEB172-04
- 739 (O); Gran, Lygna N, 11 Sept 2013, T.E. Brandrud & B. Dima, TEB530-13/DB5210 (O/BP);
- 740 Vestre Toten, Eriksrudtjern, T.E. Brandrud & B. Dima, TEB725-17/DB6542 (O). Buskerud,
- 741 Hole, Nes, T.E. Brandrud, TEB748-11 (O). Nord-Trøndelag, Levanger, Ytterøya, TEB292-13
- 742 (O). Oslo, Oslo, Østmarksetra, 13 Sept 2014, T. Berg-Hansen, F-75726 (O). Romania.
- Partium, Bihar, Pádis, 10 Oct 2010, Z. Erdei, DB4408 (BP). Spain. Catalunya, J. Ballarà, JB
- 744 6858-09. Sweden. Västergötland, Kinnekulle, Hällekis, Gössäter, T.S. Jeppesen, TSJ2004-052
- 745 (C).
- 746
- 747 Additional ITS sequence data from the public sequence repositories:
- 748 **Germany**. KJ421204/TUB020433 (as *C. riederi*), KJ420976/TUB019735 (as *C. riederi*).
- 749
- 750
- 751 Cortinarius anomaloochrascens Chevassut & Rob. Henry, Doc. Mycol. 16(63–64): 84
- 752 (1986).
- 753 MycoBank: MB 129655
- 754 Figs 8e, 8f and 8g
- 755
- 756 Holotype: France, Doubs, R. Henry 2805 (PC). GenBank: MH846269.
- 757

- 758 Synonyms: Cortinarius imbricatoides Rob. Henry, Bull. trimest. Soc. mycol. Fr. 105(1): 92
- 759 (1989). MycoBank: MB 136657. Holotype: France, Doubs, R. Henry 395 (PC).
- 760 Cortinarius stilazureus Rob. Henry, Bull. trimest. Soc. mycol. Fr. 105(2): 125 (1989).
- 761 MycoBank: MB 136688. Holotype: France, Doubs, Frasne, R. Henry 85.13 (PC).
- 762

763 Selected icones: Soop (2017, Pl. 5: 19)

764

765 *Pileus* 4–8 cm, (hemi-)spherical(-campanulate), then plano-convex (to broadly umbonate),

viscid-glutinous and glossy when dry, towards margin distinctly innately fibrillose-rivulose,

centre with a marbled structure from darker, innate fibers, sometimes with hygrophaneous

768 veins or spots towards margin; pale ochraceous brown to greybrown, sometimes with an

769 olivaceous tinge when young, the involute margin whitish. Universal veil remnants very

- 770 sparse, hardly visible.
- 771 *Lamellae* crowded (L = 70-100), 0.4–0.8 cm broad, initially deep violet to greyish violet, but
- soon fading to paler bluish grey and then greyish brown; edge even to crenulate.
- 573 Stipe $5-10 \times 0.8-1.5$ cm, with a clavate-bulbous to slightly marginate bulbous base (up to 3.5)
- cm), glossy, initially deep (greyish) violet in upper part and with a pale bluish tinge

downwards, soon fading to greyish white-pale ochraceous brown, often developing glossy

- brass brown patches, sometimes with patches with watery bluish white tomentum towards
- base. Universal veil near the bulb margin very sparse.
- 778 *Context* initially deep violet to greyish violet in (upper part) of stipe, bluish white in pileus
- and bulb, violet colour soon fading, becoming whitish with brass brown spots wheredamaged.
- 781 *Macrochemical reactions* negative, KOH somewhat brownish in context.
- 782 *Smell* none or faint, sometimes yeast-like with age.
- 783 *Exsiccata* pale brownish.
- *Basidiospores* [7, 11, 168], 10.2–12.8 × 6.3–8.1 μm (MV = 11.5 × 7.2 μm); variation of MVs:
- 785 $10.9-11.9 \times 6.9-7.6 \,\mu\text{m}; Q = 1.43-1.78 \,(\text{MV} = 1.60)$, ellipsoid to subamygdaloid, distinctly
- and densely verrucose, suprahilar plague hardly visible.
- 787 *Pileipellis* duplex. Epicutis thin. At surface of 3–5 µm wide, loosely erect-entangled,
- gelatinous hyphae, hyaline or with pale pigment, some pale verrucose to zebra-striped.
- 789 Subcutis/hypodermium hyphae 5–20 μm wide, tightly cemented, with brown parietal-

- response to the forming an amber-like embedment; in transition to epicutis with
- subparallel, interconnected bundles, with some, encrusted pigment.
- 792 Habitat: Associated with Picea abies, also in Picea plantations outside the natural distribution
- of Norway spruce; in a few cases also seen in pure *Pinus sylvestris* forests in W Norway,
- 794 possibly also associated with Abies alba in mixed stands. Preferentially in oligotrophic-
- mesotrophic spruce forests, rarely on calcareous ground (Krieglsteiner and Gminder 2010;
- 796 Jeppesen et al. 2012).
- 797 *Distribution*: Widely distributed in boreal-montane districts in Europe, but apparently
- row everywhere rather rare. Reported from C and S Fennoscandia (Soop 2011 as C.
- 799 *fulvoochrascens* p.p.; Jeppesen et al. 2012), from The Black Forest region of Germany
- 800 (Krieglsteiner and Gminder 2010) and from the Jura region of France (Henry 1989a; 1989b;
- 801 Bidaud et al. 1999). In The Black Forest, the species occurs in the montane zone (450)600–
- 802 1000 m asl., and in the lower montane mixed Picea-Abies forests it seems to be more frequent
- 803 than C. riederi (Krieglsteiner and Gminder 2010; pers. obs.). The species is further collected
- 804 by us in N Italy and the Czech Republic (see list of collections examined). Moser (1961) may
- 805 have included *C. anomaloochrascens* in his wide concept of *Phlegmacium fuscomaculatum*
- 806 which is reported not to be rare in the Alp-Prealp region.
- 807 Comments: Cortinarius anomaloochrascens, although widespread, is very little known and
- 808 described, apparently mainly included in a broad concept of *C. riederi*. The species seems
- 809 well illustrated in Bidaud et al. (1999; Pl. 213) and Soop (2017), although these specimens are
- 810 not verified by sequencing. *Cortinarius anomaloochrascens* is characterized by its rather
- 811 small and slender basidiocarps, pale ochraceous greybrown pileus, and smaller spores
- 812 compared to *C. riederi* and *C. fulvoochrascens*. The spores are usually also somewhat more
- 813 amygdaloid than those of *C. riederi* and *C. fulvoochrascens*. The contrast between the initially
- 814 bright violaceous stipe and the pale greyish ochre pileus margin is often striking. The stipe
- 815 furthermore normally has a more marginate bulb than that of *C. riederi*. Misidentifications
- 816 under the name *C. riederi* or *C. fulvoochrascens* can easily be sorted out by the smaller spores
- 817 of *C. anomaloochrascens*, which show no overlap with the spores of *C. riederi/C*.
- 818 *fulvoochrascens*, when mean values for measurement series are compared (spore MVs for *C*.
- 819 anomaloochrascens always <12 µm long). Cortinarius glaucocyanopus and C. malachioides
- 820 also have comparatively small spores, but these are associated with deciduous trees, mainly
- 821 Fagus and Betula, respectively. These also have a paler coloured pileus, and C. malachioides
- usually lacks deep violet tinges on young lamellae and stipe.
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- 823
- 824 *Collections examined*:
- 825 Czech Republic. C Bohemia, Prague region, Borovsko, Hadce u Želivky Nature Reserve,
- 826 mixed forest with Picea, Pinus, Betula on serpentinite soil, J. Borovicka 140519-29 (PRM
- 827 924382). Denmark. Northern Jutland, Faurholt hede, A. Eriksen, AE2009-65377 (C77672).
- 828 Finland. Kainuu, Paltamo, I. Kytövuori, IK92-2028 (H). France. Doubs, R. Henry 2805 (PC,
- holotype), R. Henry 2807 (PC, isotype); R. Henry 395 (PC, holotype for *C. imbricatoides*); R.
- 830 Henry 85.13 (PC, holotype for C. stilazureus). Doubs, Levier, T.G. Frøslev, T.S. Jeppesen,
- 831 TSJ2006-095 (C). Auvergne-Rhone-Alpes, Oyonnax, I. Kytövuori, IK94-1743 (H).
- 832 Germany. Baden-Württemberg, Rottweil, Dunninger Wald, T.E. Brandrud, B. Dima, G. Saar,
- 833 TEB762-12/DB4814 (O). Bayern, Sachsenkamm, Kirchsee Nord, 7 Oct 2013, M. Dondl,
- MD28/2013; Bayern, Mittenwald, Kranzbergregion, 21 Sept 2013, M. Dondl, MD 27/2013.
- 835 Italy. Trentino-Alto-Adige, St. Valentin Atzwang, 30 July 2012, F. Bellù, Cester, Bellu 30-07-
- 836 2012 (BOZ); Riscone, 14 Aug 2012, C. Rossi, Rossi 14-08-2012 (herb. Rossi); Redensberg,
- Nähe Olang, 13 Aug 2012, F. Bellù & Fuman, Bellu 13-08-2012 (BOZ). S Tyrol, Jenesien,
- 838 Remphof, 4 Sept 2017, T. Errico, Bellu 04-09-2017 (O); Bolzano, Villnösstal, Malga Zannes,
- Val di Funes, C. Rossi, Rossi 31-08-2017 (O); Mölten, Kirchebene, pr. Mettina, C. Rossi,
- Rossi 07-09-2017 (O). Norway. Oppland, Lunner, S. Oppdalen, E. Bendiksen, TEB316-10
- 841 (O). Østfold, Halden, Idd, T.E. Brandrud, TEB252-07 (O). Hordaland, Ulvik, Åsen peninsula,
- 842 T.E. Brandrud, TEB175-07 (O). Nord-Trøndelag, Nærøy, Saglivatn, E. Bendiksen & T.E.
- 843 Brandrud, EB-TEB 6-05 (O). Oslo, Oslo, Enebakkveien 866, 12 Sept 2014, T. Blakar, F-
- 844 75650 (O). Sweden. Ångermanland, Säbrå, H. Marklund, CFP 1539 (S).
- 845
- 846 Additional ITS sequence data from the public sequence repositories:
- 847 Germany. KJ420997/TUB019758 (as C. riederi); KJ421008/TUB019769 (as C. riederi);
- 848 KJ421012/TUB019770 (as *C. riederi*). Italy. JF907910/MCVE16165 (as *C. riederi*).
- 849
- 850
- 851 Cortinarius glaucocyanopus Rob. Henry, in Bidaud et al., Atlas des Cortinaires IX: 371
- 852 (1999)
- 853 MycoBank: MB 461065
- Figs 9h and 9i
- 855

Brandrud et al. 2018: Taxonomy and phylogeny of Cortinarius sect. Riederi

856	Holotype: France, Ardennes, Bois de Toges, 6 Oct 1995, P. Reumaux, GK5034 (G). GenBank:
857	MH846274.

- 858 Synonym: Cortinarius glaucocyanopus Rob. Henry, Bull. trimest. Soc. myc. Fr: 73: 32
- 859 (1957), nom. inval.
- 860
- 861

862

863 *Pileus* 2.5–7 cm, (hemi-)spherical, then plano-convex, viscid-glutinous, innately fibrillose-

864 rivulose; pale ochraceous (grey)brown to cream yellow, with an olivaceous tinge (resembling

- the colour of *C. anserinus*), sometimes with lilac tinged spots; the involute margin whitish.
- 866 Universal veil remnants very sparse, hardly visible.
- *Lamellae* crowded, initially deep violet to pale greyish violet or bluish lilac, soon fading togreyish brown; edge even.
- 869 Stipe $5-8 \times 0.7-1.5$ cm, with a slightly marginate bulbous base (up to 3 cm), glossy, initially
- pale (greyish) violet-bluish lilac, developing brass brown patches. Universal veil near the bulbmargin very sparse.
- 872 *Context* initially pale bluish violet in (upper part) of stipe, white in bulb, bluish white in
- pileus, violet colour soon fading, becoming whitish with brass brown spots where damaged.
- 874 *Macrochemical reactions* negative.
- 875 *Smell* indistinct.
- 876 *Exsiccata* pale brownish.
- 877 *Basidiospores* [2, 2, 68], $10.3-13.1 \times 6.6-7.8 \,\mu m$ (MV = $11.7 \times 7.2 \,\mu m$), variation of MVs:
- 878 $11.2-12.0 \times 7.2-7.3 \ \mu\text{m}; \ Q = 1.41-1.85 \ (MV = 1.63), \ \text{ellipsoid to subamygdaloid, distinctly}$
- and densely verrucose, suprahilar plague hardly visible.
- 880 *Pileipellis* duplex. Epicutis thin. At surface of 3–5 µm wide, loosely erect-entangled
- gelatinous hyphae, hyaline or with pale pigment. Subcutis/hypodermium hyphae 5–20 µm
- wide, tightly cemented, with brown parietal-encrusting pigment, often forming an amber-like
- 883 embedment; in transition to epicutis with subparallel, interconnected bundles, some hyphae
- 884 with weakly encrusted pigment.
- 885 Habitat: Apparently associated with Fagus, possibly also other deciduous trees; collected in
- 886 *Fagus* dominated forest with small *Picea* trees and *Betula* in Hungary, and reported from
- 887 Fagus forests in the Ardennes and French Jura (Bidaud et al. 1999).

- *Distribution:* So far known only from temperate/nemoral regions of French Ardennes-Jura
 and western Hungary. Apparently a very rare species.
- 890 *Comments:* The description of macromorphology is based on the protologue and on own
- 891 materials from Hungary. *Cortinarius glaucocyanopus* is a very rare species, so far known only
- from one site (two collections) in western Hungary, and from the type collection from the
- 893 French Ardennes, possibly also from one (not sequenced) collection from French Jura (see
- Bidaud et al. 1999). The species is genetically well-distinguished, separated from its sister
- species *C. anomaloochrascens* by 10 substitution and indel positions in the ITS regions. They
- also have different habitats; C. glaucocyanopus growing in Fagus forests, C.
- 897 anomaloochrascens in Picea forests. Morphologically, the former seems to differ by a paler,
- 898 more olive-ochre pileus, according to the protologue resembling the colour of *C. anserinus*
- 899 (=*C. amoenolens*) (Henry in Bidaud et al. 1999). However, more material is needed to
- 900 confirm the morphological-ecological characteristics of this species. *Cortinarius*
- 901 glaucocyanopus was introduced by Henry (1957), but this description is invalid since lacking
- 902 a Latin diagnosis. Moreover, the short description seems to cover one of the large-spored
- 903 *Riederi* taxa rather than the relatively small-spored *C. anomaloochrascens/C. glaucocyanopus*
- 904 ("spores.... de $15/8.8 \mu$."). In fact, the scetched specimen and the description of the pileus
- 905 ("ochracée....mate, devenant café au lait avec des plages d'un glauque bleuâtre sur le disque
- 906 plus ou moins pseudo-squamuleuse à la fin.") seems to cover *C. argenteolilacinus* better than
- 907 our species. However, the species was validated and re-described by Henry in Bidaud et al.
- 908 (1999). The description is based on two collections; the type from the Ardennes, and another
- 909 collection from the Jura (Bidaud, 27 Sept 1998). The type has been sequenced, and the type
- 910 and our Hungarian collections have identical sequences. The description (protologue) and
- 911 depicted specimens in Bidaud et al. (1999) correspond rather well with our find from
- 912 Hungary, but again the spores are given (very) large; $(11)12-15(17) \times 7.5-8.5(9) \mu m$, of the
- same size as those of *C. fulvoochrascens*, and considerably larger than our measurements.
- 914
- 915 *Collections examined*:
- 916 France. Ardennes, Bois de Toges, 06 Oct 1995, P. Reumaux, GK5034 (G, holotype).
- 917 Hungary. Vas, Farkasfa (lake Fekete-tó), 3 Oct 1999, L. Albert, AL 99/51 (BP), 1 Oct 2005,
- 918 B. Dima, DB2072 (BP, O).
- 919
- 920

921	Cortinarius argenteolilacinus var. argenteolilacinus M.M. Moser, Sydowia 6(1-4): 151
922	(1952).
923	MycoBank: MB 295797
924	Figs 9a, 9b and 9c
925	
926	Holotype: Austria, Holltal, 16 Oct 1948, M. Moser, MM 48/752 (M). GenBank: MH846277.
927	
928	Synonym: Phlegmacium argenteolilacinum (M.M. Moser) M.M. Moser, Die Gatt. Phlegm.
929	241 (1961). MycoBank: MB 302878.
930	
931	Selected icones: Moser (1961, Pl. XVII 96), Læssøe et al. (2011), Svampe 64: 57, Soop
932	(2011, Pl. 11, fig. 43)
933	
934	Pileus 4-10 cm, (hemi-)spherical, then plano-convex (to broadly umbonate), viscid-glutinous,
935	towards margin weakly, often finely (sometimes distinctly) innately fibrillose-rivulose,
936	somewhat silvery-micaceous, but hardly glossy-polished (like C. riederi), often faintly
937	fibrillose-tomentose; pale grey, greyish ochre to greyish white, often slightly more ochraceous
938	brown at centre, initially often with a bluish tinge towards margin, the involute margin
939	(bluish) whitish. Universal veil remnants sparse, sometimes with whitish veil fibers towards
940	margin when young.
941	Lamellae crowded (L = $80-110$), 0.5–1 cm broad, initially deep violet to greyish violet, but
942	(very) soon fading to paler bluish grey and then greyish brown, edge often paler crenulate-
943	serrulate.
944	Stipe 6–10(11) \times 1–2.5 cm, often robust, with a bulbous to often slightly marginate-bulbous
945	base (up to 4.5 cm), glossy to slightly fibrillose, initially greyish violet to deep violet in upper
946	part, (bluish) white downwards (sometimes distinctly violet down to bulb), bluish tinges fades
947	soon, but more slowly than that of the lamellae, and in some stages violet blue stipe apex
948	contrasts the already greyish brown lamellae; sometimes turning (spot wise) brass-brown with
949	age. Universal veil sparse to rather abundant, and then forming a girdle at or just above the
950	bulb margin; white to sometimes bluish when very young.
951	Context initially deep violet to greyish violet in upper part of stipe (especially towards
952	cortex), bluish white in pileus and bulb, violet colour soon fading, becoming whitish with
953	brass brown spots where damaged.
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- 954 *Macrochemical reactions* negative, sometimes yellow-brownish with KOH in bulb context.
- 955 *Smell* faint, indistinct.
- 956 *Exsiccata* brownish to almost whitish.
- 957 *Basidiospores* [8, 16, 319], 10.7–13.7 × 6.7–8.3 μm (MV = 12.2 × 7.5 μm), variation of MVs:
- 958 $11.6-12.9 \times 7.1-7.9 \,\mu\text{m}; Q = 1.42-1.84 \,(\text{MV} = 1.63), \text{ ellipsoid to subamygdaloid, rather}$
- strongly to very strongly and rather densely verrucose, perisporium sometimes visible,
- 960 ornaments often lense-like coalescent at apex, sometimes prominent, giving the spore a
- 961 slightly citriform outline, often also with prominent, tooth-like warts subapically; suprahilar
- 962 plague rarely visible.
- 963 *Lamellar edge* more or less fertile, sterile cells small, clavate or cylindrical.
- 964 *Pileipellis* duplex. Epicutis thin. At surface 3–5 µm wide, loosely erect-entangled, gelatinous
- 965 hyphae, hyaline or with pale, refractive pigment (especially when collapsed).
- 966 Subcutis/hypodermium weakly developed, hyphae 5–15 μm wide, tightly cemented, with pale
- 967 yellow brown parietal-encrusting pigment, often forming an amber-like embedment; in
- 968 transition to epicutis with subparallel, interconnected bundles of long and rather narrow
- 969 hyphae (5–10 μm wide), with distinct to strong zebra-striped/crustulose brown encrusting
- 970 pigment, encrustration especially prominent on some slightly wider hyphae (8–10 μm wide).
- 971 *Veil hyphae at pileus surface* scattered, mainly narrow (3–6 μm), hyaline; some hyphae
- 972 interconnected.
- 973 Habitat: Associated mainly with Fagus sylvatica, in boreonemoral region also sometimes
- 974 found under *Tilia cordata* and *Corylus avellana*, in richer to somewhat calcareous forests.
- 975 Distribution: Distributed mainly in nemoral-(sub)montane areas of C Europe and S
- 976 Scandinavia, more rare in boreonemoral areas of S Scandinavia. Apparently rare everywhere,
- 977 very rare in S Europe. Little known, recorded only from SE Norway, S Sweden and Denmark
- 978 (cf. Læssøe et al. 2011; pers. obs.), Tyrol, Austria (Moser 1961), Germany (Krieglsteiner
- 979 1991; pers. obs.) and the French Jura (see collections examined). According to Mahiques
- 980 (1999) it is found also in NE Spain (Catalonia).
- 981 Comments: Cortinarius argenteolilacinus s. lato is characterized by its pale greyish, almost
- 982 fibrillose pileus with bluish tinges when (very) young, and the initially deep violet lamellae
- 983 and stipe. The basidiocarps are often robust. The spores often possess an apical, irregular,
- 984 lense-like structure due to coalescent warts. *Cortinarius argenteolilacinus* var.
- 985 argenteolilacinus and var. dovrensis deviate genetically by only two to three nucleotide and
- 986 indel differences in the ITS region.

987 Cortinarius argenteolilacinus var. dovrensis is a N European subalpine Betula forest taxon, 988 whereas var. argenteolilacinus is a C European-S Scandinavian (boreo-)nemoral Fagus-Tilia-989 Corylus forest species. Cortinarius argenteolilacinus var. argenteolilacinus has so far never 990 been found in subalpine birch forests, and the distribution of the two varieties shows almost 991 no overlap in Scandinavia. The latter differs morphologically by the more strongly 992 ornamented spores, and the strongly zebra-striped encrusted $8-10 \,\mu m$ wide subsurface 993 pileipellis hyphae. With this combination of (i) a small, constant genetic differentiation, (ii) a 994 considerable ecogeographical differentiation, and (iii) a subtle micromorphological 995 differentiation, we have decided to treat these as two infraspecific taxa (varieties). Due to 996 geographical differences, these taxa might also be treated as subspecies, but we think it is 997 wise to stick to only one infraspecific rank (subspecies is nowadays very little applied in 998 agaric taxonomy).

999 Cortinarius argenteolilacinus var. argenteolilacinus also resembles C. malachioides, but is 1000 distinguished by the larger spores and more zebra-striped encrusted pileipellis subsurface 1001 hyphae. It usually also has a larger, more fibrillose and more (bluish) grey pileus, and initially 1002 deeper violet colours of lamellae and stipe. It is further distinguished from C. riederi and C. 1003 anomaloochrascens by the less smooth, less innately fibrillose and a paler (and bluish tinged) 1004 pileus. The present species has traditionally been classified with the C. variecolor-C. largus 1005 group, and not with C. riederi, but the micromorphology as well as genetic data show a close 1006 relationship to the latter. *Cortinarius argenteolilacinus* might look like a *C. caesiocanescens*, 1007 C. coerulescentium or a C. largus, but has larger, more ellipsoid spores. Furthermore, the bulb 1008 is more marginated on C. caesiocanescens and C. coerulescentium, and the former has more 1009 abundant (volva-like) veil remnants, and the latter has a more distinctly bluish, strongly 1010 innately fibrillose pileus when young. Cortinarius largus is distinguished by the yellow KOH-1011 reaction (context), and never marginate bulbous stipe. Cortinarius argenteolilacinus var. 1012 argenteolilacinus is so far known from a few nemoral lowland (S Sweden, Denmark), 1013 submontane (French Jura, S Germany) to upper montane (Austria and NE Hungary) Fagus 1014 sites. Furthermore, it is found in a few boreonemoral Tilia(-Quercus) forests in SE Norway, 1015 and under *Corylus* at Öland, SE Sweden. The taxon is probably distributed further East in 1016 Europe (e.g., in the Carpathians).

1017

1018 *Collections examined*:

- 1019 Austria. Holltal, M. Moser, MM 48/752 (M, holotype). Denmark. Sjælland, Vejlø skov, T.S.
- 1020 Jeppesen, TSJ2010-004 (C), TSJ2012-044 (C). France. Jura, Prénovel, 10 Oct 2008, B.
- 1021 Dima, DB3312 (BP). Germany. Baden-Württemberg, Schwaben, Ehingen a.d. Donau, T.E.
- 1022 Brandrud, G. Schmidt-Stohn et al., TEB 418-10/SSt10-174 (O). Bayern, Tegernsee,
- 1023 Riederstein, M. Dondl, MD18/2013. Hungary. Heves, Mátra Mts, L. Nagy, NL-5262 (herb.
- 1024 L. Nagy). Norway. Telemark, Drangedal, Sannes-Langen, T.E. Brandrud, TEB52-01 (O).
- 1025 Aust-Agder, Risør, Dalsvann SW, Grønnfjell, I.L. Fonneland, ILF2013-98 (O); Vestfold,
- 1026 Larvik, Kjose, T.E. Brandrud, TEB370-08 (O); T.E. Brandrud, TEB381-08 (O). Buskerud,
- 1027 Hole, Bråtåfjellet SV, T.E. Brandrud & B. Dima, TEB281-17/DB6361 (O). Sweden.
- 1028 Bohuslän, Tanum, T.E. Brandrud et al., CFP1608 (S). Öland, Borg, I. Kytövuori, IK98-2617
- 1029 (H). Skåne, Ivön, C. Ingvert, T.G. Frøslev, T.S. Jeppesen TSJ2011-025 (C)
- 1030
- 1031 Cortinarius argenteolilacinus var. dovrensis Brandrud, var. nov.
- 1032 MycoBank MB 827876
- 1033 Fig. 9d
- 1034
- Etymology: the epithet refers to the Dovre Mountains of S Norway, from where the type wascollected.
- 1037
- Holotype: Norway, Sør-Trøndelag, Oppdal, Kongsvoll, Dovre Mountains, 10 Aug 1980, T.E.
 Brandrud, TEB112-80 (O). GenBank: MH923071.
- 1040
- 1041 Selected icones: Soop 2017, Pl. 5:20 as C. "dovrensis".
- 1042
- 1043 *Pileus* 4–10 cm, (hemi-)spherical, then plano-convex (to broadly umbonate), viscid-glutinous,
- 1044 towards margin weakly, finely (sometimes distinctly)innately fibrillose-rivulose, somewhat
- silvery-micaceous, often faintly fibrillose-tomentose; pale grey, greyish ochre to greyish
- 1046 white, often slightly more ochraceous brown at centre, sometimes almost whitish, initially
- 1047 often with a bluish tinge towards margin, the involute margin (bluish) whitish. Universal veil
- 1048 remnants very sparse, sometimes with whitish veil fibers towards margin when young.
- 1049 *Lamellae* crowded (L = 80-110), 0.5-1 cm broad, initially deep violet to greyish violet, but
- 1050 (very) soon fading to paler bluish grey and then greyish brown; edge even to crenulate.

- 1051 Stipe $6-10(11) \times 1-2.5$ cm, often robust, with a bulbous to often faintly marginate-bulbous
- 1052 base (up to 4.5 cm), glossy to slightly fibrillose, initially greyish violet to deep violet in upper
- 1053 part, whitish downwards (sometimes violet down to bulb), bluish tinges soon fading,
- 1054 sometimes turning (spot wise) somewhat brass-brown with age. Universal veil near the bulb
- sparse and hardly visible; white to sometimes bluish when very young.
- 1056 *Context* initially deep violet to greyish violet in upper part of stipe (especially towards
- 1057 cortex), bluish white in pileus and bulb, violet colour soon fading, becoming whitish with1058 brass brown spots where damaged.
- 1059 *Macrochemical reactions* negative, sometimes yellow-brownish with KOH in bulb context.
- 1060 Smell faint, indistinct.
- 1061 *Exsiccata* whitish to (pale) brownish.
- 1062 *Basidiospores* [9, 16, 264], $11.2-13.8 \times 6.8-8.6 \mu m$ (MV = $12.5 \times 7.7 \mu m$); variation of MVs:
- 1063 $11.8-13.5 \times 7.3-8.2 \ \mu m; Q = 1.45-1.79 \ (MV = 1.62), ellipsoid to subamygdaloid, medium to$
- 1064 rather strongly and densely verrucose, ornaments often lense-like coalescent at apex,
- 1065 sometimes prominent; suprahilar plague rarely visible.
- 1066 *Lamellar edge* more or less fertile, sterile cells small, clavate or cylindrical.
- 1067 *Pileipellis* duplex. Epicutis thin, at surface of 3–5 µm wide, loosely erect-entangled,
- 1068 gelatinous hyphae, hyaline or with pale, refractive pigment (when collapsed).
- 1069 Subcutis/hypodermium weakly to distinctly developed, hyphae 5–15 µm wide, basally
- 1070 sometimes with inflated elements up to 20 µm wide, tightly cemented, with pale yellow
- 1071 brown parietal-encrusting pigment, often forming an amber-like embedment; in transition to
- 1072 epicutis with subparallel, interconnected bundles of 5–8 µm wide hyphae, sometimes with
- 1073 weak zebra-striped encrusting pigment (never distinct encrustrations like in the main variety).
- 1074 Veil hyphae at pileus surface scattered to almost lacking, mainly narrow (3-6 µm) and
- 1075 hyaline.
- 1076 Habitat: Associated mainly with Betula pubescens subsp. tortuosa, but also with Betula
- 1077 *pendula* and *Tilia cordata*, more rarely with *Corylus avellana*.
- 1078 *Distribution:* So far mainly known from NW Europe. Most records from subalpine/northern
- 1079 boreal areas of Fennoscandia (known north to Nordland in Norway, Umeå in Sweden, and N
- 1080 Finland (Lapland)), but also found in some boreonemoral fjord districts of W Norway.
- 1081 Apparently no finds from nemoral-montane regions of C Europe. Once collected in Alberta,
- 1082 Canada, indicating a probably wide, circumpolar distribution.

1083 Comments: Cortinarius argenteolilacinus var. dovrensis is characterized by its pale greyish-

- 1084 whitish, almost fibrillose pileus with bluish tinges when (very) young, and the initially deep
- 1085 violet lamellae and stipe. The taxon is phylogenetically very closely related to C.
- 1086 argenteolilacinus var. argenteolilacinus, and differs only in two to three substitution and indel
- 1087 positions in the ITS region. With this more or less constant differences the sequences cluster
- 1088 in two rather well-supported clades in our phylogenetic analysis (Fig. 2). Since these
- 1089 genotypes are (i) considerably differentiated ecogeographically (var. dovrensis mainly a
- 1090 northern, subalpine Betula taxon, var. argenteolilacinus a mainly nemoral-montane Fagus(-
- 1091 *Tilia*) taxon) and (ii) slightly differentiated also morphologically (*C. argenteolilacinus* var.
- 1092 dovrensis with less ornamented spores, less encrusted pileipellis hyphae and on average paler
- 1093 pileus), these are treated as separate varieties (see also comments under *C. argenteolilacinus*
- 1094 var. argenteolilacinus).
- 1095 Most of our collections of *C. argenteolilacinus* var. *dovrensis* were found under *Betula*
- 1096 (mainly subalpine *B. pubescens* subsp. tortuosa), whereas three collections were found under
- 1097 *Corylus* (-Betula?)). No collections were from *Fagus* forests, which seems to be the major
- 1098 habitat for *C. argenteolilacinus* var. *argenteolilacinus*. All the 16 DNA-analysed *C*.
- 1099 argenteolilacinus var. dovrensis collections have almost identical ITS sequences, but in some
- 1100 cases we observed single nucleotide polymorphisms in two sites. The specimen found in
- 1101 Canada differs by 3 indels from the European sequences. More material is needed to see if
- 1102 there is a phylogeographical differentiation within *C. argenteolilacinus* var. *dovrensis*. For the
- 1103 time being we keep this Canadian collection under var. *dovrensis* based on our phylogenetic
- 1104 analysis (Fig. 2).
- 1105 *Cortinarius argenteolilacinus* var. *dovrensis* may sometimes be difficult to separate
- 1106 macromorphologically from the often co-occurring *C. malachioides*. However, the var.
- 1107 *dovrensis* is distinguished by the larger spores. It usually also has a larger and more fibrillose
- 1108 pileus and brighter violet colours on lamellae and stipe.
- 1109
- 1110 *Collections examined:*
- 1111 Canada. Alberta, Hinton, 3 Sept 2011, TN11-319 (H). Finland. Perä-Pohjanmaa, Yltornio, I.
- 1112 Kytövuori, IK 97-1137 (H). Norway. Oppland, Dovre, Kongsvoll, 10 Aug 1980, T.E.
- 1113 Brandrud, TEB112-80 (O, holotype); Lunner, Skøien, T.E. Brandrud, TEB682b-11 (O).
- 1114 Hedmark, Engerdal, Femunden, T.E. Brandrud, TEB217-08 (O). Aust-Agder, Froland, Ytre
- Lauvrak, T.E. Brandrud, TEB312-09 (O); Dalsvann SW, I.L. Fonneland, 2013-119 (O); This is a post-peer-review, pre-copyedit version of an article published in Mycological Progress. The final authenticated version is available online at: http://dx.doi.org/10.1007/s11557-018-1443-0

1116	Nordland, Hattfjelldal, Storveltlia nature reserve, Ø. Weholt, KB-EB-TEB 78-10 (O), Ø.
1117	Weholt, KB-EB-TEB 79-10 (O). Hordaland, Granvin, T.S. Jeppesen, TSJ2005-042 (C).
1118	Sweden. Härjedalen, Hamrafjället, 13 Aug 2008, K. Soop, KS-CO1703 (S-F251128); 20 Aug.
1119	2008, K. Soop, KS-CO1821 (S). Dalarna, Sollerön, Klikten, 11 Sept 2013, K. Soop, KS-
1120	CO2130 (S). Öland, Gråborg, T.G. Frøslev, T.S. Jeppesen TSJ2006-010 (C).
1121	
1122	
1123	Additional ITS sequence data from the public sequence repositories:
1124	Italy. JF907933/MCVE6043 (as C. elotus). Sweden. KJ421064/KS-CO770 (as C.
1125	argenteolilacinus), Öland, Gråborg, 28 Sept 1996, K. Soop, herb. S-F44342.
1126	
1127	
1128	Cortinarius malachioides P.D. Orton, Naturalist (Suppl.): 148 (1958)
1129	MycoBank MB 295890
1130	Figs 9e, 9f and 9g
1131	
1132	Holotype: United Kingdom, Inverness-shire, Rothiemurchus, Loch-an-Eilean, P.D. Orton
1133	29/9/55 (K(M)94426). GenBank: MH846280.
1134	Icone: Soop (2011, Pl. 6, fig. 23, as C. coniferarum (M.M.Moser) Moënne-L & Reum); Soop
1135	2017, Pl. 5:18 as <i>C. jotunae</i> .
1136	
1137	
1138	Pileus 3–7(8) cm, (hemi-)spherical, then plano-convex to broadly umbonate, viscid-glutinous,
1139	towards margin finely and rather weakly innately fibrillose-rivulose, with a more net-like
1140	pattern at centre, somewhat silvery-micaceous when young, smooth to sometimes faintly
1141	fibrillose; initially rather uniformly and pale ochraceous yellow to almost ochraceous white,
1142	resembling C. talus, sometimes pale grey to greyish ochre, with a faint, fugacious, blush tinge
1143	towards margin, with age discolouring patchwise to ochraceous brown (almost redbrown).
1144	Universal veil remnants very sparse, hardly visible, sometimes with silky, white remnants at
1145	margin.
1146	<i>Lamellae</i> crowded (L = 60–90), 0.4–0.8 cm broad, initially pale greyish blue, especially
1147	towards pileus margin, very soon fading to pale greyish brown; edge even to crenulate.

- 1148 Stipe $5-10 \times (0.8)1-1.5(2)$ cm, rather slender, with a faintly to rather distinctly marginate-
- 1149 bulbous base (up to 3 cm), glossy to slightly fibrillose, initially pale greybluish white in upper
- 1150 part, but soon almost white, sometimes more distinctly, fugacious violet at apex, turning brass
- brown spotted with age (especially when bruised). Universal veil at or near the bulb very
- 1152 sparse and hardly visible; whitish. Basal mycelium sparse, white.
- 1153 Context initially pale greybluish in upper part of stipe (especially towards cortex), rarely more
- pronounced violaceous; bluish colours soon fading, otherwise whitish, often with a brownish
- 1155 line under pileipellis and with brass brown spots where damaged.
- 1156 Macrochemical reactions negative (NaOH somewhat brownish).
- 1157 *Smell* faint, indistinct to somewhat like bread dough.
- 1158 *Exsiccata* more or less brownish; pileus margin and stipe sometimes remain whitish.
- 1159 *Basidiospores* [9, 15, 276], 9.6–11.8 × 5.9–7.4 μ m (MV = 10.7 × 6.7 μ m); variation of MVs:
- 1160 $10.1-11.3 \times 6.3-7.1 \ \mu\text{m}; \ Q = 1.42-1.79 \ (MV = 1.61), \ ellipsoid to \ subamygdaloid, \ rather$
- strongly and densely verrucose, warts diffuse or sometimes lense-like coalescent at apex,
- suprahilar plague rarely visible. *Lamella edge* more or less fertile, sterile cells small, clavateor cylindrical.
- 1164 *Pileipellis* duplex. Epicutis thin, at surface of 3–5 µm wide, loosely erect-entangled,
- 1165 gelatinous hyphae, hyaline, collapsed hyphae with pale, refractive pigment.
- 1166 Subcutis/hypodermium weakly to distinctly developed, hyphae $5-15 \mu m$ wide, basally usually
- 1167 with inflated elements up to 20 µm wide, tightly cemented, with pale yellow brown parietal-
- 1168 encrusting pigment, often forming an amber-like embedment; in transition to epicutis with
- subparallel, interconnected bundles of $5-8 \mu m$ wide hyphae with pale yellow brown parietal
- 1170 pigment, sometimes a few hyphae with weak zebra-striped encrusting pigment.
- 1171 *Veil hyphae at pileus surface* scattered or absent, hyaline and narrow (3–6 μm).
- 1172 Habitat: Associated mainly with Betula pubescens subsp. tortuosa (subalpine), also with other
- 1173 Betula taxa (boreal) and probably also Tilia cordata and Corylus avellana (boreonemoral),
- 1174 possibly also with Fagus (subalpine). In somewhat richer low-herb birch forests, also found in
- 1175 more open transition to alpine heaths (with *Betula nana* and *Betula pubescens* subsp. *tortuosa*
- 1176 thickets), in fjord sides in rich *Tilia-Betula* scree forests, sometimes also in mixed boreal
- 1177 coniferous forests with some Betula pubescens (Trøndelag, C Norway). In SW France
- 1178 recorded in a subalpine Fagus-Picea forest (Henry 1981 as C. fallaceicolor "forma
- 1179 méridionale"). In N America recorded with *Betula pumila*.

1180 *Distribution:* Apparently with a mainly subalpine northern distribution in Europe. So far

- 1181 known only from Norway and Sweden (12 DNA-verified records), one record from Scotland,
- 1182 one from France (Henry 1981 as *C. fallaceicolor* "forma méridionale"), and one from Canada
- 1183 in Newfoundland. Most finds are from (middle boreal-)subalpine central parts of S Norway
- and adjacent parts of Sweden but some also from boreonemoral fjord districts of Hardanger-
- 1185 Sogn-Geiranger (three DNA-verified records). The species reaches up to the tree limit of ca.

1186 1000–1100 m asl. in Jotunheimen, S Norway. Rare to very rare.

- 1187 *Comments: Cortinarius malachioides* is characterized by its pale, uniformly ochraceous
- 1188 yellow to pale ochraceous grey pileus, pale to very pale, fugacious bluish colours, a more or
- 1189 less marginate bulbous stipe base and comparatively small spores. A few specimens with
- 1190 somewhat stronger bluish pigment are seen, but normally bluish tinges on lamellae and stipe
- are very pale and fugacious. The almost lack of bluish tinges even on very young specimens is
- a unique feature within the sect. *Riederi*. The discolouring (to brown) of the pileus on mature
- 1193 specimens, may also be a diagnostic character. The species looks much like a *C. talus*, but is

1194 distinguished from the latter by pale blue-greyish lamellae, the larger spores and lack of a

- 1195 honey smell. The co-occurring *C. argenteolilacinus* var. *dovrensis* is distinguished by the
- 1196 larger spores, less bluish-violet colours and a more fibrillose pileus. On material studied so
- 1197 far, there is hardly any overlap in spore size between these two. When occurring in mixed
- 1198 forests, *C. malachioides* may be mistaken for the coniferous forest species *C*.
- anomaloochrascens. This species also has rather small spores (slightly larger than C.
- 1200 *malachioides*), but is distinguished by the quite glossy pileus and the initially deep violet
- 1201 lamellae and stipe. Henry (1981) described a "forme méridionale" of *C. fallaceicolor* Rob.
- 1202 Henry (an otherwise invalid name according to Art. 40.1 of Melbourne Code, see Index
- 1203 Fungorum), which belongs here. This form was collected under Fagus and Picea on Mont
- 1204 Aigoual (Cévennes, Massif Central, S France), and the referred material (R. Henry 80.811)
- appears to have an identical ITS sequence to our species. However, the holotype of *C*.
- 1206 *fallaceicolor*, collected from young *Picea* forest in French Jura (R. Henry 80.800; Henry
- 1207 1981) has a different ITS sequence, a sequence which (although only partial) seems to
- 1208 correspond with that of *C. fulvoochrascens*.
- 1209 According to present data, C. malachioides seems to be a mainly northern species in Europe,
- 1210 and is one of very few phlegmacia with a major habitat in the subalpine birch forest belt of the
- 1211 Scandinavian mountain chain. Only the here described *C. argenteolilacinus* var. *dovrensis*, as
- 1212 well as *C. blattoi* and *C. durus* share this habitat preference (see Brandrud et al. 2013). Like This is a post-peer-review, pre-copyedit version of an article published in Mycological Progress. The final authenticated version is available online at: http://dx.doi.org/10.1007/s11557-018-1443-0

- 1213 C. blattoi and C. durus, C. malachioides is likely to occur also in birch forests on Iceland, and
- 1214 probably has a wider, circumpolar distribution, following the subarctic *Betula* forest range.
- 1215 The one Canadian find from Newfoundland of this little known and probably very much
- 1216 overlooked species, indicates such a wide distribution.
- 1217 The nomenclature of *C. malachioides* is a complex matter. The type of *C. malachioides*
- 1218 undoubtly represents the present taxon. That is clear both from microscope examination (pers.
- 1219 obs.) and from ITS sequencing. However, Orton (1958) in the protologue, treated C.
- 1220 *malachioides* as one of three taxa around *C. malachius*, as a complex belonging to *Telamonia*
- 1221 s. lato (*Sericeocybe* s. Orton), and not to *Phlegmacium*. So the question remains, could Orton
- 1222 (i) regard this phlegmacioid taxon belonging to sect. *Riederi* as a *Telamonia* species close to
- 1223 *C. malachius*, or (ii) did he make a mistake when he chose the type specimen selecting a
- 1224 collection belonging to another species than his real, telamonioid *C. malachioides*? In other
- 1225 words; to what extent are our species and the type of *C. malachioides* in contradiction with
- 1226 the description in the protologue?
- 1227 Although not very clear, we think that the first alternative cannot be ruled out; that Orton
- 1228 consciously included our species in his (broader?) concept of *C. malachioides*, and that he
- 1229 believed this to be a *Telamonia/Sericeocybe*. His description seems largely to cover a *C*.
- 1230 *malachius* s. lato, but our species could also be involved:
- *1. The pileus*: Orton (1958) never mentioned a sticky-viscid pileus: "pileus ... overgrown by silky-white fibrils, then, around the margin innate fibrils". This description does not fit very well with the often glabrous-glossy-sticky pileus of our species. But still, such an appearance could be within the variation spectrum. It is possible to take a *Riederi* species for a *Sericeocybe (Telamonia* s. lato) when collected in dry conditions.
- 1236 1. *The stipe*: Here the same applies; our species has a glossy, smooth stipe almost without 1237 universal veil, whereas Orton (1958) describes the malachioides-stipe as "veil forming 1238 sometimes rather vague ring zone" (which is typical for the *C. malachius* group).
- However the "sometimes ring zone" could possibly refer to elements in the protologuenot covering our species.
- 1241 2. Habitat: Orton (1958) mentiones "under conifers" whereas our species is a *Betula*(-
- 1242 *Tilia*) species. However, *Betula pubescens* is often present at the classical, Scottish
- 1243 localities of Orton, and could well have been overlooked.
- 1244 As a conclusion, we emend the description in the protologue so that it is in accordance with
- 1245 the type.

1246	
1247	Collections examined:
1248	Canada. Newfoundland, Avalon Peninsula, Cape St. Mary, 28 Sept 2007, Andrus Voitk
1249	TN07-313, H7000977 (H). France. Languedoc, Gard, Mont Aigoual, M. Chevassut 10 Oct
1250	1980, R. Henry 80.811 (PC, type of C. fallaceicolor "forme méridionale"). Norway.
1251	Hordaland, Granvin, Åsen, T.E. Brandrud, TEB176-07 (O); Granvin, Håstabbaneset V, T.S.
1252	Jeppesen, B. Dima, TSJ2005-043/TEB371-05/DB2002 (C/O/BP); Ulvik, Eddagilet T.E.
1253	Brandrud, TEB155-07 (O). Aust-Agder, Froland, Myklandvatna nature reserve, T.E.
1254	Brandrud, TEB562-08 (O). Nord-Trøndelag, Høilandet, Folldalen, E. Bendiksen, T.E.
1255	Brandrud, EB-TEB 58-05 (O); Levanger, Ytterøya, Sandstadkammen E, TEB245-13 (O).
1256	Oppland, Vågå, Jotunheimen, Gjende, TEB469-11 & TEB473-11. Sweden. Ångermanland,
1257	H. Marklund et al., CFP980 (S). Härjedalen, Tännäs, S Malmagen, S. Jacobsson, SJ 80075
1258	(GB). Medelpad, Dysjöberget, D. Laber, K. Pätzold, TEB340-16/DB6138b (O). United
1259	Kingdom. Inverness-shire, Rothiemurchus, Loch-an-Eilean, P.D. Orton 29/9/55 (holotype,
1260	K(M)94426).
1261	
1262	Additional ITS sequence data from the public sequence repositories:
1263	Sweden. KJ421177/KS-CO1708 (as Cortinarius sp.) (Holmvallen, 17 Aug 2006, J.
1264	Vesterholt, herb. S, F251129).
1265	
1266	
1267	Cortinarius parksianus A.H. Smith, Contr. Univ. Mich. Herb. 2: 16 (1939)
1268	MycoBank: MB 253834
1269	
1270	Holotype: U.S.A., California, near Trinidad, 12 Nov 1937, A.H. Smith 8695 (MICH 10393).
1271	GenBank: MH846283.
1272	
1273	Description (according to the protologue):
1274	Pileus 8–12 cm diam., convex then plane with persistently decurved margin; colour initially
1275	purplish umber, then more olivaceous brown to cinnamon brown, margin with patches of
1276	whitish fibrils.
1277	Lamellae crowded, purplish (blue) when young, finally cinnamon brown.
1278	<i>Stipe</i> $10-12 \times 2-3.5$ cm, bulbose, bulb roundish, 3–4.5 cm wide, purplish (blue), fading This is a post-peer-review, pre-copyedit version of an article published in Mycological Progress. The final authenticated version is available online at:

http://dx.doi.org/<u>10.1007/s11557-018-1443-0</u>

- 1279 rapidly to whitish lilac, silky shiny. Cortina whitish and not copious.
- 1280 *Context* lilac grey in pileus, bright purple in the stipe apex, fading to white and pale lilac,
- 1281 resp.
- 1282 Basidiospores [1, 1, 50], 9.8–11.6 × 6.3–7.1 μ m (MV = 10.7 × 6.7 μ m); Q = 1.45–1.77 (MV =
- 1283 1.61), ellipsoid, verrucose ("roughened").
- 1284 Habitat: The type was collected under redwood (Sequoia sempervirens), within the N
- 1285 Californian belt of coastal redwood forests. Redwood does not form ectomycorrhiza,
- 1286 however, the ectomycorrhizal *Pseudotsuga mensiezii*, *Tsuga heterophylla* and *Abies grandis*
- 1287 might occur in the coastal redwood forests, and our species was probably associated with one
- 1288 of these trees at the site.
- 1289 *Distribution:* only known from the type locality.
- 1290 *Comments*: The above description is based on the original description in the protologue of
- 1291 Smith (1939). According to this, the species is a robust one, with (olive) brown tinges (no
- 1292 vividly ochraceous yellow or redbrown tinges), and with a bluish ("purplish") tinge when
- 1293 young. Furthermore, the spores are described as comparatively small, resembling those of *C*.
- 1294 anomaloochrascens or C. malachioides. Also the western N American species C.
- 1295 burlinghamiae has been collected in a similar habitat (mixed Pseudotsuga-Sequoia forest of
- 1296 California), but the latter seems to have much more vivid pileus colours, and larger spores.
- 1297
- 1298 *Collections examined*: U.S.A. California, Humboldt County, Trinidad, H.E. Parks estate,
- 1299 under Sequoia sempervirens, 12 Nov 1937, A.H. Smith, AHS 8695 (MICH 10393, holotype).
- 1300
- 1301 *Cortinarius burlinghamiae* Bojantchev, sp. nov.
- 1302 MycoBank MB 827877
- 1303 Figs 10a and 10b
- 1304

1305 Etymology: in honour of the American mycologist Gertrude Simmons Burlingham, one of the

- 1306 earliest researchers to work in the Pacific region of the United States.
- 1307
- 1308 Holotype: U.S.A. Montana, Lincoln County, Cabinet Mountains, off NF-278 Rd (N48°12'23"
- 1309 W115°34'31"), elev. 3660 ft., under Picea sitchensis, P. engelmannii, Pseudotsuga menziesii,
- 1310 Tsuga mertensiana, 2 Oct 2010, D. Bojantchev, DBB37303 (UC2023490). GenBank:
- 1311 KX768115.

- 1312
- 1313 *Pileus* 7–16 cm diam., hemispherical to convex then broadly plano-convex; margin involute,
- 1314 irregular, yellow to ochraceous or ochraceous brown near the disk, paler yellow to Sulphur
- 1315 yellow towards the margin, whitish velar remnants often present.
- 1316 Lamellae crowded, 1.2–2.3 cm broad, pale bluish at first then yellow, bluish tints often
- 1317 remaining in maturity, edges even, attachment sinuate, lamellulae abundant.
- 1318 Stipe 5–12 cm long, 1.5–3 cm wide, cylindrical to subclavate, often attenuated at age, with a
- 1319 moderately emarginated bulb, white to pale bluish lilac, more so towards the apex, silky
- 1320 shiny, bruising ochraceous. Cortina whitish at first, leaving an annular zone of dense fibrils on
- 1321 the stipe.
- 1322 *Context* white to bluish, more so towards the margin of the apex. Taste mild, earthy.
- 1323 Macrochemical reactions 5% KOH on fresh material reaction variable on context, greyish to
- 1324 ochraceous orange towards the bulb, with a halo, ochraceous on pileus, on dry material
- 1325 consistently ochraceous on all parts.
- 1326 *Smell* indistinct or somewhat earthy.
- 1327 *Basidiospores* [1, 1, 21], 11.2–13.2 × 6.9–8.2 μ m (MV = 12.2 × 7.5 μ m), Q = 1.51–1.75 (MV
- 1328 = 1.63), subobovoid to amygdaliform, coarsely vertucose. Basidia $33-46 \times 8-12 \mu m$, 4-
- 1329 spored, cylindro-clavate, clamped. *Cystidia* not observed.
- 1330 *Pileipellis* an ixocutis, duplex, epicutis composed of parallel to interwoven hyphae in a dense
- 1331 gelatinous matrix 220–260 µm thick, made up of 3–10 µm wide, irregular hyphae,
- 1332 hypodermium composed of cylindrical to ventricose cells 12–28 µm wide.
- 1333 Habitat and distribution: Under conifers (Picea, Pseudotsuga, Tsuga spp.) in montane to
- 1334 submontane areas of N America.
- 1335 *Comments*: This species is a close relative of the European *C. fulvoochrascens* and differs by
- 1336 the distribution, slightly paler colours, somewhat smaller spores and several nucleotides in the
- 1337 ITS region. A common host between the two collections from Montana and California is the
- 1338 Douglas fir (*Pseudotsuga mensiezii*), but more research is needed to determine whether that
- 1339 was the primary host of the species.
- 1340 A clade without acceptable statistical support including ectomycorrhiza sequences isolated
- 1341 from *Picea* roots from Alaska, Bonanza Creek (GenBank: KF617829, Taylor et al. 2014) and
- 1342 from Canada, British Columbia, Bulkley-Nechako (UNITE: UDB031686) as well as a
- 1343 sequence originated from a basidiocarp sample (from the U.S.A.) may represent another

- 1344 complex of species in sect. *Riederi* that falls close to *C. burlinghamiae* (Fig. 2, treated as
- 1345 *Cortinarius* sp.).
- 1346
- 1347 Collections examined: U.S.A. Montana, Lincoln County, Cabinet Mountains, D. Bojantchev,
- 1348 DBB37303 (UC2023490, holotype). California, Marin County, off Bolinas Ridge Road, elev.
- 1349 1130 ft., under Pseudotsuga menziesii, Sequoia sempervirens, Notholithocarpus densiflorus,
- 1350 30 Nov 2011, D. Bojantchev, DBB57001.
- 1351

1352 Additional ITS sequence data from the public sequence repositories:

1353 U.S.A. Wyoming, AF389139/IB19970218 (as C. fulvoochrascens).

- 1354
- 1355
- 1356 Conclusion
- 1357

Section *Riederi* is a striking example of a morphologically complex group that has not been well understood, and where the taxonomy is very difficult to sort out without the aid of molecular methods. This is due to the overlapping morphological variation of many of the taxa, combined with their rarity, which makes it difficult to study the entire morphological

1362 variation and their characteristic, taxonomically relevant features.

1363 The present morphological circumscription of sect. *Riederi* is quite new, including species

- 1364 with an innately fibrillose, ochraceous red brown to grey brown or whitish pileus, most with
- 1365 initially bright violaceous blue lamellae, and a more or less bulbous to slightly marginate
- 1366 bulbous stipe. The spores are large and ellipsoid-subamygdaloid, not of the more acutely
- 1367 amygdaloid-citriform kind most frequently encountered in phlegmacioid taxa.
- 1368 Many of the species are only known from a few collections, and optimal material including
- 1369 young specimens showing well-developed characters is very scarce for some of them (e.g. C.
- 1370 *pallidoriederi* (N Italy), *C. glaucocyanopus* (France, Hungary) and the N American *C.*
- 1371 *burlinghamiae* and *C. parksianus*). The phylogenetic analyses also clearly confirmed that the
- 1372 *C. argenteolilacinus* and *C. riederi-C. fulvoochrascens* groups (formerly treated in widely
- 1373 separated sections) are intimately related. Only after the phylogenetic structure was revealed,
- 1374 we were able to re-examine morphologically our material.
- 1375
- 1376

1377	Key to the European taxa of sect. <i>Riederi</i>
1378	
1379	
1380	1 Pileus when young pale ochre yellow to greyish, sometimes with a bluish tinge, sometimes
1381	almost white (resembling C. talus or C. largus); associated with deciduous trees (mainly
1382	Fagus, Corylus, Tilia and Betula). Pileus distinctly to only weakly innately fibrillose,
1383	somewhat micaceous, but rarely glossy, sometimes becoming faintly fibrillose-tomentose;
1384	lamellae pale to deep violet when young, soon fading; stem clavate to more distinctly
1385	marginate bulbous, fugacious violet 2
1386	
1387	1* Pileus when young (pale) ochraceous brown, to warmer ochre-red brown; associated with
1388	conifers (mainly Picea). Pileus usually distinctly innately fibrillose, smooth-glossy to
1389	fibrillose; lamellae deep violet when young, soon fading; stem clavate to more distinctly
1390	marginate bulbous, fugacious violet 5
1391	
1392	2 Spores normally <11.5 μ m long (10–11.5(12) × 6–7(7.5) μ m); pileus when young pale
1393	ochraceous yellow to olivaceous ochre (like a C. talus or C. anserinus); basidiocarps rather
1394	small and slender 3
1395	
1396	2* Spores normally >11.5 μ m long (11.5–13 × 7–8(8.5) μ m); pileus when young pale (bluish)
1397	greybrown (like a <i>C. largus</i>); basidiocarps often large and robust 4
1398	
1399	3 Mainly under Betula. Known from Scotland, W Norway fjords and mountains of Norway
1400	and Sweden, recorded also from E Canada and SW France, probably circumpolar; pileus
1401	initially usually pale ochre yellow (like a C. talus); lamellae and stipe initially pale bluish,
1402	almost greyish white; stipe ± marginate bulbous <i>C. malachioides</i>
1403	
1404	3* Under Fagus. Known from France and Hungary; pileus initially usually pale ochre yellow
1405	with an olive tinge (like a pale <i>C. anserinus</i>), lamellae and stipe initially rather deep violet;
1406	stipe ± marginate bulbous <i>C. glaucocyanopus</i>
1407	

1408	4 Mainly in C Europe under Fagus, but also in boreonemoral areas of S Fennoscandia under
1409	Tilia and Corylus. Basidiocarps large and robust; spores strongly verrucose, pileipellis with
1410	strongly encrusted hyphae C. argenteolilacinus var. argenteolilacinus
1411	
1412	4* Mainly in N Europe in subalpine Betula pubescens subsp. tortuosa forests, as well as in
1413	few cases with Corylus (once collected in Canada; probably circumpolar) Basidiocarps large
1414	to small; spores moderately to fairly strongly verrucose, encrusted pigment in pileipellis
1415	hyphae weak or absent C. argenteolilacinus var. dovrensis
1416	
1417	5 Spore MV's <12 μ m long (MV's 10.9–11.9 × 6.9–7.6 μ m); basidiocarps medium and
1418	slender. Pileus pale greyish ochraceous brown or tinged olive, stipe often slightly marginate
1419	bulbous, sometimes with watery bluish tomentum towards base; pileipellis without or with
1420	weak zebra-striped encrusted pigment; so far only known from Europe
1421	C. anomaloochrascens
1422	
1423	5* Spore MV's >12 μ m long (MV's 12.1–13.6 × 6.9–8.7 μ m); basidiocarps fairly large;
1424	pileipellis with distinctly zebra-striped encrusted pigment
1425	
1426	6 Veil remnants distinct, initially rendering base of stipe \pm girdled and pileus \pm fibrillose. Pileus
1427	vivid ochre brown, fulvous to red brown, stipe more or less marginate bulbous; stipe without
1428	white to bluish white tomentum towards base. So far only in Europe
1429	C. fulvoochrascens
1430	
1431	6 Almost devoid of veil, pileus and stipe smooth-glossy. Stipe often irregularly clavate-
1432	bulbose
1433	
1434	7 Mean width of spores 7.9–8.7 μ m, pileus vivid ochraceous brown, fulvous to redbrown,
1435	stipe often with watery bluish tomentum towards base; pileipellis hyphae with strong
1436	encrustations. In Europe and western N America C. riederi
1437	7 Mean width of spores 6.9–7.7 μ m, pileus paler ochraceous brown, bluish tomentum at stipe
1438	base not seen; pileipellis hyphae with only weak encrustations. So far only known from
1439	Europe C. pallidoriederi
1440	
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1454	
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1456	
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1772	Legends
1773	
1774	Fig. 1. Compressed phylogenetic tree based on RAxML analysis of the ITS region showing
1775	the main infrageneric clades within Cortinarius in Europe. The clade Riederi is highlighted in
1776	black. The genus Hebeloma was chosen as outgroup. Bootstrap supports values (>70%) are
1777	shown at the branches. The bar indicates 0.1 expected change per site per branch.
1778	
1779	Fig. 2. Mid-point rooted phylogenetic tree of sect. Riederi inferred from ITS dataset and
1780	RAxML analysis. Bayesian posterior probabilities (>0.90) and RAxML bootstrap supports
1781	values (>70%) are indicated on the branches at nodes (PP/ML). The bar indicates 0.01
1782	expected change per site per branch.
1783	
1784	Fig. 3 Spore photos of selected specimens of each species in sect. Riederi.
1785	
1786	Fig. 4 Mean values of spore lengths of species in sect. Riederi. Bars indicate 95% confidence
1787	interval. The missing bars are due to the limited number of available collections of those
1788	species.
1789	
1790	Fig. 5 Scatter diagram of mean values of spores of each measured collection in sect Riederi.
1791	
1792	Fig. 6 95% confidence ellipses of spore variations of the conifer associated species in sect.
1793	Riederi
1794	

- 1795 Fig. 7 95% confidence ellipses of spore variations of the species associated with deciduous1796 trees in sect. *Riederi*
- 1797
- 1798 Fig. 8 Basidiocarps of Cortinarius species in sect. Riederi associated with coniferous trees: a-
- b C. riederi (a TEB258-17 Norway, b SSt10-066 Sweden); c-d C. fulvoochrascens (c
- 1800 TEB530-13 Norway, d TEB-292-13 Norway); e-g C. anomaloochrascens (e TEB762-
- 1801 12/DB4814 Germany, f SSt17-202 Germany, g PRM924382 Czech Republic); h C.
- 1802 pallidoriederi (FB30-09-2011 Italy). Photos: a, c, e B. Dima; b, f G. Schmidt-Stohn; d T.E.
- 1803 Brandrud; g J. Borovička; h E. Castellini.
- 1804
- 1805 Fig. 9 Basidiocarps of species in sect. *Riederi* associated with deciduous trees: a–c C.
- 1806 argenteolilacinus var. argenteolilacinus (a ILF2013-98 Norway, b TSJ2011-025 Sweden, c
- 1807 TEB281-17 Norway); d C. argenteolilacinus var. dovrensis (KS-CO1703 Sweden); e–g C.
- 1808 malachioides (e-f TEB245-13 Norway, g TEB340-16/DB6138 Sweden); h-i C.
- 1809 glaucocyanopus (h DB2072 Hungary, i AL99/51 Hungary). Photos: a I.-L. Fonneland; b T.S.
- 1810 Jeppesen; c, e, f T.E. Brandrud; d K. Soop; g, h B. Dima, i L. Albert.
- 1811
- 1812 Fig. 10 Basidiocarps of Cortinarius burlinghamiae (a DBB37303 USA holotype, b
- 1813 DBB57001 USA). Photos: D. Bojantchev.