



**HELMHOLTZ  
ZENTRUM FÜR  
INFEKTIONSFORSCHUNG**

**This is a pre- or post-print of an article published in  
Stielow, B., Hensel, G., Strobel, D., Makonde, H.M.,  
Rohde, M., Dijksterhuis, J., Klenk, H.-P., Göker, M.  
Hoffmannoscypa, a novel genus of brightly coloured,  
cupulate Pyronemataceae closely related to Tricharina and  
Geopora  
(2013) Mycological Progress, 12 (4), pp. 675-686**

1 ***Hoffmannoscypha*, a novel genus of brightly coloured,**  
2 **cupulate *Pyronemataceae* closely related to *Tricharina* and**  
3 ***Geopora***

4 Benjamin Stielow<sup>1</sup>, Gunnar Hensel<sup>2</sup>, Dirk Strobel<sup>3</sup>, Huxley Mae Makonde<sup>4</sup>, Manfred  
5 Rohde<sup>5</sup>, Jan Dijksterhuis<sup>1</sup>, Hans-Peter Klenk<sup>4</sup>, Markus Göker<sup>4\*</sup>

6 <sup>1</sup> *Centraalbureau voor Schimmelcultures, Uppsalalaan 8, 3584 CT Utrecht, The*  
7 *Netherlands,* <sup>2</sup> *Fungarium Gunnar Hensel, Alte Lauchstädter Straße 22, 06217 Merseburg,*  
8 *Germany,* <sup>3</sup> *Pilzberatungsstelle Altkreis Stollberg, Parkstraße 9, 09399 Niederwürschnitz,*  
9 *Germany,* <sup>4</sup> *Leibniz Institute DSMZ – German Collection of Microorganisms and Cell*  
10 *Cultures GmbH, Inhoffenstraße 7b, 38124 Braunschweig, Germany,* <sup>5</sup> *Helmholtz Centre*  
11 *for Infection Research, Inhoffenstraße 7, 38124 Braunschweig, Germany*

12 \* Corresponding author (markus.goeker@dsmz.de); +49(0)531-2616-272; Fax: +49(0)  
13 531-2616-418

14 **Abstract**

15 The rare apothecial, cupulate fungus *Geopora pellita* (*Pyronemataceae*) is characterized  
16 by a uniquely bright yellow-orange excipulum. We here re-examine its affiliations by use of  
17 morphological, molecular phylogenetic and ultrastructural analyses. *G. pellita* appears as  
18 phylogenetically rather isolated, being the sister group of a clade comprising *Phaeangium*,  
19 *Picoa*, the majority of the *Tricharina* species, and the remaining *Geopora* species. Based  
20 on its phylogenetic position and its unique combination of morphological characters, we  
21 assign *G. pellita* to *Hoffmannoscypha*, gen. nov., as *H. pellita*, comb. nov. As in a previous  
22 study, analyses of both large subunit (LSU) and internal transcribed spacer (ITS)  
23 ribosomal DNA suggest that the remaining genus *Geopora* is paraphyletic, with the  
24 hypogeous, ptychothecial type species more closely related to *Picoa* and *Phaeangium*  
25 than to the greyish-brownish cupulate and apothecial *Geopora* spp., indicating that the  
26 latter should be reassigned to the genus *Sepultaria*. The current study also shows that ITS  
27 confirm LSU data regarding the polyphyly of *Tricharina*.

28 **Keywords:** Taxonomy, phylogeny, *Pezizales*, apothecia, ascospores



## 30 **Introduction**

31 *Ascomycota* is the largest fungal phylum and includes approximately 65,000 described  
32 species (Kirk et al. 2008). The class *Pezizomycotina* includes the ecologically most  
33 specialized and morphologically most diverging species, which contribute to a variety of  
34 important ecological processes such as wood and litter decay or are plant pathogens or  
35 mutualists in mycorrhizal symbiosis. The largest and most diverse family of  
36 *Pezizomycotina*, the *Pyronemataceae*, includes approximately 80 genera and around 660  
37 species (Kirk et al. 2008; Perry et al. 2007). Their ascoma morphology is highly diverse,  
38 including cupulate, discoid and pulvinate apothecia as well as hypogeous ptychothecial  
39 and stereothecial ascomata (Burdvall 1968; Perry et al. 2007; Tamm et al. 2010).  
40 Ecological strategies within the *Pyronemataceae* vary considerably between terricolous,  
41 coprophilous, lignicolous, pyrophilous and bryophilous forms (Benkert 1994; Hansen et al.  
42 2001; Krug and Kahn 1991; Perry et al. 2007; Spooner and Butterfill 1999; Vralstad et al.  
43 2002). While most species are saprotrophic, an increasing proportion is identified as  
44 ectomycorrhizal symbionts (Laessoe and Hansen 2007; Wei et al. 2010). *Pyronemataceae*  
45 have been taxonomically controversial as they are not conjunct by common morphological  
46 characters, neither macro- nor microscopically (Perry et al. 2007). While the positioning of  
47 many genera has recently been resolved with confidence within *Pyronemataceae* (Hansen  
48 and Pfister 2006; Laessoe and Hansen 2007; Perry et al. 2007), the problem of species  
49 recognition and, hence, species-diversity estimates, in particular for the sequestrate  
50 genera, has attracted much less attention (Guevara-Guerrero et al. 2011; Tamm et al.  
51 2010).

52 Within apothecial *Geopora*, most species are characterized by a greyish-brownish  
53 excipulum such as *G. arenicola*, *G. sepulta* and *G. tenuis*. However, *Geopora pellita*  
54 (Cooke & Peck) T. Schumacher, originally described as *Peziza pellita* by Cooke and Peck  
55 (1875), strongly differs in its macromorphology from its sister species, as it displays a  
56 colourful, brightly yellow-orange excipulum. Recent records about *G. pellita* are rare (Perry  
57 et al. 2007; Schumacher 1979; Wells and Kempton 1967) and the species appears in few  
58 identification keys only (Dougoud 2007; Hansen and Knudsen 2000). Schumacher (1979)  
59 reported the species for the first time outside the USA and reassigned it from *Peziza* into  
60 *Geopora*. Perry et al. (2007), using partial 28S rDNA sequences, provided the first  
61 molecular evidence that *G. pellita* phylogenetically strongly differs from other apothecial

62 *Geopora* species, since it was positioned closer to *Tricharina* than to cupulate and  
63 ptychothecial *Geopora* species, which was confirmed by Wei et al. (2010).

64 In the present study, we analyse a recently collected specimen of *G. pellita* by  
65 macromorphological, micromorphological and ultrastructural means as well as  
66 phylogenetic analysis using complete ITS and partial D1/D2 LSU (28S) rDNA sequences.  
67 The phylogenies suggest to recognize *Geopora pellita* (Cooke & Peck) T. Schumacher as  
68 separated, novel genus. This finding is strongly supported by the species' unique yellow-  
69 orange apothecium, whose development differs from other *Geopora* species. Accordingly,  
70 we propose *Hoffmannoscypha*, gen. nov., to accommodate the species.

71

## 72 **Material and Methods**

### 73 **Collection and morphological studies**

74 In general, the methods of collection, macroscopic and microscopic studies were those of  
75 Castellano et al. (1989) and Pegler et al. (1993). Fresh and dried specimens were cut by  
76 hand and mounted in water or alternatively in 5% KOH for microscopic observation. The  
77 newly collected fungal specimen was deposited under the accession number M-0156529  
78 at the Botanische Staatssammlung München (M) (Agerer et al. 2000) and at the  
79 Fungarium Gunnar Hensel (FUNGH). Specimens designated as *Geopora pellita* deposited  
80 at New York Botanical Garden Herbarium (NYBG) and at Harvard University Herbarium  
81 (FH) (Table 1) were used for comparative morphological analysis. Additionally, the  
82 collections from the pyrophilus genus *Tricharina* deposited at NYBG and M were analysed  
83 with light microscopy and field-emission scanning electron microscopy (FESEM), like the  
84 *Geopora* collections from Guevara-Guerrero et al. (2011) and the novel ones (Table 1).  
85 Tissue measurements were made with 40x and 100x oil immersion lenses (Zeiss Axiophot)  
86 and repeated 20 times. For FESEM, spores were harvested by scratching on a cross-  
87 sectioned *G. pellita* apothecium, seated onto conductive carbon adhesive tabs and  
88 covered with a gold film by sputter coating (SCD 500, Bal-Tec, Liechtenstein) before being  
89 examined in a field-emission scanning electron microscope (Zeiss DSM 982 Gemini) using  
90 the Everhart Thornley SE detector and the in-lens detector in a 50:50 ratio at an  
91 acceleration voltage of 5 kV. Images were recorded onto MO-disk, and contrast and  
92 brightness were adjusted with Adobe Photoshop CS3 and Illustrator CS5.

### 93 **DNA isolation, PCR, cloning and sequencing**

94 Total genomic DNA was extracted from approximately 100 mg of dried apothecium  
95 material using the Masterpure® Yeast Genomic DNA Kit following the manufacturer's  
96 protocol. DNA extraction from ancient specimens obtained from NYBG and BSM (Table 1)  
97 followed a modified protocol based on the EZNA Forensic DNA kit. Between 5 and 30 mg  
98 of apothecia, depending on age and condition of the herbarium specimens, were  
99 homogenized in 1.2 ml lysis buffer containing 1% SDS, 10 mM Tris pH 8.0, 5 mM NaCl, 50  
100 mM molecular biological grade DTT, 100 µg/ml proteinase K, 10 mM EDTA and 2.5 mM  
101 PTB (N-Phenacylthiazoliumbromide) based on a modification from Erickson et al. (2005).  
102 Microtubes were incubated in a water bath at 37 °C for 24 h following centrifugation at  
103 9000 g for 10 min and transfer of 1 ml supernatant into a new microtube, precipitation with  
104 600 µl 2-propanol and 60 µl 3 M sodium acetate at 4 °C for 48 h, following the EZNA  
105 forensic DNA manufacturer's instructions with the exception of the last washing step being  
106 performed four times. The ITS nrDNA region was amplified with PCR primers ITS1/ITS4  
107 and ITS1F/ITS4 under semi-nested conditions (Gardes et al. 1993, White et al. 1990,  
108 Stielow et al. 2010). PCR conditions for amplifying the partial 28S rDNA using the standard  
109 primers LR0R and LR3 only differed in their annealing temperature (55 °C instead of 60  
110 °C). PCR for ancient specimens was performed with ITS5/ITS4 and followed by  
111 reamplification under semi-nested conditions with ITS3/ITS2 paired with ITS4/ITS5 or by  
112 direct amplification under standard conditions. PCR products were cut out or directly  
113 purified using Macherey-Nagel NucleoSpin Extract II kit (740609.50). The cycle-  
114 sequencing reaction was set up using the Beckman Coulter GenomeLab DTCS Quick  
115 Start Kit or the ABI big dye terminator v3.1 following the manufacturers' protocols, followed  
116 by bidirectional sequencing with a Beckman Coulter Genome lab capillary electrophoresis  
117 system or the Lifetechnologies (ABI) 3730XL DNA analyser. PCR products from ancient  
118 specimens that resulted in poor trace quality were cloned using the TOPO TA 2.1 cloning  
119 kit (LifeTech). Sequences were manually corrected for sequencing artefacts and forward  
120 and reverse sequences assembled using Invitrogen Vector NTI 11 or Lasergene Seqman.

### 121 **Phylogenetic inference**

122 The ITS and LSU nrDNA alignments were the ones used in Guevara-Guerrero et al.  
123 (2011). These were carefully compiled, extensively tested regarding the sensitivity of the  
124 resulting phylogenies to alignment ambiguity (which was negligible), and already used to  
125 draw taxonomic conclusions in the group. The newly obtained *G. pellita* and *Tricharina*

126 sequences that comprised both ITS1 and ITS2 were added to the ITS alignment using the  
127 POA software (version 2; Lee et al. 2002) in profile alignment mode. As in our previous  
128 study (Guevara-Guerrero et al. 2011), phylogenetic analysis under the maximum-likelihood  
129 (ML) criterion (Felsenstein 1981) was conducted with RAxML version 7.2.7, using its novel  
130 rapid bootstrap option combined with the autoMRE bootstopping criterion (Pattengale et al.  
131 2009) with subsequent search for the best tree under the GTRMIX approach (Stamatakis  
132 et al. 2008). Bootstrapping under the maximum-parsimony (MP) criterion (Fitch 1971) was  
133 again done with PAUP\* version 4.0b10 (Swofford 2002), treating gaps as missing data,  
134 collapsing branches of zero minimum length, and using 10 rounds of random sequence  
135 addition followed by TBR branch swapping per bootstrap replicate. In MP bootstrapping,  
136 1000 replicates were conducted. As before (Guevara-Guerrero et al. 2011), rooting of the  
137 resulting trees was done using the midpoint rooting method (Farris 1972; Hess and Russo  
138 2007). Sequence alignments and phylogenetic trees are included in the online  
139 supplementary material. For depicting the trees, clades comprising at least three  
140 sequences were collapsed if they were either taxonomically homogeneous or contained  
141 only environmental samples. If a clade contained environmental samples some of which  
142 had a genus annotation, these genera were indicated.

143 The additional ITS sequences from which only ITS1 or ITS2 could be amplified due to the  
144 insufficient preservation of the material were analysed separately. Here we focussed on  
145 the identity of the biological material deposited as *Geopora pellita* and thus only calculated  
146 pairwise similarities from *exact* pairwise sequence alignments using the Smith-Waterman  
147 algorithm as implemented in the EMBOSS suite (Rice et al. 2000).

148

## 149 **Results**

### 150 **Phylogenetic inference from ITS rDNA sequences**

151 The alignment comprised 250 ITS rDNA sequences and had a total length of 1813  
152 positions. The resulting best ML tree had a log likelihood of -20,735.83 and is shown in  
153 Fig. 1 together with ML (left) and MP (right) bootstrap values on each branch. The  
154 separation of the outgroup clades on the one hand, comprising operculate apothecial  
155 discomycete genera such as *Aleuria*, *Cheilymenia*, *Pseudaleuria*, *Scuttelinia*, and  
156 *Wilcoxina*, but also one of the newly generated *Tricharina* sequences (JQ824118), and the

157 ingroup clades on the other hand was strongly supported (100/97%). Note, however, that  
158 the comparison of annotations such as “fungal sp. ARIZ AZ0886” with the LSU tree (Fig. 2)  
159 indicated yet another *T. gilva* cluster, separate from all newly sequenced *Tricharina*  
160 samples. The ingroup clades included apothecial *Geopora* spp. (= *Sepultaria* spp.;  
161 Guevara-Guerrero et al. 2011), ptychothecial *Geopora* spp., *Phaeangium* spp., *Picoa* spp.  
162 and three of the newly generated *Tricharina* sequences. Strong support was achieved for  
163 the clade comprising these three sequences together with some environmental samples  
164 (96/98%) as well as for its sister group, comprising all remaining ingroup sequences  
165 (99/97%). Within the latter, *Geopora pellita* branched first, followed by environmental  
166 sequence GQ281480, whose annotation indicates the ectomycorrhizal morphotype of  
167 *Pinirhiza daqingensis*. Moderate support (78/73%) was obtained for the monophyly of all  
168 remaining ingroup sequences. The monophyly of apothecial *Geopora* spp. was strongly  
169 supported (100/99%), as well as clade comprising *Phaeangium* and *Picoa* sequences  
170 (100/99%). Moderate support was obtained for the monophyly of the hypogeous,  
171 apothecial *Geopora* spp. (92/66%) and for their sister-group relationship with *Phaeangium*  
172 and *Picoa* (72/77%).

### 173 **Phylogenetic inference from 28S (LSU) rDNA sequences**

174 The alignment comprised 630 LSU rDNA sequences and had a total length of 6711  
175 positions. The resulting best ML tree had a log likelihood of -50,155.98 and is shown in  
176 Fig. 2 together with ML (left) and MP (right) bootstrap values on each branch. Again,  
177 outgroup clades were collapsed due to their size; for the complete tree see the  
178 supplementary files. Very strong to moderate support (99/82%) was achieved for the  
179 monophyly of a group comprising the genera *Geopora*, *Phaeangium*, *Picoa* and *Tricharina*  
180 with the exception of *Tricharina gilva*, which belonged to the sister group of that clade.  
181 Strong to low (89/61%) support, depending on the phylogenetic optimality criterion  
182 (ML/MP), indicated the monophyly of a cluster comprising *Phaeangium*, *Picoa* and  
183 *Geopora* except for *G. pellita*. The sister-group relationship of hypogeous ptychothecial  
184 *Geopora*, *Phaeangium* and *Picoa* was also highly to weakly supported (95/60%), whereas  
185 the monophyly of the clade containing the majority of accessions annotated as *Tricharina*,  
186 obtained strong to moderate support (96/84%). The Genbank 28S sequence DQ220343,  
187 annotated as a *G. pellita* specimen collected in Michigan (USA) by Pfister in 1969 (Perry et  
188 al. 2007), was almost identical to the sequence obtained from our specimen (Fig. 2).



## 189 **Sequence and morphological identity of herbarium specimens**

190 The best hit of partial ITS sequence obtained from the NYBG 114 specimen (JQ062972)  
191 was to our complete *G. pellita* sequence (HQ913564), yielding 96.1% Smith-Waterman  
192 similarity. The second best hit (HM123158) corresponded to only 83.7% sequence identity.  
193 Identical results were obtained for the NYBG 301 specimen (JQ062973). The NYBG 228  
194 specimen (accession number JQ062974), however, yielded FM206460 as best hit  
195 (99.0%), followed by hits to other *Geopora arenicola* sequences. A photograph  
196 (supplementary material, image 20) attached to specimen NYBG 228 (collected in 1906),  
197 confirmed that this specimen should not be assigned to *G. pellita*, since for some of the  
198 apothecia it was obviously shown that they started in a hypogeous state, instead of  
199 developing superficially on the substrate as in all other examined specimens annotated as  
200 *G. pellita*. (The term “hypogeous” is used here as defined by Kirk et al (2008); see also  
201 Weber et al (1997) p. 156, schemes C–D and F–I, for illustrations of strictly hypogeous  
202 sporocarps in Ascomycetes. “Superficially on the substrate” refers to the appearance of  
203 sporocarps as given by Yang and Korf (1985) on p. 470, in schemes A to D). For this  
204 reason, the morphological description of *G. pellita* given below excluded NYBG 228. LSU  
205 data revealed the identity of our newly collected *G. pellita* specimen with the DHP 297  
206 specimen (DG220343), as apparent from Fig. 2.

207 Unfortunately, not all specimens could be sequenced, and in some cases the condition of  
208 the deposits was so poor that they could hardly be examined microscopically. This is  
209 particularly apparent for the holotype material (NYBG 00914741) collected by Cooke and  
210 Peck (1872), deposited at the NYBG. The type specimen of *G. pellita* available to us for  
211 comparison would most likely not allow for molecular sampling and exists, at the NYBG, as  
212 a single microscopical glass slide. Any other isotype material deposited in other institutions  
213 was not available to us for this study. The description and the drawing (supplementary  
214 material, image 21) given by Cooke and Peck, however, shows apothecium and spore  
215 characteristics identical to those of the other examined *G. pellita* specimens except NYBG  
216 228, as detailed below.

217 The partial ITS sequences obtained from herbarium deposits annotated as *Tricharina*  
218 yielded the following best hits. Specimen M-0178316 (“*Tricharina praecox*”, JQ824119)  
219 yielded as best hit HM123089 (“fungal sp. ARIZ AZ0347”), which according to the  
220 comparison with the LSU data (Fig. 2) belongs to a *Tricharina gilva* cluster. But the  
221 similarity was only 87.2%; M-0178316 was thus judged as an unknown *Tricharina* species.

222 M-0178317 ("Tricharina praecox", JQ824119) yielded a much higher similarity to this  
223 cluster (99.7% to "fungal sp. ARIZ AZ0886") and was thus regarded as a misidentification  
224 of *T. gilva*. This was confirmed by M-0178315, annotated as "Tricharina gilva", also  
225 matching this cluster (99.0–100.0%). M-0178313 ("Tricharina gilva") had as best hit  
226 "Geopora cf. cooperi SOC1051" (FJ789595), but only with a similarity of 82.4%, and thus  
227 was judged as misidentified and of uncertain affinity.

228 The size of all apothecia, as well as the size and shape of the spores of all examined  
229 specimens annotated as *Tricharina* (examples are given in the supplementary material,  
230 images 23–30) were in accordance with previously published descriptions by Yang and  
231 Korf (1985) and Ellis and Ellis (1998), e.g., much smaller in size than those of *Geopora*  
232 (details are given below). This also holds for the NYBG 1968 specimen (apothecia 1.0–1.5  
233 mm in diameter), even though its ITS sequence indicates a taxonomic affiliation to other  
234 discoid *Pyronemataceae* like *Pyronema*, *Trichopheae* and *Wilcoxina* spp. (Fig. 1). Its black  
235 excipulum, the more elongated, entirely smooth spores (supplementary material, image  
236 14, 30) containing black guttules (observed unstained in water), as well the substrate ("flat  
237 sand within a greenhouse"), do not match the description of *Tricharina* species in the  
238 literature. For this reason, ITS and morphological or ecological data appear to be in  
239 agreement regarding the investigated herbarium specimens of *Tricharina*. The following  
240 comparison between *Geopora pellita* and *Tricharina* collections thus only relies on the  
241 *Tricharina* collections that either could be approved using ITS sequencing or were  
242 collected from burned soil, even though the morphological differences to *G. pellita* would  
243 be the same for the other specimens annotated as *Tricharina*.

#### 244 **Morphological comparison of *Geopora* spp. and *Tricharina* spp. with *G.*** 245 ***pellita***

246 Results from macro- and micromorphological as well as ultrastructural examinations of the  
247 target specimens are shown in Figs. 3 and 4; further Nomarski interference contrast and  
248 SEM pictures of the investigated reference specimens are given in the supplementary  
249 material (images 1–16, 23–30).

250 The apothecia of *G. pellita* (Fig. 3, image 1) were yellow-orange and superficially attached  
251 to the substrate. They might be slightly sunken into the substrate but did not emerge from  
252 a hypogeous development. The star-like shape of the mature apothecia (Fig. 3, image 1)  
253 was similar to the one typical for other *Geopora* spp. (Fig. 3, image 5; *G. sepulta*) but

254 these nevertheless strongly differ in color. The holotype drawing of *P. pellita* Cook & Peck  
255 (1872), given in the supplementary material (image 21), shows exactly the same type of  
256 apothecium as given in Fig. 3, image 1. Mature apothecia of *Tricharina* (supplementary  
257 material, images 17–19) were considerably smaller than the ones of *G. pellita*, commonly  
258 less than 10 mm in diameter (mature apothecia are 1–5 mm in diameter for most species),  
259 but similar in color to *G. pellita*.

260 Juvenile apothecia of *Geopora* spp. tended to expand their exoperidium to very distinct  
261 lobes, thereby pulling the apothecium from a hypogeous to a position superficially on the  
262 substrate (Fig. 3, image 5). But *G. pellita* did not appear to develop in this way, even  
263 though the lobes of its exoperidium were obvious (Fig. 3, image 1). Already its juvenile  
264 apothecia were superficially attached to the substrate and its development thus appeared  
265 more like the one of *Tricharina* spp., even though the apothecia of the latter were not lobed  
266 (supplementary material, images 17–19).

267 Multiseptate, finely warted and cylindrical setae with blunted to pointed apices were found  
268 in all investigated species. *G. pellita* showed a broad basal cell connected to the cells of  
269 the ectal excipulum, from where the finely warted setae emerge (Fig. 3, images 4, 6). The  
270 setae were always arranged in fascicles nonetheless their density became narrower  
271 towards the apothecial base. We did not detect differences to the setae of other *Geopora*  
272 spp., or *Tricharina* spp. The setae of *G. pellita* did not contain globose inclusions, as  
273 known from pyrophilus genera such as *Wilcoxina* spp (Yang and Korf 1985).

274 Spore sizes were in the range of 21–27 x 10–12  $\mu\text{m}$  in *G. pellita*. Similarly, the examined  
275 *Geopora* specimens never showed spores less than 20  $\mu\text{m}$  in length. In contrast, spore  
276 sizes were 12–17 x 5–11  $\mu\text{m}$  for the investigated *Tricharina* species. Asci of all three  
277 genera were of similar shape and size, and always cylindrical, apically operculate with a  
278 narrowing base, non-amyloid, uniseriate and eight-spored. In *Tricharina* they were always  
279 less than 200  $\mu\text{m}$  in length but in *G. pellita* and the other *Geopora* spp. always longer than  
280 200  $\mu\text{m}$  (e.g., Figure 3, image 2). The paraphyses were usually slender, multiseptate and  
281 slightly clavate at the apex. In those species with a colorful hymenium, the paraphyses  
282 contained small inclusions of pigments that apparently gave rise to the overall color of the  
283 apothecium (Figure 3, images 2, 4). This character could mostly only observed on fresh,  
284 recently collected apothecia. All findings are supported by pictures assembled in the  
285 supplementary material, obtained from the examined herbarium specimens (Table 1).

286 The ascospores of *G. pellita* and *Geopora* spp. appear, when observed with Nomarski  
287 interference contrast microscopy, entirely smooth and guttulate (Fig. 3, image 2). The  
288 guttulate spores of *G. pellita* have already been indicated by Cooke & Peck in their  
289 drawing of the type specimen (supplementary material, image 21). Likewise, the spore  
290 surfaces of *Tricharina* spp. appeared smooth (supplementary material, images 23–29).

291 Mature ascospores of *Geopora pellita* showed a finely warted ornamentation when  
292 observed with FESEM (Fig. 3, image 7; Fig. 4, images 1, 2; supplementary material,  
293 images 5–9). This ornamentation was not observed in juvenile spores, which were entirely  
294 smooth (Fig. 3, image 7; Fig. 4, image 2). The same ornamentation of juvenile,  
295 intermediate and mature spores was observable on the specimens obtained from NYBG  
296 and FH (supplementary material, images 5–9), indicating the biological identity of this *G.*  
297 *pellita* material. Immature spores of *G. arenosa* and *G. sepulta* were also smooth, whereas  
298 mature ones showed a rough but not warted spore surface (supplementary material).  
299 *Tricharina* sp. M-0178316 showed juvenile smooth spores in addition to mature spores that  
300 were even more pronouncedly warted than the ones of *G. pellita* (supplementary material,  
301 images 11–13, 15–16). The same pronounced ornamentation was observed for specimen  
302 M-0178317, but not for specimen Rehm. Ascom. 456 / 1878 that showed a rough but not  
303 warted surface (supplementary material, image 10 and 11–12).

## 304 **Discussion**

### 305 **Identity of the investigated specimens**

306 There is ample evidence that our specimen corresponds to the *G. pellita* from the  
307 literature. First, the unique macro- and micromorphology of *G. pellita*, which is not found in  
308 any closely related apothecial genera, makes a misidentification rather unlikely. Second,  
309 with the exception of the specimen annotated as *G. pellita* (NYBG 228), which turned out  
310 to be affiliated to *G. arenicola*, both morphologically and regarding its ITS rDNA, the  
311 morphology and the ITS sequences were almost to entirely identical between the newly  
312 collected *G. pellita* specimen and the herbarium material. This was confirmed by  
313 observations on the ultrastructure of the spores, which revealed a finely warted  
314 ornamentation in all proper *G. pellita* specimens that was somewhat distinct from the other  
315 *Geopora* species examined, as well as from the *Tricharina* species under study. This  
316 finding is not in conflict with the smoothness of the ascospores of *G. pellita* reported by

317 Cooke and Peck (1872), Dougoud (2007), Schumacher (1979) and Wells and Kempton  
318 (1967) because the warts could not be seen in light microscopy. Accordingly, the  
319 ascospore surface of other *Geopora* spp. has also been described as entirely smooth by a  
320 variety of authors (Breitenbach and Kränzlin 1981; Dennis 1981; Hansen and Knudsen  
321 2000; Tamm et al. 2010), even though the mature ascospores are not smooth when  
322 visualized by FESEM.

323 Some herbarium deposits were too old and scarce to extract DNA. This is particularly  
324 evident for the type specimen (*Peziza pellita* Cooke & Peck 1872, *Grevillea* 1: 5, NYBG  
325 specimen ID 00914741). Accordingly, epitypification appears to be the best way to address  
326 the critical issue of a representative specimen for this species. The epitype, our novel  
327 collection, is deposited at the Botanische Staatssammlung München under the accession  
328 number described below (corresponding curator Dr. D. Triebel). Despite its overall poor  
329 condition, however, the type deposit contains a drawing, which unambiguously indicates a  
330 micro- and macromorphology identical to the proposed epitype, and thus the biological  
331 identity of the investigated specimens. The description of the *G. pellita* habitats found in  
332 the literature (Schumacher 1979), “growing in a sand accumulation in the upper inundation  
333 zone of the river on coarse sand among *Pohlia gracilis* and *Bryum* spp.”, also corresponds  
334 well to the collection site of our novel specimen, which was found in a sand pit in  
335 association with *Pinus* sp. and embedded in unidentified mosses.

336 Dried ascomata of many small discoid *Pyronemataceae* are known to be very brittle, and  
337 often cells cannot be properly hydrated again. Even later tissue observations made from  
338 herbarium specimens of our own collections of *G. pellita*, whose apothecia are much larger  
339 in size than those of *Tricharina*, were almost impossible. Thus it is difficult to identify  
340 herbarium specimens of *Tricharina* based on apothecial macromorphology due to the age  
341 of the specimens. A cross-comparison of the sequence affiliation of those specimens from  
342 which ITS sequences could be obtained with the LSU data, however, allowed us to  
343 conclude that two distinct but real *Tricharina* clades exists, one harboring the type species,  
344 *T. gilva*, and a second one containing at least *T. ochroleuca* (which occurs in a clade of  
345 comparable positioning in both the ITS and LSU trees; see also Perry et al. 2007) but  
346 probably also *T. hiemalis* (this study), *T. groenlandica* (this study) and *T. praecox* (Perry et  
347 al. 2007). Moreover, we have shown that *Tricharina* species reported to develop smooth  
348 ascospores when viewed with the light microscope (Yang and Korf 1985) have obviously  
349 warted spores when examined by scanning-electron microscopy. Our examination also

350 indicates that quite a few herbarium specimens of *Tricharina* are misidentified, and that a  
351 revision of the genus is needed.

### 352 **Classification of *G. pellita* relative to *Geopora* and *Tricharina***

353 Recently, phylogenetic studies on major genera of *Pyronemataceae* have been conducted  
354 by Perry et al. (2007). The results of this study revealed *Geopora* (except *G. pellita*) as a  
355 monophyletic group using maximum-parsimony and Bayesian analyses, but neither  
356 *Phaeangium* nor *Picoa* were included in the dataset. Using either 28S or ITS rDNA  
357 sequence data, our analyses place these two genera within a paraphyletic *Geopora* with  
358 high confidence, at least under the maximum-likelihood criterion, in agreement with the  
359 results obtained by Guevara-Guerrero et al. (2011) and Sbissi et al. (2010), and  
360 highlighting the importance of sufficient taxon sampling. Because *Sepultaria* already exists  
361 as a validly published name for the apothecial *Geopora* spp., there is little reason for not  
362 using it once again for these fungi. As shown in the present study, the other necessary  
363 measure to obtain a monophyletic *Geopora* is to exclude *G. pellita*. This species is neither  
364 phylogenetically placed within the genus nor does its macromorphology agree with the  
365 other *Geopora* species.

366 *G. pellita* forms a grade in the LSU tree together with *Tricharina*, which appears  
367 polyphyletic in the tree, subdivided into a *T. gilva* and a *T. ochroleuca/praecox* clade. The  
368 latter is more closely related to *Geopora*, *Phaeangium*, *Picoa* and *Sepultaria* than *G.*  
369 *pellita*, whereas the *T. gilva* clade is the sister group of all these taxa (Fig. 2). Apparently,  
370 including *G. pellita* into *Tricharina* would at most change the status of the latter from  
371 polyphyletic to paraphyletic (see Farris 1974 for formal definitions of these terms) and  
372 would not be an acceptable solution either. The ITS tree (Fig. 1) shows the same  
373 relationships, the main difference being the position of the *T. gilva* cluster; but this is just  
374 an issue of rooting. That only monophyletic groups can be accepted in modern taxonomic  
375 classifications can hardly be denied (Farris 1979; Hennig 1965; Wiley and Lieberman  
376 2011).

377 The presence of the prominent hyaline to brownish fascicular hairs in *Tricharina*, arising  
378 from the ectal excipulum, which are also found in *G. pellita* or other *Geopora* species  
379 (Yang and Korf 1985), cannot be used as morphological character delimiting the three  
380 genera. Clear differences in septation, size, tip shape or inclusions have neither been  
381 found. Since the apothecium and excipulum micromorphologies are not useful for the

382 delimitation of *Geopora* species, too, spore characters and excipulum macromorphology  
383 have been used instead (Dougoud 2007; Hansen and Knudsen 2000; Tamm et al. 2010;  
384 Yao and Spooner 1996). Schumacher (1979) renamed *Peziza pellita* to *Geopora pellita*  
385 based on a description in accordance with the ones given by Cooke and Peck (1872) and  
386 by Wells and Kempton (1967), which referred to a similar excipulum and spore  
387 morphology. The lobes of the excipulum fulfill an important function in *Geopora* spp. by  
388 pulling the mature apothecium above the substrate to disperse the spores; they should not  
389 be compared to possible rifts, which may occur in many small *Discomycetes* such as  
390 *Tricharina*.

391 Prominent lobes are seen in *G. pellita*, too, but cannot have the same function as in other  
392 *Geopora* species because already the juvenile apothecia of *G. pellita* are superficially  
393 attached to the substrate. Another obvious distinction of *G. pellita* from *Geopora* species  
394 such as *G. arenicola*, *G. arenosa*, *G. cervina*, *G. sepulta* and *G. sumneriana* is that these  
395 have a brownish-grayish pigmented excipulum in common, whereas the yellow-orange  
396 apothecia of *G. pellita* resemble the ones of *Tricharina* regarding their color.

397 But morphologically *G. pellita* does neither fit to *Tricharina*. Both *G. pellita* and the other  
398 *Geopora* species are characterized by a strong lobation of mature apothecia, which is  
399 neither known from *Tricharina* nor other pyrophilus genera. This is also supported by the  
400 size of the ascospores, which are within *Geopora* (and *G. pellita*) always longer than 20  
401  $\mu\text{m}$  and in *Tricharina* always shorter than 20  $\mu\text{m}$ . The specimens observed in this study  
402 analogously differed regarding the length of their asci. Apothecia of *Tricharina* are strictly  
403 cupulate, in some species even discoid (Yang and Korf 1985). The size of their apothecia  
404 is usually between 1–10 mm, only in a single species, *T. fibrillosa* (Currey) Yang & Korf, up  
405 to 20 mm, which is similar to the size of *G. pellita* (Yang and Korf 1985). Ecological  
406 differences separating *G. pellita* from *Tricharina* are less certain at the moment, since *G.*  
407 *pellita* is not definitively known to form ectomycorrhizal associations. At least, this species  
408 has not yet been found on pyrophilous sites or on decaying wood, which is typical for most  
409 *Tricharina* species (Yang and Korf 1985).

410 Based on our results, we suggest the novel, so far monotypic genus *Hoffmannoscypha* to  
411 accommodate *G. pellita* as *H. pellita*, comb. nov. An additional splitting of *Tricharina*, which  
412 is, of course, beyond the scope of the present study, might be an acceptable future  
413 solution for the remaining non-monophyly of *Tricharina* apparent in phylogenetic analyses.

414

415 **Taxonomy**

416

417 **Hoffmannoscypha** Stielow, Göker & Klenk, gen. nov.

418 **Mycobank number:** MB 561770.

419 **English description:** APOTHECIA superficially attached to the substrate, often slightly  
420 sunken into it, without stipe, roundish when juvenile and nearly entirely closed at the apex,  
421 cupulate when mature, at full maturity with ripped edges; lobes erect when juvenile,  
422 effused when mature. ECTAL EXCIPULUM (epicutis) dark yellow-orange, seldom orange-  
423 brown, pseudoparenchymatous, with angular or isodiametric, thick-walled cell  
424 agglomerates (textura angularis), cells giving rise to dark brown, thick-walled, multiple  
425 septate cylindrical finely warted setae, single or cespitose, dark brown in water and 5%  
426 KOH. ASCI inoperculate, cylindrical, eight-spored, monoseriate. PARAPHYSES slender,  
427 single, septate, with gentle orange pigmentation giving rise to the colour of the hymenium,  
428 slightly thickened and bent at the tip, arranged in palisade order.

429 **Type species:** *Hoffmannoscypha pellita* (Cooke & Peck) Stielow, Hensel, Göker & Klenk

430 **Etymology:** “*Hoffmannoscypha*” = “Hoffmann's cup”. Named in honour of the German  
431 mycologist and DSMZ curator Dr. Peter Hoffmann who dedicated the over 40 years of his  
432 working life to collection, preservation and identification of fungi.

433 **Anamorphs:** The species is unknown from pure axenic culture.

434

435 ***Hoffmannoscypha pellita*** (Cooke & Peck) Stielow, Hensel, Göker & Klenk, comb. nov.

436 = *Peziza pellita* Cooke & Peck 1872, Grevillea 1: 5

437 = *Lachnea pellita* (Cooke & W. Phillips) Sacc. 1889, Syll. fung. (Abellini) 8: 169

438 = *Sepultaria pellita* (Cooke & Peck) Seaver 1928, North American Cup-fungi,  
439 (Operculates), New York 152

440 = *Geopora pellita* (Cooke & Peck) T. Schumacher 1979, Norw. J. Bot. 26 (1): 56



441 **Mycobank number:** MB 561771.

442 **English description:** With the features of the genus. APOTHECIA 0.5–2 cm in diameter.  
443 ODOUR not distinct. ECTAL EXCIPULUM (epicutis) 325–400  $\mu\text{m}$  wide, cells 14–45  $\mu\text{m}$   
444 wide; setae on average 60–140  $\mu\text{m}$  in length, at most 500  $\mu\text{m}$  long, 4–6  $\mu\text{m}$  thick.  
445 MEDULLA 400–600  $\mu\text{m}$  wide, with prostrated interwoven hyphae (intermediate between  
446 textura intricata and textura globulosa), several hyphae forming large globose inflations;  
447 hyphae 3–14  $\mu\text{m}$  wide; globose inflations 15–30  $\mu\text{m}$  wide; hyaline, slightly orange in water  
448 when fresh; hyaline, colourless in 5% KOH. ASCOSPORES ellipsoid, smooth and hyaline  
449 when juvenile; guttulate with a large central guttule, two smaller guttules adjacently beside,  
450 spores with guttules of the same size rarely observable; entirely mature spores finely  
451 warted-punctuated (ornamentation approximately 300–800 nm in height), walls  
452 approximately 0.5–1  $\mu\text{m}$  thick, in water and 5% KOH. Ascospore size (23–)23.5–27.5(–28)  
453  $\times$  (10.5–)11–13(–13.5)  $\mu\text{m}$ , on average 25  $\times$  12  $\mu\text{m}$ , length/width quotient 1.9–2.3, on  
454 average 2.1. ASCI 220–280  $\times$  14–19(–24)  $\mu\text{m}$  in size. PARAPHYSES in water and 5%  
455 KOH 3.5–5.5  $\mu\text{m}$  in diameter at the tips, 6–7  $\mu\text{m}$  in diameter at the centre.

456 **Specimens examined:** M-0156529, Botanische Staatssammlung Munich = GH20100409,  
457 Fungarium Gunnar Hensel; Germany, Freihufen-Großräschen, sand pit, approximate  
458 location 51°34'31.22" N 13°57'49.42" E, 130 m above sea level, in sandy mineral soil,  
459 beneath pine trees (*Pinus* sp.), Dirk Strobelt & Gunnar Hensel, 4<sup>th</sup> September 2010. Four  
460 additional specimens of *G. pellita* that were investigated in this study are given in Table 1.

461 **Epitypification:** Investigated holotype of *Peziza pellita* (Cooke & Peck) T. Schumacher  
462 1979, New York Botanical Garden, specimen ID 00914741. We here designate the  
463 specimen of *Hoffmannoscypha pellita* (Cooke & Peck) Stielow, Hensel, Göker & Klenk,  
464 deposited at Botanische Staatssammlung Munich as epitype of the genus, specimen ID M-  
465 0156529.

466

## 467 **Key to the investigated genera**

468 1 Mature  
469 apothecia not lobed, usually less than 10 mm in diameter, brightly orange-red to orange-  
470 brown pigmented, growing superficially on or slightly sunken the substrate; ascospores  
471 less than 20  $\mu\text{m}$  in length; asci less than 200  $\mu\text{m}$  in length; occurring on burnt substrate or

- 472 decaying organic material *Tricharina*
- 473 1\* Mature apothecia lobed, usually more than 10 mm in diameter, either brightly  
474 orange-yellow or greyish to (light) brownish pigmented; ascospores more than 20 µm in  
475 length; asci more than 200 µm in length; neither occurring on burnt ground nor on  
476 decaying organic material.
- 477 2 Apothecia always with pronounced orange-yellow pigments; juvenile apothecia  
478 growing superficially on, at most slightly sunken into, the substrate
- 479 *Hoffmannoscypha*
- 480 2\* Apothecia never with orange, red or yellow pigments, but greyish to (light) brownish  
481 pigmented; juvenile apothecia strictly hypogeous, when mature growing superficially on  
482 the substrate *Geopora*

## 483 **Acknowledgments**

484

485 We kindly thank Dr. Ellen Bloch, New York Botanical Garden Herbarium, for the  
486 straightforward loans of *Sepultaria pellita* (*G. pellita*) and *Tricharina* spp. specimens, as  
487 well giving permission for nucleic acid extraction from specimens and digitalization of the  
488 drawings by G. Masee and F. Seaver (included in the supplementary material). Dr.  
489 Dagmar Triebel, Botanische Staatssammlung München, is kindly acknowledged for giving  
490 permission for nucleic acid extraction from *Tricharina* spp. specimens and the kind  
491 cooperation.

492

## 493 **References**

- 494 Agerer R, Ammirati J, Blanz P, Courtecuisse R, Desjardin DE, Gams W, Hallenberg N,  
495 Halling R, Hawksworth DL, Horak E, Korf RP, Mueller GM, Oberwinkler F, Rambold G,  
496 Summerbell RC, Triebel D, Watling R (2000) Open letter to the scientific community of  
497 mycologists: "Always deposit vouchers". *Mycorrhiza* 10:95–97.
- 498 Benkert D (1994) Contributions to the knowledge of bryophilous *Pezizales*. 1.  
499 *Lamprospora lubicensis*, new species from Northern Germany. *Z Mykol* 60:195–202.
- 500 Burdsall H (1968) A revision of the genus *Hydnocystis* (*Tuberales*) and of the hypogeous

501 species of *Geopora* (*Pezizales*). *Mycologia* 60:496–525.

502 Breitenbach J, Kränzlin F (1981) *Pilze der Schweiz, Band 1. Ascomyceten*. Mykologia,  
503 Luzern.

504 Castellano MA, Trappe JM, Maser Z, Maser C (1989) *Keys to spores of the genera of*  
505 *hypogeous fungi of North temperate forests with special reference to animal mycophagy*.  
506 Mad River Press, California.

507 Cooke MC, Peck CH (1872) *Pezizae americanae*. *Grevillea* 1:5–7.

508 Dennis RWG (1981) *British Ascomycetes*. J. Cramer, Vaduz.

509 Dougoud R (2007) Définition taxonomique et clé du genre *Geopora* Harkness. Ascofrance  
510 website, (<http://www.ascofrance.com/index.php>).

511 Erickson DL, Smith BD, Clarke AC, Sandweiss DH, Tuross N (2005) An Asian origin for a  
512 10.000-year-old domesticated plant in the Americas. *PNAS* 51:18315–18320

513 Ellis BM and Ellis JP (1998) *Microfungi on miscellaneous substrates: An identification*  
514 *handbook*. The Richmond Publishing and Co. Ltd., England.

515 Farris J (1972) Estimating phylogenetic trees from distance matrices. *Am Nat* 106:645–  
516 667.

517 Farris JS (1974) Formal Definitions of Paraphyly and Polyphyly. *Syst Zool* 23:548–554.

518 Farris JS (1979) The information content of the phylogenetic system. *Syst Zool* 28:483–  
519 519.

520 Felsenstein J (1981) Evolutionary trees from DNA sequences: a maximum likelihood  
521 approach. *J Mol Evol* 17:368–376.

522 Fitch WM (1971) Towards defining the course of evolution: minimal change for a specified  
523 tree topology. *Syst Zool* 20:406–416.

524 Gardes M, Bruns TD (1993) ITS primers with enhanced specificity for basidiomycetes  
525 application to the identification of mycorrhizae and rusts. *Mol Ecol* 2:113–118.

526 Guevara-Guerrero G, Stielow B, Tamm H, Cázares-Gonzales E, Göker M (2011) *Genea*  
527 *mexicana* and *Geopora toluicana*, new sequestrate *Pyronemataceae* from Mexico, and the  
528 phylogeny of *Geopora* s.l. reevaluated. *Mycol Prog* 11:711–724.

529 Hansen L, Knudsen H (2000) *Nordic Macromycetes Vol. 1 Ascomycetes*. Nordsvamp,

530 Copenhagen.

531 Hansen K, Laessoe T, Pfister DH (2001) Phylogenetics of the *Pezizaceae*, with an  
532 emphasis on *Peziza*. *Mycologia* 93:958–990.

533 Hansen K, Pfister DH (2006) Systematics of the *Pezizomycetes* the operculate  
534 discomycetes. *Mycologia* 98:1029–1040.

535 Hennig W (1965) Phylogenetic systematics. *Ann Rev Entomol* 10:97–116.

536 Hess PN, De Moraes Russo CA (2007) An empirical test of the midpoint rooting method.  
537 *Biol J Linn Soc* 92:669–674.

538 Kirk PM, Cannon PF, Minter DW, Stalpers JA (2008) Dictionary of the fungi. CABI, UK.

539 Krug JC, Khan RS (1991) *Dictyocoprotus*, a new genus of the *Pyronemataceae* with  
540 reticulated ascospores. *Mycologia* 83:103106.

541 Læssø T, Hansen K (2007) Truffle trouble: what happened to the *Tuberales*? *Mycol Res*  
542 111:1075–1099.

543 Lee C, Grasso C, Sharlow MF (2002) Multiple sequence alignment using partial order  
544 graphs. *Bioinformatics* 18:452–464.

545 Pattengale ND, Alipour M, Bininda-Emonds ORP, Moret BME, Stamatakis A (2009) How  
546 many bootstrap replicates are necessary? *Lect Notes Comput Sc* 5541:184–200

547 Pegler DN, Spooner BM, Young TWK (1993) British truffles, a revision of British  
548 hypogeous fungi. Royal Botanic Gardens, Kew.

549 Perry BA, Hansen K, Pfister DH (2007) A phylogenetic overview of the family  
550 *Pyronemataceae* (*Ascomycota*, *Pezizales*). *Mycol Res* 111:549–571.

551 Rice P, Longden I, Bleasby A (2000) EMBOSS: the European Molecular Biology Open  
552 Software Suite. *Trends Genet* 16:276–277.

553 Schumacher T (1979) Notes on taxonomy, ecology, and distribution of operculate  
554 discomycetes (*Pezizales*) from river banks in Norway. *Norw J Bot* 26:53–83.

555 Sbissi I, Neffati M, Boudabous A, Murat C, Gtari M (2010) Phylogenetic affiliation of the  
556 desert truffles *Picoa juniperi* and *Picoa lefebvrei*. *Anton Leeuw Int J G* 98:429–436.

557 Stamatakis A, Hoover P, Rougemont J (2008) A rapid bootstrap algorithm for the RAxML  
558 web servers. *Syst Biol* 75:758–771.

559 Spooner BM, Butterfill GB (1999) Coprophilous discomycetes from the azores. Kew  
560 Bulletin 54: 541–560.

561 Stielow B, Bratek Z, Orczán KA, Rudnoy S, Hensel G, Hoffmann P, Klenk H-P, Göker M  
562 (2011) Species delimitation in taxonomically difficult fungi: the case of *Hymenogaster*.  
563 PLoS ONE 6:e15614.

564 Stielow B, Bubner B, Hensel G, Münzenberger B, Hoffmann P, Klenk H-P, Göker M (2010)  
565 The neglected hypogeous fungus *Hydnotrya bailii* Soehner (1959) is a widespread sister  
566 taxon of *Hydnotrya tulasnei* (Berk.) Berk. and Broome (1846). Mycol Prog 9:195–203.

567 Swofford DL (2002) PAUP\*: Phylogenetic analysis using parsimony (\*and other methods),  
568 Version 4.0 b10. Sinauer Associates, Sunderland.

569 Tamm H, Poldmaa K, Kullman B (2010) Phylogenetic relationships in genus *Geopora*  
570 (*Pyronemataceae*, *Pezizales*). Mycol Prog 9:509–522.

571 Vralstad T, Myhre E, Schumacher T (2002) Molecular diversity and phylogenetic affinities  
572 of symbiotic root associated *Ascomycetes* of the *Heliotiales* in burnt and metal polluted  
573 habitats. New Phytol 155:131–148.

574 Weber NS, Trappe JM, Denison WC (1997) Studies on western American *Pezizales*.  
575 Collecting and describing Ascomata – macroscopic features. Mycotaxon 61: 153–176.

576 Wei J, Persoh D, Agerer R (2010) Four ectomycorrhizae of *Pyronemataceae*  
577 (*Pezizomycetes*) on chinese pine (*Pinus tabulaeformis*): a morphoanatomical and  
578 molecular phylogenetic analyses. Mycol Prog 9:267–280.

579 Wells VL, Kempton PE (1967) Studies on the fleshy fungi of Alaska. J Nat Prod 30:258–  
580 268.

581 White TJ, Bruns TD, Lee S, Taylor J (1990) Amplification and direct sequencing of fungal  
582 ribosomal RNA genes for phylogenetics. In: Innis MA, Gelfand DH, Sninsky JJ, White TJ  
583 (eds) PCR Protocols: a guide to methods and applications. Academic Press, New York, pp.  
584 315–322.

585 Wiley EO, Lieberman BS (2011) Phylogenetics. Theory and practice of phylogenetic  
586 systematics, Second Edition. Wiley-Blackwell, Hoboken.

587 Yao YJ, Spooner BM (1996) Notes on british species of *Geopora*. Mycol Res 100:72–74.

588 Yang CS, Korf RP (1985) A monograph of the genus *Tricharina* and of a new, segregate

589 genus, *Wilcoxina* (*Pezizales*). Mycotaxon 24:467–531.

590

591 **Tables**

592 **Table 1.** List of herbarium specimens and fungal strains used for molecular sequence and  
 593 comparative morphological analysis; specimens without Genbank accession number were  
 594 used for studying the morphology only. Species names given in square brackets are likely  
 595 to be misidentified (see the main text for details). Accession numbers marked with stars  
 596 are LSU sequences; the other represent the ITS. Abbreviation used in accordance with the  
 597 Index Herbariorum where applicable: CBS, Centraalbureau voor Schimmelcultures;  
 598 FUNGH, Fungarium Gunnar Hensel; FH, Harvard University Herbarium; ITCV, Instituto  
 599 Tecnológico de Ciudad Victoria; M, Botanische Staatssammlung München; NYBG, New  
 600 York Botanical Garden; S, Swedish Museum of Natural History.

Species	Collection data	Habitat details	Deposit	Accession number
<i>Geopora arenosa</i> (Fuckel) S. Ahmad	Stolberg, Hainfeld; Germany; 2006; G. Hensel & U. Täglich	Next to walking path on diabase	FUNGH: GH20090606	
<i>Geopora</i> cf. <i>cooperi</i> Harkn.	Santibanez de Valcorba, Spain	?	S: F23212	FR694203
<i>Geopora</i> cf. <i>cooperi</i> Harkn.	Gotland, Sweden	?	S. F48895	FR694202
<i>Geopora</i> cf. <i>cooperi</i> Harkn.	Sachsen-Anhalt, Germany; 2010; G. Hensel & U. Täglich	?	FUNGH: GH GH20100807	HQ184958, HQ184959
<i>Geopora pellita</i> (Cooke & Peck) T. Schumacher	USA; 1872; Cooke & Peck	?	NY: NYBG 00914741	
<i>Geopora pellita</i> (Cooke & Peck) T. Schumacher	Colorado, Peaceful Valley; USA; 1929; F. Seaver & P.F. Shope	Soil, in humus	NY: NYBG 159	JQ062972
<i>Geopora pellita</i> (Cooke & Peck) T. Schumacher	Colorado, Kingston Peak; USA; 1935; P. & V. Shope	Under aspens	NY: NYBG 301	JQ062973
[ <i>Geopora pellita</i> (Cooke & Peck) T. Schumacher]	Canada, Ontario, Toronto, High Park; 1906; J. H. Faull & J. H. Jackson	Sandy soil	NY: NYBG 228	JQ062974
<i>Geopora pellita</i> (Cooke & Peck)	Freienhufen; Germany; 2010; D.	On sandy mineral soil, embedded in	M: GH20100409	HQ913564, HQ993571*

T. Schumacher	Strobelt & G. Hensel	moss close to Pinus sp.		
<i>Geopora pellita</i> (Cooke & Peck) T. Schumacher	USA; 1969; D. Pfister	?	GH: DHP 297	DQ220343*
<i>Geopora sepulta</i> (Fr.) Korf & Burds.	Belzig; Germany; 2009; G. Hensel & U. Täglich	City centre Belzig, lawn on sand	FUNGH: GH20091122	
<i>Geopora toluquensis</i> Guevara, Göker & Stielow	Parque Nacional Nevado de Toluca, Mexico; 2009; G. Guevara	?	ITCV: ITCV 1081	HQ184960, HQ184961
<i>Tricharina gilva</i> (Boud.) Eckblad	Bavaria, Augsburg; Germany; 1878; Britzelmeyer	On calcareous soil between ash in a backyard	NY: Rehm. Ascom. 456	
[ <i>Tricharina gilva</i> (Boud.) Eckblad]	North Carolina, Durham; USA; 1968; F.A. Wolf	On flat sand in greenhouse, New York Bot. Gard. Herb. Specimen	NY: NYBG 1968	JQ824118
<i>Tricharina gilva</i> (Boud.) Eckblad	Berlin; Germany; 1885; P. Sydow	Lake shore	NY: NYBG 775	
<i>Tricharina gilva</i> (Boud.) Eckblad	USA; 1915; F. Seaver	?	NY: NYBG FJ1915	
<i>Tricharina gilva</i> (Boud.) Eckblad	USA; 1904	On sandy soil among moss	NY: NYBG 1905	
<i>Tricharina gilva</i> (Boud.) Eckblad	Hechendorf, Breitbrunn; Germany; 1963; Th. Kupka	Burned soil	M: M-0178315	JQ824122, JQ824123
<i>Tricharina gilva</i> (Boud.) Eckblad	Tirol, Nederjoch; Austria; 1948; M. Moser	Burned soil	M: M-0178314	
[ <i>Tricharina gilva</i> (Boud.) Eckblad]	Lappland, Kiruna; Sweden; 1965; A. Bresinsky	Burned soil	M: M-0178313	JQ824121
<i>Tricharina groenlandica</i> Chin S. Yang & Korf	Greenland; 1983; H. Dissing	?	CBS: CBS 237.85	JQ824125
<i>Tricharina hiemalis</i> Chin S. Yang & Korf	USA; H.K. Saksena	?	CBS: CBS 263.60	JQ824124
<i>Tricharina mikolae</i> Yang & Wilcox	Oregon; USA; 1982; Yang	Ascocarps collected on soil of a pot culture of red pine seedlings in greenhouse, Syracuse	NY: NYBG 914971	

Tricharina ochroleuca (Bers.) Eckblad	Norway; 1979; S. Silvertsen & H. Dissing	?	CBS: CBS 238.85	JQ824126
[ <i>Tricharina praecox</i> (Karst.) Boudier]	Holzkirchen, MTB 8136; Germany; 1980; A. Einhellinger	Burned soil	M: M-0178317	JQ824119
[ <i>Tricharina praecox</i> (Karst.) Boudier]	Tirol, Nederjoch; Austria; 1948; M. Moser	Burned soil	M: M-0178316	JQ824120

601

602

### 603 **Figures**

604 **Fig. 1.** Phylogenetic tree inferred under the maximum-likelihood (ML) criterion from the ITS  
605 rDNA alignment. Numbers on the branches represent support values from 1000 bootstrap  
606 replicates under the ML (left) and the maximum-parsimony criterion (right) if at least 60%.  
607 Branches are scaled in terms of the expected number of substitutions per sites. Leaf  
608 names are from their original annotations; corrections, if any, can be inferred from the  
609 group names on the right side of the vertical bars. The sequences newly generated in the  
610 course of our study are listed in Table 1. See the electronic supplementary material for the  
611 complete tree.

612 **Fig. 2.** Phylogenetic tree inferred under the ML criterion from the 28S (LSU) rDNA  
613 alignment. Numbers on the branches represent support values from 1000 bootstrap  
614 replicates under the ML (left) and the maximum-parsimony criterion (right) if at least 60%.  
615 Branches are scaled in terms of expected number of substitutions per sites. Leaf names  
616 are from their original annotations; corrections, if any, can be inferred from the group  
617 names on the right side of the vertical bars. The sequences newly generated in the course  
618 of our study are listed in Table 1. See the electronic supplementary material for the  
619 complete tree.

620 **Fig. 3.** 1: Mature yellow-orange apothecia of *H. pellita* (M: M-0156529). 2: Monoseriate  
621 asci arranged in parallel as well as the orange-yellow pigmented paraphyses causing the  
622 characteristic colour of the excipulum; bar = 10 µm. 3: Juvenile apothecium of *H. pellita*. 4:  
623 Cross-section through the excipulum of *H. pellita*; bar = 1000 µm. 5: Apothecia of *Geopora  
624 sepulta* (FUNGH: GH20091122). 6: Cells and setae of the ectal excipulum of *H. pellita*, bar  
625 = 10 µm. 7: FESEM image of a single juvenile spore in a broken ascus; bar = 10 µm.



626 **Fig. 4.** 1: FESEM image of a mature, finely warted ascospore of *H. pellita*; bar = 10  $\mu\text{m}$ . 2:  
627 FESEM image of juvenile *H. pellita* spores (smooth surface) and a single mature spore  
628 (warted surface; central position); bar = 10  $\mu\text{m}$ .

629