Psora taurensis (Psoraceae, Lecanorales), a new lichen species from Turkey

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Abstract
Herein we describe the new species, Psora taurensis, from two localities in the Taurus Mountains in Turkey at ca. 1000 m altitude. Investigations of anatomy, secondary chemistry and DNA sequences (ITS and mtSSU) of P. taurensis and presumed close relatives suggest that P. taurensis is a distinct evolutionary lineage with P. tenuifolia as its sister, although it is morphologically more similar to P. russellii and P. vallesiaca.

Key words
Anatomy, DNA, phylogeny, Lecanorales, lichenized ascomycetes, taxonomy, TLC, Turkey

Introduction
After publication of our recent paper on Psora altotibetica Timdal et al. (Timdal et al. 2016; see this paper also for a general background on the genus), we sequenced an unidentified specimen of Psora collected by one of us (ET) in the Taurus Mountains in Turkey in 1994. Based on morphology and secondary chemistry, it had been suspected to be related to the North American P. russellii (Tuck.) A.Schneider, but DNA sequence data from the internal transcribed spacer region (ITS) suggested a closer relation to the P. altotibetica–tenuifolia–vallesiaca clade, recovered by Timdal et al. (2016: fig. 1). Independently, AMK had sequenced the ITS region from a second spec-
imen, which had been collected some 150 km east-southeast in the same mountains by MGH in 2012. A preliminary comparison of the two DNA sequences suggested we had collected the same species. The aim of the present study was to further investigate the relatedness and shared distinctness of these two specimens in a broader phylogenetic context that includes presumed closely related species and an additional and more conserved genetic region, the mitochondrial ribosomal small subunit (mtSSU).

**Material and methods**

**The specimens**

This study is based on: (1) the two specimens of *Psora taurensis* referred to above, (2) the specimens with DNA sequence data in Timdal et al. (2016), (3) five additional *Psora* specimens deposited in O and sequenced for this work, and (4) two additional specimen of *Psora testacea* Hoffm., from which ITS sequences were available from GenBank. DNA sequence data for *P. elenkinii* Rass. and *P. pseudorussellii* Timdal is herein provided for the first time. Voucher data, major lichen substances, and GenBank accession numbers for these 42 specimens are given in Table 1. With the exception of three specimens, *P. himalayana* (C.Bab.) Timdal 1, *P. testacea* 2, and *P. testacea* 3, we have examined all collections listed in Table 1 by morphology and secondary chemistry during this project or previously.

**Anatomy**

Microscope sections were cut on a freezing microtome at 16 μm and mounted in water, 10% KOH (K), lactophenol cotton blue, a modified Lugol’s solution in which water was replaced by 50% lactic acid, as well as 25% sulphuric acid, and chlor-zinc-iodine. Amyloid reactions were observed in the modified Lugol’s solution after pretreatment in K. Chlor-zinc-iodine was used to locate remnants of algae in the cortex, and polarized light was used to locate crystals of secondary metabolites and calcium oxalate. Calcium oxalate was identified by adding 25% sulphuric acid to the section; the oxalate crystals dissolve and needle shaped crystals of calcium sulphate precipitate. Ascospore measurements are given as $X \pm 1.5 \times SD$ rounded to 0.5 μm, where $X$ is the arithmetic mean and SD the standard deviation.

**Secondary chemistry**

Thin-layer chromatography (TLC) was performed in accordance with the methods of Culberson (1972), modified by Menlove (1974) and Culberson and Johnson (1982). All specimens were examined by TLC.
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<table>
<thead>
<tr>
<th>Taxon, specimen</th>
<th>Voucher information</th>
<th>Major lichen substances</th>
<th>GenBank accession number</th>
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**DNA extraction, PCR and sequencing**

We performed DNA extraction, PCR amplification, PCR purification, and cycle sequencing as described by Bendiksby and Timdal (2013). DNA was extracted from apothecia of 7 specimens (Table 1; GenBank Accession Numbers KY426119–KY426132). All DNA isolates produced for the present study are deposited in the DNA collection at Natural History Museum, University of Oslo or Molecular Biology Lab of Erciyes University, Faculty of Science (only *P. taurensis* 1). We amplified and sequenced the ITS and the mtSSU using the primer pairs ITS5/ITS4 (White et al. 1990) and mtSSU1/mitSSU3R (Zoller et al. 1999), respectively.

**Data analyses**

Sequences were assembled and edited using SEQUENCHER v.4.1.4 (Gene Codes Corporation, Ann Arbor, Michigan, U.S.A.). Alignments were established in BIOEDIT IT 7.2.3 (Hall 1999) using the “ClustalW/Multiple alignment” option with subsequent manual adjustments. We analysed and summarized the data with parsimony and Bayesian phylogenetic methods, including model testing, as described in Bendiksby et al. (2015). As configuration settings in PartitionFinder v.1.1.1 (Lanfear et al. 2012),
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Results

Anatomy

The following key characters for including *P. taurensis* in *Psora* were observed in the new species: the upper cortex contained remnants of algae throughout both the lower stainable layer and the upper epinecral layer (‘Scheinrindentyp’ of Poelt 1958); the hypothecium contained calcium oxalate crystals; the epihymenium contained orange crystals which dissolved in K with a purple diffusion (assumed to be anthraquinones); and the ascus contained a well-developed, amyloid tholus with a central, deeper amyloid tube structure (*Porpidia*-type).

The following species level characters were observed in *P. taurensis*: Upper cortex composed of thick-walled hyphae with angular to rounded lumina; crystals of norstictic acid and calcium oxalate present in medulla; no crystals in upper cortex; poorly developed lower cortex; ascospores 11–16 × 5.5–7 μm.

Secondary chemistry

The results of the TLC examinations are given in Table 1. The two specimens of *P. taurensis* contained norstictic acid.

Alignments and phylogenetic analyses

Altogether 14 DNA sequences were generated from 7 specimens for the present study (7 ITS and 7 mtSSU; Table 1). The ITS end-trimmed alignment of 41 accessions was 573 basepairs long and contained 236 parsimony-informative characters. The corresponding numbers for the mtSSU matrix of 32 accessions was 794 and 22, respectively. For ITS1, ITS2 and mtSSU, the HKY+I+G model gave the best fit, whereas K80+I+G had the estimated best fit for 5.8S. Tree-topologies from both parsimony and Bayesian analyses of ITS vs mtSSU alignments were congruent but resolved to various extents (not shown). Final analyses were therefore performed on a concatenated dataset of 1367 bp. In the Bayesian analysis, the average standard deviation of split frequencies (ASDSF) had fallen to 0.004765 at termination (five million generations) and the first
1500 saved trees (i.e. 30%) were discarded as burn-in, ensuring that only generations with ASDSF below 0.01 were kept for summarizing. The Bayesian 50% majority rule consensus tree is presented as an unrooted tree with both Bayesian posterior probability (PP > 0.9) and parsimony jackknifing (JK > 90) branch support superimposed (Fig. 1). Multiple accessions of all species group to the respective species with high support. The single exception is a clade consisting of two accessions of *P. himalayana* that is nested within the *P. vallesiaca–himalayana* clade, grouping with high support with *P. vallesiaca* (Schaer.) Timdal 4−7 to the exclusion of *P. vallesiaca* 1−3. *Psora tenuifolia* Timdal is strongly supported as sister to the new species, *P. taurensis*. The *P. tenuifolia–taurensis* clade is sister to *P. altotibetica*, which in turn is sister to the *P. vallesiaca–himalayana* clade. A clade consisting of *P. hyporubescens* Timdal and *P. pacifica* Timdal is also strongly supported. The sister relationship between *P. californica* Timdal and *P. indigirkae* Timdal & Zhurb. received low support (JK=57; PP=0.51). *Psora globifera* (Ach.) A.Massal. is supported as sister to the *P. altotibetica–tenuifolia–taurensis–vallesiaca–himalayana* clade, but this relationship was not supported by parsimony jackknifing. The same accounts for the grouping of the *P. hyporubescens–pacifica* clade with the aforementioned multispecies clade. Apart from this, the molecular data support no further inter-species relationships.

**Discussion**

Our molecular data strongly support *Psora taurensis* as a distinct evolutionary unit and, given the current taxon sampling, *P. tenuifolia* is its sister (Fig. 1). *Psora tenuifolia* differs in having thinner, generally more ascending squamules containing zeorin (and often norstictic acid, as *P. taurensis*) and in having a well-developed lower cortex composed of mainly anticlinally oriented hyphae which are densely covered by calcium oxalate crystals (cf. Timdal 1986). *Psora tenuifolia* is known from winter-cold, arid sites in Alaska and arctic Canada (Timdal 1986), Yakutia (Zhurbenko 2003), and the Great Himalayas (Timdal et al. 2016).

*Psora altotibetica*, which falls out as sister to the *P. tenuifolia–taurensis* clade (Fig. 1), differs in having strictly adnate squamules which are more evenly covered by pruina and in containing gyrophoric acid (cf. Timdal et al. 2016). *Psora altotibetica* is known only from the Great Himalayas between 4230 and 5000 m altitude (Timdal et al. 2016).

Outside that clade is the complex of *P. vallesiaca*, which consists of several strongly supported subclades with varying branch lengths and with *P. himalayana* embedded (Fig. 1). *Psora himalayana* and *P. vallesiaca* are distinguished mainly on the thallus chemistry, i.e. no lichen substances in the former and norstictic acid in the latter. Timdal et al. (2016) indicated that they may be conspecific, based on only the ITS sequence of a single specimen of *P. himalayana* (from Yakutia), which rendered it as nested within a clade of seven accessions of *P. vallesiaca*. In the present study, both ITS and mtSSU sequences of a second specimen of *P. himalayana* (from Yukon) is provided. The two specimens of *P. himalayana* group with moderate support (JK=81;
Figure 1. Bayesian 50% majority rule consensus tree based on a concatenated alignment of ITS and mtSSU sequences of 42 accessions of 17 *Psora* species (see Table 1). Parsimony jackknife support values above 90% are shown below branches and Bayesian posterior probabilities above 0.9 above. The curly branch leading to *P. testacea* has been shortened to reduce the size of a broad figure.

PP=1), and the species remains nested in the *P. vallesiaca* complex (Fig. 1). A broader sampling is needed, however, especially from the Himalayas, before *P. himalayana* may be synonymised with *P. vallesiaca*. The complex differs morphologically from *P.*
taurensis mainly in forming more distinctly white-edged squamules with a more upturned margin.

*Psora elenkinii* was synonymized with *P. himalayana* by Timdal (1986). However, the ITS and mtSSU sequences provided here, from a morphologically typical specimen from Yakutia, shows that the species falls outside the *P. vallesiaca–himalayana* clade (Fig. 1). The species is hence accepted here.

The North American desert lichen *P. russellii* differs morphologically from *P. taurensis* mainly in forming closely adnate squamules with a more down-turned margin and often with a regular, central depression, and in having medium brown apothecia. The species contains norstictic acid both in the upper cortex and in the medulla and there is also sometimes a trace of gyrophoric acid (Timdal 1986). The three sequenced specimens of *P. russellii* group with high support and are not closely related to *P. taurensis* (Fig. 1).

*Psora pseudorussellii* differs from *P. russellii* mainly in lacking lichen substances and in forming smaller, more elongated squamules without a central depression (Timdal 1986). It differs from *P. taurensis* mainly in lacking lichen substances and in the medium brown colour of the apothecia. This essentially eastern North American species was reported new to Europe from Greece (Crete) by Grube et al. (2001). We have examined additional European specimens from Greece (Crete and Samos), Italy (Calabria), and Spain (Granada, Madrid, and Soria) (Timdal unpubl.), and here provide DNA sequences from the species for the first time. Phylogenetically it is not closely related to *P. taurensis* (Fig. 1).

*Psora peninsularis* Timdal, occurring in coastal scrubs and Sonoran desert in southern California and Baja California, differs morphologically mainly in forming castaneous brown, shiny, epruinose squamules. It contains norstictic acid in the medulla (Timdal 2002). Phylogenetically it is not closely related to *P. taurensis* (Fig. 1).

Two additional species are relevant for the discussion of the taxonomy of *P. taurensis*: *P. gresinonis* B.de Lesd. and *P. subrubiformis* (Vain.) Dzhur. Lack of sequence data makes this discussion purely morphological. We know the former species from c. 15 localities in Mediterranean Europe and Central Asia and the latter only from the type collection from Turkmenistan (Timdal 1984, Timdal unpubl.). *Psora gresinonis*, which often contains norstictic acid like *P. taurensis*, differs in forming smaller, thinner, more rounded and concave squamules with a non-pruinose, brown or sometimes greyish margin. The holotype of *Psora subrubiformis* lacks lichen substances and differs from *P. taurensis* in having persistently plane to only weakly convex, densely white pruinose apothecia (cf. Timdal 1984) and a thallus morphology resembling that of the *P. vallesiaca* complex. Except for the more plane apothecia, there are few arguments for regarding it as a distinct species within the the *P. vallesiaca* complex.

Hence, since *P. taurensis* is now known from two localities and its distinctness is supported by various data, we hereby describe it as a new species.
Psora taurensis (Psoraceae, Lecanorales), a new lichen species from Turkey

**Taxonomy**

*Psora taurensis* Timdal, Bendiksby, Kahraman & Halıcı, sp. nov.
Mycobank: MB 820063
Fig. 2

**Diagnosis.** Morphologically most similar to *Psora russellii*, but squamules more ascending and lacking a central depression, and apothecia brownish black. Phylogenetic sister species of *P. tenuifolia*, but having a thicker, more adnate thallus with a poorly developed lower cortex and lacking zeorin.

**TYPE.** TURKEY. Mersin: Gülnar-Silifke Highway, exit of Kayrak, 36°21'24.5"N, 33°33'08.8"E, 1000-1020 m alt., on soil on calcareous bedrock, 12 Apr 2012, M.G. Halıcı (holotype: ERCH-AMEKA 0.018!)

**Description.** Thallus squamulose; squamules up to 8 mm wide, rounded, adnate with ascending margin to imbricate, becoming deeply lobed, concave; upper surface medium brown, dull, pruinose in the outer part of the lobes, with regular fissures in the cortex; margin first concolorous with upper side, soon becoming white by pruina, straight or somewhat up-turned; upper cortex up to 130 μm thick, including an up to 20 μm thick epinecral layer, composed of thick-walled hyphae with angular to rounded lumina, not containing crystals, containing remnants of algae throughout (chlor-zinc-iodine!); algal layer continuous, 30–45 μm thick; medulla not amyloid, containing lichen substances (K+ yellow, red crystals precipitating) and calcium oxalate; lower cortex poorly developed; lower surface white to pale brown. Apothecia up to 1.5 mm diam., laminal or submarginal on the squamules, weakly convex and indistinctly marginate when young, soon becoming strongly convex and immarginate, brownish black, epruinose. Proper exciple yellowish brown in the rim, colourless in inner part, lacking crystals, composed of radiating, thick-walled hyphae; hypothecium colourless in lower part, pale brown in upper part, containing crystals of calcium oxalate; epihymenium yellowish brown, containing orange crystals dissolving in K, K+ purple; hymenium 70–90 μm high, colourless, amyloid. Paraphyses straight, thin-walled, moderately conglutinated, sparingly branched and anastomizing, with a slightly swollen apical cell. Ascus clavate, with a well-developed, amyloid tholus containing a deeper amyloid tube, lacking an ocular chamber (*Porpidia*-type); ascospores ellipsoid, non-septate, hyaline, 11–16 × 5.5–7 μm (n = 20). Conidiomata unknown.

**Chemistry.** Norstictic acid (by TLC); medulla K+ yellow turning red, C–, KC–, P+ orange.

**Habitat and distribution.** The species is known from two localities in Turkey, both at c. 1000 m altitude. Both sites are in areas with Mediterranean climate. The holotype was collected in a rocky area with scrub vegetation derived by forest degradation; the paratype grew in an open pasture. Both specimens were terricolous, the holotype grew on soil over limestone.
Figure 2. *Psora taurensis*, habitus. A, part of holotype; B, part of paratype. Scale bar = 2 mm.

**Etymology.** The name refers to its occurrence in the Taurus Mountains.

**Other specimen examined.** Turkey. Antalya: along the road a few km SE of Gündoğmuş, 36°48.1′N, 32°00.3′E, 1000 m alt., on soil in open pasture, 24 Apr 1994, E.Timdal 7908 (O L-203076, paratype).
Psora taurensis (Psoraceae, Lecanorales), a new lichen species from Turkey

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References


Resolving the genus *Phaeographina* Müll. Arg. in China

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Abstract

As part of ongoing studies of the lichen family Graphidaceae in China, the status of all taxa traditionally assigned to the genus *Phaeographina* reported from China is resolved in the present paper. Five new combinations are proposed: *Phaeographis pleiospora* (Zahlbr.) Z.F. Jia & Lücking, **comb. nov.**, *Platygramme elaeoplaca* (Zahlbr.) Z.F. Jia & Lücking, **comb. nov.**, *Platythecium maximum* (Groenh.) Z.F. Jia & Lücking, **comb. nov.**, *P. pyrrhochroa* (Mont. & Bosch) Z.F. Jia & Lücking, **comb. nov.**, *P. ptyrrochroa* (Mont. & Bosch) Z.F. Jia & Lücking, **comb. nov.**. Six new synonyms are established: *Phaeographina callospora* Zahlbr. [= *Diorygma hieroglyphicum* (Pers.) Staiger & Kalb], *P. fukiensis* Zahlbr. [= *Pallidogramme chrysenteron* (Mont.) Staiger, Kalb & Lücking], *P. fukiensis* var. *substriata* Zahlbr. [= *Pallidogramme chrysenteron* (Mont.) Staiger, Kalb & Lücking], *P. granulans* Zahlbr. [= *Platygramme platyloma* (Müll. Arg.) M. Nakan. & Kashiw.], *P. pluvisilvarum* Zahlbr. [= *Graphis alpestris* (Zahlbr.) Staiger], and *P. valida* Zahlbr. [= *Thecographa prosiliens* (Mont. & Bosch) A. Massal.]. Two additional synonyms are reported: *Phaeographina subrigida* (Nyl.) Zahlbr. is synonymized under *Platygramme platyloma* (Müll. Arg.) M. Nakan. & Kashiw., and *Platythecium dimorphodes* (Nyl.) Staiger under *Platythecium pyrrhochroa* (Mont. & Bosch) Z.F. Jia & Lücking.

Key words

Lichen, taxonomy, Graphidaceae, Ostropales
Introduction

The lichen genus *Phaeographina* Müll. Arg. is an artificial, ascospore-based genus in *Graphidaceae* Dumort. (Müller 1882), in use until after the turn of the millenium. A revised generic concept in the family, based on Staiger (2002) and further elaborated using molecular phylogenetic approaches (Staiger et al. 2006; Rivas Plata et al. 2012, 2013; Lücking et al. 2013; Lumbsch et al. 2014), resulted in subsuming the name *Phaeographina* into synonymy with *Thecographa* A. Massal., with the single species, *Thecographa prosiliens* (Mont. & Bosch) A. Massal. (Lücking et al. 2007; Lücking and Rivas Plata 2008). Following this revised generic concept, most of the species at some point included in *Phaeographina*, representing a total of 265 names, have been redispositioned into other genera based on ascoma morphology and anatomy, such as *Pallidogramme* Staiger, Kalb & Lücking, *Phaeographis* Müll. Arg., *Platygramme* Fée, *Thecaria* Fée, and the aforementioned *Thecographa* (Staiger 2002; Archer 2006, 2009).

As part of a revision of Chinese *Graphidaceae*, we attempted to resolve the status of all species reported under the name *Phaeographina* from China (Wei 1991; Aptroot and Seaward 1999; Aptroot and Sipman 2001, Aptroot and Sparrius 2003). Twenty-one species were reported under the name *Phaeographina*, which are here presented in the form of an annotated checklist and listed under or transferred to the corresponding genera, namely *Chapsa* A. Massal., *Diorygma* Eschw., *Glyphis* Ach., *Graphis* Adans., *Pallidogramme*, *Phaeographis*, *Platygramme*, *Platythecium* Staiger, *Sarcographina* Müll. Arg., *Sarcographa* Fée, *Thecographa*, and *Thecaria*.

Materials and methods

Type specimens and other material investigated for this study are deposited in BRSL, H, HMAS-L, KUN, LCU-L, PC, TNS, UPS, and W. For several of the names included here that have already been treated by Kalb et al. (2004), Staiger (2002), and Lücking et al. (2009), we do not provide full synonymies and type specimen citations but give the corresponding reference. A dissecting microscope (Olympus SZX12) and a light microscope (Olympus BX51 and Nikon Eclipse-55i) were used for the morphological and anatomical studies. Measurements were taken from manual cross sections of fruit bodies in water. Amyloidity of the ascospores was tested using Lugol’s solution. In cases where the chemistry of the type material had not been studied previously, lichen substances were identified by thin-layer chromatography (Culberson and Kristinsson 1970; Culberson and Kristinsson 1972; White and James 1985).

Taxonomy

Annotated checklist of Chinese species previously reported under the name *Phaeographina* Müll. Arg.

Figure 1A–B

Feddes Repert. 31: 220, 1933. Type: Taiwan, *Faurie* 118 (W, holotype!).


Based on the *Diorygma*-like thallus and ascomata, this taxon belongs in *Diorygma*. It has a clear hymenium, single-spored asci, hyaline, muriform ascospores 90–100 × 29–32 μm (Zahlbruckner 1933), and stictic acid. These characters, as well as the immersed lirellae with a split between excipulum and thalline margin, agree with *D. hieroglyphicum* (Pers.) Staiger & Kalb and hence, we propose this name as a synonym of the latter. The ascospores in this material are partially old and become brownish, which is the reason why it was described in *Phaeographina*. *Phaeographina callospora* Zahlbr. should not be confused with *Graphis collospora* Vain. [≡ *Graphina collospora* (Vain.) Zahlbr.], which is also a synonym of *D. hieroglyphicum* (Kalb et al. 2004). *Diorygma hieroglyphicum* is a corticolous species reported from Taiwan (type locality of *P. callospora*, Zahlbruckner 1933, 1940; Lamb 1963; Wang-Yang and Lai 1973 and 1976), Fujian, Hainan and Yunnan (Meng and Wei 2008; Wei et al. 2013; Jia and Wei 2016).

2. *Phaeographina chlorocarpoides* (Nyl.) Zahlbr.


≡ *Pallidogramme chlorocarpoides* (Nyl.) Staiger, Kalb & Lücking in Lücking et al., Fieldiana, Bot. 38: 9, 2008.

Following Lücking et al. (2008), this taxon was transferred to the genus *Pallidogramme*. It is a corticolous species reported from Guangdong, Guangxi, Hunan, Hainan and Hong Kong (Krempehlhuber 1873; Zahlbruckner 1930; Thrower 1988; Miao et al. 2007; Wei et al. 2013; Jia and Wei 2016).


Hedwigia 30: 52, 1891 [as ‘chrysentera’].

≡ *Pallidogramme chrysenteron* (Mont.) Staiger, Kalb & Lücking in Lücking et al., Fieldiana, Bot. 38: 9, 2008; *Hemithecium chrysenteron* (Mont.) Trevis., Spighe Paglie: 13, 1853.

Following Lücking et al. (2008), this taxon belongs in *Pallidogramme*. It is a corticolous species reported from Fujian, Guangdong, Guangxi, Yunnan, Hunan, Hainan and
Taiwan (Zahlbruckner 1930, 1932; Miao et al. 2007; Wei et al. 2013; Jia and Wei 2016; Aptroot and Sparrius 2003).

_Pallidogramme_ is listed as invalid name in _Index Fungorum_ (accessed 18 January 2017), presumably based on ICN Art. 41.5 (incomplete citation of the replaced synonym). However, the replaced synonym is not _Hemithecium_ subgen. _Leucogramma_ Staiger (Staiger 2002: 277), but _Leucogramma_ A. Massal. (1860: 273, 320), an illegitimate later homonym of _Leucogramma_ Meyer (1825: 331). Both _Leucogramma_ A. Massal. and its type species, _L. chrysenteron_ (Mont.) A. Massal. were established on the same page (1860: 320), and since in the protologue of _Pallidogramme_ the full reference is given for the type species (designated as holotype), _L. chrysenteron_ (Mont.) A. Massal., this fulfills the requirements of ICN Art. 41 for valid publication of the genus name.

4. _Phaeographina elaeoplaca_ Zahlbr.


≡ _Platygramme elaeoplaca_ (Zahlbr.) Z.F. Jia & Lücking, **comb. nov.** (see below).
This material is characterized by lirellae with thick, apically carbonized, exposed labia and a closed disc, an inspersed hymenium, single-spored asci producing brown ascospores about 80–110 × 20–30 μm, and the absence of secondary substances. Ascoma morphology and the brown ascospores place this material in the genus *Platygramme*. Several species have been described which belong in *Platygramme* and agree with *Phaeographina elaeoplaca* in ascoma morphology (thick labia with closed disc, single-spored asci, absence of secondary substances: *Graphis commutabilis* Kremp. (Krempelhuber 1875) [≡ *Platygramme commutabilis* (Kremp.) A. W. Archer] has ascomata with whitish thalline cover and ascospores 80–110 μm long; *Phaeographina platyloma* Müll. Arg. (Müller 1882) [≡ *Platygramme platyloma* (Müll. Arg.) M. Nakan. & Kashiw.; syn.: *Platygramme impudica* (A. W. Archer) A. W. Archer] has ascomata with distinct, lateral thalline cover and ascospores 120–180 μm long; and *Graphis subrigida* Nyl. (Nylander & Crombie, 1883) [≡ *Phaeographina subrigida* (Nyl.) Zahlbr.] has ascomata with exposed labia and ascospores 110–150 μm long. On the annotation label of the lectotype of *Phaeographina elaeoplaca*, Nakanishi noted that this name should be a synonym of *Ph. subrigida*; however, the ascospores of the latter are much longer. Archer (2009) described *Platygramme commutabilis*, which agrees with *Phaeographina elaeocarpa* in ascospore size, as having exposed labia, which would make *Ph. elaeocarpa* a synonym of *Pl. commutabilis*. However, this observation is in conflict with the type of *Pl. commutabilis*, which clearly shows the labia covered by a whitish thallus layer. Thus, we conclude that *Ph. elaeocarpa* is neither conspecific with *Ph. subrigida* nor with *Pl. commutabilis* but represents a distinct taxon, recombined below as *Platygramme elaeoplaca* (Zahlbr.) Z.F. Jia & Lücking. We suspect that the Australian material identified by Archer (2009) as *Pl. commutabilis* in reality represents *Pl. elaeocarpa*. Based on ascospore size, *Ph. subrigida* is to be placed as an additional synonym under *Pl. platyloma*.

*Platygramme elaeocarpa* is a corticolous species reported from Fujian (type locality; Zahlbruckner 1932, 1934, 1940).


Figure 2A–B


Due to the characteristics of thallus and ascomata, this taxon belongs in *Pallidogramme*. According to the annotation label by Nakanishi, it is similar to *Pallidogramme chlorocarpoides* (Nyl.) Staiger, Kalb & Lücking, but differs by having smaller ascospores (36–48 × 12–16 μm, Zahlbruckner 1930) than the latter (*P. chlorocarpoides*: 55–110 × 20–39 μm, Staiger 2002). It therefore agrees perfectly with *P. chrysenteron* (ascospores 33–60 × 10–15 μm, Staiger 2002) and represents another synonym of the latter. The distribution of this species in China is treated above.
Figure 2. A–B Phaeographina fukiensis Zahlbr. (Chung 399a) C–D Graphis glyphiza Nyl. (Nylander 6989). Scale bars = 1 mm.

Phaeographina fukiensis var. substriata Zahlbr.

Ann. Mycol. 30: 432, 1932. Type: Fujian: Chung 597 (W, holotype!).

= Pallidogramme chrysenteron (Mont.) Staiger, Kalb & Lücking, in Lücking et al., Fieldiana, Bot. 38: 9, 2008.

According to Zahlbruckner, this taxon differs from the nominal variety by its striate labia and smaller ascospores, which is odd considering that the type of the nominal variety has distinctly striate labia. The ascospores fall within the lower range of variation of Pallidogramme chrysenteron and hence this is considered another synonym of that species.

6. Phaeographina glyphiza (Nyl.) Zahlbr

Figure 2, C–D

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Following Singh and Sinha (2010), this taxon belongs in *Sarcographa* as *S. glyphiza* (Nyl.) Kr.R. Singh & G.P. Sinha. It is a corticolous species reported from Hong Kong (type location; Nylander 1863, Leighton 1869, Hue 1891 as *Graphis glyphiza*; Zahlbruckner 1923 as *Phaeographis glyphiza*; Zahlbruckner 1930, 1932, Aptroot 1999 as *Phaeographina glyphiza*). Seaward and Aptroot (2005) reported this taxon with the current name *Sarcographa glyphiza* from Hong Kong.


Figure 3, A–B


Based on the characteristics of the ascomata, excipulum and ascospores, which are about 140–160 μm long, this taxon is a further synonym of *Platygramme platyloma* (Müll. Arg.) M. Nakan. & Kashiw. (see discussion under *Pl. elaeocarpa* above), as already annotated by Nakanishi on the holotype. *Platygramme platyloma* is a corticolous species reported from Fujian (Type locality of *Phaeographina granulans*, Zahlbruckner 1932, 1934, 1940; Lamb 1963; Jia and Kalb 2013).

8. *Phaeographina heterospora* (Nyl.) Zahlbr.

≡ *Sarcographina heterospora* (Nyl.) Z.F. Jia & Lücking, comb. nov. (see below)

Staiger (2002), placed this taxon in *Gymnographa* as *G. heterospora* (Nyl.) Staiger. Restudy of the type material confirmed that this taxon is very closely related to *Sarcographina cyclospora* Müll. Arg., which was recently shown to belong in tribe *Ocellularieae* within *Graphidaceae* (Kraichak et al. 2014). Hence, *Phaeographina heterospora* is here also recombined in that genus, as *Sarcographina heterospora* (Nyl.) Z.F. Jia & Lücking. *Sarcographina heterospora* is a corticolous species reported from Taiwan (Aptroot and Sparrius 2003).

Figure 3C–D


≡ *Phaeographis pleiospora* (Zahlbr.) Z.F. Jia & Lücking, *comb. nov.* (see below).

This material belongs in a difficult group of taxa characterized by rounded to shortly elongate, erumpent to prominent ascomata with a usually fissured, thalline margin partially exposing the (dark) brown disc, and with muriform ascospores. Staiger (2002) distinguished two species in this group, viz. *Phaeographis kalbii* Staiger, with norstic-tic acid, 1–4-spored asci, and ascospores 95–125 × 20–30 μm, and *Ph. lecanographa* (Nyl.) Staiger, with virensic acid, 1–4-spored asci, and ascospores 85–150 × 25–50 μm. Lücking (2015) later found an earlier name for the latter species with virensic acid, viz. *Thelotrema spondaicum* Nyl., now *Phaeographis spondaica* (Nyl.) Lücking. *Phaeographis kalbii* was synonymized with *Ph. atromaculata* (A.W. Archer) A.W. Archer (Archer 2006), but Lendemer and Harris (2014) proposed to keep the two species separate and added a further species to this group, *Ph. oricola* Lendemer & R.C. Harris, which dif-
fers mainly in its single-spored asci. The type material of *Ph. lecanographa* var. *pleiospora* contains norstictic acid (as already noticed by Zahlbruckner) and would belong in that latter complex, but the ascospores are consistently smaller (60–80 × 20–30 μm). We therefore accept this as a distinct species, *Ph. pleiospora* (Zahlbr.) Z.F. Jia & Lücking. It is a corticolous species reported from Fujian (Type locality. Zahlbruckner 1930, 1933, 1934 and 1940).

10. *Phaeographina macrospora* (Zahlbr.) Nakanishi


Following Aptroot (2004), this taxon is a synonym of *Thecaria montagnei* (Bosch) Staiger. *Thecaria montagnei* is a corticolous species reported from Taiwan (Type locality of *Phaeographina macrospora*, Zahlbruckner 1933; Wang-Yang and Lai 1973) and Hainan (Wei et al. 2013).


in Blumea, Suppl. 5 (H. J. Lam Jubilee Vol.). 107, 1958. Type: Indonesia, Sumatra, *Groenhart 9453* (BO, holotype, not seen, but original image in protologue seen).

≡ *Platythecium maximum* (Groenh.) Z.F. Jia & Lücking, comb. nov. (see below).

According to the original description and excellent illustrations, this is a species of *Platythecium*, agreeing with *P. serpentinellum* (Nyl.) Staiger in the brown ascospores and with *P. allosporellum* (Nyl.) Staiger in the carbonized hypothecium. Groenhart (1958) beautifully illustrates the enormous extension of the thallus of the specimen in the field, with over 100 cm covering a large boulder. This is consistent with our observations that *Platythecium* species often form large thalli on rocks near streams. Since the type material is different from any known in the genus, it is here recombined as separate species, *Platythecium maximum* (Groenh.) Z.F. Jia & Lücking. It is a saxicolous species, within China reported from Hong Kong (Thrower 1988).


Figure 4A–B

Feddes Repert. 31: 219, 1933. Type: China. Taiwan, *Asahina 376* (W, holotype!).

Following Aptroot (2004), this taxon is to be treated in the genus *Fissurina*, as *F. micromma* (Zahlbr.) Aptroot. It is a corticolous species reported from Taiwan (Type locality, Zahlbruckner 1933, 1940; Lamb 1963; Wang-Yang and Lai 1973).

**13. Phaeographina mirabilis** Zahlbr.


Following Rivas Plata et al. (2010), this taxon is currently accepted in *Chapsa*, as *C. mirabilis* (Zahlbr.) Lücking. It is a corticolous species reported from Fujian (Type locality, Zahlbruckner 1930, 1932).

Flora 65: 399, 1882.

Following Staiger (2002), this taxon is to be treated in *Thecaria* as *T. montagnei* (Bosch) Staiger. The distribution of this species is treated above.

15. *Phaeographina obfirmata* (Nyl.) Zahlbr.

Figure 5A–B


Following Staiger (2002), this taxon is a synonym of *Glyphis scyphulifera* (Ach.) Staiger. The latter is a corticolous species reported from Shanghai (Type locality) and Hong Kong (Thrower 1988).


Figure 5C–D


This taxon was suggested to represent a synonym of *Graphis alpestris* by Nakanishi on an annotation label. Zahlbruckner (1930) gives the ascospores as 135–165 × 30–33 μm, somewhat larger than reported for *G. alpestris* (70–115 × 19–30 μm). Unfortunately, the remaining material has only empty lirellae, so this information cannot be checked. We therefore follow Nakanishi in treating this name as a synonym of *Graphis alpestris* (Zahlbr.) Staiger. It is a corticolous species reported from Yunnan (Type locality of *Phaeographina pluvisilvarum*, Zahlbruckner 1930, 1932).


Flora 65: 398, 1882.
Following Lücking et al. (2007), this taxon was retransferred to *Thecographa* as *T. prosiliens* (Mont. & Bosch) A. Massal. The species is a corticolous species reported from Taiwan (Apt-root and Sparrius 2003 as *Phaeographina prosiliens*; Zahlbruckner 1933, 1940; Lamb 1963; Wang-Yang and Lai 1973 as *Phaeographina valida* Zahlbr. another synonym seen below).


Figure 6A–B


≡ *Platythecium pyrrhochroon* (Mont. & Bosch) Z.F. Jia & Lücking, comb. nov. (= *Platythecium dimorphodes* (Nyl.) Staiger).

According to the lectotype in PC which we studied, this name is synonymous with *Platythecium dimorphodes* (Nyl.) Staiger and unfortunately provides an earlier epithet for it. Since the latter name has only been introduced recently and has been used few times afterwards, a conservation proposal seems not in order and we propose the new
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combination *Platythecium pyrrhochroum* (Mont. & Bosch) Z.F. Jia & Lücking, with *P. dimorphodes* (Nyl.) Staiger as synonym. Redinger (1935) and Nakanishi (1966) had a different concept of *Phaeographina pyrrhochra*, referring to a species with prominent lirellae with closed labia and large ascospores, which would correspond to what is now known as *Pallidogramme chrysenteron*. Nakanishi discovered this error when he studied the type in 1973, but this has apparently never been published. *Platythecium dimorphodes* is a corticolous species reported from Hong Kong (Aptroot and Seaward 1999) and Taiwan (Zahlbruckner 1933; Wang-Yang and Lai 1973).

≡ *Thecaria quassiicola* Fée, Essai Crypt. Exot. (Paris): xcii, tab. 1, fig. 16 (1825) [1824][as ‘quassiacola’].

Staiger (2002) retransferred this taxon to *Thecaria* as *T. quassiicola* Fée. It is a corticolous species reported from Zhejiang (Xu 1989), Fujian (Zahlbruckner 1930), Hainan
(Wei et al. 2013), Taiwan (Zahlbruckner 1933; Wang-Yang and Lai 1973) and Hong Kong (Thrower 1988; Aptroot and Seaward 1999, Aptroot and Sipman 2001).


Following Staiger (2002), this taxon was transferred to Phaeographis as P. scalpturata (Ach.) Staiger. It is a corticolous species reported from Hong Kong (Aptroot and Seaward 1999; Aptroot and Sipman 2001) and Taiwan (Aptroot and Sparrius 2003).

Figure 6C–D
Feddes Repert. 31: 217, 1933. Type: China. Taiwan, Asahina 355 (W, lectotype!).

Based on the characteristics of the lirellae, excipulum and ascospores, this taxon belongs in Thecographa. It has muriform ascospores, 140–160 × 20–28 μm in size, hence we treat it as a synonym of Thecographa prosiliens (Mont. & Bosch) A. Massal. The distribution of this species is referred above.

Nomenclatural novelties
Phaeographis pleiospora (Zahlbr.) Z.F. Jia & Lücking, comb. nov.
MycoBank No. 820372

Bas.: Phaeographina lecanographa var. pleiospora Zahlbr., in Handel-Mazzetti, Symb. Sin. 3: 61, 1930. Type: China. Fujian, Chung 596b (W, holotype!).

Description. Thallus corticolous, crustose, surface pale grey to yellowish-grey, smooth to slightly rough; apothecia rounded to shortly elongate, erumpent to prominent, usually fissured, single, rarely branched, 1–3 mm long and 0.5–2 mm wide; thalline margin partially exposing the (dark) brown disc; labia inconspicuous; discs opened, brown; proper exciple laterally carbonized; hymenium inspersed; 2–4 ascospores per ascus, brown, ellipsoid to subovate, muriform, 60–80 × 20–30 μm, I–.

Chemistry. Norstictic acid.

Notes. This species is distinguished from Phaeographis lecanographa by the smaller ascospores (60–80 × 20–30 μm vs. 85–150 × 25–50 μm).
**Platygramme elaeoplaca** (Zahlbr.) Z.F. Jia & Lücking, comb. nov.
MycoBank No. 820376


**Description.** Thallus corticolous, crustose, surface yellowish-grey, somewhat greenish, waxy, flat to slightly rough; apothecia lirelliform, sessile, black, elongate, single, not branched, 1–5 mm long and 0.2–0.4 mm wide; labia entire, exposed; discs closed, or slightly opened; proper exciple apically carbonized, thick; hymenium inspersed; 1 ascospore per ascus, brown, elongate-ellipsoid, densely muriform, 80–110 × 20–30 μm, I+ red-brown.

**Chemistry.** No lichen compounds detected.

**Notes.** This species is similar to *Platygramme commutabilis*, but the latter differs in having ascomata covered with a whitish thallus layer. *Platygramme platyloma* has a distinct, lateral thalline margin covering the ascomata, as well as larger ascospores (see above).

**Platythecium maximum** (Groenb.) Z.F. Jia & Lücking, comb. nov.
MycoBank No. 820377

Bas.: *Phaeographina maxima* Groenb., in Blumea, Suppl. 5 (H. J. Lam Jubilee Vol.). 107, 1958. Type: Indonesia, Sumatra, *Groenhart 9453* (BO, holotype, not seen, but original image in protologue seen).

**Description.** Thallus saxicolous, crustose, surface grey, smooth; apothecia lirelline, elongate, subimmersd, single and branched, 1–20 mm long and 0.2–0.3 mm wide, with lateral thalline margin; labia conspicuous; discs open, dark; proper exciple completely carbonized; hymenium clear; 8 ascospores per ascus, brown, ellipsoid, muriform, 4/1–2-locular, 14–16 × 8–9 μm, I–.

**Chemistry.** Not tested.

**Notes.** This species is similar to *Platythecium serpentinellum* (Nyl.) Staiger in the brown ascospores and to *P. allosporellum* (Nyl.) Staiger in the carbonized hypothecium. The type material is different from any known species in the genus, it is here recombined as a separate species.

**Platythecium pyrrhochroum** (Mont. & Bosch) Z.F. Jia & Lücking, comb. nov.
MycoBank No. 820382


**Description.** Thallus corticolous, crustose, surface whitish to yellowish, smooth to minutely roughened; apothecia lirelliform, sessile, elongate, single to branched, 1–5 mm long and 0.3–0.4 mm wide; labia inconspicuously striate, thalline margin whitish, concolorous with the thallus; discs closed, or slit-like; proper exciple uncarbonized, pale-brownish; hymenium clear; 2–4 ascospores per ascus, brownish, oblong, densely muriform, 28–60 × 10–17 μm.

**Chemistry.** Norstictic acid.

**Notes.** As stated above, this name is synonymous with *Platythecium dimorphodes* (Nyl.) Staiger and unfortunately provides an earlier epithet for it.

*Sarcographina heterospora* (Nyl.) Z.F. Jia & Lücking, comb. nov.
MycoBank No. 820385


**Description.** Thallus corticolous, crustose, surface yellowish to pale grey, flat to slightly rough; apothecia lirelline, immersed in stromata; stromata white, irregularly circular, 0.1–1 mm wide; lirellae thin, black, immersed, in irregular, stellate cluster, 0.05–1 mm wide; discs black, closed; proper exciple uncarbonized, pale yellow brown; hymenium clear; 8 ascospores per ascus, brown, rounded ellipsoid, irregularly 4/1, 2/2 or 3/1 –2 locular, 10–13 × 6–8 μm, I+ red-brown.

**Chemistry.** Psoromic acid and conpsoromic acid.

**Notes.** This species is very similar to *Sarcographina cyclospora* Müll. Arg., but the latter differs in having a brownish to apically carbonized excipulum.

**Acknowledgements**

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References


Unexpected high species diversity among European stalked puffballs – a contribution to the phylogeny and taxonomy of the genus Tulostoma (Agaricales)

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Abstract
A three-gene data set was generated to explore species diversity and delimitations within the stalked puffballs (Tulostoma, Agaricales) in Europe. Data on species from other parts of the world were included for comparison of species concepts and distribution ranges. Sequence data from 26 type specimens are included. The phylogenetic analyses support Tulostoma as monophyletic. Eleven major clades, 37 minor clades, and 20 single branches were recovered and found to correspond to 30 described species and 27 species without scientific names.

Five species are here described as new to science: Tulostoma calcareum, T. calongei, T. eckbladii, T. grandisporum, and T. pannonicum. In total we report 26 described, and 19 undescribed, species from Europe. An epitype for T. fimbriatum with ITS sequence data is selected to fix the name.

The recovered tree topology was not in congruence with the current infrageneric classification of Tulostoma, suggesting that many of the morphological characters used for segregation of taxa are plesiomorphic or homoplasious. Spore ornamentation and hyphal structure of the peridium are found to be reliable characters for delimitation of species.
The majority of the species occur in the dry, arid areas of southern and east central Europe but a few are shown to be restricted to humid temperate regions in the North. The study confirms that species with smooth or sub-smooth spores are restricted to dry and arid habitats whereas species with more strongly ornamented spores occur in humid habitats.

Areas with steppe vegetation in Hungary and Spain are here identified as hot spots for *Tulostoma* species diversity.

**Key words**
Gasteroid fungi, molecular systematics, species diversity hot spot, steppe vegetation, taxonomy

**Introduction**

The genus *Tulostoma* Pers. : Pers. was erected by Persoon (1794, 1801) to accommodate puffball species with a “peridium pedicellatum”, opening with an “ore cylindraceo cartilagineo”. Persoon included two species in the genus, *T. brumale* and *T. squamosum*. The former had previously been illustrated by Tournefort (1700) as *Lycoperdon parisiense* and later described by Linnaeus (1753) as *Lycoperdon pedunculatum*. Fries (1829) mentioned four species, viz. *T. mammosum* (including *T. squamosum*), *T. fimbriatum*, *T. tortuosum* and *T. laceratum* (=*Schizostoma lacerata*). However, in a later work, Fries (1849) replaced *Tulostoma* with *Tulasnodea*, honouring the Tulasne brothers, contemporary mycologists based in Paris. However, according to the rules of nomenclature, *Tulostoma* must be considered the correct name for this characteristic genus, being sanctioned by Persoon (1801). An orthographic variant, *Tylostoma*, is sometimes seen in older literature. Early contributions to the knowledge of *Tulostoma* are the works of Bottomley (1948), Coker and Couch (1928), Cunningham (1925, 1932), Hollós (1904, 1913), Lloyd (1906), Long (1944, 1946), Long and Ahmad (1947), Morgan (1891), Petri (1904, 1909), Pouzar (1958), Shvartsman and Filimonova (1970), Smith (1951), and White (1901).

Schroeter (1876) studied young and old basidiomata of *Tulostoma* and described the development of the basidiomata, in particular basidia and capillitium. Later Wright (1955) surveyed important morphological characters in the genus. This was followed by the world monograph of *Tulostoma* (Wright 1987), where he accepted 139 species. Wright’s species concept was largely based on type studies and studies of numerous additional herbarium specimens. With the introduction of scanning electron microscopy (SEM), detailed spore morphology could be used as a base for taxonomic rearrangements and introduction of new species (Altés and Moreno 1995, Altés and Moreno 1999, Altés et al. 1999, Moreno et al. 1992a, Moreno et al. 1997).

From the late 20th century the arid regions of Mexico have been intensely explored with regard to *Tulostoma* (Esqueda et al. 2004, Hernández-Navarro et al. 2015, Moreno et al. 1995b, Piña et al. 2010, Wright et al. 1972). In addition, South America, particularly Brazil, has recently been subject to detailed studies on *Tulostoma* (e.g. Baseia and Milanez 2002, Cortez et al. 2009, Silva et al. 2007). Silva’s (2006) thesis on *Tulostoma* substantially increased the knowledge of the genus in Brazil. For South Africa, Coetzee

The introduction of molecular data has led to major changes in fungal systematics and taxonomy. Hibbett et al. (1997) were the first to show that the puffball genus Tulostoma belongs in Agaricales. Although the genus has been intensively studied from a morphological point of view, so far comparatively few molecular phylogenetic studies with a focus on Tulostoma have been published, and these are mainly geographically restricted and deal with a low number of species (cfr. Hernández-Caffot et al. 2011, Hussain et al. 2016).

In the present study we use a three-gene phylogeny with the aim to explore species diversity and phylogenetic relationships within Tulostoma. The data set includes the ITS, LSU, and Tef-α regions of the majority of the European species. In addition, species described from other parts of the world are included for comparison of species concepts and distribution ranges. Retrieval of ITS sequence data from type and other herbarium specimens was attempted when permission for sequencing was given and as the condition of the material allowed.

**Material and methods**

**Morphological methods**

The genus Tulostoma is easy to identify to genus level in the field. Precise determination to species level is, however, often difficult, because the characters used are few in number and subtle in nature. According to Index Fungorum (http://www.indexfungorum.org; accessed July 1, 2016) there are well over a hundred Tulostoma names to consider, many of them known from old and scanty type materials at best. Our sampling and selection of species was compiled through field studies and collecting in various parts of Europe during the last 25 years (collections stored in GB and AH).

Additional collections and type specimens were studied as loans from herbaria around the world (BPI, BRA, C, FH, L, LD, LISU, MA, NY, O, PC, PRM, S, TRH, and UPS). Herbarium acronyms are in accordance with Index Herbariorum (http://sweetgum.nybg.org/science/ih/). Species very recently recorded from Europe (T. bruchi, T. leslei, T. palatinum, and T. vulgare) are not included in this study (Antonín and Kreisel 2008, Calonge 1992, 1998, Specht et al. 2016).
Collections were mostly photographed in situ, dried, and later studied in the laboratory under a stereo-microscope. Morphological features are named in accordance with Wright (1987). For microscopic studies, samples of mature gleba were mounted in Cotton blue + lactophenol and heated to boiling. Features of the peridium were studied in Melzer’s reagent. A light microscope equipped with a Dino-Eye Eyepiece Camera was used for studies of micro-morphological characters. Spores were measured at a magnification of 1000x, using the Dino-Lite 2.0 software (www.dino-lite.eu). A minimum of 20 spores were measured for each sample. All spore measurements are given excluding ornamentation. Studies under SEM were conducted according to the procedure of Moreno et al. (1995b).

**Molecular methods**

**Taxon sampling**

For this study, 183 ingroup specimens were sequenced, including the type specimen of 24 species. ITS sequence data of the type specimens of *T. domingueziae* (HQ667597) and *T. ahmadii* (KP738712) were taken from GenBank (Hernández-Caffot et al. 2011, Clark et al. 2016). Specimens were selected with the aim to cover all the species known to occur in Europe and from a broad geographic distribution within the area. In addition, specimens from North America and Central Asia were included for comparison of species concepts and distribution ranges. The sequenced specimens shown in Figure 1a–c are indicated with an asterisk in the lists of specimens examined and are listed with GenBank accession numbers as Suppl. material 1.

Three additional ITS and LSU sequences from Hernández-Caffot et al. (2011) were retrieved from GenBank and included in the data set (HQ667594, HQ667598-HQ667599). Based on results from earlier molecular phylogenetic studies of Agaricales (Hibbett et al. 1997, Matheny et al. 2006), *Lycoperdaceae, Cyathus, Crucibulum, and Inocybe* were selected as outgroup taxa.

**DNA extraction, PCR, and sequencing**

Sequence data from three genetic markers were generated for the study: the complete ITS region and about 1400 bases of the 5’end of the nuclear ribosomal LSU DNA, and about 1000 bases of translation elongation factor subunit 1 alpha (Tef-1a). DNA extractions were performed using DNeasy Plant Mini kit (Qiagen, Hilden, Germany), PCR reactions, and sequencing were performed as described in Larsson and Örstad (2008). Primers used to amplify the complete ITS region and the 5’ end of the LSU region were ITS1F (Gardes and Bruns 1993) and LR21, LR0R, and LR7 (Hopple and Vilgalys 1999). For Tef-1a we used EF983F and EF2218R (Rehner and Buckley 2005). Primers used for sequencing were ITS1, ITS4 (White et al. 1990), LR0R, LR5,
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Figure 1. a–c One of the most parsimonious trees obtained from the parsimony analysis based on ITS, LSU, and Tef-1α sequence data of Tulostoma, with focus on species diversity in Europe. Bootstrap values and Bayesian Posterior Probabilities are indicated on branches. The major clades (1-11) are marked with a scale bar. The clades represent species or species groups that are discussed in more detail in the text.

and LR3R (Hopple and Vilgalys 1999, Tedersoo et al. 2015), EF983F, and EF2218R. Type specimens were extracted using a modified CTAB method, and PCR and sequencing followed protocols described in Larsson and Jacobsson (2004).

Phylogenetic analyses

Sequences were edited and assembled using Sequencher 5.1 (Gene Codes, Ann Arbor). Alignment of individual genes was performed using the L-INS-i strategy as implemented in MAFFT v. 7.299 (Katoh and Standley 2013). The alignments were adjusted manually using AliView (Larsson 2014). Sequences generated for this study have been...

Separate phylogenetic analysis was done for all three genetic markers to test for overall congruity of the phylogenetic signal. The trees were found to be compatible with
Figure 1. Continue.
respect to the overall clades (results not shown), and the three genetic markers were concatenated for the final analyses. Heuristic searches for the most parsimonious trees were performed using PAUP*. All transformations were considered unordered and equally weighted. Variable regions with ambiguous alignment, mainly from the ITS region, were excluded, and gaps were treated as missing data. Heuristic searches were performed with 1,000 random-addition sequence replicates, TBR branch swapping, and the MultTrees option in effect. Relative robustness of clades was assessed by the bootstrap method using 1,000 heuristic search replicates with 100 random taxon addition sequence replicates and TBR branch swapping, the latter saving at most 25 trees in each replicate.

Bayesian phylogenetic analyses were carried out in MrBayes 3.2.6 (Ronquist et al. 2012), with a best-fit model of nucleotide evolution for the separate gene partitions supplied by MrModeltest 2.2 (Nylander 2004). Eight default-setting Metropolis-Coupled Markov Chain Monte Carlo (MCMCMC) chains were run for 10 million generations with trees sampled every 5,000 generations and an initial burn-in of 50%. After discarding the trees prior to the burn-in threshold, a 50% majority-rule consensus phylogram was computed from the remaining 1,000 trees.

Results

Molecular analysis

The aligned complete dataset, including sequences downloaded from GenBank, consisted of 198 taxa and 3,340 characters. After exclusion of ambiguous regions, mainly from the ITS1 and ITS2, 3,005 characters remained for the analysis. Of these, 1,992 were constant, 226 were variable but parsimony uninformative, and 787 (26%) were parsimony informative.

The maximum parsimony analysis yielded 3625 equally parsimonious trees (length=3,726 steps, CI= 0.3814, and RI= 0.8228). One of the trees is presented as a phylogram in Figure 1a–c.

The bootstrap analysis recovered *Tulostoma* as monophyletic with 95 % bootstrap support (BS). Thirty-seven minor clades and 20 single branches were recovered, corresponding to 30 described species and 27 without a scientific name. Five of these clades are here described as new species to science. Eleven major clades, ranging from virtually unsupported to having strong support, within the ingroup are recognized and named as Clades 1-11 in the phylogenetic tree (Figure 1a–c). These clades are further described and discussed below.

As suggested by MrModeltest, the nucleotide evolution model HKY+G was used for the ITS1 spacer; SYM was used for the 5.8S gene; HKY+G was used for the ITS2 spacer, and GTR+I+G were used for the nLSU and TEF genes in the Bayesian analysis. The MCMC analysis converged well in advance of the burn-in threshold and chain mixing was found to be satisfactory, as assessed by using Tracer v1.5 (Drummond et al. 2012). Also in the Bayesian analysis, *Tulostoma* was recovered as monophyletic with
strong support (a Bayesian posterior probability (BPP) of 1.00). The Bayesian tree topology is similar to the MP bootstrap tree. The same clades and single branches recovered in the maximum parsimony analysis were also recovered in the Bayesian analysis, with the minor differences that several clades with low or no bootstrap support received a moderate to high BPP value. BPP values are indicated on the corresponding branches in Figure 1a–c.

None of the newly generated ITS sequences were significantly similar to the “most wanted fungi” ITS dataset of Nilsson et al. (2016) as explored through BLAST.

Comments on the major clades

Clade 1 (100, 1.0) includes only one species, *T. punctatum*. The exoperidium is hyphal and the mouth is fibrillose-fimbriate. It is similar to *T. fimbriatum* in macro-morphology but has smaller and more coarsely ornamented spores without anastomosing ridges. The species was treated as a variety of *T. fimbriatum* by Wright (1987).

Clade 2 (unsupported) includes four species that all have a hyphal exoperidium. Two of the species (sp.1, sp. 2) are unidentified. The specimens of these originate from halophytic vegetation in Spain and sand steppe vegetation in Hungary, respectively. Both have a fibrillose-fimbriate mouth, thus inviting comparison to the widespread *T. fimbriatum* in morphology. *Tulostoma cyclophorum* has a typically fibrillose-fimbriate mammose mouth but differs from other European species of *Tulostoma* by having abundant mycosclereids, i.e. rounded to elongated cells on the endoperidial surface, and subreticulate spores. *Tulostoma obesum* has a smooth endoperidial surface that lacks mycosclereids and the spores are completely smooth.

Clade 3 (90, 1.0) includes four species with similar morphology. They all have a hyphal exoperidium, fibrillose-fimbriate mouth and rather stout basidiomata, features that are characteristic of *T. fimbriatum*. An ITS sequence from the lectotype of *T. campestre* (=*T. fimbriatum* var. *campestre*) was generated and forms a monophyletic clade with the sequences of *T. fimbriatum* sensu stricto. As the morphology also is in congruence, we regard it as a synonym. An epitype is selected for *T. fimbriatum* (see taxonomy section) from the same district in Sweden as the holotype to fix the name also by ITS sequence data (Nilsson et al. 2012, Ariyawansa et al. 2014). *Tulostoma winterhoffii*, a recently described species from Germany, is identical with the ITS sequence from the holotype of *T. fimbriatum* var. *heterosporum*, and thus replaces that name. We describe a new species from Spain, *T. calongei*. It is similar to *T. fimbriatum* in habit but differs in spore morphology and molecular data. The unidentified *T*. sp. 3, also with the habit of *T. fimbriatum*, collected in Hungary, indicates the occurrence of cryptic speciation within this morphologically similar group. Although having been considered synonymous with *T. fimbriatum* or mere varieties of it (Altés and Moreno 1995, Wright 1987), ITS data (not shown in the tree) of type materials of the extra-European taxa *T. egranulosum*, *T. readeri*, and *T. tuberculatum* indicate them to be distinct species. The *T. fimbriatum* complex is in need of further studies.
Clade 4 (unsupported) includes two unidentified species (Tulostoma spp. 4 and 5) collected in sandy habitats in Central Europe. Tulostoma sp. 4 is fairly similar to T. fimbriatum in habit. Tulostoma sp. 5 has a circular, tubularly protruding mouth, a hyphal exoperidium, and irregularly ornamented, verrucose-echinate spores. The two species are very different morphologically and their placements in the tree are ambiguous and without significant support.

Clade 5 (-, .73) is an unsupported clade including 11 species. The holotype of T. lusitanicum (Figure 12), a species recently (2000) described from Portugal, is placed on a single branch basal in the clade, but without significant support. However, the residual subclades form a moderately supported clade (78/.99, Figure 1b). ITS sequence data of the holotypes of T. kotlabae, T. lysocephalum (from North America), and T. lusitanicum were included, but still we have eight unidentified clades (Tulostoma spp. 6-13). This indicates a high and previously unrecognized species diversity within the group, including cryptic speciation. The clade is in need of further study, and the addition of more sequence data appear to be needed to resolve the phylogenetic relationships. Tulostoma kotlabae and spp. 6-13 share morphological characters such as the circular, more or less tubularly protruding mouth, a hyphal exoperidium, pale colours of the basidiomata, and weakly to moderately ornamented spores. Tulostoma sp. 12 is known only from the Mediterranean area, and an Italian finding (AH 16793) was published as T. kotlabae by Altés et al. (1994). Tulostoma lysocephalum differs by having a fibrillose-fimbriate mouth (Figure 22).

Clade 6 (95, 1.0) includes five species. Tulostoma aff. cretaceum forms a strongly supported clade that comprises specimens from Russia, Hungary, and Spain. In morphology they are characterized by stout basidiomata with pale colours and a hyphal or slightly membranous exoperidium, a very dark, chocolate brown colour of the mature gleba, an indistinct, irregular mouth that with age becomes lacerate, combined with totally smooth spores. However, as there is substantial sequence variation that correlates with geographic distribution within the clade (Russia, Hungary and Spain, respectively), it must be regarded as a complex of species. More data such as the sequence of the type of T. cretaceum are needed to resolve species delimitation within the Tulostoma aff. cretaceum clade. Sequence data from the holotype of T. macrocephalum from North America show this species to be closely related to Tulostoma sp. 14, which includes two specimens from Spain, collected in a halophytic habitat. Both species have a hyphal exoperidium but differ in basidiomata size and spore wall ornamentation. Tulostoma pseudopulchellum, a species characterized by a membranous exoperidium, a fibrillose-fimbriate mouth, and finely and irregularly ornamented spores comes out as a sister clade with strong support to what we describe as a new species (T. pannonicum, see taxonomy section). The new species is recognized by having small smooth spores and has previously been reported from Hungary as T. leiosporum (cfr. Jeppson et al. 2011, Rimóczi et al. 2011).

Clade 7 (86, .98) includes two species. Tulostoma submembranaceum was described from Mexico by Moreno et al. (1995b). The exoperidium is thinly membranous-verrucose, the mouth is fibrillose-fimbriate, and the spores have low verrucae that have a tendency to form crests. Tulostoma sp. 15 from Hungary and Spain, more or less
matches the macro-morphology of *T. submembranaceum* but differs in spore ornamen-
tation (irregularly rugulose) and molecular data.

Clade 8 (54.85) includes four species. Uniting morphological characters are the
presence of a fimbriate-fibrillose mouth and totally smooth spores. Two well-known
species are *T. fulvellum* and *T. lloydii*. One of the recovered clades represents a species
with large spores, characteristic of sandy habitats of East Central Europe. It is pro-
posed as a new species (*T. grandisporum*, see taxonomy section). *Tulostoma* sp. 16 is a
specimen from Siberia that in macro-morphology is similar to the newly described *T.
grandisporum* but has significantly smaller spores. Both have irregular, undulating, or
ragged inner walls of the capillitium. This striking character is also present in the lecto-
type of *T. leiosporum*, from which we were not able to obtain sequence data.

Clade 9 (100, 1.0) includes *T. pulchellum* and *T. striatum* and corresponds to sec-
tion *Poculata* Pouzar & Moravec. The exoperidium is distinctly membranous, detaching
in flakes. The endoperidium is white, velvety in young specimens, and has a mammose,
fibrillose-fimbriate mouth that sometimes is surrounded by a more or less delimited peri-
stone. They share macro-morphological features but can be readily separated by their
spores that are finely asperulate in *T. pulchellum* and distinctly striate in *T. striatum*.

Clade 10 (69, -) includes 14 species. All species in this clade have a mouth that is circular
and more or less tubular, or conically protruding. The spores are moderately ornamented.
*Tulostoma simulans*, a species described from N. America, is according to our results widely
distributed also in Europe. In some parts of Europe, *T. simulans* has been mistaken for *T.
beccarianum* and the species *T. moravecii*. The latter now appears to be a later synonym. The
well-known and widely distributed *T. brumale* is closely related, but clearly distinct from *T.
simulans*, both in molecular and morphological data. *Tulostoma beccarianum* is a rare spe-
cies described from Italy by Bresadola, but with recent records from Spain and East Central
Europe. *Tulostoma* spp. 17-21, all represented by few specimens, are similar in morphology
to *T. beccarianum* and *T. simulans*, and represent un-named cryptic species that need fur-
ther attention. The ITS of the types of *T. albicans*, *T. excentricum*, and *T. xerophilum*, three
species from North America, were generated and included, but did not match with any of
our unknown species, and the latter occurs on a single basal branch in Clade 10. *Tulostoma
giovanellae* is mainly associated with sandy, and often halophytic, habitats in southern and
central Europe. The sister clade to *T. giovanellae* represents a species from northernmost
Scandinavia that we describe as a new species (*T. eckbladii*, see taxonomy section). Within
the clade there are also *T*. sp. 21, an unidentified Spanish specimen, closely related to *T.
eckbladii*, and the characteristic bryophilous *T. niveum*.

Clade 11 (95, 1.0) includes eight species that morphologically are characterized by
having rather dark colours of the stipe, endoperidia with initially brownish colours, and
moderately to strongly ornamented, verrucose-echinate spores. The widely distributed
*T. squamosum* has a membranous-verrucose exoperidium of sphaerocysts, which upon
maturity forms a dark reticulum on the brownish endoperidium. The closely related
*T. subsquamosum* has a paler, hyphal exoperidium with only scattered sphaerocysts,
and never forms a reticulum. It was described from India and reported from Spain
by Altés et al. (1996) and is here confirmed also from East Central Europe (Hungary
and Slovakia). Sequence data indicate a close relationship with the recently described *T. ahmadii* (Hussain et al. 2016). *Tulostoma melanocyclum* is a widely distributed species in Europe. The recently described *T. domingueziae*, reported from South America (Hernández-Caffot et al. 2011), as well as *T. rufum* (lectotype; North America) come out as sister species. One of the recovered clades is described as a new species (*T. calcareum*, see taxonomy section) that has a wide European distribution. The sequence data of *T. sp. 22* represent an undescribed species from South America.

### Taxonomy

Recognized European species in *Tulostoma* and the descriptions of five new species


**Figure 2a**

**Holotype.** ITALY, Pisa: Beccari (S!)*.

The holotype material of *T. beccarianum*, described in 1904, was sequenced in this study. Identical sequences were obtained from recently collected material from Hungary, Slovakia, and Spain. The species was given a detailed description by Altés and Moreno (1993) based on the holotype. The newly collected samples have rather large and stout basidiomata, spore-sacs measuring up to 22 mm in diameter, and a stem reaching 120 mm. The exoperidium is indistinct or hyphal, and the endoperidium is smooth, dirty white-greyish, with a circular, shortly raised mouth. The stem base is widened and forms a volva-like structure. The capillitial septa are slightly widened. The capillitium seems to break up easily at the septa, leaving segments with somewhat widened, rounded ends. The type material shows irregular, undulating inner walls of the capillitium, a character that is also noted in the recently collected material. The spores are irregularly verrucose, 4–5 μm, av. 4.5 μm, ornamentation excluded (Figure 2a) and agree with those of the holotype (4.7–6 μm, ornamentation included; Altés and Moreno 1993). The synonymisation of *T. beccarianum* with *T. simulans*, as proposed by Altés and Moreno (1993), was contradicted by the molecular analyses. The name *T. beccarianum* has apparently sometimes been misapplied for *T. simulans* (syn. *T. moravecii*). *Tulostoma* sp. 19 includes a single collection from Cyprus, with more or less identical morphology. It is currently treated as an undescribed species until more material becomes available.

**Habitat and distribution.** In semi-shaded to exposed localities in dry grasslands (Hungary and Slovakia) and halophytic vegetation on sand (Spain). Apparently a very rare species.

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Figure 2. Spore ornamentation under SEM of European species: a T. beccarianum (Finy 2, GB) b T. brumale (M. Jeppson 8372, GB) c T. aff. cretaceum (M. Jeppson 3821, GB) d T. cyclophorum (AH 16885) e T. fulvellum (AH 13415) f T. giovannellae (AH 11641) g T. koltabae (M. Jeppson 5597, GB) h T. lloydii (AH 11606) i T. melanocyclum (S. Hanson 2008-247, GB) j T. niveum (M. Jeppson 7699, GB) k T. obsenum (AH 20901) l T. pulchellum (M. Jeppson 7833, GB). Scale bars: 1 μm.

Figure 2b

**Type.** FRANCE, “circa Parisios” (L); several collections by Persoon available at L, but no type seems to have been formally designated (cfr. Wright 1987).

*Tulostoma brumale* is well characterized by its membranous exoperidium, the circular-tubular mouth—which is usually surrounded by a brown peristome—and its abrupt widenings of the capillitial septa. SEM-photos of the spores show an irregular ornamentation of blunt, broad-based verrucae (Figure 2b). It can be separated from *T. simulans* by the exoperidial features, its slightly smaller spores, and the presence of irregular crystals adhering to the capillitial walls.

**Habitat and distribution.** According to Wright (1987), *T. brumale* is recorded from N and S America and Asia (Georgia). Shvartsman and Filimonova (1970) reported it from Central Asia and Rebriev and Gorbunova (2007) added findings from Siberia. In Europe it is one of the more common species of *Tulostoma*, recorded north to 60° in Fennoscandia. It occurs in all types of sandy and calcareous grasslands, sand dunes, and sand steppe vegetation, as well as on moss covered rocks and stone walls. It is often associated with mosses, particularly *Syntrichia* spp.


*Tulostoma calcareum* Jeppson, Altés, G. Moreno & E. Larss., sp. nov.

MycoBank number: MB819399
GenBank: KU519086 (ITS-LSU), KU843881 (Tef-1a)

**Figure 3**

**Holotype.** Sweden, Södermanland, Mörkön, Egelsvik, Kalkberget, on calcareous soil under *Artemisia campestris* in rocky slope, 5 Nov. 2004, M. Jeppson 6965 (GB!, isotype AH).

**Etymology.** The name refers to its habitat requirement, on calcareous sandy soil or among calcareous rocks and cliffs.

**Description.** Spore-sac subglobose, often somewhat depressed, 5–12 mm. Exoperidium hyphal-verrucose, deciduous, sometimes persisting as whitish verrucae scattered
on the endoperidial surface. Endoperidium brownish-ochraceous, initially rather dark coloured, with age fading to greyish white. Mouth circular, shortly tubular (Figure 3f), surrounded by a greyish or brownish peristome. Socket separated from the stem, inconspicuous. Stem slender, 20–50 x 2–3 mm, initially orange brown, with age warm reddish brown, longitudinally furrowed, smooth to fissured and appressed scaly, with a basal mycelial bulb (Figure 3a,e). Gleba ochraceous to ferrugineous brown. Capillitium 4–10 μm with medium-thick walls, sometimes abundantly ramified. Septa not or very slightly widened (Figure 3b). Spores subglobose, 4.0–6.0 μm (av. 4.7–5.0 μm), verrucose-echinate. SEM-photos show spines in groups with connected tips, sometimes coalescing to form ridges and crests (Figure 3g–i).
**Habitat and distribution.** Occurs in dry, exposed to semi-shaded situations in calcareous, sandy habitats and on calcareous rocks and cliffs. It is currently on record from Hungary, Norway, Spain, and Sweden.

**Notes.** *Tulostoma calcareum* was commonly misinterpreted as *T. squamosum* (cfr. Nitare 1997), under which name it is red-listed as Critically endangered (CR) in Sweden (Artdatabanken 2015). It differs from *T. squamosum* by having a hyphal to slightly verrucose exoperidium, by lacking sphaerocysts, and by never forming the reticulate pattern typical of *T. squamosum*. The stipe is also less squamulose. Spores and capillitial characters are almost identical. It differs from *T. melanocyclus*, with which it can easily be confused, by having a more robust stature, more orange-reddish brown colours of the stipe, and slightly smaller spores (av. 4.7–5.0 μm vs 5.2–5.4 μm in *T. melanocyclus*). SEM-photos of *T. melanocyclus* (Figure 2i) show similar spore ornamentation but with spines with less tendency to coalesce and form ridges. In macro-morphology *T. calcareum* is fairly similar to *T. ferrugineum* D.M. Oliver & Hosford, a species described from northwestern USA, whose type material no one has been able to locate (Wright 1987). However, according to its original description, it differs from *T. calcareum* by having distinctly widened capillitial septa.

**Tulostoma calongei** Jeppson, Altés, G. Moreno & E. Larss., sp. nov.

MycoBank number: MB819400
GenBank: KU518973 (ITS-LSU), KU844000 (Tef-1a)

Figure 4


**Etymology.** Named in honour of the Spanish mycologist Francisco D. Calonge.

**Description.** Spore-sac 8–10 mm, subglobose. Exoperidium granulose hyphal, encrusting soil, sometimes detaching in small flakes, whitish-greyish, rather persistent (Figure 4f). Endoperidium smooth, whitish. Mouth fimbriate. Stipe rather short, 15–20 x 1.5–2.5 mm, warm reddish brown, with darker appressed scales and a basal mycelial bulb (Figure 4f). Gleba ochraceous to ferrugineous brown. Capillitium 3–6 μm, thick-walled, with uneven inner walls (Figure 4c-d). Some ribbon-like hyphae present (Figure 4b). Septa rare, slightly widened. Spores globose-subglobose, 3.5–4.5 μm (av 4.1 μm) ornamentation excluded, verrucose-echinate, with spines frequently fused at their tips forming little developed pyramidal groups (Figure 4g-i).

**Habitat and distribution.** *Tulostoma calongei* has been collected only from central Spain, usually on sandy acidic soils.

**Notes.** Although usually having a less robust stature, this new species is very similar in macro-morphology to *T. fimbriatum*, with which it has been confused. The
main differences are that the spores are slightly smaller and have a conical-pyramidal ornamentation, without the characteristic crests of *T. fimbriatum* (Figure 7c–e). The molecular data also clearly support them as distinct but closely related species, with *T. calongei* occurring as a sister clade to *T. fimbriatum* (Clade 3, Figure 1a). The distribution area for the species is likely to increase after reexamination of additional samples identified as *T. fimbriatum* from Meridional Europe.


**Figures 2c, 5a–b**

**Remarks.** Clade 6 in our study encompasses some closely related taxa previously attributed to *T. volvulatum* I.G. Borschch. and *T. obesum* Cooke & Ellis (cfr. Hollós 1904, Altés et al. 1999, Rimóczi et al. 2011). The main characteristics are the rather large basidiomata with hyphal exoperidium and almost pure white endoperidium with an indistinct to fibrillose-fimbriate mouth (or irregular scar) that with age becomes lacerate. The stipe base is provided with a volva and a prominent pseudorhiza. The spores are smooth and the capillitium is fragile, breaking up in segments with rounded ends. In our material this complex is represented by collections from Hungary, Russia (Altay Republic), and Spain. They are recovered as closely related and can be considered as cryptic species with a strong geographical signal. In morphology they fit the original description of the American species *T. cretaceum* fairly well. The Spanish collections deviate from the Hungarian specimens in having a stouter habit.

**Habitats and distribution.** *Tulostoma cretaceum* was originally described from New Mexico (USA) and later reported from Argentina (Wright 1987) and Baja California, Mexico (Moreno et al 1995b). There are recent records from Mexico (Sonora) and Brazil (Esqueda et al. 2004, Silva et al. 2007). In Europe *T. aff. cretaceum* is found in sand steppe vegetation in Central Hungary and in calcareous steppe habitats in Spain.

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Figure 5. Habit of a selection of studied collections: a T. aff. cretaceum (M. Jeppson 7759, GB) b T. aff. cretaceum (M. Jeppson 6194, GB) c T. obesum (M. Jeppson 8695, GB) d T. punctatum (M. Jeppson 10058, GB) e T. simulans (M. Jeppson 7865, GB) f T. simulans (M. Jeppson 9302, GB) g T. subsquamosum (M. Jeppson 9336, GB) h T. winterhoffii (M. Jeppson 2379, GB). Scale bars: 10 mm.


_Tulostoma cyclophorum_ Lloyd, Mycol. Writings 7: 25. 1906.

Figure 2d

_Holotype._ SOUTH AFRICA, Stoneman, herb. Lloyd 4495 (BPI).

_Tulostoma cyclophorum_ is a morphologically characteristic species, originally described from South Africa, with a tomentose, white to ochraceous endoperidium covered with mycosclereids and a distinctly membranous exoperidium. The mouth is silky fibrillose-fimbriate. Its spores are unique among European _Tulostoma_ lineages, being more or less reticulate (visible under SEM, Figure 2d; asperulate under light microscope).
Habitat and distribution. Mostly found in semi-shaded localities among grasses and herbs on lawns in city parks on somewhat sandy soil. From Europe currently reported from France, Italy, Hungary, and Spain (Demoulin 1984, Jeppson et al. 2011, Moreno et al. 1990, Sarasini 2005). It appears to have a cosmopolitan world distribution.


Tulostoma eckbladii Jeppson, Altés, G. Moreno & E. Larss., sp. nov.
MycoBank number: MB819401
GenBank: KU519068 (ITS-LSU), KU843951 (Tef-1a).
Figure 6

Holotype. Norway, Nordland, Saltdal, Junkerdalsura, på sten blandt mos, 4 Sept. 1957, I. Jørstad (O F-58850!).


Description. Spore-sac subglobose to onion-shaped, 6–8 mm in diameter. Exoperidium not observed. Endoperidium smooth, initially brownish, later ochraceous white. Mouth circular, shortly tubular, with a more or less brown peristome. Socket inconspicuous. Stem stout, 25–30 x 3–4 mm, flattened, light brown, somewhat longitudinally furrowed, almost smooth to finely appressed scaly, attenuated towards the base, with whitish basal rhizomorphs attached to moss rhizomes (Figure 6d). Mature gleba ochraceous-ferrugineous. Capillitium 4–6 μm with rather thick walls, with scattered septa that are slightly widened (Figure 6b). Spores subglobose, 4.5–5.5 μm (av. 4.9 μm) ornamentation excluded, coarsely but not very densely verrucose. SEM photos show isolated, irregular verrucae with broad bases (Figure 6e–g).

Habitat and distribution. Tulostoma eckbladii is so far only known from two findings in northern Norway, in both cases growing among mosses on calcareous boulders.

Notes. The species reminds of T. niveum in its choice of habitat on moss-covered rocks. It can, however, be distinguished by its stouter basidiomata and the presence of a brown peristome. Whether or not the flattened stipe is a constant species character cannot be decided from the available material. The spores are similar, as observed from SEM photos (Figure 2j, 6e-g). It is distinguished from T. brumale by the presence of a stout stipe, the less widened capillitial septa, and the larger spores. Also T. simulans is morphologically similar but is recognized by its more slender basidiomata and different habitat and distribution. The two findings of T. eckbladii
constitute northern extremities in the distribution of the genus *Tulostoma*. It has previously been reported and discussed by Brochmann et al. (1981) as *Tulostoma* sp.

**Other specimens examined.** NORWAY, Finnmark: Porsanger, Hestnes, 17 Jul. 1993, B. Jonsell (as *T. kotlabae*, TRH 9565)*.

Figures 7, 8


**Figure 6.** *T. eckbladii*, holotype (O F-58850): a, d basidiocarps b capillitium c, e–g spores. Scale bars: a, d = 10 mm; b–c = 10 μm; e–g = 1 μm.

*Tulostoma fimbriatum* is one of the more frequently recorded species of the genus. It was described by Fries (1829) from southernmost Sweden. In the Elias Fries Herbarium at UPS there is a collection from the type locality (Lomma, S. Sweden; Fries 1829, 1849). On the label there is the handwriting of T. M. Fries (Elias Fries’ son) but the material is likely to be an authentic collection of Elias Fries. The material is scanty (one basidiome glued to a piece of cardboard) with the endoperidial mouth zone destroyed and with indistinct micro-morphological characters. Nevertheless it should be considered a holotype, but since its characteristics are ambiguous, an epitype is here chosen from recently collected and sequenced material originating from southernmost Sweden. *Tulostoma fimbriatum* can be distinguished from other species with similar macro-morphology by its spores, which have a low ornamentation of verrucae coalescing in ridges and Y-shaped crests (Figure 7c–e). The lectotype of *T. campestre* (= *T. fimbriatum* var. *campestre*) forms a monophyletic clade with *T. fimbriatum* sensu stricto. As the morphology (Figure 8) also is congruent we regard them as synonymous. See also *T. winterhoffii*, with which this species has previously been confused.

Habitat and distribution. In dry and exposed habitats; both on calcareous and acidic soils, widespread in Europe reaching 60 °N in Scandinavia. Cosmopolitan distribution but in Europe often confused with the widely distributed *T. winterhoffii*.

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Figure 7. *T. fimbriatum*, epitype (GB): a basidiocarps b detail of spore sac showing the fibrillose-fimbriate mouth c–e spores. Scale bars: a = 10 mm; b = 2 mm; c–e = 1 μm

Figure 8. *T. campestre*, lectotype (NY): a–b basidiocarps c spore ornamentation under SEM. Scale bars: a, b = 10 mm; c = 1 μm.


Figure 2e


**Holotype.** ITALY, Trento: “juxta vias”, Bresadola (K).

Our sequence data confirm *T. fulvellum* as a well-defined species in Clade 8, as a sister species to *T. lloydii*. In agreement with Wright (1987), we consider *T. fulvellum* to be the valid name for this species.

**Habitat and distribution.** *Tulostoma fulvellum* is a species of humus-rich habitats in shaded to semi-shaded habitats. Reported from Europe (France, Germany, Italy, Slovakia, Spain, and Switzerland; Bataille 1910, Candoussau 1973, Jeppson 2008, Kabát 1987, Monthoux and Röllin 1974, Wright 1987) and Japan (Asai and Asai 2008).


Figure 2f


**Holotype.** ITALY, Trento: in Horto Cappucinarum, 1880, Bresadola (S!)  

Our sequence data confirm *T. giovanellae* as a well-defined species. Morphologically it is characterized by its finely verrucose spores with verrucae arranged in radial lines, as seen under SEM (Figure 2f). Some photos of the holotype collection at S (habit and spore ornamentation under SEM) were included in Altés et al. (1999). Since *T. volvulatum* is an older synonym, it takes priority over *T. giovanellae*. Art. 57 of the Melbourne Code allows an exception to the priority rule, preventing the use of a name that incurs in serious conflict with the taxonomic concept traditionally associated with it. Also, Art. 56 allows rejection of a name in that circumstance. A formal proposition will be put forward to the General Committee (to be published in Taxon) that the name *T. volvulatum* become a nomen rejiciendum, whereas *T. giovanellae* would be a conserved name to be used in its traditional sense.

A sequence from a collection labelled *T. caespitosum* var. *acaule*, originating from Patouillard’s herbarium (BPI), proved to be identical with *T. giovanellae*. See discussion (below) for issues around interpretation of *T. caespitosum* and *T. caespitosum* var. *acaule*. 
Habitat and distribution. Africa, Asia, and Europe. It is a typical Mediterranean species found on sandy or halophytic soils in more or less exposed sites, but it is also recorded from anthropogenic habitats in Austria, Hungary, and Germany (Bohus and Babos 1977, Kreisel 1984, Rimóczi et al. 2011).


Tulostoma grandisporum Jeppson, Altés, G. Moreno & E. Larss., sp. nov.
MycoBank number: MB819402
GenBank: KU519003 (ITS-LSU), KU843924 (Tef-1a).
Figure 9


Etymology. The name refers to the large spores.

Description. Spore-sac subglobose, 3–8 mm. Endoperidium hyphal, encrusting sand. Endoperidium greyish white to pale greyish ochraceous with some adhering sand grains, smooth. Mouth fibrillose-fimbriate, slightly mammose (Figure 9f). Socket inconspicuous. Stem slender, 15–23 x 1–2 mm, whitish to pale yellowish brown towards the base, which is slightly widened, covered by sand and provided with a distinct pseudorhiza (Figure 9a,e). Capillitium 3–7 μm in diameter, with irregular, undulating inner walls (Figure 9b). Septa scarce, not or only very slightly widened. Spores subglobose, 5.5–7.0 μm (av. 6.5 μm), smooth under light microscopy and SEM (Figure 9c, g–i).

Habitat and distribution. Recorded in exposed sand steppe habitats in Hungary and Slovakia. Usually not very numerous in its localities.

Notes. The species reminds of the new species T. pannonicum in its habit, but a check under the microscope is enough to distinguish T. grandisporum. Tulostoma leiosporum R.E. Fr., described from S. America, is similar to T. grandisporum in macro-morphology but differs in spore size (lectotype: 4.5–5.5 μm according to Moreno et al. 1997). The lectotype of T. leiosporum has the same type of ragged, undulating inner walls of the capillitium.

Other specimens examined. HUNGARY, Bács-Kiskun: Kerekegyháza, Kákás-Király tó, sandy pasture along wheel track, 16 Apr. 2009, U. Andersson, T. Gunnarsson, M. Jeppson 8924 (GB)*; Kiskunhalas, sandy grassland, 19 Jan. 2014, P. Finy 10 (GB)*; Lakitelek, Szikra, 300 m SE of railway station, along railway, on sandy path,
Figure 9. *T. grandidisporum*, holotype (M. Jeppson 8907, GB): a, e basidiocarps b capillitium c, g–i spores d basidiocarps (M. Jeppson 7735, GB) f detail of spore sac showing the fibrillose-fimbriate mouth; Scale bars: a, d–e = 10 mm; b–c = 10 μm; f = 4 mm; g–i = 1 μm.


*Tulostoma kotlabae* Pouzar, in Pilát, Fl. ČSR, B-1, Gasteromycetes: 598, 815. 1958. Figures 2g, 10

Figure 10. *T. kotlabae*, holotype (PRM 704203): a basidiocarps b spore ornamentation under SEM. Scale bars: a = 10 mm; b = 1 μm.

The studied material fits the original description as well as that of Wright (1987), and it is furthermore supported by ITS sequence data from the holotype. The eight closely related, but still unidentified, species (*T.* spp. 6-13) recovered in our phylogenetic analysis (Figure 1b) are in need of further investigation before they can be described as distinct and separate species.

**Habitat and European distribution.** *Tulostoma kotlabae* occurs in dry and exposed habitats such as steppe vegetation and sand dunes, preferably on calcareous, sandy soils. It is, to date, only known from Europe, from where it is on record from Denmark, France, Germany, Hungary, Lithuania, Poland, Russia (S part), Slovakia, Spain, and Sweden (Calonge 1998, Nitare 1988, Jeppson 2006a, 2008, Rimóczi et al. 2011, Rudnicka-Jezierska 1991, Winterhoff 2000, Wright 1987).

Figures 2h, 11

Holotype. USA (S!).
A characteristic species originally described from N. America. We have not sequenced the holotype, but its macro- and micromorphology (Figure 11a) are very similar to that of the collections here studied (Figure 2h).

Habitat and distribution. Tulostoma lloydii is recorded from N. America and Europe (Wright 1987). In Europe it has a Mediterranean distribution, having been reported from Italy and Spain. According to Wright (1987) it is a species of “forest soil, among humus and plant debris”. The studied material from Italy was collected in a garden, but there are also findings from more xeric environments on calcareous soil and on littoral sand dunes in Spain.


Figure 12

Holotype. PORTUGAL, Estremadura: Pinhal do Rei, 8 Jan. 1991, G. Almeida, F. Estrella, F.D. Calonge (LISU-MGA 8!)*.

Figure 11. T. lloydii, holotype (S): a spore ornamentation under SEM. Scale bar: 1 μm.

Figure 12. T. lusitanicum, holotype (LISU-MGA 8): a basidiocarp b spore ornamentation under SEM. Scale bars: a = 10 mm; b = 1 μm.
The species was invalidly published by Calonge and Almeida (1992) and later validated by Calonge (2000). ITS sequence data confirm it as a distinct species (Figure 1b).

**Habitat and distribution.** Recorded only from sand dunes in Portugal.


Figure 2i

**Holotype.** ITALY: in glabrosis prope Tridentinum, 1902, G. Bresadola (K).

This species is closely related to *T. squamosum*, *T. subsquamosum*, and the new species *T. calcareum*. It is distinguished by its dark brown to almost black stipe and a prominent dark peristome contrasting to the pale colours of the surrounding endoperidium in old and weathered specimens. In young basidiomata the endoperidium is more or less orange brown and the stem ± orange brown. The spores are verrucose-echinate and normally lack anastomoses and crests as seen under SEM (Figure 2i). It can be readily distinguished from *T. brumale*, which often grows in the same locations, by its darker stem and the more coarsely ornamented spores. Photos of the holotype collection at K (habit and spore ornamentation under SEM) were included in Altés et al. (1996).

**Habitat and distribution.** Grows in dry sandy grasslands, on sand dunes, sand steppes and rupicolous steppe slopes. According to Wright (1987) it is a typically European species. It is, however, also recorded from Central Asia (Dörfelt and Täglich 1990). It is widespread in Europe, reaching as far north as southernmost Fennoscandia.


Figure 2j

**Holotype.** SWEDEN, **Uppland**: Djurö, Munkö, 23 Sept. 1973, L.E. Kers 4264 (S!).

*Tulostoma niveum* is a well-defined species with small and slender, whitish basidiomata with conically protruding mouth and distinctly membranous exoperidium. Molecular data suggest a close relationship with the new species *T. eckbladii* and *T. giovanellae* (Figure 1c).

**Habitat and distribution.** Occurs among mosses on ± calcareous boulders, flat rocks, or stone walls in semi-shaded localities. Recorded from Finland, Norway, Sweden, Switzerland, and the UK (Scotland). Not known from outside Europe.


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**Tulostoma obesum** Cooke & Ellis, Grevillea 6: 82. 1878.

Figures 2k, 5c

**Holotype.** U.S.A., **Colorado**, Coke 2715 (K 39158); Isotype: (NY!)*

Sequenced Spanish collections were shown to be identical with the ITS sequence of the isotype from Colorada, USA (Cooke 2715, NY). With respect to morphology, *T. obesum* reminds of a species reported under *T. obesum* or *T. volvulatum* (*T. volvulatum* var. *obesum*) by Babos (1999), Calonge (1998), Hollós (1904), and Rimóczi et al. (2011) from the Iberian Peninsula and East Central Europe. However, according to molecular data, those records can be attributed to other species despite sharing a number of morphological features with *T. obesum*. SEM-photos for *T. obesum* show completely smooth spores (Figure 2k). Photos of the holotype and isotype collections at K and NY (habit and spore ornamentation under SEM) were included in Altés et al. (1999).

**Habitat and distribution.** A psammophilous species originally described from Colorado. In Europe it is recorded from littoral sand dunes and halophytic steppe habitats in SE Spain.

**Other specimens examined.** SPAIN, **Almería**: Cabo de Gata, La Salina, NW part, among halophilous vegetation on sand, 21 Apr. 2008, M. Jeppson 8695 (GB)*; **Almería**: Tabernas, Puente de los Callejones, among halophilous vegetation on sandy
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Tulostoma pannonicum Jeppson, Altés, G. Moreno & E. Larss., sp. nov.
MycoBank number: MB819403
GenBank: KU519010 (ITS-LSU)
Figure 13


Etymology. The name refers to Pannonia, an ancient Roman province in Central Europe, where this species was first collected.

Description. Spore-sac subglobose, 4–7 mm in diameter. Exoperidium indistinct to thinly membranous, incrusting sand, remaining as a cup-like structure at the base of the spore-sac (Figure 13d). Endoperidium greyish white, smooth, or pitted from detached sand grains. Mouth indistinct to fibrillose-fimbriate (Figure 13f). Socket inconspicuous, slightly separated. Stem slender, 10–20 x 1–2 mm, ochraceous to pale greyish, smooth, somewhat longitudinally furrowed, with a slightly widened base and a distinct pseudorhiza (Figure 13a,d). Mature gleba ferrugineous. Capilltium 3–5 μm in diameter with even inner walls and abundant, strongly widened septa (Figure 13b). The capillitium easily breaks up at the septa. Spores subglobose to broadly ellipsoid, 3.5–4.5 μm (av. 4.0 μm), smooth. Under SEM the spore wall is smooth to slightly rugulose (Figure 13g-i).

Habitat and distribution. The species is recorded from exposed sand steppe habitats on calcareous soil in Hungary.

Notes. Rimóczi et al. (2011) recorded this species from Hungary as T. leiosporum and noted that the spores were slightly smaller than those of the lectotype of T. leiosporum, deposited in Herbarium S. Some photos of the lectotype collection of T. leiosporum (habit and spore ornamentation under SEM) were included in Moreno et al. (1997).


Figure 13. *T. pannonicum*, holotype (M. Jeppson 7764, GB): a, d–e basidiocarps b capillitium c, g–i spores f detail of spore sac showing the fibrillose-fimbriate mouth (M. Jeppson 7803, GB). Scale bars: a, d–e = 10 mm; b–c = 10 μm; f = 5 mm; g–i = 1 μm.

*Tulostoma pseudopulchellum* is a small and slender species with a macro-morphological resemblance to *T. pulchellum*. It differs from that species by having significantly smaller basidiomata and irregularly rugulose spores (SEM). It has a distinctly membranous exoperidium and a fibrillose-fimbriate mouth. Moreno et al. (1992b) suggested a close relationship with *T. pulchellum* but this could not be confirmed in the molecular analyses.

**Habitat and distribution.** So far only known from its type locality in dry gypsum slopes at Alcalá de Henares in Central Spain.

Figures 2l, 14, 15


**Holotype.** AUSTRALIA, Tepper (PAD).

This is a characteristic species with a distinct membranous exoperidium, a snow-white endoperidium and a ± delimited, fibrillose-fimbriate mouth. The ornamentation of the spores is composed of a dense cover of low verrucae, sometimes coalescing into ridges. Collections with a ridged type of ornamentation have been distinguished as *T. hollosii* and *T. pulchellum* var. *subfuscum* (cfr. Moreno et al. 1992b, Specht and Schubert 2012). Our sequence data based on specimens with isolated verrucae as well as on those with distinct ridges (Figure 2l), a paratype of *T. hollosii* included (Figure 15), indicate conspecificity. Our molecular data suggest *T. pulchellum* to be closely related to *T. striatum* (Figure 1b).

**Habitat and distribution.** *Tulostoma pulchellum* has a cosmopolitan distribution, occurring in exposed, dry, and warm habitats. In Europe it is on record from the Czech Republic, Germany, Hungary, Italy, Romania, Slovakia, and Spain (Kreisel 2001, Kříž 2011, Specht and Schubert 2012, Sarasini 2005). It is here confirmed also from Central Asia (Mongolia).


Figures 5d, 16a, 17, 18


The first European records of this traditionally North American species are here confirmed. The species can easily be confused with *T. fimbriatum*, but is distinguished by the somewhat smaller spores that are more ornamented with a verrucose-spiny ornamentation without ridges (Figure 16a). Study of the macro-micromorphology and sequence data of the type collections of *T. punctatum* (Figure 17) and *T. subfuscum* (Figure 18), indicate conspecificity. It is surprising to see that the spore ornamentation of the type of *T. subfuscum* is very different from the expected one (cfr. description and illustrations by Wright 1987). The presence of distinctive punctate pits on the
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**Figure 16.** Spore ornamentation under SEM of European species (cont.): a *T. punctatum* (M. Jeppson 7472, GB) b *T. simulans* (M. Jeppson 9302, GB) c *T. squamosum* (AH 15483) d *T. striatum* (AH 15543) e *T. subsquamosum* (M. Jeppson 3475, GB) f *T. winterhoffii* (M. Jeppson 2379, GB). Scale bars: 1 μm.

**Figure 17.** *T. punctatum*, lectotype (BPI 729033): a basidiocarps b spore ornamentation under SEM. Scale bars: a = 10 mm; b = 1 μm.

The endoperidium of *T. punctatum*, as described by Peck (1896), could not be observed in the European specimens. A somewhat pitted appearance of the endoperidium is a common, but inconsistent, feature among species of *Tulostoma*, and of little taxonomic value. The pits are scars or depressions caused by sand grains adhering to the peridium in early stages of development.

**Habitat and distribution.** Originally described from the USA (Kansas). In Europe on sandy soil on exposed, anthropogeneous sites (e.g. church lawns/car parks, abandoned orchards) in SW Slovakia.

**Other specimens examined.** SLOVAKIA, Záhorská nížina: Vel’ke Leváre, by the church, grassy road side on sandy soil, 2 Oct. 2005, J. Jeppson, M. Jeppson 7472
(GB)*; Stupava, in dry sandy grassland in an abandoned orchard, 16 Oct. 2014, T. Knutsson, E. Larsson, R. Bednar, V. Kautman, I. Kautmanová, A. Košuthová, M. Jeppson 10058 (GB)*.


Figures 5e–f, 16b, 19


**Holotype.** USA, Texas: Denton, Long (herb. Lloyd 13636, BPI 704611).

As was indicated already by its author (Lloyd 1906), this species “simulates” other species of *Tulostoma*, particularly *T. brumale*, from which it differs by having a hyphal exoperidium and larger spores. According to our molecular results it is known from Austria, Hungary, Spain, Sweden, and the UK (a British sequence was kindly provided by Martyn Ainsworth, Kew, but is not shown in the tree). Sequence data of the European specimens, labelled as *T. beccarianum*, *T. brumale*, and *T. moravecii* (isotype, Figure 19), cluster with a paratype in PC of *T. simulans*. Unfortunately the holotype material of *T. simulans* was not available for DNA extraction.

SEM-photos of the collections used in our study show irregular, rather dense, broadly based conical verrucae (Figure 16b), thus closely coinciding with the holotype material of *T. simulans* (cfr. Moreno and Altés 1992). Altés and Moreno (1993) synonymized *T. simulans* with *T. beccarianum* based on morphological similarities, but sequence data show that they should be regarded as separate species.

**Habitat and distribution.** N. America, S. America, Australia, New Zealand, Europe, and Asia (Kazachstan). It is here confirmed also from eastern Russia (Altay Republic). It has been found in a wide range of habitats (“forest soil, among litter, sometimes dunicolous, tree nurseries” according to Wright 1987).

European occurrences are from sand steppe habitats, sand dunes, gypsum hills, rupicolous steppe meadows, and urban plantations.

Remarks. Tulostoma squamosum is morphologically well characterized by its brown, rather tall and squarrose stem combined with a brownish warty exoperidium that upon
dehiscence leaves a distinct reticulate pattern. The exoperidium is composed of brownish sphaerocysts. The spore ornamentation under SEM (Figure 16c) shows spines fused at their tops into small pyramidal groups, sometimes coalescing to form short crests. In Scandinavia the name *T. squamosum* has previously been associated with the proposed new species *T. calcareum*, described above (cfr. ArtDatabanken 2015, Nitare 1997),

**Habitat and distribution.** Usually found among herbs in semi-shaded places, including open deciduous forests and gardens. Taking into account its recently established synonyms *T. mussooriense* and *T. verrucosum* (Moreno et al. 1992a), it has a cosmopolitan distribution. It seems to be one of the more widespread *Tulostoma* species in Europe, reaching north to Germany (Kreisel 1984).


Figure 16d


As was noted by Cunningham (1925), this species (originally described from Australia) strongly reminds of *T. pulchellum* (as *T. poculatum*), but can be readily distinguished by its striate spores (Figure 16d). The molecular results confirm it is a distinct species (Figure 1b).

**Habitat and distribution.** It is widespread in the Americas, South Africa and Australia (Wright 1987). Altés and Moreno (1991) recorded it from Europe (Spain), where it to date is only known from the city of Madrid. Asai (2004) reported it from Japan and it is now also confirmed to occur in Mongolia. According to Wright (1987) and Calonge (1998) it is a psammophilous species.


Figures 5g, 16e

**Holotype.** INDIA, Gurdaspur: Jaggatpur, Aug. 1938, S. Ahmad 258 (BPI).

*Tulostoma subsquamosum* is characterized by a hyphal exoperidium with scattered sphaerocyst-like cells. The spores are verrucose-echinate with a tendency to form a
subreticulum (Figure 16e). It seems to be one of the more common species in the sand steppe vegetation of Central Hungary, from where it was traditionally recorded as *T. squamosum* (cfr. Hollós 1904, Rimóczi et al. 2011). In the molecular analyses it comes out as a sister species to *T. squamosum* (Figure 1c).

**Habitat and distribution.** Occurs in dry grasslands on sandy-clayish soil. It has a wide distribution in Asia (India, Pakistan), N and S America and Europe (Hungary, Slovakia, Spain).


Figures 5h, 16f, 20


*Tulostoma winterhoffii* was recently described from Germany. The holotype was not available for sequencing in this study but a paratype, provided by its authors, was sequenced and found to match sequence data of the type material of *T. fimbriatum* var. *heterosporum*. In macro-morphology (Figure 5h) *T. winterhoffii* strongly reminds of *T. fimbriatum* but important features to distinguish it are the verrucose-echinate spores (no anastomoses seen under SEM; Figure 16f) and the variation in spore size (Wright 1987: up to 9.7 μm in diameter; Specht and Schubert (2013): (4.5–) 5.5–8.5 (–10.3) μm including ornamentation). Our own material shows a similar variation, averaging 5.7 – 6.1 μm, some spores reaching 9 μm. The same can be observed in the type collection of *T. fimbriatum* var. *heterosporum* (Figure 20). Based on morphology
Figure 20. *T. fimbriatum* var. *heterosporum*, holotype (L 64172): a basidiocarps b spore ornamentation under SEM; spore size variation. Scale bars: a = 10 mm; b = 1 μm.

and sequence data, *T. fimbriatum* var. *heterosporum* must thus be considered a distinct species, for which the name *T. winterhoffi* has priority.

**Habitat and distribution.** Mostly in dry and sandy habitats in exposed places, on more or less calcareous soils. Currently known from the Czech Republic, Denmark, France, Germany, Hungary, the Netherlands, Norway, Spain, and Sweden. In Scandinavia it reaches 60° N. The world distribution comprises Europe, Asia (Caucasus), and N. America (California) according to Wright (1987; as *T. fimbriatum* var. *heterosporum*).


**New species identified in this study that lack scientific names**


Tulostoma sp. 4: HUNGARY, Budapest: Bikás, sandy grassland, 2 Nov. 2011, P. Finy 12 (GB)*.
Tulostoma sp. 11: SPAIN, Madrid: Casa de Campo, 14 Nov. 1988, M. Jeppson 881114 (GB)*.

Tulostoma sp. 17: SPAIN, Madrid: Alcalá de Henares, junto al cerro Malvecino, terreno margoso con algo de yeso, 7 Nov. 1990, equipo micología UAH (AH 13674, GB)*.


Tulostoma sp. 19: CYPRUS, 14 Feb. 2014, T. Læssøe, A. Molia 140115-2 (GB)*.

Tulostoma sp. 20: SPAIN, Zaragoza: Los Monegros, Sástago, along road Escatrón-Sástago, 500 m N of Ebro-bridge, dry, sandy calcareous soil, 27 Apr. 1999, S. Jeppson, J. Jeppson, M. Jeppson 5015 (GB)*.

Tulostoma sp. 21: SPAIN, Madrid: Alcalá de Henares, cerros de Alcalá, 3 Nov. 1989, (as T. moravecii, AH 11698)*.

Type specimens of species examined, from outside of Europe

Tulostoma albicans V.S. White: USA, Texas, 1893, E.D. Cope (holotype, NY)*. Figure 21.

Tulostoma egranulosum Lloyd: AUSTRALIA, Victoria, Melbourne: D. McAlpine (holotype, herb. Lloyd 15424, BPI 704420, GenBank KC333072).


Tulostoma leiosporum R.E. Fr.: ARGENTINA, Jujuy: Pampa Blanca, 23 May 1901, R. Fries 64 (lectotype, S).

Tulostoma lysocephalum Long: USA, New Mexico: on highway 70, 10 mi. W. of Deming, 12 Sept. 1941, W.H. Long, D.J. Stouffer (holotype, herb. Long 9639, BPI 749235)*. Figure 22.


Tulostoma readeri Lloyd: AUSTRALIA, Reader (holotype, herb. Lloyd 15421, BPI, GenBank KC333076).

Tulostoma rufum Lloyd: USA, Alabama: Spring Hill, A.S. Berrolet (lectotype, herb. Lloyd 25542, BPI 704578)*. Figure 23.


Tulostoma xerophilum Long: USA, Arizona: 7 mi from Nogales, on Hwy 89, 13 Nov. 1936, Long & Stouffer (holotype, herb. Long 9688, BPI 802484)*. Figure 24.
Discussion

In this study, we found that the species diversity of *Tulostoma* was unexpectedly high, particularly from the arid areas of the world. The majority of the recovered 21 unidentified clades from the phylogenetic analysis (Figure 1a–c) comprise sequence data...
from specimens collected in dry arid areas of Europe, notably Hungary and Spain. In Europe, the sand steppe vegetation of Hungary has been shown to be exceptionally species rich in gasteroid fungi (Hollós 1904, Rimóczi et al. 2011). The sand steppes of Hungary constitute a western outpost of the Euroasian steppe zone and in this study 14 of the included identified species of *Tulostoma* occur there, along with nine of the hitherto unnamed species. Thus, we confirm the steppe vegetation as a species diversity hot spot for gasteroid fungi. Another identified area with high species diversity is the arid vegetation in the central part of the Iberian Peninsula, where 11 of the identified species and six of the hitherto unnamed species occur.

But species of *Tulostoma* are not restricted to arid regions only. Studies in the Tropics and temperate woodlands have indicated the presence of new species, growing in shaded, humus-rich, or even humid conditions (Hernández-Caffot et al. 2011). In Europe, there are also a few species that are adapted to grow in shaded and more humid conditions as *T. fulvellum*, *T. lloydii*, *T. niveum*, and *T. squamosum*.

In contrast to several recent phylogenetic studies of the Basidiomycota, where the traditional generic concepts of morphologically similar groups have shown to be polyphyletic and resulted in a number of new genera and higher taxa (e.g. Lodge et al. 2014, Örstadius et al. 2015, Miettinen et al. 2016, Petersen and Hughes 2016), *Tulostoma* is here confirmed as monophyletic. However, the recovered tree topology is not in congruence with the current infrageneric classification of *Tulostoma* (Pouzar 1958, Wright 1987), suggesting that many of the morphological characters used for delimitations are plesiomorphic or homoplasious. The endoperidial mouth region and the exoperidium, the major delimiting characters for the infrageneric classification when based on morphology, are here shown to be incongruent with the phylogeny, as the different types are distributed among the major clades. On the other hand, endo- and exoperidial features, as well as the spore morphology, are here confirmed and shown to be important characters for the delimitation of species.

Wright (1954) and Wright et al. (1972) listed morphological features in *Tulostoma* and classified their taxonomic importance as “primary” and “secondary”. Wright (1987) divided the genus in two subgenera (*Tulostoma* and *Lacerata*), based on features of the mouth. The former subgenus was divided in two series (*Brumalia* and *Fimbriata*), each of them containing five sections. In our analyses Clade 3 corresponds more or less with the section *Fimbriata*, and a supported branch in Clade 9 corresponds partly to section *Poculata*. However, species belonging to Wright’s sections *Brumalia* and *Hyphales* are distributed amongst the major clades and occur in Clades 3, 4, 5, 9, and 10 (Figure 1a–c). A new infrageneric classification of *Tulostoma* is necessary, but should be done in the light of further taxon sampling, sequence data, and analyses.

Several of the clades include groups with morphologically cryptic speciation, that are distinguished by the molecular data, but for which discriminatory morphological features are ambiguous or lacking. This is particularly evident in Clades 2, 3, 5, and 10, where a number of unidentified taxa from xerothermic habitats were recovered.

The phylogenetic analyses recovered and confirmed the classical, well-known species from Europe, namely *T. brumale*, *T. cyclophorum*, *T. fimbriatum*, *T. fulvellum*, *T.
giovanellae, T. kotlabae, T. lloydii, T. melanocyclum, T. pulchellum, and T. squamosum. In addition, one species not previously known from the continent – T. punctatum – was identified. Further Tulostoma calcareum, T. calongei and T. eckbladii are described as new to science. The former has a wide distribution in Europe whereas the two latter are hitherto only known from Central Spain and northernmost Norway respectively.

We also discovered species with misapplied names and two species new to science were identified from the continental sand steppe vegetation in East Central Europe: T. grandisporum and T. pannonicum. These had previously been misinterpreted as T. leiosporum (Jeppson et al. 2011, Rimóczi et al. 2011). The name T. obesum has in current use been applied for a species found in dry sandy habitats in East Central Europe as well as in dry steppic sites on the Iberian Peninsula (Altés et al. 1999, Rimóczi et al. 2011). It is characterized by having smooth spores and a large, whitish spore-sac that opens lacerately at maturity. In this study an ITS sequence from of the isotype of T. obesum (from Colorado, USA) was generated and included in the analyses. It did not form a clade with the European concept of T. obesum but with a hitherto unnamed species in Clade 2, known from sand dune habitats of SE Spain. Thus we here confirm T. obesum from Europe, but with a different species interpretation from that of previous European mycologists (Altés et al. 1999, Rimóczi et al. 2011).

The Hungarian and Spanish collections, previously attributed to T. obesum, are placed in Clade 6 and are here treated as T. aff. cretaceum. This clade consists of a complex of closely related cryptic species with a strong geographical signal. In East Central Europe and Central Asia the name T. volvulatum I.G. Borshch. has been misapplied to cover this species complex (e.g. Hollós 1904, Shvartsman and Filimonova 1970). As demonstrated by Altés et al. (1999), the holotype of T. volvulatum was based on a specimen of T. giovanellae, with typical ornamented spores. Under the name T. volvulatum, Hollós (1904) recorded T. giovanellae (e.g. S, F265540) but also specimens with smooth spores (Babos 1999) that closely match Sorokin’s (1890) interpretation of T. volvulatum from the Karakoum Desert (Sorokin 1890, plate XXV, Figure 353a). A number of poorly known species described from desert habitats in northern Africa (e.g. T. barbeyanum Henn., T. boissieri Kalchbr., T. giolianum Bacc., and T. ruhmerianum Henn.) as well as the American T. cretaceum Long and T. meristostoma Long, share characters typical of this clade and need to be included in an extended study, which should also deal with several recently described species from China (T. brevistipitatum B. Liu, Z.Y. Li & Du, T. sabulous B. Liu, Z.Y. Li & Du, and T. helanshanense B. Liu, Z.Y. Li & Du). Calonge (1998) identified Spanish material attributed to T. volvulatum as Schizostoma laceratum (Ehrenb.: Fr.) Lév. Based on sequence data, Gube (2009) recovered S. laceratum as closely related to the type of T. cretaceum. As pointed out by Kuhar et al. (2012), the whole complex is in need of a detailed study comprising morphological as well as molecular data. Similarly, the status of the genus Schizostoma is unclear and calls for further study.

Tulostoma simulans is here identified from Europe. According to our data it seems to have a wide distribution but appears to be rare. The species is known from East Central Europe under the name T. moravecii, and it may have been overlooked due to its resemblance with T. brumale.
The application of the name *Tulostoma caespitosum* Trab. is an unsolved problem. The species was described from northern Africa and later reported from the Americas, Asia, and Europe (Calonge 1998, Calonge and Wright 1988, Wright 1987). Among its main characteristics, according to Wright (1987), are the distorted basidiomata and the sometimes caespitose growth. Wright (1987) did not find any collection that he considered to be suitable as a holotype and suggested a neotype, collected by Trabut in Algeria, coinciding with the protologue of *T. caespitosum* (*T. caespitosum* var. *acaule*, herb. Patouillard 1422, FH; unpublished herbarium name). However, Wright did not formally designate a neotype. An authentic collection of *T. caespitosum* var. *acaule*, from Algeria (leg. Trabut), is kept in BPI (ex herb. Patouillard, herb. Lloyd 15427, BPI 704324) but was unfortunately found to be in poor condition and too scanty for DNA sequencing. A second collection from Africa (indet. region) in BPI (Patouillard 4639, herb. Lloyd 15426, BPI 704323) was sequenced for the present study and found to be identical with *T. giovanellae*, in Clade 10, Figure 1c. The spore ornamentation of this specimen is identical to *T. giovanellae*. However, in collection BPI 704324 the spores were shown to be strongly verrucose, thus differing from *T. giovanellae*. The typification of *T. caespitosum* thus remains unsettled and the unclear interpretation of this name calls for further study.

Kreisel and Al-Fatimi (2008) observed that gasteroid basidiomycetes from desert habitats tend to produce smooth spores. This stands in contrast to taxa growing in vegetation types that are regularly exposed to rainfall – these usually have more or less distinctly ornamented, verrucose–echinate spores. Kuhar et al. (2012) found that the relative abundance of species with smooth spores within the study in Argentina reached a maximum under desert conditions. The relative abundance of species with ornamented spores was shown to go up with increasing humidity. Their hypothetical explanation was that the spore wall ornamentation increases the water repellency, favouring a wind-blown dispersal (Kreisel and Al-Fatimi 2008).

In this study, species of *Tulostoma* with smooth or sub-smooth spores are found to occur mainly in arid, and exposed habitats in East Central Europe and Spain (*T. aff. cretaceum, T. grandisporum, T. obesum, T. pannonicum*). Species with moderately to strongly ornamented spores (*T. beccarianum, T. brumale, T. calongei, T. giovanellae, T. melanocyclum, T. punctatum, T. simulans, and T. subsquamosum*) are found in more steppic, dry grassland communities in southern and east central Europe. In semi-shaded or shaded to somewhat humid habitats, the genus is represented by species with ornamented spores (*T. cyclophorum, T. niveum, and T. squamosum*) but also a few with completely smooth spores (*T. fulvellum and T. lloydii*). Among the species regularly encountered in northern and north-western Europe, i.e. in sub-oceanic or oceanic climates, all have moderately to distinctly ornamented spores (*T. eckbladii, T. brumale, T. calcareum, T. fimбриatum, T. kotlabae, T. melanocyclum, T. niveum, T. simulans, T. squamosum, and T. winterhoffii*). In accord with the observations by Kreisel and Al-Fatimi (2008), *Tulostoma* species with smooth or sub-smooth spores tend to occur mainly in dry and arid habitats whereas those with distinct and strongly ornamented spores grow in more humid habitats.
Unexpected high species diversity among European stalked puffballs - a contribution...

Due to difficulties to determine *Tulostoma* specimens to species level, many published records and herbarium collections are based on misinterpretations. It is therefore problematic to evaluate species distribution data. Some seem to have wide, more or less cosmopolitan distribution ranges (e.g. *T. fimbriatum*, *T. pulchellum*, *T. striatum*, *T. squamosum*, and *T. subsquamosum*). Others appear to be restricted to certain geographical areas or habitat types (e.g. *T. grandisperum*, *T. pannonicum*, *T. niveum*, and *T. eckbladii*). In Europe most species have southern and southeastern distributions. The number of species decreases towards the north. *Tulostoma brumale*, *T. calcaratum*, *T. fimbriatum*, *T. kotlabae*, *T. melanocyclum*, *T. simulans*, and *T. winterhoffii* reach the south of Fennoscandia, whereas *T. niveum* is regularly found between 55° and 62° N, from southern Finland westwards to northern Scotland. The report herein of *T. niveum* from near Geneva in Switzerland is a significant range extension. The newly described species, *T. eckbladii*, is, as far as currently known, limited to areas above the Arctic Circle.

A number of species of *Tulostoma* occur in habitat types that are under the threat of changing agricultural methods, anthropogenic exploitation, and construction works. Hernández-Caffot et al. (2011) described *T. domingueziae* from threatened woodland ecosystems in Argentina. In Europe, several *Tulostoma* species are on the red-list of threatened species in the countries where they occur: *T. brumale*, *T. fimbriatum*, *T. kotlabae*, *T. melanocyclum*, *T. simulans* (as *T. moravecii*), *T. pulchellum*, and *T. niveum* (ArtDatabanken 2015, ArtsDatabanken 2015, Holec and Beran 2006, Rassi et al. 2010). For *T. niveum*, national action plans for its conservation have been elaborated and become effective in Great Britain and Sweden (BAP 2005, Jeppson 2005, 2006b). *Tulostoma niveum* has also been nominated for the IUCN global red-list (http://iucn.net/ekoo.se/iucn/species_view/325133) and *T. aff. cretaceum* (as *T. volvulatum*) is legally protected in Hungary (Siller et al. 2005). Many species of *Tulostoma* can be considered as indicators of valuable and threatened habitats.

In this study we show, based on molecular data, that the species diversity of *Tulostoma* is much higher than previously known. Much of this detected diversity is associated with dry, arid areas, such as the steppe vegetation in Hungary. This suggests that the number of species is likely to increase even more as other dry areas of the world are explored.

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Unexpected high species diversity among European stalked puffballs - a contribution...

Supplementary material I

Unexpected high species diversity among European stalked puffballs – a contribution to the phylogeny and taxonomy of the genus Tulostoma

Authors: Mikael Jeppson, Alberto Altés, Gabriel Moreno, R. Henrik Nilsson, Yolanda Loarce, Alfredo de Bustos, Ellen Larsson

Data type: Table/specimen data

Explanation note: Data of specimens sequenced in this study with GenBank accession numbers for the ITS/LSU and TEF-1α regions. Type specimens are marked in bold.

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Supplementary material 2

Unexpected high species diversity among European stalked puffballs – a contribution to the phylogeny and taxonomy of the genus Tulostoma

Authors: Mikael Jeppson, Alberto Altés, Gabriel Moreno, R. Henrik Nilsson, Yolanda Loarce, Alfredo de Bustos, Ellen Larsson

Data type: Table/sequence data

Explanation note: Distance matrices of the ITS alignment (uncorrected “p” distance and GTR distance) and the multiple sequence alignment of the ITS.

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