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A new species of Volvariella from India

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ABSTRACT—Volvariella sathei sp. nov., collected from Agharkar Research Institute Campus, Pune, is described, illustrated, and compared with morphologically closely related taxa of Volvariella and Volvopluteus. Diagnostic characters of the new species are the large white fruitbodies, relatively small basidiospores, and cylindric-clavate pleuro- and cheilocystidia. Neighbour-Joining analysis based on ITS sequences confirmed that it is distinct from its closest relative, V. nivea and other morphologically similar Volvariella species. A key to the Indian species of Volvariella and Volvopluteus is provided.

KEY WORDS-Agaricales, Basidiomycetes, macrofungi, taxonomy

Introduction

Volvariella Speg. (*Agaricales*) is well represented in India, and considerable work has been carried out by Pathak et al. (1978) and Pradeep et al. (1998), who have studied the genus and described and/or reported several species for the country. Agaric checklists compiled by Manjula (1983), Natarajan et al. (2005), and Kumaresan (unpublished list) cite 22 species of *Volvariella* including *Volvopluteus* Vizzini et al. as reported from India. A white agaric from Pune is here described as a new *Volvariella* species based on its morphological and molecular characters. Its phylogenetic position has been determined by ITS1+5.8S rDNA+ITS2 sequence analyses.

Materials & methods

Thin handmade sections were made from dried specimens, revived in 10% KOH, and stained in 2% Phloxine. The microscopical characters were studied using an Olympus U-CMAD3 microscope. Approximately 30 basidiospores from sections were measured; the range (extreme values in parentheses) precedes mean spore measurements (in parentheses). Camera lucida diagrams were made using Nikon E200 attached with Nikon Y-IDT prism. Colour terminology follows Kornerup & Wanscher (1978). The type specimen is deposited at Ajrekar Mycological Herbarium (AMH), MACS' Agharkar Research Institute, Pune, India.

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TABLE 1. Sequences used in the phylogenetic analyses.

Species	Voucher No.	GenBank No.
Volvariella bombycina	AJ244	нм562212.1
V. caesiotincta	MA54717	нм562211.1
V. dunensis	SCM3513	JF415140.1
V. hypopithys	TO AV137	нм246492.1
V. lepiotospora	AJ155	нм562214.1
V. nivea	gdgm25489	FJ749127.1
V. nullicystidiata	SP393639	EU920671.1
V. pusilla	AJ51(LOU)	JF415137.1
V. pusilla	TO AV139	нм246494.1
V. sathei	амн 9436	JN792550
V. strangulata	TO AV141	нм246493.1
V. surrecta	AJ55	нм562213.1
V. taylorii	AJ54	нм562210.1
V. terrea	LUG11010	JF415141.1
V. volvacea	TO AV143	нм246500.1
V. volvacea	ATCC MYA-4696*	HQ999973.1
Volvariella sp.	LOU18924	JF415139.1
Volvopluteus earlei (= V. acystidiata)	TO HG1973	нм246499.1
Vp. earlei (= V. cookei)	TO AV133	нм246496.1
Vp. earlei (= V. earlei)	to av134	нм246497.1
Vp. earlei (= V. media)	TO HG2001	нм246498.1
<i>Vp. gloiocephalus (= V. gloiocephala)</i>	to av136	нм246495.1
Schizophyllum commune	IFM 56967*	AB566277.1
*Church and		

*Strain no.

DNA isolation methodology followed Aljanabi & Martinez (1997) with slight modifications. ~300 mg of the holotype gills was washed thoroughly with 99% ethanol and water. DNA was extracted using high salt extraction buffer (NaCl, Tris. HCl and EDTA) and addition of one volume of phenol:chloroform:isoamyl alcohol (25:24:1) and finally precipitated with isopropanol, washed with 70% ethanol, and suspended in Tris-EDTA buffer. The protocol outlined by White et al. (1990) was followed for polymerase chain reaction in Eppendorf Master cycler (Eppendorf, Germany) with universal primers ITS5 and ITS4. Sequencing was performed using BigDye Terminator Cycle Sequencing Kit (Applied Biosystem, UK) as per manufacturer's instructions. The cleaned PCR product was run on ABI 3100 automated DNA sequencer. The raw sequences obtained from ABI sequencer were manually corrected for inconsistencies with the help of Chromas lite software. The DNA sequences obtained in Fasta format were subjected to BLASTn (http://www.ncbi.nlm.nih.gov) analysis.

For phylogenetic analysis, additional related sequences obtained from NCBI DNA sequence database (TABLE 1) were aligned by CLUSTAL W to generate a phylogenetic tree using MEGA v.5.0 software (Neighbour-joining using Kimura-2 parameter model). The outgroup comprised *Schizophyllum commune* and *Volvopluteus* spp. (recently transferred from *Volvariella*).



PLATE 1. *Volvariella sathei*: Basidiomata under natural conditions in Agharkar Research Institute campus. A. Surface view of basidiome. B. Gill view of basidiome. Photo Senthilarasu G.

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Taxonomy

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Differs from *Volvariella nivea* in smaller subglobose to ovoid basidiospores and smaller pleuro- and cheilocystidia.

TYPE: India, Maharashtra State, Pune, Agharkar Research Institute Campus (18°52′N 73°83′E), on ground (soil), grass field, 03.08.2009, coll. G. Senthilarasu (**Holotype** AMH 9436; GenBank JN792550).

ЕтумоLOGY: The epithet honors Dr. A.V. Sathe, a leading mycologist in western India.

PILEUS 8–9 cm diam., convex; surface white to yellowish white (4A2), fibrillose, shiny with fine radial striations in the margin; MARGIN thin, eroded; LAMELLAE free, remote, white becoming sordid, ≤ 10 mm wide, crowded with lamellulae of different lengths, margin concolorous; STIPE 80–105 × 9–13 mm, cylindric, distinctly compressed, tapering towards apex, solid; surface white to cream, glabrous; PILEAL CONTEXT 7 mm thick near stipe, white, unchanging on bruising; VOLVA free from the stipe, white (4A2), irregularly lobed, thick; ODOR fungus like; TASTE not recorded.

Basidiospores $(5.5)6-6.5 \times (4.5)5-5.5$, $(5.97 \pm 0.26 \times 5.05 \pm 0.26) \mu m$, Q = 1.1, subglobose to ovoid, hyaline with a stramineous thickened wall, smooth; BASIDIA 20-36×5.5-11 µm, cylindric clavate to clavate, tetrasporic; STERIGMATA short ($\leq 2 \mu m$), slender; LAMELLA EDGE heteromorphous; CHEILOCYSTIDIA scattered, 20–62 \times 8–33 μ m, cylindric-clavate with broadly rounded apex, thin-walled, easily collapsing; PLEUROCYSTIDIA scattered, $27-72 \times 15-34 \mu m$, cylindric-clavate, mucronate to broadly rounded apex, thin-walled; LAMELLA TRAMA convergent, consisting of thin walled, hyaline hyphae 5-18 µm diam.; SUBHYMENIUM little developed up to 20 µm wide, pseudoparenchymatous; PILEAL SURFACE a regular cutis consisting of radially repent to semi erect hyphae, thin-walled, 5–30 µm diam.; PILEAL CONTEXT consisting of tightly interwoven hyphae, 5-30 µm diam., thin-walled; VOLVA consisting of gelatinous layer, up to 50 µm thick, hyphae thin-walled, up to 33 µm diam., volval elements 20-47 \times 15–24 µm, cylindric-clavate; STIPITIPELLIS hyphae made of cylindric, septate hyphae, 5–30 μ m diam., terminal elements 30–45 × 12–13 μ m, with rounded apex, caulocystidia absent; CLAMP-CONNECTIONS absent in all hyphae.

ADDITIONAL SPECIMENS EXAMINED: INDIA, MAHARASHTRA STATE, Pune, Agharkar Research Institute Campus (18°52'N 73°83'E), on ground (soil), grass field, 21.06.2009, coll. G. Senthilarasu (AMH 9443).

Justo et al. (2010, 2011) reassessed species traditionally assigned to *Pluteaceae* Kotl. & Pouzar based on molecular data. *Volvariella* has been placed outside the pluteoid clade with a new genus, *Volvopluteus*, erected to accommodate species with an ixocutis pileipellis and spores > 11 µm long. Consequently, *Volvariella gloiocephala*, *V. earlei*, and *V. acystidiata* have been transferred to



 $\label{eq:plate 2.} PLATE 2. Volvariella sathei: a. Basidiospores. b. Basidia. c. Cheilocystidia. d. Pleurocystidia. e. Volval elements. f. Terminal elements of stipitipellis. Scale bar = 10 \mu m.$

Volvopluteus. Volvariella acystidiata is considered now a form, as *Volvopluteus earlei* f. *acystidiatus* (N.C. Pathak) Vizzini & Contu (Justo et al. 2011).

Of the 20 species described from India, two belong in *Volvopluteus* (*Vp. gloiocephalus* (DC.) Vizzini et al. and *Vp. earlei* (Murrill) Vizzini et al.), and the remaining species belong in *Volvariella. Volvariella sathei* is compared here with morphologically similar taxa belonging to both genera.

Volvopluteus earlei, which also produces white basidiomata, differs in smaller (3–5 cm diam.) size and longer (>11 µm) spores (Justo & Castro 2010; Justo et al. 2010). *Volvopluteus earlei* f. *acystidiatus*, originally described by Pathak (1975) and later by Vizzini & Contu (2010), also closely resembles *V. sathei* but clearly differs by its smaller (20–30 mm diam.) yellowish tinged pileus with pinkish margin, shorter more slender stipe (30–40 × 2–3 mm), larger spores (10.5–16.5 × 7.5–10.5 µm) and basidia (45–70 × 10–13.5), and absence of hymenial cystidia (Justo et al. 2010). Phylogenetically *Volvopluteus* species are clearly distant from *Volvariella* species, including *V. sathei*. There are certain regions within ITS including 5.8S that distinguishes *Volvariella* from *Volvopluteus*.

Volvariella sathei is similar to *Volvariella pusilla* (Pers.) Singer (Pegler 1977, 1983, as *V. parvula*) in the white basidiomata with fibrillose pileus and remote lamellae. However, *V. pusilla* differs in stature with smaller pilei (1.5–3 cm diam.) and stipes (10–40 × 2–4 mm) as well as producing smaller basidia (17–29 × 5–7.5 µm), pleurocystidia (36–53 × 11–17 µm), and cheilocystidia (25–40 × 9–16 µm).

Volvariella nivea T.H. Li & Xiang L. Chen described from China (Li et al. 2009) also produces large, white basidiomata but clearly differs in slightly larger ovoid to broadly ellipsoid spores and considerably larger cheilo- $(50-150 \times 20-46 \ \mu\text{m})$ and pleurocystidia $(60-132 \times 19-44 \ \mu\text{m})$.

Volvariella bombycina (Schaeff.) Singer, V. bombycina var. microspora Dennis, V. hypopithys (Fr.) Shaffer, V. liliputiana (Henn.) G.C. Rath, and V. delicatula (Massee) Manjula are also similar to V. sathei. However, V. bombycina is lignicolous, and the spores are larger (7.3–9.4 µm in V. bombycina; 6–7.5 µm in V. bombycina var. microspora). Volvariella hypopithys has smaller basidiomes (pileus \leq 5 cm diam.) with squamulose pilei and hairy stipes, while Volvariella liliputiana and V. delicatula are much smaller (pileus 5–30 mm diam.) than V. sathei.

A BLASTn search comparing the AMH 9436 530bp nucleotide sequences (comprising ITS1 = 210bp, 5.8S = 151bp, ITS2 = 157bp) with the Genbank DNA ITS sequence database showed maximum (84%) similarity with *V. terrea* (JF415141.1 query coverage 88%) and 83% similarity with *V. dunensis* (JF415140.1 query coverage 88%). Twenty-two ITS sequences representing 13 *Volvariella* spp., 2 *Volvopluteus* sp., and 1 *Schizophyllum* sp. obtained from Genbank database were aligned using CLUSTAL W with manual adjustments.



PLATE 3. Neighbour-joining tree showing phylogenetic position of *Volvariella sathei*, *Volvariella*, and *Volvopluteus* species. Bootstrap values > 50% are indicated.

The 5.8S region was almost perfectly aligned, but numerous gaps (indels) and misalignments among *Volvariella* species were observed in ITS1 and ITS2 regions. All analysed *Volvariella* species were divided into two main clusters (I and II, PLATE 3), with *V. sathei*, *V. nivea*, and *V. taylorii* together in subclade Ia. Although *V. sathei* and *V. nivea*, both characterized by pure white basidiocarps, are placed together with a strong (96%) bootstrap support, the branch lengths show considerable sequence divergence between them. Our analysis places the two sequences of *Volvariella volvacea* (HM246500.1, TO AV143; HQ999973.1, ATCC MYA-4696) far apart (in completely two different clusters), suggesting ambiguity in deposited sequences.

As Li et al. (2009) has indicated for *V. nivea*, we noted a high degree of variability (including considerable misalignment and several indels) throughout the ITS 1 and ITS 2 regions among *Volvariella* species, but the 5.8S region remained conserved with relatively few changes.

Key to species of Volvopluteus (Vp.) and Volvariella (V.) in India: Two newly described species - Volvariella indica Pathak et al. (Manjula 1983) and V. minuta Kumar et al. (Kumar et al. 2009–10)— and a new record of V. congolensis N.C. Pathak (Kumar et al. 2009-10) from India are poorly known and have been excluded from the key. 2. Pileus 5–15 cm diam.; spores $11-18 \times 6.5-9.5 \mu m$ (ivory white to light brownish grey, often darker on the disc)Vp. gloiocephalus 2. Pileus 3–5 cm diam.; spores 11–15.5 µm long Vp. earlei 5. Spores frequently 10-10.5 µm longV. diplasia 5. Spores not > 9.5 µm long, volva white V. bombycina 6. Volva internally rugoso-tomentose; lamellae pale flesh pink; 6. Volva smooth; lamellae with brownish tints; spores $6.3-8 \times 4-5 \ \mu m \dots V$. thwaitesii 7. Basidiome growing on other mushrooms V. surrecta 9. Pileus and stipe distinctly hairy (pileus \leq 5 cm diam., white; spores 6–7 × 4.5–5 μm) V. *hypopithys* 10. Pileus 8–9 cm diam. white, fibrillose; stipe $80-105 \times 9-13$ mm V. sathei 12. Spores $4.5-7.5 \times 4-6.5 \mu m$, pileus pure white to pale cream (silky smooth to silky fibrillose, volva white, appressed) V. pusilla 13. Pileus different shades of brown14 14. Volva, fleshy, rigid, pale to dark brown to blackish brown above, 15. Basidiome large and fleshy; pileus robust \leq 12 cm diam., tawny brown to 15. Basidioma small to medium ≤ 6.5 cm diam.; stipe 2–6 mm diam.;

16. Spores frequently > 9 μ m long (6–10.4 × 4.5–7 μ m) V. t	volvacea
16. Spores not > 9 μm long	17
17. Pileus light seal brown at the disc, paling to grayish vinaceous at the margin	L
with fine brown radial striae; spores 4.5–8.6 \times 3–5.3 μm V.	cubensis
17. Pileus bright chestnut brown, spores $5-6.5 \times 3.5-4 \ \mu m \dots V$.	castanea
18. Pileus grayish; volva grayish	19
18. Pileus brownish; volva brown or white	20
19. Pileus evenly pale to dark grey, slightly darker at the disc;	
margin not striate	taylorii
19. Pileus evenly grayish with blackish disc; margin or the entire cap	
deeply grooved	grodisca
20. Pileus vinaceous brown at the disc, paler towards the margin; stipe very long	g,
$4-10 \text{ cm} \times 3-6 \text{ mm}$; volva fuliginous brown V. pseudov	volvacea
20. Pileus ash gray to grayish brown with blackish radial fibrils; stipe short,	
$4-4.5 \text{ cm} \times 2-4 \text{ mm}$, volva whiteV. apa	ılotricha

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