<table>
<thead>
<tr>
<th><strong>Title</strong></th>
<th>Phylogenetic and morphological assessment of five new species of Thozetella from an Australian rainforest</th>
</tr>
</thead>
<tbody>
<tr>
<td><strong>Author(s)</strong></td>
<td>Paulus, B; Gadek, P; Hyde, KD</td>
</tr>
<tr>
<td><strong>Citation</strong></td>
<td>Mycologia, 2004, v. 96 n. 5, p. 1074-1087</td>
</tr>
<tr>
<td><strong>Issued Date</strong></td>
<td>2004</td>
</tr>
<tr>
<td><strong>URL</strong></td>
<td><a href="http://hdl.handle.net/10722/53372">http://hdl.handle.net/10722/53372</a></td>
</tr>
<tr>
<td><strong>Rights</strong></td>
<td>This work is licensed under a Creative Commons Attribution-NonCommercial-NoDerivatives 4.0 International License.</td>
</tr>
</tbody>
</table>
Phylogenetic and morphological assessment of five new species of *Thozetella* from an Australian rainforest

Barbara Paulus\(^1\)
Paul Gadek
School of Tropical Biology, James Cook University, Cairns, QLD 4870, Australia

Kevin Hyde
Centre for Research in Fungal Diversity, Department of Ecology and Biodiversity, University of Hong Kong, Pokfulam Road, Hong Kong, SAR, People’s Republic of China

Abstract: During an investigation of saprobic microfungi in leaf litter from an Australian rainforest, five new species of *Thozetella*, namely *T. acerosa, T. boonjiensis, T. falcata, T. gigantea* and *T. queenslandica*, were identified and these are described and illustrated here. The morphology of specimens derived from cultures grown under different conditions and from natural substrata was compared. DNA sequence data of ITS regions within nuclear rDNA confirmed the morphological species concept and indicated that *Thozetella* species are anamorphs of the ascomycete genus *Chaetosphaeria*.

Key words: anamorphic fungi, *Chaetosphaeria*, Chaetosphaeriaceae, key, microfungi, morphology, taxonomy, tropical fungi

INTRODUCTION

*Thozetella* O. Kuntze (1891) is a nomen novum for *Thozetia* Berk. et F. Mueller, which was antedated by *Thozetia* F. Mueller ex Bentham (Asclepiadaceae). Pirozynski and Hodges (1973) revised the genus and accepted four species. Since then, a further five species were added by Nag Raj (1976), Sutton and Cole (1983), Castañeda Ruiz (1984), Castañeda Ruiz and Arnold (1985) and Mercado Sierra et al (1997). The genus *Thozetella* is characterized by phialidic conidiogenesis, aseptate, hyaline, curved conidia with an unbranched setula on each pole, microawns and sporodochial, synnematous or effuse conidiomata (Pirozynski and Hodges 1973). Microawns are distinctive, sterile elements, which are produced from conidiophores and found in the spore mass. Their function is unknown but they might aid spore dispersal. The generic definition outlined above make the inclusion of *T. ciliata* (Mercado-Sierra et al 1997) doubtful, because this species lacks microawns and produces setae. Species are primarily saprobes of decaying plant material, including terrestrial litter, bark and palm fronds (Pirozynski and Hodges 1973, Yanna et al 2002), submerged wood (Sivichai et al 2002), and *T. tocklaiensis* was observed as a nonpathogenic root endophyte (Waipara et al 1996).

While revising the genus, Pirozynski and Hodges (1973) questioned the species concept in *Thozetella* because knowledge of variation with growth conditions and age was limited. Interpretation of species concepts in the genus is complicated by the fact that some species are known only from cultures and others only from fruiting bodies grown on natural substrata (Sutton and Cole 1983). Consequently, Sutton and Cole (1983) showed that characters of *T. effusa* can vary greatly depending on the substratum on which the fungus has grown.

During an investigation of the diversity of saprobic microfungi on leaf litter in an Australian rainforest (Paulus et al 2003), we isolated a number of specimens belonging in *Thozetella*. They could be assigned to five morphological groups based on microawn length and/or morphology. To test whether they were indeed separate species rather than different stages of development of the same species or artifacts of cultural conditions, sequence analysis of internal transcribed spacer regions (ITS) of the nuclear ribosomal repeat (nrDNA) was undertaken. Sequence data also were used to clarify the placement of the genus among ascomycete lineages.

METHODS

Morphology of specimens from natural substrata and from culture.—Specimens of *Thozetella* were isolated after they sporulated on decaying leaves incubated in moist chambers (Booth 1971). In addition, spore suspensions were spread on 2% tap water agar containing 50 mg L\(^{-1}\) streptomycin and after 3–4 d small colonies were subcultured (Booth...
TABLE I. Collection and isolation details of Thozetella strains and herbarium specimens and the GenBank Accession number for DNA sequences of the ITS region

<table>
<thead>
<tr>
<th>Species name</th>
<th>Collection No.</th>
<th>BRIP No.</th>
<th>Substrata</th>
<th>Collection detailsb</th>
<th>Herbarium specimen or culturec</th>
<th>GenBank accession number</th>
</tr>
</thead>
<tbody>
<tr>
<td>T. acerosa</td>
<td>5790</td>
<td>29319</td>
<td>Cm</td>
<td>Site 1</td>
<td>C (PF)</td>
<td>QY330996</td>
</tr>
<tr>
<td>T. boonjiensis</td>
<td>2334</td>
<td>29318</td>
<td>Cm</td>
<td>Site 1</td>
<td>C (PF)</td>
<td>AY330994</td>
</tr>
<tr>
<td>T. boonjiensis</td>
<td>2383</td>
<td>29316</td>
<td>Oh</td>
<td>Site 1</td>
<td>C (PF)</td>
<td>AY330995</td>
</tr>
<tr>
<td>T. falcata</td>
<td>F658</td>
<td>29178</td>
<td>Fp</td>
<td>Site 1</td>
<td>H, C (ss)</td>
<td>AY330999</td>
</tr>
<tr>
<td>T. falcata</td>
<td>F634</td>
<td>29192</td>
<td>Cm</td>
<td>Site 2</td>
<td>H, C (ss)</td>
<td>AY310000</td>
</tr>
<tr>
<td>T. falcata</td>
<td>F711</td>
<td>29201</td>
<td>Cm</td>
<td>Site 2</td>
<td>H, C (ss)</td>
<td>AY310004</td>
</tr>
<tr>
<td>T. falcata</td>
<td>F715</td>
<td>29193</td>
<td>Cm</td>
<td>Site 2</td>
<td>H, C (ss)</td>
<td>AY310003</td>
</tr>
<tr>
<td>T. gigantea</td>
<td>F709</td>
<td>29200</td>
<td>Cm</td>
<td>Site 2</td>
<td>H, C (ss)</td>
<td>AY310001</td>
</tr>
<tr>
<td>T. gigantea</td>
<td>F712</td>
<td>29202</td>
<td>Cm</td>
<td>Site 2</td>
<td>H, C (ss)</td>
<td>AY310002</td>
</tr>
<tr>
<td>T. queenslandica</td>
<td>F415</td>
<td>29164</td>
<td>Cm</td>
<td>Site 1</td>
<td>H, C (ss)</td>
<td>AY330997</td>
</tr>
<tr>
<td>T. queenslandica</td>
<td>F612</td>
<td>29188</td>
<td>Cm</td>
<td>Site 1</td>
<td>H, C (ss)</td>
<td>AY330998</td>
</tr>
</tbody>
</table>

* Cm = decaying leaves of Cryptocarya mackinnoniana, Fp = decaying leaves of Ficus pleurocarpa, Oh = decaying leaves of Opisthiolepis heterophylla.

b Site 1 = Old Boonjie Road, Topaz, Queensland, Australia; Site 2 = Brooke’s Road, Milala Millaa, Queensland, Australia.

c H = herbarium specimen, C = culture, PF = particle filtration, ss = cultures isolated via spore suspension and small colony isolation.

1971). Cultures also were isolated by particle filtration (Kirby and Webster 1990, Bills and Polishook 1994, Paulus et al 2003). Specimens and cultures are listed in Table I. Subcultures were grown on malt-yeast agar (MYA; Gams et al 1998) for colony description and on potato-carrot agar (PCA; Johnston and Booth 1983), which included small pieces of triple sterlized banana leaf (Matsushima 1971) and wheat straw to encourage sporulation. A total of 25 conidia, 25 microawns and 10 conidiophores mounted in 90% lactic acid were measured per specimen. The length of microawns of various shapes always was measured along its longest axis and at the widest point. Differences in microawn and conidial length of specimens grown on CMA under different conditions using a Kruskal-Wallis Test (SPSS 2001).

DNA extraction, amplification and sequencing.—Eleven strains of Thozetella (Table I) were incubated in 1.5 mL microfuge tubes containing 2% potato-carrot extract, to which 1% glucose had been added, for 14 d at room temperature. DNA was isolated using a protocol modified from Smith and Stanosz (1995). The resulting pellet was resuspended in 50 µl double-sterilized distilled water (ddH2O). DNA target amplification by polymerase chain reaction (PCR) was performed for the 5.8S gene and the flanking ITS1 and ITS2 regions using the primers ITS1 and ITS4 (White et al 1990). All PCR reactions employed Taq DNA polymerase (Life Technologies) following the manufacturer’s product protocol. The resulting PCR product was purified with an Ultrascreen PCR Purification Kit (Mobio Laboratories Inc.) according to manufacturer’s instructions.

Sequencing reactions were carried out with 30–90 ng of purified template DNA using the same primers mentioned above as sequencing primers and the Big Dye Terminator Cycle Sequencing Ready Reaction Kit (Perkin Elmer Applied Biosystems) according to manufacturer’s instructions. Products were sequenced on an ABI-377 Prism Automated DNA Sequencer at the University of New South Wales, Australia. The nucleotide sequences were aligned and consensus sequences constructed using Sequence Navigator, version 1.0.1. (Applied Systems). All sequences reported in this article were deposited at GenBank (2003; http://www.ncbi.nlm.nih.gov/Genbank/) and their GenBank accession numbers are given in Table I. Alignments and trees are accessible through TreeBase (2003; http://www.treebase.org/treebase/index.html) under the reference number SN1712. A BLAST search GenBank (2003; http://www.ncbi.nlm.nih.gov/Genbank/) indicated high sequence similarity with members of the Chaetosphaeriaceae and an
unidentified leaf litter ascomycete (GenBank Accession number AF502902).

**DNA sequence analysis.**—Two datasets were generated. The first aligned ITS rDNA sequences from two representative *Thozetella* species, *T. gigantea* and *T. queenslandica*, with the *Chaetosphaeria* dataset of Reblova and Winka (2000) available from TreeBase (2003; http://www.treebase.org/treebase/index.html) under the reference number M1287. As a starting point, we accepted the alignment and outgroups proposed by Reblova and Winka (2000) but were more conservative in the analysis by treating gaps as missing data as opposed to fifth characters. The second dataset of more conservative in the analysis by treating gaps as missing groups proposed by Reblova and Winka (2000) was ambiguous in places and characters had to be excluded. Based on the results of the first study, we selected *Codinaeopsis gonytrichoides* (Shearer et J.L. Crane) Morgan-Jones and *Striatosphaeria codinaeaphora* Samuels et E. Müll. outgroups. Indels initially were excluded from analysis, but to test the sensitivity of analysis to variable alignment, indels were scored as presence/absence. We were unable to use DNA of previously described *Thozetella* species for this study, which would have provided further information.

For both datasets, parsimony analysis was performed in PAUP* 4.0b10 (Swofford 2002). Gaps were treated as missing data. The computing scheme used for parsimony analysis was: heuristic search option with stepwise addition random, 1000 replications, tree bisection-reconnection branch swapping (TBR), MulTrees option in effect. A strict consensus tree was constructed from all most parsimonious trees and support for clades was estimated with a bootstrapping procedure (Felsenstein 1985) using 1000 replications. Pairwise genetic distances were calculated based on “uncorrected p” (transitions plus transversions/total number of nucleotides) as performed in PAUP*.

**RESULTS**

**Taxonomy.**—Microawn, conidioma and conidium sizes and their morphology from fruiting bodies derived from natural substrata and from cultures for new and previously described *Thozetella* species were summarized (Table II).

*Thozetella* Kuntze, *Revis. gen. pl.* (Leipzig) 2:873 (1891)

Type species: *Thozetella nivea* (Berk.) Kuntze, *Revis. gen. pl.* (Leipzig) 2:873 (1891)

*Thozetella acerosa* Paulus, P.Gadek et K.D. Hyde, sp. nov. Fig. 1A

Ad fungos conidiales, hyphomycetes pertinens. Coloniae creameae et fuscae Conidiomata sporodochiae vel effusae, 120–250 × 300–400 μm. Conidiophorae macronematae, brunnae, ramosae, 1,5–3 μm latae. Cellulae conidiigenae pallidae brunneae, 15–20 × 1,5–2,5 μm. Microaristae angulatae, 0–2 septatae ad basi, apice acerosae et basi leviter obtusae, hyalinae, 60–80 × 5–4,5 μm in PCA. Conidia lunateae, unicellulariae, hyalinae, 14–20 × 2–3 μm in PCA, laeves et utrinque solo appendice, filiformi praedita, 6–7 μm longa.

Etymology. “acerosa” referring to the needle-like apices of the microawns.

Colonies cream and light brown, flat, woolly to subfally, margin incised and indistinct. Conidiomata sporodochial or effuse, superficial, sessile, forming a convex or flat hymenium, topped by a moist spore mass, 200–1500 μm diam. Conidiophores macronematous, brown, irregularly cylindrical, branched, arising from a basal plate, 2,5–4 μm wide. Conidigenous cells monophialidic, integrated, determinate, terminal, light brown, irregularly cylindrical, with no or minute collarette, periclinal wall thickened, 12–20 × 2–4 μm. Microawns produced from conidiophores, predominantly L-shaped, basal part thin-walled and hyaline, with 0–2 septa depending on growing conditions (Table III), apical part acerosae and slightly undulating, smooth, thick-walled with the upper portion becoming solid, refractive, 60–80 × 3,5–4,5 μm in PCA. Conidia lunate, aseptate, finely guttulate, hyaline and smooth, 14–20 × 2–3 μm in PCA, with one filiform setula at each end, 6–7 μm long.

**Specimens examined.** AUSTRALIA, QUEENSLAND: Atherton Tablelands, Topaz, Old Boonjie Road, on decaying leaves of Cryptocarya mackinnoniana, 8 Sep 2001, B. Paulus and C. Pearce, BP5970, HOLOTYPE: BRIP 29319.

Commentary. *Thozetella acerosa* resembles *T. nivea* in microawn shape. Our isolate differs in having shorter and narrower conidia and longer microawns than reported for *T. nivea*. In addition, our specimens have 0–2 septa at the basal part of the microawns at 25°C while microawns of *T. nivea* are aseptate. Two other species described below, namely *T. boonjiensis* and *T. gigantea*, also have L-shaped microawns and form sporodochia. Mature specimens of *T. gigantea* are distinguished easily because they have much longer microawns. Mean values of microawn and conidial length, as well as the presence or absence of septation in microawns, clearly differentiate between *T. acerosa* and *T. boonjiensis.*

*Thozetella boonjiensis* Paulus, P.Gadek et K.D. Hyde, sp. nov. Fig. 1B–E

TABLE II. Synopsis of *Thozetella* species based on literature. All measurements are given in μm

<table>
<thead>
<tr>
<th>Species</th>
<th>Microawn morphology</th>
<th>Microawn size (μm)a</th>
<th>Conidiomatal morphology and sizea</th>
<th>Conidia [setulae] (μm)a</th>
<th>Remarksa</th>
</tr>
</thead>
<tbody>
<tr>
<td><em>T. acerosa</em></td>
<td>Predominantly L-shaped, 0–2 septate in thin-walled basal part</td>
<td>60–80 × 3.5–4.5 on PCA</td>
<td>Sporodochia, 200–1500 μm, no conidiophore proliferations</td>
<td>14–20 × 2–3 [6–7] on PCA</td>
<td>Known from culture only</td>
</tr>
<tr>
<td><em>T. boonjiensis</em></td>
<td>Predominantly L-shaped, aseptate</td>
<td>48–75 × 3–5 on PCA</td>
<td>Sporodochia, 300–400 μm, no conidiophore proliferations</td>
<td>10–15 × 2–3 [5–8] on PCA</td>
<td>Known from culture only</td>
</tr>
<tr>
<td><em>T. canadensis</em> (Nag Raj 1976)</td>
<td>Irregularly sigmoid, lunate or sickle-shaped, with flat attachment scar and medium to submedian septum</td>
<td>32–37 × 2.5–3 on ns</td>
<td>Sporodochia, sessile and pulvinate, up to 250 μm in diameter and 120 μm high, no conidiophore proliferations</td>
<td>13–16 (–18) × 2–2.5 [5–7] on ns</td>
<td>Known only from natural substrata</td>
</tr>
<tr>
<td><em>T. ciliata</em> (Mercado-Sierra, Holubová-Jechová, Mena Portales 1997)</td>
<td>None reported</td>
<td>n/a</td>
<td>Synnemata, straight or curved, 300–500 μm high and 19.5–41.5 μm wide at the base, setae present</td>
<td>11.5–14.5 × 2.3–2.7 [4.5–7] on ns</td>
<td>Known only from natural substrata</td>
</tr>
<tr>
<td><em>T. cristata</em> (Pirozynski &amp; Hodges 1973)</td>
<td>Sickle-shaped, sigmoid, aseptate</td>
<td>40–60 × 2.5–3 on ns</td>
<td>Synnemata, with conidiophores proliferating synchronously to produce ridges; 250 × 40–60 μm</td>
<td>11–17 × 2–2.5 [6–9] on ns</td>
<td>Known only from natural substrata</td>
</tr>
<tr>
<td><em>T. cubensis</em> (Castañeda &amp; Arnold 1985)</td>
<td>Sigmoid, uncinate, aseptate</td>
<td>40–100(–110) × 2.5–4 on ns</td>
<td>Sporodochia, 450–500 × 250–300 μm, no conidiophore proliferations reported, conidiogenous cells have collar-ettes</td>
<td>11–17 × 2–2.5 [6–9] on ns</td>
<td>Known only from natural substrata</td>
</tr>
<tr>
<td><em>T. effusa</em> (Sutton &amp; Cole 1983)</td>
<td>Curved, hamate or loosely once or twice coiled, aseptate</td>
<td>20–30 × 3 on ns, 29–36 × 2.5–3 on 3% malt agar</td>
<td>Discrete, rarely confluent, 10–50 mm in diameter, flat, effuse</td>
<td>16–19 × 4–4.5 [6–11.5] on ns; 20–28 × 3.5 [6–13] on 3% malt agar</td>
<td>Known from natural substrata and culture</td>
</tr>
<tr>
<td><em>T. falcata</em></td>
<td>Predominantly sickle-shaped; L-shaped, straight, sigmoid microawns also observed in subcultures, aseptate</td>
<td>40–95 × 2.5–5 on ns; 50–110 × 2–4 on PCA</td>
<td>Funnel-shaped synnemata, 40–250 μm long, 75–110 μm wide at apex on ns; 300–1800 μm or longer on PCA, branched</td>
<td>13–16 × 1.5–3 [5–8] on ns; 15–16 × 2–3 [5–8] on PCA</td>
<td>Known from natural substrata and culture</td>
</tr>
<tr>
<td><em>T. gigantea</em></td>
<td>Predominantly L-shaped, apex acerosa and straight, aseptate</td>
<td>71–280 × 2.5–8 on ns; 70–210 × 3–6 on PCA</td>
<td>Sporodochia 100–600 × 90–500 μm</td>
<td>14–18 × 2.5–3 [6–10] on ns; 13–17 × 2–3 [6–10] on PCA</td>
<td>Known from natural substrata and culture</td>
</tr>
<tr>
<td><em>T. havanensis</em> (Castañeda 1984)</td>
<td>Sigmoid, allantoid, unciform, aseptate</td>
<td>22.4–35 × 1.5–3.2 on ns; 25.6–32 × 1.5–2 on CMA</td>
<td>Effuse and pulvinate synnemata</td>
<td>11–14 × 2.3 [5–9] on ns; 12–13.2 × 1.8–3 [4.2–6] on CMA</td>
<td>Known from natural substrata and culture</td>
</tr>
<tr>
<td>Species</td>
<td>Microawn morphology</td>
<td>Microawn size (µm)a</td>
<td>Conidiomatal morphology and sizea</td>
<td>Conidia [setulae] (µm)a</td>
<td>Remarksa</td>
</tr>
<tr>
<td>---------------------------</td>
<td>--------------------------------------------</td>
<td>---------------------</td>
<td>----------------------------------</td>
<td>-------------------------</td>
<td>-------------------</td>
</tr>
<tr>
<td><em>T. nivea</em> (Kuntze 1891)</td>
<td>Predominantly L-shaped, undulate, acerose apex, aseptate</td>
<td>50–70 × 3–4</td>
<td>Sporodochium, 200–250 µm in diameter</td>
<td>(17.5–21) (–24) × 3–3.8</td>
<td>Known only from natural substrata</td>
</tr>
<tr>
<td><em>T. queenslandica</em> Paulus, P. Gadek &amp; Hyde sp. nov.</td>
<td>Predominantly hamate, aseptate</td>
<td>24–33 × 2–3.5 on ns; 21–34 × 2–4 on PCA</td>
<td>Sporodochia, up to 200–250 × 50–125 µm with conidiophores proliferating synchronously in some specimens to produce ridges</td>
<td>10–12 × 1.5–2.5 [3–6] on PCA</td>
<td>Known from natural substrata and culture</td>
</tr>
<tr>
<td><em>T. radiata</em> (Pirozynski &amp; Hodges 1973)</td>
<td>Curved, sigmoid or L-shaped, aseptate</td>
<td>30–60 × 3–4.5 in culture</td>
<td>Funnel shaped synnemata (up to 750 µm long)</td>
<td>11–13 × 2.5–3 [3]b in culture</td>
<td>Known from culture only</td>
</tr>
<tr>
<td><em>T. tocklaiensis</em> (Pirozynski &amp; Hodges 1973)</td>
<td>L-shaped and other configurations, aseptate</td>
<td>18–38 (–44) × 1.5–3 on malt agar</td>
<td>On natural substratum sporodochia (1000 µm wide × 300 µm high), synnematal in culture (1–1.5 mm long)</td>
<td>9–13 (–18) × 1.5–3 [3–8 (–13)] on malt agar</td>
<td>Known from natural substrata and culture</td>
</tr>
</tbody>
</table>

*ns = natural substrata.

b According to Pirozynski and Hodges (1973), original description 13–17 × 2.5–3 µm.
PAULUS ET AL.: FIVE NEW SPECIES OF Thozetella


nae, unicellularae, 40–95 × 2.5–5 μm in foliis dejectis, 50–110 × 2–4 μm in PCA. Conidia lunatae, unicellulariae, hyalinae, 13–16 × 1.5–3 μm, laeves et utrinque solo appendice, filiformi praeedita, 5–8 μm longa.

Etymology. “falcata” referring to the sickle-shaped microawns.

Colonies hyaline with brown, concentric rings, flat, no aerial mycelium, synnemata formed in concentric rings on agar surface and on banana leaf. Conidiomata funnel-shaped synnemata, brown to dark brown, abundant on leaf, 40–250 μm long, 12–35 μm wide at base and 75–110 μm at apex on natural substrata, in PCA much more elongated (300–1800 μm long, 12–80 μm wide at base and 120–300 μm at apex), some synnemata forming branches. Conidiophores macronematous, brown, paling toward apex, irregularly cylindrical, densely compacted along synnematal axis, branched, 1.5–3 μm wide. Conidiogenous cells monophialidic, integrated, determinate, terminal, light brown, packed into tight palisades, irregularly cylindrical with bluntly rounded apex, lacking an apical collarette, periclinal wall thickened, 6–19 × 2–3 μm. Microawns predominantly sickle-shaped in specimens on natural substrata; L-shaped, sigmoid and
Table III. Percentage of microawns in *Thozetella acerosa* that formed septa under different growing conditions

<table>
<thead>
<tr>
<th>Media</th>
<th>Temperature</th>
<th>Light</th>
<th>No. of septa</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td></td>
<td></td>
<td>0</td>
</tr>
<tr>
<td>CMA</td>
<td>20°C</td>
<td>light/dark</td>
<td>95</td>
</tr>
<tr>
<td></td>
<td></td>
<td>dark</td>
<td>40</td>
</tr>
<tr>
<td></td>
<td>25°C</td>
<td>light/dark</td>
<td>35</td>
</tr>
<tr>
<td></td>
<td></td>
<td>dark</td>
<td>25</td>
</tr>
<tr>
<td>PCA</td>
<td>20°C</td>
<td>light/dark</td>
<td>100</td>
</tr>
<tr>
<td></td>
<td></td>
<td>dark</td>
<td>90</td>
</tr>
<tr>
<td></td>
<td>25°C</td>
<td>light/dark</td>
<td>45</td>
</tr>
<tr>
<td></td>
<td></td>
<td>dark</td>
<td>40</td>
</tr>
</tbody>
</table>

straight microawns also were observed in subcultures in addition to sickle-shaped microawns, 40–95 × 2.5–5 μm on natural substrata, 50–110 × 2–4 μm on PCA. Conidia lunate, aseptate, finely guttulate or eguttulate, hyaline, smooth, 13–16 × 1.5–3 μm on natural substrata, 13–16 × 2–3 μm on PCA, provided with a single, filiform setula at either pole, 5–8 μm long.

*Specimens examined.* AUSTRALIA, QUEENSLAND: Atherton Tablelands, Millaa Millaa, Brooke’s Road, on decaying leaves of *C. mackinnoniana*, 28 May 2002, B. Paulus and I.G. Steer, BP F66, BRIP 29192.

AUSTRALIA, QUEENSLAND: Atherton Tablelands, Millaa Millaa, Brooke’s Road, on decaying leaves of *C. mackinnoniana*, 21 Jun 2002, B. Paulus and I.G. Steer, BP F711, BRIP 29201.

Commentary. *Thozetella falcata*, *T. cristata*, *T. radiata* and *T. tocklaiensis* all produce synnemata. *Thozetella falcata* is also similar to *T. cristata* and *T. radiata* in microawn shape and in conidial dimensions. However, *T. falcata* differs from these two species in having considerably longer microawns on natural substrata and in culture. In addition, *T. falcata* differs from *T. radicata* in having branched and longer synnemata. Conidiomata of *T. falcata* are also different from those of *T. cristata*, in which conidiophores proliferate simultaneously to form visible ridges. In addition, *Thozetella cristata* forms consistently cylindrical synnemata while those in *T. falcata* are funnel-shaped. The length and morphology of synnemata in *T. tocklaiensis* are similar to those of *T. falcata*, however, these two fungal taxa differ in microawn morphology and size.

*Thozetella gigantea* Paulus, P.Gadek et K.D. Hyde, sp. nov.

Ad fungos conidiales, hyphomycetes pertinens. Conidiomatae sporodochiae, 100–600 × 90–500 μm. Conidiophorae macronematae, brunnae, 2.5–4 mm latae. Cellulae conidiogenae pallidae brunneae, 12–20 × 2–4 μm. Microar-

istae angulatae, apice acerosae et basi leviter obtusae, hyalinae, unicellularae, $71 \pm 280 \times 3.25 \pm 8 \mu m$ in foliis dejectis, $70 \pm 210 \times 2-3 \mu m$ in PCA. Conidiae lunatae, unicellularae, hyalinae, $14-18 \times 2.5-3 \mu m$ in substr. nat. et $13-17 \times 2-3 \mu m$ in PCA, laeves et utrinque solo appendice, filiformi praedita, 6-10 \mu m longa.

**Etymology.** “gigantea” referring to the size of microawns.

Colonies cream with brown radial lines developing, subfelty, fast growing. Conidiomata sporodochial, superficial, sessile, forming a convex or flat hymenium, topped by a moist, white spore mass with long, straight needle-like apices of microawns protruding from it, few sporodochia produced per leaf, 100-600 $\times 90-500 \mu m$. Conidiophores macronematous, brown, irregularly cylindrical, branched, arising from a basal plate, non-proliferating, 2.5-4 \mu m wide. Conidiogenous cells monophialidic, integrated, determinate, terminal, light brown, irregularly cylindrical, collarette missing, periclinal wall thickened, $12-20 \times 2-4 \mu m$. Microawns produced from conidiophores, predominantly L-shaped, basal part thin-walled, hyaline, continuous, long apical part acerose, smooth, straight, thick-walled with the upper portion becoming solid, narrowing down to less than 0.5 \mu m wide, refractive, $71-280 \times 2.5-8 \mu m$ on natural substrata, $70-210 \times 3-6 \mu m$ on PCA. Conidia lunatae, aseptate, finely guttulate, hyaline, smooth, slightly truncate on basal pole, $14-18 \times 2.5-3 \mu m$ on natural substrata and $13-17 \times 2-3 \mu m$ on PCA, with a single, 6-10 \mu m long, filiform setula at each end.


**Commentary.** *Thozetella gigantea* resembles *T. nivea*, *T. acerosa* and *T. boonjiensis* in the production of sporodochia and in microawn shape. It differs in having considerably longer microawns on natural substrata and in culture.

**Thozetella queenslandica** Paulus, P.Gadek et K.D. Hyde, sp. nov.

Ad fungos conidiales, hyphomycetes pertinens. Conidiomata sporodochiae superficialiae, 200-250 $\times 50-125 \mu m$. Conidiophorae macronematous, brunneae, ramosae. Cel-

lulae conidiogenae pallidae brunneae vel subhyalinae, 10–

$25 \times 1.5-2.5 \mu m$. Microaristae hamatae, sigmoidae, laeves vel verrucosae ad apicem, 24–33 $\times 2-3.5 \mu m$. Conidia lunatae, unicellularae, hyalinae, 10–12 $\times 1.5-2.5 \mu m$, laeves et utrinque solo appendice, filiformi praedita, 3-6 \mu m longa.

**Etymology.** “queenslandica” referring to the state of Queensland in Australia where this fungus was first isolated.

Colonies cream to gray, woolly, reverse with gray concentric rings. Conidiomata superficial, sessile sporodochia, 200-250 $\times 50-125 \mu m$, forming a flat or convex hymenium on a dark brown, stromatic base, which bears a globose, ovoid or otherwise shaped white mass of conidia and microawns; some sporodochia produce dark brown transverse ridges, each representing an area of synchronous proliferation of conidiophores. Conidiophores macronematous, brown, irregularly cylindrical, branched, compact at base, more or less free toward of upper part of sporodochium. Conidiogenous cells monophialidic, integrated, determinate, terminal, light brown to sub-

hyaline, irregularly cylindrical, without collarettes, periclinal wall thickened, 10–25 $\times 1.5-2.5 \mu m$. Microawns produced from conidiophores, predominantly hamate, rarely sigmoid, refractive, with a smooth or verrucose apex, 24–33 $\times 2-3.5 \mu m$ on natural substrata, 21–34 $\times 2-4 \mu m$ on PCA. Conidia lu-

nate, continuous, hyaline, eguttulate, smooth, 10–12 $\times 1.5-2.5 \mu m$ on natural substrate and 9–12 $\times 1.5-
Table IV. Sporulation of type cultures after 2 wk incubation on two media and under different conditions

<table>
<thead>
<tr>
<th></th>
<th>Light/dark</th>
<th>Dark</th>
<th>Light/dark</th>
<th>Dark</th>
</tr>
</thead>
<tbody>
<tr>
<td><strong>CMA</strong></td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>T. acerosa</td>
<td>sp &amp; eff +</td>
<td>sp ++</td>
<td>sp ++</td>
<td>sp ++</td>
</tr>
<tr>
<td>T. boonjiensis</td>
<td>—</td>
<td>sp (+)</td>
<td>sp &amp; eff +</td>
<td>—</td>
</tr>
<tr>
<td>T. falcata</td>
<td>sy ++</td>
<td>sy ++</td>
<td>sy ++</td>
<td>sy ++</td>
</tr>
<tr>
<td>T. gigantea</td>
<td>sp (+)</td>
<td>—</td>
<td>sp</td>
<td>sp</td>
</tr>
<tr>
<td>T. queenslandica</td>
<td>sp (+)</td>
<td>sp +</td>
<td>—</td>
<td>—</td>
</tr>
<tr>
<td><strong>PCA</strong></td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>T. acerosa</td>
<td>sp &amp; eff ++</td>
<td>sp &amp; eff ++</td>
<td>sp ++</td>
<td>—</td>
</tr>
<tr>
<td>T. boonjiensis</td>
<td>—</td>
<td>—</td>
<td>sp &amp; eff, sy +</td>
<td>—</td>
</tr>
<tr>
<td>T. falcata</td>
<td>sy ++</td>
<td>sy +</td>
<td>sy ++</td>
<td>—</td>
</tr>
<tr>
<td>T. gigantea</td>
<td>—</td>
<td>—</td>
<td>(sp (+))</td>
<td>—</td>
</tr>
<tr>
<td>T. queenslandica</td>
<td>sp +</td>
<td>sp +</td>
<td>—</td>
<td>—</td>
</tr>
</tbody>
</table>

sp = sporodochia.
sy = synnemata.
eff = effuse layer of conidiophores.
— none, (+) = few, + = some, ++ = many, +++ = abundant sporulating structures.

2.5 μm on PCA, provided with a single filiform setula at each end, 3–6 μm long.


Commentary. Thozetella queenslandica can be differentiated from other known species of Thozetella on the basis of its distinctive, short hamate microawns and the production of sporodochia, which show proliferation ridges. The microawns of T. queenslandica overlap in length with those of T. effusa and T. hawanensis, but these species differ from T. queenslandica in both microawn and conidiomatal morphology.

Growth and sporulation of cultures under different growing conditions.—Although some variation of microawn and conidial lengths were observed in specimens grown on PCA and on natural substrata, these were not found to be significant (df = 1; P = 0.524 and df = 1; P = 0.727, respectively). Similarly, no significant differences in microawn and conidial lengths were noted for cultures grown on different media (df = 1; P = 0.457 and df = 1; P = 0.583, respectively). Microawn length also did not differ significantly for cultures grown under different light and temperature treatments (df = 3; P = 0.968). In contrast, conidial length differed significantly under different cultures conditions (df = 3; P = 0.002). Examination of raw data (not shown) revealed that conidia of T. gigantea were markedly longer at 25 C compared to 20 C (18.3 and 15.2 μm respectively). The mean diameter of type strains incubated 3 wk on CMA at different temperature and light regimes is given in Fig. 3. All cultures grew well on CMA and PCA, and generally similar patterns were observed on both media. One group, which included T. queenslandica, T. falcata and T. boonjiensis, appeared to grow faster at 25 C. In contrast, T. gigantea and T. acerosa appeared to favor 20 C with the slower growth rate observed at 25 C with light/dark cycles.

A summary of sporulation patterns after 2 wk is provided in Table IV. Thozetella acerosa and T. falcata commenced sporulation in the first week and continued to sporulate profusely on both media and under all conditions. Scant sporulation from the second week onward was observed for T. gigantea and T. boonjiensis. Thozetella queenslandica sporulated well under a relatively wide range of conditions but did not form fruiting structures at 25 C in the dark on either medium (Table IV). With some exceptions, conidiomata of all species appeared to be consistent in their basic configuration over the 3 wk observation period. Exceptions included the development of effuse conidiophores in addition to sporodochia in T. acerosa and T. boonjiensis under some conditions. Thozetella boonjiensis also developed cream to light brown, submerged synnemata at 25 C on PCA when exposed to light/dark cycles. At 25 C synnemata of T. falcata were submerged in the agar.

**KEY TO THE DESCRIBED SPECIES OF THOZETELLA**

1. microawns present .......................... 2
2. microawns present ........................ 2
3. microawns predominantly L-shaped ....... 3
4. microawns not L-shaped or variously shaped 6
5. microawns 0–2 septate, 60–80 × 3.5–4.5 μm (mean 72 × 3.9 μm) ........................ 4
6. microawns aseptate .......................... 4
7. microawns very long, 70–280 μm × 2.5–8 μm
(mean 153.8 × 5 μm on natural substrata, 132.4 × 4.2 μm on PCA) ....... T. gigantea
4. microawns shorter than 75 μm ................. 5
5. conidia (17.5–21(–24) × 3–3.8 μm, microawns 50–70 × 3–4 μm, microawn apex undulating or geniculate .......... T. niwai
6. conidia 10–15 × 2–3 μm, microawns 48–75 × 3–5 μm (mean 60.1 × 3.6 μm), microawn apex straight or slightly undulating or T. boonjiensis
7. microawns predominantly sickle-shaped, unci
8. microawns predominantly straight, sigmoid or any other shape ................. 11
9. conidiomata predominantly synnematous ......... 8
10. conidiomata predominantly sporodochial ... 10
11. conidiomata effuse, never sporodochial or synnematous, microawns 20–30 × 3 μm, conidia 16–19 × 4–4.5 μm ............... T. effusa
12. conidiomata sporodochial or synnematous ...... 12
13. conidiomata variously shaped, bulbous base, acerose apex, straight, undulate, unicinate or bent 18–38
14. microawns with ± uniform width, sigmoid, allan
toid, unicinate, verruculose, 22.4–35 × 1.5–3.2 μm, conidia 11–14 × 2.3 μm, conidiomata synnematous or effuse ............... T. havanensis

ITS DNA sequence analyses.—The aligned dataset of Thozetella and Chaetosphaeria ITS sequences contained 529 positions. Of the 503 characters included in analysis, 284 were variable and 219 were constant. Among the variable sites, 209 were parsimony infor-
mative. The heuristic search identified two equally parsimonious trees, each 872 steps long, which differed only in their placement of Chaetosphaeria dilabens Réblová et W. Gams. The strict consensus tree showed 11 clades, which were supported by bootstrap values higher than 55% (Fig. 4). The two representative taxa of Thozetella grouped next to Striatosphaeria codinaeaphora and Codinaeopsis gonytrichoides with this clade having a bootstrap support of less than 55% (Fig. 4). A branch supporting the majority of Chaetosphaeria species has high bootstrap support (92%). The genetic distances of all Chaetosphaeria species excluding outgroups and excluding Thozetella were 0.0–25.5%; the genetic distances of Thozetella and Chaetosphaeria species were 11.0–31.8% (details not shown).

The aligned dataset of specimens of Thozetella plus two outgroups contained 494 positions excluding indel scores. Of the 467 characters included in the analyses, 133 were variable and 334 were constant. Among the variable sites, 66 were parsimony informative. Intraspecific genetic distances were low (0–0.7%), whereas interspecific genetic differences were 2.3–8.5%. The lowest interspecific genetic distance was observed between T. acerosa and T. boonjiensis (2.3 and 2.4%). The next closest species was T. gigantea, which differed from these two species between 4.1 and 4.4%. The greatest genetic distance was observed between T. gigantea and T. queenslandica (8.2 and 8.4%), while T. falcata differed from all other species between 4.9 and 6.4%. When indels were excluded from analysis, three most parsimonious trees were obtained, each 189 steps long and differing only in the arrangement of terminal branches within the clade representing Thozetella falcata. The strict consensus tree of three most parsimonious trees shows two clades and five subclades (Fig. 5). Thozetella boonjiensis and T. acerosa are sister clades and group next to T. gigantea. This tree also shows T. falcata and T. queenslandica to be sister clades. Bootstrap support is generally moderate to high (82–100%) but two branches have bootstrap values of only 56 and 59% (Fig. 5). One of these branches (viz. the branch supporting T. queenslandica and T. fal-
cata) collapsed to form a polytomy when the data were re-analyzed with indel scores included (not shown).

DISCUSSION

Relationship of Thozetella and Chaetosphaeria.—Analysis of ITS sequence data associated the anamorph genus Thozetella with the ascomycete genus Chaetosphaeria (Fig. 4). Réblová and Winka (2000) discussed the phylogeny of Chaetosphaeria and used

\[\text{\textsuperscript{4}}\text{The original descriptions of } T. \text{tocklaiensis shows microawns of various shapes, whereas Pirozynski and Hodges (1973) show the microawns of } T. \text{tocklaiensis to be L-shaped. We have accepted the original description for this key.}\]
FIG. 4. Phylogenetic relationships among species of *Chaetosphaeria* and its anamorphs based on ITS rDNA sequences. The phylogram, which is rooted by the outgroup method, depicts the strict consensus tree of five most parsimonious trees. Significant bootstrap values (>55%) are given above branches.

Two separate genomic regions in their analysis. Our ITS rDNA analysis was conservative and, hence, the resulting strict consensus tree was less resolved than that of Reblova and Winka (2000). Subgroups 1A and 1B inferred by Reblova and Winka (2000) clustered together, and the two *Thozetella* sequences inserted in this clade (Fig. 4). In addition to two teleomorph genera, *Chaetosphaeria pulviscula* and *Striatosphaeria codinaeaphora*, subgroup 1A includes the anamorphs *Dictyochaeta*, *Menispora* and *Codinaeopsis*, which are characterized by producing oblong, fusiform, often curved conidia with terminal setulae (Reblova and Winka 2000). The genus *Thozetella* shares these morphological characteristics and, therefore, morphological data are consistent with the results of ITS sequence analysis, despite the low bootstrap support values obtained for those clades. The teleomorphs of individual *Thozetella* species remain unknown because cultures did not produce ascomata and nucleotide divergence among available *Thozetella* and *Chaetosphaeria* sequences consistently was greater than 11%.

*Phylogenetic analysis of Thozetella species.*—This study applied both phylogenetic and morphological species recognition (Taylor et al. 2000) and both were congruent. For example, sequence analysis of ITS rDNA supported recognition of the morphological species as five distinct clades (Fig. 5). The strict consensus tree of the *Thozetella* sequences suggests that they are a strongly supported monophyletic group (100%, Fig. 5). Two main sister clades are resolved
FIG. 5. Phylogenetic relationships among species of Thozetella based on ITS rDNA sequences. The phylogram, which is rooted by the outgroup method, depicts the strict consensus tree of three most parsimonious trees. Significant bootstrap values (>55%) are given above the branches.

with the clade grouping *T. acerosa*, *T. boonjensis* and *T. gigantea* showing a moderately high bootstrap value (82%, Fig. 5). These species share the same basic microawn shape and predominant conidiomatal configuration but differ in microawn length (TABLE II). In addition to their morphological similarity, they also have the smallest interspecific genetic differences observed in our study.

Despite its relatively low bootstrap value, the clade supporting *T. acerosa* and *T. boonjensis* remained stable when indel scores were included (not shown). The other clade, which includes *T. queenslandica* and *T. falcata*, was weakly supported (bootstrap value 59%, Fig. 5) and the branch supporting this clade collapsed when indel scores were included in analysis. Because insertions and deletions are thought to be the primary source of ITS variation in fungi (Hibbett 1992), the close relationship between *T. queenslandica* and *T. falcata* cannot be inferred with as much confidence as that of the species in the first clade. This uncertainty is mirrored in the morphology of these species and the moderately high genetic distance (6.2 and 6.4%). Although *T. queenslandica* and *T. falcata* share a similar microawn shape (i.e., curved), their basic conidiomatal configurations differ (TABLE II). Including ITS sequences of other species and testing the concordance of more than one gene genealogy might help to resolve the phylogeny of *Thozetella* in the future (Taylor et al 2000).

**Species delimitation based on morphology.**—Re-examination of morphological characters following ITS sequence analysis suggested that basic microawn morphology, such as L-shaped versus curved, microawn length and basic conidiomatal configuration, such as sporodochia versus synnemata, are key characters in species delimitation. Conidium and conidiophore morphology and dimensions, which are usually important characters in differentiating anamorphic fungi species, can play only a secondary role in *Thozetella* due to their similarity among some species (TABLE II). With one exception, each new species in our collection differs from the most similar, previously described taxa in at least one key criterion and usually by one or more secondary characteristics (TABLE II, Key). *Thozetella boonjensis* does not differ from *T. nivea* by a key taxonomic character but by three other characters, namely conidial length, conidial width and finer microawn morphology (TABLE II, Key).
Comparison of fruiting structures from cultures and from leaves.—One of the criticisms raised by Sutton and Cole (1983) in delimiting Thozetella species was that new species had been described from few specimens and in a number of instances were known only either from culture or from natural substrata. Sutton and Cole (1983) reported that there was a considerable increase in conidial size and a moderate increase in microawn length, as well as greater variation of conidiomata morphology in cultures of *T. effusa*, as compared to those developing on natural substrata. In contrast, Agnihothrudu (1958) and Castañeda Ruiz (1984) observed minimal size variation in conidia and microawns of *T. tocklaiensis* and *T. havanensis* from natural substrata and from cultures.

In our study, cultures were available for all five species and for three of these specimens also could be studied from natural substrata (Table I). A comparison of microawn and conidial dimensions revealed some variation among specimens derived in vitro and from natural substrata, but these differences were not statistically significant. Similarly, microawn length appeared to decrease in cultures of all species with increasing temperature and with dark treatment (data not shown), but no statistical significance was detected. In contrast, conidial length differed significantly among cultures grown under different conditions. This difference could be attributed to a marked increase in conidial length in cultures of *T. gigantea* when grown at 25 C rather than at 20 C. Overall, microawn and conidial dimensions did not differ significantly on CMA and PCA. However, temperature preferences as observed by different growth rates (Fig. 3) and peaks in sporulation (Table IV) might provide additional taxonomic information.

The identification of many anamorphic genera (e.g., *Cylindrocladium, Cylindrocladiella* and *Fusarium*) has been based almost solely on cultural characteristics under standard conditions (e.g., Joffe 1986, Crous et al 1992, Crous and Wingfield 1994). The same approach has been suggested for *Thozetella* (Sutton and Cole 1983) and may be particularly useful if conidiomatal configuration is to be utilized as a taxonomic character. Specimens of the same *Thozetella* species have been observed to differ in the type of conidiomata formed even when derived from the same substratum (Castañeda Ruiz pers comm), but conidiomatal configuration remained basically stable in cultures of the five species investigated (Table IV). Although long-term stability was not evaluated, we recommend that *Thozetella* species be characterized routinely on CMA that includes a sterilized banana leaf, after incubation at 20 C for 2 wk. This recommendation is based on the slow development of conidiomata in some of the species and differs from other anamorphs where assessment usually is undertaken after 7 d (e.g., Fisher et al 1982, Samuels and Brayford 1990, Crous et al 1992).

ACKNOWLEDGMENTS

We gratefully acknowledge these institutions for financing this project: the Centre for Research of Fungal Diversity at the Department of Ecology and Biodiversity; The University of Hong Kong; the Cooperative Research Centre for Rainforest Ecology and Management; and the School of Tropical Biology, James Cook University. Kylie McKenzie (Trinity Anglican School, Cairns) contributed to the assessment of cultures under different growing conditions as part of a student research scheme. B.P. thanks N. Tucker for access to one of the collection sites, C. Pearce and I. Steer for their company on collection trips, H. Leung for providing support with photography, M. Harrington for advice on phylogenetic analysis and useful comments on the manuscript and L. Jones and M. Pye for assistance with technical aspects. We especially thank R. Castañeda Ruiz for sharing his taxonomic knowledge and experience. Two anonymous reviewers are thanked for their valuable comments.

LITERATURE CITED


Transactions of the Mycological Society of Japan 33: 533–556.


