

# Systematic revision of species of *Atractilina* and *Spiropes* hyperparasitic on Meliolales (Ascomycota) in the tropics

Miguel A. Bermúdez-Cova<sup>1,2</sup>, Tina A. Hofmann<sup>3</sup>, Nourou S. Yorou<sup>4</sup>, Meike Piepenbring<sup>1</sup>

<sup>1</sup> Mycology Research Group, Faculty of Biological Sciences, Goethe University Frankfurt Am Main, Frankfurt Am Main, Germany

<sup>2</sup> Departamento de Biología de Organismos, División de Ciencias Biológicas, Universidad Simón Bolívar, Caracas, Venezuela

<sup>3</sup> Centro de Investigaciones Micológicas (CIMI), Herbario UCH, Universidad Autónoma de Chiriquí, David, Panama

<sup>4</sup> Research Unit Tropical Mycology and Plants-Soil Fungi Interactions (MyTIPS), Faculty of Agronomy, University of Parakou, BP 123, Parakou, Benin

Corresponding author: Miguel A. Bermúdez-Cova ([miguelangelbermudez11@hotmail.com](mailto:miguelangelbermudez11@hotmail.com))

## Abstract

*Atractilina* Dearn. & Barthol. and *Spiropes* Cif. are genera of asexual fungi that comprise species mainly hyperparasitic on black mildews (Meliolales, Ascomycota). Although a common group of anamorphic fungi, they have been described up to now only by morphology and their systematic position is unknown. The present study provides a morphological treatise of all known species of *Atractilina* and *Spiropes* hyperparasitic on Meliolales, including insights into their systematic position, based on DNA sequences generated here for the first time. The study was conducted, based on 33 herbarium specimens and 23 specimens recently collected in Benin and Panama. The obtained DNA sequence data (28S rDNA and ITS rDNA) of *A. parasitica* and of two species of *Spiropes* show systematic placements in the Dothideomycetes and Leotiomyces, respectively. The sequence data of the two *Spiropes* spp. do not group together. Moreover, the anamorph-teleomorph connection between *Atractilina parasitica* and *Malacaria meliolicola*, a pseudothecioid fungus, is confirmed. Three species in the genus *Spiropes* are proposed as new to science, namely *S. angycalycis*, *S. carpobolbiae* and *S. croissantiformis*. Four species are reported for Benin for the first time, three species for Panama and one species for mainland America. *Atractilina* and *Spiropes* are currently two genera with highly heterogeneous species and they might have to be split in the future, once the taxonomic concepts are validated by morphology and molecular sequence data.

**Key words:** Anamorph-teleomorph connection, Benin, Dothideomycetes, Hyperparasitism, Leotiomyces, Panama



Academic editor: Gerhard Rambold

Received: 27 November 2023

Accepted: 8 March 2024

Published: 11 April 2024

Citation: Bermúdez-Cova MA, Hofmann TA, Yorou NS, Piepenbring M (2024) Systematic revision of species of *Atractilina* and *Spiropes* hyperparasitic on Meliolales (Ascomycota) in the tropics. MycoKeys 103: 167–213. <https://doi.org/10.3897/mycokeys.103.115799>

Copyright: © Miguel A. Bermúdez-Cova et al. This is an open access article distributed under terms of the Creative Commons Attribution License (Attribution 4.0 International – CC BY 4.0).

## Introduction

Meliolales (Sordariomycetes, Ascomycota) form a large order of biotrophic, obligate plant parasitic fungi in the Tropics and subtropics (Piepenbring et al. 2011; Hongsanan et al. 2015; Zeng et al. 2017). The order comprises two families, Armatellaceae and Meliolaceae, with *Armatella* Theiss. & Syd. and *Meliola* Fr. being the most species-rich genera of each family, respectively (Hosagoudar 2003; Jayawardena et al. 2020). They are commonly known as “black mildews”,

because they produce black colonies that are composed of dark, thick-walled, branched, superficial hyphae (Rodríguez Justavino et al. 2015).

Approximately 200 species of hyperparasitic fungi, i.e. fungi parasitic on other parasites, have been reported to grow on colonies of Meliolales (Bermúdez-Cova et al. 2022, 2023a). These hyperparasites mainly belong to the Dothideomycetes and the Sordariomycetes, although the systematic positions of a large number of these fungi still remain unknown (Bermúdez-Cova et al. 2022; Bermúdez-Cova et al. 2023a). Hyperparasitic fungi frequently overgrow entire colonies of black mildews, so the meliolalean host may be detected only by careful search with a light microscope (Stevens 1918; Ciferri 1955; Bermúdez-Cova et al. 2023a).

Amongst the hyperparasitic fungi, species of the anamorphic genera *Atractilina* Dearn. & Barthol. and *Spiropes* Cif. are common hyperparasites of black mildews in the tropics. In the past, they were regarded as conidial stages of Meliolales (Ciferri 1955; Bermúdez-Cova et al. 2023b) and nowadays as incertae sedis in the Ascomycota (Bermúdez-Cova et al. 2022). The genus *Atractilina* includes six species of mostly hyperparasitic hyphomycetes with true synnematata, denticulate conidiogenous loci and pale pluriseptate conidia (Deighton and Pirozynski 1972; Mel'nik and Braun 2013). On the other hand, the genus *Spiropes* comprises 34 species of dematiaceous, mostly hyperparasitic hyphomycetes with mononematous, fasciculate or synnematous conidiophores (Ellis 1968, 1971, 1976; Seifert and Hughes 2000; Bánki et al. 2023). Species of *Spiropes* are characterised by the presence of conidiogenous cells with conspicuous, flat and numerous scars, as well as pigmented conidia with 1–9 septa or pseudosepta (Ellis 1968).

*Arthrobotryum* Ces., *Cercospora* Fresen. ex Fuckel, *Helminthosporium* Link, *Pleurophragmium* Costantin and *Podosporium* Schwein. are only a few of the many genera to which species of *Atractilina* and *Spiropes* have been assigned in the past, although they were not congeneric with the type specimens of those genera (Ellis 1968; Deighton and Pirozynski 1972; Alcorn 1988). This resulted in taxonomic uncertainty with species being transferred from one genus to another. This problem was initially addressed by Ellis (1968) and Deighton and Pirozynski (1972), as they did an extensive morphological revision of taxa now assigned to *Atractilina* or *Spiropes*. For example, all the synnematous fungi, hyperparasitic on Meliolales formerly assigned to the genus *Arthrobotryum*, were transferred to the genus *Spiropes* by Ellis (1968), with the exception of *A. parasiticum* (Winter) Hansf., which was transferred to the genus *Atractilina* by Deighton and Pirozynski (1972).

There is currently one valid species of *Atractilina*, namely *A. parasitica* (G. Winter) Deighton & Piroz. and 19 species of the genus *Spiropes* known to be hyperparasitic on colonies of Meliolales (Ellis 1968; Deighton and Pirozynski 1972; Mel'nik and Braun 2013; Bermúdez-Cova et al. 2022). However, species delimitation within these two genera has up to now been done by morphology only, as species were described in the past before the molecular era and because of the challenges of isolating DNA from mixed infections (Bermúdez-Cova et al. 2022, 2023a, 2023b). As a result, the systematic position of both genera within the Ascomycota remained unknown. The present study revises the morphology of the species of *Atractilina* and *Spiropes* and provides the first insights into their systematic position according to molecular sequence data, with emphasis on the species hyperparasitic on Meliolales.

## Materials and methods

### Sample collection and morphological characterisation

Samples of leaves infected with black mildews were opportunistically collected in western Panama from January-March 2020 and in Benin in February as well as September-October 2022. For the present study, colonies of Meliolales hyperparasitised by *Atractilina parasitica* and species of *Spiropes* were considered. Infected leaves were dried in a plant press and deposited in the Herbarium at the Universidad Autónoma de Chiriquí (UCH, specimens from Panama) or in the Mycological Herbarium of the University of Parakou (UNIPAR) in Benin. Duplicates of large-sized samples were deposited in the Botanische Staatssammlung München (M). In some cases, fungal tissue was collected prior to drying of the specimens and preserved in CTAB buffer for subsequent DNA extraction.

Dried specimens were observed by stereomicroscopy and by light microscopy (LM). Measurements of at least 20 conidia and other structures have been made for each specimen at magnifications of 600× and 1000×. Measurements are presented as mean value ± standard deviation with extreme values in parentheses. Line drawings were made freehand on scaled paper. Scars on conidiophores are drawn in surface view although further cells of the conidiophore are drawn in optical sections. Images and drawings were edited with Photoshop (Adobe, San Jose, California). Specimens were also analysed morphologically by scanning electron microscopy (SEM). Materials used for SEM were prepared according to Hofmann et al. (2010).

### Host plant identification

Host plants were identified by morphological characteristics and, in some cases, by molecular sequence data. Morphological identifications were made by comparison with herbarium specimens, literature (e.g. Akoègninou et al. (2006); Condit et al. (2011)) and with the help of local botanists. Molecular sequence data for species identifications were obtained by polymerase chain reaction (PCR) for the amplification of the partial region of chloroplast *rbcL* with the primer pairs *rbcLa-F* (Levin et al. 2003) and *rbcLa-R* (Kress et al. 2009). DNA was extracted from approx. 0.05 g of leaf tissue dried with silica gel using the innuPREP Plant DNA Kit (Analytik Jena, Germany) and following the manufacturer's instructions. Protocols for PCR were carried out as described by Fazekas et al. (2012).

### DNA extraction, PCR amplification and sequencing of fungal DNA

DNA was isolated from the synnemata and hyphae of specimens using the E.Z.N.A Forensic DNA Extraction Kit, following the manufacturer's instructions. To extract total genomic DNA, a small amount of clean synnemata or single conidiophores were transferred into a sterile Eppendorf tube with approx. 200 µl of distilled water using sterilised tweezers and trying to avoid picking cells of any other organism associated with the leaves and the colonies of black mildews. For example, for the synnemata of *Atractilina parasitica* and *Spiropes melanoplaca*, only the upper parts were used for DNA extraction, in order to avoid the basal parts that are in direct contact with cells of other organisms.

The samples were frozen for 24 h at -20 °C, and later homogenised for 10–12 min. using a Retsch Mixer Mill MM301 with TL buffer and 2.5 mm Zirconia beads. Isolated DNA was re-suspended in elution buffer and stored at -20 °C.

Two partial nuclear gene regions (ribosomal loci) were amplified and sequenced: For the large subunit nuclear ribosomal DNA (nrLSU, 28S rDNA), the primers LR0R (Wagner and Ryvarden 2002) and LR5 (Vilgalys and Hester 1990) were used. For the internal transcribed spacer region of ribosomal DNA (ITS), the primers ITS5 and ITS4 (White et al. 1990) were used. The PCR mixtures consisted of 1 µl genomic DNA, 15× MgCl<sub>2</sub> reaction buffer (Bioline, Luckenwalde, Germany), 25 mM MgCl<sub>2</sub>, 25 µM of each dNTP, 10 µM of each primer and 5 U Taq DNA polymerase (VWR) in a total volume of 30 µl. Cycling parameters of the PCR were as follows: initial denaturation at 94 °C for 3 min, followed by 35 cycles of amplification [denaturation at 94 °C for 30 s, primer annealing at 52 °C for 30 s and primer extension at 72 °C for 45 s] and a final extension at 72 °C for 5 min, followed by storage at 8 °C. PCR-products were checked on 1.5% agarose electrophoresis gels containing HDGreenPlus DNA stain. Amplified PCR products were purified with the Cycle Pure Kit (VWR-Omega, USA). Sequencing was performed at SeqLab GmbH, Germany.

## Phylogenetic analyses

Consensus sequences of trace files were generated with Geneious 10.2.2 (<https://www.geneious.com>, Kearse et al. 2012) and searched against GenBank (<https://www.ncbi.nlm.nih.gov/>, Benson et al. 2014) with MegaBLAST. Ambiguous and miscalled bases were corrected, when possible, after examination of the corresponding chromatogram files. Sequences with a high similarity were aligned with MAFFT v. 7 using the L-INS-i algorithm (Nakamura et al. 2018). The alignments were manually checked by using MEGA v. 7 (Kumar et al. 2016). Gblocks v. 0.91b (Talavera and Castresana 2007) was used to remove poorly-aligned positions and divergent regions from the DNA alignment. Phylogenetic analyses of this study were conducted by applying Maximum Likelihood (ML) in RAxML-HPC2 v.8.2.12 (Stamatakis 2014) on XSEDE (Miller et al. 2010) and Bayesian phylogenetic inference with the programme MrBayes 3.2.6. (Ronquist et al. 2012) on XSEDE (Miller et al. 2010), available on the CIPRES Science Gateway web portal ([http://www.phylo.org/sub\\_sections/portal/](http://www.phylo.org/sub_sections/portal/)). The alignment and tree are included in Suppl. material 1.

We also used T-BAS 2.1 (Carbone et al. 2019) and the “Place Unknowns” tool to place newly-generated ITS sequences on to the Pezizomycotina tree version 2. Two FASTA files of the newly-generated ITS sequences of *Spiropes* were uploaded to the T-BAS interface. We selected the “de novo” option for the RAxML placement, with 500 bootstrap replicates.

## Results

### Taxonomy

Based on morphological evidence, the hyperparasitic fungi collected in Panama and Benin are assigned to the genera *Atractilina* or *Spiropes*. Amongst these, three species are proposed as new to science, all in the genus *Spiropes*.

Four species represent new reports for Benin and three for Panama. We also present a revision from herbarium material of 17 of the 19 known species of the genus *Spiropes* and one species of *Atractilina* hyperparasitic on Meliolales. All species synonyms, unless specified, are taken from Deighton and Pirozynski (1972) for *Atractilina parasitica* and from Ellis (1968) for species of *Spiropes*.

***Atractilina* Dearn. & Barthol., Mycologia 16: 175, 1924.**

***Atractilina parasitica* (G. Winter) Deighton & Piroz., Mycol. Pap. 128: 34, 1972  
Fig. 1**

- ≡ *Arthrosporium parasiticum* G. Winter, Hedwigia 25: 103, 1886.
- ≡ *Arthrobotryum parasiticum* (G. Winter) Hansf., Proc. Linn. Soc. Lond. 155: 64, 1943.
- = *Isariopsis penicillata* Ellis & Everh., Bull. Torrey bot. Club 22: 438, 1895.
- ≡ *Phaeoisariopsis penicillata* (Ellis & Everh.) S.C. Jong & E.F. Morris, Mycopath. Mycol. appl. 34: 271, 1968.
- = *Arthrobotryum tecomae* Henn., Hedwigia 43: 397, 1904.
- = *Arthrobotryum caudatum* Syd. & P. Sydow, Etudes sur la Flore du Bas et Moyen Congo 3(1): 22, 1909.
- = *Arthrobotryum dieffenbachiae* F. Stevens, Bot. Gaz. 65: 237, 1918.
- = *Atractilina callicarpae* Dearn. & Barthol., Mycologia 16: 175, 1924.
- = *Podosporium pallidum* Pat., Scient. Surv. P. Rico 8(1) Bot.: 103, 1926.
- = *Eriomycoopsis bosquieae* Hansf., Bothalia 4(2): 466, 1942.
- = *Arthrobotryum deightonii* Hansf., Mycol. Pap. 15: 218, 1946.
- = *Malacaria meliolicola* Syd., Annls. Mycol. 28(1/2): 69, 1930. New synonym proposed in this study.
- = *Paranectria flagellata* Hansf., Proc. Linn. Soc. London 153(1): 28, 1941. New synonym proposed in this study.
- ≡ *Malacaria flagellata* (Hansf.) Hansf., Mycol. Pap. 15: 128, 1946. New synonym proposed in this study.

**Description.** **Colonies** effuse, rust brown or pale brown, with hyphae that form large, erect, dark synnemata clearly visible under the stereomicroscope, but sometimes only loose unstalked tufts around the tips of the setae of the meliolalean host. **Hyphae** superficial, branched, septate, thin-walled, 1–2.5 µm wide, smooth. **Conidiophores** may form straw-coloured or pale olivaceous synnemata up to 1.5 mm long, 40 µm wide at the basal stalk-like part. Sometimes the synnemata grow around and up the setae of the meliolalean host. Individual conidiophores straight or sometimes flexuous, cylindrical, 2.5–5 µm thick towards the apex, pale olivaceous brown, with denticles. **Conidia** solitary, straight or slightly curved, fusiform, truncate at the base, tapering towards the apex and often terminating in a little bulbous swelling, 1 to mostly 3 septate, thin-walled, variable in size, (17–)30–37(–80) × (3.5–)7–8.5 µm, at first more or less colourless, at maturity becoming pale straw coloured, minutely rough-walled. As seen by SEM, the ornamentation of the surface of the conidia is distinctly reticulated, with thin networks and no ridges.

**Specimens examined.** On *Meliola* sp. on living leaves of *Opilia celtidifolia* (Opiliaceae), Benin, Campus University of Abomey-Calavi, botanical garden, 6°25'7"N,



**Figure 1.** *Atractilina parasitica* (MB127, MB136) **a** synnemata (gold spots) on colonies of *Meliola* sp. (black spots) on a leaf of *Opilia celtidifolia* **b** synnemata of (gold spots) on colonies of *Meliola clerodendricola* (black spots) on a leaf of *Clerodendrum capitatum* **c** synnemata **d** conidiophores drawn in optical section. The thickness of the wall is indicated only in the drawing in the middle **e** conidia shown in optical section **f–i** as seen by SEM **f** conidiophores with denticles **g** a denticle at the tip of a conidiophore **h** conidium **i** bulbous swelling at the tip of a conidium. Scale bars: 1.5 mm (**b**); 1 mm (**c**); 5  $\mu$ m (**d,e,i**); 8  $\mu$ m (**f**); 1  $\mu$ m (**g**); 6  $\mu$ m (**h**).

2°20'34"E, 24 m a.s.l., 9 February 2022, M. A. Bermúdez-Cova, A. Tabé, D. Dongnima, O.P. Agbani, M. Piepenbring, N.S. Yorou, MB127 (UNIPAR, M); on *Meliola clerodendricola* on living leaves of *Clerodendrum capitatum* (Lamiaceae), Benin, Abomey-Calavi, Zopah, 6°30'8"N, 2°20'24"E, 37 m a.s.l., 12 February 2022, M. A. Bermúdez-Cova, A. Tabé, D. Dongnima, O.P. Agbani, M. Piepenbring, N.S. Yorou, MB133; on *Meliola clerodendricola* on living leaves of *Clerodendrum capitatum*, Benin, Allada, Sékou, 6°38'56"N, 2°11'38"E, 48 m a.s.l., 12 February 2022, M. A. Bermúdez-Cova, A. Tabé, D. Dongnima, O.P. Agbani, M. Piepenbring, N.S. Yorou, MB136 (UNIPAR, M, GenBank accession number: OR804686); on *Meliola* sp. on living leaves of *Pterocarpus santalinoides* (Fabaceae), Benin, Lokoli, border of forest, 7°3'41"N, 2°15'26"E, 22 m a.s.l., 20 February 2022, M. A. Bermúdez-Cova, A. Tabé, D. Dongnima, L. Konetche, M. Piepenbring, R. Hounkarin, MB160 (M);

on *Meliola* sp. on living leaves of *Coffea arabica* (Rubiaceae), Benin, Attogon, Niaouli, CRA-Sud center, 6°44'24"N, 2°8'25"E, 122 m a.s.l., 28 February 2022, M. A. Bermúdez-Cova, A. Tabé, I. Agonglo, M. Piepenbring, N.S. Yorou, O.P. Agbani, MB178 (UNIPAR, M, GenBank accession numbers: OR804685 and OR804687); on *Meliola* sp. on living leaves of *Coffea arabica*, Benin, Atlantique, Attogon, Niaouli Forest, 6°44'23"N, 2°8'26"E, 119 m a.s.l., 19 September 2022, A. Krauß, A. Tabé, O. Koukol, N.S. Yorou, AK06H (UNIPAR, M, GenBank accession number: OR804684); on *Meliola* sp. on living leaves of *Clerodendrum capitatum*, Benin, Atlantique, Attogon, Pahou Forest, 6°22'56"N, 2°9'35"E, 13 m a.s.l., 6 October 2022, A. Krauß, A. Tabé, O. Koukol, N.S. Yorou, AK61.

**Additional specimens examined.** On *Meliola lasiotricha* on leaves of unknown plant host, Puerto Rico, 1926, M.B. Ellis (IMI 130722, type specimen of *Podosporium pallidum*); On *Meliola clerodendri* on leaves of *Clerodendrum cyrtophyllum*, Taiwan, 1938, W. Yamamoto (IMI 31921b, type specimen of *Atractilina parasitica*).

**Illustrations.** This species was illustrated by Deighton and Pirozynski (1972).

**Known hosts and distribution.** On colonies of *Amazonia* spp., *Asteridiella* spp., *Irenopsis* spp. and *Meliola* spp. on living leaves of various plants in Congo, Ghana, Guinea, India, Mauritius, Nigeria, Perú, Philippines, Puerto Rico, Sierra Leone, St. Thomé, Taiwan, Tanzania, Uganda, U.S.A. and Venezuela. Only one single collection on *Balladyna* sp. (Balladynaceae, Dothideomycetes) as a fungal host (Deighton and Pirozynski 1972). *Atractilina parasitica* is reported here for the first time for Benin.

**Notes.** Only two species of the genus *Atractilina* with hyperparasitic lifestyle are known, namely *A. asterinae* and *A. parasitica* (Deighton and Pirozynski 1972). *Atractilina asterinae* differs from *A. parasitica* by the presence of 3–10 septate, thick-walled conidia.

The specimens of *A. parasitica* collected on leaves of *Coffea arabica* (MB 178, AK4H, AK06H) were found growing together with pseudothecia of *Malacaria meliolicola* Syd. (Tubeufiales, Dothideomycetes). According to Hansford (1941, as *Paranectria flagellata*; 1946), *M. flagellata* is most probably the perfect state of *A. parasitica*. The specimens collected by Hansford were also growing on coffee leaves. The latter and the fact that the DNA sequences we obtained from *A. parasitica* (GenBank accession numbers: OR804684, OR804686, OR804685 and OR804687) and *M. meliolicola* (GenBank accession numbers: OR805247 and OR805248) clustered together in one single strongly-supported clade (Fig. 22), confirm the anamorph-teleomorph connection between both species. For an updated species description of *M. meliolicola*, see Bermúdez-Cova et al. (2023b).

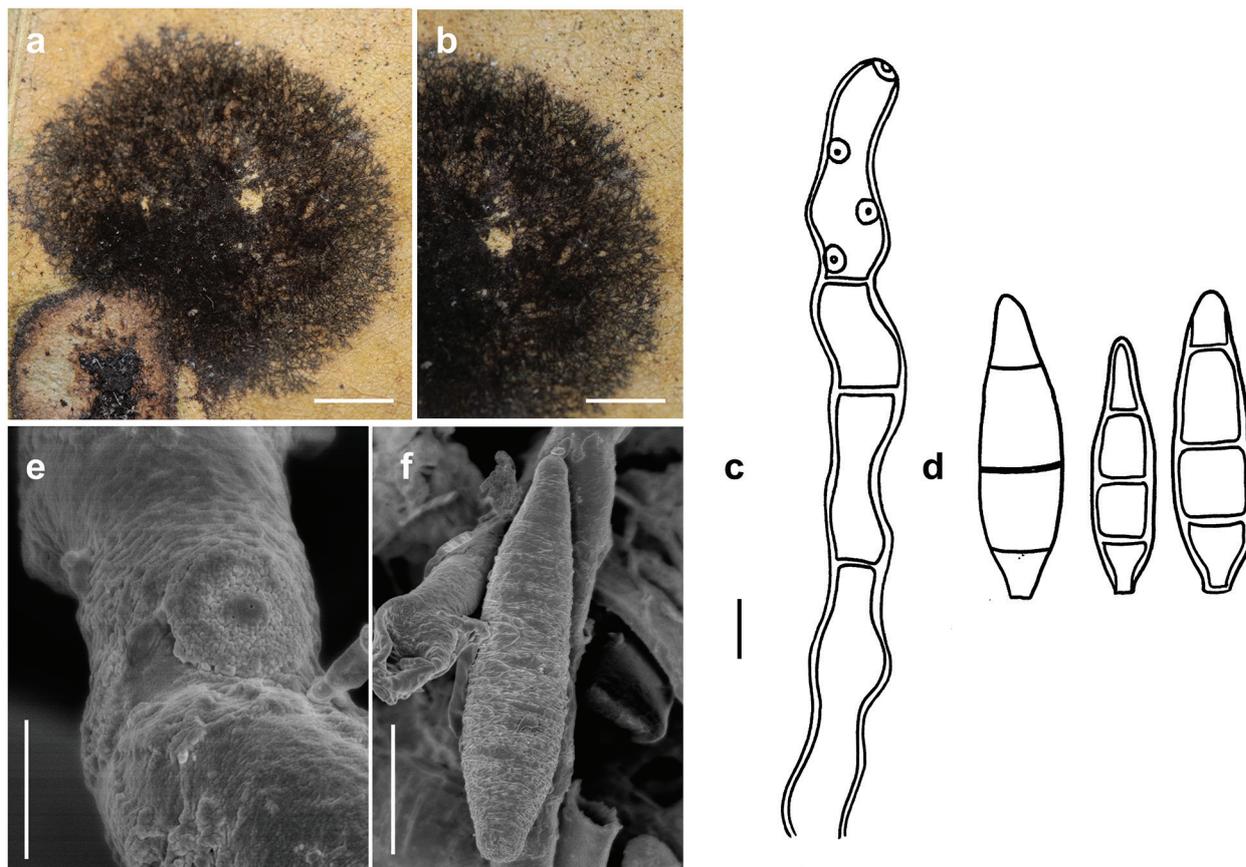
### ***Spiropes* Cif., *Sydowia* 9(1–6): 302, 1955**

#### ***Spiropes angylocalycis* Berm.-Cova & M. Piepenbr., sp. nov.**

MycoBank No: 850990

Fig. 2

**Holotype.** On *Meliola* sp. on living leaves of *Angylocalyx oligophyllus* (Fabaceae), Benin, Atlantique, Attogon, Niaouli Forest, 6°44'42"N, 2°7'50"E, 69 m a.s.l.,



**Figure 2.** *Spiropes armatellae* (MB 167) **a, b** conidiophores growing intermingled with hyphae of *Meliola* sp. on leaves of *Angylocalyx oligophyllus* **c** conidiophore with scars **d** conidia shown in optical section. The thickness of the wall is shown in the two drawings on the right-hand side **e, f** as seen by SEM **e** part of a conidiophore with scar **f** conidium. Scale bars: 0.3 mm (**a**); 0.2 mm (**b**); 5  $\mu$ m (**c, d**); 2  $\mu$ m (**e**); 7  $\mu$ m (**f**).

28 February 2022, M.A. Bermúdez, A. Tabé, D. Dongnima, I. Agonglo, O.P. Agbani, M. Piepenbring, N.S. Yorou, MB167 (M).

**Etymology.** Named after the genus of the host plant.

**Description.** **Colonies** effuse, dark brown to black, velvety to hairy. **Hyphae** superficial, branched, anastomosing, septate, 0.5–2  $\mu$ m wide, straw-coloured, smooth. **Conidiophores** arising singly, erect or ascending, straight to flexuous, mostly flexuous at the tips, septate, up to 350  $\mu$ m long, 4–6  $\mu$ m thick, pale olivaceous-brown to brown, with rough surface, with scattered scars mostly in upper parts of the conidiophores. **Conidia** solitary, straight or slightly curved, fusiform to obclavate, 3-septate, (15–)17–25(–30)  $\times$  5–6.5  $\mu$ m, 2–3  $\mu$ m wide at the base, brown, the cells at each end pale brown, septa darker in colour, verrucose. As seen by SEM, the ornamentation of the spores is distinctly reticulated, with thin to thick networks and no ridges.

**Known distribution.** On colonies of *Meliola* sp. on living leaves of *Angylocalyx oligophyllus* in Benin.

**Notes.** *Spiropes angylocalycis* is similar to *S. clavatus* by the presence of 3-septate mostly fusiform conidia, with a similar size range (Ellis 1968). However, the conidiophores of *S. clavatus* are synnematous, while they are mononematous in *S. angylocalycis*.

***Spiropes armatellae* M.B. Ellis, Mycol. Pap. 125: 15, 1971**

Fig. 3

**Type.** On *Armatella cinnamomicola* on leaves of *Cinnamomum* sp. (Lauraceae), Sri Lanka, Ceylon, 1971, M.B. Ellis (IMI134405b. The type specimen was not available for loan).

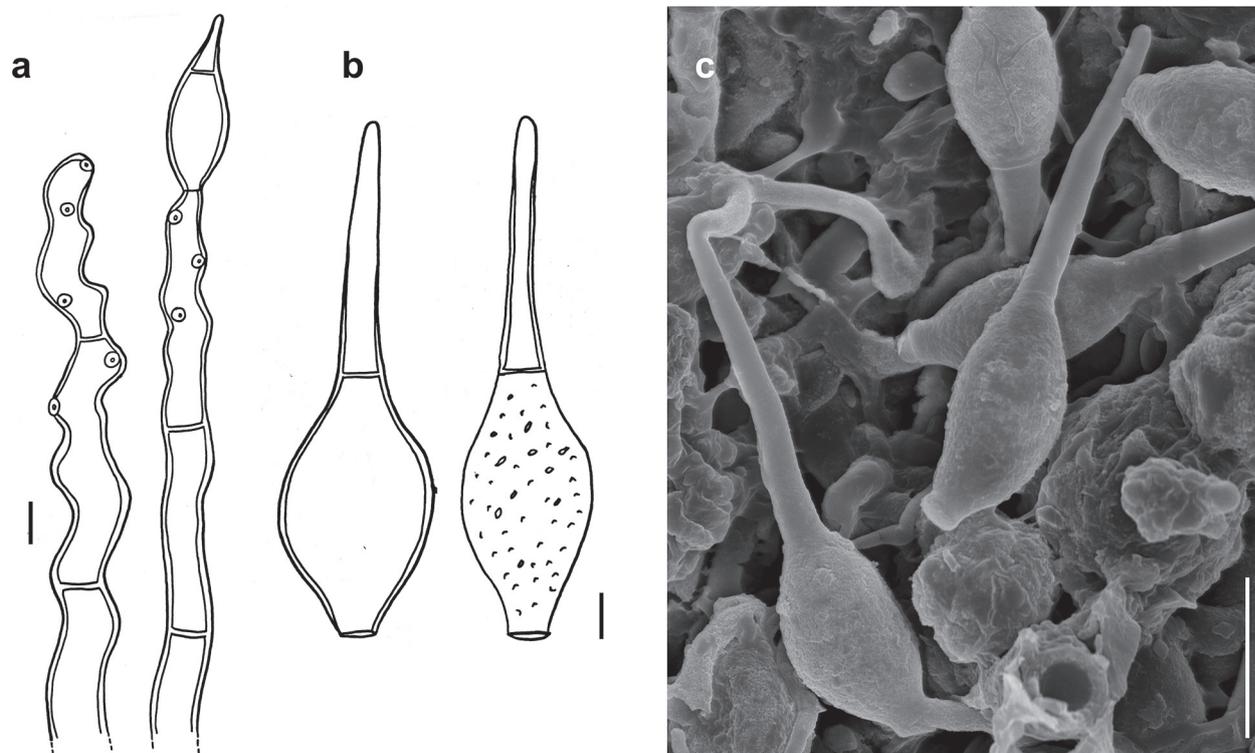
**Description.** **Colonies** effuse, dark brown to black, hairy. **Hyphae** superficial, branched, septate, 1–3 µm wide, straw-coloured, smooth. **Conidiophores** arising singly, erect or ascending, straight to flexuous, mostly flexuous at their tips, septate, up to 300 µm long, 5–8 µm thick, brown to dark brown, paler towards the apex, with rough surface, with scattered scars in upper parts of the conidiophores. **Conidia** solitary, straight or slightly curved, obclavate to obpyriform, mostly 1-septate, (20–)30–42(–50) × (6–)7–8(–10) µm, 2–3.5 µm wide at the base, brown, paler towards the ends, verrucose when seen by LM and SEM.

**Specimen examined.** On *Armatella litseae* on leaves of *Daphnidium pulcherrimum* (Lauraceae), India, west Bengal, 1967, M.K. Maity (IMI 136371); on *Armatella cinnamomicola* on leaves of *Cinnamomum* sp., Myanmar, Thaton, 1971, M.M. Thaug, (IMI 161265).

**Known hosts and distribution.** On colonies of *Armatella* spp. on various plants in India, Myanmar and Sri Lanka (Ellis 1971).

**Illustrations.** This species was illustrated by Ellis (1971).

**Notes.** Two known species of *Spiropes* are hyperparasitic on species of the genus *Armatella* (Meliolales, Armatellaceae), namely *S. armatellae* and *S. armatellicola* (Ellis 1971, Hosagoudar et al. 2002). According to Hosagoudar et



**Figure 3.** *Spiropes armatellae* (IMI 161265) **a** conidiophores with young conidium **b, c** conidia **b** shown in optical section. The thickness of the wall is indicated only in the drawing on the left-hand side **c** as seen by SEM. Scale bars: 5 µm (**a**); 2.5 µm (**b**); 10 µm (**c**).

al. (2002), both species are similar, but differ by the ornamentation of the conidia. The conidia of *S. armatelicola* are smooth, while those of *S. armatellae* are distinctly reticulated. However, it is sometimes difficult to observe the surface of the conidia by LM. Therefore, we recommend to analyse the ornamentation of the spores of *S. armatelicola* by SEM. The scars of *S. armatellae* could not be observed by SEM and it is necessary to collect fresh specimens of this fungus for further morphological analysis.

***Spiropes armatelicola* M.B. Ellis, Mycol. Pap. 125: 15, 1971**

**Type.** On *Armatella* sp. on leaves of *Actinodaphne* sp. (Lauraceae), Banasuran Hills, Wyanad, Kerala, India, 16 April 1999, C.K. Biju (HCIO 43621. The type specimen was not available for loan by HCIO).

**Species description.** This species was described by Hosagoudar et al. (2002).

**Known hosts and distribution.** On colonies of *Armatella* sp. on living leaves of *Actinodaphne* sp. in India (Hosagoudar et al. 2002).

**Illustrations.** This species was illustrated by Hosagoudar et al. (2002).

**Notes.** This species is only known from the type specimen.

***Spiropes capensis* (Thüm.) M.B. Ellis, Mycol. Pap. 114: 5, 1968**

Fig. 4

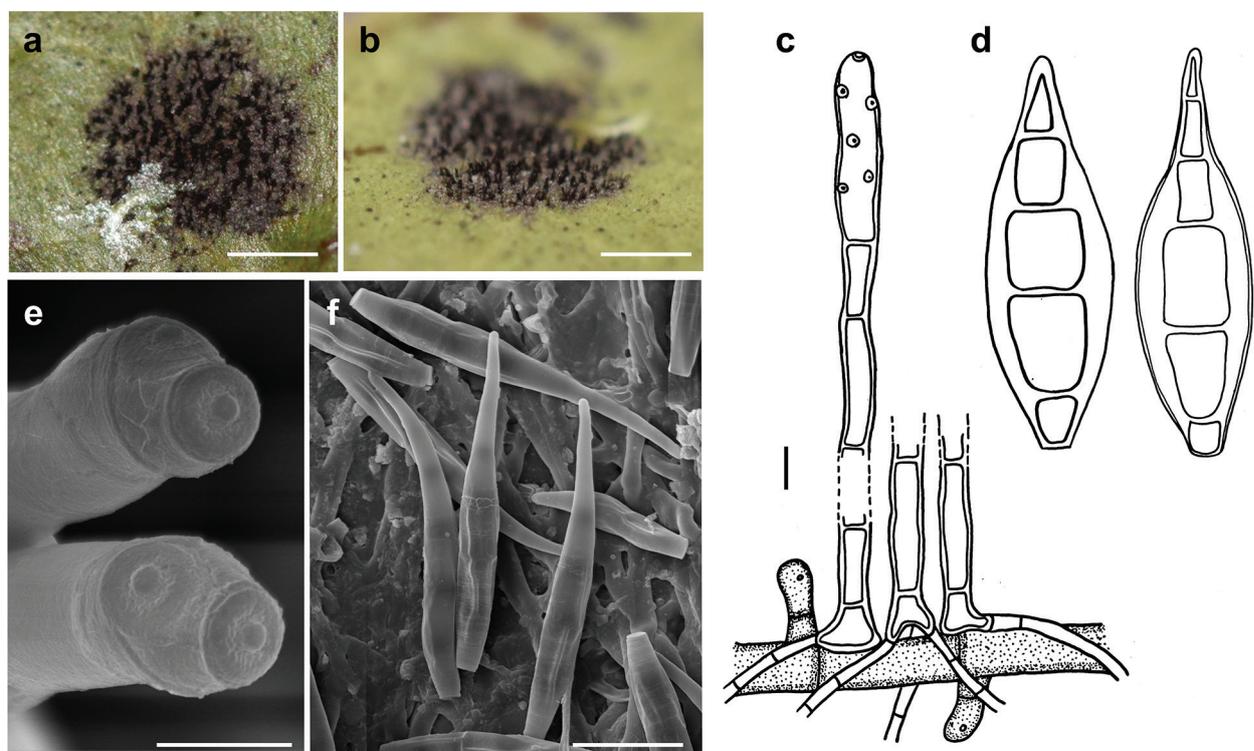
- ≡ *Cercospora capensis* (Thüm.) Sacc., Syll. fung. 4: 469, 1886.
- ≡ *Helminthosporium capense* (Thüm.) [as '*Helmisporium*'], Flora, Regensburg 59: 570, 1876.
- ≡ *Pleurophragmium capense* (Thüm.) S. Hughes, Can. J. Bot. 36: 796, 1958.
- = *Helminthosporium carpocrinum* Cif. [as '*Helmisporium*'], Annls. Mycol. 36(2/3): 236, 1938.
- = *Helminthosporium coffeae* Masee [as '*Helmisporium*'], Bull. Misc. Inf., Kew: 167, 1901.
- ≡ *Sporhelminthium coffeae* (Masee) Speg., Physis, Rev. Soc. Arg. Cienc. Nat. 4(no. 17): 292, 1918.
- = *Helminthosporium fici* H.S. Yates [as '*ficuum*'], Philipp. J. Sci. (Bot.) 13: 382, 1918.
- = *Helminthosporium ficinum* Sacc. [as '*Helmisporium*'], Atti Accad. Sci. Ven.-Trent.-Istr., Sér. 3, 10: 90, 1919.
- = *Helminthosporium fumagineum* Sacc. [as '*Helmisporium*'], Atti Accad. Sci. Ven.-Trent.-Istr., Sér. 3, 10: 90, 1919.
- = *Helminthosporium filicicola* Henn., Hedwigia 44: 71, 1905.
- = *Helminthosporium glabroides* F. Stevens [as '*Helmisporium*'], Bot. Gaz. 65(3): 240, 1918.
- = *Helminthosporium meliolooides* Sacc. [as '*Helmisporium*'], Atti Accad. Sci. Ven.-Trent.-Istr., Sér. 3, 10: 89, 1919.
- = *Helminthosporium orbiculare* Lév., Annls. Sci. Nat., Bot., Sér. 3, 5: 299, 1846.
- = *Helminthosporium philippinum* Sacc. [as '*Helmisporium*'], Atti Accad. Sci. Ven.-Trent.-Istr., Sér. 3, 10: 89, 1919.

- = *Helminthosporium subsimile* Sacc., Boll. Orto bot., Napoli 6: 23, 1921.
- = *Helminthosporium tapuræ* Allesch., Hedwigia 36(4): 245, 1897.
- = *Napicladium portoricense* Speg., Boln Acad. nac. Cienc. Córdoba 26(2–4): 363, 1921.
- ≡ *Helminthosporium portoricense* (Speg.) Cif., Sydowia 9(1–6): 298, 1955.
- = *Nascimentoa pseudoendogena* Cif. & Bat., Publicações Inst. Micol. Recife 44:4, 1956.

**Description.** **Colonies** effuse, dark brown to black, hairy (Ellis 1968). **Hyphae** superficial, branched, septate, 2–4 µm wide, pale olive to olivaceous-brown, smooth. **Conidiophores** arising singly or in groups, sometimes in large groups of 50–100 conidiophores, terminally or laterally from the hyphae, erect or ascending, straight or flexuous, septate, up to 600 µm long, 5–9 µm thick along most of their length, brown to dark brown, paler closer to the apex, with terminal and lateral scars. **Conidia** solitary, straight or curved, fusiform to obclavate, truncate at the base, 3–6 (usually 4 or 5) pseudosepta, (33–)50–60(–78) × (5.5–)6–11(–16) µm, 1–4 µm wide at the base, light brown to brown, smooth.

**Specimen examined.** On *Meliola* sp. on living leaves of *Coffea arabica*, Benin, Atlantique, Attogon, Niaouli Forest, 6°44'23"N, 2°8'26"E, 119 m a.s.l., 19 September 2022, A. Krauß, A. Tabé, O. Koukol, N.S. Yorou, AK06H.

**Additional specimens examined.** – On leaves of *Ficus ulmifolia* (Moraceae), Philippines, Los Baños, 1915, C.F. Baker, 451 (IMI 130940, type of *Helminthosporium fumagineum*); on *Meliola compositarum* on leaves of *Eupatorium portoricense*



**Figure 4.** *Spiropes capensis* (AK06H) **a, b** groups of conidiophores growing on hyphae of *Meliola* sp. **c** conidiophores growing on hyphae of *Meliola* sp. shown in optical section **d** conidia shown in optical section. The thickness of the outer wall layer is indicated only in the drawing on the right-hand side **e, f** as seen by SEM **e** conidiophores with scars **f** conidia. Scale bars: 1 mm (**a, b**); 8.5 µm (**c**); 5 µm (**d**); 5 µm (**e**); 20 µm (**f**).

(Asteraceae), Puerto Rico, Bega Vaja, 1921, no. 1753 (IMI 100331a, type of *Napicladium portoricense*).

**Known hosts and distribution.** On colonies of *Appendiculella* spp., *Asteridiella* spp., *Irenopsis* spp. and *Meliola* spp. on living leaves of various plants in Amboina, Bolivia, Brazil, Cameroon, Congo, Dominican Republic, Ghana, India, Jamaica, Malaya, Peru, Philippines, Puerto Rico, Sabah, Sierra Leone, South Africa, Tanzania, Trinidad, Uganda and Venezuela (Ellis 1968); on *Meliola* sp. on living leaves of *Coffea arabica* in Benin (this study). *Spiropes capensis* is reported here for the first time for Benin.

**Illustrations.** This species was illustrated by Ellis (1968).

**Notes.** According to the nomenclatural and taxonomic database Index Fungorum (<http://www.IndexFungorum.org>), the current name of the *Spiropes capensis* is *Pleurophragmium capense* (Thüm.) S. Hughes. The genus *Pleurophragmium* (*incertae sedis*, Ascomycota) was established by Costantin (1888) and it comprises species with brown to dark brown conidiophores and sympodially proliferating, denticulate conidiogenous cells producing holoblastic, simple, mostly 3-septate, brown to dark brown conidia (Abarca et al. 2007). According to Ellis (1968), the flat double scar is a good taxonomic character to distinguish species of *Spiropes* from *Pleurophragmium*, since, in the latter, the conidia are borne at the tips of tapered denticles. The morphological analysis of our samples and the type specimens (AK06H, IMI 100331a and IMI 130940) revealed the presence of flat double scars (Fig. 4e) and no denticles. We think that the examined species differs morphologically from species in the genus *Pleurophragmium* and, therefore, it should be retained in the genus *Spiropes*.

### ***Spiropes caribensis* Hol.-Jech., Česká Mykol. 38(2): 113, 1984**

Fig. 5

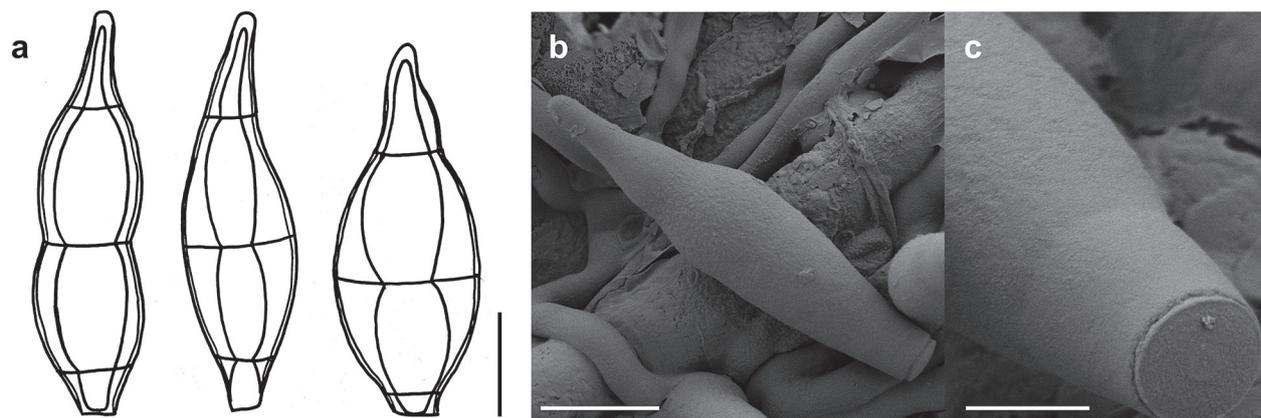
**Description.** **Colonies** effuse, dark brown to black, velvety to hairy. **Hyphae** superficial, branched, septate, 1.5–3.5 µm wide, pale olivaceous-brown, smooth. **Conidiophores** arising singly, erect or ascending, straight or flexuous, septate, up to 240 µm long, 4–8 µm thick, pale brown to brown, smooth, with few scars. **Conidia** solitary, straight or slightly curved, obclavate, central cells barrel-shaped, 3-septate, (30–)36–48(–41.5) × (7.5–)9.5–11.5 µm, 4.5–6 µm wide at the truncate base, the central cells pale brown, the cells at the ends paler and almost hyaline, smooth.

**Specimen examined.** On *Meliola* sp. on leaves of an unknown palm-tree, Cuba, Isla de La Juventud (= Isla de Pinos), Los Indios, south-west of La Cañada, 1981, V. Holubová-Jechová (PRM 831531, holotype).

**Known hosts and distribution.** On *Meliola* sp. on living leaves of an unidentified palm tree in Cuba (Holubová-Jechová and Sierra 1984).

**Illustrations.** This species was illustrated by Holubová-Jechová and Sierra (1984).

**Notes.** *Spiropes caribensis* is similar to *S. helleri*, but differs from the latter by paler conidia, with wider truncate base (*S. helleri* has conidia with a truncate base 3–4 µm wide) and shorter conidiophores (up to 600 µm long in *S. helleri*; Holubová-Jechová and Sierra (1984)). As seen by SEM, conidia of *S. caribensis* are smooth (Fig. 5b), while conidia of *S. helleri* are distinctly



**Figure 5.** *Spiropes caribensis* (PRM 8311531) **a** conidia shown in optical section **b, c** as seen by SEM **b** conidium **c** basis of a conidium with a flat scar. Scale bars: 10 µm (**a**); 9 µm (**b**); 4 µm (**c**).

reticulated (Fig. 13e). The scars could not be observed by SEM and it is, therefore, necessary to collect fresh specimens of this fungus for further morphological analyses. *S. caribensis* is only known from the type specimen.

***Spiropes carpolobiae* Berm.-Cova & M. Piepenbr., sp. nov.**

Mycobank No: 850987

Fig. 6

**Holotype.** On *Meliola* cf. *carpolobiae* on living leaves of *Carpolobia lutea* (Polygalaceae), Benin, Atlantique, Atogon, Niaouli Forest, 6°44'41"N, 2°7'52"E, 68 m a.s.l., 28 February 2022, M.A. Bermúdez, A. Tabé, D. Dongnima, I. Agonglo, O.P. Agbani, M. Piepenbring, N.S. Yorou, MB166 (M).

**Etymology.** Named after the genus of the host plant.

**Description.** **Colonies** effuse, dark brown to black, velvety to hairy. **Hyphae** superficial, branched, anastomosing, septate, 1–2 µm wide, straw-coloured, smooth. **Conidiophores** arising singly, erect or ascending, straight to flexuous, septate, up to 250 µm long, 2–5 µm thick, sometimes thicker at the apex, brown, not smooth, with scattered scars mostly in the upper parts of the conidiophores. **Conidia** solitary, straight or slightly curved, ovate to slightly fusiform, 3–septate, (12.5–)13–16(–19) × 5–7 µm, 2–2.5 µm wide at the base, brown, the cells at each end pale brown, septa darker, surface verrucose. As seen by SEM, the ornamentation of the conidia is distinctly reticulated, with thin to thick networks that can form ridges.

**Known distribution.** On colonies of *Meliola* cf. *carpolobiae* on living leaves of *Carpolobia lutea* in Benin.

**Notes.** *S. carpolobiae* is the only known species of *Spiropes* with ovate to slightly fusiform conidia.

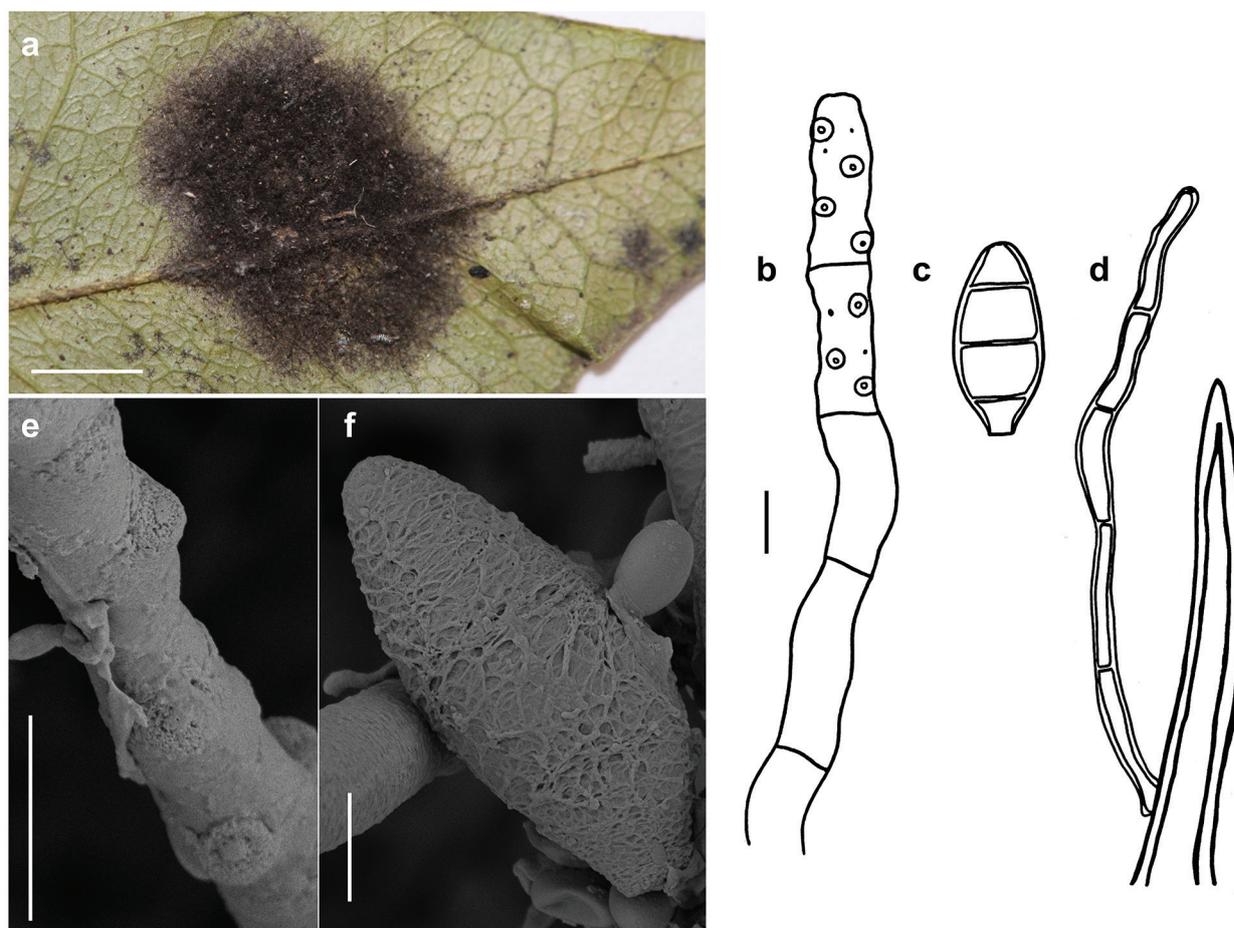
***Spiropes clavatus* (Ellis & Martin) M.B. Ellis, Mycol. Pap. 114: 25, 1968**

Fig. 7

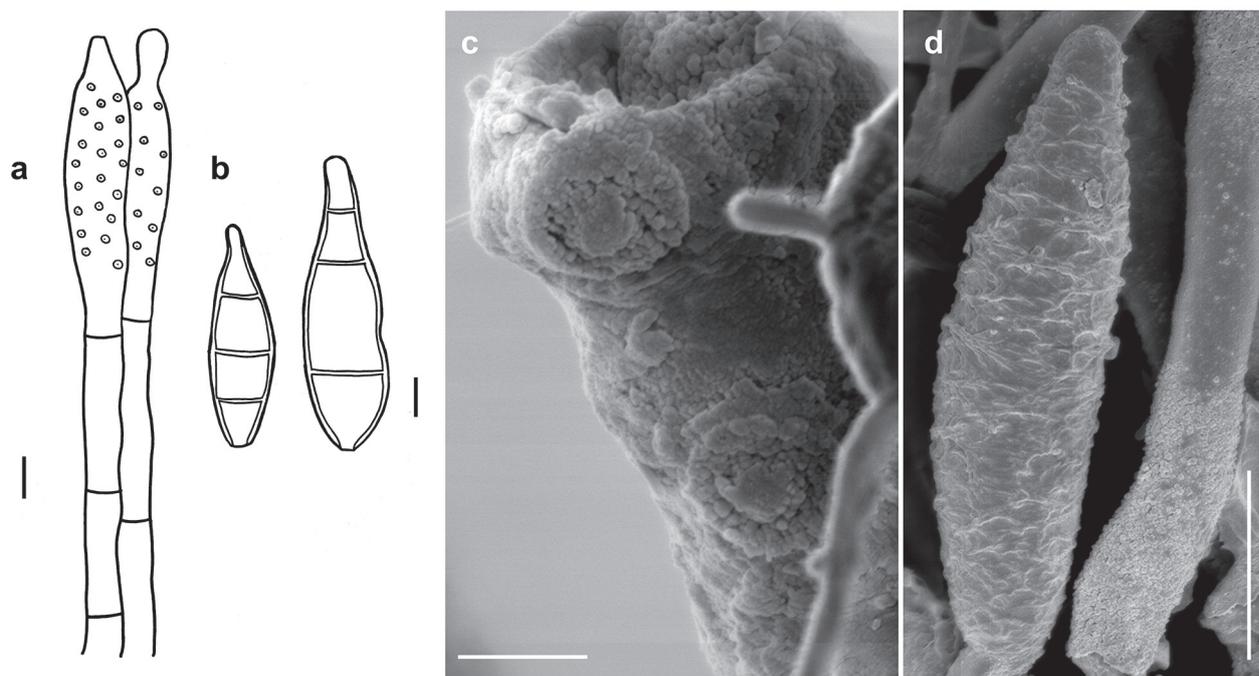
≡ *Isariopsis clavata* Ellis & Martin, Am. Nat. 18: 188, 1884.

- ≡ *Arthrotryum clavatum* (Ellis & Martin) Höhn, Sber. Akad. Wiss. Wien, Math.-naturw. Kl., Abt. 1, 125: 120, 1916.
- ≡ *Bitunicostilbe clavata* (Ellis & Martin) M. Morelet, Bull. Soc. Sci. nat. Arch. Toulon et du Var 7: 195, 1971.
- = *Podosporium chlorophaeum* Speg., An. Mus. nac. Hist. nat. B. Aires 20: 450, 1910.
- = *Arthrotryum noz-moscatae* Bat. & J. Silva, Anais IV Congr. Soc. bot. Brasil: 144, 1953.

**Description.** **Colonies** effuse, brown to dark brown or black. **Hyphae** superficial, branched, anastomosing, septate, 1–3 µm wide, pale olivaceous-brown. **Conidiophores** tightly packed to form dark brown to blackish synnemata up to 700 µm long, 20–40 µm thick, often splaying out to a width of up to 110 µm at the apex. Individual hyphae straight or flexuous, cylindrical, 1–3 µm thick near the base, 4–7 µm thick near the apex, dark brown, paler towards the apex, verrucose, with numerous conidial scars. **Conidia** solitary, fusiform to obclavate, mostly 3–, rarely 1–, 2– or 4–septate, (13–)18–25(–33) × (4–)5–7(–8) µm, tapering to about 1–1.5 µm at the apex and at the base, pale brown to brown,



**Figure 6.** *Spiropes carpolobiae* (MB 166) **a** conidiophores growing intermingled with hyphae of *Meliola* sp. on a leaf of *Carpolobia lutea* **b** conidiophore with scars **c** Conidia shown in optical section. The thickness of the wall is shown in the left-hand drawing **d, e** as seen by SEM **d** conidiophore with scar **e** conidium. Scale bars: 0.3 mm (**a**); 5 µm (**b, c**); 5 µm (**d**); 3 µm (**e**).



**Figure 7.** *Spiropes clavatus* (IMI 102772) **a** conidiophores with scars **b** conidia shown in optical section **c, d** as seen by SEM **c** conidiophore with scars **d** conidium. Scale bars: 5 µm (**a**); 2.5 µm (**b**); 1 µm (**c**); 5 µm (**d**).

the cells at each end paler, wrinkled. As seen by SEM, the ornamentation of the spores is distinctly reticulated, with thin to thick networks and no ridges.

**Specimens examined.** On *Meliola panici* on leaves of *Panicum glutinosa*, Puerto Rico, El Alto de la Bandera, 1913, F.L. Stevens & W.E. Hess, n°4368 (IMI 130764); on *Meliola* sp. on leaves of *Raphia monbuttorum*, Uganda, 1915, R. Dümmer, (IMI 102772); on *Meliola thouinia* on leaves of an unknown plant, Brasil, São Paulo, 1940, A.R. Campos (IMI 130975, type of *Arthrobotryum noz-moscatae*).

**Illustrations.** This species was illustrated by Ellis (1968).

**Known hosts and distribution.** On colonies of Meliolales on living leaves of various plants in Argentina, Brazil, Ghana, Malaysia, Puerto Rico, Sierra Leone, Trinidad and Uganda (Ellis 1968).

**Notes.** In the nomenclatural and taxonomic database Index Fungorum (<http://www.IndexFungorum.org>), the current name of the *Spiropes clavatus* is *Bitunicostilbe clavata* (Ellis & Martin) M. Morelet. The genus *Bitunicostilbe* (*incertae sedis*, Ascomycota) was proposed by Morelet (1971) to accommodate two species, namely *B. clavata* and *B. linderiae*, that were previously cited in other genera. Although the publication by Morelet was not available for this study, the morphological analysis of the herbarium specimens (IMI 130764, 130975) revealed that the features of these specimens are consistent with the description of *Spiropes clavatus* by Ellis (1968). The species has typical characteristics of the genus *Spiropes*, such as flat double scars (Fig. 7c) and, therefore, it should be classified in this genus. De Beer et al. (2013) analysed the type and additional specimens of *B. linderiae* (as *Graphium linderiae*) and concluded that this species should be also classified in the genus *Spiropes*.

***Spiropes croissantiformis* Berm.-Cova & M. Piepenbr., sp. nov.**

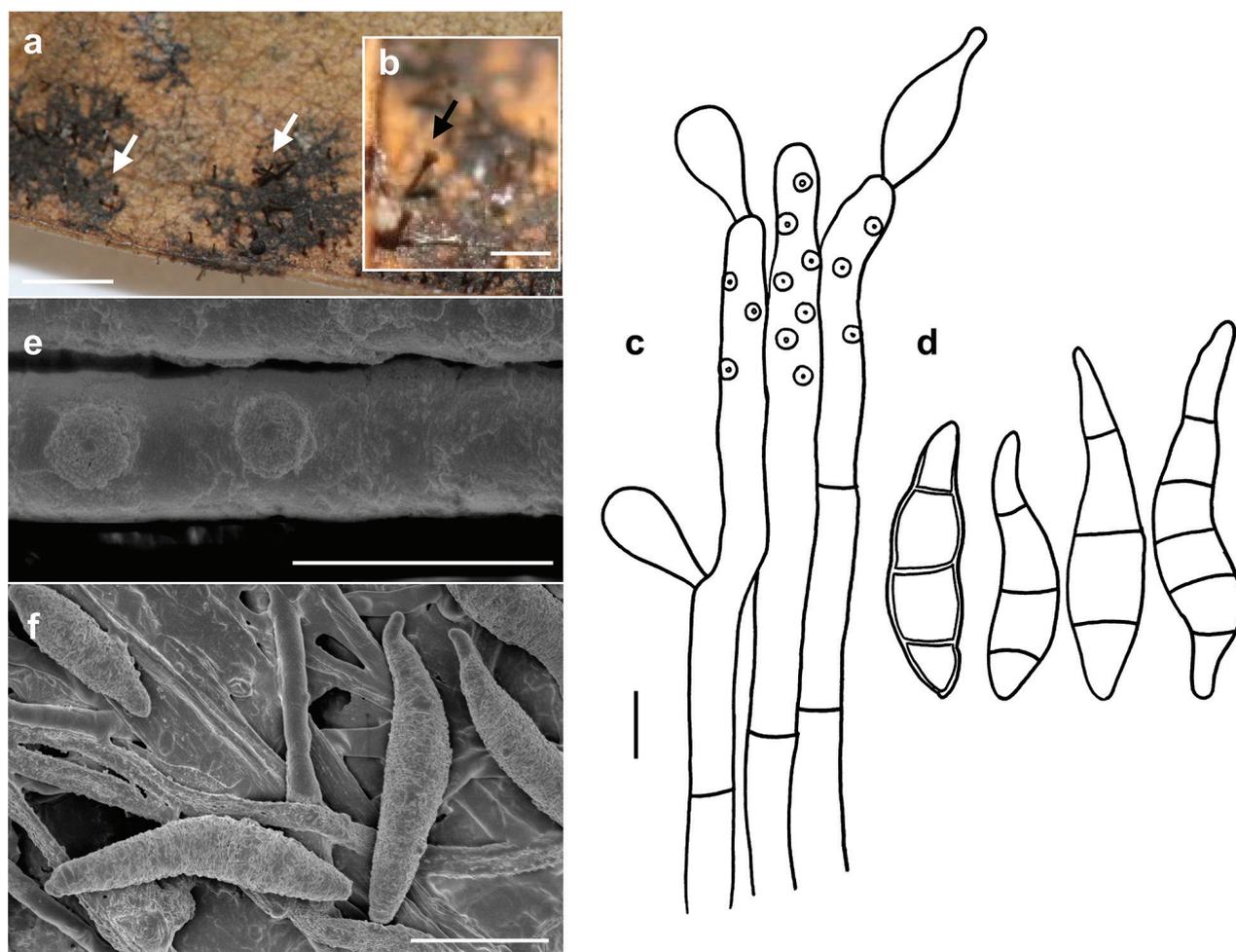
MycoBank No: 850984

Fig. 8

**Holotype.** On *Meliola* cf. *xylopii* on living leaves of *Xylopia frutescens*, Panama, Chiriquí Province, Cochea, Cochea River Trail, 8°32'37"N, 82°23'03"W, 181 m a.s.l., 26 February 2020, M.A. Bermúdez, A. Sanjur, A. Villarreal, MB110 (UCH).

**Etymology.** Named after the shape of the conidia.

**Description.** **Colonies** effuse, dark brown to black, with tightly packed hyphae that form erect, dark synnemata clearly visible under the stereomicroscope. **Hyphae** superficial, branched, septate, 1–2 µm wide, straw-coloured, smooth. **Conidiophores** tightly packed to form dark brown to blackish synnemata up to 400 µm high, spreading out at the apex, up to 80 µm diam. Individual hyphae mostly straight, cylindrical, 3–5 µm thick, with numerous small scars, brown, paler towards the apex, rough. **Conidia** straight or curved, mostly crescent-shaped, sometimes fusiform, mostly 3(–5)–septate, (14–)20–24(–33) × (3.5–)5–6.5 µm, with two golden brown middle cells and paler cells at each. As



**Figure 8.** *Spiropes croissantiformis* (MB 110) **a** synnemata (indicated by white arrows) growing on colonies of *Meliola* cf. *xylopii* **b** synnema (indicated by a black arrow) **c** conidiophores with scars and young conidia, shown in optical section **d** conidia shown in optical section. The thickness of the wall is only shown for the first spore from the left **e**, **f** as seen by SEM **e** part of a conidiophore with scars **f** conidia. Scale bars: 160 µm (**a**); 400 µm (**b**); 5 µm (**c**, **d**); 5 µm (**e**); 10 µm (**f**).

seen by SEM, the ornamentation of the spores is distinctly reticulated, with thin to thick networks and no ridges.

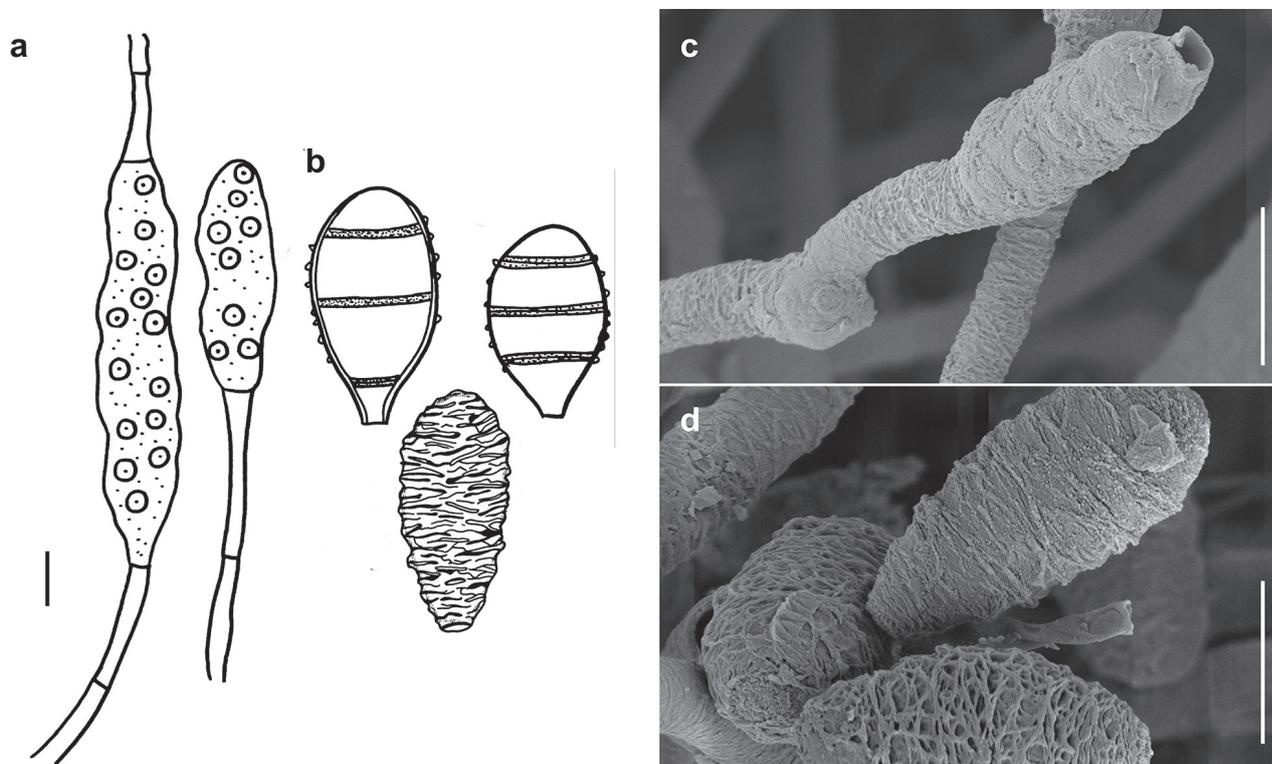
**Known distribution.** On colonies of *Meliola* cf. *xylopieae* on living leaves of *Xylopiea frutescens* (Annonaceae) in Panama.

**Notes.** *Spiropes xylopieae* is a synnematosus hyperparasitic species of *Spiropes* with the shortest synnemata (up to 400 µm), when compared to other synnematosus species, such as *S. melanoplaca* with synnemata that can reach up to 1.5 mm and *S. penicillium* with synnemata up to 700 µm high. In addition to this, the new species has crescent-shaped conidia, a feature that is not present in any other known species of the genus.

***Spiropes deightonii* M.B. Ellis, Mycol. Pap. 114: 18, 1968**

Fig. 9

**Description.** **Colonies** effuse, olive to olivaceous-brown, velvety or hairy. **Hyphae** superficial, branched, septate, 0.5–2 µm wide, pale olive to olivaceous-brown, smooth. **Conidiophores** arising singly or in groups terminally or laterally from the hyphae, erect or ascending, straight or flexuous, septate, up to 400 µm long, 2–4 µm thick along most of their length, swollen towards the apex, 5–8 µm thick, brown, reticulate as seen by SEM, with scattered cylindrical scars. **Conidia** solitary, straight or slightly curved, obovate to clavate, truncate at their base, 3–septate, (10–)12–14(–15) × (5–)6–8 µm, 1.5–2 µm wide at the base, the cells



**Figure 9.** *Spiropes deightonii* (IMI48956a) **a** conidiophores **b** conidia, as seen by LM (two upper spores; the thickness of the wall is indicated only in the drawing on the left-hand side) and by SEM (bottom spore) **c, d** as seen by SEM **c** conidiophore **d** conidia. Scale bars: 5 µm (**a, b**); 8 µm (**c**); 5 µm (**d**).

at each end of a conidium subhyaline or pale brown, intermediate cells brown, ornamented. As seen by SEM, the ornamentation of the spores is distinctly reticulated, with thin to thick networks that can form ridges.

**Specimen examined.** On *Meliola borneensis* on *Uvaria chamae*, Sierra Leone, 1951, F.C. Deighton, (IMI 48956a, type of *S. deightonii*).

**Illustrations.** This species was illustrated by Ellis (1968).

**Known hosts and distribution.** On colonies of *Meliola borneensis* on living leaves of *Uvaria chamae* (Annonaceae) in Sierra Leone (Ellis 1968).

**Notes.** *Spiropes deightonii* and *Spiropes intricatus* are the only known species of the genus that present conidiophores that swell in the areas where conidia are formed (Figs 9, 14; Ellis (1968)). *Spiropes intricatus* differs from *S. deightonii* by the presence of larger conidia (16–23 µm long) that are more oblong-ellipsoid (Ellis 1968), rather than obovate or clavate. *S. deightonii* is only known from the type specimen.

### ***Spiropes dorycarpus* (Mont.) M.B. Ellis, Mycol. Pap. 114: 27, 1968**

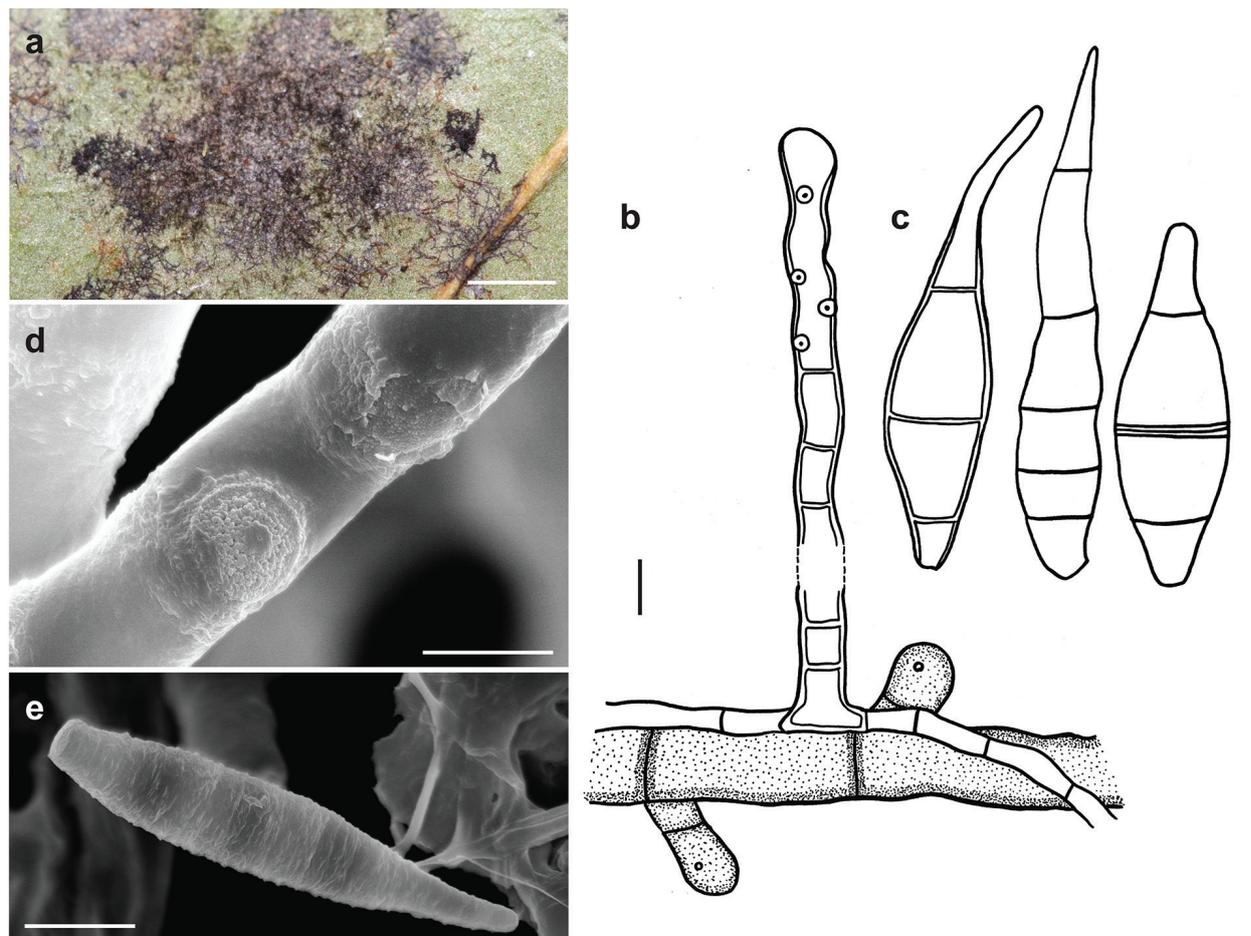
Fig. 10

- ≡ *Helminthosporium dorycarpum* Mont., Anns Sci. nat., 2 Sér., 17: 120, 1842.
- ≡ *Pleurophragmium dorycarpum* (Mont.) Hughes, Can. J. Bot. 36: 797, 1958.
- = *Helminthosporium orbiculare* Lév., Anns Sci. nat., 3 Sér., 5: 299, 1846.
- = *Napicladium myrtacearum* Speg., An. Soc. cient. Argent. 26: 71, 1888.
- ≡ *Sporhelminthium myrtacearum* (Speg.) Speg., Physis 4(17): 292, 1918.
- = *Helminthosporium conspicuum* McAlpine, Proc. Linn. Soc. N.S.W. 22: 40, 1897.
- = *Podosporium densum* Pat., J. Bot. Paris 11: 373, 1897.
- = *Helminthosporium asterinoides* Sacc. & P. Syd., apud Saccardo, Rc. Congr. Bot. Palermo, May 1902: 58, 1902.
- ≡ *Sporhelminthium asterinoides* (Sacc. & Syd.) Speg., Physis 4(17): 292, 1918.
- = *Helminthosporium melastomacearum* F. Stevens, Bot. Gaz. 65: 242, 1918.
- = *Helminthosporium panici* F. Stevens, Bot. Gaz. 65: 242, 1918.
- = *Helminthosporium parathesicola* [as '*parathesicolum*'] F. Stevens, Bot. Gaz. 65: 242, 1918.

**Description.** **Colonies** effuse, brown to dark brown, hairy. **Hyphae** superficial, branched, septate, 1–3 µm wide, straw-coloured, pale brown, smooth. **Conidiophores** arising singly or in groups, terminally or laterally from the hyphae, erect or ascending, straight or flexuous, septate, up to 700 µm long, 3–7 µm thick, straw-coloured, pale brown to brown, with scattered cylindrical scars towards the apex. **Conidia** solitary, straight or slightly curved, variable in shape, but mostly obclavate to fusiform, truncate at the base, mostly 3–septate, but sometimes with 4 to 5 septa, (16–)20–35(–40) × (4.5–)5–7 µm, straw-coloured to pale brown, middle cells slightly darker, wrinkled or verrucose. As seen by SEM, the ornamentation of the spores is distinctly reticulated, with thin to thick networks and no ridges.

**Specimen examined.** On *Meliola* sp. on living leaves of *Coffea arabica*, Benin, Atlantique, Attogon, Niaouli Forest, 6°44'23"N, 2°8'26"E, 119 m a.s.l., 19 September 2022, A. Krauß, A. Tabé, O. Koukol, N.S. Yorou, AK06H.

**Additional specimens examined.** On *Eugenia pungens*, Brasil, Guarapí, 1883, B. Balansa, 3939, (IMI 100322, type of *Napicladium myrtacearum*); on *Meliola* sp.



**Figure 10.** *Spiropes dorycarpus* (AK06H) **a** superficial hyphae growing on a colony of *Meliola* sp. on a leaf of *Coffea arabica* **b, c** in optical section **b** conidiophore growing on a hypha of *Meliola* sp. **c** conidia. The thickness of the wall is indicated only in the drawing on the left-hand side **d, e** As seen by SEM **d** conidiophore with a scar **e** conidium. Scale bars: 1 mm (**a**); 5  $\mu$ m (**b**); 3.5  $\mu$ m (**c**); 3  $\mu$ m (**d**); 7  $\mu$ m (**e**).

on leaves of an unknown plant, Cuba, R. de la Sagra (IMI 10002, type of *Helminthosporium dorycarpum*).

**Illustrations.** This species was illustrated by Ellis (1968).

**Known hosts and distribution.** On colonies of *Appendiculella* spp., *Asteridiella* spp., *Clypeolella* spp., *Irenopsis* spp., *Meliola* spp. and *Schiffnerula* spp., on living leaves of various plants in Australia, Brazil, Chile, Congo, Cuba, Dominican Republic, Ghana, Guyana, India, Malaysia, Nigeria, Puerto Rico, Sierra Leone, South Africa, Taiwan, Tanzania and Uganda (Ellis 1968). *Spiropes dorycarpus* is reported here for the first time for Benin.

**Notes.** *Spiropes dorycarpus* is similar to *S. effusus* and *S. helleri* by the presence of non-synnematous conidiophores and conidia mostly with three true septa. However, conidia of *S. effusus* are narrower (3–5  $\mu$ m) than those of *S. helleri* (7–13  $\mu$ m).

***Spiropes effusus* (Pat.) M.B. Ellis, Mycol. Pap. 114: 10, 1968**

Fig. 11

$\equiv$  *Podosporium effusum* Pat., Scient. Surv. P. Rico 8(1): 103, 1926.

= *Helminthosporium dorycarpum* var. *amazoniae* Hughes [as '*Helmisporium*'], Mycol. Pap. 50: 24, 1953.

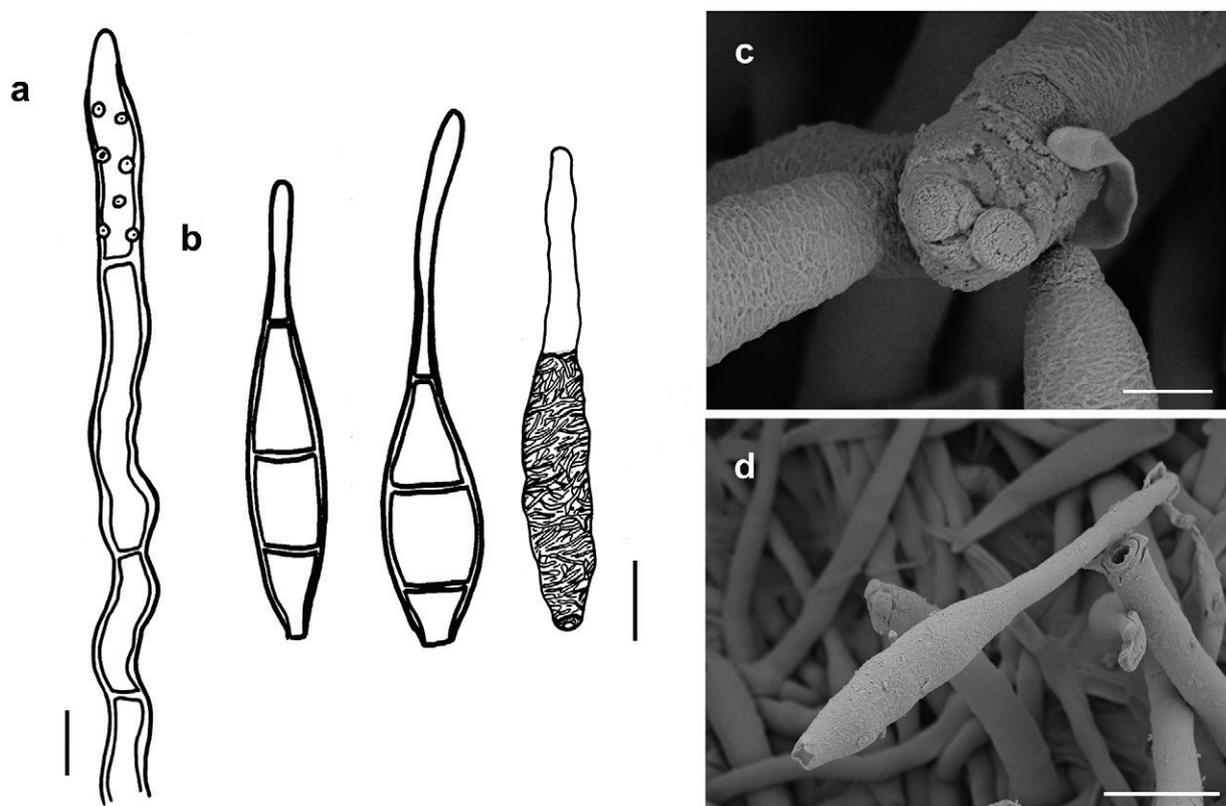
≡ *Pleurophragmium dorycarpum* var. *amazoniae* (S. Hughes) S. Hughes, Can. J. Bot. 36: 797, 1958.

**Description.** *Colonies* effuse, olive to brown, hairy. *Hyphae* superficial, branched, septate, 1–2 µm wide, yellowish, olive or pale brown, smooth. *Conidiophores* arising singly or in groups, as terminal and lateral branches on the hyphae, erect, straight or flexuous, septate, up to 300 µm long, 3–4 µm thick, slightly reticulated when seen by SEM, with few or many small conidial scars towards the apex. *Conidia* solitary, narrowly obclavate to fusiform, truncate at the base, mostly 3(–5)–septate, (15–)20–36 × (3–)3.8–4.5(–5) µm, pale brown, the central cells slightly darker, verruculose. As seen by SEM, the ornamentation of the spores is distinctly reticulated, with thin networks and no ridges.

**Specimen examined.** On meliolalean fungus on leaves of *Piper* sp., Puerto Rico, Río Piedras, 1926, Heller, 142 (IMI 130721, type of *Podosporium effusum*); on *Amazonia psychotriae* on leaves of *Psychotria warneckeii*, Ghana, Togoland, 1938, F.C. Deighton M1617B (IMI 9996a).

**Illustrations.** This species was illustrated by Ellis (1968).

**Known hosts and distribution.** On colonies of Meliolales, especially *Amazonia* spp., on living leaves of various plants in Ghana, Puerto Rico, Sierra Leone and Venezuela. One record on *Asterina* sp. (Asterinales, Ascomycota) in Uganda (Ellis 1968).



**Figure 11.** *Spiropes effusus* (IMI 130721) **a** conidiophore shown in optical section **b** conidia. The first two drawings show spores in optical section. The right-hand drawing shows a conidium as seen by SEM **c**, **d** as seen by SEM **c** conidiophore with scars and conidia **d** conidium. Scale bars: 5 µm (**a**); 8 µm (**b**); 2 µm (**c**); 8 µm (**d**).

**Notes.** *Spiropes effusus* has conidia similar in size to those of *S. dorycarpus*. However, conidia of *S. dorycarpus* are wider (5–7 µm) than in *S. effusus*.

***Spiropes fumosus* (Ellis & Martin) M.B. Ellis, Mycol. Pap. 114: 20, 1968.**

≡ *Helminthosporium fumosum* Ellis & Martin, Am. Nat. 18: 70, 1884.

≡ *Brachysporium fumosum* (Ellis & Martin) Sacc., Syll. Fung. 4: 428, 1886.

**Type.** On *Meliola* sp. on leaves of *Persea palustris* (Lauraceae), Florida, U.S.A, 1883, G. Martin (NY 830274. The type specimen was not available for loan by NY).

**Species description.** This species was described by Ellis (1968).

**Known hosts and distribution.** On colonies of *Meliola* sp. on living leaves of *Persea palustris* in the U.S.A. (Ellis 1968).

**Specimen examined.** On Meliolales on living leaves of *Persea palustris*, U.S.A, Florida, Cove Springs, 1890, G. Martin, (IMI 16307).

**Illustrations.** This species was illustrated by Ellis (1968).

**Notes.** The specimen IMI 16307 was analysed, but no fungal cells were seen.

***Spiropes guareicola* (F. Stevens) Cif., Sydowia 9(1–6): 302, 1955**

Fig. 12

≡ *Helminthosporium guareicola* F. Stevens [as '*Helmisporium guareicolum*'], Bot. Gaz. 65(3): 241, 1918.

≡ *Pleurophragmium guareicola* (F. Stevens) S. Hughes, Can. J. Bot. 36: 797, 1958.

= *Cladosporium elegans* var. *singaporensis* Sacc., Bull. Orto Bot. Regia Univ. Napoli 6: 60, 1921.

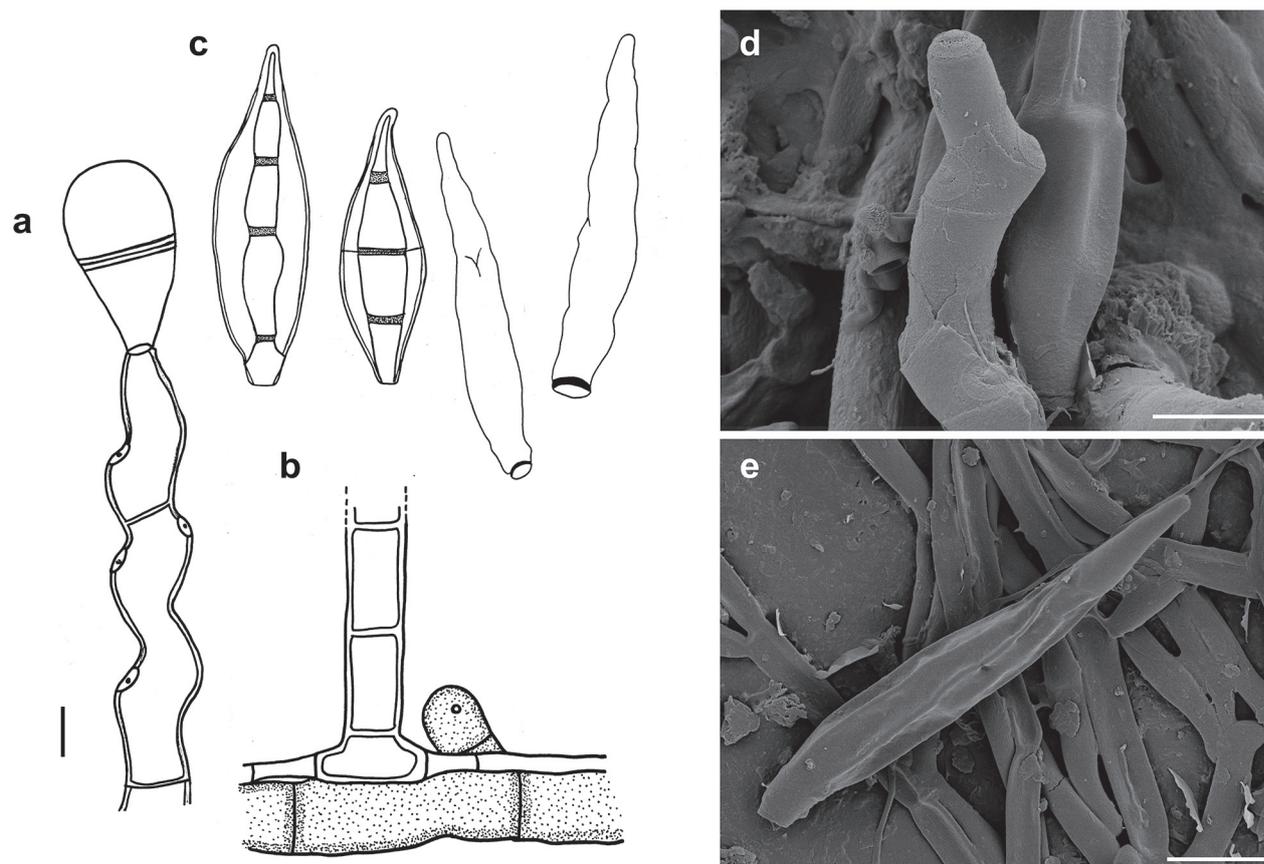
= *Helminthosporium flagellatum* H.S. Yates [as '*Helmisporium*'], Philipp. J. Sci. (Bot.) 13: 383, 1918.

= *Helminthosporium spirotrichum* Sacc. [as '*Helmisporium*'], Boll. Orto bot. 6: 61, 1921.

**Description.** **Colonies** effuse, dark brown to black, hairy. **Hyphae** superficial, branched, septate, 2–4 µm wide, pale olivaceous-brown, smooth. **Conidiophores** arising singly or in groups, as lateral branches on the hyphae, erect, sterile lower part straight or flexuous, upper fertile part in zigzag shape, septate, up to 400 µm long, 6–9 µm thick, brown to dark brown, paler towards the apex, more or less smooth, with numerous well-defined, dark conidial scars. **Conidia** solitary, broadly fusiform, truncate at the base, with 3 to 5 pseudosepta, (25–)35–52(–60) × (7–)8–10(–13) µm, 3.5–5 µm wide at the base, pale to dark brown or olivaceous-brown, smooth as seen by SEM.

**Specimen examined.** On leaves of *Cyrtophyllum fragrans* (Gentianaceae), Singapore, 1921, Baker (IMI 49160, type of *Helminthosporium spirotrichum*); on *Meliola* sp. on leaves of *Daniellia thurifera* (Fabaceae), Sierra Leone, 1936, F.C. Deightonii M1267 (IMI 10010).

**Illustrations.** This species was illustrated by Ellis (1968).



**Figure 12.** *Spiropes guareicola* (IMI 10010) **a** conidiophore with scars and a young conidium shown in optical section **b** base of a conidiophore growing on a hypha of *Meliola* sp. shown in optical section **c** conidia shown in optical section (two drawings on the left-hand side) and as seen by SEM (two drawings on the right-hand side) **d, e** as seen by SEM **d** zigzag-shaped conidiophore with scars **e** conidium. Scale bars: 5 µm (**a–c**); 8 µm (**d**); 10 µm (**e**).

**Known hosts and distribution.** On colonies of *Asteridiella* spp., *Irenopsis* spp. and *Meliola* spp. on living leaves of various plants in Bougainville Islands, Ghana, India, Indonesia, Malaysia, Philippines, Puerto Rico, Sabah, Sierra Leone, Solomon Islands and Uganda (Ellis 1968).

**Notes.** *Spiropes guareicola* is the type species of the genus *Spiropes* and it differs from other species of the genus by the presence of zigzag-shaped conidiophores in the fertile upper parts (Ellis 1968). *S. guareicola* presents smooth conidia, a feature that is only evident by SEM.

***Spiropes helleri* (F. Stevens) M.B. Ellis, Mycol. Pap. 114: 14, 1968**

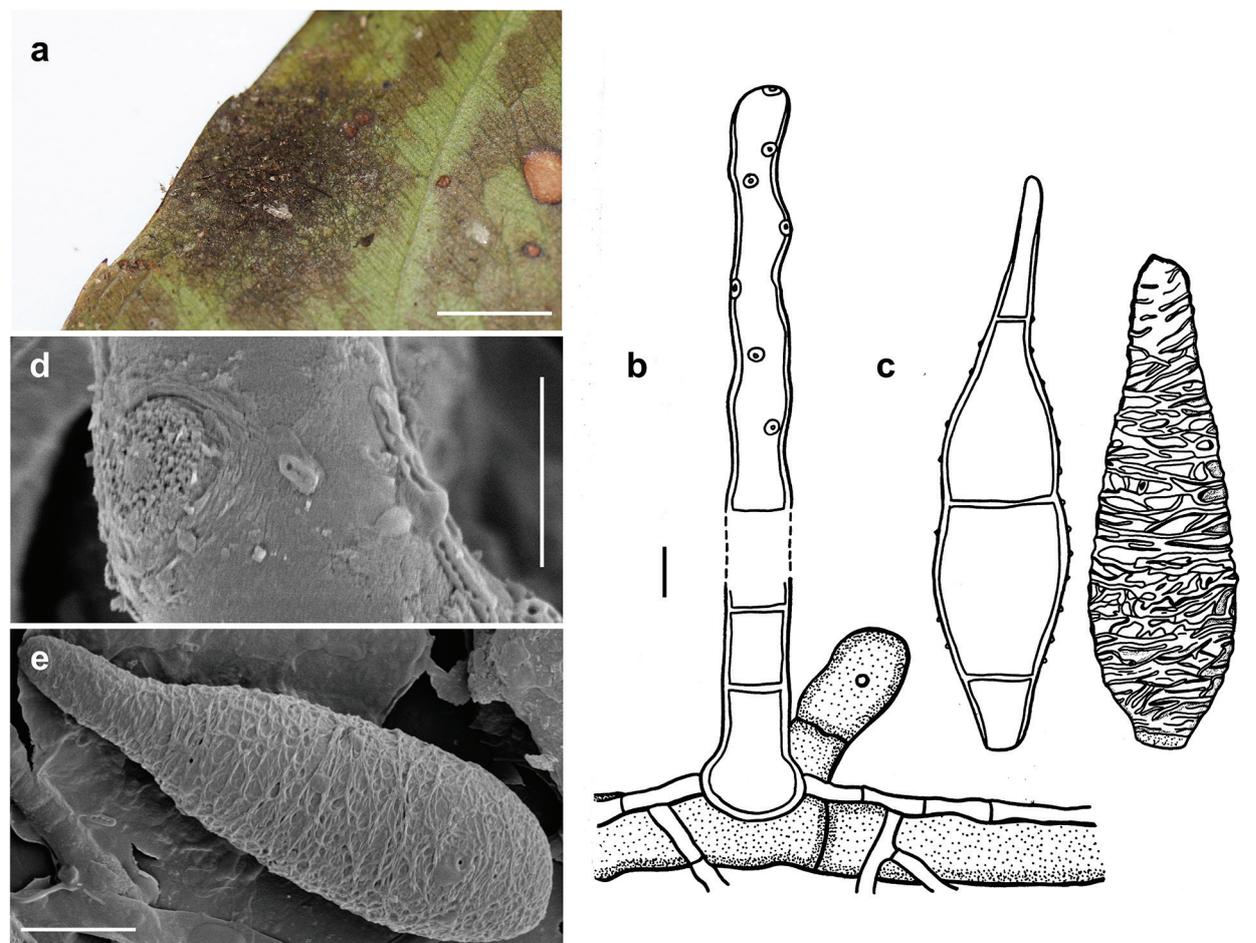
Fig. 13

≡ *Helminthosporium helleri* F. Stevens [as '*Helmisporium*'], Bot. Gaz. 65(3): 242, 1918.

= *Helminthosporium leucosykes* H.S. Yates [as '*Helmisporium leucosykeae*'], Philipp. J. Sci., C, Bot. 13(6): 382, 1918.

= *Helminthosporium maculosum* Sacc. [as '*Helmisporium*'], Atti Accad. Sci. Ven.-Trent.-Istr. 10: 91, 1919 [1917].

≡ *Pleurophragmium maculosum* (Sacc.) S. Hughes, Can. J. Bot. 36: 797, 1958.



**Figure 13.** *Spiropes helleri* (IMI130940) **a** superficial hyphae growing on a colony of *Meliola* sp. on a leaf of *Cupania guatemalensis* **b** conidiophore growing on a hypha of *Meliola* sp. shown in optical section **c** conidia shown in optical section (drawing on the left-hand side) and as seen by SEM (drawing on the right-hand side) **d, e** as seen by SEM **d** part of a conidiophore with a scar **e** conidium. Scale bars: 1 mm (**a**); 5  $\mu\text{m}$  (**b**); 6  $\mu\text{m}$  (**c**); 4  $\mu\text{m}$  (**b**); 5  $\mu\text{m}$  (**c**).

**Description.** **Colonies** effused, dark brown to black, hairy. **Hyphae** superficial, branched, septate, 1–3  $\mu\text{m}$  wide, straw-coloured or pale brown, smooth. **Conidiophores** arising singly as terminal or lateral branches on the hyphae, erect, straight or flexuous, septate, up to 600  $\mu\text{m}$  long, 5–8  $\mu\text{m}$  wide, brown to dark brown, paler towards the apex, smooth, with scattered conidial scars. **Conidia** solitary, obclavate, frequently rostrate, 3(–4)–septate, (26–)36–43(–50)  $\times$  (6–)7–10(–13)  $\mu\text{m}$ , 3–4  $\mu\text{m}$  wide at the truncate base, pale brown to brown, verruculose. As seen by SEM, the ornamentation of the spores is clearly reticulated, with thin networks and no ridges.

**Specimens examined.** On *Meliola* sp. on leaves of *Cupania guatemalensis* (Sapindaceae), Panama, Chiriquí Province, Botanical Garden of the Autonomous University of Chiriquí (UNACHI), 8°25'55"N, 82°27'03"W, 34 m a.s.l., 11 February 2020, M. A. Bermúdez-Cova, A. Sanjur MB92 (UCH15489, M); on *Meliola* sp. on living leaves of *Pterocarpus santalinoides* (Fabaceae), Benin, Atlantique, Attogon, Niaouli Forest, 6°44'40"N, 2°7'53"E, 72 m a.s.l., 20 September 2022, A. Krauß, A. Tabé, O. Koukol, N.S. Yorou, AK15 (M).

**Additional specimens examined.** On Meliolales on living leaves of an undetermined plant, Gold Coast Colony, Banau, 1949, S.J. Hughes 1141 (IMI44564);

on *Meliola* sp. on leaves of *Myrcia deflexa*, Puerto Rico, El Alto de la Bandera, F.L. Stevens 8268 (IMI9991, type of *Helminthosporium helleri*).

**Illustrations.** This species was illustrated by Ellis (1968).

**Known hosts and distribution.** On colonies of *Asteridiella* spp., *Irenopsis* spp. and *Meliola* spp. on living leaves of various plants in Ghana, Malaysia, New Caledonia, Philippines, Puerto Rico, Sabah, Sierra Leone and Uganda (Ellis 1968). *Spiropes helleri* is reported here for the first time for Benin and for mainland America (Panama).

**Notes.** *Spiropes helleri* is similar to *S. effusus*, *S. dorycarpus* and *S. leonensis* by the presence of obclavate to sometimes fusiform conidia, but differs from the first two by wider conidia (3.8–4.5 µm in *S. effusus* and 5–7 µm in *S. dorycarpus*) and from the last one by narrower ones (10–11 µm).

### ***Spiropes intricatus* (Sacc.) M.B. Ellis, Mycol. Pap. 114: 9, 1968**

Fig. 14

≡ *Brachysporium intricatum* Sacc., Atti Accad. scient. Veneto-trent.-istriana, Ser. 3, 10: 88, 1919.

= *Spiropes pirozynskii* M.B. Ellis, Mycol. Pap. 114: 19, 1968. New synonym proposed in this study.

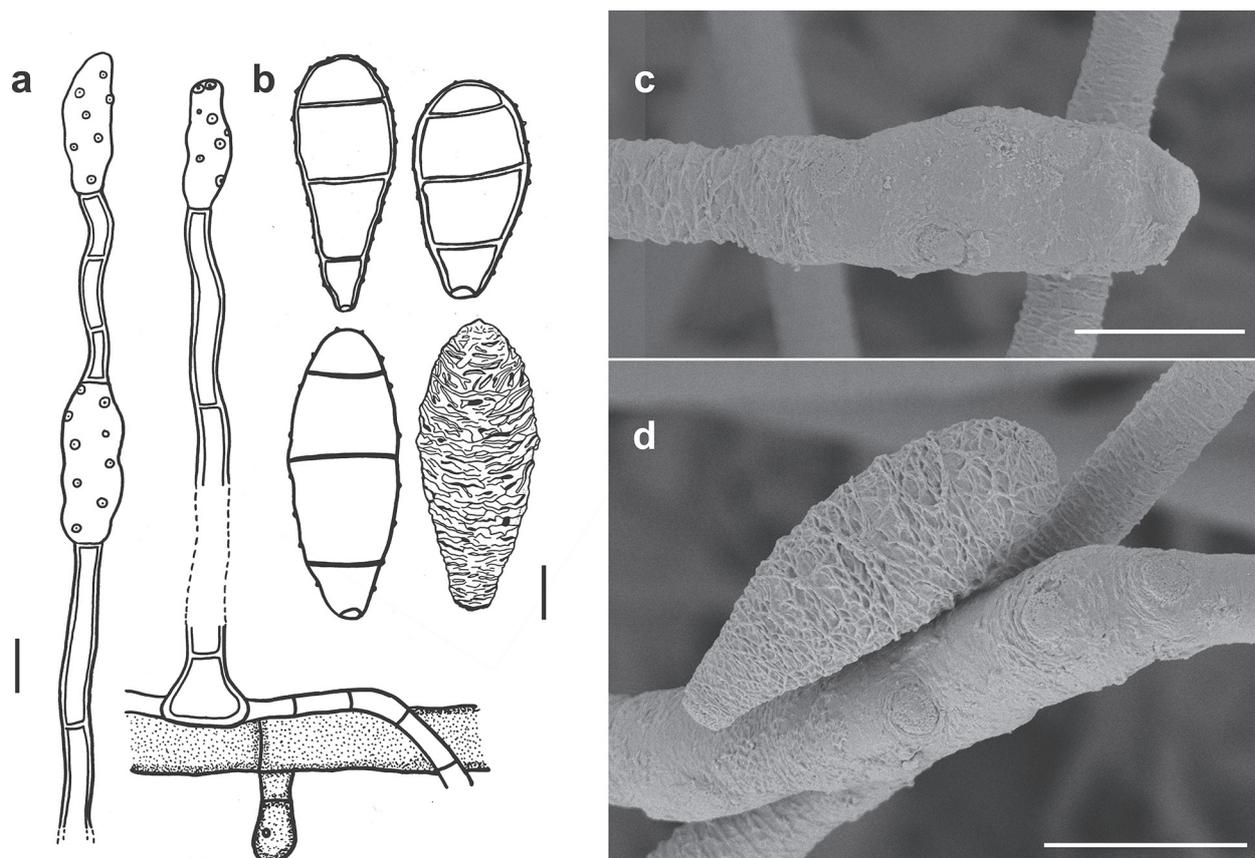
**Description.** **Colonies** effuse, straw-coloured, olive or olivaceous-brown, velvety or hairy. **Hyphae** superficial, branched, anastomosing, septate, 1–2 µm wide, pale olivaceous brown, smooth. **Conidiophores** arising singly or in groups, terminally or laterally from the hyphae, erect or ascending, straight or flexuous, septate, up to 900 µm long, 2–5 µm thick along most of their length, swollen to 4–9 µm towards the apex and in intercalary parts that produce conidia, pale olivaceous-brown to brown, reticulate as seen by SEM, with scattered cylindrical scars. **Conidia** solitary, straight or slightly curved, oblong-ellipsoid or obovate to clavate, truncate at the base, mostly 3-septate, (13–)16–23(–25) × (4.5–)6–8 µm, 1.5–3 µm wide at the base, the cells at each end of a conidium pale brown, intermediate cells brown, ornamented. As seen by SEM, the ornamentation of the spores is distinctly reticulated, with thin to thick networks that can form ridges.

**Specimens examined.** On *Irenopsis* sp. on *Lindackeria bukobensis* (Achariaceae), Tanzania, Kigoma, 1964, K.A. Pirozynski M418 b&c (IMI 106645b-c, type of *Spiropes pirozynskii*); on leaves of *Camellia drupifera* (Theaceae), Nepal, Kathmandu, Godawari, 1986, U. Budathoki KU294 (IMI323287).

**Illustrations.** This species was illustrated by Ellis (1968).

**Known hosts and distribution.** On colonies of Meliolales on living leaves of various plants in Ghana, Philippines and Tanzania (Ellis 1968).

**Notes.** *Spiropes intricatus* and *S. deightonii* are the only known species of the genus that present conidiophores that swell in the areas where conidia are formed (Figs 9, 14; Ellis (1968)). *Spiropes deightonii* differs from *S. intricatus* by the presence of smaller conidia (12–14 µm long) that are more obovate or clavate rather than oblong-ellipsoid. The type specimen of *S. pirozynskii* (IMI 106645b-c) is morphologically similar to *S. intricatus*. Both species present oblong-ellipsoid conidia with a similar size range (Fig. 15). Therefore, we propose *S. pirozynskii* as a synonym of *S. intricatus*.



**Figure 14.** *Spiropes intricatus* (IMI 106645b-c) **a** conidiophores, growing on a hypha of *Irenopsis* sp., shown in optical section **b** conidia shown in optical section (the thickness of the wall is indicated only in the drawings on the upper row) and as seen by SEM (second row right) **c, d** as seen by SEM **c** conidiophore with scars **d** conidium. Scale bars: 5 μm (**a**); 3 μm (**b**); 7 μm (**c**); 8 μm (**d**).

***Spiropes japonicus* (Henn.) M.B. Ellis, Mycol. Pap. 114: 22, 1968**

Fig. 16

≡ *Podosporium japonicum* Henn., Bot. Jb. 29: 152, 1900.

= *Helminthosporium insigne* Gaillard ex Sacc. [as '*Helmisporium*'], Atti Accad. Sci. Ven.-Trent.-Istr. 10: 89, 1917.

**Description.** **Colonies** effuse, amphigenous, sometimes dense, dark brown to black, with tightly packed hyphae that form large, erect, dark synnemata clearly visible under the stereomicroscope. **Hyphae** superficial, branched, septate, 1–4 μm wide, pale olivaceous-brown, smooth. **Conidiophores** tightly packed to form dark brown to blackish synnemata up to 1 mm high, spreading out at the apex and upper half of the synnemata; conidiophores individually flexuous or straight, thick-walled, septate, 6–8 μm thick, brown to dark brown at the base, paler towards the apex, smooth, with scattered cylindrical scars. **Conidia** solitary, fusiform to obclavate, with 4(–6) pseudosepta, (50–)67–80 × (7–)8–14 μm, 2–3 μm wide at the apex, 3–5 μm at the truncate base, pale brown to brown, striate.

**Specimens examined.** On *Meliola* sp. on living leaves of Asteraceae, Panama, Chiriquí Province, Boquerón District, Chuspa Hydroelectric, 8°32'20"N, 82°36'21"W, 281 m a.s.l., 6 March 2020, M. A. Bermúdez-Cova, A. Sanjur, S.

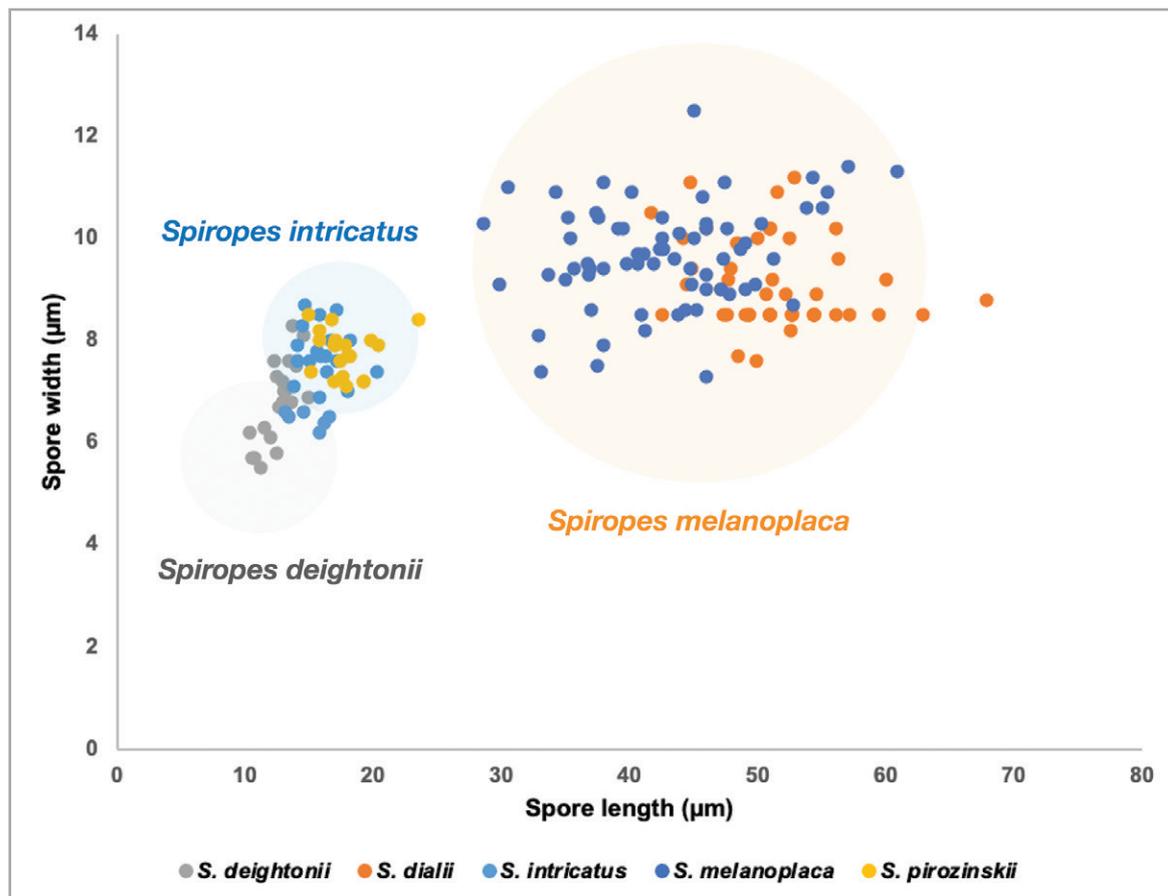


Figure 15. Scatter plot of spore size (width and length) of species of *Spiropes*.

Samaniego, MB120 (UCH15492); on *Meliola* sp. on living leaves of Fabaceae, Panama, Chiriquí Province, Bugaba District, area around Gariché River, 8°38'38.1"N, 82°41'19.6"W, 566 m a.s.l., 8 March 2020, M. A. Bermúdez-Cova, A. Sanjur, A. Villarreal, MB123 (UCH15493, M).

**Additional specimens examined.** On *Irenina entebbeensis* on *Alchornea hirtella* (Euphorbiaceae), Sierra Leone, 1939, Makump, M1774 (IMI 38813); on *Asteridiella aucubae* on *Aucuba japonica* (Garryaceae), Japan, Ise, 1899, P. Hennings (IMI 130973, type of *Podosporium japonicum*).

**Illustrations.** This species was illustrated by Ellis (1968).

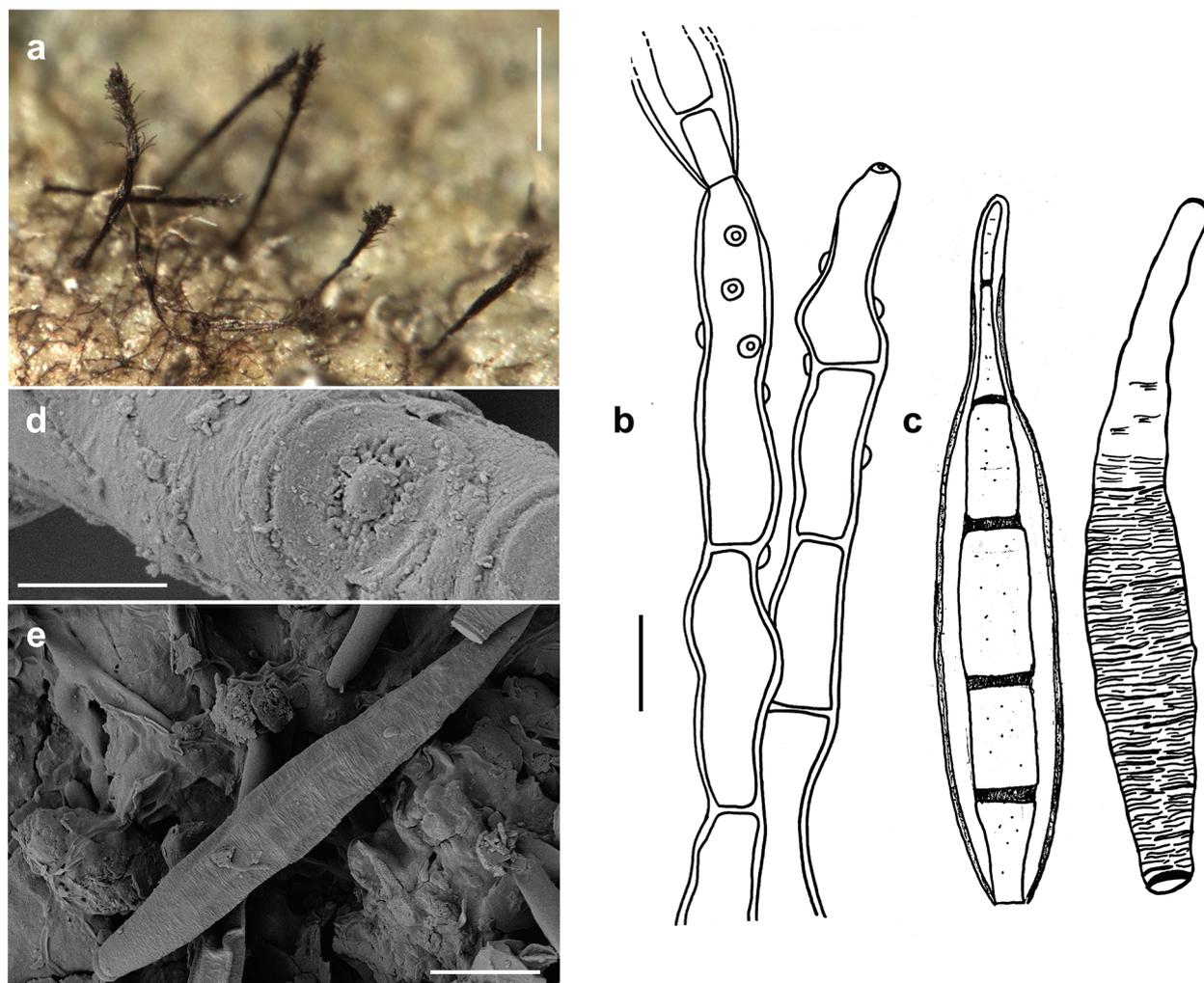
**Known hosts and distribution.** On colonies of Meliolales on living leaves of various plants in the Cook Islands, Japan, Malaysia, Papua New Guinea and Sierra Leone (Ellis 1968). *Spiropes japonicus* is reported here for the first time for Panama.

**Notes.** *Spiropes japonicus* is the only known synnematosus species of *Spiropes* that produces conidia with 4–6 pseudosepta, as well as synnemata that splay out at the apex and upper half (Ellis 1968).

### *Spiropes leonensis* M.B. Ellis, Mycol. Pap. 114: 15, 1968

Fig. 17

**Description.** **Colonies** effuse, grey to dark blackish-brown, hairy. **Hyphae** superficial, branched, septate, 2–6 µm wide, pale brown, smooth. **Conidiophores**



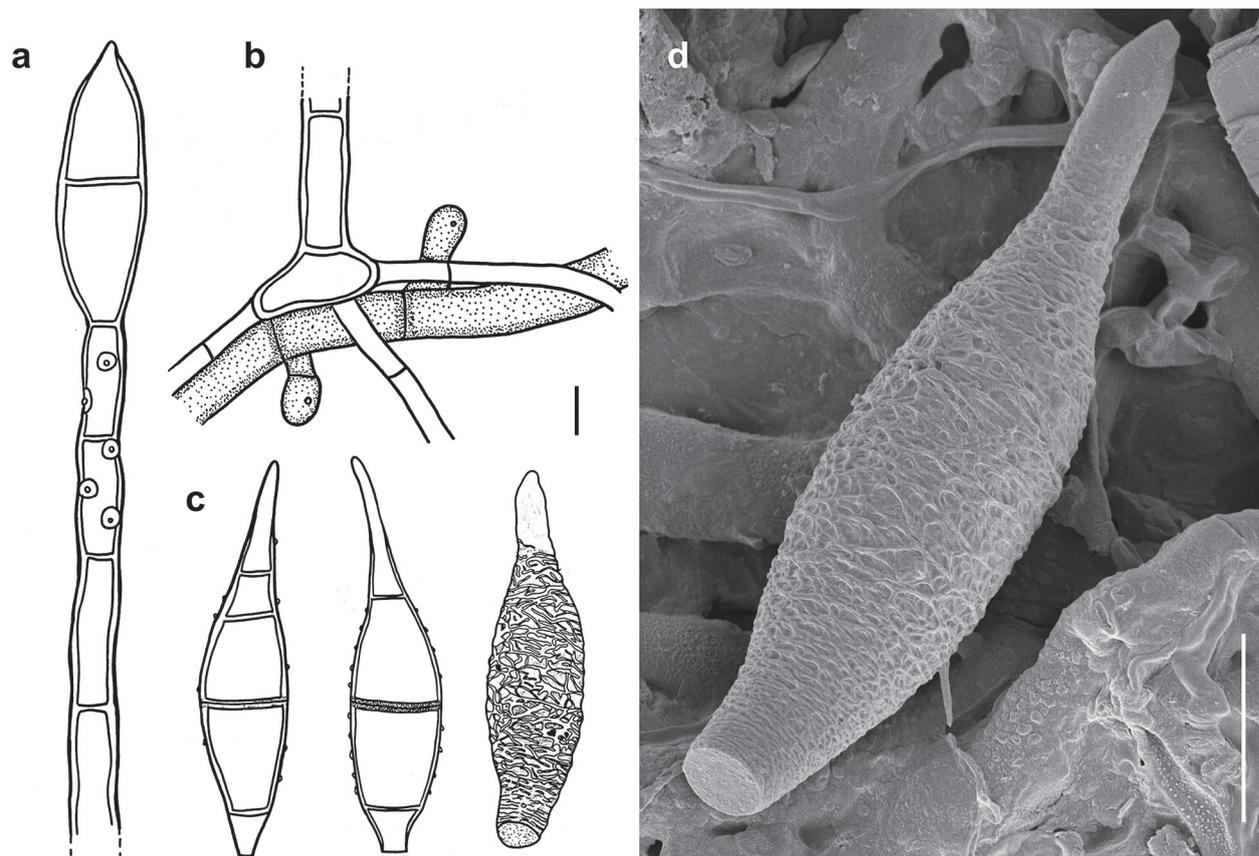
**Figure 16.** *Spiropes japonicus* (MB120, 123) **a** synnemata growing on a colony of *Meliola* sp. **b** conidiophores with scars and a young conidium, shown in optical section **c** a conidium shown in optical section (drawing on the left) and as seen by SEM (drawing on the right) **d, e** as seen by SEM **d** conidiophore with a scar **e** conidium. Scale bars: 1 mm (**a**); 10  $\mu$ m (**b, c**); 3  $\mu$ m (**d**); 9  $\mu$ m (**e**).

arising singly, as terminal and lateral branches on the hyphae, erect, straight or flexuous, septate, up to 700  $\mu$ m long, 8–12  $\mu$ m thick, sometimes swollen to 16–17  $\mu$ m at the base, dark brown to dark blackish-brown, paler towards the apex, smooth, with scattered conidial scars. **Conidia** solitary, obclavate, rostrate, 3(–4)–septate, (38–)40–54(–63)  $\times$  (8–)10–11(–13)  $\mu$ m, 4–6  $\mu$ m wide at the truncate base, pale brown to brown, verruculose. As seen by SEM, the ornamentation of the spores is distinctly reticulated, with thin networks and no ridges. It was not possible to see the scars by SEM.

**Specimen examined.** On *Meliola garciniae* on leaves of *Pentadesma butyracea* (Clusiaceae), Sierra Leone, Rokupr, 1951, F.C. Deighton M3920 (IMI 46589b, holotype); on *Meliola garciniae* on *Pentadesma butyracea*, Sierra Leone, near Rokupr, 1939, F.C. Deighton (IMI 9992a, type of *Spiropes leonensis*).

**Illustrations.** This species was illustrated by Ellis (1968).

**Known hosts and distribution.** On colonies of *Meliola garciniae* on living leaves of *Pentadesma butyracea* (Clusiaceae) in Sierra Leone (Ellis 1968).



**Figure 17.** *Spiropes leonensis* (IMI 46589b) **a** conidiophore with scars and a young conidium, shown in optical section **b** part of a conidiophore growing on a hypha of *Meliola* sp., shown in optical section **c** conidia shown in optical section (first two drawings, from left to right) and as seen by SEM **d** conidium as seen by SEM. Scale bars: 8.5 µm (**a–c**); 7 µm (**d**).

**Notes.** *Spiropes leonensis* is similar to *S. helleri* by the presence of rostrate, obclavate, 3-septate conidia (Ellis 1968). However, conidia in *S. helleri* are smaller (36–43 µm).

***Spiropes melanoplaca* (Berk. & M.A. Curtis) M.B. Ellis, Mycol. Pap. 114: 28, 1968**  
Fig. 18

= *Arthrobotryum melanoplaca* Berk. & M.A. Curtis, J. Linn. Soc. Bot. 10(46): 360, 1868.

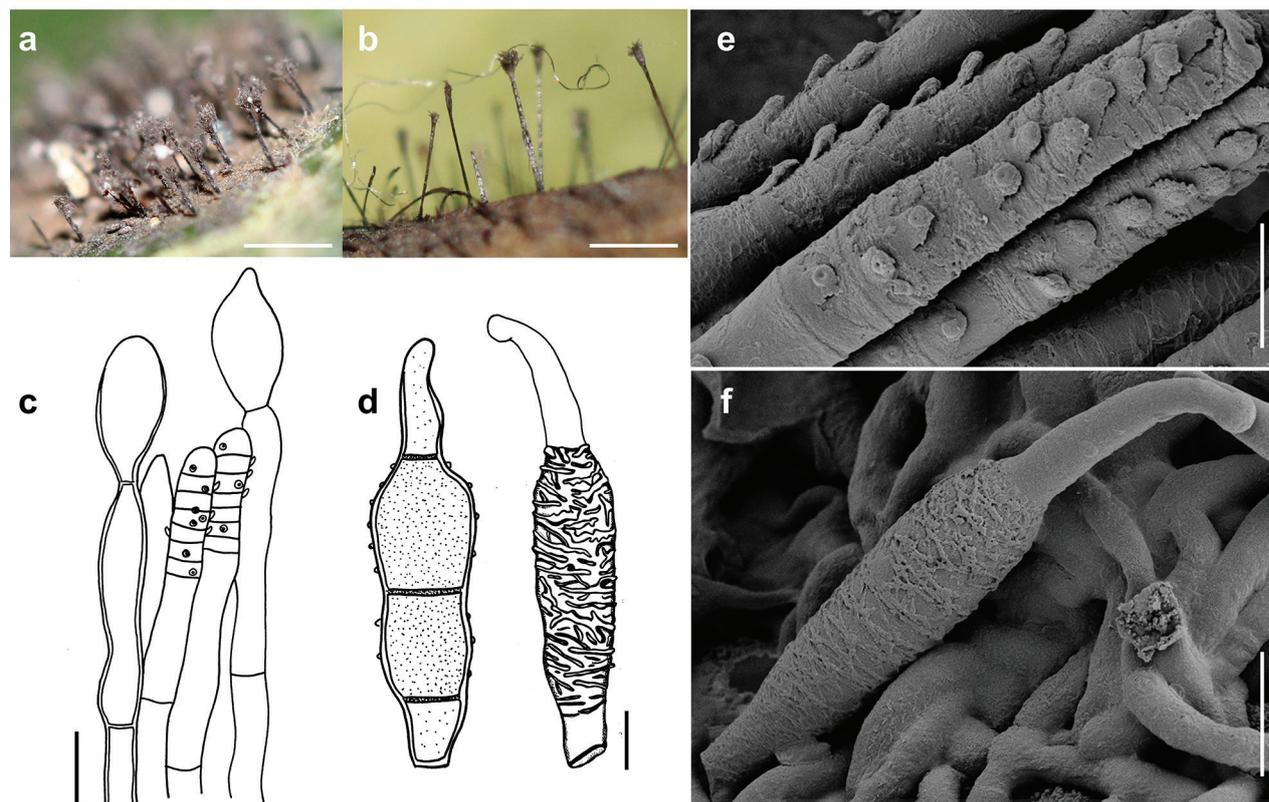
≡ *Podosporium melanoplaca* (Berk. & M.A. Curtis) Cif., Sydowia 9(1–6): 310, 1955.

= *Podosporium dialii* Bat. [as '*dialiumii*'], Atas Inst. Micol. 1: 266, 1960. New synonym proposed in this study.

≡ *Spiropes dialii* (Bat.) M.B. Ellis, Mycol. Pap. 114: 27, 1968. New synonym proposed in this study.

= *Arthrobotryum scoparium* Henn., Hedwigia 43(6): 397, 1904. New synonym proposed in this study.

**Description.** **Colonies** effuse, dark brown to black, hairy, with tightly packed hyphae that form large, erect, dark synnemata clearly visible under the



**Figure 18.** *Spiropes melanoplaca* (MB81, MB119, IMI189570a) **a, b** synnemata growing on hyphae of *Meliola mangiferae* on living leaves of *Mangifera indica* **c** conidiophores with scars and young conidia shown in optical section. The thickness of the wall is only shown in the first conidiophore, from left to right **d** conidia, shown in optical section (left-hand drawing) and as seen by SEM (right-hand drawing) **e, f** as seen by SEM **e** parts of conidiophores with scars **f** conidium. Scale bars: 1.5 mm (**a**); **b**); 0.9 mm (**c**); 8  $\mu$ m (**d**); 7  $\mu$ m (**e**); 8  $\mu$ m (**f**).

stereomicroscope. **Hyphae** superficial, branched, septate, 1.5–6  $\mu$ m wide, pale olivaceous, smooth. **Conidiophores** tightly packed to form dark brown to blackish synnemata up to 1.5 mm high, spreading out at the apex, 20–80  $\mu$ m thick, splaying out at the apex. Individual hyphae straight or flexuous, cylindrical, 2–6  $\mu$ m thick along most of their length, 5–8  $\mu$ m thick near the apex, with numerous small scars that may overlap like scales. As evident by SEM, the scales are produced by the peeling of the outer wall layers where the scars are located. **Conidia** straight or curved, fusiform to obclavate, 3-septate, (30–)40–52(–68)  $\times$  (7–)9–11(–14)  $\mu$ m, with the two middle cells usually golden brown or brown, warty and the cells at each end paler. As seen by SEM, the ornamentation of the spores is distinctly reticulated, with thin to thick networks and no ridges.

**Specimens examined.** On *Meliola mangiferae* on living leaves of *Mangifera indica* (Anacardiaceae), Panama, Chiriquí Province, Los Algarrobos, 8°31'05"N, 82°25'25"W, 168 m a.s.l., 20 January 2020, M. A. Bermúdez-Cova, MB81; same fungal and plant host, Panama, Chiriquí Province, Universidad Autónoma de Chiriquí (UNACHI), 8°25'57"N, 82°27'02"W, 37 m a.s.l., 23 January 2020, M. A. Bermúdez-Cova, MB85 (UCH15487); same fungal and plant host, Panama, Chiriquí Province, Los Algarrobos, Majagua River Trail, 8°28'56"N, 82°24'47"W, 101 m a.s.l., 23 January 2020, M. A. Bermúdez-Cova, MB89 (UCH15488, M); same fungal and plant host, Panama, Chiriquí Province, Meseta de Chorcha, 8°24'19"N, 82°13'26"W, 94 m a.s.l., 16 February 2020, M. A. Bermúdez-Cova, A. Sanjur, MB101 (UCH);

same fungal and plant host, Panama, Chiriquí Province, Boquerón District, Hidroeléctrica Chuspa, 8°33'37"N, 82°36'22"W, 331 m a.s.l., 6 March 2020, M. A. Bermúdez-Cova, A. Sanjur, S. Samaniego, MB119 (UCH15491); On *Meliola* sp. on living leaves of *Angylocalyx oligophyllus* (Fabaceae), Benin, Attogon, Niaouli, Niaouli Forest, 6°44'42"N, 2°7'50"E, 69 m a.s.l., 28 February 2022, M.A. Bermúdez-Cova, A. Tabé, I. Agonglo, O.P. Agbani, M. Piepenbring, N.S. Yorou, MB173 (M); on *Meliola mangiferae* on living leaves of *Mangifera indica*, Benin, Attogon, Niaouli, Niaouli Forest, 6°44'44"N, 2°7'49"E, 65 m a.s.l., 28 February 2022, M.A. Bermúdez-Cova, A. Tabé, I. Agonglo, O.P. Agbani, M. Piepenbring, N.S. Yorou, MB180 (M).

**Additional specimens examined.** On *Meliola mangiferae* on *Mangifera indica*, Brunei, 1974, W.T.H. Peregrine (IMI189570a); on *Meliola* sp. on *Psychotria* sp. (Rubiaceae), Cuba, 1879, C. Wright (IMI 105348 and IMI 105349, syntypes of *Arthrobotryum melanoplaca*).

**Illustrations.** This species was illustrated by Ellis (1968).

**Known hosts and distribution.** On colonies of Meliolales, especially *Meliola* spp., on living leaves of various plants in Brazil, Cuba, China, Dominican Republic, Ghana, Guadalcanal, India, Malaysia, Peru, Philippines, Sierra Leone, Tanzania, Trinidad and Uganda (Ellis 1968; Zhao et al. 1996; Dubey and Moonnambeth 2013). *Spiropes melanoplaca* is reported here for the first time for Benin and Panama.

**Notes.** According to Ellis (1968), the main difference between *Spiropes melanoplaca* and *S. dialii* is the range of spore width, with *S. melanoplaca* having wider spores (9–14 µm wide) than *S. dialii* (7–9 µm wide). However, after revision of several specimens and herbarium material from both species, we noticed that the aspect of the colonies, morphological features (both as seen in LM and by SEM) are similar between the species and both species present conidia with a similar size range (Fig. 15). Therefore, we propose *S. dialii* as a synonym of *S. melanoplaca*.

### ***Spiropes palmetto* (W.R. Gerard) M.B. Ellis, Mycol. Pap. 114: 16, 1968**

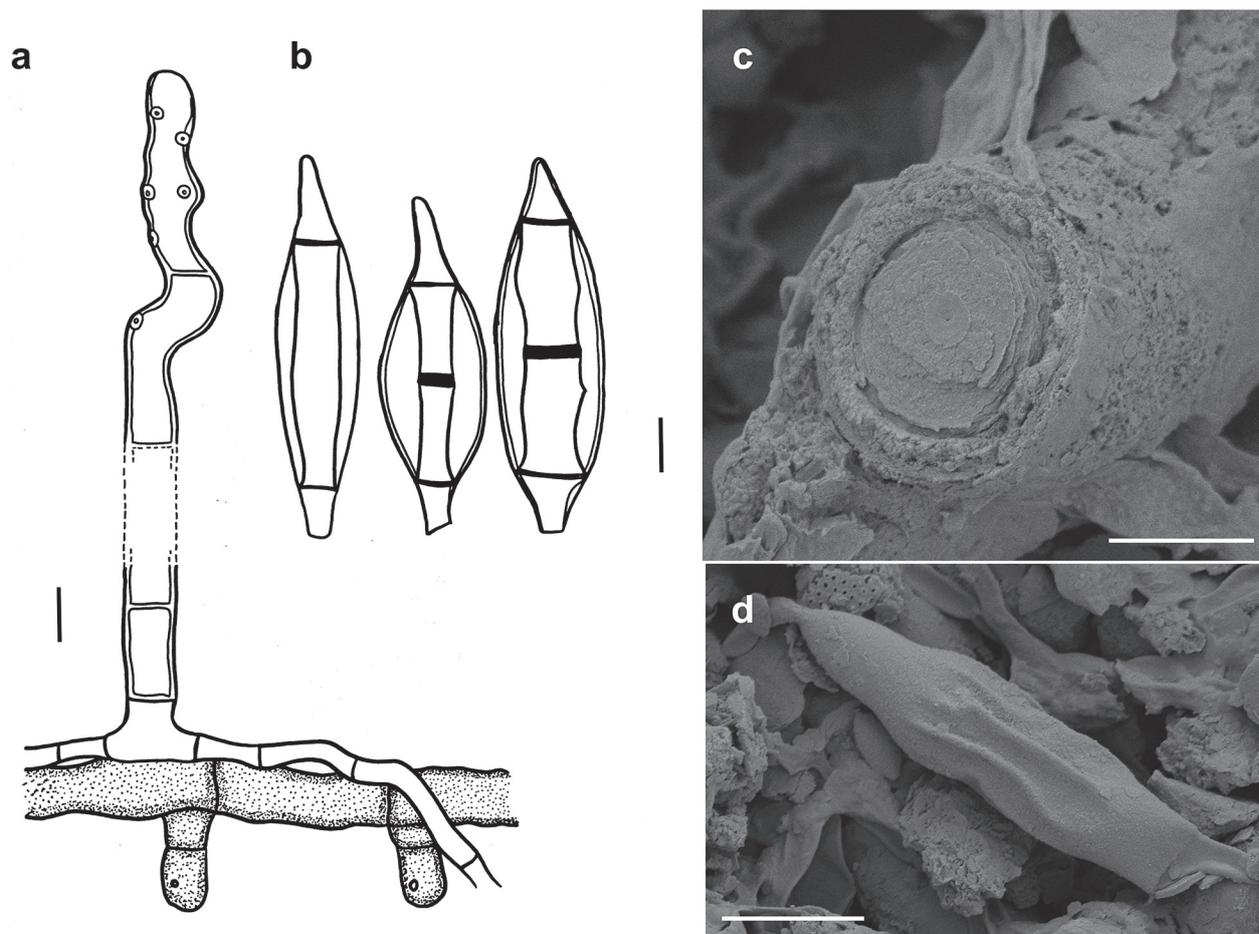
Fig. 19

≡ *Helminthosporium palmetto* W.R. Gerard, Grevillea 17(83): 68, 1889.

≡ *Pleurophragmium palmetto* (W.R. Gerard) S. Hughes, Can. J. Bot. 36: 778, 1958.

**Description.** **Colonies** effuse, dark brown to black, hairy. **Hyphae** superficial, branched, anastomosing, septate, 1–4 µm wide, pale olivaceous-brown, smooth. **Conidiophores** arising singly or in groups, as terminal and lateral branches on the hyphae, erect, straight or flexuous, septate, up to 400 µm long, 6–10 µm thick, dark brown, paler towards the apex, smooth, with scattered conidial scars. **Conidia** solitary, obclavate to fusiform, rostrate, with 2 septa delimiting a barrel-shaped central cell and often with an additional dark central pseudoseptum, (27–)30–46 × (7–)9–12(–15) µm, 3–5 µm wide at the truncate base, brown, middle cells pale brown, smooth as seen by LM and SEM.

**Specimens examined.** On *Meliola* sp. on leaves of *Elaeis guineensis* (Arecaceae), Ghana, Apremodo, 1949, S.J. Hughes 534 (IMI 38617); on *Meliola* sp. on leaves of *Sabal palmetto* (Arecaceae), U.S.A, Louisiana (IMI 10032, type of *Helminthosporium palmetto*).



**Figure 19.** *Spiropes palmetto* (IMI 10032) **a** conidiophore growing on a hypha of *Meliola* sp., shown in optical section **b** conidia shown in optical section. The thickness of the walls is only shown in the two last drawings **c**, **d** as seen by SEM **c** part of a conidiophore with a scar **d** conidium. Scale bars: 7 µm (**a**); 5 µm (**b**); 6 µm (**c**); 7 µm (**d**).

**Illustrations.** This species was illustrated by Ellis (1968).

**Known hosts and distribution.** On colonies of *Irenopsis* spp. and *Meliola* spp. on living leaves of various plants in Ghana, Malaysia, New Zealand, Puerto Rico, Sierra Leone and the U.S.A. (Ellis 1968).

**Notes.** *Spiropes palmetto* can be easily recognised by the presence of conidia with two septa that delimit a barrel-shaped central cell and with a dark central pseudoseptum (Ellis 1968).

***Spiropes penicillium* (Speg.) M.B. Ellis, Mycol. Pap. 114: 23, 1968**

Fig. 20

≡ *Podosporium penicillium* Speg., Boln. Acad. nac. Cienc. Córdoba 11: 618, 1889.

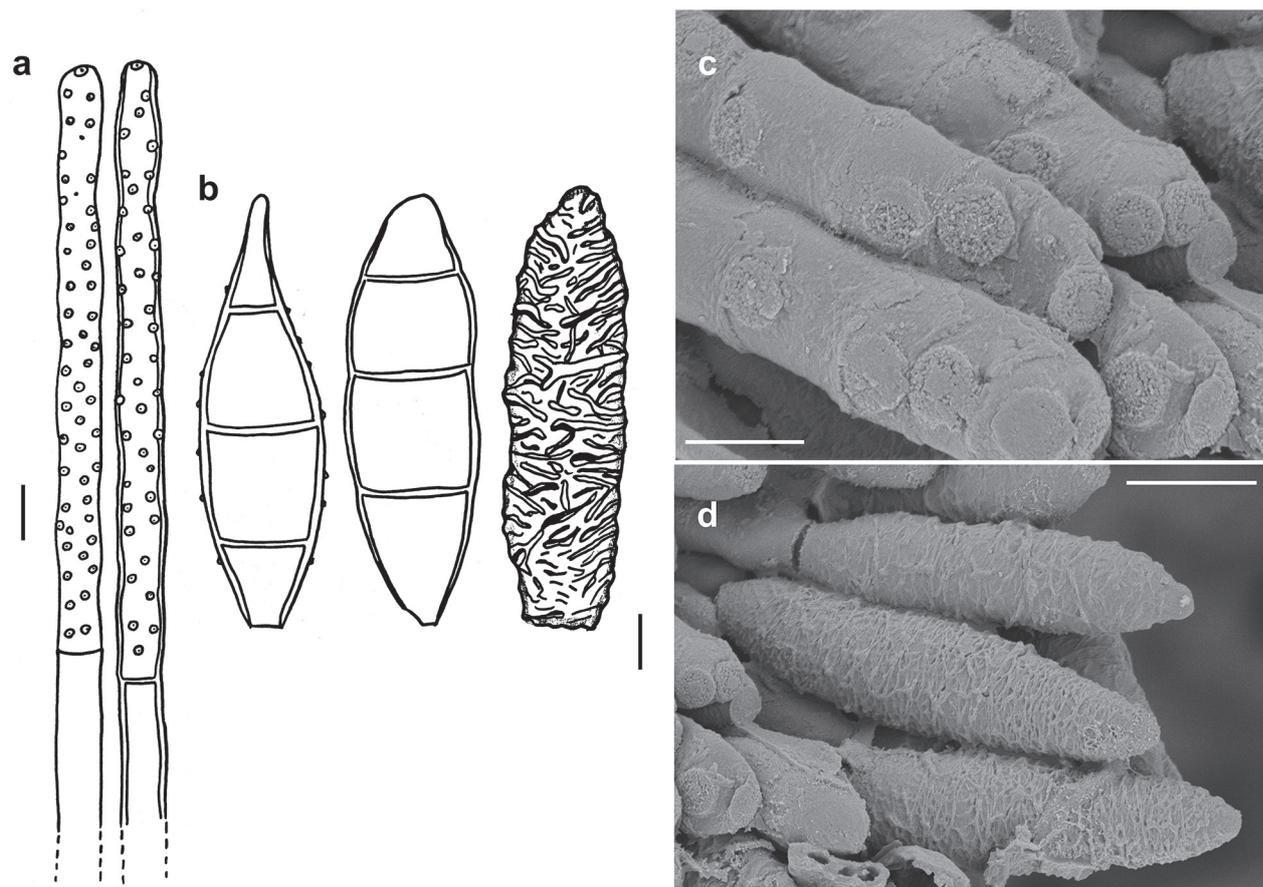
≡ *Arthrobotryum penicillium* (Speg.) F. Stevens, Bot. Gaz. 65: 238, 1918.

= *Arthrobotryum strychni* Henn., Hedwigia 43: 397, 1904.

≡ *Podosporium strychni* (Henn.) Cif., Sydowia 9: 311, 1955.

= *Arthrobotryum glabroides* F. Stevens, Bot. Gaz. 65: 237, 1918.

≡ *Podosporium glabroides* (F. Stevens) Cif., Sydowia 9: 309, 1955.



**Figure 20.** *Spiropes penicillium* (IMI 51664) **a** conidiophores with scars (the thickness of the wall is shown on the right-handed drawing) **b** conidia shown in optical section (first two left-hand drawings) and as seen by SEM **c, d** as seen by SEM **c** tips of conidiophores with scars **d** conidia. Scale bars: 5  $\mu\text{m}$  (**a**); 2.5  $\mu\text{m}$  (**b**); 3  $\mu\text{m}$  (**c**); 5  $\mu\text{m}$  (**d**).

**Description.** **Colonies** effuse, yellowish to dark olivaceous-brown, velvety, with tightly packed hyphae that form large, erect, dark synnemata clearly visible under the stereomicroscope. A bright yellow pigment diffuses out when colonies are mounted in lactic acid or lacto-phenol. **Hyphae** superficial, branched, septate, 1–2  $\mu\text{m}$  wide, yellowish, pale olive, smooth. **Conidiophores** tightly packed to form dark brown to blackish synnemata up to 650  $\mu\text{m}$  long, 10–40  $\mu\text{m}$  thick, often splaying out to a width of 100  $\mu\text{m}$  at the apex. Individual hyphae straight or flexuous, cylindrical, 1–2  $\mu\text{m}$  thick near the base, 2–3.5  $\mu\text{m}$  thick near the apex, pale olivaceous-brown, smooth, with numerous small conidial scars. **Conidia** solitary, fusiform or occasionally almost cylindrical, mostly 3(–5)–septate, 16–23(–37)  $\times$  (3–)3.5–5(–7)  $\mu\text{m}$ , tapering to about 1  $\mu\text{m}$  at the apex and base, middle cells pale brown, the cells at each end paler, surface wrinkled or verruculose. As seen by SEM, the ornamentation of the spores is distinctly reticulated, with thin to thick networks that can form ridges-like structures.

**Specimen examined.** On *Meliola calva* on leaves of Lauraceae, Brasil, S. Paulo, Apiahy, 1881, J. Puiggari 1483 (IMI 131184, type of *Podosporium penicillium*); on *Meliola* sp. on leaves of *Oxyanthus* sp. (Rubiaceae), Sierra Leone, 1951, D.S. Rennis (IMI 51664).

**Illustrations.** This species was illustrated by Ellis (1968).

**Known hosts and distribution.** On colonies of *Asteridiella* spp. and *Meliola* spp. on living leaves of various plants in Brazil, Congo, Costa Rica, Ghana, Ivory Coast, Nigeria, Sierra Leone and Uganda (Ellis 1968).

**Notes.** *Spiropes penicillium* is easily distinguishable from other known synnematosus species of the genus *Spiropes* by the presence of fusiform to cylindrical conidia without rostra. In addition, a bright yellow pigment diffuses out of the cells when colonies are mounted in lactic acid or lacto-phenol (Ellis 1968).

### Key to species of *Atractilina* and *Spiropes* hyperparasitic on Meliolales

- 1 Conidiophores synnematosus.....2
- Conidiophores single or in groups .....7
- 2 Synnemata straw-coloured to pale olivaceous; conidiophores with denticulate conidiogenous loci; pale multiseptate conidia.....**A. parasitica**
- Synnemata dark brown to black; conidiophores with cicatrised conidiogenous loci; conidia pigmented and multiseptate .....3
- 3 Synnemata up to 400 µm long; conidia mostly crescent shape .....**S. croissantiformis**
- Synnemata longer, from 700 µm to 1.5 mm long; conidia fusiform to obclavate, occasionally cylindrical .....4
- 4 Conidia fusiform to almost cylindrical; a yellow pigment diffuses out when colonies are mounted in lactic acid or lacto-phenol ..... **S. penicillium**
- Conidia fusiform to obclavate; no yellow pigment.....5
- 5 Conidia always 4–6 septate ..... **S. japonicus**
- Conidia always 3–septate .....6
- 6 Conidia 17–25 × 5–6.5 µm .....**S. clavatus**
- Conidia 40–52 × 9–11 µm ..... **S. melanoplaca**
- 7 Conidia with 3–6 pseudosepta.....8
- Conidia 1–3–septate .....10
- 8 Conidiophores in larger groups; conidia with 3–6 (usually 4 or 5) pseudosepta .....**S. capensis**
- Conidiophores single or in small groups; conidia with 3–5 pseudosepta ....9
- 9 Conidiophores with zigzag shape; conidia with 3–5 pseudosepta, fusiform to obclavate ..... **S. guareicola**
- Conidiophores without zigzag shape; conidia with 3–4 pseudosepta, obovate..... **S. fumosus**
- 10 Conidia 1–septate .....11
- Conidia 3–septate .....12
- 11 Conidia obpyriform, verrucose ..... **S. armatellae**
- Conidia obpyriform, smooth .....**S. armatellicola**
- 12 Conidia oblong-ellipsoid .....**S. intricatus**
- Conidia of various shapes, not oblong-ellipsoid .....13
- 13 Conidia obovate to clavate; conidiophores swollen towards the apex or in areas where conidia are produced ..... **S. deightonii**
- Conidia ovate or fusiform to obclavate; conidiophores not swollen towards the apex or in areas where conidia are produced .....14

- 14 Conidia obclavate; central cells barrel-shaped..... 15
- Conidia ovate or fusiform to obclavate; without central barrel-shaped cells ..... 16
- 15 Conidia with 3 true septa ..... ***S. caribensis***
- Conidia with 2 septa and a dark central pseudoseptum ..... ***S. palmetto***
- 16 Conidia ovate ..... ***S. carpolobiae***
- Conidia fusiform to obclavate ..... 17
- 17 Conidia 3–4.5 µm wide ..... ***S. effusus***
- Conidia wider ..... 18
- 18 Conidia 17–25 µm long ..... ***S. angylocalycis***
- Conidia longer ..... 19
- 19 Conidia 20–35 µm long ..... ***S. dorycarpus***
- Conidia longer ..... 20
- 20 Conidia 36–48 µm long ..... ***S. helleri***
- Conidia 40–54 µm long ..... ***S. leonensis***

In Fig. 21, we propose a visual key to the known species of *Spiropes* hyperparasitic on Melioidales.

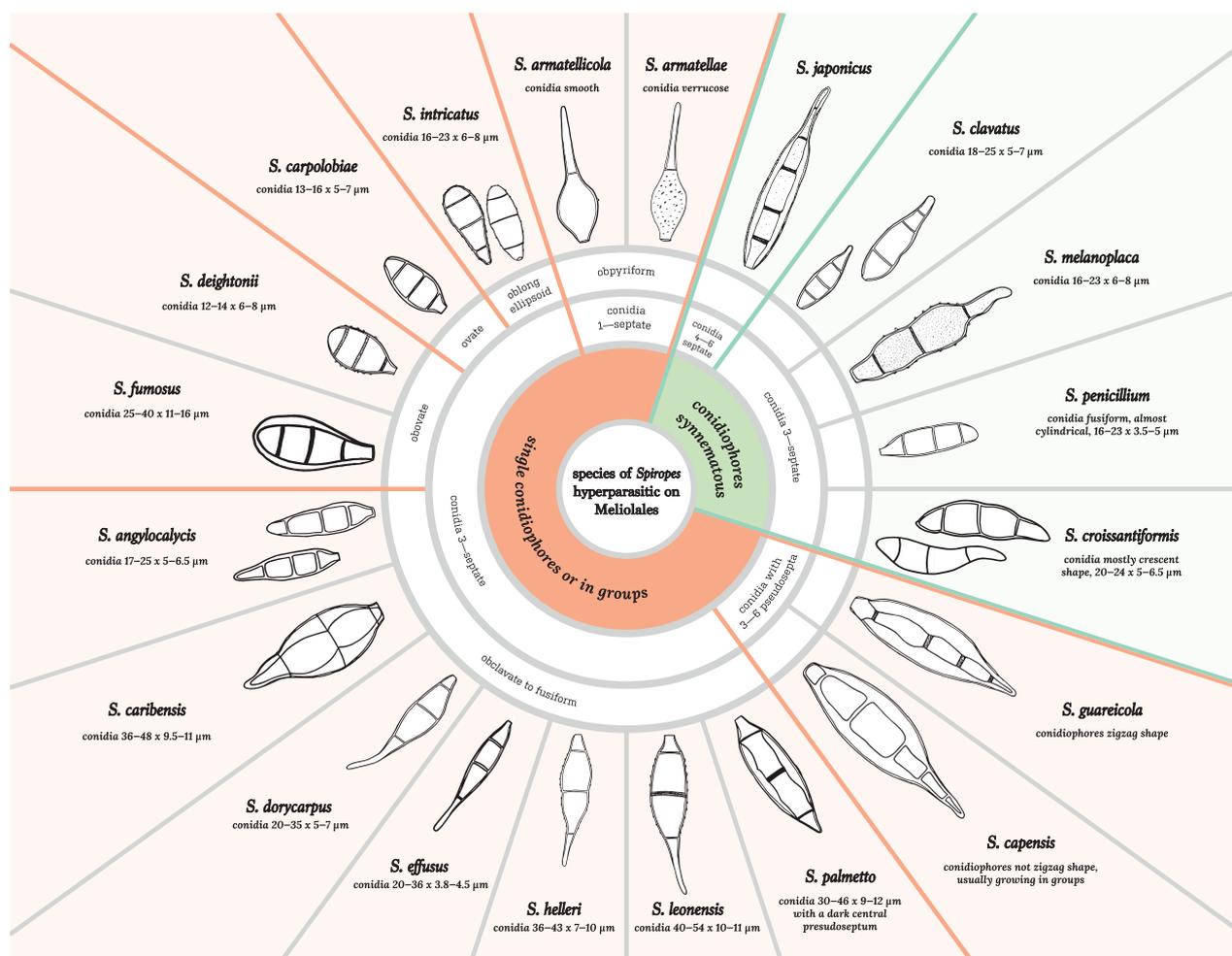
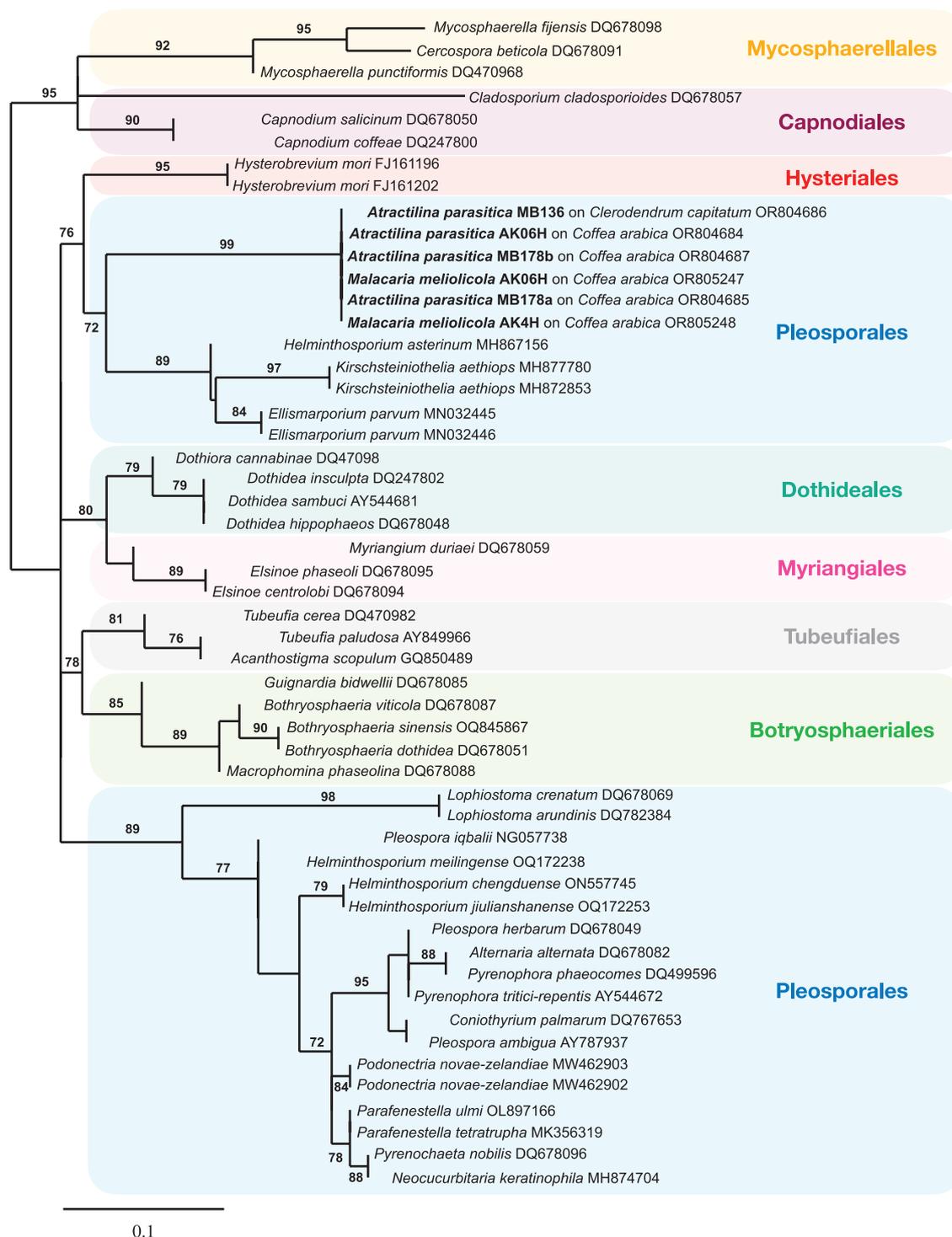


Figure 21. Visual key to known species of *Spiropes* hyperparasitic on Melioidales.



**Figure 22.** Phylogenetic tree inferred from a Maximum Likelihood analysis of nuc LSU rDNA sequences of members of the Dothideomycetes, including new sequences of *Atractilina parasitica* and *Malacaria meliolicola* (written with bold letters). The tree is rooted with sequences of species of the orders Capnodiales and Mycosphaerellales. Bootstrap values are indicated above the branches. Sequences downloaded from GenBank are given with accession numbers.

### Molecular position of species of *Atractilina* and *Spiropes*

In order to know the systematic positions of species of *Atractilina* and *Spiropes* hyperparasitic on Meliolales, new sequences of recently-collected specimens were obtained.

The BLAST query revealed that the nrLSU sequences of *Atractilina parasitica* (specimens MB136 and MB178) show approximately 82% similarity with sequences of species of the Dothideomycetes, such as *Botryosphaeria* spp., *Helminthosporium asterinum* Cooke, *Hysterobrevium mori* (Schwein.) E. Boehm & C.L. Schoch and *Neoheleiosa lincangensis* Mortimer, amongst others. In the tree inferred from the analysis of LSU sequences of 45 specimens of several orders of Dothideomycetes (Fig. 22), the sequences of *A. parasitica* are located in a well-supported clade that comprises species of Pleosporales, such as *Ellismarsporium parvum* R.F. Castañeda & W.B. Kendr., *Kirschsteiniothelia aethiops* (Sacc.) D. Hawksw. and *Helminthosporium asterinum*. In addition, the sequences of *A. parasitica* cluster together in a strongly-supported clade with two DNA sequences we obtained from *Malacaria meliolicola* (specimens AK4H and AK06H), a hyperparasitic perithecioid fungus that usually grows amongst the synnemata of *A. parasitica* on coffee leaves (see Bermúdez-Cova et al. (2023b) for the updated species description of *M. meliolicola*).

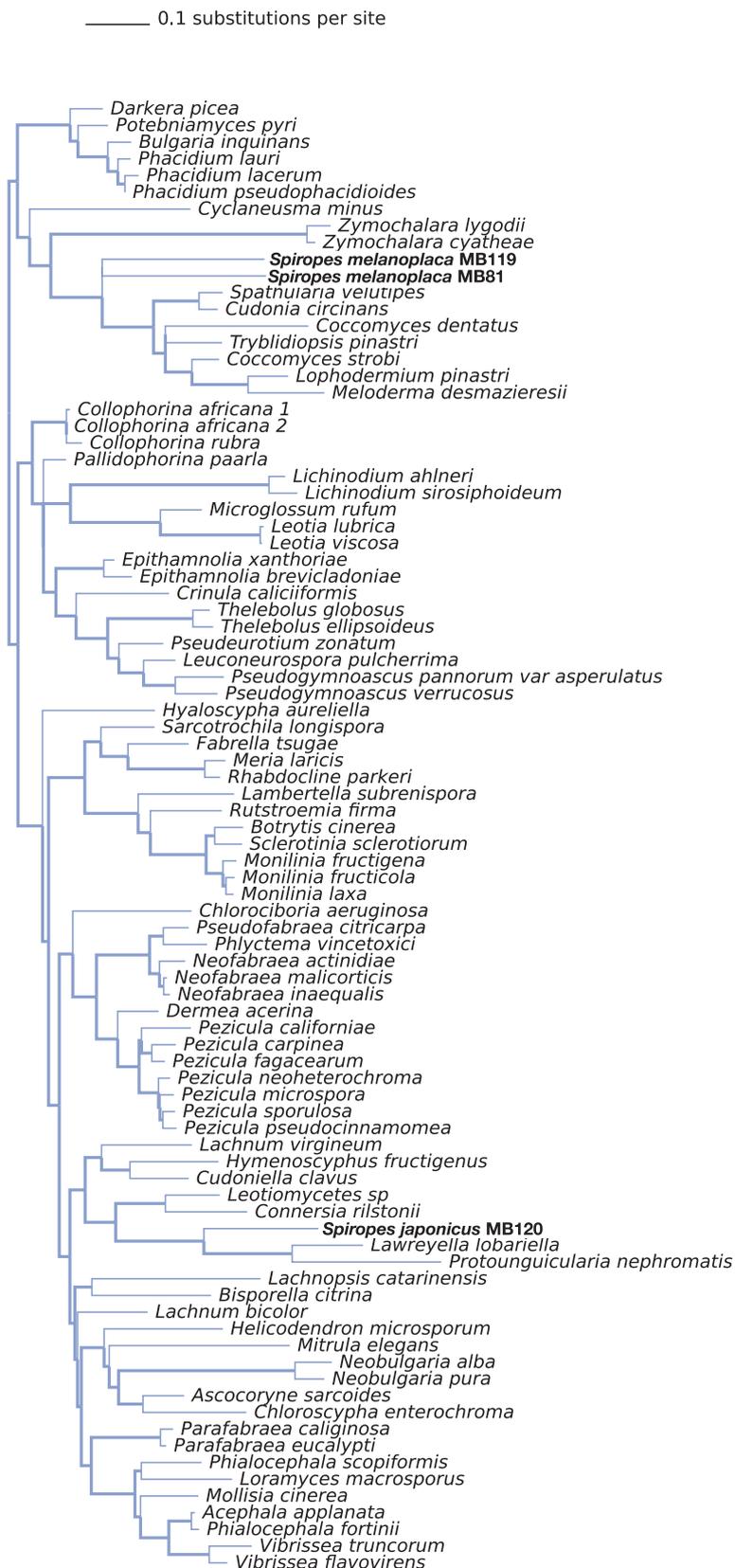
As for species of *Spiropes*, the BLAST query revealed that the nrITS sequences of *Spiropes melanoplaca* (specimens MB81 and MB119) and *Spiropes japonicus* (specimen MB 120) are not closely related to each other (60% similarity) and show between 88 and 90% similarity with species of the Leotiomycetes, such as *Lophodermium actinothyrium* Fuckel and *Hypoderma* spp., amongst others. Placement on to the Pezizomycotina tree version 2 in T-BAS confirmed that the newly-generated ITS sequences for the two species of *Spiropes* are placed in the Leotiomycetes (Fig. 23).

## Discussion

### ***Atractilina* and *Spiropes*, two genera with heterogeneous species**

Morphology-based identification of a species can be very difficult, especially amongst asexual or non-sporulating fungi (Jeewon et al. 2002; Promputtha et al. 2005, 2007). However, it continues to be an essential tool, especially for understudied groups of fungi and when DNA sequences are not available or scarce (Raja et al. 2017). The morphological analyses and the literature review of specimens of *Atractilina* and *Spiropes* revealed that both genera include highly heterogeneous species that are not necessarily congeneric with the type species of each genus.

The type species of *Atractilina*, *Atractilina callicarpae* Dearn. & Barthol. (= *Atractilina parasitica* (G. Winter) Deighton & Piroz.), has consistently true synnematos conidiophores, denticulate conidiogenous loci, pale pluriseptate (phragmoseptate) conidia and a hyperparasitic lifestyle (Deighton and Pirozynski 1972; Mel'nik and Braun 2013). Based on these characteristics, only three species of the genus are congeneric with *A. parasitica*, namely *A. alinae* Melnik & U. Braun, *A. biseptata* R.F. Castañeda and *A. calycini* T.K. Jana, S.N. Ghosh & A.K. Das (Castañeda-Ruiz 1986; Jana et al. 2006; Mel'nik and Braun 2013). The remaining two species present non-synnematos conidiophores and are probably not congeneric. *Atractilina asterinae* (Hansf.) Deighton & Piroz. is a species hyperparasitic on Asterinales and presents single conidiophores and distoseptate conidia (Deighton and Pirozynski 1972). *Atractilina hymenaeae* Bat. & J.L. Bezerra (introduced as *Atractina hymenaeae* by the authors) is hyperparasitic



**Figure 23.** Placement of *Spiropes japonicus* and *S. melanoplaca* on to Pezizomycotina reference tree version 2 in T-Bas. Only the Leotiomyces clade is shown. The tree is the result of RAxML analysis of nuc ITS rDNA with 500 bootstraps replicates. For each node, the Maximum Likelihood bootstrap ( $\geq 70\%$ ) is presented as thick branches. Names of *Spiropes* species with newly-generated sequence data are written in bold.

on Meliolales, but also with non-synnematous conidiophores and conidia with a variable number of septa (Batista and Bezerra 1961). Therefore, we believe that both species have been incorrectly assigned to the genus *Atractilina*.

The description of *A. parasitica* introduced by Deighton and Pirozynski (1972) is very broad. As a result, specimens with significant morphological variations are grouped into a single species concept. For example, Chen and Tzean (2007) described a parasitic fungus from Taiwan growing on decaying leaves of *Liquidambar* sp. (Altingiaceae), with conidia that resemble those of *A. parasitica*. However, conidiophores of this fungus are non-synnematous and very short (less than 15 µm long), a feature that has never been reported before for *A. parasitica*. It is necessary to re-evaluate this and other identifications, to narrow the species concept of *A. parasitica*, as well as to complement it with DNA sequence data.

The DNA molecular analyses of the nrLSU rDNA region of the specimens of *A. parasitica* from Benin revealed that this species belongs to the Dothideomycetes. The Dothideomycetes are the largest and most diverse class of fungi and comprise species that exhibit a broad range of lifestyles, including saprotrophs, plant pathogens, mycoparasites and hyperparasites, as well as lichenised and lichenicolous fungi (Pem et al. 2021). They typically produce flask-like structures called pseudothecia, though apothecial, hysterothecial and cleistothecoid ascomata also exist (Hessen and Jahns 1973; Valenzuela-Lopez et al. 2019). Bitunicate asci are one of the diagnostic characters for Dothideomycetes taxonomy (Von Arx and Müller 1975; Pem et al. 2021). Asexual stages are frequent amongst pathogenic genera in the families *Cladosporiaceae*, *Mycopsphaerellaceae*, *Pleosporaceae* and *Tubeufiaceae*, amongst others (Hyde et al. 2013; Wanasinghe et al. 2018; Hongsanan et al. 2020). Conidiophores in these anamorphic species are usually solitary or in groups forming synnemata (Thambugala et al. 2017). The sequences of *A. parasitica* showed 98% similarity with sequences of *Malacaria meliolicola* (Dothideomycetes, Ascomycota), a pseudothecoid hyperparasite that was found repeatedly amongst the synnemata of *A. parasitica* (Bermúdez-Cova et al. 2023b). The pseudothecia of *M. meliolicola* were also found to be growing without the presence of synnemata of *A. parasitica*. These colonies were used to extract the DNA of *M. meliolicola*. Therefore, the systematic position of *A. parasitica* in the Dothideomycetes and the anamorph-teleomorph connection between these two species are confirmed. This connection has been proposed in the past for these fungi on leaves of *Coffea arabica* (Hansford 1941, 1946; Bermúdez-Cova et al. 2023b). Here, a DNA sequence from a specimen of *A. parasitica* on *Meliola* sp. on leaves of *Clerodendrum capitatum* clustered with the aforementioned sequences in a highly-supported clade. The phylogenetic analysis of the nrLSU DNA locus showed that sequences of *A. parasitica* are located in a well-supported subclade together with other species of Pleosporales s.l., such as *Ellismarsporium parvum* (Zhang et al. 2020). Many species of the Dothideomycetes, especially the asexual genera, are known to be polyphyletic (Schoch et al. 2009). To confirm the systematic hypothesis and to determine the placement of *A. parasitica* at family level, the use of multi-loci phylogenies is necessary in the future.

As for the genus *Spiropes*, the generic diagnosis given by Ellis (1968, 1971) allows us to include in this genus all species with cicatrised conidiogenous cells and conspicuous, flat and numerous scars, as well as pigmented, mostly

obclavate phragmoconidia with 1–9 septa or pseudosepta. Seifert and Hughes (2000) proposed an amendment of this generic concept to also include species with dictyoconidia. As a result, *S. dictyosporus* is the only known species of the genus with muriform conidia. However, this morphological diagnosis allows for species with a wide range of types of conidiophores, conidiogenesis and conidia to be included in *Spiropes* (McTaggart et al. 2007). For example, the type species of the genus, *Spiropes guareicola* (F. Stevens) Cif., has distinctly sympodial-geniculate (zigzag-shaped) conidiophores, a character that is not present in any other known species of the genus (Ellis 1968). This species, in addition, presents distoseptate conidia, i.e. conidia with pseudosepta, a morphological feature that is present only in four species, namely *S. capensis*, *S. fumosus*, *S. guareicola* and *S. japonicus*. The remaining species of the genus present euseptate conidia (Ellis 1968, 1971). It is also possible to find a wide range of conidial shapes, such as obpyriform, obovate, ovate and oblong ellipsoid, to obclavate and fusiform (see the visual key to species of *Spiropes* in Fig. 21). Therefore, *Spiropes* is currently a genus with morphologically highly heterogeneous species and probably polyphyletic.

Identifying species of *Spiropes*, based on morphology alone, is not always easy. The most comprehensive key to species of the genus was proposed by Ellis (1968). However, this key is mainly based on the differences in the size range of the conidia of the species and, in some cases, these size differences are very subtle. Particular attention should be paid to herbarium specimens, as they may include immature or not well-preserved spores that can affect measurement results (Ordynets et al. 2021). We believe that other morphological characteristics that are not visible using standard light microscopy techniques should be considered when identifying species of *Spiropes* (e.g. Lutzoni et al. (2004)). Scanning electron Microscopy (SEM), for example, allowed us to observe for the first time the surface of the conidia of species of *Spiropes*. *Spiropes diallii* and *S. melanoplaca* were considered as different species by Ellis (1968). However, both species have overlapping spore-size ranges and the morphological analysis by SEM revealed that these species also have similar conidiogenesis and ornamentation patterns on conidia. This situation is similar for *S. intricatus* and *S. pirozynskii*. Therefore, we propose both groups of species as synonyms.

As for the molecular-based identification of species of *Spiropes*, there are currently no DNA sequences available in publicly-accessible databases. Species of the genus remain “incertae sedis” for many taxonomic ranks and it is difficult to assign new DNA sequences to species concepts (Bermúdez-Cova et al. 2022, 2023a). The DNA sequences generated for the first time in the context of this study suggest that species of *Spiropes* hyperparasitic on Meliolales may be polyphyletic in the Leotiomyces. Fungi in the class Leotiomyces are ecologically diverse and have been described as aquatic hyphomycetes, ectomycorrhizal parasites, endophytes, fungal parasites, mycorrhizal fungi, nematode-trapping fungi and plant-pathogens, amongst others (Wang et al. 2006a; Johnston et al. 2019). Many fungi have been suggested to belong to this class without any clear teleomorphic connection (Wang et al. 2006b). Up to date, no sexual stages have been linked to any species of *Spiropes* (Bermúdez-Cova et al. 2022). There is one genus with species morphologically similar to species of *Spiropes*, namely *Pseudospiropes* M.B. Ellis (Helotiales, Leotiomyces; Ellis (1971)). Species of this genus differ from species of *Spiropes* by broadly enlarged, thickened, protuberant, strongly

melanised conidiogenous loci and distoseptate conidia only (Castañeda-Ruiz et al. 2001; McTaggart et al. 2007). Species of *Pseudospiropes* have *Strossmayeria* Schulzer (Helotiales, Leotiomycetes) teleomorphs (Iturriaga and Korf 1984, 1990; Castañeda-Ruiz et al. 2001). Thus, there is a possibility that species of the genus *Spiropes* also belong to the Leotiomycetes. It is necessary to continue generating new DNA sequences from the different species of the genus in order to confirm this hypothesis, especially from those species that form part of mixed infections.

It is difficult to obtain molecular sequence data from hyperparasites especially because of the fact that they develop intermingled with the primary parasite and many other organisms and, as a result, no specific set of molecular methods has been developed to study hyperparasites (Bermúdez-Cova et al. 2022; Bermúdez-Cova et al. 2023a). As a consequence, isolating and sequencing hyperparasitic fungi is a challenging task. There is also a lack of sequences of hyperparasitic fungi in public. Therefore, the sequences obtained can be related to existing species concepts only based on morphology databases (Bermúdez-Cova et al. 2023b). For hyperparasitic fungi on Meliolales, for example, it is advised to obtain the same or very similar DNA sequences repeatedly from a given morphospecies in order to be sure to have the correct DNA sequence of that morphospecies. Despite many attempts, it was not possible to obtain DNA sequences from some of the species included in this study. However, this research provides valuable information that lays the foundation for future research on hyperparasites in Meliolales, highlighting the importance of field work paired with molecular for the study of challenging fungal groups. Further methodologies, such as metabarcoding, could represent another way to try to isolate the DNA of these organisms.

### **The need for re-evaluation, resampling and epitypification**

Applications of names based on morphological characteristics without DNA data is a challenge, resulting in the description of an excessive number of species or, in contrast, in the overlooking of cryptic species that can only be detected through molecular analyses (Hibbett et al. 2007; Crous et al. 2014; Jayasiri et al. 2015). The knowledge of morphological characteristics, however, is important to understand the evolution of fungal diversity (Raja et al. 2017). Instead of describing new species as part of *Atractilina* and *Spiropes*, a re-evaluation of the natural concepts of both genera is needed. Here we propose a list of actions that are necessary to carry out such a re-evaluation:

- Restudy the type species of each genus. When the type specimens of the type species are not in good condition or there is no more fungal material available for examination, it is necessary to recollect them. Epitypes and neotypes should be designated in these cases.
- After redefining the type species, all species belonging to the two genera need to be recollected, re-analysed morphologically and compared to the type species.
- The DNA of all existing species should be extracted, amplified and sequenced, in order to confirm or propose new concepts of genera and species. Multi-loci phylogenetic analyses are necessary to validate or propose new systematic hypotheses.

*Atractilina* and *Spiropes* are currently two repository genera of highly heterogeneous species and they may be split in the future, once species and genus concepts are validated respectively by morphology and molecular methods.

## Acknowledgements

We are grateful to the University of Parakou and the University of Abomey-Calavi, Benin, for the support and facilities made available for this study. We acknowledge help by Dr. Pierre Agbani (Botanical Garden of the Université d'Abomey-Calavi) for his assistance with the identification of host plants and help by Daouda Dongnima during fieldwork. Special thanks to Affoussatou Tabé, Alicia Sanjur and Anna Krauss for their support and collecting efforts throughout the whole research process. We acknowledge the support and facilities made available by Orlando Cáceres and the Universidad Autónoma de Chiriquí (UNACHI) in Panama. The Environmental Ministry of Panama (MiAmbiente) is thanked for issuing the collection and export permits (SE/APHO-1-2019, SEX/H-5-2020, PA-01-ARG-049-2021). We are grateful to the Ministry of Environment of the Benin Republic for issuing the collecting permits and for the elaboration of the ABS Nagoya Protocol documents n° 636/DGEFC/ANC-APA/DCPRN/PF-APA.

## Additional information

### Conflict of interest

The authors have declared that no competing interests exist.

### Ethical statement

No ethical statement was reported.

### Funding

The first author acknowledges support from the German Academic Exchange Service (DAAD), within the framework of the scholarship programme for doctoral studies in Germany (Ref. no.: 91726217).

### Author contributions

MB-C compiled and analyzed the data and wrote the first draft of the manuscript. MB-C, TH, NY and MP contributed to writing and editing the manuscript. All authors contributed to the article and approved the submitted version.

### Author ORCIDs

Miguel A. Bermúdez-Cova  <https://orcid.org/0000-0002-7712-7347>

Tina A. Hofmann  <https://orcid.org/0000-0003-1124-402X>

Nourou S. Yorou  <https://orcid.org/0000-0001-6997-811X>

Meike Piepenbring  <https://orcid.org/0000-0002-7043-5769>

### Data availability

All of the data that support the findings of this study are available in the main text or Supplementary Information.

## References

- Abarca GH, Ruiz RC, Arias RM, Saikawa M, Stadler M (2007) Anamorphic fungi from submerged plant material: *Acumisporea verruculosa*, *Pleurophragmium aquaticum* and *P. miniumbonatum*. *Mycotaxon* 101: 89–97.
- Akoègninou A, Van der Burg WJ, Van der Maesen LJG (2006) Flore Analytique du Bénin. Backhuys Publishers, Wageningen, 1034 pp.
- Alcorn JL (1988) The taxonomy of “*Helminthosporium*” species. *Annual Review of Phytopathology* 26(1): 37–56. <https://doi.org/10.1146/annurev.py.26.090188.000345>
- Bánki O, Roskov Y, Döring M, Ower G, Vandepitte L, Hobern D, Remsen D, Schalk P, DeWalt RE, Keping M, Miller J, Orrell T, Aalbu R, Abbott J, Adlard R, Adriaenssens EM, Aedo C, Aesch E, Akkari N, et al. (2023) Catalogue of Life Checklist [Version 2023-07-10]. Catalogue of Life. <https://doi.org/10.48580/dfry>
- Batista AC, Bezerra JL (1961) Novos ou raros Deuteromycetes. *Memórias da Sociedade Broteriana* 14: 73–82.
- Benson DA, Clark K, Karsch-Mizrachi I, Lipmanbn DJ, Ostell J, Sayers EW (2014) GenBank. *Nucleic Acids Research* 42(D1): D32–D37. <https://doi.org/10.1093/nar/gkt1030>
- Bermúdez-Cova MA, Cruz-Laufer AJ, Piepenbring M (2022) Hyperparasitic fungi on black mildews (Meliolales, Ascomycota): Hidden fungal diversity in the tropics. *Frontiers in Fungal Biology* 3: 885279. <https://doi.org/10.3389/ffunb.2022.885279>
- Bermúdez-Cova MA, Haelewaters D, de Bekker C, Piepenbring M, Schoutteten N, Quandt CA (2023a) Hyperparasitic fungi – definitions, diversity, ecology, and research. *Authorea*, June 27, 2023. <https://doi.org/10.22541/au.168787020.07281183/v1>
- Bermúdez-Cova MA, Krauß A, Sanjur A, Tabé A, Hofmann TA, Yorou NS, Piepenbring M (2023b) Diversity of hyperparasitic fungi on Meliolales (Sordariomycetes, Ascomycota): New species, records, and molecular data from Benin and Panama. *Mycological Progress* 22(9): 65. <https://doi.org/10.1007/s11557-023-01913-5>
- Carbone I, White JB, Miadlikowska J, Arnold AE, Miller MA, Magain N, U'Ren JM, Lutzoni F (2019) T-BAS version 2.1: Tree-Based Alignment Selector toolkit for evolutionary placement of DNA sequences and viewing alignments and specimen metadata on curated and custom trees. *Microbiology Resource Announcements* 8(29): e00328–e19. <https://doi.org/10.1128/MRA.00328-19>
- Castañeda-Ruiz RF (1986) *Fungi Cubenses*, 20 pp.
- Castañeda-Ruiz RF, Heredia G, Reyes M, Arias RM (2001) A revision of the genus *Pseudospiropes* and some new taxa. *Cryptogamie. Mycologie* 22(1): 3–18. [https://doi.org/10.1016/S0181-1584\(01\)01057-0](https://doi.org/10.1016/S0181-1584(01)01057-0)
- Chen J-L, Tzean S-S (2007) Hyphomycetes from Taiwan. *Akanthomyces* and allied species. *Fungal Science* 22(3, 4): 63–77.
- Ciferri R (1955) Observations on meliolicolous Hyphales from Santo Domingo. *Sydowia* 9(1–6): 296–335.
- Condit R, Pérez R, Daguerre N (2011) *Trees of Panama and Costa Rica*. Princeton University Press, Princeton NJ, 481 pp. <https://doi.org/10.1515/9781400836178>
- Costantin JN (1888) *Les Mucédinées Simples. Histoire, Classification, Culture et Rôle des Champignons Inférieurs dans les Maladies des Végétaux et des Animaux*. Klincksieck, Paris, i-viii, 210 pp. [190 figs]
- Crous PW, Giraldo A, Hawksworth DL, Rober V, Kirk PM, Guarro J, Robbertse B, Schoch CL, Damm U, Trakunyingcharoen T, Groenewald JZ (2014) The Genera of Fungi: Fixing the application of type species of generic names. *IMA Fungus* 5(1): 141–160. <https://doi.org/10.5598/imafungus.2014.05.01.14>

- De Beer ZW, Seifert KA, Wingfield MJ (2013) A nomenclator for ophiostomatoid genera and species in the Ophiostomatales and Microascales. *Biodiversity Series* 12: 245–322.
- Deighton FC, Pirozynski KA (1972) Microfungi. V. More hyperparasitic hyphomycetes. *Mycological Papers* 128: 110.
- Dubey R, Moonnambeth NA (2013) Hyperparasitism of *Isthmospora spinosa* Stevens and *Spiropes melanoplaca* (Berk. & Curtis) Ellis on *Meliola tylophorae* – *indicae* Hosag. parasitizing *Tylophora indica* (Burm. f.) Merrill from India- A New Record. *Journal on New Biological Reports* 2(1): 64–66.
- Ellis MB (1968) Dematiaceous Hyphomycetes. IX. *Spiropes* and *Pleurophragmium*. *Mycological Papers* 114: 1–44.
- Ellis MB (1971) Dematiaceous Hyphomycetes. X. *Mycological Papers* 125: 1–30. <https://doi.org/10.1079/9780851986180.0000>
- Ellis MB (1976) More Dematiaceous Hyphomycetes. Kew, Commonwealth Mycological Institute, 507 pp. <https://doi.org/10.1079/9780851983653.0000>
- Fazekas AJ, Kuzmina ML, Newmaster SG, Hollingsworth PM (2012) DNA barcoding methods for land plants. In: Kress WJ, Erickson DL (Eds) *DNA Barcodes. Methods in Molecular Biology*. Humana, Totowa, 223–252. [https://doi.org/10.1007/978-1-61779-591-6\\_11](https://doi.org/10.1007/978-1-61779-591-6_11)
- Hansford CG (1941) Contribution towards the fungus flora of Uganda. III. *Proceedings of the Linnean Society of London* 153: 4–52. <https://doi.org/10.1111/j.1095-8312.1941.tb01378.x>
- Hansford CG (1946) The foliicolous ascomycetes, their parasites and associated fungi. *Mycological Papers* 15.
- Hessen A, Jahns HM (1973) '1974' lichens. Eine Einführung in die Flechtenkunde. Thieme, Stuttgart.
- Hibbett DS, Binder M, Bischoff JF, Blackwell M, Cannon PF, Eriksson OE, Huhndorf S, James T, Kirk PM, Lücking R, Lumbsch HT, Lutzoni F, Matheny PB, McLaughlin DJ, Powell MJ, Redhead S, Schoch CL, Spatafora JW, Stalpers JA, Vilgalys R, Aime MC, Aptroot A, Bauer R, Begerow D, Benny GL, Castlebury LA, Crous PW, Dair YC, Gams W, Geisers DM (2007) A higher-level phylogenetic classification of the fungi. *Mycological Research* 111(5): 509–547. <https://doi.org/10.1016/j.mycres.2007.03.004>
- Hofmann TA, Kirschner R, Piepenbring M (2010) Phylogenetic relationships and new records of Asterinaceae (Dothideomycetes) from Panama. *Fungal Diversity* 43(1): 39–53. <https://doi.org/10.1007/s13225-010-0042-4>
- Holubová-Jechová V, Sierra AM (1984) Studies on hyphomycetes from Cuba II. Hyphomycetes from the Isla de la Juventud. *Česká Mykologie* 38: 96–120.
- Hongsanan S, Tian Q, Persoh D, Zeng X-Y, Hyde KD, Chomnunti P, Boonmee S, Bahkali AH, Wen T-C (2015) Meliolales. *Fungal Diversity* 74(1): 91–141. <https://doi.org/10.1007/s13225-015-0344-7>
- Hongsanan S, Hyde KD, Phookamsak R, Wanasinghe DN, McKenzie EHC, Sarma VV, Boonmee S, Lücking R, Bhat DJ, Liu NG, Tennakoon DS, Pem D, Karunarathna A, Jiang SH, Jones EBG, Phillips AJL, Manawasinghe IS, Tibpromma S, Jayasiri SC, Sandamali DS, Jayawardena RS, Wijayawardene NN, Ekanayaka AH, Jeewon R, Lu YZ, Dissanayake AJ, Zeng XY, Luo ZL, Tian Q, Phukhamsakda C, Thambugala KM, Dai DQ, Chethana KWT, Samarakoon MC, Ertz D, Bao DF, Doilom M, Liu JK (2020) Refined families of Dothideomycetes: Dothideomycetidae and Pleosporomycetidae. *Mycosphere: Journal of Fungal Biology* 11(1): 1553–2107. <https://doi.org/10.5943/mycosphere/11/1/13>
- Hosagoudar VB (2003) Armatellaceae, a new family segregated from the Meliolaceae. *Sydowia* 55: 162–167.

- Hosagoudar VB, Biju CK, Abraham TK, Agarwal DK (2002) *Spiropes armatelicola* sp. nov. from Kerala, India. *Journal of Economic and Taxonomic Botany* 26(3): 603–604.
- Hyde KD, Jones E, Liu JK, Ariyawansa H, Boehm E, Boonmee S, Braun U, Chomnunti P, Crous PW, Dai D-Q, Diederich P, Dissanayake A, Doilom M, Doveri F, Hongsanan S, Jayawardena R, Lawrey JD, Li Y-M, Liu Y-X, Lücking R, Monkai J, Muggia L, Nelsen MP, Pang K-L, Phookamsak R, Senanayake IC, Shearer CA, Suetrong S, Tanaka K, Thambugala KM, Wijayawardene NN, Wikee S, Wu H-X, Zhang Y, Aguirre-Hudson B, Alias SA, Aptroot A, Bahkali AH, Bezerra JL, Bhat DJ, Camporesi E, Chukeatirote E, Gueidan C, Hawksworth DL, Hirayama K, De Hoog S, Kang J-C, Knudsen K, Li W-J, Li X-H, Liu Z-Y, Mapook A, McKenzie EHC, Miller AN, Mortimer PE, Phillips AJL, Raja HA, Scheuer C, Schumm F, Taylor JE, Tian Q, Tibpromma S, Wanasinghe DN, Wang Y, Xu J-C, Yacharoen S, Yan J-Y, Zhang M (2013) Families of Dothideomycetes. *Fungal Diversity* 63(1): 1–313. <https://doi.org/10.1007/s13225-013-0263-4>
- Iturriaga T, Korf RP (1984) Studies in the genus *Strossmayeria* (Helotiales). 1. generic delimitation. 2. Two lost species. 3. Three excluded taxa. *Mycotaxon* 20: 169–178.
- Iturriaga T, Korf RP (1990) A monograph of the Discomycete genus *Strossmayeria* (Leotiaceae), with comments on its anamorph, *Pseudospiropes* (Dematiaceae). *Mycotaxon* 36: 383–454.
- Jana TK, Das AK, Ghosh SN (2006) Studies on foliicolous fungi – I. *Geobios* (Jodhpur) 33(1): 9–16.
- Jayasiri SC, Hyde KD, Ariyawansa HA, Bhat J, Buyck B, Cai L, Dai Y-C, Abd-Elsalam KA, Ertz D, Hidayat I, Jeewon R, Jones EBG, Bahkali AH, Karunarathna SC, Liu J-K, Luangsa-ard JJ, Lumbsch HT, Maharachchikumbura SSN, McKenzie EHC, Moncalvo J-M, Ghobad-Nejhad M, Nilsson H, Pang K-L, Pereira OL, Phillips AJL, Raspé O, Rollins AW, Romero AI, Etayo J, Selçuk F, Stephenson SL, Suetrong S, Taylor JE, Tsui CKM, Vizzini A, Abdel-Wahab MA, Wen T-C, Boonmee S, Dai DQ, Daranagama DA, Dissanayake AJ, Ekanayaka AH, Fryar SC, Hongsanan S, Jayawardena RS, Li W-J, Perera RH, Phookamsak R, de Silva NI, Thambugala KM, Tian Q, Wijayawardene NN, Zhao R-L, Zhao Q, Kang J-C, Promputtha I (2015) The Faces of Fungi database: Fungal names linked with morphology, phylogeny and human impacts. *Fungal Diversity* 74(1): 3–18. <https://doi.org/10.1007/s13225-015-0351-8>
- Jayawardena RS, Hyde KD, Chen YJ, Papp V, Palla B, Papp D, et al. (2020) One stop shop IV: taxonomic update with molecular phylogeny for important phytopathogenic genera: 76–100 (2020). *Fungal Diversity* 103: 87–218. <https://doi.org/10.1007/s13225-020-00460-8>
- Jeewon R, Liew ECY, Hyde KD (2002) Phylogenetic relationships of *Pestalotiopsis* and allied genera inferred from ribosomal DNA sequences and morphological characters. *Molecular Phylogenetics and Evolution* 25(3): 378–392. [https://doi.org/10.1016/S1055-7903\(02\)00422-0](https://doi.org/10.1016/S1055-7903(02)00422-0)
- Johnston PR, Quijada L, Smith CA, Baral HO, Hosoya T, Baschien C, Pärtel K, Zhuang WY, Haelewaters D, Park D, Carl S, López-Giráldez F, Wang Z, Townsend JP (2019) A multigene phylogeny toward a new phylogenetic classification of Leotiomyces. *IMA Fungus* 10(1): 1. <https://doi.org/10.1186/s43008-019-0002-x>
- Kearse M, Moir R, Wilson A, Stones-Havas S, Cheung M, Sturrock S, Buxton S, Cooper A, Markowitz S, Duran C, Thierer T, Ashton B, Meintjes P, Drummond A (2012) Geneious basic: An integrated and extendable desktop software platform for the organization and analysis of sequence data. *Bioinformatics* (Oxford, England) 28(12): 1647–1649. <https://doi.org/10.1093/bioinformatics/bts199>
- Kress WJ, Erickson DL, Jones FA, Swenson NG, Perez R, Sanjurjo O, Bermingham E (2009) Plant DNA barcodes and a community phylogeny of a tropical forest dynamics plot in

- Panama. Proceedings of the National Academy of Sciences of the United States of America 106(44): 8621–18626. <https://doi.org/10.1073/pnas.0909820106>
- Kumar S, Stecher G, Tamura K (2016) MEGA7: Molecular evolutionary genetics analysis version 7.0 for bigger datasets. Molecular Biology and Evolution 33(7): 1870–1874. <https://doi.org/10.1093/molbev/msw054>
- Levin RA, Wagner WL, Hoch PC, Nepokroeff M, Pires JC, Zimmer EA, Sytsma KJ (2003) Family-level relationships of Onagraceae based on chloroplast rbcL and ndhF data. American Journal of Botany 90(1): 107–115. <https://doi.org/10.3732/ajb.90.1.107>
- Lutzoni F, Kauff F, Cox CJ, McLaughlin D, Celio G, Dentinger B, Padamsee M, Hibbett D, James TY, Baloch E, Grube M, Reeb V, Hofstetter V, Schoch C, Arnold AE, Miadlikowska J, Spatafora J, Johnson D, Hambleton S, Crockett M, Shoemaker R, Sung GH, Lücking R, Lumbsch T, O'Donnell K, Binder M, Diederich P, Ertz D, Gueidan C, Hansen K, Harris RC, Hosaka K, Lim YW, Matheny B, Nishida H, Pfister D, Rogers J, Rossman A, Schmitt I, Sipman H, Stone J, Sugiyama J, Yahr R, Vilgalys R (2004) Assembling the fungal tree of life: Progress, classification, and evolution of subcellular traits. American Journal of Botany 91(10): 1446–1480. <https://doi.org/10.3732/ajb.91.10.1446>
- McTaggart AR, Shivas RG, Braun U (2007) *Annelosympodia orbiculata* gen. et sp. nov. and *Scolecostigmia flagellariae* sp. nov. from Australia. Australasian Plant Pathology 36(6): 573–579. <https://doi.org/10.1071/AP07061>
- Mel'nik VA, Braun U (2013) *Atractilina alinae* sp. nov. and *Neosporidesmium vietnamense* sp. nov. – two synnematosous hyphomycetes from Vietnam. Mycobiota 3: 1–9. <https://doi.org/10.12664/mycobiota.2013.03.01>
- Miller MA, Pfeiffer W, Schwartz T (2010) Creating the CIPRES science gateway for inference of large phylogenetic trees. 2010 Gateway Computing Environments Workshop (GCE). IEEE, New Orleans, 8. <https://doi.org/10.1109/GCE.2010.5676129>
- Morelet M (1971) De aliquibus in mycologia novitatibus (5<sup>e</sup> note). Bulletin de la Société de Sciences Naturelles et d'Archéologie de Toulon et du Var 195: 7–8. <https://doi.org/10.2113/gssgfbull.S7-XIII.1-2.195>
- Nakamura T, Yamada KD, Tomii K, Katoh K (2018) Parallelization of MAFFT for large-scale multiple sequence alignments. Bioinformatics (Oxford, England) 34(14): 2490–2492. <https://doi.org/10.1093/bioinformatics/bty121>
- Ordynets A, Keßler S, Langer E (2021) Geometric morphometric analysis of spore shapes improves identification of fungi. PLOS ONE 16(8): e0250477. <https://doi.org/10.1371/journal.pone.0250477>
- Pem D, Jeewon R, Chethana KWT, Hongsanan S, Doilom M, Suwannarach N, Hyde KD (2021) Species concepts of Dothideomycetes: Classification, phylogenetic inconsistencies and taxonomic standardization. Fungal Diversity 109(1): 283–319. <https://doi.org/10.1007/s13225-021-00485-7>
- Piepenbring M, Hofmann TA, Kirschner R, Mangelsdorff R, Perdomo O, Rodríguez Justavino D, Trampe T (2011) Diversity patterns of Neotropical plant parasitic microfungi. Ecotropica (Bonn) 17: 27–40.
- Promptuttha I, Jeewon R, Lumyong S, McKenzie EHC, Hyde KD (2005) Ribosomal DNA fingerprinting in the identification of non-sporulating endophytes from *Magnolia liliifera* (Magnoliaceae). Fungal Diversity 20: 167–186.
- Promptuttha I, Lumyong S, Vijaykrishna D, McKenzie EHC, Hyde KD, Jeewon R (2007) A phylogenetic evaluation of whether endophytes become saprotrophs at host senescence. Microbial Ecology 53(4): 579–590. <https://doi.org/10.1007/s00248-006-9117-x>

- Raja HA, Miller AN, Pearce CJ, Oberlies NH (2017) Fungal Identification Using Molecular Tools: A Primer for the Natural Products Research Community. *Journal of Natural Products* 24 80(3): 756–770. <https://doi.org/10.1021/acs.jnatprod.6b01085>
- Rodríguez Justavino D, Kirschner R, Piepenbring M (2015) New species and new records of Meliolaceae from Panama. *Fungal Diversity* 70(1): 73–84. <https://doi.org/10.1007/s13225-014-0292-7>
- Ronquist F, Teslenko M, van der Mark P, Ayres DL, Darling A, Höhna S, Larget B, Liu L, Suchard MA, Huelsenbeck JP (2012) MrBayes 3.2: Efficient bayesian phylogenetic inference and model choice across a large model space. *Systematic Biology* 61(3): 539–542. <https://doi.org/10.1093/sysbio/sys029>
- Schoch CL, Crous PW, Groenewald JZ, Boehm EWA, Burgess TI, de Gruyter J, de Hoog GS, Dixon LJ, Grube M, Gueidan C, Harada Y, Hatakeyama S, Hirayama K, Hosoya T, Huhndorf SM, Hyde KD, Jones EBG, Kohlmeyer J, Kruys Å, Li YM, Lücking R, Lumbsch HT, Marvanová L, Mbatchou JS, McVay AH, Miller AN, Mugambi GK, Muggia L, Nelsen MP, Nelson P, Owensby CA, Phillips AJL, Phongpaichit S, Pointing SB, Pujade-Renaud V, Raja HA, Plata E Rivas, Robbertse B, Ruibal C, Sakayaroj J, Sano T, Selbmann L, Shearer CA, Shirouzu T, Slippers B, Suetrong S, Tanaka K, Volkmann-Kohlmeyer B, Wingfield MJ, Wood AR, Woudenberg JHC, Yonezawa H, Zhang Y, Spatafora JW (2009) A class-wide phylogenetic assessment of Dothideomycetes. *Studies in Mycology* 64: 1–15. <https://doi.org/10.3114/sim.2009.64.01>
- Seifert KA, Hughes SJ (2000) *Spiropes dictyosporus*, a new synnematosus fungus associated with sooty moulds. *New Zealand Journal of Botany* 38(3): 489–492. <https://doi.org/10.1080/0028825X.2000.9512698>
- Stamatakis A (2014) RAxML version 8: A tool for phylogenetic analysis and post-analysis of large phylogenies. *Bioinformatics (Oxford, England)* 30(9): 1312–1313. <https://doi.org/10.1093/bioinformatics/btu033>
- Stevens FL (1918) Some meliolicolous parasites and commensals from Porto Rico. *Botanical Gazette (Chicago, Ill.)* 65(3): 227–249. <https://doi.org/10.1086/332230>
- Talavera G, Castresana J (2007) Improvement of phylogenies after removing divergent and ambiguously aligned blocks from protein sequence alignments. *Systematic Biology* 56(4): 564–577. <https://doi.org/10.1080/10635150701472164>
- Thambugala KM, Wanasinghe DN, Phillips AJL, Camporesi E, Bulgakov TS, Phukhamsakda C, Ariyawansa HA, Goonasekara ID, Phookamsak R, Dissanayake A, Tennakoon DS, Tibpromma S, Chen YY, Liu ZY, Hyde KD (2017) Mycosphere notes 1–50: Grass (Poaceae) inhabiting Dothideomycetes. *Mycosphere : Journal of Fungal Biology* 8(4): 697–796. <https://doi.org/10.5943/mycosphere/8/4/13>
- Valenzuela-Lopez N, Magaña-Dueñas V, Cano-Lira JF, Wiederhold N, Guarro J, Stchigel AM (2019) Two new species of *Gloniopsis* (Hysteriales, Ascomycota) from clinical specimens: Morphological and molecular characterization. *Mycoses* 62(12): 1164–1173. <https://doi.org/10.1111/myc.13006>
- Vilgalys R, Hester M (1990) Rapid genetic identification and mapping of enzymatically amplified ribosomal DNA from several *Cryptococcus* species. *Journal of Bacteriology* 172(8): 4238–4246. <https://doi.org/10.1128/jb.172.8.4238-4246.1990>
- Von Arx JA, Müller E (1975) A re-evaluation of the bitunicate ascomycetes with keys to families and genera. *Studies in Mycology* 9: 1–159.
- Wagner T, Ryvarden L (2002) Phylogeny and taxonomy of the genus *Phylloporia* (Hymenochaetales). *Mycological Progress* 1(1): 105–116. <https://doi.org/10.1007/s11557-006-0009-8>

- Wanasinghe DN, Phukhamsakda C, Hyde KD, Jeewon R, Lee HB, Jones EBG, Tibpromma S, Tennakoon DS, Dissanayake AJ, Jayasiri SC, Gafforov Y, Camporesi E, Bulgakov TS, Ekanayake AH, Perera RH, Samarakoon MC, Goonasekara ID, Mapook A, Li W-J, Senanayake IC, Li J, Norphanphoun C, Doilom M, Bahkali AH, Xu J, Mortimer PE, Tibell L, Tibell S, Karunarathna SC (2018) Fungal diversity notes 709–839: Taxonomic and phylogenetic contributions to fungal taxa with an emphasis on fungi on Rosaceae. *Fungal Diversity* 89(1): 1–236. <https://doi.org/10.1007/s13225-018-0395-7>
- Wang Z, Johnston PR, Takamatsu S, Spatafora JW, Hibbett DS (2006a) Towards a phylogenetic classification of Leotiomyces based on rDNA data. *Mycologia* 98(6): 1065–1075. <https://doi.org/10.1080/15572536.2006.11832634>
- Wang Z, Binder M, Schoch CL, Johnston PR, Spatafora JW, Hibbett DS (2006b) Evolution of helotialean fungi (Leotiomyces, Pezizomycotina): A nuclear rDNA phylogeny. *Molecular Phylogenetics and Evolution* 41(2): 295–312. <https://doi.org/10.1016/j.ympev.2006.05.031>
- White TJ, Brun T, Lee S, Taylor J (1990) Amplification and direct sequencing of fungal ribosomal RNA genes for phylogenetics. In: Innis MA, Gelfand DH, Sninsky JJ, White TJ (Eds) *PCR Protocols: a guide to methods and applications*, Academic Press, San Diego. <https://doi.org/10.1016/B978-0-12-372180-8.50042-1>
- Zeng XY, Zhao JJ, Hongsanan S, Chomnunti P, Boonmee S, Wen TC (2017) A checklist for identifying Meliolales species. *Mycosphere: Journal of Fungal Biology* 8(1): 218–359. <https://doi.org/10.5943/mycosphere/8/1/16>
- Zhang K, Guo W, Sosa D, Magdama F, Serrano L, Malosso E, Li D-W, Castañeda-Ruiz RF (2020) Phylogeny and morphology of *Ellismarsporium parvum* and the new combination *E. varium*. *Mycotaxon* 135(2): 443–452. <https://doi.org/10.5248/135.443>
- Zhao G, Wu Y, Li N (1996) Six fungi species of hyperparasite on Meliolaceae. *Journal of Beijing Forestry University* 18(1): 99–103.

## Supplementary material 1

### Alignments and tree generated during the analysis of the DNA sequences of *Atractilina parasitica*, *Malacaria meliolicola* and other members of the Dothideomycetes

Authors: Miguel A. Bermúdez-Cova, Tina A. Hofmann, Nourou S. Yorou, Meike Piepenbring  
Data type: docx

Explanation note: Alignment is shown in NEXUS format. The tree is shown in Newick format.

Copyright notice: This dataset is made available under the Open Database License (<http://opendatacommons.org/licenses/odbl/1.0/>). The Open Database License (ODbL) is a license agreement intended to allow users to freely share, modify, and use this Dataset while maintaining this same freedom for others, provided that the original source and author(s) are credited.

Link: <https://doi.org/10.3897/mycokeys.103.115799.suppl1>