



New asexual morph taxa in *Phaeosphaeriaceae*

Li WJ^{1,2,3,4}, Bhat DJ⁵, Camporesi E⁶, Tian Q^{3,4}, Wijayawardene NN^{3,4}, Dai DQ^{3,4}, Phookamsak R^{3,4}, Chomnunti P^{3,4}, Bahkali AH⁷ & Hyde KD^{1,2,3,7*}

¹World Agroforestry Centre, East and Central Asia, 132 Lanhei Road, Kunming 650201, China

²Key Laboratory of Economic Plants and Biotechnology, Kunming Institute of Botany, Chinese Academy of Sciences, Lanhei Road No 132, Panlong District, Kunming, Yunnan Province, 650201, PR China

³Center of Excellence in Fungal Research, Mae Fah Luang University, Chiang Rai 57100, Thailand

⁴School of Science, Mae Fah Luang University, Chiang Rai 57100, Thailand

⁵Formerly, Department of Botany, Goa University, Goa 403206, India

⁶A.M.B. Gruppo Micologico Forlivese “Antonio Cicognani”, Via Roma 18, Forlì, Italy

⁷Botany and Microbiology Department, College of Science, King Saud University, Riyadh, KSA 11442, Saudi Arabia

Li WJ, Bhat DJ, Camporesi E, Tian Q, Wijayawardene NN, Dai DQ³, Phookamsak R, Chomnunti P, Bahkali AH, Hyde KD 2015 – New asexual morph taxa in *Phaeosphaeriaceae*. *Mycosphere* 6(6), 681–708, Doi 10.5943/mycosphere/6/6/5

Abstract

Species of *Phaeosphaeriaceae*, especially the asexual taxa, are common plant pathogens that infect many important economic crops. Ten new asexual taxa (*Phaeosphaeriaceae*) were collected from terrestrial habitats in Italy and are introduced in this paper. In order to establish the phylogenetic placement of these taxa within *Phaeosphaeriaceae* we analyzed combined ITS and LSU sequence data from the new taxa, together with those from GenBank. Based on morphology and molecular data, *Poaceicola* gen. nov. is introduced to accommodate the new species *Po. arundinis* (type species), *Po. bromi* and *Po. elongata*. The new species *Parastagonospora dactylidis*, *P. minima*, *P. italica*, *P. uniseptata* and *P. allouniseptata*, *Septoriella allojunci* and *Wojnowicia spartii* are also introduced with illustrated accounts and compared with similar taxa. We also describe an asexual morph of a *Nodulosphaeria* species for the first time.

Key words – Asexual morphs – *Phaeosphaeriaceae* – Phylogeny – Taxonomy

Introduction

The order Pleosporales has been of great research interest in the recent years and has undergone considerable revision (Zhang et al. 2009, 2012, Hyde et al. 2013, Ariyawansa et al. 2014, 2015b). Hyde et al. (2013) provided a taxonomic account for 41 families in Pleosporales, in which *Phaeosphaeriaceae* is a large and important family, comprising more than 300 species, and has also undergone considerable revision (Hyde et al. 2013, Phookamsak et al. 2014, Liu et al. 2015). Species in *Phaeosphaeriaceae* can be saprobic or endophytic, but many are pathogenic on economically

important plants and crops (Kirk et al. 2008, Zhang et al. 2009, Quaedvlieg et al. 2013, Hyde et al. 2013, Wijayawardene et al. 2014, Phookamsak et al. 2014). For example, *Parastagonospora* species are pathogenic on *Poaceae* and are directly or indirectly responsible for significant crop losses in wheat, barley and rye worldwide (Quaedvlieg et al. 2013). In addition, many genera, such as, *Phaeosphaeriopsis*, *Setophoma*, *Xenoseptoria* and *Wojnowicia* have been found associated with leaf spot disease of various hosts (Arzanlou and Crous 2006, Quaedvlieg et al. 2013; Wijayawardene et al. 2013; Phookamsak et al. 2014).

Barr (1979a) introduced *Phaeosphaeriaceae* based on the type genus *Phaeosphaeria*, and the history has been reviewed by Phookamsak et al. (2014). The latter authors also carried out a revision of *Phaeosphaeriaceae*, based on multi-gene (ITS, LSU, SSU, RPB2 and TEF1) analyses coupled with morphological data. They included 30 genera in the family, in which 17 genera are asexual morphs, Liu et al. (2015) and Ariyawansa et al. (2015c) added four additional genera in *Phaeosphaeriaceae*, and to date, 37 genera are accommodated in the family (Table 1).

In the present study we introduce ten new asexual taxa in *Phaeosphaeriaceae* using combined ITS and LSU sequence data, as well as morphological data.

Material & methods

Collection and examination of specimens

Fresh specimens were collected from terrestrial habitats in Italy. Samples were examined and pure cultures obtained by single spore isolation following the method described in Chomnunti *et al.* (2014). The growing colonies were transferred to 2 % potato-dextrose agar (PDA) and incubated at 25°C. The colony characters and growth rates were determined after one to four weeks. The pure cultures from our study are deposited at Mae Fah Luang University Culture Collection (MFLUCC). Duplicate cultures are deposited in the International Collection of Microorganisms from Plants, Landcare Research, New Zealand (ICMP) and Kunming Institute of Botany Culture Collection (KUMCC). The holotype is deposited at the herbarium of Mae Fah Luang University (MFLU), Chiang Rai, Thailand, and the isotype specimens are deposited at the herbarium of Kunming institute of Botany Chinese Academy of Sciences (KUN). Faces of Fungi numbers and Index Fungorum numbers were obtained as explained in Jayasiri *et al.* (2015). The asexual morphs were established in culture using the method of Phookamsak *et al.* (2015).

DNA extraction, PCR amplification and sequencing

Isolates were grown on PDA plates in the darkness at 25°C until completely covering the agar surface. The mycelium (about 50 mg) was scraped off and collected in a 1.5 ml micro centrifuge tube. Genomic DNA was extracted from fresh mycelium, following the specification of Biospin Fungus Genomic DNA Extraction Kit (BioFlux®). For *Poaceicola arundinis* sp. nov. and *Septoriella allojunci* sp. nov., despite several attempts we could not isolate them into culture. Therefore, DNA was extracted directly from the conidiomata of the dried specimens following the method of Li *et al.* (2015). The primer pairs LROR and LR5 as defined by Vilgalys and Hester (1990) were used to amplify a segment of the large subunit rDNA. Primer pairs ITS4 and ITS5 as defined by White *et al.* (1990) were used to amplify the internal transcribed spacers (ITS). DNA amplification was performed by polymerase chain reaction (PCR). The sequencing of PCR products were carried at Shang Hai Biological Engineering Technology Co. Ltd (Shang Hai, P. R. China).

DNA sequence data analysis

Sequences were obtained from GenBank mostly following previous literature (Phookamsak *et al.* 2014, Ariyawansa *et al.* 2014, 2015c, Crous *et al.* 2015b, Liu *et al.* 2015) and are listed in Table 2.

Multiple sequences were aligned using Bioedit v. 7.0.9 (Hall, 1999) and Clustal X v. 1.83 (Thompson *et al.*, 1997). The alignments were checked visually and improved manually wherever necessary. A maximum likelihood (ML) analysis was performed with raxmlGUI version 1.3 (Silvestro & Michalak 2011). The optimal ML tree search was conducted with 1000 separate runs, using the default algorithm of the program from a random starting tree for each run. The final tree was selected among suboptimal trees from each run by comparing likelihood scores under the GTRGAMMA substitution model. The resulting trees were printed with TreeView v. 1.6.6 (Page, 1996).

Results

Phylogenetic analyses

ITS and LSU sequence data were used to resolve the generic placement of the collections of *Phaeosphaeriaceae*. The alignment datasets included 83 taxa of which *Didymella exigua* was used as the outgroup taxon. The combined datasets comprised 1435 characters including gaps. The best scoring RAxML tree was chosen as the backbone tree and is shown in Figure 1.

The phylogenetic analyses show that *Poaceicola elongata* comb. nov. and two other species, *Po. arundinis* and *Po. bromi*, are represented as a lineage distinct from any other genera in *Phaeosphaeriaceae* (Fig. 1). The other strains are distributed in *Nodulosphaeria* (one species), *Parastagonospora* (five species), and *Septoriella* (one species). Based on the morphology and sequence data, these fresh collections are introduced as new asexual taxa in *Phaeosphaeriaceae*. The new taxa are five species of *Parastagonospora*, i.e. *P. allouniseptata*, *P. dactylidis*, *P. minima*, *P. talica* and *P. uniseptata*; *Septoriella allojunci* sp. nov., and an asexual morph of *Nodulosphaeria*, *N. digitalis*.

Taxonomy

Nodulosphaeria digitalis W.J. Li, Camporesi, D.J. Bhat & K.D. Hyde, **sp. nov.**

Fig. 2

Index Fungorum number: IF551664

Facesoffungi number: FoF 01302,

Etymology – Named after the host genus, *Digitalis*.

Holotype – MFLU 15–2716

Saprobic on dead stems of *Digitalis* sp., forming conspicuous, rounded, black fruiting bodies.

Sexual morph – Undetermined. **Asexual morph** – Coelomycetous. *Conidiomata* 130–150 µm high, 240–270 µm diam., pycnidoid, brown, solitary, aggregated or confluent, semi-immersed, subglobose, unilocular, papillate. *Ostiole* centrally located, short. *Wall of conidiomata* 20–60 µm wide, composed thick-walled, brown outer cells of *textura angularis*, and an inner layers of hyaline *textura prismatica*. *Conidiophores* cylindrical, short, hyaline, arising from the inner layer of conidioma. *Conidiogenous cells* 7–15 × 2–4 µm, entoblastic, phialidic, cylindrical to subcylindrical, hyaline, smooth-walled. *Conidia* 9–18 × 2–9 µm (\bar{x} = 15 × 3 µm; n = 30), hyaline, falcate, fusiform, strongly curved towards the apices, rounded at both ends, 0–1-septate, smooth and thick-walled.

Material examined – ITALY, Province of Arezzo [AR], Bagno di Cetica, on dead stem of *Dactylis* sp., 1 October 2012, Erio Camporesi, IT-753 (MFLU 15–2716); *ibid.* (KUN! HKAS 90731).

Notes – *Nodulosphaeria digitalis* clustered with *Nodulosphaeria* species with high bootstrap support (99%), but *N. digitalis* formed a separate branch basal to *N. modesta* (Rabenh.) Munk ex L. Holm) and *N. italica* (Fig. 1). This is the first recorded asexual morph for *Nodulosphaeria*.

Table 1 Genera of *Phaeosphaeriaceae*, types species and morphs

Genus name	Types species	Sexual/Asexual morph	Reference
<i>Allophaeosphaeria</i>	<i>Allophaeosphaeria muriformia</i>	Asexual morph: Coelomyctous	Liu et al. 2015
<i>Amarenographium</i>	<i>Amarenographium metableticum</i>	Sexual morph unknown	Phookamsak et al. 2014
<i>Amarenomyces</i>	<i>Amarenomyces ammophilae</i>	Asexual morph unknown	Phookamsak et al. 2014
<i>Ampelomyces</i>	<i>Ampelomyces quisqualis</i>	Sexual morph unknown	Phookamsak et al. 2014
<i>Bricookea</i>	<i>Bricookea sepalorum</i>	Asexual morph unknown	Phookamsak et al. 2014
<i>Chaetosphaeronema</i>	<i>Chaetosphaeronema hispidulum</i>	Sexual morph unknown	De Gruyter et al. 2010
<i>Dematiopleospora</i>	<i>Dematiopleospora mariaae</i>	Asexual morph unknown	Wanasinghe et al. 2014
<i>Diederichomyces</i>	<i>Diederichomyces xanthomendozae</i>	Sexual morph unknown	Trakunyingcharoe et al. 2014
<i>Dothideopsella</i>	<i>Dothideopsella agminalis</i>	Asexual morph unknown	Wijayawardene et al. 2014b
<i>Entodesmium</i>	<i>Entodesmium rude</i>	Asexual morph unknown	Zhang et al. 2009
<i>Eudarluca</i>	<i>Eudarluca australis</i>	Asexual morph unknown	Wijayawardene et al. 2014b
<i>Galliicola</i>	<i>Galliicola pseudophaeosphaeria</i>	Asexual morph unknown	Ariyawansa et al. 2015 b
<i>Leptospora</i>	<i>Leptospora porphyrogona</i>	Asexual morph unknown	Crous et al. 2006
<i>Loratospora</i>	<i>Loratospora aestuari</i>	Asexual morph unknown	Schoch et al. 2009
<i>Neosetophoma</i>	<i>Neosetophoma samararum</i>	Sexual morph unknown	De Gruyter et al. 2010
<i>Neostagonospora</i>	<i>Neostagonospora caricis</i>	Sexual morph unknown	Quaedvlieg et al.2013
<i>Nodulosphaeria</i>	<i>Nodulosphaeria hirta</i>	Asexual morph: Coelomyctous	from this study
<i>Ophiobolus</i>	<i>Ophiobolus disseminans</i>	Asexual morph unknown	Phookamsak et al. 2014
<i>Ophiosphaerella</i>	<i>Ophiosphaerella graminicola</i>	Asexual morph unknown	Zhang et al. 2012
<i>Paraphoma</i>	<i>Paraphoma radicina</i>	Sexual morph unknown	Gruyter et al. 2010
<i>Parastagonospora</i>	<i>Parastagonospora nodorum</i>	Sexual morph = phaeosphaeria-like	Quaedvlieg et al. 2013
<i>Phaeosphaeria</i>	<i>Phaeosphaeria oryzae</i>	Asexual morph = Phaeoseptoria	Quaedvlieg et al. 2013
<i>Phaeosphaeriopsis</i>	<i>Phaeosphaeriopsis glaucopunctata</i>	Asexual morph Coelomycetous	Phookamsak et al. 2014
<i>Phaeostagonospora</i>	<i>Phaeostagonospora nolinae</i>	Sexual morph unknown	Phookamsak et al. 2014
<i>Poaceicola</i>	<i>Poaceicola arundinis</i>	Sexual morph= Phaeosphaeria-like	from this study
<i>Populocrescentia</i>	<i>Populocrescentia forlicesenensis</i>	Sexual morph unknown	Ariyawansa et al. 2015 b
<i>Sclerostagonospora</i>	<i>Sclerostagonospora heraclei</i>	Sexual morph = phaeosphaeria-like	Quaedvlieg et al. 2013
<i>Scolecosporiella</i>	<i>Scolecosporiella typhae</i>	Asexual morph unknown	Phookamsak et al. 2014
<i>Scolicosporium</i>	<i>Scolicosporium fagi</i>	Sexual morph unknown	Wijayawardene et al. 2013
<i>Septoriella</i>	<i>Septoriella phragmitis</i>	Sexual morph unknown	Crous et al. 2015

Genus name	Types species	Sexual/Asexual morph	Reference
<i>Setomelanomma</i>	<i>Setomelanomma holmii</i>	Sexual morph unknown	Phookamsak et al. 2014
<i>Setophoma</i>	<i>Setophoma terrestris</i>	The sexual morph of <i>Setophoma</i> is similar to <i>Phaeosphaeria</i> species	Phookamsak et al. 2014
<i>Stagonospora</i>	-	Sexual morph unknown	Phookamsak et al. 2014
<i>Sulcispora</i>	<i>Sulcispora pleurospora</i>	Asexual morph unknown	Senanayake et al. 2014
<i>Tiarospora</i>	<i>Tiarospora westendorpii</i>	Sexual morph unknown	Phookamsak et al. 2014
<i>Vrystaatia</i>	<i>Vrystaatia aloecicola</i>	Sexual morph unknown	Quaedvlieg et al. 201
<i>Wojnowicia</i>	<i>Wojnowicia hirta</i>	Sexual morph unknown	Crous et al. 2015
<i>Wojnowiciella</i>	<i>Wojnowiciella eucalypti</i>	Sexual morph unknown	Crous et al. 2015a
<i>Xenophoma</i>	<i>Xenophoma puncteliae</i>	Sexual morph unknown	Phookamsak et al. 2014
<i>Xenoseptoria</i>	<i>Xenoseptoria neosaccardoii</i>	Sexual morph unknown	Quaedvlieg et al. 2013

Table 2 Collection details and GenBank accession number of isolates includes in this study. The newly generated sequences are indicated in bold, T signifies ex-type/ex-epitype isolates.

Species name	strain	Host	Country	GenBank accession number	
				ITS	LSU
<i>Allophaeosphaeria muriformia</i>	MFLUCC 13-0349 ^T	-	Italy	KP765680	KP765681
<i>Allophaeosphaeria subcylindrospora</i>	MFLUCC 13-0380 ^T	<i>Dactylis glomerata</i>	Italy	KT314184	KT314183
<i>Ampelomyces quisqualis</i>	CBS 129.79	-	Canada	HQ108038	JX681064
<i>Chaetosphaeronema coonsii</i>	CBS 559.78	<i>Malus sylvestris</i>	Japan	-	EU754196
<i>Dematiopleospora mariae</i>	MFLUCC 13-0612 ^T	<i>Ononis spinosa</i>	Italy	-	KJ749653
<i>Didymella exigua</i>	CBS 183.55 ^T	<i>Rumex arifolius</i>	France	GU237794	EU754155
<i>Diederichomyces caloplacae</i>	CBS 129338	<i>Caloplaca cerina</i>	Canada	KP170639	JQ238643
<i>Diederichomyces cladoniicola</i>	CBS 128026	<i>Cladonia</i> sp.	Spain	KP170642	JQ238628
<i>Diederichomyces ficuzzae</i>	CBS 128019	<i>Ramalina fastigiata</i>	France	KP170647	JQ238616
<i>Entodesmium rude</i>	CBS 650.86	Unknown	-	-	GU301812
<i>Galliicola pseudophaeosphaeria</i>	MFLUCC 14-0527 ^T	<i>Galium</i> sp.	Italy	KT326692	KT326693
<i>Leptospora rubella</i>	CPC 11006	<i>Eucalyptus</i> sp.	Colombia	DQ195780	DQ195792
<i>Loratospora aestuarii</i>	JK 5535B	<i>Juncus roemerianus</i>	USA	-	GU301838
<i>Neosetophoma samarorum</i>	CBS 138.96 ^T	<i>Phlox paniculata</i>	Netherlands	FJ427061	KF251664
<i>Neostagonospora caricis</i>	CBS 135092/S616 ^T	<i>Carex acutiformis</i>	Netherlands	KF251163	KF251667
<i>Neostagonospora elegiae</i>	CBS 135101 ^T	<i>Elegia cuspidata</i>	South Africa	KF251164	KF251668

Species name	strain	Host	Country	GenBank accession number	
				ITS	LSU
<i>Nodulosphaeria aconiti</i>	MFLUCC 13-0728	<i>Aconitum vulparia</i>	Italy		
<i>Nodulosphaeria digitalis</i>	-	<i>Dactylis</i> sp.	Italy	KU058710	KU058720
<i>Nodulosphaeria dolioloides</i>	MFLUCC 15-0065	<i>Achillea</i> sp.	Italy	-	-
<i>Nodulosphaeria hirta</i>	MFLUCC 13-0867	<i>Achillea</i> sp.-	Italy		
<i>Nodulosphaeria sanguisorbae</i>	MFLUCC 13-0745	<i>Sanguisorba minor</i>	Italy		
<i>Nodulosphaeria scabiosae</i>	MFLUCC 14-1111	<i>Scabiosa</i> sp.	Italy		
<i>Nodulosphaeria senecionis</i>	MFLUCC 15-1297	<i>Senecio</i> sp.	Italy	KT290257	KT290257
<i>Nodulosphaeria spectabilis</i>	MFLUCC 14-1112	<i>Peucedanum cervaria</i>	Italy	-	-
<i>Ophiobolus cirsii</i>	MFLUCC 13-0218	-	Italy	KM014664	KM014662
<i>Ophiobolus disseminans</i>	AS2L14-6	-	-	KP117305	-
<i>Ophiosphaerella agrostidis</i>	MFLUCC 12-0007	grass	Thailand	KM434272	KM434282
<i>Ophiosphaerella agrostidis</i>	MFLUCC 11-0152	grass	Thailand	KM434271	KM434281
<i>Paraphoma chrysanthemicola</i>	CBS 522.66	<i>Chrysanthemum morifolium</i>	United Kingdom	FJ426985	KF251670
<i>Paraphoma radicina</i>	CBS 111.79 ^T	<i>Malus sylvestris</i>	Netherlands	KF251172	KF251676
<i>Parastagonospora allouniseptata</i>	MFLUCC 13-0386^T	<i>Dactylis glomerata</i>	Italy	KU058711	KU058721
<i>Parastagonospora avenae</i>	CBS 289.69	<i>Lolium perenne</i>	Germany	KF251174	KF251678
<i>Parastagonospora avenae</i>	CBS 290.69	<i>Lolium perenne</i>	Germany	KF251175	KF251679
<i>Parastagonospora caricis</i>	CBS 135671/S615 ^T	<i>Carex acutiformis</i>	Netherlands	KF251176	KF251680
<i>Parastagonospora dactylidis</i>	MFLUCC 13-0375^T	<i>Dactylis</i> sp.	Italy	KU058712	KU058722
<i>Parastagonospora italica</i>	MFLUCC 13-0377^T	<i>Dactylis</i> sp.	Italy	KU058714	KU058724
<i>Parastagonospora minima</i>	MFLUCC 13-0376^T	<i>Dactylis</i> sp.	Italy	KU058713	KU058723
<i>Parastagonospora nodorum</i>	CBS 110109	<i>Lolium perenn</i>	Denmark	KF251177	KF251681
<i>Parastagonospora poae</i>	CBS 135091	<i>Poa</i> sp.	Netherlands	KF251179	KF251683
<i>Parastagonospora poae</i>	CBS 135089 ^T	<i>Poa</i> sp.	Netherlands	KF251178	KF251682
<i>Parastagonospora poagena</i>	CBS 136776 ^T	<i>Poa</i> sp.	Netherlands	KJ869116	KJ869174
<i>Parastagonospora uniseptata</i>	MFLUCC 13-0387^T	<i>Daucus</i> sp.	Italy	KU058715	KU058725
<i>Phaeosphaeria papayae</i>	CBS 135416/S528 ^T	<i>Carica papaya</i>	Brazil	KF251187	KF251690
<i>Phaeosphaeria alpina</i>	CBS 456.84	<i>Phleum alpinum</i>	Switzerland	KF251181	KF251684
<i>Phaeosphaeria chiangraina</i>	MFLUCC 13-0231 ^T	<i>Oryza sativa</i>	Thailand	KM434270	KM434280
<i>Phaeosphaeria eustoma</i>	CBS 573.86	<i>Dactylis glomerata</i>	Switzerland	AF439479	-
<i>Phaeosphaeria eustoma</i>	CBS 573.86	<i>Dactylis glomerata</i>	Switzerland	AF439479	-
<i>Phaeosphaeria musae</i>	CBS 120026 ^T	Unknown	Unknown	DQ885894	GU301862
<i>Phaeosphaeria nigrans</i>	CBS 307.79	<i>Zea mays</i>	Switzerland	KF251184	KF251687
<i>Phaeosphaeria oryzae</i>	CBS 110110 ^T	<i>Oryza sativa</i>	South Korea	KF251186	KF251689
<i>Phaeosphaeria thysanolaenicola</i>	MFLUCC 10-0563 ^T	<i>Thysanolaena maxima</i> Kuntze	Thailand	KM434266	KM434276

Species name	strain	Host	Country	GenBank accession number	
				ITS	LSU
<i>Phaeosphaeria typharum</i>	CBS 296.54	<i>Nardus stricta</i>	Switzerland	KF251192	KF251695
<i>Phaeosphaeriopsis glaucopunctata</i>	MFLUCC 13-0265 ^T	<i>Ruscus aculeatus</i>	Italy	KJ522473	KJ522477
<i>Phaeosphaeriopsis glaucopunctata</i>	MFLUCC 13-0220	<i>Ruscus hypoglossum</i>	Italy	KJ522474	KJ522478
<i>Phaeosphaeriopsis triseptata</i>	MFLUCC 13-0271 ^T	<i>Ruscus aculeatus</i>	Italy	KJ522475	KJ522479
<i>Poaceicola arundinis</i>	-	reed (<i>Juncaceae</i>)	Italy	KU058716	KU058726
<i>Poaceicola bromi</i>	MFLUCC 13-0739^T	<i>Bromus sterilis</i>	Italy	KU058717	KU058727
<i>Poaceicola elongate</i>	MFLUCC 12-4444^T	Unknown	Italy	KM491546	KM491548
<i>Populocrescentia forlicesenensis</i>	MFLUCC 15-0651 ^T	<i>Populus nigra</i>	Italy	KT306948	KT306952
<i>Sclerostagonospora cycadis</i>	CBS 123538 ^T	<i>Cycas revoluta</i>	Japan	FJ372393	FJ372410
<i>Scolicosporium minkeviciusii</i>	MFLUCC 12-0089 ^T	<i>Quercus pubescens</i>	Italy	-	KF366382
<i>Septoriella allojunci</i>	-	<i>Juncus</i> sp.	Italy	KU058718	KU058728
<i>Septoriella hubertusii</i>	CBS 338.86 ^T	<i>Phragmites australis</i>	France	KF251230	KF251733
<i>Septoriella leuchtmannii</i>	CBS 459.84	<i>Phragmites australis</i>	Switzerland	KF251188	KF251691
<i>Septoriella oudemansii</i>	CPC 24116 ^T	<i>Phragmites australis</i>	Netherlands	-	KJ869224
<i>Septoriella phragmitis</i>	CPC 24118 ^T	<i>Phragmites</i> sp.	Netherlands	KR873251	KR873279
<i>Septoriella poae</i>	CBS 136766 ^T	<i>Poa</i> sp.	Netherlands	KJ869111	KJ869169
<i>Setomelanomma holmii</i>	CBS 110217	Unknown	-	-	GU301871
<i>Setophoma chromolaena</i>	CBS 135105 ^T /CPC 18553	<i>Chromolaena odorata</i>	Brazil	KF251244	KF251747
<i>Setophoma sacchari</i>	CBS 333.39	<i>Saccharum officinarum</i>	Brazil	KF251245	KF251748
<i>Setophoma sacchari</i>	MFLUCC 11-0154	<i>Saccharum officinarum</i>	Thailand	KJ476144	KJ476146
<i>Stagonospora neglecta</i>	CBS 343.86	<i>Phragmites australis</i>	France	AJ496630	EU754218
<i>Sulcisporea pleurospora</i>	MFLUCC 13-0796	Unknown	Italy	KP271443	KP271444
<i>Vagicola vagans</i>	CBS 604.86 ^T	<i>Calamagrostis arundinacea</i>	Sweden	KF251193	KF251696
<i>Vrystaatia aloecicola</i>	CBS 135107/CPC 20617 ^T	<i>Aloe maculata</i>	South Africa	KF251278	KF251781
<i>Wojnowicia dactylidicola</i>	MFLUCC 13-0738 ^T	<i>Dactylis glomerata</i>	Italy	KP744469	KP684147
<i>Wojnowicia loniceriae</i>	MFLUCC 13-0737 ^T	<i>Lonicerasp</i>	Italy	KP744471	KP684151
<i>Wojnowicia spartii</i>	MFLUCC 13-0402^T	<i>Spartium</i> sp.	Italy	KU058719	KU058729
<i>Wojnowicia dactylidis</i>	MFLUCC 13-0735 ^T	<i>Dactylis glomerata</i>	Italy	KP744470	KP684149
<i>Wojnowiciella eucalypti</i>	CPC 25024 ^T	<i>Eucalyptus grandis</i>	Colombia	KR476741	KR476774
<i>Wojnowiciella viburni</i>	MFLUCC 12-0733 ^T	<i>Viburnum utile</i>	China	KC594286	KC594287
<i>Xenophoma puncteliae</i>	CBS 128022 ^T	<i>Punctelia rudecta</i>	USA	JQ238617	JQ238619
<i>Xenoseptoria neosaccardoii</i>	CBS 128665 ^T	<i>Lysimachia vulgaris</i> var. <i>Davurica</i>	South Korea	KF251281	KF251784
<i>Xenoseptoria neosaccardoii</i>	CBS 120.43	<i>Cyclamen persicum</i>	Netherlands	KF251280	KF251783

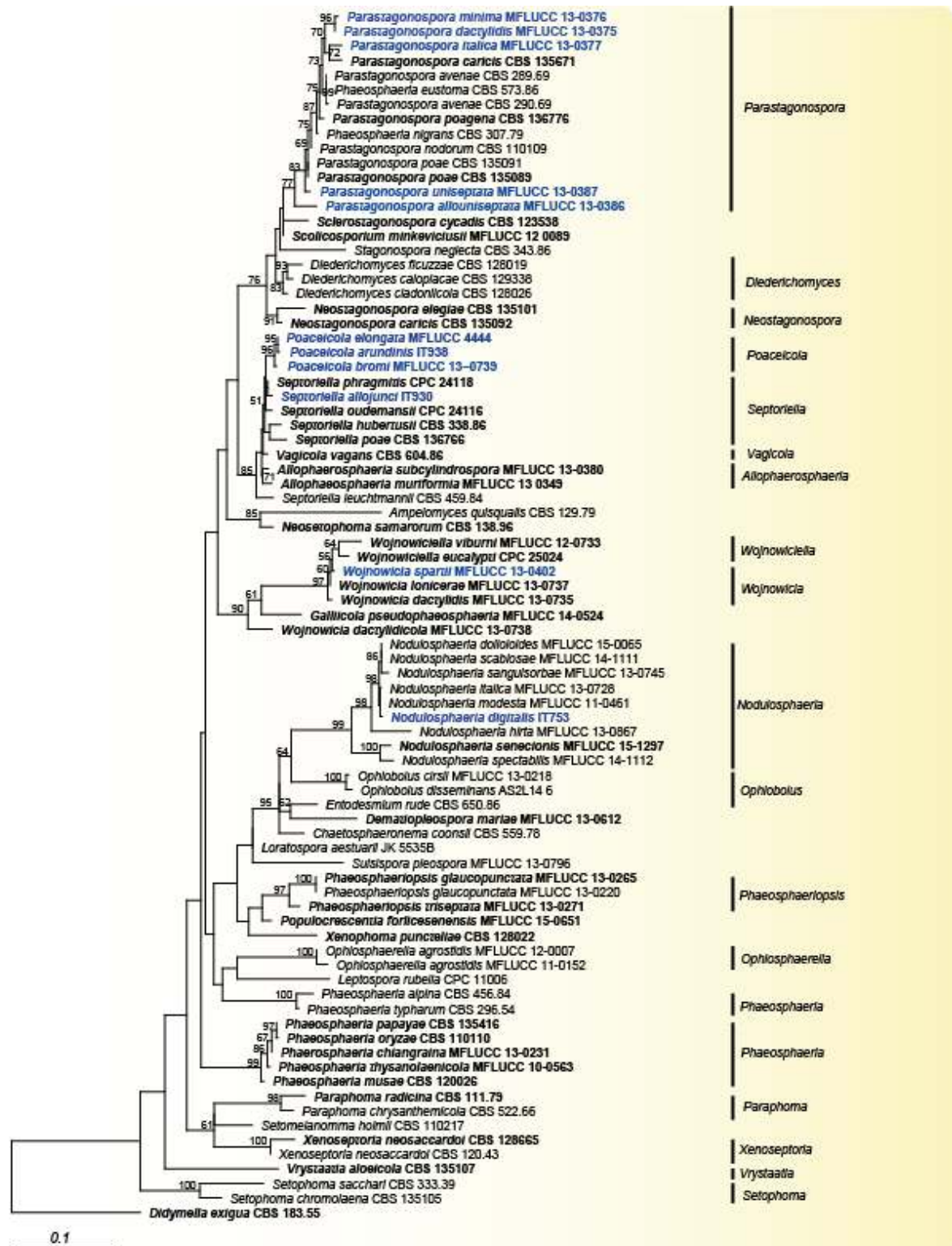


Fig. 1 – Best scoring RAxML tree of *Phaeosphaeriaceae* strains obtained from analysis of ITS and LSU sequence data. RAxML bootstrap support values (equal to or greater than 50% based on 1,000 replicates) are shown at the nodes. The ex-type strains are in bold; the new isolates are in blue. The tree is rooted to *Didymella exigua* CBS 183.55.

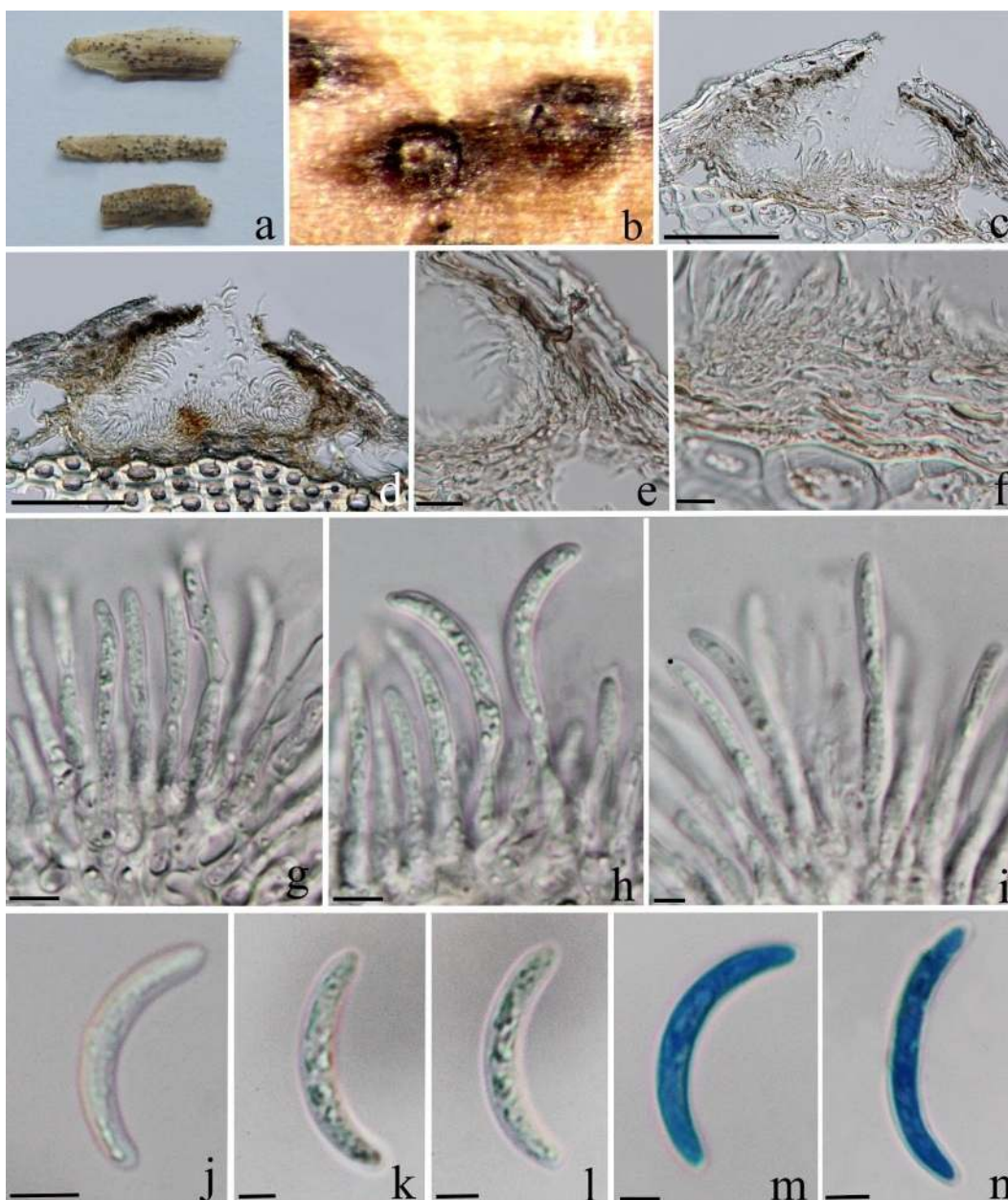


Fig. 2 – *Nodulosphaeria digitalis* (MFLU 15–2716, **holotype**) a Herbarium specimen b Appearance of black conidiomata on the host. c, d Vertical section of conidioma. e, f Section of peridium. g–i Conidiogenous cells and developing conidia. g–n Conidia. – Bars: c–d = 100 μm ; e–f = 20 μm ; g–i = 5 μm ; j–n = 5 μm .

Parastagonospora allouniseptata W.J. Li, Camporesi, D.J. Bhat & K.D. Hyde, **sp. nov.**

Fig. 3

Index Fungorum number: IF551665

Facesoffungi number: FoF 01307

Etymology – Named after its morphological similarity to *Parastagonospora uniseptata*

Holotype – MFLU 15–0698

Saprobic on dead stem of *Dactylis glomerata* L. (*Poaceae*), forming conspicuous, rounded, black fruiting bodies. **Sexual morph** – Undetermined. **Asexual morph** – Coelomycetous. *Conidiomata* 60–90 μm high, 70–90 μm diam., pycnidial, black, solitary or gregarious, semi-

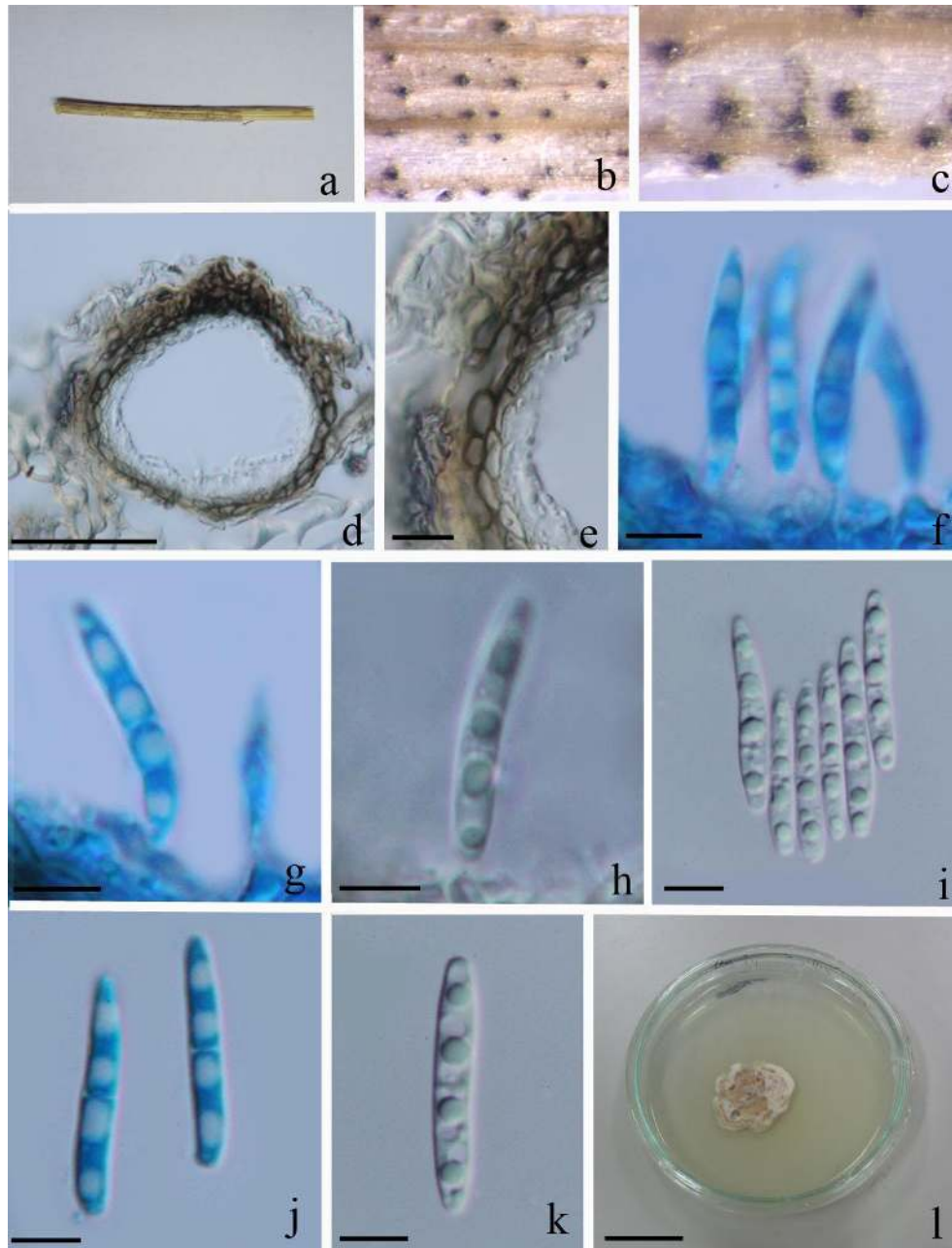


Fig. 3 – *Parastagonospora allouniseptata* (MFLU 15–0698, **holotype**) a Herbarium specimen. b–c Appearance of black conidiomata on the host. d Vertical section of conidioma. e Section of peridium. f–h Conidiogenous cells and developing conidia. i–k Conidia. l Culture on PDA. – Bars: d = 50 μ m; e = 10 μ m; f–k = 5 μ m; l = 25 mm.

immersed, unilocular, globose to subglobose, ostiolate. *Ostiole* centrally located, papillate. *Wall of conidiomata* 5–10 μ m wide, comprising outer layer of brown to dark brown, thin-walled cell of *textura globosa*, and an inner layer of light brown to hyaline cell of *textura angularis*. *Conidiophores* reduced to conidiogenous cells, arising from inner layers of stroma. *Conidiogenous cells* 3–5 μ m long \times 3–5.5 μ m wide, phialidic, integrated, short, hyaline, smooth. *Conidia* 16–22 \times 2.5–3.5 μ m (\bar{x} = 19.5 \times 3 μ m; n = 30), hyaline, subcylindrical, 1-septate, with narrow and subobtuse apex and truncate base, granular, smooth and thick-walled.

Culture characteristics – Colonies slow growing on PDA, reaching 15 mm diam. after 4 weeks at 20–25°C, circular to irregular, with uneven margins, whitened to pink, flattened, filamentous, sparse, aerial mycelium on the surface, reverse similar in colour.

Material examined – ITALY, Province of Forlì-Cesena [FC], Galeata, Passo delle Forche, on dead stem of *Dactylis glomerata* L. (*Poaceae*), 15 October 2013, Erio Camporesi, IT-719 (MFLU 15–0698, **holotype**); ex-type living culture, MFLUCC 13–0386, KUMCC 15–0134; *ibid.* IT-719A (KUN! HKAS 90735, **isotype**).

Notes – *Parastagonospora allouniseptata* is similar to *P. uniseptata* in having pycnidial, globose to subglobose, conidiomata, with hyaline, subcylindrical, 1-septate, granular conidia, but is distinct in having shorter and narrower conidia than *P. uniseptata* (14–18 × 2–3 µm). Phylogenetically, *P. allouniseptata* clusters with *Parastagonospora* species with 88% bootstrap support. Furthermore, *P. allouniseptata* forms a distinct branch in the *Parastagonospora* phylogeny (Fig. 1).

Parastagonospora dactylidis W.J. Li, Camporesi, D.J. Bhat & K.D. Hyde, **sp. nov.**

Fig. 4

Index Fungorum number: IF551666

Facesoffungi number: FoF 01303

Etymology – Named after the host genus, *Dactylis*.

Holotype – MFLU 15–0693

Saprobic on dead stem of *Dactylis* sp., forming numerous, conspicuous, dark brown fruiting bodies in a linear series on the surface. **Sexual morph** – Undetermined. **Asexual morph** – Coelomycetous. *Conidiomata* 50–100 µm high, 100–150 µm diam., pycnidial, brown, separate, ampulliform or globose, immersed, unilocular, thick-walled, ostiolate. *Wall of conidiomata* 10–15 µm wide, composed of 3–4 layers, with outer 1–2-layers of dark brown and inner 1–2-layers of hyaline cells of *textura angularis*. *Ostiole* circular, papillate. *Conidiophores* reduced to conidiogenous cells. *Conidiogenous cells* 2–6 µm long × 3–8 µm wide, phialidic, ampuliform, determinate, hyaline, unbranched, and developing only at the base, formed from the inner cells of the pycnidial wall. *Conidia* 25–40 × 4–5.5 µm (\bar{x} = 30 × 4.5 µm; n = 30), hyaline, fusiform, curved, rounded at both ends, slightly narrower at the base, 3-septate, constricted at the septa, with distinctly granular cytoplasm.

Culture characteristics – Colonies on PDA, reaching 20 mm after 2 wk at 20–25°C, with dense, white, flat, aerial mycelium, with rounded, smooth, margins, reverse similar in colour.

Material examined – ITALY, Province of Arezzo [AR], Passo della Consuma, on dead stem of *Dactylis* sp., 19 June 2012, Erio Camporesi, IT-448 (MFLU 15–0693, **holotype**); ex-type living culture, MFLUCC 13–0375, ICMP 20774, KUMCC15-0131; *ibid.* (KUN! HKAS 90738, **isotype**).

Notes – *Parastagonospora dactylidis* is most closely related to *P. minima* (Fig. 1). However, the individual sequence of *P. dactylidis* as compared with *P. dactylidis* has seven different substitutions in the LSU sequence data. Morphologically, these two species also can be easily distinguished in form of conidia. *P. dactylidis* has fusiform conidia with a slightly narrower base, and distinctly granular cytoplasm, whereas *P. minima* has subcylindrical conidia which are wider in the basal half, and narrow at the apex. In addition, the conidiomata and conidia of *P. dactylidis* are larger and longer than those of *P. minima* (conidiomata: 40–70 µm high, 50–100 µm diam., conidia: 20–28 × 3.5–4.5 µm). Thus *P. dactylidis* is introduced as a novel species.

Parastagonospora minima W.J. Li, Camporesi, D.J. Bhat & K.D. Hyde, **sp. nov.**

Fig. 5

Index Fungorum number: IF551667

Facesoffungi number: FoF 01304

Etymology – Named after the small conidiomata.

Holotype – MFLU 15–0694

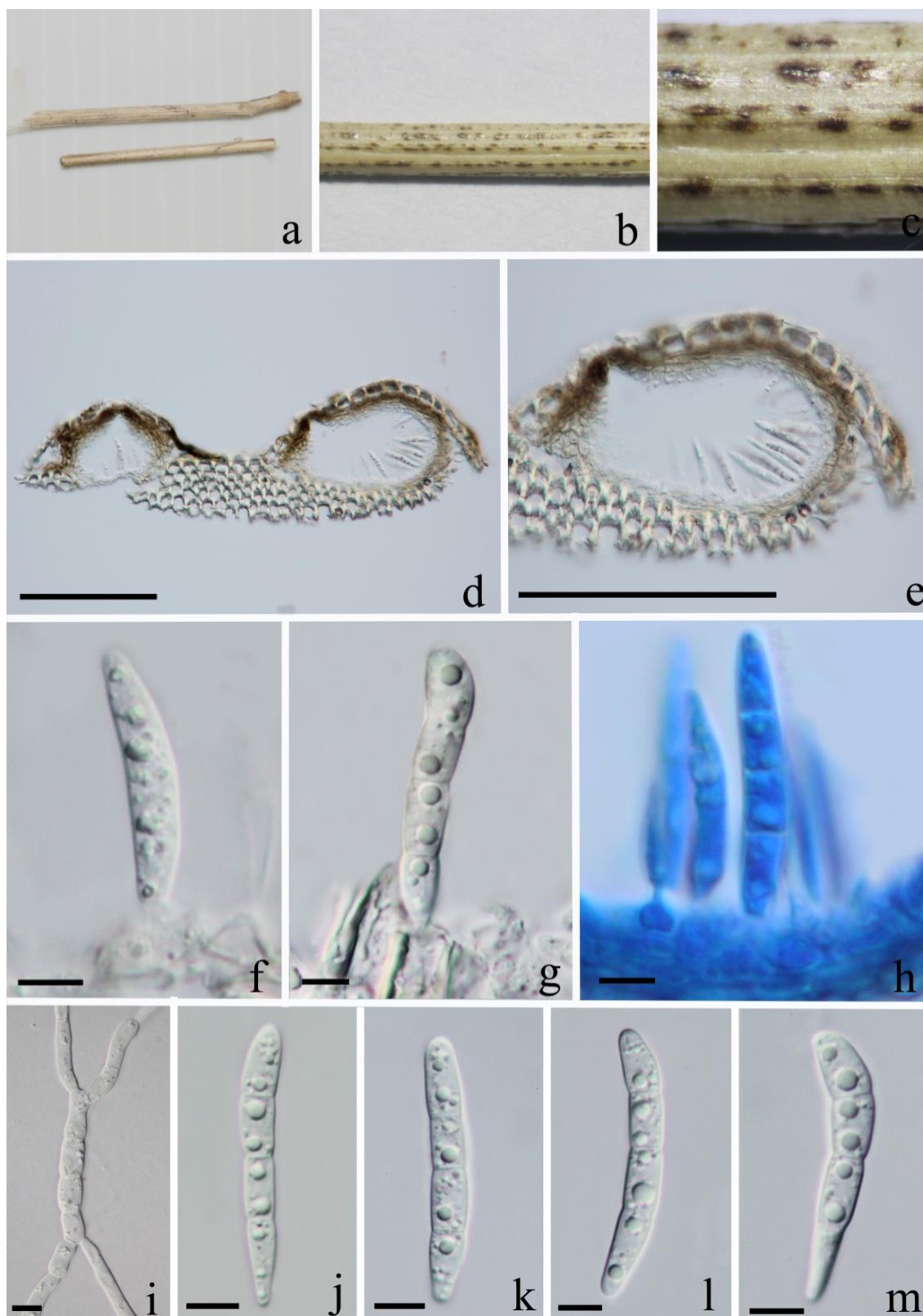


Fig. 4 – *Parastagonospora dactylidis* (MFLU 15–0693, **holotype**) a Herbarium specimen. b, c Appearance of brown conidiomata on the host. d–e Vertical section of conidioma. f–h Conidiogenous cells and developing conidia. i Germinating spore. j–m Conidia. – Bars: d–e = 50 μ m; f–m = 5 μ m.

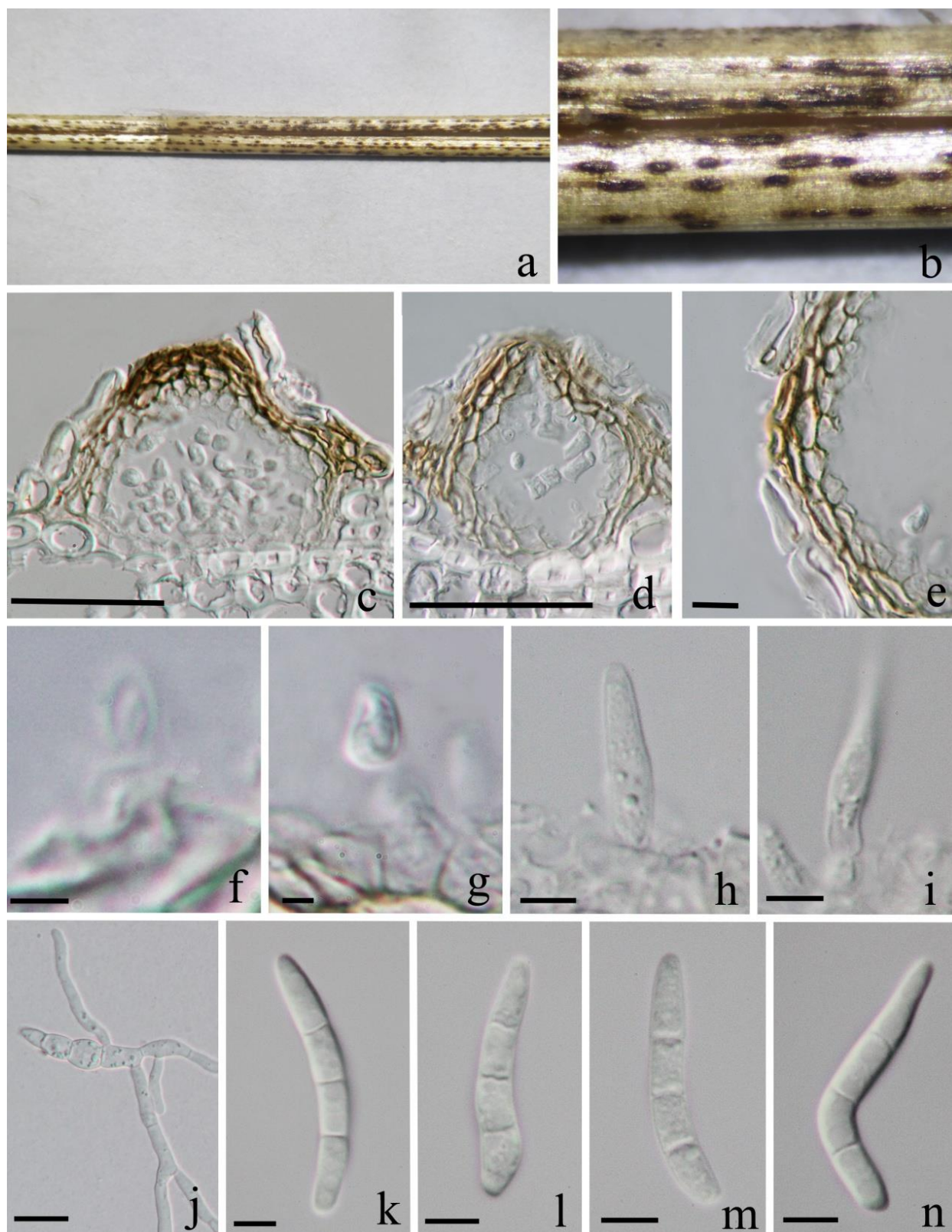


Fig. 5 – *Parastagonospora minima* (MFLU 15–0694, **holotype**) a Herbarium specimen. b Appearance of black conidiomata on the host. c–d Vertical section of conidioma. e Section of peridium. f–i Conidiogenous cells and developing conidia. j Germinating spore. k–n Conidia. – Bars: c–d = 100 μm ; e = 20 μm ; f–i, k–m = 5 μm ; j = 20 μm .

Saprobic on dead stem of *Dactylis* sp., with dark brown fruiting bodies, in a linear series. **Sexual morph** – Undetermined. **Asexual morph** – Coelomycetous. *Conidiomata* 40–70 µm high, 50–100 µm diam., pycnidial, brown to black, amphigenous, separate, gregarious or confluent, obpyriform or globose, subepidermal, unilocular, ostiolate, thick-walled. *Wall of conidiomata* 6–10 µm wide, composed of dark brown to pale brown, thick-walled cells of *textura angularis*. *Ostiole* centrally located, papillate, circular. *Conidiophores* reduced to conidiogenous cells. *Conidiogenous cells* 3–6.5 µm long × 3–7 µm wide, phialidic, hyaline, smooth. *Conidia* 20–28 × 3.5–4.5 µm (\bar{x} = 25.5 × 4 µm; n = 30), hyaline, subcylindrical, wider at the basal half, narrow and rounded at both ends, slightly curved, 3-euseptate, smooth-walled.

Culture characteristics – Colonies on PDA, reaching 25 mm after 2 wk at 20–25°C, with moderate aerial mycelium, flat, dense, spreading, with rounded, smooth margins.

Material examined – ITALY, Province of Arezzo [AR], Passo della Consuma, on dead stem of *Dactylis* sp., 19 June 2012, Erio Camporesi IT-451 (MFLU 15–0694, **epitype**); ex-type living culture, MFLUCC 13–0376, ICMP 20776, KUMCC15–0132; *ibid.* (KUN! HKAS 90734, **isotype**).

Notes – *Parastagonospora minima* is morphologically similar with the generic type, *P. nodorum*, in having subcylindrical, hyaline conidia, with an obtuse apex, as well as being of similar size (*P. nodorum* 13–28 × 2.8–4.6 µm) (Quaedvlieg et al 2013). However, recognizable differences between these two species are in the septation. *Parastagonospora minima* has uniformly 3-septate conidia, whereas *P. nodorum* has (0–)1(–3)-septate conidia. Phylogenetically, *P. minima* is distinct from any other species of *Parastagonospora* (Fig.1). Based on the above morphological characters together with molecular sequence data, *P. minima* is introduced as a new species.

Parastagonospora italica W.J. Li, Camporesi, D.J. Bhat & K.D. Hyde, **sp. nov.**

Fig. 6

Index Fungorum number: IF551668

Facesoffungi number: FoF 01305

Etymology – Named after the country of collection.

Holotype – MFLU 15–0696

Saprobic on dead stem of *Dactylis* sp.. **Sexual morph** – Undetermined. **Asexual morph** – Coelomycetous. *Conidiomata* 65–80 µm high, (40–) 100–150 µm diam., pycnidial, brown to black, solitary or gregarious, subepidermal, immersed, unilocular, globose to subglobose, thick-walled, black. *Wall of conidiomata* 10–20 µm wide, composed of 2–4 layers, thick-walled cells of *textura angularis*, brown. *Ostiole* centrally located, circular, short, papillate. *Conidiophores* reduced to conidiogenous cells, arising from inner layers of pycnidia. *Conidiogenous cells* short, broadly cylindrical, phialidic, hyaline, smooth. *Conidia* 25–32 × 3–4 µm (\bar{x} = 29.5 × 3–5 µm; n = 20), hyaline, cylindrical-fusiform, with narrow and obtuse apex, truncate at base, slightly curved, 3-euseptate, smooth and thick-walled.

Culture characteristics – Colonies on PDA, reaching 20 mm after 10 d at 20–25°C, with crenate, white margins, pink, flattened, with dense, aerial mycelium on the surface; reverse pale-yellowish to pale brown in the middle, whitened at the margins.

Material examined – ITALY, Province of Forli-Cesena [FC], Santa Sofia, Campigna, on dead stem of *Dactylis* sp., 24 June 2012, Erio Camporesi, IT-471 (MFLU 15–0696, **holotype**); ex-type living culture, MFLUCC 13–0377, ICMP 20777, KUMCC. *ibid.* 15 May 2013, Erio Camporesi, IT-471A (KUN! HKAS 90736, **isotype**).

Notes – In the multi-locus phylogeny, *Parastagonospora italica* clusters with *P. caricis* Quaedv. et al., but this grouping is not well-supported. Morphologically, these two species can easily be distinguished by dimension and septation of conidia; *P. italica* has 3-septate conidia which are shorter and narrower than the 7–15-septate, (50–)60–70(–75) × (5–) 6 µm conidia in *P. caricis*. In addition, the conidiomata of *P. italica* are smaller than those of *P. caricis* which are up to 250 µm.



Fig. 6 – *Parastagonospora italica* (MFLU 15–0696, **holotype**) a Herbarium specimen. b Appearance of black conidiomata on the host surface. c–d Vertical section of conidioma. e Section of peridium. f–g Conidiogenous cells and developing conidia. h Germinating spore. i–k Conidia. l Culture on PDA. – Bars: c = 100 μ m; d = 25 μ m; e = 20 μ m; f–g = 5 μ m; h = 10 μ m; i–k = 5 μ m; l = 25 mm.

Parastagonospora uniseptata W.J. Li, Camporesi, D.J. Bhat & K.D. Hyde, **sp. nov.**

Fig. 7

Index Fungorum number: IF551669

Facesoffungi number: FoF 01306

Etymology – Named after the fact that conidia are 1-septate.

Holotype – MFLU 15–0699

Saprobic on dead stem of *Daucus* sp., forming conspicuous, to oval, black, fruiting bodies.

Sexual morph – Undetermined. **Asexual morph** – Coelomycetous. *Conidiomata* 60–100 µm high, 70–100 µm diam, pycnidial, brown to black, appearing on upper surface of the stem, solitary, sometimes gregarious or confluent, immersed to semi-immersed, globose or subglobose, unilocular, with a blackened tip above, ostiolate at the centre, thick-walled. *Wall of conidiomata* 10–20 µm wide, composed of 6-layers, outer 4–5-layers brown, inner 2–3-layers hyaline, upper wall region dark brown, with periclinal and basal wall composed of thick-walled cells of *textura angularis*. *Conidiophores* reduced to conidiogenous cells, arising all around the basal region of the conidiomata. *Conidiogenous cells* 3–6 µm long × 3–6.5 µm wide, ampulliform to broadly conical, phialidic, hyaline, smooth. *Conidia* 14–18 × 2–3 µm (\bar{x} = 16 × 2.5; n = 30), hyaline, subcylindrical, 1-septate, guttulate, truncate at the base, with obtuse apex, smooth-walled.

Culture characteristics – Colonies slow growing on PDA, reaching 20 mm diam. after 4 weeks at 20–25°C, with a glabrous, circular margin, pale yellow, flattened, filamentous, sparse, with aerial mycelium on the surface, similar in colour from below.

Material examined – ITALY, Province of Arezzo [AR], Modigliana, Bagno di Cetica, on dead stems of *Daucus* sp., 18 September 2012, Erio Camporesi, IT-727 (MFLU 15–0699, **holotype**); ex-type living culture, MFLUCC 13-0387, KUMCC 15–0134; *ibid.* IT-727A (KUN! HKAS 90730, **isotype**).

Notes – *Parastagonospora uniseptata* should be morphologically compared with stagonospora-like asexual morphs. The results show that *P. uniseptata* is more similar to the type of *Neostagonospora*, *N. caricis* Quaedvl. et al. than to other species within *Parastagonospora*. However, phylogenetic tree reconstruction based on multigene (LSU and ITS) sequence analyses shows that *P. uniseptata* clusters together with type *P. nodorum*, as well as other species of *Parastagonospora* (BS= 88%) (Fig. 1). *Parastagonospora uniseptata* however, forms a discrete branch basal to *P. poae* Quaedvl. et al. (Fig. 1). Thus *P. uniseptata* is introduced as a new species in *Parastagonospora*.

Poaceicola W.J. Li, Camporesi, D.J. Bhat & K.D. Hyde, **gen. nov.**

Index Fungorum number: IF551658

Facesoffungi number: FoF 01298

Etymology – Named after inhabiting grass (*Poaceae*).

Type species – ***Poaceicola arundinis*** W.J. Li, Camporesi, D.J. Bhat & K.D. Hyde, **sp. nov.**

Saprobic on dead stems of plant host. **Sexual morph** – *Ascomata* solitary to gregarious, black, immersed to semi-immersed, subepidermal, uniloculate, glabrous, globose to subglobose, papillate. *Ostiole* centrally located, black, smooth, with an ostiole, filled with hyaline periphyses. *Peridium* thick-walled, brown, composed of cells of *textura angularis*. *Hamathecium* comprising cellular, septate, broad, dense pseudoparaphyses. *Asci* 8-spored, bitunicate, fissitunicate, cylindrical, with a short, broad pedicel. *Ascospores* tetraseriate or partially overlapping, reddish-brown, fusiform, straight or inequilaterally curved, echinulate, 10-septate, fourth cell from apex swollen towards middle and slightly longer than adjacent cells, with a conspicuous sharply delimited sheath. **Asexual morph** – Coelomycetous. *Conidiomata* pycnidoid, solitary or aggregated, immersed or semi-immersed, unilocular, globose, papillate, dark brown. *Wall of conidiomata* comprising inner layers of *textura angularis* cells, gradually merging with the outer surrounding layers of brown, *textura oblita*. *Ostiole*

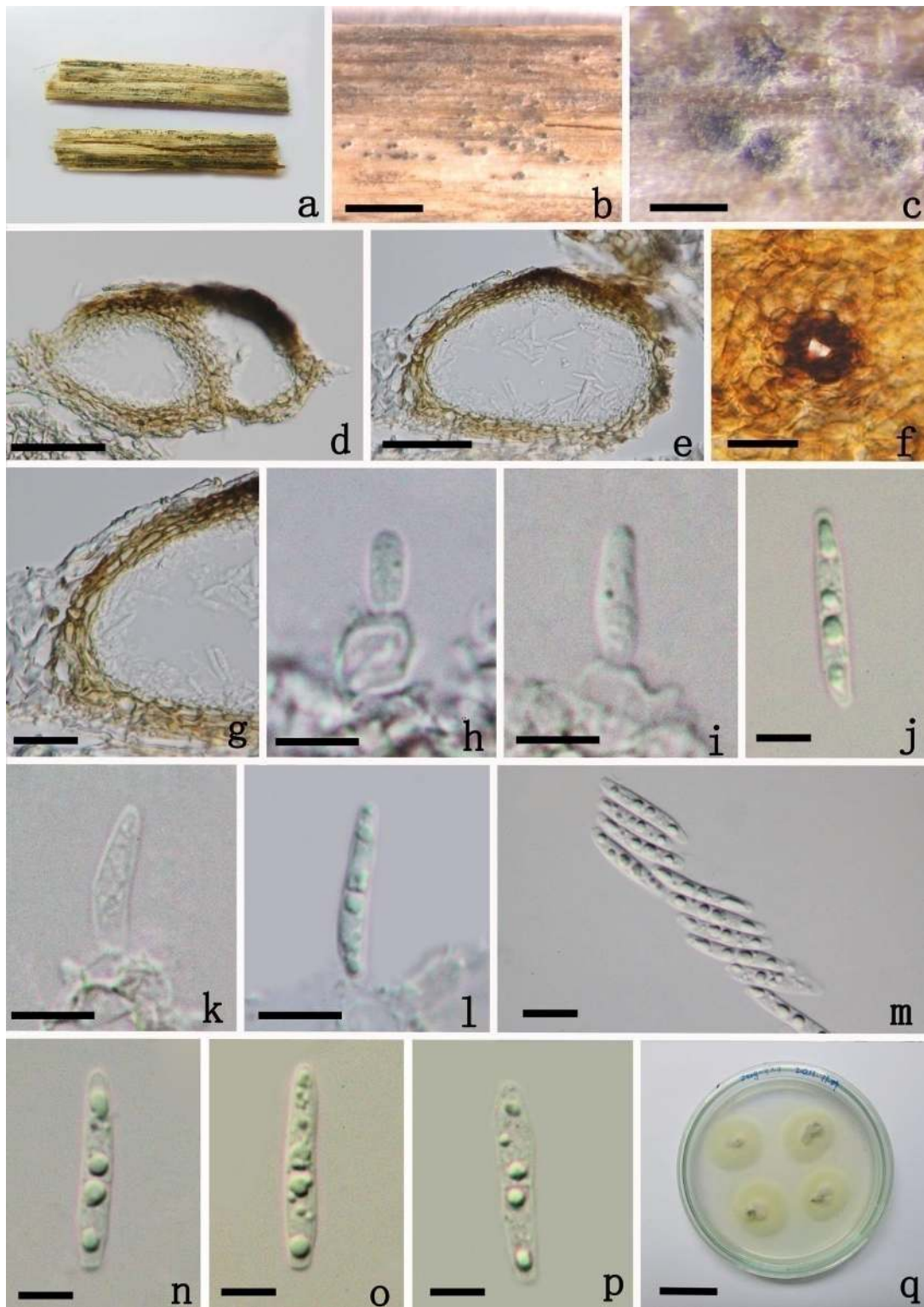


Fig. 7 – *Parastagonospora uniseptata* (MFLU 15–0699, **holotype**) a Herbarium specimen. b–c Appearance of black conidiomata on the host. d–e Vertical section of conidiomata. f Ostiole. g Section of peridium. h–i, k–l Conidiogenous cells and developing conidia. j, m–p Conidia. q Culture on PDA. – Bars: b–c = 200 μ m; d–e = 50 μ m; f = 15 μ m; g = 20 μ m; h–i, k–l = 5 μ m; j, m–p = 5 μ m; q = 25 mm.

central, circular to oval, papillate. *Conidiophores* reduced to conidiogenous cells. *Conidiogenous cells* holoblastic, phialidic, hyaline, smooth-walled, discrete, formed from the inner cells of the stroma. *Conidia* pale brown, cylindrical or sub-cylindrical, up to 7-euseptate, straight or slightly curved, smooth-walled,.

Notes – The phylogeny of the family *Phaeosphaeriaceae* is reconstructed based on combined gene (LSU and ITS) analysis, showing that *Poaceicola* species cluster away from any other genera in *Phaeosphaeriaceae* (Fig. 1). Thus the genus *Poaceicola* is introduced to accommodate *Poaceicola elongata* comb. nov. and two asexual morphs introduced in this study i.e. *Po. arundinis* sp. nov. and *Po. bromi* sp. nov.

Poaceicola arundinis W.J. Li, Camporesi, D.J. Bhat & K.D. Hyde, **sp. nov.**

Fig. 8

Index Fungorum number: IF551659

Facesoffunginumber: FoF 01299

Etymology – Named after the host genus *Arundo*.

Holotype – MFLU 15–0702

Saprobic on dead stems of *Arundo plinii* Turra, forming conspicuous, rounded, black fruiting bodies. **Sexual morph** – Undetermined. **Asexual morph** – Coelomycetous. *Conidiomata* 100–150 µm high, 100–200 µm diam., pycnidoid, dark brown, solitary or aggregated, semi-immersed, unilocular, globose, papillate. *Wall of conidiomata* 10–30 µm wide, comprising inner layers of cells of *textura angularis*, gradually merging with the outer, surrounding layers of brown, *textura oblita*. *Ostiole* central, circular to oval, papillate. *Conidiophores* reduced to conidiogenous cells. *Conidiogenous cells* 3–6 µm wide, holoblastic, phialidic, hyaline, smooth-walled, discrete, formed from the inner cells of the stroma. *Conidia* 30–40 × 6.5–10 µm (\bar{x} = 35 × 8 µm; n = 30), pale brown, cylindrical, flexuous, up to 8-euseptate, slightly curved, smooth-walled, with middle cells wider than end cells, guttulate, with an acute apex, truncate at the base.

Material examined – ITALY, Province of Arezzo [AR], Montemezzano, on dead on dead stems of *Arundo plinii* Turra, 25 August 2013, Erio Camporesi, IT-938 (MFLU 15–0702); *ibid.* (KUN! HKAS 90732).

Notes – *Poaceicola arundinis* is distinct from *Po. bromi* in conidiomata and conidia form. *Poaceicola arundinis* has globose conidiomata with circular to oval, short ostioles, while *Po. bromi* has pyriform conidiomata with cylindrical, long ostioles. In addition, the conidia of *Po. arundinis* with up to 8-septa are longer and wider than those of *Po. bromi* which are 7-septate and 15–23 × 3–5 µm.

Poaceicola bromi Wijayawardene, W.J. Li, Camporesi, D.J. Bhat & K.D. Hyde, **sp. nov.**

Fig. 9

Index Fungorum number: IF551660

Facesoffungi number: FoF 01300

Etymology – Named after the host genus, *Bromus*.

Holotype – MFLU 15–2719

Saprobic on dead twigs of *Bromus sterilis* L. (*Poaceae*). **Sexual morph** – Undetermined. **Asexual morph** – Coelomycetous. *Conidiomata* 310–350 µm high, 240–280 µm diam., pycnidoid, dark brown, solitary, semi-immersed, unilocular, globose, papillate. *Wall of conidiomata* 29–49 µm wide, thick-walled, composed of thick-walled cells of *textura angularis* at the outside, with inner layers lighter and flattened. *Setae* 20–30 × 5–6 µm, lateral, sparse, subcylindrical, 1–2-septate, dark brown. *Ostiole* centrally located, cylindrical to subcylindrical, papillate. *Conidiophores* reduced to conidiogenous cells. *Conidiogenous cells* 1–2 × 1–3 µm, holoblastic, phialidic, hyaline, smooth-walled, discrete, formed from the inner cells of the stroma. *Conidia* 15–23 × 3–5 µm (\bar{x} = 20.5 × 4 µm; n = 30), pale brown to dark brown, cylindrical, straight or slightly curved, obtuse at both ends, up to 7-euseptate, constricted at the septa, smooth-walled.

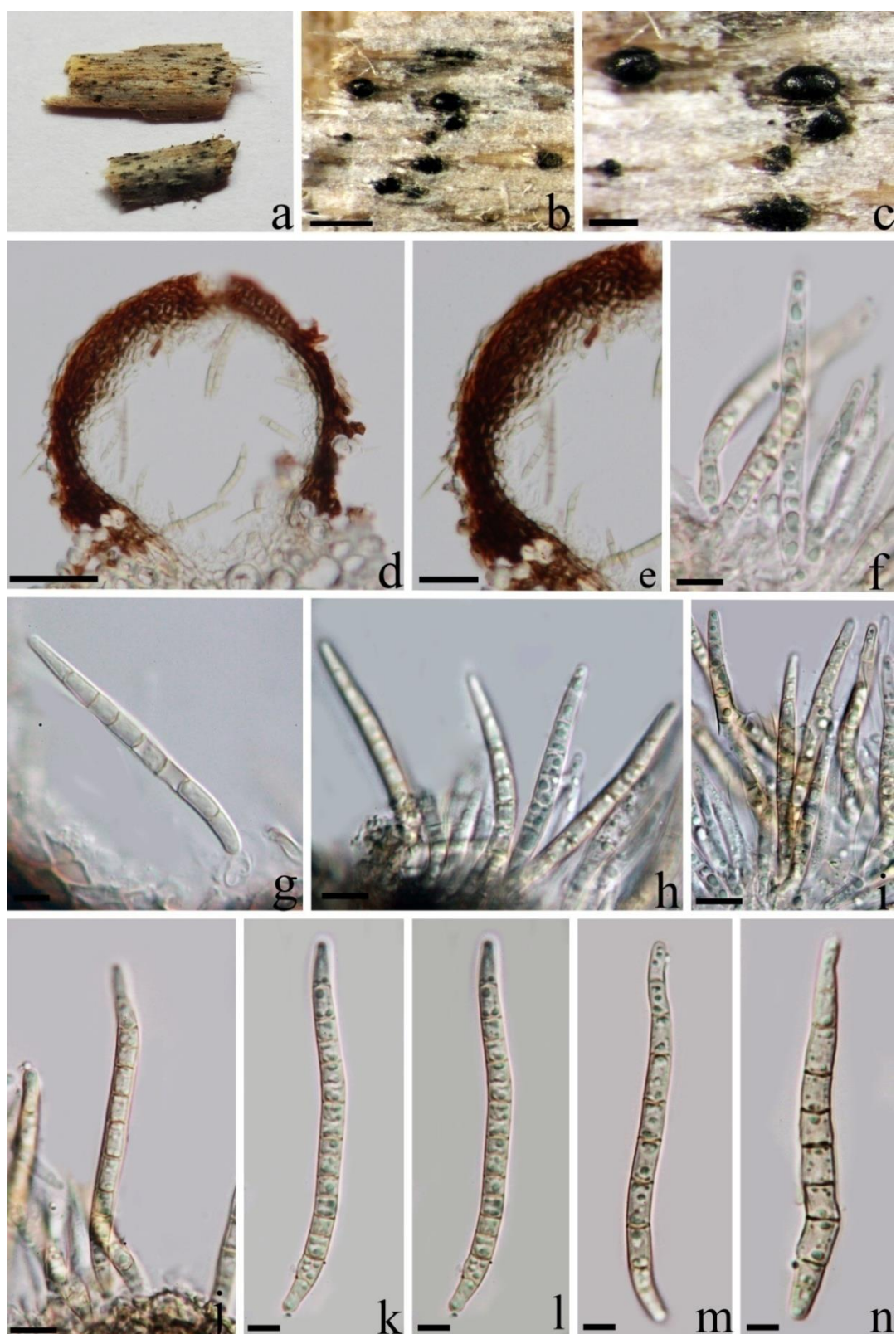


Fig. 8 – *Poaceicola arundinis* (MFLU 15–0702, **holotype**) a Herbarium specimen. b–c Appearance of black conidiomata on the host. d Vertical section of conidioma. e Section of peridium. f–i Conidiogenous cells and developing conidia. j–n Conidia. m Germinating spore. n Culture on PDA. – Bars: b = 200 μ m; c = 20 μ m; d = 100 μ m; e = 50 μ m; f–i = 10 μ m; j–n = 10 μ m.

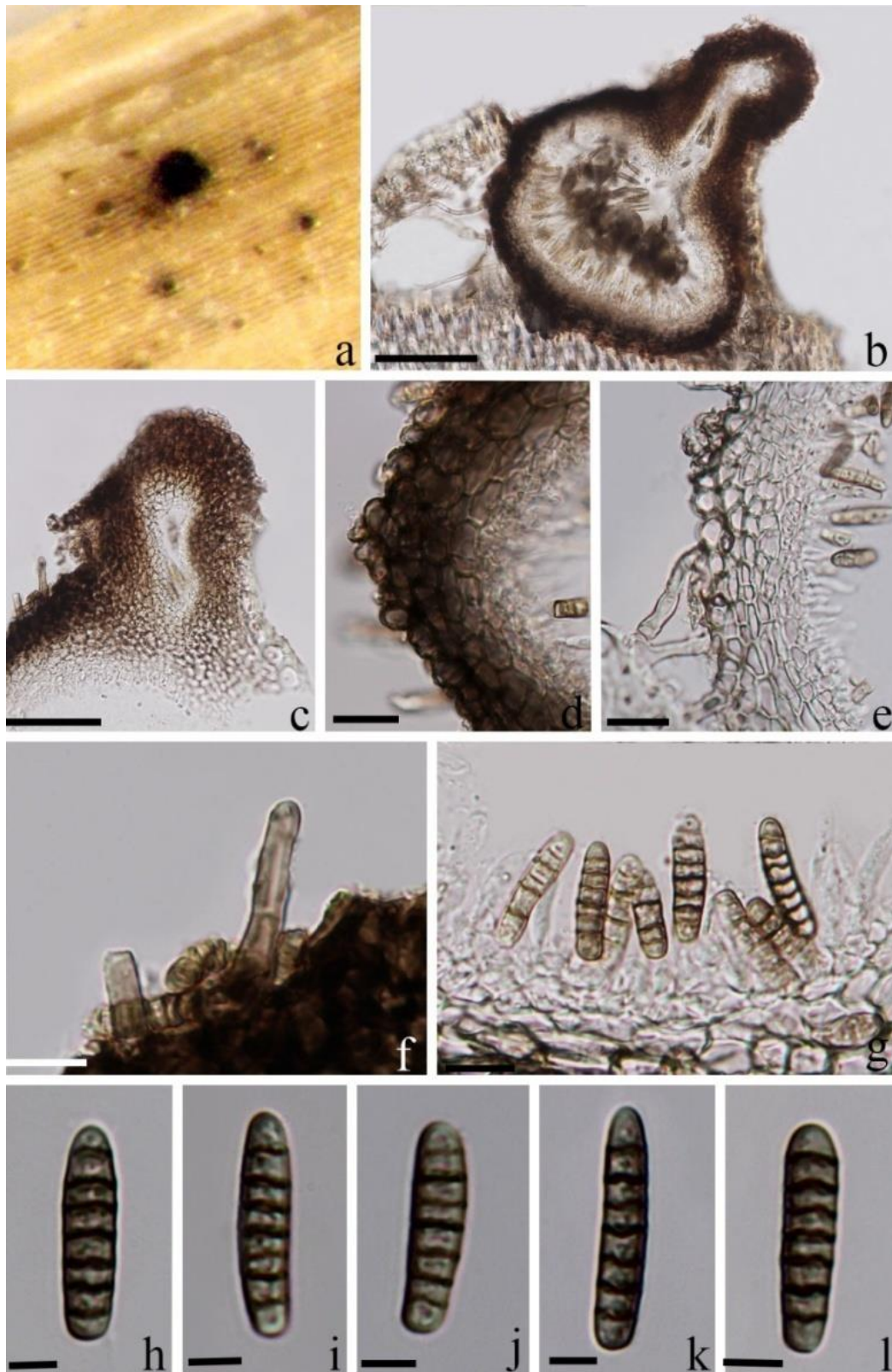


Fig. 9 – *Poaceicola bromi* (MFLU MFLU 15–2719, **holotype**) a Appearance of black conidioma on host. b Vertical section of conidioma. c Ostiole. d–e Section of peridium. f–g Conidiogenous cells and developing conidia. h–l Conidia. – Bars: b = 100 μ m; c = 50 μ m; d–e = 20 μ m; f–g = 10 μ m; h–l = 5 μ m.

Culture characteristics – On PDA slow growing, attaining 2 cm diam. in 7 days, with circular to slightly wavy margin, greyish white from above, grey from below, with thin mycelium.

Material examined – ITALY, Province of Forlì-Cesena [FC], on leaves of *Bromus sterilis* L. (*Poaceae*), 19 July 2013, Erio Camporesi, NNW IT-1389 (MFLU 15–2719, **holotype**). (KUN! HKAS 90729, isotype), living cultures, MFLUCC 13–0739, GUCC 1389

Notes – *Poaceicola bromi* is morphologically and phylogenetically distinct from *Po. arundinis*; see notes under *Po. arundinis*.

Poaceicola elongata (Wehm.) Shoemaker & C.E. Babc.) W.J. Li, Camporesi & K.D. Hyde, **comb. nov.**

Index Fungorum number: IF551661

Facesoffungi number: FoF 01301

Holotype – MFLU14–0635

Basionym – *Phaeosphaeria elongata* (Wehm.) Shoemaker & C.E. Babc., *Can. J. Bot.* 67(5): 1540 (1989)

Synonym – *Leptosphaeria elongata* Wehm., *Mycologia* 44: 633 (1952). 633 (1952).

Notes – This specimen was collected from dead wood in Italy. Ariyawansa et al. (2014) designated the collection as a reference specimen of *Phaeosphaeria elongata*. However, *P. elongata* clusters close to *Po. cylindrispora* and *Po. bromi* with high bootstrap support (BS = 96%) (Fig. 1), and is best assigned to *Poaceicola*.

Septoriella allojunci W.J. Li, Camporesi, D.J. Bhat & K.D. Hyde, **sp. nov.**

Fig. 10

Index Fungorum number: IF551662

Facesoffungi number: FoF 01308

Etymology – Named after its morphological similar to *Septoriella junci*.

Holotype – MFLU 15–0701

Saprobic on dead stem of *Juncus* sp.. **Sexual morph** – Undetermined. **Asexual morph** – Coelomycetous. *Conidiomata* 90–130 µm high, 100–150 µm diam., stromatic, pycnidial, dark brown, solitary or aggregated, linearly disposed with only the dark spots visible in surface view, immersed, unilocular, globose, papillate. *Ostiole* centrally located, circular to oval, apapillate. *Wall of conidiomata* 10–20 µm wide, composed of brown, thick-walled cells of *textura angularis*, at the base, thin-walled, dark brown cells in upper part and surrounding the ostiole. *Conidiophores* reduced to conidiogenous cells. *Conidiogenous cells* 3–6 µm wide, holoblastic, discrete, phialidic, ampulliform, hyaline, smooth-walled, formed from the inner cells of the conidiomata. *Conidia* 48–70 × 3–6.6 µm (\bar{x} = 60 × 4 µm; n = 30), pale brown, subcylindrical, flexuous, with obtuse ends, 3–5-euseptate, smooth-walled, guttulate, with mucoid cap at the apex.

Material examined – ITALY, Province of Forlì-Cesena [FC], Galeata, Strada San Zeno, on dead stems of *Juncus* sp. (*Juncaceae*), 27 November 2012, Erio Camporesi, IT-930 (MFLU 15–0701, **holotype**); *ibid.* (KUN! HKAS 90733, **isotype**).

Notes – Crous et al. (2015b) fixed the application of the type species of generic name (*Septoriella phragmitis* Oudem.), and confirmed the placement of *Septoriella* in *Phaeosphaeriaceae*. In the present study, we introduced an additional taxon, *Septoriella allojunci*. Morphologically, *S. allojunci* shows similarities with *S. junci* and *S. canadensis* in the form of conidiomata and conidiogenous cells, but differs in the dimension, and septation of conidia. *Septoriella allojunci* has 3–5-septate conidia which are longer and wider than those of *S. canadensis* which have 3–4-septate conidia (36–56 × 2.5–3.5 µm), as well as shorter and wider than *S. junci* (6–7-septate, 49–90 × 2–3 µm). Phylogenetically, *S. allojunci* is close to *S. phragmitis* (Fig. 1).

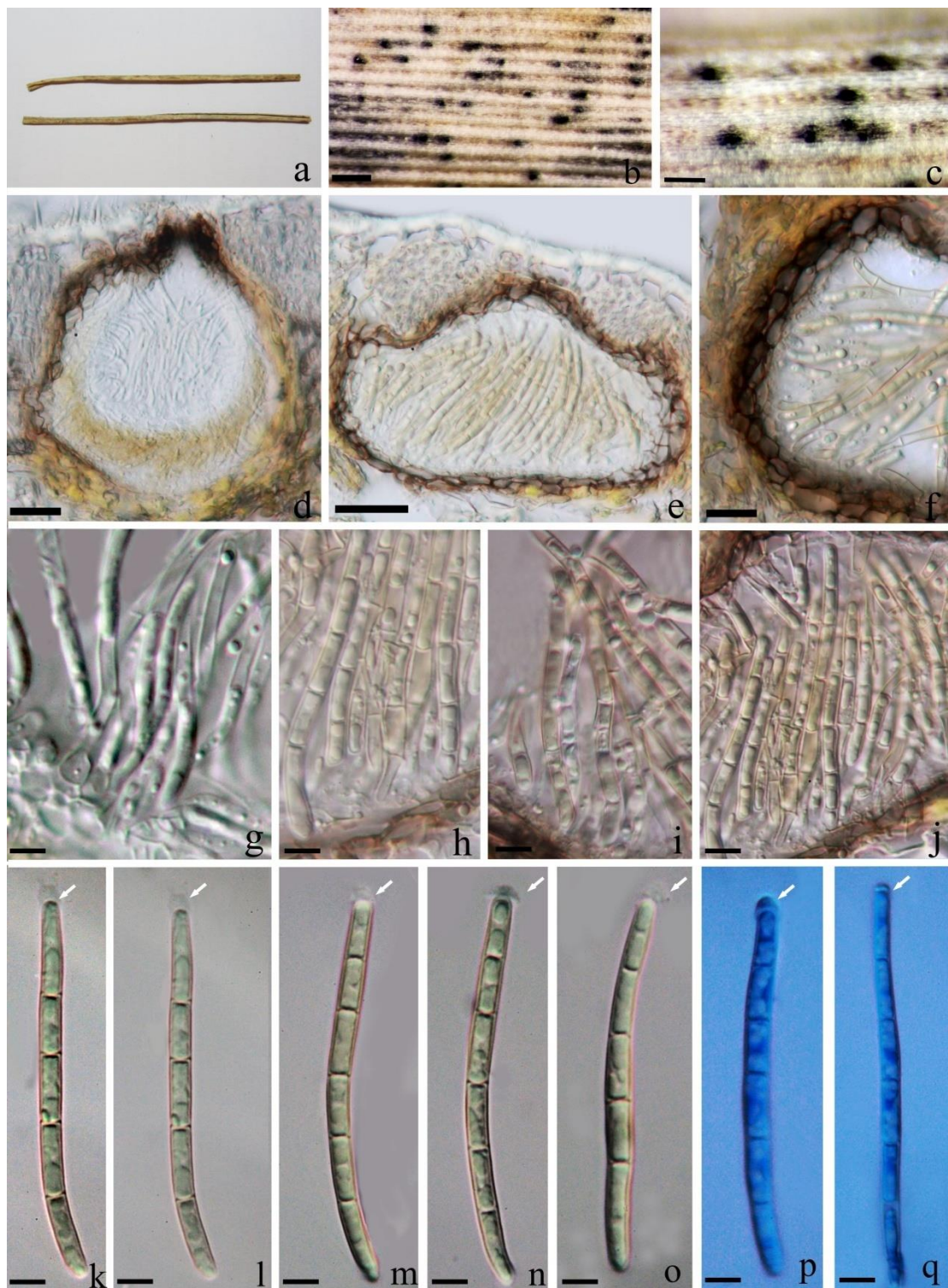


Fig. 10 – *Septoriella allojunci* (MFLU 15–0701, **holotype**) a Herbarium specimen. b–c Appearance of black conidiomata on the host surface. d–e Vertical section of conidioma. f Section of conidioma. g–j Conidiogenous cells and developing conidia. k–q Conidia. – Bars: b–c = 200 μm ; d–e = 100 μm ; f = 50 μm ; g–j = 10 μm ; k–q = 10 μm .

Wojnowicia spartii W.J. Li, Camporesi & K.D. Hyde, **sp. nov.**

Fig. 11

Index Fungorum number: IF551663

Facesoffungi number: FoF 01309

Etymology – Named after the host genus, *Spartium*.

Holotype – MFLU 15–0700

Saprobic on dead stem of *Spartium* sp., forming conspicuous, round, black fruiting bodies.

Sexual morph – Undetermined. **Asexual morph** – Coelomycetous. *Conidiomata* 120–180 µm high, 140–200 µm diam., pycnidial, dark brown, scattered, immersed to semi-immersed, unilocular, globose to subglobose, glabrous, ostiolate. *Ostiole* centrally located, papillate, cylindrical. *Wall of conidiomata* 13–28 µm wide, composed of 4–5-cell layers, of thick-walled cells of *textura angularis*, and inner layers of light brown to hyaline *textura angularis*. *Conidiophores* reduced to conidiogenous cells. *Conidiogenous cells* 2.8–6.5 µm long × 3–7 µm wide, phiallidic, hyaline, integrated, flask-shaped, thick-walled. *Conidia* 28–36 × 5–6 (\bar{x} = 28 × 5 µm; n = 30), dark brown, fusiform or cylindrical, straight or slightly curved, wide in the middle, 7–12-septate, constricted at septa, subobtuse at each end, thick-walled, smooth-walled, guttulate.

Culture characteristics – Colonies slow growing on PDA, reaching 10 mm diam. after 5d at 20–25°C, circular, flattened, with sparse, grey to olivaceous aerial mycelium, surface patches of olivaceous-grey in the older parts; reverse olivaceous-black.

Material examined – ITALY, Province of Forlì-Cesena [FC], Bagno di Romagna, Passo del Carnaio, on dead stem of *Spartium* sp., 17 November 2012, Erio Camporesi, IT-889 (MFLU 15–0700, **holotype**); ex-type living culture, MFLUCC 13–0402, ICMP 20790, KUMCC 15–0136; *ibid.* (KUN! HKAS 90737, **isotype**).

Notes – *Wojnowicia* was introduced by Saccardo (1892) with *W. hirta* as the type species. Subsequently, the genus was expanded to include two more species, viz. *W. colluvium* D.F. Farr & Bills and *W. ephedrae* Hollós (Sutton 1980, Farr & Bills 1995). According to *Index Fungorum* (2015) there are 14 species epithets in *Wojnowicia*. However, *W. graminis* (McAlpine) Sacc. & D. Sacc. and *W. tenella* Pat. were synonymised under *W. hirta* by Sutton (1980). *Wojnowicia buxi* Bertault & Malençon was synonymised under *W. ephedrae* Hollós by Farr & Bills (1995). *Wojnowicia bryophila* Racov., *W. exilis* (Corda) Sacc. & Traverso and *W. lophostoma* (Höhn.) Sacc. were excluded from *Wojnowicia* (Sutton 1980, Farr & Bills 1995, Wijayawardene et al. 2013). *Wojnowicia viburni* Wijayaw. et al. was transferred to *Wojnowiciella* (Crous et al. 2015a). Presently, seven species are accepted in *Wojnowicia*, including *W. spartii* described in this paper (Sutton 1980, Wijayawardene et al. 2013, Liu et al. 2015).

Wojnowicia spartii should be compared with *W. dactylidis* Wijayaw. et al. and *W. lonicerae* Wijayaw. et al. However, conidia of *W. spartii* are shorter than in *W. dactylidis* (35–40 × 4–5.5 µm) and *W. lonicerae* (38–49 × 5–6 µm). *Wojnowicia spartii* has similar conidial dimensions as *W. dactylidicola* Wijayaw. et al. (25–35 × 3.5–6.5 µm (\bar{x} = 28.38 × 4.87 µm), but can be distinguished using septation (7–12 septa in *W. spartii* and 3–5 septa in *W. dactylidicola*). Multigene analyses shows that *W. spartii* is distinct from any other *Wojnowicia* species (Fig. 1).

Discussion

Many genera of ascomycetes are pleomorphic and propagate through sexual or asexual reproduction in different geographical locations and at different times, and it is often difficult to understand the whole life cycle of a species (Hyde et al. 2011, Wijayawardene et al. 2012). Observation of sporulating structures from single ascospores on artificial media, the occurrence of two morphs on the same substrate, or mating studies was previously the only way to link morphs (Wijayawardene et al. 2014). However, the occurrence of an asexual morph adjacent to the sexual morph on a host cannot equivocally show they are linked; they may be unrelated species. DNA



Fig. 11 – *Wojnowicia spartii* (MFLU 15–0700, **holotype**) a Herbarium specimen. b–c Appearance of black conidiomata on the host. d Vertical section of conidioma. e Ostiole. f Section of peridium. j–i Conidiogenous cells and developing conidia. j Germinating spore. k–n Conidia. o Culture on PDA. – Bars: b = 200 μ m; c = 200 μ m; d = 100 μ m; e = 20 μ m; f = 20 μ m; g–i = 5 μ m; j–n = 10 μ m; o = 25 nm.

sequence analysis and phylogenetic inference can irrefutably establish asexual and sexual morph connections, for example, the *Mycosphaerella arbuticola* / *Septoria unedonis*, *Phaeosphaeria papaya* / *Phaeoseptoria oryzae*, *Leptosphaeria maculans* / *Plenodomus lingam*, and also provide phylogenetic placements for asexual taxa within the modern taxonomic classification schemes (Quaedvlieg et al. 2013; Ariyawansa et al. 2015b, Crous et al. 2015b). In this study, sequence data together with morphology are used to delimit and propose several new species and one new genus in *Phaeosphaeriaceae* and link the sexual and asexual morph of *Nodulosphaeria*.

The observations presented in this study show that the LSU and ITS sequence data can be used to distinguish most genera in *Phaeosphaeriaceae* such as *Parastagonospora* and *Nodulosphaeria*. Nevertheless, the circumscription of *Septoriella* is still unclear, and our phylogenetic analysis (Fig. 1) suggests that *Septoriella leuchtmanii* should be excluded from the genus. Future studies should use more isolates and gene regions to resolve the genetic concepts of *Septoriella*.

Acknowledgments

The authors extend their sincere appreciations to the Deanship of Scientific Research at King Saud University for its funding this Prolific Research group (PRG-1436–09). We would like to thank Humidtropics, a CGIAR Research Program that aims to develop new opportunities for improved livelihoods in a sustainable environment, for partially funding this work. KD Hyde thanks The Chinese Academy of Sciences, project number 2013T2S0030, for the award of Visiting Professorship for Senior International Scientists at Kunming Institute of Botany. The authors also thank MFU Dothideomycetes grant for supporting this study. The authors would like to thank Shaun Pennycook for assistance in checking the name of new taxa.

References

- Ariyawansa HA, Hawksworth DL, Hyde KD, Jones EBG, Maharachchikumbura SSN, Camporesi E, Manamgoda DS, Thambugala KM, Udayanga D, Daranagama A, Jayawardena R, Liu JK, McKenzie EHC, Phookamsak R, Senanayake IC, Shivas RG, Tian Q, X JC. 2014 – Epitypification and neotypification: guidelines with appropriate and inappropriate examples. *Fungal Diversity* 69, 57–91. <http://dx.doi.org/10.1007/s13225-014-0315-4>
- Ariyawansa HA, Hyde KD, Jayasiri SC, Buyck B, Chethana KWT, Cui YY, Dai DQ, Dai YC, Daranagama DA, Jayawardena RS, Lücking R, Ghobad-Nejhad M, Niskanen T, Thambugala KM, Voigt K, Zhao RL, Boonmee S, Bahkali AH, Chen J, Cui BK, Dayarathne MC, Dissanayake AJ, Ekanayaka AH, Hashimoto A, Hongsanan S, Jones EBG, Larsson E, Lewis D, Li WJ, Li QR, Liu JK, Luo ZL, Maharachchikumbura SSN, Mapook A, McKenzie EHC, Norphanphoun C, Pang KL, Perera RH, Phookamsak R, Phukhamsakda C, Randrianjohany E, Senanayake IC, Singtripop C, Shang QJ, Tanaka K, Tian Q, Tian CM, Tibpromma S, Verbeken A, Abdel-Wahab MA, Wanasinghe D, Wijayawardene NN, Zhang JF, Zhang H, Abdel-Aziz FA, Adamčík S, Ammirati JF, Bulgakov T, Cabral AL, Callaghan TM, Callac P, Chang CH, Coca LF, Dal-Forno M, Dollhofer V, Fliegerová K, Greiner K, Griffith GW, Ho HM, Hofstetter V, Jeewon R, Kang JC, Kirk PM, Kytövuori I, Lawrey JD, Li XH, Liu ZY, Liu XZ, Liimatainen K, Lumbsch HT, Matumura M, Moncada B, Nuankaew S, Parnmen S, de A. Santiago M, Sato G, Sommai S, Song Y, de Souza CAF, de Souza-Motta CM, Su HY, Suetrong S, Wang Y, Wei SF, Wen TC, Yuan HS, Zhou LW, Reblova M, Fournier J, Camporesi E. 2015c – Fungal Diversity Notes 111–252 – Taxonomic and phylogenetic contributions to fungal taxa. *Fungal Diversity* (In press) <http://dx.doi.org/10.1007/s13225-015-0346-5>
- Ariyawansa HA, Thambugala KM, Manamgoda DS, Jayawardena R, Camporesi E, Boonmee S, Wanasinghe DN, Phookamsak R, Hongsanan S, Singtripop C, Chukeatirote E, Kang JC, Jones EBG, Hyde KD. 2015a – Towards a natural classification and backbone tree for *Pleosporaceae*. *Fungal Diversity* 71, 85–139. <http://dx.doi.org/10.1007/s13225-015-0323-z>
- Ariyawansa HA, Phukhamsakda C, Thambugala KM, Bulgakov TS, Wanasinghe DN, Perera RH, Mapook A, Camporesi E, Kang JC, Jones EBG, Bahkali AH, Jayasiri SC, Hyde KD, Liu ZY, Bhat JD. 2015b – Revision and phylogeny of *Leptosphaeriaceae*. *Fungal Diversity* 75, (online). <http://dx.doi.org/10.1007/s13225-015-0349-2>

- Arzanlou M, Crous PW. 2006 – *Phaeosphaeriopsis musae*. Fungal Planet 9. CBS- KNAW Fungal Biodiversity Centre, Utrecht, The Netherlands.
- Barr ME 1979 – A classification of Loculoascomycetes. *Mycologia* 71, 935–957
- Chomnunti P, Hongsanan S, Aguirre-Hudson B, Tian Q, Peršoh D, Dhami MK, Alias AS, Xu JC, Liu XZ, Stadler M, Hyde KD 2014) The sooty moulds. *Fungal Diversity* 66,1–36. <http://dx.doi.org/10.1007/s13225-014-0278-5>
- Crous PW, Slippers B, Wingfield MJ, Rheeder J, Marasas WFO, Phillips AJL, Alves A, Burgess T, Barber P, Groenewald JZ 2006 – Phylogenetic lineages in the *Botryosphaeriaceae*. *Studies in Mycology* 55, 235– 253. <http://dx.doi.org/10.3767/003158515X685841>
- Crous PW, Carris LM, Giraldo A, Groenewald JZ, Hawksworth DL, Hernández-Restrepo M, Jaklitsch WM, Lebrun MH, Schumacher RK, Stielow JB, van der Linde EJ, Vilcāne J, Voglmayr H, Wood AR. 2015b – The Genera of Fungi - fixing the application of the type species of generic names – G 2: *Allantophomopsis*, *Latorua*, *Macrodiplodiopsis*, *Macrohilum*, *Milospium*, *Protostegia*, *Pyricularia*, *Robillarda*, *Rotula*, *Septoriella*, *Torula* and *Wojnowicia*. *IMA Fungus* 6,163–198. <http://dx.doi.org/10.5598/imafungus.2015.06.01.11>.
- Crous PW, Wingfield MJ, Guarro J, Hernández-Restrepo M, Sutton DA, Acharya K, Barber PA, Boekhout T, Dimitrov RA, Dueñas M, Dutta AK, Gené J, Gouliamova DE, Groenewald M, Lombard L, Morozova OV, Sarkar J, Smith MT, Stchigel AM, Wiederhold NP, Alexandrova AV, Antelmi I, Armengol J, Barnes I, Cano-Lira JF, Castañeda Ruiz RF, Contu M, Courtecuisse PrR, da Silveira AL, Decock CA, de Goes A, Edathodu J, Ercole E, Firmino AC, Fourie A, Fournier J, Furtado EL, Geering ADW, Gershenzon J, Giraldo A, Gramaje D, Hammerbacher A, He XL, Haryadi D, Khemmuk W, Kovalenko AE, Krawczynski R, Laich F, Lechat C, Lopes UP, Madrid H, Malysheva EF, Marín-Felix Y, Martín MP, Mostert L, Nigro F, Pereira OL, Picillo B, Pinho DB, Popov ES, Rodas Peláez CA, Rooney-Latham S, Sandoval-Denis M, Shivas RG, Silva V, Stoilova-Disheva MM, Telleria MT, Ullah C, Unsicker SB, van der Merwe NA, Vizzini A, Wagner HG, Wong PTW, Wood AR, Groenewald JZ. 2015a – Fungal Planet description sheets: 320–370. *Persoonia* 34, 167–266. <http://dx.doi.org/10.3767/003158515X688433>
- De Gruyter J, Woudenberg JHC, Aveskamp MM, Verkley GJM, Groenewald JZ, Crous PW 2010 – Systematic reappraisal of species in *Phoma* section *Paraphoma*, *Pyrenochaeta* & *Pleurophoma*. *Mycologia* 102, 1066–1081. <http://dx.doi.org/10.3852/09-240>
- Farr DF, Bills GF. 1995 – *Wojnowicia colluvium* sp. nov. isolated from conifer litter. *Mycologia* 87(4), 518–524.
- Hall TA 1999 – BioEdit: a user-friendly biological sequence alignment editor and analysis program for Windows 95/98/NT. In: *Nucleic Acids Symposium Series*, pp 95–98
- Hyde KD, Jones EBG, Liu JK, Ariyawansa H, Boehm E, Boonmee S, Braun U, Chomnunti P, Crous PW, Dai DQ, Diederich P, Dissanayake A, Doilom M, Doveri F, Hongsanan S, Jayawardena R, Lawrey JD, Li YM, Liu YX, Lücking R, Monka J, Muggia L, Nelsen MP, Pang KL, Phookamsak R, Senanayake IC, Shearer CA, Suetrong S, Tanaka K, Thambugala KM, Wijayawardene NN, Wikee S, Wu HX, Zhang Y, Begoña AH, Alias SA, Aptroot A, Bahkali AH, Bezerra JL, Bhat DJ, Camporesi E, Chuksa E, Gueidan C, Hawksworth DL, Hirayama K, Hoog SD, Kang JK, Knudsen K, Li WJ, Li XH, Liu ZY, 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 JC, Yacharoen S, Yan JY, Zang M 2013 – Families of Dothideomycetes. *Fungal Diversity* 63, 1–313. <http://dx.doi.org/10.1007/s13225-013-0263-4>
- Hyde KD, McKenzie EHC, KoKo TW 2011 – Towards incorporating anamorphic fungi in a natural classification – checklist and notes for 2010. *Mycosphere* 2, 1–88.

<http://dx.doi.org/10.5943/mycosphere/3/2/5>

Index Fungorum (2015) <http://www.indexfungorum.org/names/names.asp>

Jayasiri SC, Hyde KD, Ariyawansa HA, Bhat DJ, Buyck B, Cai L, Dai YC, Abd-Elsalam KA, Ertz D, Gibertoni TB, Hidayat I, Jeewon R, Jones EBG, Karunarathna SC, Kirk P, Li WJ, Liu JK, Luangsa-ard JJ, Maharachchikumbura SSN, Manamgoda DS, McKenzie EHC, Moncalvo JM, Ghobad-Nejhad M, Nilsson H, Pang KL, Pereira OL, Raspé O, Rollins AW, Romero AI, Salazar JAE, Stephenson S, Suetrong S, Taylor JE, Tsui CKM, Vizzini A, Abdel-Wahab MA, Wen TC, Boonmee S, Dai DQ, Daranagama DA, Dissanayake AJ, Ekanayaka AH, Hongsanan S, Jayawardena RS, Perera RH, Phookamsak R, De Silva NI, Thambugala KM, Tian Q, Wanasinghe DN, Wijayawardene NN, Zhao RL, Kang JC, Promputtha I 2015 – The Faces of Fungi database-A unique perspective: Fungal names linked with morphology, phylogeny and human impacts. *Fungal Diversity* (in press). <http://dx.doi.org/10.1007/s13225-015-03518-8>

Kirk PM, Cannon PF, Minter DW 2010 – *Dictionary of the Fungi* 10th ed.-CABI (2008)2.

Li WJ, Liu JK, Bhat DJ, Camporesi E, Dai DQ, Mortimer PE, Xu JC, Hyde KD, Chomnunti P 2015 – Molecular phylogenetic analysis reveals two new species of *Discosia*. *Phytotaxa* 203, 37–46. <http://dx.doi.org/10.11646/phytotaxa.181.1.1>

Liu JK, Hyde KD, Jones EBG, Ariyawansa HA, Bhat DJ, Boonmee Maharachchikumbura SSN, McKenzie EHC, Phookamsak R, Phukhamsakda C, Shenoy BD, Abdel-Wahab MA, Buyck B, Chen J, Chethana KWT, Singtripop C, Dai DQ, Dai YC, Daranagama DA, Dissanayake AJ, Doliom M, D'Souza MJ, Fan XL, Goonasekara ID, Hirayama K, Hongsanan S, Jayasiri SC, Jayawardena RS, Karunarathna SC, Li WJ, Mapook A, Norphanphoun C, PANG KL, Perera RH, Peršoh D, Pinruan U, Senanayake IC, Somrithipol S, Suetrong S, Tanaka K, Thambugala KM, Tian Q, Tibpromma S, Udayanga D, Wijayawardene NN, Wanasinghe D, Wisitrassameewong K, Abdel-Aziz FA, Adamčík S, Bahkali AH, Boonyuen N, Bulgakov T, Callac P, Chomnunti p, Greiner K, Hashimoto A, Hofstetter V, Kang JC, Lewis D, Li XH, Liu XX, Liu ZY, Matumura M, Mortimer PE, Rambold G, Randrianjohany E, Sato G, Sri-Indrasutdhi V, Tian CM, Verbeken A, Von Brackel W, Wang Y, Wen TC, Xu JC, Yan JY, Zhao RL, Camporesi E. 2015 – Fungal diversity notes 1–110: taxonomic and phylogenetic contributions to fungal species. *Fungal Diversity* 72 (1), 1–197. <http://dx.doi.org/10.1007/s13225-015-0324-y>

Page RDM 1996 – TreeView: an application to display phylogenetic trees on personal computers. *Computer Applications in the Biosciences* 12, 357–358.

Phookamsak R, Liu JK, McKenzie EHC, Manamgoda DS, Ariyawansa H, Thambugala KM, Dai DQ, Camporesi E, Chukeatirote E, Wijayawardene NN, Bahkali AH, Mortimer PE, Xu JC, Hyde KD. 2014 – Revision of *Phaeosphaeriaceae*. *Fungal Diversity* 68, 159–238. <http://dx.doi.org/10.1007/s13225-014-0308-3>

Phookamsak R, Norphanphoun C, Tanaka K, Dai DQ, Luo ZL, Liu JK, Su HY, Bhat DJ, Bahkali Ah, Mortimer PE, Xu JC, Hyde 2015 – Towards a natural classification of *Astrosphaeriella*-like species; introducing *Astrosphaeriellaceae* and *Pseudoastrosphaeriellaceae* fam. nov. and *Astrosphaeriellopsis*, gen. nov. 2015 – *Fungal Diversity* (in press)

Quaedvlieg W, Verkley GJM, Shin H-D, Barreto RW, Alfenas AC, Swart WJ, Groenewald JZ, Crous PW 2013 – Sizing up *Septoria*. *Studies in Mycology* 75, 307–390. <http://dx.doi.org/10.3114/sim0017>.

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, Rivas Plata E, 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. <http://dx.doi.org/10.3114/sim.2009.64.01>
- Silvestro D, Michalak I. 2011 – raxmlGUI: a graphical front-end for RAxML. *Organisms Diversity and Evolution* 12: 335–337. <http://dx.doi.org/10.1007/s13127-011-0056-0>
- Sutton BC 1980 – *The Coelomycetes-Fungi imperfecti with pycnidia, acervuli and stromata*. Commonwealth Mycological Institute, Kew, UK, 496 pp.
- Thompson JD, Gibson TJ, Plewniak F, Jeanmougin F, Higgins DG 1997 – The CLUSTAL_X windows interface: flexible strategies for multiple sequence alignment aided by quality analysis tools. *Nucleic Acids Research* 25(24), 4876–4882.
- Trakunyingcharoen T, Lombard L, Groenewald JZ, Cheewangkoon R, Toanun C, Alfenas AC, Pedro W Crous. 2014 – Mycoparasitic species of *Sphaerellopsis*, and allied lichenicolous and other genera. *IMA Fungus* 5(2), 391–414. <http://dx.doi.org/10.5598/imafungus.2014.05.02.05>
- Vilgalys R, Hester M. 1990 – Rapid genetic identification and mapping of enzymatically amplified ribosomal DNA from several *Cryptococcus* species. *Journal of Bacteriology* 172:4238–4246.
- Wanasinghe DN, Jones EBG, Camporesi E, Boonmee S, Karunaratna SC, Thines M, Mortimer PE, Xu JC, Hyde KD 2014 – *Dematiopleospora mariaegen*. sp. nov., from *Ononis spinosa* in Italy. *Cryptogamie, Mycologie* 35 (2), 105–117. <http://dx.doi.org/10.7872/crym.v35.iss2.2014.105>
- White TJ, Bruns 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, San Diego, pp 315–322
- Wijayawardene DNN, Crous PW, Kirk PM, Hawksworth DL, Boonmee S, Braun U, Dai DQ, D'souza MJ, Diederich P, Dissanayake A, Doilom M, Hongsanan S, Jones EBG, Johannes Z, Groenewald, Jayawardena R, Lawrey JD, Liu JK, Lücking R, Madrid H, Manamgoda DS, Muggia L, Nelsen MP, Phookamsak R, Suetrong S, Tanaka K, Thambugala KM, Wanasinghe DN, Wikee S, Zhang Y, Aptroot A, Ariyawansa HA, Bahkali AH, Bhat DJ, Gueidan C, Chomnunti P, De Hoog GS, Knudsen K, Li WJ, McKenzie EHC, Miller AN, Phillips AJL, Piątek M, Raja HA, Shivas RS, Slippers B, Taylor JE, Tian Q, Wang Y, Woudenberg JHC, Cai L, Jaklitsch WM, Hyde KD 2014 – Naming and outline of Dothideomycetes – 2014 including proposals for the protection or suppression of generic names. *Fungal Diversity* 69, 1–55. <http://dx.doi.org/10.1007/s13225-014-0309-2>
- Wijayawardene DNN, McKenzie EHC, Hyde KD. 2012b – Towards incorporating anamorphic fungi in a natural classification–checklist and notes for 2011. *Mycosphere* 3(2): 157–228. <http://dx.doi.org/10.5943/mycosphere/3/2/5>
- Wijayawardene DNN, Song Y, Bhat DJ, McKenzie EHC, Chukeatirote E, Wang Y, Hyde KD. 2013 – *Wojnowicia viburni* sp. nov., from China and its phylogenetic placement. *Sydowia* 65:129–138.
- Zhang Y, Crous PW, Schoch CL, Hyde KD. 2012 – Pleosporales. *Fungal Diversity* 53:1–221. <http://dx.doi.org/10.1007/s13225-011-0117-x>
- Zhang Y, Schoch CL, Fournier J, Crous PW, de Gruyter J, Woudenberg JHC, Hirayama K, Tanaka K, Pointing SB, Spatafora JW, Hyde KD 2009 – Multi-locus phylogeny of Pleosporales: a taxonomic, ecological and evolutionary re-evaluation. *Studies in Mycology* 64:85–102. <http://dx.doi.org/10.3114/sim.2009.64.04>