

Families of *Diaporthales* based on morphological and phylogenetic evidence

I.C. Senanayake^{1,2,3}, P.W. Crous⁴, J.Z. Groenewald⁴, S.S.N. Maharachchikumbura⁵, R. Jeewon⁶, A.J.L. Phillips⁷, J.D. Bhat^{8,9}, R.H. Perera³, Q.R. Li¹⁰, W.J. Li^{1,2,3}, N. Tangthirasunun^{11,12}, C. Norphanphoun³, S.C. Karunarathna^{1,2*}, E. Camporesi^{13,14,15}, I.S. Manawasighe¹⁶, A.M. Al-Sadi⁵, and K.D. Hyde^{1,2,3}

¹Key Laboratory for Plant Diversity and Biogeography of East Asia, Kunming Institute of Botany, Chinese Academy of Science, Kunming 650201, Yunnan, China; ²East and Central Asia, World Agroforestry Centre, Kunming 650201, Yunnan, China; ³Center of Excellence for Fungal Research, Mae Fah Luang University, Chiang Rai, Thailand; ⁴Westerdijk Fungal Biodiversity Institute, Uppsalalaan 8, 3584 CT Utrecht, The Netherlands; ⁵Department of Crop Sciences, College of Agricultural and Marine Sciences, Sultan Qaboos University, P.O. Box 34, Al-Khod 123, Omari, ⁶Department of Health Sciences, Faculty of Science, University of Mauritius, Reduit, 80837, Mauritius; ⁷Faculty of Sciences, Biosystems and Integrative Sciences Institute (BioISI), University of Lisbon, Campo Grande, 1749-016 Lisbon, Portugal; ⁸Department of Botany, Goa University, Goa 403 206, India; ⁹No. 128/1-J, Azad Housing Society, Curca, P.O. Goa Velha 403108, India; ¹⁰Engineering Research Center of Southwest Bio-Pharmaceutical Resources, Ministry of Education, Guizhou University, Guiyang, Guizhou 550025, China; ¹¹Univ Paris Diderot, Sorbonne Paris Cité, Institut des Energies de Demain (IED), Paris 75205, France; ¹²Univ Paris Sud, Institut de Génétique et Microbiologie, UMR8621, Orsay 91405, France; ¹³A.M.B. Gruppo Micologico Forlivese, Antonio Cicognani, Via Roma 18, Forli, Italy; ¹⁴A.M.B. Circolo Micologico, Giovanni Carini, 314 Brescia, Italy; ¹⁵Società per gliStudiNaturalisticidella Romagna, 144 Bagnacavallo, RA, Italy; ¹⁶Institute of Plant and Environment Protection, Beijing Academy of Agriculture and Forestry Sciences, No. 9 of ShuGuangHuaYuanZhongLu, Haidian District, Beijing 100097, China

*Correspondence: S.C. Karunarathna, samanthakarunarathna@gmail.com

Abstract: Diaporthales is an important ascomycetous order comprising phytopathogenic, saprobic, and endophytic fungi, but interfamilial taxonomic relationships are still ambiguous. Despite its cosmopolitan distribution and high diversity with distinctive morphologies, this order has received relativelyiaceae, *Macrohilaceae*, *Melanconidaceae*, *Pseudoplagiostomaceae*, *Schizoparmaceae*, *Stilbosporaceae* and *Sydowiellaceae*. Taxonomic uncertainties among genera are also clarified and recurrent discrepancies in the taxonomic position of families within the *Diaporthales* are discussed. An updated outline and key to families and genera of the order is presented.

Key words: Multi-gene DNA phylogeny, New taxonomic arrangement, Phytopathogenic fungi, Sordariomycetes, Systematics.

Taxonomic novelties: New families: Apiosporopsidaceae Senan. Maharachch. & K.D. Hyde, Apoharknessiaceae Senan. Maharachch. & K.D. Hyde, Asterosporiaceae Senan. Maharachch. & K.D. Hyde, Auratiopycnidiellaceae Senan. Maharachch. & K.D. Hyde, Erythrogloeaceae Senan. Maharachch. & K.D. Hyde, Melanconiellaceae Senan. Maharachch. & K.D. Hyde, Prosopidicolaceae Senan. & K.D. Hyde, New genera: Marsupiomyces Senan. & K.D. Hyde, Microascospora Senan., Camporesi & K.D. Hyde, Phaeoappendicospora Senan., Q.R. Li & K.D. Hyde, Paradiaporthe Senan., Camporesi, & K.D. Hyde, Hyaliappendispora Senan., Camporesi & K.D. Hyde, Chiangraiomyces Senan. & K.D. Hyde; New species: Chiangraiomyces bauhiniae Senan. & K.D. Hyde, Coniella pseudokoreana Senan., Tangthir. & K.D. Hyde, Cytospora centrivillosa Senan., Camporesi & K.D. Hyde, Cytospora rosae Senan., Camporesi & K.D. Hyde, Gnomoniopsis agrimoniae Senan., Camporesi & K.D. Hyde, Hyaliappendispora galii Senan., Camporesi & K.D. Hyde, Marsupiomyces epidermoidea R.H. Perera, Senan., Bulgakov & K.D. Hyde, Marsupiomyces quercina Senan., Camporesi & K.D. Hyde, Melanconis italica Senan., Camporesi & K.D. Hyde, Microascospora rubi Senan., Camporesi & K.D. Hyde, Phaeoappendicospora thailandensis Senan., Q.R. Li & K.D. Hyde, Plagiostoma jonesii Senan., & K.D. Hyde, Phaeoappendicospora thailandensis Senan., Q.R. Li & K.D. Hyde, Plagiostoma jonesii & K.D. Hyde, Phaeoappendicospora thailandensis Senan., Q.R. Li & K.D. Hyde, Plagiostoma jonesii & K.D. Hyde, Phaeoappendicospora thailandensis Senan., Q.R. Li & K.D. Hyde, Plagiostoma jonesii Senan., & K.D. Hyde, Plagiostoma salicicola Senan., Camporesi & K.D. Hyde, Microascospora rubi Senan., Camporesi & K.D. Hyde, Plagiostoma salicicola Senan., Camporesi & K.D. Hyde, Phaeoappendicospora thailandensis Senan., Q.R. Li & K.D. Hyde, Plagiostoma jonesii Senan., & K.D. Hyde, Plagiostoma salicicola Senan., Camporesi &

Available online 1 August 2017; http://dx.doi.org/10.1016/j.simyco.2017.07.003.

INTRODUCTION

The *Diaporthales* is a distinct order in the subclass *Diaporthomycetidae* (*Sordariomycetes*) and it includes pathogens, saprobes and endophytes, with no known coprophilous, hypersaprobes or mycophylic species (Barr 1978, Rossman *et al.* 2007, Vasilyeva *et al.* 2007, Maharachchikumbura *et al.* 2015, 2016). Taxa of this order inhabit a wide diversity of hosts and substrates, including most economically and ecologically important trees and crops, soil and living animal and human tissues (Barr 1978, Gryzenhout *et al.* 2006c). Species in *Diaporthales* form solitary or aggregated, immersed to erumpent, rarely superficial, orange, brown to black

perithecial ascomata, with short or long necks, that are located in stromatic tissues or substrates, with a centrum (or hamathecium) lacking or with few paraphyses (Alexopoulos & Mims 1978, Barr 1978, Castlebury *et al.* 2002). Asci are unitunicate with a conspicuous refractive ring (Hawksworth *et al.* 1995, Rossman *et al.* 2007). Ascospore morphology is diverse, ranging from short to elongate and aseptate or septate with hyaline or pigmented walls. The asexual morphs of *Diaporthales* are generally coelomycetous (Rossman *et al.* 2007), producing acervuli or pycnidial conidiomata, with or without a well-developed stroma. Conidiogenesis is phialidic or rarely annellidic and conidia are usually unicellular or 1-septate (Rossman *et al.* 2007).

Peer review under responsibility of Westerdijk Fungal Biodiversity Institute.

^{© 2017} Westerdijk Fungal Biodiversity Institute. Production and hosting by ELSEVIER B.V. This is an open access article under the CC BY-NC-ND license (http://creativecommons.org/licenses/by-nc-nd/4.0/).

Fungal taxa placed in "Diaporthaceae sensu lato" were divided into two groups (von Höhnel 1917) as "Eu-Diaportheen", to accommodate genera without allantoid ascospores and "Valseen" to accommodate genera with allantoid ascospores. Nannfeldt (1932) introduced the order Diaporthales to accommodate von Höhnel's Eu-Diaportheen group. Luttrell (1951) described Diaporthales as an order comprising species that have a "Diaporthe-type centrum" and "Endothia-type ascus". Chadefaud (1960) analysed characters of stromatic tissues in diaporthoid taxa and recognised families as Diaporthaceae or Cytosporaceae (= Valsaceae), Melanconidaceae and Gnomoniaceae. Wehmeyer (1975) classified the Diaporthales to include three families: Diaporthaceae, Gnomoniaceae and Cytosporaceae. Barr (1978) revised the order Diaporthales accepting Gnomoniaceae and Cytosporaceae in the suborder Gnomoniineae. Melanconidaceae and Pseudovalsaceae were accommodated in the suborder Melanconidineae. To differentiate genera, Barr (1978) used characters such as presence or absence of stromata, stromatic development and tissue types, the position of the perithecia and perithecial necks relative to the substrate, as well as ascospore shape; and Monod (1983) distinguished genera within Gnomoniaceae based on characters of the stromatic tissues, asexual morphs and ascospores. Three families were recognised in Diaporthales by Eriksson (2001), including Cytosporaceae, Melanconidaceae and Vialaeaceae. Based on analysis of LSU nrDNA sequence data, Castlebury et al. (2002) accepted Diaporthaceae, Gnomoniaceae, Melanconidaceae and Cytosporaceae in Diaporthales. Gnomoniaceae was revised by several recent studies and new taxa were introduced (Sogonov et al. 2008, Walker et al. 2010, 2012, Mejía1 et al. 2011). Castlebury et al. (2002) did not confirm Vialaeaceae as a family in Diaporthales and therefore excluded it from Diaporthales. Réblová et al. (2004) introduced Togniniaceae to this order based on small subunit (SSU) nrDNA; however, Mostert et al. (2006) concluded that its placement was ambiguous based on large subunit (LSU) nrDNA. Maharachchikumbura et al. (2015) excluded Togniniaceae from Diaporthales and accommodated it in Togniniales. Gryzenhout et al. (2006c) introduced the Cryphonectria-Endothia complex as the family Cryphonectriaceae. Sydowiellaceae and the Schizoparme-Pilidiella complex with the genus Coniella were introduced as Schizoparmaceae in Diaporthales (Rossman et al. 2007, Alvarez et al. 2016). Harknessiaceae was introduced into Diaporthales accommodating Harknessia with wuestneia-like sexual morphs (Crous et al. Pseudoplagiostomaceae was introduced 2012b). by Cheewangkoon et al. (2010) to accommodate Pseudoplagiostoma. Voglmayr & Jaklitsch (2014) resurrected Stilbosporaceae in Diaporthales based on phylogenetic analysis of LSU nrDNA sequence data and transferred the genera Stegonsporium and Stilbospora to this family. Macrohilaceae was introduced by Crous et al. (2015), based on an analysis of LSU nrDNA to accommodate Macrohilum. Suetrong et al. (2015) introduced Tirisporellaceae into Diaporthales; however, Jones et al. (2015) excluded this family from Diaporthales. Norphanphoun et al. (2016) introduced Lamproconiaceae to accommodate Lamproconium and Hercospora. Juglanconidaceae was introduced in the Diaporthales by Voglmayr et al. (2017). However, molecular data suggest that additional families still remain to be elucidated

(Gryzenhout *et al.* 2006c, Crous *et al.* 2012a, 2015, Voglmayr *et al.* 2017). Currently there are 14 families accepted in the *Diaporthales*.

Given the taxonomic discrepancies within *Diaporthales*, the present study uses a combined taxonomic approach based on morphology and DNA sequence analyses of the partial 28S nrDNA (LSU), the internal transcribed spacer regions and intervening 5.8S nrDNA (ITS), DNA-directed RNA polymerase II second largest subunit (*rpb2*), and translation elongation factor 1-alpha (*tef1*) gene regions to investigate phylogenetic relationships of all genera in *Diaporthales* to update their classification. All taxonomic novelties and present taxonomic families are redescribed and illustrated where necessary. We also present new data on each family to provide a better taxonomic understanding.

MATERIALS AND METHODS

Isolates and specimens

Specimens were collected from Germany, Italy, Russia, Thailand and the UK. They were placed in paper bags and collection details noted. Specimens were brought to the laboratory in Ziplock plastic bags and examined with a Motic SMZ 168 stereomicroscope. Rehydrated fruiting bodies were used to observe morphological characteristics of ascomata, asci, ascospores and other tissues and characters were photographed with a Canon 550D digital camera fitted to the Nikon ECLIPSE 80i compound microscope. Photomicrographs were arranged with Adobe Photoshop v. CS6 and all measurements were made with Tarosoft v. 0.9.0.7. Specimens were preserved and are deposited at the BBH and MFLU fungaria. Taxonomic novelties and descriptions were deposited in MycoBank (Crous et al. 2004), and new species were established using modern criteria and standards (Taylor et al. 2000, Seifert & Rossman 2010, Jeewon & Hyde 2016).

Sporocarps were removed from the substrate using a sterilised needle and placed in a few drops of sterilised distilled water on a sterilised cavity slide and a spore suspension was prepared as described in Chomnunti *et al.* (2014). Germinating ascospores were aseptically transferred to Petri dishes containing Potato Dextrose Agar (PDA) or Malt Extract Agar (MEA) (Crous *et al.* 2009). Colonies were photographed and characters were noted. Colony colour on PDA and MEA was determined with the colour charts of Rayner (1970). Living cultures are deposited at Mae Fah Luang University (MFLU) and the Westerdijk Fungal Biodiversity Institute (CBS) culture collections. Autoclaved pine needles were placed on water agar (WA) to observe conidiomatal development and sporulating (Crous *et al.* 2009).

Types and other relevant authentic specimens were loaned from accessible fungaria [New York State Museum (NY), Naturhistorisches Museum Wien (W), Swedish Museum of Natural History (S), Royal Botanic Gardens, Kew (K), Universität Wien (WU)]. A small part of the fungarium specimen was cut and rehydrated in water or 5 % KOH. Micro-morphological characters were observed from rehydrated ascomata and photography was done as previously described. Download English Version:

https://daneshyari.com/en/article/5740983

Download Persian Version:

https://daneshyari.com/article/5740983

Daneshyari.com