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Impacts of the International Code of Nomenclature for Algae, Fungi, and Plants (Melbourne Code) on the Scientific Names of Plant Pathogenic Fungi

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Introduction

The largest group of plant pathogens belongs to the Kingdom Fungi, a group of organisms estimated to consist of 1.5 million to millions of species (6,27). Communication about fungal plant pathogens is primarily through the use of scientific names that appear in journals and books where the research of plant pathologists is published. These names encapsulate taxonomic (and recently phylogenetic) knowledge, and keeping the distinction between nomenclature (the formal naming of taxa) and taxonomy (the science of classifying organisms) is fraught with challenges. Our human attempts to create coherent nomenclatural units that are unambiguous and universal for taxa that are variable and evolving will run inexorably into inherent logical and semantic problems (31). An important point to remember is that the purpose of a name is to act as a symbol for communication. Scientific names are abstract concepts that help us to talk about the diversity we encounter in the natural world, used to make reliable generalizations about groups of related organisms, and for information storage and retrieval. Names are excellent tools, but as we continue to refine and revise their application, it is essential to remember Hey's observations that, "we tend not to notice that our categories lie largely within us ..." and that, "... utility, and not ontological standing, is the key to our appreciation of higher taxa" (31). Modern taxonomists now include many biochemical, ecological, physiological, and molecular characters to delineate new species and genera, which provides more insight into evolutionary history, but often requires a reassessment of nomenclature. Despite the desire for stability, disagreements about species and genus delimitation will persist. This means that scientific names will continue to change, and frustration by the users of names will also continue.

Historically, rules that govern how fungi are named are provided in the international codes of botanical nomenclature, which are revised at each International Botanical Congress (held every six years). Recently, four major changes in the rules that affect the naming of fungi were made at the XVIII International Botanical Congress held in

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[taxon.igmp.net](#). The Melbourne Code has significant implications for the scientific names of fungi especially those that cause plant diseases and therefore are of concern to plant pathologists. In this article, we introduce the major changes in fungal nomenclature. We also explain the impacts of these changes in fungal names by giving examples of plant-associated fungi. Finally, we discuss possible approaches on how to deal with fungal names in this transitional period.

Major Changes in Melbourne Code

(i) Starting on January 1, 2012, electronic publication is permitted of new scientific names in Portable Document Format (PDF) in online publications with an International Standard Serial Number (ISSN) or International Standard Book Number (ISBN) (Art. 29-31).

It is no longer necessary for new names to appear in printed matter in order to be effectively published. However, publication of names at public meetings, in collections or gardens open to the public, or by the issue of microfilm made from manuscripts is not considered effective publication (48).

(ii) Starting on January 1, 2012, English may be used as an alternative to Latin for the descriptions or diagnoses of new taxa (Art. 39).

Changes (i) and (ii) apply to plants and algae in addition to fungi (48).

(iii) Starting on January 1, 2013, all new fungal names, including new taxa, new combinations, names at new ranks, and replacement names, must have an identifier issued by a recognized repository (Art. 42).

This new requirement took effect on January 1, 2013, after which new fungal names published without an identifier from a recognized repository are not considered to be validly published. In the past ten years, an increasing number of mycologists have been using MycoBank (www.mycobank.org) to register new fungal names. A unique number (e.g. MB 802972) is issued by MycoBank for each registered fungal name, which serves as an identifier to be cited in the publication where the name is proposed. Recently it was decided that three repositories may serve as official repositories of fungal names, i.e., MycoBank, Fungal Names (fungalinfo.im.ac.cn/fungalname/fungalname.html) and Index Fungorum (www.indexfungorum.org) (59).

(iv) Starting on January 1, 2013, the dual naming system for fungi is replaced with one scientific name for each species based on priority.

In previous International Codes of Botanical Nomenclature, Article 59 permitted non-lichen forming fungi to apply different names to different states (pleomorphic fungi). The Sydney Code approved in 1981 introduced specialized terms for the different

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preferentially (20). These practices for fungi are now discontinued. According to the Melbourne Code, all legitimate fungal names published prior to January 1, 2013 compete equally for priority, and the sole correct name is now the earliest legitimate name, regardless of the life history state of the type (48). However, in order to maintain nomenclatural stability in pleomorphic fungi, Article 57.2 states that "... in cases where, prior to 1 January 2013, both ... names were widely used for a taxon, an asexual state-typified name that has priority is not to displace the sexual name(s) unless and until a proposal to reject the former ... or to deal with the latter ... has been submitted and rejected" (48).

Changing to One Scientific Name for Each Species of Fungus

Moving to one scientific name for each species of fungus aligns the fungi with the other groups of organisms governed by codes of nomenclature including the International Code of Zoological Nomenclature (ICZN) and the International Code of Nomenclature of Bacteria. For no other group of organisms has two scientific names for one species been allowed except for the fossils and with the Melbourne Code that changed as well. Also in agreement with the other groups of organisms, the correct scientific name for fungi is based on priority, meaning that the first scientific name to be applied to a species is the name that should be used. The Melbourne Code covers algae, plants and all groups traditionally treated as fungi, i.e. the Kingdom Fungi, straminopiles such as the Peronosporales, and the Myxomycetes. Despite their relationship to the true Fungi, it was decided that the Microsporidia should continue to be governed by the ICZN (48).

Integrating the scientific names of fungi into one system allows all entities with clear phylogenetic relatedness within a genus to be called by the same generic name. Previously separate genera were used for the sexual and asexual states of a species. For example, the sexual state species *Cochliobolus miyabeanus* www.google.co.za/search?hl=en&q=Cochliobolus+miyabeanus produces an asexual state, called *Bipolaris oryzae*. To those not familiar with the taxonomic history of these fungi, it is not obvious that these names apply to the same species, nor is it apparent that all species described in *Cochliobolus* have an asexual state in *Bipolaris* or *Curvularia* (Fig. 1). As molecular phylogenetic studies increasingly reveal clear relationships among fungi, the use of different generic names for closely related i.e. congeneric species causes confusion. In some cases it results in the redundant description of new generic and species names when the relationship to named genera and species is already known. For example, when an asexually reproducing species with no known sexual state was discovered and determined to belong phylogenetically in the sexual state genus *Chrysosporthe*, the previous nomenclatural code (Vienna Code) required that a separate genus be described for this species even though its



a sexual state genus and vice versa thus eliminating the need, for example, for the genus *Chrysoporthella*.

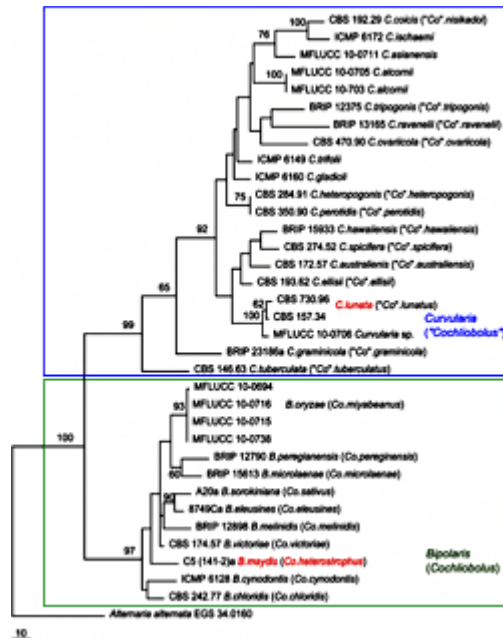


Fig. 1. Phylogram generated from the parsimony analysis based on combined genes of rDNA ITS and GPDH sequence data. Bootstrap values greater than 60 are shown. CBS 730.96 is the exepitype of *Curvularia lunata* that represents the generic type of *Curvularia*. The sequences of *Bipolaris maydis* C5 (141-2) were obtained from Berbee et al. 1999. The tree is rooted with *Alternaria alternata* EGS 34.0160.

For some groups of plant-associated fungi, there will essentially be no changes in scientific names because they do not have pleomorphic life cycles, or the alternative states were not named. These include the straminopiles, Blastocladiomycota, Chytridiomycota, Mucoromycotina, Entomophthoromycotina, Glomeromycota, and most members of the Basidiomycota with the exception of some rust fungi and *Rhizoctonia* relatives. Even among Ascomycota there are groups in which the scientific names will remain unchanged as for Taphrinomycotina, most of Pezizales, and even some of the pleomorphic ascomycetes. For example, the generic name *Alternaria* and most species in it were described long before any of the sexual state genera or species were established, thus the scientific names in *Alternaria* will remain unchanged (71). *Anisogramma anomala*, the eastern filbert blight pathogen, and *Cryphonectria parasitica*, the chestnut blight fungus, will not change names.

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geneticists refer to the causal agent of southern corn leaf blight by the name derived from the sexual state, *Cochliobolus heterostrophus*. Under previous codes of nomenclature this name was the correct name to refer to the complete species (holomorph). Plant pathologists who frequently encounter only the asexual state preferentially continue to prefer *Bipolaris maydis*. Although considerable disruption will result from changing scientific names for a familiar organism, the end result will be improved communication and a better appreciation for the whole fungus no matter how it manifests itself. In addition, these changes are already stimulating the broadening of research into fungi with little genetic and phylogenetic data, resulting in a clarification of evolutionary relationships. It is our fervent belief that, although this period of transition will be challenging, the end goal of a more stable and simplified naming system will benefit all fungal researchers working in various disciplines.

Following are examples of important plant pathogenic fungi for which integration of scientific names has been initiated. These represent different scenarios and demonstrate the range of issues that are being encountered in fulfilling this initiative.

Well-known genera and species with lists of accepted scientific names forthcoming: *Aspergillus*, *Penicillium* and *Talaromyces*

Aspergillus, *Penicillium* and *Talaromyces* (Eurotiomycetes, Eurotiales) are well-known mould genera in plant pathology, with hundreds of species in each genus, including many important plant pathogens, mycotoxin producers, industrial fungi, and human and animal pathogens. The International Commission on *Penicillium* and *Aspergillus*, a commission of the International Union of Microbiological Sciences, has sponsored a series of workshops and associated proceedings (63,64,65), culminating in a formal publication of a list of names in current use (54). The names *Aspergillus* and *Penicillium* both predate the names of the multiple sexual state genera associated with each group.

Aspergillus is important because of its age and because of the role that one of its species played in the origins of dual nomenclature, and has been the focus of several books [e.g., (2,25,45)]. *Aspergillus* was first described by Micheli in 1729 (49), making it one of the oldest named fungal genera. The distinctive morphology of the aspergillum (the asexual spore head) remains the principal diagnostic character. By 1854, De Bary (18) noticed that an *A. glaucus* mycelium produced sexual spores in a cleistothecium, which had been observed before and given its own name, *Eurotium herbararium* (Fig. 2). When De Bary realized that the two forms were different reproductive phases of the same organism, *Aspergillus* became the poster child for the nomenclatural predicament that has bedeviled mycologists for over 150 years. Subsequently, many other species named in *Aspergillus* for their asexual states were connected ("linked") to

considerable bewilderment outside taxonomic circles, interfering with efficient information retrieval (1). Currently there are approximately 250 named species of *Aspergillus* (23), many documented in the series of monographs based on cultural and morphological characters by Thom & Church in 1926 (74), Thom & Raper in 1954 (75) and Raper & Fennell in 1965 (57), and a more recent of multigene driven revisions by Houbraken et al. (35) (66,77).

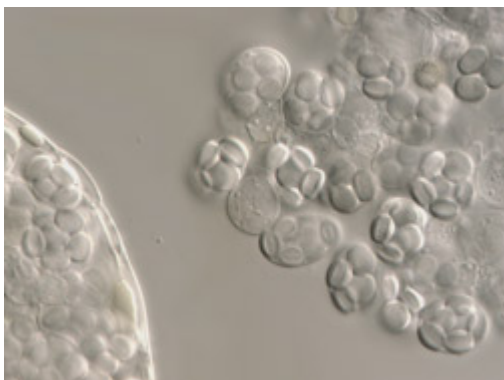


Fig. 2. *Eurotium herbariorum* asci: sexual state of *Aspergillus glaucus*.

Table 1. Synonyms and important species in *Aspergillus*, *Penicillium*, and *Talaromyces*.

| Single generic name | Alternate state synonyms | Other synonyms | Important species | | Reference |
|----------------------------|--------------------------|----------------|---------------------------------|----------------------------|---|
| <i>Aspergillus</i> 1729 | <i>Chaetosartorya</i> | | | | |
| | <i>Emericella</i> | | Genetic model | <i>A. nidulans</i> | Pontecorvo, 1956; Martinelli and Kinghorn, 1994 |
| | <i>Eurotium</i> | | Grain and food spoilage | <i>A. glaucus</i> complex | Bennett and Klich, 2003 |
| | <i>Fennellia</i> | | | | |
| | <i>Hemicarpentales</i> | Koenigstein | | | |
| | <i>Neocarpentales</i> | | | | |
| | <i>Neopetromyces</i> | | Plant pathogens, and mycotoxins | <i>A. flavus</i> relatives | |



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| | | | | Steinbach, 2008 |
| | <i>Petromyces</i> | | Mycotoxins | A. <i>ochraceus</i> Bennett and Klich, 2003 |
| | <i>Sclerocleista</i> | | | |
| | <i>Warcupiella</i> | | | |
| | | <i>Phialosimplex</i> | | |
| | | <i>Polypaecilum</i> | | |
| <i>Penicillium</i> | <i>Eupenicillium</i> | | Mycotoxins | many species |
| | | | Storage pathogens | <i>P. digitatum</i> and <i>P. italicum</i> (Citrus), <i>P. expansum</i> (pome fruit), <i>P. hirsutum</i> complex (root crops) |
| <i>Talaromyces</i> | <i>Penicillium</i> subgenus <i>Biverticillium</i> | | Human pathogen | <i>T. marneffe</i> |
| | | | Biocontrol of soilborne pathogens | <i>T. flavus</i> complex |

Because of the ecological and sexual state diversity, the best approach for developing a single name system for *Aspergillus* is controversial. The molecular and genomic data examined to date suggests that each of the former sexual genera is monophyletic, but the larger *Aspergillus* clade is also monophyletic, albeit with weak basal structure and with the inclusion of some aberrant asexual genera (*Phialosimplex*, *Polypaecilum*) among the basal clades (34,52). In an attempt to maintain broad usage of the oldest available name in *Aspergillus* for **all** of the species of economic significance, the

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system is adopted, but how stable it will be in the long term remains to be seen. The economic and historical importance of *Aspergillus* makes it likely that this taxon will remain at center stage in future discussions about nomenclature and mycological diversity.

Penicillium (Fig. 3) also has a long taxonomic history, since it was first described in 1809 by Link (43), named for its brush-like spore producing heads. Its historical association with named sexual genera is simpler, with two morphological distinct groups called *Eupenicillium* (with hard, sclerotium-like fruiting bodies with cell walls of angular cells) and *Talaromyces* (with soft fruiting bodies with an outer layer of hyphae). More than 250 species of *Penicillium* are now accepted, presented in series of monographs based on cultural and morphological characters by Thom in 1910 (73), Raper & Thom in 1949 (58), Pitt in 1979 (53) and Ramírez in 1982 (56), and a recent series of molecular revisions of individual sections and series by Samson et al. (67) and Houbraken et al. (36,37). In the sense of Pitt (53), *Penicillium* was determined to be polyphyletic (5), and the delimitation of new monophyletic generic concepts resulted in the adoption of the name *Penicillium* for the former subgenera *Aspergilloides*, *Furcatum*, and *Penicillium* (34). The sexual *Eupenicillium* is now considered a synonym of *Penicillium*; although the former name was used frequently in taxonomic literature, it was rarely applied in experimental or functional literature. The originally sexual genus *Talaromyces* was redefined to include many asexual species classified in *Penicillium* subgenus *Biverticillium* (68). The impact of these changes for plant pathologists is minimal, but the important human pathogen *Penicillium marneffe* is now classified in *Talaromyces*. Although the latter name change is unfortunate, a satisfactory single name system for *Penicillium* and *Talaromyces* is now in place (39).



Fig. 3. *Penicillium expansum*, the type species of *Penicillium*, on an apple.



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The asexual genus *Colletotrichum* contains many well-known plant pathogens that cause anthracnose and black spot diseases on economic crops and ornamental plants (12,38). Around 30 species of *Colletotrichum* have been linked to their sexual morphs assigned to the genus *Glomerella* (12). The International Subcommittee on the Taxonomy of *Colletotrichum* (ISTC) has been established and the inaugural meeting was held on 9 August 2012 in Beijing, China. All the ISTC members present at the meeting supported using the asexual name *Colletotrichum* over *Glomerella*. This decision was not only due to priority (16,78), but also because *Colletotrichum* is a more commonly used name in the applied sciences. Database searching done by the ISTC on 9 August 2012 showed generally 80% greater use of *Colletotrichum* over *Glomerella* in most mainstream search engines and databases such as Google, Google Scholar, Science Direct, and Scopus, and this trend has been historically consistent. In addition, fewer name changes would result by using *Colletotrichum* as the name for species in the genus. Although the genus *Vermicularia* 1790 is an earlier name for *Colletotrichum*, any move to establish *Vermicularia* as the genus name for this group would result in unbearable chaos for taxonomists and plant pathologists (12). The first meeting of the ISTC also reached an agreement that an "accepted list of names" will be provided in the light of recent reviews on this topic, such as Hyde et al. (38) "*Colletotrichum* – names in current use" and Cannon et al. (12) "*Colletotrichum* – current status and future directions." However, the criteria for making a "list of rejected names" need more consideration.

Sexual genus has priority with agreement on name changes: *Epichloë* 1865 vs. *Neotyphodium* 1996

The type species of the genus *Epichloë* was initially described by Persoon in 1798 (51) as the sexual species *Sphaeria typhina*. In 1849, Fries classified *Sphaeria typhina* in the genus *Cordyceps* subgenus *Epichloë* (22). The Tulasne brothers, in 1865 (76), elevated the subgenus *Epichloë* to generic status with *E. typhina* as the type species. From 1881 until 1982 the asexual state of the broadly defined *Epichloë typhina* was classified by Saccardo as *Sphacelia typhina* (62). However, Diehl indicated in 1950 (19) that the conidial fructifications of *Epichloë* were significantly different from those of *Claviceps* and did not apply the form-genus *Sphacelia* to the asexual state of *Epichloë*. Morgan-Jones and Gams reclassified this species in 1982 in the genus *Acremonium* as *A. typhinum* in their section *Albo-lanosa* (50). Based on their 1996 phylogenetic analysis, Glenn et al. (24) reclassified species of *Acremonium* in section *Albo-lanosa* in a new form-genus *Neotyphodium*. The type species of the genus *Neotyphodium* is *N. coenophialum* (24) based on *Acremonium coenophialum* (50) and includes *N. typhinum*.

In addition to their similarity in morphological and ecological characters, phylogenetic analyses of the sexual *Epichloë* species and the asexual *Neotyphodium* species show that these fungi form a monophyletic group and suggest that the asexual species with



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these fungi as a result of transfer of *Neotyphodium* species to *Epimeria*. Considerable efforts are already underway to make new combinations in *Epichloë*.

Asexual genus to be proposed for conservation, few name changes: *Bipolaris* 1959 vs *Cochliobolus* 1934

The genera *Bipolaris* and *Cochliobolus* are fungi with species that cause diseases of economically important grass crops. *Bipolaris* is typified by *B. maydis* (basionym *Helminthosporium maydis*) while *Cochliobolus* is typified by *C. heterostrophus* (basionym *Ophiobolus heterostrophus*). These type species are synonyms i.e. they are the asexual and sexual states of the same species (Fig. 1). This species is the cause of southern leaf blight of maize, a serious disease that occurs in temperate, warm temperate and tropical regions throughout the world (20,72).

At present 115 names have been included in *Bipolaris* while 54 names were described in *Cochliobolus* (40). Preliminary molecular phylogenetic studies suggest that some species described in *Bipolaris* and *Cochliobolus* actually belong in the closely related genus *Curvularia* (Fig. 1) (46). Both *Bipolaris* and *Curvularia* have sexual states that have been placed in *Cochliobolus* (Fig. 1). Although the sexual genus *Cochliobolus* 1934 is an older name than the asexual genus *Bipolaris* 1959, the asexual states are more commonly encountered in nature, thus the name *Bipolaris* has been used more frequently than *Cochliobolus*. In general the sexual states for these species were discovered and named after the asexual states. Thus, the generic name *Bipolaris* may be proposed for conservation or protection. The conservation of *Bipolaris* would result in only one name change; all other names of *Bipolaris* have priority. However, if priority is followed and *Cochliobolus* is retained for this genus, 46 names of *Bipolaris* would need to be transferred to *Cochliobolus* and seven names of *Bipolaris* that have priority would replace names currently in *Cochliobolus*.

Three species of this genus, specifically *Cochliobolus heterostrophus* (= *Bipolaris maydis*), *C. carbonum* (= *B. zeicola*), and *C. sativus* (= *B. sorokiniana*), have been studied extensively as model organisms. In most cases the publications concerning their genomics and genetics have referred to these fungi using the name in *Cochliobolus*, thus it is regrettable to change these important scientific names. However, two of these three species names would need to be changed even if the generic name *Cochliobolus* were used because the oldest epithet is currently placed in *Bipolaris*. For example *Cochliobolus sativus* based on *Ophiobolus sativus* 1929 would be replaced by the older name *Helminthosporium sorokinianum* 1890 transferred to *Cochliobolus*. Similarly *Cochliobolus carbonum* 1959 would be replaced by the older name *Helminthosporium zeicola* 1930.

Given the number of name changes required if *Cochliobolus* were used and the frequency with which *Bipolaris* is employed especially by plant pathologists, it seems

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Nectria 1849 vs. *Tubercularia* 1790

Since it was initially described over one hundred years ago, the sexual genus name *Nectria* was used for non-black, fleshy, uniloculate, perithecial ascomycetes. Booth (8) established one group within *Nectria* called the *N. cinnabarina* group that corresponded to the concept of *Nectria* in the narrow sense. This genus was further restricted to only 29 species by Hirooka et al. (33). Of the 1104 names described in *Nectria*, most have been placed in other genera, such as *Bionectria*, *Haematonectria*, *Lanatonectria*, *Leuconectria*, *Neonectria*, and *Sphaerostilbella*.

The type species of the genus *Nectria* is *N. cinnabarina* with its asexual state *Tubercularia vulgaris*, a species that is well known as the cause of coral spot of hardwood trees. Similarly the asexual genus *Tubercularia* is typified by *T. vulgaris* (Fig. 4), the asexual state of *N. cinnabarina*. Because these two names apply to the same species that are types of *Nectria* and *Tubercularia*, these genera are synonyms. Within the genus *Tubercularia* includes pale-colored sporodochial fungi with slimy aseptate conidia for which 247 names have been described. Seifert (60) examined the type specimens of many of these names and showed them belong in other genera. This genus has never been monographed.



Fig. 4. *Tubercularia vulgaris* on a twig, asexual form of *Nectria cinnabarina*.

The generic concepts of *Nectria* and *Tubercularia* have varied considerably over time but are now narrowly defined and still synonyms (32,33). The generic name *Tubercularia* was described earlier than *Nectria* and thus has priority and should be used when moving to one name for fungi. If the name *Tubercularia* were used, most of the 29 names in *Nectria* would require transfer to that genus. If the generic name *Nectria* were protected against *Tubercularia*, only three species would require name changes. In addition to the many fewer name changes, the name *Nectria* is more precisely defined and familiar to most plant pathologists. As a result it has been

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Working Committee of the International Commission for the Taxonomy of Fungi has been published (60).

Sexual genus proposed for conservation over a broadly defined asexual genus: *Neonectria* 1917 vs. *Cylindrocarpon* 1913

The genus *Cylindrocarpon* is based on the type species *C. cylindroides*. In the broad sense, *Cylindrocarpon* included 143 names for species having elongated, multi-septate conidia with broadly rounded ends. When Booth monographed this genus in 1966 (7), he demonstrated that many of these species have nectria-like sexual states. The sexual states of species of *Cylindrocarpon* were placed in the genus *Neonectria* by Rossman et al. (61). Using a multigene phylogenetic analysis of species representing the breadth of *Neonectria-Cylindrocarpon*, Chaverri et al. (15) demonstrated that several major clades existed within this genus and established new generic names for them. Although several segregate genera were recognized, the type species of *Neonectria*, *N. ramulariae*, and *Cylindrocarpon*, *C. cylindroides*, still belong to the same genus and thus are considered synonyms (13,15). In the strict sense *Neonectria* includes the cause of European beech bark disease, *N. coccinea*; American beech bark disease, *N. faginata*; and hardwood canker disease, *N. ditissima* (13). Although a number of important plant pathogenic fungi are included in *Cylindrocarpon*, many of these are no longer considered in the restricted genus *Neonectria-Cylindrocarpon*. "*Cylindrocarpon*" *destructans*, the cause of many root rot diseases, is now placed in a segregate genus as *Ilyonectria radicolica* (basonym: *Nectria radicolica*) (10). The concept of the genus *Cylindrocarpon* based solely on conidial shape is ill-defined including five genera; many of the plant pathogens previously referred to as *Cylindrocarpon* have already been placed in more phylogenetically defined genera. On the other hand the genus *Neonectria* is well-circumscribed and includes a number of plant pathogenic species. Based on these arguments, it is recommended that the generic name *Neonectria* be protected against *Cylindrocarpon*.

Sexual genus not monophyletic, controversy over use of asexual genera: *Pyricularia* 1880 not a synonym of *Magnaporthe* 1972 and *Nakataea* 1939 vs. *Magnaporthe* 1972

The pathogens causing rice blast and stem rot of rice have had various scientific names applied to them due to difficulty in resolving both taxonomic and nomenclatural issues associated with these fungi.

In 1877, Cattaneo (14) first recorded stem rot, a new disease of rice in Italy and described the causal fungus as *Sclerotium oryzae* based on its sclerotial state. About a century later, Krause and Webster (41) established a new sexual genus *Magnaporthe* to accommodate this species as *Magnaporthe salvinii*. The genus *Sclerotium* belongs in

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which is the combination of the oldest legitimate genus and species epithet.

Recently the rice blast fungus, *Magnaporthe oryzae* (synonym *Pyricularia oryzae*) (17), was shown to be not congeneric to the type species of *Magnaporthe*, *M. salvinii*, based on phylogenetic analysis, morphology, and ecological characters (Figs. 5 and 6) (44,80). Therefore, the rice blast fungus does not belong to *Magnaporthe* as typified by *M. salvinii* and should be placed in another genus. The generic name *Pyricularia* has been widely used for the rice blast fungus as *P. oryzae* and this species is congeneric with the type species of *Pyricularia*, *P. grisea*, with these two species shown to be distinct (17). Thus, the scientific name of the rice blast fungus could revert to the previously used name *Pyricularia oryzae*. Alternatively, the fungal name *Magnaporthe oryzae* could be considered for conservation. This is allowed under the Melbourne Code but would require a proposal to conserve the generic name *Magnaporthe* with a different type species, namely *M. oryzae*. Such a proposal would be published in the journal *Taxon*, discussed and voted on by the Nomenclature Committee for Fungi of the International Association of Plant Taxonomy (IAPT), and finally voted on at the next Nomenclature Session of the International Botanical Congress in 2017. This issue is a taxonomic one because the generic names *Magnaporthe* and *Pyricularia* do not compete for priority, i.e., they are not congeneric.

**A****B**

Fig. 5. Conidiophore (**A**) and conidium (**B**) of *Nakataea oryzae* (*Magnaporthe salvinii*, rice stem rot pathogen).



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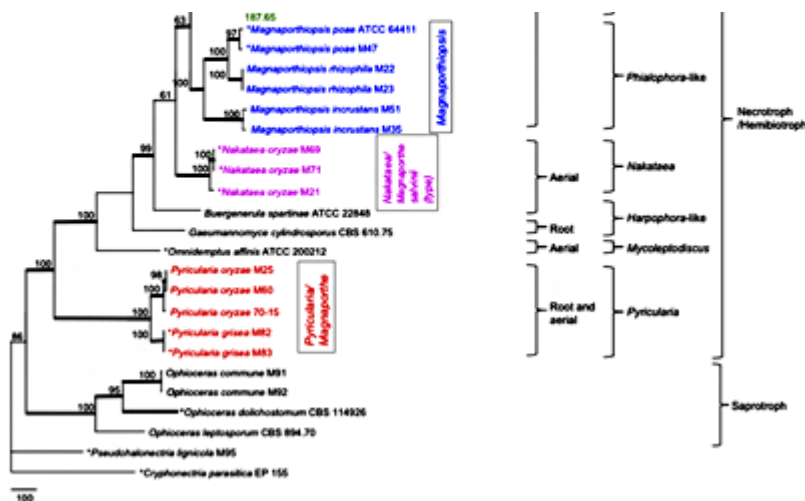
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Fig. 6. The most parsimonious tree inferred from the combined SSU, ITS, LSU, *MCM7*, *RPB1*, and *TEF1* sequence datasets of Magnaporthaceae taxa. MP bootstrap values $\geq 50\%$ are indicated above internodes. Branches in bold have ML bootstrap values $\geq 90\%$ and BI posterior probabilities ≥ 0.95 . Mode of nutrition, anamorph (asexual state) and associated host part are mapped on the tree. Type species of the corresponding genera are indicated with asterisks (44).

One argument against conserving the name *Magnaporthe* with a new type species over *Pycularia* is the excessive number of name changes that would be required for those species currently placed in *Pycularia*. Few name changes would be required if *Pycularia* were accepted. However, 75 names have been placed in *Pycularia*, many of which are plant pathogens of important grass hosts (9). Conserving *Magnaporthe* for the rice blast fungus would require changing many names of species currently in *Pycularia* to *Magnaporthe*.

Both *Pycularia* and *Magnaporthe* are widely used generic names. A discussion and poll are ongoing regarding which name to suggest/reject for the rice blast fungus (<http://magnaporthe.blogspot.com>). The formal list of accepted/rejected names for the blast fungus and related taxa will be generated by the *Pycularia/Magnaporthe* working group (www.fungaltaxonomy.org/subcommissions) and submitted to the General Committee after the issue is fully discussed in the user community.

How Should Plant Pathologists Cope with the New Fungal Nomenclature?

The change to the nomenclature for fungi will result in the use of some scientific names that are unfamiliar to the user community. In order to minimize this and achieve nomenclatural stability, plant pathologists, mycologists, geneticists, and the broad user community are working together to determine which names to use for pleomorphic fungi. A number of subcommissions of the International Commission on

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These lists and proposals will be submitted to the Nomenclature Committee for Fungi and General Committee of the IAPT and are subject to the decision of the next International Botanical Congress (29,30). We encourage everyone who works with fungi to get involved in the decision-making process by actively participating in working groups or subcommissions that have been established (www.fungaltaxonomy.org/subcommissions), or to propose a new working group.

As scientific names of plant pathogenic fungi are integrated, lists of accepted names will be published as has been done for well-studied groups such as *Aspergillus* and *Penicillium*. As the process results in accepted lists, accurate scientific names will be placed on websites such as the USDA-ARS SMML Fungal Nomenclature (nt.ars-grin.gov/fungaldatabases/nomen/nomenclature.cfm) that emphasizes plant-associated fungi, and Index Fungorum (www.indexfungorum.org/Names/Names.asp) and MycoBank www.mycobank.org that list scientific names of all groups of fungi. These databases will be updated as decisions are made. The National Center for Biotechnology Information (NCBI) manages a taxonomy database (21), which is the standard nomenclature and classification repository for the International Nucleotide Sequence Database Collaboration (INSDC), comprising the GenBank, ENA (EMBL) and DDBJ databases. This widely used resource can be an important aid in transitioning between different Codes. Currently several specialist taxonomists examine new names as they are added to the database, but dual fungal names are still treated inconsistently. An overhaul of fungal names is underway but advice and interaction from the broader fungal research community will be essential.

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