MINIREVIEW

Where Are All the Undescribed Fungi?

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ABSTRACT

The hypothesis that there are 1.5 million fungal species on Earth, of which only about 70,000 are described, implies that 1.43 million remain undescribed. The recognition that many new species have yet to be found is of fundamental importance to plant pathologists, agronomists, and plant regulatory officials, among others, who continue to encounter diseases caused by previously unknown or understudied fungi. Unexplored habitats with their arsenal of unknown fungi are also of interest to those searching for novel organisms for use in biological control or for their pharmaceutical attributes. This paper presents data on the expected numbers of fungi in some relatively unexplored habitats, such as tropical forests, and those obligately associated with plants, lichens, and insects.

In addition to undiscovered species, many have been collected but remain lost or hidden as named species and ignored for lack of modern characterization; others have been collected and recognized as new species but remain undescribed. Some fungal species are unrecognized within erroneously circumscribed species, often based on presumed host specificity, while others exist as biological species but remain buried within those broadly defined species for lack of gross morphological characterization. From these data, one must conclude that enormous numbers of unrecognized fungi can be found almost everywhere, including one’s own backyard.

Additional keywords: biodiversity, biosystematics, diversity, emerging diseases, inventory, lichenized fungi, plant pathogens.

TROPICAL FORESTS

Many biodiversity specialists, particularly those with expertise in macroorganisms, have determined that tropical forests are more species-rich than temperate forests; thus, this is generally assumed to be true for fungi as well. However, this hypothesis has not been rigorously tested for fungi, although a few comprehensive studies have been undertaken that provide supportive evidence (8,19). For example, more than 500 fungi were identified in intensive studies of litter from only five tree species in Panama (11). After 1 year of intensive collecting in one forest in Kenya, 59 of 75 (79%) of the leaf-dwelling ascomycetes encountered were new species (40). Recent investigations of fungi on palms yielded 1,580 species, of which 75% of the species collected were new to science (27). Data on the primarily tropical ascomycete family Phyllachoraceae, which 75% of the species collected were new to science (27).

In exploring tropical regions for fungi, it is anticipated that the most widespread and common species in a site will tend to be found first. Most likely, such species have been collected before and are described already. However, the longer the time spent in intensive exploration, the more species are discovered, as evidenced by the continuing accounts of new species, primarily of mitotically active fungi, that have been discovered in Taiwan during the past 16 years (29–37). A 2-month visit to Papua New Guinea yielded 6 genera and 89 species of lichen-forming and lichenicolous fungi new to science (2). Only through an intense examination of all fungi in a defined area will the finite number of fungal species be determined. The results of a recent workshop to plan an all-taxa biodiversity inventory in the Área de Conservación Guanacaste, Costa Rica, estimated the number of fungal species remaining to be encountered in the 120,000-ha reserve at around 50,000 (6).

An overall indication of whether the tropics are more species-rich in fungi than temperate regions might be expected based on an analysis of the origin of newly described fungi. The following data were extracted from the Index of Fungi database maintained at the International Mycological Institute (IMI), Egham, U.K. Tropical countries were the source of 49% of the 16,013 fungal species, including fossils, described as new during a 10-year period, 1981 to 1990. However, the United States was the origin of the largest number of new taxa, with 1,623 (10.1%) species, followed closely by India, with 1,554 (9.7%) species. Of the remaining countries with more than 1% of the total, 8 of 22 were tropical. Rather than indicating whether tropical countries are the richest
source of new fungal species, these data suggest that all countries are inadequately known mycologically. Those yielding the most species new to science are merely the ones that receive the most intense scrutiny rather than those that are inherently the most species-rich. This view is supported by the situation in the British Isles, the most comprehensively studied region for fungi in the world. Here critical or intensive studies invariably yield species that have never been described: 460 species new to science were described from Great Britain and Ireland during the same 10-year period.

UNEXPLORED HABITATS

The number of habitats that potentially support specialized and unique fungi is enormous. The fungi described as new to science during 1981 to 1990 (mentioned above) were associated with 1,982 host genera or substrata. Some previously unexplored substrata and habitats from which these fungi were found include the rumens of herbivorous mammals; algae, lichens, and mosses; marine plants, including mangroves and driftwood; rocks; and insect scales. Soil and plant litter also continue to be a rich source of unusual fungi. Selected examples from a growing number of data sets are indicative of our current lack of knowledge.

Plants

The 200,000 species of vascular plants continue to serve as the major reservoir of novel fungi. It has been estimated that there may be around 270,000 species of plant-pathogenic fungi in the tropics (47). As in the case of geographic analysis, it is intensity of study that best explains the observations. For example, during the 70-year period from 1920 to 1990, about three times as many new species of fungi were described from the relatively well-studied Poaceae compared to the related but less-studied Cyperaceae (8). Within the Poaceae, the number of fungi per potential host species ranged from a low of 0.3:1 in the genus Poa to 35:1 in the genus Zea, with 387 reported on Zea mays alone. Even among grasses in the United States, there are no reports of fungi on 45% of the grass hosts (15). After analyzing these data (15), Clay (10) reported that among the 20 grasses with the highest reported numbers of fungi in the United States, only 3 are native grasses. These numbers reflect the intensity of study, i.e., economic importance, rather than the biology of the host plants or the true fungal biodiversity of these two related plant families.

The same situation is found when one plant species becomes the focus of attention. In a search for potential fungal biocontrol agents, the pernicious weedy shrub Lantana camara was scrutinized carefully and is now known to support 55 fungi, of which 28 appear to be restricted to that one host species (3). Three new species, one representing a new genus, were discovered during one study of L. camara in Brazil (3). The maritime rush Juncus roemerianus, found along the eastern coast of the United States, was not known to support any host-specific fungi until two researchers began investigating its mycota in 1991. From this one plant species, Kohlmeyer et al. (28) have now described 20 new species, including 8 new genera and 1 new family. Even on this single-host plant, each of the numerous microhabitats are occupied by different fungal species.

In the tropics, it is particularly difficult to ascertain patterns of species richness. A. C. Batista and coworkers (12) recorded 3,340 fungi from Brazil associated with 523 plant species, equivalent to an average of 6.4 fungi per host plant studied. Such data were not collected for an analysis of species richness and are difficult to assess, because not all plants were studied and plant species without fungi were not listed. Further, although several new species could occur on a single leaf, no information is available from which to make sound extrapolations. How many leaves on the one tree species had novel fungi was not specifically reported nor was how many of those fungi occurred on different and even unrelated trees nearby. Well-designed studies are needed to provide reliable data on this important issue.

Data on fungi reported on Eucalyptus recently have been compiled (46). Of the known 450 species in Eucalyptus, 1,350 species of fungi are known from 150 host species. The number of fungi from each host species ranges from 1 to 282. The highest number of fungi were found on E. globulus, and 150 of the 282 fungi on that host were not listed on any other species of Eucalyptus. Analyses of these and other data sets suggest that 5.3 unique fungi per host plant species is a reasonable working number (20).

Lichens

Lichenicolous fungi, i.e., fungi growing on and in lichens, are a largely undiscovered group of organisms, many of which are novel genera and species. The number catalogued has increased from 457 species in 1976 to 894 in January 1996, a rise of 96% during 20 years. These totals are set to increase even more steeply, as evidenced by the following four studies. In a recent preliminary account of heterobasidiomycetes growing on lichens, 53 species were recognized, of which 46 (92%) were previously undescribed (13). An examination of fissitunicate pyrenomycetes on tropical foliicolous lichens yielded 49 species, of which 36 (73%) were new to science (38). In examining fungi from the relatively well-studied lichen Peltigera, 96 species are now known, 61 of which occur only on this lichen host; 7 of these species were described only within the past year (22). A study of the fungi isolated from thalli of two fruticose lichen genera from one site in Germany revealed 506 strain types (42).

Insects

In preparing estimates of the global numbers of fungi, Hawksworth (17) adopted a conservative approach to both insect numbers and the extent of host specificity of the insectivorous fungi. New data have been presented on the extent of host specificity in the insect-infecting Laboulbeniales, a group of fungi that occur on the exoskeletons of beetles and flies. Detailed comparisons were made between rich collections from sites in the United Kingdom and Sulawesi in Indonesia (50). This analysis suggests that, although only 1,855 species of Laboulbeniales are currently known, the actual number is likely to be between 10,000 and 50,000 species. This implies that only 3.7 to 18.5% of the species in the order are known. The situation may be even more extreme in the case of the Trichomycetes, a group of enigmatic fungi occurring in the hindguts of insects and other invertebrates. These organisms are so poorly studied in most regions of the world that generalizations are hazardous. However, it is clear that these are a major source of undiscovered fungi even in the United Kingdom (41).

New Techniques Applied to Previously Studied Substrata

Soil is one of the substrates from which fungi have been studied for many years, resulting in some relatively comprehensive accounts of soil fungi (14,44). Yet, the results of application of selective methods for isolating fungi from soil, such as selective isolation of low-temperature microfungi (9) or basidiomycetes (49), demonstrates that entire groups of fungi have been overlooked, even in relatively well-studied substrata. Species occurring at lower frequencies and those that are technically difficult to detect are less likely to have been discovered.

The increased use of techniques that favor detection of rare species, specifically particle filtration, has resulted in the isolation of several hundred fungi from one substrate without reaching an asymptote in the species-discovery curve (4). This technique, in which washed, pulverized, and filteredsubstrates are placed on a weak agar medium and closely observed, allows elimination of fast-growing fungi, so slower growing species can be detected. Using this technique, the number of fungal species isolated from the decaying plant parts of a single tropical plant exceeded 200
In comparing species diversity from one plant host determined by two detection methods, the particle filtration technique yielded 10 times the number of fungal species detected by moist-chamber incubation (43). Thus, methods of isolation and observation play an important role in determining fungal diversity.

**LOST OR HIDDEN SPECIES**

Many fungi are lost or hidden within existing constructs.

**Cryptic Biological Species**

Fungi traditionally have been distinguished on the basis of morphological characters. However, genetic and molecular data suggest there are many functional biological species masquerading under a single species name or recognized only at an infraspecific rank or as a “special form.” It is striking that almost every fungus critically studied by population biologists is found to comprise a number of reproductively isolated “biological species” (5). In distinguishing such biological species as formally described taxa, the biological information characteristics of that entity, e.g., its host range, pathogenicity, and growth conditions, likewise may be distinguished. If a biological species concept were adopted for all fungi, a case could be made for multiplying the existing total number of species in some groups by a factor of five or more.

**Lost Within Described Species**

In some groups of fungi, there has been a tradition of describing species as new if the fungus is found on a new host plant. This can lead to the unnecessary proliferation of species names, but it also can mask situations in which more than one species of a fungal genus occurs on a single host. When critically examined, many of these fungi can be keyed out based on morphological criteria alone, without reference to the host. This was the case in a recent study on Meliola in Kenya (40). The net effect of this tradition is unknown, but experience at IMI suggests that in groups such as Cercospora and Meliola too few, rather than too many, species are currently being recognized. Recent work on species of Alternaria on Citrus exemplifies the situation in which isolates from one host are uncritically identified as a single, widely reported species considered to be host specific, while, in reality, a number of distinct fungal species occur on that host (48). Such simplifications have serious plant quarantine implications.

**Named and Orphaned Species**

Many fungi have already been named but remain lost amongst the 250,000 species names that have not been characterized by modern standards. These species are not reported in the recent literature and are not included in the total number of known species. Many of these obscure names are probably synonyms of known species. Based on 15 fungal monographs (18), one can calculate a ratio of 2.5 synonyms for each “good” species. This ratio suggests that of the 250,000 species names in existence 100,000 of these are accepted species. Because only 72,000 are treated in the modern literature (21), there are 28,000 currently undopted “orphan” species. Although this is not a major portion of the absentees, it does make the total number of expected species considerably more realistic. When genera are studied on a world scale, comprehensive monographs reveal spectacular numbers of previously undescribed species. Of 39 species accepted in Meliola on tropical plants, 26 (67%) were new (26), and of 20 in Lichenothelia on rocks, 18 (90%) were new (24). In some cases, the number of species known in one fungal genus is reduced significantly when species prove to be plurivorous. This is exemplified by an account of Didymosphaeria, in which it was determined that the 7 accepted species had more than 100 synonyms (1). However, the usual pattern for little-studied fungi is that when investigated a large percentage of undescribed species are discovered.

**Collected and Unidentified**

Most mycologists concentrate on one or two major groups of fungi that they collect and study throughout their lifetime. With the opportunity to visit previously unexplored regions of the world, each scientist bears the responsibility of collecting specimens of all groups of fungi and distributing them to experts. Yet, for many groups of fungi, including plant pathogens, no expert exists to study the specimens. At best, the specimens are deposited in the world’s herbaria and identified only to the genus or not at all. Almost every mycological collection has folders, drawers, or boxes of specimens that represent a bonanza of as-yet unstudied and, possibly, undescribed species. Experts are needed who can undertake the rigorous study required to provide a formal description and place each new species in its taxonomic framework. In the case of major mycological institutions, these may amount to several thousand specimens and represent a significant portion of the undiscovered fungal biodiversity. To speculate, the number of undescribed fungal species represented by these specimens is probably more than 20,000.

**OPPORTUNITY AND CHALLENGE**

The impression is often given that fungi in general have wider geographic distributions than vascular plants. In practice, the situation varies markedly between different groups. Particularly widespread are ancient fungi from phyla extending back to the first colonization of land in the Silurian region, some 300 million years before the first flowering plants. Fungi known from the early fossil record include endomycorrhizas, chytrids, myxomycetes, and some lichen-forming fungi. Also found in the category of fungi with wide geographic distributions are certain nonspecific, wood-rotting bracket fungi, opportunistic r-strategy spoilage molds, coprophilous fungi, and rapidly growing soil saprobes. Detailed comparisons between national checklists of fungi, although incomplete, would give some indication of the extent of overlap. However, when new species abound everywhere, it is difficult to know whether these discoveries are restricted, endemic species, or cosmopolitan taxa.

The extent of our ignorance poses major problems for mycologists and plant pathologists working in little-studied countries or habitats. What is already known? Which names are correct? How does one store information on new collections? How can one begin to identify newly collected specimens? How does one ascertain what is new? How does one gain access to available information when important publications no longer in print may be impossible to obtain? Conversely, how does one make new information accessible to others? These are complex problems that must be addressed (23). Answers are far from easy because of the lack of attention devoted to the myriad organisms involved.

Some may say that molecular approaches are the sole answer to describing fungal biodiversity. We do not. With around 1,800 fungi new to science being described every year, and not more than 100 species sequenced each year, the gap widens rather than narrows. Other than for fungi of medical and agricultural significance, the technological developments, including automated molecular and biochemical procedures, are unlikely to solve this problem within the foreseeable future. We suggest that those working in descriptive mycology must be well supported in the essential task of characterizing newly discovered fungal species. With increased electronic resources and use of the Internet to disseminate information, the products of every mycologist describing fungi can be made available rapidly along with user-friendly keys for identification linked to diagrams explaining cryptic features and illustrations of the species. Ideally, the two approaches to understanding fungal species would be combined, such that newly described species would be routinely sequenced and organisms representing the ends of molecular cladograms would be fully described. The result would be a whole that equaled more than the sum of its parts.
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Literature Cited