

## **Rapid publication of taxonomic and nomenclatural novelties**

The advantages of modern methods in species identification and delimitation has started a new era in the exploration of fungal diversity. The overload of newly discovered taxa has resulted in the need to publish them in a simple way (see e.g. Fungal Planet sheets). Parallel to this the novel nomenclature rules enable effective publication of new names in electronic PDF format in the absence of printed hard copy since 2012. These two steps have strongly accelerated the publication of taxonomic novelties, but also started a race against time in naming and combining new taxa, as can be seen in Index Fungorum.

As a reaction to these recent changes, the Editorial Board of Czech Mycology has decided to shorten the publication process of studies and researches, and to accelerate valid publication of taxonomic novelties in order to ensure priority nomenclature proposals.

A new type of papers for simple and fast publication of taxonomic and nomenclatural novelties is introduced since Volume 67 (2015). The manuscripts have a reduced form, but are peer-reviewed by external reviewers and should be submitted in the same way as other manuscripts. This reduced form is only intended for fast publication of novelties, ensuring the priority of proposed changes. Papers containing more detailed data (e.g. intraspecific variability, ecology, phylogeny) should be submitted as an ordinary paper.

A reduced-form manuscript should contain:

- Title of the paper.
- Author's/s' name(s) and author's/s' address(es).
- Brief abstract (up to 50 words).
- Name(s) of the newly described taxon (taxa) including its/their registration numbers (MycoBank, Index Fungorum, Fungal Names).
- Short diagnosis and/or detailed description with complete type citation.
- Notes, including the most important information usually presented in the Introduction, Methods and Discussion parts; this part should include accession numbers of sequences acquired by the author(s) and discuss their similarity to published sequences.
- Figures, phylogeny trees (facultative).
- References.