

## Special Issue

# Comparative and Functional Genomics of Toxigenic Fungi

### Message from the Guest Editors

**Dear Colleagues,** Fungi can produce a wide array of secondary metabolites (SMs), some of which have beneficial properties, while others can be harmful toxins that pose risks to human and animal health, as well as negatively impact agriculture and related industries. The advent of whole-genome sequencing has initiated a new phase in the study of fungal biological systems. This has been made possible by the significant increase in the catalogue of genome sequences of fungi and the development of new reference genomes.

Advancements in omics techniques have facilitated identification, characterization, variation, and comparative studies of fungi, providing new insights into the relationship between secretome and fungal lifestyles and helping to identify gene clusters for both known and previously undiscovered cryptic SMs. This Special Issue is focused on the recent advances in mycotoxin research, with an emphasis on the genetics and biology of fungal toxin production. The topics of interest will include comparative genomics and transcriptomics addressing genome-wide variations, pathogenicity mechanisms, secretome analysis, and the regulation of secondary metabolite gene clusters.

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### Guest Editors

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### Deadline for manuscript submissions

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The International Journal of Molecular Sciences (*IJMS*, ISSN 1422-0067) is an open access journal, which was established in 2000. The journal aims to provide a forum for scholarly research on a range of topics, including biochemistry, molecular and cell biology, molecular biophysics, molecular medicine, and all aspects of molecular research in chemistry. *IJMS* publishes both original research and review articles, and regularly publishes special issues to highlight advances at the cutting edge of research. We invite you to read recent articles published in *IJMS* and consider publishing your next paper with us.

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