Special Issue

Diversity of *Mycobacterium tuberculosis*

Message from the Guest Editors

Although it has gained less attention than COVID-19, tuberculosis (TB) caused by the Mycobacterium tuberculosis (Mtb) complex remains among the most prevalent and deadly infectious diseases worldwide. Infections result in diverse clinical outcomes, from latent asymptomatic infection to pulmonary or extrapulmonary manifestations of disease, with an array of severity symptoms. The origin of the diversity of these clinical presentations remains poorly understood.

As Co- of this Special Issue, we invite you to submit research articles, review articles, and short communications related to the diversity, microdiversity, and adaptation of the Mycobacterium tuberculosis complex. Reviews should focus on present knowledge on the current diversity of Mtb and its functional consequences. Research articles and short communications may present new exploratory studies or new methods to unravel microdiversity, mechanisms that control diversification, and/or relate diversity to clinical or geographical features of the bacillus.

Guest Editors

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Deadline for manuscript submissions

closed (28 February 2023)

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Message from the Editor-in-Chief

"Microorganism" merges the idea of the very small with the idea of the evolving reproducing organism is a unifying principle for the discipline of microbiology. Our journal recognizes the broadly diverse yet connected nature of microorganisms and provides an advanced publishing forum for original articles from scientists involved in high-quality basic and applied research on any prokaryotic or eukaryotic microorganism, and for research on the ecology, genomics and evolution of microbial communities as well as that exploring cultured microorganisms in the laboratory.

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