

Special Issue

Molecular Epidemiology of Leishmaniasis

Message from the Guest Editor

The molecular epidemiology of leishmaniasis is a developing field that has been revolutionised by the emergence of genomic epidemiology and population genetics, as well as deep sequencing applications to study population diversity within hosts, which also inform the development of improved diagnostic and genotyping methods. This Special Issue will accept Original Research Articles and Reviews that cover all aspects related to the molecular epidemiology of leishmaniasis, including, but not limited to, genomic epidemiology, the development, application, and validation of improved genetic markers in diagnosis and genotyping, and studies that use molecular markers and genome sequences to elucidate transmission, association with clinical presentation, new reservoir hosts or vectors, and the taxonomy of leishmaniasis.

Guest Editor

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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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