



Molecular Epidemiology of Leishmaniasis

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Deadline for manuscript
submissions:

31 July 2024

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.





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Message from the Editor-in-Chief

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