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New Resources and Methods for Reconstructing Human-Pathogen Co-evolution

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Deadline for manuscript
submissions:

closed (30 April 2022)

Message from the Guest Editors

This Special Issue will take a multidisciplinary approach geared towards utilising the currently available clinical and historical human infectious agent data to systematically explore the molecular histories of human infectious diseases. The aim of this Special Issue is to bring clinicians, evolutionary biologists, archaeologists and data scientists together to provide readers with an up-to-date and state-of-the-art summary of sampling resources, electronic data archiving and sharing and computational and statistical methods, to provide a comprehensive analysis of human pathogen exposure and the molecular histories of human infectious diseases. Research articles, review articles as well as short communications are invited. We ask you to send us an abstract prior to submission to ensure that your work fits within the scope of this Special Issue.



mdpi.com/si/78449

Special Issue



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Message from the Editorial Board

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