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Hemorrhagic Fever Viruses

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submissions:

closed (31 December 2020)

Message from the Guest Editors

Dear Colleagues,

Viral hemorrhagic fever (VHF) refers to a severe multisystem syndrome, which can be caused by a wide range of viruses, including arenaviruses, bunyaviruses, filoviruses, and flaviviruses. Examples of high-consequence hemorrhagic fever viruses include Ebola virus, Lassa virus, Hantavirus, and Crimean–Congo Hemorrhagic fever virus, but also Alkhurma virus, Chapare virus, Guanarito virus, Kyasanur Forest disease virus, Lujo virus, Omsk hemorrhagic fever virus, and Sabiá virus. These viruses are found across multiple continents and continue to put large portions of the global population at risk.

This Special Issue will cover a wide range of topics focusing on viral hemorrhagic fevers and aims to help to enhance current knowledge of these viruses and their respective diseases. All types of articles will be considered for publication, including short reports, primary research articles, and reviews.

We look forward to your contribution.



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Special Issue



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Editor-in-Chief

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Message from the Editor-in-Chief

The worldwide impact of infectious disease is incalculable. The consequences for human health in terms of morbidity and mortality are obvious and vast but, when infections of animals and plants are also taken into account, it is hard to imagine any other disease that has such a significant impact on our lives—on healthcare systems, on agriculture and on world economics. *Pathogens* is proud to continue to serve the international community by publishing high quality studies that further our understanding of infection and have meaningful consequences for disease intervention.

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