

Special Issue

Hepatitis E Virus Molecular Epidemiology and Evolution

Message from the Guest Editor

Even if much progress has been achieved in our understanding of hepatitis E virus (HEV), many gaps in our knowledge remain. HEV has a broad host range being isolated from fish, birds, and mammals, such as bats, rodents, ungulates, and humans. Most likely, other vertebrate hosts still remain to be identified. Viruses belonging to the same genus have the potential to cross species barriers, apart from HEV1 and HEV2, which only infect humans. These two types, which cause large waterborne outbreaks with high mortality rates in the pregnant female, are also associated with pancreatitis.

Guest Editor

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Message from the Editor-in-Chief

Viruses (ISSN 1999-4915) is an open access journal which provides an advanced forum for studies of viruses. It publishes reviews, regular research papers, communications, conference reports and short notes. Our aim is to encourage scientists to publish their experimental and theoretical results in as much detail as possible. There is no restriction on the length of the papers. The full experimental details must be provided so that the results can be reproduced. We also encourage the publication of timely reviews and commentaries on topics of interest to the virology community and feature highlights from the virology literature in the 'News and Views' section.

Electronic files or software regarding the full details of the calculation and experimental procedure, if unable to be published in a normal way, can be deposited as supplementary material.

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