

In this issue

Research Article

[Open Access](#) [Research Article](#) PTZAID:AHR-8-132

The dark proteome of rodent hepatitis E virus: Analysis of intrinsically disordered regions

Published On: February 18, 2022 | Pages: 005 - 011

Author(s): Zoya Shafat, Anwar Ahmed, Mohammad K Parvez, Asimul Islam and Shama Parveen*

Hepatitis E virus (HEV) is the causative agent of Hepatitis E infections across the world. Intrinsically disordered protein regions (IDPRs) or Intrinsically Disordered Protein (IDPs) are regions or proteins that are characterized by a lack of definite structure. These regions or proteins play significant roles in a wide range of biological processes, such as cell cycl

...

[Abstract View](#) [Full Article View](#) [DOI: 10.17352/ahr.000032](#)

[Open Access](#) [Research Article](#) PTZAID:AHR-8-131

Prevalence of hepatitis B virus infection among persons with hepatitis D virus and diabetes mellitus in Pakistan, 2019-2021

Published On: February 11, 2022 | Pages: 001 - 004

Author(s): Ahmad Raza, Muhammad Waqar Mazhar*, Saira Saif, Mudasara Sikandar, Iram Shahzadi, Javaria Mehmood, Wajeeha Iram, Hira Tahir, Shanzab Noor, Fatima Mazhar

Introduction: The HBV virus has its enveloped protein that surrounds nucleic acid for its protection. Hepatitis B DNA virus belongs to the Hepadnaviridae family and is similar to retroviruses. Globally, HBV infected people approximately 2 billion in the world, about 350 million people were chronic carriers. One million deaths are caused by the HBV virus every year.

Ea ...

[Abstract View](#) [Full Article View](#) [DOI: 10.17352/ahr.000031](#)

Opinion

Bio-informatics approaches in liver disease: Incomplete without experimental validation

Published On: September 27, 2022 | Pages: 012 - 013

Author(s): Pratibha Gaur*

In the search for the treatment of hepatic diseases, multiple approaches have been used frequently such as bioinformatics and algorithm-based systems biology for the screening of already published data that leads to the identification of promising drug candidates at molecular levels. These studies provide a large data set of information and claim to identify many gene ...

[Abstract View](#)

[Full Article View](#)

[DOI: 10.17352/ahr.000033](#)