Reviewer's report

Title: Molecular epidemiology of hepatitis B virus infection in Norway

Version: 0 Date: 13 Dec 2018

Reviewer: Zhi Chen

Reviewer's report:

This paper analyzed 1,157 HBV small S-gene sequences collected from patients in Norway, and investigated epidemiological data from the Norwegian surveillance system for communicable diseases. The authors found the distribution of HBV genotypes in the Norwegian population is dominated by genotype D (32%) followed by genotype A (22%), B and C (18% and 18% respectively), and E (7%). They showed that the patients in Norway were primarily to have been infected with HBV of genotype A and D. There are some grammar mistakes in the manuscript. Please carefully modify the whole manuscript.

Are the methods appropriate and well described?
If not, please specify what is required in your comments to the authors.

Yes

Does the work include the necessary controls?
If not, please specify which controls are required in your comments to the authors.

Unable to assess

Are the conclusions drawn adequately supported by the data shown?
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Yes

Are you able to assess any statistics in the manuscript or would you recommend an additional statistical review?
If an additional statistical review is recommended, please specify what aspects require further assessment in your comments to the editors.

I am able to assess the statistics

Quality of written English
Please indicate the quality of language in the manuscript:

Needs some language corrections before being published
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