Reviewer’s report

Title: Clinical and whole-exome sequencing findings in two siblings from Hani ethnic minority with congenital glycosylation disorders

Version: 3 Date: 23 Sep 2019

Reviewer: Kamwing Jair

Reviewer's report:

More detail information of how they perform the whole-exome sequencing followed by analytical strategy have been be provided so other researchers will be benefit from this study.

Unfortunately, a functional assay, a measurement by qRT-PCR, had performed unsuccessfully due to the poor quality of participant samples. However, those data are critical to link the mutation to its mechanism so that this finding can achieve the level of Research Article, not a Case Report.

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.

Yes

Are the conclusions drawn adequately supported by the data shown?
If not, please explain in your comments to the authors.

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:

Acceptable
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