Reviewer’s report

Title: Mitochondrial neurogastrointestinal encephalopathy as a mimic of Crohn's disease: a case report

Version: 3 Date: 22 Dec 2018

Reviewer: Holm Uhlig

Reviewer’s report:

The authors have made minor adjustments (page 4, lines 92-97):

"The previously reported whole genome sequencing of Crohn's disease produced median coverage 4x genome-wide. At this sequencing depth it is not possible to exclude ultra-rare variants in TYMP in the 2513 patients who were analysed. This would require high coverage sequencing. Their absence does, however, make it unlikely that TYMP variation is a significant cause of Crohn's disease or that MNGIE phenocopying Crohn's disease is a common problem." This broad statement is correct although not unexpected.

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.

Not relevant to this manuscript

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

Acceptable
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