Reviewer’s report

Title: Whole-exome Sequencing Identifies a Novel Missense Variant within LOXHD1 Causing Rare Hearing Loss in a Chinese Family

Version: 0 Date: 22 Dec 2018

Reviewer: Jianjun Xiong

Reviewer's report:

The authors identified a novel missense variant at LOXHD1 gene exome associated with a rare NSHL from a Chinese family. The results demonstrate the effectiveness of whole-exome sequencing for molecular diagnosis of rare diseases, and expand the genotypic spectrum of DFNB77. The manuscript is well written and recommended to publish.

Suggestion: please list the primers for validation the variant in LOXHD.

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?
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Yes

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

Yes

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Not relevant to this manuscript

Quality of written English
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