Reviewer's report

Title: Whole-exome sequencing identifies MYO15A mutations as a cause of autosomal recessive nonsyndromic hearing loss in Korean families

Version: 2 Date: 12 June 2013

Reviewer: Tom Walsh

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If the Real-time QPCR analysis of exon 42 will be included in the Supplement please provide hg19 location of the probe in the legend.

Level of interest: An article whose findings are important to those with closely related research interests

Quality of written English: Acceptable

Statistical review: No, the manuscript does not need to be seen by a statistician.

Declaration of competing interests:

I declare that I have no competing interests'