Reviewer’s report

Title: A very early diagnosis of Alström syndrome by next generation sequencing

Version: 2 Date: 01 Jun 2020

Reviewer: Milad Gholami

Reviewer's report:

1-Add the methods and table of filtering strategy for variant detection to manuscript.

2-What software and database did you use to analyze the novelty of mutation and which showed your mutation is pathogenic? You can refer to the article "Whole exome sequencing identified two homozygous ALMS1 mutations in an Iranian family with Alström syndrome" for more information.

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:

Needs some language corrections before being published

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