Author’s response to reviews

Title: Whole exome sequencing highlights variants in association with Keratoconus in Jordanian Families

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Response to Editor
22 July 2020

1. Abbreviations

All abbreviations were given at the end of the manuscript

2. Consent to Participate

Amended accordingly and indicated in the manuscript body as well.

3. Availability of data and materials

The data is in the process of uploading to SRA-NCBI and the accession number will be added upon the completion of upload.

4. Authors’ contributions

The co-authors have significant contribution; they contributed in designing the methodology as well as writing the clinical methodology. They also have read and approve the manuscript.

5. Figure 1

a) Please amend the in-text citation “Figure 1J” to “Figure 1G” in your manuscript. Please also amend the label of “J)” to “G)” in Figure 1.

It is amended accordingly.
b) Please ensure that you include figure legends for each of the sub-figures (A, B, C, D, E, F, G, H) of Figure 1 in your manuscript.

This is was checked.

6. Overlap

The overlapping was rechecked and amended in the methodology as well as in the first paragraph of the introduction.

Many thanks
Tawfiq Froukh