Author’s response to reviews

Title: TNF-α−308 G/A and IFN-γ+874 A/T gene polymorphisms in Saudi patients with cutaneous leishmaniasis

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Author’s response to reviews:

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March 10, 2020

Dr. Kamlesh Guleria /Dr. Matteo Pasini
Editors,
BMC Medical Genetics

Dear Dr. Guleria/Pasini:

Thank you for giving us the opportunity to RE-Re-revise our manuscript MGTC-D-19-00560R2, entitled “TNF-α−308 G/A and IFN-γ+874 A/T gene polymorphisms in Saudi patients with cutaneous leishmaniasis”. The manuscript has been RE-Re-revised in light of the editor’s comments for publication in BMC Medical Genetics. The material contained in the manuscript is original and has not been submitted for publication elsewhere.
We have addressed all the concerns raised by editors as detailed under the enclosed point-by-point reply to the reviewers.

We continue to RE-express our gratitude to the editors as well as reviewer’s constructive criticisms, which have improved the quality of the manuscript. We hope that the editors and the reviewers will find the answers satisfactory.

Sincerely,

Zafar Rasheed, MS., Ph.D., PGDCA.,
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POINT-BY-POINT REPLY TO THE EDITORS TECHNICAL COMMENTS

Manuscript Reference: MGTC-D-19-00560R4
Manuscript Title: TNF-α−308 G/A and IFN-γ+874 A/T gene polymorphisms in Saudi patients with cutaneous leishmaniasis
Journal: BMC Medical Genetics

We are grateful to the editors for their thorough review and for the constructive comments, which enabled us to further RE-improve our manuscript. Concerns raised have been addressed again. Please note that all the changes in the Re-revised manuscript are highlighted.

Editor Comments:

Comment 1:
Technical Comments:
Editor Comments:

The figures have not been modified as suggested. Replace the photograph of Figure 1 and Figure 2 as suggested below:
As mentioned that ARMS PCR was performed once with Common Forward primer and with G specific primer second with same forward and A specific reverse primer for the same sample. So for GG genotype, amplification will be seen only in one combination (Common Forward and G allele specific Reverse Primer (as mentioned in lane 1). For AA genotype, amplification will be seen in combination (Common Forward and A allele specific Reverse Primer as mentioned in lane 4) and for GA genotype, amplification will be seen in both the combinations (as mentioned in lane 5 and 6).

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<th>GG</th>
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<td>Sample 1</td>
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Author’s Response:
We highly appreciate you for your thorough review again, which definitely enabled us to further RE-improve our manuscript. As instructed, figure 1 and figure 2 have now been replaced with new figures and the details have been given in the figure legends on page 18 in the re-revised manuscript.

We believe that the RE-revised manuscript will now meet the high standards of the journal and suitable for “BMC Medical Genetics”.

Dr. Zafar Rasheed
Corresponding Author