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

Authors:

Ahmed Ahmed (ahmedali@qummed.edu.sa)
Zafar Rasheed (zafarrasheed@qummed.edu.sa)
Tarek Salem (tareksalem@qummed.edu.sa)
Mohammed Al-Dhubaibi (m.aldubaibi@qummed.edu.sa)
Ahmad Al Robaee (arobaee@gmail.com)
Abdullateef Alzolibani (azolibani@yahoo.com)

Version: 3 Date: 04 Mar 2020

Author’s response to reviews:

Department of Dermatology/Medical Biochemistry,
College of Medicine,
Qassim University,
Buraidah, KSA.

February 28, 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.,
ORCID: 0000-0002-8651-4218
Corresponding Author

POINT-BY-POINT REPLY TO THE EDITORS TECHNICAL COMMENTS

Manuscript Reference: MGTC-D-19-00560R2
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: Replace figure 1 and figure 2 as mentioned below

<table>
<thead>
<tr>
<th></th>
<th>1</th>
<th>2</th>
<th>3</th>
<th>4</th>
<th>5</th>
<th>6</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>GG</td>
<td>AA</td>
<td>GA</td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
</tbody>
</table>

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 same sample. So for GG genotype amplification will be seen only in first combination, for AA genotype with second combination and for GA genotype amplification will be seen in both combinations. Therefore, label Figure 1 as mention in above figure. Similarly change the figure 2 on same pattern.

Author’s Response:
We highly appreciate you for your thorough review again, which definitely enabled us to further RE-improve our manuscript. As suggested, levels in figure 1 and figure 2 have further been modified 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