

PEER-REVIEW REPORT

Name of journal: *World Journal of Clinical Infectious Diseases*

Manuscript NO: 79603

Title: Three-dimensional model of antigens with serodiagnosis potential for leprosy: an in silico study

Provenance and peer review: Unsolicited Manuscript; Externally peer reviewed

Peer-review model: Single blind

Reviewer's code: 06325500

Position: Peer Reviewer

Academic degree: PhD

Professional title: Academic Research

Reviewer's Country/Territory: China

Author's Country/Territory: Brazil

Manuscript submission date: 2022-08-28

Reviewer chosen by: Dong-Mei Wang

Reviewer accepted review: 2022-11-09 00:18

Reviewer performed review: 2022-11-21 13:15

Review time: 12 Days and 12 Hours

Scientific quality	<input type="checkbox"/> Grade A: Excellent <input type="checkbox"/> Grade B: Very good <input type="checkbox"/> Grade C: Good <input checked="" type="checkbox"/> Grade D: Fair <input type="checkbox"/> Grade E: Do not publish
Language quality	<input type="checkbox"/> Grade A: Priority publishing <input type="checkbox"/> Grade B: Minor language polishing <input checked="" type="checkbox"/> Grade C: A great deal of language polishing <input type="checkbox"/> Grade D: Rejection
Conclusion	<input type="checkbox"/> Accept (High priority) <input type="checkbox"/> Accept (General priority) <input type="checkbox"/> Minor revision <input checked="" type="checkbox"/> Major revision <input type="checkbox"/> Rejection
Re-review	<input checked="" type="checkbox"/> Yes <input type="checkbox"/> No

Peer-reviewer statements	Peer-Review: [<input checked="" type="radio"/>] Anonymous [<input type="radio"/>] Onymous
	Conflicts-of-Interest: [<input type="radio"/>] Yes [<input checked="" type="radio"/>] No

SPECIFIC COMMENTS TO AUTHORS

The manuscript entitled " Three-dimensional model of antigens with serodiagnosis potential for leprosy: an insilico study" reports the three-dimensional structure prediction results of four proteins (ML2038, ML0286, ML0050 and Antigen 85B). (1) The method part is too simple. What is the standard for template search? (2) The template Identity of one protein is only 40%, which is too low. (3) The simulated structure obtained by the authors should be optimized, and molecular dynamics simulation is necessary. (4) The AlphaFold Protein Structure Database also has prediction results for these proteins, whether the author's results are better than them. (5) The language of the manuscript needs to be improved.

RE-REVIEW REPORT OF REVISED MANUSCRIPT

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Professional title: Academic Research

Reviewer's Country/Territory: China

Author's Country/Territory: Brazil

Manuscript submission date: 2022-08-28

Reviewer chosen by: Ji-Hong Liu

Reviewer accepted review: 2022-12-30 07:46

Reviewer performed review: 2022-12-30 08:06

Review time: 1 Hour

Scientific quality	<input type="checkbox"/> Grade A: Excellent <input type="checkbox"/> Grade B: Very good <input checked="" type="checkbox"/> Grade C: Good <input type="checkbox"/> Grade D: Fair <input type="checkbox"/> Grade E: Do not publish
Language quality	<input type="checkbox"/> Grade A: Priority publishing <input checked="" type="checkbox"/> Grade B: Minor language polishing <input type="checkbox"/> Grade C: A great deal of language polishing <input type="checkbox"/> Grade D: Rejection
Conclusion	<input type="checkbox"/> Accept (High priority) <input checked="" type="checkbox"/> Accept (General priority) <input type="checkbox"/> Minor revision <input type="checkbox"/> Major revision <input type="checkbox"/> Rejection
Peer-reviewer	Peer-Review: <input checked="" type="checkbox"/> Anonymous <input type="checkbox"/> Onymous



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statements

Conflicts-of-Interest: [] Yes [Y] No

SPECIFIC COMMENTS TO AUTHORS

The authors have addressed all my concerns.