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PEER-REVIEW REPORT

Name of journal: *World Journal of Clinical Cases*

Manuscript NO: 70362

Title: Pulmonary Cladosporium infection coexisting with subcutaneous Corynespora cassiicola infection in a patient: A case report and literature review

Provenance and peer review: Unsolicited Manuscript; Externally peer reviewed

Peer-review model: Single blind

Reviewer's code: 00188507
Position: Editorial Board
Academic degree: MD, PhD
Professional title: Doctor, Full Professor
Reviewer's Country/Territory: Japan
Author's Country/Territory: China

Manuscript submission date: 2021-07-31

Reviewer chosen by: AI Technique

Reviewer accepted review: 2021-08-01 00:50

Reviewer performed review: 2021-08-01 01:02

Review time: 1 Hour

Scientific quality	[ ] Grade A: Excellent [ ] Grade B: Very good [Y] Grade C: Good [ ] Grade D: Fair [ ] Grade E: Do not publish
Language quality	[ ] Grade A: Priority publishing [ ] Grade B: Minor language polishing [Y] Grade C: A great deal of language polishing [ ] Grade D: Rejection
Conclusion	[ ] Accept (High priority) [ ] Accept (General priority) [ ] Minor revision [Y] Major revision [ ] Rejection
Re-review	[Y] Yes [ ] No

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Peer-reviewer statements	Peer-Review: [Y] Anonymous [ ] Onymous Conflicts-of-Interest: [ ] Yes [Y] No

SPECIFIC COMMENTS TO AUTHORS

The authors reported the case of Cladosporium infection with subcutaneous Corynespora cassiicola infection having CARD9 mutation. It seems the case was in the rare situation and therefore, can be considered for further review in this journal, however, there are several concerns to be clarified prior for the further review. 1. While the authors reported the history and laboratory exams, they should include the information regarding inflammation and immune deficient conditions (like, WBC value, CRP, and HIV antibodies, etc). 2. The authors may want to include the images of subcutaneous lesions for showing the clinical courses. 3. The information regarding the detecting the mutation is poorly described. They should add the description regarding the methods to detect the mutation.