

PEER-REVIEW REPORT

Name of journal: *World Journal of Clinical Cases*

Manuscript NO: 70331

Title: Mixed Porokeratosis With a Novel MVK Gene Mutation: A Case Report and Literature Review

Provenance and peer review: Unsolicited manuscript; externally peer reviewed

Peer-review model: Single blind

Reviewer's code: 00731613

Position: Editorial Board

Academic degree: MD, PhD

Professional title: Associate Professor

Reviewer's Country/Territory: India

Author's Country/Territory: China

Manuscript submission date: 2021-07-30

Reviewer chosen by: Ze-Mao Gong

Reviewer accepted review: 2021-09-03 15:01

Reviewer performed review: 2021-09-12 13:42

Review time: 8 Days and 22 Hours

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 Grade E: Do not publish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 type="checkbox"/> Accept (General priority) <input checked="" type="checkbox"/> Minor revision <input type="checkbox"/> Major revision <input type="checkbox"/> Rejection
Re-review	<input checked="" type="checkbox"/> Yes <input type="checkbox"/> No

**Peer-reviewer
statements**

Peer-Review: [☒] Anonymous [☐] Onymous

Conflicts-of-Interest: [☐] Yes [☒] No

SPECIFIC COMMENTS TO AUTHORS

1. The authors have mentioned in the manuscript that genomic DNA from 100 normal individuals were extracted to act as normal controls. Can the authors elaborate on this? Why was the genomic data from 100 individuals needed? Were they taken from the archives or the genomic DNA analysis was done for this case only? 2. A discussion on the various therapies, the use of genetic analysis in affected families to predict the occurrence of these lesions in other individuals of the same family can be added in the manuscript. 3. Does the variation in the genetic involvement affect the prognosis? 4. A follow-up data in the present case, if available can be added.