

PEER-REVIEW REPORT

Name of journal: *World Journal of Clinical Cases*

Manuscript NO: 71782

Title: Targeted next-generation sequencing identifies a novel nonsense mutation in ANK1 for hereditary spherocytosis: a case report

Provenance and peer review: Unsolicited Manuscript; Externally peer reviewed

Peer-review model: Single blind

Reviewer's code: 03353662

Position: Peer Reviewer

Academic degree: PhD

Professional title: Senior Researcher

Reviewer's Country/Territory: United Kingdom

Author's Country/Territory: China

Manuscript submission date: 2021-09-22

Reviewer chosen by: AI Technique

Reviewer accepted review: 2021-09-30 21:02

Reviewer performed review: 2021-10-10 13:03

Review time: 9 Days and 16 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 Grade E: Do not publish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 type="checkbox"/> Yes <input checked="" type="checkbox"/> No



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**Peer-reviewer
statements**

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

Conflicts-of-Interest: [☐] Yes [☒] No

SPECIFIC COMMENTS TO AUTHORS

The following to be reviewed 1. p3 line 61 hematopathy p77 unsaturated iron.
Please review terminology and spelling More details on the instrumentation used for
NGS

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Reviewer's code: 01221812

Position: Peer Reviewer

Academic degree: MPhil, PhD

Professional title: Academic Research, Doctor, Research Scientist, Teacher

Reviewer's Country/Territory: Pakistan

Author's Country/Territory: China

Manuscript submission date: 2021-09-22

Reviewer chosen by: Xin Liu (Online Science Editor)

Reviewer accepted review: 2021-12-17 10:20

Reviewer performed review: 2021-12-24 17:13

Review time: 7 Days and 6 Hours

Scientific quality	<input type="checkbox"/> Grade A: Excellent <input checked="" type="checkbox"/> Grade B: Very good <input 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: [<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

Targeted next-generation sequencing identifies a novel nonsense mutation in ANK1 for hereditary spherocytosis: a case report and a brief review of literature Fu et al. recruit an infant with hereditary spherocytosis (HS) which is characterized by anemia, jaundice, splenomegaly and cholelithiasis. The authors carried out targeted next-generation sequencing and identified a novel nonsense mutation in ANK1. This is an interesting study, however, the manuscript would benefit from the following changes/corrections:

1. Please mention OMIM number with each malformation and gene reported in the text.
2. Please mention the prevalence estimate of Hereditary spherocytosis in the global populations and the study populations.
3. Unclear statement, "a suspicious positive family history, and then finally confirmed by genetic sequencing.". Please elaborate the statement. What is the phenotypic presentation in the father of the index person.
4. The protocol of next-generation sequencing is not described. Please give detail of library preparation, controls, cut-off values, assembly used, etc.
5. The variant filtration strategy has not been given.
6. It is not clear who many rare variants were left in the final filtration scheme.
7. Please give the allele frequency of the variant as reported in public databases like 1000G and ExAC.
8. Discussion part is weak and redundant. Second paragraph of Disucussion may be shifted to Introduction.
9. Strengths and limitations of the study are not given.

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Peer-review model: Single blind

Reviewer's code: 00069130

Position: Editorial Board

Academic degree: BM BCh, PhD

Professional title: Academic Fellow, Assistant Professor

Reviewer's Country/Territory: India

Author's Country/Territory: China

Manuscript submission date: 2021-09-22

Reviewer chosen by: Xin Liu (Online Science Editor)

Reviewer accepted review: 2021-12-21 06:25

Reviewer performed review: 2021-12-27 03:45

Review time: 5 Days and 21 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 checked="" type="checkbox"/> Grade A: Priority publishing <input 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
Re-review	<input checked="" type="checkbox"/> Yes <input type="checkbox"/> No



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Peer-reviewer statements	Peer-Review: <input checked="" type="checkbox"/> Anonymous <input type="checkbox"/> Onymous Conflicts-of-Interest: <input type="checkbox"/> Yes <input checked="" type="checkbox"/> No
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SPECIFIC COMMENTS TO AUTHORS

The manuscript is well written and may be accepted for publication. The authors may include in the discussion a section on the mutation-nonsense mutation (exon23:c.G2467T;p.E823X) of ANK1. This would help the general readers about the location, impact and consequences of the mutation. Unsaturated iron on page 77.