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PEER-REVIEW REPORT

Name of journal: World Journal of Clinical Cases

Manuscript NO: 88220

Title: Mental retardation, seizures and language delay caused by new SETD1B

mutations: Three case reports

Provenance and peer review: Unsolicited manuscript; Externally peer reviewed

Peer-review model: Single blind

Reviewer's code: 01221812 Position: Peer Reviewer

Academic degree: MPhil, PhD

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

Reviewer's Country/Territory: Pakistan

Author's Country/Territory: China

Manuscript submission date: 2023-09-19

Reviewer chosen by: Yu-Lu Chen

Reviewer accepted review: 2023-11-09 15:18

Reviewer performed review: 2023-11-12 15:55

Review time: 3 Days

	[] Grade A: Excellent [Y] Grade B: Very good [] Grade C:
Scientific quality	Good
	[] Grade D: Fair [] Grade E: Do not publish
Novelty of this manuscript	[] Grade A: Excellent [Y] Grade B: Good [] Grade C: Fair [] Grade D: No novelty
Creativity or innovation of	[] Grade A: Excellent [Y] Grade B: Good [] Grade C: Fair
this manuscript	[] Grade D: No creativity or innovation



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Scientific significance of the conclusion in this manuscript	[] Grade A: Excellent [Y] Grade B: Good [] Grade C: Fair [] Grade D: No scientific significance
Language quality	[Y] Grade A: Priority publishing [] Grade B: Minor language polishing [] Grade C: A great deal of language polishing [] Grade D: Rejection
Conclusion	[] Accept (High priority) [] Accept (General priority) [Y] Minor revision [] Major revision [] Rejection
Re-review	[Y]Yes []No
Peer-reviewer statements	Peer-Review: [Y] Anonymous [] Onymous Conflicts-of-Interest: [] Yes [Y] No

SPECIFIC COMMENTS TO AUTHORS

Clinical analysis of three cases with mental retardation, seizures and language delay caused by new SETD1B mutations This is an interesting study. The manuscript would however, benefit from the following corrections: 1. Please italicize the names of genes. 2. Please provide the OMIM number of the malformations given in the text. 3. Please mention the variant filtration strategy for WES. 4. It is not clear how many variants were filtered in the last step. 5. Fig. 2. Please translate the Chinese legends/labels of the figures. 6. Please give the allele frequencies of the detected variants as reported in public databases. 7. The clinical symptoms of the patients should be presented in a comparative table showing all cases side by side. 8. It is not clear if there was parental consanguinity was present in these cases. 9. Please give the differential diagnosis of the phenotypes in the patients. 10. Please give the basic demographic information of the study subjects.