Author’s response to reviews

Title: Case reports: three Novel variants in PCCA and PCCB Genes in Chinese Patients with propionic acidemia

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Author’s response to reviews:

Dear Editor:

We greatly appreciate this opportunity to revise our manuscript. We would like to thank the reviewers for their kind comments regarding our work and for their efforts in reviewing this paper. We have carefully read the editor’s comments regarding our submission (MGTC-D-19-00561R3), and have responded point by point to each of the editor’s comments as listed below, and the revised portions in the paper. We feel that the editor’s comments have greatly helped us to strengthen our manuscript and we hope that the revised version will be acceptable for publication in BMC Medical Genetics.

Sincerely yours,

QI Yang.

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The Maternal and Child Health Hospital of Guangxi Zhuang Autonomous Region
Editor Comments:

1. Consent for publication

Please remove the following statement from the Consent for publication section of your manuscript: “This study was approved by the Institutional Review Boards and Ethics Committees of Guangxi Maternal and Child Health Hospital.”

Response: Thanks for the excellent suggestion. We have revised it.

2. Availability of Data and Materials

Please note that all data must be made public prior to publication and we cannot proceed to publish your manuscript if your data is not deposited and made publicly available. We are an open access journal and submission of a manuscript to a BMC journal implies that materials described in the manuscript, including all relevant raw data, will be freely available to any scientist wishing to use them for non-commercial purposes. As your data falls under the list of data types which must be deposited (listed below), we ask that you please provide the relevant accession numbers, and the name of the database in which your data is stored, if your data has been deposited into a database or the appropriate web links if the data has been uploaded into a repository. If the data has not yet been deposited, please do so and provide the relevant information needed to access it in the ‘Availability of data and materials’ section in your manuscript. Please ensure all data is already released and that any accession numbers/web links are in the ‘Availability of Data and Materials’ in their final form, and that all web links and accessions allow access to public data.

**Please note that BMC mandates data deposition for the following:

- Protein sequences – can be deposited in Uniprot
- DNA and RNA sequences – can be deposited in Genbank, DNA DataBank of Japan (DDBJ), or EMBL or Nucleotide Sequence Database (ENA)
- DNA and RNA sequencing data – can be deposited in NCBI Trace Archive or NCBI Sequence Read Archive (SRA)
• Genetic polymorphisms – can be deposited in NCBI dbSNP, NCBI dbVar, or European Variation Archive (EVA)

• Linked genotype and phenotype data – can be deposited in NCBI dbGAP or The European Genome-phenome Archive (EGA)

• Macromolecular structure – can be deposited in the Worldwide Protein Data Bank (wwPDB) or the Biological Magnetic Resonance Data Bank (BMRB)

• Microarray data (must be MIAME compliant) – can be deposited in Gene Expression Omnibus (GEO) or ArrayExpress

• Crystallographic data for small molecules – can be deposited in Cambridge Structural Database

**Please also ensure that you include the accession numbers and the full names of the data banks/repositories corresponding to all of the data generated and/or analysed in your study in the Availability of Data and Materials section. If accession numbers are listed in a table or additional file, please include an in-text reference to the file or table in the Availability of Data and Materials section.

Response:The datasets generated and/or analysed during the current study are available in the [https://pan.baidu.com/] repository, [https://pan.baidu.com/s/1vDtsnFRet8E50AFIlvvGmw], access code:ssqh. Usename: shengwjqiji; password: SHENGWUQIIq12.

3. Authors' Contributions

Please amend the initials “XH” to “HX”.

Response: Thanks for the excellent suggestion. We have revised it.

4. At this stage, please upload your manuscript as a single, final, clean version that does not contain any tracked changes, comments, highlights, strikethroughs or text in different colours. All relevant tables/figures/additional files should also be clean versions. Should you wish to respond to these revision requests, please put your responses to the reviewers'/editors’ comments in the Response to Reviewers box in Editorial Manager. Please do not upload a separate letter.

Response: Thanks for the excellent suggestion. We have uploaded our manuscript as necessary.