Author’s response to reviews

Title: Association analysis between the tag single nucleotide polymorphisms of DENND1A and the risk of polycystic ovary syndrome in Chinese Han women

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Author’s response to reviews:

Dear Editor,

Re: MGTC-D-19-00211R5 (Research Article) Zhu et al "Association analysis between the tag single nucleotide polymorphisms of DENND1A and the risk of polycystic ovary syndrome in Chinese Han women".

We would like to thank the editor for their final requests. Editor’s requests are responded as follows. We hereby resubmit our revised manuscript to BMC Medical Genetics.

Editor’s suggestions are responded as follows. We hope that the manuscript can now be accepted for publication in this journal. We are looking forward to your favorable decision.

1. Availability of data and materials
Response: As the editor’s comments, we have deposited the data in NCBI BioSample & BioProject. The accession numbers are in the ‘Availability of Data and Materials’ in the revised manuscript.
"The datasets generated and analysed during the current study are available in the NCBI BioSample & BioProject database repository. The data are accessible via the accession number: SAMN13640133 & PRJNA596822: human Phenotype or genotype (TaxID: 9606)."

2. 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:
We've gone through the manuscript as required. Now the manuscript is a single, final, clean version that does not contain any tracked changes, comments, highlights, strikethroughs or text in different colours. If there is any dirty or something else, please let us know. Thank you very much.

Yours sincerely,
On behalf of my coauthors
Yong Wang Ph.D.