Author’s response to reviews

Title: Contiguous 22.1-kb deletion embracing AVPR2 and ARHGAP4 genes at novel breakpoints leads to nephrogenic diabetes insipidus in a Chinese pedigree

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1. Thank you for providing an Availability of data and materials heading for the Declarations section. Please include the full name of the database where raw sequences have been deposited.

Response: The statement has been added as following:

“The raw sequences are saved in the private human mutation database of the First Affiliated Hospital, Zhengzhou University.” (Page 9, Line 8-9)

2. Thank you for including a Funding heading for the Declarations section. Please also state the role of the funding body in the design of the study and collection, analysis, and interpretation of data and in writing the manuscript.

Response: The statement has been added as following:

“None of the funding bodies participated in the design of the study and collection, analysis, and interpretation of data and in writing the manuscript.” (Page 9, Line 16-17)