

CORRECTION

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# Correction: Genome-wide allele and haplotype-sharing patterns suggested one unique Hmong–Mein-related lineage and biological adaptation history in Southwest China

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Following publication of the original article [1], the authors reported that the name of Guanglin He needed to be moved from the last to the first of the authorship.

The correct authorship has been provided in this correction.

The original article [1] has been corrected.

## Reference

1. He G, Wang J, Yang L, et al. Genome-wide allele and haplotype-sharing patterns suggested one unique Hmong–Mein-related lineage and biological adaptation history in Southwest China. *Hum Genomics*. 2023;17:3. <https://doi.org/10.1186/s40246-023-00452-0>.

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