

CORRECTION

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Correction: Evaluation of in silico pathogenicity prediction tools for the classification of small in-frame indels

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Correction: *BMC Med Genomics* 16, 36 (2023)

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Following publication of the original article [1], the authors reported errors in additional file 2. Some of the HGVS annotations in this file were incorrect, affecting two columns in the table. These annotations were not used for variant classification or pathogenicity prediction, so would not have affected the results presented in the paper. The updated additional file is supplied in this correction article. The original article [1] has been corrected.

Supplementary Information

The online version contains supplementary material available at <https://doi.org/10.1186/s12920-023-01732-3>.

Additional file 2: Table S1. Inframe-indels and pathogenicity predictions from tools used in this study

References

1. Cannon S, Williams M, Gunning AC, et al. Evaluation of in silico pathogenicity prediction tools for the classification of small in-frame indels. *BMC Med Genomics*. 2023;16:36. <https://doi.org/10.1186/s12920-023-01454-6>.

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