Genome Biology

AUTHOR CORRECTION

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Author Correction: Cross-phyla protein annotation by structural prediction and alignment

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The original article can be found online at https://doi.org/10.1186/s13059-023-02942-9.

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Following publication of the original article [1], the authors identified two errors in the citations.

First, in citation 109, the predicted *S. lacustris* protein structures are not deposited under https://doi.org/10.5452/ma-yoep2 but rather https://doi.org/10.5452/ma-coffe-slac.

Second, the Biological Structure Model Archive was cited incorrectly in place of the ModelArchive; citation 11 should therefore point to:

Schwede T, Sali A, Honig B, Levitt M, Berman HM, Jones D, Brenner SE, Burley SK, Das R, Dokholyan NV, Dunbrack RL Jr, Fidelis K, Fiser A, Godzik A, Huang YJ, Humblet C, Jacobson MP, Joachimiak A, Krystek SR Jr, Kortemme T, Kryshtafovych A, Montelione GT, Moult J, Murray D, Sanchez R, Sosnick TR, Standley DM, Stouch T, Vajda S, Vasquez M, Westbrook JD, Wilson IA. Outcome of a workshop on applications of protein models in biomedical research. Structure. 2009 Feb 13;17(2):151–9. https://doi.org/10.1016/j.str.2008.12.014. PMID: 19217386; PMCID: PMC2739730.

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Reference
1. Ruperti F, Papadopoulos N, Musser JM, et al. Cross-phyla protein annotation by structural prediction and alignment. Genome Biol. 2023;24:113. https://doi.org/10.1186/s13059-023-02942-9.

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