

CORRECTION

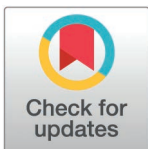
Correction: Computational analysis of the deleterious non-synonymous single nucleotide polymorphisms (nsSNPs) in TYR gene impacting human tyrosinase protein and the protein stability

Wei Fan, Heng Li Ji, Mohibullah Kakar, Shabbir Ahmed, Hussah M. Alobaid, Yasmeen Shakir

There are errors in the Funding statement. The correct Funding statement is as follows: This work is funded by the Research Supporting Project Number (RSP2025R174), King Saud University, Riyadh, Saudi Arabia.

Reference

1. Fan W, Ji HL, Kakar M, Ahmed S, Alobaid HM, Shakir Y. Computational analysis of the deleterious non-synonymous single nucleotide polymorphisms (nsSNPs) in TYR gene impacting human tyrosinase protein and the protein stability. PLoS One. 2024;19(11): e0308927. <https://doi.org/10.1371/journal.pone.0308927> PMID: [39541331](https://pubmed.ncbi.nlm.nih.gov/39541331/)



OPEN ACCESS

Citation: Fan W, Ji HL, Kakar M, Ahmed S, Alobaid HM, Shakir Y (2025) Correction: Computational analysis of the deleterious non-synonymous single nucleotide polymorphisms (nsSNPs) in TYR gene impacting human tyrosinase protein and the protein stability. PLoS ONE 20(3): e0321098. <https://doi.org/10.1371/journal.pone.0321098>

Published: March 25, 2025

Copyright: © 2025 Fan et al. This is an open access article distributed under the terms of the [Creative Commons Attribution License](https://creativecommons.org/licenses/by/4.0/), which permits unrestricted use, distribution, and reproduction in any medium, provided the original author and source are credited.