

Corrigenda and Addenda

Correction: Structural and Functional Impacts of SARS-CoV-2 Spike Protein Mutations: Insights From Predictive Modeling and Analytics

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In “Structural and Functional Impacts of SARS-CoV-2 Spike Protein Mutations: Insights From Predictive Modeling and Analytics” [1], the authors noted four errors.

Author JAJ has now been marked as an equal contributor, as indicated by the asterisk added to their name.

Affiliation 1 has been changed from the following:

Department of Mathematics and Information Communication Technology, School of Physical Sciences, Dambai College of Education, Dambai, Ghana

The affiliation now reads:

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Affiliation 11 has been changed from the following:

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The corrections will appear in the online version of the paper on the JMIR Publications website, together with the publication of this correction notice. Because these were made after submission to PubMed, PubMed Central, and other full-text repositories, the corrected article has also been resubmitted to those repositories.

References

1. Netsey EK, Naandam SM, Asante Jnr J, et al. Structural and functional impacts of SARS-CoV-2 spike protein mutations: insights from predictive modeling and analytics. JMIR Bioinform Biotech. 2025;6:e73637. [doi: [10.2196/73637](https://doi.org/10.2196/73637)] [Medline: [12685290](https://pubmed.ncbi.nlm.nih.gov/3685290/)]

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