



Correction

Correction: Oluwagbemi et al. Bioinformatics, Computational Informatics, and Modeling Approaches to the Design of mRNA COVID-19 Vaccine Candidates. *Computation* 2022, 10, 117

Olugbenga Oluseun Oluwagbemi ^{1,2,3,*}, Elijah K. Oladipo ^{4,5}, Olatunji M. Kolawole ⁶, Julius K. Oloke ⁷, Temitope I. Adelusi ⁸, Boluwatife A. Irewolede ⁵, Emmanuel O. Dairo ^{5,9}, Ayodele E. Ayeni ^{5,10}, Kehinde T. Kolapo ⁵, Olawumi E. Akindiya ^{5,11}, Jerry A. Oluwasegun ⁵, Bamigboye F. Oluwadara ⁵ and Segun Fatumo ^{12,13,*}

- Department of Computer Science and Information Technology, Faculty of Natural and Applied Sciences, Sol Plaatje University, Kimberley 8301, South Africa
- Department of Mathematical Sciences, Stellenbosch University, Stellenbosch 7602, South Africa
- National Institute of Theoretical and Computational Sciences (NiThECs), Stellenbosch 7602, South Africa
- ⁴ Laboratory of Molecular Biology, Immunology and Bioinformatics, Department of Microbiology, Adeleke University, Ede 232104, Nigeria
- Genomics Unit, Helix Biogen Institute, Ogbomoso 210214, Nigeria
- ⁶ Department of Microbiology, University of Ilorin, Ilorin 234031, Nigeria
- Department of Natural Science, Precious Cornerstone University, Ibadan 200223, Nigeria
- ⁸ Computational Biology/Drug Discovery Laboratory, Biochemistry Department, Ladoke Akintola University of Technology, (LAUTECH), Ogbomoso 210214, Nigeria
- Department of Virology, College of Medicine, University of Ibadan, Ibadan 200132, Nigeria
- Department of Medical Microbiology and Parasitology, University of Ibadan, Ibadan 200132, Nigeria
- Microbiology Programme, Department of Biological Science, Olusegun Agagu University of Science and Technology, Okitipupa 350113, Nigeria
- The African Computational Genomics (TACG) Research Group, MRC/UVRI and LSHTM, Entebbe 7545, Uganda
- Department of Non-Communicable Disease Epidemiology (NCDE), London School of Hygiene and Tropical Medicine, London WC1E 7HT, UK
- * Correspondence: olugbenga.oluwagbemi@fulbrightmail.org (O.O.O.); segun.fatumo@lshtm.ac.uk (S.F.); Tel.: +27-663926506 (O.O.O.)

check for **updates**

Citation: Oluwagbemi, O.O.;
Oladipo, E.K.; Kolawole, O.M.; Oloke, J.K.; Adelusi, T.I.; Irewolede, B.A.;
Dairo, E.O.; Ayeni, A.E.; Kolapo, K.T.;
Akindiya, O.E.; et al. Correction:
Oluwagbemi et al. Bioinformatics,
Computational Informatics, and
Modeling Approaches to the Design
of mRNA COVID-19 Vaccine
Candidates. Computation 2022, 10,
117. Computation 2022, 10, 211.
https://doi.org/10.3390/
computation10120211

Received: 4 November 2022 Accepted: 7 November 2022 Published: 2 December 2022

Publisher's Note: MDPI stays neutral with regard to jurisdictional claims in published maps and institutional affiliations.



Copyright: © 2022 by the authors. Licensee MDPI, Basel, Switzerland. This article is an open access article distributed under the terms and conditions of the Creative Commons Attribution (CC BY) license (https://creativecommons.org/licenses/by/4.0/).

Missing Funding

The following information was missing in the original manuscript [1]: "This work was partly supported by the Wellcome Trust grant number 220740/Z/20/Z to Segun Fatumo".

Acknowledgments

The acknowledgement section of the original manuscript was missing a section. We have updated the corrected manuscript with this missing information as follows: "This work was partly supported by the Wellcome Trust grant number 220740/Z/20/Z to Segun Fatumo".

Update of An Author

In the original publication, the author's name "Oluwasegun J. Ayobami" was wrong. The corrected Author's name should be "Jerry A. Oluwasegun". The corrected Author Contributions statement updates here:

Conceived and conceptualized: O.O.O.; study design: O.O.O., E.K.O. and S.F.; methodology: O.O.O., E.K.O., B.A.I., E.O.D., A.E.A., K.T.K., O.E.A., J.A.O. and B.F.O.; curation of data: O.O.O., E.K.O., B.A.I., E.O.D., O.M.K., J.K.O., T.I.A., A.E.A., K.T.K., O.E.A., J.A.O. and B.F.O.; data analysis: O.O.O., E.K.O., B.A.I., E.O.D., O.M.K., J.K.O., T.I.A., A.E.A., K.T.K., O.E.A., J.A.O. and B.F.O.; resources: O.O.O.; writing—original draft preparation: O.O.O.; writing—critical review, revised version, and re-editing: O.O.O. and S.F.; project

Computation **2022**, 10, 211 2 of 2

administration: O.O.O. and E.K.O.; funding acquisition: O.O.O.; principal investigator: O.O.O. The funders had no role in study design, data collection and analysis, decision to publish, or preparation of the manuscript. All authors have read and agreed to the published version of the manuscript.

References

Oluwagbemi, O.O.; Oladipo, E.K.; Kolawole, O.M.; Oloke, J.K.; Adelusi, T.I.; Irewolede, B.A.; Dairo, E.O.; Ayeni, A.E.; Kolapo, K.T.; Akindiya, O.E.; et al. Bioinformatics, Computational Informatics, and Modeling Approaches to the Design of mRNA COVID-19 Vaccine Candidates. Computation 2022, 10, 117. [CrossRef]