CORRECTION

Correction: Immunoinformatics approach of epitope prediction for SARS-CoV-2

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Correction: J Genet Eng Biotechnol 20, 60 (2022)
https://doi.org/10.1186/s43141-022-00344-1

Following publication of the original article [1], the authors identified an error in the HTML version of Fig. 1. The publisher apologise for this error. The correct figure is given hereafter.

The original article [1] has been corrected.

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Reference
1. Awad N, Mohamed RH, Ghoneim NI et al (2022) Immunoinformatics approach of epitope prediction for SARS-CoV-2. J Genet Eng Biotechnol 20:60. https://doi.org/10.1186/s43141-022-00344-1

The original article can be found online at https://doi.org/10.1186/s43141-022-00344-1.

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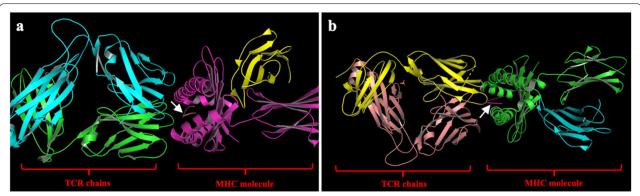Full list of author information is available at the end of the article

Fig. 1 The crystal structures of 5YXN and 4PRP. **a** 5YXN MHC molecule on the right side and TCR chains on the left side. **b** 4PRP MHC molecule on the right side, and TCR chains are on the left side. (white arrows indicate the co-crystalized epitopes)