

CORRECTION

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Correction: Rapid prediction of possible inhibitors for SARS-CoV-2 main protease using docking and FPL simulations

Minh Quan Pham,^{ab} Khanh B. Vu,^{cd} T. Ngoc Han Pham,^e Le Thi Thuy Huong,^{ab} Linh Hoang Tran,^{df} Nguyen Thanh Tung,^{ag} Van V. Vu,^h Trung Hai Nguyen^{ij} and Son Tung Ngo^{*ij}

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In the original article at the time of publication, ref. 44 had only been uploaded to ChemRxiv (DOI: <https://doi.org/10.26434/chemrxiv.12771068.v1>) and had not been published in a finalised format, so the full reference information was not included. The research described in ref. 44 has been published, and the full details of this reference are included below:

44 S. T. Ngo, H. M. Nguyen, L. T. Thuy Huong, P. M. Quan, V. K. Truong, N. T. Tung and V. V. Vu, Assessing potential inhibitors of SARS-CoV-2 main protease from available drugs using free energy perturbation simulations, *RSC Adv.*, 2020, **10**, 40284–40290.

The Royal Society of Chemistry apologises for these errors and any consequent inconvenience to authors and readers.



^aGraduate University of Science and Technology, Vietnam Academy of Science and Technology, Hanoi, Vietnam

^bInstitute of Natural Products Chemistry, Vietnam Academy of Science and Technology, Hanoi, Vietnam

^cSchool of Biotechnology, International University, Ho Chi Minh City, Vietnam

^dVietnam National University, Ho Chi Minh City, Vietnam

^eFaculty of Pharmacy, Ton Duc Thang University, Ho Chi Minh City, Vietnam

^fFaculty of Civil Engineering, Ho Chi Minh University of Technology (HCMUT), Ho Chi Minh, Vietnam

^gInstitute of Materials Science, Vietnam Academy of Science and Technology, Hanoi, Vietnam

^hNTT Hi-Tech Institute, Nguyen Tat Thanh University, Ho Chi Minh City, Vietnam

ⁱLaboratory of Theoretical and Computational Biophysics, Ton Duc Thang University, Ho Chi Minh City, Vietnam. E-mail: ngosontung@tdtu.edu.vn

^jFaculty of Applied Sciences, Ton Duc Thang University, Ho Chi Minh City, Vietnam