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Correction: Rosetta custom score functions accurately predict $\Delta\Delta G$ of mutations at protein–protein interfaces using machine learning

Sumant R. Shringari, † Sam Giannakoulis, † John J. Ferrie * and E. James Petersson *

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

Department of Chemistry, University of Pennsylvania, 231 South 34th Street, Philadelphia, PA 19104, USA. E-mail: jferrie@sas.upenn.edu, ejpetersson@sas.upenn.edu

† These authors contributed equally.

