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CORRECTION

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Correction: A critical comparison of coarse-grained structure-based approaches and atomic models of protein folding

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Correction for 'A critical comparison of coarse-grained structure-based approaches and atomic models of protein folding' by Jie Hu *et al.*, *Phys. Chem. Chem. Phys.*, 2017, **19**, 13629–13639.

It has come to our attention that there is a typographical error in the legend of Fig. 2: "The black solid curve and the shaded band represent $R_g = R_0 \times N^{\nu}$ with $R_0 = 1.330$ and $\nu = 0.598 \pm 0.028$ obtained from fitting experimental data on an extensive set of proteins (ref. 38)" should read "The black solid curve and the shaded band represent $R_g = R_0 \times N^{\nu}$ with $R_0 = 0.1927$ and $\nu = 0.598 \pm 0.028$ obtained from fitting experimental data on an extensive set of proteins (ref. 38)". The value $R_0 = 0.1927$ nm was what we actually used to produce Fig. 2. This value follows the Correction to ref. 38 published in *Proc. Natl. Acad. Sci. U. S. A.*, 2005, **102**, 14475, which states that "the prefactor R_0 is 1.927 Å". This typographical error has no effect on the results presented in our paper.

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

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