

## Correction to Balanced Protein—Water Interactions Improve Properties of Disordered Proteins and Non-Specific Protein Association

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In our paper we did not properly acknowledge the contribution by Head-Gordon and co-workers<sup>2</sup> (ref 41 of our manuscript). These authors used a similar modification of the mixing rules for protein-water interactions, although the correction was applied on a per-atom basis to both the characteristic distance (sigma) and energy (epsilon) parameters rather than using a single scaling factor for epsilon as we did. Head-Gordon and co-workers used extensive solvation free energy data on model organic compounds to fit their parameters, whereas we used FRET efficiencies for a single unfolded protein (cold shock protein) to estimate our scaling parameter. Overall, we reached qualitatively similar conclusions regarding the solvation free energies of force fields with Amber 94<sup>3</sup> Lennard-Jones parameters being too unfavorable, as reported previously by these authors.<sup>2</sup> The more systematic approach of Nerenberg et al. showed a mean signed error of solvation free energies close to zero for their optimized force field.2

We would also like to make a specific correction to the sentence on page 5115, which is inaccurate: "Recently, a full parametrization of protein Lennard-Jones parameters to reproduce solvation free energies showed some promising results, but unfortunately resulted in very unstable folded proteins". Instead, it should read as follows: "Recently, a rescaling of protein—water Lennard-Jones parameters to reproduce solvation free energies showed some promising results for correcting protein—protein interactions using concentrated solutions of glycine and leucine dipeptides. In conjunction with a 12-10 van der Waals (vdW) term to replace the standard 12-6 vdW potential to model the nonelectrostatic part of the hydrogen bonding between the backbone carbonyl oxygens and amide protons, this modification resulted in stable folded proteins."

## **■ REFERENCES**

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- (2) Nerenberg, P. S.; Jo, B.; Tripathy, A.; Head-Gordon, T. J. Phys. Chem. B 2012, 116, 4524.
- (3) Cornell, W. D.; Cieplak, P.; Bayly, C. I.; Gould, I. R.; Merz, K. M.; Ferguson, D. M.; Spellmeyer, D. C.; Fox, T.; Caldwell, J. W.; Kollman, P. A. J. Am. Chem. Soc. 1995, 117, 5179.

