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Erratum

Erratum to "Molecular simulation study of cooperativity in hydrophobic association: clusters of four hydrophobic particles" [Biophys. Chem. 105 (2003) 339–359]

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In the original published article, Figs. 2, 3, 5, 6, and 7 were inadvertently published in black and white with legends referring to colour.

These figures are reproduced here with the correct legends. The publisher apologises for this error.

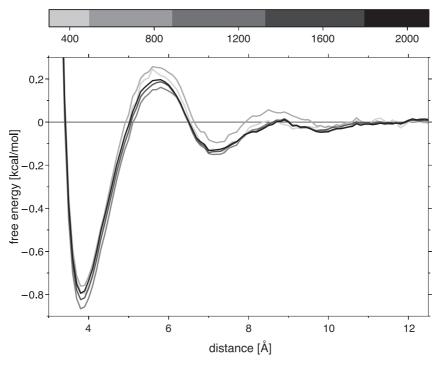


Fig. 2. PMF curves for increasing number of data for the m+m system. The gray-scale in the upper bar denotes the duration of a run (in picoseconds) in each window.

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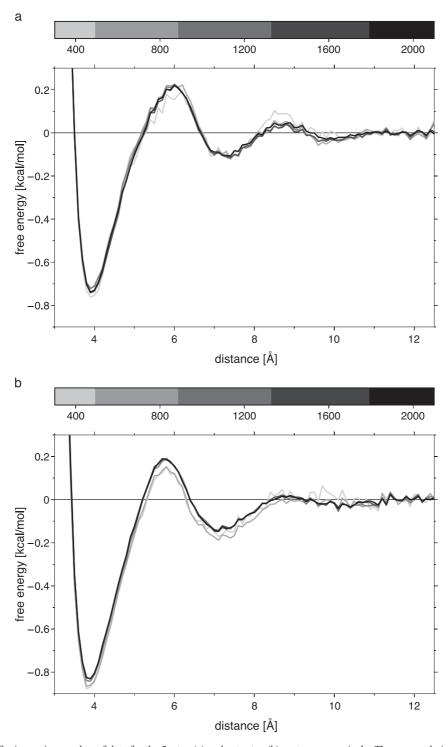


Fig. 3. The PMF curves for increasing number of data for the 2m+m (a) and m+m+m (b) system, respectively. The gray-scale in the upper bar denotes the duration of a run (in picoseconds) in each window.

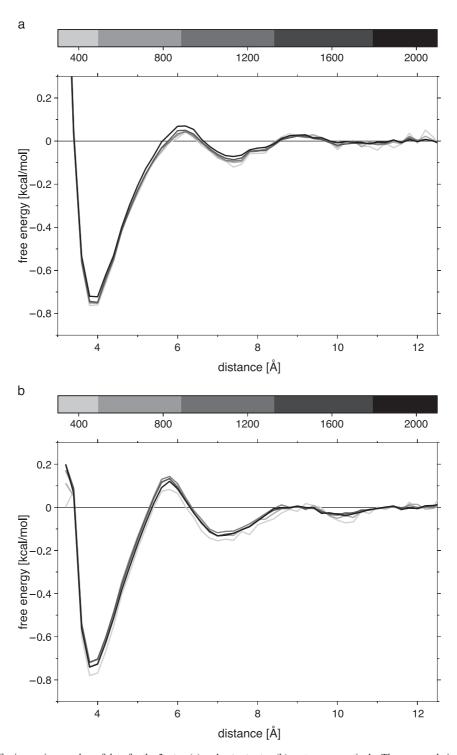


Fig. 5. The PMF curves for increasing number of data for the 3m+m (a) and m+m+m+m (b) system, respectively. The gray-scale in the upper bar denotes the duration of a run (in picoseconds) in each window.

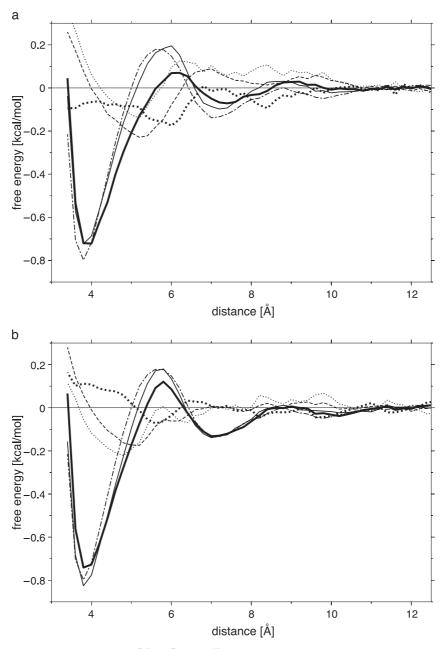


Fig. 6. Curves for the PMF and the cooperative terms $\delta F^{(3,4)}$, $\delta F^{(3)}$ and $\delta F^{(4)}$ for the 3m+m (a) and m+m+m+m (b) system, respectively. Dash–dotted line: dimer PMF, thin solid line: tetramer PMF, dashed line: total cooperative contribution ($\delta F^{(3,4)}$) to the tetramer PMF, thin dotted line: three-body contribution ($\delta F^{(3)}$) to the tetramer PMF, thick dotted line: four-body contribution ($\delta F^{(4)}$) to the tetramer PMF.

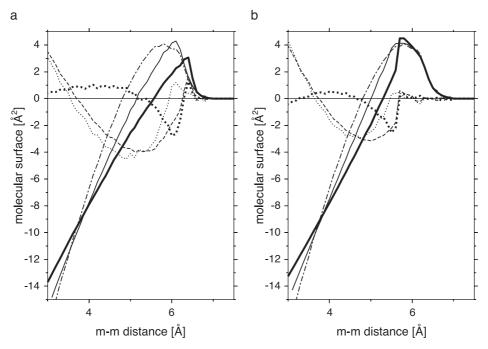


Fig. 7. Distance dependence of the molecular surface area and its decomposition into multibody terms for the 3m+m (a) and m+m+m+m (b) system, respectively. The line scheme is the same as for Fig. 6.