

Supplementary materials for

Calculation of Protein-Ligand Binding Affinities Based on Fragment Quantum Mechanical Method

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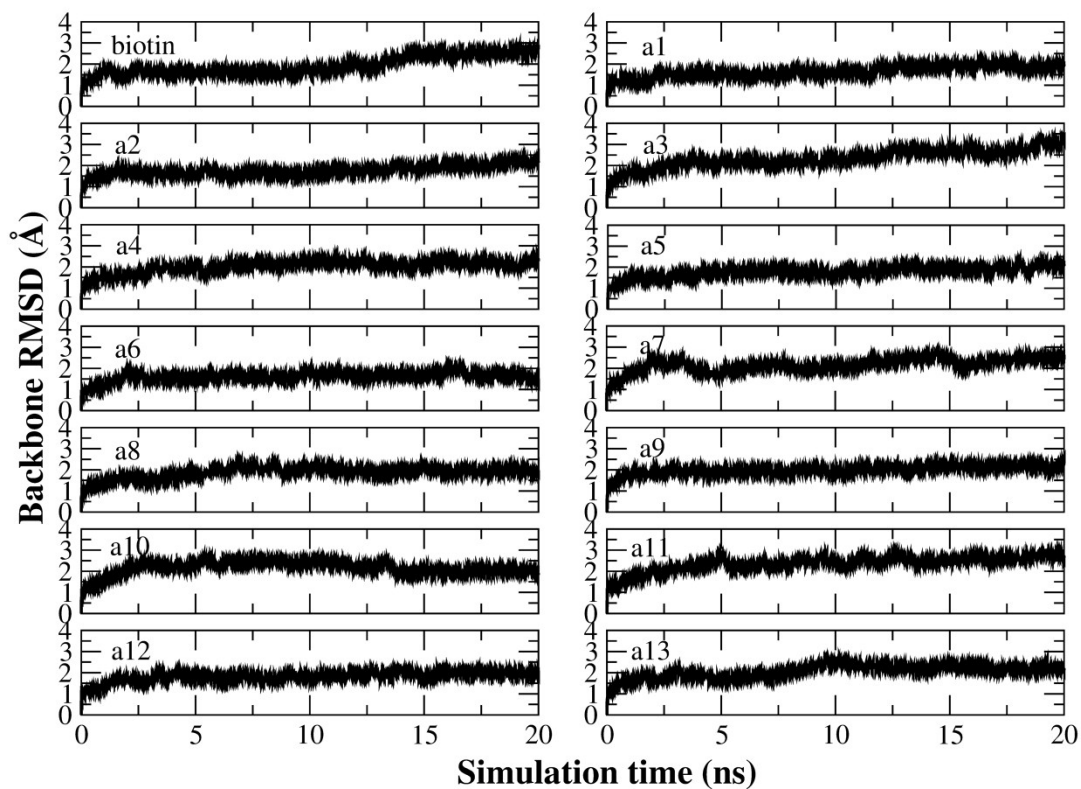


Figure S1. The backbone RMSD of 14 avidin-biotin (and biotin analogues) binding complexes as a function of simulation time.