

Supplementary Materials

Metadynamics supports molecular dynamics simulation-based binding affinities of eucalyptol and beta-cyclodextrins inclusion complexes

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Figure Caption

Fig. S1 RMSD plots of all atoms in the inclusion complex for the five eucalyptol/ β CDs: (a-b) eucalyptol/ β CD, (c-d) eucalyptol/2,6-DM β CD, (e-f) eucalyptol/2-HP β CD, (g-h) eucalyptol/6-HP β CD and (i-j) eucalyptol/2,6-DHP β CD obtained from the other two independent simulations.

Fig. S2 Distances between the centers of mass of eucalyptol and the CD molecules in (a-b) eucalyptol/ β CD, (c-d) eucalyptol/2,6-DM β CD, (e-f) eucalyptol/2-HP β CD, (g-h) eucalyptol/6-HP β CD and (i-j) eucalyptol/2,6-DHP β CD inclusion complexes obtained from the other two independent simulations.

Fig. S3 The distributions of (a) the intramolecular hydrogen bond distances, $O3_{(n)}-O2_{(n+1)}$, and (b) the distances between the adjacent glycosidic oxygens, $O4_{(n)}-O4_{(n+1)}$, of β CDs in ligand bound form.

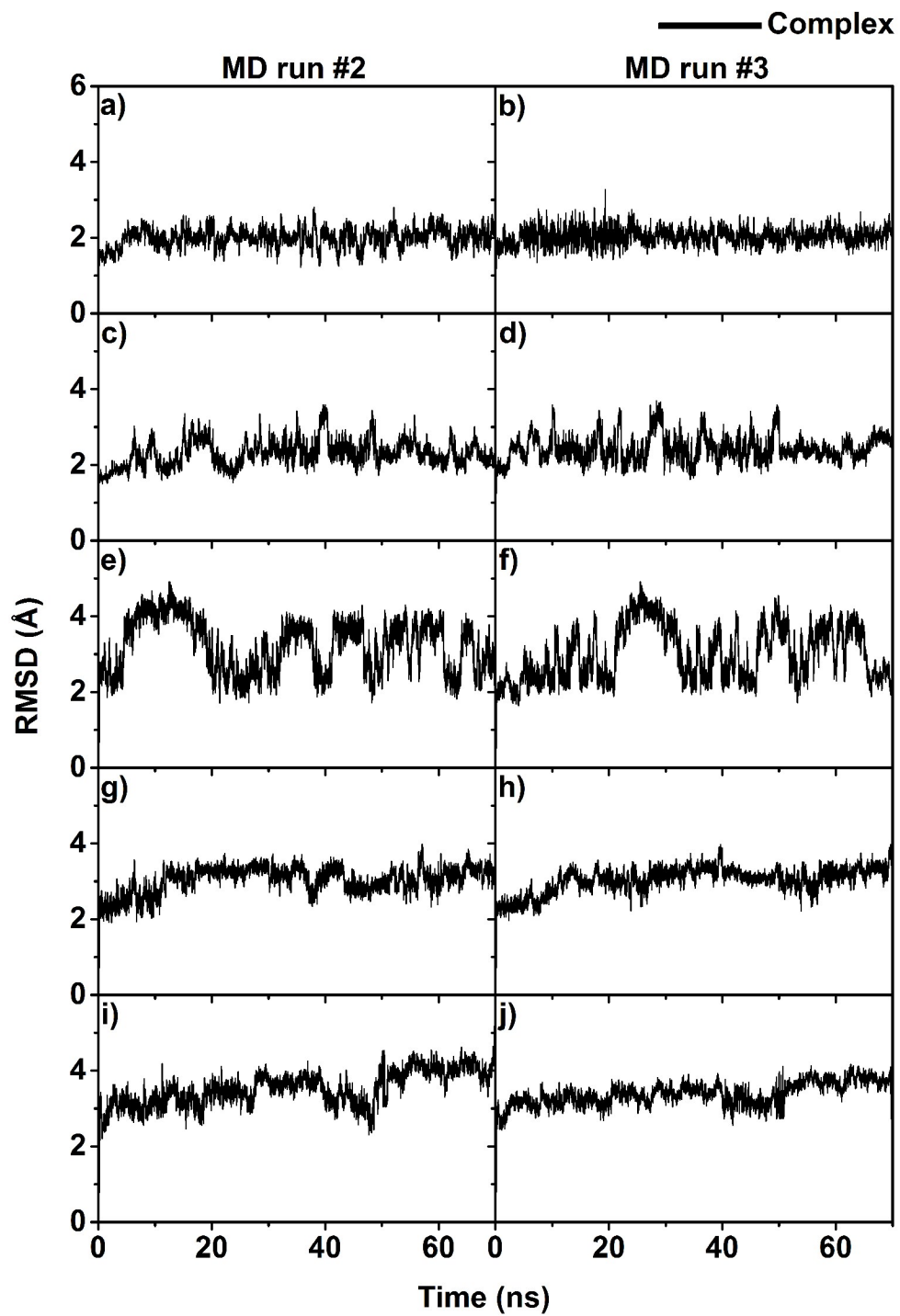


Figure S1

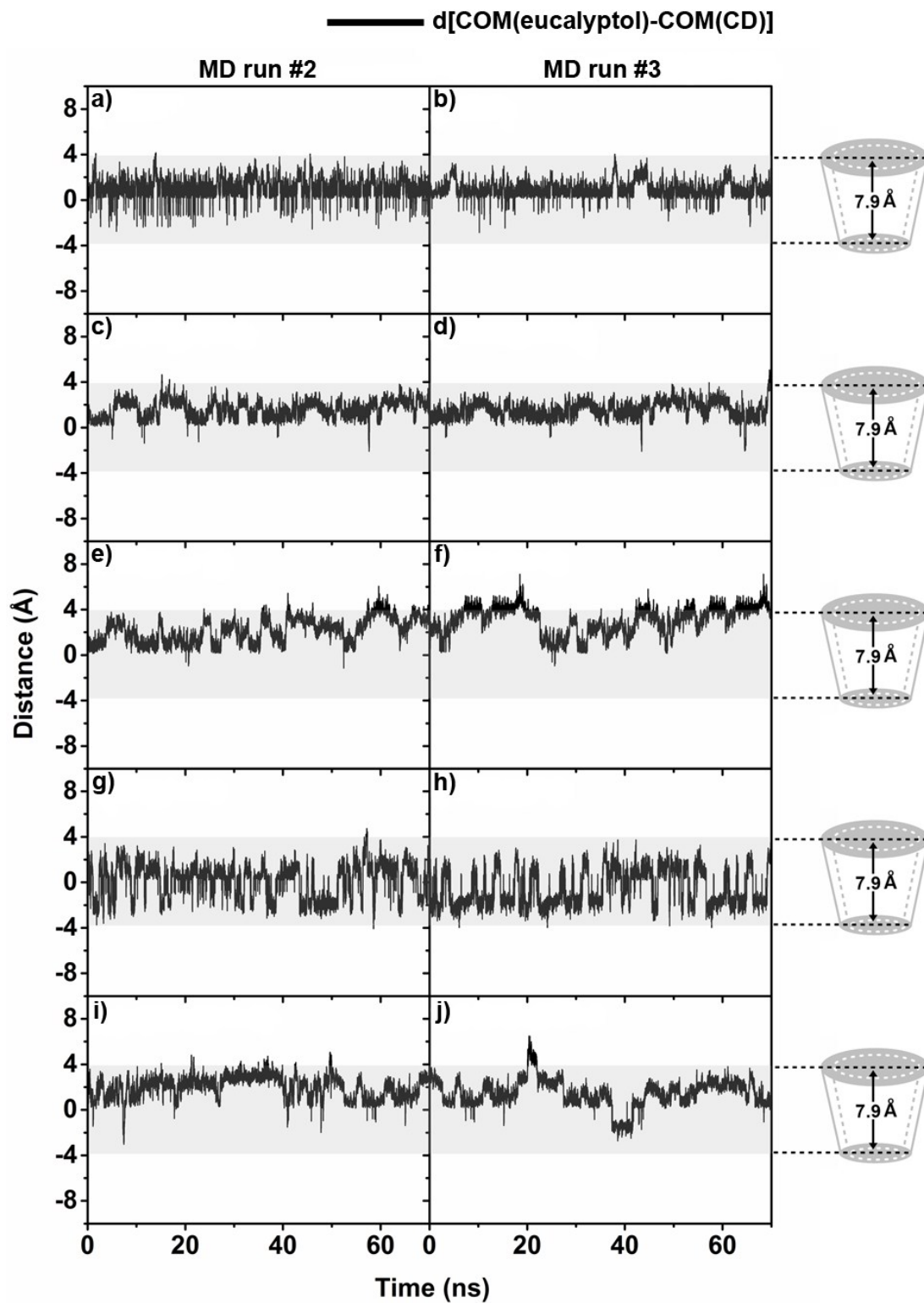


Figure S2

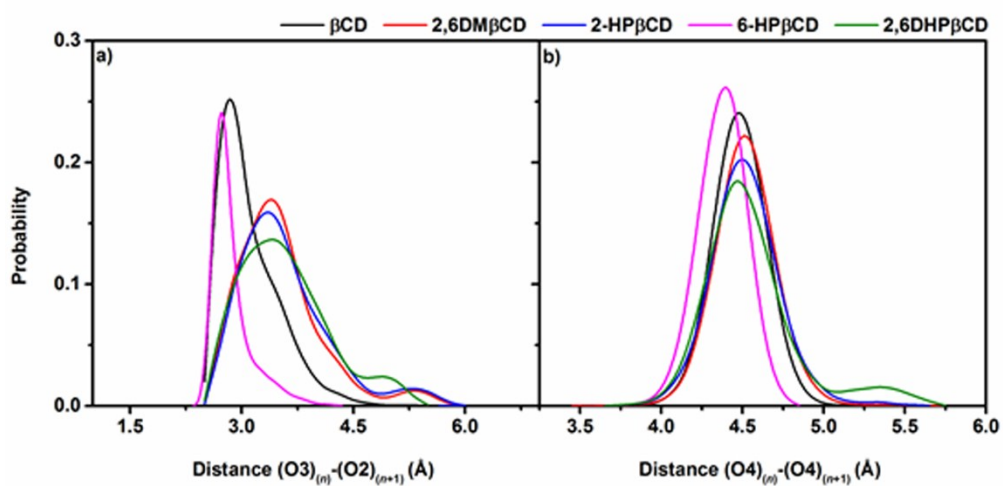
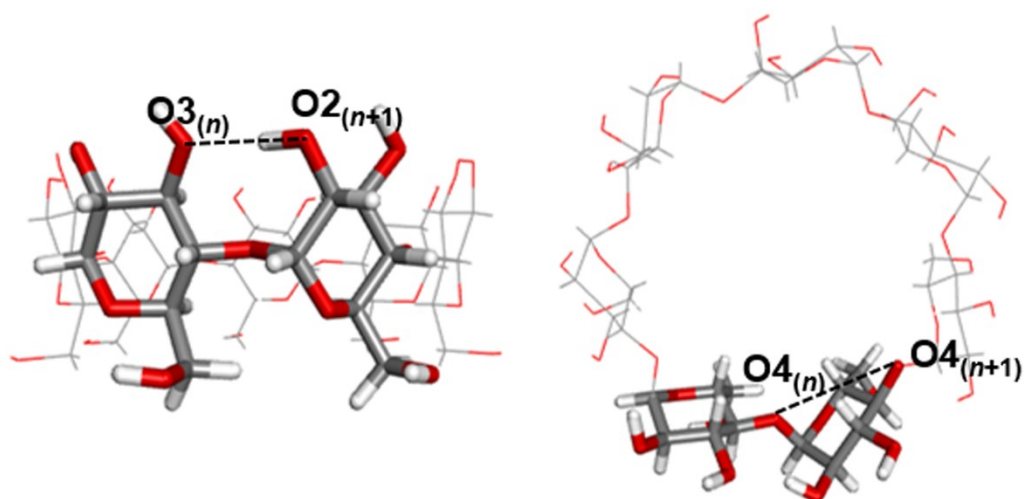


Figure S3