Supplementary Information

Quercetin, Catechin, and Diosmin as Modulators of Haloperidol–HSA Interactions: A Biophysical and Computational Study

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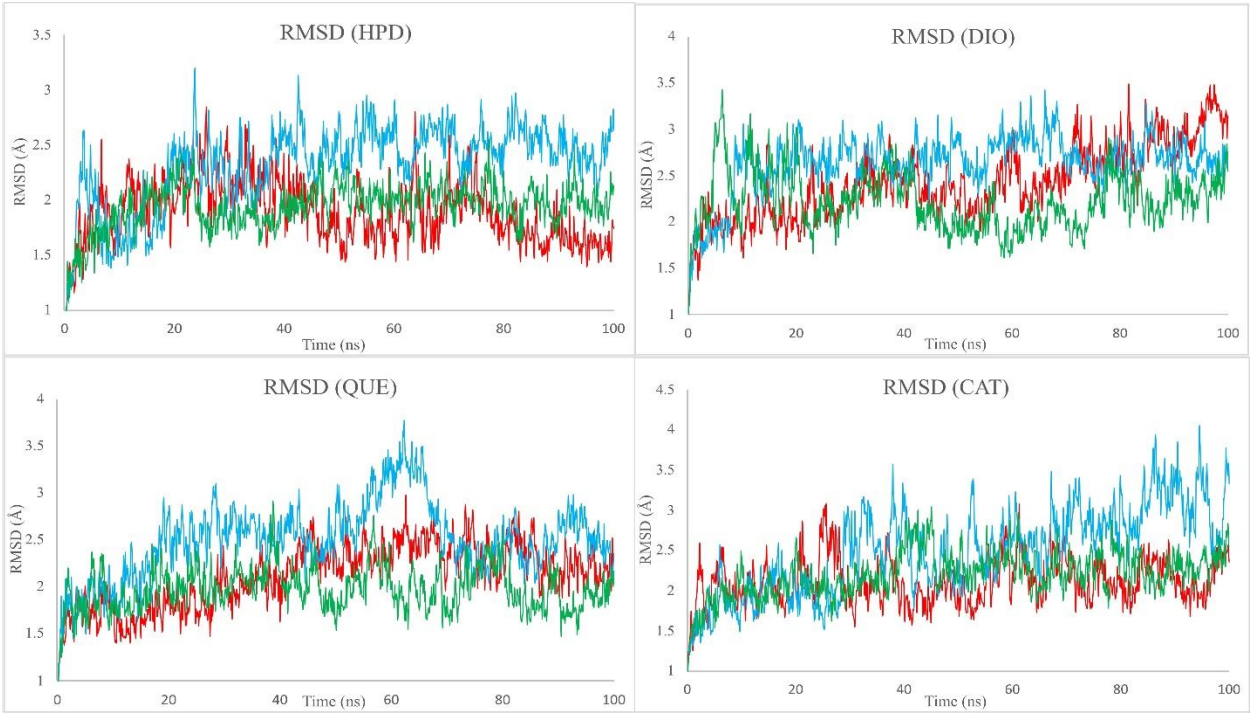

Figure S1. Root Mean Square Deviation (RMSD) profiles for three independent 100 ns MD simulations of investigated complexes.

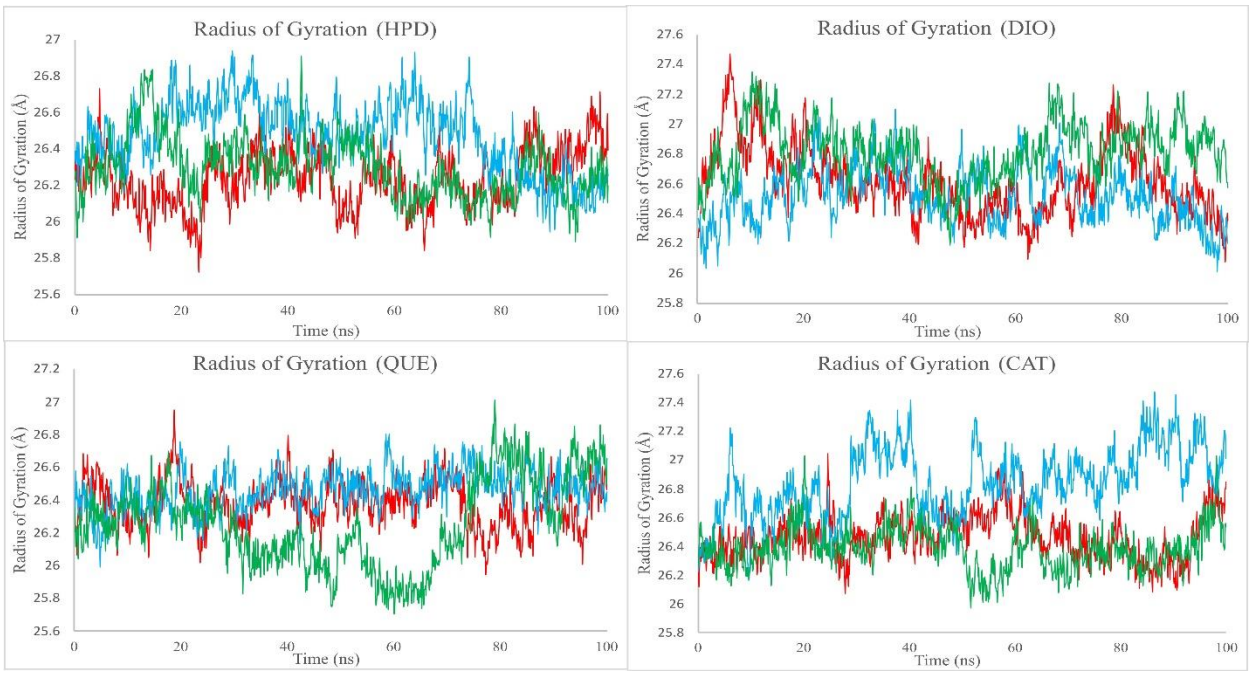

Figure S2. Radius of Gyration (Rg) profiles for three independent 100 ns MD simulations of investigated complexes.

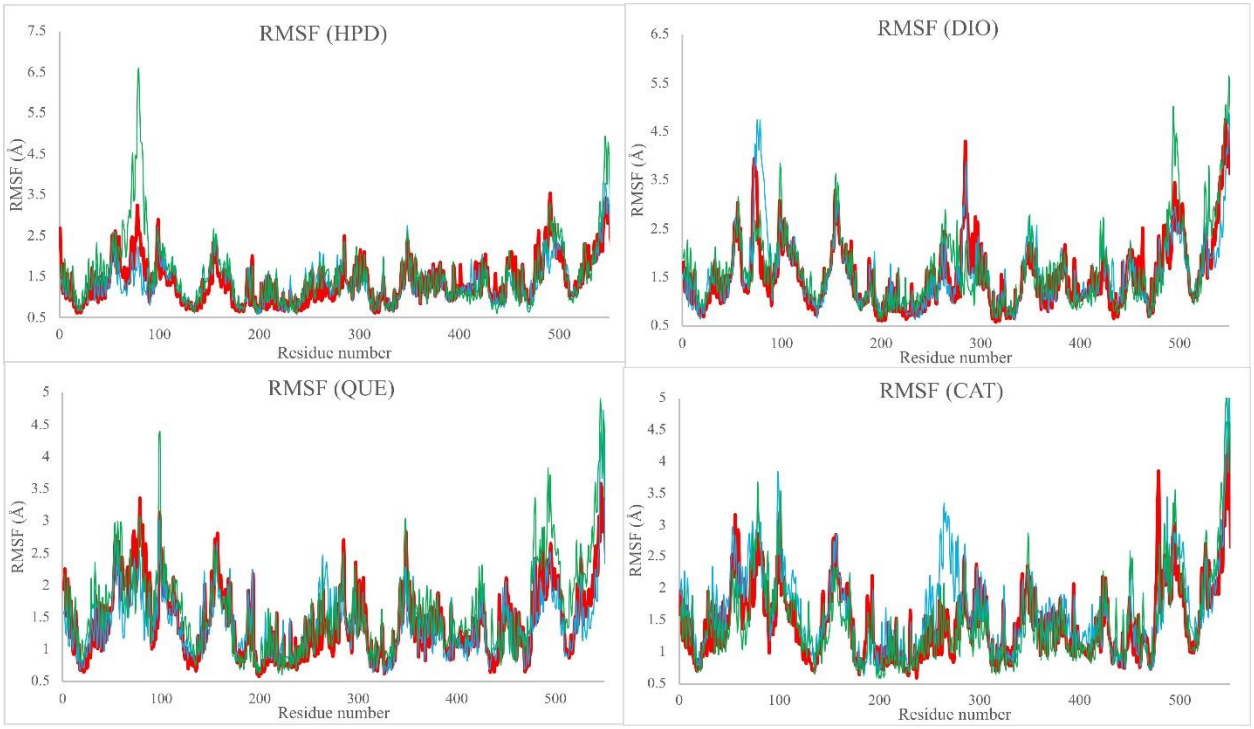

Figure S3. Root Mean Square Fluctuations (RMSF) profiles for three independent 100 ns MD simulations of investigated complexes.

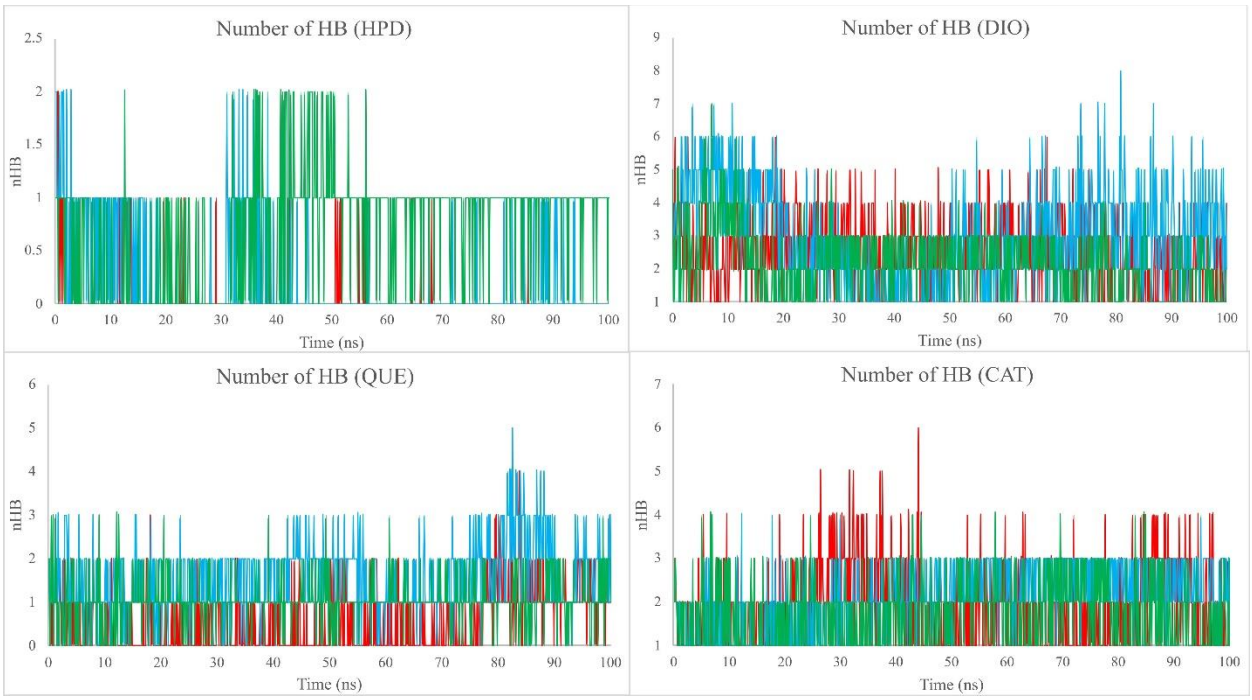

Figure S4. Number of Hydrogen Bonds (nHB) profiles for three independent 100 ns MD simulations of investigated complexes.