

Supporting Information for:

Do Double Hybrid Functionals Benefit from Regularization in the PT2
Term? Observations from an Extensive Benchmark

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Table S1: Optimized parameters, total WTMA2 and its decomposition into major subcategories for κ DSD-PBEP86-D3BJ, κ DSD-PBEPBE-D3BJ, κ DSD-BLYP-D3BJ, and κ xDSD₇₅-PBEP86-D3BJ. Heatmapping is from red (worst) via yellow to green (best). Thermo=Small Molecule Thermochemistry; Barrier=barrier heights; Large=reaction energies for large systems; Conf=conformer/intramolecular interactions; Intermol=intermolecular interactions; and NCI= Total Non-Covalent Interaction (Conformer/Intramolecular + Intermolecular).

Functionals	WTMA2	Parameters ^[a]					Δ WTMA2 (kcal/mol)					
	(kcal/mol)	C _{C,DFT}	C _{2ab}	C _{2ss}	S ₆	κ	Thermo	Barrier	Large	Conf	Intermol	NCI
κ DSD-PBEP86-D3BJ ($c_{X,HF}=0.69$)	3.404	0.1713	0.8220	0.3588	0.5073	1.1	1.021	0.557	0.683	0.617	0.527	1.144
	2.807	0.2339	0.7410	0.2935	0.4331	1.45	0.842	0.386	0.576	0.555	0.449	1.004
	2.477	0.3176	0.6534	0.2193	0.3953	2.0	0.665	0.301	0.520	0.508	0.482	0.990
	2.428	0.3397	0.6336	0.2003	0.3950	2.2	0.628	0.291	0.512	0.498	0.500	0.998
	2.383	0.3647	0.6148	0.1740	0.3960	2.5	0.589	0.281	0.507	0.486	0.519	1.006
	2.352	0.3906	0.5974	0.1474	0.3989	3.0	0.554	0.277	0.505	0.474	0.542	1.016
	2.346	0.4075	0.5864	0.1220	0.4111	3.5	0.543	0.270	0.513	0.469	0.552	1.021
	2.348	0.4164	0.5824	0.1085	0.4175	4.0	0.534	0.269	0.519	0.466	0.560	1.026
	2.367	0.4312	0.5761	0.0816	0.4339	∞	0.521	0.268	0.552	0.457	0.569	1.026
κ DSD-PBEPBE-D3BJ ($c_{X,HF}=0.68$)	3.467	0.1708	0.8221	0.3865	0.5489	1.1	1.070	0.544	0.693	0.644	0.516	1.160
	2.996	0.2431	0.7297	0.3071	0.5075	1.45	0.906	0.409	0.599	0.619	0.463	1.082
	2.760	0.3305	0.6601	0.2026	0.5136	2.0	0.755	0.342	0.535	0.620	0.508	1.127
	2.722	0.3506	0.6391	0.1983	0.5037	2.2	0.726	0.337	0.522	0.612	0.526	1.137
	2.686	0.3760	0.6182	0.1790	0.5061	2.5	0.693	0.329	0.511	0.608	0.546	1.154
	2.659	0.4068	0.6076	0.1376	0.5206	3.0	0.658	0.320	0.507	0.606	0.568	1.174
	2.650	0.4239	0.5985	0.1161	0.5347	3.5	0.644	0.314	0.506	0.607	0.578	1.185
	2.648	0.4293	0.5933	0.1095	0.5394	4.0	0.636	0.313	0.507	0.608	0.583	1.191
	2.668	0.4518	0.5893	0.0661	0.5739	∞	0.640	0.299	0.532	0.610	0.586	1.197
κ DSD-BLYP-D3BJ ($c_{X,HF}=0.71$)	3.670	0.2647	0.8008	0.3807	0.6462	1.1	0.977	0.607	0.697	0.642	0.746	1.389
	2.978	0.3479	0.7033	0.3226	0.5770	1.45	0.789	0.425	0.543	0.564	0.657	1.221
	2.494	0.4197	0.6303	0.2878	0.5245	2.0	0.640	0.309	0.434	0.513	0.598	1.111
	2.418	0.4387	0.6166	0.2724	0.5236	2.2	0.612	0.292	0.426	0.506	0.583	1.089
	2.370	0.4621	0.5905	0.2728	0.5094	2.5	0.584	0.279	0.434	0.493	0.579	1.072
	2.355	0.4900	0.5714	0.2478	0.5181	3.0	0.566	0.274	0.464	0.483	0.568	1.051
	2.364	0.5092	0.5585	0.2327	0.5219	3.5	0.556	0.275	0.493	0.474	0.566	1.040
	2.377	0.5168	0.5551	0.2196	0.5279	4.0	0.556	0.274	0.518	0.468	0.561	1.030
	2.430	0.5342	0.5467	0.1924	0.5489	∞	0.556	0.275	0.575	0.468	0.556	1.024
κ xDSD ₇₅ -PBEP86-D3BJ ($c_{X,HF}=0.75$)	3.781	0.1118	0.9307	0.3960	0.4170	1.1	1.062	0.690	0.725	0.663	0.641	1.303
	3.011	0.1774	0.8383	0.3231	0.3412	1.45	0.864	0.467	0.593	0.570	0.518	1.088
	2.459	0.2553	0.7557	0.2471	0.3112	2.0	0.680	0.327	0.487	0.516	0.449	0.965
	2.358	0.2709	0.7405	0.2319	0.3067	2.2	0.642	0.303	0.465	0.506	0.442	0.948
	2.261	0.2878	0.7201	0.2253	0.2932	2.5	0.601	0.284	0.442	0.491	0.442	0.933
	2.178	0.3151	0.7066	0.1905	0.2922	3.0	0.552	0.265	0.438	0.470	0.453	0.923
	2.146	0.3328	0.6929	0.1738	0.2913	3.5	0.525	0.256	0.446	0.453	0.466	0.919
	2.135	0.3382	0.6850	0.1772	0.2829	4.0	0.510	0.257	0.450	0.446	0.472	0.918
	2.145	0.3590	0.6777	0.1389	0.3071	∞	0.494	0.250	0.487	0.432	0.481	0.914

^[a]All results are with fixed $a_1 = 0$; $s_8 = 0$; and $a_2 = 5.5$ (for κ DSD-PBEP86-D3BJ and κ DSD-PBEPBE-D3BJ), 5.2 (for κ DSD-BLYP-D3BJ), and 5.6 (for κ xDSD₇₅-PBEP86-D3BJ).

Table S2: Optimized parameters, total WTMAP2 and its decomposition into major subcategories for κ DSD-PBEP86-D4, κ DSD-PBEPBE-D4, κ DSD-BLYP-D4, and κ DSD₇₅-PBEP86-D4. Heatmapping is from red (worst) via yellow to green (best). Thermo=Small Molecule Thermochemistry; Barrier=barrier heights; Large=reaction energies for large systems; Conf=conformer/intramolecular interactions; Intermol=intermolecular interactions; and NCI= Total Non-Covalent Interaction (Conformer/Intramolecular + Intermolecular).

Functionals	WTMAP2 (kcal/mol)	Parameters ^[a]							Δ WTMAP2 (kcal/mol)					
		$c_{c,DFT}$	c_{2ab}	c_{2ss}	s_6	a_1	a_2	κ	Thermo	Barrier	Large	Conf	Intermol	NCI
κ DSD-PBEP86-D4 ($c_{X,HF}=0.69$)	3.431	0.1687	0.8473	0.3131	0.6038	0.012	5.751	1.1	1.008	0.560	0.693	0.588	0.581	1.169
	2.796	0.2270	0.7527	0.2577	0.5507	0.158	5.000	1.45	0.847	0.387	0.581	0.494	0.487	0.980
	2.410	0.2919	0.6841	0.1822	0.5461	0.262	4.461	2	0.700	0.303	0.508	0.447	0.452	0.899
	2.349	0.3153	0.6629	0.1645	0.5469	0.282	4.394	2.2	0.663	0.289	0.505	0.439	0.453	0.892
	2.294	0.3440	0.6400	0.1444	0.5458	0.312	4.327	2.5	0.624	0.279	0.507	0.432	0.453	0.885
	2.252	0.3719	0.6229	0.1170	0.5483	0.360	4.096	3	0.588	0.275	0.512	0.418	0.459	0.877
	2.239	0.3882	0.6130	0.0995	0.5558	0.369	4.103	3.5	0.572	0.272	0.521	0.414	0.460	0.874
	2.233	0.4012	0.6069	0.0816	0.5657	0.375	4.099	4	0.564	0.267	0.532	0.410	0.462	0.871
	2.248	0.4224	0.5935	0.0566	0.5917	0.371	4.201	∞	0.545	0.260	0.573	0.406	0.463	0.869
κ DSD-PBEPBE-D4 ($c_{X,HF}=0.68$)	3.472	0.1540	0.8696	0.3069	0.6543	0.052	5.351	1.1	1.060	0.544	0.717	0.562	0.589	1.151
	2.924	0.2309	0.7709	0.2302	0.6350	0.180	4.752	1.45	0.904	0.407	0.620	0.492	0.502	0.994
	2.580	0.3246	0.6785	0.1603	0.6421	0.304	4.108	2	0.761	0.341	0.555	0.447	0.475	0.922
	2.521	0.3456	0.6604	0.1425	0.6501	0.311	4.089	2.2	0.732	0.331	0.540	0.444	0.474	0.918
	2.462	0.3607	0.6437	0.1296	0.6568	0.350	3.849	2.5	0.708	0.327	0.519	0.432	0.476	0.908
	2.400	0.3955	0.6257	0.0953	0.6706	0.353	3.883	3	0.669	0.315	0.508	0.425	0.483	0.908
	2.369	0.4108	0.6181	0.0769	0.6852	0.359	3.796	3.5	0.651	0.312	0.499	0.423	0.485	0.908
	2.354	0.4184	0.6169	0.0608	0.6997	0.381	3.745	4	0.643	0.310	0.501	0.419	0.481	0.900
	2.363	0.4371	0.6111	0.0318	0.7226	0.384	3.756	∞	0.635	0.301	0.523	0.420	0.484	0.904
κ DSD-BLYP-D4 ($c_{X,HF}=0.71$)	3.745	0.2819	0.8015	0.3617	0.7394	-0.015	5.689	1.1	0.957	0.609	0.743	0.626	0.810	1.436
	3.027	0.3480	0.7151	0.3112	0.6836	0.020	5.577	1.45	0.776	0.425	0.574	0.549	0.702	1.251
	2.512	0.4207	0.6480	0.2549	0.6576	0.072	5.335	2	0.633	0.315	0.450	0.499	0.615	1.114
	2.430	0.4426	0.6250	0.2527	0.6486	0.095	5.227	2.2	0.601	0.300	0.437	0.490	0.603	1.092
	2.370	0.4671	0.5978	0.2541	0.6379	0.112	5.163	2.5	0.573	0.286	0.435	0.483	0.592	1.075
	2.352	0.4823	0.5821	0.2372	0.6484	0.213	4.539	3	0.557	0.287	0.448	0.468	0.592	1.060
	2.351	0.5078	0.5654	0.2204	0.6528	0.157	4.897	3.5	0.544	0.282	0.475	0.466	0.584	1.050
	2.365	0.5135	0.5608	0.2145	0.6567	0.147	4.950	4	0.541	0.281	0.496	0.464	0.583	1.047
	2.428	0.5231	0.5578	0.1891	0.6808	0.156	4.878	∞	0.545	0.282	0.554	0.468	0.579	1.047
κ DSD ₇₅ -PBEP86-D4 ($c_{X,HF}=0.75$)	3.768	0.1191	0.9210	0.4375	0.5258	-0.337	8.486	1.1	1.042	0.689	0.735	0.696	0.607	1.302
	2.997	0.1799	0.8374	0.3464	0.4116	-0.335	8.236	1.45	0.848	0.467	0.597	0.567	0.519	1.085
	2.456	0.2403	0.7745	0.2404	0.3934	0.072	5.702	2	0.698	0.330	0.484	0.479	0.465	0.944
	2.354	0.2600	0.7572	0.2241	0.3910	0.095	5.623	2.2	0.658	0.307	0.464	0.468	0.457	0.925
	2.252	0.2864	0.7340	0.2013	0.3944	0.114	5.592	2.5	0.609	0.286	0.450	0.454	0.453	0.907
	2.177	0.3013	0.7203	0.1816	0.3985	0.331	4.212	3	0.573	0.273	0.434	0.429	0.468	0.897
	2.126	0.3220	0.7069	0.1637	0.4057	0.190	5.193	3.5	0.544	0.259	0.441	0.430	0.452	0.882
	2.112	0.3348	0.6972	0.1518	0.4114	0.191	5.203	4	0.529	0.254	0.449	0.423	0.457	0.880
	2.118	0.3517	0.6623	0.1168	0.4246	0.283	4.720	∞	0.508	0.252	0.493	0.409	0.457	0.866

^[a]All results are with fixed $c_{ATM} = 1$ and $s_8 = 0$.

Table S3: Optimized parameters, total WTMAP2 and its decomposition into major subcategories for κ DOD-PBEP86-D4, κ DOD-PBEPBE-D4, κ DOD-BLYP-D4, and κ xDOD₇₅-PBEP86-D4. Heatmapping is from red (worst) via yellow to green (best). Thermo=Small Molecule Thermochemistry; Barrier=barrier heights; Large=reaction energies for large systems; Conf=conformer/intramolecular interactions; Intermol=intermolecular interactions; and NCI= Total Non-Covalent Interaction (Conformer/Intramolecular + Intermolecular).

Functionals	WTMAP2 (kcal/mol)	Parameters ^[a]							Δ WTMAP2 (kcal/mol)					
		C _{C,DFT}	C _{2ab}	C _{2ss}	S ₆	a ₁	a ₂	κ	Thermo	Barrier	Large	Conf	Intermol	NCI
κ DOD-PBEP86-D4 (C _{X,HF} =0.69)	3.632	0.2174	0.9390	0.0	0.7101	0.001	5.570	1.1	1.002	0.655	0.777	0.572	0.627	1.199
	3.004	0.2653	0.8332	0.0	0.6707	0.124	4.927	1.45	0.856	0.442	0.676	0.495	0.536	1.031
	2.585	0.3200	0.7413	0.0	0.6408	0.247	4.403	2	0.738	0.318	0.602	0.455	0.473	0.927
	2.501	0.3363	0.7187	0.0	0.6350	0.264	4.345	2.2	0.710	0.300	0.580	0.444	0.468	0.911
	2.419	0.3604	0.6921	0.0	0.6251	0.293	4.225	2.5	0.677	0.285	0.560	0.428	0.470	0.898
	2.336	0.3892	0.6638	0.0	0.6043	0.320	4.194	3	0.639	0.271	0.550	0.411	0.465	0.876
	2.297	0.4013	0.6456	0.0	0.6076	0.352	4.033	3.5	0.618	0.267	0.544	0.402	0.467	0.869
	2.280	0.4093	0.6341	0.0	0.6139	0.357	4.024	4	0.604	0.264	0.543	0.400	0.469	0.869
	2.271	0.4301	0.6131	0.0	0.6158	0.344	4.243	∞	0.575	0.253	0.581	0.399	0.464	0.862
κ DOD-PBEPBE-D4 (C _{X,HF} =0.68)	3.664	0.2310	0.9378	0.0	0.7777	0.045	5.323	1.1	1.032	0.644	0.793	0.578	0.617	1.195
	3.084	0.2885	0.8294	0.0	0.7367	0.180	4.596	1.45	0.898	0.441	0.703	0.503	0.540	1.043
	2.697	0.3553	0.7302	0.0	0.7255	0.296	4.039	2	0.778	0.341	0.618	0.466	0.495	0.961
	2.616	0.3682	0.7134	0.0	0.7218	0.309	3.990	2.2	0.752	0.322	0.594	0.463	0.485	0.948
	2.528	0.3868	0.6886	0.0	0.7231	0.344	3.822	2.5	0.722	0.306	0.569	0.450	0.481	0.931
	2.441	0.4059	0.6624	0.0	0.7263	0.347	3.921	3	0.690	0.294	0.539	0.436	0.482	0.918
	2.398	0.4179	0.6463	0.0	0.7358	0.381	3.745	3.5	0.672	0.289	0.519	0.437	0.481	0.918
	2.375	0.4240	0.6383	0.0	0.7380	0.397	3.592	4	0.662	0.291	0.517	0.425	0.480	0.905
	2.369	0.4465	0.6175	0.0	0.7410	0.377	3.770	∞	0.641	0.289	0.531	0.423	0.485	0.908
κ DOD-BLYP-D4 (C _{X,HF} =0.71)	3.996	0.3470	0.9431	0.0	0.8170	-0.013	5.545	1.1	1.058	0.732	0.779	0.609	0.819	1.428
	3.323	0.3884	0.8559	0.0	0.7746	0.004	5.498	1.45	0.916	0.521	0.631	0.549	0.707	1.255
	2.846	0.4519	0.7574	0.0	0.7557	0.067	5.144	2	0.794	0.385	0.523	0.508	0.635	1.143
	2.751	0.4643	0.7379	0.0	0.7556	0.085	5.029	2.2	0.768	0.360	0.501	0.500	0.623	1.123
	2.667	0.4801	0.7203	0.0	0.7512	0.100	4.966	2.5	0.750	0.336	0.483	0.495	0.604	1.099
	2.615	0.5029	0.6866	0.0	0.7592	0.133	4.777	3	0.718	0.321	0.490	0.484	0.601	1.085
	2.606	0.5334	0.6657	0.0	0.7469	0.107	4.996	3.5	0.708	0.314	0.510	0.479	0.595	1.074
	2.600	0.5442	0.6501	0.0	0.7530	0.127	4.892	4	0.692	0.311	0.527	0.477	0.593	1.070
	2.622	0.5548	0.6297	0.0	0.7661	0.146	4.779	∞	0.667	0.301	0.576	0.480	0.598	1.078
κ xDOD ₇₅ -PBEP86-D4 (C _{X,HF} =0.75)	4.055	0.1649	1.0712	0.0	0.6211	-0.090	6.100	1.1	1.072	0.815	0.835	0.647	0.686	1.333
	3.295	0.2213	0.9558	0.0	0.5503	-0.015	5.776	1.45	0.900	0.575	0.702	0.543	0.576	1.119
	2.721	0.2732	0.8599	0.0	0.5136	0.010	5.625	2	0.753	0.389	0.572	0.482	0.525	1.007
	2.597	0.2776	0.8399	0.0	0.5261	0.131	4.899	2.2	0.726	0.345	0.544	0.470	0.512	0.982
	2.476	0.2994	0.8129	0.0	0.5133	0.144	4.917	2.5	0.691	0.319	0.515	0.454	0.497	0.952
	2.361	0.3175	0.7862	0.0	0.5160	0.286	4.201	3	0.657	0.288	0.497	0.437	0.482	0.919
	2.284	0.3431	0.7654	0.0	0.4943	0.138	5.143	3.5	0.624	0.276	0.477	0.426	0.483	0.908
	2.248	0.3538	0.7555	0.0	0.4903	0.169	5.064	4	0.610	0.267	0.480	0.419	0.473	0.892
	2.215	0.3695	0.7092	0.0	0.4865	0.199	4.952	∞	0.582	0.251	0.507	0.402	0.473	0.875

^[a]All results are with fixed c_{ATM} = 1 and s₈ = 0.

Table S4: Optimized parameters, total WTAD2 and its decomposition into major subcategories for κ DSD_X-PBEP86-D3BJ. Heatmapping is from red (worst) via yellow to green (best). Thermo=Small Molecule Thermochemistry; Barrier=barrier heights; Large=reaction energies for large systems; Conf=conformer/intramolecular interactions; Intermol=intermolecular interactions; and NCI= Total Non-Covalent Interaction (Conformer/Intramolecular + Intermolecular).

WTAD2 (kcal/mol)	Parameters ^[a]							Δ WTAD2 (kcal/mol)					
	C _{X,HF}	C _{X,DFT}	C _{C,DFT}	C _{2ab}	C _{2ss}	S ₆	K	Thermo	Barrier	Large	Conf	Intermol	NCI
3.085	0.50	0.50	0.3329	0.6187	0.0368	0.7936	1.1	0.943	0.370	0.599	0.616	0.558	1.174
3.024	0.50	0.50	0.4029	0.5240	0.0100	0.7458	1.45	0.864	0.390	0.595	0.586	0.590	1.176
3.087	0.50	0.50	0.4816	0.4590	-0.0633	0.7483	2.0	0.825	0.419	0.621	0.573	0.649	1.222
3.113	0.50	0.50	0.5014	0.4402	-0.0734	0.7499	2.2	0.817	0.425	0.629	0.568	0.674	1.242
3.146	0.50	0.50	0.5262	0.4156	-0.0825	0.7485	2.5	0.806	0.430	0.642	0.565	0.703	1.268
3.189	0.50	0.50	0.5481	0.3941	-0.0964	0.7596	3.0	0.805	0.429	0.663	0.563	0.730	1.293
3.218	0.50	0.50	0.5604	0.3818	-0.1007	0.7608	3.5	0.802	0.428	0.679	0.564	0.744	1.309
3.239	0.50	0.50	0.5671	0.3753	-0.1079	0.7673	4.0	0.806	0.422	0.695	0.567	0.748	1.315
3.307	0.50	0.50	0.5887	0.3581	-0.1237	0.7774	∞	0.816	0.409	0.739	0.572	0.772	1.344
2.978	0.55	0.45	0.2843	0.6742	0.1262	0.7143	1.1	0.938	0.367	0.571	0.597	0.505	1.102
2.795	0.55	0.45	0.3536	0.5687	0.0998	0.6675	1.45	0.836	0.334	0.540	0.561	0.524	1.085
2.790	0.55	0.45	0.4322	0.5045	0.0169	0.6654	2.0	0.763	0.356	0.549	0.539	0.583	1.122
2.807	0.55	0.45	0.4554	0.4877	-0.0026	0.6660	2.2	0.749	0.361	0.557	0.532	0.608	1.139
2.831	0.55	0.45	0.4799	0.4696	-0.0234	0.6678	2.5	0.737	0.365	0.570	0.527	0.633	1.160
2.863	0.55	0.45	0.5063	0.4471	-0.0423	0.6764	3.0	0.728	0.365	0.585	0.521	0.664	1.185
2.888	0.55	0.45	0.5233	0.4324	-0.0505	0.6777	3.5	0.720	0.365	0.599	0.519	0.685	1.204
2.905	0.55	0.45	0.5340	0.4236	-0.0593	0.6854	4.0	0.719	0.361	0.609	0.518	0.699	1.216
2.962	0.55	0.45	0.5554	0.4077	-0.0799	0.6975	∞	0.724	0.345	0.654	0.521	0.718	1.239
2.989	0.58	0.42	0.2537	0.6938	0.2093	0.6625	1.1	0.953	0.388	0.562	0.591	0.495	1.085
2.711	0.58	0.42	0.3202	0.5965	0.1598	0.6102	1.45	0.831	0.315	0.523	0.557	0.485	1.041
2.658	0.58	0.42	0.4059	0.5330	0.0658	0.6041	2.0	0.727	0.328	0.526	0.527	0.550	1.077
2.665	0.58	0.42	0.4288	0.5163	0.0390	0.6123	2.2	0.714	0.327	0.533	0.520	0.570	1.090
2.678	0.58	0.42	0.4546	0.4969	0.0185	0.6149	2.5	0.696	0.333	0.540	0.510	0.600	1.110
2.703	0.58	0.42	0.4848	0.4744	-0.0035	0.6184	3.0	0.680	0.335	0.552	0.500	0.635	1.135
2.722	0.58	0.42	0.5024	0.4615	-0.0188	0.6251	3.5	0.674	0.332	0.565	0.497	0.654	1.151
2.736	0.58	0.42	0.5130	0.4524	-0.0296	0.6346	4.0	0.674	0.326	0.575	0.497	0.665	1.161
2.761	0.58	0.42	0.5277	0.4438	-0.0446	0.6411	∞	0.667	0.307	0.614	0.495	0.679	1.173
3.116	0.63	0.37	0.2117	0.7499	0.2922	0.5855	1.1	0.980	0.447	0.593	0.597	0.499	1.096
2.672	0.63	0.37	0.2820	0.6549	0.2282	0.5260	1.45	0.819	0.327	0.526	0.546	0.454	1.001
2.506	0.63	0.37	0.3638	0.5852	0.1344	0.5112	2.0	0.690	0.287	0.506	0.517	0.505	1.023
2.490	0.63	0.37	0.3878	0.5691	0.1113	0.5111	2.2	0.661	0.288	0.506	0.506	0.528	1.033
2.484	0.63	0.37	0.4092	0.5497	0.0877	0.5189	2.5	0.646	0.284	0.510	0.499	0.545	1.044
2.488	0.63	0.37	0.4395	0.5319	0.0543	0.5260	3.0	0.624	0.283	0.522	0.486	0.572	1.058
2.494	0.63	0.37	0.4593	0.5192	0.0335	0.5386	3.5	0.612	0.280	0.530	0.478	0.594	1.072
2.501	0.63	0.37	0.4678	0.5118	0.0226	0.5470	4.0	0.610	0.275	0.540	0.477	0.601	1.078
2.544	0.63	0.37	0.4892	0.5012	-0.0043	0.5670	∞	0.601	0.270	0.575	0.470	0.628	1.098
3.404	0.69	0.31	0.1713	0.8220	0.3588	0.5073	1.1	1.021	0.557	0.683	0.617	0.527	1.144
2.808	0.69	0.31	0.2339	0.7410	0.2935	0.4331	1.45	0.842	0.386	0.576	0.555	0.449	1.004
2.476	0.69	0.31	0.3176	0.6534	0.2193	0.3953	2.0	0.665	0.301	0.520	0.508	0.482	0.990
2.428	0.69	0.31	0.3397	0.6336	0.2003	0.3950	2.2	0.628	0.291	0.512	0.498	0.500	0.998
2.383	0.69	0.31	0.3647	0.6148	0.1740	0.3960	2.5	0.589	0.281	0.507	0.486	0.519	1.006
2.352	0.69	0.31	0.3906	0.5974	0.1474	0.3989	3.0	0.554	0.277	0.505	0.474	0.542	1.016
2.346	0.69	0.31	0.4075	0.5864	0.1220	0.4111	3.5	0.543	0.270	0.513	0.469	0.552	1.021
2.348	0.69	0.31	0.4164	0.5824	0.1085	0.4175	4.0	0.534	0.269	0.519	0.466	0.560	1.026
2.367	0.69	0.31	0.4312	0.5761	0.0816	0.4339	∞	0.521	0.268	0.552	0.457	0.569	1.026
3.804	0.75	0.25	0.1295	0.9038	0.4260	0.4209	1.1	1.069	0.698	0.815	0.644	0.578	1.221
3.104	0.75	0.25	0.1922	0.8159	0.3559	0.3507	1.45	0.875	0.489	0.685	0.572	0.484	1.056
2.634	0.75	0.25	0.2699	0.7330	0.2826	0.3121	2.0	0.681	0.374	0.579	0.527	0.474	1.000
2.551	0.75	0.25	0.2888	0.7143	0.2639	0.3091	2.2	0.638	0.357	0.558	0.517	0.481	0.998
2.472	0.75	0.25	0.3118	0.6921	0.2496	0.2955	2.5	0.590	0.342	0.541	0.501	0.498	0.999

2.398	0.75	0.25	0.3335	0.6729	0.2281	0.2943	3.0	0.540	0.328	0.526	0.489	0.515	1.004
2.368	0.75	0.25	0.3498	0.6647	0.2053	0.3045	3.5	0.514	0.321	0.525	0.482	0.525	1.008
2.357	0.75	0.25	0.3601	0.6582	0.1917	0.3086	4.0	0.500	0.319	0.529	0.476	0.532	1.008
2.358	0.75	0.25	0.3755	0.6514	0.1638	0.3267	∞	0.481	0.317	0.556	0.466	0.537	1.003
6.195	1.00	0.00	-0.0601	1.2348	0.7766	0.0267	1.1	1.437	1.381	1.526	0.915	0.936	1.851
5.257	1.00	0.00	-0.0136	1.1566	0.6961	-0.0592	1.45	1.175	1.109	1.355	0.796	0.822	1.619
4.568	1.00	0.00	0.0745	1.0584	0.5820	-0.0650	2.0	0.952	0.970	1.182	0.729	0.735	1.464
4.429	1.00	0.00	0.0977	1.0359	0.5582	-0.0651	2.2	0.906	0.945	1.144	0.713	0.720	1.434
4.282	1.00	0.00	0.1172	1.0262	0.5181	-0.0548	2.5	0.855	0.922	1.099	0.705	0.701	1.406
4.141	1.00	0.00	0.1405	1.0178	0.4743	-0.0443	3.0	0.805	0.901	1.058	0.693	0.684	1.377
4.068	1.00	0.00	0.1580	1.0165	0.4339	-0.0315	3.5	0.779	0.894	1.034	0.685	0.678	1.363
4.029	1.00	0.00	0.1643	1.0162	0.4240	-0.0325	4.0	0.765	0.887	1.025	0.678	0.675	1.353
3.968	1.00	0.00	0.1710	1.0175	0.4025	-0.0194	∞	0.736	0.880	1.015	0.675	0.661	1.337

^[a]All results are with fixed $a_1 = 0$; $s_8 = 0$; and $a_2 = 5.5$

Table S5: Optimized parameters, total WTAD2 and its decomposition into major subcategories for κ DH_X-PBEP86-D3BJ. Heatmapping is from red (worst) via yellow to green (best). Thermo=Small Molecule Thermochemistry; Barrier=barrier heights; Large=reaction energies for large systems; Conf=conformer/intramolecular interactions; Intermol=intermolecular interactions; and NCI= Total Non-Covalent Interaction (Conformer/Intramolecular + Intermolecular).

WTAD2 (kcal/mol)	Parameters ^[a]							Δ WTAD2 (kcal/mol)					
	C _{X,HF}	C _{X,DFT}	C _{C,DFT}	C _{Zab}	C _{Zss}	S ₆	K	Thermo	Barrier	Large	Conf	Intermol	NCI
3.399	0.50	0.50	0.2913	0.4083	0.4083	0.7315	1.1	1.220	0.377	0.569	0.645	0.588	1.232
3.366	0.50	0.50	0.3603	0.3288	0.3288	0.7018	1.45	1.136	0.403	0.591	0.621	0.616	1.237
3.454	0.50	0.50	0.4261	0.2578	0.2578	0.7110	2.0	1.108	0.421	0.640	0.616	0.669	1.285
3.489	0.50	0.50	0.4392	0.2437	0.2437	0.7159	2.2	1.108	0.424	0.655	0.618	0.683	1.301
3.533	0.50	0.50	0.4583	0.2260	0.2260	0.7244	2.5	1.108	0.426	0.670	0.618	0.711	1.329
3.590	0.50	0.50	0.4708	0.2090	0.2090	0.7396	3.0	1.126	0.419	0.697	0.625	0.723	1.348
3.629	0.50	0.50	0.4896	0.1980	0.1980	0.7354	3.5	1.115	0.429	0.705	0.626	0.753	1.379
3.654	0.50	0.50	0.4966	0.1924	0.1924	0.7373	4.0	1.117	0.432	0.716	0.628	0.763	1.391
3.730	0.50	0.50	0.5113	0.1776	0.1776	0.7525	∞	1.141	0.422	0.751	0.632	0.785	1.416
3.231	0.55	0.45	0.2366	0.4767	0.4767	0.6656	1.1	1.214	0.366	0.511	0.625	0.515	1.140
3.080	0.55	0.45	0.3158	0.3862	0.3862	0.6319	1.45	1.096	0.344	0.521	0.588	0.531	1.119
3.129	0.55	0.45	0.3825	0.3194	0.3194	0.6195	2.0	1.027	0.371	0.556	0.579	0.596	1.176
3.162	0.55	0.45	0.4001	0.3036	0.3036	0.6184	2.2	1.015	0.380	0.568	0.579	0.620	1.198
3.206	0.55	0.45	0.4222	0.2839	0.2839	0.6219	2.5	1.008	0.386	0.585	0.579	0.648	1.227
3.264	0.55	0.45	0.4446	0.2654	0.2654	0.6272	3.0	1.001	0.397	0.602	0.580	0.685	1.265
3.304	0.55	0.45	0.4598	0.2505	0.2505	0.6356	3.5	1.011	0.389	0.619	0.585	0.700	1.285
3.331	0.55	0.45	0.4690	0.2423	0.2423	0.6389	4.0	1.015	0.387	0.630	0.588	0.710	1.298
3.405	0.55	0.45	0.4880	0.2236	0.2236	0.6574	∞	1.040	0.374	0.664	0.591	0.736	1.326
3.214	0.58	0.42	0.2161	0.5090	0.5090	0.6271	1.1	1.211	0.375	0.507	0.621	0.499	1.121
2.981	0.58	0.42	0.2782	0.4333	0.4333	0.5729	1.45	1.082	0.342	0.491	0.583	0.485	1.067
2.979	0.58	0.42	0.3625	0.3509	0.3509	0.5668	2.0	0.991	0.341	0.521	0.563	0.564	1.127
3.005	0.58	0.42	0.3814	0.3326	0.3326	0.5703	2.2	0.981	0.343	0.533	0.562	0.586	1.147
3.044	0.58	0.42	0.4041	0.3153	0.3153	0.5666	2.5	0.962	0.358	0.545	0.560	0.620	1.180
3.098	0.58	0.42	0.4237	0.2988	0.2988	0.5688	3.0	0.951	0.372	0.562	0.561	0.652	1.214
3.137	0.58	0.42	0.4344	0.2875	0.2875	0.5749	3.5	0.957	0.372	0.579	0.565	0.665	1.230
3.165	0.58	0.42	0.4468	0.2769	0.2769	0.5799	4.0	0.961	0.368	0.591	0.567	0.679	1.246
3.214	0.58	0.42	0.4597	0.2626	0.2626	0.5968	∞	0.984	0.347	0.623	0.565	0.695	1.260
3.323	0.63	0.37	0.1741	0.5767	0.5767	0.5487	1.1	1.212	0.433	0.543	0.626	0.510	1.135
2.931	0.63	0.37	0.2364	0.4966	0.4966	0.4911	1.45	1.064	0.350	0.486	0.578	0.453	1.032
2.820	0.63	0.37	0.3131	0.4153	0.4153	0.4772	2.0	0.963	0.323	0.482	0.551	0.501	1.052
2.826	0.63	0.37	0.3396	0.3931	0.3931	0.4752	2.2	0.938	0.320	0.493	0.545	0.530	1.075
2.844	0.63	0.37	0.3594	0.3767	0.3767	0.4687	2.5	0.915	0.328	0.504	0.543	0.555	1.098
2.886	0.63	0.37	0.3796	0.3596	0.3596	0.4698	3.0	0.898	0.338	0.521	0.543	0.586	1.129
2.919	0.63	0.37	0.3992	0.3430	0.3430	0.4749	3.5	0.895	0.334	0.537	0.544	0.610	1.154
2.945	0.63	0.37	0.4078	0.3346	0.3346	0.4833	4.0	0.897	0.333	0.547	0.543	0.625	1.168
3.015	0.63	0.37	0.4220	0.3220	0.3220	0.4906	∞	0.900	0.339	0.586	0.543	0.647	1.190
3.609	0.69	0.31	0.1296	0.6531	0.6531	0.4548	1.1	1.234	0.533	0.646	0.645	0.551	1.196
3.070	0.69	0.31	0.1875	0.5744	0.5744	0.3849	1.45	1.059	0.406	0.554	0.584	0.466	1.050
2.808	0.69	0.31	0.2703	0.4911	0.4911	0.3582	2.0	0.926	0.355	0.500	0.545	0.482	1.027
2.784	0.69	0.31	0.2881	0.4737	0.4737	0.3575	2.2	0.902	0.352	0.495	0.540	0.495	1.036
2.773	0.69	0.31	0.3064	0.4567	0.4567	0.3543	2.5	0.877	0.353	0.493	0.536	0.514	1.050
2.779	0.69	0.31	0.3290	0.4370	0.4370	0.3547	3.0	0.852	0.358	0.497	0.530	0.542	1.072
2.796	0.69	0.31	0.3418	0.4259	0.4259	0.3586	3.5	0.840	0.363	0.505	0.527	0.561	1.088
2.814	0.69	0.31	0.3493	0.4196	0.4196	0.3579	4.0	0.834	0.368	0.516	0.526	0.571	1.097
2.872	0.69	0.31	0.3661	0.4073	0.4073	0.3629	∞	0.824	0.378	0.550	0.522	0.600	1.121
4.004	0.75	0.25	0.0607	0.7559	0.7559	0.3614	1.1	1.267	0.655	0.789	0.689	0.603	1.292
3.363	0.75	0.25	0.1295	0.6625	0.6625	0.2861	1.45	1.075	0.499	0.675	0.614	0.501	1.115
2.989	0.75	0.25	0.2170	0.5721	0.5721	0.2505	2.0	0.920	0.431	0.583	0.566	0.489	1.055
2.933	0.75	0.25	0.2425	0.5476	0.5476	0.2516	2.2	0.894	0.422	0.565	0.554	0.497	1.052
2.888	0.75	0.25	0.2650	0.5300	0.5300	0.2371	2.5	0.860	0.419	0.552	0.545	0.512	1.057

2.863	0.75	0.25	0.2874	0.5096	0.5096	0.2404	3.0	0.834	0.420	0.543	0.539	0.527	1.066
2.864	0.75	0.25	0.3019	0.4990	0.4990	0.2329	3.5	0.816	0.425	0.546	0.535	0.542	1.077
2.870	0.75	0.25	0.3070	0.4934	0.4934	0.2359	4.0	0.810	0.429	0.550	0.533	0.548	1.081
2.900	0.75	0.25	0.3190	0.4846	0.4846	0.2384	∞	0.796	0.442	0.575	0.523	0.564	1.087
6.317	1.00	0.00	-0.1264	1.1149	1.1149	-0.0507	1.1	1.567	1.290	1.541	0.955	0.965	1.919
5.438	1.00	0.00	-0.0722	1.0278	1.0278	-0.1395	1.45	1.306	1.062	1.395	0.849	0.826	1.675
4.858	1.00	0.00	0.0222	0.9244	0.9244	-0.1831	2.0	1.119	0.967	1.259	0.769	0.745	1.513
4.758	1.00	0.00	0.0404	0.9010	0.9010	-0.1835	2.2	1.095	0.952	1.222	0.751	0.739	1.490
4.665	1.00	0.00	0.0617	0.8795	0.8795	-0.1840	2.5	1.063	0.942	1.196	0.741	0.723	1.464
4.585	1.00	0.00	0.0832	0.8599	0.8599	-0.1978	3.0	1.034	0.936	1.165	0.722	0.728	1.450
4.553	1.00	0.00	0.0957	0.8474	0.8474	-0.1952	3.5	1.027	0.935	1.152	0.716	0.724	1.439
4.540	1.00	0.00	0.1008	0.8398	0.8398	-0.1857	4.0	1.027	0.937	1.146	0.715	0.716	1.431
4.534	1.00	0.00	0.1075	0.8306	0.8306	-0.1746	∞	1.024	0.943	1.150	0.713	0.705	1.417

^[a]All results are with fixed $a_1 = 0$; $s_8 = 0$; and $a_2 = 5.5$

Table S6: Optimized parameters, total WTAD2 and its decomposition into major subcategories for κ DOD χ -PBEP86-D3BJ. Heatmapping is from red (worst) via yellow to green (best). Thermo=Small Molecule Thermochemistry; Barrier=barrier heights; Large=reaction energies for large systems; Conf=conformer/intramolecular interactions; Intermol=intermolecular interactions; and NCI= Total Non-Covalent Interaction (Conformer/Intramolecular + Intermolecular).

WTAD2 (kcal/mol)	Parameters ^[a]							Δ WTAD2 (kcal/mol)					
	C _{X,HF}	C _{X,DFT}	C _{C,DFT}	C _{Zab}	C _{2ss}	S ₆	K	Thermo	Barrier	Large	Conf	Intermol	NCI
3.090	0.50	0.50	0.3449	0.6243	0.0	0.8050	1.1	0.932	0.371	0.610	0.616	0.562	1.178
3.024	0.50	0.50	0.4044	0.5270	0.0	0.7511	1.45	0.864	0.388	0.597	0.586	0.589	1.176
3.099	0.50	0.50	0.4697	0.4360	0.0	0.7209	2.0	0.819	0.438	0.613	0.571	0.658	1.230
3.135	0.50	0.50	0.4842	0.4161	0.0	0.7173	2.2	0.814	0.448	0.622	0.570	0.680	1.250
3.180	0.50	0.50	0.5059	0.3915	0.0	0.7094	2.5	0.803	0.463	0.633	0.567	0.714	1.281
3.240	0.50	0.50	0.5234	0.3599	0.0	0.7185	3.0	0.820	0.458	0.657	0.569	0.736	1.305
3.284	0.50	0.50	0.5332	0.3424	0.0	0.7239	3.5	0.832	0.456	0.676	0.573	0.748	1.321
3.316	0.50	0.50	0.5407	0.3323	0.0	0.7242	4.0	0.837	0.457	0.688	0.575	0.758	1.333
3.417	0.50	0.50	0.5581	0.3024	0.0	0.7367	∞	0.876	0.442	0.735	0.582	0.784	1.365
3.021	0.55	0.45	0.3164	0.7010	0.0	0.7613	1.1	0.915	0.378	0.617	0.606	0.506	1.112
2.817	0.55	0.45	0.3754	0.5964	0.0	0.7131	1.45	0.829	0.332	0.570	0.564	0.522	1.087
2.792	0.55	0.45	0.4340	0.5133	0.0	0.6711	2.0	0.765	0.355	0.552	0.540	0.580	1.120
2.807	0.55	0.45	0.4552	0.4862	0.0	0.6651	2.2	0.749	0.362	0.557	0.531	0.609	1.140
2.834	0.55	0.45	0.4759	0.4620	0.0	0.6547	2.5	0.731	0.373	0.566	0.526	0.638	1.164
2.873	0.55	0.45	0.4992	0.4327	0.0	0.6524	3.0	0.717	0.382	0.579	0.522	0.674	1.196
2.906	0.55	0.45	0.5123	0.4139	0.0	0.6527	3.5	0.717	0.382	0.594	0.523	0.691	1.213
2.931	0.55	0.45	0.5204	0.4020	0.0	0.6572	4.0	0.718	0.383	0.603	0.522	0.705	1.227
3.020	0.55	0.45	0.5348	0.3731	0.0	0.6747	∞	0.752	0.368	0.647	0.527	0.726	1.253
3.080	0.58	0.42	0.3083	0.7364	0.0	0.7310	1.1	0.916	0.413	0.650	0.608	0.493	1.101
2.778	0.58	0.42	0.3642	0.6358	0.0	0.6782	1.45	0.814	0.323	0.590	0.557	0.495	1.053
2.677	0.58	0.42	0.4188	0.5541	0.0	0.6368	2.0	0.740	0.318	0.545	0.528	0.547	1.075
2.673	0.58	0.42	0.4352	0.5316	0.0	0.6318	2.2	0.722	0.322	0.541	0.521	0.568	1.089
2.679	0.58	0.42	0.4569	0.5043	0.0	0.6252	2.5	0.702	0.328	0.543	0.511	0.597	1.107
2.703	0.58	0.42	0.4837	0.4729	0.0	0.6178	3.0	0.680	0.335	0.552	0.500	0.635	1.135
2.725	0.58	0.42	0.4977	0.4545	0.0	0.6166	3.5	0.671	0.337	0.563	0.499	0.654	1.153
2.742	0.58	0.42	0.5038	0.4458	0.0	0.6181	4.0	0.668	0.338	0.573	0.500	0.664	1.164
2.786	0.58	0.42	0.5151	0.4258	0.0	0.6281	∞	0.677	0.320	0.609	0.500	0.680	1.179
3.305	0.63	0.37	0.2856	0.8226	0.0	0.6731	1.1	0.962	0.508	0.725	0.613	0.499	1.111
2.844	0.63	0.37	0.3336	0.7191	0.0	0.6219	1.45	0.823	0.354	0.639	0.568	0.460	1.028
2.601	0.63	0.37	0.3953	0.6234	0.0	0.5773	2.0	0.717	0.283	0.571	0.523	0.508	1.030
2.563	0.63	0.37	0.4111	0.6020	0.0	0.5675	2.2	0.694	0.279	0.556	0.511	0.523	1.035
2.525	0.63	0.37	0.4280	0.5782	0.0	0.5634	2.5	0.668	0.275	0.538	0.502	0.543	1.044
2.506	0.63	0.37	0.4525	0.5491	0.0	0.5533	3.0	0.640	0.276	0.535	0.484	0.572	1.056
2.504	0.63	0.37	0.4641	0.5326	0.0	0.5537	3.5	0.626	0.273	0.539	0.480	0.587	1.066
2.507	0.63	0.37	0.4724	0.5237	0.0	0.5519	4.0	0.615	0.274	0.543	0.475	0.600	1.075
2.544	0.63	0.37	0.4878	0.4993	0.0	0.5660	∞	0.600	0.271	0.575	0.471	0.628	1.098
3.702	0.69	0.31	0.2639	0.9147	0.0	0.6046	1.1	1.034	0.661	0.861	0.626	0.521	1.146
3.113	0.69	0.31	0.3128	0.8131	0.0	0.5470	1.45	0.875	0.460	0.739	0.570	0.468	1.039
2.720	0.69	0.31	0.3598	0.7291	0.0	0.4979	2.0	0.742	0.326	0.635	0.536	0.481	1.017
2.642	0.69	0.31	0.3733	0.7080	0.0	0.4861	2.2	0.711	0.304	0.610	0.526	0.490	1.017
2.561	0.69	0.31	0.3941	0.6790	0.0	0.4764	2.5	0.672	0.287	0.584	0.510	0.509	1.018
2.481	0.69	0.31	0.4193	0.6480	0.0	0.4651	3.0	0.630	0.273	0.561	0.485	0.532	1.017
2.439	0.69	0.31	0.4308	0.6281	0.0	0.4694	3.5	0.605	0.264	0.548	0.479	0.542	1.021
2.421	0.69	0.31	0.4362	0.6176	0.0	0.4725	4.0	0.592	0.259	0.548	0.476	0.547	1.023
2.410	0.69	0.31	0.4443	0.6047	0.0	0.4790	∞	0.567	0.253	0.569	0.466	0.556	1.022
4.201	0.75	0.25	0.2344	1.0233	0.0	0.5365	1.1	1.137	0.832	1.011	0.663	0.558	1.221
3.514	0.75	0.25	0.2837	0.9252	0.0	0.4543	1.45	0.969	0.598	0.869	0.579	0.499	1.077
2.992	0.75	0.25	0.3419	0.8202	0.0	0.4143	2.0	0.791	0.431	0.746	0.534	0.491	1.025
2.887	0.75	0.25	0.3507	0.8018	0.0	0.4105	2.2	0.752	0.398	0.711	0.532	0.494	1.026
2.770	0.75	0.25	0.3610	0.7832	0.0	0.4035	2.5	0.710	0.364	0.669	0.528	0.500	1.027

2.653	0.75	0.25	0.3796	0.7560	0.0	0.3921	3.0	0.662	0.337	0.629	0.512	0.513	1.025
2.583	0.75	0.25	0.3897	0.7394	0.0	0.3919	3.5	0.633	0.322	0.603	0.504	0.521	1.025
2.541	0.75	0.25	0.3978	0.7279	0.0	0.3916	4.0	0.615	0.315	0.588	0.496	0.526	1.023
2.492	0.75	0.25	0.4068	0.7124	0.0	0.3975	∞	0.586	0.302	0.591	0.483	0.530	1.013
6.871	1.00	0.00	0.1163	1.4450	0.0	0.2693	1.1	1.654	1.651	1.737	0.958	0.871	1.829
5.964	1.00	0.00	0.1691	1.3564	0.0	0.1453	1.45	1.453	1.345	1.564	0.812	0.790	1.601
5.211	1.00	0.00	0.1992	1.2903	0.0	0.1137	2.0	1.254	1.108	1.372	0.790	0.686	1.477
5.043	1.00	0.00	0.2085	1.2770	0.0	0.0980	2.2	1.213	1.057	1.324	0.771	0.678	1.449
4.858	1.00	0.00	0.2246	1.2563	0.0	0.0814	2.5	1.171	1.004	1.269	0.741	0.673	1.414
4.664	1.00	0.00	0.2389	1.2288	0.0	0.0755	3.0	1.111	0.965	1.202	0.716	0.671	1.386
4.558	1.00	0.00	0.2478	1.2093	0.0	0.0733	3.5	1.070	0.950	1.165	0.697	0.676	1.372
4.496	1.00	0.00	0.2547	1.2013	0.0	0.0698	4.0	1.056	0.938	1.145	0.684	0.673	1.357
4.395	1.00	0.00	0.2593	1.1853	0.0	0.0800	∞	1.012	0.927	1.110	0.678	0.668	1.346

^[a]All results are with fixed $a_1 = 0$; $s_8 = 0$; and $a_2 = 5.5$

Table S7: Optimized parameters, total WTAD2 and its decomposition into major subcategories for knoDispSDx-PBEP86. Heatmapping is from red (worst) via yellow to green (best). Thermo=Small Molecule Thermochemistry; Barrier=barrier heights; Large=reaction energies for large systems; Conf=conformer/intramolecular interactions; Intermol=intermolecular interactions; and NCI= Total Non-Covalent Interaction (Conformer/Intramolecular + Intermolecular).

WTAD2 (kcal/mol)	Parameters ^[a]							Δ WTAD2 (kcal/mol)					
	C _{X,HF}	C _{X,DFT}	C _{C,DFT}	C _{2ab}	C _{2ss}	S ₆	K	Thermo	Barrier	Large	Conf	Intermol	NCI
5.721	0.50	0.50	0.3345	0.5009	0.6659	0.0	1.1	1.308	0.734	0.880	1.475	1.325	2.800
5.410	0.50	0.50	0.3764	0.4575	0.5529	0.0	1.45	1.204	0.869	0.848	1.325	1.163	2.488
5.342	0.50	0.50	0.4651	0.3712	0.4429	0.0	2.0	1.041	0.932	0.891	1.314	1.164	2.478
5.355	0.50	0.50	0.4813	0.3566	0.4270	0.0	2.2	1.020	0.959	0.907	1.308	1.162	2.469
5.389	0.50	0.50	0.5129	0.3385	0.3854	0.0	2.5	0.976	0.961	0.932	1.329	1.191	2.519
5.448	0.50	0.50	0.5429	0.3141	0.3504	0.0	3.0	0.930	0.956	0.968	1.365	1.228	2.593
5.497	0.50	0.50	0.5623	0.3010	0.3294	0.0	3.5	0.909	0.956	0.993	1.388	1.251	2.639
5.536	0.50	0.50	0.5746	0.2893	0.3242	0.0	4.0	0.900	0.964	1.010	1.401	1.262	2.662
5.663	0.50	0.50	0.6049	0.2585	0.2977	0.0	∞	0.857	0.926	1.080	1.475	1.326	2.801
5.260	0.55	0.45	0.2682	0.5895	0.7012	0.0	1.1	1.299	0.637	0.787	1.295	1.242	2.537
4.860	0.55	0.45	0.3296	0.5234	0.5665	0.0	1.45	1.138	0.722	0.749	1.165	1.086	2.251
4.747	0.55	0.45	0.4166	0.4404	0.4628	0.0	2.0	0.985	0.807	0.776	1.128	1.052	2.180
4.751	0.55	0.45	0.4392	0.4218	0.4374	0.0	2.2	0.948	0.821	0.791	1.132	1.059	2.191
4.771	0.55	0.45	0.4680	0.3989	0.4091	0.0	2.5	0.905	0.835	0.813	1.144	1.074	2.218
4.819	0.55	0.45	0.4961	0.3722	0.3848	0.0	3.0	0.862	0.846	0.846	1.166	1.099	2.265
4.865	0.55	0.45	0.5143	0.3537	0.3737	0.0	3.5	0.839	0.855	0.870	1.184	1.117	2.301
4.903	0.55	0.45	0.5299	0.3406	0.3600	0.0	4.0	0.819	0.847	0.889	1.208	1.140	2.348
5.020	0.55	0.45	0.5636	0.3129	0.3272	0.0	∞	0.778	0.815	0.949	1.275	1.203	2.478
5.040	0.58	0.42	0.2460	0.6318	0.7052	0.0	1.1	1.279	0.589	0.761	1.210	1.202	2.411
4.568	0.58	0.42	0.3045	0.5583	0.5757	0.0	1.45	1.097	0.643	0.700	1.081	1.048	2.129
4.417	0.58	0.42	0.3854	0.4835	0.4680	0.0	2.0	0.944	0.723	0.716	1.032	1.003	2.035
4.413	0.58	0.42	0.4144	0.4630	0.4406	0.0	2.2	0.906	0.741	0.729	1.033	1.004	2.037
4.426	0.58	0.42	0.4378	0.4406	0.4188	0.0	2.5	0.868	0.759	0.750	1.039	1.011	2.050
4.465	0.58	0.42	0.4693	0.4115	0.3947	0.0	3.0	0.821	0.774	0.780	1.058	1.032	2.090
4.507	0.58	0.42	0.4870	0.3927	0.3838	0.0	3.5	0.796	0.781	0.804	1.076	1.049	2.126
4.542	0.58	0.42	0.4986	0.3849	0.3709	0.0	4.0	0.783	0.783	0.823	1.090	1.064	2.154
4.600	0.58	0.42	0.5262	0.3599	0.3501	0.0	∞	0.737	0.754	0.872	1.130	1.107	2.237
4.782	0.63	0.37	0.1936	0.7027	0.7474	0.0	1.1	1.277	0.567	0.741	1.071	1.127	2.198
4.174	0.63	0.37	0.2611	0.6231	0.5981	0.0	1.45	1.060	0.567	0.641	0.941	0.964	1.905
3.934	0.63	0.37	0.3448	0.5467	0.4738	0.0	2.0	0.872	0.597	0.641	0.890	0.933	1.823
3.917	0.63	0.37	0.3670	0.5230	0.4569	0.0	2.2	0.835	0.615	0.650	0.888	0.930	1.818
3.913	0.63	0.37	0.3939	0.5060	0.4264	0.0	2.5	0.796	0.634	0.662	0.888	0.932	1.821
3.933	0.63	0.37	0.4309	0.4774	0.3956	0.0	3.0	0.745	0.646	0.687	0.905	0.950	1.855
3.963	0.63	0.37	0.4503	0.4603	0.3805	0.0	3.5	0.718	0.651	0.707	0.920	0.965	1.886
3.992	0.63	0.37	0.4572	0.4489	0.3814	0.0	4.0	0.710	0.664	0.724	0.927	0.968	1.895
4.090	0.63	0.37	0.4764	0.4312	0.3695	0.0	∞	0.695	0.677	0.781	0.948	0.988	1.937
4.618	0.69	0.31	0.1444	0.8036	0.7424	0.0	1.1	1.263	0.568	0.783	0.924	1.078	2.003
3.864	0.69	0.31	0.2065	0.7231	0.5995	0.0	1.45	1.031	0.516	0.646	0.785	0.887	1.672
3.480	0.69	0.31	0.2959	0.6381	0.4827	0.0	2.0	0.829	0.516	0.585	0.731	0.821	1.551
3.439	0.69	0.31	0.3145	0.6186	0.4659	0.0	2.2	0.791	0.526	0.585	0.724	0.813	1.537
3.408	0.69	0.31	0.3383	0.5953	0.4451	0.0	2.5	0.746	0.536	0.592	0.722	0.813	1.535
3.403	0.69	0.31	0.3680	0.5649	0.4214	0.0	3.0	0.693	0.537	0.612	0.731	0.829	1.560
3.415	0.69	0.31	0.3872	0.5464	0.4082	0.0	3.5	0.665	0.542	0.628	0.740	0.841	1.580
3.433	0.69	0.31	0.4049	0.5331	0.3922	0.0	4.0	0.641	0.535	0.643	0.753	0.861	1.614
3.509	0.69	0.31	0.4265	0.5149	0.3757	0.0	∞	0.618	0.540	0.693	0.773	0.885	1.658
4.592	0.75	0.25	0.1094	0.8806	0.7575	0.0	1.1	1.256	0.625	0.877	0.823	1.011	1.835
3.742	0.75	0.25	0.1750	0.8003	0.5994	0.0	1.45	0.999	0.512	0.723	0.681	0.828	1.509
3.224	0.75	0.25	0.2502	0.7250	0.4890	0.0	2.0	0.793	0.474	0.603	0.613	0.740	1.353
3.144	0.75	0.25	0.2680	0.7094	0.4661	0.0	2.2	0.751	0.473	0.582	0.604	0.733	1.337
3.071	0.75	0.25	0.2979	0.6836	0.4393	0.0	2.5	0.700	0.469	0.570	0.600	0.733	1.333

3.026	0.75	0.25	0.3250	0.6611	0.4161	0.0	3.0	0.654	0.472	0.573	0.597	0.732	1.328
3.018	0.75	0.25	0.3397	0.6422	0.4107	0.0	3.5	0.626	0.476	0.582	0.600	0.736	1.335
3.023	0.75	0.25	0.3481	0.6305	0.4089	0.0	4.0	0.612	0.479	0.591	0.602	0.739	1.341
3.068	0.75	0.25	0.3598	0.6192	0.3993	0.0	∞	0.595	0.488	0.633	0.603	0.749	1.353
6.195	1.00	0.00	-0.0676	1.2333	0.8180	0.0	1.1	1.442	1.359	1.535	0.913	0.948	1.860
5.266	1.00	0.00	-0.0069	1.1538	0.6469	0.0	1.45	1.168	1.133	1.350	0.828	0.786	1.615
4.585	1.00	0.00	0.0743	1.0693	0.5295	0.0	2.0	0.946	0.988	1.181	0.776	0.693	1.469
4.445	1.00	0.00	0.0925	1.0532	0.5064	0.0	2.2	0.901	0.961	1.141	0.765	0.678	1.443
4.296	1.00	0.00	0.1153	1.0329	0.4797	0.0	2.5	0.851	0.937	1.095	0.747	0.667	1.414
4.150	1.00	0.00	0.1438	1.0272	0.4267	0.0	3.0	0.801	0.915	1.048	0.719	0.667	1.386
4.074	1.00	0.00	0.1529	1.0229	0.4133	0.0	3.5	0.774	0.902	1.027	0.710	0.661	1.371
4.033	1.00	0.00	0.1600	1.0173	0.4073	0.0	4.0	0.759	0.896	1.018	0.702	0.658	1.360
3.971	1.00	0.00	0.1662	1.0183	0.3938	0.0	∞	0.731	0.886	1.008	0.691	0.655	1.346

^[a]All results are with fixed $a_1 = 0$; $s_8 = 0$; and $a_2 = 5.5$

Table S8: Optimized parameters, total WTMAD2 and its decomposition into major subcategories for κ DSD₆₉-PBEP86-D3BJ and κ DSD₆₉-PBEPBE-D3BJ@PBE0P86. Heatmapping is from red (worst) via yellow to green (best). Thermo=Small Molecule Thermochemistry; Barrier=barrier heights; Large=reaction energies for large systems; Conf=conformer/intramolecular interactions; Intermol=intermolecular interactions; and NCI= Total Non-Covalent Interaction (Conformer/Intramolecular + Intermolecular).

Functionals	WTMAD2 (kcal/mol)	Parameters ^[a]					Δ WTMAD2 (kcal/mol)					
		C _{C,DFT}	C _{2ab}	C _{2ss}	S ₆	κ	Thermo	Barrier	Large	Conf	Intermol	NCI
κ DSD ₆₉ -PBEP86-D3BJ@ PBE0P86	3.821	0.1812	0.7392	0.2924	0.6121	1.1	1.136	0.638	0.727	0.680	0.641	1.321
	3.081	0.2537	0.6253	0.2330	0.5253	1.45	0.960	0.424	0.577	0.600	0.519	1.120
	2.594	0.3415	0.5054	0.1953	0.4683	2.0	0.800	0.305	0.479	0.547	0.463	1.010
	2.515	0.3597	0.4833	0.1880	0.4509	2.2	0.767	0.298	0.455	0.533	0.462	0.995
	2.436	0.3845	0.4579	0.1780	0.4331	2.5	0.726	0.297	0.425	0.516	0.473	0.989
	2.366	0.4305	0.4330	0.1417	0.4255	3.0	0.670	0.296	0.409	0.484	0.507	0.991
	2.326	0.4587	0.4165	0.1165	0.4325	3.5	0.644	0.290	0.401	0.465	0.526	0.991
	2.299	0.4757	0.4088	0.0977	0.4409	4.0	0.627	0.287	0.393	0.456	0.537	0.993
	2.276	0.5368	0.3811	0.0381	0.4883	∞	0.590	0.242	0.434	0.419	0.591	1.010
κ DSD ₆₉ -PBEP86-D3BJ	3.404	0.1713	0.8220	0.3588	0.5073	1.1	1.021	0.557	0.683	0.617	0.527	1.144
	2.807	0.2339	0.7410	0.2935	0.4331	1.45	0.842	0.386	0.576	0.555	0.449	1.004
	2.477	0.3176	0.6534	0.2193	0.3953	2.0	0.665	0.301	0.520	0.508	0.482	0.990
	2.428	0.3397	0.6336	0.2003	0.3950	2.2	0.628	0.291	0.512	0.498	0.500	0.998
	2.383	0.3647	0.6148	0.1740	0.3960	2.5	0.589	0.281	0.507	0.486	0.519	1.006
	2.352	0.3906	0.5974	0.1474	0.3989	3.0	0.554	0.277	0.505	0.474	0.542	1.016
	2.346	0.4075	0.5864	0.1220	0.4111	3.5	0.543	0.270	0.513	0.469	0.552	1.021
	2.348	0.4164	0.5824	0.1085	0.4175	4.0	0.534	0.269	0.519	0.466	0.560	1.026
	2.367	0.4312	0.5761	0.0816	0.4339	∞	0.521	0.268	0.552	0.457	0.569	1.026

^[a]All results are with fixed $a_1 = 0$; $s_8 = 0$; and $a_2 = 5.5$.

Table S9: Total WTMAD2 and its decomposition into major subcategories for selected nonempirical DHDFs. Thermo=Small Molecule Thermochemistry; Barrier=barrier heights; Large=reaction energies for large systems; Conf=conformer/intramolecular interactions; Intermol=intermolecular interactions; and NCI= Total Non-Covalent Interaction (Conformer/Intramolecular + Intermolecular).

Functionals	κ	WTMAD2 (kcal/mol)	Δ WTMAD2 (kcal/mol)					
			Thermo	Barrier	Large	Conf	Intermol	NCI
PBE0-DH-D3BJ ^{1,2}	1.1	5.745	1.469	0.629	1.282	1.096	1.270	2.366
	1.45	5.637	1.415	0.641	1.237	1.052	1.291	2.344
	2.0	5.545	1.357	0.657	1.183	1.027	1.320	2.347
	∞	5.439	1.273	0.698	1.100	1.003	1.365	2.368
SOS1-PBE-QIDH-D3BJ ^{3,4}	1.1	5.984	1.389	0.912	1.354	1.233	1.096	2.329
	1.45	5.233	1.223	0.760	1.234	1.024	0.993	2.016
	∞	3.838	0.874	0.456	0.873	0.768	0.867	1.635
SOSO-PBE0-2-D3BJ ^{5,6}	1.1	6.401	1.429	1.217	1.452	1.016	1.287	2.303
	1.45	5.347	1.214	0.994	1.293	0.825	1.021	1.846
	∞	3.461	0.736	0.561	0.846	0.672	0.647	1.320
PBE0-DH ¹	1.1	9.225	1.396	0.591	1.545	2.814	2.879	5.693
	1.45	8.822	1.330	0.568	1.495	2.678	2.750	5.428
	2.0	8.466	1.261	0.563	1.440	2.565	2.638	5.203
	∞	8.080	1.163	0.568	1.361	2.465	2.524	4.989
SOS1-PBE-QIDH ³	1.1	9.611	1.374	1.011	1.534	2.638	3.055	5.693
	1.45	8.716	1.193	0.852	1.411	2.416	2.843	5.259
	∞	7.057	0.820	0.515	1.074	2.122	2.525	4.648
SOSO-PBE0-2 ⁵	1.1	9.686	1.392	1.345	1.577	2.409	2.964	5.373
	1.45	8.521	1.176	1.122	1.419	2.121	2.683	4.805
	∞	6.460	0.700	0.664	0.998	1.774	2.325	4.099

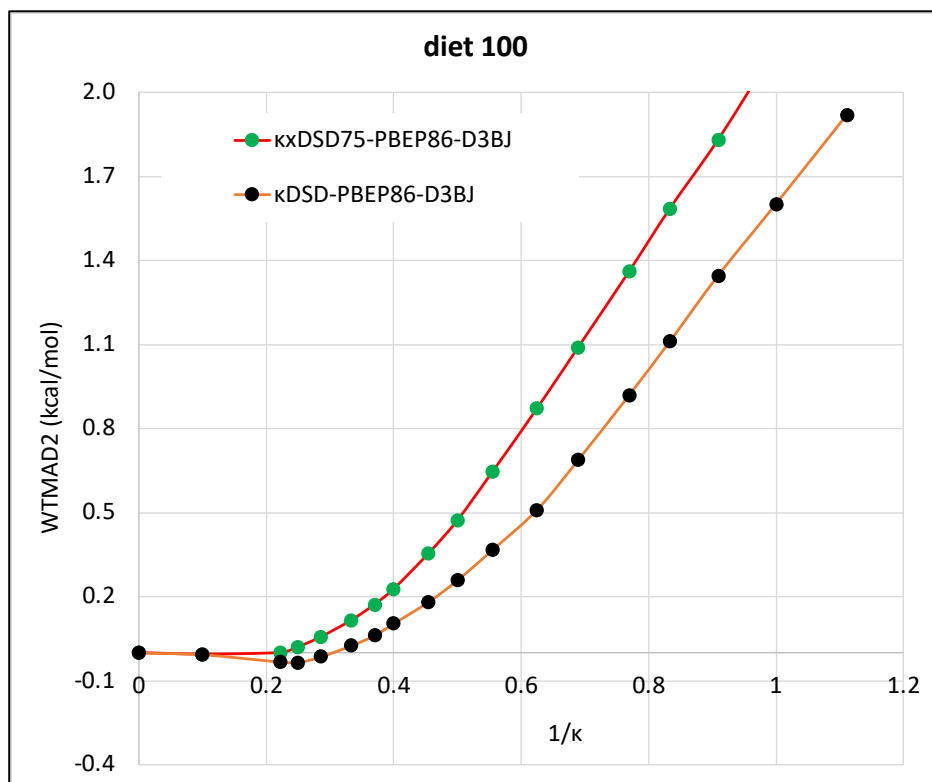


Figure S1: Dependence of WTMAD2 (kcal/mol) on reciprocal κ for κDSD -PBEP86-D3BJ and $\kappa xDS75$ -PBEP86-D3BJ.

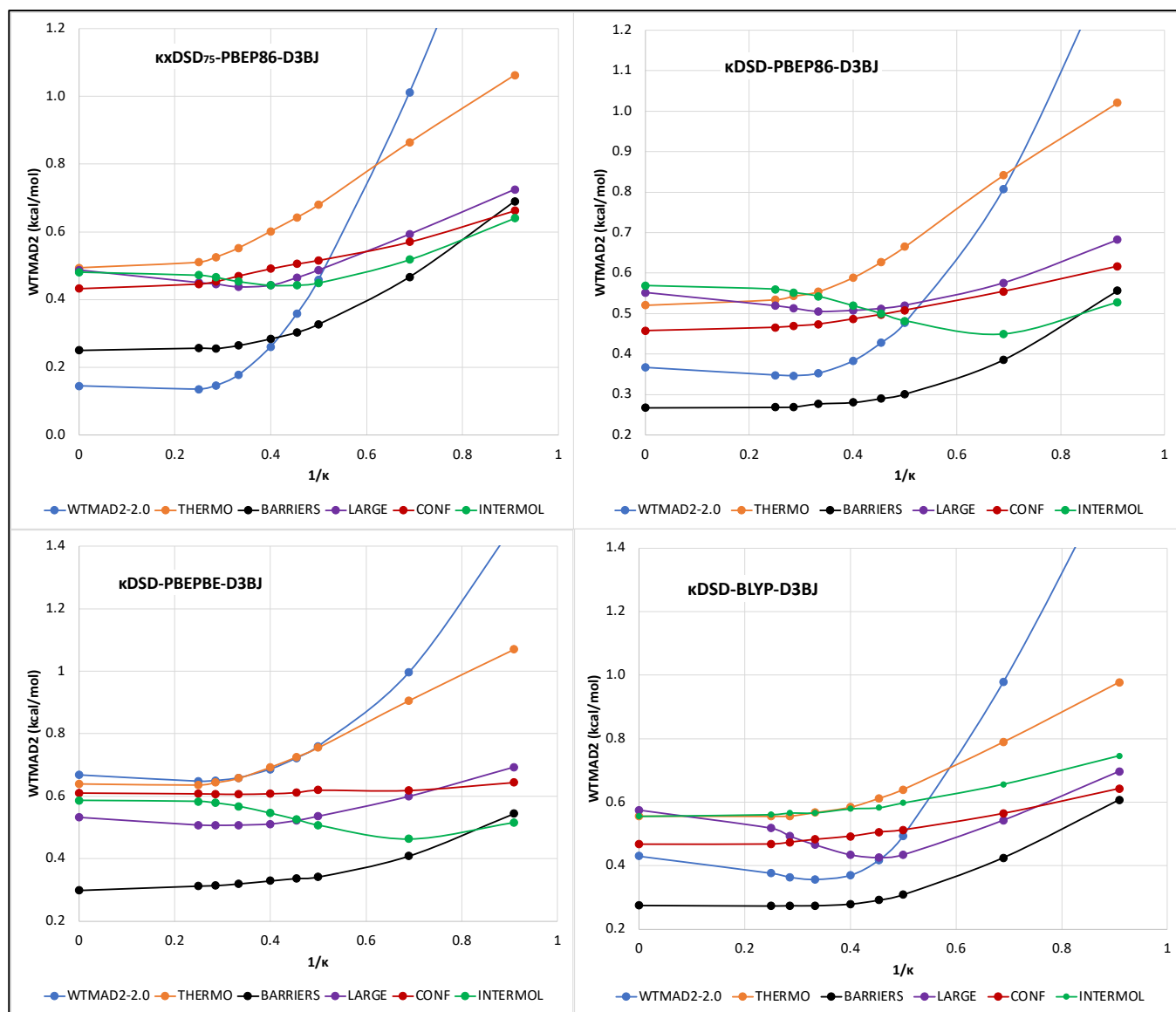


Figure S2: Dependence of total WTMA2 (kcal/mol) and contribution (Δ WTMA2 in kcal/mol) from five major subcategories on reciprocal κ for $\kappa \times \text{DSD}_{75}\text{-PBEP86-D3BJ}$, $\kappa \text{DSD-PBEP86-D3BJ}$, $\kappa \text{DSD-PBEPBE-D3BJ}$, $\kappa \text{DSD-BLYP-D3BJ}$, and $\kappa \times \text{DSD}_{75}\text{-PBEP86-D3BJ}$. THERMO= Small Molecule Thermochemistry; BARRIER= barrier heights; LARGE= reaction energies for large systems; CONF= conformer/intramolecular interactions; and INTERMOL= intermolecular interactions.

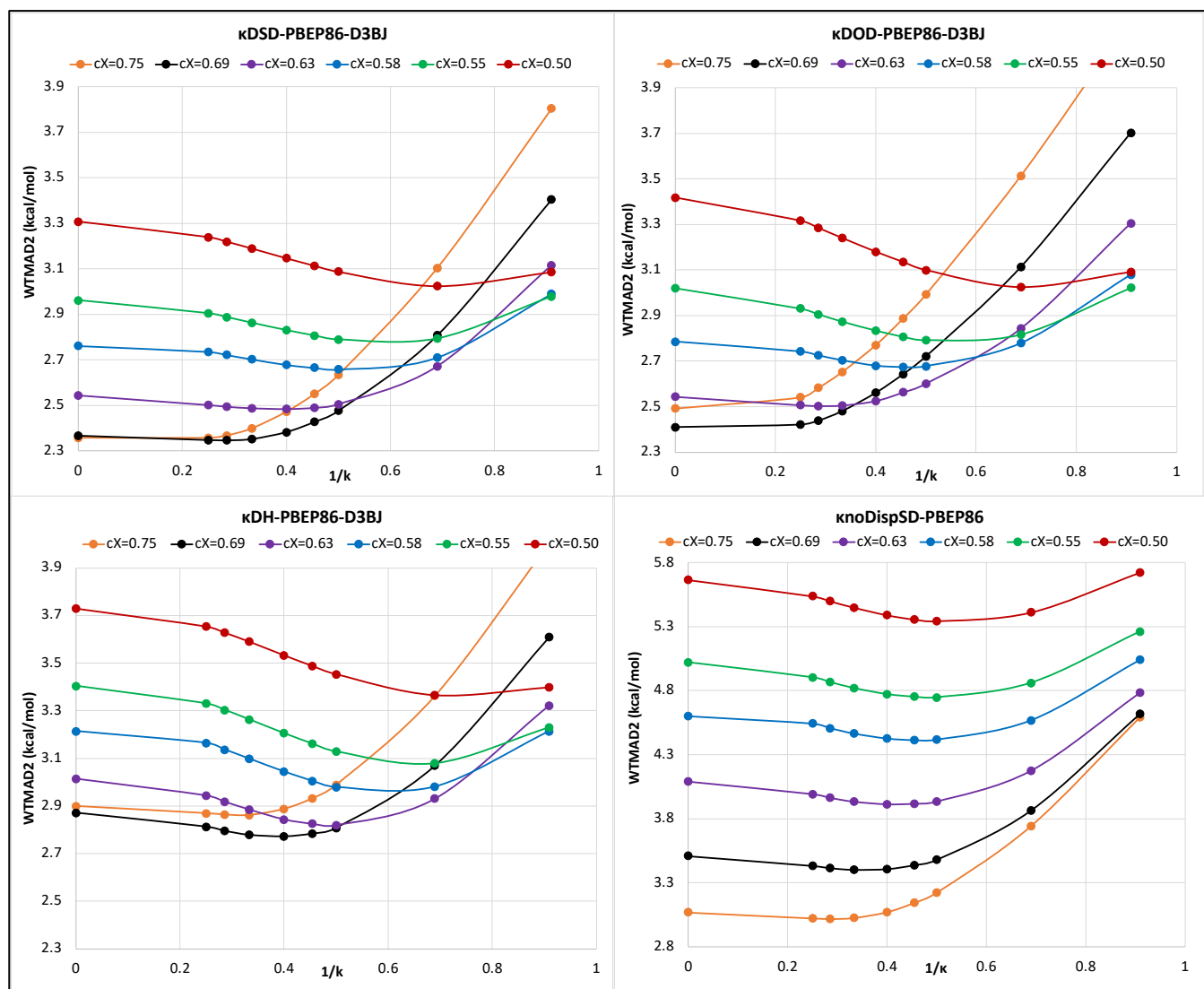


Figure S3: Dependence of WTMAD2 (kcal/mol) on reciprocal κ for κ DSD-PBEP86-D3BJ, κ DOD-PBEP86-D3BJ, κ DH-PBEP86-D3BJ, and κ noDispSD-PBEP86 functionals. Six colors represent six different fractions of exact exchange (c_X) ranging from 0.75 to 0.50

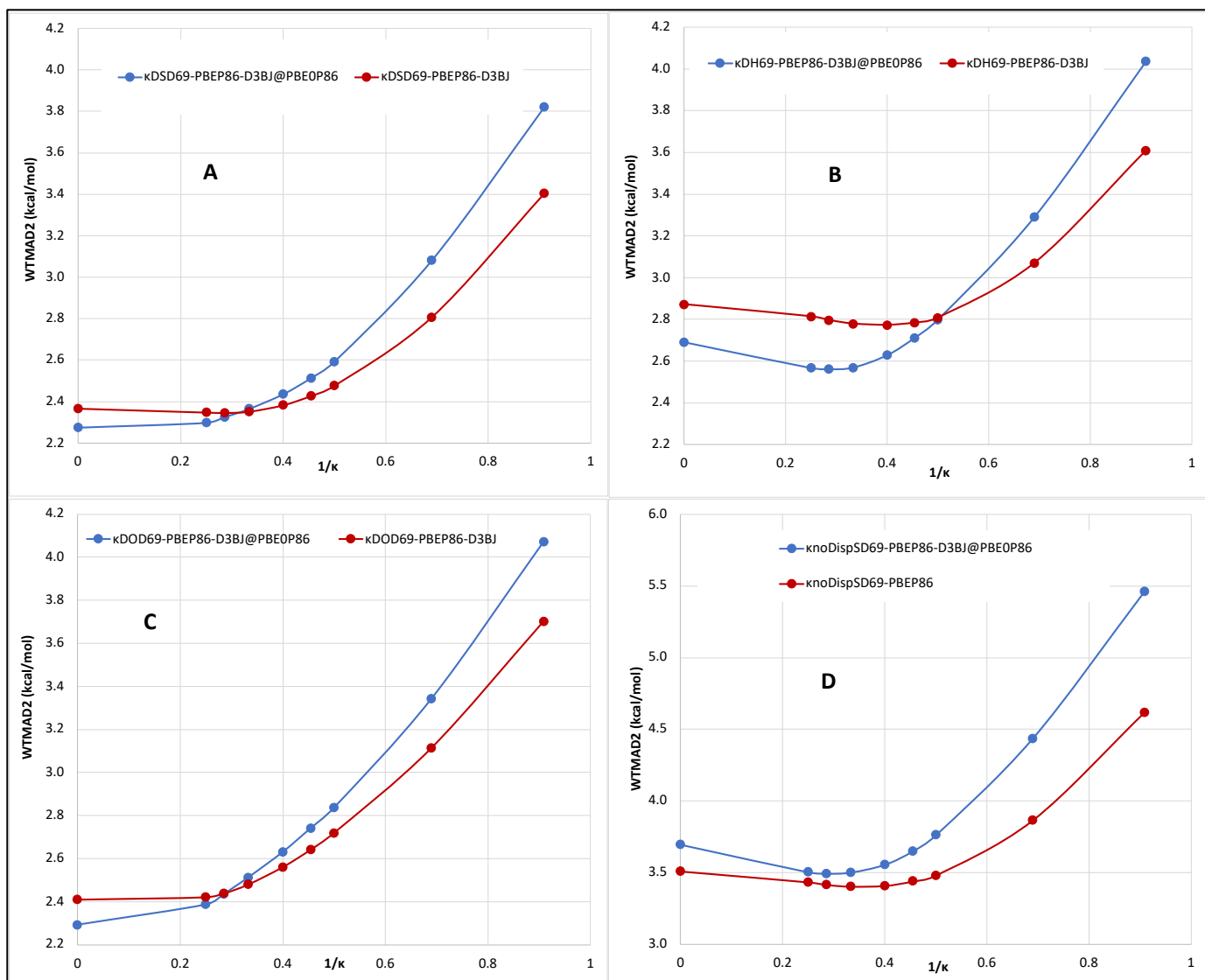


Figure S4: Dependence of total WTMA2 (kcal/mol) on reciprocal κ for κ DSD₆₉-PBEP86-D3BJ@PBE0P86 and κ DSD₆₉-PBEP86-D3BJ (A); their κ DH counterparts (B); κ DOD variants (C); and the dispersion-*uncorrected* forms (D).

Sample Input:

κ DSD-BLYP-D3BJ ($\kappa=2.2$)

```
$rem
SYM_IGNORE           TRUE
SYMMETRY             false
BASIS                 def2-QZVPP
exchange             GEN
XC_GRID              3
ECP                  def2-ECP
MAX_SCF_CYCLES       1000
SCF_CONVERGENCE       7
THRESH               12
MEM_STATIC            2000
SET_ITER             100
MOLDEN_FORMAT        false
PRINT_ORBITALS        10
SCF_FINAL_PRINT       1
DFT_D                D3_BJ
DFT_D3_S6            52360
DFT_D3_S8            0
DFT_D3_A2            520000
DFT_D3_A1            0
$end

$xc_functional
C LYP 0.4387
X HF 0.7100
X B88 0.2900
$end

$molecule
0 1
C          0.000000000000    0.000000000000    0.000000000000
O          0.000000000000    0.000000000000    1.131400000000
$end

@@@

$molecule
read
$end

$rem
SCF_GUESS            read
basis                def2-QZVPP
aux_basis            rimp2-def2-QZVPPD
correlation          rimp2
SCS                  3
SSS_FACTOR           272400
SOS_FACTOR           616600
SYMMETRY             FALSE
SYM_IGNORE           TRUE
gen_scfman           true
REGULARIZED_O2       2
REG_VARIABLE         2200
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MEM_TOTAL	64000
MEM_STATIC	2000
n_frozen_core	FC
SCF_CONVERGENCE	7
THRESH	12
MAXSCF	0
ECP	def2-ECP

\$end

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