

Supporting Information for: “Standard grids for high-precision integration of modern density functionals: SG-2 and SG-3”

Saswata Dasgupta and John M. Herbert*

Department of Chemistry and Biochemistry, The Ohio State University, Columbus, Ohio 43210 USA

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The following data can be found below:

- Table S1: Atomization energies computed for the G2 data set at the M06-2X/aug-cc-pVTZ level of theory, including both pruning errors and grid errors for SG-2 and SG-3. Pruning errors remain small (MUD = 0.09 kcal/mol for SG-2 and 0.03 kcal/mol for SG-3), albeit slightly larger than for ω B97X-V/aug-cc-pVTZ (*cf.* Table II).
- Table S2: Atomization energies computed for the G2 data set at the ω B97M-V/aug-cc-pVTZ level of theory, including both pruning errors and grid errors for SG-2 and SG-3. Pruning errors remain small (MUD = 0.09 kcal/mol for SG-2 and 0.03 kcal/mol for SG-3), albeit slightly larger than for ω B97X-V/aug-cc-pVTZ (*cf.* Table II).
- Table S3 Isomerization energies for organic reactions at the ω B97M-V/TZV(d,p) level using different integration grids. Both the pruning and grid errors are observed to be very small for SG-2 and SG-3.
- Table S4: Comparison of geometrical parameters optimized at the ω B97M-V/aug-cc-pVTZ level using various integration grids. Both the pruning and the grid errors for bond lengths and bond angles are quite small, for both SG-2 and SG-3.
- Table S5: Harmonic vibrational frequencies computed at the ω B97M-V/aug-cc-pVTZ level using SG-2 and SG-3 grids. Both pruning and grid errors are quite small.
- Table S6: Harmonic vibrational frequencies at the ω B97X-D/aug-cc-pVTZ level using SG-2 and SG-3. In this case, analytic Hessians are available in Q-CHEM so we compare these analytic results to frequencies computed via finite difference of analytic gradients. Results show that there is very little difference between the two approaches, which validates other finite-difference frequencies reported in this work. Errors in ω B97X-D frequencies are comparable to those observed using M11, both of which exhibit larger errors as compared to ω B97X-V or ω B97M-V.
- Tables S8, S7 and S9: Interaction energies for the S66 data set of non-covalent dimers (using benchmark S66 geometries), computed at the ω B97X-V/, M06-2X/, and ω B97M-V/aug-cc-pVTZ levels of theory, respectively. Both pruning errors and grid errors are quite small, for all three methods and for both SG-2 and SG-3.
- Table S10: Tests of rotational invariance. Isomerization energies are computed for three different reactions upon various rigid rotations of the product and reactant species. Rotational invariance is preserved to better than 0.01 kcal/mol in each case.
- Figure S1: Potential energy curves for the “sandwich” isomer of $(\text{C}_6\text{H}_6)_2$ along the face-to-face distance coordinate, computed at the M06-2X/6-311++G(3df,3pd) level using the low-quality SG-0 and SG-1 integration grids. Very large, spurious oscillations in the potential energy curves are obtained.

*herbert@chemistry.ohio-state.edu

TABLE S1: Atomization energies^a (in kcal/mol) at the M06-2X/aug-cc-pVTZ level of theory.

Molecule	Energy		Error		Energy		Error	
	SG-2	(75,302)	pruning ^b	grid ^c	SG-3	(99,590)	pruning ^b	grid ^c
BeH	678.62	678.62	0.00	-0.01	678.61	678.61	0.01	0.00
CH ₂ CH ₂	561.22	561.05	-0.17	0.01	561.12	561.06	-0.06	0.00
CH ₂ O	372.04	371.98	-0.06	0.00	372.00	371.98	-0.03	0.00
CH ₂ singlet	179.13	179.09	-0.04	0.01	179.11	179.10	-0.01	0.00
CH ₂ triplet	192.48	192.35	-0.13	0.01	192.44	192.36	-0.08	0.00
CH ₃ CH ₃	709.84	709.78	-0.06	0.03	709.82	709.81	-0.01	0.00
CH ₃ Cl	394.31	394.20	-0.11	0.00	394.22	394.20	-0.02	0.00
CH ₃ OH	512.02	511.99	-0.03	0.00	512.02	511.99	-0.02	0.00
CH ₃	306.26	306.14	-0.12	0.01	306.20	306.15	-0.05	0.00
CH ₃ SH	472.40	472.23	-0.17	0.06	472.33	472.28	-0.05	0.00
CH ₄	417.83	417.95	0.12	0.00	417.84	417.95	0.11	0.00
CH	83.19	83.19	0.00	0.01	83.20	83.20	0.00	0.00
Cl ₂	58.64	58.45	-0.19	0.01	58.43	58.46	0.03	0.00
ClF	60.83	60.74	-0.09	-0.02	60.89	60.72	-0.17	0.00
ClO	62.35	62.34	-0.01	0.00	62.35	62.34	-0.01	0.00
CN	175.70	175.63	-0.07	0.03	175.66	175.66	0.00	0.00
CO ₂	387.31	387.33	0.02	-0.01	387.32	387.32	0.01	0.00
CO	257.02	257.02	0.00	0.00	257.02	257.02	0.00	0.00
CS	167.28	167.24	-0.04	0.03	167.29	167.27	-0.02	0.00
F ₂	32.32	32.38	0.06	-0.01	32.28	32.37	0.10	0.00
H ₂ O	230.62	230.65	0.03	0.00	230.65	230.66	0.01	0.00
HCCH	403.04	402.98	-0.06	0.02	403.00	402.99	-0.01	0.00
HCl	105.56	105.51	-0.05	-0.04	105.52	105.48	-0.04	0.00
HCN	309.34	309.27	-0.07	0.03	309.30	309.30	0.00	0.00
HCO	277.03	277.03	0.00	0.00	277.02	277.03	0.01	0.00
HF	139.51	139.48	-0.03	0.00	139.50	139.48	-0.02	0.00
HOCl	164.23	164.14	-0.09	0.00	164.18	164.14	-0.03	0.00
HOOH	265.96	265.99	0.03	0.01	265.97	266.00	0.03	0.00
Li ₂	24.39	24.25	-0.14	0.08	24.32	24.33	0.00	0.00
LiF	137.34	137.29	-0.05	0.03	137.29	137.32	0.02	0.00
LiH	57.02	56.96	-0.06	0.03	56.99	56.99	0.00	0.00
N ₂	222.65	222.53	-0.12	0.03	222.56	222.56	0.00	0.00
Na ₂	19.90	19.88	-0.02	0.01	19.89	19.90	0.00	0.00
NaCl	98.30	98.26	-0.04	0.01	98.29	98.27	-0.01	0.00
NH ₂ NH ₂	433.09	433.03	-0.06	0.04	433.04	433.07	0.03	0.00
NH ₂	180.89	180.74	-0.15	0.02	180.78	180.76	-0.01	0.00
NH ₃	295.48	295.44	-0.04	0.02	295.46	295.46	0.00	0.00
NH	83.18	83.06	-0.12	0.02	83.07	83.08	0.01	0.00
NO	150.50	150.53	0.03	0.02	150.55	150.55	0.00	0.00
O ₂	118.15	118.17	0.02	-0.01	118.13	118.16	0.03	0.00
OH	106.45	106.44	-0.01	0.00	106.45	106.44	-0.01	0.00
P ₂	117.10	117.01	-0.09	0.09	117.10	117.09	-0.01	0.00
PH ₂	155.96	155.91	-0.05	0.04	155.95	155.95	0.00	0.00
PH ₃	241.69	241.66	-0.03	0.04	241.68	241.70	0.02	0.00
S ₂	102.59	102.35	-0.24	-0.02	102.46	102.34	-0.12	0.00
SH ₂	181.23	181.14	-0.09	0.02	181.15	181.15	0.01	0.00
Si ₂	74.31	74.09	-0.22	0.06	74.18	74.14	-0.03	0.00
SiH ₂ singlet	153.56	153.38	-0.18	0.01	153.39	153.38	-0.01	0.00
SiH ₂ triplet	134.21	134.04	-0.17	0.03	134.08	134.06	-0.03	0.00
SiH ₃	227.24	227.02	-0.22	0.02	227.07	227.04	-0.02	0.00
SiH ₃ SiH ₃	529.09	528.64	-0.45	0.03	528.69	528.68	-0.01	0.00
SiH ₄	320.57	320.44	-0.13	0.02	320.42	320.47	0.05	0.00
SiO	188.07	187.85	-0.22	0.03	187.90	187.88	-0.02	0.00
SO ₂	239.46	239.40	-0.06	-0.01	239.37	239.38	0.01	0.00
SO	121.43	121.50	0.07	-0.02	121.48	121.49	0.01	0.00
MUD ^d			0.09	0.03			0.03	0.00

^aUsing MP2/6-31G* geometries. ^bDifference between pruned and unpruned results.^cDifference with respect to a (250,974) grid. ^dMean unsigned deviation.

TABLE S2: Atomization energies^a (in kcal/mol) at the ω B97M-V/aug-cc-pVTZ level of theory.

Molecule	Energy		Error		Energy		Error	
	SG-2	(75,302)	pruning ^b	grid ^c	SG-3	(99,590)	pruning ^b	grid ^c
BeH	670.94	670.94	0.00	0.00	670.94	670.94	0.00	0.00
CH ₂ CH ₂	562.69	562.58	-0.10	0.00	562.61	562.58	-0.03	0.00
CH ₂ O	373.34	373.28	-0.06	0.00	373.30	373.28	-0.03	0.00
CH ₂ singlet	181.19	181.15	-0.04	0.00	181.18	181.15	-0.03	0.00
CH ₂ triplet	190.09	190.07	-0.02	0.00	190.07	190.07	0.00	0.00
CH ₃ CH ₃	712.00	711.92	-0.07	0.00	711.95	711.92	-0.03	0.00
CH ₃ Cl	395.41	395.34	-0.08	0.00	395.36	395.34	-0.02	0.00
CH ₃ OH	514.31	514.33	0.02	0.00	514.33	514.33	0.00	0.00
CH ₃	307.89	307.88	-0.01	-0.03	307.90	307.88	-0.02	-0.03
CH ₃ SH	474.99	474.92	-0.07	0.00	474.94	474.92	-0.02	0.00
CH ₄	419.55	419.47	-0.08	0.00	419.51	419.47	-0.04	0.00
CH	84.18	84.19	0.00	0.00	84.18	84.19	0.00	0.00
Cl ₂	58.92	59.04	0.12	0.00	59.07	59.04	-0.02	0.00
ClF	62.16	62.17	0.01	0.00	62.14	62.17	0.03	0.00
ClO	64.53	64.50	-0.04	0.00	64.50	64.50	0.00	0.00
CN	174.31	174.28	-0.03	0.00	174.29	174.28	-0.01	0.00
CO ₂	388.11	388.06	-0.05	0.00	388.08	388.06	-0.03	0.00
CO	258.70	258.65	-0.05	0.00	258.68	258.65	-0.02	0.00
CS	169.78	169.76	-0.02	0.00	169.79	169.76	-0.03	0.00
F ₂	37.11	37.13	0.01	0.00	37.16	37.13	-0.03	0.00
H ₂ O	231.45	231.42	-0.03	0.00	231.44	231.42	-0.02	0.00
HCCH	402.77	402.76	-0.01	0.00	402.73	402.76	0.03	0.00
HCl	106.78	106.79	0.01	0.00	106.79	106.79	0.01	0.00
HCN	311.38	311.38	-0.01	0.00	311.36	311.38	0.01	0.00
HCO	278.82	278.77	-0.05	0.00	278.80	278.77	-0.03	0.00
HF	139.51	139.52	0.01	0.00	139.52	139.52	0.00	0.00
HOCl	165.22	165.24	0.02	0.00	165.24	165.24	0.00	0.00
HOOH	268.43	268.39	-0.05	0.00	268.42	268.39	-0.03	0.00
Li ₂	23.82	23.81	0.00	0.00	23.82	23.81	-0.01	0.00
LiF	138.02	138.02	0.00	0.00	138.01	138.02	0.01	0.00
LiH	60.47	60.47	0.00	0.00	60.48	60.47	0.00	0.00
N ₂	226.89	226.85	-0.04	0.00	226.85	226.85	0.00	0.00
Na ₂	17.06	17.06	0.00	0.00	17.06	17.06	0.00	0.00
NaCl	97.08	97.09	0.01	0.00	97.08	97.09	0.00	0.00
NH ₂ NH ₂	431.33	431.33	0.00	0.00	-431.33	-431.33	0.00	0.00
NH ₂	181.51	181.52	0.01	0.00	-181.52	-181.52	0.00	0.00
NH ₃	299.06	299.06	-0.01	0.00	299.06	299.06	0.00	0.00
NH	85.01	85.00	-0.01	0.00	84.98	85.00	0.02	0.00
NO	154.00	153.99	-0.01	0.00	154.00	153.99	0.00	0.00
O ₂	120.56	120.53	-0.03	0.00	120.55	120.53	-0.02	0.00
OH	107.27	107.30	0.03	0.00	107.28	107.30	0.02	0.00
P ₂	115.72	115.75	0.03	0.00	115.77	115.75	-0.02	0.00
PH ₂	156.17	156.14	-0.03	0.00	156.15	156.14	-0.01	0.00
PH ₃	243.57	243.55	-0.01	0.01	243.58	243.55	-0.03	0.01
S ₂	102.41	102.39	-0.01	0.00	102.42	102.39	-0.03	0.00
SH ₂	183.68	183.64	-0.04	0.00	183.65	183.64	-0.01	0.00
Si ₂	68.08	68.08	0.00	0.00	68.09	68.08	-0.01	0.00
SiH ₂ singlet	154.34	154.20	-0.14	-0.01	154.21	154.20	-0.01	-0.01
SiH ₂ triplet	130.15	130.04	-0.10	0.00	130.06	130.04	-0.02	0.00
SiH ₃	225.60	225.46	-0.14	-0.01	225.46	225.46	0.00	-0.01
SiH ₃ SiH ₃	530.90	530.61	-0.29	-0.01	530.62	530.61	-0.01	-0.01
SiH ₄	322.20	322.05	-0.15	0.00	322.07	322.05	-0.02	0.00
SiO	189.54	189.40	-0.14	0.00	189.42	189.40	-0.02	0.00
SO ₂	245.05	245.02	-0.03	0.00	245.04	245.02	-0.02	0.00
SO	123.19	123.13	-0.06	0.00	123.15	123.13	-0.02	0.00
MUD ^d			0.04	0.00			0.01	0.00

^aUsing MP2/6-31G* geometries. ^bDifference between pruned and unpruned results.^cDifference with respect to a (250,974) grid. ^dMean unsigned deviation.

TABLE S3: Isomerization energies (in kcal/mol) for organic reactions at the ω B97M-V/TZV(d,p) level.

Rxn.	Energy		Error		Energy		Error	
	SG-2	(75,302)	pruning ^a	grid ^b	SG-3	(99,590)	pruning ^a	grid ^b
1	-0.35	-0.34	0.01	0.02	-0.32	-0.32	0.00	0.00
2	21.61	21.60	-0.01	0.03	21.63	21.63	0.00	0.00
3	5.78	5.80	0.02	0.06	5.86	5.86	0.00	0.00
4	0.99	1.05	0.06	0.04	1.10	1.10	0.00	0.00
5	1.22	1.17	-0.05	-0.05	1.11	1.11	0.00	0.00
6	2.65	2.68	0.03	0.03	2.71	2.71	0.00	0.00
7	9.58	9.62	0.03	0.09	9.71	9.71	0.00	0.00
8	21.84	21.81	-0.03	-0.03	21.78	21.78	0.00	0.00
9	6.09	6.11	0.02	0.11	6.21	6.22	0.00	0.00
10	3.64	3.54	-0.10	-0.04	3.56	3.51	-0.05	0.00
11	0.96	1.05	0.08	0.14	1.18	1.19	0.01	0.00
13	37.87	37.73	-0.13	0.09	37.82	37.83	0.01	0.00
14	22.53	22.52	-0.02	0.01	22.52	22.52	0.00	0.00
15	7.98	7.98	0.01	-0.01	7.98	7.98	0.00	0.00
16	8.93	8.95	0.02	0.04	9.00	8.99	0.00	0.00
17	27.46	27.49	0.02	0.07	27.56	27.56	0.00	0.00
18	11.33	11.31	-0.02	-0.02	11.29	11.29	0.00	0.00
19	4.70	4.67	-0.03	0.05	4.71	4.72	0.01	0.00
20	18.56	18.58	0.02	-0.04	18.55	18.55	-0.01	0.00
21	1.18	1.16	-0.03	0.03	1.18	1.18	0.00	0.00
22	3.68	3.65	-0.03	-0.01	3.64	3.64	0.00	0.00
23	5.36	5.35	-0.01	-0.01	5.33	5.35	0.01	0.00
24	11.76	11.76	0.00	0.00	11.75	11.76	0.00	0.00
25	25.82	25.84	0.02	0.04	25.88	25.88	0.00	0.00
26	17.41	17.40	0.00	0.02	17.42	17.42	0.00	0.00
27	64.81	64.83	0.02	0.05	64.88	64.88	0.00	0.00
28	30.61	30.62	0.01	-0.01	30.61	30.61	0.00	0.00
29	13.11	13.13	0.02	0.03	13.15	13.16	0.01	0.00
30	10.32	10.31	-0.01	0.00	10.31	10.31	0.00	0.00
31	15.67	15.70	0.03	-0.03	15.67	15.67	0.00	0.00
32	6.92	6.96	0.04	0.06	7.02	7.02	0.00	0.00
33	8.78	8.79	0.00	0.04	8.82	8.83	0.01	0.00
34	7.33	7.30	-0.03	0.01	7.30	7.30	0.00	0.00
MUD ^d			0.03	0.04			0.01	0.00

^aDifference between pruned and unpruned results. ^bDifference with respect to a (250,974) grid. ^dMean unsigned deviation.

TABLE S4: Comparison of geometrical parameters^a optimized at the ω B97M-V/aug-cc-pVTZ level.

Molecule	Parameter	Value		Error		Value		Error	
		SG-2	(75,302)	pruning ^b	grid ^c	SG-3	(99,590)	pruning ^b	grid ^c
H ₂	$r(\text{HH})$	0.74073	0.74029	-0.00044	0.00000	0.74058	0.74029	-0.00029	0.00000
Li ₂	$r(\text{LiLi})$	2.62504	2.62474	-0.00031	-0.00006	2.62465	2.62458	-0.00007	0.00009
N ₂	$r(\text{NN})$	1.08990	1.08994	0.00004	-0.00001	1.08993	1.08993	0.00000	0.00000
O ₂	$r(\text{OO})$	1.19868	1.19864	-0.00004	-0.00001	1.19862	1.19863	0.00000	0.00000
F ₂	$r(\text{FF})$	1.38170	1.38196	0.00026	0.00001	1.38190	1.38198	0.00008	-0.00001
Si ₂	$r(\text{SiSi})$	1.97313	1.97384	0.00071	-0.00001	1.97381	1.97381	0.00001	0.00002
P ₂	$r(\text{PP})$	1.88464	1.88563	-0.00001	-0.01618	1.86971	1.86958	-0.00013	-0.00013
S ₂	$r(\text{SS})$	1.88586	1.88563	-0.00023	0.00001	1.88586	1.88563	-0.00023	0.00001
Cl ₂	$r(\text{ClCl})$	1.99929	1.99982	0.00053	0.00006	1.99960	1.99987	0.00027	0.00001
SiH ₂	$r(\text{SiH})$	1.52015	1.52024	0.00009	-0.00006	1.52015	1.52015	0.00000	0.00003
	$\angle(\text{HSiH})$	91.27604	91.27937	0.00333	-0.00383	91.27460	91.27417	-0.00043	0.00137
CH ₃ OH	$r(\text{CH}_a)$	1.08689	1.08680	-0.00009	0.00002	1.08686	1.08681	-0.00005	0.00001
	$r(\text{CH}_b)$	1.09245	1.09247	0.00002	0.00002	1.09246	1.09248	0.00002	0.00001
	$r(\text{OH})$	0.95914	0.95913	-0.00001	-0.00001	0.95914	0.95912	-0.00002	0.00000
	$\angle(\text{OCH}_a)$	106.85461	106.83938	-0.01523	0.00152	106.84965	106.84074	-0.00891	0.00016
	$\angle(\text{COH})$	108.41613	108.42132	0.00519	-0.00057	108.41589	108.42065	0.00476	0.00010
	$\angle(\text{H}_b\text{CH}_b)$	109.01194	109.04461	0.03267	-0.00073	109.02502	109.04367	0.01865	0.00021
	$r(\text{CO})$	1.41980	1.41995	0.00015	0.00001	1.41987	1.41995	0.00008	0.00001
HCO	$r(\text{CO})$	1.17061	1.17059	-0.00002	0.00001	1.17060	1.17059	-0.00001	0.00001
	$r(\text{OH})$	1.12209	1.12223	0.00014	-0.00003	1.12211	1.12220	0.00009	0.00000
	$\angle(\text{OCH})$	124.24198	124.24365	0.00167	0.00348	124.24808	124.24689	-0.00119	0.00024
NH ₂ NH ₂	$r(\text{NN})$	1.42977	1.43023	0.00046	-0.00001	1.42991	1.43022	0.00031	0.00000
	$r(\text{NH}_a)$	1.00962	1.00968	0.00006	-0.00001	1.00965	1.00967	0.00002	0.00000
	$r(\text{NH}_b)$	1.01260	1.01266	0.00006	-0.00002	1.01263	1.01264	0.00001	0.00000
	$\angle(\text{NNH}_b)$	112.10811	112.02791	-0.08020	-0.00118	112.07649	112.02601	-0.05048	0.00072
	$\angle(\text{NNH}_a)$	107.80480	107.74268	-0.06212	0.00225	107.78490	107.74458	-0.04032	0.00035

^aBond lengths in Ångstroms and bond angles in degrees. ^bDifference between pruned and unpruned results. ^cDifference with respect to a (250,974) grid.

TABLE S5: Harmonic vibrational frequencies (ν , in cm^{-1}), and differences ($\Delta\nu$) engendered by pruning and grid errors, computed at the $\omega\text{B97M-V/aug-cc-pVTZ}$ level.

Molecule	(75,302) Grid			(99,590) Grid		
	ν	Error, $\Delta\nu$		ν	Error, $\Delta\nu$	
		pruning ^a	grid ^b		pruning ^a	grid ^b
BH ₃	1133.5	18.6	0.0	1151.5	0.6	0.0
	1202.5	-0.4	-0.4	1201.1	1.0	-0.4
	1203.0	-0.9	-0.4	1201.3	0.8	-0.4
	2564.9	18.0	2.2	2585.6	-2.8	2.3
	2697.6	10.4	2.3	2710.7	-2.8	2.3
	2697.8	10.2	2.3	2710.8	-2.9	2.3
Cl ₂	578.3	1.2	0.8	582.3	-2.9	0.9
CO ₂	679.5	-0.3	0.5	679.6	-0.4	0.4
	679.5	-0.3	0.5	679.6	-0.4	0.4
	1388.2	-0.6	0.3	1388.0	-0.4	0.4
	2411.5	-1.3	0.7	2411.0	-0.8	0.8
F ₂	1107.9	-1.1	-0.5	1107.3	-0.7	-0.4
NH ₂ NH ₂	435.0	-0.1	-2.3	433.3	1.5	-2.2
	824.1	1.8	-0.8	824.4	1.4	-0.7
	973.8	2.1	-0.7	974.4	1.4	-0.6
	1127.7	0.0	-0.1	1127.6	0.1	-0.2
	1301.1	-0.2	-0.3	1300.7	0.2	-0.2
	1328.5	0.1	0.0	1328.6	0.1	0.0
	1670.5	0.7	0.1	1670.8	0.4	0.1
	1684.4	0.3	-0.1	1684.4	0.3	-0.1
	3498.9	0.1	-1.1	3498.3	0.8	-1.2
	3507.0	0.3	-1.1	3506.5	1.0	-1.2
	3596.6	0.4	-1.5	3595.9	1.2	-1.6
	3601.0	0.4	-1.6	3600.3	1.3	-1.7
H ₂ O ₂	392.8	-7.2	4.6	394.6	-9.1	4.7
	1011.3	-0.2	0.1	1011.4	-0.3	0.1
	1345.8	-1.1	0.2	1345.8	-1.1	0.2
	1451.2	0.4	0.2	1451.6	0.1	0.1
	3798.9	-3.0	1.7	3797.9	-2.1	1.7
	3800.0	-3.1	1.7	3799.0	-2.1	1.7
H ₂	4451.2	1.5	0.0	4452.0	0.8	0.0
H ₂ S	1204.9	-2.0	1.5	1202.8	0.4	1.3
	2722.4	4.6	0.1	2726.9	0.2	0.0
	2734.4	5.1	-0.1	2739.1	-0.8	1.0
HCCH	691.4	0.8	0.6	690.5	1.7	0.6
	691.5	0.8	0.6	690.5	1.7	0.6
	770.5	1.0	0.6	769.5	2.1	0.6
	770.6	1.0	0.6	769.5	2.1	0.6
	2091.3	-0.3	0.3	2091.2	-0.2	0.2
	3419.3	1.5	-1.3	3418.7	2.1	-1.4
3530.1	1.3	-1.2	3529.5	1.9	-1.3	
N ₂	2484.8	0.1	-0.4	2484.6	0.4	-0.5
O ₂	1693.3	-0.5	0.6	1693.4	-0.5	0.5
PH ₂	1129.4	-3.4	-2.0	1123.4	2.6	-2.0
	2414.4	-3.2	2.8	2414.6	-4.3	3.7
	2421.1	-2.9	2.9	2421.5	-4.1	3.7
S ₂	759.2	0.9	-2.0	758.1	1.9	-1.9
SiH ₂	1034.0	-4.1	-5.3	1026.3	3.6	-5.3
	2074.4	-9.6	5.2	2070.1	-5.0	4.9
	2075.2	-10.1	5.7	2070.9	-5.4	5.3
MUD ^c		2.8	1.3		1.6	1.3

^aDifference between pruned and unpruned results.

^bDifference with respect to a (250,974) grid.

^cMean unsigned deviation.

TABLE S6: Harmonic vibrational frequencies (ν , in cm^{-1}), and differences ($\Delta\nu$) engendered by pruning and grid errors, at the $\omega\text{B97X-D/aug-cc-pVTZ}$ level, comparing analytic and finite-difference results.

Molecule	Analytic Hessian						Finite Difference of Analytic Gradients					
	(75,302) Grid			(99,590) Grid			(75,302) Grid			(99,590) Grid		
	ν	Error, $\Delta\nu$		ν	Error, $\Delta\nu$		ν	Error, $\Delta\nu$		ν	Error, $\Delta\nu$	
	pruning ^a	grid ^b		pruning ^a	grid ^b		pruning ^a	grid ^b		pruning ^a	grid ^b	
BH ₃	1151.18	-1.96	-0.91	1149.89	-1.58	0.38	1147.69	0.06	2.60	1150.38	-0.13	-0.09
	1204.32	-0.83	-0.08	1203.82	-0.88	0.42	1204.40	-0.14	0.01	1204.70	0.28	-0.29
	1204.43	-2.15	-0.01	1204.43	-1.13	-0.01	1204.47	-0.20	-0.06	1204.79	0.26	-0.38
	2544.31	-17.08	15.49	2552.43	-11.39	7.37	2544.18	-17.77	15.64	2553.41	-9.72	6.41
	2676.60	-17.59	16.00	2684.63	-11.88	7.97	2676.57	-18.54	16.14	2686.04	-10.14	6.67
	2676.81	-18.11	15.97	2685.71	-11.45	7.07	2676.62	-18.57	16.10	2686.15	-10.09	6.57
Cl ₂	586.22	3.27	0.16	587.82	-0.37	-1.44	586.22	3.27	0.16	587.82	-0.37	-1.44
CO ₂	686.22	-0.08	0.19	686.26	-0.34	0.15	686.22	-0.08	0.18	686.26	0.66	0.14
	686.23	-0.08	0.19	686.27	0.66	0.15	686.22	-0.08	0.19	686.26	0.66	0.15
	1395.62	-2.49	0.45	1396.57	0.76	-0.50	1395.62	-2.49	0.45	1396.57	0.75	-0.50
	2441.00	-4.40	0.37	2442.22	1.30	-0.85	2441.01	-4.40	0.37	2442.22	1.29	-0.84
F ₂	1112.02	17.52	-15.20	1091.40	-6.37	5.42	1112.01	17.51	-15.19	1091.40	-6.38	5.42
NH ₂ NH ₂	443.95	1.06	2.63	446.31	1.92	0.27	443.93	1.06	2.62	446.30	1.92	0.25
	819.20	2.30	2.05	820.91	3.01	0.34	819.21	2.29	2.30	820.92	3.01	0.59
	969.78	1.33	2.31	971.17	1.90	0.92	969.78	1.33	2.32	971.16	1.89	0.94
	1141.33	-0.17	0.46	1142.00	0.34	-0.21	1141.33	-0.17	0.73	1142.00	0.34	0.06
	1305.85	-0.39	2.41	1307.03	0.47	1.23	1305.83	-0.40	1.11	1307.01	0.47	-0.07
	1337.09	-3.83	-0.52	1336.79	-2.73	-0.22	1337.07	-3.83	-0.16	1336.78	-2.72	0.13
	1680.82	-1.34	0.40	1681.37	-0.99	-0.15	1680.84	-1.34	0.32	1681.38	-0.99	-0.22
	1694.00	0.13	1.52	1695.21	0.74	0.31	1694.00	0.12	0.98	1695.21	0.73	-0.23
	3508.30	-2.99	0.09	3508.37	-2.06	0.02	3508.30	-3.00	1.09	3508.37	-2.07	1.02
	3516.69	-3.47	0.35	3516.93	-1.90	0.11	3516.69	-3.45	1.28	3516.92	-1.90	1.05
	3611.58	-5.75	-0.02	3611.75	-2.55	-0.19	3611.59	-5.75	1.26	3611.75	-2.55	1.10
	3616.24	-6.27	0.27	3616.44	-2.56	0.07	3616.27	-6.27	1.31	3616.47	-2.57	1.11
	H ₂ O ₂	397.28	1.11	1.47	401.15	-0.35	-2.40	397.22	1.11	1.67	401.09	-0.35
1025.57		7.22	-8.08	1016.35	-0.83	1.14	1025.56	7.21	-8.08	1016.35	-0.82	1.13
1360.43		-0.27	0.65	1361.58	-0.74	-0.50	1360.36	-0.28	0.65	1361.51	-0.75	-0.50
1468.17		0.94	0.74	1469.30	0.33	-0.39	1468.26	0.95	0.73	1469.39	0.34	-0.40
3837.40		-3.14	2.53	3839.96	-0.29	-0.03	3837.42	-3.14	2.53	3839.98	-0.29	-0.03
3838.80		-3.23	2.52	3841.35	-0.32	-0.03	3838.79	-3.23	2.52	3841.34	-0.32	-0.03
H ₂	4438.49	-0.34	-0.01	4438.48	-0.51	0.00	4438.49	-0.35	0.00	4438.49	-0.50	0.00
H ₂ S	1219.51	4.26	-5.34	1211.79	2.29	2.38	1219.48	4.25	-5.33	1211.77	2.30	2.38
	2709.66	6.65	29.90	2748.10	16.01	-8.54	2709.62	6.59	29.94	2748.09	16.00	-8.53
	2721.14	5.33	31.53	2760.68	15.50	-8.01	2721.13	5.27	31.56	2760.70	15.49	-8.01
HCCH	690.45	4.36	3.06	693.77	6.25	-0.26	690.43	4.36	3.05	693.75	6.25	-0.27
	690.45	4.36	3.06	693.77	6.25	-0.26	690.46	4.37	3.06	693.79	6.27	-0.27
	777.15	4.97	2.86	780.14	6.51	-0.13	777.17	4.97	2.86	780.16	6.51	-0.13
	777.15	4.97	2.86	780.14	6.51	-0.13	777.18	4.97	2.87	780.17	6.51	-0.12
	2087.58	0.02	2.90	2089.14	-0.53	1.34	2087.59	0.02	2.89	2089.15	-0.53	1.33
	3418.57	-7.42	3.80	3421.24	-0.83	1.13	3418.59	-7.41	3.80	3421.27	-0.82	1.12
	3527.41	-6.77	3.78	3529.98	-0.88	1.21	3527.41	-6.76	3.78	3529.99	-0.87	1.20
	N ₂	2488.37	0.98	0.54	2489.63	1.53	-0.72	2488.37	0.97	0.55	2489.64	1.53
O ₂	1701.67	-1.71	2.06	1702.98	-0.26	0.75	1701.68	-1.71	2.05	1702.99	-0.25	0.74
PH ₂	1136.66	-2.40	-4.94	1131.47	-0.41	0.25	1136.64	-2.40	-4.94	1131.46	-0.40	0.24
	2388.99	-22.50	18.21	2388.87	-31.04	18.33	2388.99	-22.48	18.20	2388.87	-31.02	18.32
	2398.12	-22.94	18.89	2399.05	-30.21	17.96	2398.18	-22.92	18.87	2399.11	-30.18	17.94
S ₂	752.50	-3.50	-1.97	750.27	-2.09	0.26	752.50	-3.51	-1.96	750.27	-2.09	0.27
SiH ₂	1033.32	-6.18	-2.97	1032.60	0.52	-2.25	1033.31	-6.18	-2.97	1032.59	0.52	-2.25
	2061.09	-37.16	15.16	2099.31	31.51	-23.06	2061.10	-37.15	15.16	2099.31	31.50	-23.05
	2063.95	-35.04	13.44	2099.84	30.57	-22.45	2063.96	-35.03	13.43	2099.84	30.56	-22.45
MUD ^c		6.25	5.22		5.25	2.99		6.19	5.32		5.10	2.99

^aDifference between pruned and unpruned results. ^bDifference with respect to a (250,974) grid. ^cMean unsigned deviation.

TABLE S7: Interaction energies (in kcal/mol) for the S66 data set computed at the M06-2X/aug-cc-pVTZ level.

Dimer	Energy		Error		Energy		Error	
	SG-2	(75,302)	pruning ^a	grid ^b	SG-3	(99,590)	pruning ^a	grid ^b
AcNH ₂ -AcNH ₂	-16.31	-16.18	-0.13	-0.01	-16.20	-16.19	-0.01	0.00
AcNH ₂ -Uracil	-19.33	-19.07	-0.26	-0.03	-19.12	-19.10	-0.02	0.00
AcOH-AcOH	-19.98	-19.89	-0.09	0.01	-19.88	-19.88	0.00	0.00
AcOH-Uracil	-19.79	-19.72	-0.07	-0.01	-19.71	-19.73	0.02	0.00
Benzene-AcNH ₂ (NH- π)	-4.33	-4.48	0.15	0.00	-4.33	-4.48	0.15	0.00
Benzene-AcOH (OH- π)	-5.28	-5.11	-0.17	-0.02	-5.10	-5.13	0.03	0.00
Benzene-AcOH	-4.09	-4.11	0.02	0.01	-3.95	-4.09	0.14	-0.01
Benzene-Benzene (π -stacked)	-2.65	-2.51	-0.14	-0.04	-2.48	-2.54	0.06	-0.01
Benzene-Benzene (T-shaped)	-2.75	-2.56	-0.19	-0.03	-2.59	-2.59	0.00	0.00
Benzene-Cyclopentane	-3.69	-3.52	-0.17	-0.03	-3.51	-3.55	0.04	0.00
Benzene-Ethene	-1.27	-1.44	0.17	0.00	-1.43	-1.45	0.02	0.01
Benzene-Ethyne (CH- π)	-3.07	-3.03	-0.04	0.01	-3.03	-3.02	-0.01	0.00
Benzene-MeNH ₂ (NH- π)	-3.40	-3.46	0.06	0.02	-3.42	-3.43	0.01	-0.01
Benzene-MeOH (OH- π)	-4.77	-4.60	-0.17	0.00	-4.60	-4.61	0.01	0.01
Benzene-Neopentane	-2.73	-2.68	-0.05	-0.02	-2.68	-2.67	-0.01	-0.03
Benzene-Peptide (NH- π)	-5.55	-5.43	-0.12	0.00	-5.43	-5.45	0.02	0.02
Benzene-Pyridine (π -stacked)	-3.21	-3.15	-0.06	-0.09	-3.14	-3.21	0.07	-0.03
Benzene-Pyridine (T-shaped)	-3.08	-3.07	-0.01	-0.04	-3.04	-3.11	0.07	0.00
Benzene-Uracil (π -stacked)	-6.28	-5.97	-0.31	-0.02	-5.94	-5.98	0.04	-0.01
Cyclopentane-Cyclopentane	-2.43	-2.42	-0.01	0.01	-2.39	-2.42	0.03	0.01
Cyclopentane-Neopentane	-1.98	-2.03	0.05	0.02	-2.02	-2.00	-0.02	-0.01
Ethene-Pentane	-1.69	-1.95	0.26	-0.03	-1.86	-1.98	0.12	0.00
Ethyne-AcOH (OH- π)	-5.53	-5.45	-0.08	0.00	-5.48	-5.45	-0.03	0.00
Ethyne-H ₂ O (CH-O)	-2.94	-2.92	-0.02	0.02	-2.90	-2.90	0.00	0.00
Ethyne-Pentane	-1.75	-1.68	-0.07	-0.03	-1.71	-1.71	0.00	0.00
H ₂ O-H ₂ O	-5.19	-5.15	-0.04	0.02	-5.16	-5.13	-0.03	0.00
H ₂ O-MeNH ₂	-7.11	-6.93	-0.18	-0.01	-7.05	-6.94	-0.11	0.00
H ₂ O-MeOH	-5.66	-5.73	0.07	0.00	-5.66	-5.73	0.07	0.00
H ₂ O-Peptide	-8.89	-8.63	-0.26	0.03	-8.71	-8.60	-0.11	0.00
H ₂ O-Pyridine	-6.83	-6.69	-0.14	0.00	-6.75	-6.69	-0.06	0.00
MeNH ₂ -H ₂ O	-7.50	-7.51	0.01	0.00	-7.51	-7.51	0.00	0.00
MeNH ₂ -MeNH ₂	-4.11	-4.12	0.01	0.00	-4.18	-4.12	-0.06	0.00
MeNH ₂ -MeOH	-2.69	-2.79	0.10	0.00	-2.75	-2.79	0.04	0.00
MeNH ₂ -Peptide	-5.55	-5.57	0.02	0.00	-5.58	-5.57	-0.01	0.00
MeNH ₂ -Pyridine	-3.83	-3.88	0.05	-0.03	-3.87	-3.91	0.04	0.00
MeOH-H ₂ O	-5.11	-5.10	-0.01	0.00	-5.11	-5.10	-0.01	0.00
MeOH-MeNH ₂	-7.45	-7.50	0.05	0.00	-7.49	-7.50	0.01	0.00
MeOH-MeOH	-5.62	-5.70	0.08	0.00	-5.63	-5.70	0.07	0.00
MeOH-Peptide	-8.51	-8.45	-0.06	-0.01	-8.47	-8.46	-0.01	0.00
MeOH-Pyridine	-7.21	-7.02	-0.19	-0.03	-7.06	-7.05	-0.01	0.00
Neopentane-Neopentane	-1.26	-1.26	0.00	0.03	-1.23	-1.23	0.00	0.00
Neopentane-Pentane	-2.06	-2.12	0.06	-0.09	-2.23	-2.21	-0.02	0.00
Pentane-AcNH ₂	-3.53	-3.52	-0.01	-0.06	-3.56	-3.58	0.02	0.00
Pentane-AcOH	-2.27	-2.86	0.59	-0.03	-2.38	-2.89	0.51	0.00
Pentane-Pentane	-3.55	-3.56	0.01	-0.07	-3.63	-3.63	0.00	0.00
Peptide-Ethene	-2.86	-3.00	0.14	-0.03	-3.02	-3.03	0.01	0.00
Peptide-H ₂ O	-5.04	-5.08	0.04	-0.01	-5.10	-5.09	-0.01	0.00
Peptide-MeNH ₂	-7.25	-7.23	-0.02	0.00	-7.25	-7.23	-0.02	0.00
Peptide-MeOH	-6.08	-6.11	0.03	-0.01	-6.08	-6.12	0.04	0.00
Peptide-Pentane	-4.05	-4.44	0.39	-0.04	-4.18	-4.48	0.30	0.00
Peptide-Peptide	-8.49	-8.55	0.06	0.01	-8.45	-8.53	0.08	-0.01
Pyridine-Ethene	-1.92	-1.85	-0.07	-0.06	-1.91	-1.91	0.00	0.00
Pyridine-Ethyne	-3.83	-3.74	-0.09	0.00	-3.76	-3.75	-0.01	0.01
Pyridine-Pyridine (CH-N)	-3.30	-3.60	0.30	0.03	-3.50	-3.57	0.07	0.00
Pyridine-Pyridine (π -stacked)	-3.47	-3.64	0.17	-0.01	-3.52	-3.66	0.14	0.01
Pyridine-Pyridine (T-shaped)	-3.38	-3.24	-0.14	0.00	-3.16	-3.24	0.08	0.00
Pyridine-Uracil (π -stacked)	-7.41	-7.02	-0.39	0.01	-6.89	-7.01	0.12	0.00
Uracil-Cyclopentane	-4.38	-3.96	-0.42	-0.01	-3.94	-3.96	0.02	-0.01
Uracil-Ethene	-3.87	-3.61	-0.26	-0.03	-3.53	-3.64	0.11	0.00
Uracil-Ethyne	-4.08	-4.05	-0.03	-0.01	-3.99	-4.07	0.08	0.01
Uracil-Uracil (Watson-Crick)	-4.47	-3.60	-0.87	0.02	-3.87	-3.58	-0.29	0.00
Uracil-Pentane	-4.95	-4.91	-0.04	-0.08	-4.89	-5.00	0.11	0.01
Uracil-Uracil (Watson-Crick)	-17.03	-16.85	-0.18	-0.01	-16.86	-16.86	0.00	0.00
Uracil-Uracil (π -stacked)	-10.98	-10.34	-0.64	-0.02	-10.32	-10.36	0.04	0.00
MUD ^c			0.11	0.02			0.03	0.00

^aDifference between pruned and unpruned results. ^bDifference with respect to a (250,974) grid. ^cMean unsigned deviation.

TABLE S8: Interaction energies (in kcal/mol) for the S66 data set computed at the ω B97X-V/aug-cc-pVTZ level.

Dimer	Energy		Error		Energy		Error	
	SG-2	(75,302)	pruning ^a	grid ^b	SG-3	(99,590)	pruning ^a	grid ^b
AcNH ₂ -AcNH ₂	-16.31	-16.33	0.02	0.00	-16.33	-16.33	0.00	0.00
AcNH ₂ -Uracil	-19.28	-19.34	0.06	-0.01	-19.35	-19.34	-0.01	0.01
AcOH-AcOH	-20.10	-20.03	-0.07	0.01	-20.02	-20.03	0.01	-0.01
AcOH-Uracil	-19.89	-19.98	0.09	0.00	-19.98	-19.98	0.00	0.00
Benzene-AcNH ₂ (NH- π)	-4.45	-4.43	-0.02	0.00	-4.43	-4.40	-0.03	0.03
Benzene-AcOH (OH- π)	-4.98	-4.97	-0.01	0.00	-4.97	-4.95	-0.02	0.02
Benzene-AcOH	-4.06	-4.01	-0.05	0.00	-4.01	-3.98	-0.03	0.03
Benzene-Benzene (π -stacked)	-2.97	-2.92	-0.05	-0.01	-2.93	-2.90	-0.03	0.03
Benzene-Benzene (T-shaped)	-2.96	-2.89	-0.07	0.00	-2.89	-2.89	0.00	0.00
Benzene-Cyclopentane	-3.74	-3.74	0.00	-0.01	-3.75	-3.74	-0.01	0.01
Benzene-Ethene	-1.41	-1.44	0.03	0.00	-1.44	-1.44	0.00	0.00
Benzene-Ethyne (CH- π)	-3.04	-3.02	-0.02	0.00	-3.02	-3.02	0.00	0.00
Benzene-MeNH ₂ (NH- π)	-3.30	-3.30	0.00	0.00	-3.30	-3.29	-0.01	0.01
Benzene-MeOH (OH- π)	-4.35	-4.31	-0.04	-0.01	-4.32	-4.32	0.00	0.00
Benzene-Neopentane	-3.16	-3.15	-0.01	0.01	-3.14	-3.13	-0.01	0.01
Benzene-Peptide (NH- π)	-5.48	-5.45	-0.03	-0.01	-5.46	-5.45	-0.01	0.01
Benzene-Pyridine (π -stacked)	-3.51	-3.49	-0.02	-0.01	-3.50	-3.48	-0.02	0.02
Benzene-Pyridine (T-shaped)	-3.41	-3.37	-0.04	0.00	-3.37	-3.37	0.00	0.00
Benzene-Uracil (π -stacked)	-5.76	-5.83	0.07	0.00	-5.82	-5.83	0.01	-0.01
Cyclopentane-Cyclopentane	-3.08	-3.11	0.03	-0.01	-3.12	-3.09	-0.03	0.03
Cyclopentane-Neopentane	-2.62	-2.62	0.00	0.01	-2.61	-2.62	0.01	-0.01
Ethene-Pentane	-2.03	-2.03	0.00	0.00	-2.03	-2.03	0.00	0.00
Ethyne-AcOH (OH- π)	-5.20	-5.20	0.00	0.01	-5.19	-5.19	0.00	0.00
Ethyne-H ₂ O (CH-O)	-2.98	-2.99	0.01	0.01	-2.98	-2.98	0.00	0.00
Ethyne-Pentane	-1.85	-1.84	-0.01	0.00	-1.84	-1.84	0.00	0.00
H ₂ O-H ₂ O	-5.01	-5.00	-0.01	0.01	-4.99	-5.00	0.01	-0.01
H ₂ O-MeNH ₂	-7.10	-7.08	-0.02	-0.01	-7.09	-7.10	0.01	-0.01
H ₂ O-MeOH	-5.60	-5.61	0.01	0.00	-5.61	-5.60	-0.01	0.01
H ₂ O-Peptide	-8.29	-8.29	0.00	0.01	-8.28	-8.28	0.00	0.00
H ₂ O-Pyridine	-7.04	-7.01	-0.03	0.00	-7.01	-7.02	0.01	-0.01
MeNH ₂ -H ₂ O	-7.38	-7.39	0.01	0.00	-7.39	-7.39	0.00	0.00
MeNH ₂ -MeNH ₂	-4.01	-4.07	0.06	0.00	-4.07	-4.06	-0.01	0.01
MeNH ₂ -MeOH	-3.03	-3.04	0.01	0.00	-3.04	-3.04	0.00	0.00
MeNH ₂ -Peptide	-5.34	-5.41	0.07	0.00	-5.41	-5.39	-0.02	0.02
MeNH ₂ -Pyridine	-3.83	-3.84	0.01	-0.01	-3.85	-3.83	-0.02	0.02
MeOH-H ₂ O	-5.06	-5.06	0.00	0.00	-5.06	-5.06	0.00	0.00
MeOH-MeNH ₂	-7.69	-7.67	-0.02	0.00	-7.67	-7.68	0.01	-0.01
MeOH-MeOH	-5.73	-5.75	0.02	0.00	-5.75	-5.73	-0.02	0.02
MeOH-Peptide	-8.37	-8.38	0.01	0.00	-8.38	-8.37	-0.01	0.01
MeOH-Pyridine	-7.55	-7.50	-0.05	0.00	-7.50	-7.51	0.01	-0.01
Neopentane-Neopentane	-2.00	-2.01	0.01	0.01	-2.00	-2.00	0.00	0.00
Neopentane-Pentane	-2.82	-2.82	0.00	-0.01	-2.83	-2.85	0.02	-0.02
Pentane-AcNH ₂	-3.68	-3.69	0.01	-0.01	-3.70	-3.70	0.00	0.00
Pentane-AcOH	-3.10	-3.15	0.05	0.01	-3.14	-3.09	-0.05	0.05
Pentane-Pentane	-4.01	-4.01	0.00	-0.01	-4.02	-4.03	0.01	-0.01
Peptide-Ethene	-3.01	-3.03	0.02	0.00	-3.03	-3.03	0.00	0.00
Peptide-H ₂ O	-5.13	-5.14	0.01	0.01	-5.13	-5.14	0.01	-0.01
Peptide-MeNH ₂	-7.43	-7.45	0.02	0.00	-7.44	-7.45	0.01	-0.01
Peptide-MeOH	-6.12	-6.16	0.04	0.01	-6.15	-6.14	-0.01	0.01
Peptide-Pentane	-4.43	-4.46	0.03	-0.01	-4.47	-4.44	-0.03	0.03
Peptide-Peptide	-8.64	-8.72	0.08	0.01	-8.71	-8.67	-0.04	0.04
Pyridine-Ethene	-1.85	-1.83	-0.02	-0.01	-1.84	-1.84	0.00	0.00
Pyridine-Ethyne	-4.19	-4.17	-0.02	0.00	-4.17	-4.17	0.00	0.00
Pyridine-Pyridine (CH-N)	-3.91	-3.97	0.06	0.01	-3.96	-3.96	0.00	0.00
Pyridine-Pyridine (π -stacked)	-3.93	-3.92	-0.01	0.00	-3.92	-3.91	-0.01	0.01
Pyridine-Pyridine (T-shaped)	-3.49	-3.47	-0.02	0.01	-3.46	-3.45	-0.01	0.01
Pyridine-Uracil (π -stacked)	-6.93	-6.94	0.01	0.00	-6.94	-6.91	-0.03	0.03
Uracil-Cyclopentane	-4.32	-4.33	0.01	0.00	-4.32	-4.33	0.01	-0.01
Uracil-Ethene	-3.46	-3.45	-0.01	0.00	-3.46	-3.44	-0.02	0.02
Uracil-Ethyne	-3.83	-3.86	0.03	0.00	-3.86	-3.84	-0.02	0.02
Uracil-Uracil (Watson-Crick)	-17.30	-17.36	0.06	0.00	-17.36	-17.35	-0.01	0.01
MUD			0.03	0.01			0.01	0.01



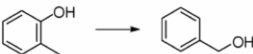
^aDifference between pruned and unpruned results. ^bDifference with respect to a (250,974) grid. ^cMean unsigned deviation.

TABLE S9: Interaction energies (in kcal/mol) for the S66 data set computed at the ω B97M-V/aug-cc-pVTZ level.

Dimer	Energy		Error		Energy		Error	
	SG-2	(75,302)	pruning ^a	grid ^b	SG-3	(99,590)	pruning ^a	grid ^b
AcNH ₂ -AcNH ₂	-16.48	-16.42	0.06	0.00	-16.46	-16.42	0.04	0.00
AcNH ₂ -Uracil	-19.61	-19.44	0.17	-0.01	-19.51	-19.45	0.06	0.00
AcOH-AcOH	-19.99	-19.95	0.03	0.01	-19.93	-19.95	-0.01	0.00
AcOH-Uracil	-20.11	-20.03	0.07	0.00	-20.03	-20.03	0.00	0.00
Benzene-AcNH ₂ (NH- π)	-4.45	-4.50	-0.05	0.00	-4.53	-4.51	0.02	0.00
Benzene-AcOH (OH- π)	-5.08	-5.09	0.00	0.00	-5.13	-5.09	0.05	0.00
Benzene-AcOH	-4.15	-4.17	-0.02	0.00	-4.19	-4.17	0.02	0.00
Benzene-Benzene (π -stacked)	-3.09	-3.16	-0.07	0.00	-3.24	-3.16	0.08	0.00
Benzene-Benzene (T-shaped)	-3.01	-3.00	0.01	0.00	-3.03	-3.01	0.02	0.00
Benzene-Cyclopentane	-3.90	-3.91	-0.01	-0.02	-3.94	-3.93	0.02	0.00
Benzene-Ethene	-1.47	-1.59	-0.12	-0.01	-1.63	-1.59	0.04	0.00
Benzene-Ethyne (CH- π)	-3.06	-3.09	-0.03	0.01	-3.09	-3.08	0.01	0.00
Benzene-MeNH ₂ (NH- π)	-3.27	-3.43	-0.16	0.01	-3.40	-3.42	-0.03	0.00
Benzene-MeOH (OH- π)	-4.37	-4.44	-0.07	-0.01	-4.44	-4.45	-0.01	0.00
Benzene-Neopentane	-3.18	-3.24	-0.06	0.00	-3.26	-3.24	0.02	0.00
Benzene-Peptide (NH- π)	-5.65	-5.63	0.03	0.00	-5.65	-5.63	0.02	0.00
Benzene-Pyridine (π -stacked)	-3.72	-3.71	0.01	-0.03	-3.82	-3.74	0.08	0.00
Benzene-Pyridine (T-shaped)	-3.50	-3.47	0.03	-0.01	-3.47	-3.48	-0.01	0.00
Benzene-Uracil (π -stacked)	-6.22	-6.22	0.01	0.00	-6.21	-6.21	0.00	0.00
Cyclopentane-Cyclopentane	-3.20	-3.12	0.08	-0.02	-3.16	-3.14	0.02	0.00
Cyclopentane-Neopentane	-2.57	-2.61	-0.04	0.01	-2.61	-2.60	0.01	0.00
Ethene-Pentane	-1.99	-2.01	-0.02	0.00	-2.03	-2.01	0.02	-0.01
Ethyne-AcOH (OH- π)	-5.31	-5.26	0.05	0.00	-5.32	-5.26	0.06	0.00
Ethyne-H ₂ O (CH-O)	-2.98	-2.97	0.01	0.00	-2.97	-2.97	0.00	0.00
Ethyne-Pentane	-1.83	-1.83	0.00	0.00	-1.83	-1.83	0.00	0.00
H ₂ O-H ₂ O	-4.94	-4.99	-0.05	0.00	-4.96	-4.99	-0.04	0.00
H ₂ O-MeNH ₂	-6.91	-6.99	-0.08	0.00	-6.94	-7.00	-0.06	0.00
H ₂ O-MeOH	-5.60	-5.60	0.00	0.00	-5.60	-5.60	0.00	0.00
H ₂ O-Peptide	-8.34	-8.27	0.07	0.00	-8.29	-8.27	0.02	0.00
H ₂ O-Pyridine	-6.88	-6.91	-0.03	0.00	-6.88	-6.91	-0.04	0.00
MeNH ₂ -H ₂ O	-7.30	-7.31	-0.02	0.00	-7.30	-7.31	-0.01	0.00
MeNH ₂ -MeNH ₂	-4.05	-4.11	-0.05	0.00	-4.09	-4.10	-0.02	0.00
MeNH ₂ -MeOH	-2.97	-3.01	-0.04	0.00	-2.99	-3.01	-0.03	0.00
MeNH ₂ -Peptide	-5.54	-5.53	0.01	0.00	-5.54	-5.53	0.01	0.00
MeNH ₂ -Pyridine	-3.94	-3.96	-0.01	-0.01	-3.97	-3.97	0.00	0.01
MeOH-H ₂ O	-5.05	-5.08	-0.03	0.00	-5.06	-5.08	-0.02	0.00
MeOH-MeNH ₂	-7.52	-7.58	-0.07	0.00	-7.54	-7.58	-0.04	0.00
MeOH-MeOH	-5.77	-5.77	0.00	0.00	-5.76	-5.77	0.00	0.00
MeOH-Peptide	-8.47	-8.43	0.04	0.00	-8.44	-8.44	0.00	0.00
MeOH-Pyridine	-7.46	-7.43	0.02	-0.01	-7.43	-7.44	-0.01	0.00
Neopentane-Neopentane	-1.93	-1.93	0.00	0.01	-5.38	-1.92	0.00	0.00
Neopentane-Pentane	-2.74	-2.79	-0.06	-0.02	-2.81	-2.81	0.00	0.00
Pentane-AcNH ₂	-3.72	-3.72	0.00	0.00	-3.71	-3.72	-0.01	0.00
Pentane-AcOH	-3.22	-3.11	0.12	0.00	-3.12	-3.11	0.01	0.00
Pentane-Pentane	-3.99	-4.03	-0.03	0.01	-4.01	-4.02	-0.01	0.00
Peptide-Ethene	-2.98	-3.06	-0.08	0.00	-3.04	-3.06	-0.02	0.00
Peptide-H ₂ O	-5.20	-5.15	0.05	0.00	-5.15	-5.15	0.01	0.00
Peptide-MeNH ₂	-7.45	-7.46	-0.01	0.00	-7.46	-7.46	0.00	0.00
Peptide-MeOH	-6.24	-6.21	0.03	0.00	-6.22	-6.20	0.02	0.00
Peptide-Pentane	-4.52	-4.50	0.02	-0.01	-4.50	-4.52	-0.02	0.00
Peptide-Peptide	-8.96	-8.78	0.18	0.01	-8.86	-8.78	0.08	0.00
Pyridine-Ethene	-1.95	-1.98	-0.04	-0.01	-2.00	-1.99	0.01	0.00
Pyridine-Ethyne	-4.20	-4.17	0.03	0.00	-4.18	-4.17	0.01	0.00
Pyridine-Pyridine (CH-N)	-3.90	-3.95	-0.04	0.01	-3.87	-3.93	-0.06	0.00
Pyridine-Pyridine (π -stacked)	-4.14	-4.15	-0.01	0.00	-4.17	-4.15	0.02	0.00
Pyridine-Pyridine (T-shaped)	-3.60	-3.57	0.03	0.01	-3.56	-3.55	0.00	0.00
Pyridine-Uracil (π -stacked)	-7.35	-7.28	0.07	0.00	-7.30	-7.28	0.02	0.00
Uracil-Cyclopentane	-4.56	-4.45	0.11	-0.01	-4.52	-4.46	0.06	0.00
Uracil-Ethene	-3.71	-3.61	0.10	-0.01	-3.73	-3.62	0.01	0.00
Uracil-Ethyne	-4.10	-4.03	0.07	0.00	-4.01	-4.03	-0.02	0.00
Uracil-Uracil (Watson-Crick)	-4.09	-4.07	0.02	0.00	-4.73	-4.07	0.01	0.00
Uracil-Pentane	-5.22	-5.21	0.01	-0.04	-5.63	-5.25	-0.02	0.00
Uracil-Uracil (Watson-Crick)	-17.58	-17.50	0.08	-0.02	-17.52	-17.51	0.01	0.00
Uracil-Uracil (π -stacked)	-10.85	-10.62	0.23	-0.03	-10.69	-10.65	0.04	0.00
MUD			0.05	0.01			0.02	0.00

^aDifference between pruned and unpruned results. ^bDifference with respect to a (250,974) grid. ^cMean unsigned deviation.

TABLE S10: Test of rotational invariance for organic isomerization reactions at the ω B97X-V/aug-cc-pVTZ level. Isomerization energies are reported for four different rotation angles θ about each of three different axes.

Reaction	Rotation Axis	Grid	$\Delta E / \text{kcal mol}^{-1}$			
			$\theta = 0^\circ$	$\theta = 30^\circ$	$\theta = 45^\circ$	$\theta = 60^\circ$
	<i>x</i>	SG-2	19.87	19.87	19.87	19.87
		(75,302)	19.88	19.88	19.88	19.88
		SG-3	19.88	19.88	19.88	19.88
		(99,590)	19.88	19.88	19.88	19.88
	<i>y</i>	SG-2	19.87	19.87	19.87	19.87
		(75,302)	19.88	19.88	19.88	19.88
		SG-3	19.88	19.88	19.88	19.88
		(99,590)	19.88	19.88	19.88	19.88
	<i>z</i>	SG-2	19.87	19.87	19.87	19.87
		(75,302)	19.88	19.88	19.88	19.88
		SG-3	19.88	19.88	19.88	19.88
		(99,590)	19.88	19.88	19.88	19.88
	<i>x</i>	SG-2	7.68	7.69	7.69	7.69
		(75,302)	7.68	7.69	7.69	7.69
		SG-3	7.68	7.69	7.69	7.69
		(99,590)	7.68	7.68	7.68	7.68
	<i>y</i>	SG-2	7.68	7.69	7.69	7.69
		(75,302)	7.68	7.69	7.69	7.69
		SG-3	7.68	7.69	7.69	7.69
		(99,590)	7.68	7.68	7.68	7.68
	<i>z</i>	SG-2	7.68	7.69	7.69	7.69
		(75,302)	7.68	7.69	7.69	7.69
		SG-3	7.68	7.69	7.69	7.69
		(99,590)	7.68	7.68	7.68	7.68
	<i>x</i>	SG-2	7.22	7.22	7.22	7.22
		(75,302)	7.21	7.21	7.21	7.21
		SG-3	7.22	7.22	7.22	7.22
		(99,590)	7.21	7.21	7.21	7.21
	<i>y</i>	SG-2	7.22	7.22	7.22	7.22
		(75,302)	7.21	7.21	7.21	7.21
		SG-3	7.22	7.22	7.22	7.22
		(99,590)	7.21	7.21	7.21	7.21
	<i>z</i>	SG-2	7.22	7.22	7.22	7.22
		(75,302)	7.21	7.21	7.21	7.21
		SG-3	7.22	7.22	7.22	7.22
		(99,590)	7.21	7.21	7.21	7.21

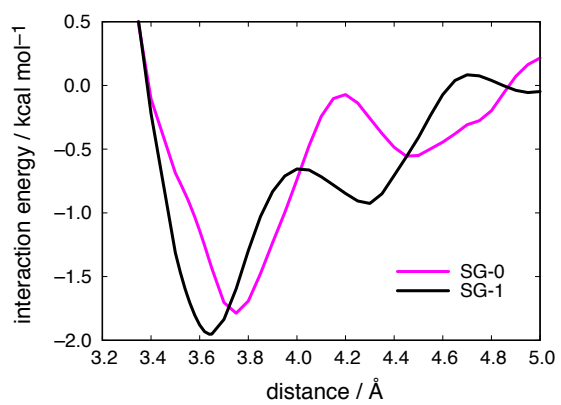


FIG. S1: Potential energy curves for the “sandwich” isomer of $(C_6H_6)_2$ along the face-to-face distance coordinate, computed at the M06-2X/6-311++G(3df,3pd) level with low-quality integration grids.