

Supplementary Information for

DSD-PBEP86: In search of the best double-hybrid DFT
with spin-component scaled MP2 and dispersion corrections

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Theoretical Method

For the W4-08 and DBH24 sets we used the aug'-pc3 basis set of Jensen,¹⁻⁵ with diffuse functions on non-hydrogen atoms and adding a high exponent d function for the W4-08 in order to properly account for high oxidation states. For the rest we used the Weigend and Ahlrichs Def2 bases:^{6,7} Def2-TZVP for S22 and Grubbs, as the size of the molecules made it arduous to climb to a quadruple- ζ type of functions. Following past practice, we added half the counterpoise correction in the S22 set to correct for the basis set superposition error, exploiting the well-known finding that uncorrected weak interaction energies converge from above, and corrected ones from below, in function of the basis set^{8,9}); Def2-QZVP was applied to MB, and Def2-QZVPP to the Pd set.

The Def2-QZVP basis set was used throughout the S130 set for the D3BJ parametrization. In addition, we applied one-half the counterpoise correction for the S22 and RG6 sets. The W4-11 set was calculated with aug-pc3+d; HFREQ27 with aug-pc2+2d and aug-pc3+d.

For the validation sets, we used the aug'-pc3+d basis set for NHTBH38, HTBH38, H-Bonds and vdW. For Pericyclic the aug'-pc2+2d basis set was used, and for Alkanes pc3.

The MP2 energies were calculated with the default "frozen core" correlation calculation, with only the valence electrons in the active space. For the transition metals, also the n-1 s and p atomic orbitals were included in the active space.

All the DH calculations including D2 dispersion corrections were carried out using an in-house modified Gaussian09 version,¹⁰ which includes the dispersion correction. D3BJ dispersion corrections were calculated with the DFTD3 software.¹¹

LDA Alternatives

We briefly considered LDA correlation parametrizations other than VWN5. Perdew-Zunger,¹² while yielding slightly different total energies, yields essentially indistinguishable reaction energies, and it can reasonably be assumed that the same will hold true for the Perdew-Wang parametrization¹³ (not implemented in Gaussian). For the VWN3 parametrization we do find a degradation in performance: unlike the others which are fitted to quantum Monte Carlo simulations of the uniform electron gas,¹⁴ VWN3 is fitted to RPA (random phase approximation¹⁵) data for the same: RPA can be regarded as a crude approximation to CCSD.¹⁶

Examples of DSD-PBEP86 input for Gaussian09, Gaussian 03, Molpro, ORCA, and Q-CHEM.

- *Gaussian09* [10]

```
#p b2plyp/cc-pVTZ
iop(3/125=0250005300,3/78=0430004300,3/76=0300007000,3/74=1004)
```

Example G09 input

```
0 1
o
h 1 r
h 1 r 2 th
```

```
r=0.957
th=104.5
```

IOP(3/74) selects the functional (1004 corresponds to PBE-P86), overriding the default b-lyp; IOP(3/125) chooses c_S and c_O (0250005300 means 25% same spin and 53 % opposite spin MP2); IOP(3/78) gives the DFT correlation (in this case 43% P86); and IOP(3/76) the amount of DFT and exact exchange (30% PBE and 70% HF for this example).

- *Gaussian03* [17]

```
#P pbep86/cc-pVTZ int(grid=fine)
IOP(3/74=1004,3/78=0430004300,3/76=0300007000) extraoverlay
```

```
8/10=4/1;9/16=-3/6;
```

Example G03 input

```
0 1
o
h 1 r
h 1 r 2 th
```

```
r=0.957
th=104.5
```

Total energy has to be extracted as SCF energy + 0.25*(E2alpha-alpha + E2beta-beta) + 0.53*E2alpha-beta.

- *Molpro [18]*

```
gthresh,energy=1d-9,grid=1d-6
geom={
o;
h,1,r;
h,1,r,2,th
}
r=0.957 ang
th=104.5 deg
basis=vtz
percenthf=70.0
percentDFTc=43.0
percentPTab=53.0
percentPTss=25.0
{rks,pbex,p86
exchange,percenthf*0.01
dftfactor,1.00-percenthf*0.01,percentDFTc*0.01}
ekohnsham=energy
mp2
eab=EMP2_SING+EMP2_TRIP/3d0
ess=2d0*EMP2_TRIP/3d0
ecorrmp2val=(eab*percentPTab+ess*percentPTss)*0.01
edsdpbep86val=ekohnsham+ecorrmp2val
```

- *Orca [19]*

```
* xyz 0 1
C 0.0 0.0 0.0
O 0.0 0.0 1.1314
*
! DFT aug-cc-pVTZ
%method
Exchange x_PBE
Correlation c_P86
LDAOpt C_VWN5
ScalHFX 0.70
ScalDFX 0.30
ScalGGAC 0.43
ScalLDAC 0.43
ScalMP2C 1.00
FrozenCore FC_ELECTRONS
end
%mp2
DoSCS True
Ps 0.53
Pt 0.25
end
```

- Q-CHEM [20]

```
$comment
DSD-PBEP86
$end
$molecule
O 1
O
H 1 R
H 1 R 2 TH
R=0.957
TH=104.5
$end
$rem
BASIS = aug-cc-pVTZ
CORRELATION = MP2
DH = 1
DH_OS = 510000
DH_SS = 230000
EXCHANGE = general
GUI = 0
JOB_TYPE = SP
SCF_FINAL_PRINT = 1
$end
$xc_functional
X HF 0.68
X PBE 0.32
C P86 0.45
$end
```

Optionally one might add

```
$empirical_dispersion
S6 0.29
$end
```

in this case the following line should be added to the \$rem block

```
DFT_D = EMPIRICAL_GRIMME
```

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Table 1: List of abbreviations.

Abbreviation	Significance
Methods:	
DH	Double Hybrid
DSD	Dispersion corrected, SCS, DH
DOD	Dispersion corrected, SOS, DH
SCS	Spin Component Scaled
SOS	Scaled Opposite Spin
D2	2 nd generation dispersion correction
D3BJ	3 rd generation dispersion correction, with Becke-Johnson damping function.
Coefficients	
c_0	Opposite spin MP2 coefficient.
c_s	Same spin MP2 coefficient.
c_x	DFT exchange coefficient.
c_c	DFT correlation coefficient.
s_6	Linear scaling of the r^{-6} component in D2 or D3BJ.
s_8	Linear scaling of the r^{-8} component in D3BJ.
a_1, a_2	D3BJ damping function parameters.
Training sets	
W4-08	Total atomization energies
DBH24	Main group reaction barriers
S22	van der Waals and H-bonded systems
Pd	Oxidative addition reactions on a Pd atom
Grubbs	Metathesis reactions on Ru complexes
MB	Mindless Benchmark (main group random thermochemistry).
Training sets for D3BJ optimization (S130)	
RG6	Rare gases dimers.
ACONF	N-alkane conformers.
ADIM6	Alkane dimers from ethane to n-heptane.
S22	van der Waals and H-bonded systems.
CYCONF	Cysteine conformers.
SCONF	Sugar conformers.
PCONF	Small peptide conformers.
S22+	Extended S22 set (at two stretched geometries).
Validation sets	
NHTBH38	Non Hydrogen Transfer Barrier Heights
HTBH38	Hydrogen Transfer Barrier Heights
H-Bonds	H-bonded dimers
vdW	van der Waals dimers
Pericyclic	Pericyclic reactions.
Alkanes	Alkanes thermochemistry and isomerization.
HFREQ27	Harmonic frequencies of small molecules.
W4-11	Atomization energies (extension of W4-08).

Table 2: Statistical errors (in kcal/mol) over the training sets for the DSD-DFT with different functional combinations, in their optimal parameters.^a

Exchange	S				B88					PBE				
Correlation	VWN5	P86	LYP	PBE	VWN5	LYP	PBE	P86	PW91	VWN5	LYP	PBE	P86	PW91
Grubbs	0.96	1.45	1.06	1.63	1.60	1.10	1.37	1.10	1.29	1.64	1.13	1.18	0.99	1.12
W4-08 ^b	2.58	3.16	3.03	3.61	3.55	2.69	2.61	2.39	2.54	3.31	2.67	2.64	2.33	2.57
non MR	2.30	3.02	2.82	3.58	3.20	2.25	2.43	2.03	2.34	3.04	2.36	2.56	2.07	2.44
MR	3.93	3.90	4.06	3.74	5.21	4.57	3.55	3.94	3.56	4.65	4.07	3.09	3.52	3.26
DBH24	1.38	2.31	1.85	2.76	1.96	1.09	1.34	1.09	1.31	1.57	1.17	1.33	1.11	1.29
s22 ^c	0.33	0.42	0.33	0.51	0.93	0.44	0.61	0.45	0.58	1.01	0.64	0.48	0.40	0.45
Pd	1.27	1.71	1.36	1.74	1.53	0.95	1.55	1.54	1.48	1.34	0.88	1.37	1.37	1.22
MB	4.47	5.12	5.63	4.83	6.05	4.51	3.51	3.50	3.50	5.63	4.41	3.49	3.51	3.59
Mean	1.83	2.36	2.21	2.51	2.60	1.80	1.83	1.68	1.78	2.42	1.82	1.75	1.62	1.71

Exchange	mPW				HSE				O	X	TPSS	B98	BMK	τHCTH	HCTH407
Correlation	LYP	PW91	P86	PBE	PBE	PW91	P86	LYP	LYP	LYP					
Grubbs	1.12	1.15	1.02	1.21	1.09	1.05	0.91	1.08	1.63	1.11	1.45	0.77	3.36	1.69	1.99
W4-08 ^b	2.64	2.48	2.28	2.55	2.63	2.56	2.34	2.70	2.50	2.66	2.50	2.92	3.50	2.98	2.83
non MR	2.27	2.32	1.97	2.40	2.53	2.44	2.05	2.36	2.07	2.25	2.24	2.52	2.96	3.01	2.82
MR	4.30	3.31	3.65	3.31	3.21	3.21	3.68	4.21	4.29	4.43	3.76	4.72	5.83	2.84	2.88
DBH24	1.11	1.26	1.12	1.29	1.32	1.29	1.10	1.17	1.30	1.10	1.43	1.55	4.01	2.50	2.20
s22 ^c	0.52	0.48	0.40	0.51	0.44	0.43	0.36	0.63	0.68	0.47	0.65	0.54	1.15	1.57	2.35
Pd	0.92	1.42	1.51	1.48	1.24	1.19	1.28	0.83	1.27	0.93	2.07	2.33	2.24	1.31	1.02
MB	4.53	3.50	3.43	3.51	3.53	3.52	3.61	4.57	4.29	4.50	4.47	4.10	3.93	4.93	4.90
Mean	1.81	1.72	1.63	1.76	1.71	1.67	1.60	1.83	1.94	1.80	2.09	2.04	3.03	2.50	2.55

^a Including D2 dispersion correction.

^b Full W4-08 set, and discrimination between high- and low- multireference systems.

^c Including one half the counterpoise correction.

Table 3: Parameters of the DSD-DFT with different functional combinations, with D2 dispersion correction.

Exchange	S				B88					PBE				
Correlation	VWN5	P86	LYP	PBE	VWN5	LYP	PBE	P86	PW91	VWN5	LYP	PBE	P86	PW91
c _S	0.11	0.00	0.00	0.00	0.72	0.43	0.13	0.24	0.15	0.67	0.40	0.12	0.23	0.14
c _O	0.58	0.74	0.68	0.77	0.48	0.46	0.51	0.49	0.50	0.47	0.50	0.53	0.51	0.51
s ₆	0.28	0.14	0.17	0.17	0.48	0.35	0.54	0.41	0.53	0.44	0.28	0.42	0.29	0.40
c _C	0.34	0.22	0.29	0.22	0.48	0.55	0.56	0.49	0.56	0.48	0.5	0.51	0.45	0.52
c _X	0.29	0.23	0.25	0.22	0.24	0.29	0.35	0.33	0.35	0.26	0.28	0.34	0.32	0.34

Exchange	mPW				HSE				O	X	TPSS	B98	BMK	τHCTH	HCTH407
Correlation	LYP	PW91	P86	PBE	PBE	PW91	P86	LYP	LYP	LYP					
c _S	0.41	0.14	0.24	0.12	0.12	0.15	0.23	0.40	0.38	0.42	0.30	0.34	0.35	0.41	0.43
c _O	0.48	0.51	0.51	0.51	0.52	0.52	0.50	0.49	0.47	0.47	0.57	0.56	0.79	0.56	0.55
s ₆	0.31	0.47	0.34	0.48	0.39	0.39	0.27	0.26	0.58	0.33	0.28	0.00	0.14	0.30	0.52
c _C	0.53	0.54	0.46	0.54	0.52	0.52	0.46	0.51	0.55	0.54	0.45	0.55	0.31	0.46	0.52
c _X	0.29	0.35	0.32	0.35	0.34	0.34	0.32	0.28	0.31	0.29	0.3	0.22	0.1	0.29	0.29

Table 4: Optimal parameters and statistical errors (in kcal/mol) for DOD-DFT.

	B-LYP	PBE-PBE
c _S	0	0
c _O	0.54	0.51
s ₆	0.98	0.53
c _C	0.59	0.56
c _X	0.36	0.38
Grubbs	1.50	1.18
W4-08	4.53	2.92
non MR	4.42	2.85
MR	5.13	3.31
DBH24	1.11	1.10
s22	2.06	0.52
Pd	1.56	1.35
MB	6.15	3.95
Mean	2.82	1.84

Table 5: RMSD for the S130 set of long range interactions with DSD-PBEP86 (in kcal/mol), with different dispersion correction methods.

	Raw ^a	D2 ^b	D3BJ ^c
RG6	0.24	0.09	0.06
ACONF	0.30	0.12	0.02
ADIM6	1.62	0.19	0.12
S22	1.29	0.26	0.21
CYCONF	0.14	0.11	0.18
SCONF	0.51	0.23	0.26
PCONF ^d	0.98	0.30	0.25
S22+	0.70	0.14	0.11
RMSD ^e	0.63	0.16	0.13

^a No dispersion correction.

^b With $s_6=0.29$.

^c With $s_6=0.418$, $a_2=5.65$ (optimal parameters).

^d With TZVPP basis set.

^e With a weight of 20 for the RG6 set.

Table 6: RMSD for the validation sets (in kcal/mol). Unless specified, all the functionals include D2 dispersion correction.

DSD-DFT	B98	B LYP	B P86	B PBE	B PW91	HSE PBE	mPW LYP	MPW PW91	O LYP	PBE LYP	PBE PBE	S VWN5	TPSS	X LYP	PBE P86	PBE-P86 D3BJ
NHTBH38	2.74	2.08	2.09	2.12	2.06	2.06	2.07	2.00	2.32	2.14	2.06	2.21	2.47	2.04	1.99	2.18
HTBH38	1.29	1.14	1.13	1.07	1.05	1.15	1.14	1.12	1.05	1.10	1.14	1.28	1.07	1.08	1.21	1.12
H-Bonds	0.41	0.45	0.28	0.34	0.31	0.33	0.56	0.28	0.31	0.62	0.30	0.62	0.24	0.49	0.42	0.38
vdW	0.49	0.43	0.31	0.38	0.36	0.33	0.53	0.30	0.29	0.60	0.30	0.45	0.32	0.45	0.42	0.47
Pericyclic																
ref W1	2.76	3.15	3.22	2.97	2.99	2.59	2.85	2.76	3.45	2.72	2.58	1.93	3.38	2.87	2.06	2.05
ref CBS	1.87	2.06	2.01	2.01	1.98	1.79	1.85	1.82	2.23	1.74	1.80	1.71	2.07	1.85	4.16	1.58
ref Exp	3.52	3.65	3.73	3.87	3.82	3.50	3.40	3.60	4.02	3.29	3.56	3.06	3.86	3.40	2.82	2.51
Alkanes																
TAEs	10.89	3.48	3.18	2.24	2.64	1.18	0.80	1.57	7.28	1.09	1.74	2.48	4.10	0.42	0.99	1.07
Isom.	0.07	0.40	0.45	0.50	0.47	0.29	0.31	0.37	0.52	0.32	0.32	0.22	0.21	0.33	0.25	0.16

Table 7: RMSD for the complete W4-11 validation set (140 molecules), and discriminated between non multi-reference (124 molecules) and multi-reference (16 molecules) character systems, in kcal/mol. Unless specified, all the functionals include D2 dispersion correction.

	B98	B LYP	B P86	B PBE	B PW91	HSE PBE	mPW LYP	mPW PW91	mPW P86	mPW PBE	PBE PW91	PBE LYP	PBE PBE	S VWN5	TPSS	X LYP	PBE P86	PBE-P86 D3BJ
All	3.16	2.95	2.41	2.75	2.66	2.82	2.80	2.62	2.41	2.73	2.73	2.83	2.81	2.93	2.77	2.78	2.44	2.48
non-MR	2.65	2.55	1.91	2.49	2.27	2.59	2.31	2.28	1.81	2.40	2.42	2.32	2.51	2.61	2.03	2.10	1.92	1.92
MR	2.67	2.45	2.50	2.20	2.40	2.14	2.58	2.29	2.66	2.29	2.27	2.61	2.26	2.31	2.92	2.83	2.55	2.61

Table 8: RMSD (in cm^{-1}) for the HFREQ27 set with selected methods.

DSD Basis set	S-VWN5 apc2+2d	B-LYP apc2+2d	B-P86 apc2+2d	PBE-P86 apc2+2d	mPW-P86 apc2+2d	B-P86 apc3+d	PBE-P86 apc3+d
Unscaled:							
RMSD	18.93	31.78	21.02	19.65	20.64	17.85	16.71
MSD	10.78	20.31	9.45	8.8	9.84	7.5	6.9
Scaled:							
RMSD	15.08	20.48	17.47	16.56	16.85	15.53	14.82
MSD	0.85	-0.99	-0.85	-0.36	-0.66	-0.13	0.23
Scaling factor	0.9948	0.9889	0.9946	0.9952	0.9945	0.9960	0.9965