

Supplementary data for article:.

Blagojević, J. P.; Zarić, S. D. Stacking Interactions of Hydrogen-Bridged Rings-Stronger than the Stacking of Benzene Molecules. *Chemical Communications* **2015**, *51* (65), 12989–12991. <https://doi.org/10.1039/c5cc04139b>

Cite this: DOI: 10.1039/c0xx00000x

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ARTICLE TYPE

## -Supplementary material-

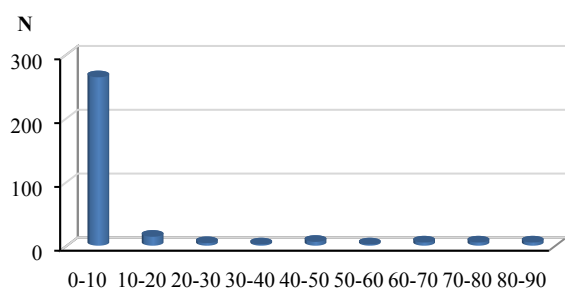
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Received (in XXX, XXX) Xth XXXXXXXXXX 20XX, Accepted Xth XXXXXXXXXX 20XX

DOI: 10.1039/b000000x

### 5 Distribution of interplanar angle and absolute of torsion angles $H_1\Omega_1\Omega_2H_2$ ( $\theta_1$ ) and $A_1\Omega_1\Omega_2A_2$ ( $\theta_2$ )

Interplanar angle and absolute torsion angle  $H_1\Omega_1\Omega_2H_2$  ( $\theta_1$ ) and  $A_1\Omega_1\Omega_2A_2$  ( $\theta_2$ ) distributions, indicating the preferred antiparallel orientation of the middle planes of saturated five-membered hydrogen-bridged rings with planar surroundings of all atoms in the ring, are given in Fig. S1 and S2, respectively.



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Fig. S1. Interplanar angle distributions

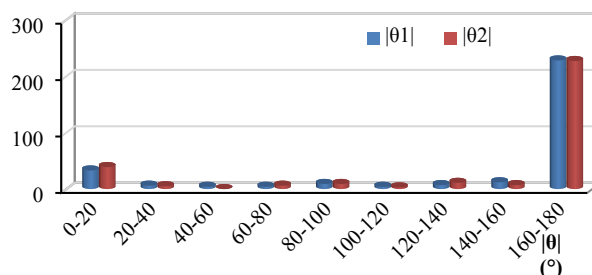


Fig. S2. Distributions of absolute values of torsion angles  $H_1\Omega_1\Omega_2H_2$  ( $|\theta_1|$ ) and  $A_1\Omega_1\Omega_2A_2$  ( $|\theta_2|$ ). Centroid and atom labelling is consistent with the scheme given in Fig 1.

### Evaluation of methods for calculating potential curves

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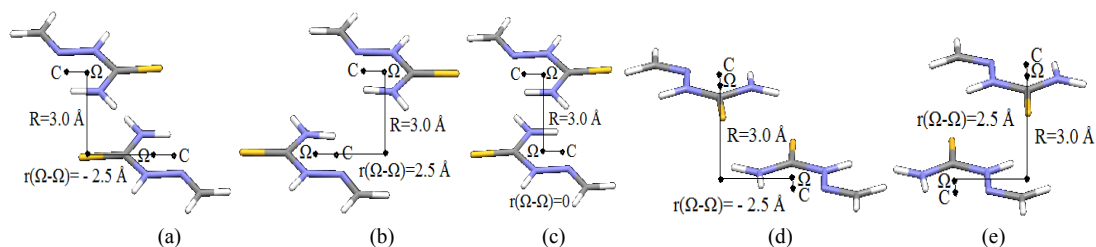


Figure S3. Model systems of 2-methylidenehydrazinecarbothioamide, used for the evaluation of different methods: a) orientation 1; b) orientation 2; c) orientation 3; d) orientation 4; e) orientation 5

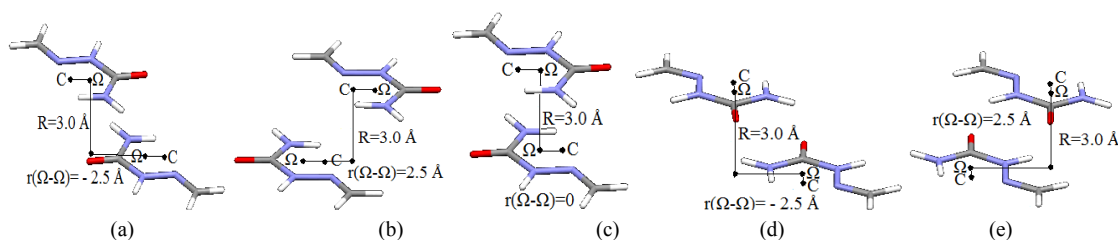


Figure S4. Model systems of 2-methylidenehydrazinecarboxamide, used for the evaluation of different methods: a) orientation 1; b) orientation 2; c) orientation 3; d) orientation 4; e) orientation 5

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Several model systems (Figures S3 and S4) were chosen for the estimation of interaction energies at CCSD(T) level at complete basis set, by performing the method of Mackie and DiLabio<sup>1</sup>. Model systems are composed of dimers, having antiparallel middle molecular planes and offset values of -2.5 Å and 2.5 Å along Ω-C direction (orientations 1 and 2, respectively- Figures S3 and S4), 0.0 Å (orientation 3- Figures S3 and S4), -2.5 Å and 2.5 Å along the direction orthogonal to Ω-C direction (orientations 4 and 5, respectively- Figures S3 and S4). Distances between middle molecular planes in all model systems are 3.0 Å. Calculated interaction energies at CCSD(T)/CBS level are given in Tables S1-S5. Møller-Plesett perturbation theory of the second order (MP2)<sup>2</sup> and several D3 corrected<sup>3</sup> DFT functionals were tested in order to find less time-consuming methods for the calculation of potential curves, which approximately reproduce CCSD(T)/CBS interaction energies. Basis set effect was also tested by applying several basis sets: cc-pVDZ<sup>4</sup>, cc-pVTZ<sup>5</sup>, aug-cc-pVDZ<sup>4,6</sup> and 6-31++G\*\*<sup>7</sup>. The results are given in Tables S1-S5 for all five orientations. with the values obtained by methods that were chosen for calculation of potential curves pointed out within a box. The selected method for calculating potential curves in the Ω-C direction is BLYP-d3/cc-pVDZ, since this method gives good results in orientations 2 (Table S1) and 3 (Table S3), while M06HF-d3/cc-pVDZ and M052X-d3/6-31++G\*\* methods were chosen for calculating potential curves for 2-methylidenehydrazinecarbothioamide and 2-methylidenehydrazinecarboxamide, respectively, in the orthogonal direction, since they give good results in orientations 3 (Table S3), 4 (Table S4), and 5 (Table S5). Neither of the selected methods gives particularly good results for orientations 1, but this is of no great importance, since energies are not high.

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Table S1. Interaction energies in kcal/mol, obtained by MP2 method and different D3 corrected DFT functionals, by using several basis sets and energies at CCSD(T)/CBS level, calculated on model systems corresponding to orientation 1 in Figures S3 and S4

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orientation 1	2-methylidenehydrazinecarbothioamide				2-methylidenehydrazinecarboxamide			
	cc-pVDZ	cc-pVTZ	aug-cc-pVDZ	6-31++G**	cc-pVDZ	cc-pVTZ	aug-cc-pVDZ	6-31++G**
MP2	4.99	0.22	-0.58	3.63	1.63	-0.48	-1.13	0.58
TPSS-D3	1.77	1.65	0.96	1.18	-0.48	-0.77	-0.95	-0.74
BLYP-D3	2.62	2.04	1.90	2.12	-1.17	-1.36	-1.39	-1.07
BP86-D3	0.60	-0.15	-0.49	-0.21	-0.85	-1.43	-1.54	-1.21
PBE1PBE-D3	2.01	1.66	1.20	1.36	-0.37	-0.87	-1.01	-0.80
M05-D3	3.51	3.45	2.75	3.00	-0.68	-1.02	-1.45	-1.29
M06-D3	1.18	0.29	-0.22	0.05	-1.13	-1.98	-2.21	-2.11
M052X-D3	2.05	1.06	0.56	0.79	-0.63	-1.47	-1.67	-1.55
M06HF-D3	-0.36	-0.96	-2.43	-1.61	-1.10	-1.68	-2.34	-2.56
<b>CCSD(T)/CBS</b>			<b>-0.72</b>				<b>-1.92</b>	

Table S2. Interaction energies in kcal/mol, obtained by MP2 method and different D3 corrected DFT functionals, by using several basis sets and energies at CCSD(T)/CBS level, calculated on model systems corresponding to orientation 2 in Figures S3 and S4

orientation 2	2-methylidenehydrazinecarbothioamide				2-methylidenehydrazinecarboxamide			
	cc-pVDZ	cc-pVTZ	aug-cc-pVDZ	6-31++G**	cc-pVDZ	cc-pVTZ	aug-cc-pVDZ	6-31++G**
MP2	-0.27	-1.59	-1.90	-0.70	0.55	-0.72	-1.03	0.08
TPSS-D3	-2.16	-2.19	-2.23	-2.13	-1.48	-1.54	-1.60	-1.50
BLYP-D3	-2.28	-2.24	-2.13	-2.03	-1.60	-1.58	-1.51	-1.41
BP86-D3	-2.03	-2.26	-2.25	-2.12	-1.36	-1.62	-1.64	-1.50
PBE1PBE-D3	-2.04	-2.21	-2.25	-2.16	-1.37	-1.57	-1.63	-1.54
M05-D3	-2.15	-2.22	-2.40	-2.38	-1.55	-1.65	-1.85	-1.83
M06-D3	-1.60	-2.03	-2.25	-2.34	-1.02	-1.52	-1.74	-1.78

M052X-D3	-1.69	-2.12	-2.20	-2.18	-1.06	-1.50	-1.60	-1.56
M06HF-D3	-1.36	-1.73	-2.02	-2.22	-0.68	-1.07	-1.37	-1.60
<b>CCSD(T)/CBS</b>	<b>-2.23</b>				<b>-1.54</b>			

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Table S3. Interaction energies in kcal/mol, obtained by MP2 method and different D3 corrected DFT functionals, by using several basis sets and energies at CCSD(T)/CBS level, calculated on model systems corresponding to orientation 3 in Figures S3 and S4

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orientation 3	2-methylidenehydrazinecarbothioamide				2-methylidenehydrazinecarboxamide			
	cc-pVDZ	cc-pVTZ	aug-cc-pVDZ	6-31++G**	cc-pVDZ	cc-pVTZ	aug-cc-pVDZ	6-31++G**
MP2	0.74	-2.76	-3.49	-0.58	2.53	-0.25	-0.83	1.28
TPSS-D3	-2.29	-2.36	-2.72	-2.86	-0.41	-0.58	-0.93	-0.99
BLYP-D3	-2.89	-3.37	-3.14	-3.18	-1.16	-1.53	-1.42	-1.38
BP86-D3	-3.64	-4.24	-4.35	-4.40	-1.41	-1.96	-2.10	-2.03
PBE1PBE-D3	-2.07	-2.34	-2.59	-2.75	-0.31	-0.65	-0.89	-0.97
M05-D3	-1.67	-1.73	-2.14	-2.19	-0.23	-0.46	-0.86	-0.92
M06-D3	-3.33	-3.94	-4.54	-4.71	-1.23	-2.00	-2.33	-2.42
M052X-D3	-2.07	-2.68	-3.22	-3.45	-0.30	-0.99	-1.40	-1.50
M06HF-D3	-3.58	-3.73	-5.20	-5.53	-1.25	-1.65	-2.77	-3.28
<b>CCSD(T)/CBS</b>	<b>-3.64</b>				<b>-1.54</b>			

Table S4. Interaction energies in kcal/mol, obtained by MP2 method and different D3 corrected DFT functionals, by using several basis sets and energies at CCSD(T)/CBS level, calculated on model systems corresponding to orientation 4 in Figures S3 and S4

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orientation 4	2-methylidenehydrazinecarbothioamide				2-methylidenehydrazinecarboxamide			
	cc-pVDZ	cc-pVTZ	aug-cc-pVDZ	6-31++G**	cc-pVDZ	cc-pVTZ	aug-cc-pVDZ	6-31++G**
MP2	-1.57	-2.74	-3.07	-2.03	0.35	-0.75	-1.01	-0.24
TPSS-D3	-3.39	-3.29	-3.37	-3.45	-1.70	-1.72	-1.85	-1.94
BLYP-D3	-3.37	-3.26	-3.14	-3.19	-1.73	-1.74	-1.69	-1.75
BP86-D3	-3.29	-3.35	-3.33	-3.41	-1.63	-1.79	-1.83	-1.89
PBE1PBE-D3	-3.27	-3.32	-3.40	-3.48	-1.57	-1.72	-1.83	-1.92
M05-D3	-3.12	-3.18	-3.41	-3.48	-1.50	-1.64	-1.85	-1.95
M06-D3	-3.03	-3.50	-3.68	-3.86	-1.21	-1.68	-1.81	-1.96
M052X-D3	-2.83	-3.23	-3.32	-3.50	-1.17	-1.59	-1.69	-1.80
M06HF-D3	-3.28	-3.69	-4.16	-4.36	-1.34	-1.78	-2.22	-2.47
<b>CCSD(T)/CBS</b>	<b>-3.27</b>				<b>-1.57</b>			

Table S5. Interaction energies in kcal/mol, obtained by MP2 method and different D3 corrected DFT functionals, by using several basis sets and energies at CCSD(T)/CBS level, calculated on model systems corresponding to orientation 5 in Figures S3 and S4

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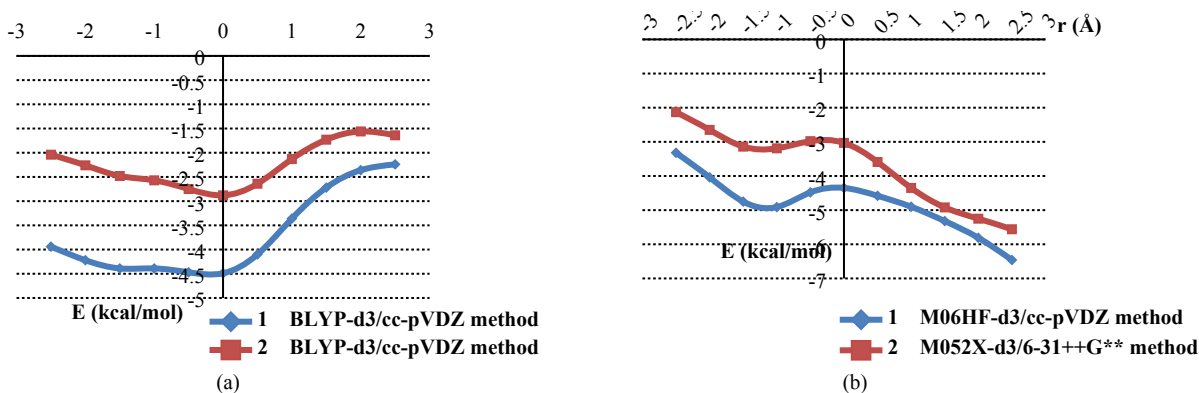
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orientation 5	2-methylidenehydrazinecarbothioamide				2-methylidenehydrazinecarboxamide			
	cc-pVDZ	cc-pVTZ	aug-cc-pVDZ	6-31++G**	cc-pVDZ	cc-pVTZ	aug-cc-pVDZ	6-31++G**
MP2	0.20	-4.37	-5.05	-1.19	-0.14	-3.50	-4.27	-2.02
TPSS-D3	-2.95	-3.29	-3.71	-3.79	-2.60	-3.02	-3.39	-3.51
BLYP-D3	-2.71	-3.34	-3.43	-3.35	-3.15	-3.74	-3.83	-3.79
BP86-D3	-4.37	-5.15	-5.42	-5.38	-3.60	-4.42	-4.69	-4.67
PBE1PBE-D3	-3.08	-3.56	-3.90	-4.01	-2.80	-3.41	-3.72	-3.86
M05-D3	-2.03	-2.16	-2.87	-2.92	-2.70	-3.05	-3.71	-3.88
M06-D3	-4.08	-4.91	-5.77	-6.05	-3.56	-4.60	-5.23	-5.52
M052X-D3	-3.14	-4.35	-4.63	-4.82	-3.14	-4.38	-4.66	-4.87
M06HF-D3	-5.30	-6.45	-7.32	-7.33	-4.56	-5.64	-6.49	-7.07
<b>CCSD(T)/CBS</b>	<b>-5.31</b>				<b>-4.89</b>			

## Potential curves

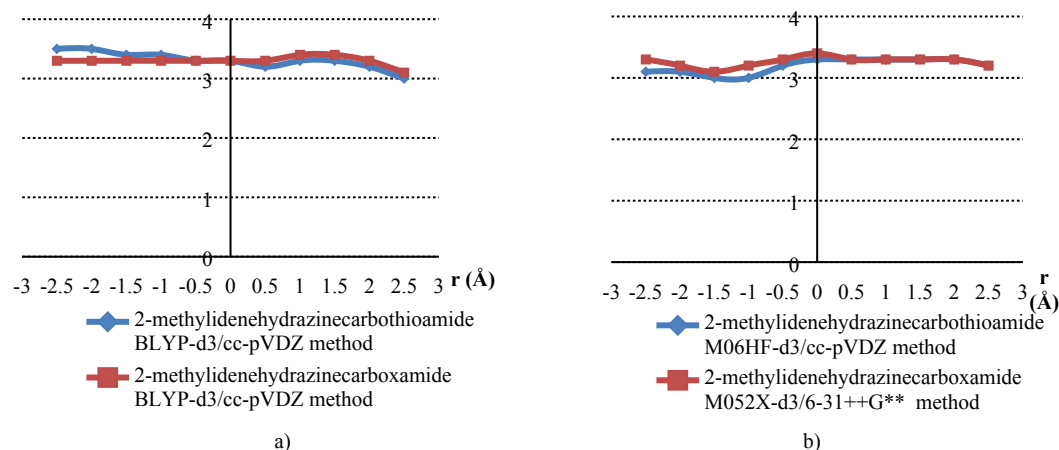
Potential curves obtained by moving molecules along  $\Omega$ -C direction and along the orthogonal direction are shown in Fig S5.



5 Fig S5. (a) Potential curves for offset values varied in the  $\Omega$ -C direction; (b) Potential curves for offset values varied orthogonal to  $\Omega$ -C direction

## Normal distances

Figure S6 shows normal distance dependencies on offset values.



10 Figure S6. Normal distance dependencies on offset values; a)  $\Omega$ -C direction; b) orthogonal to  $\Omega$ -C direction

Normal distances are approximately constant, in the range between 3.0 and 3.5 Å, which is in accord with the results of crystal structure analysis (Fig. 2).

## Electrostatic potentials

15 Maps of electrostatic potentials for 2-methylidenehydrazinecarbothioamide and 2-methylidenehydrazinecarboxamide molecules, calculated and visualized from wavefunction files using the Wavefunction Analysis Program (WFA-SAS)<sup>8,9</sup> are shown in Figure S7. The wavefunctions were calculated on mp2/cc-pVTZ level of theory using Gaussian09 series of programs<sup>10</sup>. Electronic density is more localized on oxygen atom in case of 2-methylidenehydrazinecarboxamide than it is localized on sulphur atom in case of 2-methylidenehydrazinecarbothioamide.

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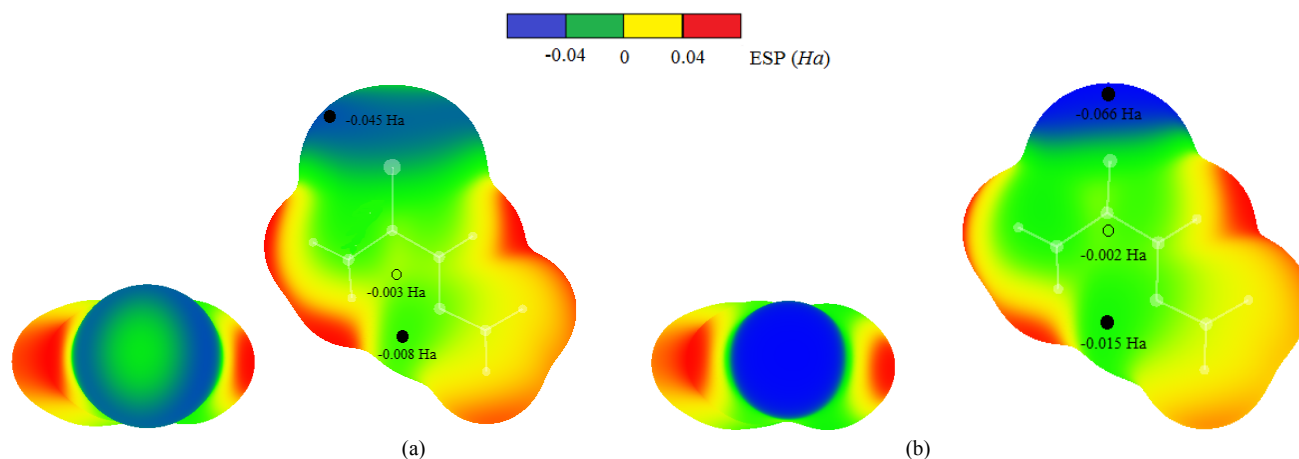


Figure S7. Side and top views of electrostatic potential maps for (a) 2-methylidenehydrazinecarbothioamide and (b) 2-methylidenehydrazinecarboxamide molecule; Values of some important ESP maxima (empty circles) and minima (filled circles) in hartrees ( $H_a$ ) are indicated onto the surfaces of the maps.

## Notes and references

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