

COMMENTS

Comment on “Strength of the N–H···O=C Bonds in Formamide and N-Methylacetamide Dimers”

Jann A. Frey and Samuel Leutwyler*

Departement für Chemie und Biochemie, Universität Bern,
Freiestrasse 3, CH-3012 Bern, Switzerland

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Cis-amides are common in biologically important molecules. They provide neighboring hydrogen bonding N–H donor and C=O acceptor sites, thereby significantly influencing the structures of nucleic acids, proteins, and peptides. The formamide dimer, (FA)₂, is the simplest cis-amide dimer and has therefore been widely studied experimentally and theoretically in the gas and liquid phases.^{1–5} The results obtained are essential for understanding and modeling the structures and interactions between more extended biopolymers.

Vargas et al. calculated the binding energies of five different H-bonded (FA)₂ dimers at the MP2 level of theory, using the aug-cc-pVXZ (X = D, T, and Q) basis sets and extrapolated them to the complete basis set (CBS) limit.¹ The aug-cc-pVXZ series of basis sets was designed⁶ for frozen-core MP2 calculations; i.e., the 1s electrons should not be correlated. However, Vargas et al. employed the aug-cc-pVXZ basis sets for fully correlated calculations.¹ We address several points, comparing their CBS limit binding energy of the most stable (formamide)₂ isomer with values obtained using frozen-core MP2 calculations. The binding energies D_e and the counterpoise (CP) corrected binding energies D_e^{CPC} obtained by frozen-core and fully correlated¹ calculations are compared in Table 1 and plotted in Figure 1.

First, in ref 1 only half of the necessary CP correction was applied to the aug-cc-pVDZ binding energy, giving rise to a CBS extrapolation curve with an unusual shape. The correctly calculated D_e^{CPC} , marked by a diamond in Figure 1b, is given in parentheses in Table 1 and leads to a monotonically decreasing CBS extrapolation curve, as is usually found. Second, the CBS binding energy $D_{e,\text{CBS}} = -14.35$ kcal/mol given in ref 1 corresponds to an extrapolation⁷ of the CP-uncorrected D_e values *only*. No CBS limit binding energy for the CP-corrected D_e^{CPC} energies was given in ref 1, which is unusual, given that both extrapolations should, in principle, lead to nearly the same value. It is also a priori unclear why the uncorrected and not the corrected binding energies should be extrapolated. We calculated the CP-corrected value to be $D_{e,\text{CBS}}^{\text{CPC}} = -14.82$ kcal/mol (using the correct aug-cc-pVDZ CP-corrected binding energy). This value is about 0.5 kcal/mol larger than the $D_{e,\text{CBS}}$ value; see also Figure 1b. This rather large difference can be traced to the use of the fully correlated calculation: Extrapolation⁸ of the *frozen core* corrected and uncorrected binding energies result in $D_{e,\text{CBS}}(\text{FC}) = -14.86$ kcal/mol and $D_{e,\text{CBS}}^{\text{CPC}}(\text{FC}) = -14.72$ kcal/mol. The much smaller difference between these two extrapolated binding energies shows that the CBS procedure works reliably if the basis sets are used as intended.⁷ The average frozen core CBS value $D_e^{\infty}(\text{FC}) =$

TABLE 1: Calculated Frozen Core and Fully Correlated MP2 Binding Energies of (formamide)₂ (kcal/mol), Using the aug-cc-pVXZ (X = D, T, Q) Basis Sets. the Fully Correlated Binding Energies in Parentheses Are Recalculated

	fully correlated		frozen core	
	D_e	D_e^{CPC}	D_e	D_e^{CPC}
aug-cc-pVDZ	-15.80 ^a	-14.68 ^a (-13.39) ^b	-15.598 ^b	-13.388 ^b
aug-cc-pVTZ	-16.83 ^a	-13.98 ^a	-15.362 ^b	-14.109 ^b
aug-cc-pVQZ	-15.37 ^a	-14.49 ^a	-15.073 ^b	-14.482 ^b
CBS limit	-14.35 ^{a,c}	(-14.82) ^{b,c}	-14.86 ^{b,d}	-14.72 ^{b,d}

^a Values from ref 1. ^b This work. ^c Extrapolation to the complete basis set limit according to ref 7. ^d Extrapolation to the complete basis set limit according to ref 8.

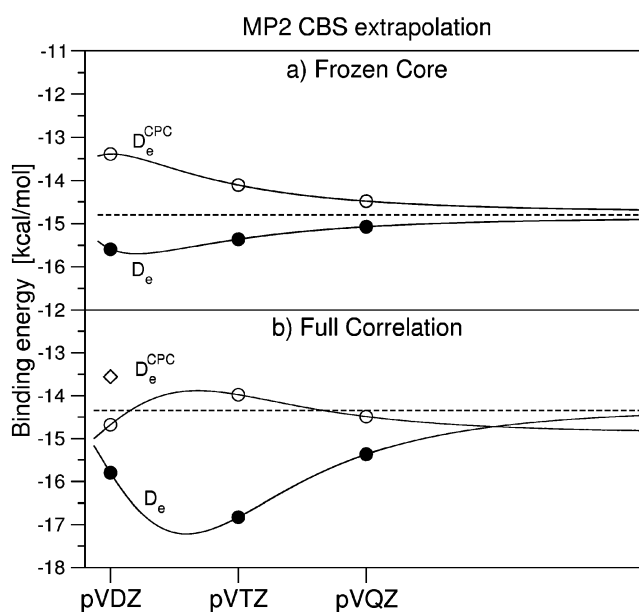


Figure 1. Complete basis set (CBS) extrapolations of the binding energies D_e (●) and counterpoise-corrected binding energies D_e^{CPC} (○) of the most stable isomer of (formamide)₂ using (a) frozen core, this work, and (b) fully correlated MP2 calculations, reported in ref 1. The aug-cc-pVDZ point in (b) labeled by ○ should be corrected to the value ◇; see Table 1 and the text. The dashed lines correspond to the CBS limits: (a) frozen core, this work; (b) fully correlated, reported in ref 1.

-14.79 kcal/mol is shown as a dashed line in Figure 1a. This binding energy is 0.51 kcal/mol larger than that reported in ref 1.

References and Notes

- Vargas, R.; Garza, J.; Friesner, R. A.; Stern, H.; Hay, B. P.; Dixon, D. A. *J. Phys. Chem. A* **2001**, *105*, 4963.
- Watson, T. M.; Hirst, J. D. *J. Phys. Chem.* **2002**, *106*, 7858.
- Ohtaki, H. *J. Mol. Liq.* **2003**, *103*, 3.
- Ireta, J.; Neugebauer, J.; Scheffler, M. *J. Phys. Chem. A* **2004**, *108*, 5692.
- Müller, A.; Losada, M.; Leutwyler, S. *J. Phys. Chem. A* **2004**, *108*, 157.
- Kendall, R. A.; Dunning, T. H., Jr.; Harrison, R. J. *J. Chem. Phys.* **1992**, *96*, 6796.
- Peterson, K. A.; Woon, D. E.; Dunning, T. H. *J. Chem. Phys.* **1994**, *100*, 7410.
- Klopper, W. *J. Chem. Phys.* **1995**, *102*, 6168.