C7790 Introduction to Molecular Modelling TSM Modelling Molecular Structures

Lesson 17 Reaction Energy II (QM specific)

PS/2020 Distant Form of Teaching: Rev1

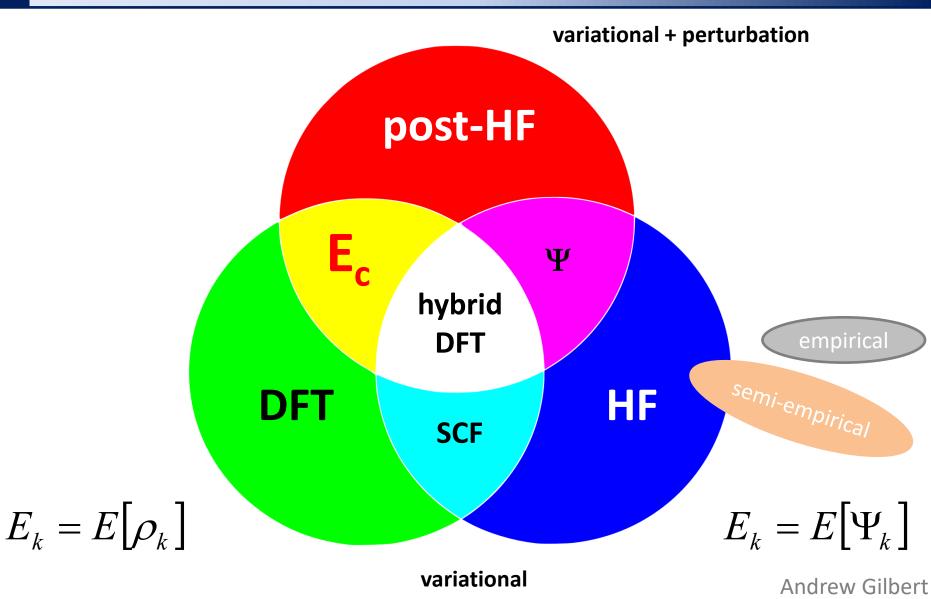
Petr Kulhánek

kulhanek@chemi.muni.cz

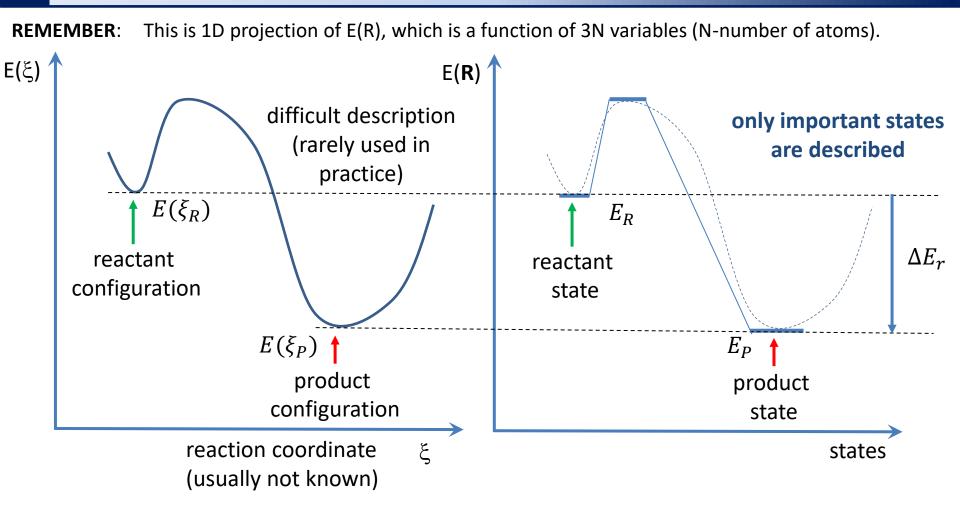
National Centre for Biomolecular Research, Faculty of Science Masaryk University, Kamenice 5, CZ-62500 Brno

C7790 Introduction to Molecular Modelling

Revision: QM method classification



Reaction Energy



reaction energy

the sign convention:

 $\Delta E_r = E(\xi_P) - E(\xi_R) = E_P - E_R$

always use the thermodynamics convention

QM - two approches

I. supermolecular approach:

each component is characterized individually

$$\Delta E_r = E(\xi_P) - E(\xi_R) = E_P - E_R$$

II. energy decomposition (EDA):

- only interaction energy is accessible for non-covalent interactions
- no deformation energy is available
- > typical methods:
 - SAPT (Symmetry-adapted perturbation theory)

$$E_{int}^{SAPT0}$$

Supermolecular approach

I. supermolecular approach:

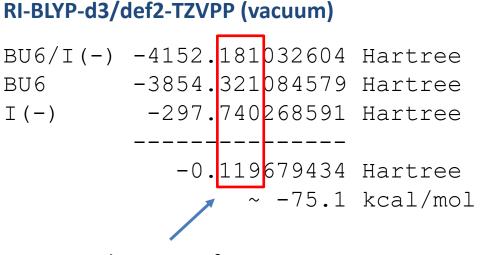
each component is characterized individually

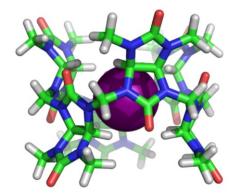
$$\Delta E_r = E(\xi_P) - E(\xi_R) = E_P - E_R$$

It seems to be SIMPLE. But it is NOT because ...

Supermolecular approach, cont

Problem 1: Small numbers from big numbers





bambus[6]uril/anion interaction (139 atoms)

chemistry of interest

Requirements:

- robust (numerically stable) algorithms
- well optimized geometries
- well converged WF and energy

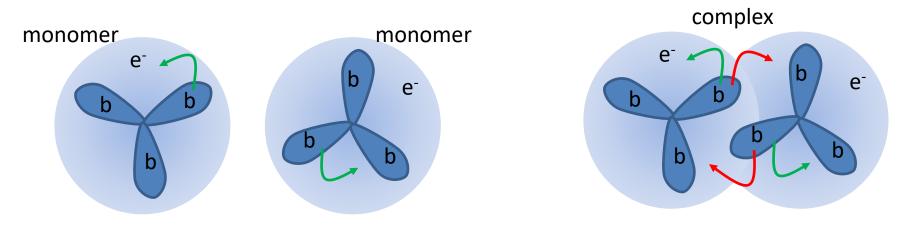
Supermolecular approach, cont.

Problem 2: Basis set superposition error

This error is consequence of **finite atom centered basis sets** and **variational nature of employed theory** (HF, DFT). It also influences non-variational post-HF methods.

Cause:

- As the atoms of interacting molecules (or different parts of the same molecule) approach one another, their basis functions overlap.
- Each monomer "borrows" functions from other nearby components, effectively increasing its basis set and improving the calculation of derived properties such as energy.

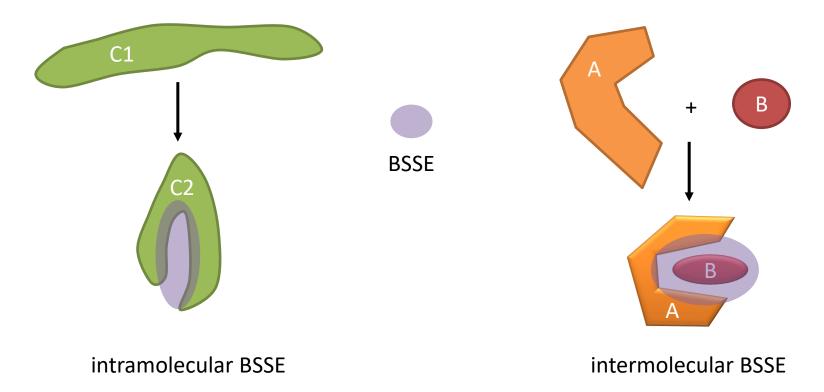


long-distance separation https://en.wikipedia.org/wiki/Basis_set_superposition_error

Basis set superposition error (BSSE)

Types of basis set superposition error:

- intramolecular (conformation changes)
- intermolecular (interaction)

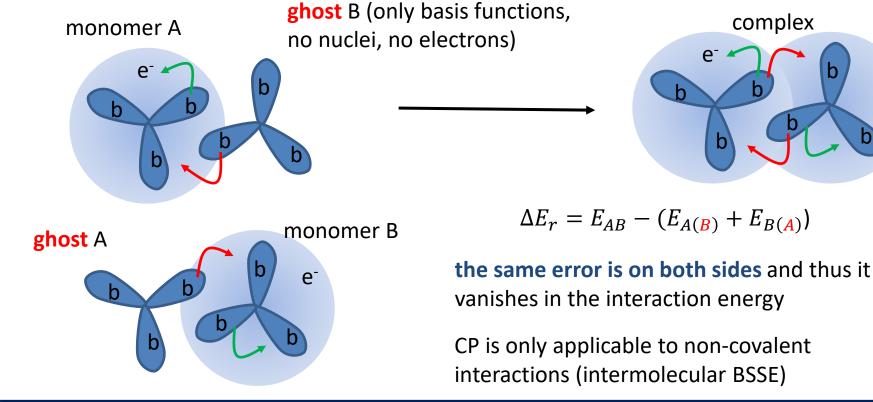


BSSE corrections

BSSE can be avoided or corrected by:

- the chemical Hamiltonian approach (CHA) a priori correction
- the counterpoise method (CP) a posteriori correction
- extrapolation to CBS limit a posteriori correction
- > space centered basis functions such as plane waves *a priori* correction

Counterpoise method



C7790 Introduction to Molecular Modelling

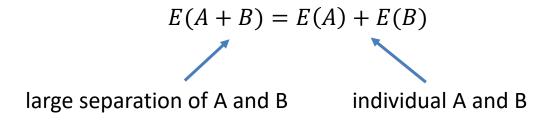
e

Supermolecular approach, cont.

Problem 3: Size consistency

Size consistency is a concept relating to how the behavior of quantum chemistry calculations changes with size.

Size consistency (or strict separability) is a property that guarantees the consistency of the energy behavior when interaction between the involved molecular system is nullified (for example, by distance).



For example:

The Restricted Hartree–Fock model (RHF, a single reference method) is not able to correctly describe the dissociation curves of H_2 and therefore all post HF methods that employ HF as a starting point will fail in that matter.

The solution would be to use multi-reference methods, which are however more computationally demanding.

Supermolecular approach, cont.

Problem 4: Deficient description of long-range interactions

HF and DFT method provide no or incomplete treatment of dispersion interaction. Dispersion interaction is a weak attractive long-range force.

This can be problematic for studying systems, in which these forces dominates:

- noble gases interaction
- supramolecular and biomolecular systems

Solution:

dispersion corrected methods:

- ➢ HF-3c
- DFT-D3, DFT-D4, etc.

Further readings:

Grimme, S.; Hansen, A.; Brandenburg, J. G.; Bannwarth, C. Dispersion-Corrected Mean-Field Electronic Structure Methods. *Chem. Rev.* **2016**, *116* (9), 5105–5154. <u>https://doi.org/10.1021/acs.chemrev.5b00533</u>.

Symmetry-adapted perturbation theory

SAPT (Symmetry-adapted perturbation theory)

- only interaction energy is accessible for non-covalent interactions
- no deformation energy is available
- it employs the perturbation theory (the method is not variational)
- interaction energy is composed from several contributions:
 - electrostatic
 - exchange repulsion
 - ➤ induction
 - ➤ dispersion
 - and other contributions ...

$$E_{int}^{SAPT0} = E_{ele}^{(1)} + E_{exch}^{(1)} + E_{ind}^{(2)} + E_{exch-ind}^{(2)} + E_{disp}^{(2)} + E_{exch-disp}^{(2)}$$

- no susceptible to BSSE, but the accuracy strongly depends on basis set
- accuracy might depend on cancellation of errors (low order SAPT + specially tuned basis sets)
- high accuracy requires higher orders (SAPT2, ...), which are computationally demanding

Summary

- While QM provides very sophisticated methods, their use is practically difficult due to several QM specific problems:
 - numerical stability
 - basis set dependence of calculated properties including energy
 - size consistency
 - deficiency in proper description of long-range interactions
- Therefore, a special care must be taken when QM is utilized for calculation of reaction, binding, and interaction energies.