# C7790 Introduction to Molecular Modelling TSM Modelling Molecular Structures

Lesson 17
Reaction Energy II (QM specific)

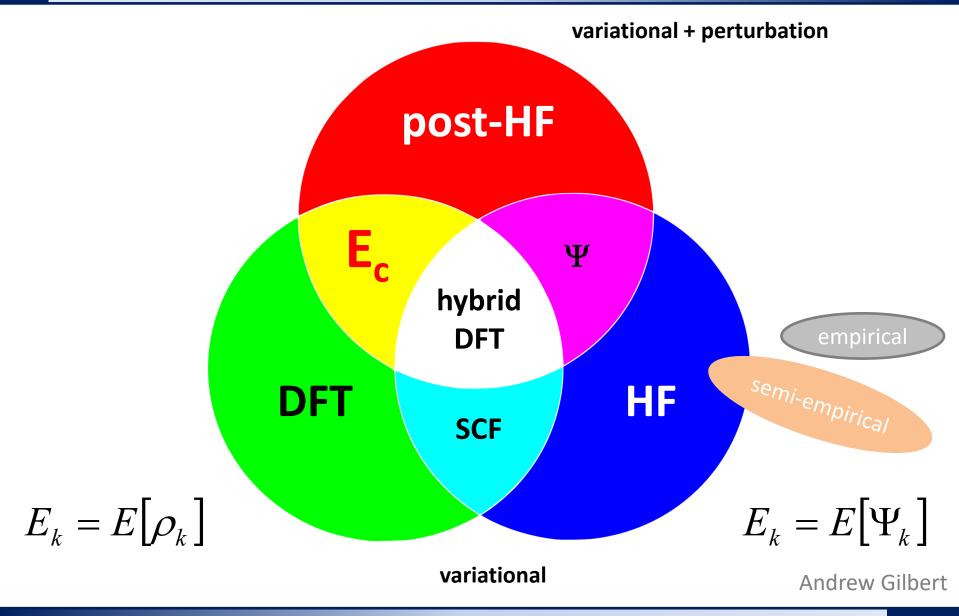
JS/2022 Distant Form of Teaching: Rev1

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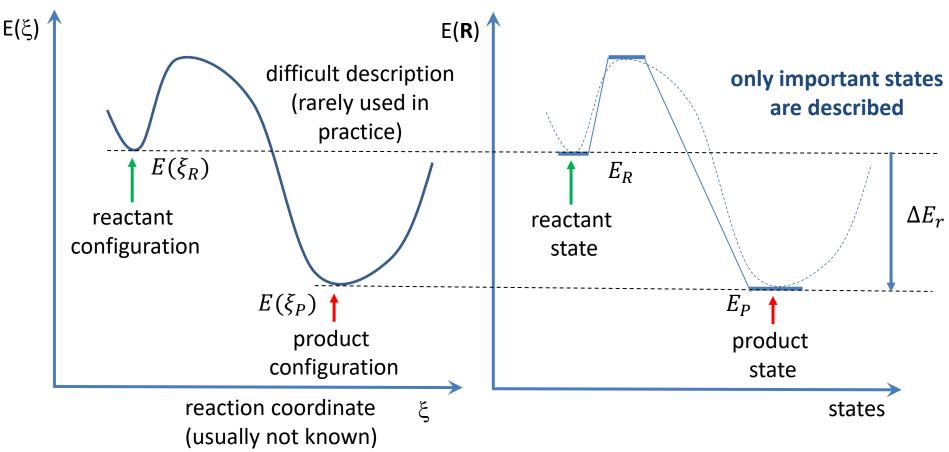
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### Revision: QM method classification



### **Reaction Energy**

**REMEMBER**: This is 1D projection of E(R), which is a function of 3N variables (N-number of atoms).



### reaction energy

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

the sign convention: 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

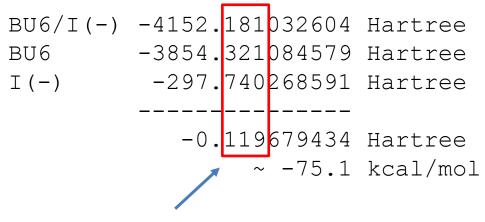
$$\Delta E_r = E(\xi_P) - E(\xi_R) = E_P - E_R$$

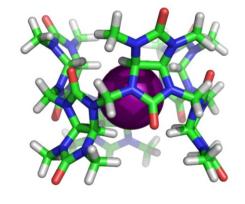
## It seems to be SIMPLE. But it is NOT because ...

### Supermolecular approach, cont

### **Problem 1: Small numbers from big numbers**

#### RI-BLYP-d3/def2-TZVPP (vacuum)





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

chemistry of interest

#### **Requirements:**

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

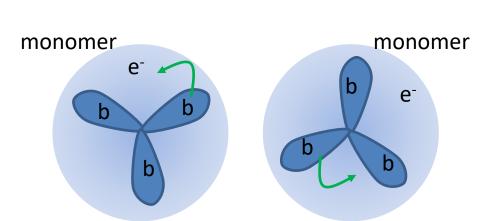
### Supermolecular approach, cont.

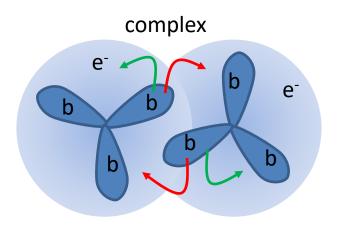
### **Problem 2: Basis set superposition error**

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.





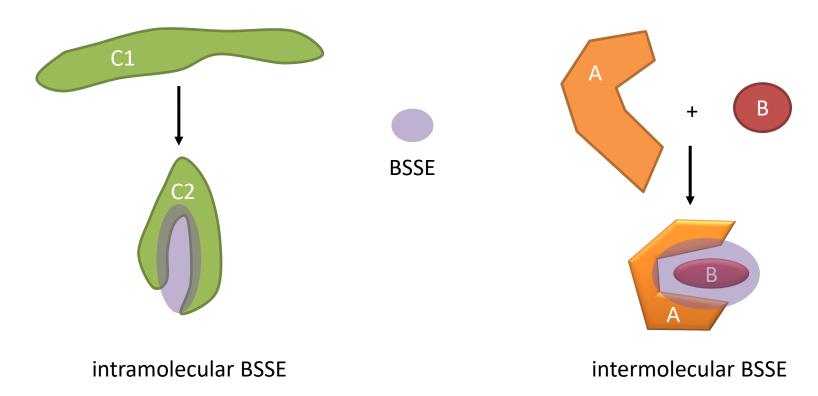
long-distance separation

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)

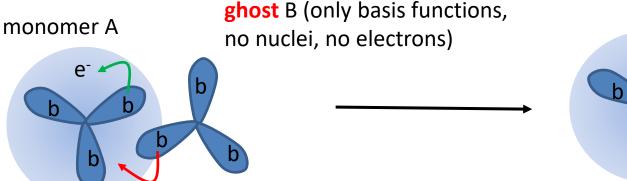


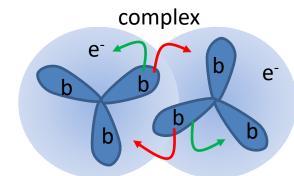
### **BSSE** corrections

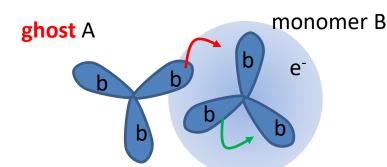
BSSE can be avoided or corrected by:

- 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**







$$\Delta E_r = E_{AB} - (E_{A(B)} + E_{B(A)})$$

the same error is on both sides and thus it vanishes in the interaction energy

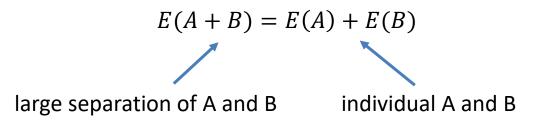
CP is only applicable to non-covalent interactions (intermolecular BSSE)

### 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<sub>2</sub> 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. <a href="https://doi.org/10.1021/acs.chemrev.5b00533">https://doi.org/10.1021/acs.chemrev.5b00533</a>.

### 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.