

Extrapolation techniques to improve the scaling of electronic structure

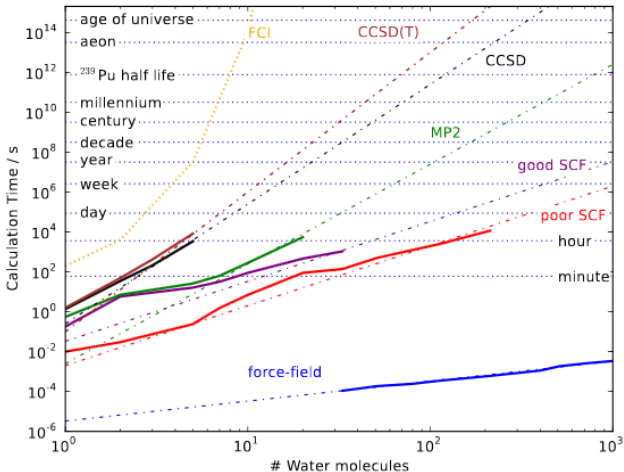
Ankit Mahajan

Department of Physics,
IIT Bombay

Outline

- 1 Introduction
 - More is difficult!
 - What can we do?
 - Tasks
- 2 PIE-ONIOM
 - What is PIE-ONIOM?
 - How is it implemented?
 - BOMD results
- 3 π interactions
 - Introduction
 - Dimers
 - Trimer and tetramer
- 4 DMRG
 - MPS
 - Extrapolation

More is difficult!



What can we do?

- Come up with new methods: DMRG

What can we do?

- Come up with new methods: DMRG
- Improve existing ones: novel DFT functionals, improvements to MP2

What can we do?

- Come up with new methods: DMRG
- Improve existing ones: novel DFT functionals, improvements to MP2
- Composite approaches: make a series of systematic approximations and assemble to extrapolate to an accurate result

What can we do?

- Come up with new methods: DMRG
- Improve existing ones: novel DFT functionals, improvements to MP2
- Composite approaches: make a series of systematic approximations and assemble to extrapolate to an accurate result

Tasks

- Doing dynamics with DMRG: Analytic DMRG gradients available in ORCA
Focus on strongly correlated systems

Tasks

- Doing dynamics with DMRG: Analytic DMRG gradients available in ORCA
Focus on strongly correlated systems

Tasks

- Doing dynamics with DMRG: Analytic DMRG gradients available in ORCA
Focus on strongly correlated systems
Have done static DMRG

Tasks

- Doing dynamics with DMRG: Analytic DMRG gradients available in ORCA
Focus on strongly correlated systems
Have done static DMRG
- π -stacking systems: Accounting for non-covalent bonded interactions
Benzene trimers, tetramers, etc.

Tasks

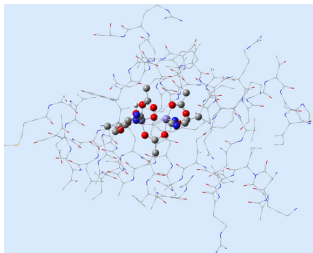
- Doing dynamics with DMRG: Analytic DMRG gradients available in ORCA
Focus on strongly correlated systems
Have done static DMRG
- π -stacking systems: Accounting for non-covalent bonded interactions
Benzene trimers, tetramers, etc.

Outline

- 1 Introduction
 - More is difficult!
 - What can we do?
 - Tasks
- 2 PIE-ONIOM
 - What is PIE-ONIOM?
 - How is it implemented?
 - BOMD results
- 3 π interactions
 - Introduction
 - Dimers
 - Trimer and tetramer
- 4 DMRG
 - MPS
 - Extrapolation

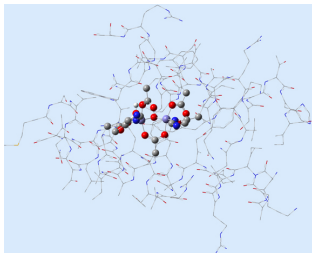
What is PIE-ONIOM?

Hybrid methods
Divide into layers



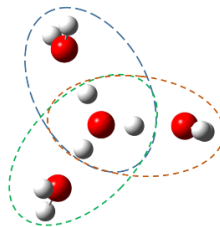
What is PIE-ONIOM?

Hybrid methods
Divide into layers



+

Frgament methods
Divide into fragments

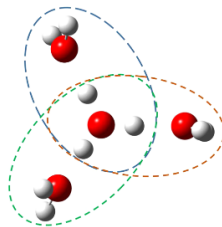
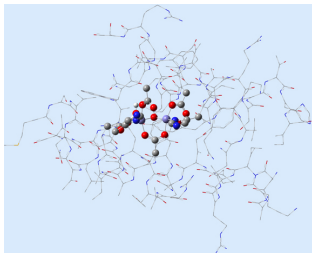


What is PIE-ONIOM?

Hybrid methods
Divide into layers

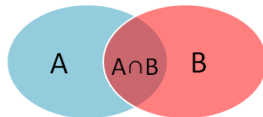
+

Frgament methods
Divide into fragments



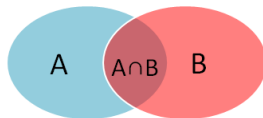
PIE-ONIOM
Divide into layered fragments

What is PIE-ONIOM?



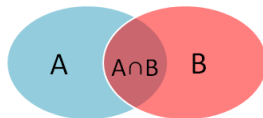
$$E_{PIE-ONIOM} = [E^{high}(A) + E^{high}(B) - E^{high}(A \cap B)]$$

What is PIE-ONIOM?



$$E_{PIE-ONIOM} = [E^{high}(A) + E^{high}(B) - E^{high}(A \cap B)] \\ + E^{low}(A \cup B) - [E^{low}(A) + E^{low}(B) - E^{low}(A \cap B)]$$

What is PIE-ONIOM?



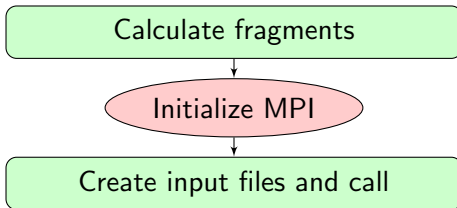
$$E_{PIE-ONIOM} = [E^{high}(A) + E^{high}(B) - E^{high}(A \cap B)] \\ + E^{low}(A \cup B) - [E^{low}(A) + E^{low}(B) - E^{low}(A \cap B)]$$

Generalize with the principle of inclusion-exclusion

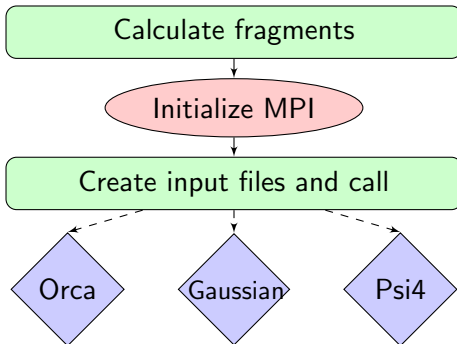
How is it implemented?

Calculate fragments

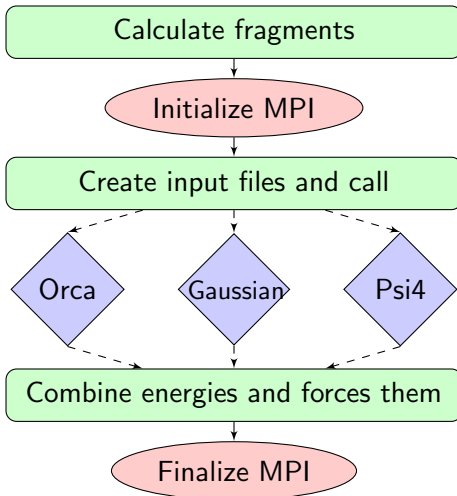
How is it implemented?



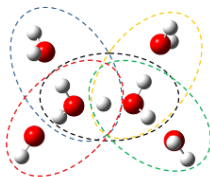
How is it implemented?



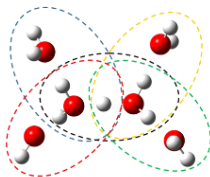
How is it implemented?



Born Oppenheimer Molecular Dynamics



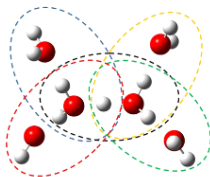
Born Oppenheimer Molecular Dynamics



$$M_I \ddot{\mathbf{R}}_I(t) = -\nabla_I \min \left\{ \int \Psi^* \hat{H} \Psi d\mathbf{r} \right\}$$

$$\hat{H}\Psi_0 = E_0\Psi_0$$

Born Oppenheimer Molecular Dynamics

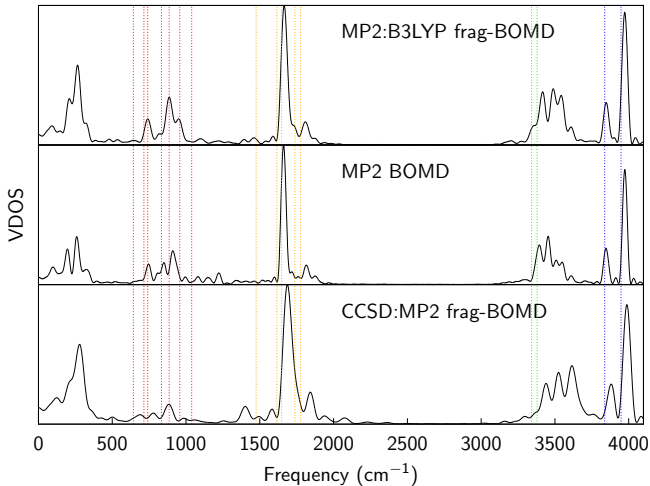


$$M_I \ddot{\mathbf{R}}_I(t) = -\nabla_I \min \left\{ \int \Psi^* \hat{H} \Psi d\mathbf{r} \right\}$$

$$\hat{H}\Psi_0 = E_0\Psi_0$$

$$I_V(\omega) = \lim_{T \rightarrow \infty} \int_{t=0}^{t=T} \exp(-i\omega t) \langle \mathbf{V}(0) \cdot \mathbf{V}(t) \rangle dt$$

Vibrational density of states

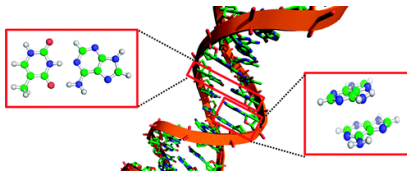


Outline

- 1 Introduction
 - More is difficult!
 - What can we do?
 - Tasks
- 2 PIE-ONIOM
 - What is PIE-ONIOM?
 - How is it implemented?
 - BOMD results
- 3 π interactions
 - Introduction
 - Dimers
 - Trimer and tetramer
- 4 DMRG
 - MPS
 - Extrapolation

Introduction

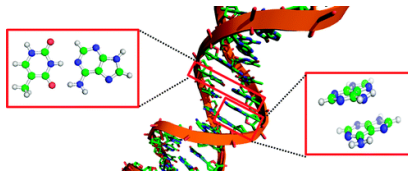
- Weak but important: found in proteins, DNA molecules, drugs, enzymes, etc., govern processes like condensation, crystallization, catalysis, solvation, etc.



Source: <http://www.chem.purdue.edu/Slipchenko/research/>

Introduction

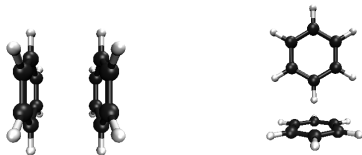
- Weak but important: found in proteins, DNA molecules, drugs, enzymes, etc., govern processes like condensation, crystallization, catalysis, solvation, etc.



Source: <http://www.chem.purdue.edu/Slipchenko/research/>

- Not amenable to mean field treatments; non-local, correlated methods necessary

Dimers



(a) Sandwich (D_{2h})

(b) T-shaped (C_{2v})

Dimers



(c) Sandwich (D_{2h})

(d) T-shaped (C_{2v})

Rigid monomers

Method dependence

$$E_{int} = E(AB; AB) - E(A; A) - E(B; B)$$

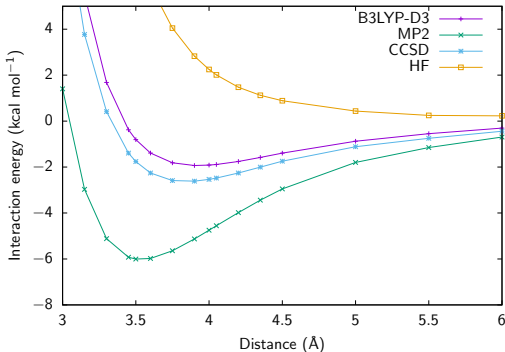


Figure: Sandwich Dimer PES scans (using aug-cc-pVDZ basis set, not CP corrected)

Basis dependence

$$E_{int}^{CP} = E(AB; AB) - E(A; AB) - E(B; AB)$$

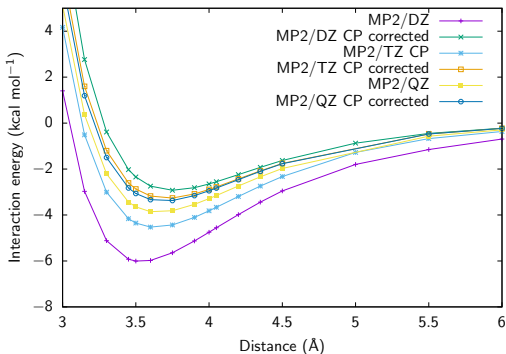


Figure: Sandwich Dimer PES scans ('XZ' denotes aug-cc-pVXZ basis set)

T-shaped

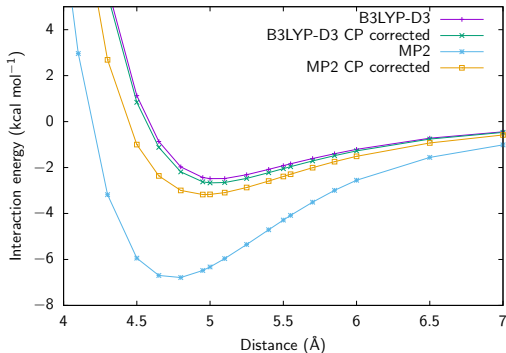


Figure: T-shaped Dimer PES scans (using aug-cc-pVDZ basis set)

Many body decomposition

Interaction energy of the cluster $f_1 f_2 \dots f_n$:

$$E_{int} = [E_1 + E_2 + \dots + E_n] - \sum_{i=1}^N E[f_i]$$

Many body decomposition

Interaction energy of the cluster $f_1 f_2 \dots f_n$:

$$E_{int} = [E_1 + E_2 + \dots + E_n] - \sum_{i=1}^N E[f_i]$$

$$E_1 = \sum_{i=1}^n E[f_i^*]$$

Many body decomposition

Interaction energy of the cluster $f_1 f_2 \dots f_n$:

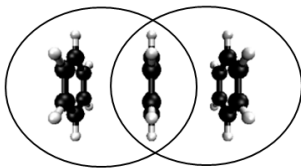
$$E_{int} = [E_1 + E_2 + \dots + E_n] - \sum_{i=1}^N E[f_i]$$

$$E_1 = \sum_{i=1}^n E[f_i^*]$$

$$E_2 = \sum_{i=1}^{n-1} \sum_{j>i}^n \Delta_2 E[f_i f_j^*]$$

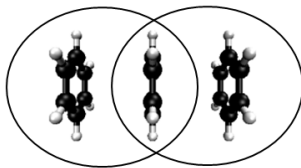
$$\Delta_2 E[f_i f_j^*] = E[f_i f_j^*] - (E[f_i^*] + E[f_j^*])$$

Trimer



- Tauer et al. showed that in benzene π clusters neighboring two-body terms dominate the interaction energy

Trimer



- Tauer et al. showed that in benzene π clusters neighboring two-body terms dominate the interaction energy
- PIE-ONIOM approach with overlapping dimer fragments could be effective

Trimer PES

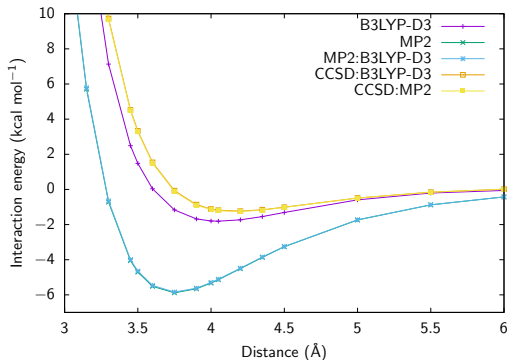


Figure: Trimer PES scans for the sandwich geometry

Tetramer

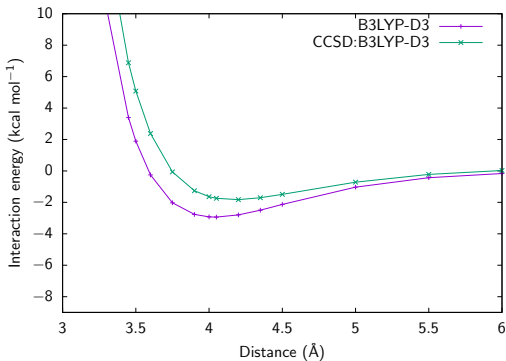


Figure: Tetramer PES scans for the sandwich geometry

Outline

- 1 Introduction
 - More is difficult!
 - What can we do?
 - Tasks
- 2 PIE-ONIOM
 - What is PIE-ONIOM?
 - How is it implemented?
 - BOMD results
- 3 π interactions
 - Introduction
 - Dimers
 - Trimer and tetramer
- 4 DMRG
 - MPS
 - Extrapolation

Matrix product states

- A general state of a multi-electron system can be written as

$$|\psi\rangle = \sum_{n_{j\sigma}} C_{n_{1\uparrow}n_{1\downarrow}\dots n_{L\uparrow}n_{L\downarrow}} |n_{1\uparrow}n_{1\downarrow}\dots n_{L\uparrow}n_{L\downarrow}\rangle$$

Matrix product states

- A general state of a multi-electron system can be written as

$$|\psi\rangle = \sum_{n_{j\sigma}} C_{n_{1\uparrow}n_{1\downarrow}\dots n_{L\uparrow}n_{L\downarrow}} |n_{1\uparrow}n_{1\downarrow}\dots n_{L\uparrow}n_{L\downarrow}\rangle$$

- Following multiple SVD's

$$C_{n_{1\uparrow}n_{1\downarrow}\dots n_{L\uparrow}n_{L\downarrow}} = \sum_{\alpha_1, \dots, \alpha_{L-1}} A[1]_{\alpha_1}^{n_{1\uparrow}n_{1\downarrow}} A[2]_{\alpha_1; \alpha_2}^{n_{2\uparrow}n_{2\downarrow}} \dots A[2]_{\alpha_{L-1}}^{n_{L\uparrow}n_{L\downarrow}}$$

Matrix product states

- A general state of a multi-electron system can be written as

$$|\psi\rangle = \sum_{n_{j\sigma}} C_{n_{1\uparrow}n_{1\downarrow}\dots n_{L\uparrow}n_{L\downarrow}} |n_{1\uparrow}n_{1\downarrow}\dots n_{L\uparrow}n_{L\downarrow}\rangle$$

- Following multiple SVD's

$$C_{n_{1\uparrow}n_{1\downarrow}\dots n_{L\uparrow}n_{L\downarrow}} = \sum_{\alpha_1, \dots, \alpha_{L-1}} A[1]_{\alpha_1}^{n_{1\uparrow}n_{1\downarrow}} A[2]_{\alpha_1; \alpha_2}^{n_{2\uparrow}n_{2\downarrow}} \dots A[2]_{\alpha_{L-1}}^{n_{L\uparrow}n_{L\downarrow}}$$

- DMRG variationally optimizes this matrix product with truncated virtual dimensions

Extrapolation

- Large virtual dimension (D) prohibitive

Extrapolation

- Large virtual dimension (D) prohibitive

-

$$E_D - E_{FCI} = C W_D^{disc}$$

Energy is a linear functional on the reduced density matrix

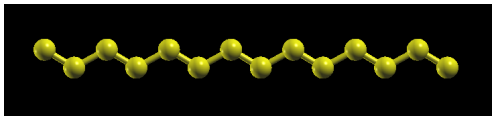
Extrapolation

- Large virtual dimension (D) prohibitive

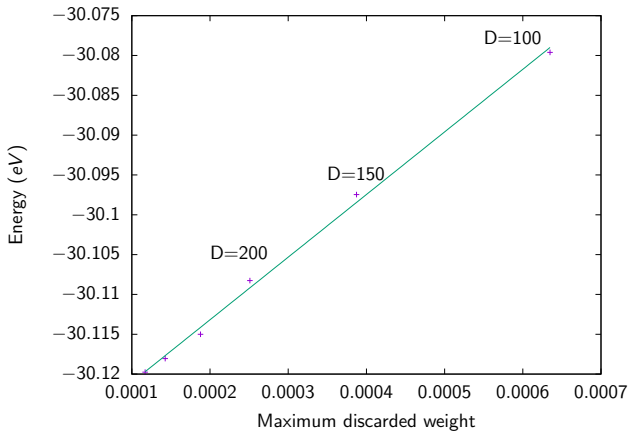
-

$$E_D - E_{FCI} = C W_D^{disc}$$

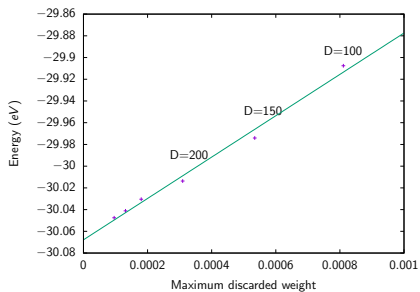
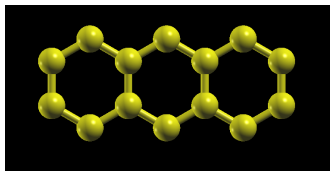
Energy is a linear functional on the reduced density matrix



TPA



Acene



TZGND2

