

# Large-scale coupled cluster calculations on supermolecular wires

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## Chemistry :

- Modeling's and simulations integrated part of experimental investigation
- Interplay between theory and experiment more important in the future

### Example : *Molecular manufacturing*

- Synthesize materials of desired properties by manipulating individual atoms
- Embraces the potential of altering the future of technology

# Foresight Nanotech Institute and Battelle

Report 2007 : Technology road map for productive Nano systems

p.17 “*Extending the **scale, scope, and accuracy** of atomistic modeling techniques is **a high priority** and can greatly facilitate atomically precise technologies design and implementation*”

Today's modeling on Nano systems use models of low accuracy

- Density functional theory, Force field

p.152 ‘*While quantum mechanical methods exist that approach the absolute limits of accuracy, the use of these methods is currently limited to diminutive chemical systems (< 20 atoms)*’

**What have the high accuracy calculation on small systems shown ?**

## Small Molecules: *Accomplishments using high accuracy models*

- Calculations changed how many experimental investigation are carried out,  
*A theoretical investigation often precede an experimental*
- Calculations broadening the understanding of the investigated phenomena
- In some cases calculations have replaced experiment :  
*Who would nowadays try to determine equilibrium geometries for small molecules experimentally?*

## Large molecules: *Challenges for high accuracy models*

- The scaling in conventional implementations prohibit applications
- Restructure calculations for lower scaling
- Even low scaling algorithms require large computational resources
- Development of low scaling methods must go hand in hand with access to increased computational resources
- Expect similar development with respect to interplay with experiment as for small molecules

# Development of code for the future

## Theoretical challenges

- Restructure high scaling models to become lower scaling
- The low scaling models have to be massively parallel
- Parallelism at several levels (Coarse, medium and fine grained)

## Challenges in connection with implementations

- Memory available
- Data traffic
- Many many more, refer to next talk by P. Ettenhuber

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Testing, testing and testing !!!!!

## Granting agency :

- Which chemical problem can you solve (using available methods)
- Equally important: Development of the methods for the future

For the future : *Close contact between developers of tomorrows code and tomorrows supercomputer software and hardware*

- Open new horizon we have not even dreamed about

# 30 years of research in electronic structure theory for small molecules have shown

a) Coupled cluster theory is the method of choice

b) Hierarchies of coupled models with improved accuracy

MP2 : Second order Møller Plesset theory ,  $N^5$  scaling

CCSD : Coupled cluster singles and doubles model ,  $N^6$  scaling

CCSD(T) : CCSD with perturbative triples corrections ,  $N^7$  scaling

(N denote system size)

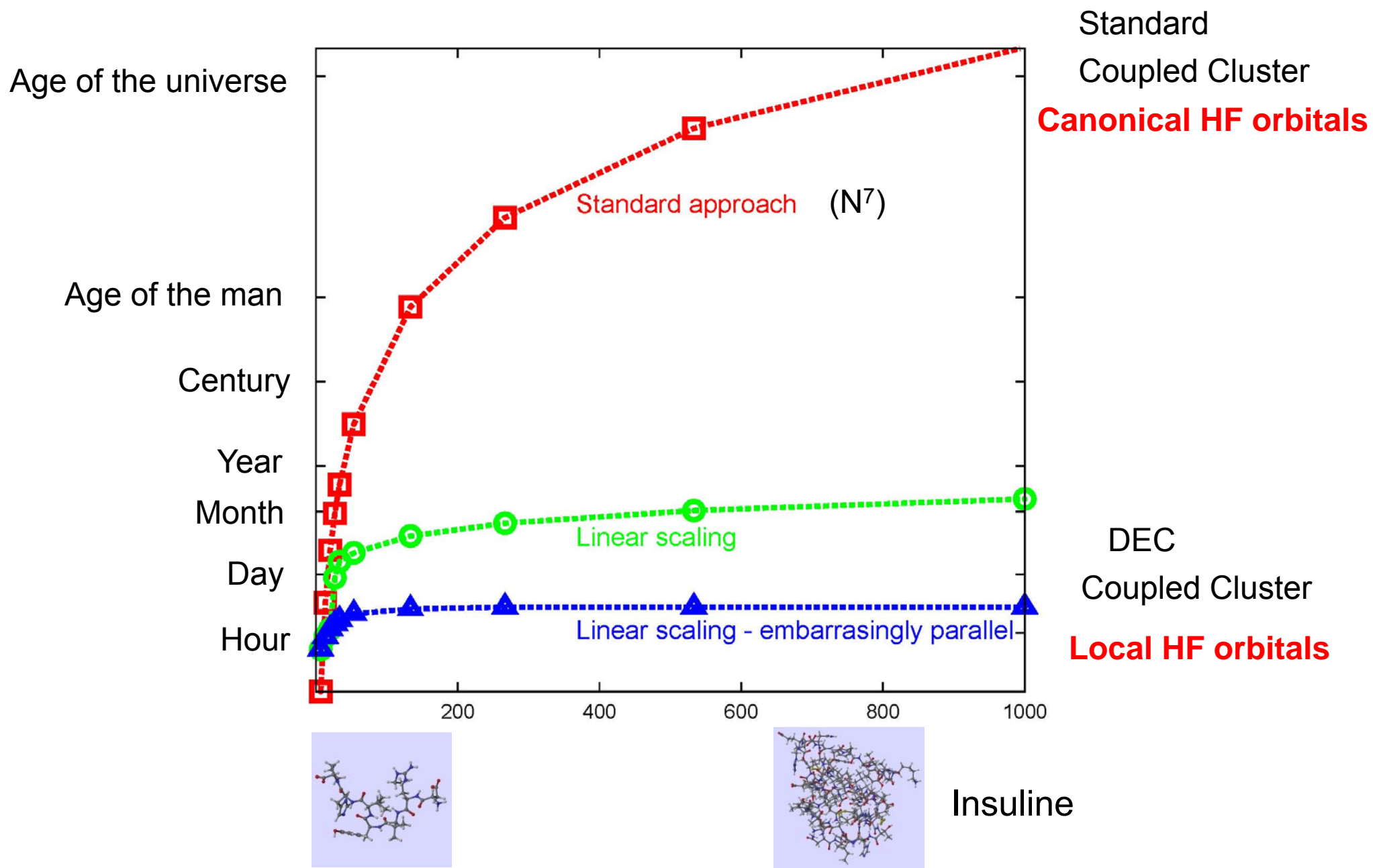
***CCSD(T) is the golden standard of quantum chemistry***

Many molecular properties are described to experimental accuracy or better e.g. equilibrium geometries and reaction enthalpies

***Problem: The scaling in standard implementations***



# Scaling in Coupled Cluster theory CCSD(T)



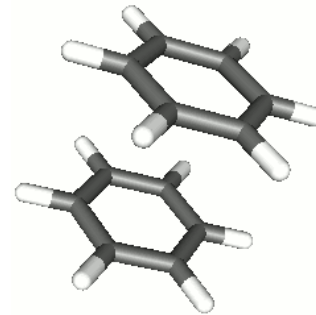
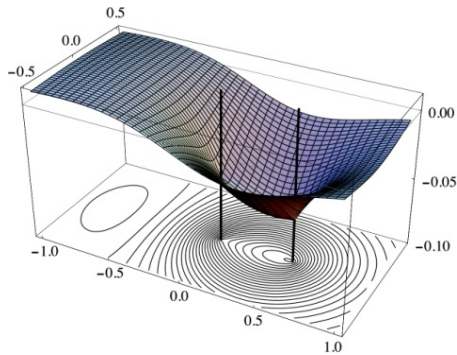
# Coupled cluster wave function calculation

*Hartree-Fock calculation determines the reference state*

- Electrons move in an averaged field of the others
- Long range potential described to high accuracy
- **Give local orbitals for both occupied and virtual space**

*Coupled cluster calculation describes local electron correlation effects*

- Coulomb hole, Short range
- Dispersion forces,  $R^{-6}$



Local phenomena described in **local basis** give linear scaling

The divide-expand-consolidate (DEC) coupled cluster method

# The DEC coupled cluster (CC) method

## Assumption

- **Local occupied** and **virtual** HF orbitals can be determined

## Method

- Assign **local occupied** and **virtual** HF orbitals to atomic sites,  $P, Q, \dots$
- Partition correlation energy into atomic fragment  $E_P$  and pair  $\Delta E_{PQ}$  energies

Replace  $(i,j)$  or  $(a,b)$  summations in correlation energy with  $P$  and  $PQ$  summations

- Evaluate  $E_P$  and  $\Delta E_{PQ}$  from small orbital fragment spaces adjusted to give the energies to a predefined fragment optimization threshold (FOT)

## Features

- Adjustable orbital spaces give error control for energies and amplitudes
- Black box, linear scaling, and embarrassingly parallel

# Precision of Coupled Cluster calculation

## *Standard Coupled Cluster*

- Residual norm ( **$R$** ) of amplitude equation

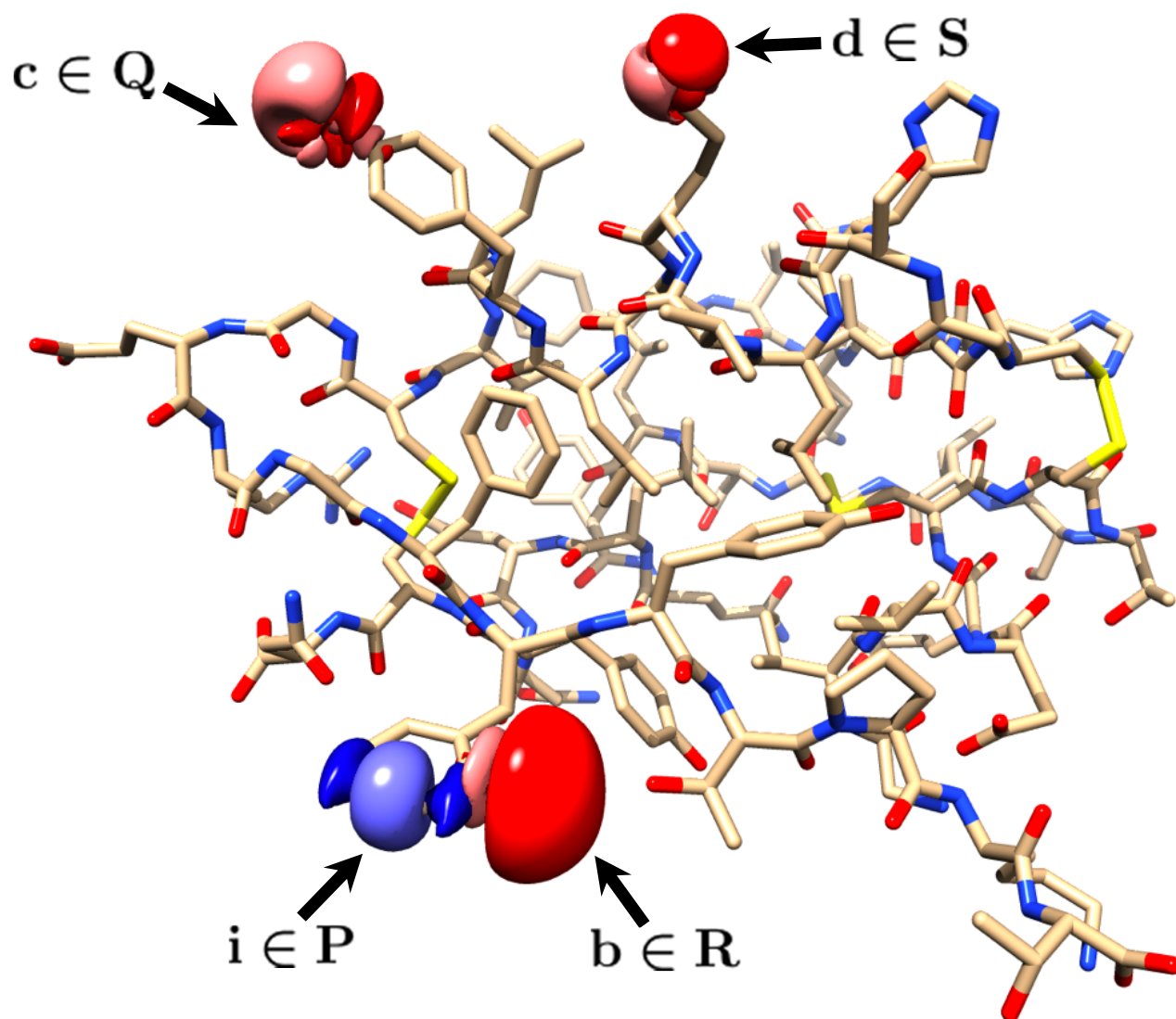
## *DEC Coupled Cluster* ( new strategy which exploit locality efficiently)

- Fragment optimization threshold ( **$FOT$** ) for atomic fragment energies

## *Standard and DEC Coupled Cluster on par with respect to precision*

- Single parameter (  **$R$**  or  **$FOT$** ) define the precision of energy, density, ...

# Determine local Hartree-Fock orbitals



Occupied orbitals:

$i, j, k, l$

Virtual orbitals:

$a, b, c, d$

Atomic sites:

$P, Q, R, S$

***Assign orbitals  
to atomic sites***

INSULIN MOLECULE (HYDROGENS OMITTED)

# Correlation energy MP2 and CCSD

$$E_{corr} = \sum_{ijab} (t_{ij}^{ab} + t_i^a t_j^b) (2g_{iajb} - g_{ibja})$$

## CCSD(T) energy correction

$$\Delta E^{CCSD(T)} = \sum_{AI} {}^*t_I^A {}^*T_I^A + \frac{1}{4} \sum_{IJAB} {}^*t_{IJ}^{AB} {}^*T_{IJ}^{AB}$$

## DEC Strategy

*Assign local orbitals to atomic sites  $P, Q, \dots$*

*Replace summation over two occupied  $i, j$  (virtual  $a, b$ ) orbitals with summations over sites  $P$  and pair sites  $P, Q$  and summations over orbitals belonging to these sites*

# Occupied space partitioning of $E_{corr}$

*No approximation*      Coulomb hole      Dispersion energy      Quadratic scaling

$$E_{corr} = \sum_P E_P^O + \sum_{P>Q} \Delta E_{PQ}^O$$

Occupied atomic fragment energy

$$E_P^O = \sum_{\substack{ij \in P \\ ab}} \left( t_{ij}^{ab} + t_i^a t_j^b \right) \left( 2g_{iajb} - g_{ibja} \right)$$

Occupied pair interaction energy

$$\Delta E_{PQ}^O = \sum_{\substack{i \in P, j \in Q \\ ab}} \left( t_{ij}^{ab} + t_i^a t_j^b \right) \left( 2g_{iajb} - g_{ibja} \right) + \quad P \leftrightarrow Q \text{ term}$$

No approximations have so far been made, only a reorganization

# Atomic fragment energy $E_P$

$$E_P^O = \sum_{ij \in P} \sum_{ab \in [P]} \left( t_{ij}^{ab} + t_i^a t_j^b \right) \left( 2g_{iajb} - g_{ibja} \right)$$

Determine  $[P]$  in a black box manner  
such that errors in  $E_P$  is smaller than  
**Fragment Optimization Threshold (FOT)**

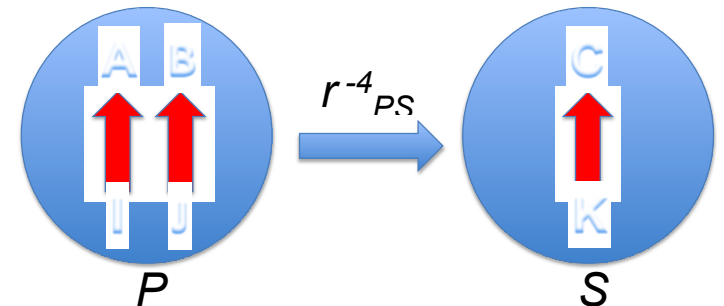
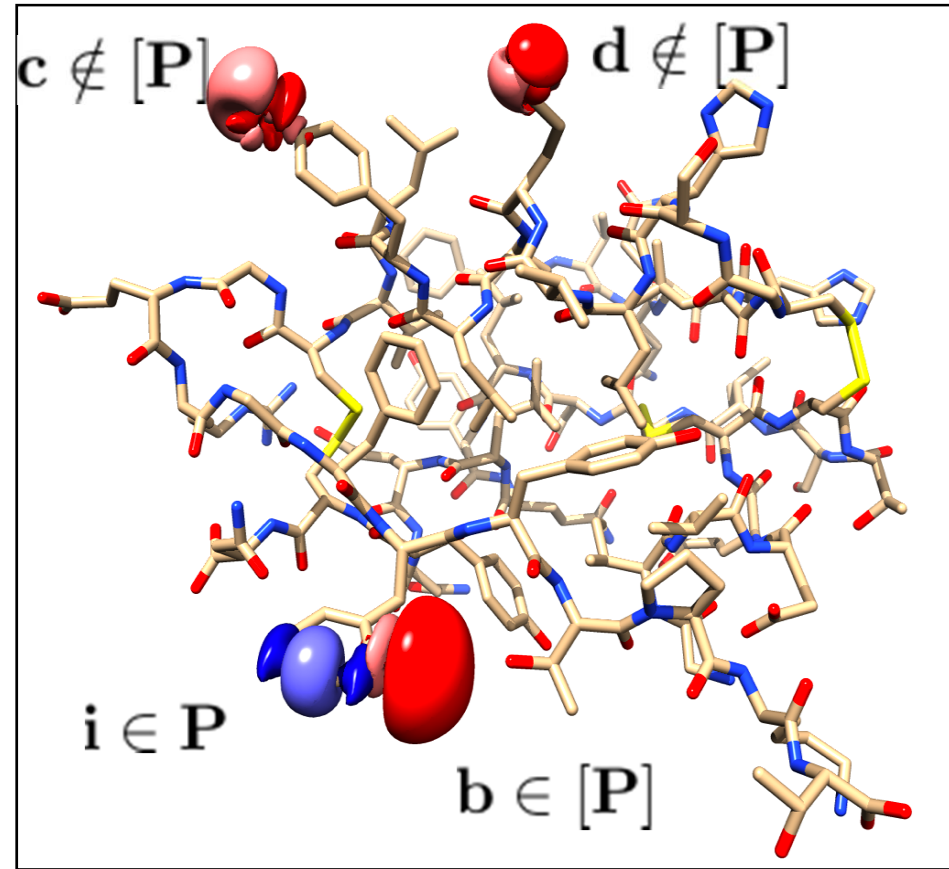
## CCSD and MP2

Charge distributions in integrals  
determine distance decay from site P

$$g_{iajb} \quad ; \quad i, j \in P \quad , \quad a, b \in [P]$$

## CCSD(T)

Charge distributions together with third  
party excitations to site S determine  
distance decay from site P





# Occupied space partitioning of $E_{corr}$

Occupied atomic fragment energy

$$E_P^O = \sum_{\substack{ij \in P \\ ab \in [P]}} \left( t_{ij}^{ab} + t_i^a t_j^b \right) \left( 2g_{iajb} - g_{ibja} \right)$$

(atomic fragment orbital space)

Occupied atomic pair interaction energy

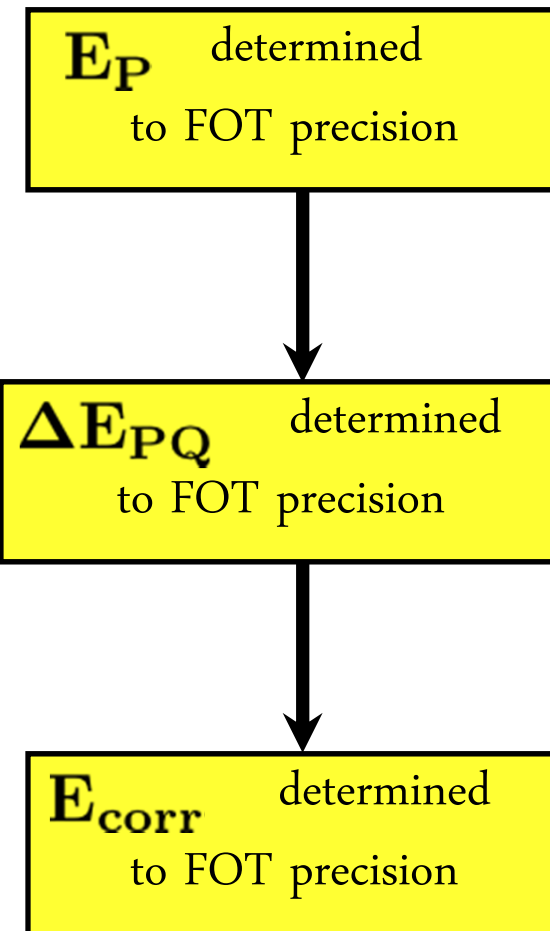
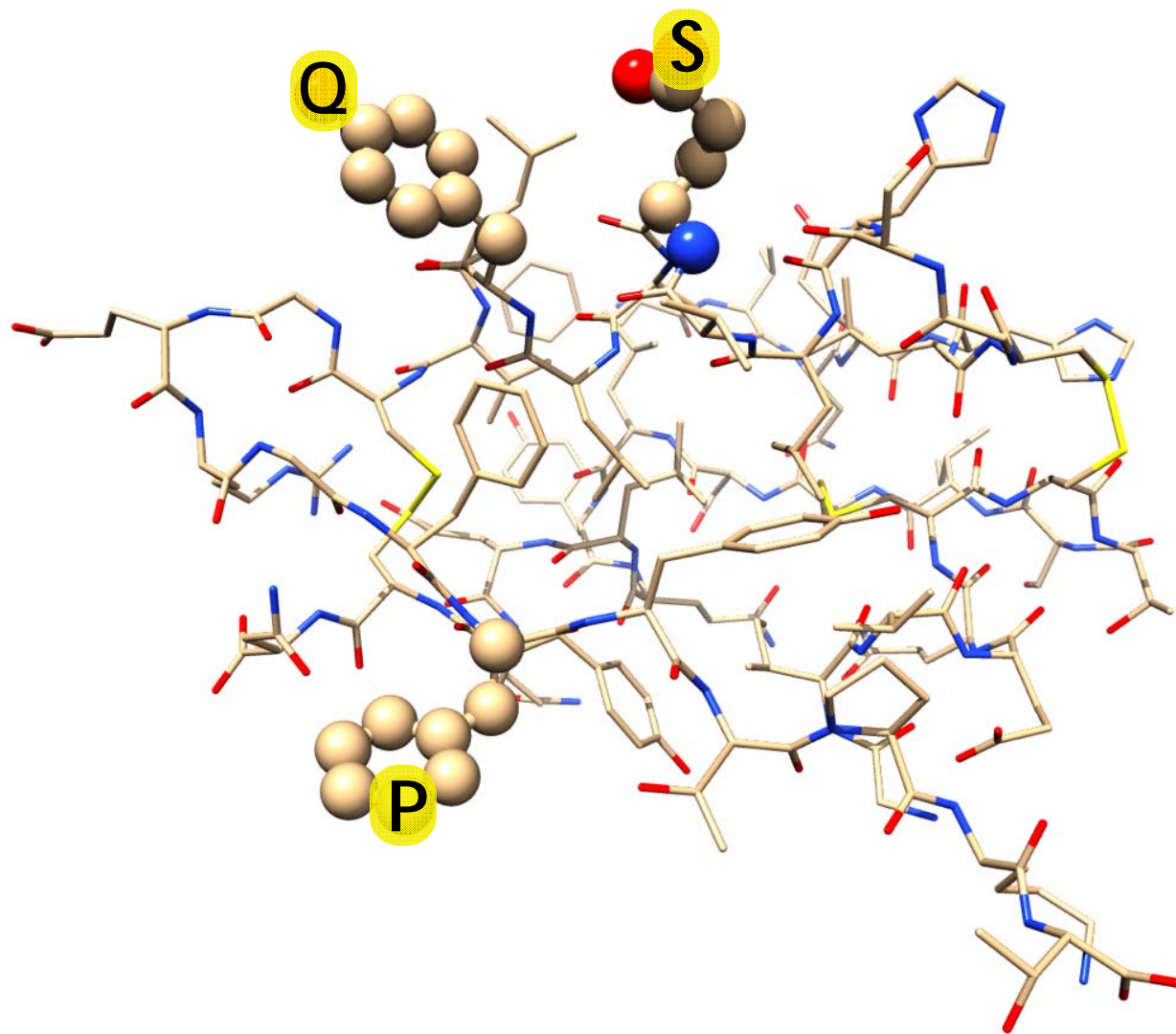
$$\Delta E_{PQ}^O = \sum_{\substack{i \in P, j \in Q \\ ab \in [P] \cup [\Theta]}} \left( t_{ij}^{ab} + t_i^a t_j^b \right) \left( 2g_{iajb} - g_{ibja} \right) + \quad P \leftrightarrow Q \text{ term}$$

(union of atomic fragment orbital spaces)

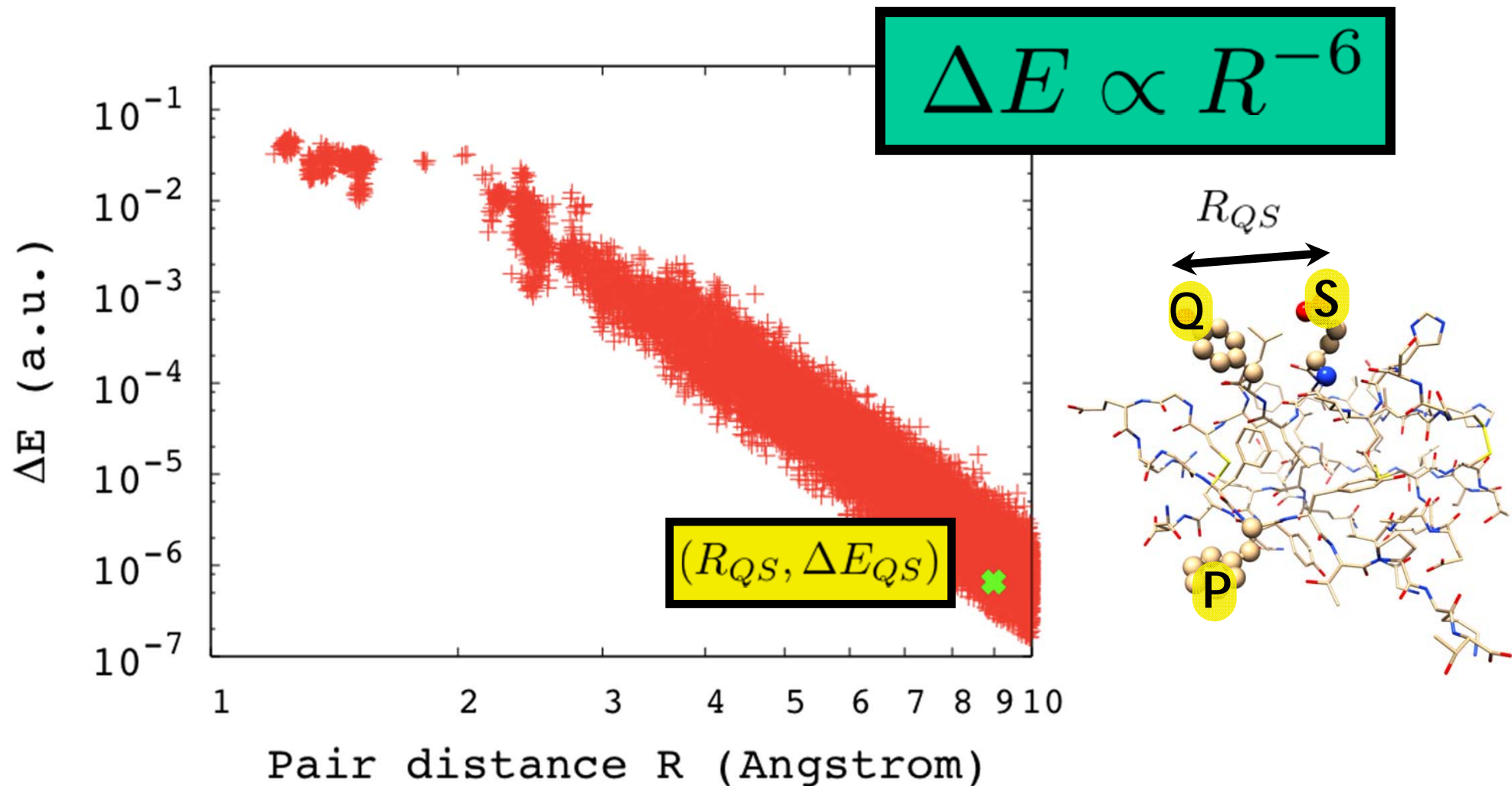
## Insulin - three examples of atomic fragments

$$E_{\text{corr}} = \sum_P E_P + \sum_{P>Q} \Delta E_{PQ}$$

Quadratic scaling



# Insulin - pair interaction energies $\Delta E_{PQ}$



Pairs separated by more than  $\sim 10$  Å can be neglected  
without affecting the precision of the calculation

**Linear scaling  
algorithm**

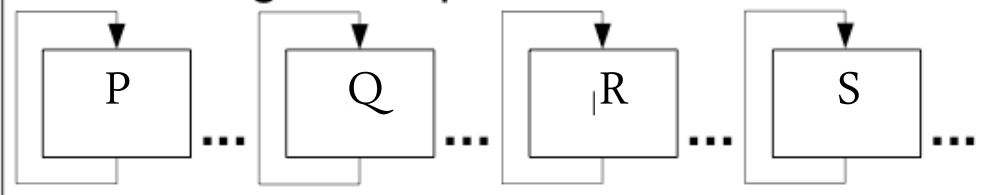
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# Summary of the DEC scheme

Local HF orbitals

Atomic fragment optimization

$N$  atomic fragments

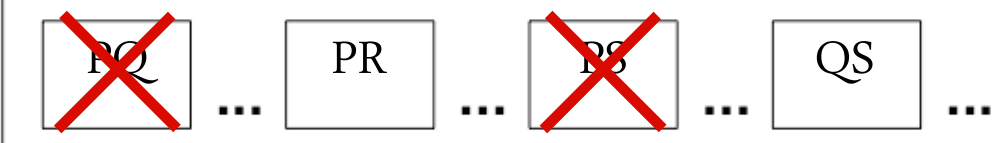


Pair fragments

(pair distance  $< 10\text{\AA}$ )

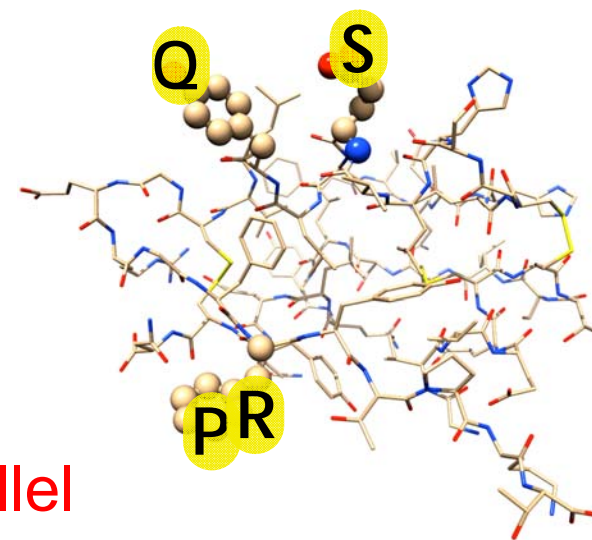
Formally:  $N(N-1)/2$  pair fragments

Use **cut-off**:  $\text{const} \cdot N$  pair fragments

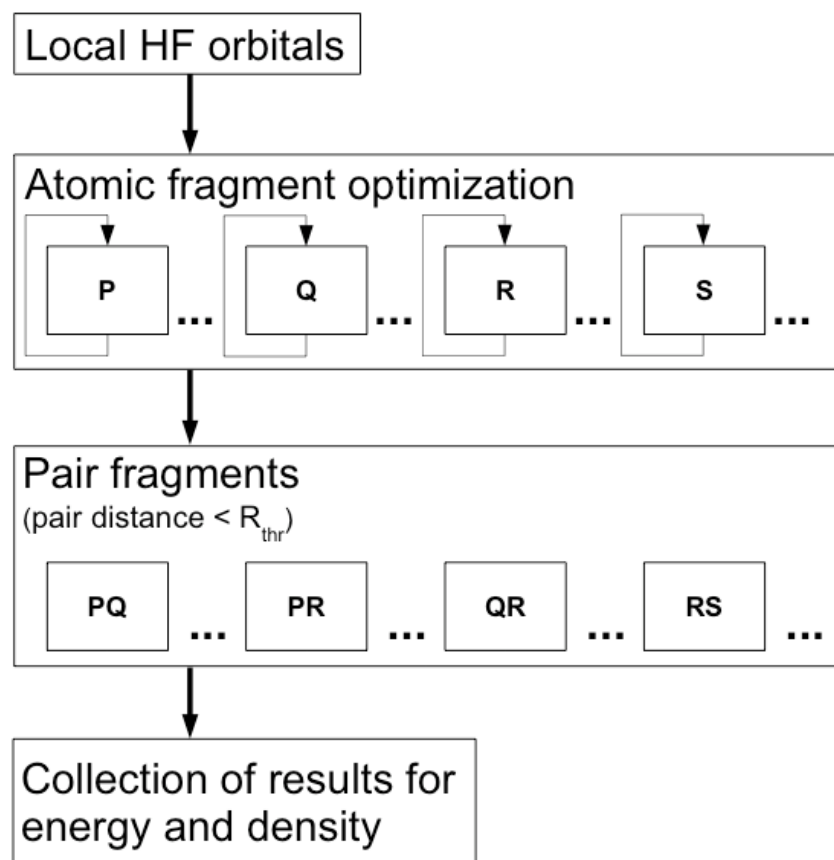


Collection of results for energy and density

DEC is linear scaling and embarrassingly parallel



# Parallelism in DEC calculation



Parallelity at three levels

***Coarse grained parallelism***

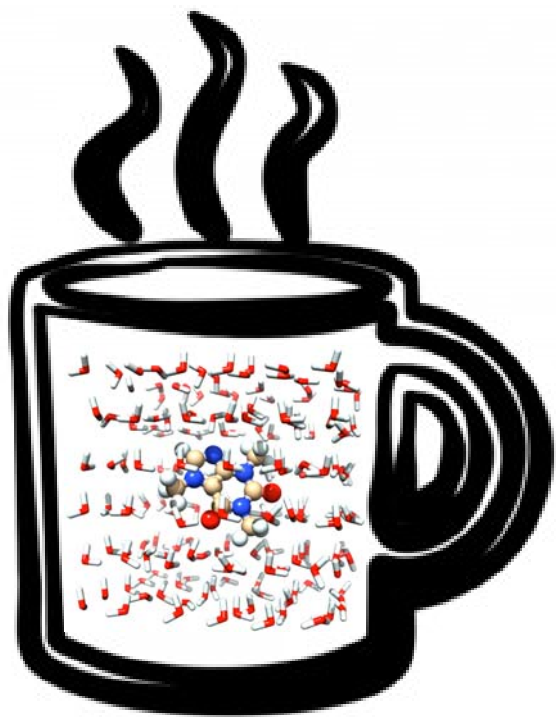
All fragment calculations  $E_P$  and  $\Delta E_{PQ}$  carried out independently

***Medium and fine grained parallelism***

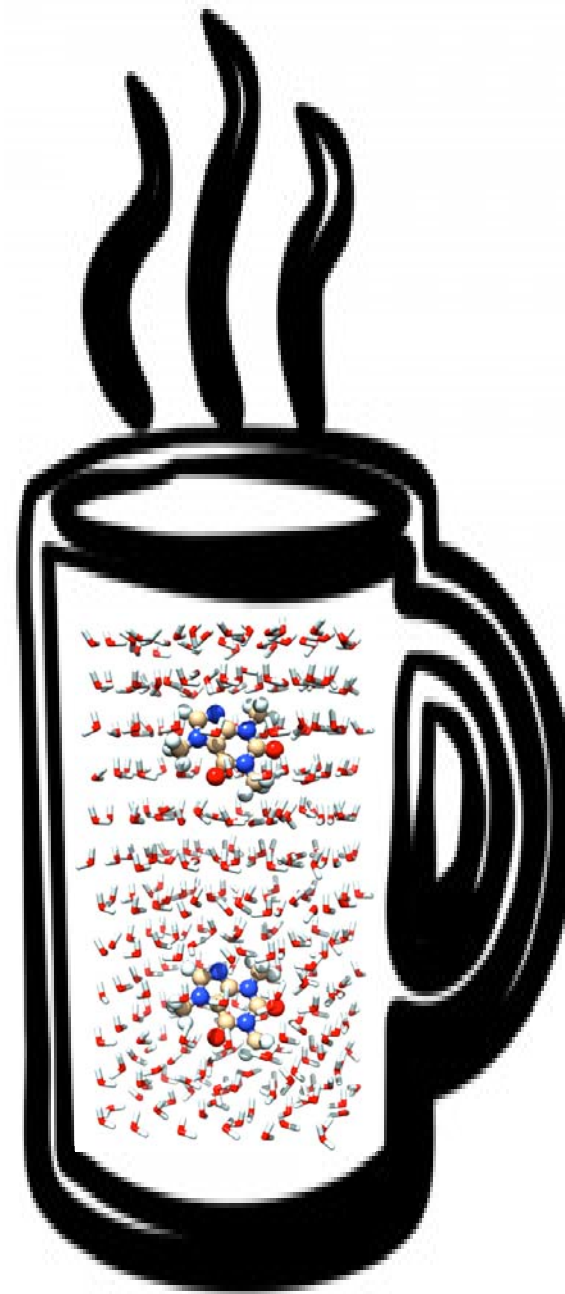
Individual fragment calculations parallelized at two levels (MPI and OpenMP)  
( as in conventional implementations )

# “A nano-cup of coffee...”

- Calculate MP2 correlation energy and density using cc-pVDZ basis.
- Nanospresso (system 1): 528 atoms (4278 BF.)
- Nanospresso Doppio (system 2): 1056 atoms (8556 BF.)



Nanospresso



Nanospresso doppio

# Time to solution (TTS)

System	#fragments	#nodes	TTS(hours)
1	7136	5890	1.66
1	7136	11780	0.93
2	16151	5890	4.49
2	16151	11780	2.37

- Calculations on Titan, Oak Ridge National Laboratory\* using 8 OpenMP threads per MPI process. Nodes refer to NUMA nodes

## ■ Scaling with system size?

$$\#frags(2) / \#frags(1) = 2.26 \quad (\text{ideal})$$

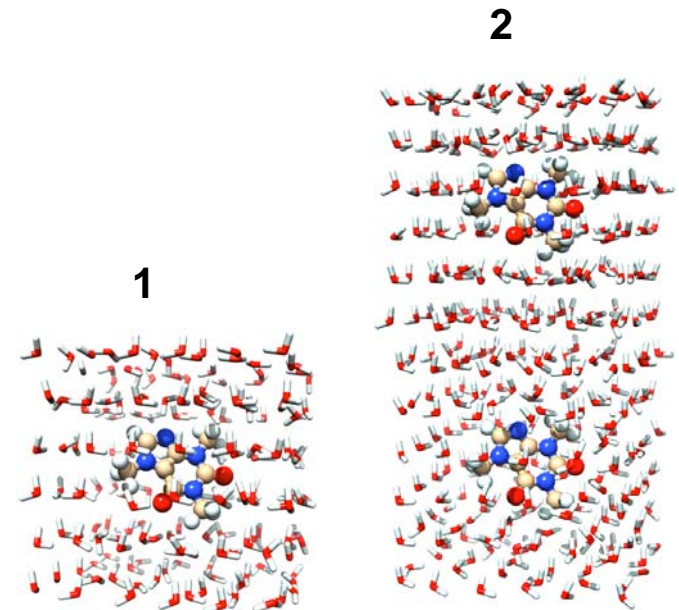
$$TTS(2) / TTS(1) = 2.70 \quad (5890 \text{ nodes})$$

$$TTS(2) / TTS(1) = 2.55 \quad (11780 \text{ nodes})$$

## ■ Parallel (strong) scaling (ideal: 0.5)

$$TTS(11780) / TTS(5890) = 0.56 \quad (\text{system 1})$$

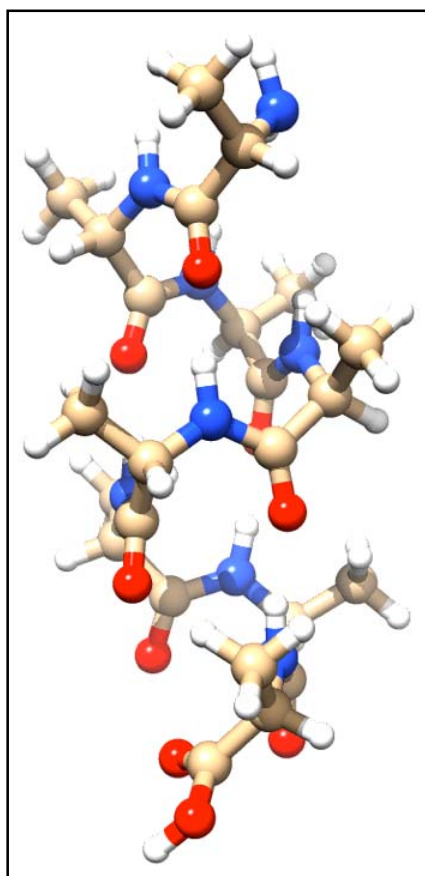
$$TTS(11780) / TTS(5890) = 0.53 \quad (\text{system 2})$$





# Test molecule: Energy errors vs. FOT

**MP2/cc-pVDZ calculation: Errors in DEC correlation energies (a.u.)**



**Alanine(8)**

FOT	Occupied energy error	Virtual energy error	Lagrangian energy error	Average relative error*	Average percentage of $E_{\text{corr}}$
$10^{-2}$	$1.38 \cdot 10^{-1}$	$2.51 \cdot 10^{-1}$	$2.00 \cdot 10^{-1}$	$3.1 \cdot \text{FOT}$	96.9%
$10^{-3}$	$2.83 \cdot 10^{-2}$	$2.76 \cdot 10^{-2}$	$1.87 \cdot 10^{-2}$	$3.9 \cdot \text{FOT}$	99.6%
$10^{-4}$	$3.20 \cdot 10^{-3}$	$1.87 \cdot 10^{-3}$	$1.95 \cdot 10^{-3}$	$3.6 \cdot \text{FOT}$	99.96%
$10^{-5}$	$2.69 \cdot 10^{-4}$	$2.56 \cdot 10^{-4}$	$1.78 \cdot 10^{-4}$	$3.7 \cdot \text{FOT}$	99.996%
$10^{-6}$	$3.25 \cdot 10^{-5}$	$1.09 \cdot 10^{-5}$	$2.01 \cdot 10^{-5}$	$3.3 \cdot \text{FOT}$	99.9997%



# DEC MP2 insulin calculation: Energy

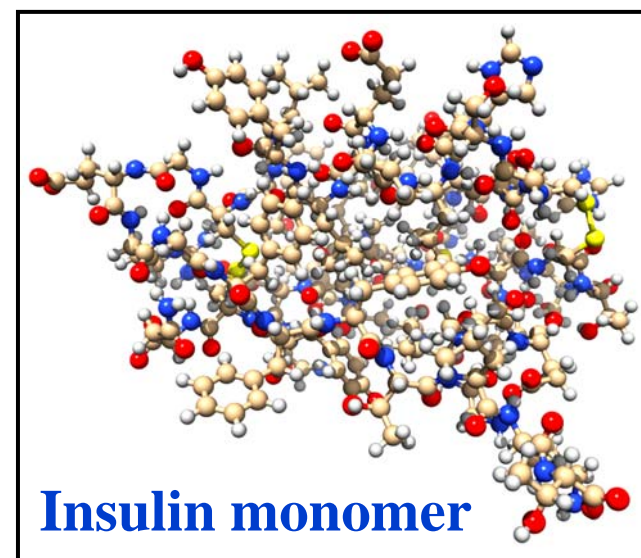
cc-pVDZ: 7604 basis functions

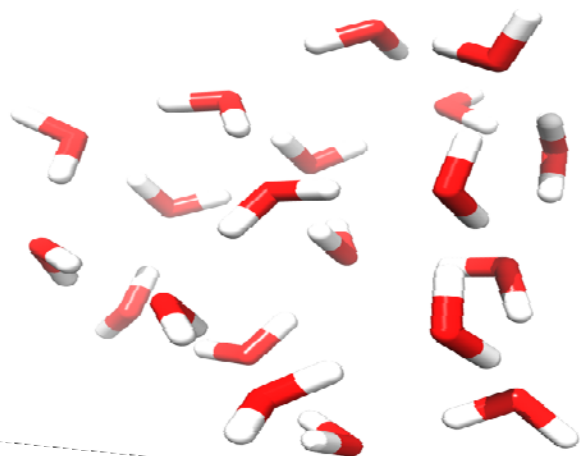
FOT	Occupied correlation energy	Virtual correlation energy	Lagrangian correlation energy	Estimated absolute error*	Estimated relative error**	Estimated percentage of $E_{\text{corr}}$
$10^{-4}$	-61.755	-61.783	-61.763	0,028	$4.5 \cdot \text{FOT}$	99.95%

- ✱ Consistent with errors for smaller test molecules
- ✱ E.g. alanine(8) for  $\text{FOT}=10^{-4}$ : Relative error =  $3.6 \cdot \text{FOT}$  (99.96% of  $E_{\text{corr}}$ )

DEC is a black box method:

The relative errors are system-independent, regardless of system size





## 20 water cluster

6-31G\*

FOT	MP2	CCSD	(T)	CCSD(T)
$10^{-3}$	$2.1 \cdot 10^{-2}$	$1.4 \cdot 10^{-2}$	$7.0 \cdot 10^{-3}$	$2.2 \cdot 10^{-2}$
$10^{-4}$	$2.3 \cdot 10^{-3}$	$4.4 \cdot 10^{-4}$	$2.4 \cdot 10^{-3}$	$2.8 \cdot 10^{-3}$
$10^{-5}$	$2.3 \cdot 10^{-4}$	$4.7 \cdot 10^{-5}$	$3.1 \cdot 10^{-4}$	$3.6 \cdot 10^{-4}$
$10^{-6}$	$2.3 \cdot 10^{-5}$	$-3.5 \cdot 10^{-5}$	$9.0 \cdot 10^{-5}$	$5.5 \cdot 10^{-5}$

## 20 water cluster cc-pVTZ

FOT	CCSD	(T)	CCSD(T)
$10^{-3}$	$2.8 \cdot 10^{-2}$	$1.6 \cdot 10^{-2}$	$4.4 \cdot 10^{-2}$
$10^{-4}$	--	--	--

Largest conventional CCSD(T) calculation  $(\text{H}_2\text{O})_{20}$

E. Apra *et al.*, **SC09**, submission for Gordon Bell prize (2009)

**Also:**

***CCSD(T) calculation carried out on one “Nanospresso”  
with  $\text{FOT}=10^{-3}$ .***

***4278 basis functions on 3201 nodes in 4 hours and 20 minutes***

## *Conclusion for DEC*

- New strategy for CC energy and properties which explore locality efficiently
- Full system in terms of CC calculations on small fragments of total orbital space
- Linear scaling and massive parallel algorithm
- Full control of errors in energy and cluster amplitudes
- Black box method
- Performance (speed) depends on locality of HF orbitals
- DEC on par with standard CC method,  
FOT defines precision as residual norm in standard CC

# *Acknowledgements*

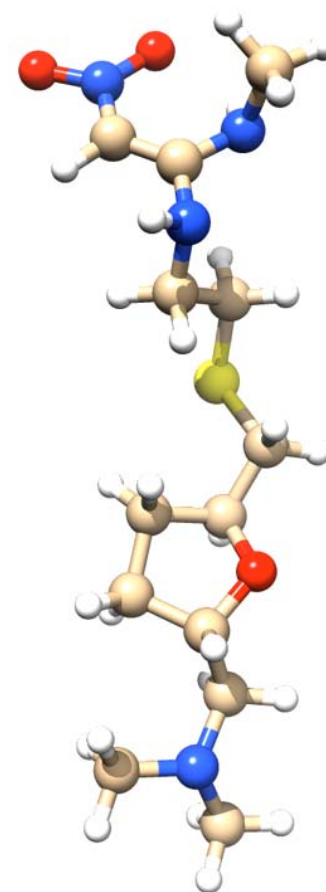
## Work supported by:

- US Department of Energy
- Oak Ridge Leadership Computing Facility
- European Research Council
- Danish Council for Independent Research – Natural Sciences
- Aarhus University

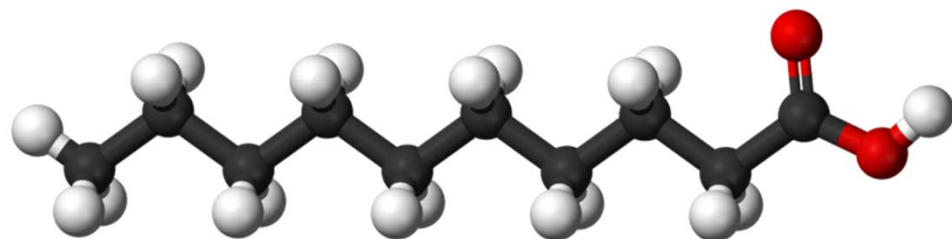
# DEC MP2 molecular gradient errors

cc-pVTZ basis, all errors are given in a.u.

FOT	Energy error	RMS error for gradient
$10^{-3}$	$9,8 \cdot 10^{-3}$	$1,5 \cdot 10^{-3}$
$10^{-4}$	$9.1 \cdot 10^{-4}$	$4,5 \cdot 10^{-4}$
$10^{-5}$	$6.9 \cdot 10^{-5}$	$2,7 \cdot 10^{-5}$
$10^{-6}$	$7.5 \cdot 10^{-6}$	$8,3 \cdot 10^{-6}$



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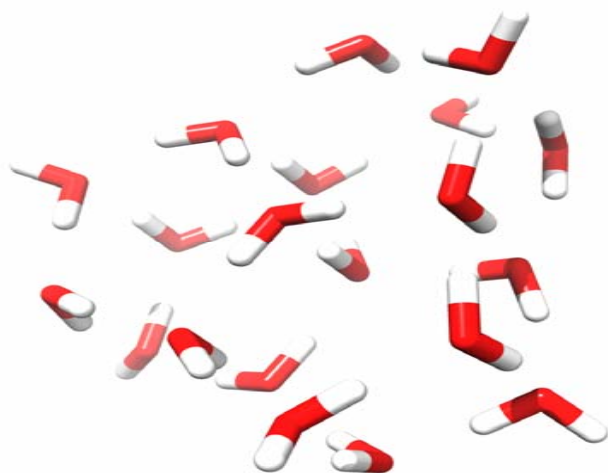
**Decanoic acid**

**6-31G\***

<b>FOT</b>	<b>MP2 occ</b>	<b>MP2 vir</b>	<b>CCSD occ</b>	<b>CCSD vir</b>	<b>CCSD(T) occ</b>	<b>CCSD(T) vir</b>
$10^{-3}$	$1.9 \cdot 10^{-2}$	$1.5 \cdot 10^{-2}$	$-2.0 \cdot 10^{-3}$	$-1.1 \cdot 10^{-2}$	$5.1 \cdot 10^{-3}$	$-4.4 \cdot 10^{-3}$
$10^{-4}$	$1.3 \cdot 10^{-3}$	$1.5 \cdot 10^{-3}$	$-3.8 \cdot 10^{-3}$	$-3.5 \cdot 10^{-3}$	$-1.6 \cdot 10^{-3}$	$-2.0 \cdot 10^{-3}$
$10^{-5}$	$1.2 \cdot 10^{-4}$	$1.4 \cdot 10^{-4}$	$-1.2 \cdot 10^{-3}$	$-1.3 \cdot 10^{-3}$	$-3.6 \cdot 10^{-4}$	$-9.0 \cdot 10^{-4}$
$10^{-6}$	$1.2 \cdot 10^{-5}$	$1.3 \cdot 10^{-5}$	$-3.3 \cdot 10^{-4}$	$-3.6 \cdot 10^{-4}$	$-1.1 \cdot 10^{-4}$	$-2.3 \cdot 10^{-4}$

Largest conventional CCSD(T) calculation (H<sub>2</sub>O)<sub>20</sub>

E. Apra *et al.*, **SC09**, submission for Gordon Bell prize (2009)



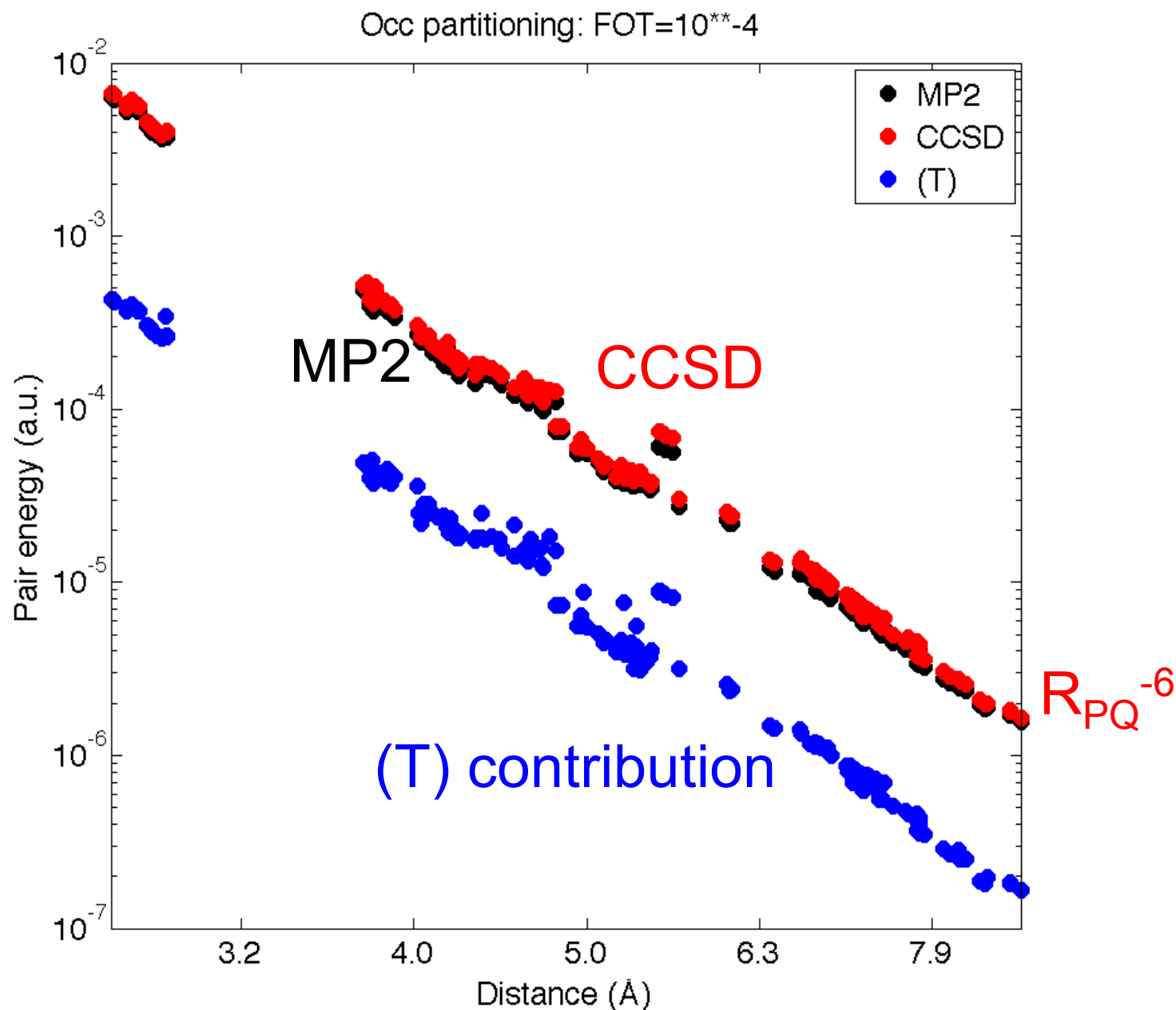
cc-pVTZ

Model Error	MP2	CCSD(T)
FOT=10 <sup>-3</sup>	1.5e-2	1.5e-2
FOT=10 <sup>-4</sup>	1.6e-3	4.6e-4



# MP2, CCSD, and CCSD(T) pair interaction energies

$(\text{H}_2\text{O})_{20}$   
cc-pVTZ



# DEC perspective

## MP2

- Energy, density, molecular gradient and geometry optimizer ***Done***
- F12 energies, Nuclear shifts, ... ***In progress***

## CCSD

- Energy implemented ***Ready to be tested for massive parallelism***
- Future work, density, molecular gradient, F12 energy ...

## CCSD(T)

- Energy implemented ***Ready to be tested for massive parallelism***
- Future work, density, molecular gradient, ...

# DEC CCSD(T) energy

$$\Delta E^{CCSD(T)} = \sum_{AI} {}^*t_I^A {}^*T_I^A + \frac{1}{4} \sum_{IJAB} {}^*t_{IJ}^{AB} {}^*T_{IJ}^{AB}$$

Two occupied and two virtual indices

$$\Delta E^{CCSD(T)} = \sum_P {}^{(T)}E_{\underline{P}} + \sum_{P>Q} {}^{(T)}E_{\underline{PQ}}$$

Triples fragment and pair fragment energies

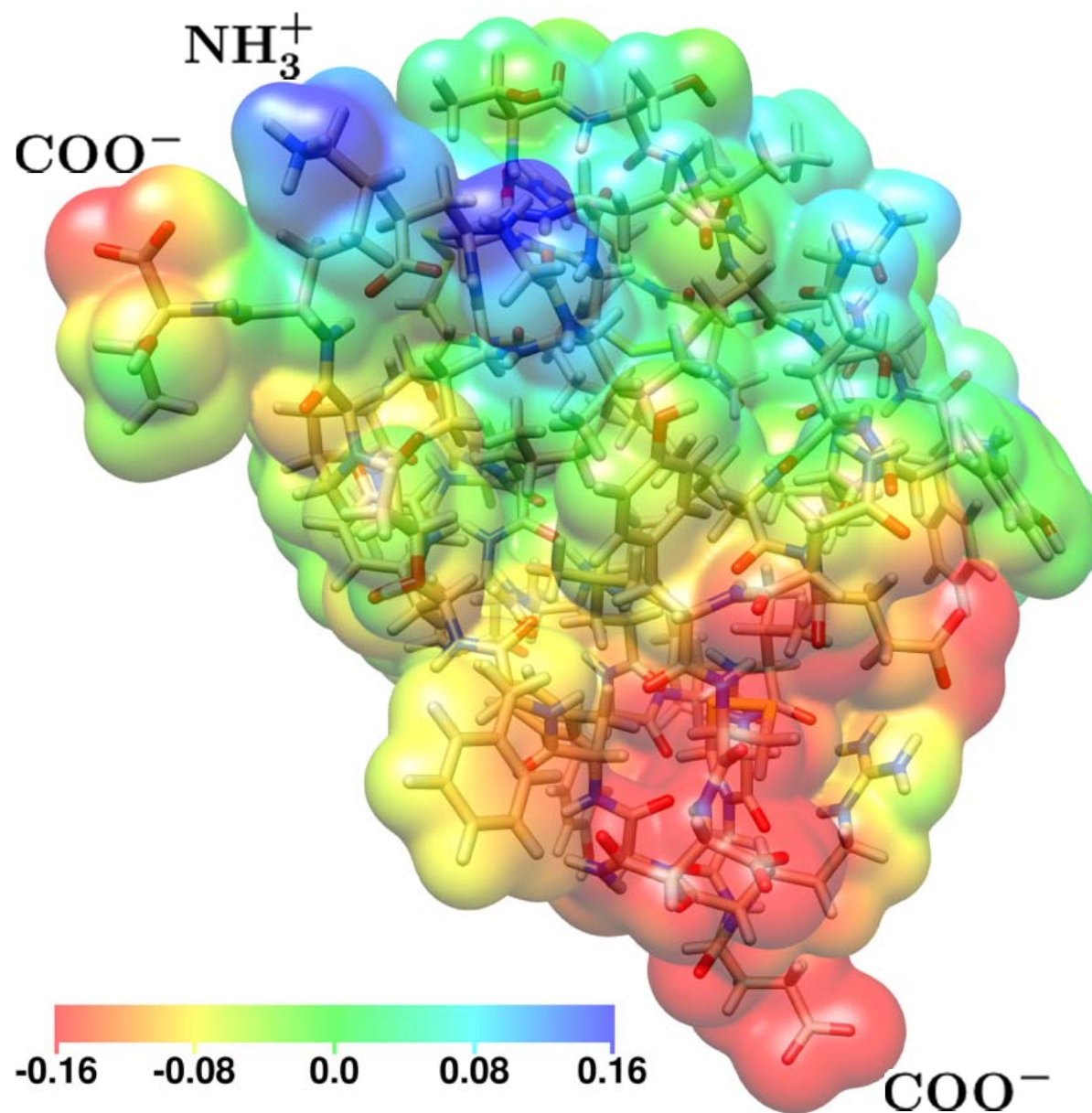
$${}^*T_I^A = \langle AI[\Phi, {}^*T_3] \| HF \rangle = \frac{1}{12} \sum_{\substack{CD \\ KL}} ({}^*t_{IKL}^{ACD} - 2 {}^*t_{LKI}^{ACD}) g_{KCLD}^a$$

$${}^*T_{IJ}^{AB} = \langle ABIJ[\Phi, {}^*T_3] \| HF \rangle = \frac{2}{3} \left[ \sum_{CDK} [g_{DKBC}^a {}^*t_{IJK}^{ACD} - g_{DKBC} {}^*t_{KJI}^{ACD}] - \sum_{CKL} [g_{LCKJ}^a {}^*t_{IKL}^{ABC} - g_{LCKJ} {}^*t_{LKI}^{ABC}] \right]$$

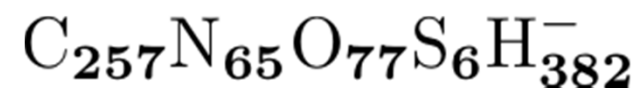
$$\mathcal{E}_{IJK}^{ABC} {}^*t_{IJK}^{ABC} = -P(IJK)P(ABC) \left[ \sum_D g_{CKBD}^a {}^*t_{IJ}^{AD} - \sum_L g_{CKLJ}^a {}^*t_{IL}^{AB} \right]$$

$$P(QRS)f(Q,R,S) = f(Q,R,S) + f(S,Q,R) + f(R,S,Q)$$

## DEC MP2 electrostatic potential for insulin



Insulin monomer:



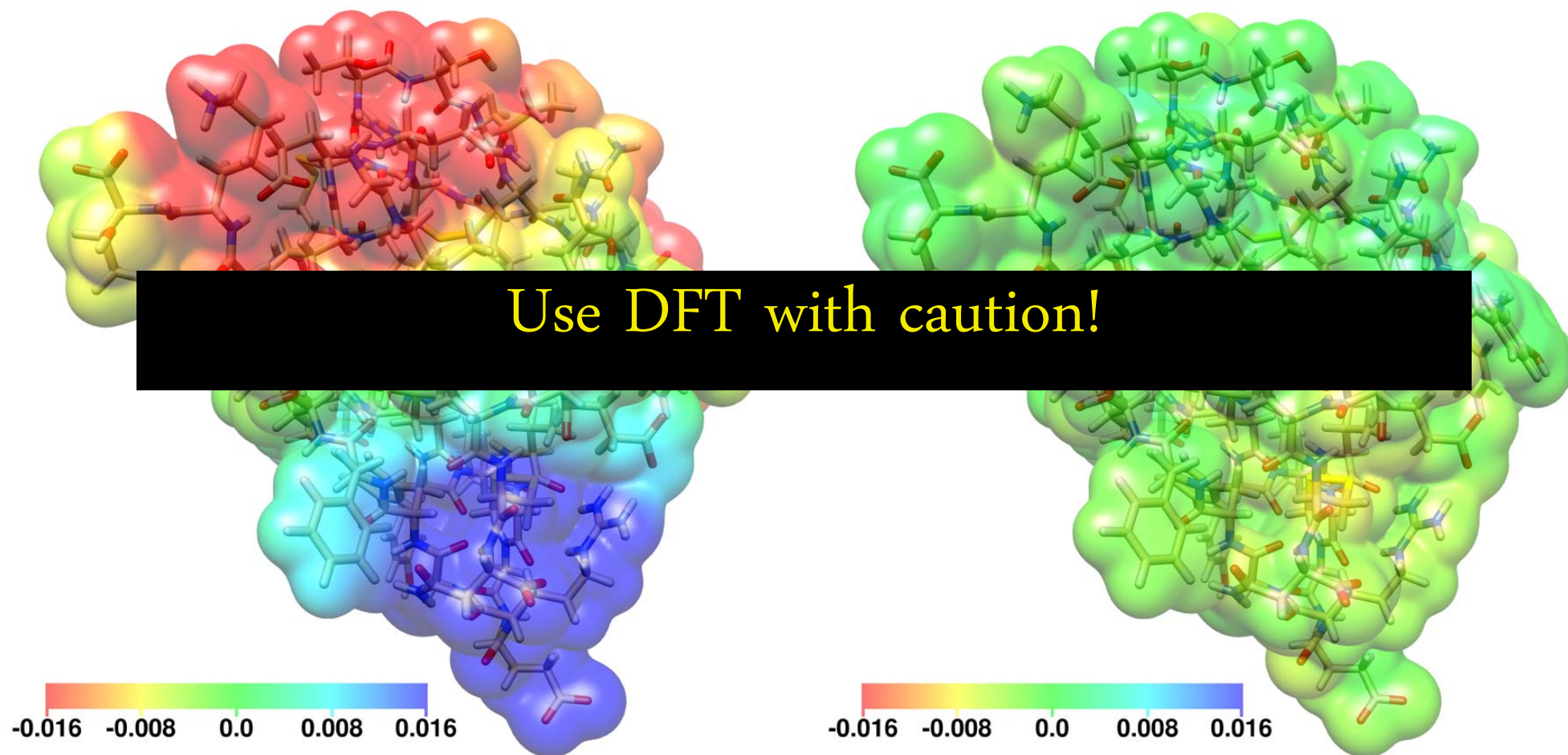
Red/blue regions indicate high/low potential energy for a positive point charge

Benchmarking other models:  
DFT - MP2 difference in electrostatic potential

B3LYP - MP2  
(no long-range correction)

CAMB3LYP - MP2  
(long-range correction)

Use DFT with caution!



Red/blue regions correspond to increased/decreased  
electrostatic potential for DFT compared to MP2