

# HPC Challenges in Quantum Chemistry

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# Outline

- What is Quantum Chemistry?
- Why is Quantum Chemistry important?
- Some HPC Challenges in Quantum Chemistry:
  - Molecular Size
  - Molecular Motion
  - Molecular Stability
- A Scandinavian approach: Dalton
- Concluding remarks



# The fundamental equation

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$$H = -\frac{e^2}{2m_e} \sum_i \nabla_i^2 + e^2 \sum_{iK} \frac{Z_K}{R_{iK}} + \frac{e^2}{2} \sum_{ij} \frac{1}{r_{ij}} + V_{\text{NUC}}$$



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- The electronic wave function is a function of 3N variables



# The Hartree–Fock approximation

- We assume an independent particle approximation

$$F(\phi_j) \phi_i = E_i \phi_i$$

- Iterative procedure involving matrix diagonalizations
- $F$  built from one- and two-electron integrals



# More sophisticated methods

- The HF approximation ignores electron correlation
- The HF approximation scales as  $N^{2.3}$
- More sophisticated approaches scale worse
- Electron correlation important for chemical reactions
- Only *ab initio* methods can describe bond breaking and bond formation



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- Enables understanding of chemical reactions at a molecular level



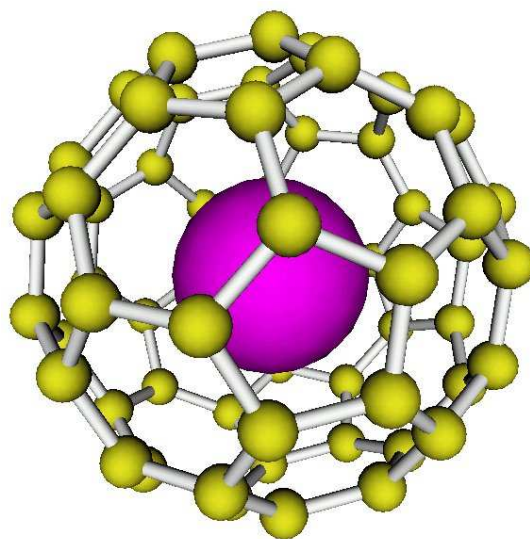
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- Enables understanding of chemical reactions at a molecular level
- Isolates individual contributions to a molecular property



# Challenge 1: Molecular size

- By chemical standards, *ab initio* methods still work on small molecules

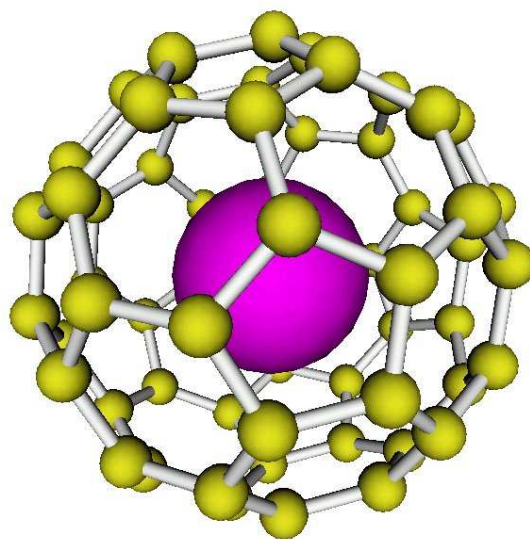


- The reason:



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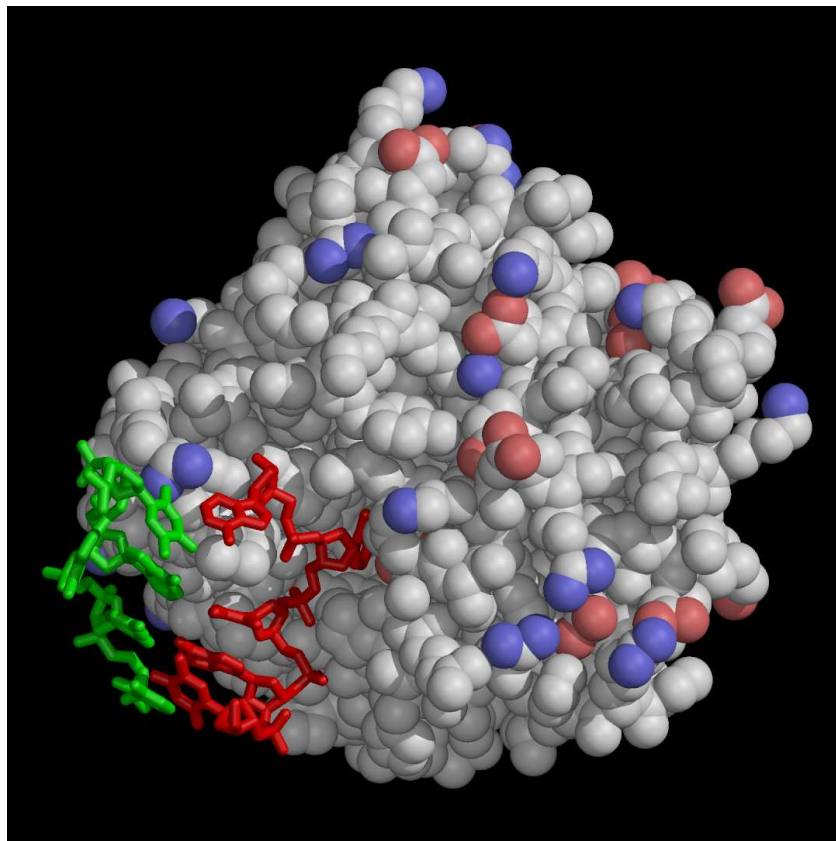
- By chemical standards, *ab initio* methods still work on small molecules



- The reason: Unfavorable scaling (at least  $N^2$ )



# What we would like to do:



The hydrogen atoms  
have been omitted

Picture by courtesy of Magne Olufsen (NORSTRUCT)

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- The approach parallelizes very well
- New (parallel) bottleneck: matrix diagonalization
- Convergence problems a challenge
- Better functionals



# HPC challenges/needs

- A lot of computer power



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- A lot of computer power
- More work on linear scaling technology
- Efficient, parallel matrix multiplication and matrix diagonalization routines
- Routines for operations on sparse matrices





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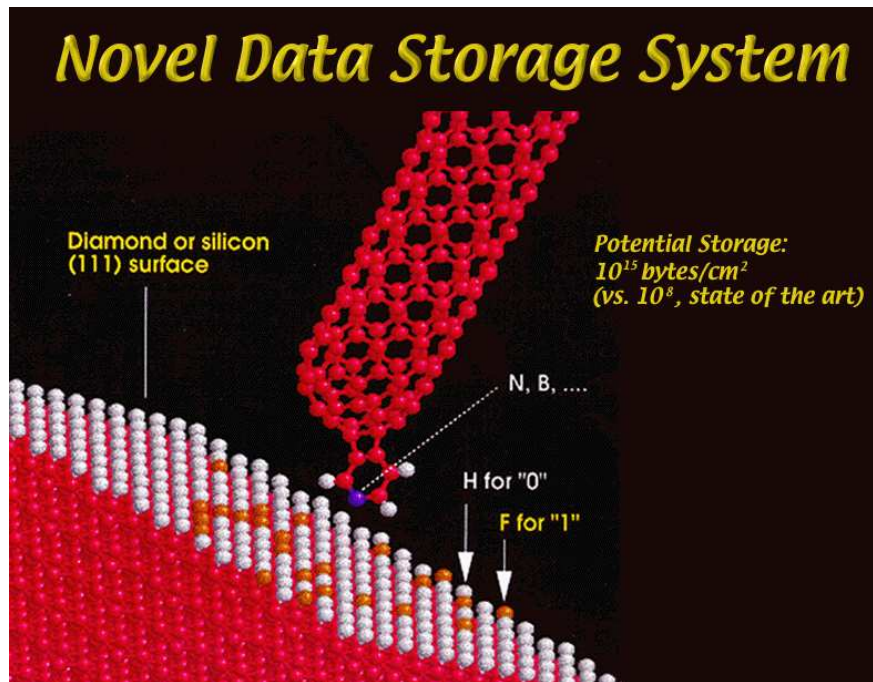


# Challenge 2: Molecular motion

- Chemistry has two parts:
  - Properties of molecules
  - Reaction of molecules
- Currently no computationally efficient method that can correctly break bonds
- This remains a quantum-chemical challenge

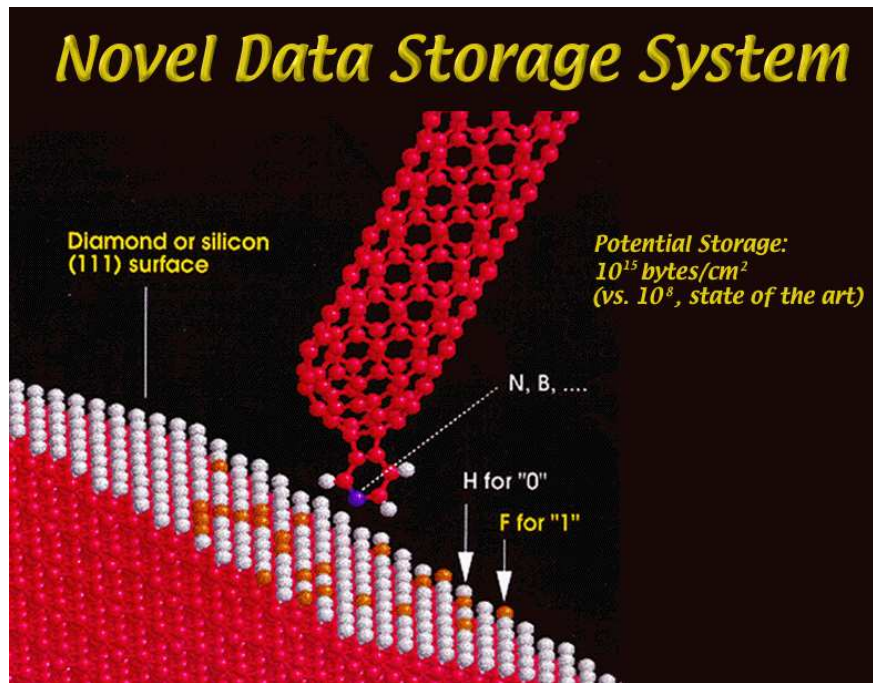


# How do molecules react and interact?



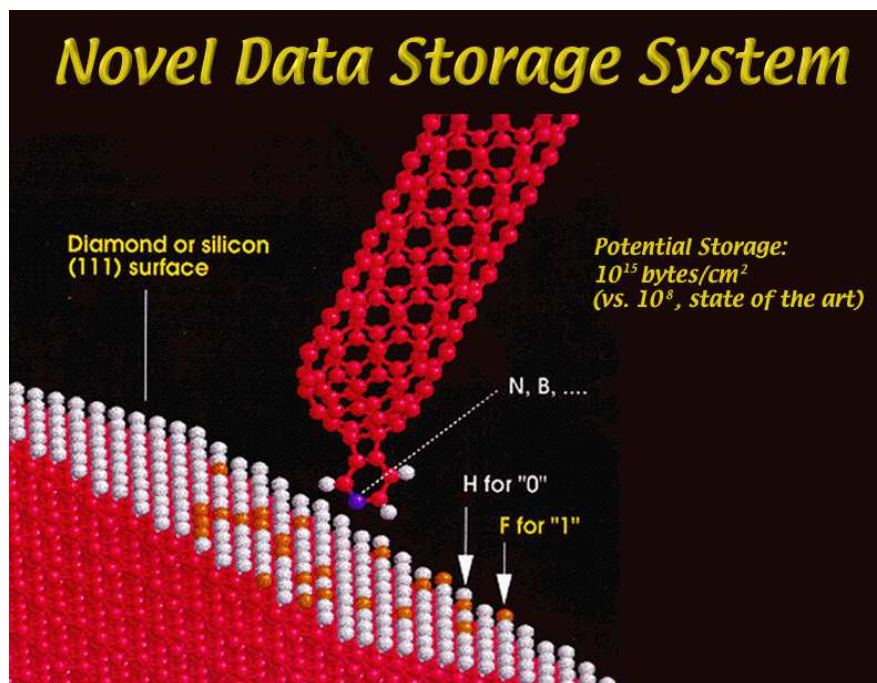
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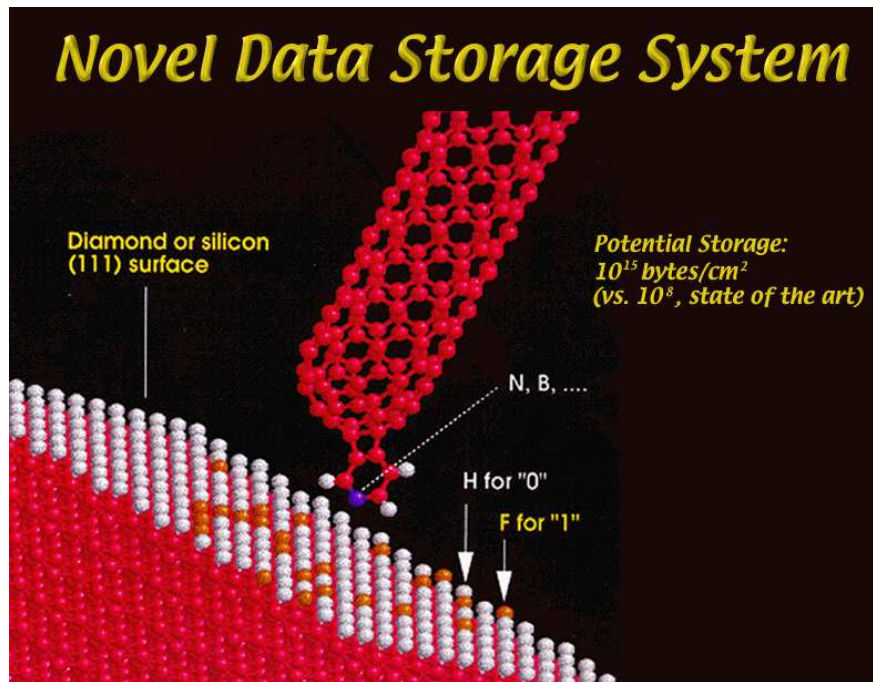
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- 3M-6 vibrational coordinates
- Easily parallelizable
- Analytic approaches possible

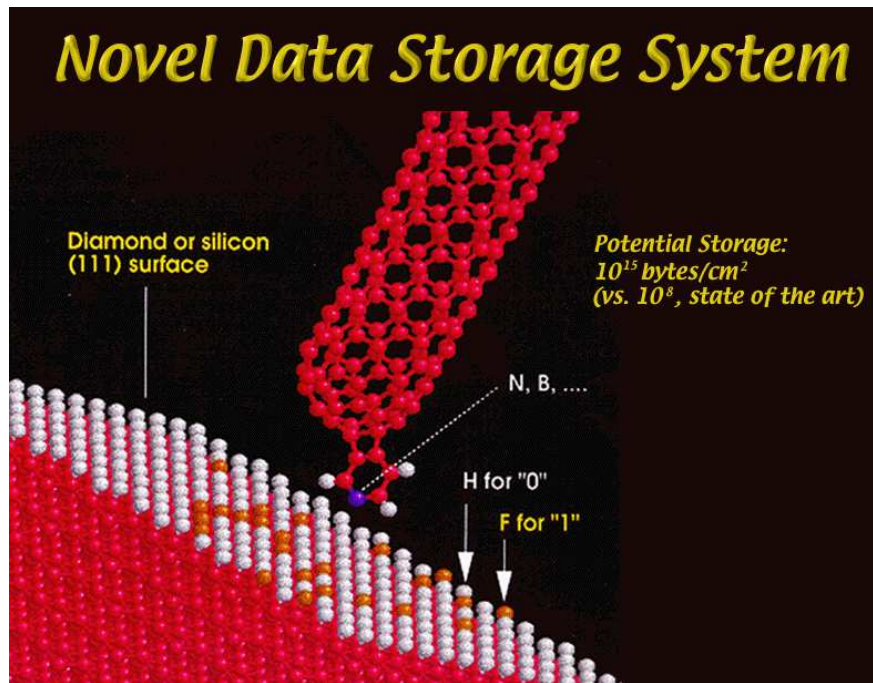


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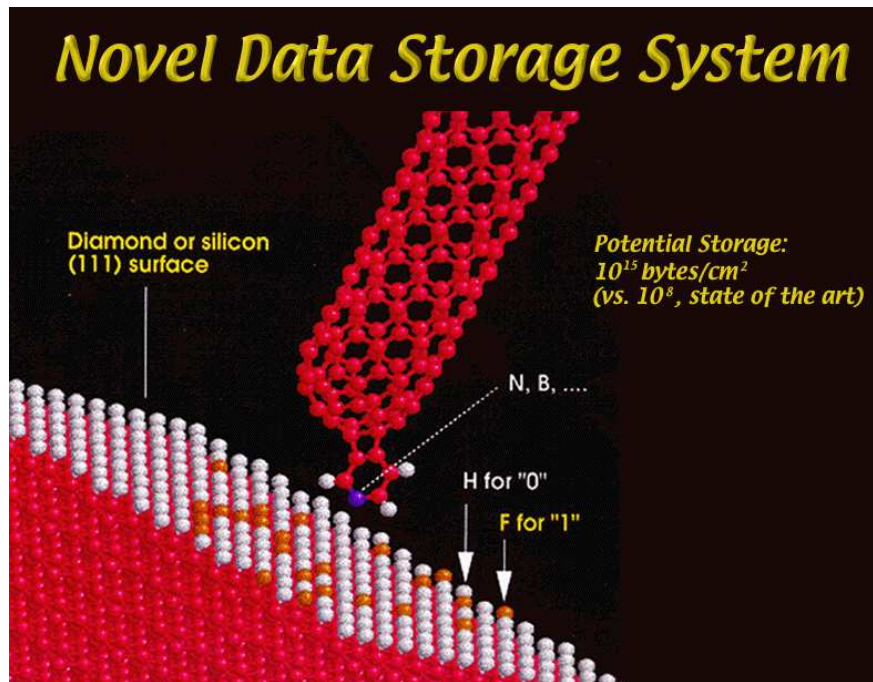
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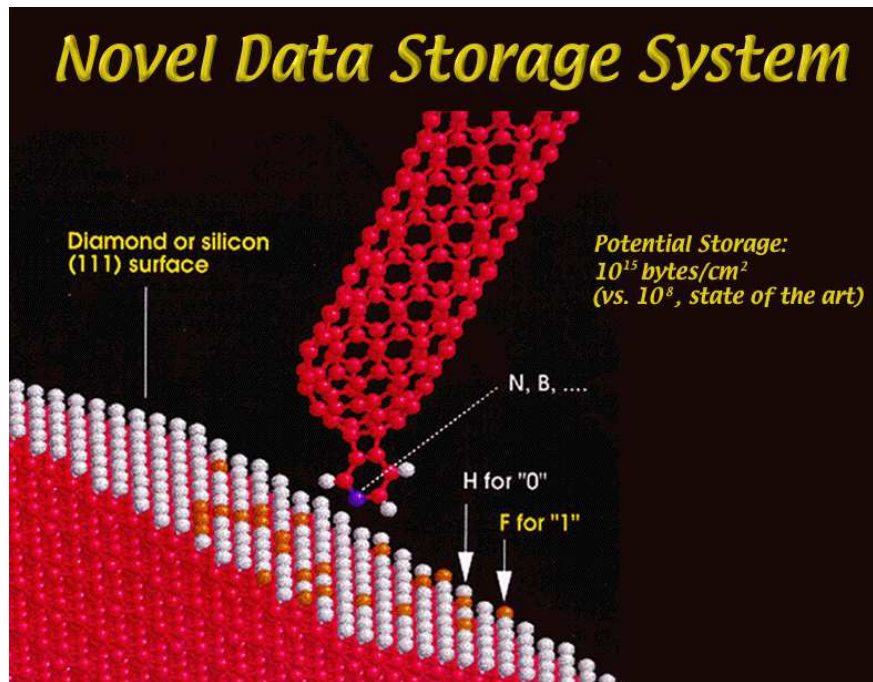
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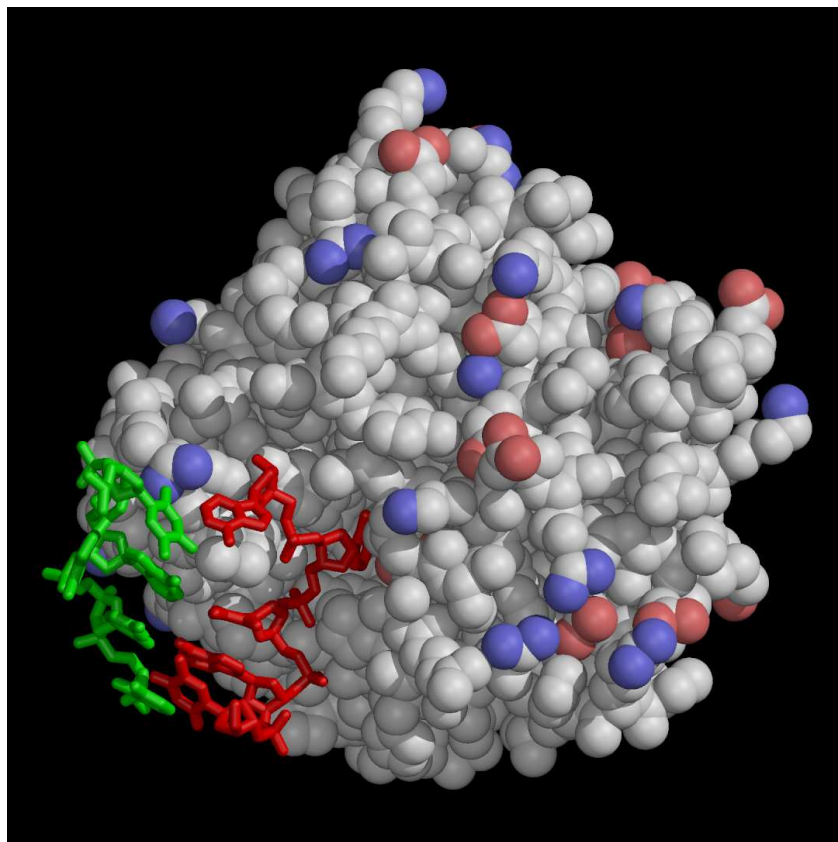
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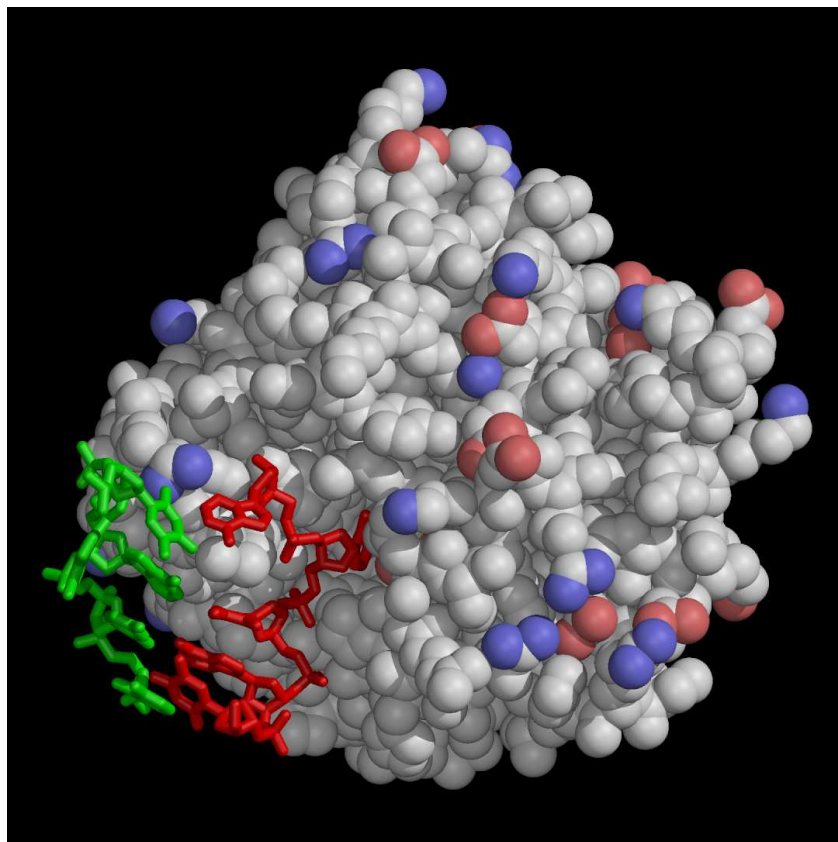
- 3M-6 vibrational coordinates
- How do the molecules interact?
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- Parallelization difficult
- Long time scales

# Challenge 3: Molecular stability



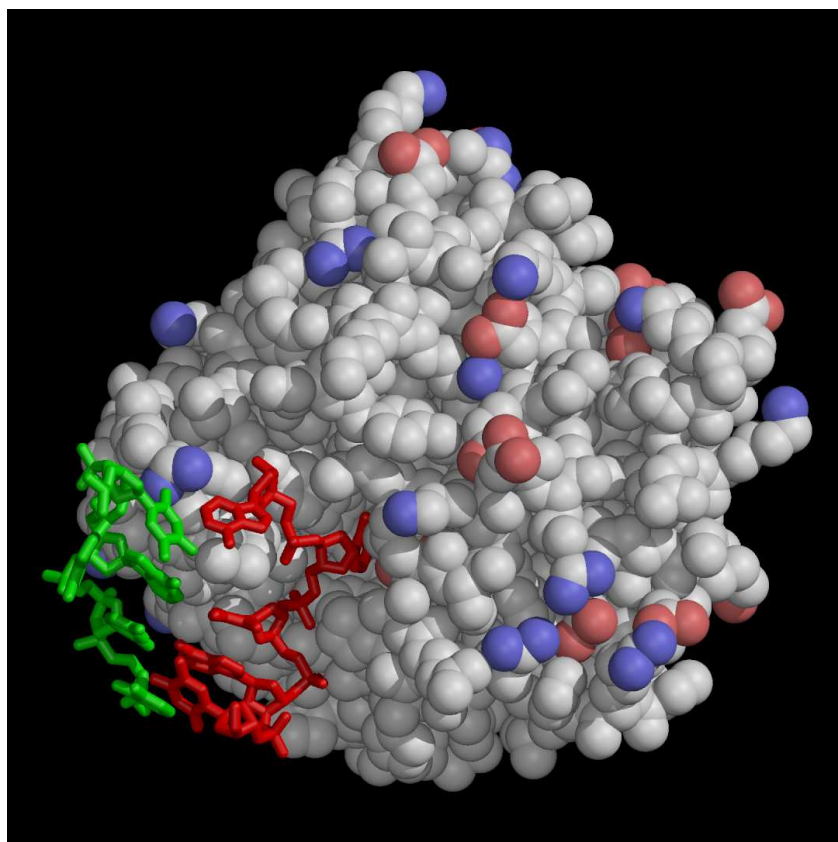
- Is this the most stable structure?

# Challenge 3: Molecular stability



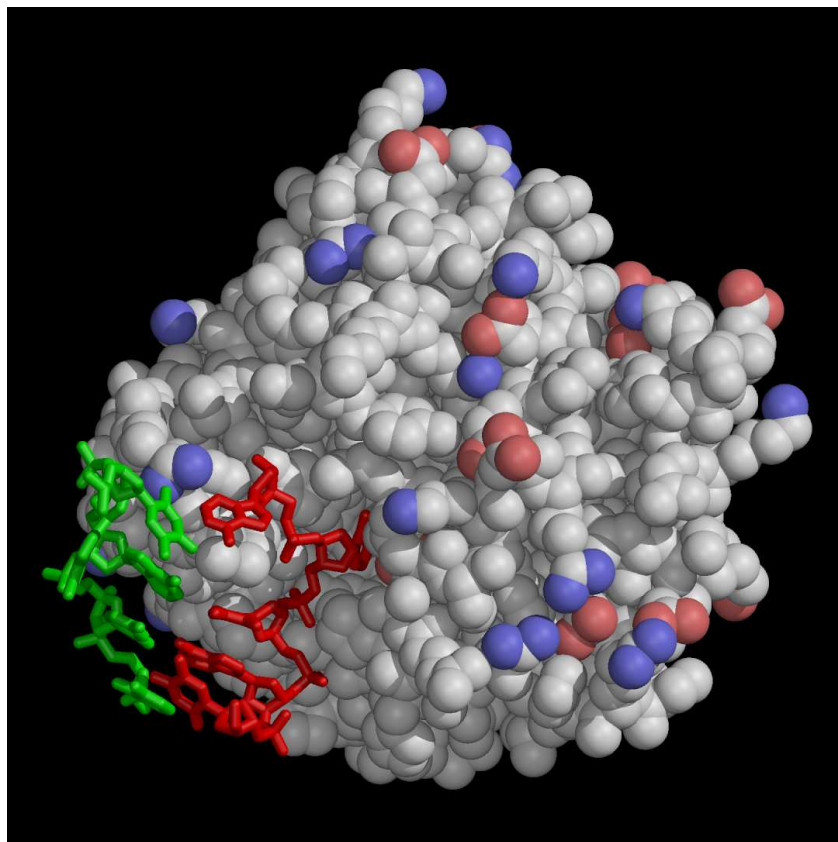
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- Is this the most stable structure?
- Which other structures are accessible?
- Global minimization
- Decoupled search (parallelizable)



# The Dalton program

C. Angeli, K. L. Bak, V. Bakken, O. Christiansen, R. Cimiraglia, S. Coriani, P. Dahle, E. K. Dalskov, T. Enevoldsen, B. Fernandez, C. Hättig, K. Hald, A. Halkier, H. Heiberg, T. Helgaker, H. Hettema, H. J. Aa. Jensen, D. Jonsson, P. Jørgensen, S. Kirpekar, W. Klopper, R. Kobayashi, H. Koch, O. B. Lutnæs, K. V. Mikkelsen, P. Norman, J. Olsen, M. J. Packer, T. B. Pedersen, Z. Rinkevicius, E. Rudberg, T. A. Ruden, K. Ruud, P. Sałek, A. Sanchez de Meras, T. Saue, S. P. A. Sauer, B. Schimmelpfennig, K. O. Sylvester-Hvid, P. R. Taylor, O. Vahtras, D. J. Wilson, H. Ågren.

<http://www.kjemi.uio.no/software/dalton/dalton.html>



# Current status of Dalton

- Dalton 1.2.1 distributed to more than 1200 research groups
- Distributed free of charge (but signed license required)
- Dalton 2.0 released March 1, and now distributed to more than 400 research groups
- Can be downloaded from <http://www.kjemi.uio.no/software/dalton/dalton.html>
- Largest developer sites: Stockholm, Linköping, Oslo, Trondheim, Tromsø, Århus, Odense, København



# Methodological advances

- As for most other HPC research fields, improvements in algorithms and programs as important as improvements in hardware
- Perhaps even more so for chemistry since hardware not adapted to our problems, and our problems are 'non-standard'
- Not all these methodological advances can lead to publications (developed by others): How do we ensure they are implemented and used?
- Two Dalton examples:
  - Incore storage of two-electron integrals
  - Density fitting

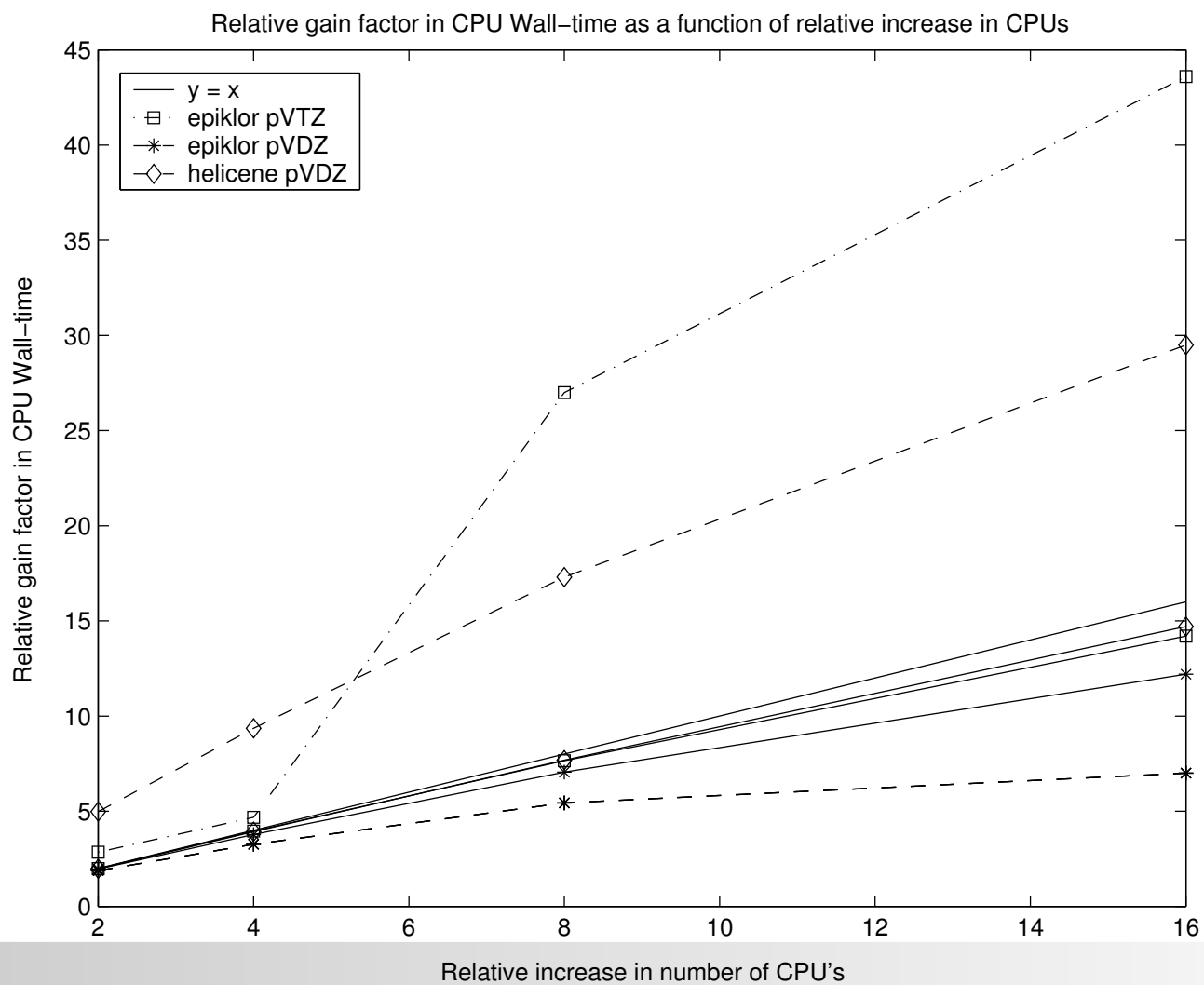


# In-core integral calculations

- Much more memory available on parallel machines
- Can be used to store two-electron integrals instead of recalculating these
- Particularly effective with density fitting (three-index integrals)
- Need to know where integrals are stored
- Master keeps track of stored integral batches ( $ij \mid **$ )
- Send batches stored back to the same slave to be used in the (partial) construction of a Fock matrix



# HF energy calculations: scaling



# Density-fitting

- One of our four-index quantities is the Coulomb-interaction

$$J_{\mu\nu\rho\sigma} = \int \int \phi_{\mu}^* (\mathbf{r}_1) \phi_{\nu} (\mathbf{r}_1) \frac{1}{|\mathbf{r}_1 - \mathbf{r}_2|} \phi_{\rho}^* (\mathbf{r}_2) \phi_{\sigma} (\mathbf{r}_2) d\mathbf{r}_1 d\mathbf{r}_2$$

- Interaction between two densities  $\Rightarrow$  we fit one of the densities to a single basis

$$J_{\mu\nu\rho\sigma} \approx \sum_{\tau} C_{\tau} \int \int \phi_{\mu}^* (\mathbf{r}_1) \phi_{\nu} (\mathbf{r}_1) \frac{1}{|\mathbf{r}_1 - \mathbf{r}_2|} \Lambda_{\tau} (\mathbf{r}_2) d\mathbf{r}_1 d\mathbf{r}_2$$

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- Reduces the formal scaling from  $N^4$  to  $N^3$
- For a calculation of 1000 basis functions, integral time reduced from 62 minutes to 5 minutes



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- Requires man-power to implement better algorithms developed by others (and develop new algorithms)
- Requires incentives for users to switch to Dalton

