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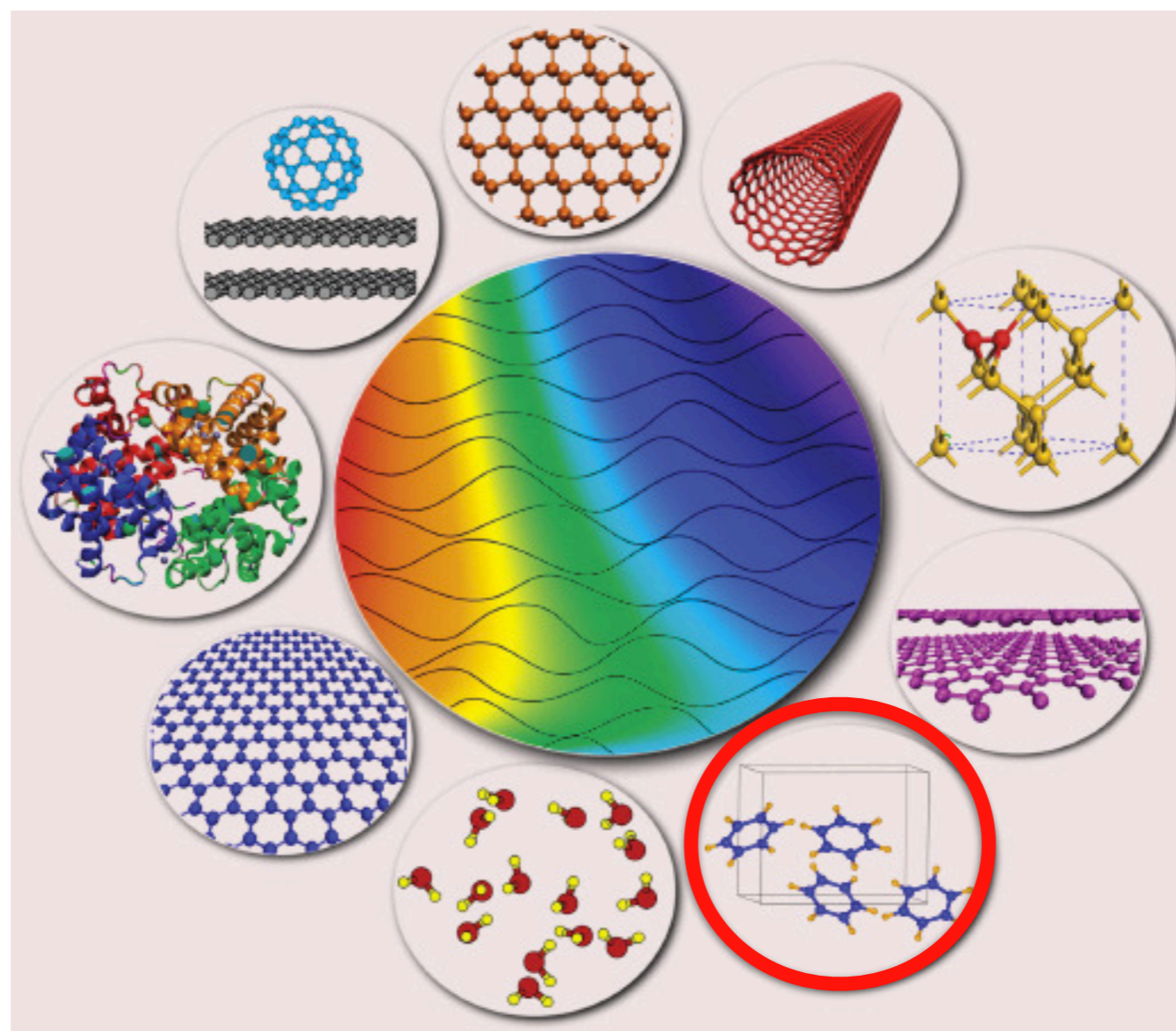
UL HPC School - June 2017

HPC in Theoretical Chemical Physics: Molecular Crystal Structure Prediction

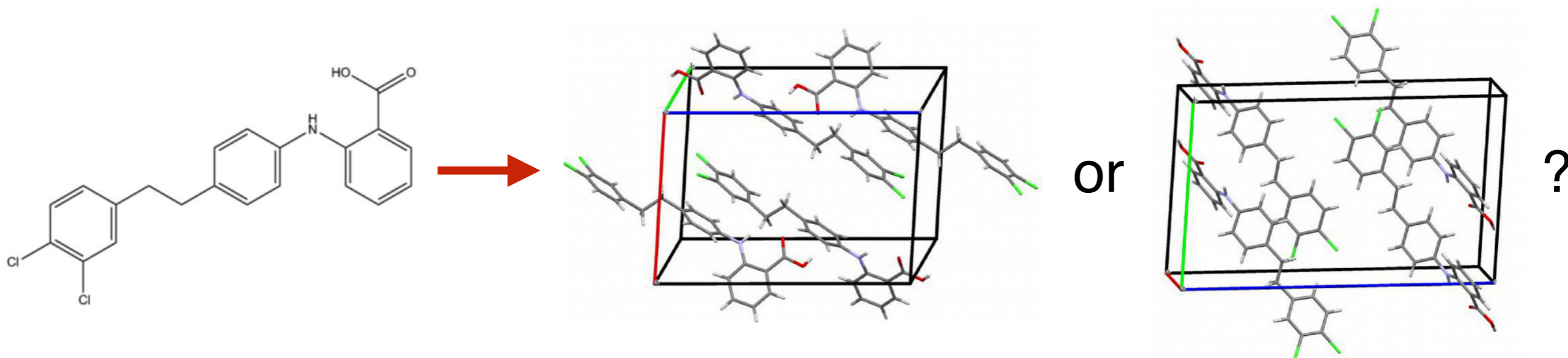
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We develop advanced quantum-mechanical first-principles methods and apply them to achieve increasingly reliable description of complex molecules and materials.



Treating van der Waals interactions as coupling between waves is a paradigm shift for chemistry and materials science © University of Luxembourg



- Solids composed of molecular moieties
- Held together by intermolecular interactions
- Different crystal-packing motifs (polymorphs) possible
- Used as pharmaceuticals, organic electronics, explosives, ...
- Energy difference between polymorphs $\sim 1 - 4 \text{ kJ/mol}$
 - ↳ $\sim 1 - 2 \%$ of lattice energy
 - ↳ High accuracy required

Polymorphs can exhibit completely different

- Kinetic stabilities
- Solubilities
- Densities
- Vibrational Spectra (THz)
- NMR chemical shifts
- Melting Points
- Conductivities
- Refractive Indices
- Vapor pressure
- Elastic constants
- Heat capacities
- ...

Vibrational free energy
— polymorph stability at
finite temperatures

Lattice energy
— relative stability of
polymorphs

Structure
— molecular structure
— crystal packing

Gibbs free energy
— phase diagram

**Crystal
structure
prediction
(CSP)**

Vibrational spectra
(IR/THz, Raman, INS)

Thermal expansion

Heat capacity

NMR chemical shifts

Elastic constants
— mechanical stability

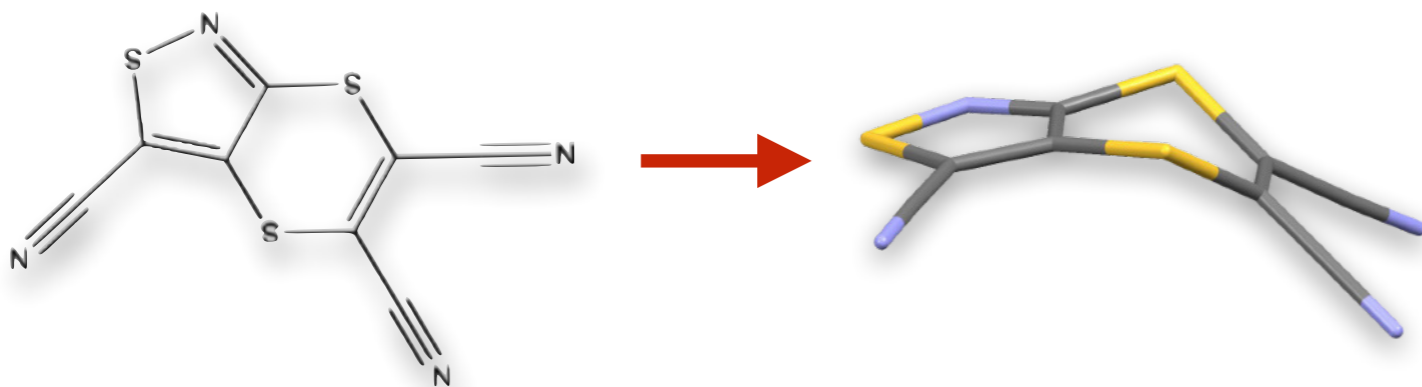
Vibrational modes

2D structural formula

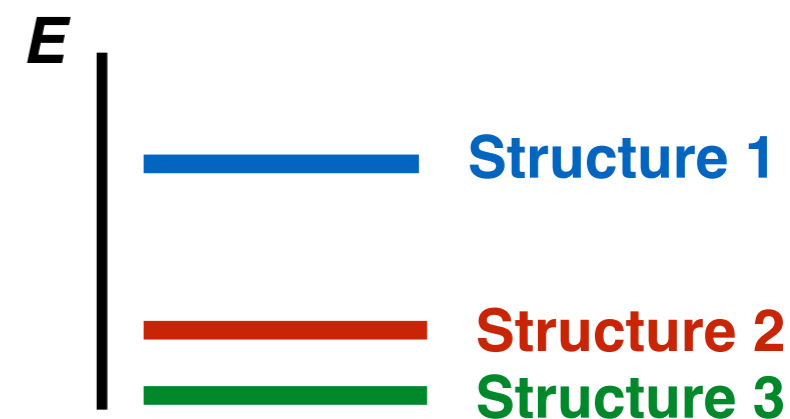


molecular crystal structure

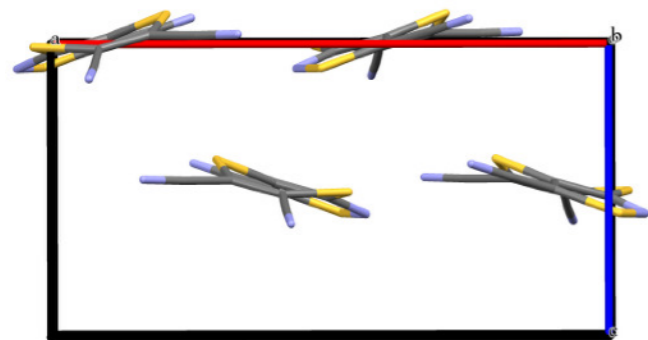
Step 1: Generating Molecular Conformations



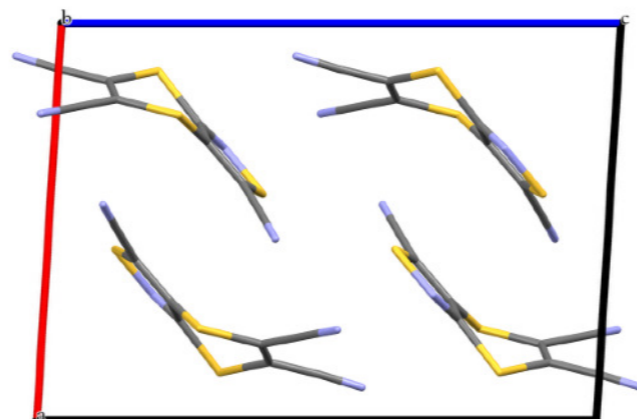
Step 3: Stability Ranking



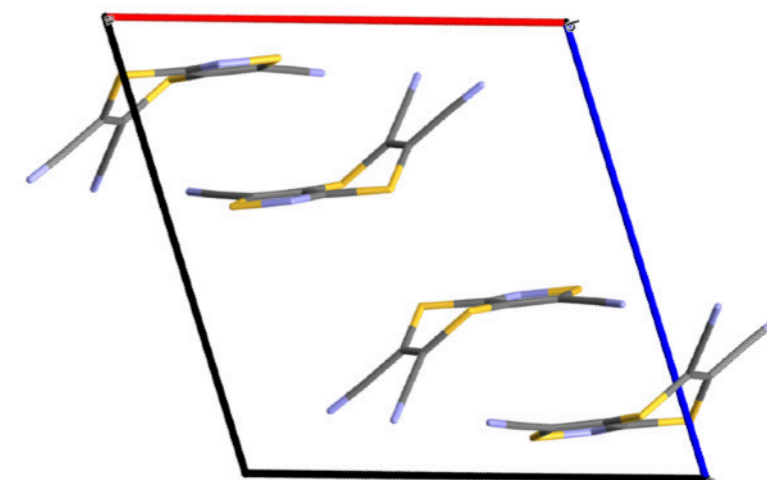
Step 2: Crystal-Packing Arrangement



Structure 1



Structure 2



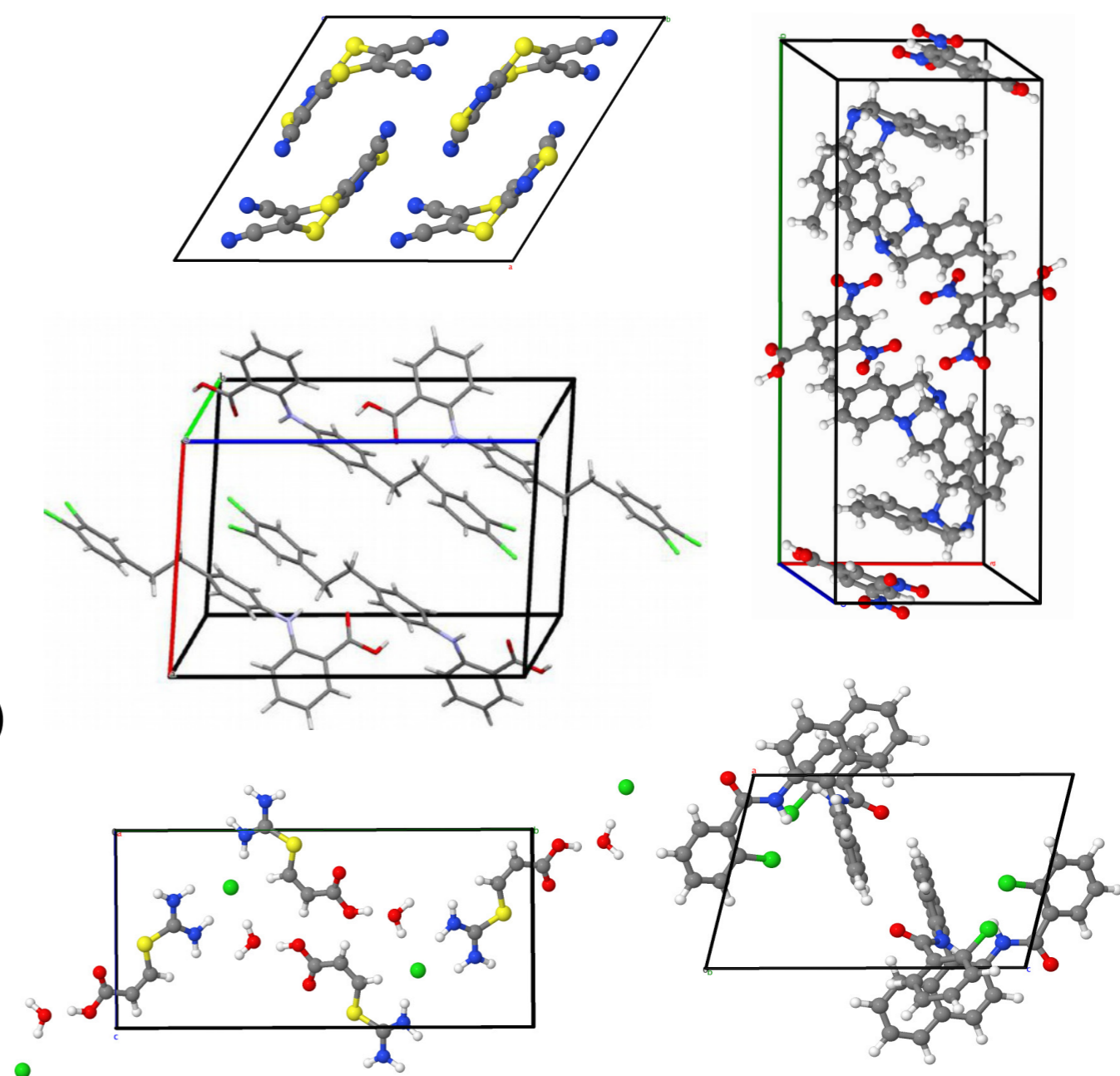
Structure 3

- Pre-screening with first-principles-derived force field
- Hierarchical first-principles approach
- vdW-inclusive DFT + harmonic vibrational free energy

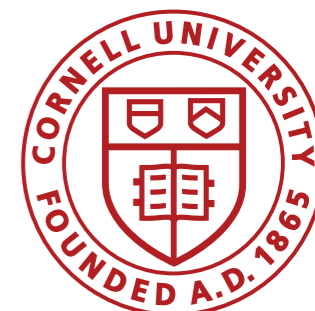
Number and size of calculations for 5 model systems:

- 500 initial structures
- 15 - 992 atoms per unit cell (average 193)
- For free energies on average
 - ~ 300 finite displacements
 - ~ 350 atoms in supercell

↳ **Large computational workload**



MAX-PLANCK-GESELLSCHAFT



Fritz Haber Institute *ab initio* molecular simulations

- Accurate and efficient quantum mechanics for molecules and materials
- All-electron, numeric atom-centered basis functions
- Non-periodic and periodic structures within same framework
- Density Functional Theory (DFT)
- Various van der Waals models, e.g. many-body dispersion [2]
- Beyond DFT: RPA, GW, MP2, ...
- Scalable: 1,000s of atoms on up to (ten)thousands of CPUs

<http://aims.fhi-berlin.mpg.de>



MAX-PLANCK-GESELLSCHAFT

[1] Blum, Gehrke, Hanke, Havu, Havu, Ren, Reuter, Scheffler, *Comput. Phys. Commun.* 180, 2175 (2009).

[2] A. Tkatchenko, R.A. DiStasio Jr., R. Car, and M. Scheffler, *Phys. Rev. Lett.* 108, 236402 (2012).

LAPACK — Linear Algebra PACKage

- solving systems of linear equations
- linear least squares
- eigenvalue problems
- matrix factorizations
- singular value decomposition
- ...

ScaLAPACK — Scalable LAPACK

- LAPACK routines redesigned for distributed memory

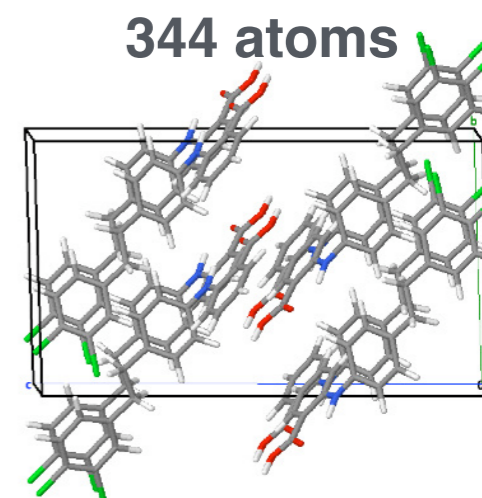
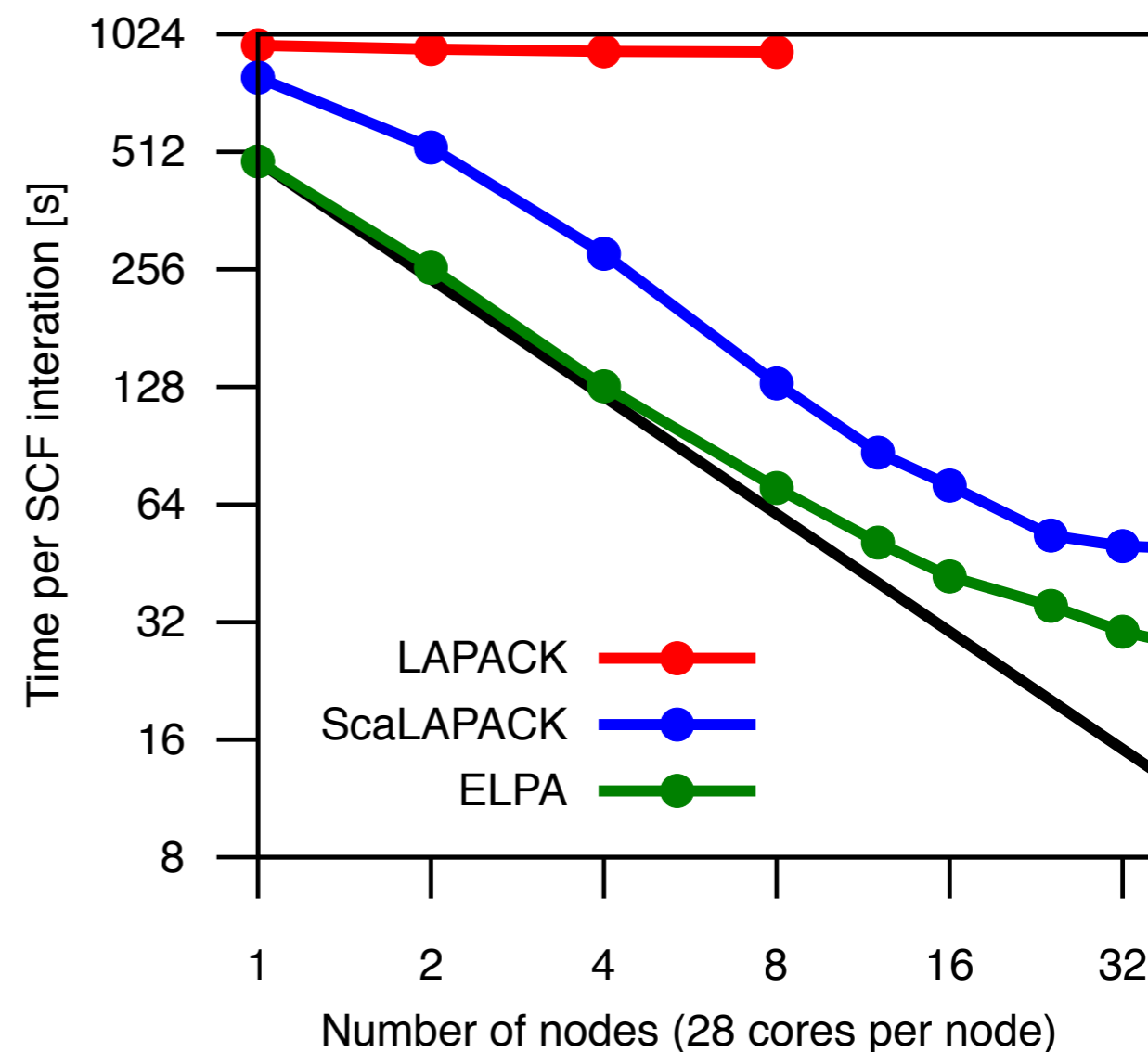
ELPA - Eigenvalue SoLvers for Petaflop-Applications

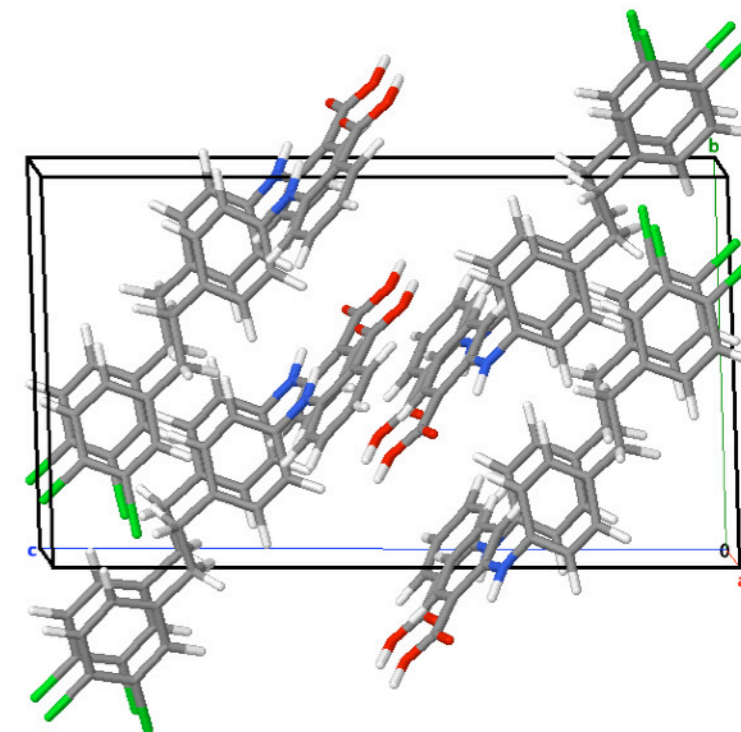
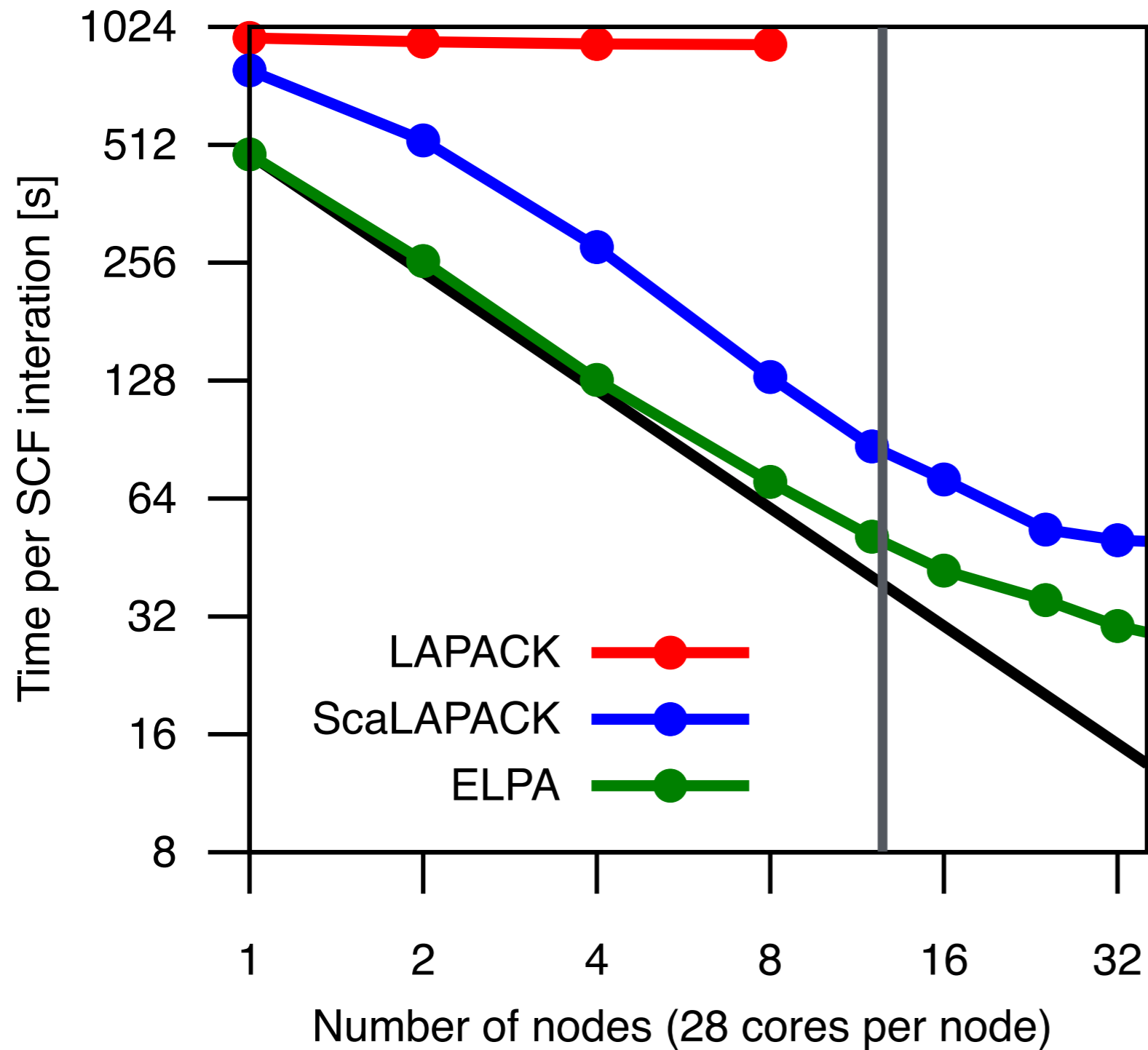
- highly efficient and highly scalable direct eigensolvers for symmetric matrices

<https://gitlab.mpcdf.mpg.de/elpa/elpa>



MAX-PLANCK-GESELLSCHAFT





344 atoms

Memory per core [GB]

	1 N	2 N	8 N
LAPACK	~4.0	~4.0	~4.0
ScaLAPACK	~1.6	~0.8	~0.3
ELPA			