



**GPAW**

**Jussi Enkovaara**

**CSC – IT Center for Science, Finland**



# Outline

- Introduction to GPAW
  - Projector-augmented wave method
  - Basis sets in GPAW
- Using Atomic Simulation Environment and GPAW
- Hands-on exercises
  
- Lunch
  
- Time-dependent DFT with GPAW
- Parallel calculations ( + other features by request)
- Hands-on exercises

# GPAW



- Implementation of projector augmented wave method on
  - uniform real-space grids, atomic orbital basis, plane waves
- Density-functional theory, time-dependent DFT, many-body perturbation theory, ...
- Massively parallelized

[wiki.fysik.dtu.dk/gpaw](http://wiki.fysik.dtu.dk/gpaw)

[gpaw-users@listserv.fysik.dtu.dk](mailto:gpaw-users@listserv.fysik.dtu.dk)

**J. J. Mortensen et al., Phys. Rev. B 71, 035109 (2005)**

**J. Enkovaara et al., J. Phys. Condens. Matter 22, 253202 (2010)**

# GPAW



- Open source software licensed under GPL
  - 20-30 developers in Europe and USA



- The main GPAW references cited ~350 times
- ~300 subscribers in users mailing list



# GPAW features

- Total energies, forces, structural optimization
  - analysis of electronic structure
- Excited states, optical spectra
  - Non-adiabatic electron-ion dynamics
- Wide range of XC-potentials (thanks to libxc!)
  - LDAs, GGAs, meta-GGAs, hybrids, DFT+U, vdW, RPA
- Electron transport
- GW-approximation, Bethe-Salpeter equation
- ...

# Projector-augmented wave method



- Exact all-electron formalism
- “Pseudopotentials done right”
- Good description over the whole periodic table
- Access to full wave functions and density
- Norm-conserving and ultrasoft pseudopotentials can be derived as approximations to PAW



# PAW transformation

- Linear transformation from pseudo wave functions to all-electron wave functions

$$|\psi_k\rangle = \hat{T}|\tilde{\psi}_k\rangle$$

- Transformation operator

$$\hat{T} = 1 + \sum_{a,i} \left( |\phi_i^a\rangle - |\tilde{\phi}_i^a\rangle \right) \langle \tilde{p}_i^a |$$

$|\phi_i^a\rangle$     **all-electron atomic orbital**

$|\tilde{\phi}_i^a\rangle$     **pseudo atomic orbital**

$|\tilde{p}_i^a\rangle$     **projector function**

- Core electrons are frozen

# Properties of PAW transformation



- Projector functions are localized inside the augmentation spheres

$$p_i^a(r) = 0, r > R_a$$

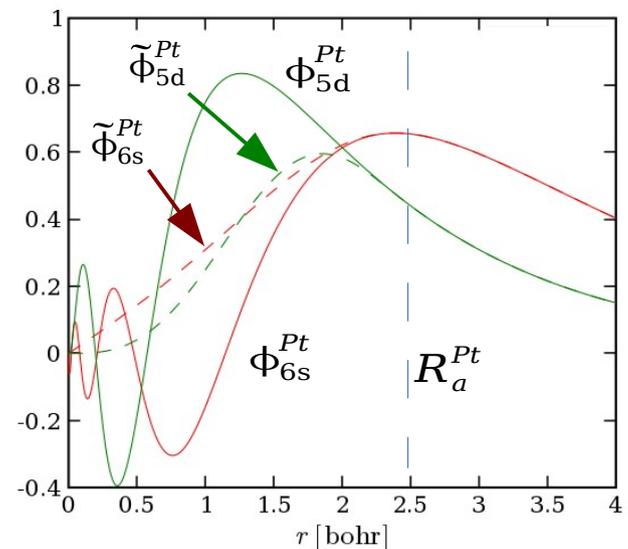
- All-electron orbitals and pseudo orbitals are equal outside the augmentation spheres

$$\phi_i^a(r) = \tilde{\phi}_i^a(r), r > R_a$$

- Projector functions are orthogonal to pseudo orbitals

$$\langle \tilde{p}_i^a | \tilde{\phi}_j^a \rangle = \delta_{ij}$$

Example: Platinum





# PAW transformation

- The projector functions and partial waves are constructed from all-electron calculation for spherical symmetric atom
- To be exact, infinite number of projectors and partial waves is needed
  - In practice, 1-2 functions per angular momentum is enough



# One center expansion

- Inside augmentation spheres one can define one-center expansions of AE and PS state

$$\psi_n^a(\mathbf{r}) = \sum_i P_{in}^a \phi_i^a(\mathbf{r})$$

$$\tilde{\psi}_n^a(\mathbf{r}) = \sum_i P_{in}^a \tilde{\phi}_i^a(\mathbf{r})$$

with expansion coefficients

$$P_{in}^a = \langle \tilde{\phi}_i^a | \tilde{\psi}_n \rangle$$

- All electron wave function can now be written as

$$\psi_n = \tilde{\psi}_n + \sum_a (\psi_n^a - \tilde{\psi}_n^a)$$



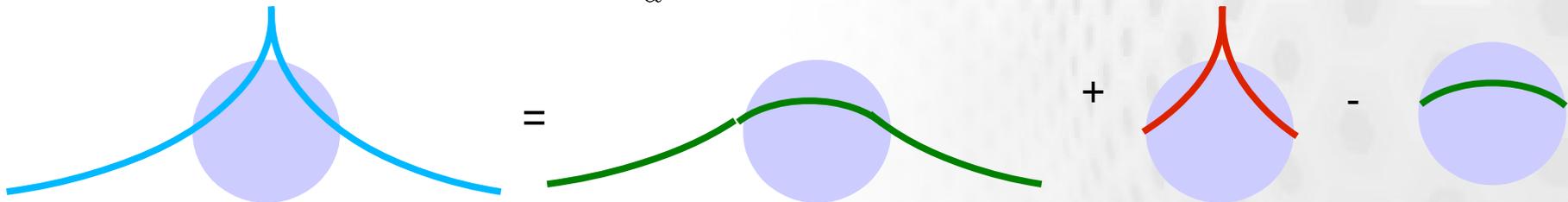
# PAW expectation values

- Within frozen core approximation expectation value of operator  $\hat{O}$  is

$$\langle \hat{O} \rangle = \sum_n^{val} f_n \langle \psi_n | \hat{O} | \psi_n \rangle + \sum_a \sum_{\alpha} \langle \phi_{\alpha}^{a,core} | \hat{O} | \phi_{\alpha}^{a,core} \rangle$$

- By inserting the PAW expression, one obtains for (semi)local operators

$$\begin{aligned} \langle \psi_n | \hat{O} | \psi_n \rangle &= \langle \tilde{\psi}_n | \hat{O} | \tilde{\psi}_n \rangle + \sum_a \left( \langle \psi_n^a | \hat{O} | \psi_n^a \rangle - \langle \tilde{\psi}_n^a | \hat{O} | \tilde{\psi}_n^a \rangle \right) \\ &= \tilde{O} + \sum_a \left( O^a - \tilde{O}^a \right) \end{aligned}$$





# PAW Hamiltonian

- PAW Hamiltonian can be written as

$$\hat{H} = -\frac{1}{2}\nabla^2 + \tilde{v}(\mathbf{r}) + \sum_a \sum_{i_1 i_2} |\tilde{p}_{i_1}^a\rangle \Delta H_{i_1 i_2}^a \langle \tilde{p}_{i_2}^a|$$

- Pseudo wave functions are orthonormal only with respect to overlap operator

$$\langle \psi_i | \psi_j \rangle = \langle \tilde{\psi}_i | \hat{T}^\dagger \hat{T} | \tilde{\psi}_j \rangle = \langle \tilde{\psi}_i | \hat{S} | \tilde{\psi}_j \rangle = \delta_{ij}$$

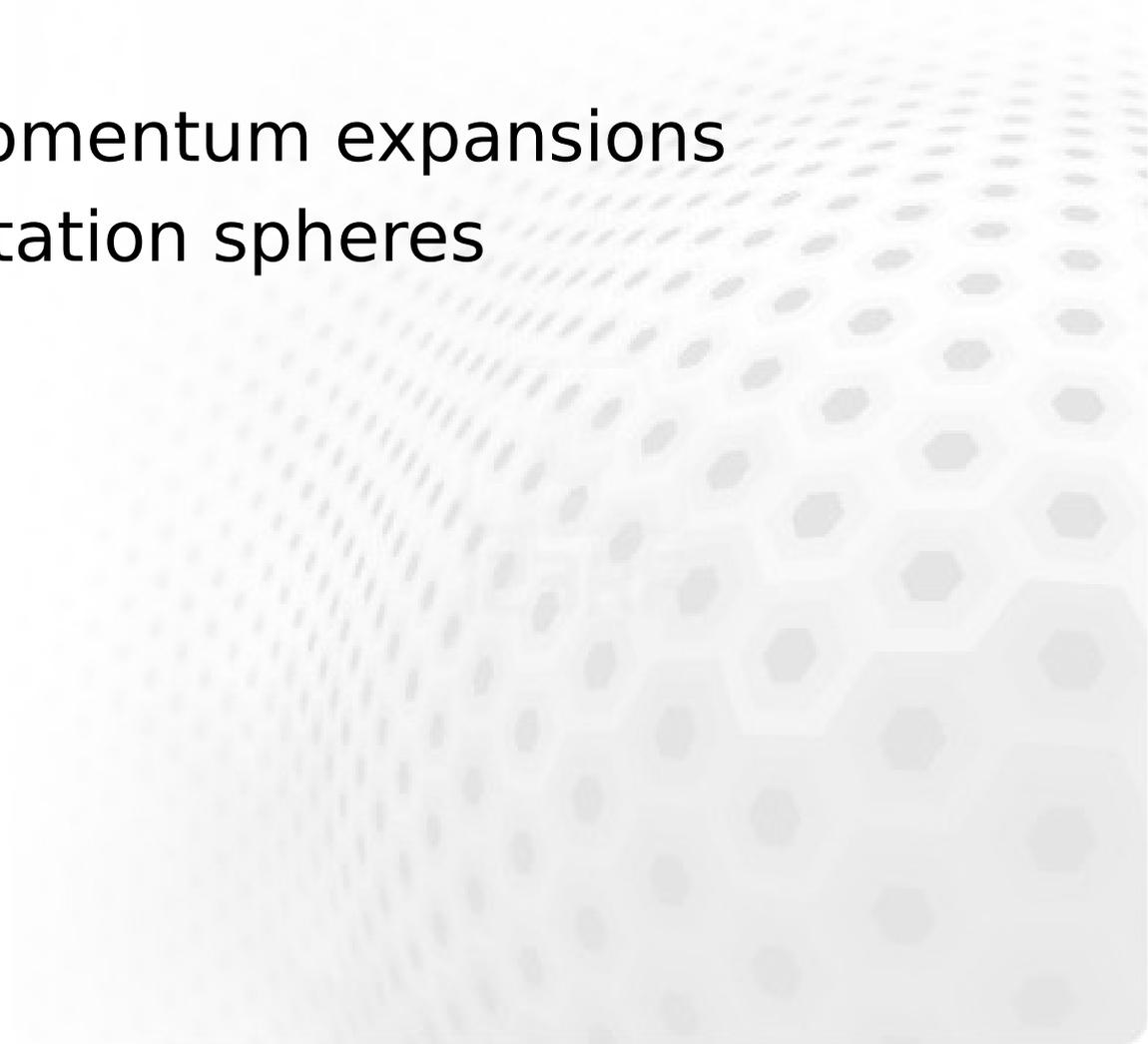
- Generalized eigenvalue equation

$$\hat{H} \tilde{\psi}_n = \epsilon_n \hat{S} \tilde{\psi}_n$$



# Approximations in PAW

- Finite number of projectors
  - typically two projectors per angular momentum are used
- Truncated angular momentum expansions
- Overlapping augmentation spheres
- Frozen core





# PAW setup

➤ A set of

$$\phi^a, \tilde{\phi}^a, \tilde{p}^a, n_c^a, \tilde{n}_c^a, \bar{v}^a$$

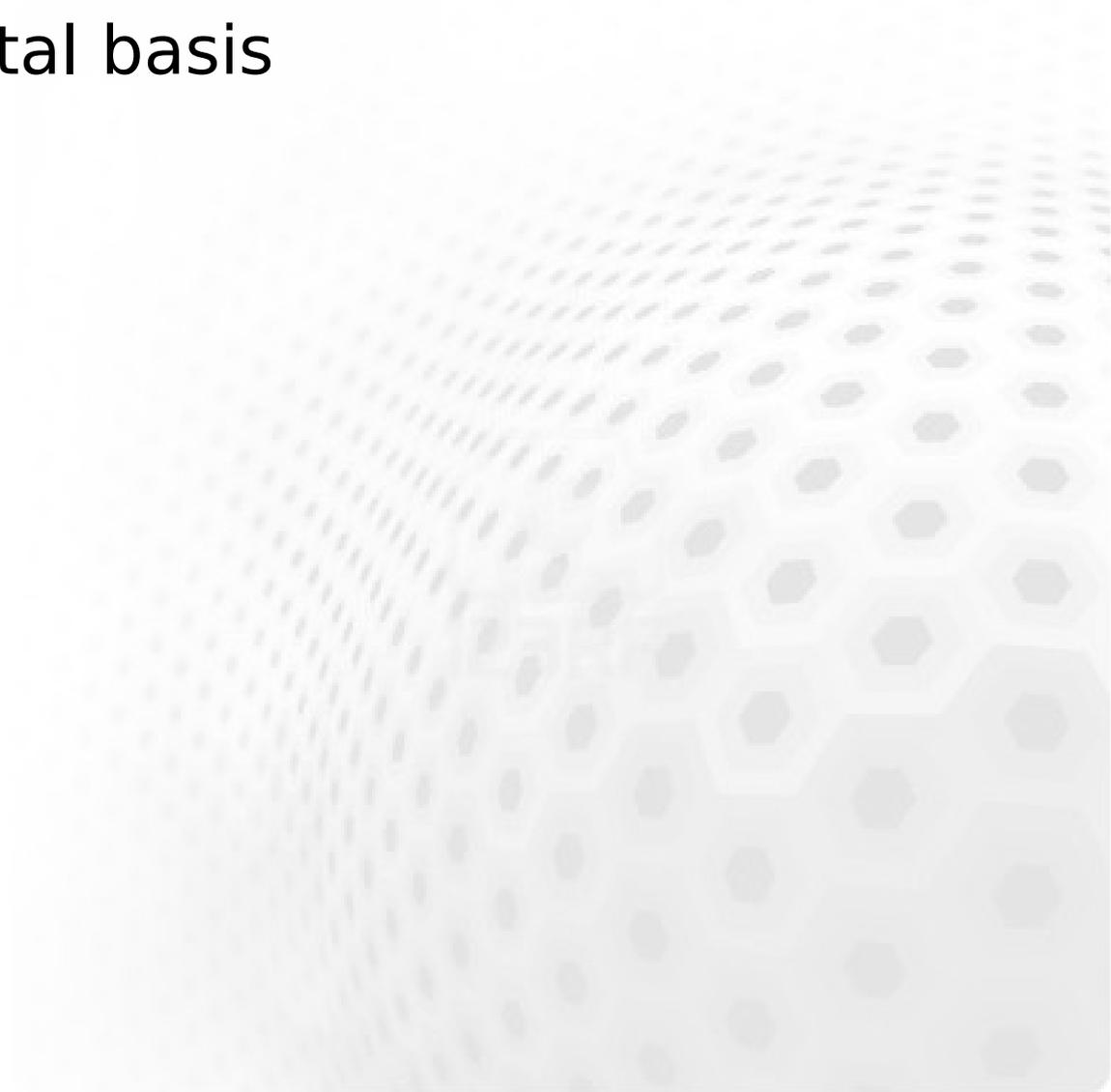
for a single atom constitutes a PAW setup

- Setups are generated for individual atoms
- The actual PAW calculations use pregenerated setups
- Setup testing is non-trivial and time-consuming
  - correct properties for the particular atom
  - transferability in different molecules and solids



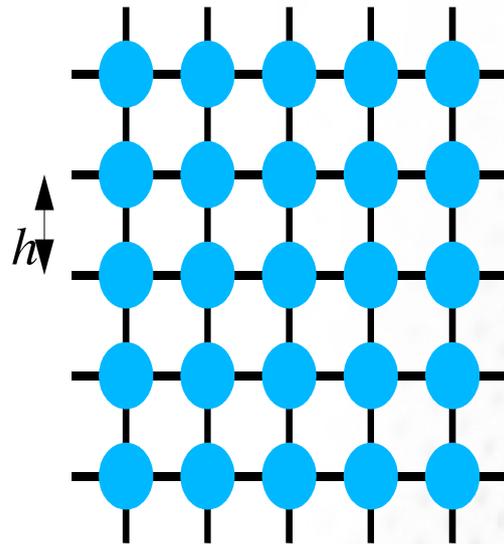
# Basis sets in GPAW

- Real-space grids
- Localized atomic orbital basis
- Plane waves



# Real-space grids

- Wave functions, electron densities, and potentials are represented on grids.
- Single parameter, grid spacing  $h$



- Accuracy of calculation can be improved systematically
- Derivatives by finite differences



# Boundary conditions

- Real-space description allows flexible boundary conditions
- Zero boundary conditions (finite systems)
  - Useful especially in charged systems
- Periodic boundary conditions (bulk systems)
- Boundary conditions can be mixed
  - periodic in one dimension (wires)
  - periodic in two dimensions (surfaces)



# Atomic orbital basis

- Linear combination of atomic orbitals (LCAO) provide compact basis set

$$\tilde{\psi}_n = \sum_{\nu} C_{n\nu} \Phi_{\nu}(\mathbf{r})$$

$$\Phi_{nlm}^a(\mathbf{r}) = R_{nl}^a(|\mathbf{r} - \mathbf{R}^a|) Y_{lm}(\mathbf{r} - \mathbf{R}^a)$$

- The atomic orbitals are obtained from a free atom in a confining potential well
- Systematic improvement of accuracy is non-trivial
- Possible to switch between localized basis and real-space grids



# Plane wave basis

- Functions which are periodic with respect to unit cell can be written as sum of plane waves

$$\psi_{n\mathbf{k}}(\mathbf{r}) = \frac{1}{\Omega^{1/2}} \sum_{\mathbf{G}} C_{\mathbf{G},n\mathbf{k}} e^{i(\mathbf{G}+\mathbf{k})\cdot\mathbf{r}}$$

$$n(\mathbf{r}) = \sum_{\mathbf{G}} n_{\mathbf{G}} e^{i\mathbf{G}\cdot\mathbf{r}} \quad V(\mathbf{r}) = \sum_{\mathbf{G}} V_{\mathbf{G}} e^{i\mathbf{G}\cdot\mathbf{r}}$$

- The expansion is truncated according to

$$\frac{1}{2} |\mathbf{G} + \mathbf{k}|^2 < E_{cutoff}$$

- Only periodic cells (supercells)



# Comparison of basis sets

## ➤ Real-space grids

- systematic convergence with single parameter
- good parallelization prospects
- some integrals complicated in real-space



## ➤ Localized basis set

- compact basis
- systematic convergence can be difficult



## ➤ Plane waves

- systematic convergence with single parameter
- some integrals simplified in reciprocal space
- very efficient in small to medium size systems
- parallelization more limited due FFTs





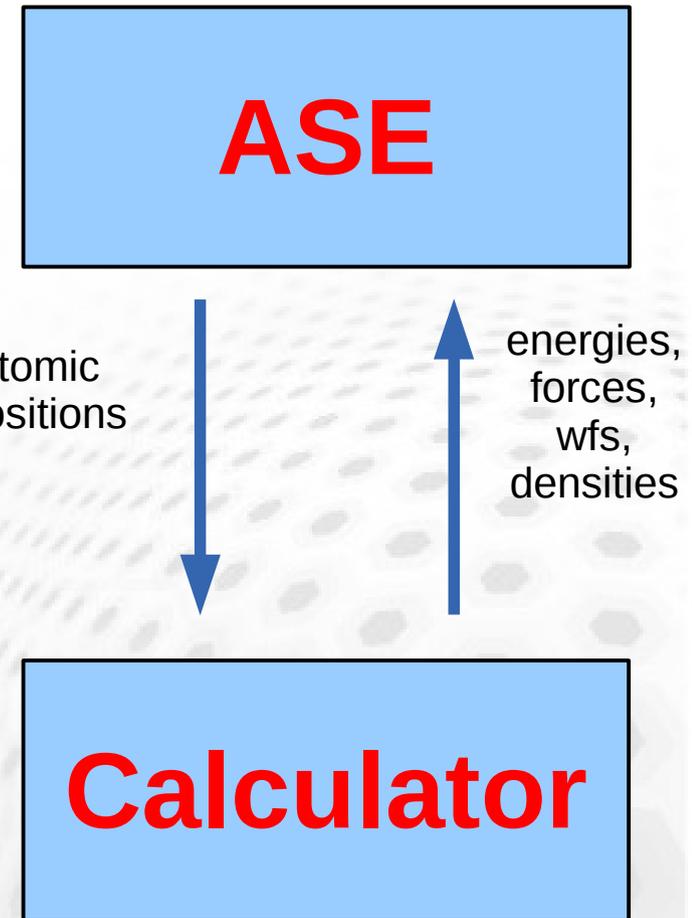
# Using the Atomic Simulation Environment



# Atomic Simulation Environment



- ASE is a Python package for
  - building atomic structures
  - structure optimization and molecular dynamics
  - analysis and visualization
- ASE relies on external software which provides total energies, forces, etc.
  - GPAW, Abinit, Siesta, Vasp, Castep, ...
- Input files are Python scripts
  - calculations are run as “`python input.py`”
  - simple format, no knowledge of Python required
  - knowledge of Python enables great flexibility
- Simple graphical user interface



[wiki.fysik.dtu.dk/ase](http://wiki.fysik.dtu.dk/ase)



# Setting up the atoms

## ➤ Specifying atomic positions directly

```
from ase.all import * # Setup the atomic simulation environment

d0 = 1.10
x = d0 / sqrt(3)
atoms = Atoms('CH4',
              positions=[(0.0, 0.0, 0.0), # C
                          (x, x, x),     # H1
                          (-x, -x, x),   # H2
                          (-x, x, -x),    # H3
                          (x, -x, -x)]    # H4
              )
view(atoms)
```

## ➤ Reading atomic positions from a file

```
...
atoms = read('CH4.xyz')
view(atoms)
```

- Several file formats supported



# Setting up the unit cell

- By default, the simulation cell of an Atoms object has zero boundary conditions and edge length of 1 Å
- Unit cell can be set when constructing Atoms

```
atoms = Atoms(..., # positions must be now in absolute coordinates  
              cell=(1., 2., 3.), pbc=True) # or pbc=(True, True, True)
```

or later on

```
atoms = Atoms(...) # positions in relative coordinates  
atoms.set_cell((2.5, 2.5, 2.5), scale_atoms=True)  
atoms.set_pbc(True) # or atoms.set_pbc((True, True, True))
```

```
atoms = ...  
atoms.center(vacuum=3.5) # finite system 3.5 Å empty space around atoms
```

```
atoms = ...  
atoms.set_pbc((False, True, True)) # surface slab  
atoms.center(axis=0, vacuum=3.5) # 3.5 Å empty space in x-direction
```



# Units in ASE

➔ Length: Å

➔ Energy: eV

➔ Easy conversion between units:

```
from ase.units import Bohr, Hartree  
  
a = a0 * Bohr      # a0 in a.u., a in Å  
E = E0 * Hartree  # E0 in Hartree, E in eV
```

– also Rydberg, kcal, nm, ...

# Pre-defined molecules and structures



## ➤ Database of small molecules (G2-1 and G2-2 sets)

```
from ase.structure import molecule

mol = molecule('C6H6') # coordinates from MP2 calculation
mol.center(3.5)        # molecule() returns unit cell of 1 Å
```

## ➤ Bulk structures of elemental materials

```
from ase.lattice import bulk

atoms = bulk('Si') # primitive (2-atom) unit cell with exp. lattice constant
atoms_conv = bulk('Si', cubic=True) # cubic 8-atom unit cell
atoms_my_a = bulk('Si', a=5.4) # User specified lattice constant
```



# Supercells and surfaces

- Existing Atoms objects can be “repeated” and individual atoms removed

```
from ase.lattice import bulk

atoms = bulk('Si', cubic=True) # cubic 8-atom unit cell
supercell = atoms.repeat((4, 4, 4)) # 512 atom supercell
del supercell[0] # remove first atom, e.g. create a vacancy
```

- Utilities for working with surfaces

```
from ase.lattice.surface import fcc111, add_adsorbate

slab = fcc111('Cu', size=(3,3,5)) # 5-layers of 3x3 Cu (111) surface
# add O atom 2.5 Å above the surface in the 'bridge' site
add_adsorbate(slab, 'O', 2.5, position='bridge')
```



# Performing a calculation

- In order to do calculation, one has to define a calculator object and attach that to Atoms

```
from ase.structure import molecule # Setup the atomic simulation environment
from gpaw import GPAW # Setup GPAW

atoms = molecule('CH4')
atoms.center(3.5)
calc = GPAW() # Use default parameters
atoms.set_calculator(calc)
atoms.get_potential_energy() # Calculate the total energy
```

- Specifying calculator parameters

```
...
calc = GPAW(h=0.18, nbands=6, # 6 bands and grid spacing of 0.20 Å
            kpts=(4,4,4), # 4x4x4 Monkhorst-Pack k-mesh
            xc='PBE', txt='out.txt') # PBE and print text output to file
...
```

- See [wiki.fysik.dtu.dk/gpaw/documentation/manual.html](http://wiki.fysik.dtu.dk/gpaw/documentation/manual.html) for all parameters



# Performing a calculation

- Serial calculations and analysis can be carried out with normal Python interpreter

```
[jenkovaa@flamingo ~]$ python input.py
```

- Parallel calculations with **gpaw-python** executable

```
#PBS -N gpaw_test  
#PBS -l select=4  
#PBS -l walltime=00:20:00  
...  
aprun -n 96 gpaw-python input.py
```



# Structural optimization

```
from ase.all import *           # Setup the atomic simulation environment
from gpaw import GPAW          # Setup GPAW

atoms = ...
calc = GPAW(...)
atoms.set_calculator(calc)

opt = BFGS(atoms, trajectory='file.traj') # define an optimizer
opt.run(fmax=0.05) # optimize the structure until forces smaller than 0.05 eV / Å
```

- See [wiki.fysik.dtu.dk/ase/ase/optimize.html](http://wiki.fysik.dtu.dk/ase/ase/optimize.html) for supported optimizers
- “Best” optimizer is case-dependent



# Simple Python scripting

```
atoms = ...
calc = GPAW(...)
atoms.set_calculator(calc)

# Check convergence with grid spacing
for h in [0.35, 0.30, 0.25, 0.20, 0.18]:
    txtfile = 'test_h' + str(h) + '.txt'
    calc.set(h=h, txt=txtfile)
    e = atoms.get_potential_energy()
    print h, e
```

```
import numpy as np
atoms = ...
calc = GPAW(...)
atoms.set_calculator(calc)

# lattice constant for different XC-functionals
for xc in ['LDA', 'PBE']:
    for a in np.linspace(3.8, 4.3, 5):
        txtfile = 'test_xc_' + xc + '_a' + str(s) + '.txt'
        atoms.set_cell((a, a, a), scale_atoms=True)
        calc.set(xc=xc, txt=txtfile)
        e = atoms.get_potential_energy()
```



# Saving and restarting

- Saving full state of calculation: .gpw-files (or .hdf5-files)

```
...
calc = GPAW(...)
atoms.set_calculator(calc)
atoms.get_potential_energy() # Calculate the total energy
calc.write('myfile.gpw')    # Atomic positions, densities, calculator parameters
```

```
...
calc.write('myfile.gpw', mode='all') # Save also wave functions (larger files)
```

```
...
calc.write('myfile.hdf5', mode='all') # If GPAW is build with HDF5 support
```

- Restarting

```
from ase.all import * # Setup the atomic simulation environment
from gpaw import restart # Setup GPAW

atoms, calc = restart('file.gpw')
e0 = atoms.get_potential_energy() # no calculation needed
calc.set(h=0.20)
e1 = atoms.get_potential_energy() # calculation total energy with new grid
```



# Saving and restarting

## ➤ Trajectories: atomic positions, energies, forces

```
...
calc = GPAW(...)
atoms.set_calculator(calc)
traj = PickleTrajectory('file.traj', 'w', atoms) # define a trajectory file
for a in np.linspace(3.8, 4.3, 5):
    txtfile = 'test_xc_' + xc + '_a' + str(s) + '.txt'
    atoms.set_cell((a, a, a), scale_atoms=True)
    atoms.get_potential_energy()
    traj.write() # write cell and energy to trajectory file
```

## ➤ Reading atomic positions

```
from ase.all import * # Setup the atomic simulation environment
from gpaw import GPAW # Setup GPAW

atoms = read('file.traj') # read the last image
first = read('file.traj', 0) # first image
calc = GPAW(...)
atoms.set_calculator(calc) # calculator has to be attached
```

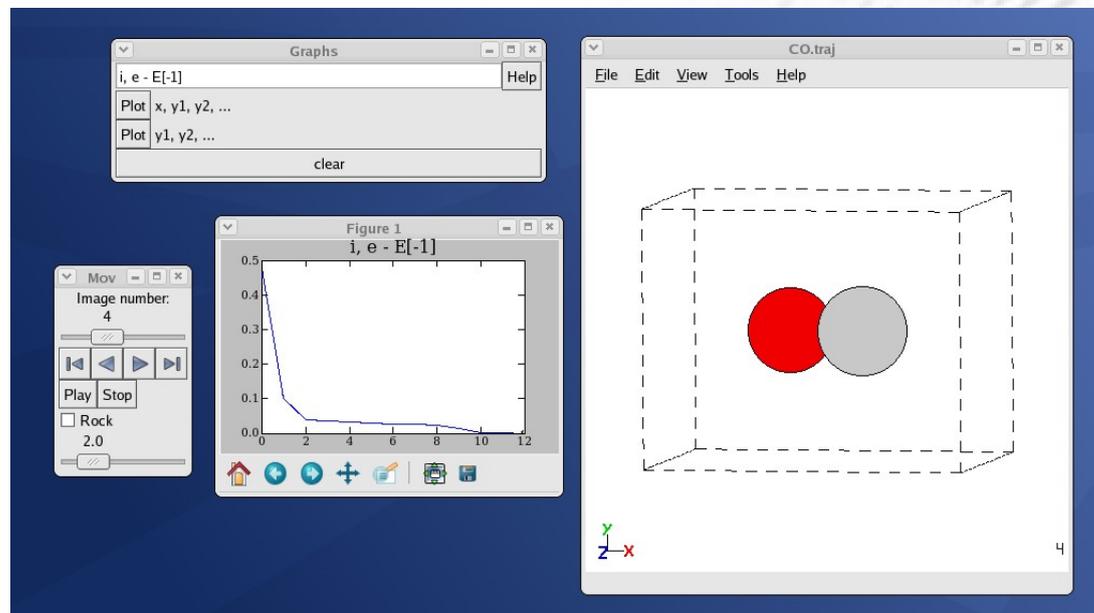
# Simple graphical interface (ase-gui)



- Trajectory can be investigated with **ase-gui** tool

```
[jenkovaa@flamingo ~]$ ase-gui file.traj
```

- Investigate how total energy, forces, bond lengths etc. vary during simulation



# Exercises



➔ Go to

[wiki.fysik.dtu.dk/gpaw/exercises/exercises.html](http://wiki.fysik.dtu.dk/gpaw/exercises/exercises.html)

and get your hands dirty!





# Time-dependent DFT with GPAW





# Time-dependent DFT

- Generalization of density-functional theory also to time-dependent cases
- Runge-Gross theorem PRL 52 (1984)
  - one-to-one mapping between the time-dependent potential and the density
- Excited state properties
  - excitation energies, optical spectra, ...
- Time-dependent Kohn-Sham equations

$$i \frac{\partial \psi_i(r, t)}{\partial t} = H \psi_i(r, t)$$
$$H = -\frac{\nabla^2}{2} + V_{eff}(r, t)$$
$$n(r, t) = \sum_i |\psi_i(r, t)|^2$$



# Real-time propagation

- Direct integration of time-dependent Kohn-Sham equations in time-domain

$$i \frac{\partial \psi_i(r, t)}{\partial t} = H(t) \psi_i(r, t)$$

$$n(r, t) = \sum_i |\psi_i(r, t)|^2$$

- Integration is done with Crank-Nicholson type scheme:

$$[1 + iH(t)\Delta t] \psi_i(t + \Delta t) = [1 - iH(t)\Delta t] \psi_i(t)$$

- Initial value problem, the starting value  $\psi_i(t = 0)$  is obtained from ground state calculation

# Optical absorption spectra from real-time TDDFT



- Excite the system with delta pulse  $E(t) = \epsilon \mathbf{k}^0 \delta(t)$
- Wave functions change instantaneously to

$$\psi(t = 0^+) = \exp(i\epsilon \mathbf{k}^0 \cdot \mathbf{r}) \psi(t = 0)$$

- Time-propagate wave functions and record the time-dependent dipole moment
- Spectra can be obtained via Fourier transform of the time-dependent dipole-moment



# Practice

- Perform ground state calculation and save the wave functions

```
from ase.all import *
from gpaw import GPAW

atoms = ...
atoms.center(5.0) # may need to be larger in real calculations
calc = GPAW(h=0.30)
atoms.set_calculator(calc)
atoms.get_potential_energy()
calc.write('gs.hdf5', mode='all')
```

- Time-propagate

```
from gpaw.tddft import TDDFT

time_step = 16.0 # as
iters = 650 # 650 x 16 as 10.4 fs
kick = [0,0,1e-3] # Weak delta kick to z-direction

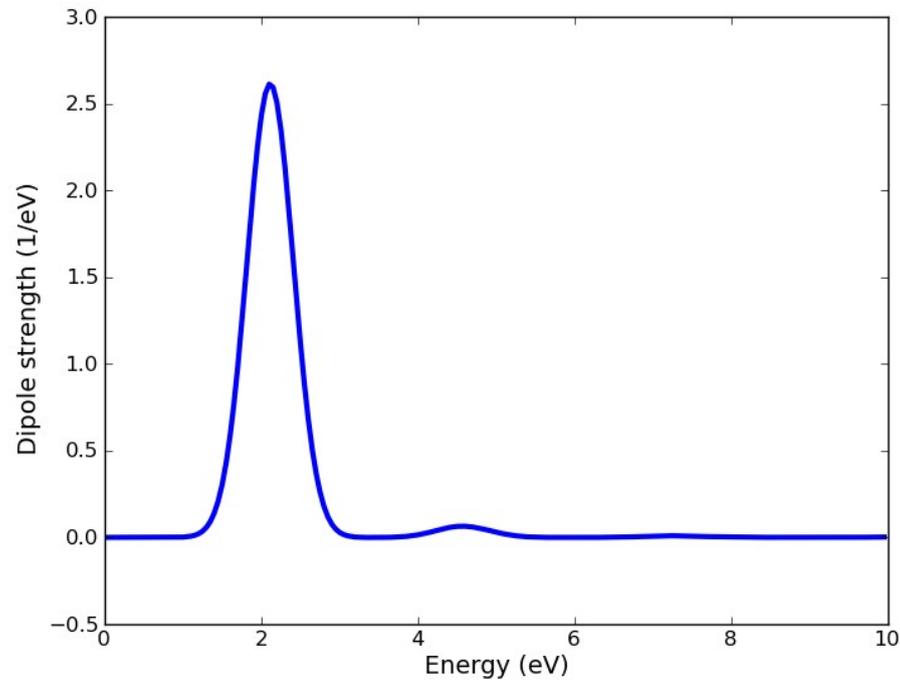
td_calc = TDDFT('gs.hdf5')
td_calc.absorption_kick(kick)
td_calc.propagate(time_step, iters, 'dmz.dat')
```

# Practice



## ➔ Calculate the optical spectra

```
from gpaw.tddft import photoabsorption_spectrum  
photoabsorption_spectrum('dmz.dat', 'spectrum_z.dat', width=0.2)
```

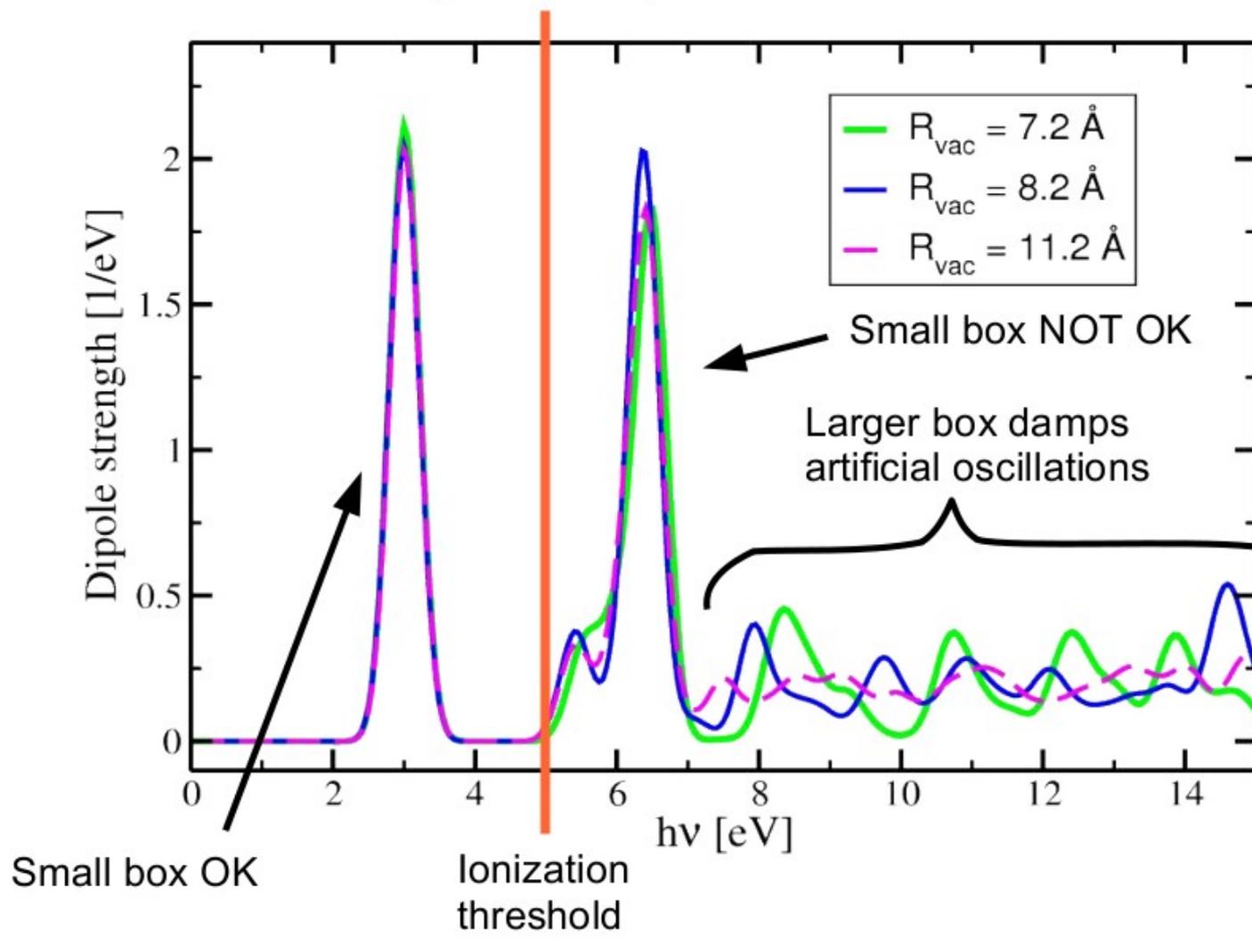




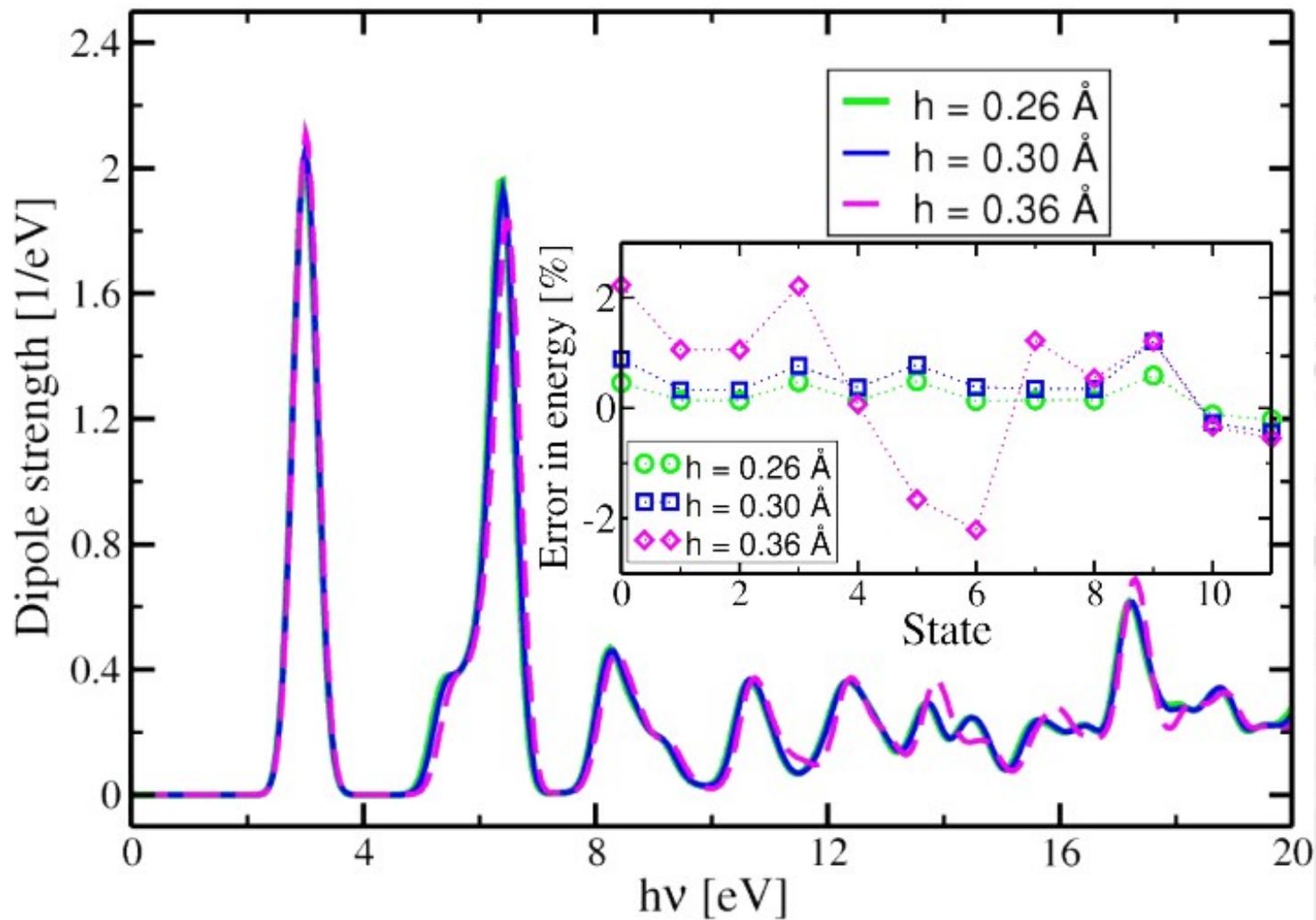
# Practical details

- Depending on the symmetry of system, separate kick in all  $x, y, z$  directions may be needed.
- Normally, the size of simulation box has to be larger than in ground state calculations
- Grid spacing can often be larger than in ground state calculations
- Time-step affects the accuracy of spectra (too large time-step may lead into unstable propagation)
- Total simulation time affects the resolution of spectra.

# Box size

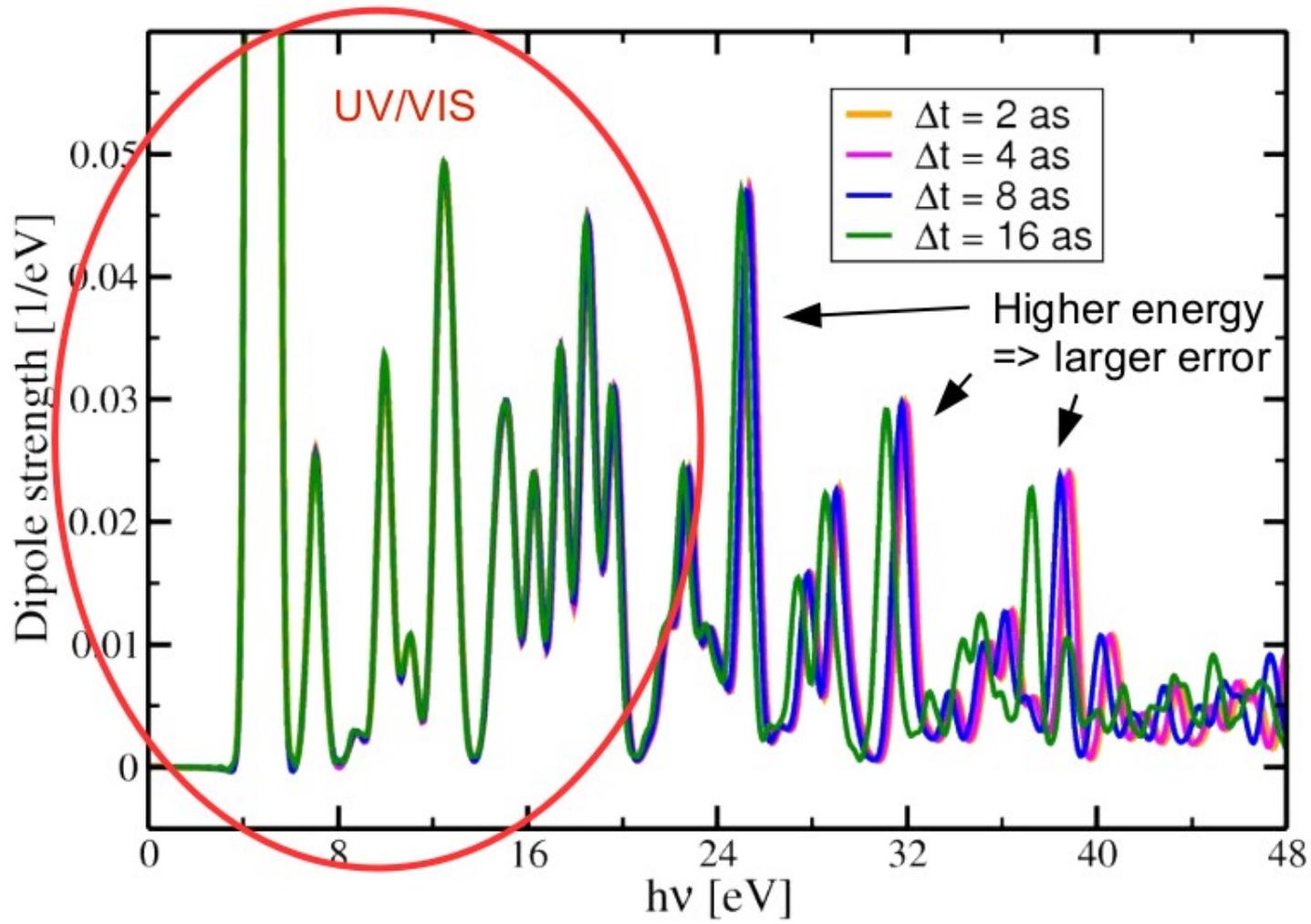


# Grid spacing

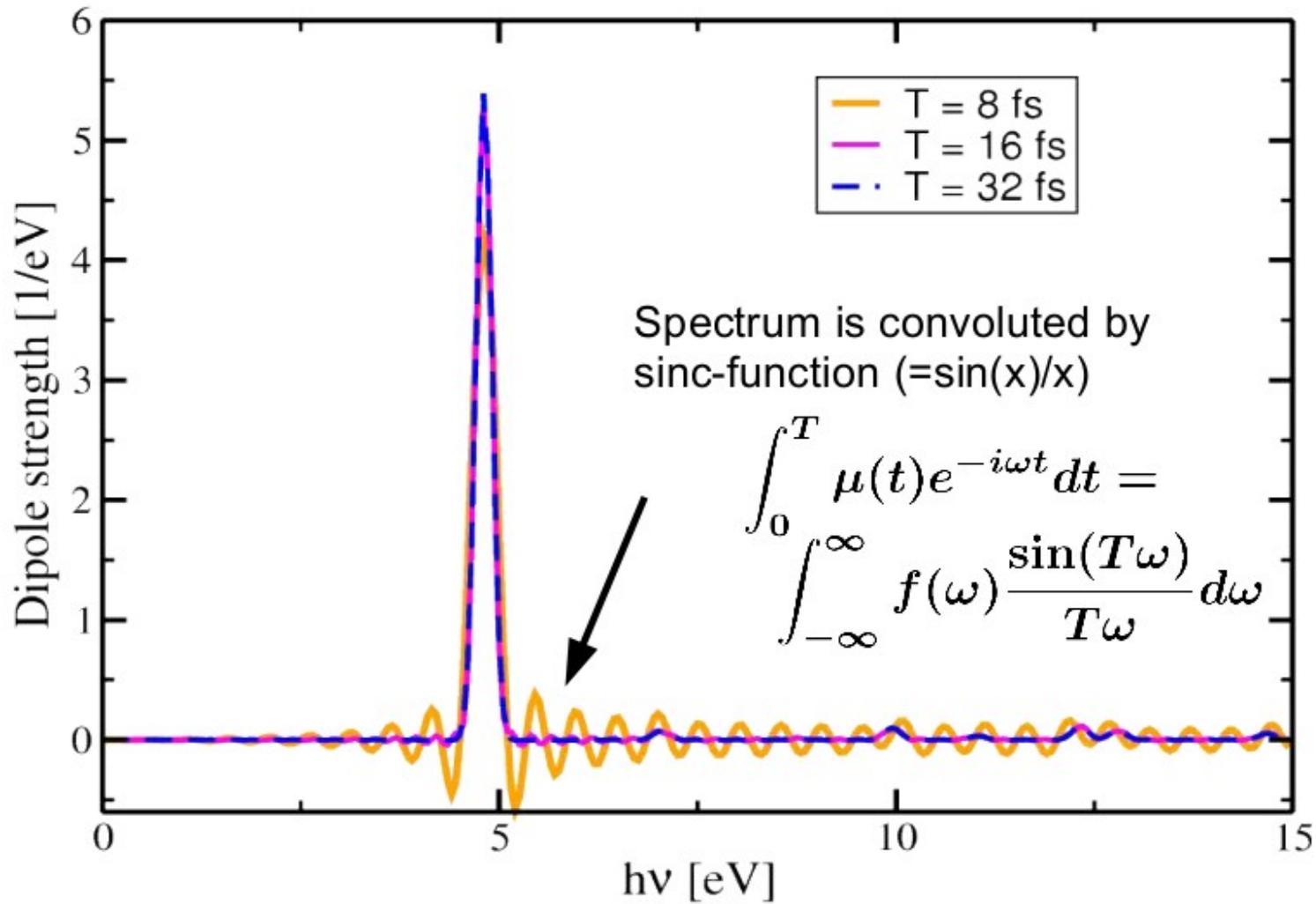


Eigenvalues are important, not forces or total energy!

# Time step



# Total simulation time



Double simulation time => double resolution



# Linear response TD-DFT

- Small perturbation to ground state potential

$$V_{ext} = V_{ext,0} + \delta V_{ext}(t)$$

results a change in density

$$n(r, t) = n_0(r) + n_1(r, t)$$

- Within linear response:

$$n_1(r, \omega) = \int d^3r' \chi(r, r', \omega) \delta V_{ext}(r', \omega)$$

- TD-DFT:

$$n_1(r, \omega) = \int d^3r' \chi_0(r, r', \omega) \delta V_{eff}(r', \omega)$$

- The interacting response function can be obtained from the non-interacting one



# Response function

- Dyson-like equation for density-density response:

$$\chi(\mathbf{r}, \mathbf{r}', \omega) = \chi_0(\mathbf{r}, \mathbf{r}', \omega) + \iint_{\Omega} d\mathbf{r}_1 d\mathbf{r}_2 \chi_0(\mathbf{r}, \mathbf{r}_1, \omega) K(\mathbf{r}_1, \mathbf{r}_2) \chi(\mathbf{r}_2, \mathbf{r}', \omega)$$

- Non-interacting response function is constructed from ground state Kohn-Sham orbitals

- Coupling kernel is:  $K(\mathbf{r}_1, \mathbf{r}_2) = \frac{1}{|\mathbf{r}_1 - \mathbf{r}_2|} + f_{xc}[n](\mathbf{r}_1, \mathbf{r}_2)$

with XC kernel  $f_{xc}(r_1, r_2) = \frac{\partial v_{xc}[n](r_1)}{\partial n(r_2)}$



# Response function

- Non-interacting response function is given by

$$\chi^0(\mathbf{r}, \mathbf{r}', \omega) = \sum_{\mathbf{k}, \mathbf{q}} \sum_{n, n'}^{\text{BZ}} \frac{f_{n\mathbf{k}} - f_{n'\mathbf{k}+\mathbf{q}}}{\omega + \epsilon_{n\mathbf{k}} - \epsilon_{n'\mathbf{k}+\mathbf{q}} + i\eta} \psi_{n\mathbf{k}}^*(\mathbf{r}) \psi_{n'\mathbf{k}+\mathbf{q}}(\mathbf{r}) \psi_{n\mathbf{k}}(\mathbf{r}') \psi_{n'\mathbf{k}+\mathbf{q}}^*(\mathbf{r}')$$

- Occupation numbers, eigenvalues and ground state orbitals of occupied and unoccupied states
- In extended systems one solves the Dyson equation in plane wave basis
  - ground state calculation can be done in any basis



# Dielectric matrix

- Dielectric matrix is related to the interacting response function

$$\epsilon_{\mathbf{G}\mathbf{G}'}^{-1}(\mathbf{q}, \omega) = \delta_{\mathbf{G}\mathbf{G}'} + \frac{4\pi}{|\mathbf{q} + \mathbf{G}|^2} \chi_{\mathbf{G}\mathbf{G}'}(\mathbf{q}, \omega)$$

- Macroscopic dielectric function

$$\epsilon_M(\mathbf{q}, \omega) = \frac{1}{\epsilon_{00}^{-1}(\mathbf{q}, \omega)}$$

- Optical spectra

$$\text{ABS} = \text{Im}\epsilon_M(\mathbf{q} \rightarrow 0, \omega)$$

- Electron-energy loss spectra

$$\text{EELS} = -\text{Im}\frac{1}{\epsilon_M(\mathbf{q}, \omega)}$$

# Excitation energies in finite systems



- Excitation energies can be calculated from eigenvalue equation:

$$\Omega F_I = \omega_I^2 F_I,$$

where

$$\Omega_{ij\sigma,kl\tau} = \delta_{ik}\delta_{jl}\delta_{\sigma\tau}\varepsilon_{ij\sigma}^2 + 2\sqrt{\varepsilon_{ij\sigma}\varepsilon_{kl\tau}}K_{ij\sigma,kl\tau},$$

$$\varepsilon_{ij} = \varepsilon_i - \varepsilon_j$$

with the coupling kernel

$$K_{ij\sigma,kl\tau} = \int dr_1 dr_2 n_{ij\sigma}^*(r_1) \left[ \frac{1}{|r_1 - r_2|} + f_{xc}(r_1, r_2) \right] n_{kl\tau}(r_2)$$

$$n_{ij\sigma}(r) = \psi_{i\sigma}(r)^* \psi_{j\sigma}(r)$$

- $i$  and  $j$  indexes go through occupied and unoccupied states, respectively

# Optical spectra in finite systems



- Dipole oscillator strengths of excitations can be calculated from the eigenvectors

$$f_{I\alpha} = \left| \sum_{ij\sigma} \langle \psi_{i\sigma} | r_{\alpha} | \psi_{j\sigma} \rangle \sqrt{\varepsilon_{ij\sigma}} F_I \right|^2$$

- Spectra with finite peak widths are obtained by folding the oscillators strengths e.g. with Gaussian

$$S_{\alpha}(\omega) = \sum_I f_{I\alpha} g(\omega - \omega_I)$$

# Practice



- Perform ground state calculation and include also unoccupied states

```
from ase.all import * # Setup the atomic simulation environment
from gpaw import GPAW # Setup GPAW
from gpaw.eigensolvers import CG # Conjugate gradient eigensolver

atoms = ...
atoms.center(vacuum=5.0) # More vacuum might be needed in reality...
calc = GPAW(nbands=1, h=0.30, txt='Na2_gs.txt')
atoms.set_calculator(calc)
e = atoms.get_potential_energy()

# Calculate also unoccupied states with the fixed density
eig = CG() # unoccupied states converge often better with cg
calc.set(nbands=20, convergence={'bands': 'all'}, # converge unoccupied states
        eigensolver=eig,
        fixdensity=True)
e = atoms.get_potential_energy()
# write the wave functions to a file
calc.write('na2_gs.gpw', 'all')
```

# Practice



## ➔ Calculate $\Omega$ matrix

```
from gpaw import * # Setup GPAW
from gpaw.lrtddft import *

atoms, calc = restart('na2_gs.gpw') # read in a ground state calculation

# Calculate the omega matrix
lr = LrTDDFT(calc, xc='LDA')
# Save the omega matrix
lr.write('Omega_Na2.gz')
```

## ➔ Diagonalization can be performed in separate step

```
from gpaw.lrtddft import *

# Read the omega matrix from a file
lr = LrTDDFT(filename='Omega_Na2.gz')
# Diagonalize the matrix
lr.diagonalize()
# Print out five lowest excitations
lr.analyse(range(5))
# Calculate the absorption spectrum and save it to a file
photoabsorption_spectrum(lr, 'Na2_spectrum.dat', e_min=0.0, e_max=10, width=0.1)
```



# Practical details

- In addition to the box size and grid spacing, accuracy is controlled by the number of electron-hole pairs
- The computational intensity of the calculation is  $O(N_{eh}^3)$
- The size of electron-hole basis can be reduced:

```
lr = LrTDDFT(calc, xc='LDA',  
            istart=5,      # first occupied state to consider  
            jend=20)     # last unoccupied state to consider
```

- The proper number of eh-pairs is very system dependent

# Real-time vs. linear response



**GPAW!**

## ➤ Real-time

- only excitations corresponding to given perturbation
- non-linear effects
- scales  $O(N^2)$  with the system size, large prefactor
- time step controls the accuracy relatively straightforwardly

## ➤ Linear response

- all excitations (within linear response)
- scales  $O(N^6)$  with the system size, small prefactor
- control of accuracy by the electron-hole basis size can be complex



# Parallel calculations with GPAW





# Parallelization levels

$$H_{k,s}\psi_{i,k,s}(\mathbf{r}) = e_i S \psi_{i,k,s}(\mathbf{r})$$

$$i\partial_t\psi_{i,s}(\mathbf{r}, t) = H_s\psi_{i,s}(\mathbf{r}, t)$$

$$n(\mathbf{r}) = \sum_{i,k,s} |\psi_{i,k,s}(\mathbf{r})|^2$$

- Parallelization over all degrees of freedom
  - real-space grid
  - k-points and spin
  - electronic states
- Additional trivial parallelizations possible
  - Electron-hole pairs
  - different atomic configurations or unit cells

# Parallelization over k-points and spin



$$H_{\mathbf{k},s}\psi_{i,\mathbf{k},s}(\mathbf{r}) = e_i\psi_{i,\mathbf{k},s}(\mathbf{r})$$

$$n(\mathbf{r}) = \sum_{i,\mathbf{k},s} |\psi_{i,\mathbf{k},s}(\mathbf{r})|^2$$

- Spin and k-points are treated equivalently
- Trivial parallelization
- Limited scalability
  - k-points only in (small) periodic systems
  - spin only in magnetic systems
- All basis sets

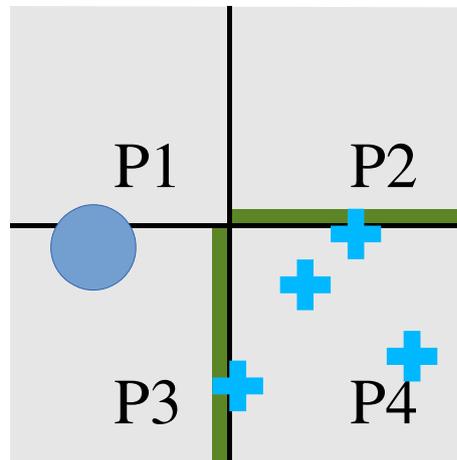
# Parallelization over real-space grid



$$H_{k,s}\psi_{i,k,s}(\mathbf{r}) = e_i\psi_{i,k,s}(\mathbf{r})$$

$$n(\mathbf{r}) = \sum_{i,k,s} |\psi_{i,k,s}(\mathbf{r})|^2$$

## Domain decomposition



Finite difference Laplacian



PAW augmentation sphere



- Only local communication
- Good parallel scalability down to domain sizes  $\sim 16 \times 16 \times 16$
- Not available in plane wave mode

# Parallelization over electronic states



$$H_{k,s}\psi_{i,k,s}(r) = e_i\psi_{i,k,s}(r)$$

$$n(r) = \sum_{i,k,s} |\psi_{i,k,s}(r)|^2$$

- Nearly trivial parallelization in real-time TDDFT
  - (similar to k-points)
  - good scalability down to 20 states per process
- Orthonormalizations are complicated in ground state DFT
  - communication of all wave functions to all processes
  - parallel scalability down to 150-250 states per process
  - all basis sets

# Parallelization over electron-hole pairs



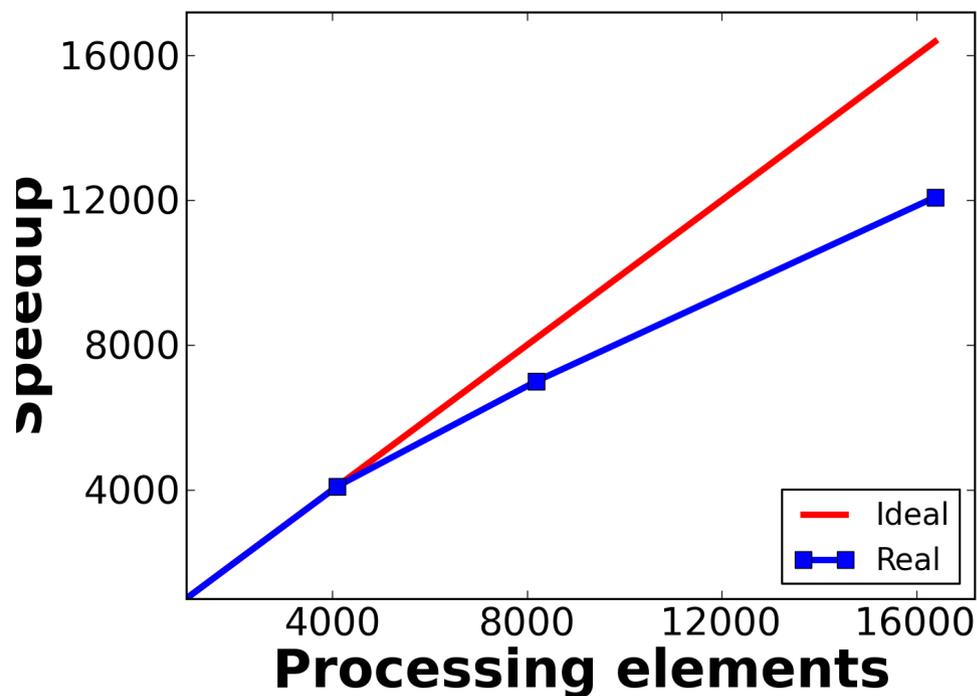
- Casida equation in linear response TD-DFT:

$$\Omega F_I = \omega_I^2 F_I,$$

$$\Omega_{ij\sigma,kl\tau} = \delta_{ik}\delta_{jl}\delta_{\sigma\tau}\varepsilon_{ij\sigma}^2 + 2\sqrt{\varepsilon_{ij\sigma}\varepsilon_{kl\tau}}K_{ij\sigma,kl\tau}$$

- Matrix elements can be calculated independently
- Nearly trivial parallelization over electron-hole pairs **ij**
- Domain decomposition for individual matrix elements

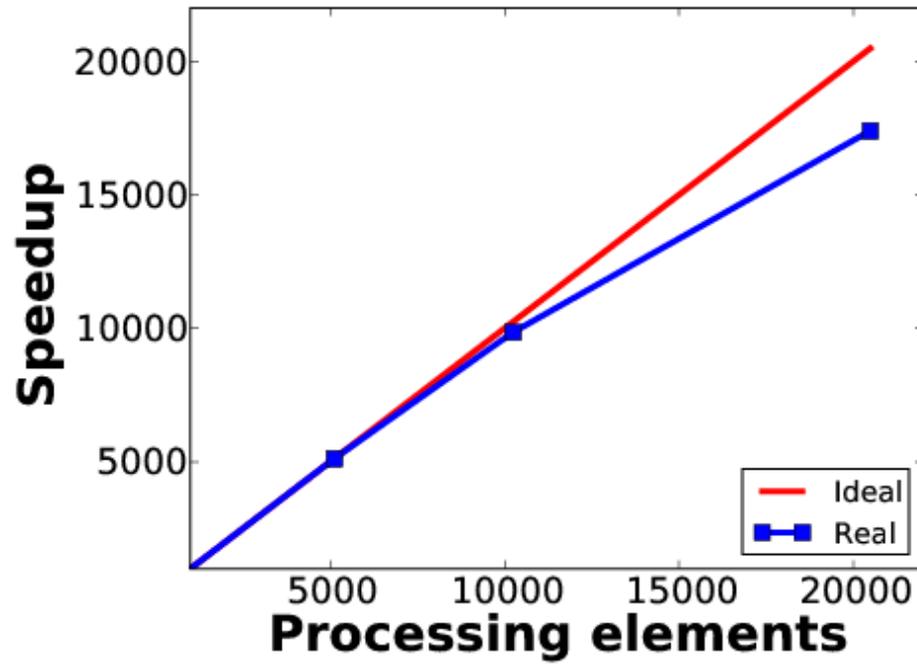
# Parallel scalability



- Ground state DFT
  - 561 Au atom cluster
  - ~6200 electronic states
  - Blue Gene P, Argonne



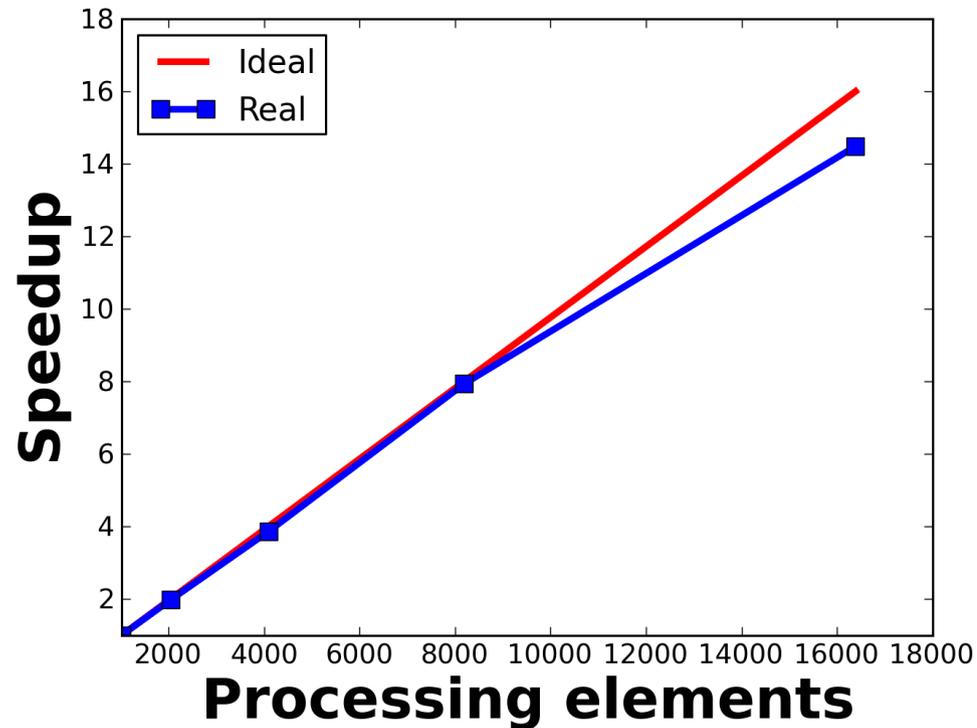
# Parallel scalability



- Real-time TD-DFT
  - 702 Si atom cluster
  - ~2800 electronic states
  - Cray XT5 Jaguar, Oak Ridge

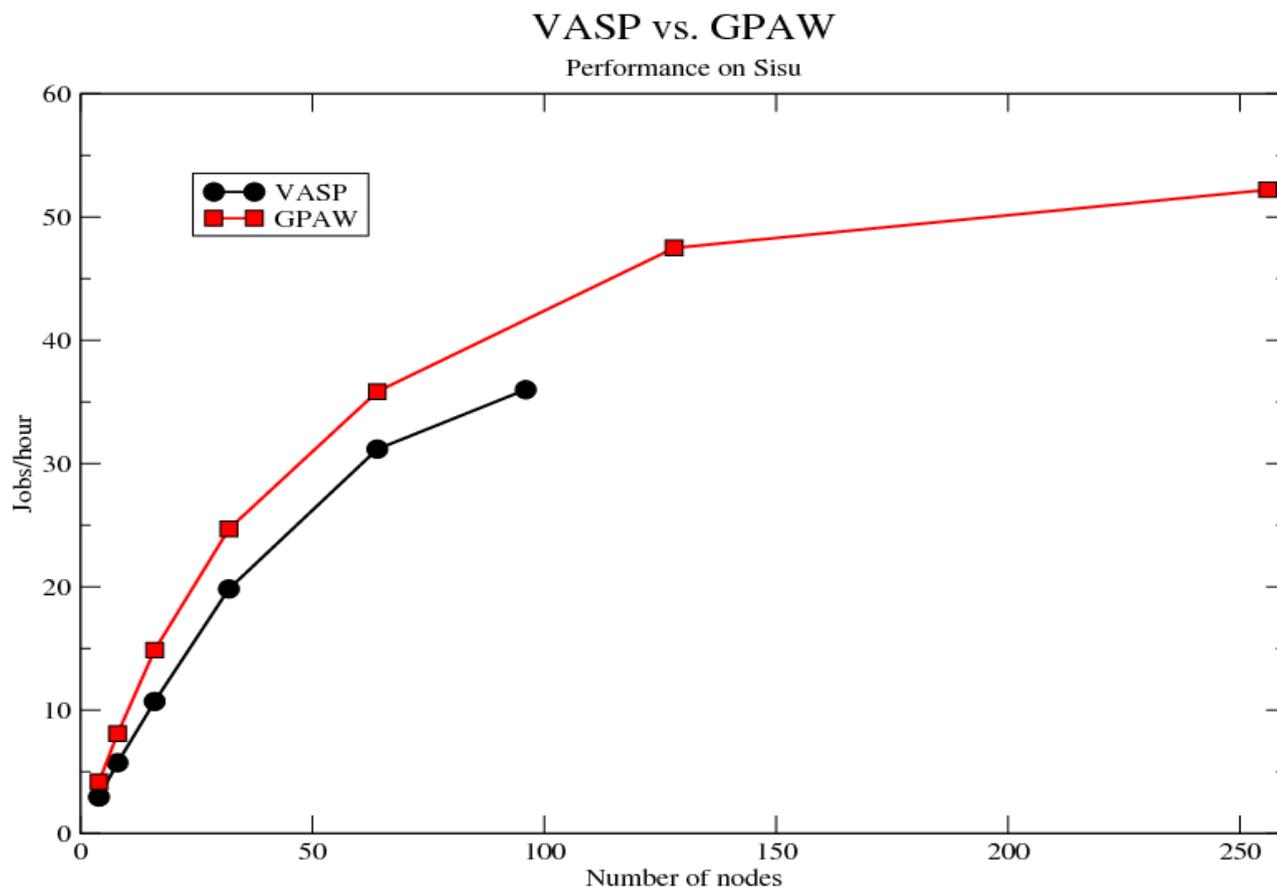


# Parallel scalability



- Linear-response TD-DFT
  - Au<sub>38</sub>(SCH<sub>3</sub>)<sub>24</sub> cluster ~160 atoms
  - ~680 electronic-hole pairs
  - Cray XE6 Hermit, HLRS, Germany

# Parallel performance



## ➤ Ground state DFT

- Large MgH<sub>2</sub> cell, 1296 atoms
- Cray XC30, CSC (node=16 cores)



# Practical details

- k-points have to be distributed evenly
  - same number of k-points in each process
- Electronic states have to be distributed evenly
  - same number of states in each process
- In principle, arbitrary number of processes can be used for domain decomposition
  - recommended to use cubic domains e.g.  $4 \times 4 \times 4$
  - recommended to use domain dimensions which factor the number of grid points



# Practical details

- By default, k-point and spin are distributed first, and the remaining processors are used for domain decomposition
- Example: magnetic system, 5 k-points
  - with 20 processors: 10 (=2x5) processors for spin/k-point and 2 processors for domain decomposition (2x1x1 layout)
  - with 24 processors: 2 processors for spin, 12 processors for domain decomposition (3x2x2 layout)



# Practical details

- Electronic state parallelization has to be specified explicitly

```
aprun -n 512 gpaw-python input.py --state-parallelization=2
```

- 2 processors for states, 256 to k-points/spin/domains

- For large calculations (> 1000 states) one more command line argument:

```
aprun -n 512 gpaw-python input.py --sl_default=4,4,64
```

- some large matrix diagonalizations are done in parallel with 16 (=4x4) processes
- 4x4 or 8x8 are typically good values (block size 64 has only small effect)



# Practical details

- Parallelization options can be given also as GPAW calculator parameters

```
...
calc = GPAW(...
    parallel={kpt : 4,          # k-point parallelization with 4 MPI tasks
              domain : 8,     # domain decomposition with 8 MPI tasks
              band : 2        # band parallelization with 2 MPI tasks
              sl_default : (4,4,64)} # parallel matrix diagonalization
    ...
)
```

- Command line arguments precede calculator parameters



# “dry-run” mode

- Often, it is desirable to check system parameters without an actual calculation
- GPAW offers a dry-run mode

```
[cscuser@cobol ~]$ python input.py --state-parallelization=2 --dry-run=512
```

- only unexpensive initializations, can be run serially
- simulates parallel calculations and shows the parallelization scheme
- estimates the memory usage