

THE UNIVERSITY *of York*

Geometry Optimization

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- Background Theory
 - Hellman-Feynman theorem
 - BFGS and other algorithms
- CASTEP details
 - Useful keywords for geometry optimization
 - Variable cell – additional considerations

Background Theory

- Classically we have the force $\mathbf{F}$ at position $\mathbf{R}$ is determined from the potential energy as

$$\mathbf{F} = -\frac{\partial U(\mathbf{R})}{\partial \mathbf{R}}$$

- Quantum mechanically we therefore have

$$\mathbf{F} = -\frac{\partial \langle E \rangle}{\partial \mathbf{R}} = -\frac{\partial}{\partial \mathbf{R}} \frac{\langle \Psi | H | \Psi \rangle}{\langle \Psi | \Psi \rangle}$$

- For a position-independent basis set

$$\mathbf{F} = -\frac{\partial \langle E \rangle}{\partial \mathbf{R}} = -\left\langle \Psi \left| \frac{\partial H}{\partial \mathbf{R}} \right| \Psi \right\rangle$$

- In DFT we have the Kohn-Sham Hamiltonian:

$$\hat{H}(\mathbf{r}, \mathbf{R}) = -\frac{1}{2} \nabla_{\mathbf{r}}^2 + V_{e-e}(\mathbf{r}) + V_{ion-e}(\mathbf{r}, \mathbf{R}) + V_{XC}(\mathbf{r}) + V_{ion-ion}(\mathbf{R})$$

- Therefore we only get contributions to the forces from the electron-ion (pseudo)potential and the ion-ion Coulomb interaction (the Ewald sum).
- As we do not have a complete basis, the wavefunction will not be exact even within DFT.
- Need to test for convergence carefully wrt cut-off energy and k-points etc.

- With a variational minimization of the total energy, the energy and wavefunction will be correct to second order errors.
 - Forces are given by energy differences and hence get some error cancellation
- With a non-variational minimization technique, such as *density mixing*, then no upper-bound guaranteed on true E_0
 - So need good convergence if using non-variational forces and stresses

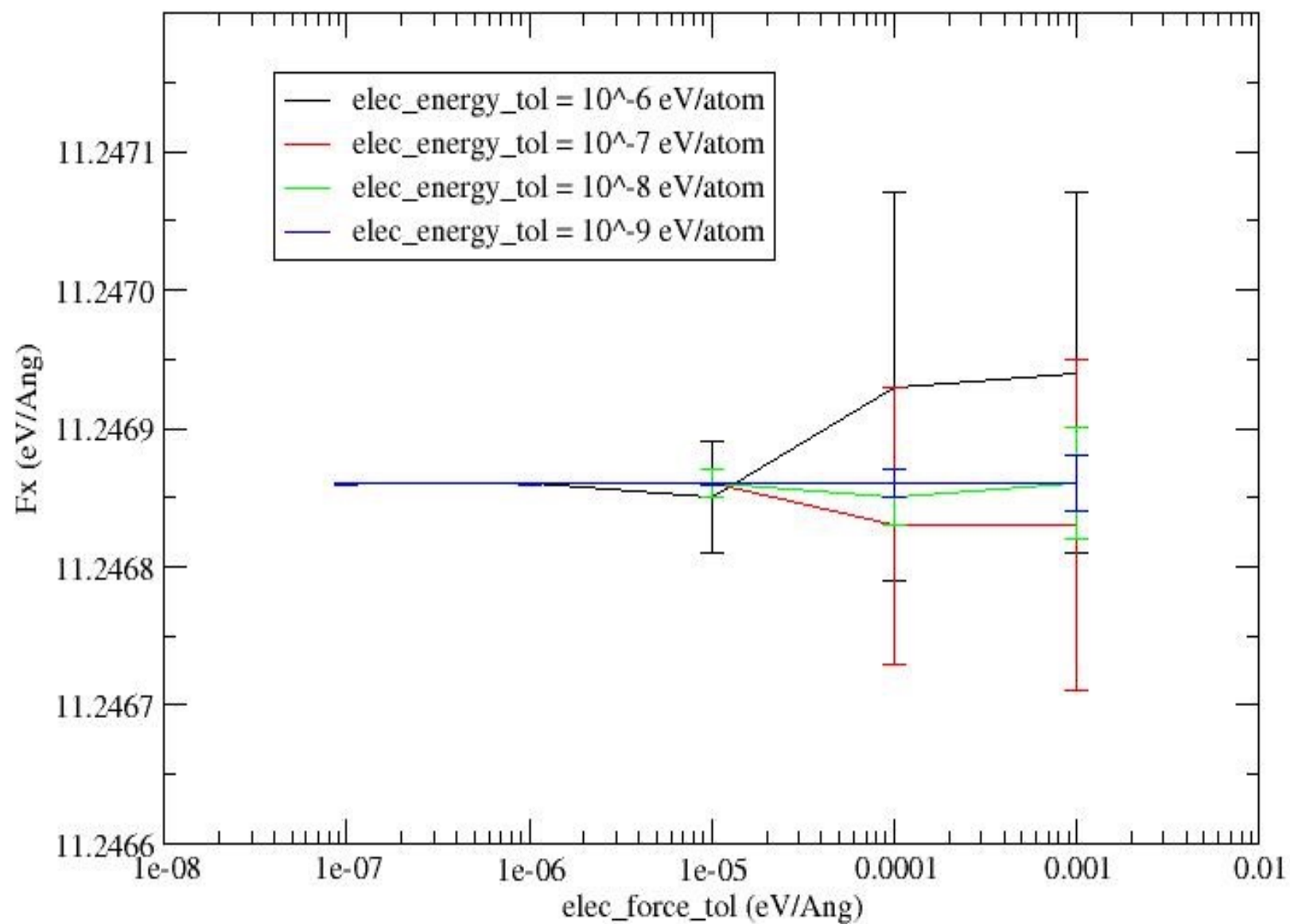
- Can only find a good structure if have reliable forces/stresses
- Stresses converge more slowly than energies as increase number of plane waves
 - CONVERGENCE!
- Should also check degree of SCF convergence

elec_energy_tol (fine quality $\leq 10^{-6}$ eV/atom)
can also set **elec_force_tol** – useful with DM

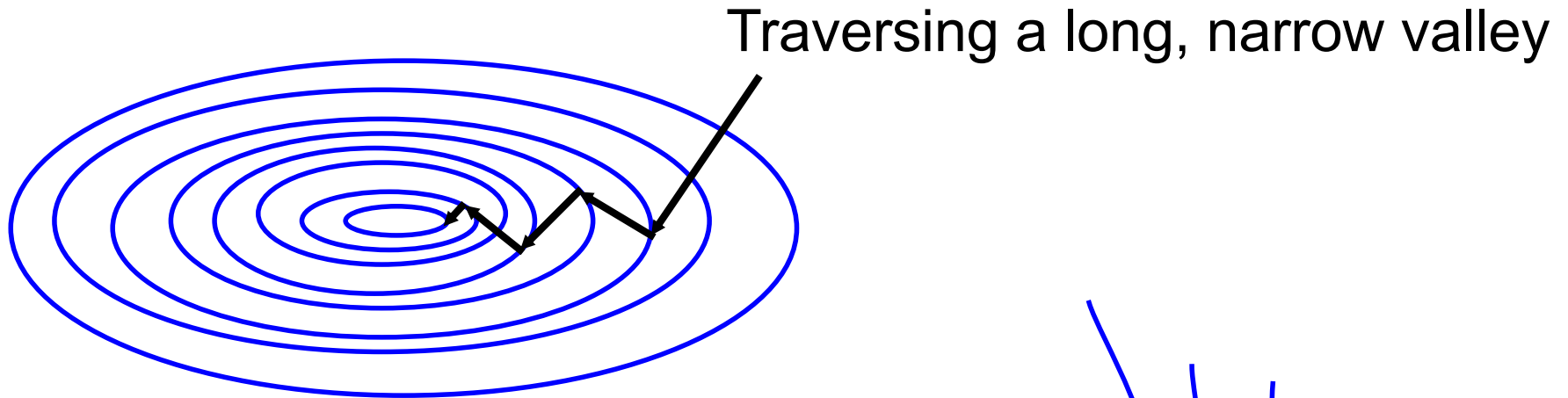
Stresses converge slower than forces!

Force on stretched N2 dimer

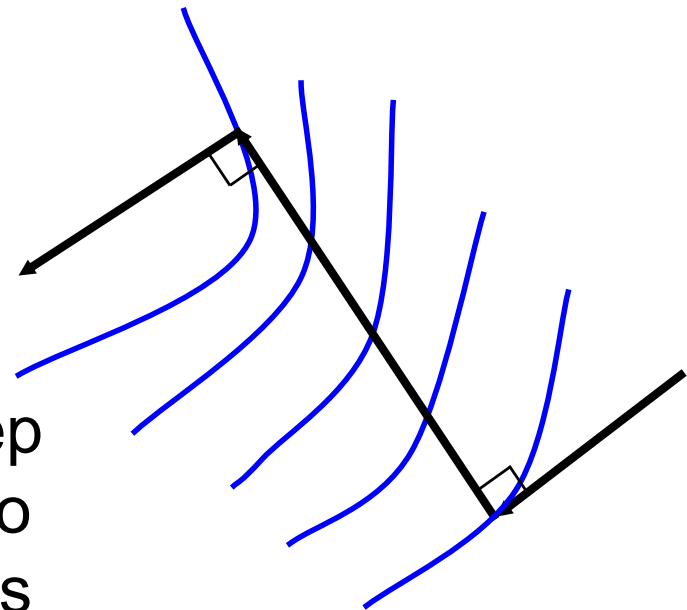
DM, grid_scale=1.5, Ecut=500 eV, recpot, LDA

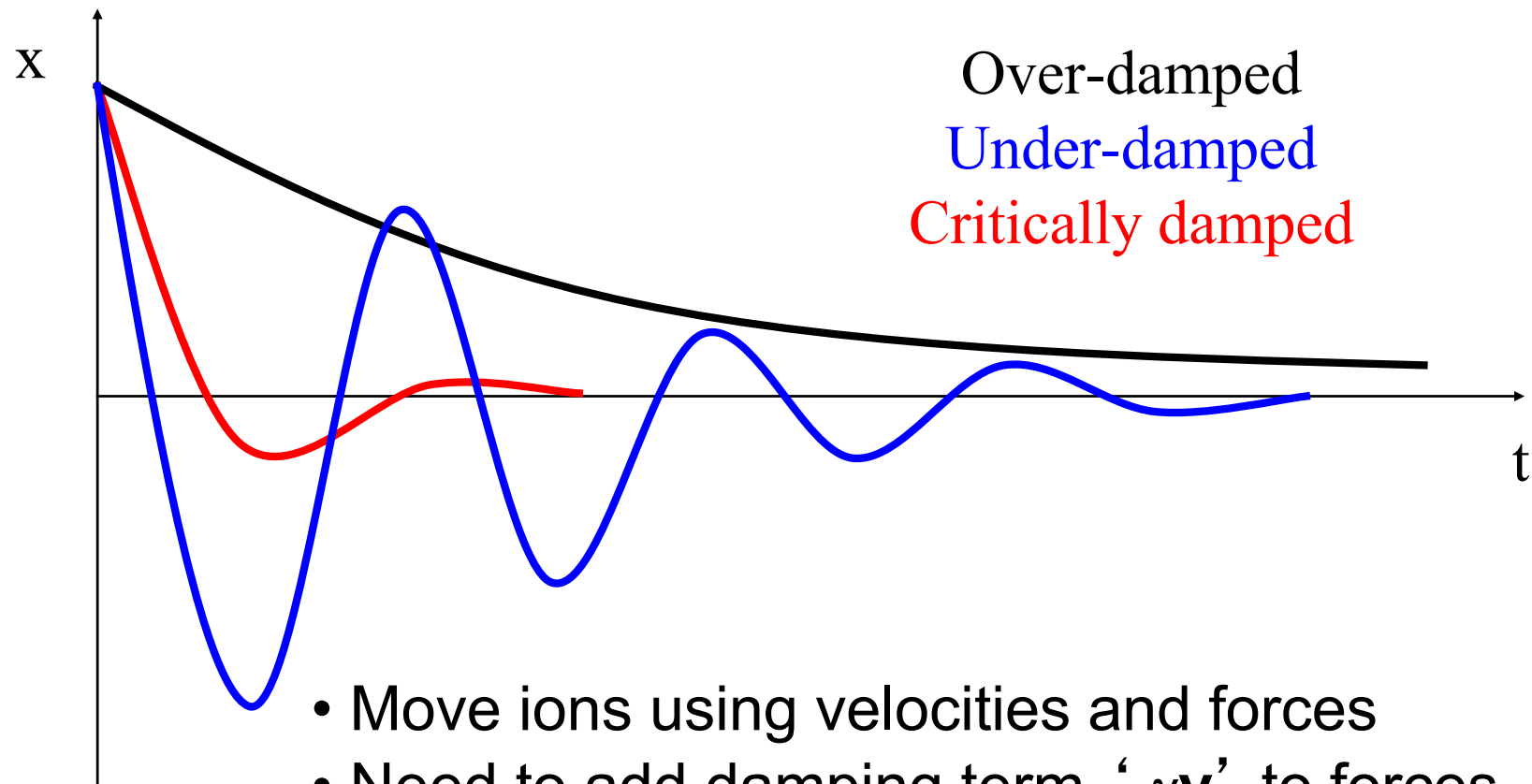


- Electrons adjust instantly to position of ions
 - Multi-dimensional potential energy surface and want to find global minimum
- Treat as an optimisation problem
 - Simplest approach is steepest descents
 - More physical approach is damped MD
 - More sophisticated approaches are conjugate gradients or BFGS
 - All of these can get stuck in local minima
 - Hence recent research in how best to find global minimum

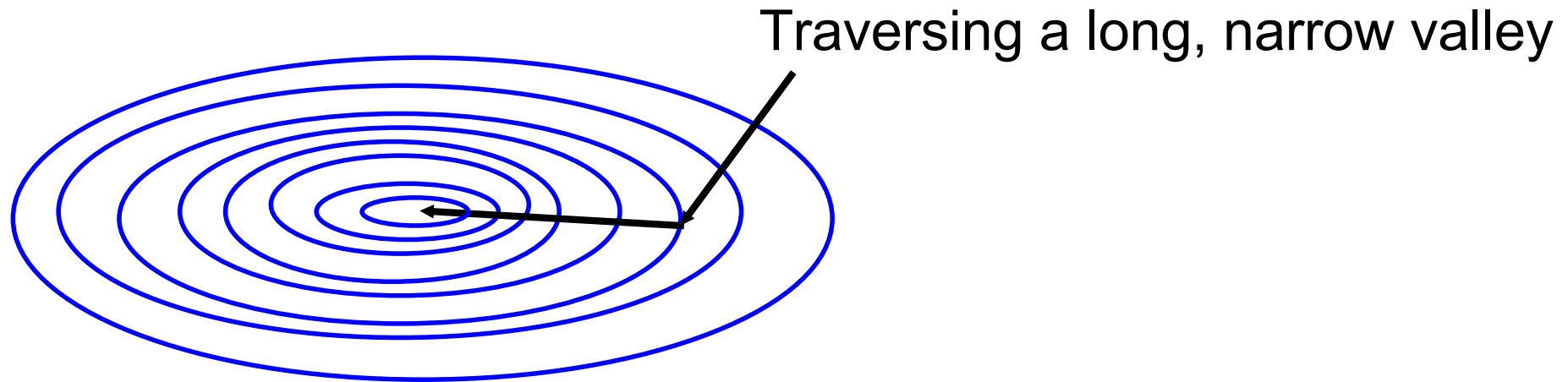


Enlargement of a single step showing the line minimisation in action – the step continues until the local energy starts to rise again whereupon a new direction is selected which is orthogonal to the previous one





- Move ions using velocities and forces
- Need to add damping term ' $-\gamma \mathbf{v}$ ' to forces
- Algorithm to set 'optimal damping' factor
- Can also adjust δt for max. convergence
- More efficient than Steepest Descents



- The initial search direction is given by steepest descents.
- Subsequent search directions are constructed to be orthogonal to the previous *and* all prior search directions.
- No explicit Hessian required or generated.

- Basic idea:

- Energy surface around a minima is quadratic in small displacements and so is totally determined by the Hessian matrix **A** (the matrix of second derivatives of the energy):

$$\mathbf{A} = \begin{pmatrix} \frac{\partial^2 E}{\partial x_1 \partial x_1} & & \frac{\partial^2 E}{\partial x_1 \partial x_N} \\ & \ddots & \\ \frac{\partial^2 E}{\partial x_N \partial x_1} & & \frac{\partial^2 E}{\partial x_N \partial x_N} \end{pmatrix}$$
$$\delta E = \frac{1}{2} (\mathbf{X} - \mathbf{X}_{\min})^T \bullet \mathbf{A} \bullet (\mathbf{X} - \mathbf{X}_{\min})$$

- so if we knew **A** then could move from any nearby point to the minimum in 1 step!

■ The Problem

- We do not know $\mathbf{A}$ *a priori*
- Therefore we build up a progressively improving approximation to $\mathbf{A}$ (or the inverse Hessian $\mathbf{H}=\mathbf{A}^{-1}$) as we move the ions according to the BFGS (Broyden-Fletcher-Goldfarb-Shanno) algorithm.
- Also known as a *quasi-Newton* method.
- Positions updated according to:
$$\mathbf{X}_{i+1} = \mathbf{X}_i + \lambda \Delta \mathbf{X}_i$$
$$\Delta \mathbf{X}_i = \mathbf{H}_i \mathbf{F}_i$$

CASTEP details

- Just put

task = Geometry Optimisation

in your `.param` file

- Is that all?
- What is going on behind the scenes?
- What might go wrong?
- What can you control?

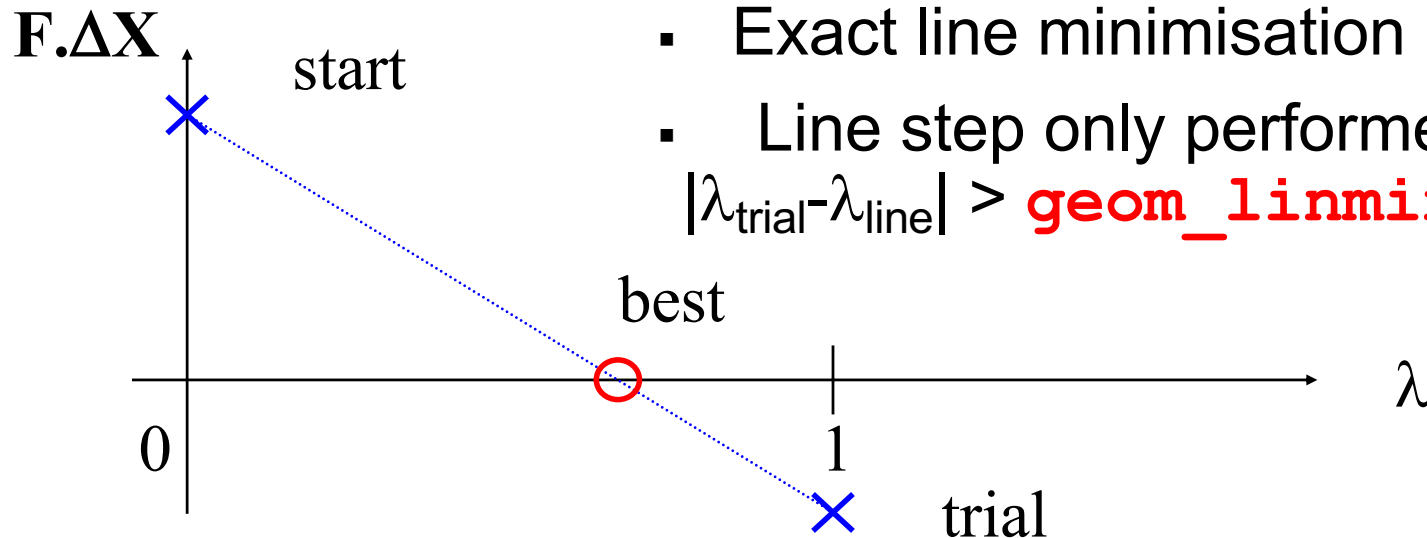
-
- **geom_method = LBFGS** (default)
 - variable ions and/or cell
 - uses fractional ionic coordinates and strains as basic variables
 - improved estimates based analysis of approx. **H** built up over convergence path printed at end
 - a “low memory” version of **geom_method = BFGS** as it only stores a limited number of updates and not full Hessian

- **geom_method = TPSD**
 - Very low memory requirements
 - No line search and no history
 - Ought to be a lot worse than (L)BFGS but we have a smart preconditioner & so not so bad
 - Much better than (L)BFGS with constraints
 - Much better if start a long way from the harmonic region
 - Can use **geom_tpsd_iterchange** to switch to BFGS after given number of TPSD steps
 - Can use **geom_tpsd_init_stepsize** to control size of initial step

$$\mathbf{X}_{i+1} = \mathbf{X}_i + \lambda \Delta \mathbf{X}_i$$

$$\Delta \mathbf{X}_i = \mathbf{H}_i \mathbf{F}_i$$

- Trial step performed (default) if **geom_use_linmin=true**
- A quadratic system has optimal $\lambda=1$
- Exact line minimisation has $\mathbf{F} \cdot \Delta \mathbf{X} = 0$
- Line step only performed if predicted $|\lambda_{\text{trial}} - \lambda_{\text{line}}| > \text{geom_linmin_tol}$



No restriction on the value of λ as long as does not cause too large an ionic displacement or change in lattice parameters – big advantage of doing a ‘line step’

- Will continue for **geom_max_iter** steps (default=30) unless converged:
geom_energy_tol (default= 2×10^5 eV/atom)
for **geom_convergence_win** steps

Changing ions:

geom_force_tol (default = 0.05 eV/Å)

geom_disp_tol (default=0.001 Å)

Changing cell:

geom_stress_tol (default=0.1 GPa)

Variable cell calculations

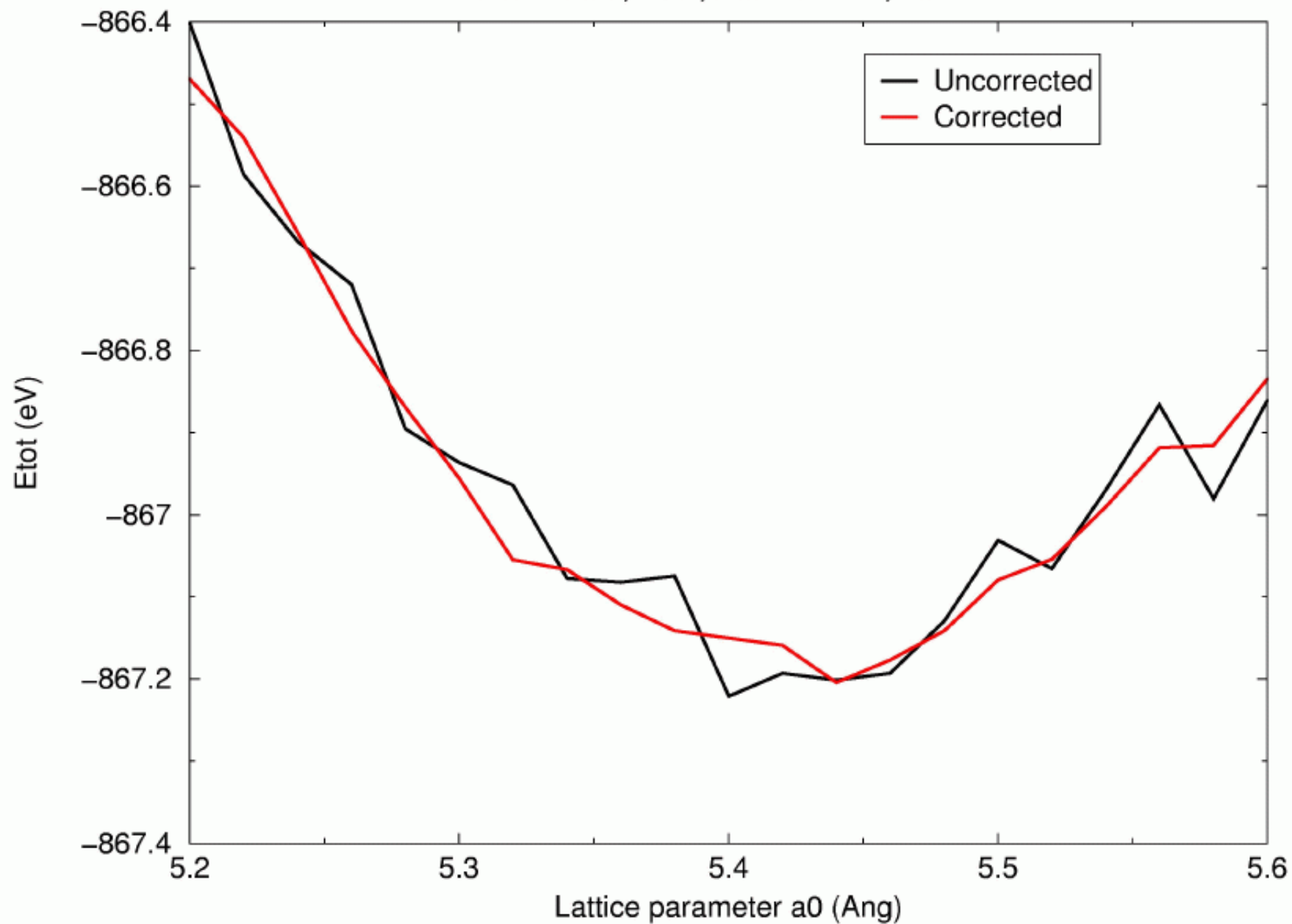
- If can calculate internal stress then can use this to adjust the cell parameters
- Convention: $\mathbf{P}_{\text{ext}} + \sigma = 0$ at equilibrium
 - i.e. $P > 0$ = compression, $P_{\text{int}} = 1/3 \text{Tr}(\sigma)$
 - Look for a state of minimum enthalpy
 - $H = E + P_{\text{ext}}V$
- With BFGS we use an augmented Hessian
 - Work in space of “fractional positions and strains” with “fractional forces and stresses”
 - Then possible to mix cell & ion terms

- Plane wave basis is independent of ionic positions but NOT of the unit cell size/shape
- **fix_NPW=true**
 - the number of plane waves is fixed
 - hence the variational principle applies and we can search for the minimum of enthalpy as cell changes
 - but the effective cut-off energy varies!
- **fix_NPW=false**
 - number of plane waves varies to keep E_{cut} constant
 - breaks variational principle
 - search for zero force & stress not minimum enthalpy
 - but physically more reasonable

- Change unit cell at constant **cut_off_energy** :
 - changes number of plane-waves
 - change in total energy due to variational principle
 - hence difficult to compare results at different cell sizes.
- Use the Finite Basis Set Correction
 - **finite_basis_corr = automatic/manual/none**
 - calculates the total energy **finite_basis_npoints** times with **finite_basis_spacing** change to **cut_off_energy** at fixed cell
 - hence calculates and prints **basis_de_dloge**
 - CASTEP can then use this to correct the total energy and stress at nearby cell sizes

Finite-Basis Set Correction

Perfect Si8, USP, Ecut=120 eV, MP=2



fix_all_ions (default = false)

fix_all_cell (default = false)

fix_com (default = NOT `fix_all_ions`)

symmetry_generate

snap_to_symmetry

%block external_pressure

[units]

P_{xx} P_{xy} P_{xz}

P_{yy} P_{yz}

P_{zz}

%endblock external_pressure

Hence hydrostatic P is $P_{xx}=P_{yy}=P_{zz}=P$ and $P_{xy}=P_{xz}=P_{yz}=0$

```
%block cell_constraints
```

```
|a| |b| |c|
```

```
 $\alpha$   $\beta$   $\gamma$ 
```

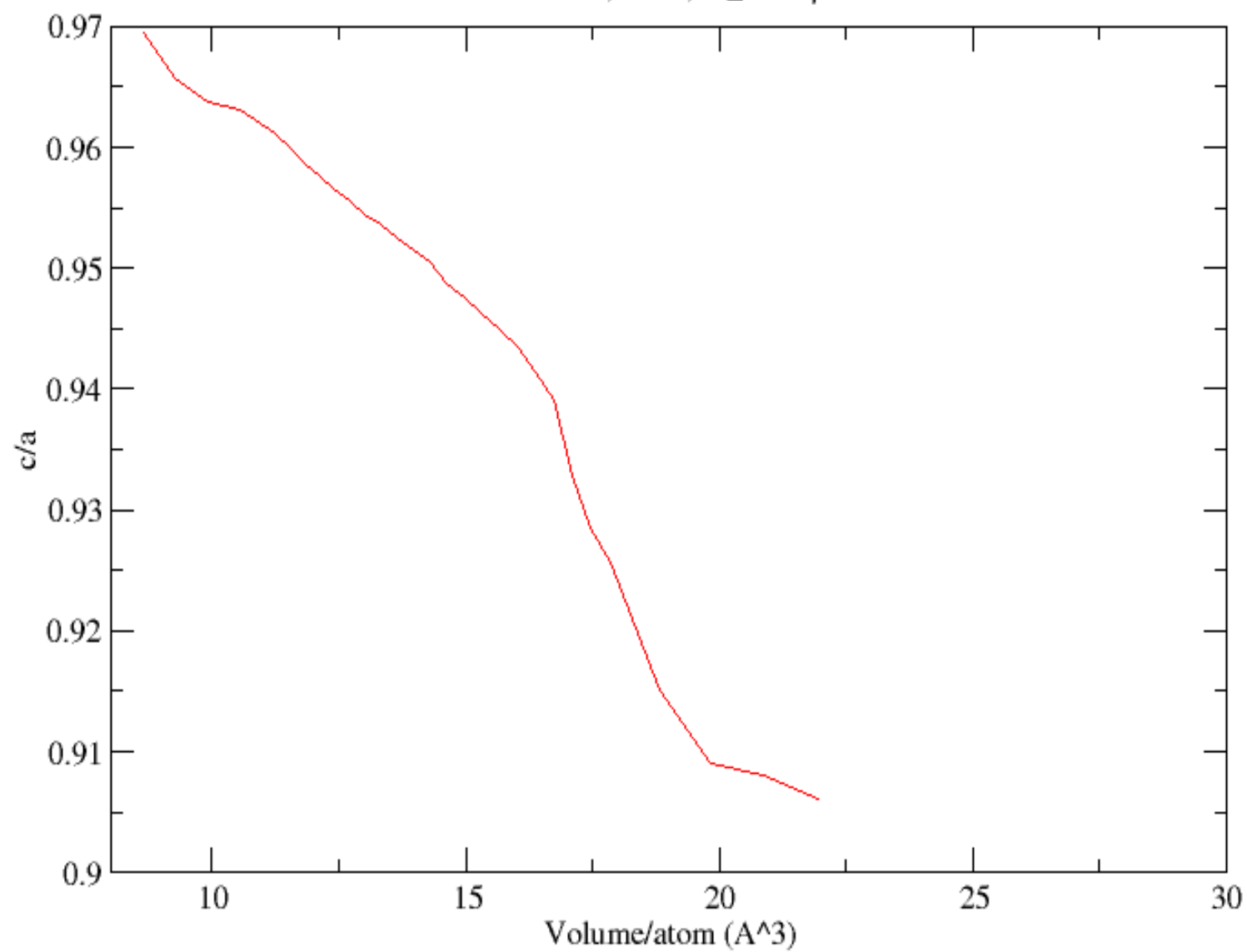
```
%endblock cell_constraints
```

- Any length (angle) can be held constant (=0), or tied to one or both of the others (=same)
 - e.g. cell optimisation of 2D structures or keeping subset of symmetry etc
 - also **fix_vol** (= false by default)

fix_vol in action

P6mmm Optimal c/a ratio

180 eV, LDA, Si_00.usp



- Records the final configuration after each step:

```
10
-7.93316351E+000 -7.85316331E+000 <-- E
0.00000000E+000 5.13126785E+000 5.13126785E+000 <-- h
5.13126785E+000 0.00000000E+000 5.13126785E+000 <-- h
5.13126785E+000 5.13126785E+000 0.00000000E+000 <-- h
-3.56997760E-003 0.00000000E+000 -3.33783917E-013 <-- S
0.00000000E+000 -3.56997760E-003 8.32597229E-013 <-- S
-3.33783917E-013 8.32597229E-013 5.93008591E-004 <-- S
Si 1 0.00000000E+000 0.00000000E+000 0.00000000E+000 <-- R
Si 2 7.56877069E+000 2.52292356E+000 7.56877069E+000 <-- R
Si 1 -5.22300739E-003 6.43530285E-003 -1.71774942E-003 <-- F
Si 2 5.22300739E-003 -6.43530285E-003 1.71774942E-003 <-- F
```

- Uses Cartesian coordinates and atomic units throughout
- Designed for other analysis programs, visualisation tools, etc.

- Final geom “trajectory” can be visualized using jmol – just drag & drop `.geom` file
- Or can add `write_cell_structure=true` or `write_cif_structure=true` to output final structure in cell/cif format
- Or can use utilities such as `geom2xyz` to convert `.geom` file to `.xyz` format for visualization by many packages

- Can specify any arbitrary number of *linear constraints* on the atomic coordinates, up to the number of degrees of freedom.
 - E.g. fixing an atom, constraining an atom to move in a line or plane, fixing the relative positions of pairs of atoms, fixing the centre of mass of the system, etc.
- Non-linear constraints
 - Much more difficult to apply and specify in general, e.g. fixing a bond length
 - Only supported by Delocalised Internals

```
%block ionic_constraints
```

```
con spec atom Ax Ay Az
```

```
. . .
```

```
%endblock ionic_constraints
```

where each constraint fixes 1 degree of freedom

- `con` is the number of the constraint (can have multiple atoms/constraint) and multiple constraints/calculation
- `spec` is the species label, `atom` is the number of the atom for this species
- A_x, A_y, A_z specify the constraint coefficient such that
$$\sum \mathbf{A} \cdot \mathbf{r} = \text{const}$$

General format:

```
%block ionic_constraints
```

```
1 Si 1 0 0 1
```

```
%endblock ionic_constraints
```

fixes the z-coordinate of the 1st Si atom

Shortcut for fixing 1 or more atoms:

```
%block ionic_constraints
```

```
fix: Si 1
```

```
%endblock ionic_constraints
```

Can also do things like `fix: all unfix: H`
to fix all atoms except H etc.

Summary

- Need accurate forces and stresses
- Pre-requisite for many property calculations
 - e.g. phonons, NMR, etc.
- Can do optimisation of ions and/or cell
 - (L)BFGS or TPSD with Cartesian coords
- Can also for fixed cell use
 - BFGS with Cartesian or delocalised internal coordinates, or
 - damped MD or FIRE with Cartesians

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