How to run VASP Part 1: Influential settings

Quick summary

- Processor cores ≤ atoms
- NCORE = cores per node
 NPAR = compute nodes
- **NSIM** = 1 (Intel), 4 (AMD)
- **KPAR** = compute nodes
- **ALGO** = Fast (or VeryFast)
- **NBANDS** = *N**cores
- Use the right VASP binary (*NGZhalf*, and *wNGZhalf* for gamma-point calculations)

Q: What's missing?

A: Read the VASP manual!

Efficiency:

Running as many jobs as possible within a given allocation of computer time.

Speed:

The amount of time (in real time, "human time") to run a specific simulation from the time it starts.

Time-to-solution:

The amount of time (in real time, "human time") it takes to get the results = runtime + the time waiting in the queue.

Efficiency

Using as few cores as possible

Speed

As many cores as possible (up to a limit)

Time-to-solution

- Finding the "right" size which fits the cluster.
- Not using up all your allocated time (in order to get higher priority in the queue).



Safe & Efficient: cores < n(atoms)

Daring & not as efficient: cores ≈ 2 x n(atoms)

> Crash! cores > n(bands)

Why should I care?





We regret to inform you that we have decided to cut your VSC time allocation from 100,000 core hours/month to 90,000 core hours/month.

Does 10% matter?

Best regards, VSC Staff

NPAR & NCORE

- Applies to the Davidson/RMM-DIIS algorithms (not hybrid calculations)
- Controls/activates band parallelization
- NPAR = 1: no parallelization over bands = BAD
- NPAR = (1-2)x number of compute nodes
- NCORE = cores working on one band

NCORE = cores per node (or socket)

Li2FeSiO4 (128 atoms) on Triolith

Standard DFT / 4 compute nodes / 64 cores



Speed-up with optimal NPAR FeCo alloy (53 atoms)



NPAR: caveat

Number of SCF iterations vs NPAR *FeCo alloy (53 atoms) on 48 cores*



K-point parallelization ("KPAR")

- Controls parallelization over k-points (version 5.3.2+)
- Great for hybrid-DFT jobs
- Not so great for metals with 100-1000s of k-points...

KPAR = "number of k-points treated in parallel" **cores / KPAR** = "group size" = cores/node?

KPAR = min(nodes, #kpts)

MgO (63 atoms) on Triolith HSE06 hybrid calculation with 4 k-points

KPAR=1 KPAR=2 KPAR=4



NSIM

- Blocking mode for RMM-DIIS algorithm (so only applies for ALGO=Fast/VeryFast)
- Effect depends on BLAS library. Less important nowadays?
- NSIM = 1 for single-node jobs on Intel Sandy Bridge
- NSIM = 4 for older systems (incl. AMD). This is VASP's default.

Li2FeSiO4 (128 atoms) on Triolith

Standard DFT / 4 compute nodes / 64 cores



PbSO4 (24 atoms) on Triolith

Standard DFT / 1 compute node / 16 cores



Core binding / pinning

- It is important to lock an MPI process to a specified processor core. Otherwise they can jump around...
- Intel MPI binds by default.
- Older versions of OpenMPI does not. Try e.g.:

mpirun -np 32 --display-map —bind-to-core ...

 Core binding on AMD Opteron 6300-series "Interlagos" processors is not intuitive. [Cf. one of my blog posts]