



Using VASP at NERSC

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Outline

- Getting started with VASP at NERSC
- Running VASP on Perlmutter
- Troubleshooting VASP
- VASP performance on Perlmutter
- Some VASP sample jobs
- Compiling VASP for Perlmutter
- Summary

Links (also in invite e-mail):

Q&A:

<https://tinyurl.com/3byu44c8>

Survey:

<https://forms.gle/gdBKo2LM9E7hnFE98>



Getting Started with VASP at NERSC

VASP Access at NERSC

- Confirm your VASP license to access the pre-compiled VASP binaries at NERSC
 - Instructions: <https://docs.nersc.gov/applications/vasp/#access>
 - Use VASP License Confirmation Request Form at: <https://bit.ly/3Ca8kXc>
- VASP access is controlled by unix groups
 - **vasp5** for all VASP 5
 - **vasp6** for VASP 6 through 6.4.3
 - Type the **groups** command to check your access
 - **Future**: vasp 6.5+ separate from earlier VASP 6
- No need to confirm licenses if you build your own

Available VASP Modules on Perlmutter

- Type “`module avail vasp`” to see the available VASP modules

Build Type*	Pure MPI	MPI+OpenMP	MPI+OpenMP+OpenACC
Standard Distributions	vasp/5.4.4-cpu (<i>default std</i>)	vasp/6.4.3-cpu vasp/6.4.2-cpu	vasp/6.4.3-gpu vasp/6.4.2-gpu
Builds with third party codes : VTST, VASPSol	vasp-tpc/5.4.4-cpu (<i>default tpc</i>)	vasp-tpc/6.4.2-cpu	vasp-tpc/6.4.2-gpu

- All builds include precompiler-enabled plugins when possible[†]:
Wannier90, **DFTD4**, **LibXC**, **vdw-BEEF**

*) **-cpu**: build for CPUs; **-gpu**: build for GPUs; **-tpc** stands for ‘third party codes’

†) Some plugins work only in **cpu-only** builds or do not function with **certain versions** of VASP

Using VASP Modules

- Type “`module show vasp/<version-str>`” to see what a module does

```
perlmutter@login34:~> module show vasp/6.4.3-gpu
```

```
-----  
/global/common/software/nersc/pe/modulefiles_hotfixes/vasp/6.4.3-gpu.lua:  
-----
```

```
depends_on("gpu")
```

```
depends_on("nccl/2.18.3-cu12")
```

```
whatis("The Vienna Ab initio Simulation Package (VASP) is a computer program for atomic scale materials modelling.")
```

```
add_property("arch","gpu")
```

```
...
```

```
setenv("PSEUDOPOTENTIAL_DIR", "/global/common/software/nersc9/vasp/dependencies/pseudopotentials")
```

```
setenv("VDW_KERNEL_DIR", "/global/common/software/nersc9/vasp/dependencies/vdw_kernel")
```

```
setenv("MAKEFILE_INCLUDE_PATH", "/global/common/software/nersc9/vasp/vasp/6.4.3-gpu")
```

```
setenv("NO_STOP_MESSAGE", "1")
```

```
setenv("MPICH_NO_BUFFER_ALIAS_CHECK", "1")
```

```
setenv("OMP_NUM_THREADS", "1")
```

```
prepend_path("PATH", "/global/common/software/nersc9/vasp/dependencies/vtstscripts-1033")
```

```
prepend_path("PATH", "/global/common/software/nersc9/vasp/dependencies/bader")
```

```
prepend_path("PATH", "/global/common/software/nersc9/vasp/vasp/6.4.3-gpu/bin")
```

path to **makefile.include** used to build this module

path to **POTCAR** files

some useful utilities

Using VASP Modules (Cont.)

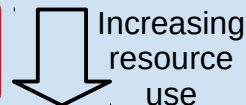
- Type “`ls -l <bin directory>`” to see available VASP binaries

```
perlmutter@login34:~> ls -l /global/common/software/nersc9/vasp/vasp/6.4.3-gpu/bin
```

```
total 152068
```

```
-rwxr-x--- 1 swowner vasp6 51307464 Oct 7 2024 vasp_gam  
-rwxr-x--- 1 swowner vasp6 52229992 Oct 7 2024 vasp_ncl  
-rwxr-x--- 1 swowner vasp6 52178704 Oct 7 2024 vasp_std
```

`vasp_gam`: the gamma point only version
`vasp_std`: the standard kpoint version
`vasp_ncl`: the non-collinear version



```
ERROR: noncollinear calculations require that VASP is compiled without the flag -DNGXhalf and -DNGZhalf
```

This error occurs when one attempts to run a noncollinear calculation using the `vasp_std` or `vasp_gam` executable. To solve, use the `vasp_ncl` executable instead.

- Type “`module load <vasp/version>`” to access VASP binaries

```
perlmutter@login34:~> module load vasp/6.4.3-gpu
```

```
perlmutter@login34:~> which vasp_std  
/global/common/software/nersc9/vasp/vasp/6.4.3-gpu/bin/vasp_std
```

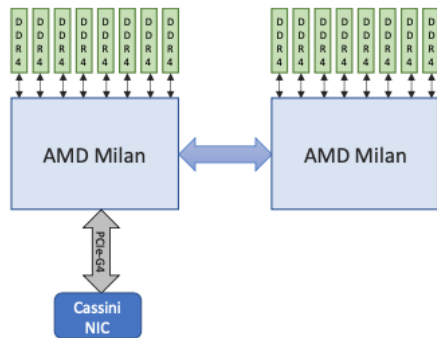


Running VASP on Perlmutter

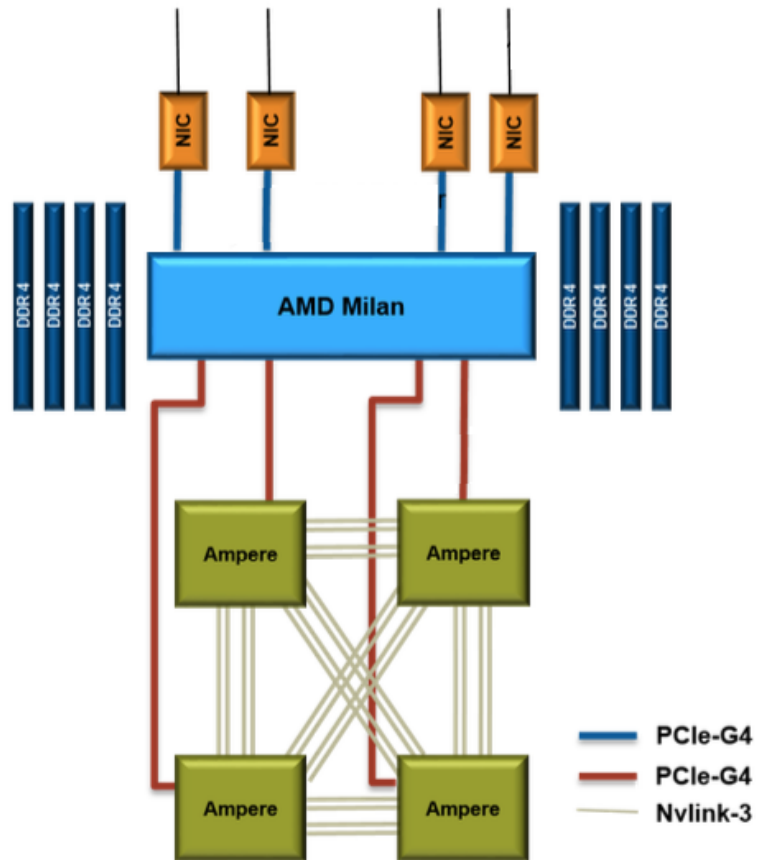
Perlmutter System Configuration

System		# of cores/node; # of gpus/node	# of CPUs per core	# of sockets per node	# of NUMA domains	Base Clock Speed (GHz)	Memory per node; per GPU (GB)
GPU accelerated nodes	GPU (A100) (1536 nodes)	64 4	2	1	4	2.45	256 40
	GPU (A100) (256 nodes)	64 4	2	1	4	2.45	256 80
CPU only nodes	CPU (Milan) (3072 nodes)	128	2	2	8	2.45	512
GPU login nodes	GPU (A100) (40 nodes)	128 1	2	2	2	2.45	512 40

Compute Nodes



AMD EPYC 7763 (Milan)



NVIDIA A100 GPU

Perlmutter Queues

- **regular** QOS – for routine use
 - Max nodes entire system, max walltime is now 48 hours
- **interactive** QOS – for real-time testing
 - Can use up to 4 nodes for 4 hours; run limit is 2
 - Starts jobs immediately or cancels in 5 minutes
- **shared** QOS – for jobs using < 1 node
- **preempt** QOS – for long, interruptible jobs
- Two job aging policy:
 - Only two jobs per QOS per user accrue priority in the queue
- Note: Perlmutter queue limits are subject to change

QOS	Max nodes	Max time (hrs)	Min time (hrs)	Submit limit	Run limit	Priority
regular	-	48	-	5000	-	medium
interactive ¹	4	4	-	2	2	high
shared_interactive	0.5	4	-	2	2	high
jupyter	4	6	-	1	1	high
debug	8	0.5	-	5	2	medium
shared ²	0.5	48	-	5000	-	medium
preempt ³	128	48 (preemptible after two hours)	2	5000	-	medium
debug_preempt	2	0.5 (preemptible after five minutes)	-	5	2	medium
premium	-	48	-	5	-	high
<u>overrun</u>	-	48 (preemptible after two hours)	-	5000	-	very low
shared_overrun	0.5	48 (preemptible after two hours)	-	5000	-	very low
<u>realtime</u>	custom	custom	custom	custom	custom	very high

Running VASP Interactively on Perlmutter Nodes

- The **interactive** QOS allows quick access to compute nodes
 - use **salloc** command to request interactive session
 - use **srun** to launch run from inside session

```
pthomas@perlmutter:login31:/global/cfs/cdirs/nstaff/pthomas/run> salloc -N 2 -q interactive -C cpu -t 30:00  
salloc: Granted job allocation 38170327  
salloc: Waiting for resource configuration  
salloc: Nodes nid[004254-004255] are ready for job
```

```
pthomas@nid004254:~> module load vasp/6.4.3-cpu  
pthomas@nid004254:~> export OMP_NUM_THREADS=8  
pthomas@nid004254:~> srun -u -n 32 -c16 --cpu-bind=cores vasp_std
```

```
-----  
  OOO PPPP EEEEE N  N M  M PPPP  
  O  O P  P E   NN N MM MM P  P  
  O  O PPPP EEEEE N N N M M M PPPP  -- VERSION  
  O  O P   E   N NN M  M P  
  OOO P    EEEEE N  N M  M P  
-----
```

```
running 32 mpi-ranks, with 8 threads/rank
```

```
...
```

Sample Job Script to Run VASP 5 on CPU Nodes

- Start by creating a small script with the job configurations, e.g. **runvasp.sh**:

```
#!/bin/bash
#SBATCH -N 1
#SBATCH -C cpu
#SBATCH -q regular
#SBATCH -t 1:00:00
```

1 node

```
module load vasp/5.4.4-cpu
```

```
srun -n 128 -c 2 --cpu-bind=cores vasp_std
```

```
#!/bin/bash
#SBATCH -N 2
#SBATCH -C cpu
#SBATCH -q regular
#SBATCH -t 1:00:00
```

2 nodes

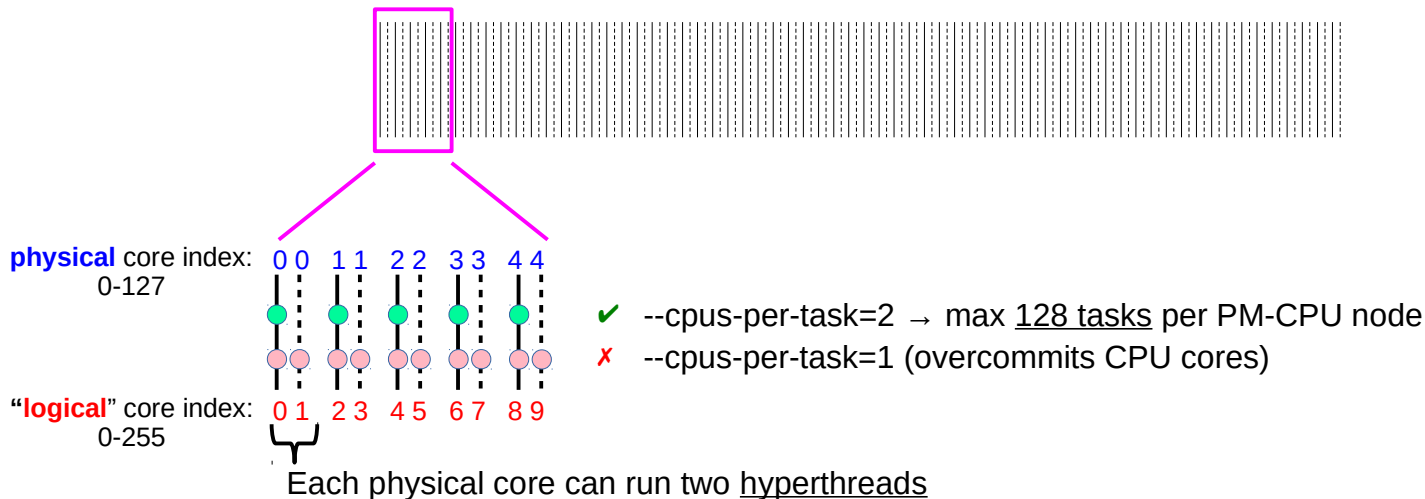
```
module load vasp/5.4.4-cpu
```

```
srun -n 256 -c 2 --cpu-bind=cores vasp_std
```

- Total MPI tasks indicated by '**-n <ntasks>**'
- The '**-c <ncpus>**' (**--cpus-per-task=<ncpus>**) parameter controls spacing of MPI tasks
- Use the **--cpu-bind=cores** option to bind tasks to CPUs
- Submit the batch script with the `sbatch` command e.g.,
sbatch runvasp.sh

Optimizing task spacing using --cpus-per-task

- The optimal MPI task spacing is 1 task per physical core



Sample Job Script to Run VASP 6 on CPU Nodes

```
#!/bin/bash
#SBATCH -N 1
#SBATCH -q regular
#SBATCH -t 1:00:00
#SBATCH -C cpu
```

1 node

```
module load vasp/6.4.3-cpu
```

```
export OMP_NUM_THREADS=4
export OMP_PLACES=threads
export OMP_PROC_BIND=spread
```

```
# launch 1 task every 16 logical cores
srun -n 32 -c 8 --cpu-bind=cores vasp_std
```

```
#!/bin/bash
#SBATCH -N 2
#SBATCH -q regular
#SBATCH -t 1:00:00
#SBATCH -C cpu
```

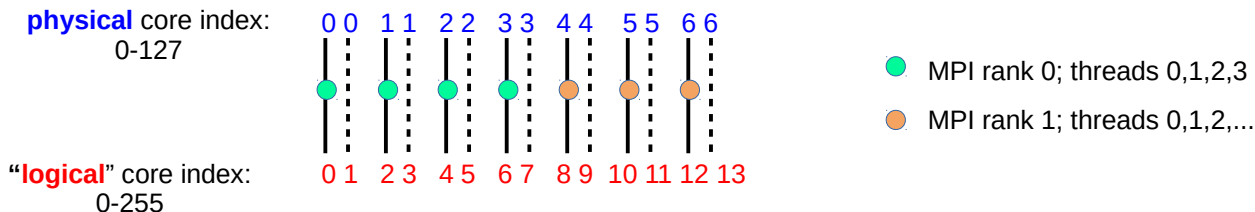
2 nodes

```
module load vasp/6.4.3-gpu
```

```
export OMP_NUM_THREADS=4
export OMP_PLACES=threads
export OMP_PROC_BIND=spread
```

```
# launch 1 task every 16 logical cores
srun -n 64 -c 8 --cpu-bind=cores vasp_std
```

- Each MPI rank spawns number of threads specified by OMP_NUM_THREADS
- Use larger **spacing** between MPI tasks to accommodate extra processes



Sample Job Script to Run VASP 6 on GPU Nodes

```
#!/bin/bash
#SBATCH -N 1
#SBATCH -q regular
#SBATCH -t 1:00:00
#SBATCH -C gpu
```

1 node

```
module load vasp/6.4.3-gpu

export OMP_NUM_THREADS=16
export OMP_PLACES=threads
export OMP_PROC_BIND=spread
```

Ensures that all GPUs are visible to all MPI ranks

```
srun -n4 -c32 -G4 --cpu-bind=cores --gpu-bind=none vasp_std
```

```
#!/bin/bash
#SBATCH -N 2
#SBATCH -q regular
#SBATCH -t 1:00:00
#SBATCH -C gpu
```

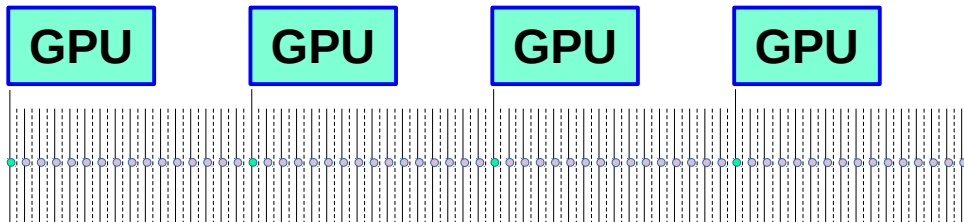
2 nodes

```
module load vasp/6.4.3-gpu

export OMP_NUM_THREADS=16
export OMP_PLACES=threads
export OMP_PROC_BIND=spread
```

```
srun -n8 -c32 -G8 --cpu-bind=cores -gpu-bind=none vasp_std
```

- **Always** use 4 ranks per node with '**-c 32**' to select 1:1 MPI-rank-to-GPU mapping
- To use additional CPU cores, use up to **16** OpenMP threads per MPI rank



NERSC Job Script Generator

https://my.nersc.gov/script_generator.php

Jobscript Generator

CPU

Job Information

This tool generates a batch script template which also realizes specific process and thread binding configurations.

Machine

Select the machine on which you want to submit your job.

Perlmutter - CPU

Application Name

Specify your application including the full path.

vasp_std

Job Name

Specify a name for your job.

test_vasp

Email Address

```
#!/bin/bash
#SBATCH -N 4
#SBATCH -C cpu
#SBATCH -q regular
#SBATCH -J test_vasp
#SBATCH --mail-user=pthomas@lbl.gov
#SBATCH --mail-type=ALL
#SBATCH -t 01:00:00
```

```
#OpenMP settings:
export OMP_NUM_THREADS=8
export OMP_PLACES=threads
export OMP_PROC_BIND=spread
```

```
#run the application:
srun -n 64 -c 16 --cpu_bind=cores vasp_std
```

Jobscript Generator

GPU

Job Information

This tool generates a batch script template which also realizes specific process and thread binding configurations.

Machine

Select the machine on which you want to submit your job.

Perlmutter - GPU

Application Name

Specify your application including the full path.

vasp_std

Job Name

Specify a name for your job.

test_vasp

Email Address

Specify your email address to get notified when the job enters a certain state.

pthomas@lbl.gov

```
#!/bin/bash
#SBATCH -N 4
#SBATCH -C gpu
#SBATCH -G 16
#SBATCH -q regular
#SBATCH -J test_vasp
#SBATCH --mail-user=pthomas@lbl.gov
#SBATCH --mail-type=ALL
#SBATCH -t 01:00:00
#SBATCH -A m1234
```

```
#OpenMP settings:
export OMP_NUM_THREADS=1
export OMP_PLACES=threads
export OMP_PROC_BIND=spread
```

```
#run the application:
#applications may perform better with --gpu-
bind=none instead of --gpu-bind=single:1
srun -n 16 -c 32 --cpu_bind=cores -G 16 --
gpu-bind=single:1 vasp_std
```

VASP-specific samples at:

<https://docs.nersc.gov/applications/vasp/#sample-job-scripts>

N.B. use `--gpu-bind=none`

Bundle Jobs with Multiple Sruns

- Sample job script bundling multiple VASP jobs

```
#!/bin/bash
#SBATCH --qos=debug
#SBATCH --nodes=10
#SBATCH --time=30:00
#SBATCH --constraint=cpu

module load vasp/5.4.4-cpu

#Assume 5 VASP jobs in run1, run2, ..., run5 directories
for j in {1..10}; do
  cd run$j
  srun -N 1 -n 128 -c 2 --cpu_bind=cores vasp_std &
  cd ..
done

wait
```

- The number of sruns in the job script should be small
 - Slurm is **not** good at handling multiple sruns in a single job script
 - Nodes with earlier completed jobs will be idle
- For many jobs, consider using a workflow tool such as Fireworks

Bundle Jobs with Multiple Sruns + Job Resizing

Sample job script to bundle VASP jobs with job resizing

```
#!/bin/bash
#SBATCH -J job_resizing
#SBATCH -q debug
#SBATCH -N 10
#SBATCH -t 30:00
#SBATCH -C cpu

module load vasp/5.4.4-cpu
nodelist="`scontrol show hostname`"

#cd to each run directory, and launch srun on each node
for node in $nodelist ;do
    cd run$j
    ../run_jobstep.sh $node &
    cd ..
done

wait
```

```
cat run_jobstep.sh
#!/bin/bash
mynode=$1
srun -N1 -w $mynode -n128 -c2 --cpu-bind=cores -o %x-%j-${mynode}.out vasp_std

#release $mynode from nodelist if $mynode is not the batch host (head node) or it is not the last remaining node
batchHost=`scontrol show hostname |head -1`
if [[ $mynode != $batchHost ]]; then
    nodelist=`squeue -h -j $SLURM_JOB_ID -o %N`
    remaining_nodelist=`scontrol show hostname $nodelist |grep -v $mynode`
    if [[ -n $remaining_nodelist ]]; then
        scontrol update job $SLURM_JOB_ID Nodelist="`echo $remaining_nodelist`"
    fi
fi
```

- **Release the nodes** as soon as the job running on them completes while the rest of the jobs are still running
- You should not release the batch host (head node)
- Make the longest job run on the head node (the first job)
- You can modify this script for your needs, e.g., running each job instance on multiple nodes

Running Long VASP Jobs

- Some VASP runs can be restarted from CONTCAR + {CHGCAR, WAVECAR}
- Use the **preempt** QOS to run at a discount after first 2 h!

```
#!/bin/bash
#SBATCH -N 1
#SBATCH -C cpu
#SBATCH -J vasp_job
#SBATCH -o %x-%j.out
#SBATCH -e %x-%j.err
#SBATCH --qos=preempt
#SBATCH --comment=96:00:00
#SBATCH --time=2:00:00
#SBATCH --time-min=0:30:00
#SBATCH --signal=B:USR1@60
#SBATCH --requeue
#SBATCH --open-mode=append
```

```
if [ -f "STOPCAR" ]; then
rm STOPCAR
fi
```

```
module load vasp/5.4.4-cpu
```

```
srun -n 128 -c 2 --cpu_bind=cores vasp_std &
```

checkpoint command

total job length

job segment length

time needed to checkpoint

```
ckpt_vasp() {
set -x
restarts=`squeue -h -O restartcnt -j $SLURM_JOB_ID`
echo checkpointing the ${restarts}-th job
```

```
restarts_num=$(echo $restarts | sed -e 's/^[ \t]*//')
echo "Restart number: ==${restarts_num}=="
```

```
echo LABORT = .TRUE. > STOPCAR
srun_pid=$(ps -file | grep srun | head -1 | awk '{print $4}')
echo srun pid is $srun_pid
wait $srun_pid
```

```
folder="ckpt-$SLURM_JOB_ID-${restarts_num}"
mkdir $folder
echo "In directory $folder"
```

```
cp -p CONTCAR POSCAR
cp -p CONTCAR "$folder/POSCAR"
cp -p OUTCAR "$folder/OUTCAR-${restarts_num}"
cp -p OSZICAR "$folder/OSZICAR-${restarts_num}"
cp -p vasprun.xml "vasprun-${restarts_num}.xml"
```

```
set +x
}
```

```
ckpt_command=ckpt_vasp
max_timelimit=7200
ckpt_overhead=60
. /global/common/sw/cray/cnl7/haswell/nersc_cr/19.10/etc/env_setup.sh
requeue_job func_trap USR1
```

```
wait
```

A Few Useful Commands

- Commonly used Slurm commands:
 - `sbatch, salloc, scancel, srun, squeue, sinfo, sqs, scontrol, sacct` (check man page)
 - `sinfo --format='%F %b'` for available features of nodes, or `sinfo --format='%C %b'`
 - `scontrol show node <nid>` for node info



Troubleshooting VASP

Common Runtime Issues

- Most common issue: hanging due to bad CPU or GPU task binding
- Executable not found (for new users with unconfirmed licenses)
- Out of Memory
- Catastrophic failure (“segmentation fault”, “double linked list corrupted”, “PMPI_Allreduce: Invalid communicator”) due to solver crash
- VASP fails to read input files created with MS-DOS endline characters
- Intermittent issues with bad nodes or file system sometimes occur

Troubleshooting Tips and Tricks

- Check job script, INCAR for mistakes
- Check OUTCAR, **standard out, err**
- **stdout output=L** releases message in output buffer
- Increase **resources**
- Run **different version** or **{cpu, gpu} build**
- **ssh nid<xxxx>** to log into node
 - **top** to check CPU activity
 - **nvidia-smi** to check GPU
- Repeat run on different node(s)

```
#!/bin/bash
#SBATCH -N 1
## SBATCH -N 2
#SBATCH -C cpu
#SBATCH -j vasp_job
#SBATCH -o %x-%j.out
#SBATCH -e %x-%j.err

module load vasp/5.4.4-cpu
## module load vasp/6.4.3-cpu
## module load vasp/6.4.3-gpu

srun -n 128 -c 2 -cpu_bind=cores stdout --output=L vasp_std
## srun -n 256 -c 2 -cpu_bind=cores vasp_std
```

Check NERSC MOTD for possible outage:
<https://www.nersc.gov/live-status/motd/>

Troubleshooting Resources

- Start with NERSC Troubleshooting Guide:
<https://docs.nersc.gov/applications/vasp/#troubleshooting-vasp>
- VASP user forums:
<https://www.vasp.at/forum/>
- File a support ticket:
<https://help.nersc.gov/>
 - Describe issue in detail, including error message, which module, steps attempted to fix
 - Provide path-to-job, not attachment

Troubleshooting VASP 📄

Solutions to frequently encountered issues

📄 General Tips >

📄 My VASP job just sits in the queue and does not start. >

📄 My VASP job immediately crashes with the message: `<vasp_executable>: No such file or directory:` >

📄 My VASP job crashes with an error when trying to read { POSCAR | INCAR | KPOINTS }, but the file appears to be correct. >

📄 My VASP job immediately crashes with an error about not being compiled for noncollinear calculations >

📄 My VASP job runs but produces no output or incomplete output >

📄 My VASP job hangs or stops making progress >

📄 My VASP job runs out of memory >

📄 My VASP job crashes with a 'BUG!' message in my OUTCAR file >

📄 My VASP jobs are running but there are problems with convergence in the SCF or relaxation >



VASP Performance on Perlmutter

VASP - #1 Production Code at NERSC

- VASP uses more than 15% of NERSC computing cycles each year
- 849 active VASP users on Perlmutter as of April, 15, 2025
- VASP was a Tier 2 NESAP application, and NVIDIA has been heavily involved with its OpenACC porting
- All major code paths in VASP have been ported to GPUs except the portion using Random Phase Approximation (GW and Adiabatic Connection Fluctuation Dissipation Theorem)
- N.B.: Perlmutter is undergoing continual changes and improvements, thus performance is subject to change

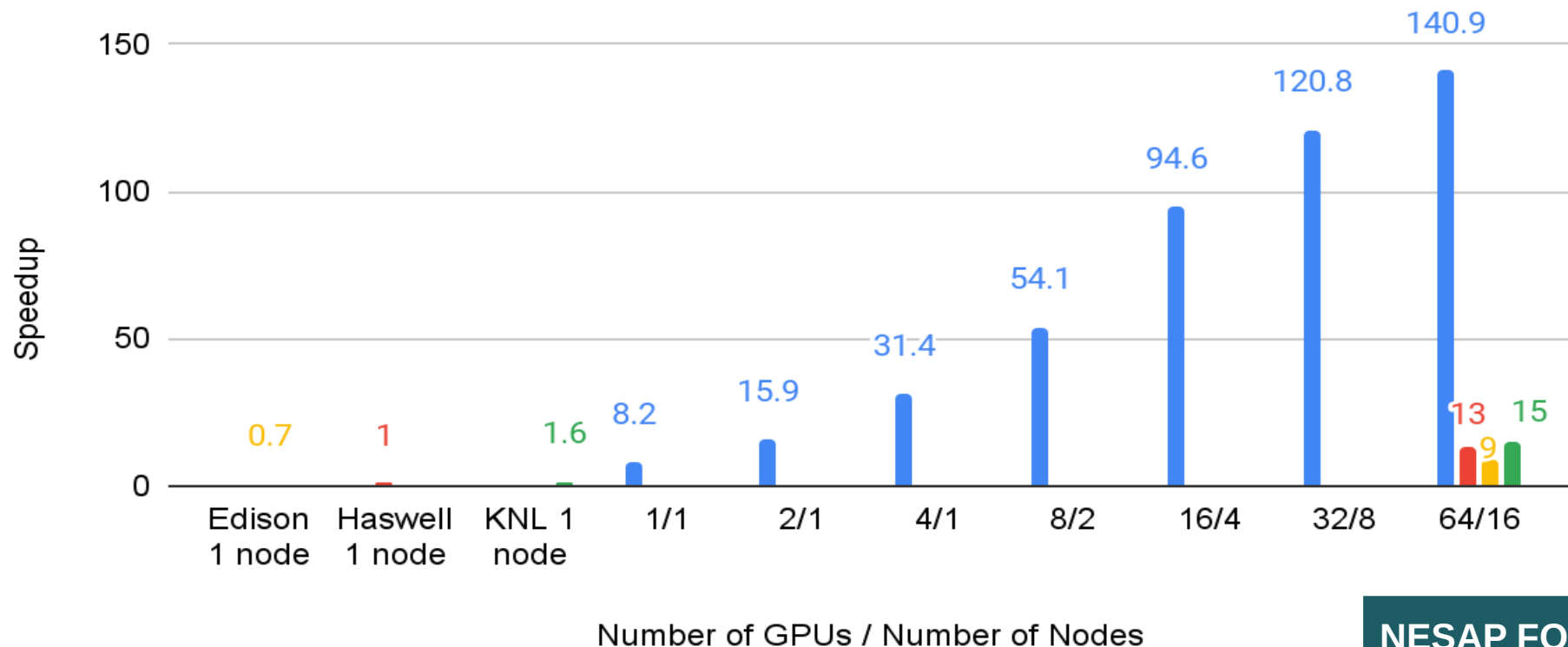
Benchmarks Were Chosen to Cover the Representative VASP Workloads and to Exercise Different Code Paths

	PdO4	GaAsBi -64	CuC_vdw	Si256_hse	B.hR105_hse	PdO2
Electrons (Ions)	3288 (348)	266 (64)	1064 (98)	1020 (255)	315 (105)	1644(174)
Functional	DFT	DFT	VDW	HSE	HSE	DFT
Algo	RMM (VeryFast)	BD+RMM (Fast)	RMM (VeryFast)	CG (Damped)	CG (Damped)	RMM (VeryFast)
NEML(NELMDL)	5 (3)	8 (0)	10 (5)	3(0)	10 (5)	10 (4)
NBANDS	2048	192	640	640	256	1024
FFT grids	80x120x54 160x240x108	70x70x70 140x140x140	70x70x210 120x120x350	80x80x80 160x160x160	48x48x48 96x96x96	80x60x54 160x120x108
NPLWV	518400	343000	1029000	512000	110592	259200
IRMAX	1445	4177	3797	1579	1847	1445
IRDMAX	3515	17249	50841	4998	2358	3515
LMDIM	18	18	18	18	8	18
KPOINTS	1 1 1	4 4 4	3 3 1	1 1 1	1 1 1	1 1 1

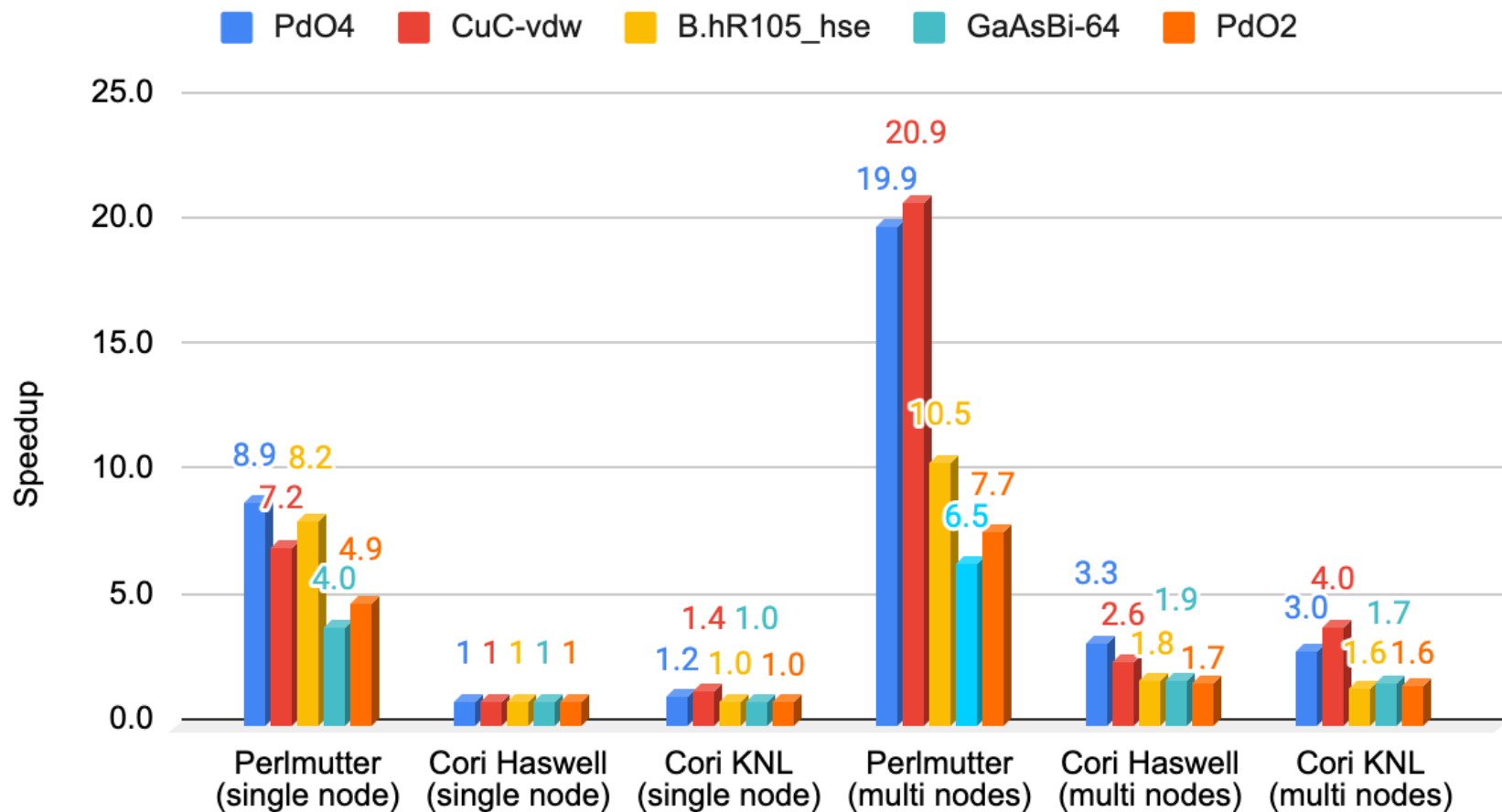
VASP Performance on Perlmutter

(Si256_hse)

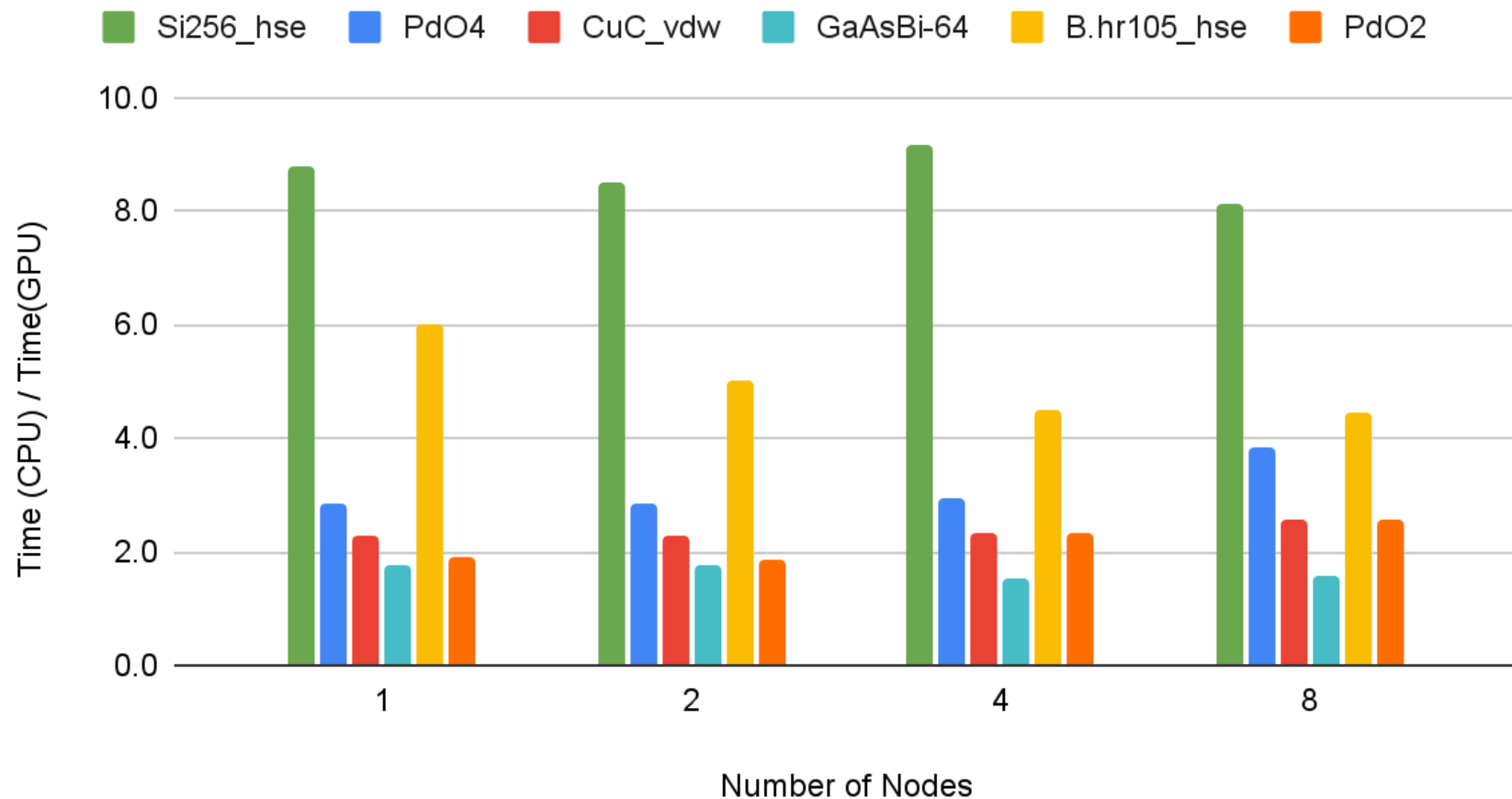
Perlmutter Cori Haswell Edison Cori KNL



VASP Performance on Perlmutter GPUs



GPU Speedup vs Milan



Tips for Maximizing Performance

- VASP performance can be affected by many parameters, including system dimensions/symmetries and linked software libraries.

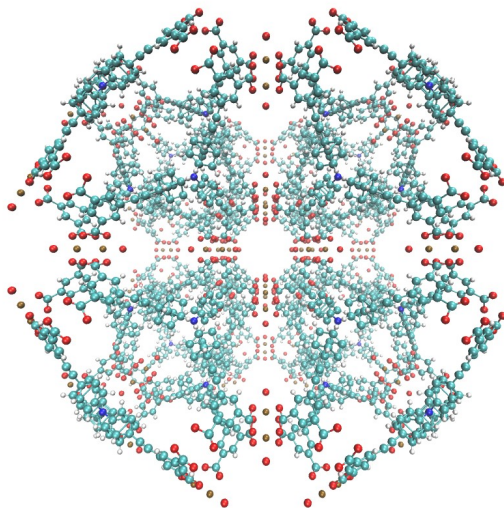
“Guess nothing, measure everything” –Anon

- Use a **GPU** build if your application allows
- Use the newest version of VASP allowed by your license
- Do not run on hyperthreads
- (CPU only): measure NCORE=4, 8, 16, ... vs. MPI-rank/OpenMP-thread ratios to find optimal values
- Set KPAR to parallelize over irreducible KPOINTS



Some VASP Sample Jobs

The Materials Genome (K. Peterson, LBNL)



MOF structure (**3584 atoms**)

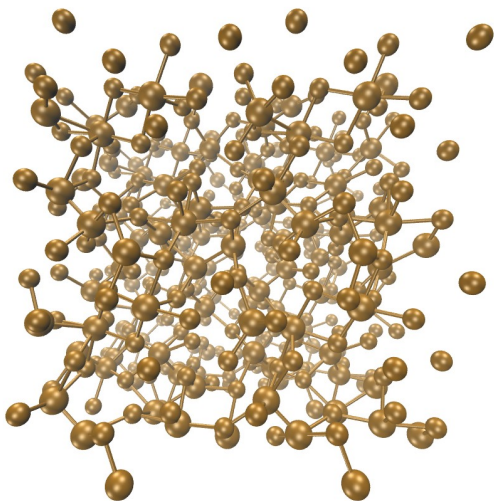
VASP Input files provided by Shyam
Dwaraknath at LBNL

- Job description (K. Peterson):

“We have a large MOF structure that was sent to the Materials Project by a user, and it was too large for us to handle through our normal workflows”
- Computation details:

Electrons(Ions): 13408 (3584)
Functional: DFT
Algo: CG (BD+RMM)
NBANDS= 9600
FFT grids: 378x378x378; 756x756x756
NPLWV: 54,010,152
KPOINTS: 1 1 1
- Completed 99 ionic steps in 32.8 hours using 32 nodes (128 GPUs) on Perlmutter

Discovery of Materials for Energy Applications (C. Wolverton, Northwestern Univ.)

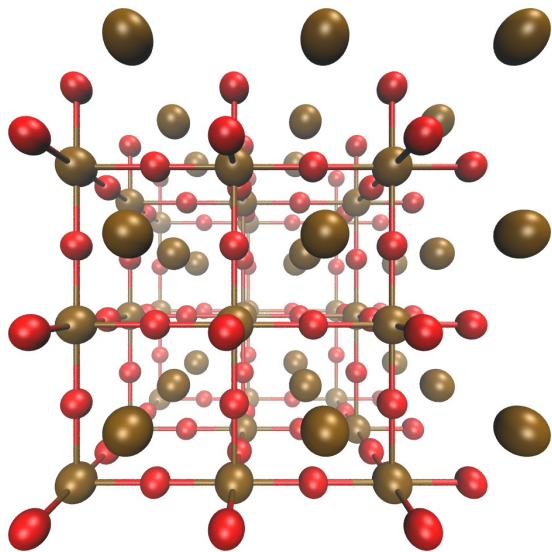


Amorphous VF₃ (416 atoms)

VASP Input files provided by
Shiqiang Hao Northwestern Univ.

- Job description (S. Hao):
“This job calculates Li diffusion in an amorphous phase of metal fluorides VF₃ using ab initio molecular dynamics methods to investigate if the amorphous VF₃ is a good Li conductor.”
- Computation details:
Electrons (Ions): 2592 (416)
Functional: DFT
Algo: CG (BD+RMM)
NBANDS: 1920
NPLWV: 1905120
FFT grids: 126x126x120; 192x180x180
KPOINTS: 7 7 7 (172 irreducible kpoints)
- Completed 66 ionic steps in 17.2 hours using 160 nodes (640 GPUs) on Perlmutter

Simulating Core-level Spectroscopy of Energy-relevant Materials (D. Prendergast, LBNL)

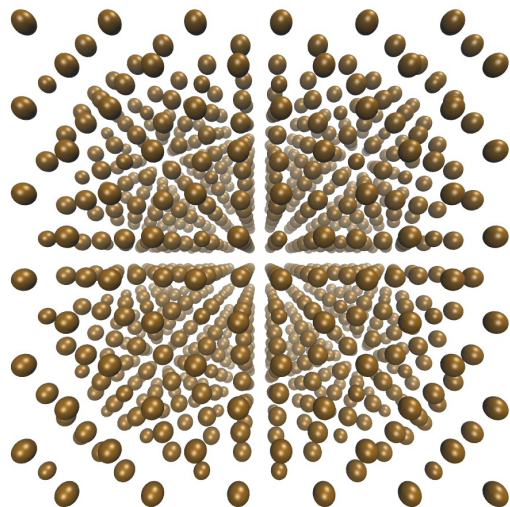


SrHfO₃ (135 atoms)

VASP Input files provided by Ella
Banyas at UCB

- Job description:
A large supercell (135 atoms) **phonon calculations with DFPT**
- Computation details:
Electrons (Ions): 1026 (135)
Functional: DFT
Algo: BD, RMM
IBRION: 7; LEPSILON = .TRUE.
NBANDS= 616
NPLWV: 1259712
FFT grids: 108x108x109; 216x216x216
Support grids: 432x432x432
KPOINTS: 3 3 3 (27 irreducible kpoints)
- Completed in 37 hours using 27 nodes (108 GPUs) on Perlmutter

Material Simulations in Joint Center for Artificial Photosynthesis (L. Wang, LBNL)



NiMoNi (864 atom) alloy

VASP Input files provided by Lin-Wang Wang LBNL

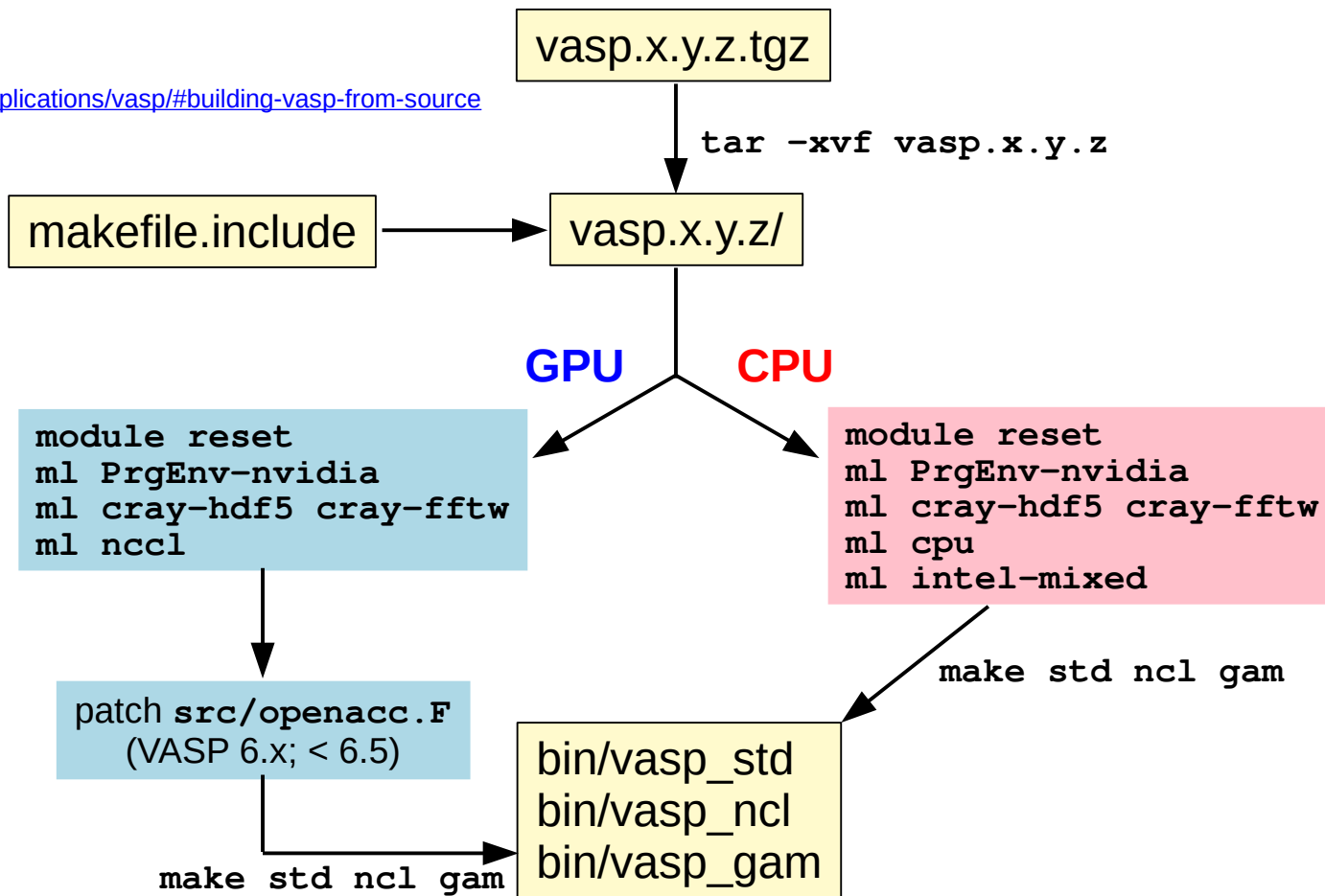
- Job description (L. Wang):
“This is an MD simulation for a BiMoNi alloy system which is used for catalysis to reduce Methane to carbon.”
- Computation details
Electrons (Ions): 4984 (864)
Functional: DFT
Algo: RMM
MDALGO = 3
NBANDS= 3008
NPLWV: 941192
FFT grids: 65x65x65; 196x196x196
KPOINTS: 1 1 1
- Completed **3000 MD steps** in 43.3 hours using 16 nodes (64 GPUs) on Perlmutter
 - 52 seconds per MD step



Compiling VASP for Perlmutter

VASP Installation Workflow on Perlmutter

<https://docs.nersc.gov/applications/vasp/#building-vasp-from-source>



Configuring VASP with **makefile.include**

- Sample 'makefile.include' for existing NERSC modules are at **`$MAKEFILE_INCLUDE_PATH/makefile.include`**
- Each programming environment requires its own 'makefile.include'! NERSC-provided ones are specific to 'PrgEnv-nvidia'.
- Samples for GNU, Intel can be found at <https://www.vasp.at/wiki/index.php/Makefile.include> (some modifications required)
- Can construct for other compilers but may require effort!

Anatomy of a sample `makefile.include`

Precompiler options

```
CPP_OPTIONS= -DHOST=\"LinuxNV_CrayMPICH\" \  
-DMPI -DMPI_BLOCK=8000 -Duse_collective \  
-DscalAPACK \  
-DCACHE_SIZE=4000 \  
-Davoidalloc \  
-Dvasp6 \  
-Duse_bse_te \  
-Dtbdyn \  
-Dqd_emulate \  
-Dflock_dblbuf \  
-D_OPENMP \  
-D_OPENACC \  
-DUSENCCL \  
-DVASP_HDF5 \  
-DVASP2WANNIER90 \  
-DDFTD4 \  
-DPROFILING
```

Disabled for GPU build:

```
# -Dsysv \  
# -DUSELIBXC \  
# -Dlibbeef \
```

```
CPP    = nvfortran -Mpreprocess -Mfree -Mextend -E $(CPP_OPTIONS) $$$(FUUFFIX) >  
FC      = ftn -mp -acc=gpu -gpu=cc80  
FCL     = ftn -v -mp -acc=gpu -gpu=cc80 -c++libs  
FREE    = -Mfree
```

- **-D_OPENMP** to add OpenMP parallelism
- **-D_OPENACC** to build for GPU; use **-DUSENCCL** to include NCCL for MPI+GPU performance
- **-DVASP2WANNIER90**, **-DDFTD4**, **-DUSELIBXC**, **-DLIBBEEF** to include optional plugins
- **-DPROFILING** to build version which prints timings

The compiler wrapper, **ftn**, is used for automatic inclusion of process specific optimizations and Cray libraries

Anatomy of a sample `makefile.include` (cont'd)

```
# use cray-fftw module for FFTs
# optional packages:
```

```
# NCCL (GPU builds only)
```

```
LLIBS    += -L$(NCCL_DIR)/lib -Wl,-rpath=$(NCCL_DIR)/lib
INCS     += -I$(NCCL_DIR)/include
```

Load NCCL module and include this section to use in GPU builds

```
# HDF5 (vasp >6.2.0 only)
```

```
LLIBS    += -L$(HDF5_ROOT)/lib -lhdf5_fortran
INCS     += -I$(HDF5_ROOT)/include
```

Uncomment Intel MKL and FFTlib for use in CPU builds for performance

```
# Use Intel MKL libraries (CPU builds only)
```

```
#MKLROOT  ?= /global/common/software/nersc/pm-stable/sw/intel/oneapi/mkl/latest
```

```
#LLIBS    += $(EXTLIBDIR)/mkl-interfaces/2022.1.0/milan/nvidia-22.5/libfftw3xf_nvidia.a
```

```
#LLIBS    += -L${MKLROOT}/lib/intel64 -lmkl_scalapack_lp64 -lmkl_intel_lp64 -lmkl_pgi_thread -lmkl_core -lmkl_blacs_intelmpi_lp64 -mp -lpthread
```

```
#LLIBS    += -Wl,-rpath=$(MKLROOT)/lib/intel64
```

```
#INCS     += -I$(MKLROOT)/include/fftw
```

```
# fftlib (with OpenMP only; useful for CPU builds)
```

```
#FCL      += fftlib.o
```

```
#CXX_FFTLIB = nvc++ -mp --no_warnings -std=c++11 -DFFTLIB_THREADSAFE
```

```
#INCS_FFTLIB = -I./include -I$(FFTW_ROOT)/include
```

```
#LIBS     += fftlib
```

```
#LLIBS     += -ldl
```

This directory contains other optional third party codes builds (must be specified to use)

```
# PATH FOR PLUGIN BUILDS
```

```
EXTLIBDIR = $(CFS)/omp/local
```

```
# libbeef (CPU builds only)
```

```
LIBBEEF_ROOT = $(EXTLIBDIR)/libbeef/2020-11-01/nvidia-22.5/milan/lib
```

```
LLIBS     += -L$(LIBBEEF_ROOT) -lbeef
```

```
...
```



Summary

Summary and Looking Ahead

- While speed-ups depend on job sizes and types, VASP achieves **4x-30x** faster node-to-node performance on Perlmutter-GPU vs Cori Haswell with representative VASP workloads
- VASP achieves **1.5x-9x** faster node-to-node performance on Perlmutter GPUs vs Perlmutter CPUs for selected VASP workloads
- VASP 6.5+ builds are planned! If you have access to source then NERSC staff can assist you with building.
- VASP 5.4.4 support to continue through Perlmutter's life cycle, with expected deprecation in N10
- Containerized VASP planned for Perlmutter, likely will replace module environment in N10

Thank You!



Questions? We are here to help!

Q&A link:

<https://tinyurl.com/3byu44c8>

Survey link:

<https://forms.gle/gdBKo2LM9E7hnFE98>