

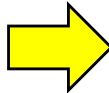
On Assignment 5

- Q. Part 1 (programming):** Since we are starting with `pmd_irecv.c` in assignment 4, do we only need to mark the OpenMP changes or both MPI & OpenMP changes?
- A.** Please only mark the OpenMP changes.
- Q. Part 3 (strong scaling):** Should we plot & submit the runtime, speedup & parallel efficiency as a function of the number of threads, as in slide 20 of “Hybrid MPI+OpenMP MD” lecture, <https://aiichironakano.github.io/cs596/05HMD.pdf>?
- A.** No, please submit only the efficiency plot.

On Assignment 5, Part 3 (Scaling)

- **Goal:** Measure **multithread parallel efficiency** on multiple cores within a single computing node
- **CPU** in standard output is total runtime (in seconds) including computing & communication; use it as $T(N, P)$ in the speedup formula, where N is the fixed problem size (proportional to the total # of atoms, $nglob = 55296$, but doesn't enter in efficiency calculation) & P is the # of cores (or threads, remember one thread per core seen using 'top' command)

```
[[liuhaora@discovery2 hw5]$ cat hmd-scale.out
8 threads
al = 4.103942e+01 4.103942e+01 4.103942e+01
lc = 16 16 16
rc = 2.564964e+00 2.564964e+00 2.564964e+00
nglob = 55296
CPU & COMT = 8.350226e+00 2.620383e-02
4 threads
al = 4.103942e+01 4.103942e+01 4.103942e+01
lc = 16 16 16
rc = 2.564964e+00 2.564964e+00 2.564964e+00
nglob = 55296
CPU & COMT = 3.940460e+00 2.325014e-02
2 threads
al = 4.103942e+01 4.103942e+01 4.103942e+01
lc = 16 16 16
rc = 2.564964e+00 2.564964e+00 2.564964e+00
nglob = 55296
CPU & COMT = 9.614403e+00 2.323241e-02
1 thread
al = 4.103942e+01 4.103942e+01 4.103942e+01
lc = 16 16 16
rc = 2.564964e+00 2.564964e+00 2.564964e+00
nglob = 55296
CPU & COMT = 1.492553e+01 2.285458e-02
```



| | |
|---------|----------------------|
| $P = 1$ | $T(N, 1) = 14.92553$ |
| $P = 2$ | $T(N, 2) = 9.614403$ |
| $P = 4$ | $T(N, 4) = 3.940460$ |
| $P = 8$ | $T(N, 8) = 8.350226$ |

$$S_2 = \frac{T(N, 1)}{T(N, 2)}$$

Strong-scaling (fixed problem size):

Speedup: $S_P = \frac{T(N, 1)}{T(N, P)}$

Efficiency: $E_P = \frac{S_P}{P}$; **only plot this!**

Why Dip in Runtime for $P = 4$?

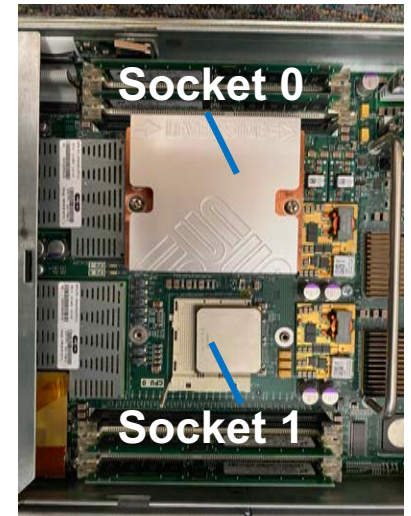
- Each of the two processors (or sockets) with multiple cores has fast local memory called cache (to be discussed in performance optimization lecture)

In prior architectures (such as the Intel® Xeon® E5 v4 Processor family):

- The mid-level cache (MLC or also known as L2) was 256 KB per core.
- The last level cache (also known as L3) was a shared inclusive cache with 2.5 MB per core.

In the architecture of the Intel® Xeon® Scalable Processor family, the cache hierarchy has changed to provide a larger MLC of 1 MB per core and a smaller shared non-inclusive 1.375 MB LLC per core. A larger MLC increases the hit rate into the MLC resulting in lower effective memory latency and also lowers demand on the mesh interconnect and LLC. The shift to a non-inclusive cache for the LLC allows for more effective utilization of the overall cache on the chip versus an inclusive cache.

<https://www.intel.com>

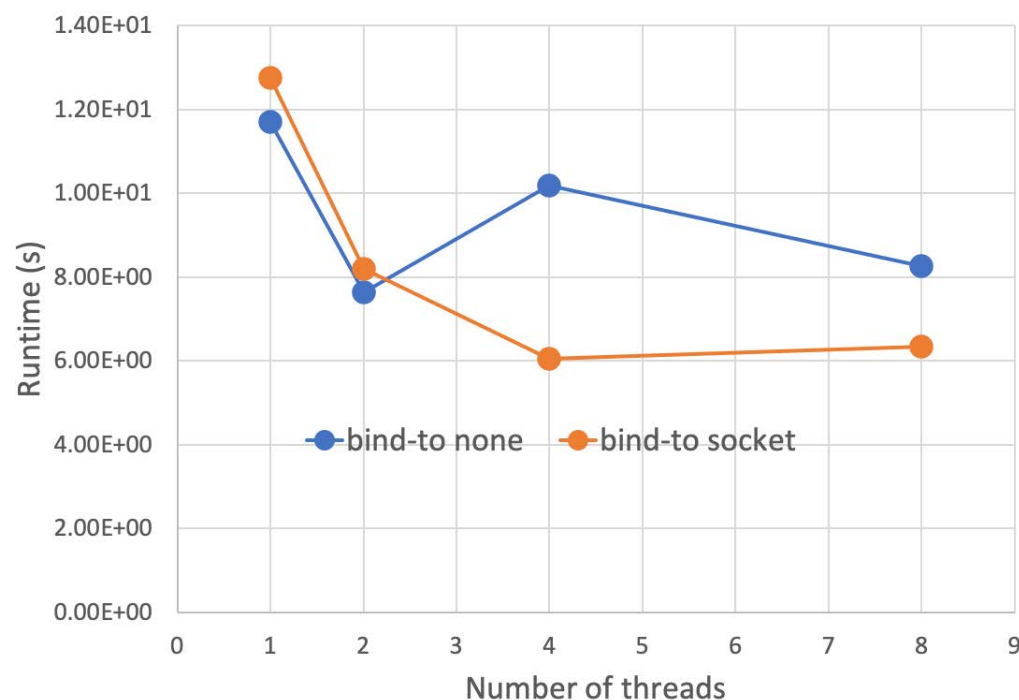


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- In addition to more arithmetic-logic operations, multiple cores provide larger caches to improve memory-access speed
- If threads are placed on different sockets, however, memory performance degrades
- Non-uniform memory access (NUMA):** Memory design, where memory access time depends on memory location relative to the processor
- Again, there also is interference with other users in the same computing node

Affinity

- **Processor (task) affinity:** Controls binding (*i.e.*, pinning) of a process to a core or socket (`mpirun -bind-to none` unbinds a rank from single core or socket, while `mpirun -bind-to socket` pins all threads within one socket)
https://en.wikipedia.org/wiki/Processor_affinity
- Binding can improve cache performance but degrade load balancing



- “There still is not an easy way for pinning MPI processes & OpenMP threads to CPU sockets & cores.” *How to gain hybrid MPI-OpenMP code performance without changing a line of code a.k.a. dealing with task affinity:*

<https://aciref.org/how-to-gain-hybrid-mpi-openmp-code-performance-without-changing-a-line-of-code-a-k-a-dealing-with-task-affinity/>

- Don't worry about nonmonotonic behavior & submit what your got (again not runtime but efficiency)
- Will revisit false sharing & affinity in performance-optimization lecture

About srun

- While **mpirun** is a command in MPI system to execute a parallel program, some versions of Slurm job-scheduling system & OpenMPI implementation of MPI support **srun** to execute an MPI program.

2. Does Open MPI support "srun -n X my_mpi_application"?

Yes, if you have configured OMPI `--with-pmi=foo`, where `foo` is the path to the directory where `pmi.h/pmi2.h` is located. Slurm (> 2.6, > 14.03) installs PMI-2 support by default.

<https://www.open-mpi.org/faq/?category=slurm#slurm-direct-srun-mpi-apps>

- Consult a system administrator about specific settings after each upgrade (following descriptions on discovery are somewhat outdated)

`srun` Launch parallel tasks (i.e., job steps) (typically for MPI jobs)

<https://carc.usc.edu/user-information/user-guides/hpc-basics/running-jobs>

```
module purge
module load gcc/8.3.0
module load openmpi/4.0.2
module load pmix/3.1.3
```

```
ulimit -s unlimited
```

```
export OMP_NUM_THREADS=$SLURM_CPUS_PER_TASK
```

Automatically bind tasks to
NUMA locality domains

```
srun --mpi=pmix_v2 --cpu-bind=ldoms -n $SLURM_NTASKS ./mpi_plus_openmp_program.x
```

<https://carc.usc.edu/user-information/user-guides/hpc-basics/slurm-templates>

Recap: Slide 20 in Hybrid MPI+MD Lecture

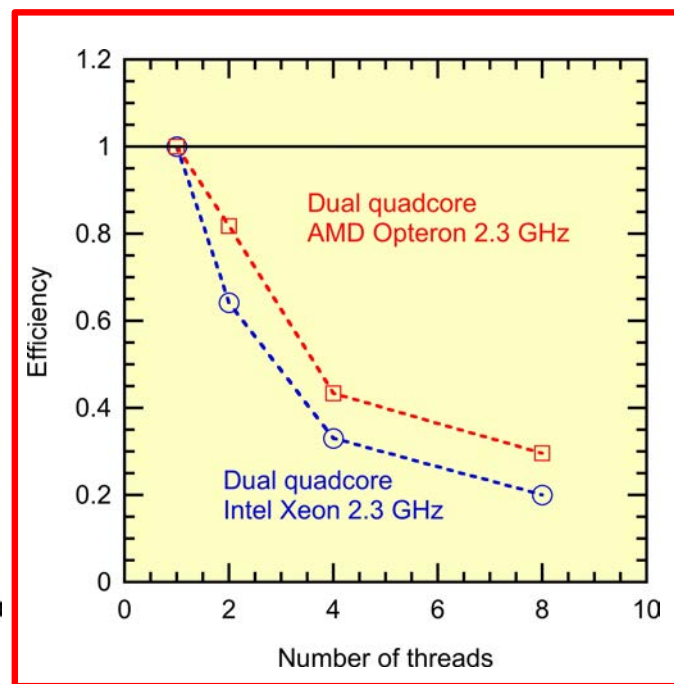
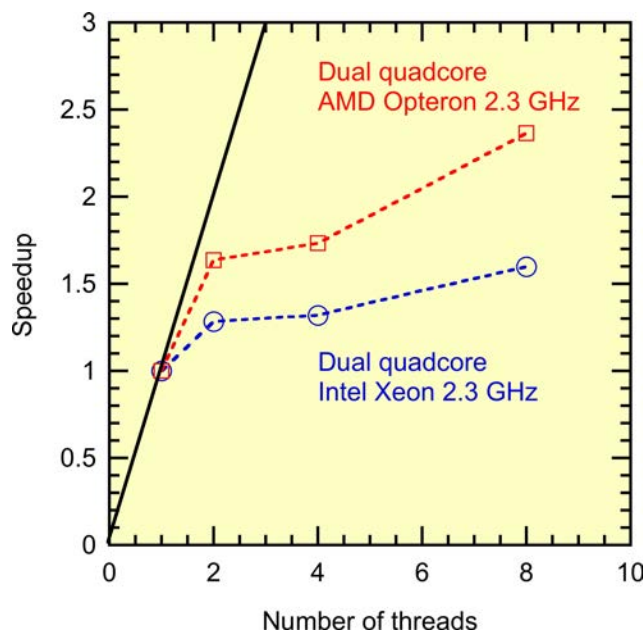
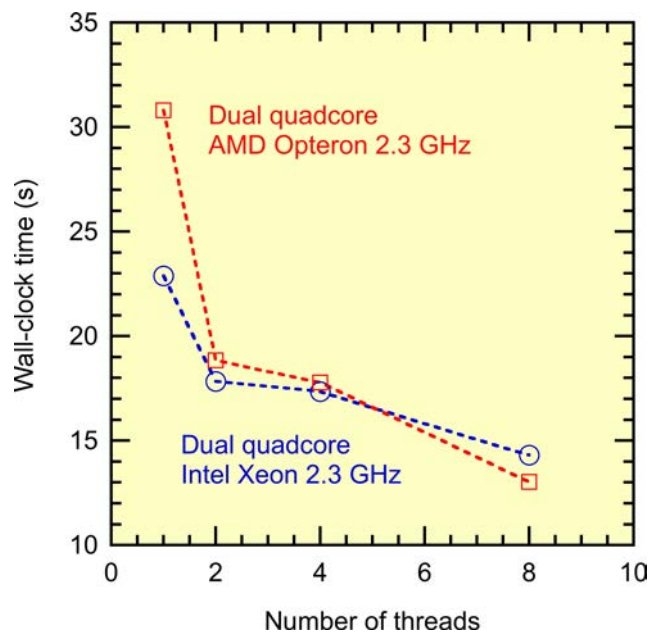
1 MPI process; 1-8 threads

In `hmd.h`:

```
vproc = {1,1,1}, nproc = 1;
vthrd = {1,1,1}, nthrd = 1;
        2 1 1          2
        2 2 1          4
        2 2 2          8
```

`pmd.in`

```
24 24 24 InitUcell[3]
0.8      Density
1.0      InitTemp
0.005    DeltaT
100      StepLimit
101      StepAvg
```



`InitUcell[] = {24,24,24}`

$N = 4 \times 24^3$
= 55296 atoms

$$S_P = \frac{T(N,1)}{T(N,P)}$$

P : Number of cores

$$E_P = \frac{S_P}{P}$$

Just one curve (no need to compare different nodes)