THE UNIVERSITY OF ARIZONA Intro to Parallel Computing on HPC

Summer 2024

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access these slides: https://bit.ly/4eZEQwD

Outline

Section 1: Background and Theory

- a. What is parallel computing? Why should we use it?
- b. Terminology and Theory

Section 2: Practical Parallel Computing on UA HPC

- a. Use cases, and user archetypes
- b. Guidelines for parallel computing

Section 3: Examples

- a. Array Jobs
- b. GNU Parallel
- c. Python multiprocessing and mpi4py
- d. Resources for R

Final Slide: References and Recommended Reading

Why Do We Care?

MacBook Pro 2021

Apple M1 Pro Chip 8-core CPU 16GB unified memory 512GB SSD \$1999



Penguin Altus XE2242

4in1 chassis. Each compute node has:

96 cores dual socket AMD EPYC 7642 512GB DDR4 3200MHz ECC memory 2TB SSD NVMe \$8000



Why Do We Care?

MacBook Pro 2021 Apple M1 Pro Chip 8-core CPU 16GB unified memory 512GB SSD \$1999



Puma Cluster 269 compute nodes 25,824 cores dual socket AMD EPYC 7642 137,728GB DDR4 3200MHz ECC memory 2PB SSD NVMe shared storage 538TB local storage \$2.7M



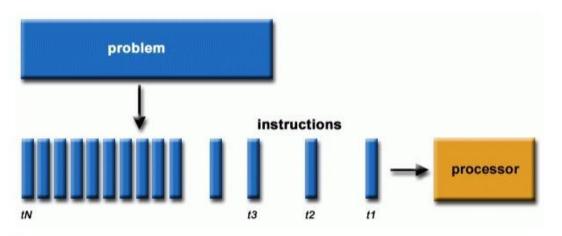


Serial Computing

Problem is broken into a *discrete series* of instructions

Instructions executed sequentially on a single processor (core)

Only **one** instruction can execute in each time step



Serial computing generic example





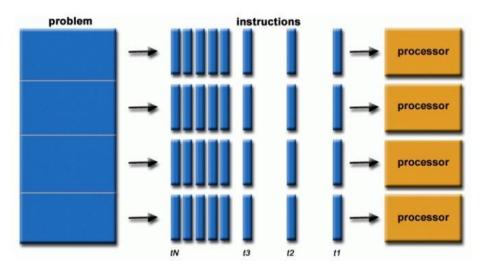
Parallel Computing

Problem divided into **discrete** parts that can be solved *concurrently*

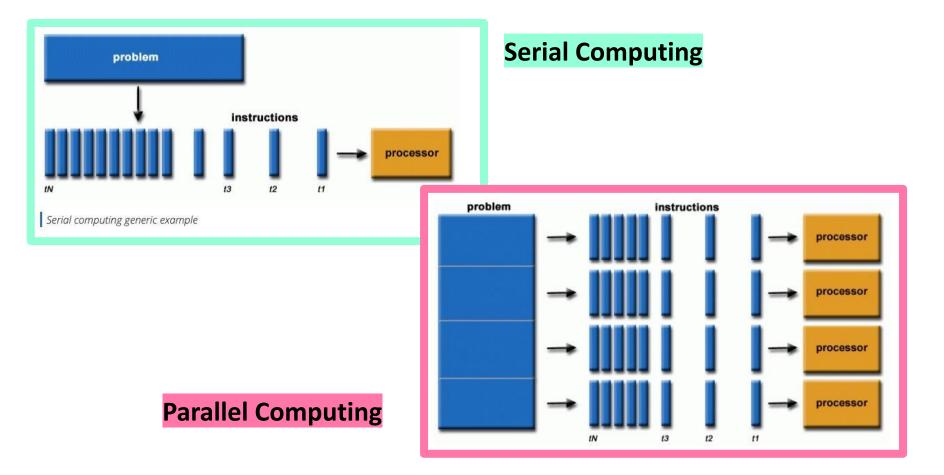
 \rightarrow Further divided to series of instructions

Instructions from each section execute simultaneously on *different* processors

Need to employ some overall coordination method



awrence Livermore lational Laboratory



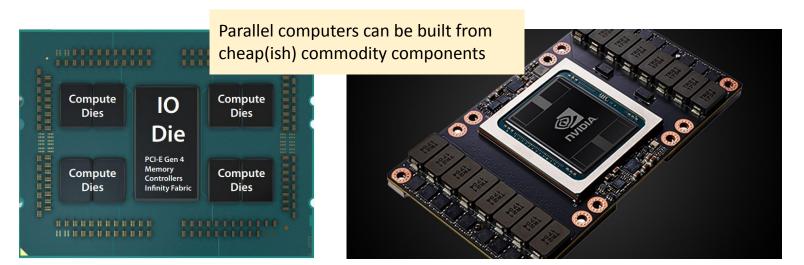
The **majority** of stand-alone computers today are *parallel* from a hardware perspective:

Multiple **functional** units

Multiple execution units/cores

Multiple hardware threads

(L1 cache, L2 cache, branch, prefetch, decode, floating-point, graphics processing (GPU), integer, etc.)



AMD EPYC Rome CPU Nvidia V100 GPU



Parallel Computers

Networks connect multiple stand-alone computers (*nodes*) to make larger parallel computer clusters.

Each <u>compute node</u> is a **multi**processor **parallel** computer in itself

 \rightarrow connected via a high-speed network

Special purpose nodes (also multiprocessor)

- \rightarrow GPU nodes
- \rightarrow high memory nodes

RACK 1 (STANDARD)		RACK 2 (STANDARD)		RACK 3 (STANDARD)		RACK 4 (STANDARD)		RACK 5 (GPU/MEM)	RACK 6 (NET/EXPANSION)
4x460P9 PDU, 68.8kW MAX		4x460P9 PDU, 68.8kW MAX		4x460P9 PDU, 68.8kW MAX		4x460P9 PDU, 68.8kW MAX		4x460P9 PDU, 68.8kW MAX	4x460P9 PDU, 68.8kW MAX
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1u41n1	r1u41n2	r2u41n1	r2u41n2	r3u41n1	r3u41n2	r4u41n1	r4u41n2	r5u41n1	spine1
1u40n1	r1u40n2	r2u40n1	r2u40n2	r3u40n1	r3u40n2	r4u40n1	r4u40n2	r6u40n1	cable management space
103961	r1u39n2	r2u39n1	r2u39n2	r3u39n1	r3u39n2	r4u39n1	r4u39n2	r5u39n1	spine2
	r1u38n2	r2u38n1	r2u38n2	r3u38n1	r3u38n2	r4u38n1	r4u38n2		cable management space
S	r1u37n2	r2u37n1	r2u37n2	r3u37n1	r3u37n2	r4u37n1	r4u37n2	Sectors 1	spine3
	r1u36n2	r2u36n1	r2u36n2	r3u36n1	r3u36n2	r4u36n1	r4u36n2		cable management space
	r1u35n2	r2u35n1	r2u35n2	r3u35n1	r3u35n2	r4u35n1	r4u35n2	ALC: NOT ALC	spine4
	r1u34n2	r2u34n1	r2u34n2	r3u34n1	r3u34n2	r4u34n1	r4u34n2		cable management space
	r1u33n2	r2u33n1	r2u33n2	r3u33n1	r3u33n2	r4u33n1	r4u33n2	resulting	edge1
	r1u32n2	r2u32n1	r2u32n2	r3u32n1	r3u32n2	r4u32n1	r4u32n2		cable management space
	r1u31n2	r2u31n1	r2u31n2	r3u31n1	r3u31n2	r4u31n1	r4u31n2	r5u31n1	mgmt
	r1u30n2	r2u30n1	r2u30n2	r3u30n1	r3u30n2	r4u30n1	r4u30n2	preserver.	switch mgmt gigabit switch
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		r2m2	1	r3m2		r4m2		r5u24n1	r6u24n1 r6u24n2
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	int space								
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	r1u17n2	r2u17n1	r2u17n2	r3u17n1	r3u17n2	r4u17n1	r4u17n2	r5u17n1	cable management space
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1u03n1	r1u03n2	r2u03n1	r2u03n2	r3u03n1	r3u03n2	r4u03n1	r4u03n2	r5u03n1	r6u03n1 r6u03n2
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								Trescold II.	1990 ALTI
-	64 nodes		64 nodes		64 nodes		64 nodes	16	40
	144 cores	6	144 cores	6	144 cores	6	144 cores	1536	3840

Puma Rack layout





Parallelization Accommodates Complexity

Natural processes can be accurately modeled with **high resolution** simulations or models

- large *number* of components
- multiple *types* of components
- interactions
- temporal sequence

Example: Natural Language Processing models have billions of parameters



Galaxy Formation

Planetary Movments

Climate Change

Real world phenomena can be simulated with parallel computing



Rush Hour Traffic

Plate Tectonics

Weather





Parallelization Decreases Time to Result

Tasks with a greater number of *independent calculations* will benefit from **dividing the load** between more processors



Working in parallel shortens completion time





Parallelization Provides Concurrency

A single processor can only perform one operation at a time.

Coordinating multiple processors allows for many operations to be performed in one clock cycle.

Example: The "shotgun" technique sequences a genome by breaking a long string of information into shorter segments, then reassemble

	Short-Insert Paired End Reads	De Novo Assembly	
	Read 1		
	Read 2		
	Long-Insert Paired End Reads (Mate Pair)		
R	tead 1		
-	_		
	Read 2		
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		National	e Livermore Laboratory

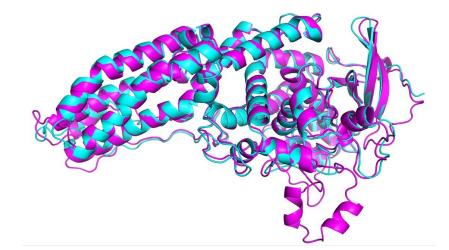


Parallelization Takes Advantage Of Non-Local Resources

- Users don't need to manage complex hardware
- Access powerful computing from anywhere with an internet connection
- Distributed computing allows for another meta-level parallelization

Example:

- Folding@Home is a distributed computing project to simulate protein dynamics.
- Supercomputers at the **three public Arizona universities** have contributed







<u>Node</u> –

an **individual computer**. A collection of them comprises a supercomputer

<u>CPU</u> –

AKA socket or **processor**. A physical device mounted on the motherboard. Puma nodes have two CPU's

<u> Core</u> –

- Part of CPU capable of conducting independent work.
- Puma CPU's have 48 cores for a total of 96 per node.
- 94 cores/node are usable
- However, in Slurm, "CPU" = "Core"







Process -

instance of a program, with access to its own memory, state and file descriptors

<u> Task</u> –

a *logically discrete* section of computational work. By default, Slurm allocates one CPU per task

s.

<u>Thread</u> –

highest level of code executed by a processor. Each process has at least one thread



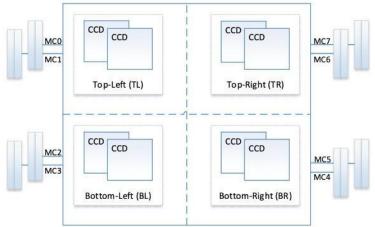


NUMA: Non-Uniform Memory Access.

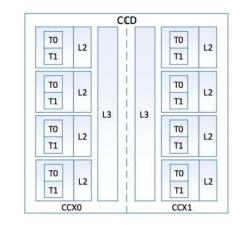
- \rightarrow Global address space shared by all cores.
- \rightarrow Memory is local to each processor or remote, which is slower.

Cache memory:

- \rightarrow memory that is much faster but smaller and expensive.
- \rightarrow L1 cache is on the core, L2 is next to each core and L3 is shared between 4 cores.



AMD Rome Core Complex





Types of HPC computation

Serial

- Computation runs on one core on one node
- Sometimes called High Throughput Computing

Shared Memory (AKA *multi-threading*)

- Single process with multiple threads
- Cores on single node work together
- Low level coordination
- Threads access shared memory space.

Distributed Memory (AKA *multi-node*)

- Cores on *multiple nodes* work *independently*
- High level coordination
- Coordination by passing messages over network.
- Supports large memory or many CPU workloads.





Massively Parallel –

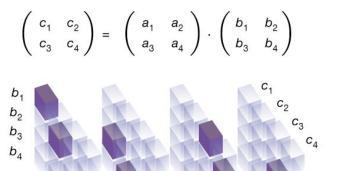
workloads that use many hundreds or thousands of cores

Embarrassingly Parallel –

A task that contains perfectly independent computations, e.g. matrix multiplication. Achieves ideal scaling.

MPI – Message Passing Interface

- Standard defining multi-node communication for distributed memory computing
- Implementations:
 - \rightarrow OpenMPI, Intel MPI, MPICH, MVAPICH
- OpenMPI and Intel MPI are encouraged on UA HPC clusters



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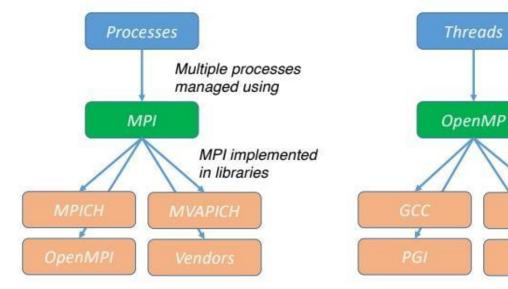
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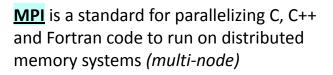
 a_1



 a_4

MPI and OpenMP big picture





OpenMP is an application programming interface (API) for shared-memory parallel programming in C, C++ and Fortran (*single node*)

Multiple threads managed using

by compilers

OpenMP implemented





von Neumann Computer Architecture

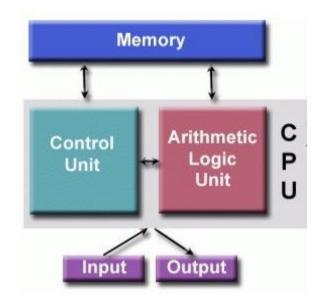
John von Neumann

- Hungarian mathematician
- authored the *general requirements for an electronic computer* in 1945

"stored-program computer"

- both program instructions and data are kept in electronic memory.
- earlier computers programmed through "hard wiring"

Since then, basically all computers have followed this basic design:





John von Neumann circa 1940s (Source: LANL archives)





Flynn's Classical Taxonomy (1966)

Scheme for classifying parallel computers.

- Distinguishes types of **multi-processor** computer architectures
- Classifies based on multiplicity of *Instruction* versus *Data* Streams
- Each of these can be *Single* or *Multiple*

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P1	P2	Pn
next instruct	next instruct	next instruct
store C(1)	call sub1(i,j)	10 continue
C(1)=A(1)*B(1)	sum=x*2	zeta=C(i)
load B(1)	x=y*z	alpha=w**3
load A(1)	call funcD	do 10 i=1,N
prev instruct	prev instruct	prev instruct

$\leftarrow \mathsf{Data} \rightarrow$

SISD	SIMD
Single Instruction stream	Single Instruction stream
Single Data stream	Multiple Data stream
MISD	MIMD
Multiple Instruction stream	Multiple Instruction stream
Single Data stream	Multiple Data stream

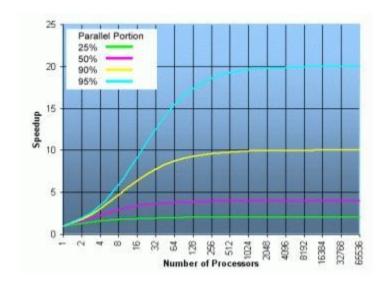
MIMD: HPC



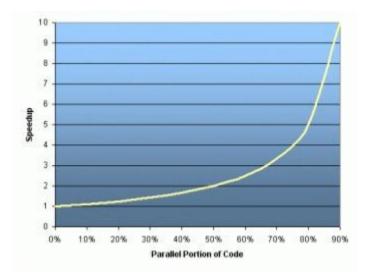


Amdahl's Law

theoretical maximum speedup is determined by the fraction of code that can be run in parallel.



Speedup when introducing more processors



Amdahl's law





Serialized code

A naïve inner product algorithm of two vectors of one million elements each

- All multiplications can be done in one time unit (parallel)
- Additions to a single accumulator in one million time units (serial)

Amdahl's Law

 If a fraction X of a computation is run in serial, the parallel speedup cannot be more than 1/X

Exercise

- what **fraction of the code** for the operation to the left is **parallelizable**?
- what is the expected fractional speedup compared to serial?



Scaling

Strong scaling (Amdahl):

Total problem size stays fixed as more processors are added.

Goal is to run the same problem size faster

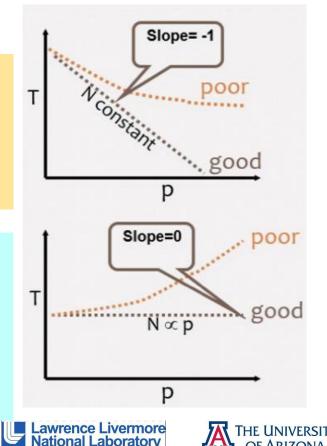
Perfect scaling means problem is solved in 1/P time (compared to serial)

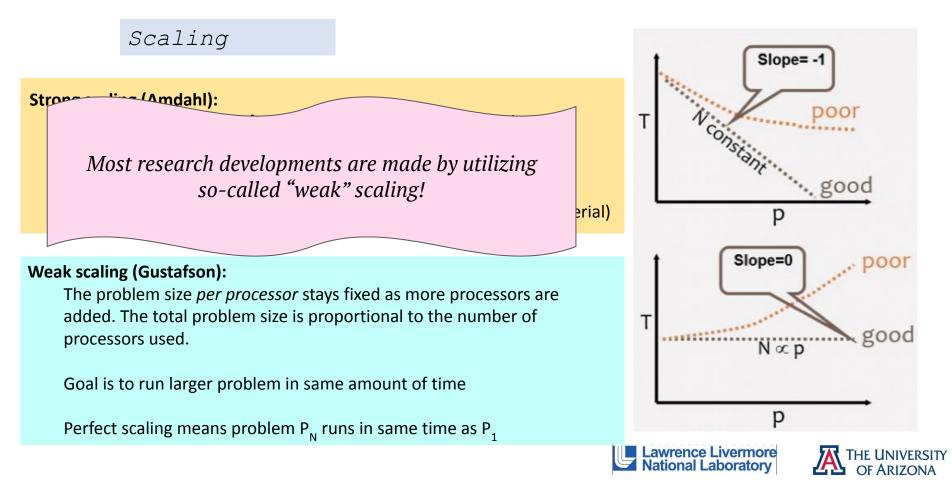
Weak scaling (Gustafson):

The problem size *per processor* stays fixed as more processors are added. The total problem size is proportional to the number of processors used.

Goal is to run larger problem in same amount of time

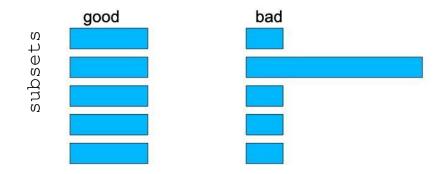
Perfect scaling means problem P_N runs in same time as P_1





Load Balance

The total amount of time to complete a parallel job is limited by the thread that takes the longest to finish



Computation per subset



Load Imbalance

Caused by non-uniform data distributions

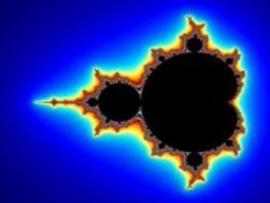
- large regions of very low density
- small regions of very high density

Occurs in astronomy, medical imaging, rendering, etc.

If the space is divided evenly across threads

- some threads will do very little work
 → low density = few elements
- some threads will do a *lot of work* → high density = many elements







MPI Implementations

Julia has anMPI language wrapper

MATLAB has its own parallel extension library implemented using MPI and PVM

Python Implementations of MPI include pyMPI, mpi4py, para, and MYMPI Boost C++ Libraries acquired Boost:MPI which include MPI Python Bindings.

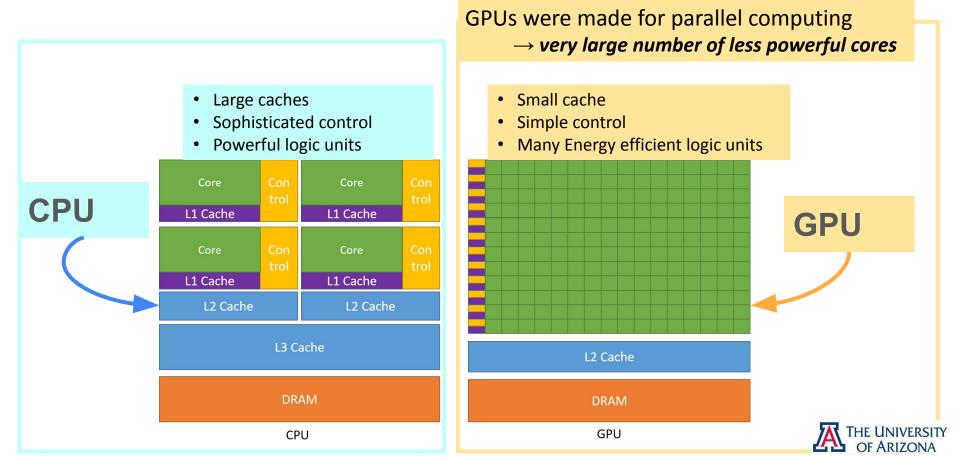
R Bindings of MPI include Rmpi and pbdMPI.

On HPC we support OpenMPI and Intel MPI. OpenMPI is a default module that is loaded with GCC 8.3

Intel MPI is provided when you unload OpenMPI and GCC, and then load the Intel compiler. By default, modules on HPC are compiled with OpenMPI

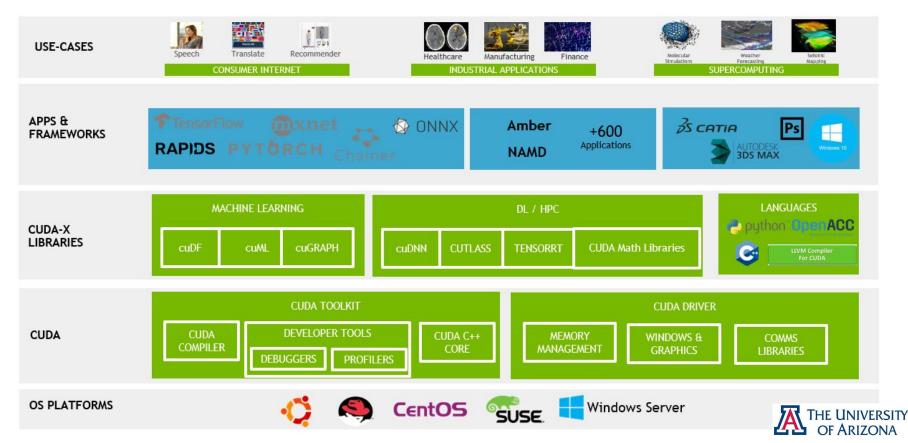


Parallel Computing CPU vs GPU



Parallel Computing GPU

Nvidia has an elaborate and growing ecosystem based on CUDA which provides parallel support



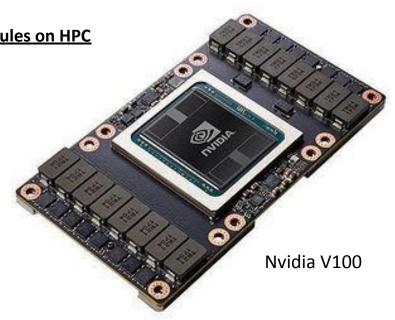
Parallel Computing GPU

CPUs for **sequential** code where latency matters

GPUs can be >20X faster for parallel code

Most of these applications are installed as modules on HPC

- Tensorflow
- PyTorch
- Matlab
- NAMD
- LAMMPS
- Quantum ESPRESSO
- Gromacs
- Relion
- Nvidia RAPIDS
- Julia
- Folding@home
- Caffe2
- Schrodinger





Parallel Programming

The art of designing parallel algorithms, such as to calculate part of the Fibonacci series ..

$$F_n=rac{arphi^n-(-arphi)^{-n}}{\sqrt{5}}=rac{arphi^n-(-arphi)^{-n}}{2arphi-1}.$$
 where

$$arphi = rac{1+\sqrt{5}}{2}pprox 1.61803\,39887\dots$$

... is beyond the scope of this workshop.

A more extensive overview of parallel programming can be found at:

https://hpc.llnl.gov/documentation/tutorials/introduction-parallel-computing-tutorial







Practical Parallel Computing on the UA HPC





Parallelization Use Cases

Research Code End User

- Expert in domain science
- Some segreeming experience
- Inter We can do this! h insights, writing publications
- Analysis consists of scripting and prewritten software packages
- **Needs functional understanding of** parallelization to speed up analysis and produce results more quickly

Research Software Engineer

- Ilm Heach you anything! Can't teach you anything! Majority of training in computer science

Realities of Developing Parallel Algorithms

- it can become very complex
- difficult to get right for non-trivial problems (weak scaling/strong scaling)

therefore..

- parallel programs for research developed by specialists
- implemented by researchers



Typical Research Workflow

- 1. Develop research question
- 2. Determine dataset

- 3. Find software that performs desired analysis
- 4. Download and learn software
- 5. Implement for research
 - a. simulations generate data by using theoretical, empirical principles
 - b. analysis software helps researchers extract useful information out of their experimental datasets using statistics, modeling, theory, etc



What does this mean?

Researchers often do not develop their own high performance analysis software from scratch

- tend to use existing modules and packages
- research software developers tackle the difficulties of implementing advanced computing algorithms

There are aspects of parallelization that are not always obvious or wellcommunicated to researchers

-> many programs <mark>need to be told how</mark> to run in parallel

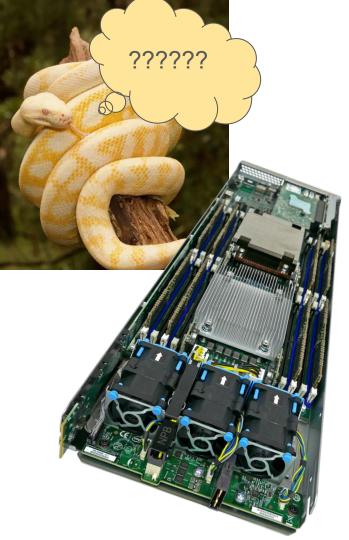
-> it does not happy automagically!

Check your software

- some programs are natively parallel
- many are not!

MPI is necessary to facilitate multi-node parallelization

- programming languages like python, R, etc do not automatically have information about the number of processors available, nor how to communicate between nodes
- must implement proper packages to enable these features



Adapting workloads, algorithms and code to parallel resources

- Simply dividing up a dataset and running independent serial analyses is generally more efficient than complex parallelization schemes
- Use **shared memory parallelism** when available (avoids internode communication overhead)
- Use **distributed memory parallelism** when one node does not provide enough memory or cores

Adapting workloads, algorithms and code to parallel resources

When writing or changing parallel code

- Do your homework
 - identify code hotspots
 - consider load balancing
- Depending on language, algorithm, and type of parallel resources, efficiently parallelizing an algorithm can range between
 - Adding a few lines of code
 - Complete algorithm redesign

Other factors: not all numerical operations are equally fast

- integer < single precision FP < double precision FP
- addition < multiplication < division

Performance Analysis and Tuning

Installed as a module

HPCToolkit/ hpctoolkit



HPCToolkit performance tools: measurement and analysis components

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Contributors	Issues	Discussions	Stars	Forks	

Installed in operating system



An integrated suite of tools for measurement and analysis of program performance Tools that can automatically detect many memory management and threading bugs, and profile your programs in detail.

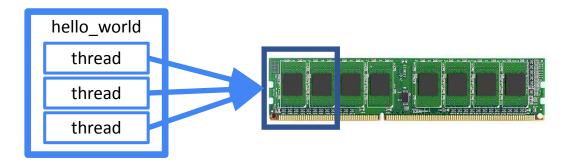


Multithreaded program schematic

1 process (task) with multiple threads

shared memory space

SLURM DIRECTIVES: #SBATCH --nodes=1 #SBATCH --ntasks=1 #SBATCH --cpus-per-task=3



./hello world <command line args>

Thread # control at program level:

- **OpenMP**:export OMP_NUM_THREADS=3
- Command line argument

OpenMP is probably the easiest (but not only) method for creating multithreaded programs

MPI schematic

mpirun -np 3 ./hello_world

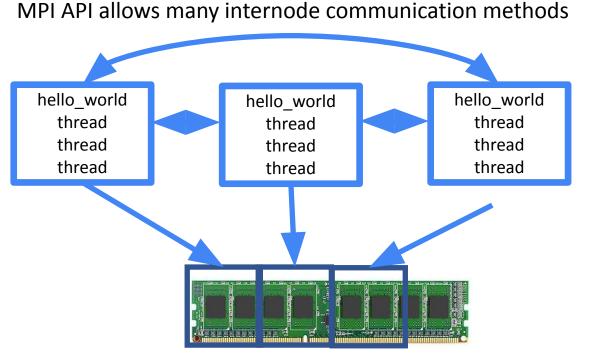
• 3 MPI processes (tasks)

- Each potentially multithreaded
- Independent memory spaces
- May be on different nodes

RELEVANT SLURM DIRECTIVES

#SBATCH --ntasks=3
#SBATCH --cpus-per-task=3

- #SBATCH --nodes=#
- #SBATCH --tasks-per-node=#





Array Jobs

Array jobs allow for meta-level parallelization.

Array jobs are useful if you have to run the same analysis on many different data sets, and if the order of completion does not matter

DO NOT USE FOR LOOPS TO SUBMIT JOBS – USE ARRAY JOBS

Parallel Computing on HPC – job arrays

Using an array to submit multiple independent jobs

Instead of this:



The above assumes that you have input files named input_file_1.in, input_file_2.in, etc

https://ua-researchcomputing-hpc.github.io/Array-and-Parallel/Basic-Array-Job/



Parallel Computing on HPC - MPI

MPI job submission - Hello World

This example Slurm script runs the Hello World executable on 10 cores on each of 3 nodes

```
#!/bin/bash
#SBATCH --job-name=Multi-Node-MPI-Job
#SBATCH --ntasks=30
#SBATCH --nodes=3
#SBATCH --ntasks-per-node=10
#SBATCH --time=00:01:00
#SBATCH --partition=standard
#SBATCH --account=YOUR_GROUP
```

module load gnu8 openmpi3

mpicc -o hello_world hello_world.c

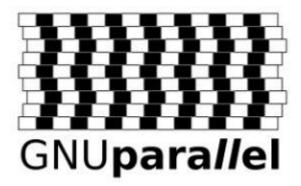
mpirun -np \$SLURM_NTASKS ./hello_world

more commentations and note: this is just an example. there is not really a good reason to use less than the maximum number of cores per CPU MPI is used for multi-node communication. and it is not necessary when running single-node, multithreaded jobs

<u>https://ua-researchcomputing-hpc.github.io/MPI-</u> <u>Examples/Multi-Node-MPI-Job/</u>

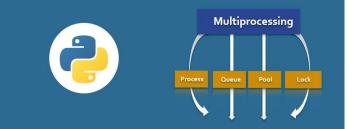






For people who live life in the parallel lane

Python Multiprocessing



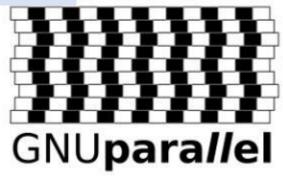
Parallel Computing on HPC - GNU parallel

Using GNU parallel to parallelize multiple tasks within one command

Access Compute Node

Use either batch job or interactive

```
$ elgato
$ interactive -a <your_group> -n 8
$ module load parallel
$ seq 1 100 | parallel 'DATE=$( date +"%T" ) && sleep 0.{} && echo \
                "Host: $(hostname) ; Date: $DATE; {}"'
```



For people who live life in the parallel lane

<u>Output</u>

Host:	junonia.hpc.arizona.edu	;	Date:	15:47:06;	1
Host:	junonia.hpc.arizona.edu	;	Date:	15:47:06;	2
Host:	junonia.hpc.arizona.edu	;	Date:	15:47:06;	3
Host:	junonia.hpc.arizona.edu	;	Date:	15:47:06;	4
Host:	junonia.hpc.arizona.edu	;	Date:	15:47:06;	5
Host:	junonia.hpc.arizona.edu	;	Date:	15:47:06;	6
Host:	junonia.hpc.arizona.edu	;	Date:	15:47:07;	10
Host:	junonia.hpc.arizona.edu	;	Date:	15:47:06;	7
Host:	junonia.hpc.arizona.edu	;	Date:	15:47:07;	11

https://ua-researchcomputinghpc.github.io/Array-and-Parallel /Basic-Parallel-Job/



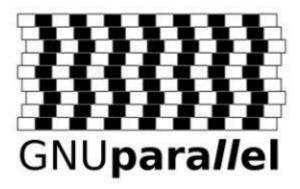
Parallel Computing on HPC - GNU parallel

Parallel will create as many jobs as inputs:

```
parallel echo {#} ::: A.txt B.txt C.txt D.txt E.txt
1
2
3
4
5
```

Limit number of jobs:

parallel -j 2 echo {%} ::: A.txt B.txt C.txt D.txt E.txt
1
2
1
2
1
2
1

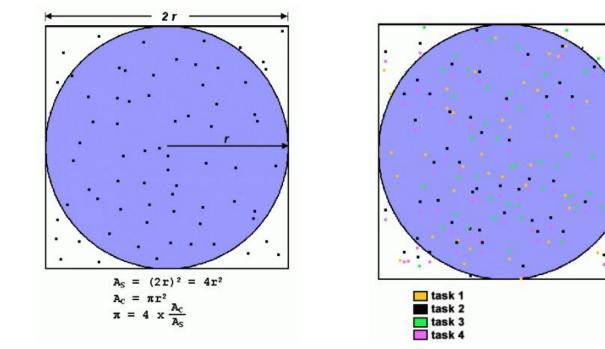


For people who live life in the parallel lane

Tons of examples online: https://www.gnu.org/software/parallel/parallel examples.html

A not-completely-trivial example of a parallelized calculation in Python

We can approximate Pi with a Monte-Carlo Simulation to guess area of circle

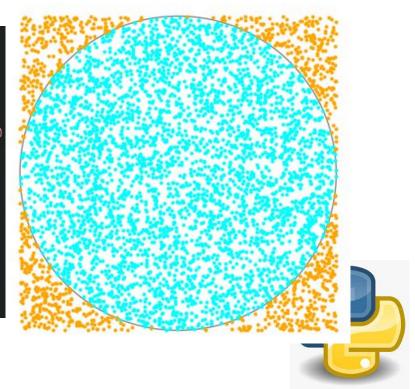




source: https://hpc.llnl.gov/documentation/tutorials/introduction-parallel-computing-tutorial##ExamplesPI

Serial Version:

```
n_points = 5000
circle_count = 0
points = np.zeros((n_points,2))
for i in np.arange(n_points):
    new_point = np.array([2.*(np.random.random()-0
    points[i] = new_point
    d = np.linalq.norm(new_point)
    if d < radius:</pre>
        circle_count += 1
pi_est = 4.0*circle_count/n_points
percent_diff = (pi_est - np.pi)/np.pi * 100
```



```
multiprocessing
 port numpy as np
 port time, os
                                                                                    Python "multiprocessing"
def monte_carlo_simulation(num):
                                                                                    library implementation
   circle_count = 0
                                                                                    \rightarrow enables single-node
   for i in np.arange(num):
       new_point = np.array([2.*(np.random.random()-0.5),2*(np.random.random()-0.5)])
                                                                                         parallelization
       if np.linalg.norm(new_point) < 1:</pre>
          circle_count += 1
                                        def master_worker_pi_calculation(num points, num tasks):
                                             from multiprocessing.pool import Pool
   return circle_count
                                            pool = Pool()
                                            batch_size = num_points // num_tasks
                                            pool = multiprocessing.Pool()
                                            results = []
                                             for _ in range(num_tasks):
                                                 task_count = pool.apply(monte_carlo_simulation,args=(batch_size,)
                                                 results.append(task_count)
                                            pool.close()
                                            pool.join()
                                             return sum(results)
```

from mpi4py import MPI import numpy as np import time, os

```
def monte_carlo_simulation(num):
    circle_count = 0
```

```
for i in np.arange(num):
    new_point = np.array([2. * (np.random.
```

```
if np.linalg.norm(new_point) < 1:
    circle_count += 1
```

return circle_count

```
def master_worker_pi_calculation(num_
    comm = MPI.COMM_WORLD
    rank = comm.Get_rank()
    size = comm.Get_size()
```

```
if rank == 0:
    total_circle_count = 0
    batch_size = num_points // (size - 1)
```

```
for i in range(1, size):
    comm.send(batch_size, dest=i)
```

```
for i in range(1, size):
    total_circle_count += comm.recv(source=i)
```

return total_circle_count
e:
betch circ _ comm recu(cou)

batch_size = comm.recv(source=0)
task_count = monte_carlo_simulation(batch_size)
comm.send(task_count, dest=0)

Python "**mpi4py**" library implementation

→ enables multi-node parallelization



Example batch script

#!/bin/bash *#SBATCH --job-name=picalc* #SBATCH --ntasks=8 #SBATCH --nodes=1 *#SBATCH --mem-per-cpu=4gb* #SBATCH --time=01:00:00 *#SBATCH --partition=standard #SBATCH --account=ejahn* #SBATCH --output=picalc.out #SBATCH --error=picalc.err module load openmpi3 python export TOTAL_NUM_POINTS=10000000 python picalc_serial.py mpirun -np 8 python picalc_parallel.py



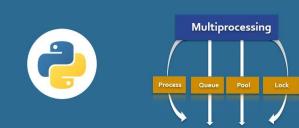
Parallel Computing on HPC -Python Multiprocessing Library

Estimate pi using a monte carlo simulation: <u>https://github.com/gelatinous-astronaut/picalc_example</u>

On HPC: Start an interactive session elgato interactive -a <your_group> -n 8 D Do II It Yourself

Set up environment
git clone <u>https://github.com/gelatinous-astronaut/picalc example.git
cd picalc_example
module load python
python3 -m venv --system-site-packages </path/to/env>
python3 -m pip install -upgrade pip
python3 -m pip install mpi4py multiprocessing</u>

Python Multiprocessing



Parallel Computing on HPC -Python Multiprocessing Library

Estimate pi using a monte carlo simulation: https://github.com/gelatinous-astronaut/picalc_example

On HPC:

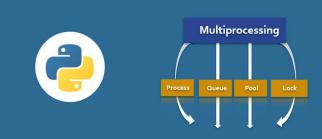
Run the code python3 picalc_serial.py

python3 picalc_multiprocessing.py

mpirun -n 8 python3 picalc_mpi4py.py



Python Multiprocessing



Parallel Computing on HPC - R

Quick Intro to Parallel Computing in R

https://nceas.github.io/oss-lessons/parallel-computing-in-r/parallel-computing-in-r.html

Using an Array with an R script

You can create an R script that generates 1000 randomized 1s and 0s, store them as a dataframe, then save the dataframe to an output file. Then run this R script as an array job.

https://ua-researchcomputing-hpc.github.io/R-Examples/R-Array-Jobs/

Check out the tidyverse – an opinionated collection of R packages designed for data science install.packages("tidyverse")

For an excellent hands-on Parallel Analysis in R tutorial:

https://github.com/ljdursi/beyond-single-core-R It covers these packages: parallel, foreach, bigmemory, Rdsm, pbdR

R Programming Language



Parallel Computing References

Introduction to Parallel Computing Tutorial

Author: Blaise Barney, Livermore Computing (retired), Donald Frederick, LLNL <u>https://hpc.llnl.gov/documentation/tutorials/introduction-parallel-computing-tutorial##Overview</u>

Recommended reading

"Introduction to Parallel Computing", Ananth Grama, Anshul Gupta, George Karypis, Vipin Kumar.

University of Oregon - Intel Parallel Computing Curriculum https://ipcc.cs.uoregon.edu/curriculum.html

An Introduction to Linux - <u>https://cvw.cac.cornell.edu/Linux/</u>

Linux Tutorial for Beginners: Introduction to Linux Operating System (link)

"Introduction to Linux" - Boston University (link)

"Parallel Processing in Python: A Practical Guide with Examples", Selva Prabhakaran (link)

