

# Python Parallelization

Summer 2025

Research Computing Services  
IS & T

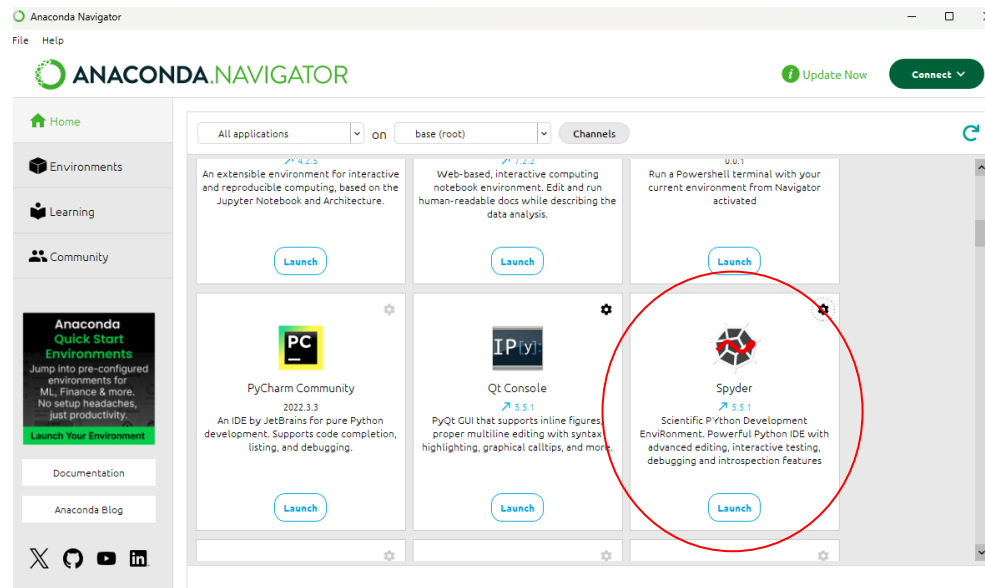
Download files:

<https://scv.bu.edu/examples/python/tutorials/PythonPar/>



# Start Spyder

- Your own computer:
  - Start Anaconda Navigator
  - Find Spyder and launch it.



- On SCC OnDemand:
  - Under Interactive Apps choose *Spyder*.
  - Load python3/3.12.4 and select 4 cores, then click the Launch button.

**Spyder**  
This app will launch an interactive Spyder desktop on a compute node.

List of modules to load (space separated)  
python3/3.12.4 Select Modules

Pre-Launch Command (optional)

Working Directory  
 Select Directory

The directory to start in. (Defaults to home directory.)

Number of hours  
4

Number of cores  
4

Number of gpus  
0

# Introduction

- Many programs can perform simultaneous operations, given multiple processors to perform the work.
- Generally speaking, the burden of managing this lies on the programmer.
- In this tutorial we'll go over a variety of ways to achieve parallelism in Python code.

# Limits on Program Speed

- **Input/Output (I/O):** The rate at which data can be read from a disk, a network file server, a remote server, a sensor, a user's physical inputs, etc. limits the performance of the program.
- **Memory:** The quantity of memory on the system limits performance.
- **CPU** (or compute): The speed of the processor is the limit on performance.
  - This is most commonly the case for scientific computing.

# Types of Parallelization

- On the SCC: queue parallelization.
  - You have N files to process. Submit N jobs.
  - Or, one [\*job array\*](#) that launches N jobs.
  - This often requires little to no changes to your code...
- Multiple Processes
  - Your program launches several copies of itself (or other programs) to solve the computational problem.
- Multiple Threads
  - Your program creates *threads*, which are parts of the **same** program that can execute independently of each other.
- Parallel Libraries
  - Use a library that internally implements some kind of parallelization.

# Performance Considerations

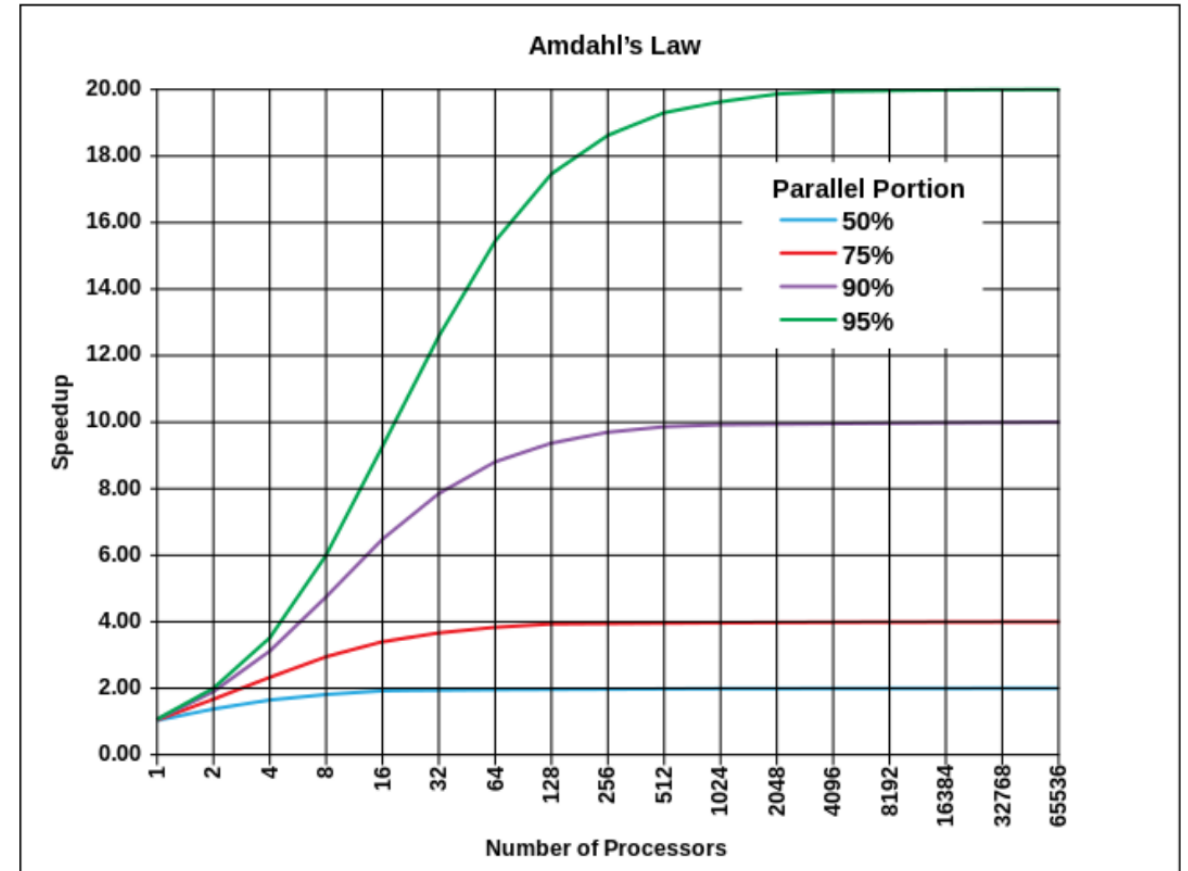
- Not every part of a program can benefit from parallelization.
- Some parts of program are inherently serial.
- Even for a function that can be done in parallel...
  - Is it worth the programming effort?
  - Is it worth the reduction in readability and ability to debug?
  - Does the function use up enough program time to make parallel computation worth the overhead?
- Parallelization is a form of optimization. Profile your code.
  - For more on profiling – see our Python Optimization tutorial.

## Amdahl's Law

- The speedup ratio  $S$  is the ratio of time between the serial code ( $T_1$ ) and the time when using  $N$  workers ( $T_N$ ):

$$S = \frac{T_1}{T_N} = \frac{T_1}{\left(f + \frac{1-f}{N}\right) T_1}$$

$N$  = number of threads or processes  
 $f$  = fraction of program that is serial



- This is the **theoretical** best speedup achievable with parallelization.

Figure from [Wikipedia](#).

# A word of caution

- When using the Python *multiprocessing* library, **always** use the “`if __name__`” convention in your main script:
- This will make your script work in interactive Python like Spyder.

```
import multiprocessing
# ...
# python script here with functions
# defined
# ...
def script_function():
    # do python stuff here
    with multiprocessing.Pool(4) as p:
        # code block etc ...

if __name__ == '__main__':
    script_function()
```

- It is **required** on Windows even in Jupyter notebooks.



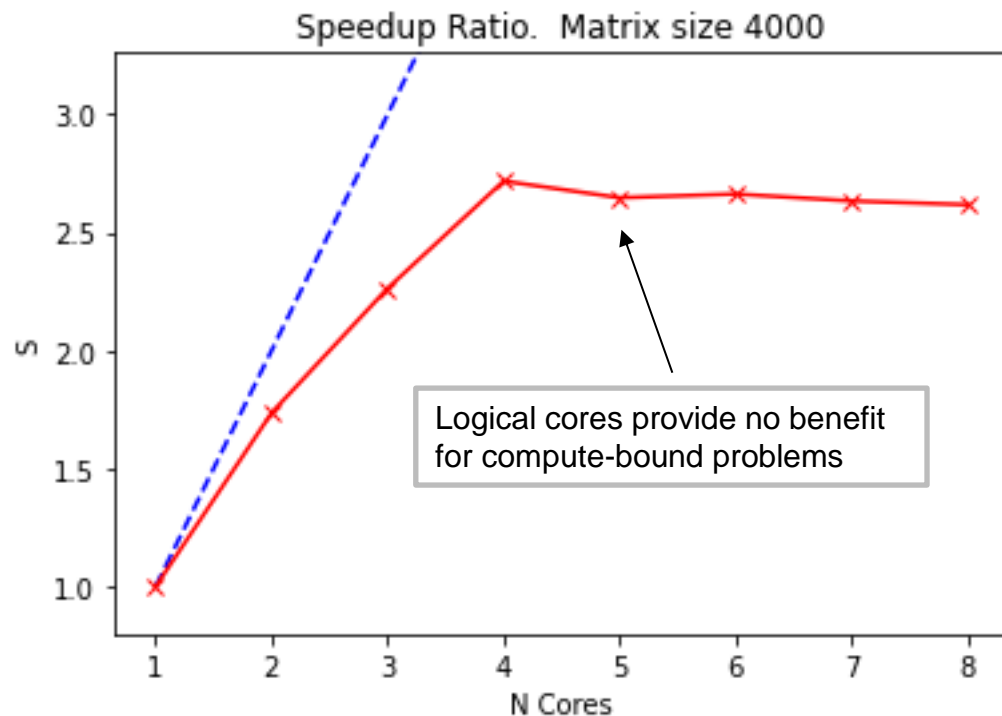
# How many cores should Python use?

- The example file *get\_n\_cores.py* provides a function that checks how many cores have been assigned to an SCC job.
  - Based on the common Python library *psutil*
- It will also work on your own computers and will choose the number of installed cores.
- Feel free to use this in your own code.

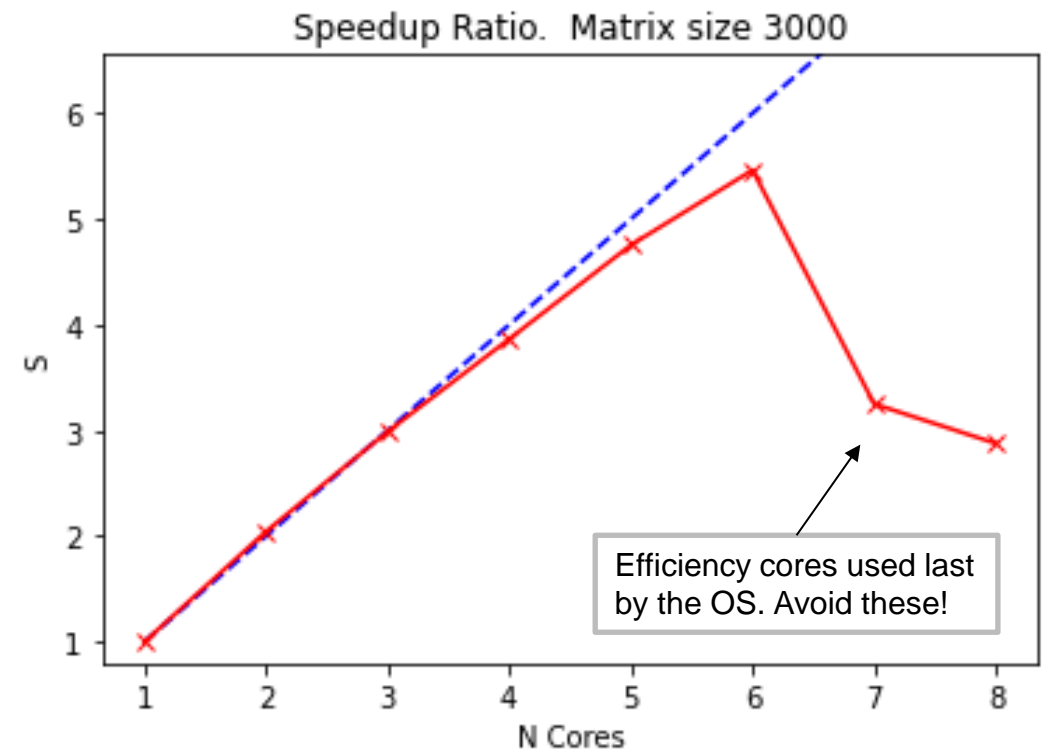
# Let's Try!

- In Spyder, open the file *lin\_alg.py*
- The computation: a linear algebra matrix-matrix multiplication.
  - Completely CPU-bound, scales well to multiple threads.
- How does your computation scale with the number of threads?
- It plots the speedup ratio. What did you expect? What if you change the size of the matrices?

# Logical, Physical, and Efficiency Cores



- Intel Core i7-1165G7
  - 4 real cores, 4 logical cores



- Macbook Pro (from 2021)
  - Apple M1 Pro CPU
    - 6 performance cores, 2 efficiency
      - About This Mac → More Info → System Report
    - `get_n_cores()` → reports 8 cores

# Python Language Parallelism

- Python provides a number of ways to perform parallel (aka concurrent) computations.
- Read the [official docs](#).

Library	Common Usage
<i>threading</i> and <i>asyncio</i>	I/O-bound programs. Example: web server, network service
<i>multiprocessing</i>	CPU-bound parallel execution.
<i>concurrent.futures</i>	Modern-style wrapper on top of threading & multiprocessing. Useful for GUIs or porting code to Python that uses this approach.
<i>subprocess</i>	Launching external processes.

# The Global Interpreter Lock

- The GIL limits the amount of multi-threading in the Python interpreter.
  - Originally introduced as part of Python's memory management system.
  - For more details, see [this explanation](#).
- Pure Python code runs in one thread only.
  - This is unlike languages like Java, C#, C++, Fortran, Matlab, or R where threads are easily used by the programmer.
- Multi-threaded code in Python is mostly implemented in external libraries.

# Python Threading

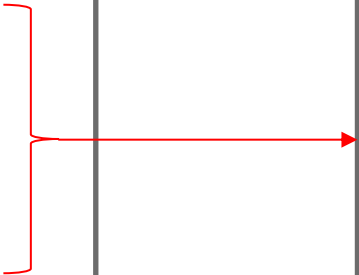
- The Python *threading* library allows for multiple threads to be created.
- Only 1 can actually execute at a time: **do not use this** for CPU-bound problems.
- This works well for I/O-bound problems.
- Each thread runs as soon as it has received data
  - Most of the threads are waiting for data from the disk, the network, the user, etc.
  - Application examples: Python web servers, file servers, network service, calling a web server API...

# Python Multiprocessing

- For CPU-bound problems multiple Python processes can be launched to do computations in parallel.
  - If you just want to parallelize a *for* loop, start here.
- The multiprocessing library handles inter-process communication automatically.
- Most convenient interface: the **Pool**, which provides a set of Python processes that divide work between them.

# Convert a loop

```
def xyz(x):  
    ''' here we do something that takes  
        some time OR gets called a lot  
        of times '''  
    # this is a regular function,  
    # nothing special  
    result = etc...  
    return result  
  
# somewhere else in your program:  
results = []  
# This loop takes a long time to  
# run:  
for elem in big_list:  
    results.append(xyz(elem))
```

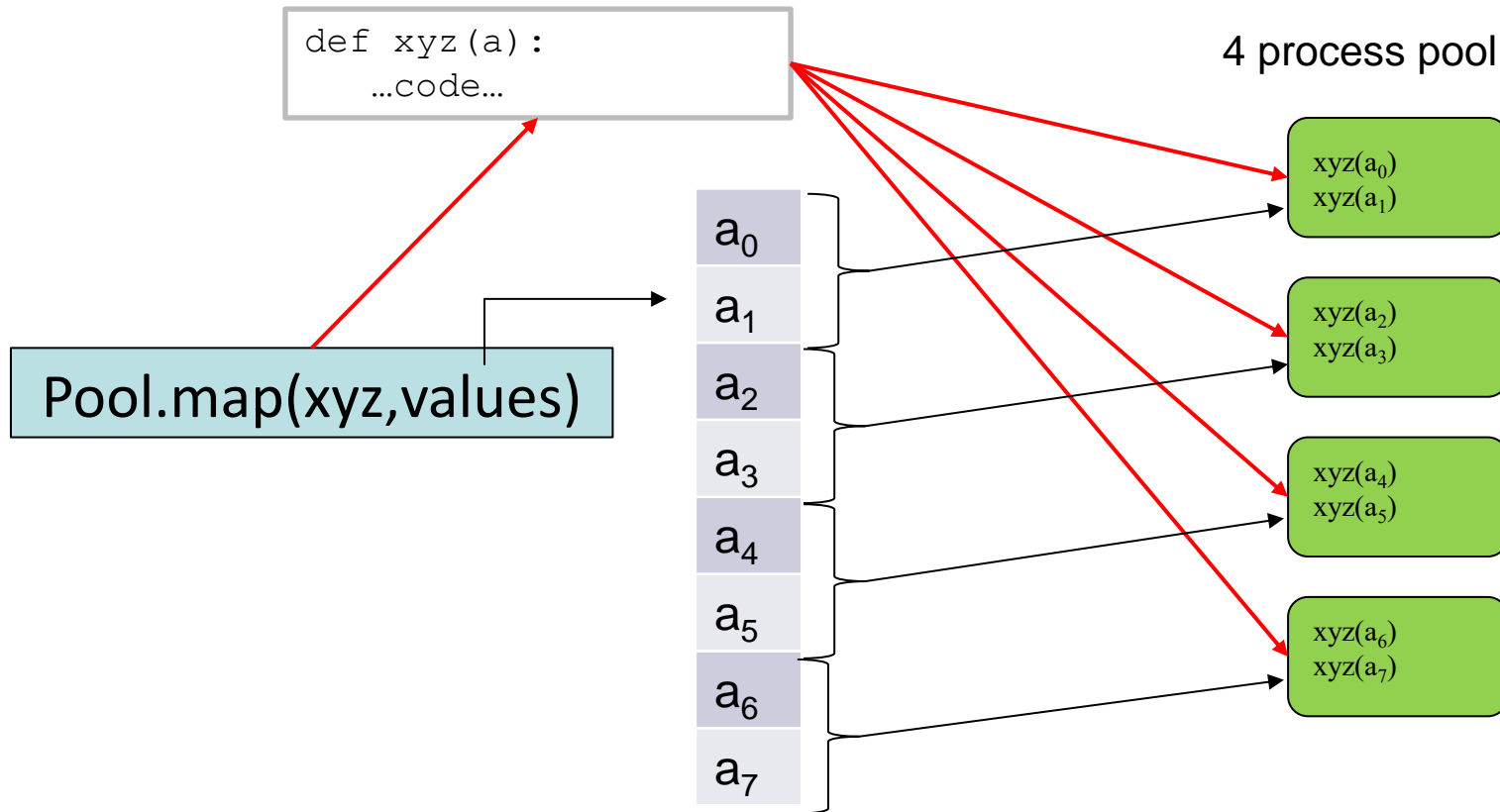


- xyz() can be easily parallelized much of the time.
- Watch for problems:
  - Printing to the screen → randomly interleaved output to the screen
  - Writing to a file → Same as the screen issue...don't parallel write to the same file!
  - Memory allocations → if the input & outputs use a lot of memory maybe limit parallelism to avoid excess memory usage.

```
import multiprocessing as mp  
  
# Replace the loop with a Pool and a map  
with mp.Pool(processes=nprocs) as pool:  
    results = pool.map(xyz, big_list)  
  
# xyz() now runs in parallel over the  
# elements of big_list.
```



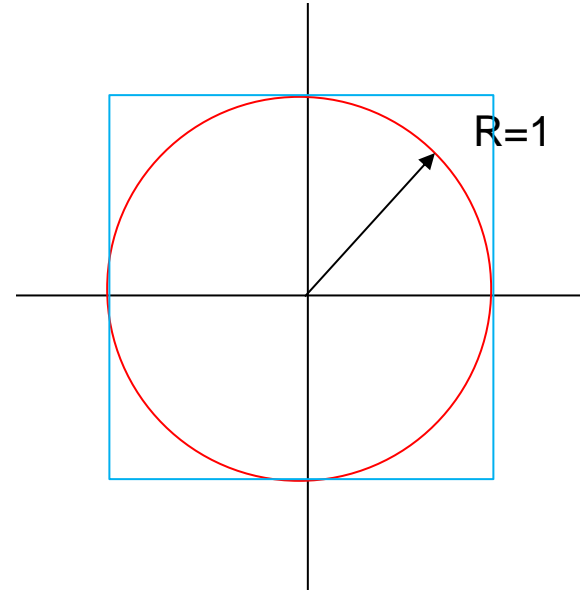
# How the Pool.map() Works



- A function is [pickled](#) and sent to each pool worker.
- The collection of data is split up, pickled, and sent to each worker.
- Each worker unpickles the function & data, runs the function on each element of the collection, pickles the result, and sends it back.
- The main process unpickles the results and puts them into a list.

# Example

- Open *pool\_basics.py*
- This calculates the value of  $\pi$



# multiprocessing.pool.Pool.map() options


- The Pool is the simplest way to add parallelism to Python code.
- Arguments: `map(function, iterable, chunksize)`
- **function**: the function to be applied to each element of the iterable
- **iterable**: a list, set, generator, dictionary, i.e. something that can be looped over
- **chunksize**: “This method chops the iterable into a number of chunks which it submits to the process pool as separate tasks. The (approximate) size of these chunks can be specified by setting *chunksize* to a positive integer.”

# Your turn to parallelize a problem...

- Open the file *my\_pool.py*
  - The problem: count the characters in 1M English words
  - You'll implement a Pool to parallelize the solution.


# Multiple iterables – Pool.starmap()

- To pass multiple arguments use starmap()



```
def xyz(a,b):  
    return a+b  
  
vals = [(1,2), (3,4)]  
  
with mp.Pool(processes=2) as pool:  
    sums = pool.starmap(xyz,vals)  
  
# 2 function calls happen in parallel:  
#     xyz(1,2)  
#     xyz(3,4)
```

- If you have 1 object and a list, try this to create a list for starmap:



```
import itertools  
a='arg1'  
b=range(3)  
  
list(zip(b,itertools.repeat(a)))  
# --> [(0, 'arg1'),  
#       (1, 'arg1'),  
#       (2, 'arg1')]
```

# Pool.imap() and Pool.imap\_unordered()

- *map()* has a disadvantage in that the iterable must be fully **in memory** before it can be distributed.
- *imap()* is lazier. It will assign chunks of work to each worker and pull them as needed from the iterable.
  - Generators can be used to save RAM in the main process.
- *imap\_unordered()* is similar but it does not guarantee the output order matches the input order.
  - Good for when computations take a varying amount of time.

# imap()

```
def xyz(a,b):  
    return a+b  
  
# A generator function  
def gen_vals(N):  
    for i in range(N):  
        # yield evens and odds  
        yield 2 * i, 2 * i + 1  
  
with mp.Pool(processes=2) as pool:  
    sums = pool.imap(xyz,gen_vals(1000),chunksize = 4)
```

- For pool worker 1, 4 calls to gen\_vals() are completed → [(0,1),(2,3),(4,5),(6,7)]
- This list is sent to worker 0.
  - Worker 0 calls xyz(0,1),then xyz(2,3) etc and returns the results in a list to the main Python process.
- Four more calls are done and that list goes to worker 1.
- When worker 0 is completed another 4 calls to gen\_vals() are done to create the next chunk, etc.
- The generator *gen\_vals()* never creates all 1000 sets of numbers in memory.

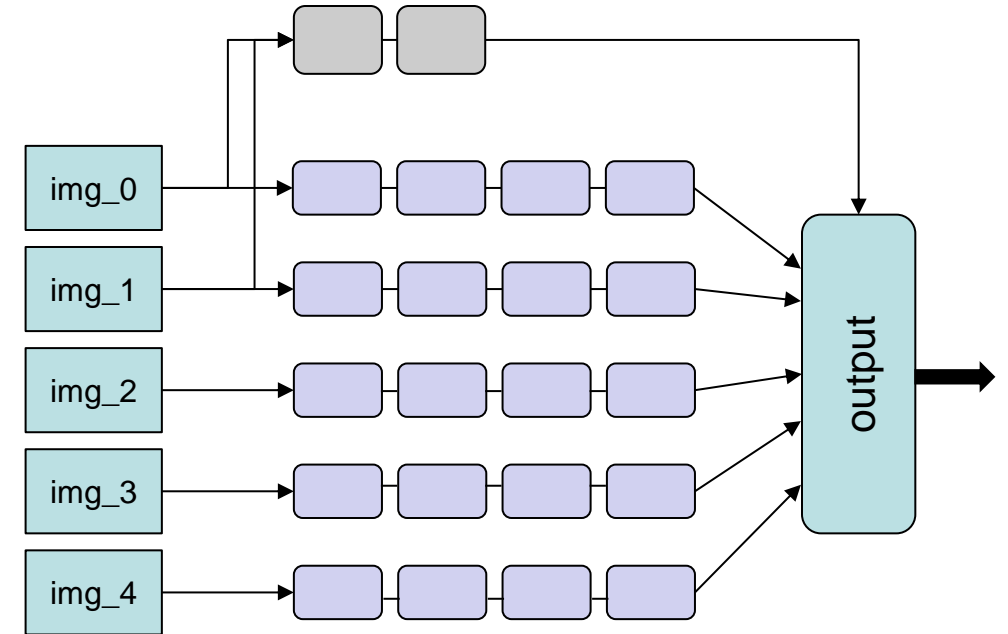
# Using map, starmap, imap, imap\_unordered

- If:
  - You have function calls being applied to some iterable (e.g. list of data objects, set of files, sets of simulation parameters, etc.)
  - The function call is *computationally expensive* – it takes a while to run.
  - Each function call is independent of the others.
    - Ex. Each input file in a list is read and processed separately.
- Then:
  - The multiprocessing.Pool is worth investigating for your code.
- Else:
  - Try the multiprocessing.Process code. This can be used to build more sophisticated parallelization strategies. Or investigate some other libraries...

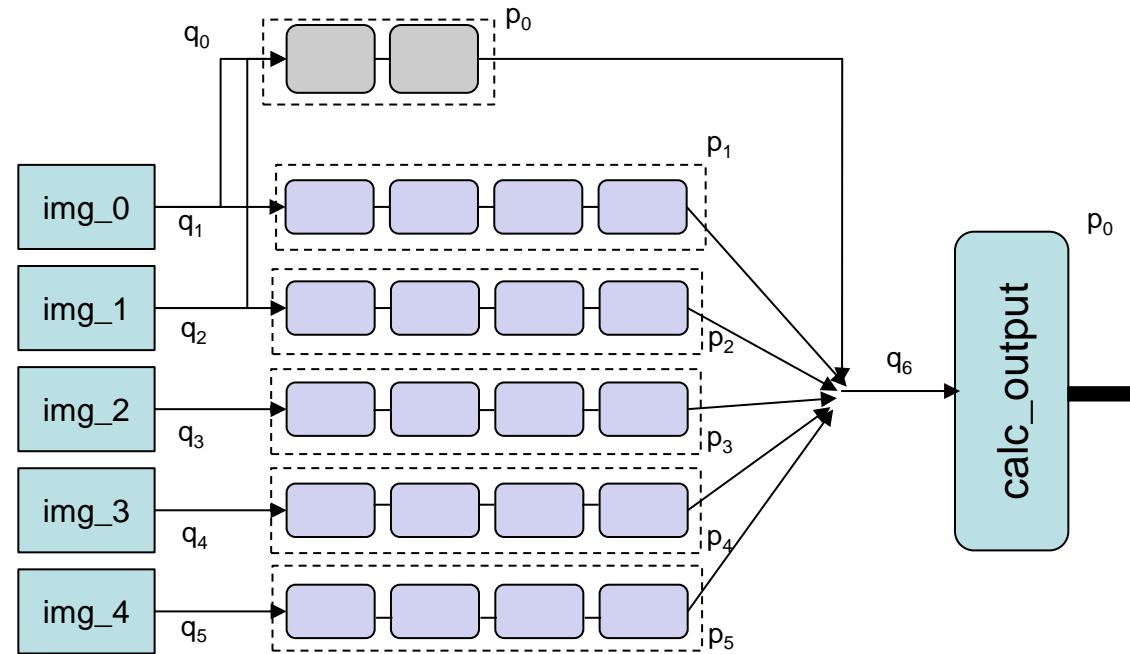


# More complex algorithms

- multiprocessing.Pool applies easily to for-loop parallelization. What about more complex patterns?
- Other tools in multiprocessing:
  - Start & stop processes that execute functions.
  - pipes, queues → send data between processes
  - Shared memory → processes access data without it being copied to them
  - locks, semaphores → protect serial-only resources from parallel access
    - Writing to a common file, updating shared memory, etc.



A pipeline where multiple images go through a series of filters. 2 images get copied to a separate set of filters. The *output* stage aggregates these results.



- Create Queue() objects for data transfer (`q0...q6`)
- Launch Process() objects to run functions (`p0...p5`)
  - Queues are connected in the Process() call
- Add data to the queues → processing starts
- Wait for data to return from the “output” process
- Shut down processes, destroy queues.
- This can all be accomplished in your Python code with the multiprocessing library.
  - How would you scale this to more cores?
  - How about fewer?
- It's a lot of work.
  - External libraries make this significantly easier.

# Parallelization with External Libraries

- When to look outside of standard Python:
  - Your dataset is greater than the amount of RAM you have available
    - You are dealing with large Pandas dataframes, numpy arrays, CSV files, database fetches, etc.
  - You have numpy-centered numeric calculations
    - Ex. A custom image processing algorithm
  - You want to scale past a single compute node
  - *mp* is causing problems due to RAM usage or poor scaling due to its multi-process nature
  - You want to implement more complex parallel algorithms

# Pandas in Parallel

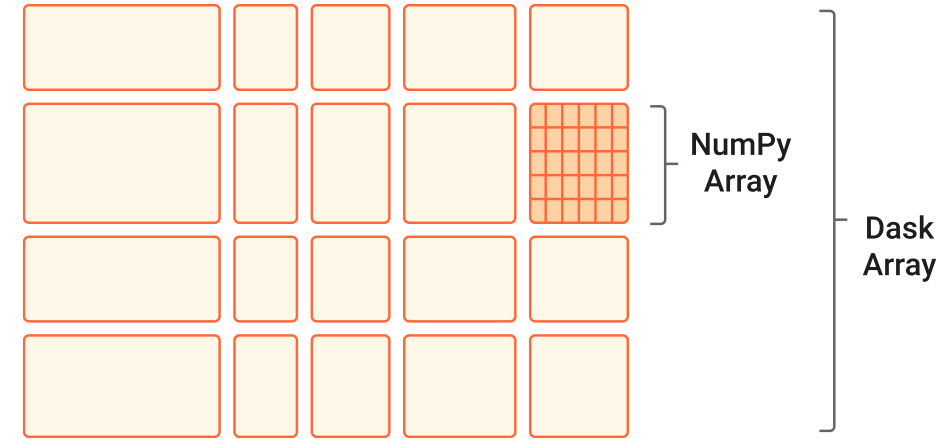


- [Modin](#)
  - Implements ~90% of the Pandas DataFrame API.
  - Autoscales Pandas calculations onto available cores.
  - Developed by UC Berkeley since 2018
- [parallel-pandas](#)
  - A simpler auto-parallelizing library for DataFrames.
- [Dask DataFrames](#)
  - Can autoscale and use available cores.
  - Built on top of Pandas.
- “[Polars](#) is a lightning fast DataFrame library/in-memory query engine.”
  - 2-20x faster than Pandas, for many operations
  - Efficiently uses memory and multiple cores
  - This is a relatively recent library, developed at RPI in 2020.
  - This is **not** compatible with your existing Pandas code. Using it requires a re-write of your code.



# Dask

- Dask supports parallelism beyond Pandas.
- [Dask Array](#): parallel numpy arrays
  - Includes efficient shared-memory access to these arrays
- [Dask Bag](#): parallelize generic functions like *map* or *groupby* on large collections
  - Example: read a file where each line is a JSON string. Convert to a format that can be converted to a DataFrame.
- [Dask Delayed](#): parallelize things that don't work with the other approaches.
  - For example, the image processing graph from a few slides ago.
- RCS is offering a Dask tutorial this summer – see the tutorial schedule.



# Ray Core



- [Ray](#) is a system for scaling up and parallelizing machine learning applications.
- [Ray Core](#) is its underlying distributed, parallel computation system.
  - You can use Ray Core to implement a wide variety of [parallel patterns](#).
  - They have an [example of using Ray to compute  \$\pi\$](#)  using the same algorithm we used earlier.
  - This is very useful if you're interested in *concurrent* programming
    - more generalized parallel programming
    - different parts of your program perform tasks in parallel
    - Example: your text editor auto-saves your file while running a spelling check while displaying text as you type.

# Common Parallel Libraries

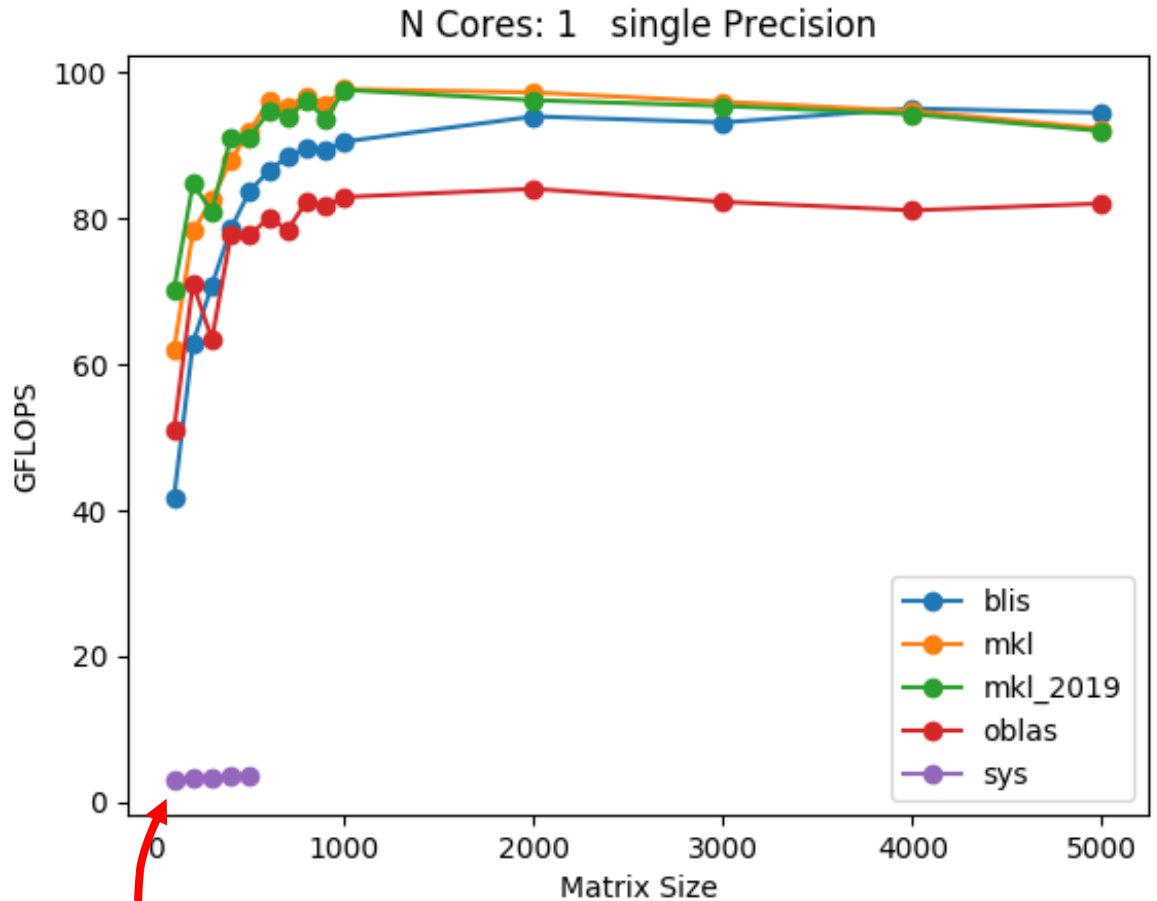
Python Library	Application	Underlying Library	Threading Lib.
numpy	Numeric algorithms	BLAS/LAPACK or MKL (C, usually)	OpenMP or MKL
cv2	Image processing	OpenCV (C++)	OpenMP or pthreads
Tensorflow, PyTorch, Jax	Machine learning	CUDA or OpenCL	OpenMP, pthreads, or GPU threads
numba	Compile/accelerate Python functions	numba C++ libs	Intel TBB
numexpr	Compile numpy code (older, not common)	numexpr libs	OpenMP

- Using Python for scientific computing naturally leads to the use of several libraries that support parallel computation using multiple threads. Those are built on top of a small set of threading libraries. Lots of other Python libraries use these “behind the scenes”.

- Glossary
  - BLAS: Basic Linear Algebra Subprograms
  - LAPACK: Linear Algebra Package
  - MKL: Intel Math Kernel Library
  - TBB: Intel Thread Building Blocks

# BLAS

- The **B**asic **L**inear **A**lgebra **S**ubprograms library provides a variety of functions for linear algebra type calculations.
- This underlies a staggering number of algorithms and computations including much of numpy and scipy.
- High performance threaded BLAS libraries continue to be an active area of computer science research.



- SCC benchmark.
- Note poor performance of default Linux system BLAS library!



# Numpy BLAS library

Anaconda, Windows

```
In [4]: np.show_config()
blas_mkl_info:
  libraries = ['blas', 'cblas', 'lapack', 'blas', 'cblas', 'lapack']
  library_dirs = ['D:\\bld\\numpy_1595523081734\\_h_env\\Library\\lib']
  define_macros = [('SCIPY_MKL_H', None), ('HAVE_CBLAS', None)]
  include_dirs = ['D:\\bld\\numpy_1595523081734\\_h_env\\Library\\include']
blas_opt_info:
  libraries = ['blas', 'cblas', 'lapack', 'blas', 'cblas', 'lapack', 'blas', 'cblas', 'lapack']
  library_dirs = ['D:\\bld\\numpy_1595523081734\\_h_env\\Library\\lib']
  define_macros = [('SCIPY_MKL_H', None), ('HAVE_CBLAS', None)]
  include_dirs = ['D:\\bld\\numpy_1595523081734\\_h_env\\Library\\include']
lapack_mkl_info:
  libraries = ['blas', 'cblas', 'lapack', 'blas', 'cblas', 'lapack']
  library_dirs = ['D:\\bld\\numpy_1595523081734\\_h_env\\Library\\lib']
  define_macros = [('SCIPY_MKL_H', None), ('HAVE_CBLAS', None)]
  include_dirs = ['D:\\bld\\numpy_1595523081734\\_h_env\\Library\\include']
lapack_opt_info:
  libraries = ['blas', 'cblas', 'lapack', 'blas', 'cblas', 'lapack', 'blas', 'cblas', 'lapack']
  library_dirs = ['D:\\bld\\numpy_1595523081734\\_h_env\\Library\\lib']
  define_macros = [('SCIPY_MKL_H', None), ('HAVE_CBLAS', None)]
  include_dirs = ['D:\\bld\\numpy_1595523081734\\_h_env\\Library\\include']
```

- You can see the exact libraries that Numpy is using with the command. The output will depend on the Python installation:

`numpy.show_config()`

python3/3.10.12 module on SCC

```
>>> np.show_config()
blas_armpl_info:
  NOT AVAILABLE
blas_mkl_info:
  NOT AVAILABLE
blis_info:
  libraries = ['blis', 'blis']
  library_dirs = ['/share/pkg.8/blis/0.9.0/install/lib']
  define_macros = [('HAVE_CBLAS', None)]
  include_dirs = ['/share/pkg.8/blis/0.9.0/install/include/blis']
  language = c
  runtime_library_dirs = ['/share/pkg.8/blis/0.9.0/install/lib']
blas_opt_info:
  libraries = ['blis', 'blis']
  library_dirs = ['/share/pkg.8/blis/0.9.0/install/lib']
  define_macros = [('HAVE_CBLAS', None)]
  include_dirs = ['/share/pkg.8/blis/0.9.0/install/include/blis']
  language = c
  runtime_library_dirs = ['/share/pkg.8/blis/0.9.0/install/lib']
lapack_armpl_info:
  NOT AVAILABLE
lapack_mkl_info:
  NOT AVAILABLE
openblas_lapack_info:
  NOT AVAILABLE
openblas_clapack_info:
  NOT AVAILABLE
flame_info:
  NOT AVAILABLE
accelerate_info:
  NOT AVAILABLE
lapack_info:
  libraries = ['lapack', 'lapack']
  library_dirs = ['/share/pkg.8/blis/0.9.0/install/lib']
  language = f77
  runtime_library_dirs = ['/share/pkg.8/blis/0.9.0/install/lib']
  extra_link_args = ['-L/share/pkg.8/blis/0.9.0/install/lib', '-llapack']
lapack_opt_info:
  libraries = ['lapack', 'lapack', 'blis', 'blis']
  library_dirs = ['/share/pkg.8/blis/0.9.0/install/lib']
  language = c
  runtime_library_dirs = ['/share/pkg.8/blis/0.9.0/install/lib']
  extra_link_args = ['-L/share/pkg.8/blis/0.9.0/install/lib', '-llapack']
  define_macros = [('HAVE_CBLAS', None), ('NO_ATLAS_INFO', 1)]
  include_dirs = ['/share/pkg.8/blis/0.9.0/install/include/blis']
Supported SIMD extensions in this NumPy install:
  baseline = SSE,SSE2,SSE3,SSSE3,SSE41,POPCNT,SSE42,AVX
  found = F16C,FMA3,AVX2,AVX512F,AVX512CD,AVX512_SKX,AVX512_CLX
  not found = AVX512_CNL,AVX512_ICL
```

# Enabling Threaded Libraries on the SCC

- Many libraries on the SCC that use multiple cores are built on the OpenMP or MKL threading libraries.
- The SCC disables this threading by default when you load Python or miniconda modules by setting environment variables.
  - Why? Because most jobs are single-threaded, and automatic threading leads to jobs using more cores than they should...and then the jobs are killed by the process reaper.
- In a compute job or at the command line you can enable these threads and they will automatically be used.

# Threading Environment Variables on the SCC

Variable	Threading Library
OMP_NUM_THREADS	OpenMP, MKL, numexpr
MKL_NUM_THREADS	MKL
NUMBA_NUM_THREADS	numba
NUMEXPR_NUM_THREADS	numexpr

- Setting these variables to a value >1 will enable automatic threading for code that uses the matching threading library.
- These should be set **before** running Python.
- Some libraries have their own internal mechanism can be used in place of the variable.
  - OpenCV example: `cv2.setNumThreads(integer_val)`

# Enable OpenMP Threading in a Job

Example qsub script:

- Request a multi-core job:
  - `qsub -pe omp 4`
- SCC jobs automatically set the variable `NSLOTS` to the number of requested cores.
- Environment variables can be set in various ways on different operating systems. Here is a [guide for Windows, Linux, and Mac OSX](#).

```
#!/bin/bash -l
# Ask for 4 cores.
#$ -pe omp 4

module load python3/3.10.5

# This sets the number of
# allowed threads to 4.
export OMP_NUM_THREADS=$NSLOTS

# Run your Python script, as
# this uses a lot of numpy code
# and might benefit from threads:
python myscript.py

#....did it run faster?
```

# numba

- [numba](#): auto-compiler for Python code.
  - Can compile code for GPU execution.
- Supports [auto-parallelization](#). Their *prange* function creates a parallelized loop.
- This lets you do low-level threading via Python.
- Thread control variable:  
NUMBA\_NUM\_THREADS
- Numba can also compile Python code so it is callable from C or C++.
- Read the [User Manual](#) and the [Reference Manual](#)
- Check out the assortment of [environment variables](#) that can be set to influence Numba behavior.

# numba usage

- Use the decorators  
*@numba.jit* or *@numba.njit*
- There are 2 modes:
  - object: Python types are used. numba must call out to Python to retrieve values.
  - nopython – no Python types are used, numba accesses values directly.
    - This is faster. Try to do this.



```
@numba.njit(parallel=True, fastmath=True)
def numba_jit_loop(mat):
    ''' A parallel double for loop over
        a 2D numpy ndarray '''
    rows, cols = mat.shape
    for i in numba.prange(rows):
        for j in numba.prange(cols):
            mat[i,j] = 2.0 * mat[i,j] - 1.0
    return mat
```

- *@numba.jit(nopython=True)*
- *@numba.njit*
  - These force nopython mode.
- *fastmath=True*: allows the compiler to use special CPU instructions.

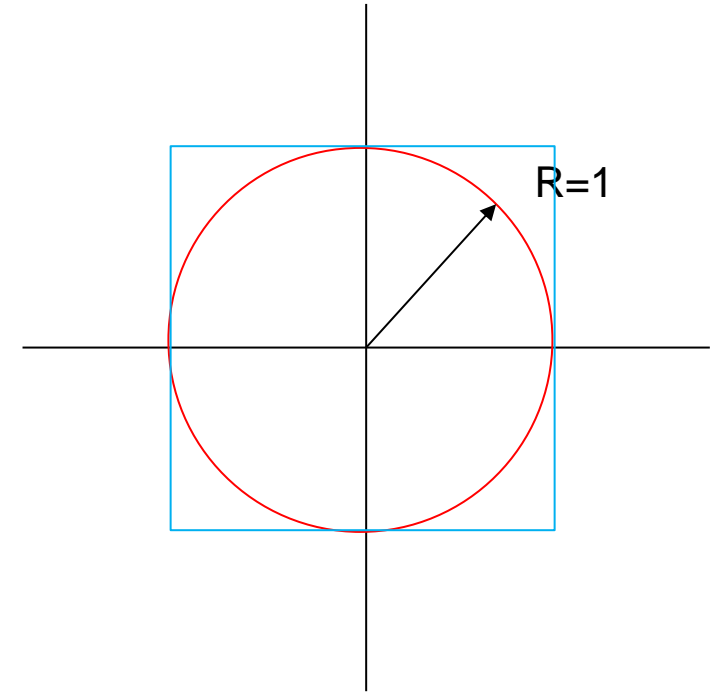
open *numba\_par.py*

# numba usage

- In general, use numpy ndarrays and functions with numba for the best performance.
  - Avoid calls to Python functions and sub-libraries
- numba'd functions should only call other numba'd functions
- This is a large library – test, profile, read the docs!



Let's calculate  $\pi$  with Python and numba



Open *numba\_pi.py*

# When is this useful?

- If your Python code heavily uses numpy data structures then it **may** benefit from automatic threading or compilation from *numba*.
- *numba* has been implementing a growing number of Python data types, [see their docs](#) for the latest.
- **Read the Numba docs.**
  - Numba is under continuous rapid development - new features appear all the time.
- Experiment! more threads is not always better.
  - The overhead of launching threads and distributing work can easily exceed the parallel execution speedup for small problems.



# End-of-course Evaluation Form

- Please visit this page and fill in the evaluation form for this course.
- Your feedback is highly valuable to the RCS team for the improvement and development of tutorials.
- If you visit this link later please make sure to select the correct tutorial – name, time, and location.

[http://scv.bu.edu/survey/tutorial\\_evaluation.html](http://scv.bu.edu/survey/tutorial_evaluation.html)