## Python Parallelization

v0.5

#### Research Computing Services IS & T



# Run Spyder

- Start the Anaconda Navigator
- Click on Spyder's Launch button
- Be patient...it takes a while to start.





## 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 <u>job array</u> 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

BOSTON

UNIVERSITY

 The speedup ratio S is the ratio of time between the serial code (T<sub>1</sub>) and the time when using N workers (T<sub>N</sub>):

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





• 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 ...
            == ' main ':
if
     name
      script function()
```

 It is <u>required</u> 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



The Python *psutil* library can't yet <u>auto-detect efficiency cores</u>. It will report them as physical 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 <u>official docs</u>.

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



# Python Language Parallelism

• There are many external libraries available.

| Library      | Common Usage   |
|--------------|--|
| <u>numba</u> | Function compiler, automatic multithreading  |
| <u>dask</u>  | Scalable auto-parallelizing library for data science, including scalable Pandas dataframes (and <b>much</b> more). Can use multiple compute nodes. |
| polars       | An auto-multithreaded alternative for Pandas.  |
| joblib       | A popular library for straightforward parallelization.   |
| <u>ray</u>   | Library that's popular in machine learning applications.   |
| <u>mpire</u> | A newer library, syntax is deliberately very similar to <i>multiprocessing</i> with higher performance.  |



## 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.





# How the Pool.map() Works



- A function is <u>pickle</u>d 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.



iterable (list, tuple, generator, set, etc.)

## 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()

If you have 1 object and a list, try this to create a list for 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)
```

```
BOSTON
UNIVERSITY
```

```
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  $\rightarrow$  [(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.



## *multiprocessing* is quite extensive...

#### More functionality exists for the Pool method.

- Shared memory between workers (avoids copies in interprocess communication)
- Asynchronous methods *map\_async*, *starmap\_async* 
  - These let the main process keep running after dispatching work.
- Process control:
  - Launch Python processes, do a calculation, wait for one or more processes to finish.
  - Interprocess communication using Queue and Pipe classes.
  - Synchronization using the Barrier, Lock, and Semaphore classes.
  - This can be used to implement much more elaborate parallelization strategies than the Pool at the expense of more programmer labor.



## 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...



## Parallelization with External Libraries

- Python *multiprocessing*: built into Python, works well on a broad array of problems, performs pretty well.
- When to look elsewhere:
  - 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



## Parallel Pandas?

- It's possible to do some parallel calculations with Pandas and multiprocessing but it's not straightforward.
- The strategy would be to send columns of dataframes to different processes and merge the results.
- This does not work with common Pandas operations like agg, groupby, query, etc.



## Dask <u>https://dask.org</u>

- Parallelizing pandas operations can be complex.
  - What if your data is too large to even read into a pandas DataFrame?
- Dask provides an equivalent DataFrame class that natively supports parallel computations.
  - Most pandas code can be handled via Dask just by importing the dask library instead of pandas.
  - Specific tutorial: <u>https://tutorial.dask.org/01\_dataframe.html</u>
  - Installed with SCC python3 modules.

BOSTON

UNIVERSITY

- Parallel computations can be run on a single computer or on a cluster using MPI communication.
- Large data sets can be loaded piecemeal to work within the memory limits of the computer.







## Dask

- Dask supports parallelism beyond Pandas.
- Dask Array: parallel numpy arrays
  - Includes efficient shared-memory access to these arrays
- <u>Dask Bag</u>: parallelize generic functions like *map* or *groupby* on large collections
  - Example: reformat every line of a CSV file so it can be converted to a DataFrame
- <u>Dask Delayed</u>: parallelize things that don't work with the other approaches.
  - This can be used in place of *multiprocessing* and can be applied to wider variety of programs than a *multiprocessing.Pool*



## **Polars**



- "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.
- If you are working with DataFrame style programs and Pandas:
  - Polars benchmarks as significantly faster than Pandas or Dask (which uses Pandas)
  - A conversion from Pandas to Polars is essentially a re-write of your program due to significant differences in syntax
- Parallelize/scale up existing Pandas codebase  $\rightarrow$  try dask
- New or smaller project  $\rightarrow$  try Polars to see if you like it



## Parallelization via Underlying Libraries

- Enabling parallelism in compiled code (C, C++, etc.) libraries that are being used by your Python code is very convenient.
- For many Python codes, this can be sufficient to achieve good parallel speedups without re-writing your code around multiprocessing.
- This is particularly true for codes that make heavy use of Pandas, numpy, and scipy data structures and routines.



## **Common Parallel Libraries**

| Python Library              | Underlying Lib. | Threading Lib.        |
|-----------------------------|-----------------|-----------------------|
| numpy (scipy, pandas, etc.) | BLAS or MKL     | OpenMP or MKL         |
| cv2                         | OpenCV (C++)    | OpenMP or pthreads    |
| Tensorflow, Keras, PyTorch  | CUDA or OpenCL  | OpenMP or GPU threads |
| numba                       | numba C++ libs  | Intel TBB             |
| numexpr                     | 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".
  - BLAS: Basic Linear Algebra Subprograms
  - MKL: Intel Math Kernel Library
  - TBB: Intel Thread Building Blocks



# BLAS

- The <u>Basic Linear Algebra</u>
   <u>S</u>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.





# 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 = cruntime 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 = f77runtime 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.

BOSTON

UNIVERSITY

- Some libraries have their own internal mechanism can be used in place of the variable.
  - OpenCV example: cv2.setNumThreads(integer\_val)

On Macs with the Accelerate library use VECLIB\_MAXIMUM\_THREADS instead of OMP\_NUM\_THREADS

# Enable OpenMP Threading in a Job

- Request a multi-core job:
  - qrsh -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 <u>guide for Windows</u>, <u>Linux, and Mac OSX</u>.

Example qsub script:

```
#!/bin/bash -1
# 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:
python myscript.py
#....did it run faster?
```



## numba

- <u>numba</u>: auto-compiler for Python code.
  - Can compile code for GPU execution.
- Supports <u>auto-parallelization</u>. 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 <u>User Manual</u> and the <u>Reference Manual</u>
- Check out the assortment of <u>environment</u> <u>variables</u> 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

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



## 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!





Open *numba\_pi.py* 



## numba

- Profiling is necessary with numba. Make sure numba provides a speedup before trying it in parallel.
- Open *numba\_par.py* for some examples of applying numba.

 Then we'll look at *numba\_convert.py* to see how an existing function might be converted to run faster under numba.



## When is this useful?

 If your Python code heavily uses numpy data structures then it may benefit from automatic threading or compilation from numba.

#### Read the Numba docs.

- Numba is under continuous rapid development new features appear all the time.
- Eperiment! more threads is not always better.
  - The overhead of launching threads and distributing work can easily exceed the parallel execution speedup.



## 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

