Parallel Computing On Pegasus: Python + Dask

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A Little Bit About Myself

- 1. Senior @ Arts & Science
- 2. Math & Computer Science majors
- 3. Physics & Economics Minor
- 4. Intern and future hire @ Coursera
- 5. 2 years @ Dr. Amelung's lab
- 6. Catch me surfing in the Bay Area starting next January!

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Wet Lab

The Problem

- 1. You write Python code, but it's slow. You think, "it must be easy to parallelize!"
- 2. You write Python code, but the data doesn't fit on 1 server. You think, "it must be easy to distribute the data across multiple servers!"
- 3. But after spending hours trying out different options, nothing works easily on Pegasus and it doesn't provide significant speedup
- 4. Enter Dask

A Possible Solution: Why not submit multiple Pegasus jobs?

- 1. It is possible to break up 1 large job on Pegasus into a lot of smaller jobs.
- 2. However, there are downsides:
 - a. What happens when a Pegasus job fails? You need to write retry code logic
 - b. What happens when an exception is thrown? You need to parse job error files to find the issue
 - You need to then write code to glue a bunch of output data back into 1 large output file

What is Dask?

- 1. "Dask provides advanced parallelism for analytics, enabling performance at scale for the tools you love"
- 2. Dask solves all the issues that the "multiple Pegasus job" solution created
 - a. Dask has smart retry
 - b. Code is executed from within Python and exceptions are thrown in the main Python script
 - c. You need minimal code glue
- 3. Dask can be run on 1 server or on High Performance Computing cluster
- 4. Dask implements close equivalents to NumPy and Pandas DataFrames
- 5. Dask has built in Diagnostics (So cool!)

The Main Parts of Dask

1. Dask Collections

a. Array, Bag, DataFrame

2. dask.delayed

 Build a sequence computations that are not executed immediately. Then execute all computation at once

3. dask.distributed

- a. "Submit" individual "futures" (think jobs) to a a scheduler that handles scheduling of execution
- b. This scheduler can operate on 1 machine or many
- c. What we will focus on

How This Talk Fits into the BIG Picture

- 1. Using Dask as a parallel framework
 - a. Tim Norris gave a talk on using Dask on a single machine. Notes are <u>here</u>
- 2. Using Dask on Pegasus
 - a. This talk
- 3. Using Dask Collections on Pegasus
 - a. Future work. Ideally we can write Numpy code and then just change import numpy to import dask.array and it would just work.
- 4. Using Dask as *the* way to run parallel code on Pegasus
 - a. Future work. This would require CCS to support Dask for all Pegasus users.

Dask: Clusters & Clients

- 1. Dask manages resources and scheduling with the Cluster abstraction
 - a. A Cluster handles all of the job execution, scheduling, and resource management
 - b. As a writer of parallel code, clusters are only configured
- 2. A Client is then how you interact with the Cluster
 - a. Once the Cluster is configured, you only need to interact with the client

Locally,

```
from dask.distributed import Client, LocalCluster
cluster = LocalCluster(n_workers=2, threads_per_worker=1)
client = Client(cluster)
```

dask.distributed: futures

- future = client.submit(f):
 - returns a Future, which refers to a remote result. This result may not yet be completed.
 Eventually it will complete
- res = future.result()
 - Blocks until task completes and data arrives

```
from dask.distributed import as_completed

futures = []
for i in range(10):
    future = client.submit(function, param1, param2, i)
    futures.append(future)

for future, result in as_completed(futures, with_results=True):
    # do something with output of each future
```

Using Dask: Let's Code!

1. Make sure you have the following packages installed:

```
a. pip install dask[complete]
```

- b. pip install numpy
- c. Pip install dask_jobqueue

Sequential Code

```
import numpy as np
import math
from timeit import timeit
Y DIM = 2000
def sequential main():
    two d array = np.random.rand(10000, Y DIM)
    vectorized sqrt = np.vectorize(lambda x: math.sqrt(x))
   output = vectorized sqrt(two d array)
    total = sum(sum(output))
    return total
print(timeit(stmt=sequential main, number=1))
```

Parallel Code

```
# My least favorite slide
# Sends data over the network
from dask.distributed import as completed
from dask.distributed import LocalCluster, Client
import numpy as np
import math
from timeit import timeit
Y DIM = 2000
NUM JOBS = 4
def parallel func(array):
    vectorized sqrt = np.vectorize(lambda x: math.sqrt(x))
    output = vectorized sqrt(array)
    total = sum(sum(output))
    return total
```

```
# Submit parallel jobs
def distributed main():
    two d array = np.random.rand(10000, Y DIM)
    futures = []
    for i in range(NUM JOBS):
        start = (i * Y DIM) // NUM JOBS
        end = ((i + 1) * Y DIM) // NUM JOBS
        future = client.submit(parallel_func,
                               two d array[:, start:end])
        futures.append(future)
    total = 0
    for future in as completed(futures):
        total += future.result()
    print(total)
    client.close()
    return total
if name == " main ":
    cluster = LocalCluster(n workers = 2,
                                     threads per worker = 1)
    client = Client(cluster)
    print(timeit(stmt=distributed main, number=1))
```

Parallel Code 2

```
# My 2nd least favorite slide
# Doesn't send data over the network
from dask.distributed import as completed
from dask.distributed import LocalCluster, Client
import numpy as np
import math
from timeit import timeit
Y DIM = 2000
NUM JOBS = 4
def parallel func2(dim):
    two_d_array = np.random.rand(10000, dim)
    vectorized_sqrt = np.vectorize(lambda x: math.sqrt(x))
    output = vectorized sqrt(two d array)
    total = sum(sum(output))
    return total
```

```
# Submit parallel jobs
def distributed main2():
    two-d-array = np.random.rand(10000, Y-DIM)
    futures = []
    for i in range(NUM JOBS):
        start = (i * Y DIM) // NUM JOBS
        end = ((i +-1) * Y DIM) // NUM JOBS
        future = client.submit(parallel func,
                               Y DIM // NUM JOBS)
        futures.append(future)
    total = 0
    for future in as completed(futures):
        total += future.result()
    print(total)
    client.close()
    return total
if name == " main ":
    cluster = LocalCluster(n workers = 2,
                                     threads per worker = 1)
    client = Client(cluster)
    print(timeit(stmt=distributed main, number=1))
```

Dask on Pegasus: dask_jobqueue

- dask_jobqueue allows you to execute Dask clusters on High Performance Computing supercomputers
 - a. LSF, PBS, and other clusters are supported
- 2. dask_jobqueue allows you to define a worker (which is realized as a job that is submitted to the HPC queue)
 - a. You can then replicate that worker many times

Dask on Pegasus: dask_jobqueue

The code barely changes:

```
cluster = LocalCluster()
```

becomes

```
from dask_jobqueue import LSFCluster
cluster = LSFCluster()
```

...well almost. The big idea is the same though. Only the cluster changes, but the client interface is the same

Dask on Pegasus: dask_jobqueue

```
from dask jobqueue import LSFCluster
import sys
cluster = LSFCluster(
              name='worker bee',
              queue='general',
                                        # the queue on Pegasus
              project='insarlab',
                                        # your project name
              cores=2,
              memory='2GB',
                                        # unused by Pegasus but a required param
              walltime='00:30',
                                         # how long the worker will run for
                                        # which network to use. NECESSARY PARAM
              interface='ib0',
              job extra=['-R "rusage[mem=2500]"', # how to define memory usage
              "-o WORKER-%J.out"], # where to write worker output files
              python = sys.executable,  # Where to look for Python executable
              config_name = 'lsf') # define your own config in a .yaml file
cluster.scale(20) # Create 20 workers with the parameters above
```

But we can hide this configuration in a .yaml config file

Dask on Pegasus: Diagnostics

- dask_jobqueue has a great way to visualize job execution. Here's how to do it:
 - a. ssh -L 8787:localhost:8787 dwg11@pegasus.ccs.miami.edu
 - b. Run your dask_jobqueue program
 - c. Locally, open up http://localhost:8787/status in your browser
- 2. Check out the link at the bottom of the slide for a great tutorial

dask_jobqueue: Configuration

- 1. You can configure your **LSFCluster** paramaters in a **.yaml** file. These parameters can be used multiple times by all programs using Dask
 - a. <u>Example</u>
- Currently, dask_jobqueue can only be configured once. For example, if instantiating an LSF cluster, the "Isf" configuration in your .yaml file will be used
- 3. After the next release of **dask_jobqueue**, you will be able to define configurations custom to your specific script, not just your specific cluster
 - a. The idea and Updates on when the code will be released

Dask on Pegasus: Knowing the hardware

- 1. Each **general** queue node has the following properties
 - a. 32 GB of memory
 - b. 16 cores
 - c. Try 1-2 GB per core
 - i. Optimal parameters that I found are 2 cores per "worker" and about 2-4 GB of memory per worker
- 2. Pegasus supports the **ib0** network. Use it!!!
 - a. This is the network over which data is transferred. The default is eth0 which is significantly slower (10x or more)
- 3. I found that I can only scale to about 40 workers before some of those worker's jobs stay in a pending state

Dask on Pegasus: Let's Try it Out!

Check out the code here:

https://github.com/2gotgrossman/dask-rsmas-presentation/blob/master/running_dask_on_pegasus.py

Dask on Pegasus is in Use: PySAR

- 1. Dask is being used by PySAR today on Pegasus
 - a. We have made a 5-15x speedup by running with Dask on Pegasus
- 2. The <u>code</u> and the <u>configuration</u>
- 3. The two relevant commits
- @YunjunZhang is defending his thesis on PySAR today at 2pm. Go check it out!