Introduction to IPCluster

Sponsored by AWS

Continuum Analytics



Unofficial Python Slogan

"CPU time is cheap, my time is expensive"



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 - 97% of the time, it's fast enough every time
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- But scientific workflows can get big
 - My expensive time spent waiting instead of doing science
 - Let's replace that with more cheap CPU time



 We have horsepower on-demand (AWS, Azure, Heroku, University Clusters)



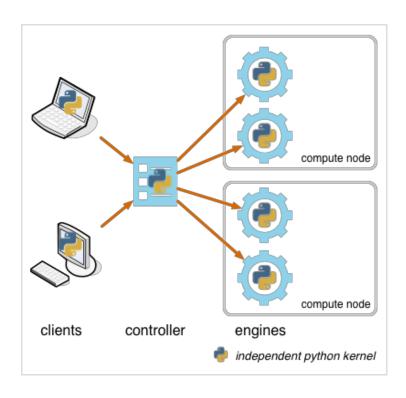


 We need a nice way to wrangle the data/code between nodes





IPCluster Overview





The Basic Use Case

- An embarassingly (pleasingly) parallel workflow
 - Little/no communication between jobs
 - Parameter scans, multiple BLAST queries, genetic algorithms, etc.



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- An embarassingly (pleasingly) parallel workflow
 - Little/no communication between jobs
 - Parameter scans, multiple BLAST queries, genetic algorithms, etc.
- Can utilize the CPU/GPU/RAM of many nodes simultaneously
- Moves relatively small amounts of data
 - Aggregates/subsets are good enough
 - The data is already on the nodes



Create a client object

```
from IPython.parallel import Client
client = Client()
```



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Construct a view

```
balanced = client.load_balanced_view()
# or
direct = client[:]
```



Define a function

```
def hello():
    return "Hello World!"
```



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Run it remotely

```
resp = direct.apply(hello)
resp.get()
```

```
Out[33]: ['Hello World!', 'Hello World!', 'Hello World!']
```

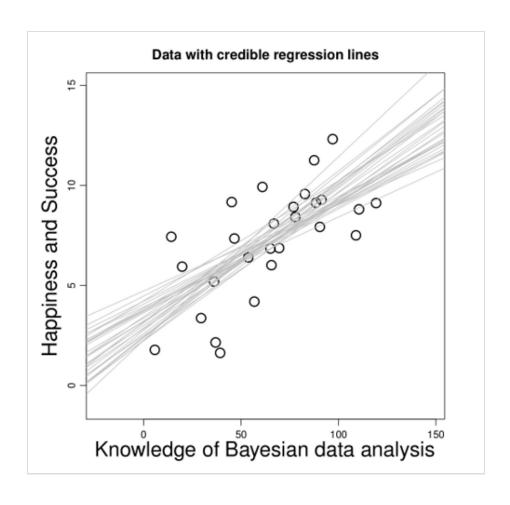


Easy Demo





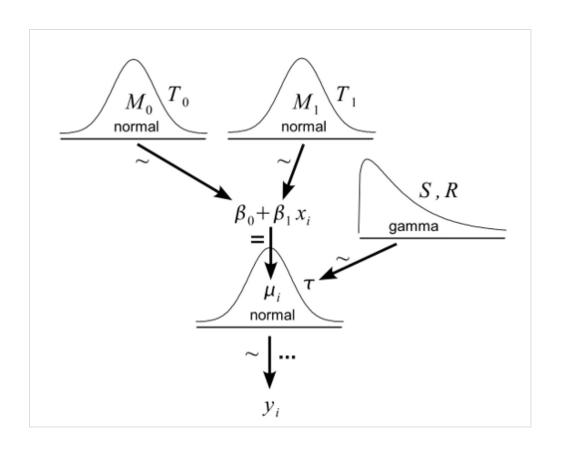
Use Case: Bayesian Estimation



Images: John K Kruschke

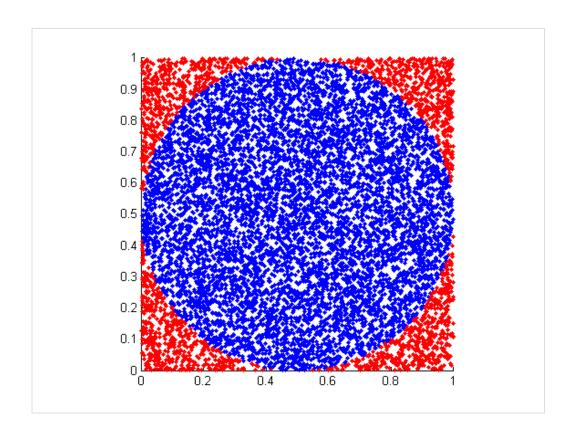


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Bayesian Demo

