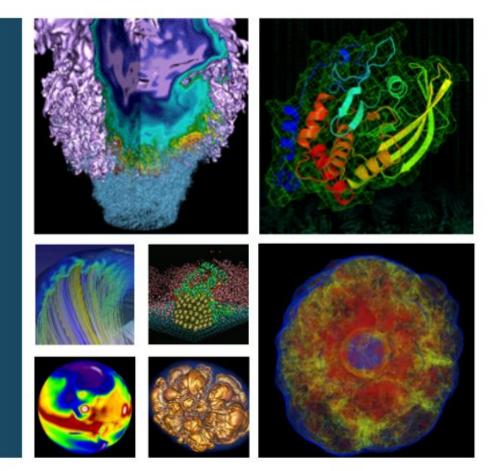
## An Introduction to Python at NERSC

### NERSC Data Day 2016





#### Rollin Thomas Data & Analytics Services Group 2016-08-22







- 1. Do you use Python?
- 2. Do you use Python 3 (yet)?
- 3. Do you use Anaconda Python?
- 4. Have you ever used numpy/scipy?
- 5. ... multiprocessing?
- 6. ... mpi4py?
- 7. ... IPython/Jupyter?
- 8. ... let's make it 8 questions.



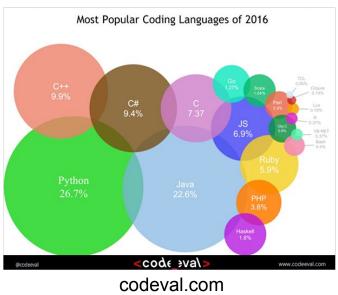




### Python is Popular

Aug 2016	Aug 2015	Change	Programming Language	Ratings	Change
1	1		Java	19.010%	-0.26%
2	2		С	11.303%	-3.43%
3	3		C++	5.800%	-1.94%
4	4		C#	4.907%	+0.07%
5	5		Python	4.404%	+0.34%
6	7	^	PHP	3.173%	+0.44%
7	9	^	JavaScript	2.705%	+0.54%
8	8		Visual Basic .NET	2.518%	-0.19%
9	10	^	Perl	<b>2.511%</b>	+0.39%
10	12	~	Assembly language	2.364%	+0.60%

#### www.tiobe.com/tiobe-index





<text><image><section-header><section-header><section-header><section-header><section-header><section-header>

#### bestprogramminglanguagefor.me



## Why Python?



Clean, clear syntax makes it very easy to learn.

Multi-paradigm interpreted language.

Extremely popular language for teaching beginners...

... but stays useful beyond the beginner phase of programming:

Powerful data structures and constructs built into the language and standard libraries. Leveraging of C/C++/Fortran.

Huge collection of useful open source packages to re-use and extend.

```
interface import Model
class BasicModel ( Model ) :
    def __init__( self, gaussian_process, training_data, upda
        self.gaussian_process = gaussian_process
        self training data
                              training_data
        training size
                            = len( self training data )
        self._input_diffs
                           = ( self.training data inputs[ No
                self.training data inputs[ :, None ] )
        self. gram
                            = numpy.zeros( ( training size, t
        self. log gram det =
                              None
        self, inv gram
                            = numpy_zeros like( self, gram )
        self_residuals
                            = numpy.zeros( training size )
        self, inv gram resp = numpy.zeros( training size )
        if update :
            self._update()
    @property
    def log p( self ) :
        return -0.5 * ( numpy.dot( self._residuals, self._inv
                self._log_gram_det + len( self.training_data
                numpy.log( 2.0 * numpy.pi ) )
    property
    def hyperparameters( self ) :
        deque = self.gaussian_process.mean_function.hyperpara
        deque.extend( self.gaussian_process.covariance_functi
        return deque
    Mhyperparameters.setter
    def hyperparameters( self, iterable ) :
        deque = collections.deque( iterable )
        self gaussian process mean function take hyperparame
```





### Python at NERSC



Supporting Python is no longer optional at HPC centers like NERSC.

Maximizing Python performance on systems like Cori and Edison can be **challenging**:

- Interpreted, dynamic languages are harder to optimize.
- Python's global interpreter lock is an issue for thread-level parallelism.
- Language design and implementation choices made without considering an HPC environment.

At the same time, users want NERSC to provide a familiar and portable Python environment.

from interface import Model	
class BasicModel ( Model ) :	
<pre>definit( self, gaussian_process, training_data, upda self.gaussian_process = gaussian_process self.training_data = training_data</pre>	
<pre>training_size = len( self training_data ) self _input_diffs = ( self training_data inputs[ No</pre>	
<pre>self_gram = numpy.zeros((training_size, t self_log_gram_det = None self_inv_gram = numpy.zeros_like(self_gram) self_residuals = numpy.zeros(training_size) self_inv_gram_resp = numpy.zeros(training_size)</pre>	
<pre>if update :     self _update()</pre>	and the second
<pre>@property def log_p( self ) :     return -0.5 * ( numpy dot( self 'residuals, self _inv         self _log_gram_det + len( self training_data         numpy log( 2.0 * numpy pi ) )</pre>	
<pre>@property def hyperparameters( self ) : deque _ self gaussian_process mean_function hyperpara deque extend( self.gaussian_process covariance_functi return deque</pre>	
<pre>Whyperparameters.setter def hyperparameters( self, iterable ) :     deque collections_deque( iterable )     self gaussian_process_mean_function_take_hyperparamet</pre>	





Python Modules at NERSC

Environment modules: Environment modules project: http://modules.sourceforge.net/

Always\* "module load python" Don't use /usr/bin/python. Using #!/usr/bin/env python: OK!

What is there? module avail python

\* Unless you install your own Python somehow.









### "NERSC-Built" Python

- Python "base" module.
- Add-on modules as desired.
- Meta-module simplifies setup.



https://docs.continuum.io/anaconda/

### **Anaconda Python**

- "Distribution" for large-scale data analytics, and scientific computing.
- ~200 packages but there is also "miniconda" bare-bones starter.
- Simplified package management and deployment (conda tool).
- Monolithic module, some add-on modules (h5py-parallel).







### Python Modules on Edison

### NERSC-built:

module load python[/2.7.9]

python\_base/2.7.9
numpy/1.9.2
scipy/0.15.1
matplotlib/1.4.3
ipython/3.1.0

Anaconda:

module load python/2.7-anaconda
module load python/3.5-anaconda

Above are the only currently recommended Python modules for Edison.







(default)



# GE

There aren't any.

**NERSC-built:** 

Python Modules on Cori

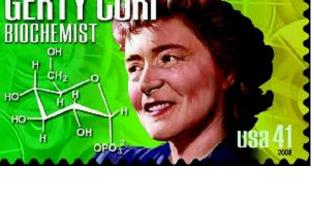
## Anaconda:

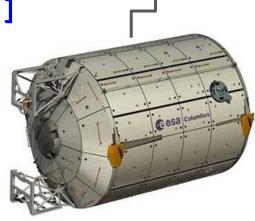
module load python[/2.7-anaconda]
module load python/3.5-anaconda

Above are the only currently recommended Python modules for Cori.

#### BIOCHEMIST

python/2.7-anaconda









## **Do-It-Yourself Python at NERSC**



Anaconda Environment under Modules: module load python/2.7-anaconda conda create -p \$PREFIX numpy... conda create -n myenv numpy... (won't work for users without .condarc defining "envs\_dirs") conda install basemap yt...

#### Your own Anaconda or Miniconda installation:

module unload python
wget https://repo.continuum.io/miniconda/Miniconda2-latest-Linux-x86\_64.sh
/bin/bash Miniconda2-latest-Linux-x86\_64.sh -b -p \$PREFIX
export PATH=\$PREFIX/bin:\$PATH
conda install basemap yt...

#### Tips:

- Conda environments do *not* mix with virtualenv.
- Several ML environments via Anaconda at NERSC.





\*https://github.com/ContinuumIO/mkl-service



## Node Parallelism: Threaded Libraries

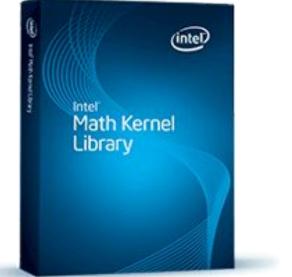
### Anaconda Python provides access to Intel Math Kernel Library (MKL) *for free:*

numpy scipy scikit-learn numexpr

### MKL Service functions\*:

```
>>> import mkl
>>> mkl.get_max_threads()
2
>>> mkl.set_num_threads(1)
>>> mkl.get_max_threads()
1
```







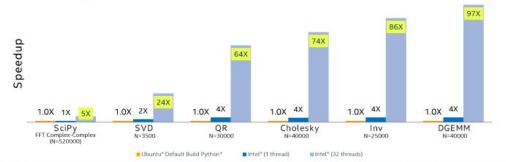


#### Available through Anaconda as well:

conda create -p \$SCRATCH/idp \
 -c intel intelpython2\_core python=2
source activate \$SCRATCH/idp

#### **Features:**

Leveraging Intel MKL, MPI, TBB, DAAL. Intel-specific enhancements (FFT, threaded RNG, etc).



Configuration info: - Versions: Intel<sup>®</sup> Distribution for Python 2.7.11 2017, Beta (Mar 08, 2016), Ubuntu<sup>+</sup> built Python<sup>+</sup>, Python 2.7.11, NumPy 1.10.4, SciPy 0.17.0 built with gcc 4.8.4; Hardware: Intel<sup>®</sup> Xeon<sup>+</sup> (PU E5-2698 us @ 2.30GHz (2 sockets, 16 cores each, HT=0FF), 64 GB of RAM, 8 DIMMS of 8GB@2133MHz; Operating System: Ubuntu 14.04 LTS; MKL version 11.3.2 for Intel Distribution for Python 2017, Beta

Software and workloads used in performance tests may have been optimized for performance only on Intel microprocessors. Performance tests, such as SYSmark and MobileMark, are measured using specific computer systems, components, software, operations and functions. Any change to any of those factors may cause the results to vary. You should consult other information and performance tests to assist you in fully evaluating your contemplated purchases, including the performance of that product when combined with other products. \* Other brands and names are the property of their respective owners. Benchmark Source: Intel Corporation

Optimization Notice: Intel's compilers may or may not optimize to the same degree for non-Intel microprocessors for optimizations that are not unique to Intel microprocessors. These optimizations include SSE2, SSE3, and SSSE3 instruction sets and other optimizations. Intel does not guarantee the availability, functionality, or effectiveness of any optimization on microprocessors not manufactured by Intel. Microprocessor-dependent optimizations in this product are intended for use with Intel microprocessors. Certain optimizations not specific to Intel microarchitecture are reserved for Intel microprocessors. Please refer to the applicable product User and Reference Guides for more information regarding the specific instruction sets covered by this notice. Notice revision #201018064.







### Multi-Node Parallelism: mpi4py

### MPI support via mpi4py (2.0.0) Added earlier this year. Includes MPI-3 features.

Compiled against Cray libraries.

Built into Anaconda modules on Edison and Cori.

Non-Anaconda route: module load mpi4py

DIY mpi4py builders... see me.

#### from mpi4py import MPI

#### # Initialize MPI.

comm = MPI.COMM\_WORLD
mpi\_rank = comm.Get\_rank()
mpi\_size = comm.Get\_size()

# Take command Line arguments.

```
seed = int(sys.argv[1])
size = int(sys.argv[2])
```

# Have root (rank 0) task confirm MPI size.

```
if mpi_rank == 0 :
    start = time.time()
    print "MPI size", mpi_size
    print
```

# Different random number per rank.

numpy.random.seed(seed + mpi\_rank \* 1000)

# Count points in Q1 of unit circle.

```
x = numpy.random.uniform(size=size)
y = numpy.random.uniform(size=size)
r2 = x * x + y * y
in_circle = numpy.sum(r2 < 1.0, dtype=float)</pre>
```

Reduce count to root MPI task (rank  $\theta$ ) by summing them all. MPI wants things in numpy arrays.

```
in_circle = numpy.array([in_circle])
total_in_circle = numpy.zeros(1, dtype=float)
comm.Reduce(in_circle, total_in_circle, op=MPI.SUM)
```

Reduce total number of points tried in the same fashion.
Note that this step is actually unnecessary.

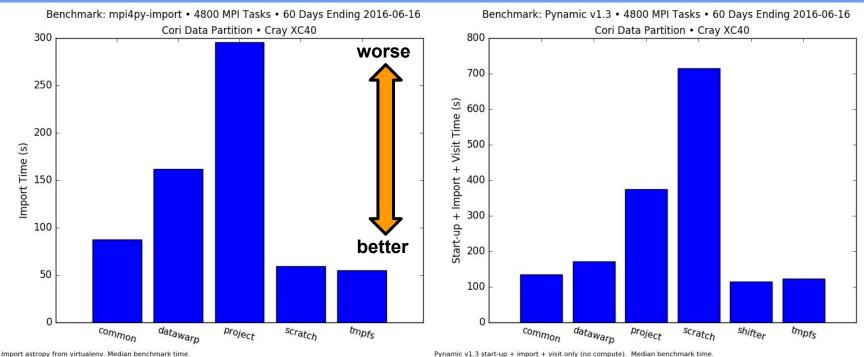
in\_square = numpy\_array([size], dtype=float)
total\_in\_square = numpy\_zeros(1, dtype=float)
comm\_Reduce(in\_square, total\_in\_square, op=MPI\_SUM)







### MPI Start-up in Python Apps at Scale



• Python's "import" statement is file metadata intensive (.py, .pyc, .so open/stat calls).

- Becomes more severe as the number of Python processes trying to access files increases.
- Result: Very slow times to just start Python applications at larger concurrency (MPI).
- BEST POSSIBLE PERFORMANCE IS SHIFTER:
  - Eliminates metadata calls off the compute nodes.
  - Paths to .so libraries can be cached via Idconfig.
- Other approaches:
  - Pack up software to compute nodes (python-mpi-bcast).
  - Install software to \$SCRATCH or /global/common.



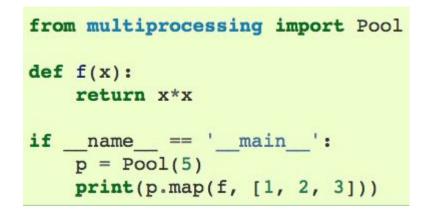




## Multiprocessing and Process Spawning

You can use multiprocessing for on-node throughput jobs.

Combining multiprocessing with mpi4py, mixed results.



### **Combining mpi4py and subprocess?** Works to spawn serial, compiled executables. Just don't compile those with Cray wrappers cc, CC, ftn. Do module load gcc and use gcc, g++, gfortran.







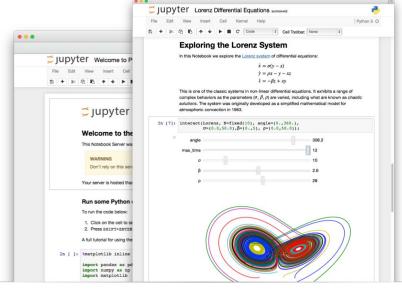
#### Jupyter Notebook: "Literate Computing." Code, text, equations, viz in a narrative.

### New way to interact with NERSC HPC resources:

Old: Use ssh or NX to get to command line. New: Open a notebook, create a narrative.

#### Move to Cori:

- Access to \$SCRATCH.
- Integration with SLURM.
- Eventually Burst Buffer.
- New ways of using Cori.
  - DASK, PySpark, IJulia...







#### Live Demo



DANGER













### **SLURM Magic Commands**



0	Jul	py <sup>.</sup>	te	r a	wes	som	e La	st Ch	neckp	oint: 03/25/	2016 (auto	osave	d)					Control	Panel	Logout
F	le	Edit	١	/iew	Ins	sert	Ce	ell	Ker	nel Hel	ip						Notebook s	aved 🥒	Pytl	non 2.7 O
B	+	»	ළු	ß	•	Ψ	H		C	Code	\$		CellToolbar	4	Ũ	0				

In [1]:	squeue -u rthomas													
Out[1]:		JOBID	USER	ACCOUNT	NAME	PARTITION	QOS	NODES	TIME_LIMIT	TIME	ST	PRIORITY	SUBMIT_TIME	START_TIME
	0	2875563	rthomas	mpccc	try.sh	regular	normal_reg	150	1:00	0:00	PD	64832	2016-08-19T15:33:09	NaN

#### In [2]: %sprio -j 2875563

Out[2]:		JOBID	PRIORITY	AGE	FAIRSHARE	QOS
	0	2875563	64832	32	0	64800

#### In [3]: %sacct -u rthomas

0		A 1	r つ '	۱.
U	u	L	10	
		1.1.1		

	JobID	JobName	Partition	Account	AllocCPUS	State	ExitCode
0							
1	2873623	try.sh	debug	mpccc	64	FAILED	1:0
2	2873623.bat+	batch	mpccc	64	FAILED	1:0	NaN
3	2873623.ext+	extern	mpccc	64	COMPLETED	0:0	NaN
4	2873623.0	shifter	mpccc	32	FAILED	1:0	NaN
5	2873630	try.sh	debug	mpccc	64	FAILED	1:0
6	2873630.bat+	batch	mpccc	64	FAILED	1:0	NaN







#### Knights Landing (KNL) to: 2x cores per node independent program Slower clock rate units; explicit Increase Thread Parallelism Less memory/core. independent execution units within the program; generally explicit Exploit Data Parallelism Single-thread or flat MPI - Same operation on multiple elements Python won't be great.

### Advice:

ENERGY Office of Science - 7 -Leverage threaded, vectorized math/specialized libraries. Consider writing Cython/C extensions you can vectorize? Learn about Intel Python and Intel profiling tools.

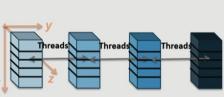




To run effectively on Cori users will have

Manage Domain Parallelism

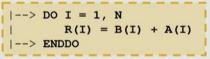
- Improve data locality
  - Cache blocking; Use on-package memory



MPI

MPI

MPI





Python is an integral element of NERSC's Data Intensive Science portfolio.

We want users to have a:familiarPython environmentproductivePython experienceperformantPython software stack

Pursuing new ways to empower Python & data users.

Always looking for feedback, advice, and even help: <u>consult@nersc.gov</u> or https://help.nersc.gov rcthomas@lbl.gov







## National Energy Research Scientific Computing Center



