



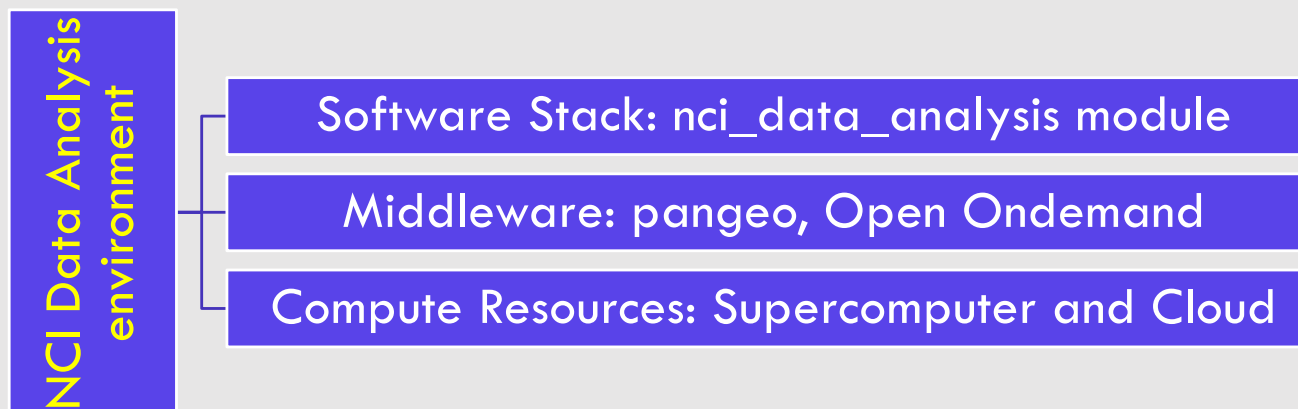
# NCI'S JUPYTER-PANGEO ENVIRONMENT FOR DATA ANALYSIS

*DATA SCIENCE WEEK 2021*

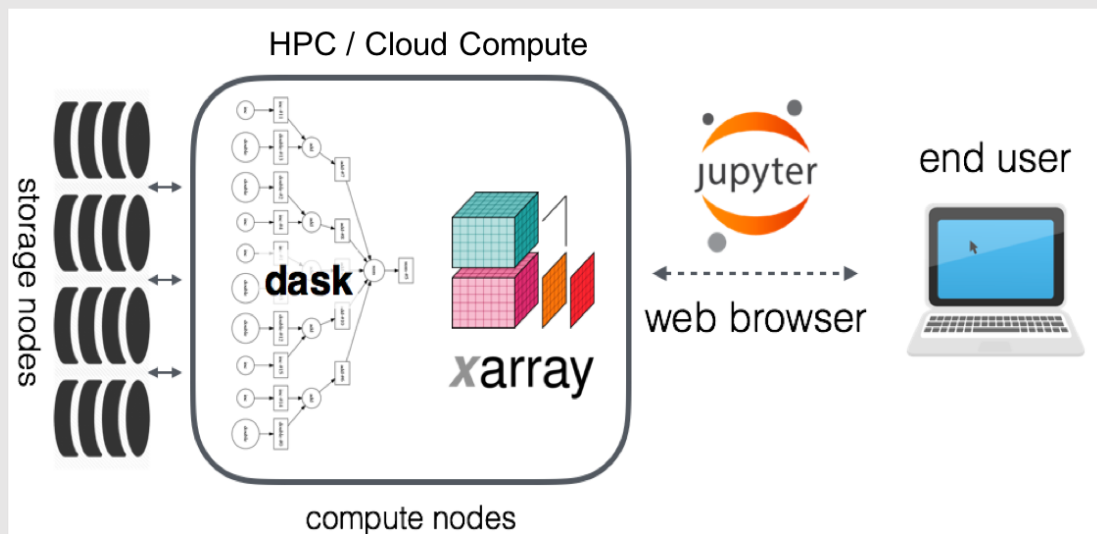


# ANALYSING NCI DATA COLLECTIONS

- The NCI Reference Data Collections are organised in a systematic way to enable fast programmatic access for analysis across multiple domains.
- NCI data collections are available for use on NCI's core computing resources like NCI supercomputer and NCI cloud-based platforms.
- NCI established the data analysis environment to help users accessing the data collections in a scalable programming way.



# TYPICAL NCI DATA ANALYSIS WORKFLOW



<https://pangeo.io>

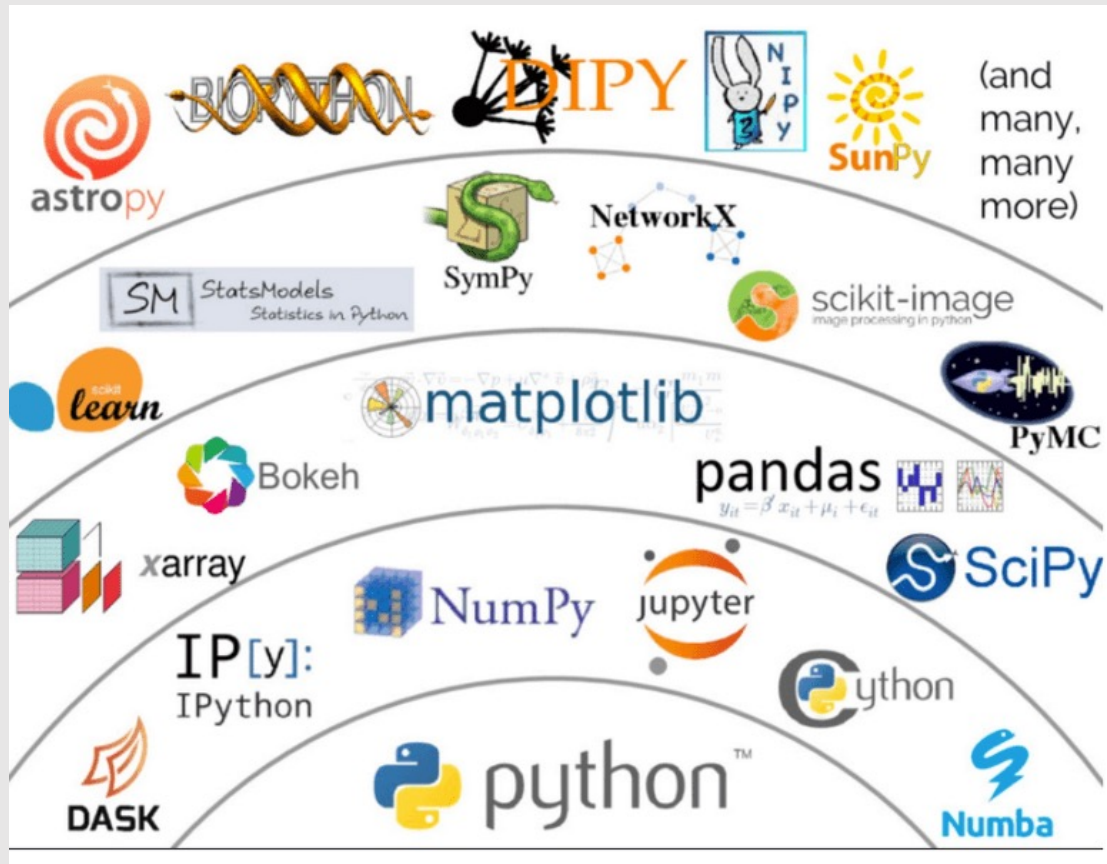
Remote Jupyter job is executing at the NCI scalable resources.

The job utilizes Dask as the parallel computing engine.

Users work with their own desktop web browser to access the remote Jupyter server.

<https://opus.nci.org.au/display/Help/Data+Analysis+Environments>

# SOFTWARE STACK FOR NCI DATA ANALYSIS

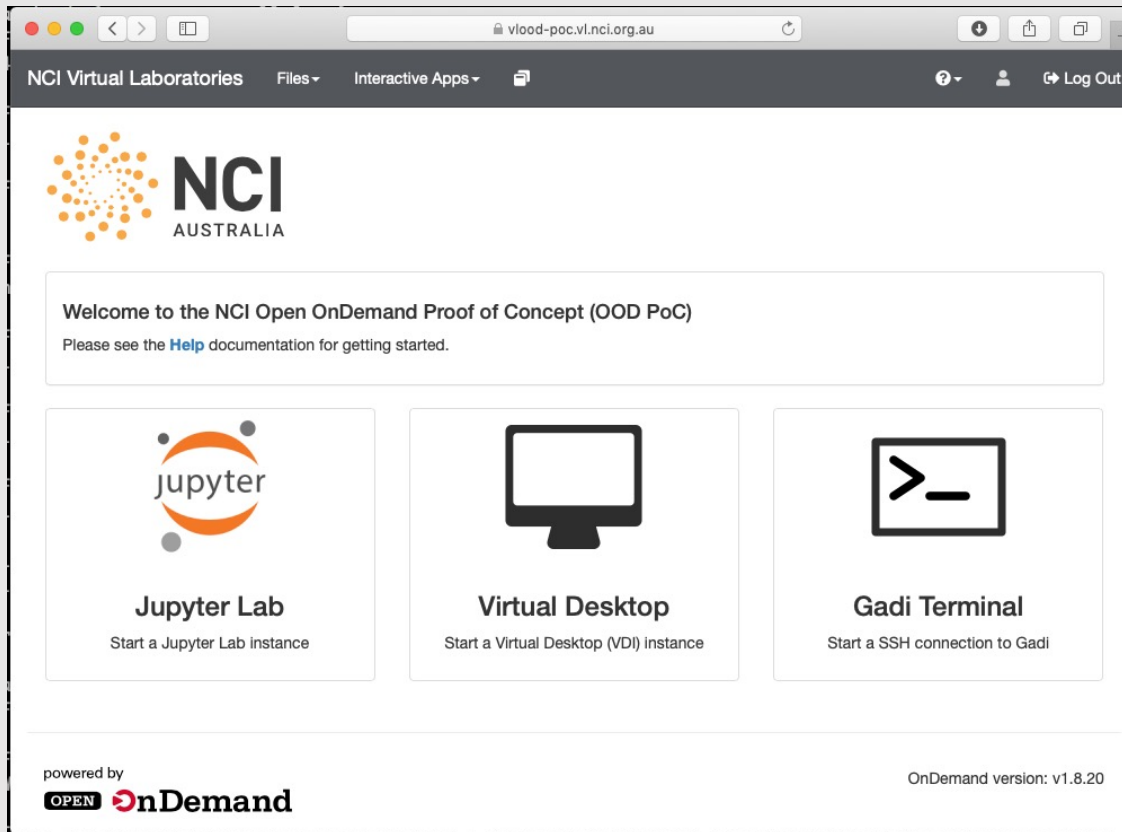


- nci-data-analysis/2021.03 module
- 485 general-purpose python libraries including Jupyter, xarray and Dask etc.
  - periodically updating (~3 months).
  - adding more libs per user's request.

# COMPUTE RESOURCES FOR NCI DATA ANALYSIS

	<b>GADI</b>	<b>VDI</b>	<b>JupyterLab</b>
<b>Description</b>	Australia's peak research supercomputer with 4000+ compute nodes including 640 NVIDIA V100 GPUs	Graphical desktop-like interface based on the NCI cloud resource	Jupyter native interface which supports scalable dask cluster jobs
<b>Hardware/Node</b>	<b>48 CascadeLake CPU cores, 192 GB memory (normal queue)</b>	16 vCPUs (SandyBridge), 32GB memory	16 vCPUs (SandyBridge), 32GB memory
<b>Job resources</b>	<b>Multiple nodes</b>	Single node	<b>Multiple nodes</b>
<b>File System</b>	<b>Lustre</b>	NFS access to Lustre	NFS access to Lustre
<b>Internet Connection</b>	No	<b>Yes</b>	<b>Yes</b>
<b>Web Browser</b>	No	<b>Yes</b>	<b>Yes</b>
<b>Typical job</b>	<b>compute intensive work</b>	<b>data analysis</b> code development <b>visualisation</b>	<b>data analysis</b>

# OPEN ONDEMAND (OOD) PLATFORM



<https://vlood-poc.vl.nci.org.au>

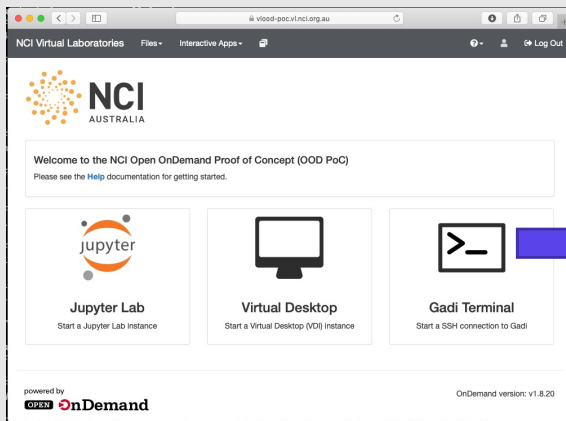
OOD is an online portal giving you access to compute resources at NCI such as VDI, Gadi, or Jupyter Lab. It provides an app based infrastructure which we will extend as new new apps are developed.

3 apps on the OOD:

- [Gadi Terminal](#): simple browser based access to Gadi; the session closes when you close your browser window (or change to another page). Useful for checking Gadi jobs from a web-browser etc.
- [VDI](#): Web-browser based VNC connection to a VDI session (i.e. alternative to Strudel).
- [Jupyter Lab](#): dedicated Jupyter Lab session which can be used to run Dask workload among others.

Currently at the stage of “Proof of Concept”.

# ACCESS GADI VIA OOD



```
#####
# Welcome to the NCI National Facility! #
# This service is for authorised clients only. It is a criminal #
# offence to: #
# - Obtain access to data without permission #
# - Damage, delete, alter or insert data without permission #
# Use of this system requires acceptance of the Conditions of Use #
# published at http://nci.org.au/users/nci-terms-and-conditions-access #
#####
| gadi.nci.org.au - 185,032 processor InfiniBand x86_64 cluster |
#####

Mar 26 2021 New Cascade Lake Megamem Now Available
We have added 4 new Cascade Lake megamem (3TB) nodes to Gadi. These nodes
serve the "megamem" queue. Note that specialised queues should only be used
by jobs that require the specialised resources of that queue.

For more information about queues see: https://opus.nci.org.au/x/2AB1BQ

Last login: Tue May 4 12:25:27 2021 from 130.56.246.173
[rxxy900@gadi-login-05 ~]$ cd /scratch/fp0/rxy900/PROJECT/data_science_week/
[rxxy900@gadi-login-05 data_science_week]$ qsub sub_batch_job.dk92.sh
21928232.gadi-pbs
[rxxy900@gadi-login-05 data_science_week]$ qstat -u rxy900

gadi-pbs:
Job ID Username Queue Jobname SessID NDS TSK Memory Req'd Req'd Elap
Time S Time
-----
21928232.gadi-pbs rxy900 normal-* pangeo_job 14188* 2 96 384gb 01:00 R 00:00
[rxxy900@gadi-login-05 data_science_week]$ cat client_cmd
ssh -N -L 8319:gadi-cpu-clx-0956.gadi.nci.org.au:8319 rxy900@gadi.nci.org.au
ssh -N -L 8723:gadi-cpu-clx-0956.gadi.nci.org.au:8723 rxy900@gadi.nci.org.au
[rxxy900@gadi-login-05 data_science_week]$
```

Code cell content:

```
[1]: from dask.distributed import Client, LocalCluster
import time

[2]: client = Client(scheduler_file='./scheduler.json')

[3]: client

[3]: Client Cluster
Scheduler: tcp://10.6.46.20:8702 Workers: 96
Dashboard: http://10.6.46.20:8723/status Cores: 96
Memory: 404.96 GB

[4]: import xarray as xr
import dask.array as da
import netCDF4 as nc
import hvplot.xarray
```

Dashboard content:

Status Workers Tasks System Profile Graph Info

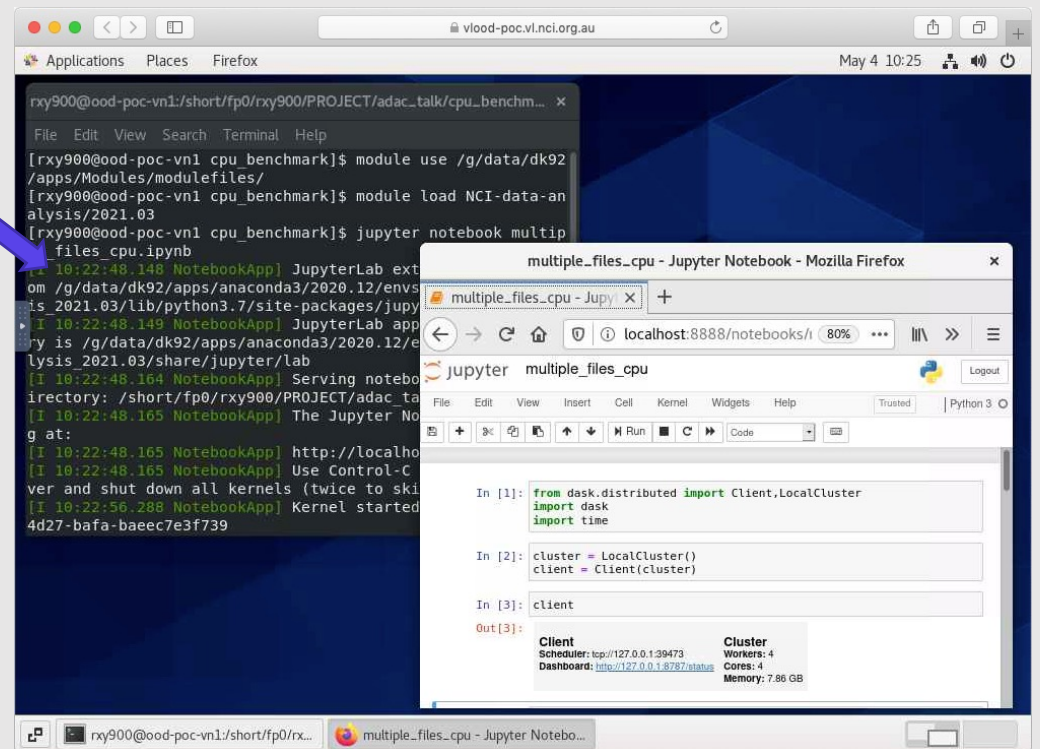
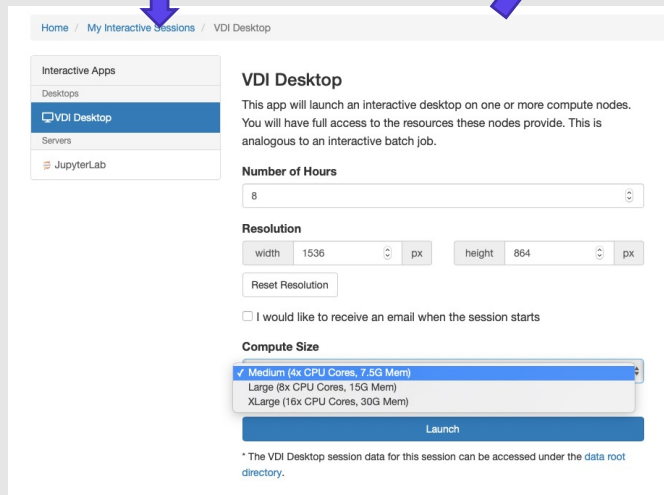
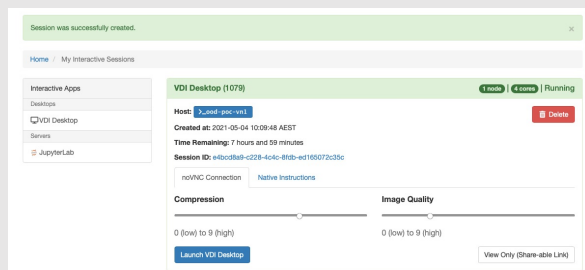
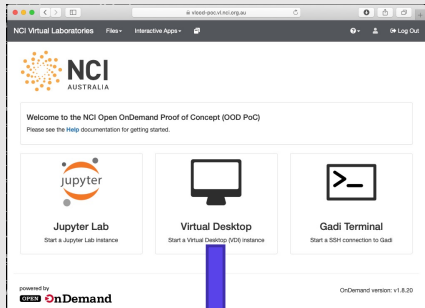
Task Stream

Progress -- total: 0, in-memory: 0, processing: 0, waiting: 0, error: 0

Pangeo Manual

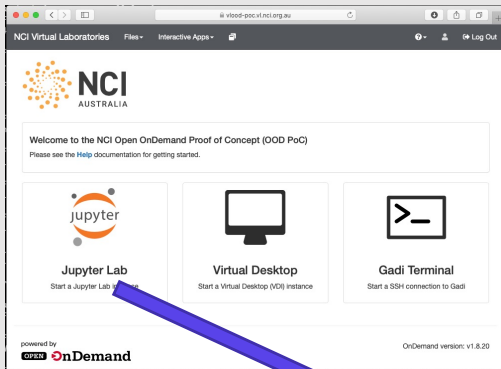
<https://opus.nci.org.au/display/Help/5.+Pangeo+on+Gadi>

# ACCESS VDI VIA OOD





# ACCESS JUPYTERLAB VIA OOD



**JupyterLab** version: 635d2c5  
This app will launch a dedicated Jupyter Lab Instance

**Number of Hours**

- Small (1x CPU Core, 1.875G Mem)
- Medium (2x CPU Cores, 3.75G Mem)
- Large (4x CPU Cores, 7.5G Mem)
- XLarge (8x CPU Cores, 15G Mem)
- XXLarge (16x CPU Cores, 30G Mem)

The size of resources available within your Jupyter Lab session. NOTE: select small if you are using Dask to offload processing to Gadi/Cloud

I would like to receive an email when the session starts

Advanced options ...

**Module directories**

module use /g/data/dk92/apps/Modules/modulefiles

Include module directories i.e. module use DIR ...

**Modules**

module load NCI-data-analysis/2021.03

Includes modules i.e. module load MOD ...

**Conda environment**

Activates a conda environment i.e. conda activate ENVIRONMENT

**Python virtual environment**

Activates a Python virtual environment i.e. source ENVIRONMENT/bin/activate

**Launch**

\* The JupyterLab session data for this session can be accessed under the data root directory.

**Jupyter Lab: Cloud (2648)** 1 node | 16 cores | Running

Host: >\_rei-n1 Delete

Created at: 2021-04-20 15:43:42 AEST

Time Remaining: 3 hours and 58 minutes

Session ID: 551d5eba-9484-4ae1-acad-d810e851a255

[Open Jupyter Lab](#) [Open Notebook from Catalogue](#)

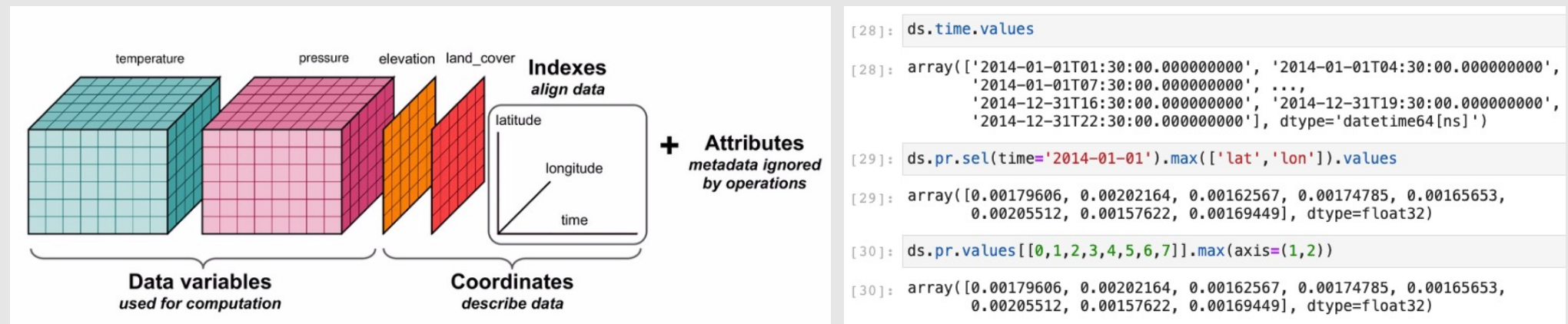
```
[5]: from dask.distributed import Client, Scheduler
      from dask_jobqueue import SLURMCluster
      cluster = SLURMCluster(cores=4, memory="29GB")
      client = Client(cluster)
      cluster.scale(cores=16)

[7]: client

[7]: Client                               Cluster
      Scheduler: tcp://10.0.16.1:34565    Workers: 16
      Dashboard: http://10.0.16.1:8787/status  Cores: 16
                                              Memory: 116.00 GB
```

# AN EXAMPLE OF SCALABLE ANALYSIS: XARRAY AND LABELS

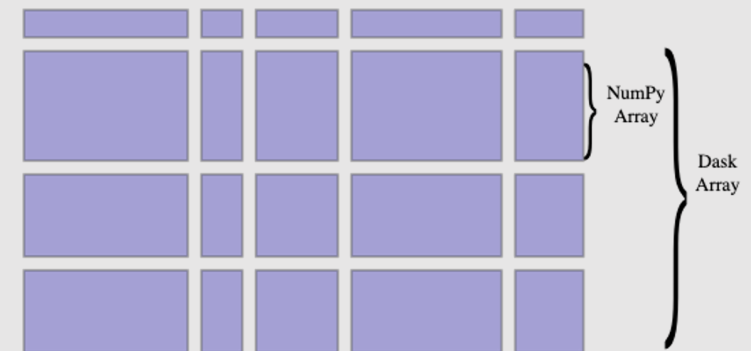
- NumPy provides the fundamental data structure and API for working with raw ndarrays.
- Xarray uses metadata in the form of labelled dimensions (e.g., 'latitude' or 'frequency') and coordinate values (e.g., the date '2021-02-05') to enable a suite of expressive, label based operations.
- Can apply operations over dimensions by name: `x.sum('time')`
- Can select values by label instead of integer location: `x.loc['2021-02-05']` or `x.sel(time='2021-02-05')`
- Mathematical operations (e.g. `x - y`) vectorize across multiple dimensions based on dimension names, not shape
- split-apply-combine paradigm with `groupby`, Database-like alignment based on coordinate labels, keep track of arbitrary metadata in the form of a Python dictionary: `x.attrs`, etc.



# AN EXAMPLE OF SCALABLE ANALYSIS: XARRAY - PARALLEL COMPUTING WITH DASK

Xarray integrates with Dask to support parallel computations and streaming computation on datasets that don't fit into memory

- Dask divides arrays into many small pieces (chunks), each of which is presumed to be small enough to fit into memory so it can provide multi-core and distributed parallel execution on larger-than-memory datasets
- Dask scales up (to a cluster) and down (to a single machine).
- High level collections like Array, Bag, and DataFrame that mimic NumPy, lists, and Pandas but can operate in parallel on datasets that don't fit into memory.
- Low Level schedulers with low-latency to execute task graphs in parallel.







# AN EXAMPLE OF SCALABLE ANALYSIS: BENCHMARK ON A CMIP6 DATA COLLECTION

```
import xarray as xr
ds = xr.open_mfdataset(allfiles, chunks={"time":753}, combine='by_coords', parallel=True)
len(allfiles)=166 # total number of source NetCDF files.
```

xarray.Dataset

► Dimensions: (bnds: 2, lat: 192, lon: 384, time: 482120)

▼ Coordinates:

<b>time</b>	(time)	datetime64[ns]	1850-01-01T01:30:00 ... 2014-12-...	 
<b>lat</b>	(lat)	float64	-89.28 -88.36 ... 88.36 89.28	 
<b>lon</b>	(lon)	float64	0.0 0.9375 1.875 ... 358.1 359.1	 

▼ Data variables:

<b>time_bnds</b>	(time, bnds)	datetime64[ns]	dask.array<chunksize=(753, 2), meta=np...	 
<b>lat_bnds</b>	(time, lat, bnds)	float64	dask.array<chunksize=(2920, 192, 2), me...	 
<b>lon_bnds</b>	(time, lon, bnds)	float64	dask.array<chunksize=(2920, 384, 2), me...	 
<b>pr</b>	(time, lat, lon)	float32	dask.array<chunksize=(753, 192, 384), m...	 

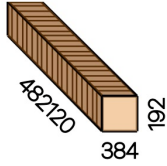
► Attributes: (44)

A model ran by the Alfred Wegener Institute, Helmholtz Centre for Polar and Marine Research, Am Handelshafen 12, 27570 Bremerhaven, Germany (AWI) in native nominal resolutions: atmos: 100 km, land: 100 km, ocean: 25 km, sea ice: 25 km.

# AN EXAMPLE OF SCALABLE ANALYSIS: PRECIPITATION FLUX VARIABLE

xarray.DataArray 'pr' (time: 482120, lat: 192, lon: 384)

	Array	Chunk
Bytes	132.42 GiB	211.78 MiB
Shape	(482120, 192, 384)	(753, 192, 384)
Count	1485 Tasks	660 Chunks
Type	float32	numpy.ndarray



▼ Coordinates:

name	units	dtype	values	actions
time	(time)	datetime64[ns]	1850-01-01T01:30:00 ... 2014-12-...	📄 🗑️
lat	(lat)	float64	-89.28 -88.36 ... 88.36 89.28	📄 🗑️
lon	(lon)	float64	0.0 0.9375 1.875 ... 358.1 359.1	📄 🗑️

▼ Attributes:

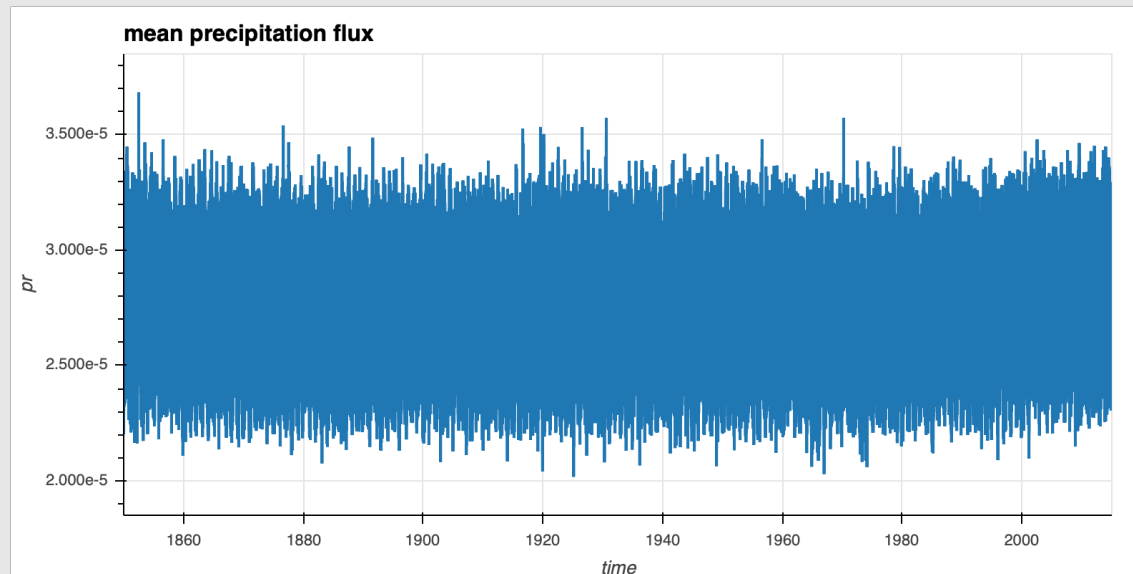
standard\_name : precipitation\_flux  
long\_name : Precipitation  
comment : includes both liquid and solid phases  
units : kg m-2 s-1  
original\_name : pr  
cell\_methods : area: time: mean  
cell\_measures : area: areacella  
history : 2019-05-02T11:50:26Z altered by CMOR: replaced missing value flag (-9e+33) with standard missing value (1e+20). 2019-05-02T11:50:26Z altered by CMOR: Inverted axis: lat.

```
ds.pr.nbytes/1e9=142.1829734#size (GB)
```

The precipitation flux variable has three dimensions. It is a dask.array concatenated over all 166 files in this directory with the total size of 132GiB. The precipitation flux variable is recorded every three hours according to the time stamps above. It's 660 chunks could be processed in parallel over the DASK cluster.

# AN EXAMPLE OF SCALABLE ANALYSIS: TIMESERIES OF MEAN PRECIPITATION FLUX

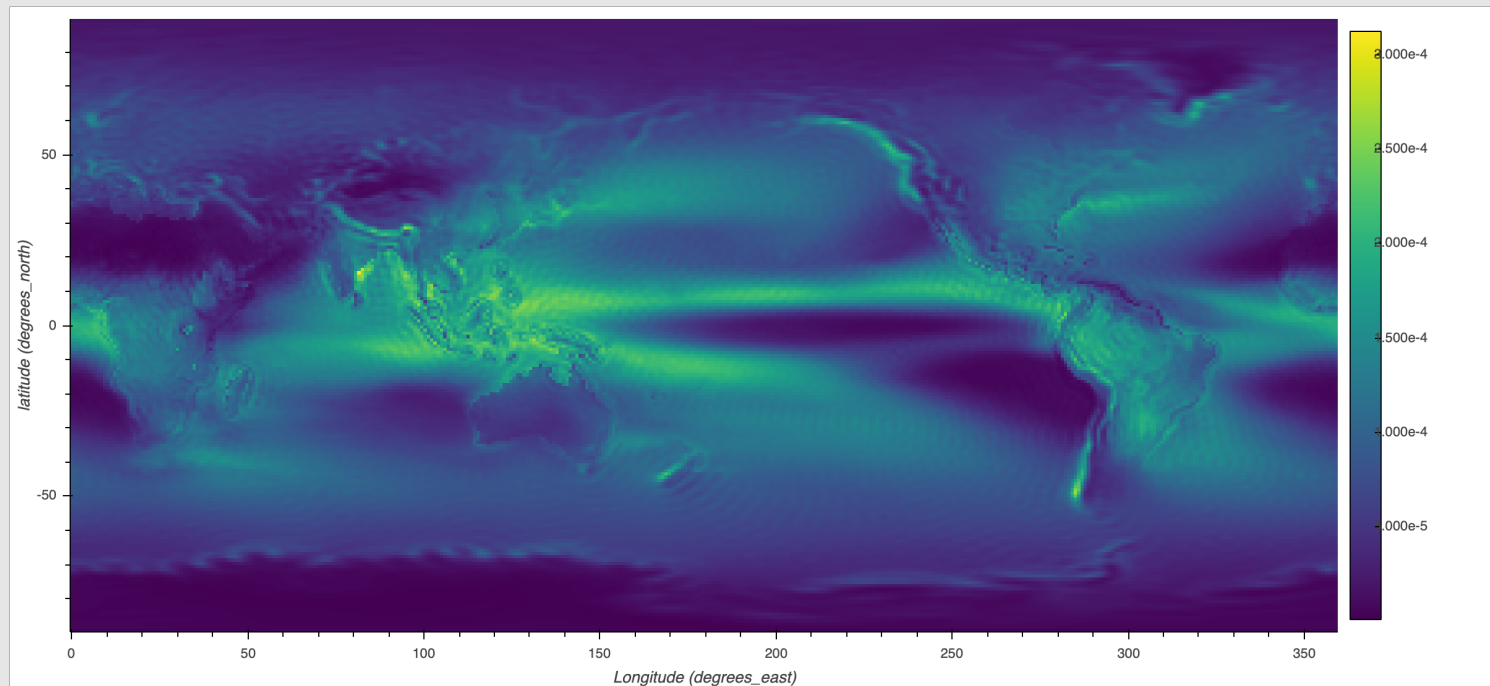
```
pr_mean = ds.pr.mean(dim=('lat','lon'))  
pr_full = pr_mean.hvplot(label='full time series dataset from 1850 to  
2014', grid=True, title='mean precipitation flux', width=800, height=400)
```



# AN EXAMPLE OF SCALABLE ANALYSIS: PRECIPITATION FLUX VARIABILITY

```
pr_std = ds.sel(time=slice("1900-01-01", "2000-12-31")).pr.std(dim='time')  
pr_std.hvplot(colormap='viridis', width=1200, height=550, rasterize=True)
```

We can examine the natural variability in precipitation flux by looking at its standard deviation over time.



## AN EXAMPLE OF SCALABLE ANALYSIS: CODE BLOCK TO ENABLE BENCHMARK

```
from dask.distributed import Client
client = Client(scheduler_file='./scheduler.json')

ds = xr.open_mfdataset(allfiles, chunks={"time":753},
combine='by_coords',parallel=True)

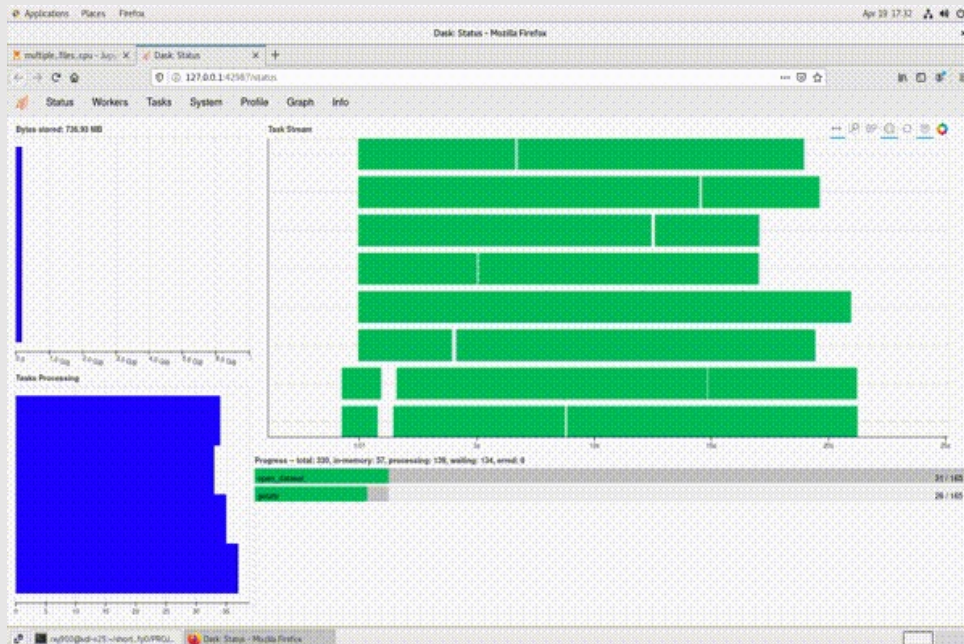
subds=ds.isel(time=slice(slice_beg,slice_end))

pr_mean=subds.pr.mean(dim=('lat','lon'))
pr_mean.compute()

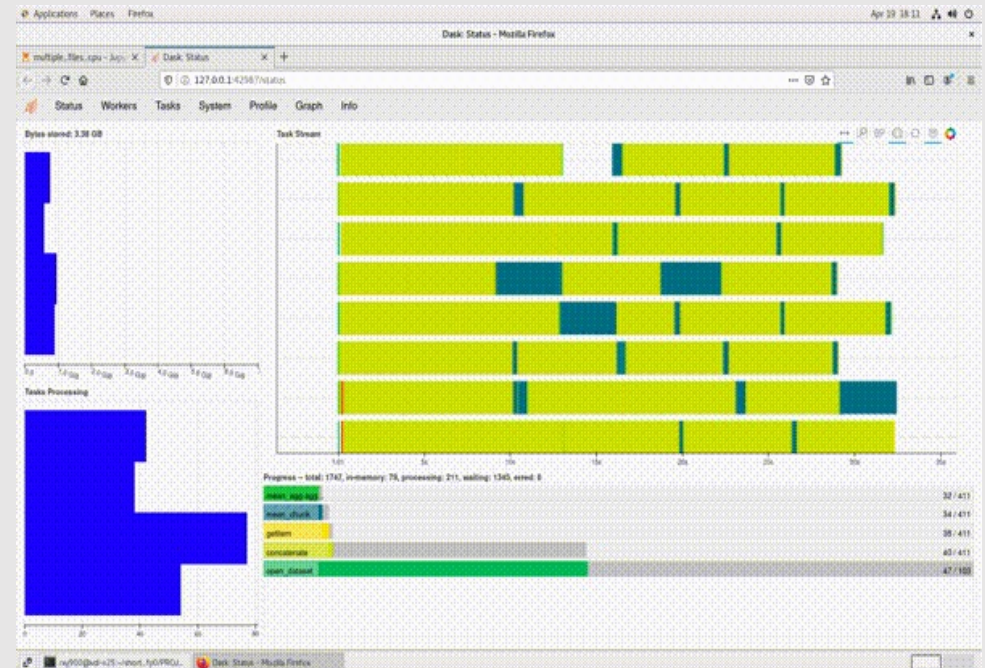
pr_std = subds.pr.std(dim='time')
pr_std.compute()
```



# AN EXAMPLE OF SCALABLE ANALYSIS: PARALLEL TASKS WITH THE DASK CLUSTER



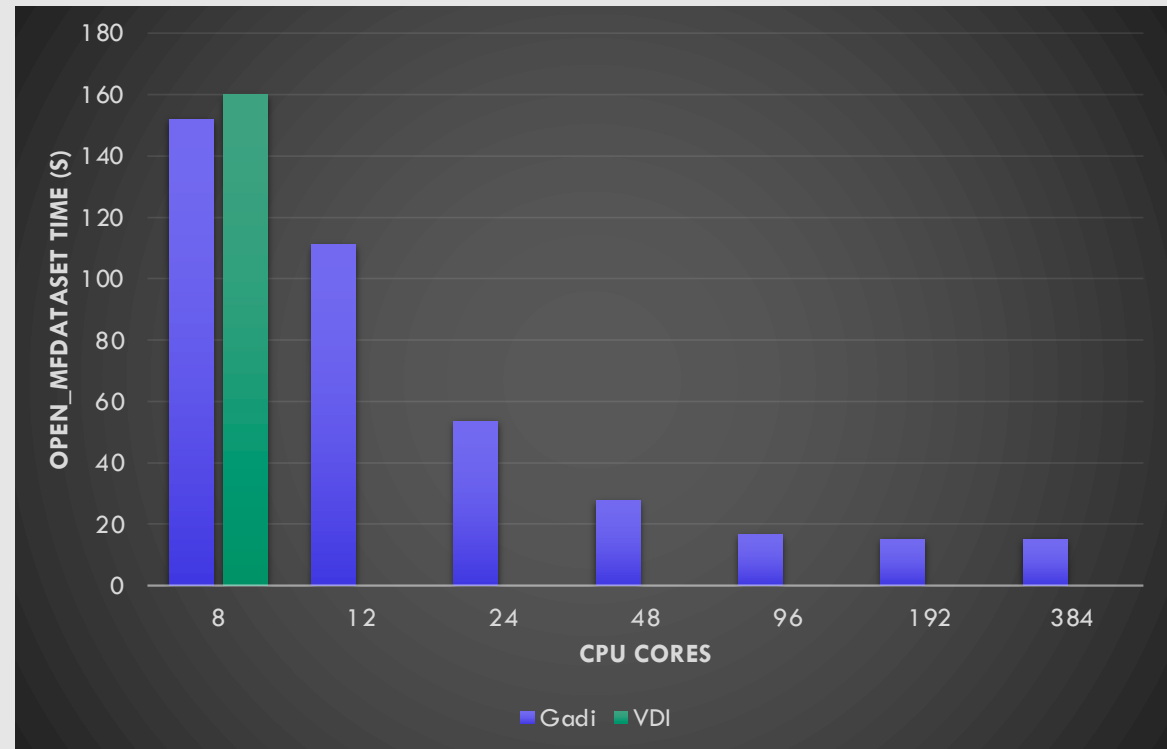
`xr.open_mfdataset()`



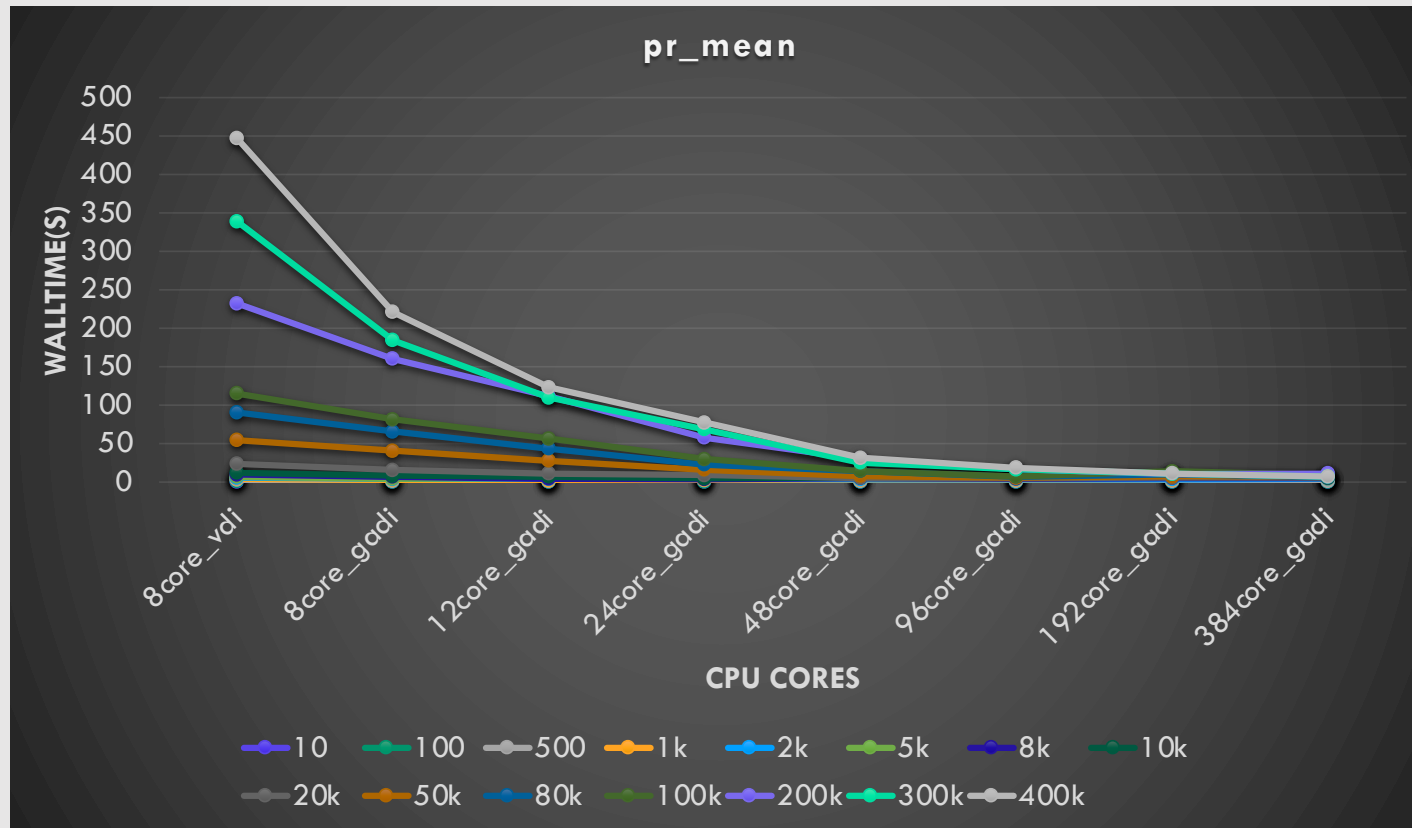
`pr_mean.compute()`

# AN EXAMPLE OF SCALABLE ANALYSIS: PERFORMANCE RESULTS OF OPEN\_MFDATASET

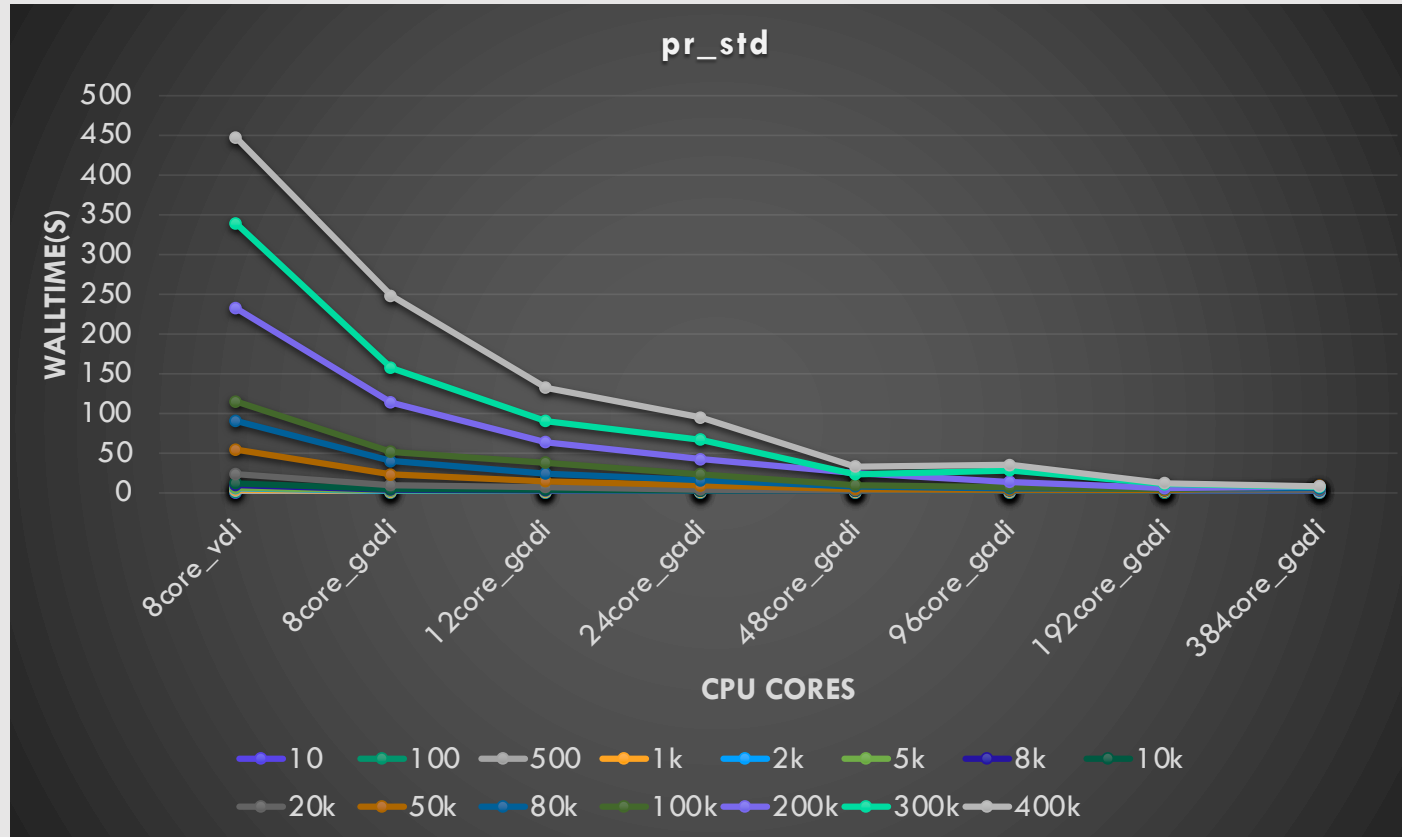
- 2 Threads per Dask worker
- $N_{\text{threads}} = N_{\text{cores}}$
- CPU resources:
  - VDI: 8 cores in a single node.
  - Gadi: 48 cores/node.



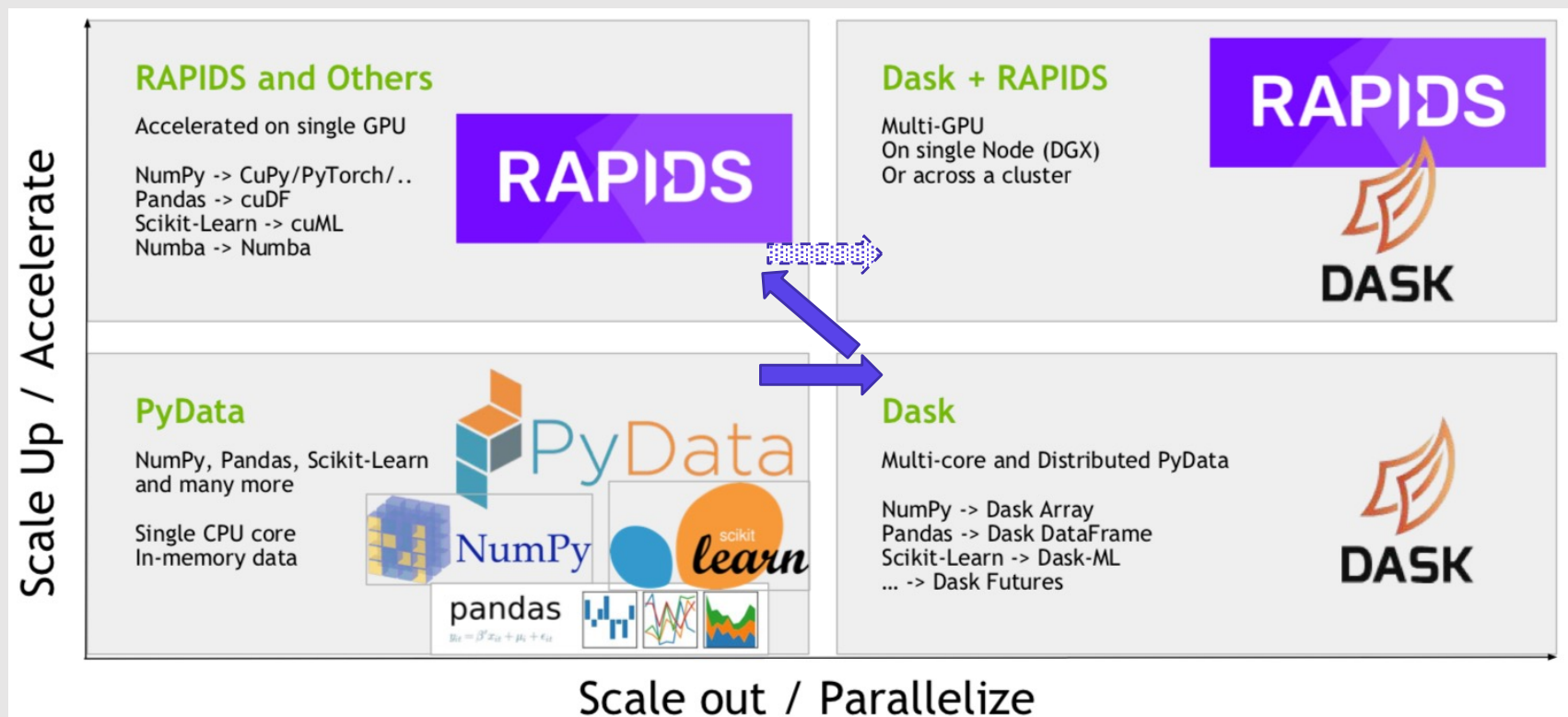
# AN EXAMPLE OF SCALABLE ANALYSIS: PR\_MEAN SCALABILITY ON THE SIZE OF TIME SLICES AND CPU



# AN EXAMPLE OF SCALABLE ANALYSIS: PR\_STD SCALABILITY ON THE SIZE OF TIME SLICES AND CPU



# GPU SUPPORT IN DEVELOPMENT



<https://developer.download.nvidia.com/video/gputechconf/gtc/2019/presentation/s9797-dask-extensions-and-new-developments-with-rapids.pdf>

# AN EXAMPLE OF SCALABLE ANALYSIS: SINGLE GPU UTILIZATION

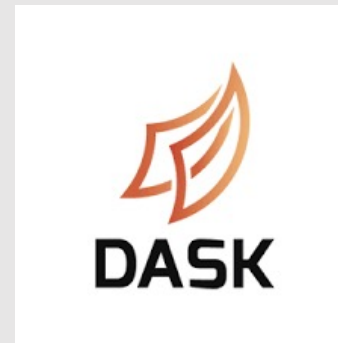
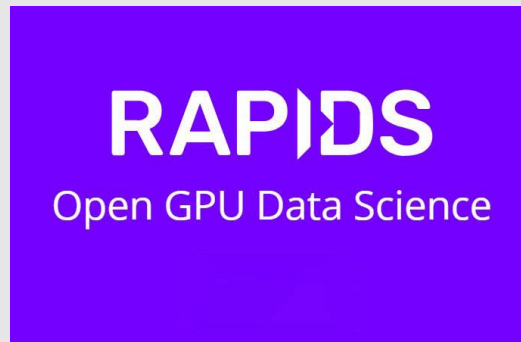
```
■ Import cupy as cp
■ from dask.distributed import Client
■ client = Client(scheduler_file='./scheduler.json')
■ ds = xr.open_mfdataset(allfiles,
  chunks={"time":753},combine='by_coords',parallel=True)
■ subds=ds.isel(time=slice(slice_beg,slice_end))
■ # Additional step to convert data type from numpy to cupy.
  subcds.pr.data = cp.asarray(subcds.pr.data) # extra convert time cvttime.
■ pr_mean=subds.pr.mean(dim=('lat', 'lon'))
■ pr_mean.compute() # meantime
■ pr_std = subds.pr.std(dim='time')
■ pr_std.compute() #stdtime
```

# AN EXAMPLE OF SCALABLE ANALYSIS: GPU BENCHMARK RESULTS

Walltime(s)	12 CPUs			1 GPU		1 GPU		
size	meantime	stdtime		meantime	stdtime	cvtime	N <sub>pr_mean</sub>	N <sub>pr_std</sub>
10	0.14	0.08		1.63	0.15	<b>0.10</b>	1	1
100	0.33	0.34		0.00	0.00	<b>0.29</b>	1	1
500	1.31	1.29		0.00	0.01	<b>2.28</b>	2	2
1000	2.79	2.17		0.00	0.04	<b>2.52</b>	1	1
2000	2.63	2.42		0.00	0.08	<b>3.70</b>	1	2
5000	2.77	2.38		0.00	0.21	<b>8.59</b>	3	4
8000	2.98	2.20		0.31	0.03	<b>9.35</b>	3	4
10000	4.14	4.39		0.00	0.40	<b>11.14</b>	3	3
20000	6.28	6.15		0.00	0.81	<b>23.47</b>	4	4
50000	14.88	12.02		0.00	2.02	<b>51.44</b>	3	4
80000	23.95	19.54		0.00	3.22	<b>102.01</b>	4	5
100000	30.44	29.22		1.17	3.65	<b>142.17</b>	5	5

Single GPU can accelerate the data analysis workflow with heavy compute operations.

# PATHWAY TOWARDS MULTIPLE GPUS



- Set up Dask LocalCUDACluster via dask-cuda library to distribute tasks over multiple GPUs.
- No direct way to read NetCDF files to xarray.Dataset crossing multiple GPUs.
- Convert xarray.Dataset to Dask-cuDF ( can read csv, json, orc, parquet directly).
- New implementations & developments in DASK+RAPIDS are on the way.



## SUMMARY

NCI set up a data analysis environment by integrating both python software stack and NCI compute resources. It could help users to efficiently process large scale datasets in a scalable way with powerful NCI compute resources.

NCI Gadi User Guide

<https://opus.nci.org.au/display/Help/Gadi+User+Guide>

NCI VDI User Guide

<https://opus.nci.org.au/display/Help/VDI+User+Guide>

NCI Data Analysis Environments

<https://opus.nci.org.au/display/Help/Data+Analysis+Environments>

Xarray tutorial

<https://github.com/NCI-data-analysis-platform/examples-dask.git>

Dask tutorial

<https://github.com/NCI-data-analysis-platform/examples-xarray.git>

*Please connect various NCI projects to access the data sources for above tutorials.*

Thank you.