# Dask for Parallel Processing Adrian Jackson, EPCC, The University of Edinburgh a.jackson@epcc.ed.ac.uk archer2 eocc

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#### **Partners**









Natural Environment Research Council

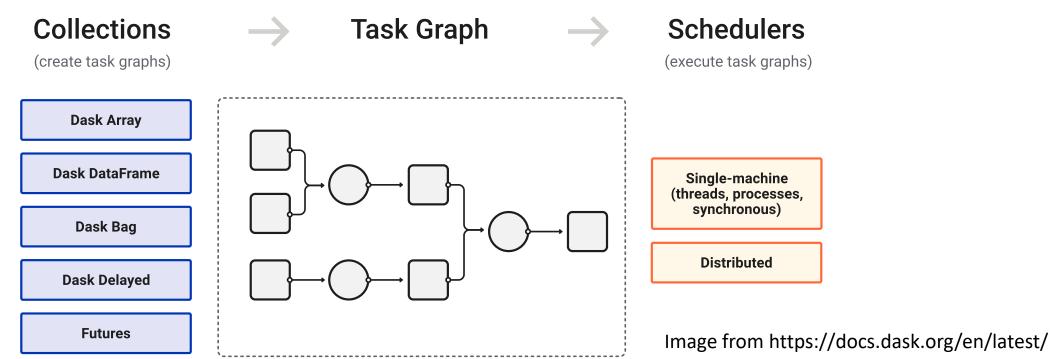




### Dask



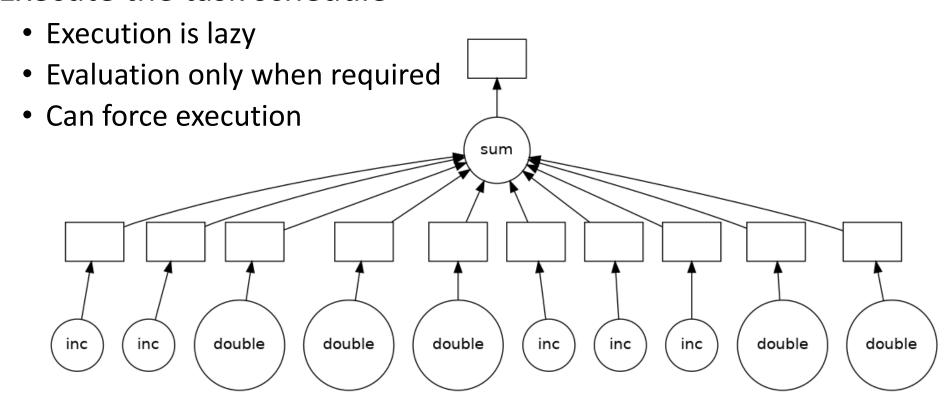
- Dask is a library to do parallel computing in Python
- Two main components
  - Data collections/types
  - Task scheduling functionality with parallel backends



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# Parallel computing

- Split work into tasks to be undertaken
- Schedule those tasks on available compute resources
- Execute the task schedule



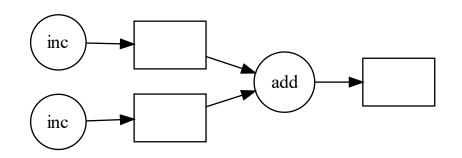
# Collections/data types



- Array
  - n-dimensional array (similar to numpy arrays)
  - Out-of-memory functionality
- DataFrame
  - pandas DataFrame functionality
  - 2-d table/spreadsheet
- Bag
  - Unstructured data types
  - Python iterators
- Dask functionality allows all of these to be distributed across compute nodes but still processed as if local

# Direct parallelisation

- dask.delayed interface
  - Construct custom parallelization
  - Dask task graph functions
- dask Futures functionality
  - Enable immediate task generation
  - Sidesteps delayed evaluation
- Data movement functions
  - gather, scatter, or realise data from futures
- Coordination/synchronisation functionality
  - Queues, Variables, Locks, etc...



# Dask Array

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- Create array with chunk size
  - import dask.array as da
  - x = da.random.random((10000, 10000), chunks=(1000, 1000))
- Many ways to create arrays
  - Random

```
random.binomial(n, p[, size, chunks])
random.normal([loc, scale, size, chunks])
random.poisson([lam, size, chunks])
random.random([size, chunks])
```

- <a href="https://docs.dask.org/en/stable/array-api.html#random">https://docs.dask.org/en/stable/array-api.html#random</a>
- numpy arrays

```
import numpy as np
import dask.array as da
np_array = np.ones((10000,10000))
x = da.from_array(np_array, chunks=(1000, 1000))
```

- Input must have a .shape, .ndim, .dtype and support numpy-style slicing
  - https://docs.dask.org/en/stable/generated/dask.array.from\_array.htm l#dask.array.from\_array

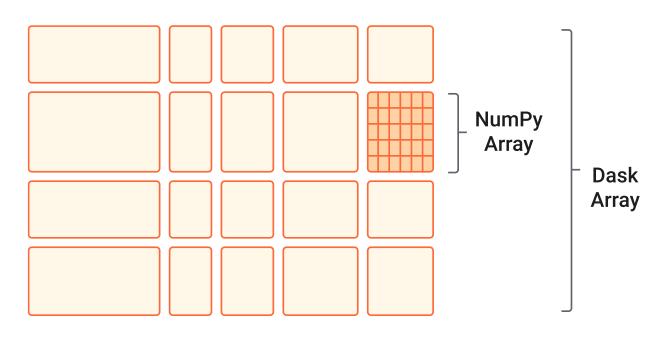


Image from https://docs.dask.org/en/stable/array.html/

# Dask Array



- From files
  - .npy are numpy binary files
  - .zarr are binary files designed for blocked/chunked and compressed data
  - Load groups of files into a single array
  - dask.array.from\_npy\_stack(dirname, mmap\_mode='r')
  - dask.array.from\_zarr(url, component=None, storage\_options=None, chunks=None, name=None, inline\_array=False, \*\*kwargs)
- From dask arrays
  - concatenate: create single dimension array from existing arrays
  - **stack**: create new dimension of data with existing arrays
- delayed
  - From delayed functions that return things that dask arrays can be constructed from
- Remember, tasking means everything is lazy

# **Array**



- Dask supports a range of numpy like functionality:
  - Arithmetic and scalar mathematics: +, \*, exp, log, ...
  - Reductions along axes: sum(), mean(), std(), sum(axis=0), ...
  - Tensor contractions / dot products / matrix multiply: tensordot
  - Axis reordering / transpose: transpose
  - Slicing: x[:100, 500:100:-2]
  - Indexing along single axes with lists or NumPy arrays: x[:, [10, 1, 5]]
  - Array protocols like \_\_array\_\_ and \_\_array\_ufunc\_\_
  - Some linear algebra: svd, qr, solve, solve\_triangular, lstsq
- Dask Array lacks the following features:
  - Much of np.linalg has not been implemented
  - Arrays with unknown shapes do not support all operations
  - Sorts are not fully supported
  - tolist
  - Iterators can be inefficent
- Full API is at: <a href="https://docs.dask.org/en/stable/array-api.html">https://docs.dask.org/en/stable/array-api.html</a>

### Dask DataFrame

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Create like a standard pandas dataframe

```
import dask.dataframe as dd
df = dd.read_csv('mydata.csv')
```

Can also specify blocksize (chunks)

dask.dataframe.read\_csv(urlpath, blocksize='default',
lineterminator=None, compression='infer', sample=256000,
sample\_rows=10, enforce=False, assume\_missing=False,
storage\_options=None, include\_path\_column=False, \*\*kwargs)

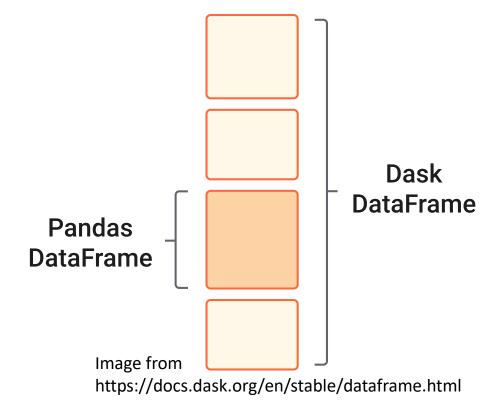
- Many ways to create dataframes
- https://docs.dask.org/en/stable/dataframe-create.html#
- From files/data sources

```
read_csv(urlpath[, blocksize, ...])
read_parquet(path[, columns, filters, ...])
read_hdf(pattern, key[, start, stop, ...])
read_orc(path[, engine, columns, index, ...])
read_json(url_path[, orient, lines, ...])
read_sql_table(table_name, con, index_col..)
read_sql_query(sql, con, index_col[, ...])
read_sql(sql, con, index_col, **kwargs)
read_table(urlpath[, blocksize, ...])
read_fwf(urlpath[, blocksize, ...])
```

- https://docs.dask.org/en/stable/array-api.html#random
- From dask objects

```
from_delayed(dfs[, meta, divisions, prefix, ...])
from_dask_array(x[, columns, index, meta])
dask.bag.core.Bag.to_dataframe([meta, ...]
```

- From other objects
  - from\_bcolz(x[, chunksize, categorize, ...])
  - from\_array(x[, chunksize, columns, meta])



### **DataFrames**

- DataFrames cover part of the pandas API:
  - Independent operations:
    - Element-wise operations: df.x + df.y, df \* df
    - Row-wise selections: df[df.x > 0]
    - Loc: df.loc[4.0:10.5]
    - Common aggregations: df.x.max(), df.max()
    - Is in: df[df.x.isin([1, 2, 3])]
    - Date time/string accessors: df.timestamp.month
  - Group operations:
    - groupby-aggregate (with common aggregations): df.groupby(df.x).y.max(), df.groupby('x').max()
    - groupby-apply on index: df.groupby(['idx', 'x']).apply(myfunc), where idx is the index level name
    - value counts: df.x.value counts()
    - Drop duplicates: df.x.drop duplicates()
    - Join on index: dd.merge(df1, df2, left index=True, right index=True)
    - Join with Pandas DataFrames: dd.merge(df1, df2, on='id')
    - Element-wise operations with different partitions / divisions: df1.x + df2.y
    - Date time resampling: df.resample(...)
    - Rolling averages: df.rolling(...)
    - Pearson's correlation: df[['col1', 'col2']].corr()
  - Group operations requiring data reordering
    - Set index: df.set\_index(df.x)
    - groupby-apply not on index (with anything): df.groupby(df.x).apply(myfunc)
    - Join not on the index: dd.merge(df1, df2, on='name')

### **DataFrames**



- DataFrame has the following limitations:
  - Setting a new index from an unsorted column is expensive
  - Many operations like groupby-apply and join on unsorted columns require setting the index, which as mentioned above, is expensive
  - The Pandas API is very large. Dask DataFrame does not attempt to implement many Pandas features or any of the more exotic data structures like NDFrames
  - Operations that were slow on Pandas, like iterating through row-by-row, remain slow on Dask DataFrame
- Full API is <a href="https://docs.dask.org/en/stable/dataframe-api.html">https://docs.dask.org/en/stable/dataframe-api.html</a>

# Dask Bag



- Bag is like a list or set
  - Unordered collection of data with repeats, i.e. {1, 2, 2, 3}
  - Immutable
- Operations
  - map, groupby, filter, fold, etc...
  - Full API <a href="https://docs.dask.org/en/latest/bag-api.html">https://docs.dask.org/en/latest/bag-api.html</a>
- Parallelise simple computations
  - unstructured or semi-structured data
  - i.e. text data, log files, JSON records, or user defined Python objects.
- Implemented using multi-processing (not threads)
  - Reduces communication efficiency between bag elements

### Bag

```
    Creating bags:

    from_sequence(seq[, partition_size, npartitions])
    from_delayed(values)
    from_url(urls)
    range(n, npartitions)
    read_text(urlpath[, blocksize, compression, ...])
    read_avro(urlpath[, blocksize, ...])

    DataFrame.to_bag([index, format])

  Bag operations

    Bag.accumulate(binop[, initial])

    import dask.bag as db
    from operator import add
    b = db.from_sequence([1, 2, 3, 4, 5], npartitions=2)
    b.accumulate(add).compute()
       Bag.all(split_every=None)

    Bag.any(split_every=None)

    Bag.count([split_every])
    Bag.max([split_every])
      Bag.min([split_every])
      Bag.msum([split_every])
    import dask.bag as db
    bool_bag = db.from_sequence([True, True, False])
    bool_bag.all().compute()
```

### Bag

```
Bag.reduction(perpartition, aggregate[, ...])
   Bag.random_sample(prob, random_state=None)

    Bag.filter(predicate)

def iseven(x):
    return x % 2 == 0
import dask.bag as db
b = db.from_sequence(range(5))
list(b.filter(iseven))
• Bag.groupby(grouper[, method, npartitions, ...])
import dask.bag as db
b = db.from_sequence(range(10))
iseven = lambda x: x \% 2 == 0
dict(b.groupby(iseven))
   Bag.foldby(key, binop[, initial, combine, ...])

    Combined reduction and groupby

      • Efficient parallel split-apply-combine tasks.
   Bags can't be changed

    Immutable

   Arrays and DataFrame are faster than Bags
   Bag groupby is slow.
      • foldby faster alternative if possible
```

# delayed

- Direct task graph creation
  - Delay python functions dask.delayed(func)(inputs...)
  - Code annotation@dask.delayed
  - compute still required to complete
- Delay function specifics matter
  - dask.delayed(f(x, y))
    - Calculates **f** first then delays the output
  - dask.delayed(f)(x, y)
    - Enables lazy evaluation
- Don't delay other dask functionality

```
def inc(x):
                                  import dask
    return x + 1
                                  @dask.delayed
def double(x):
                                  def inc(x):
    return x * 2
                                      return x + 1
def add(x, y):
                                  @dask.delayed
    return x + y
                                  def double(x):
                                      return x * 2
data = [1, 2, 3, 4, 5]
                                  @dask.delayed
output = []
                                  def add(x, y):
for x in data:
                                      return x + y
    a = inc(x)
    b = double(x)
                                  data = [1, 2, 3, 4, 5]
    c = add(a, b)
    output.append(c)
                                  output = []
                                  for x in data:
total = sum(output)
                                      a = inc(x)
                                      b = double(x)
                                      c = add(a, b)
import dask
                                      output.append(c)
                                  total = dask.delayed(sum)(output)
                                  total.compute()
```

```
output = []
for x in data:
    a = dask.delayed(inc)(x)
    b = dask.delayed(double)(x)
    c = dask.delayed(add)(a, b)
    output.append(c)

total = dask.delayed(sum)(output)
total.compute()
```

#### **Futures**



- Futures provide more complex functionality to build arbitrary task graphs
  - Similar to delayed, but tasks executed as soon as available
- Create task:

```
Client.submit(func, *args[, key, workers, ...])
Client.map(func, *iterables[, key, workers, ...])
Future.result([timeout])
```

• Move data:

```
Client.gather(futures[, errors, direct, ...])
Client.scatter(data[, workers, broadcast, ...])
```

#### **Futures**



```
from dask.distributed import Client
client = Client() # start local workers as processes
# or
client = Client(processes=False) # start local workers as threads
def inc(x):
   return x + 1
def add(x, y):
    return x + y
a = client.submit(inc, 10)
b = client.submit(inc, 20)
c = client.submit(add, a, b)
c.result()
futures = client.map(inc, range(1000))
```

### **Parallelisation**



- Dask generally defaults to threaded parallelisation
  - Single node
  - Maximum cores available
  - Dask array and dataframe
- Bag uses the multiprocessing scheduler by default
- Threads are lightweight workers for the main process (program)
  - Easy to parallelise
  - Share data easily between workers
  - Often doesn't scale up to all cores efficiently
- Processes are heavier weight workers (copies of the main program)
  - More scope for independent work
  - More heavy weight in startup costs
  - Explicit communication needed between workers if required

# Dask scheduling

- Scheduling choices can be modified by the user
- Default is generally threading:

```
import dask
dask.config.set(scheduler='threads')
```

- Works well for Array, DataFrame, and Delayed
- Relies on threading, so native python code will be restricted by the GIL
- Can change to processes:

```
import dask
dask.config.set(scheduler='processes')
```

- No GIL issues but slower for inter-task communications
- More advanced scheduling can be done using distributed from dask.distributed import Client client = Client()
  - Defaults to processesclient = Client(processes=False)
  - Might require distributed to be installed (not part of core dask install)
- Distributed can do single node or multi node
  - asynchronous API (Futures)
  - Has a dashboard
  - Improved data locality functionality for multi process work





# Dask scheduling

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- Can customise amount of resources and scheduler
  - Per run basis:

```
x.sum().compute(scheduler='processes')
```

As a context:

```
with dask.config.set(scheduler='threads'):
    x.compute()
```

• Globally:

```
dask.config.set(scheduler='threads')
```

Number of workers

```
from multiprocessing.pool import ThreadPool
dask.config.set(pool=ThreadPool(8))
```

Distributed scheduler

```
client = Client(processes=False, n_workers=4)
```

### Distributed scheduler

```
    Distributed Client

      from dask.distributed import Client
      client = Client(...)
      df.x.sum().compute()

    Local cluster (single node)

      from dask.distributed import Client, LocalCluster
      cluster = LocalCluster()
      client = Client(cluster)
      df.x.sum().compute()
• dask_jobqueue provides batch system interfaces
      HTCondorCluster([n_workers, loop, security, ...])
      LSFCluster([n_workers, loop, security, ...])
      MoabCluster([n_workers, loop, security, ...])
      OARCluster([n_workers, loop, security, ...])
      PBSCluster([n_workers, loop, security, ...])
      SGECluster([n_workers, loop, security, ...])
      SLURMCluster([n_workers, loop, security, ...])
• dask_mpi provides MPI launch interfacing
      from dask_mpi import
      initialize
      initialize()
      from dask.distributed import Client
      Client = Client()
      mpirun -np 4 python my_client_script.py
      mpirun -np 4 dask-mpi --scheduler-file ~/dask-scheduler.json
      from dask.distributed import Client
```

client = Client(scheduler\_file='~/dask-scheduler.json')



# Guiding principles



- If you don't need Dask, don't use it
  - Numpy array, panda dataframe, etc... all faster for small scale, in-memory
- Chunking/granularity important for performance
  - Too big chunks -> not enough parallelism
  - Too small chunks -> large parallelisation overhead
- compute as infrequently as possible
  - compute forces evaluation of the task graph
  - Might need multiple compute calls if task graph gets too big
- Mix threads and processes if doing larger parallelisation
  - Threads good for small scale parallelisation
- Load data with dask
- Persist datasets to memory when reduced

### Dask on ARCHER2



- Dask is available in the cray-python
  - Threading/shared memory backends
- Distributed dask needs to be installed

```
module load cray-python/3.9.7.1
export PYTHONUSERBASE=/work/t01/t01/auser/.local
export PATH=$PYTHONUSERBASE/bin:$PATH
python -m pip install --user dask distributed --upgrade
python -m pip install --user dask-jobqueue --upgrade
```

- Currently need to submit from compute nodes
  - Dask runs a scheduler that needs connection from the workers

### Threaded Dask on ARCHER2



- Running single node dask as normal job works fine
  - Default mode is threading
  - Requires process binding and thread placement to be sensible

```
#!/bin/bash
#SBATCH --job-name=my_job
#SBATCH --nodes=1
#SBATCH --tasks-per-node=1
#SBATCH --cpus-per-task=128
#SBATCH --partition=standard
#SBATCH --qos=short
#SBATCH --account=z19
#SBATCH --time=0:10:0
python dask-program.py
```

Can also do an interactive run:

srun --nodes=1 --tasks-per-node=1 --cpus-per-task=128 --exclusive --partition=standard --qos=short --reservation=shortqos --account=z19 --time=0:20:0 python dask-program.py

### Distributed Dask on ARCHER2



```
from dask_jobqueue import SLURMCluster
cluster = SLURMCluster(cores=128,
                       processes=128,
                       memory='256GB',
                       queue='standard',
                       header_skip=['--mem'],
                       job_extra=['--qos="standard"'],
                       python='srun python',
                       project='t01',
                       walltime="01:00:00",
                       shebang="#!/bin/bash --login",
                       local_directory='$PWD'
                       interface='hsn0',
                       env_extra=['module load cray-python',
                                   'export PYTHONUSERBASE=/work/t01/t01/auser/.local/',
                                   'export PATH=$PYTHONUSERBASE/bin:$PATH',
                                   'export PYTHONPATH=$PYTHONUSERBASE/lib/python3.8/site-
packages:$PYTHONPATH'])
cluster.scale(jobs=2)
                         # Deploy two single-node jobs
from dask.distributed import Client
client = Client(cluster) # Connect this local process to remote workers
import dask.array as da
x = da.random.random((10000, 10000), chunks=(1000, 1000))
mean = x.mean().compute()
```

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