

- Log into scinet with your account:

- ssh -Y hpcsi0XX@login.scinet.utoronto.ca
- ssh -Y gpc01
- qsub -X -I -l nodes=1:ppn=8,walltime=4:00:00
- cd \${SCRATCH}
- cp -r /scinet/course/parIO .
- cd parIO
- source parallellibs



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Parallel I/O

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Agenda

- Intro to I/O
- MPI-IO
- HDF5, NetCDF4
- Parallel HDF5/NetCDF4
- ADIOS (?)



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Data is getting bigger

- Increase in computing power makes simulations larger/more frequent
- Increase in sensor technology makes experiments/observations larger
- Data sizes that used to be measured in MB/GB now measured in TB/PB.
- Easier to generate the data than to store it:



Economist, 27 Feb 2010



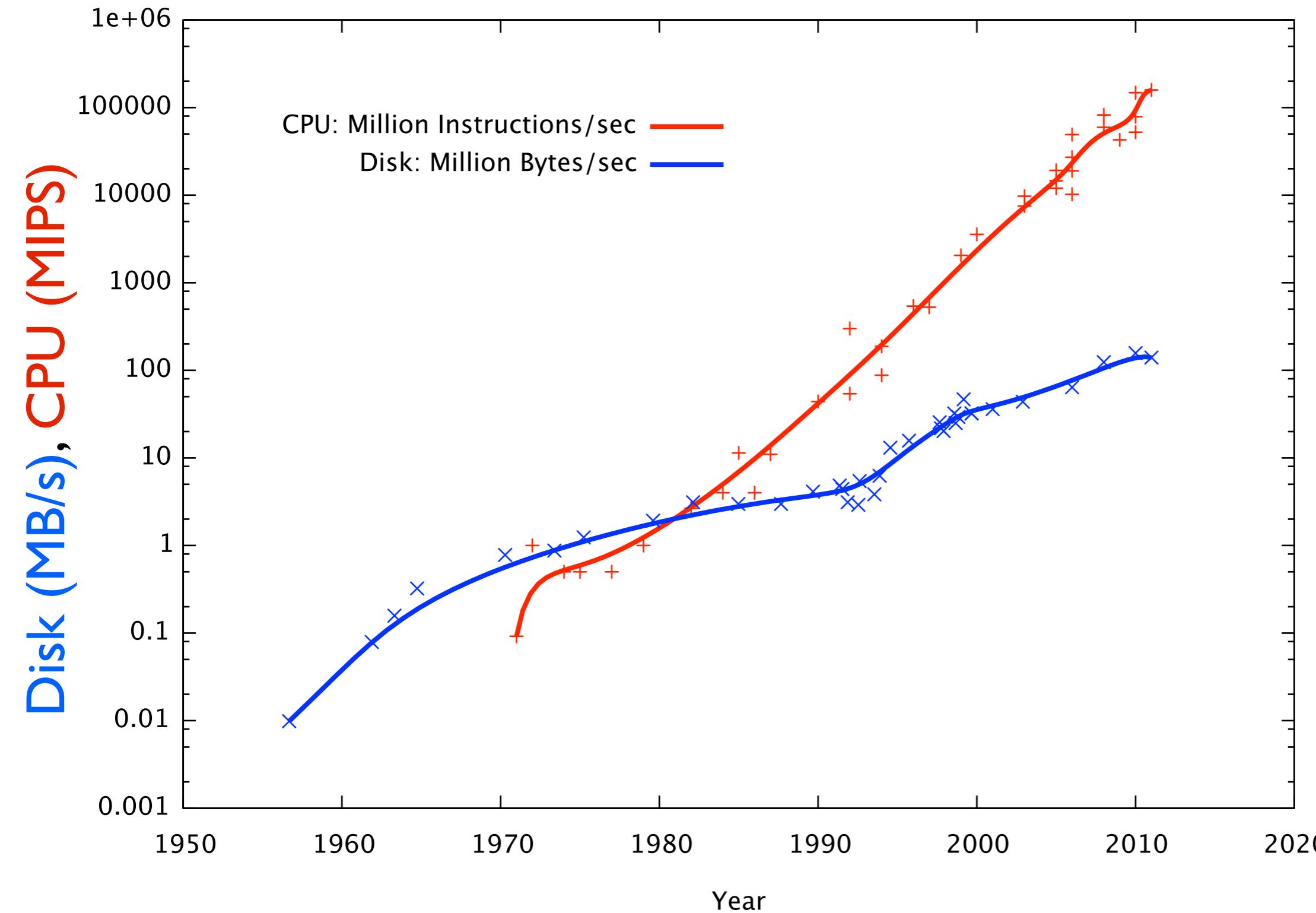
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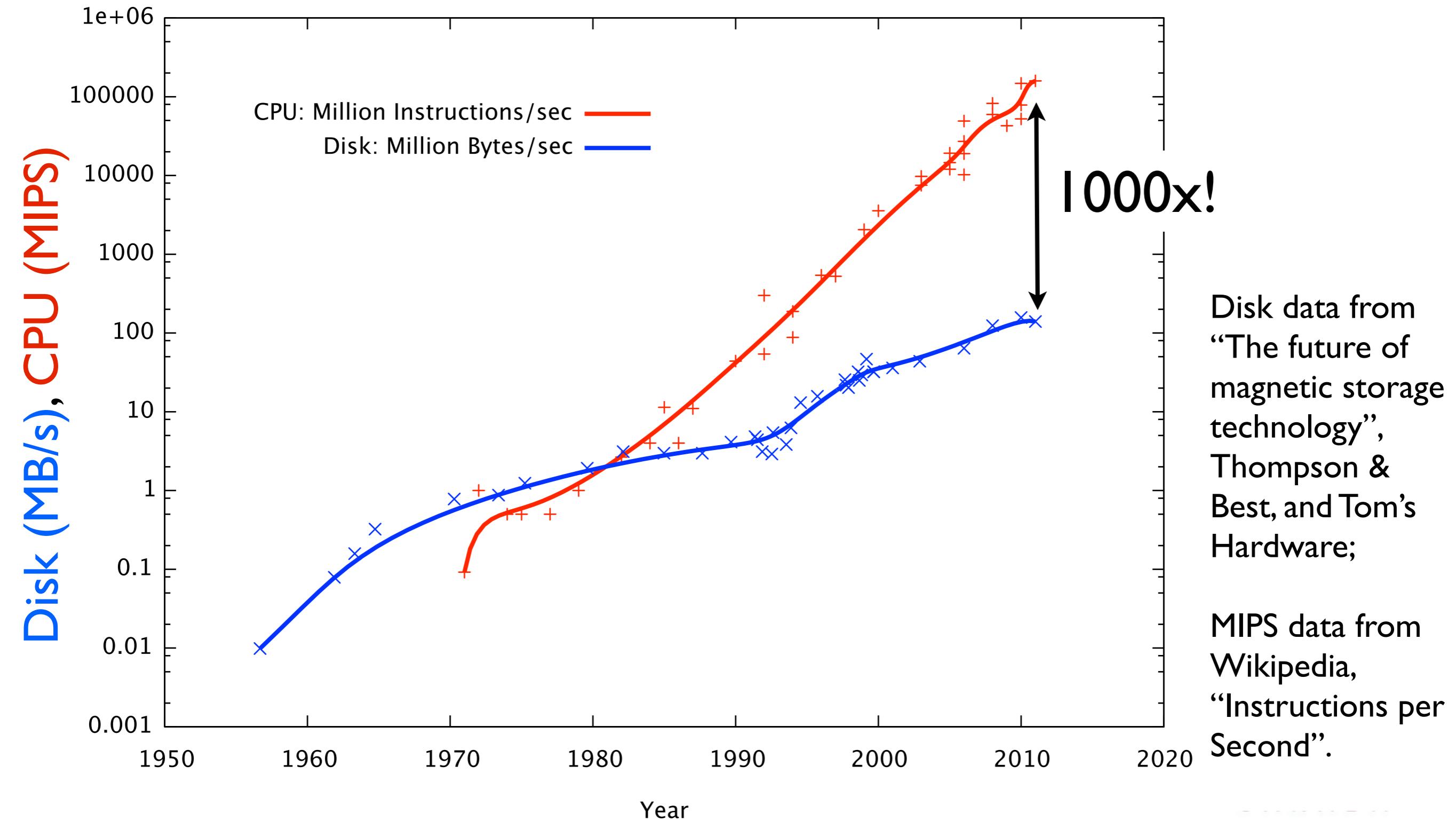
Disks are slower than CPUs (and getting slower)



Disk data from
“The future of
magnetic storage
technology”,
Thompson &
Best, and Tom’s
Hardware;

MIPS data from
Wikipedia,
“Instructions per
Second”.

Disks are slower than CPUs (and getting slower)



Understanding storage performance

- Data rate: MB/s
 - Peak or sustained
 - Write faster than read
 - Network analogy: bandwidth
- IOPS: I/O Operations Per Second
 - open, close, seek, read, write
 - Network analogy: I latency



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Understanding storage performance

Device	Bandwidth (MB/s)	IOPS
SATA HDD	100	100
SSD	250	10000

HD:

Open, Write, Close 1000x1kB files: 30.01s (eff: 0.033 MB/s)

Open, Write, Close 1x1MB file: 40ms (eff: 25 MB/s)



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Understanding storage performance

Device	Bandwidth (MB/s)	IOPS
SATA HDD	100	100
SSD	250	10000

SSD:

Open, Write, Close 1000x1kB files: 300ms (eff: 3.3 MB/s)

Open, Write, Close 1x1MB file: 4ms (eff: 232 MB/s)



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Understanding storage performance

- SSDs are much faster at IOPS:
 - No physical mechanisms that must move to file position
 - Read ~ as fast as write
- But still latency at controller, system calls, etc.
- SSDs will always be much more expensive than disk per unit storage - disk isn't going away.



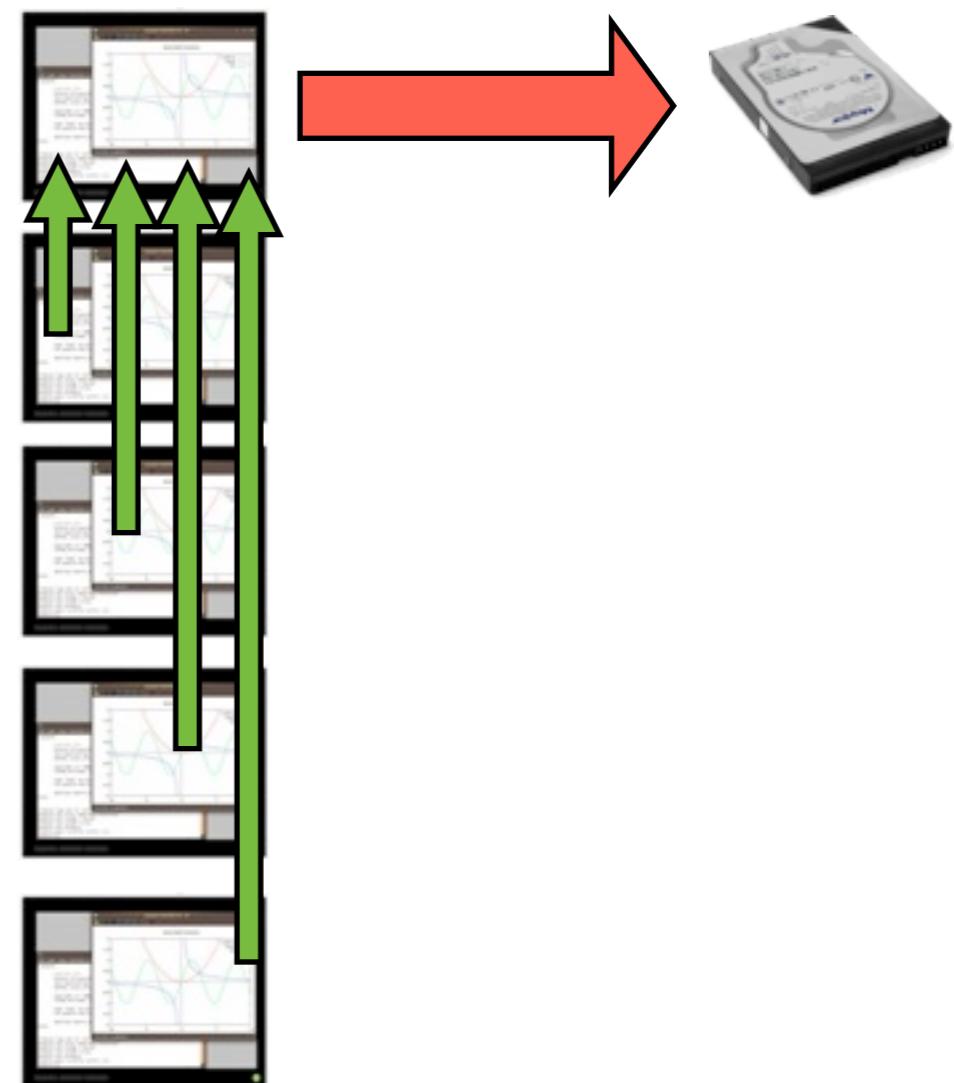
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Planning your I/O

- Parallel computation, several options.
- Everyone sends data to process 0
- Process 0 writes.
- Serialize I/O - huge bottleneck.

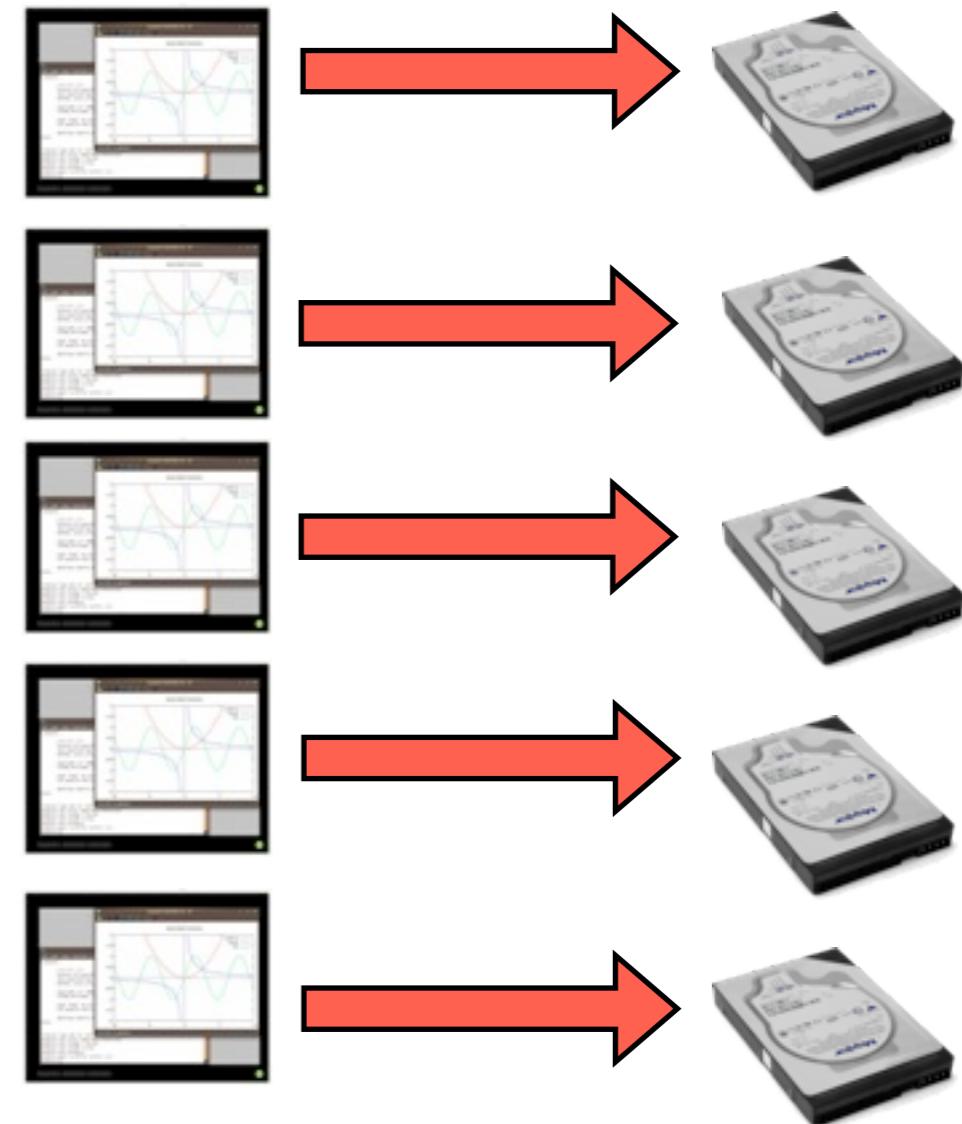


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Planning your I/O

- Parallel computation, several options.
- Each process writes a file, possibly to local disk.
- Postpones the problem
 - how do you analyze, or restart with different # of procs?



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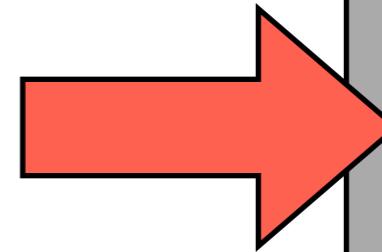
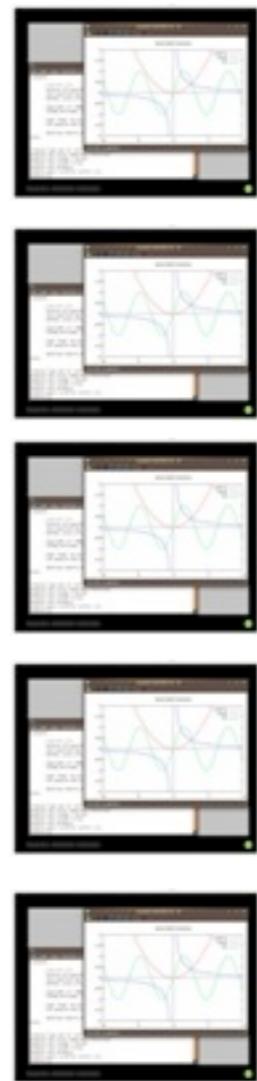


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Parallel FS

Planning your I/O

- Parallel computation, several options.
- We're going to learn to avoid doing this by using Parallel I/O
- **Coordinated** single output of multiple processes.



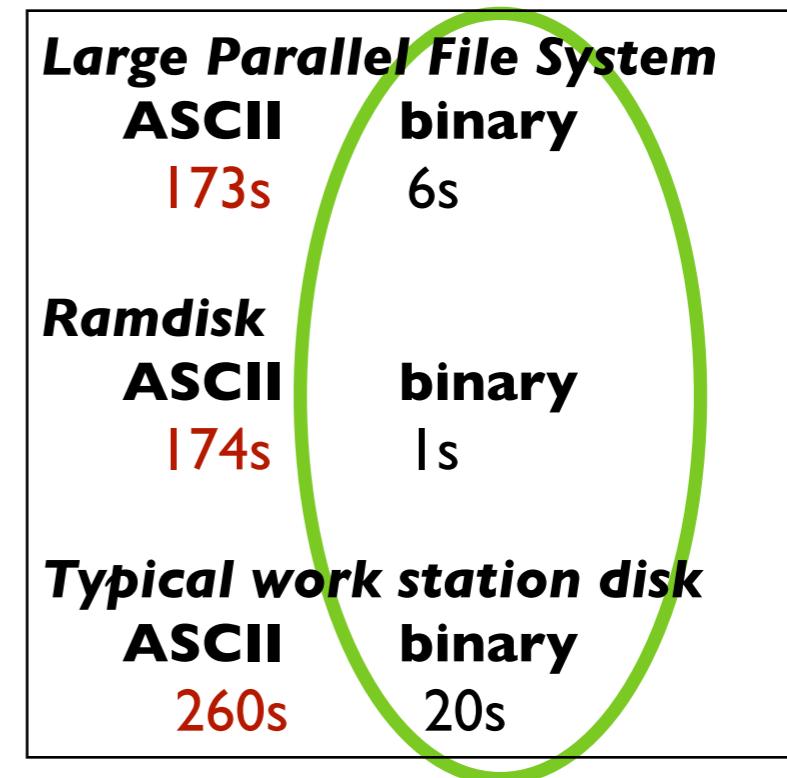
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Where, how you do I/O matters.

- Binary - smaller files, much faster to read/write.
- You're not going to read GB/TB of data yourself; don't bother trying.
- Write in 1 chunk, rather than a few #s at a time.



Timing data: writing 128M double-precision numbers



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Where, how you do I/O matters.

- All disk systems do best when reading/writing large, contiguous chunks
- I/O operations (IOPS) are themselves expensive
- moving around within a file
- opening/closing
- Seek - 3-15ms - enough time to read 0.75 MB!

Typical work station disk
binary - one large read
14s

binary - 8k at a time
20s

binary - 8k chunks, lots of seeks
150s

binary - seeky + open and closes
205s

Timing data: reading 128M double-precision numbers



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Where, how you do I/O matters.

- RAM is much better for random accesses
- Use right storage medium for the job!
- Where possible, read in contiguous large chunks, do random access in memory
- Much better *if* you use most of data read in

Large Parallel File System

ASCII	binary
173s	6s

Ramdisk

ASCII	binary
174s	1s

Typical work station disk

ASCII	binary
260s	20s

Ramdisk

binary - one large read
1s

binary - 8k at a time
1s

binary - 8k chunks, lots of seeks
1s

binary - seeky + open and closes
1.5s



Parallel I/O and large file systems

- Large disk systems featuring many servers, disks
- Can serve files to many clients concurrently
- Parallel File Systems -
- Lustre, Panasas, GlusterFS, Ceph, GPFS...



SciNet ~2k drives



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SciNet's File Systems

- 2x DCS9900 couplets
- 1,790 1TB SATA disk drives
- 1.4 PB of storage
- Single GPFS domain, accessed by all machines (TCS and GPC).
- Data to compute nodes via IB



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SciNet's File Systems

- Designed for HPC workloads
- High bandwidth to large files - big data.
- Does not do well with millions of little files:
 - wastes disk space (4MB block size)
 - lots of small scattered access is *terrible* for performance, even on desktop; multiply by hundreds of processors, can be disastrous



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Understanding storage performance

Device	Bandwidth (MB/s)	IOPS
SATA HDD	100	100
SSD	250	10000
SciNet GPFS	5000	30000

(SciNet GPFS used by ~3000 nodes.)



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Where, how you do I/O matters.

- Well built parallel file systems can greatly increase bandwidth
- Many pipes to network (servers), many spinning disks (bandwidth off of disks)
- But typically even **worse** penalties for seeky/IOPSy operations (coordinating all those disks.)
- Parallel FS can help with big data in two ways

Large Parallel File System		
ASCII		binary
173s		6s
Ramdisk		
ASCII		binary
174s		1s
Typical work station disk		
ASCII		binary
260s		20s

Large Parallel File System
binary - one large read
7.5s

binary - 8k at a time
62 s

binary - 8k chunks, lots of seeks
428 s

binary - seeky + open and closes
2137 s



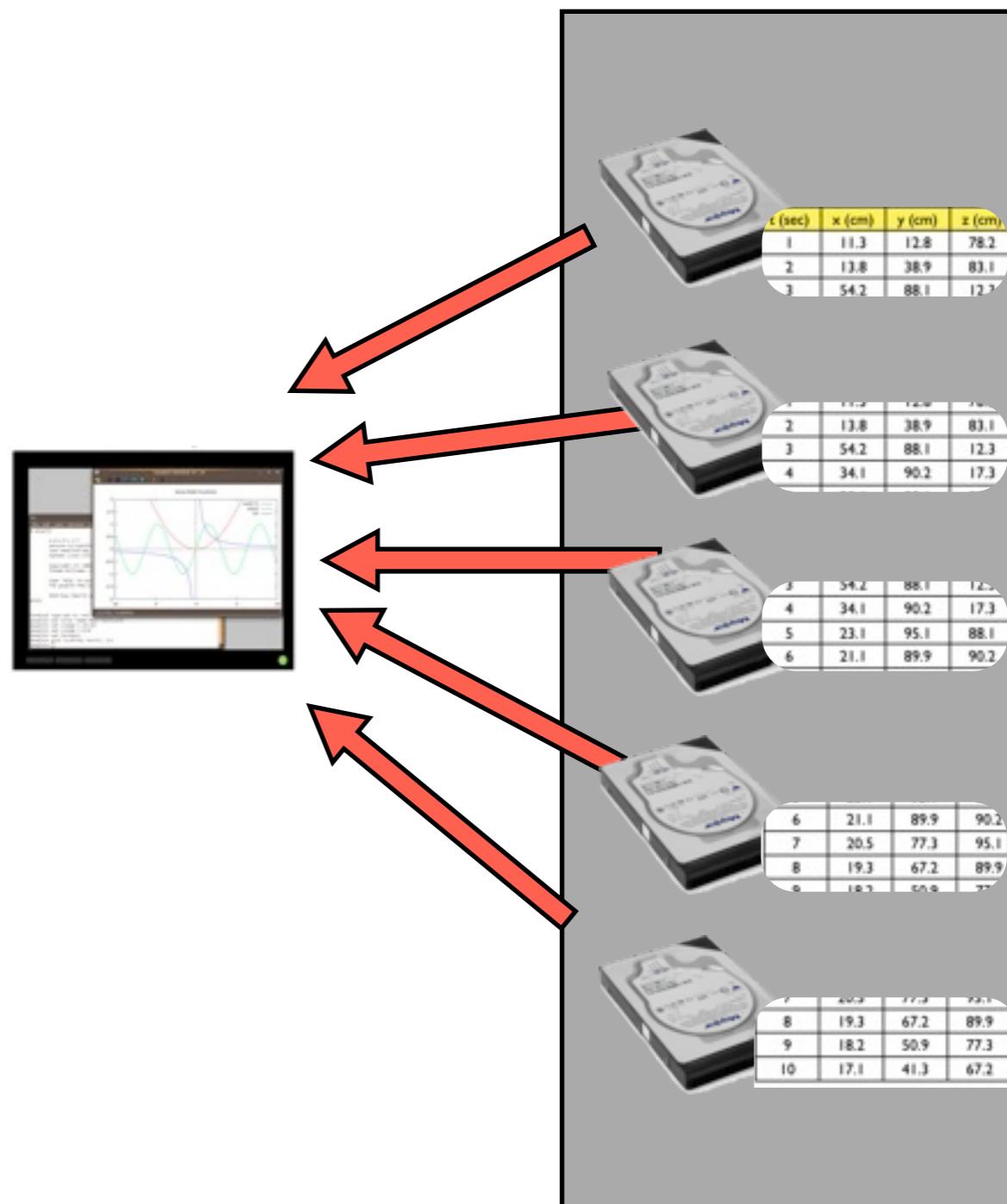
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Striping data across disks

- Single client can make use of multiple disk systems simultaneously
- “Stripe” file across many drives
- One drive can be finding next block while another is sending current block



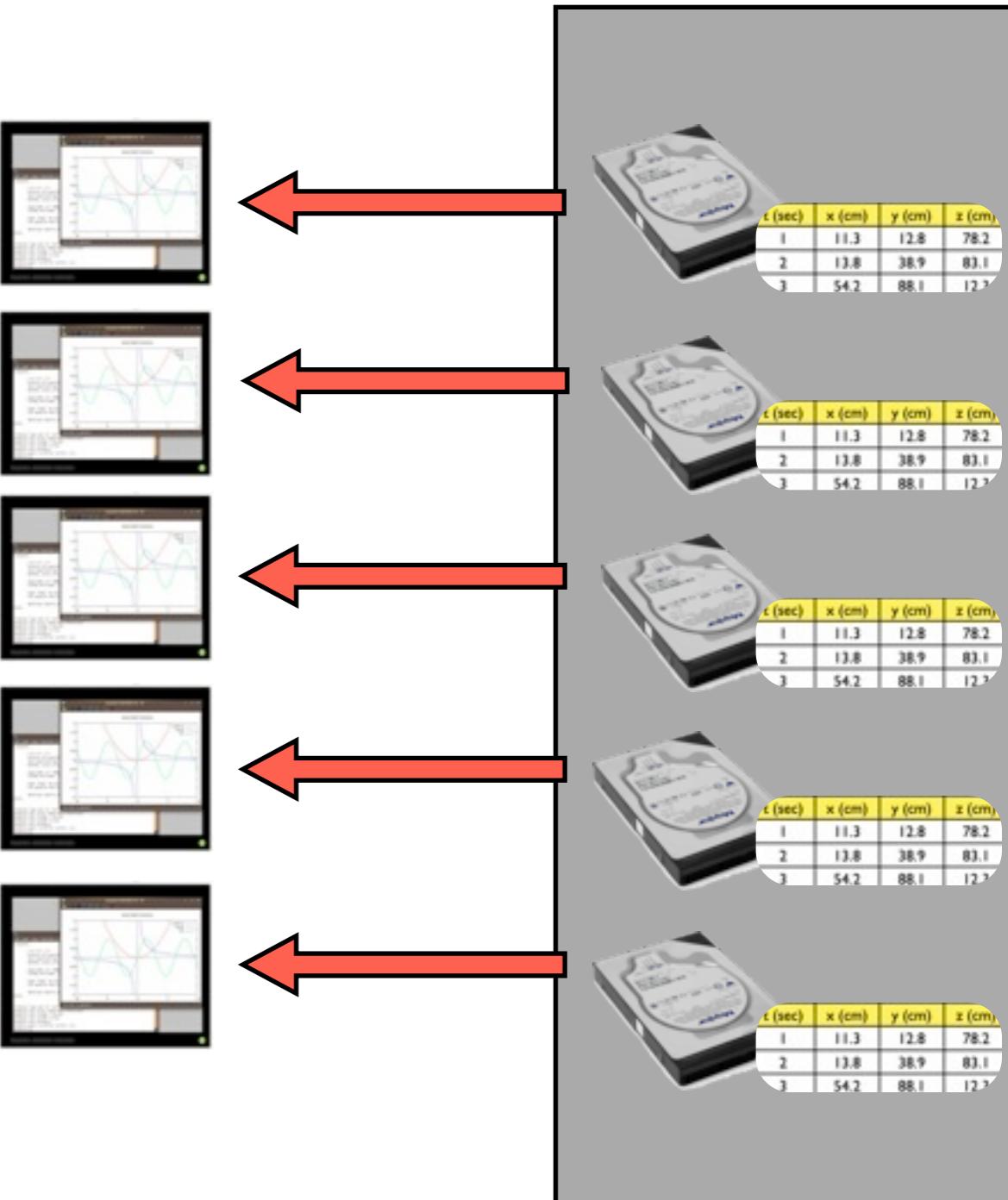
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Parallel operations on separate data

- Or can do truly parallel operations
- multiple clients doing independent work
- Easy parallelism (good for lots of small data) - process many small files separately



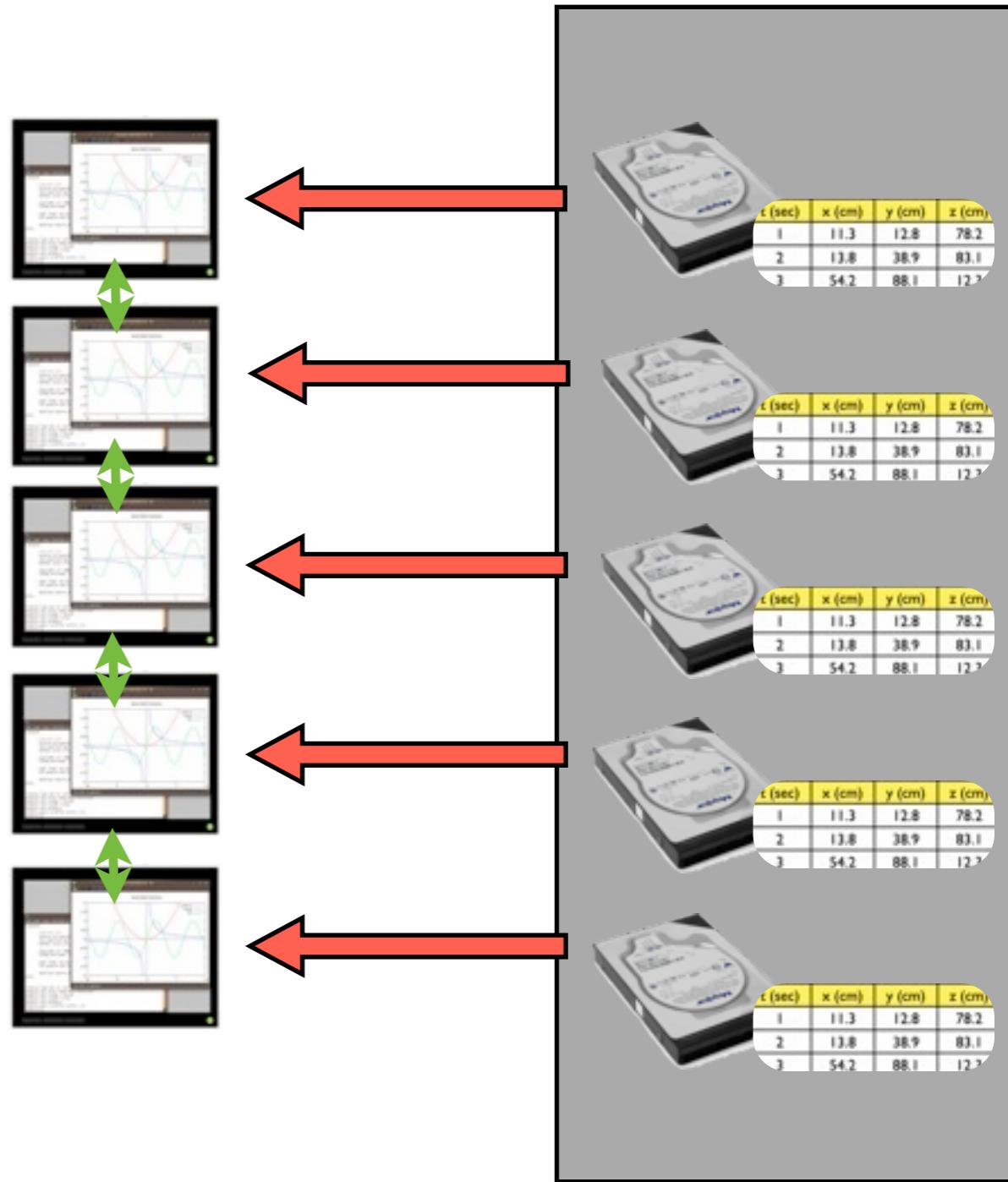
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Parallel operations on separate data

- Or can do truly parallel operations
- multiple clients doing independent work
- Easy parallelism (good for lots of small data) - process many small files separately
- Harder parallelism - each does part of a larger analysis job on a big file.



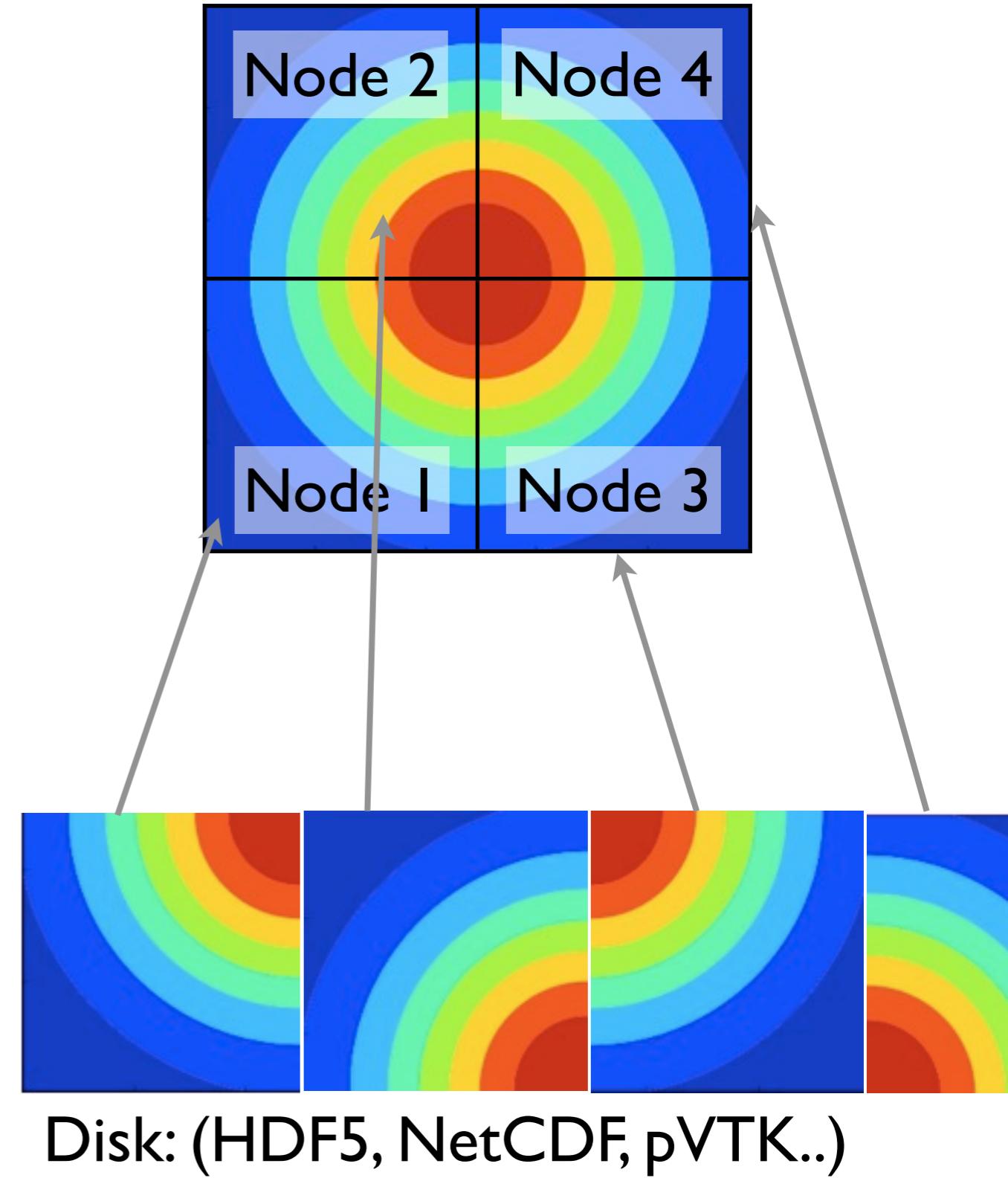
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Data files must take advantage of parallel I/O

- For parallel operations on single big files, parallel filesystem isn't enough
- Data must be written in such a way that nodes can efficiently access relevant subregions
- HDF5, NetCDF formats typical examples for scientific data



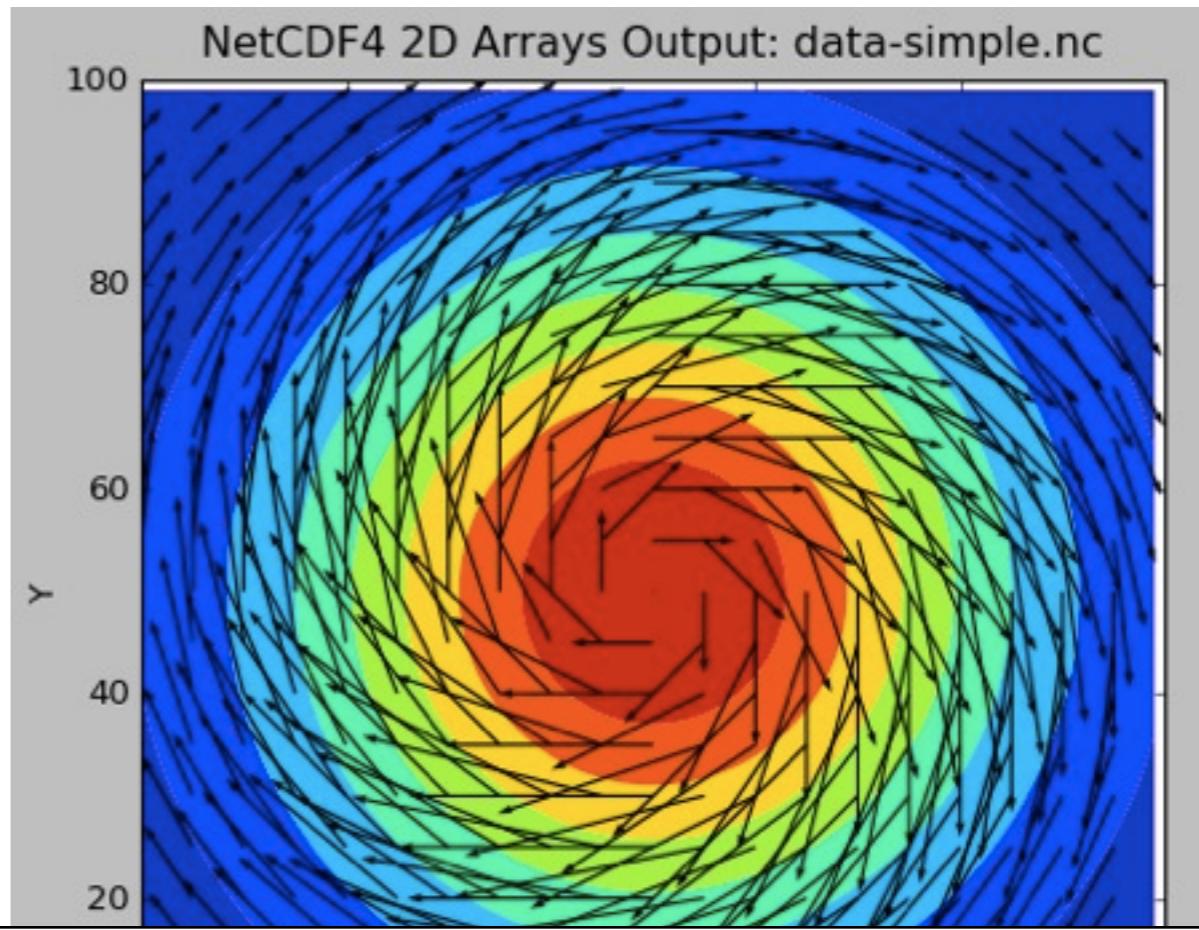
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These formats are *self-* *describing*

- HDF5, NetCDF have other advantages anyway
- Binary
- Self describing - contains not only data but names, descriptions of arrays, etc
- Many tools can read these formats
- Big data - formats matter



```
$ ncdump -h data-simple-fort.nc
netcdf data-simple-fort {
dimensions:
    x = 100 ;
    y = 100 ;
    velocity components = 2 ;
variables:
    double Density(y, x) ;
    double Velocity(y, x, velocity components) ;
}
```



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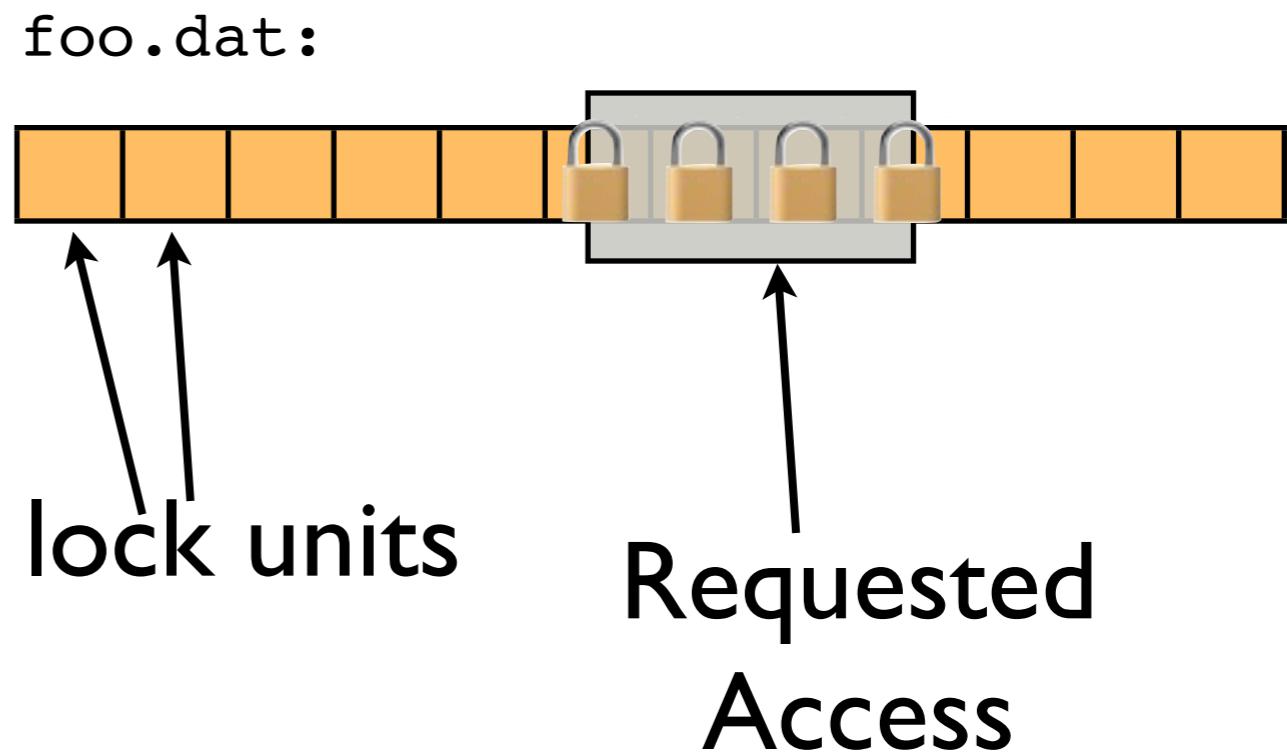
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Coordinating I/O

- Multiple nodes all accessing same filesystem.
- To prevent anarchy, locks for some requested accesses.
- File broken up into lock units, locks handed out upon request.
- “False sharing”, etc, possible.
- Files **and** directories.
- Makes (some) IOPS even more expensive



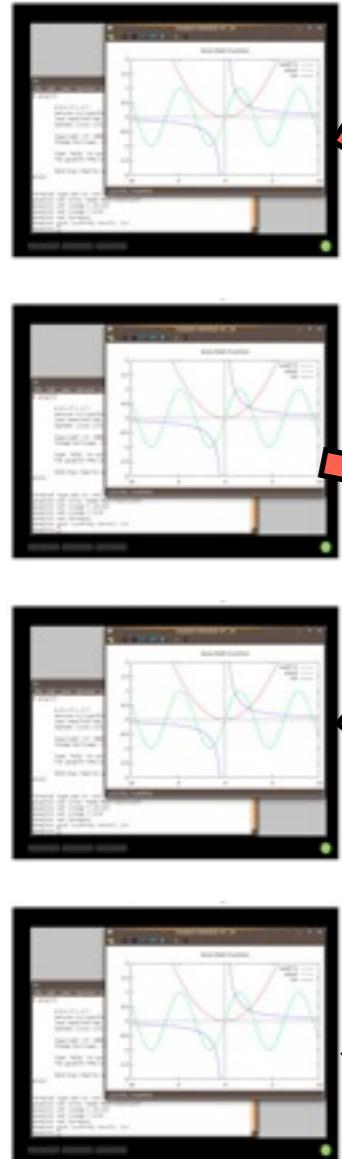
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Application

High-level Library

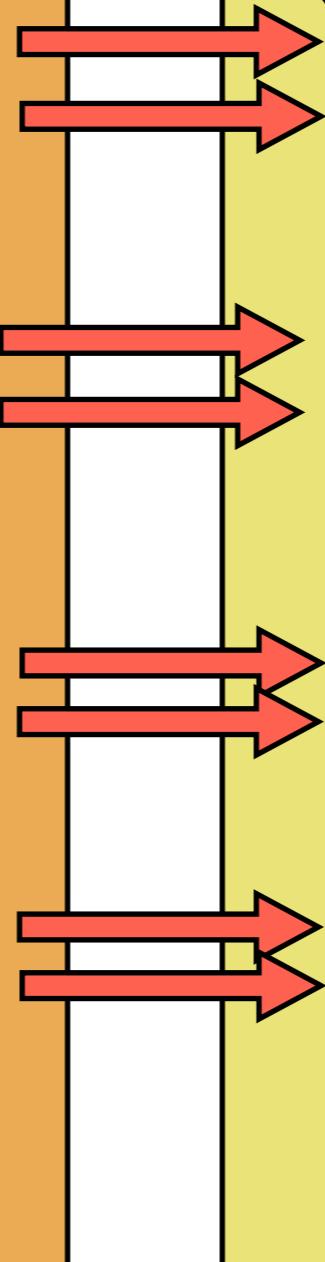


(HDF5, NetCDF,
ADIOS)

I/O

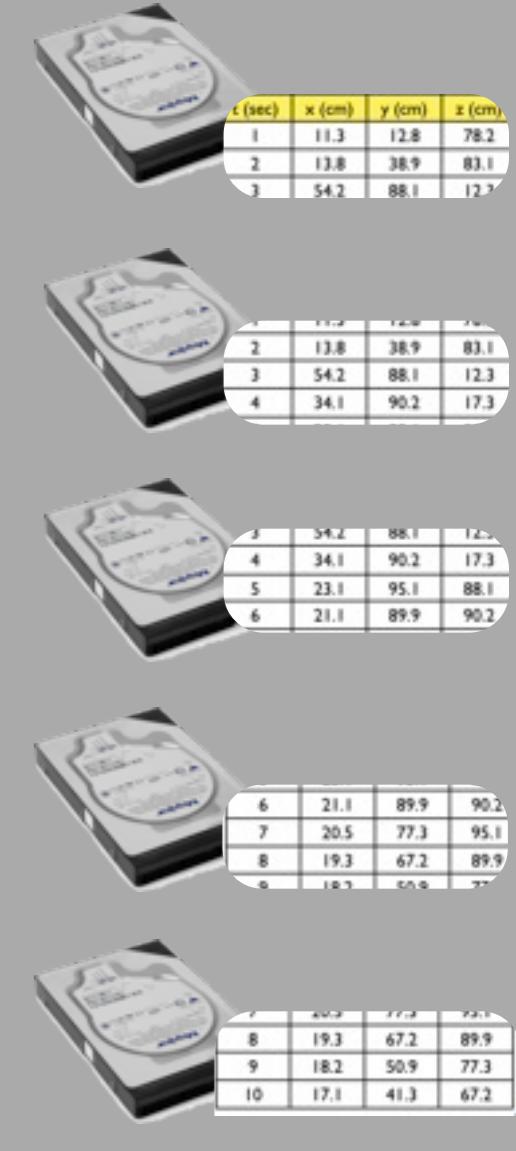
Middleware

(MPIIO)



Parallel FS

(GPFS, PVFS..)



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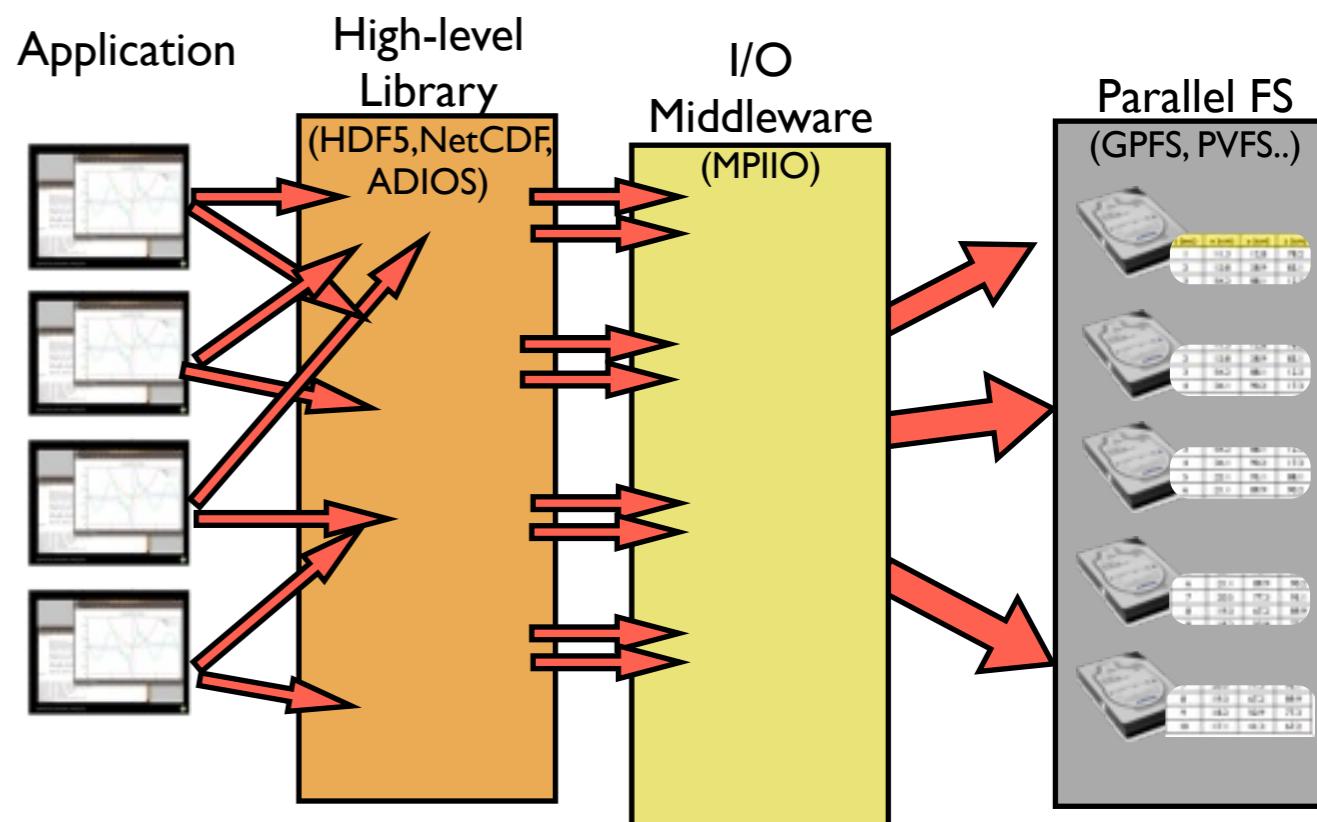
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Abstraction Layers

- High Level libraries can simplify programmers tasks
 - Express I/O in terms of the data structures of the code, not bytes and blocks
 - I/O middleware can coordinate, improve performance
 - Data Sieving
 - 2-phase I/O



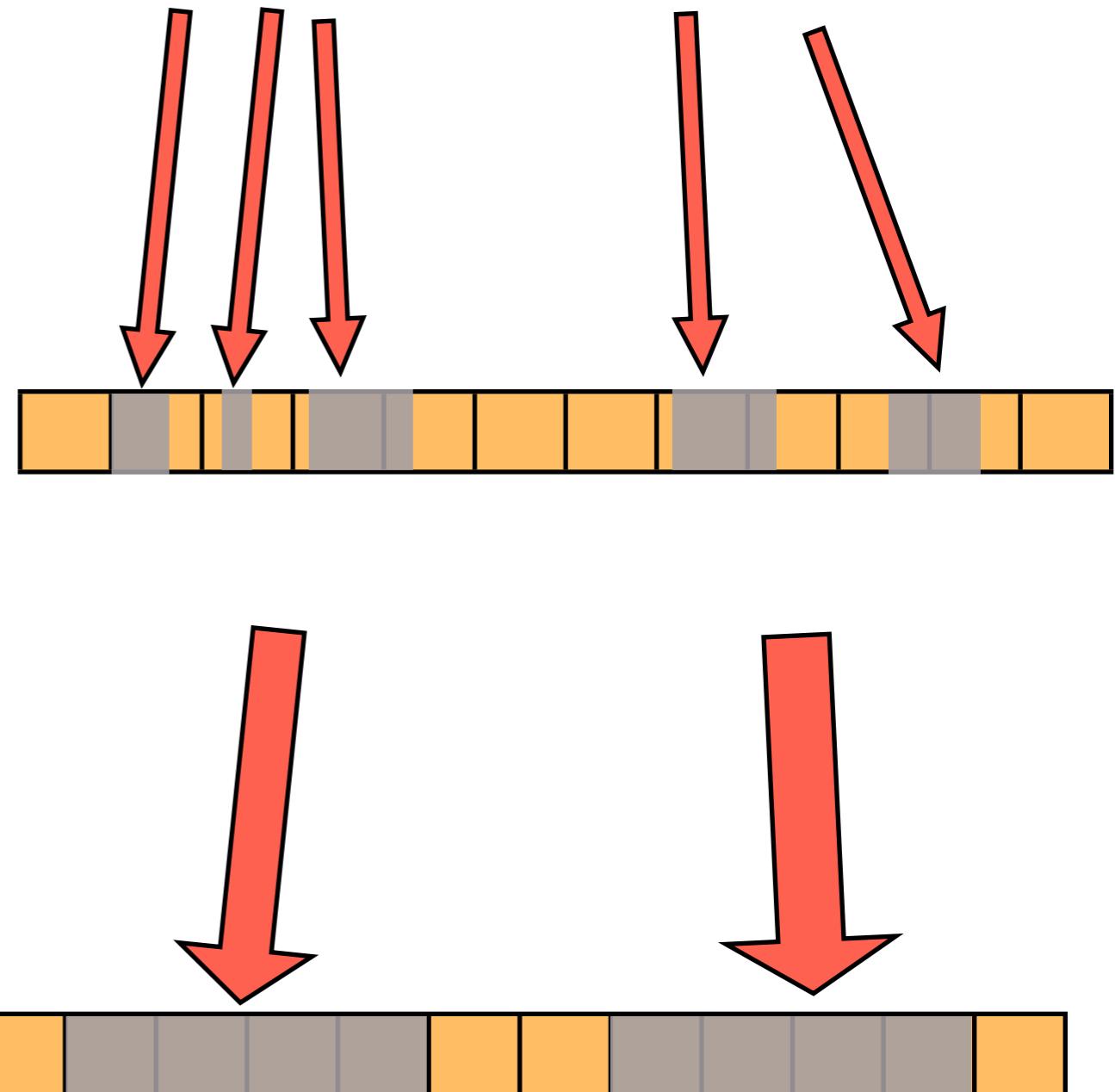
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Data Sieving

- Combine many non-contiguous IO requests into fewer, bigger IO requests
- “Sieve” unwanted data out
- Reduces IOPS, makes use of high bandwidth for sequential IO



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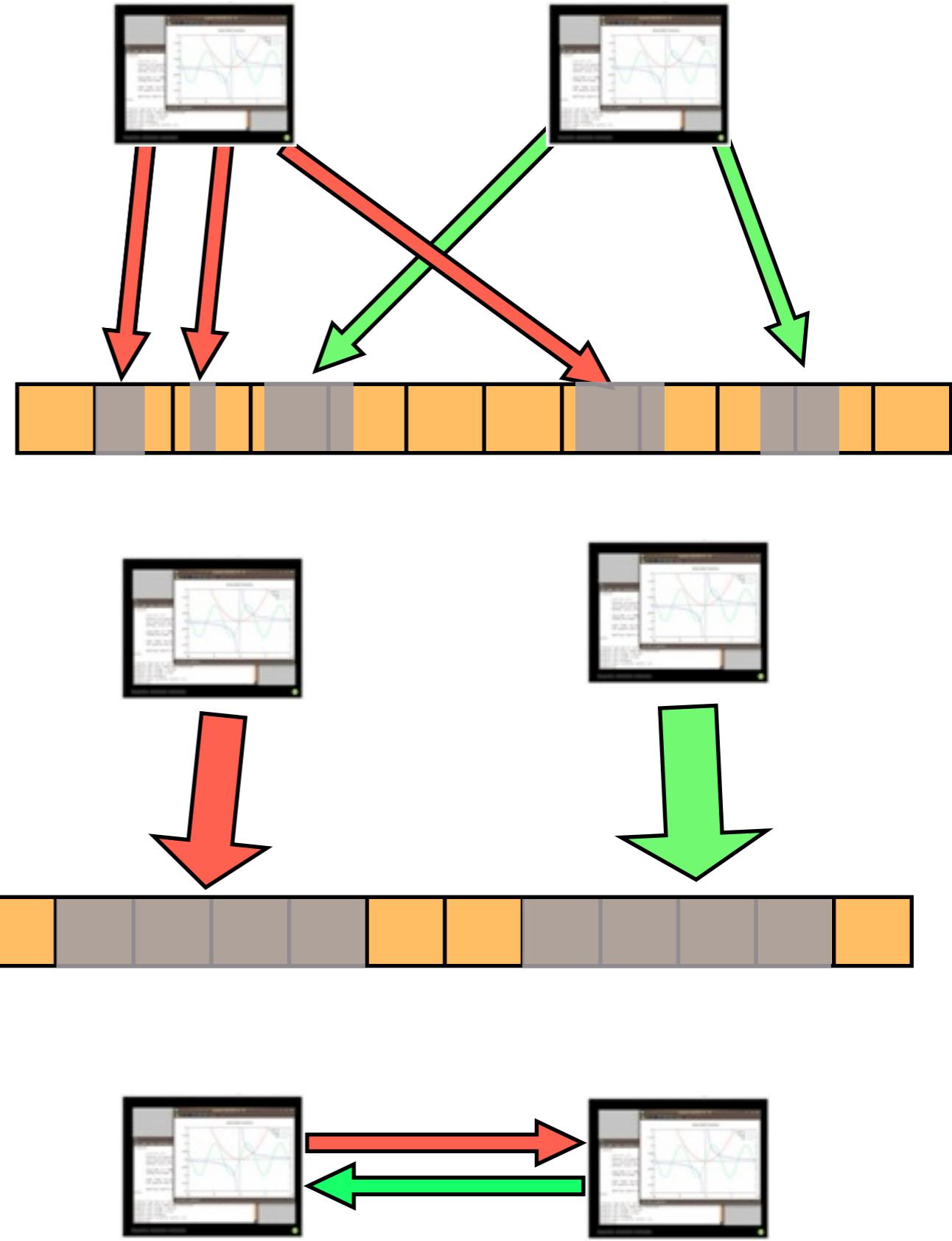
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Two-Phase IO

- Collect requests into larger chunks
- Have individual nodes read big blocks
- Then use network communications to exchange pieces
- Fewer IOPS, faster IO
- Network communication usually faster



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MPI-IO

- Part of MPI-2 standard
- Started at IBM Watson
- Maps I/O reads and writes to message passing
- ROMIO is the implementation found in MPICH2, OpenMPI
- Really only widely-available scientific computing parallel I/O middleware



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```
$ cd parIO
$ source parallellibs
$ cd samples/mpio
$ make
.....
$ mpiexec -n 4 ./helloworldc
Rank 0 has message <Hello >
Rank 1 has message <World!>
Rank 2 has message <Hello >
Rank 3 has message <World!>
Rank 4 has message <Hello >

$ cat helloworld.txt
Hello World!Hello World! $
```



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helloworldc.c

```
#include <stdio.h>
#include <string.h>
#include <mpi.h>

int main(int argc, char **argv) {
    int ierr, rank, size;
    MPI_Offset offset;
    MPI_File file;
    MPI_Status status;
    const int msgsize=6;
    char message[msgsize+1];

    ierr = MPI_Init(&argc, &argv);
    ierr |= MPI_Comm_size(MPI_COMM_WORLD, &size);
    ierr |= MPI_Comm_rank(MPI_COMM_WORLD, &rank);

    if ((rank % 2) == 0) strcpy (message, "Hello ");
    else strcpy (message, "World!");

    offset = (msgsize*rank);

    MPI_File_open(MPI_COMM_WORLD, "helloworld.txt", MPI_MODE_CREATE|MPI_MODE_WRONLY,
                  MPI_INFO_NULL, &file);
    MPI_File_seek(file, offset, MPI_SEEK_SET);
    MPI_File_write(file, message, msgsize, MPI_CHAR, &status);
    MPI_File_close(&file);

    MPI_Finalize();
    return 0;
}
```



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```
program MPIIO_helloworld
  use mpi
  implicit none

  integer(MPI_OFFSET_KIND) :: offset
  integer, dimension(MPI_STATUS_SIZE) :: wstatus
  integer, parameter :: msgsize=6
  character(msgsize) :: message
  integer :: ierr, rank, comsize, fileno

  call MPI_Init(ierr)
  call MPI_Comm_size(MPI_COMM_WORLD, comsize, ierr)
  call MPI_Comm_rank(MPI_COMM_WORLD, rank, ierr)

  if (mod(rank,2) == 0) then
    message = "Hello "
  else
    message = "World!"
  endif

  offset = rank*msgsize

  call MPI_File_open(MPI_COMM_WORLD, "helloworld.txt", ior(MPI_MODE_CREATE,MPI_MODE_WRONLY), &
                    MPI_INFO_NULL, fileno, ierr)
  call MPI_File_seek (fileno, offset, MPI_SEEK_SET, ierr)
  call MPI_File_write(fileno, message, msgsize, MPI_CHARACTER, wstatus, ierr)
  call MPI_File_close(fileno, ierr)

  call MPI_Finalize(ierr)

end program MPIIO_helloworld
```

helloworldf.f90



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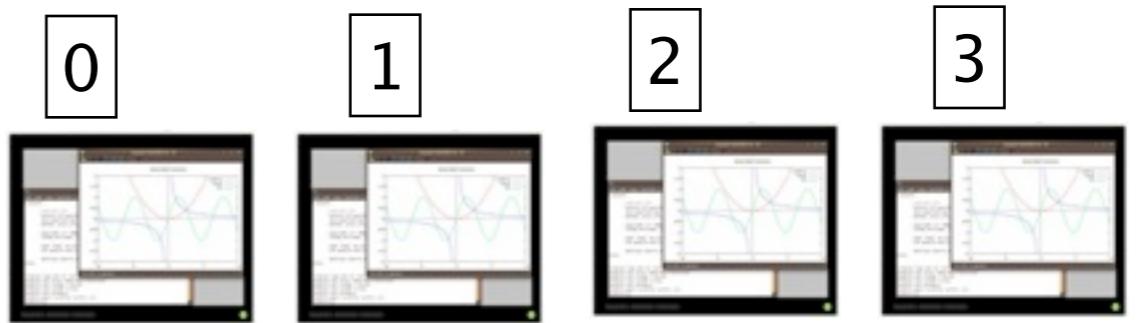


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MPI-IO

Hello

World



```
mpiexec -n 4 ./helloworldc
```



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MPI-IO

Hello

World



```
if ((rank % 2) == 0)
    strcpy (message, "Hello ");
else
    strcpy (message, "World!");
```



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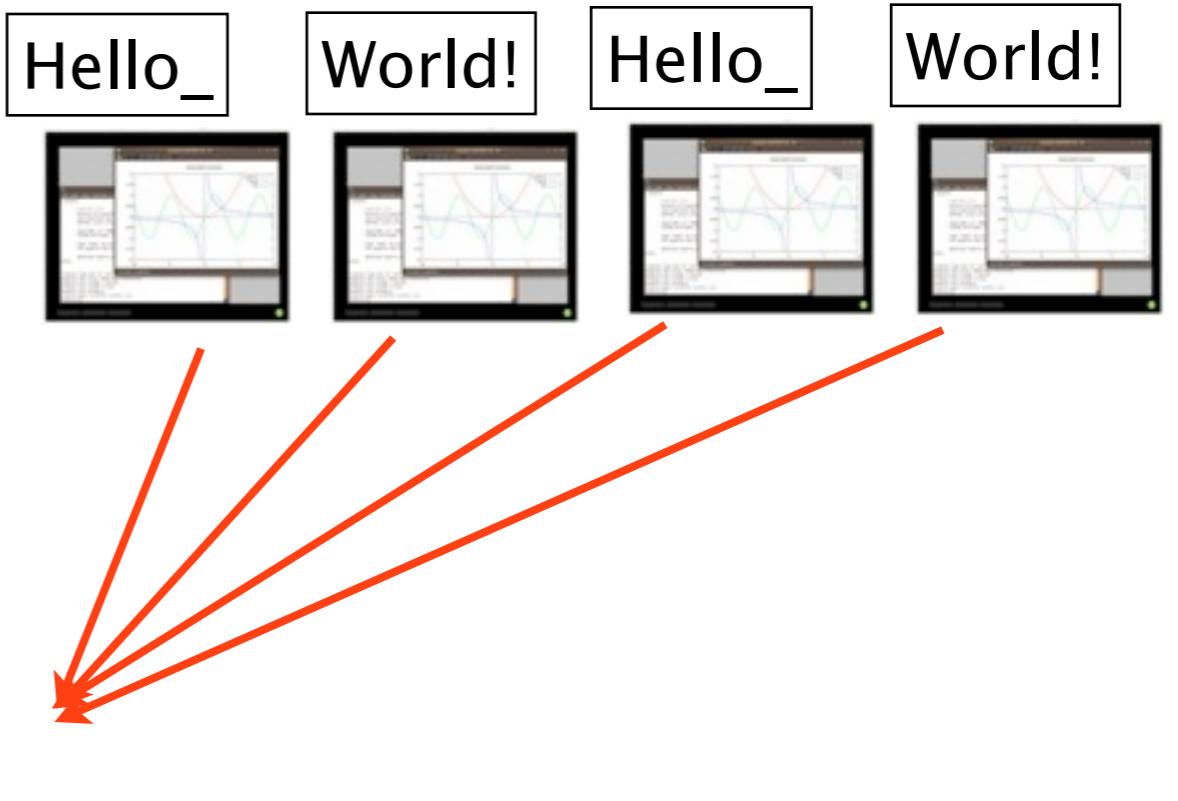


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MPI-IO

Hello

World



```
MPI_File_open(MPI_COMM_WORLD, "helloworld.txt", MPI_MODE_CREATE|MPI_MODE_WRONLY,  
MPI_INFO_NULL, &file);
```



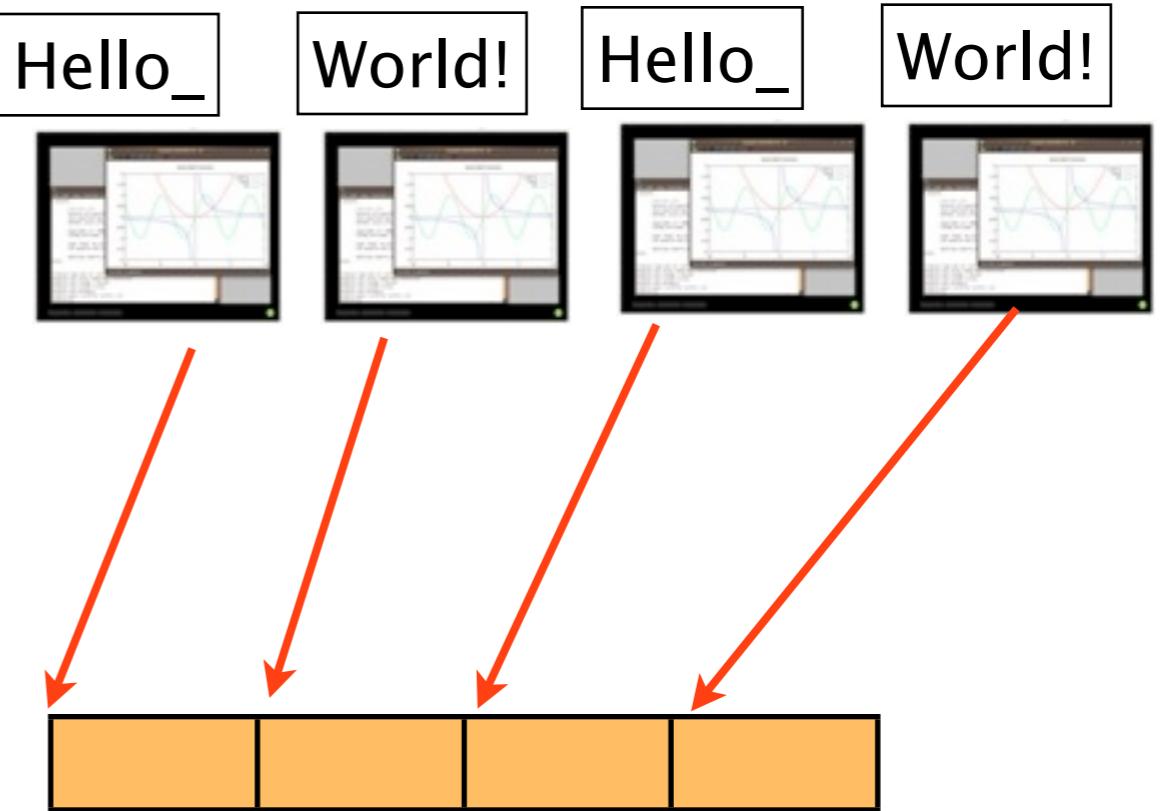
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MPI-IO

Hello World

helloworld.txt:



```
offset = (msgsize*rank);
```

```
MPI_File_seek(file, offset, MPI_SEEK_SET);
```



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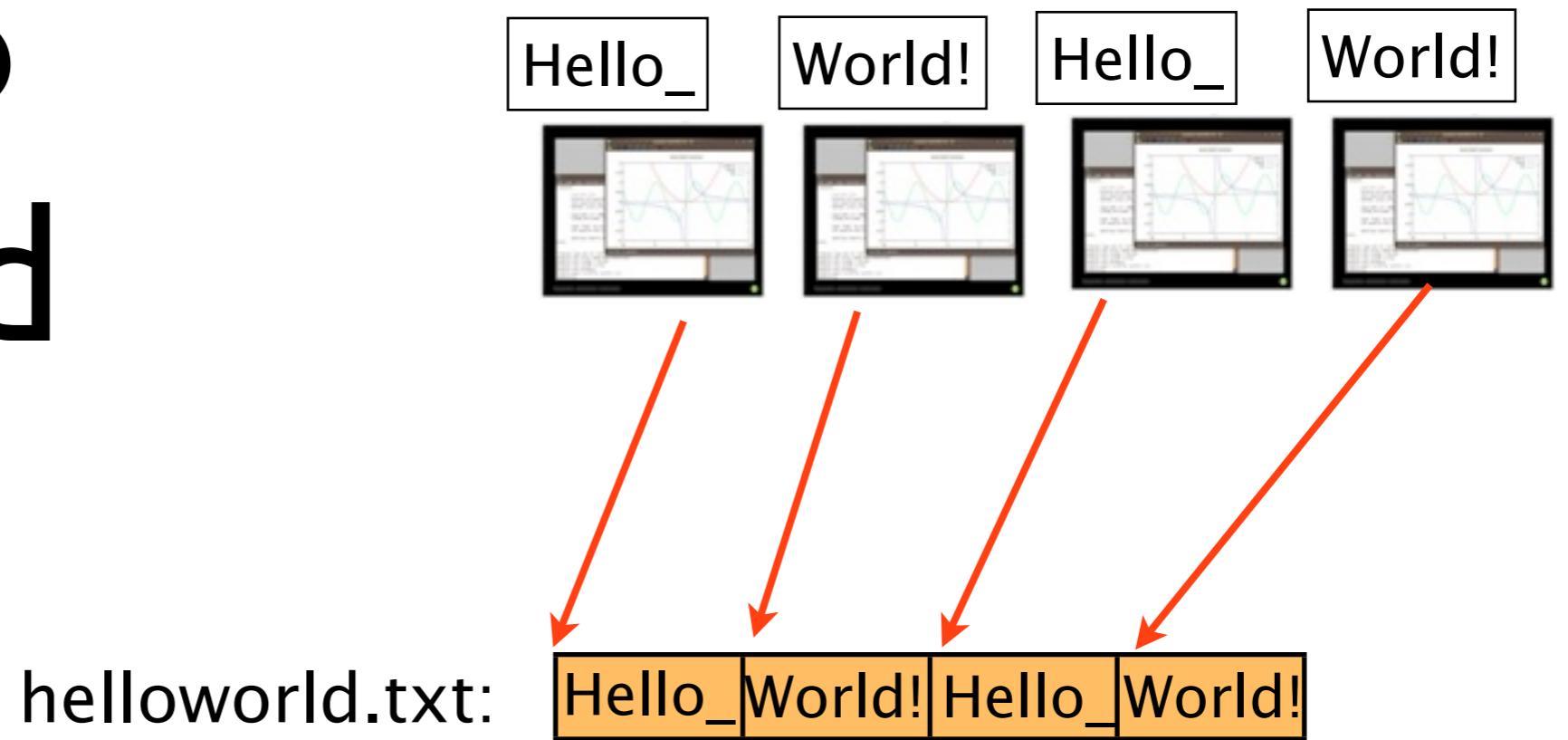
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MPI-IO

Hello World



```
MPI_File_write(file, message, msgsize, MPI_CHAR, &status);
```



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MPI-IO

Hello

World



helloworld.txt: Hello_World!Hello_World!

```
MPI_File_close(&file);
```



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MPI-IO

Hello

World

Usual MPI
startup/
teardown
boilerplate

```
#include <stdio.h>
#include <string.h>
#include <mpi.h>

int main(int argc, char **argv) {
    int ierr, rank, size;
    MPI_Offset offset;
    MPI_File file;
    MPI_Status status;
    const int msgsize=6;
    char message[msgsize+1];

    ierr = MPI_Init(&argc, &argv);
    ierr |= MPI_Comm_size(MPI_COMM_WORLD, &size);
    ierr |= MPI_Comm_rank(MPI_COMM_WORLD, &rank);

    if ((rank % 2) == 0) strcpy (message, "Hello ");
    else strcpy (message, "World");

    offset = (msgsize*rank);

    MPI_File_open(MPI_COMM_WORLD, "helloworld.txt", MPI_MODE_CREATE
                  MPI_INFO_NULL, &file);
    MPI_File_seek(file, offset, MPI_SEEK_SET);
    MPI_File_write(file, message, msgsize, MPI_CHAR, &status);
    MPI_File_close(&file);

    MPI_Finalize();
    return 0;
}
```



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MPI_File_Open

Communicator;
collective
operation.

```
call MPI_File_Open( integer      communicator,  
                   character(*) *filename,  
                   integer      mode,  
                   integer      info,  
                   integer      handle,  
                   integer      ierr);
```

```
int MPI_File_Open( MPI_Comm   communicator,  
                   char       *filename,  
                   int        mode,  
                   MPI_Info   info,  
                   MPI_File  *handle);
```



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MPI_File_Open

Info allows us to send extra hints to MPI-IO layer about file(performance tuning, special case handling)

```
call MPI_File_Open( integer      communicator,  
                   character(*) *filename,  
                   integer      mode,  
                   integer      info,  
                   integer      handle,  
                   integer      ierr);
```

```
int MPI_File_Open( MPI_Comm   communicator,  
                   char       *filename,  
                   int        mode,  
                   MPI_Info   info,  
                   MPI_File  *handle);
```

MPI_INFO_NULL: no extra info.



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Modes for MPI File Open():

<code>MPI_MODE_RDONLY</code>	read-only
<code>MPI_MODE_RDWR</code>	read-write
<code>MPI_MODE_WRONLY</code>	write-only
<code>MPI_MODE_CREATE</code>	Create if doesn't exist.
<code>MPI_MODE_APPEND</code>	On open, file pointers at end of file.
<code>MPI_MODE_EXCL</code>	Fail if try to create, does exist.
<code>MPI_MODE_UNIQUE_OPEN</code>	No one else is using this.
<code>MPI_MODE_SEQUENTIAL</code>	Will be sequential access only.
<code>MPI_MODE_DELETE_ON_CLOSE</code>	Delete when done. (OOC/scratch).



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MPI_File_Seek:

```
int MPI_File_seek( MPI_File mpi_fh,  
                   MPI_Offset offset,  
                   int mode);
```

```
call MPI_File_seek( integer :: mpi_fh,  
                    integer(mpi_offset_kind) :: offset,  
                    integer :: mode  
                    integer :: ierr)
```

MPI_SEEK_SET	Set file pointer to position <i>offset</i>
MPI_SEEK_CUR	$\text{pointer} \leftarrow \text{current position} + \text{offset}$
MPI_SEEK_END	$\text{pointer} \leftarrow \text{end of file} - \text{offset}$

Not collective; each adjusts its own local file pointer



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MPI_File_write:

```
int MPI_File_write(MPI_File fh,  
                   void *buf,  
                   int count,  
                   MPI_Datatype datatype,  
                   MPI_Status *status)
```

```
call MPI_File_write( integer :: mpi_fh,  
                     buffer,  
                     integer :: count  
                     integer :: datatype  
                     integer :: status(MPI_STATUS_SIZE)  
                     integer :: ierr)
```

Not collective; each writes.



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MPI_File_write:

- MPI File write is very much like a MPI_Send.
- “Sending” count of datatype from buf “to” the file.
- Here, writing 6 MPI_CHARs.
- Contiguous in memory starting in buffer.
- Status like a receive -- contains info about amount of data actually written, etc.

```
int MPI_File_write(MPI_File fh,  
                   void *buf,  
                   int count,  
                   MPI_Datatype datatype,  
                   MPI_Status *status)
```



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MPI_File_write:

- To write out data that is non-contiguous in memory, same as MPI_Sending non-contig data:
 - Create type that describes data layout in memory
 - “Send” in units of that type.
- Noncontiguous data in memory is written out contiguously to the file, starting at the current (local) file pointer.

stride = 2



blocklen = 1

count = 6

```
call MPI_Type_vector(count, blocklen, stride,  
MPI_CHARACTER,  
everyother, ierr)
```



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```
integer, parameter :: msgsize=6, strsize=12
character(strsize) :: message
integer :: everyother
```

```
!....
```

```
if (mod(rank,2) == 0) then
    message = "H@e#l*l^o* A"
else
    message = "WFoQr#l>d@!_"
endif
```

```
!...
```

```
call MPI_Type_vector(msgsize, 1, 2, MPI_CHARACTER, everyother, ierr)
call MPI_Type_commit(everyother, ierr)
```

```
call MPI_File_open(MPI_COMM_WORLD, "helloworld-nc.txt", ior(MPI_MODE_CREATE,MPI_MODE_WRONLY),&
                  MPI_INFO_NULL, fileno, ierr)
```

```
call MPI_File_seek (fileno, offset, MPI_SEEK_SET, ierr)
call MPI_File_write(fileno, message, 1, everyother, wstatus, ierr)
call MPI_File_close(fileno, ierr)
```

helloworld-noncontigf.f90



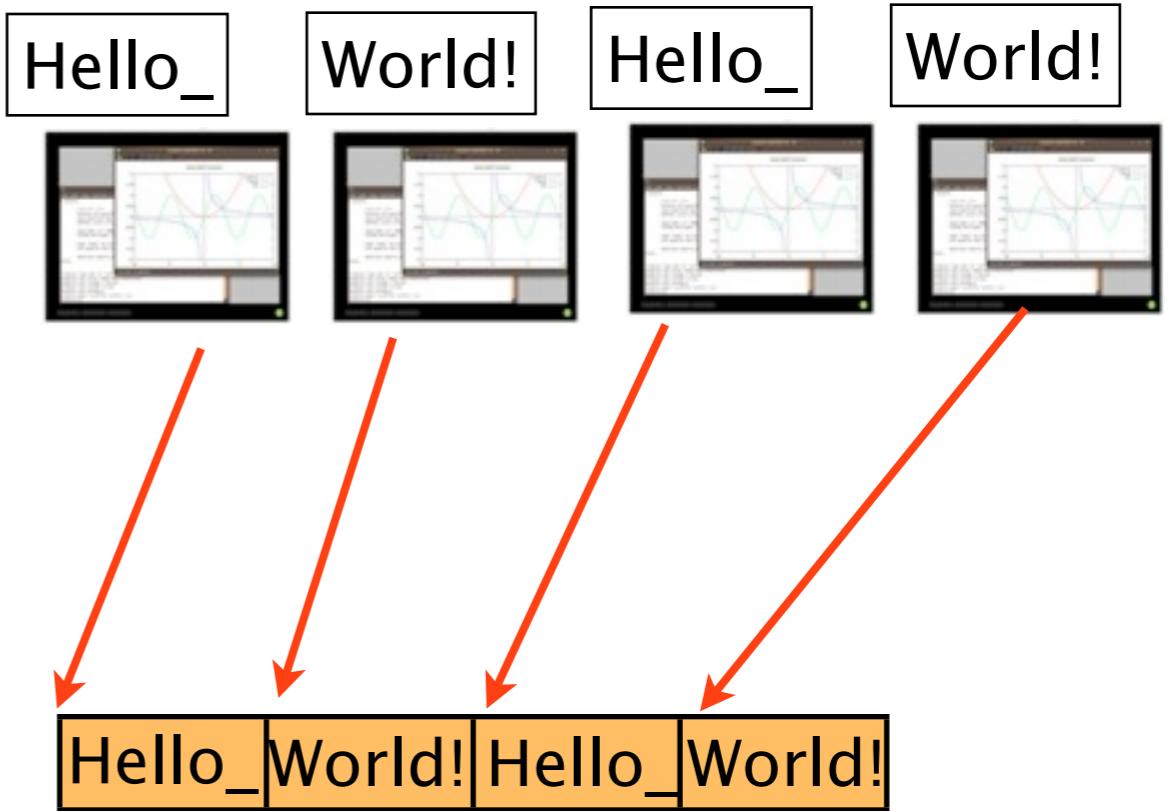
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Seek + Write

- Works, but:
 - Very low level (gets complicated for less trivial data layouts)
 - Completely independent operations (seek, write)
 - Hard for any middleware to coordinate, improve anything.
 - Could do this with POSIX I/O.



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MPI_File_write_at:

```
offset = rank*msgsize
```

```
call MPI_File_open(MPI_COMM_WORLD, "helloworld-at.txt",  
                   ior(MPI_MODE_CREATE,MPI_MODE_WRONLY), &  
                   MPI_INFO_NULL, fileno, ierr)  
  
call MPI_File_write_at(fileno, offset, message, msgsize, &  
                      MPI_CHARACTER, wstatus, ierr)  
call MPI_File_close(fileno, ierr)
```

writeatf.f90

```
offset = (msgsize*rank);
```

```
MPI_File_open(MPI_COMM_WORLD, "helloworld-at.txt",  
              MPI_MODE_CREATE|MPI_MODE_WRONLY,  
              MPI_INFO_NULL, &file);  
  
MPI_File_write_at(file, offset, message, msgsize, MPI_CHAR, &status);  
MPI_File_close(&file);
```

writeatc.c



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MPI_File_write_at:

```
int MPI_File_write_at(MPI_File fh,  
                      MPI_Offset offset ←  
                      void *buf,  
                      int count,  
                      MPI_Datatype datatype,  
                      MPI_Status *status)
```

```
call MPI_File_write_at( integer :: mpi_fh,  
                        integer(MPI_OFFSET_KIND) :: offset, ←  
                        buffer,  
                        integer :: count  
                        integer :: datatype  
                        integer :: status(MPI_STATUS_SIZE)  
                        integer :: ierr)
```

Writes at a given offset



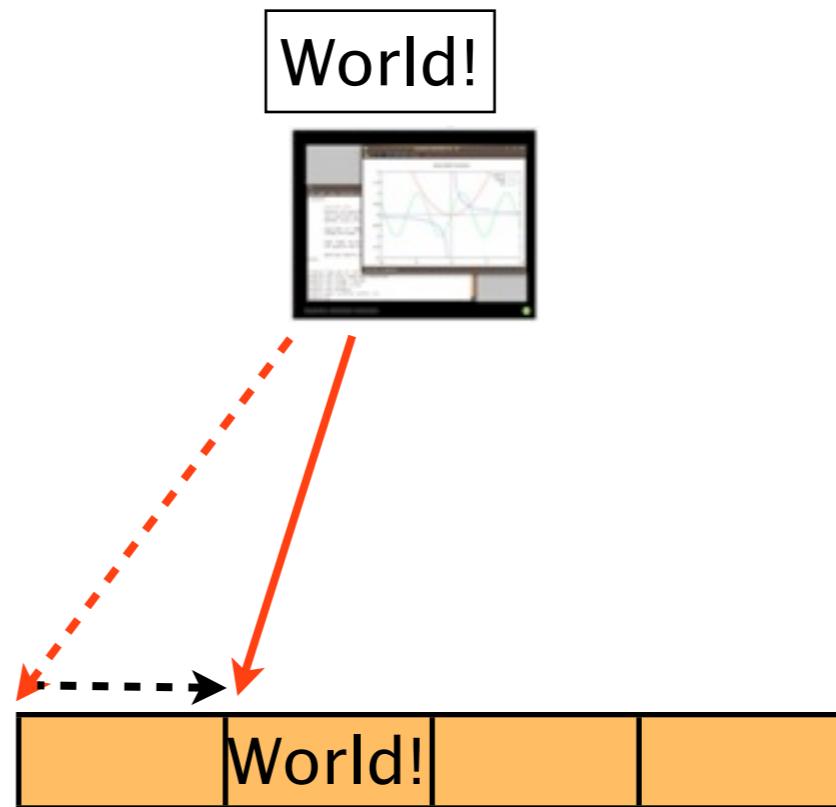
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Write_at

- Seek (relative to current, local, file pointer) + write in one operation.
- More explicit about what is going to happen - some opt. possible.
- But actions of individual processors still independent - no collective optimization possible.



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MPI_File_write_at_all:

```
offset = rank*msgsize  
  
call MPI_File_open(MPI_COMM_WORLD, "helloworld-at.txt", &  
                   ior(MPI_MODE_CREATE,MPI_MODE_WRONLY), &  
                   MPI_INFO_NULL, fileno, ierr)  
  
call MPI_File_write_at_all(fileno, offset, message, msgsize, &  
                           MPI_CHARACTER, wstatus, ierr)  
call MPI_File_close(fileno, ierr)
```

writeatallf.f90

```
offset = (msgsize*rank);  
  
MPI_File_open(MPI_COMM_WORLD, "helloworld-at.txt",  
              MPI_MODE_CREATE|MPI_MODE_WRONLY,  
              MPI_INFO_NULL, &file);  
  
MPI_File_write_at_all(file, offset, message, msgsize,  
                      MPI_CHAR, &status);  
MPI_File_close(&file);
```

writeatallc.c



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MPI_File_write_at_all:

```
int MPI_File_write_at_all(MPI_File fh,  
                           MPI_Offset offset  
                           void *buf,  
                           int count,  
                           MPI_Datatype datatype,  
                           MPI_Status *status)
```

```
call MPI_File_write_at_all( integer :: mpi_fh,  
                           integer(MPI_OFFSET_KIND) :: offset,  
                           buffer,  
                           integer :: count  
                           integer :: datatype  
                           integer :: status(MPI_STATUS_SIZE)  
                           integer :: ierr)
```

Writes at a given offset - Collective!



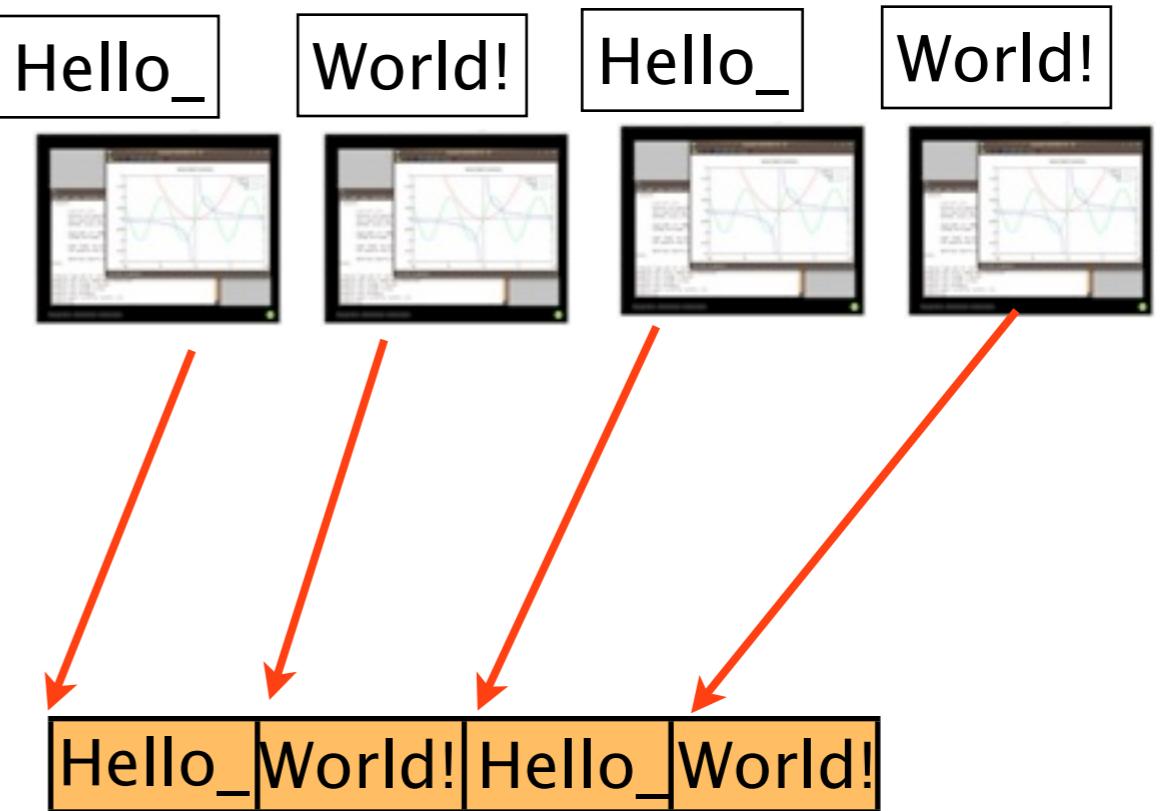
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Write_at_all

- Much more explicit about what is going to happen globally.
- Collective operation.
- All processors participate.
- Higher order optimizations possible.



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Non-contiguous in file

- Imagine having to write out a 2d file as to the right, with rank 0 “owning” the yellow data, etc.
- (eg, an image, or a complete checkpoint of a 2d domain).
- Would have to do repeated seeks, writing out one row at a time...

0	0	0	0	0	0	1	1	1	1	1	1
0	0	0	0	0	0	1	1	1	1	1	1
0	0	0	0	0	0	1	1	1	1	1	1
0	0	0	0	0	0	1	1	1	1	1	1
0	0	0	0	0	0	1	1	1	1	1	1
0	0	0	0	0	0	1	1	1	1	1	1
2	2	2	2	2	2	3	3	3	3	3	3
2	2	2	2	2	2	3	3	3	3	3	3
2	2	2	2	2	2	3	3	3	3	3	3
2	2	2	2	2	2	3	3	3	3	3	3



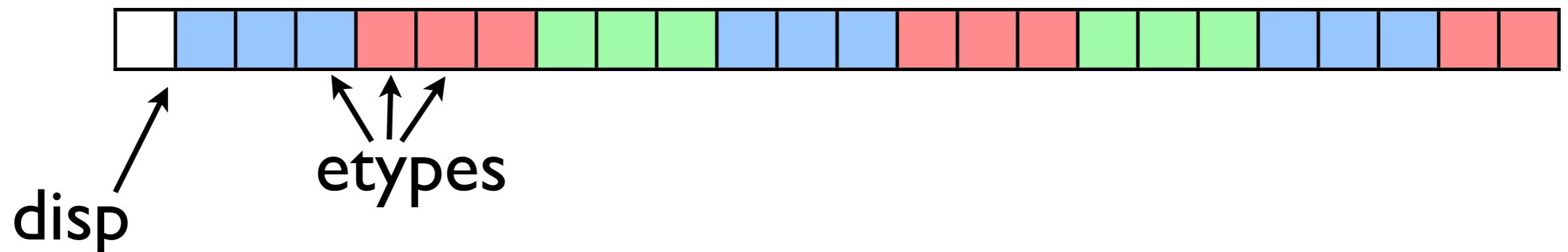
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MPI-IO File View

- int MPI_File_set_view(
 MPI_File fh,
 MPI_Offset disp,
 MPI_Datatype etype,
 MPI_Datatype filetype,
 char *datarep,
 MPI_Info info)
 /* displacement in bytes from start */
 /* elementary type */
 /* file type; prob different for each proc */
 /* ‘native’ or ‘internal’ */
 /* MPI_INFO_NULL */



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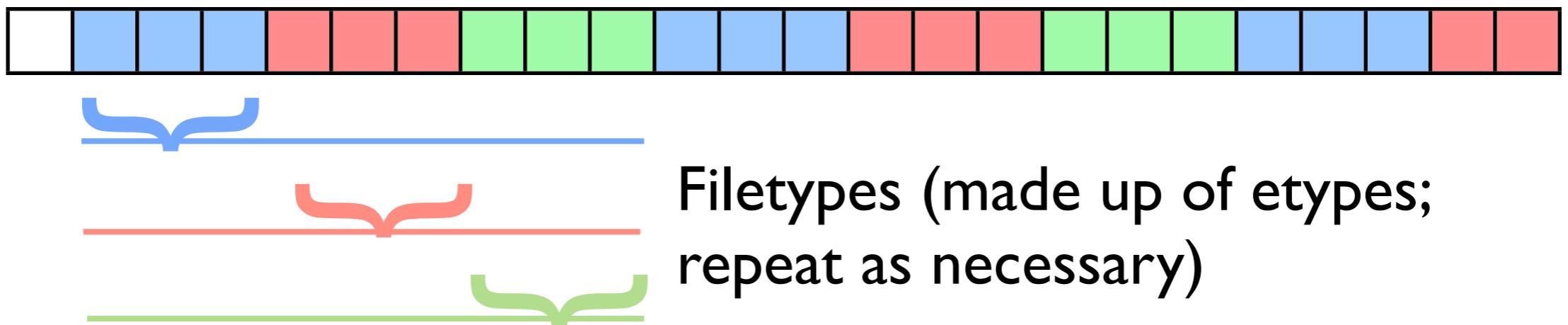


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MPI-IO File View

- ```
int MPI_File_set_view(
 MPI_File fh,
 MPI_Offset disp,
 MPI_Datatype etype,
 MPI_Datatype filetype,
 char *datarep,
 MPI_Info info)
```

/\* displacement in bytes from start \*/  
/\* elementary type \*/  
/\* file type; prob different for each proc \*/  
/\* 'native' or 'internal' \*/  
/\* MPI\_INFO\_NULL \*/



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# MPI-IO File Write

- `int MPI_File_write_all(  
 MPI_File fh,  
 void *buf,  
 int count,  
 MPI_Datatype datatype,  
 MPI_Status *status)`

Writes (`_all`: collectively) to part of file **within view.**



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# MPI\_Type\_create\_subarray

- `MPI_Type_create_subarray` ; piece of a multi-dimensional array.
- Much more convenient for higher-dimensional arrays
- (Otherwise, need vectors of vectors of vectors...)
- Here - starts = [0,0], subsizes=[5,5], sizes=[10,10].

|   |   |   |   |   |   |   |   |   |   |   |
|---|---|---|---|---|---|---|---|---|---|---|
| 0 | 0 | 0 | 0 | 0 | 0 | 1 | 1 | 1 | 1 | 1 |
| 0 | 0 | 0 | 0 | 0 | 0 | 1 | 1 | 1 | 1 | 1 |
| 0 | 0 | 0 | 0 | 0 | 0 | 1 | 1 | 1 | 1 | 1 |
| 0 | 0 | 0 | 0 | 0 | 0 | 1 | 1 | 1 | 1 | 1 |
| 0 | 0 | 0 | 0 | 0 | 0 | 1 | 1 | 1 | 1 | 1 |
| 2 | 2 | 2 | 2 | 2 | 2 | 3 | 3 | 3 | 3 | 3 |
| 2 | 2 | 2 | 2 | 2 | 2 | 3 | 3 | 3 | 3 | 3 |
| 2 | 2 | 2 | 2 | 2 | 2 | 3 | 3 | 3 | 3 | 3 |
| 2 | 2 | 2 | 2 | 2 | 2 | 3 | 3 | 3 | 3 | 3 |

```
int MPI_Type_create_subarray(
 int ndims, int *array_of_sizes,
 int *array_of_subsizes,
 int *array_of_starts,
 int order,
 MPI_Datatype oldtype,
 MPI_Datatype &newtype);

call MPI_Type_create_subarray(
 integer ndims, [array_of_sizes],
 [array_of_subsizes],
 [array_of_starts],
 order, oldtype,
 newtype, ierr)
```



## fileviewc.c

```
MPI_Type_create_subarray(2, globalsize, subsize, start, MPI_ORDER_C,
 MPI_CHAR, &viewtype);
MPI_Type_commit(&viewtype);

offset = 0;

MPI_File_open(MPI_COMM_WORLD, "viewtype.txt",
 MPI_MODE_CREATE|MPI_MODE_WRONLY,
 MPI_INFO_NULL, &file);

MPI_File_set_view(file, offset, MPI_CHAR, viewtype,
 "native", MPI_INFO_NULL);

MPI_File_write_all(file, &(mydata[0][0]), locnrows*locncols, MPI_CHAR, &status);
MPI_File_close(&file);
```



write locnrows\*locncols contiguous MPI\_CHARS to  
the (non-contiguous) view in file.



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# MPI-IO hands-on

- Fill in the blanks in sine.c or sinef.f90 to use MPI-IO to write out the 1-d array of  $\sin(x)$ .
- Already written: decomposing the array, doing the calculation, `MPI_File_open` and `MPI_file_close` calls.
- Make `sine` (`sinf`) and make `sineplot` (`sinefplot`) to build the code, and run it and plot the results.
- Can use any of the approaches above
- ~15 minutes.



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# Formats for Scientific Data Management

NetCDF4, HDF5, and whatnot



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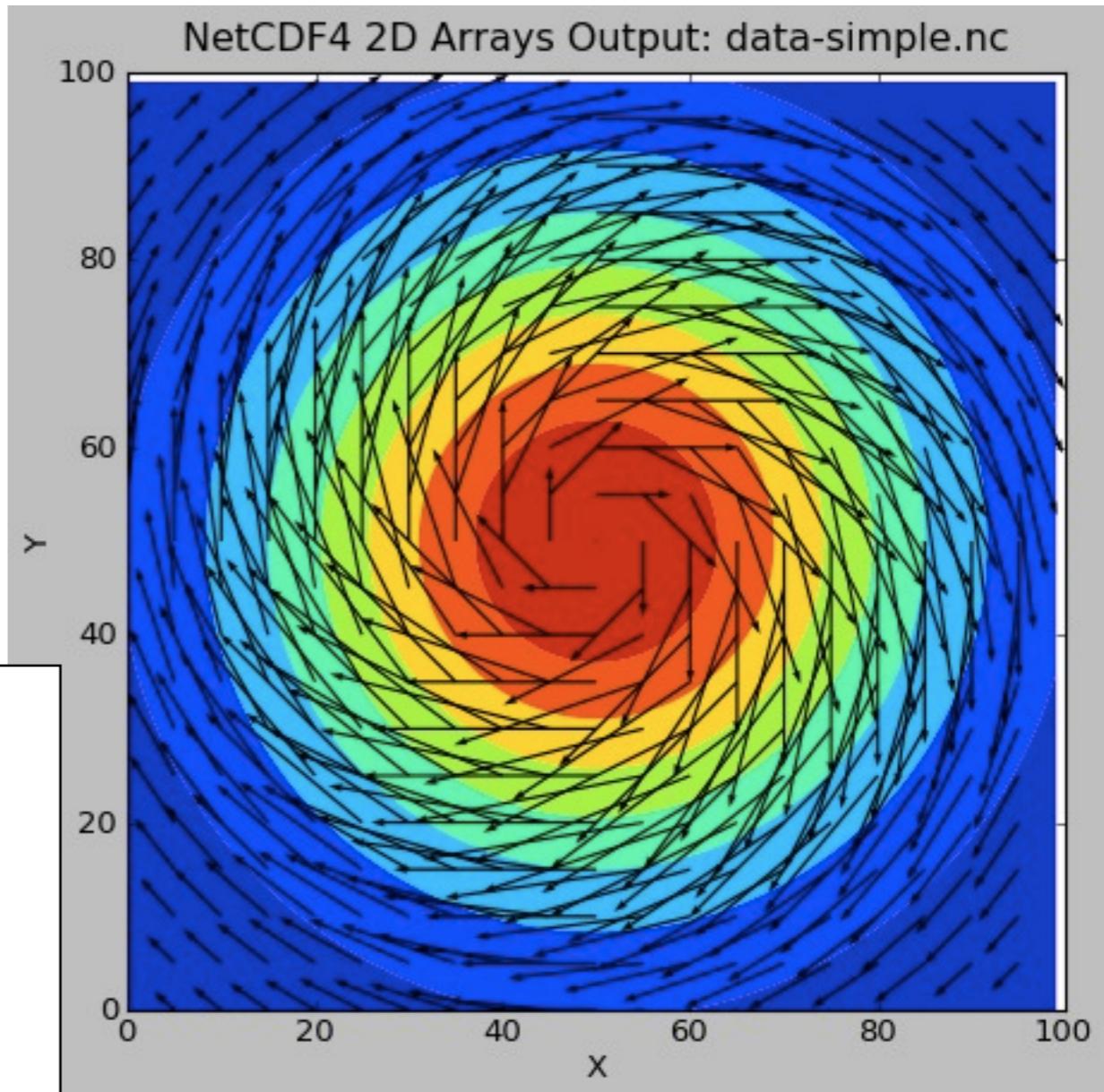
# Sample Code

```
$ cd parIO/netcdf

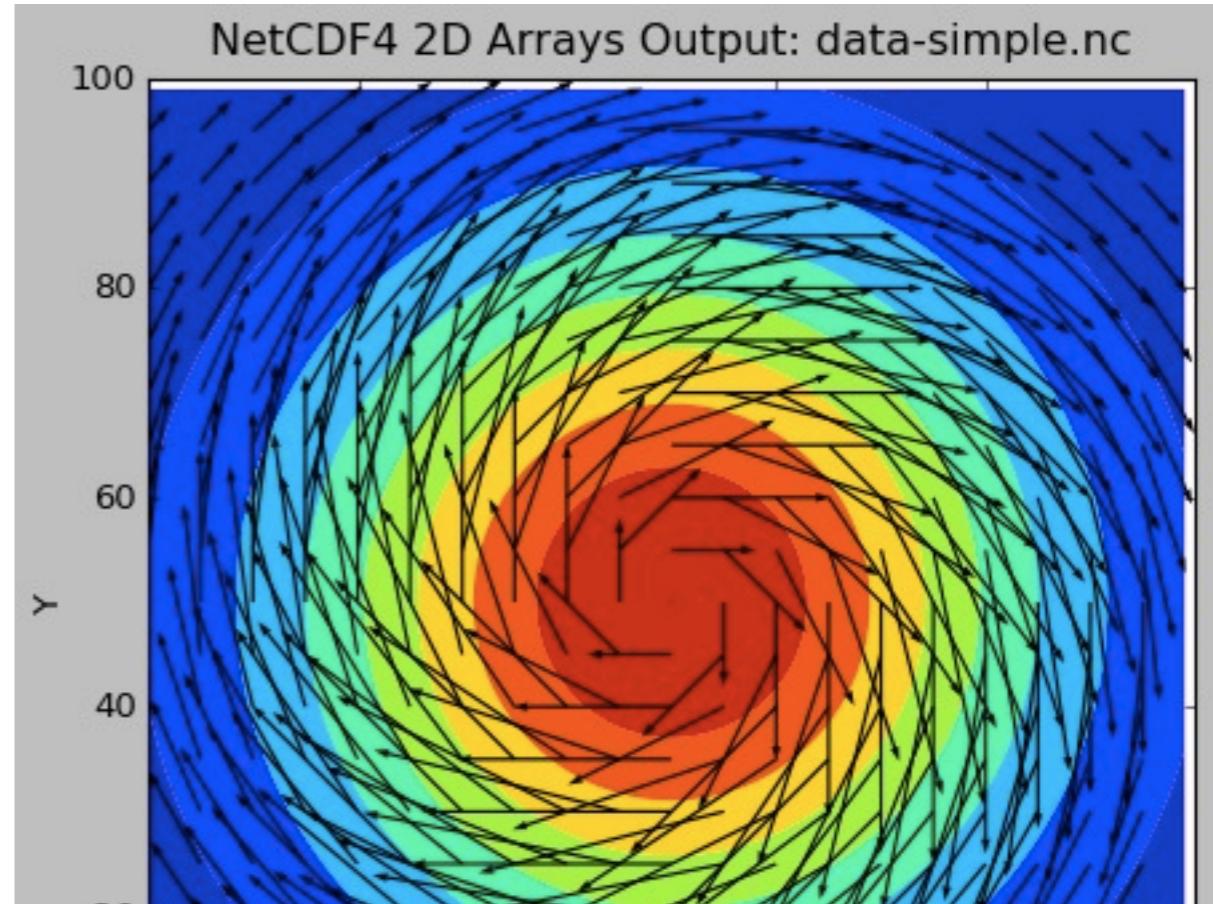
$ make 2darray-simple (C), or
$ make f2darray-simple (F90)

$./{f,}2darray-simple

$ ls *.nc
$../../plots.py *.nc
```



# Sample Code



```
$./2darray-simple --help
```

Options:

- nx=N (-x N): Set the number of grid cells in x direction.
- ny=N (-y N): Set the number of grid cells in y direction.
- filename=S (-f S): Set the output filename.

```
$./f2darray-simple --help
```

Usage: f2darray-simple [--help] [filename [nx [ny]]]

where filename is output filename, and

nx, ny are number of points in x and y directions.



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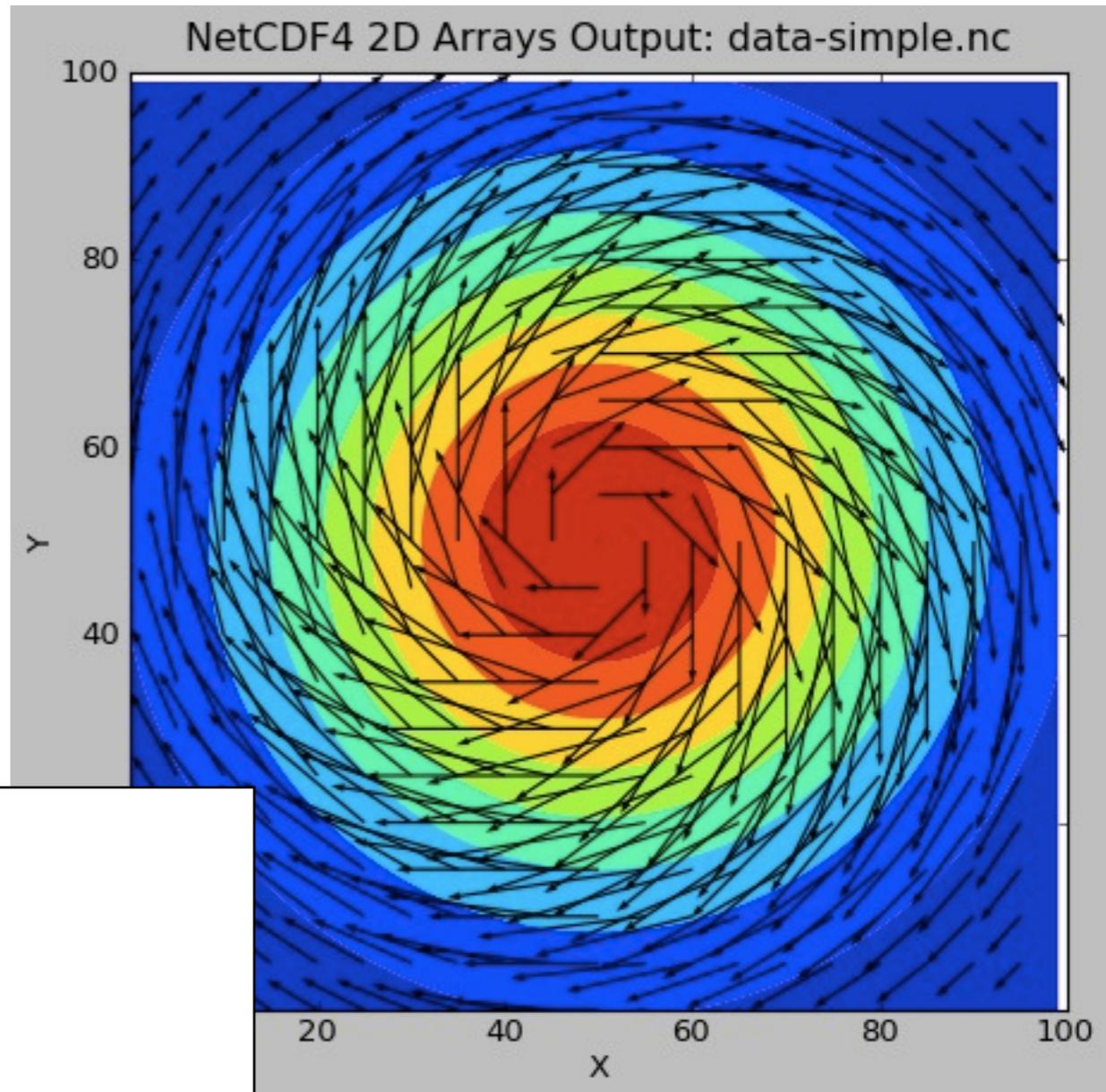
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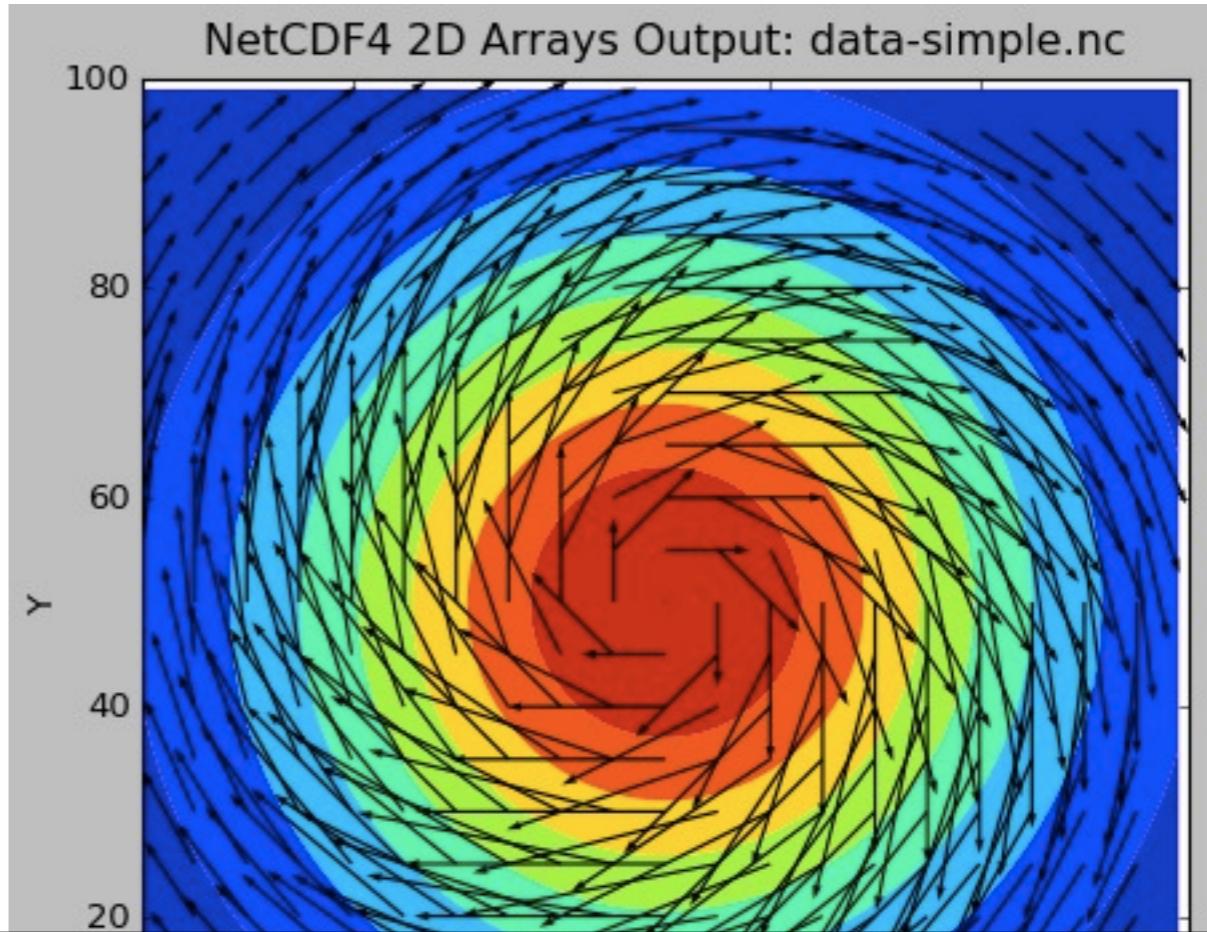
# What is this .nc file?

```
$ ncdump -h data-simple-fort.nc
netcdf data-simple-fort {
dimensions:
 X = 100 ;
 Y = 100 ;
 velocity\ components = 2 ;
variables:
 double Density(Y, X) ;
 double Velocity(Y, X, velocity\
```



# NetCDF

- NetCDF is a set of libraries and formats for:
  - portable,
  - efficient
  - “self-describing”
- way of storing and accessing large arrays (eg, for scientific data)
- Current version is NetCDF4



```
$ ncdump -h data-simple-fort.nc
netcdf data-simple-fort {
dimensions:
 X = 100 ;
 Y = 100 ;
 velocity\ components = 2 ;
variables:
 double Density(Y, X) ;
 double Velocity(Y, X, velocity\
components) ;
}
```



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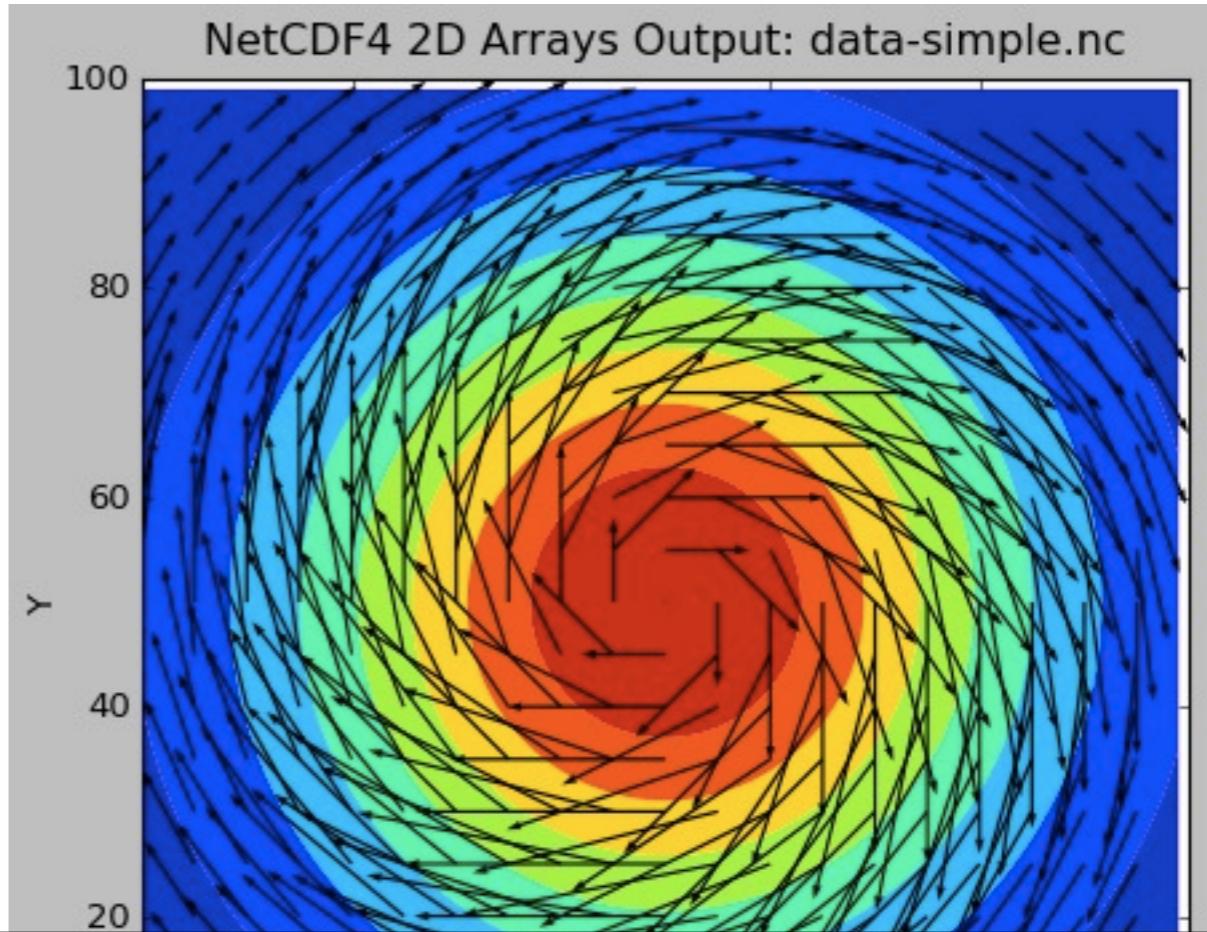
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# NetCDF: *Portable*

- Binary files, but common output format so that different sorts of machines can share files.
- Libraries accessible from C, C++, Fortran-77, Fortran 90/95/2003, python, etc.



```
$ ncdump -h data-simple-fort.nc
netcdf data-simple-fort {
dimensions:
 X = 100 ;
 Y = 100 ;
 velocity\ components = 2 ;
variables:
 double Density(Y, X) ;
 double Velocity(Y, X, velocity\ components) ;
}
```



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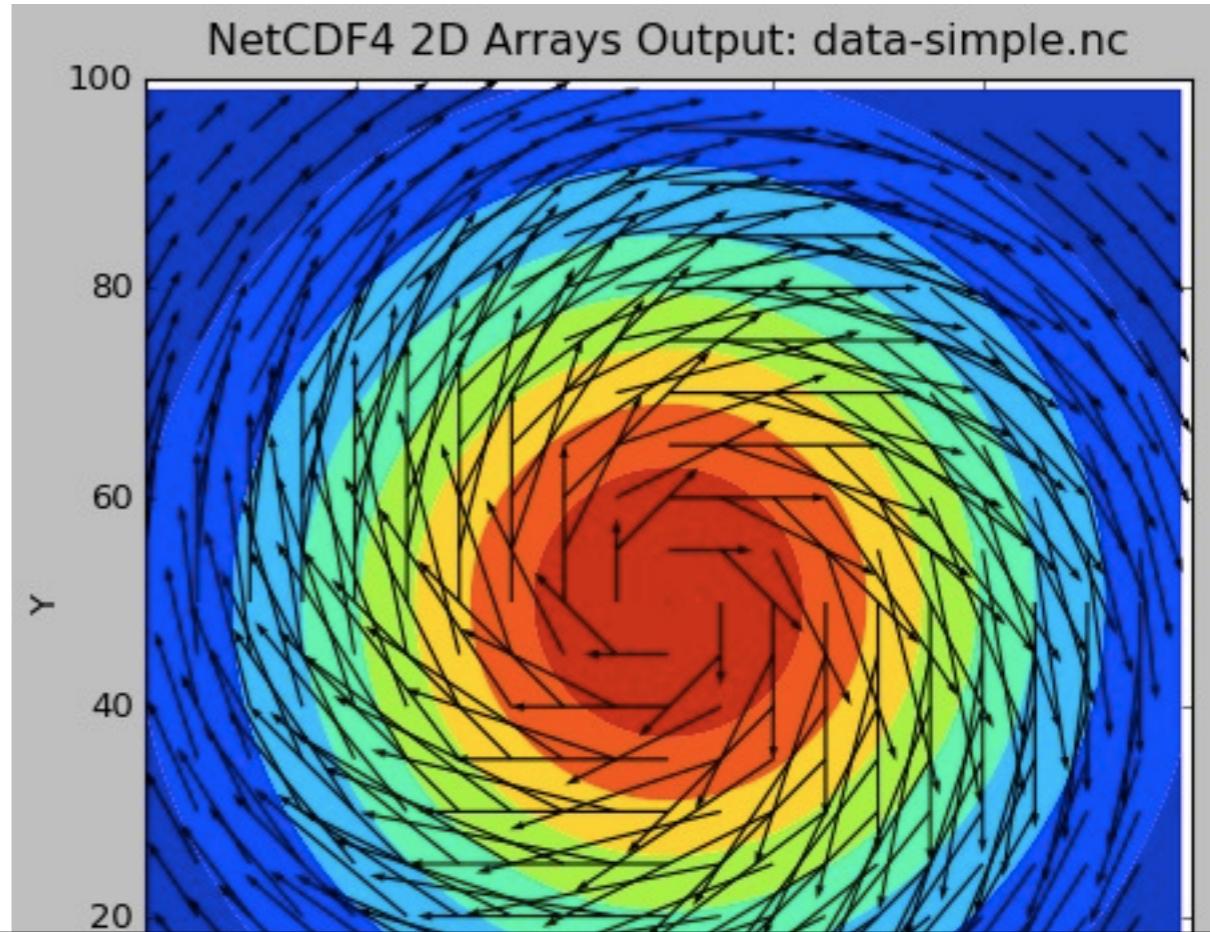
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# NetCDF: Self-Describing

- Header contains the metadata to describe the big data
- Lists:
  - Array names
  - Dimensions
  - *shared dimensions* - information about how the arrays relate
  - Other, related information



```
$ ncdump -h data-simple-fort.nc
netcdf data-simple-fort {
dimensions:
 X = 100 ;
 Y = 100 ;
 velocity\ components = 2 ;
variables:
 double Density(Y, X) ;
 double Velocity(Y, X, velocity\
components) ;
}
```



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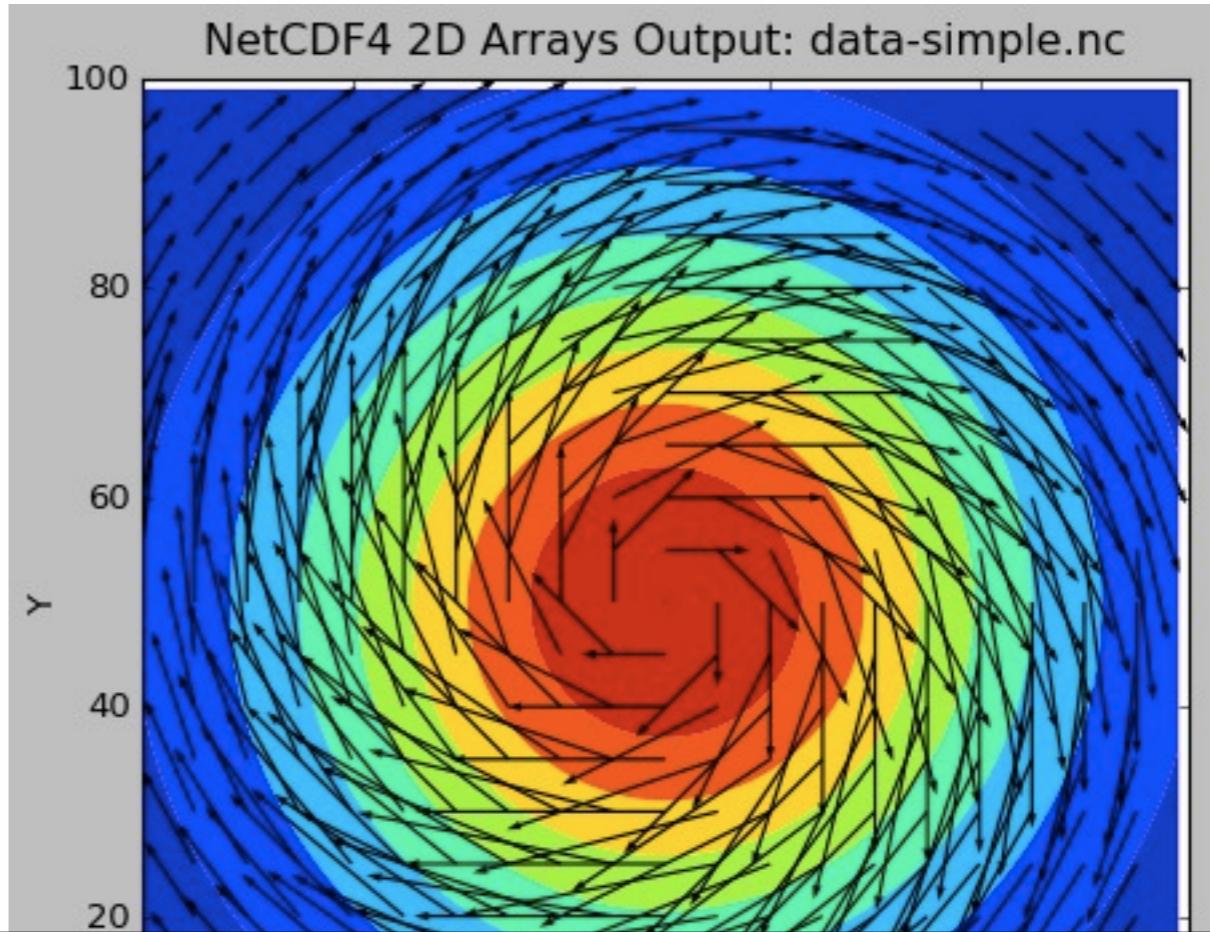
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# NetCDF: *Efficient*

- Binary, so less translation (as little is used as possible)
- IO libraries themselves are written for performance
- API, data format makes it easy to efficiently read, write subregions of arrays (slices, or ‘hyperslabs’)
- Still possible to make things slow
  - lots of metadata queries,



```
$ ncdump -h data-simple-fort.nc
netcdf data-simple-fort {
dimensions:
 X = 100 ;
 Y = 100 ;
 velocity\ components = 2 ;
variables:
 double Density(Y, X) ;
 double Velocity(Y, X, velocity\
components) ;
}
```

# 2darray-simple.c

```
#include "netcdf.h"
...

void writenetcdffile(rundata_t rundata, double **dens,
 double ***vel) {
 /* identifiers */
 int file_id;

 ...
 /* return status */
 int status;

 /* Create a new file - clobber anything existing */
 status = nc_create(rundata.filename, NC_CLOBBER, &file_id);
 /* netCDF routines return NC_NOERR on success */
 if (status != NC_NOERR) {
 fprintf(stderr, "Could not open file %s\n", rundata.filename);
 }
}
```



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Include function definitions

# 2darray-simple.c

```
#include "netcdf.h"

...
void writenetcdffile(rundata_t rundata, double **dens,
 double ***vel) {
 /* identifiers */
 int file_id;
...
 /* return status */
 int status;

 /* Create a new file, clobber anything existing */
 status = nc_create(rundata.filename, NC_CLOBBER, &file_id);
 /* netCDF routines return NC_NOERR on success */
 if (status != NC_NOERR) {
 fprintf(stderr, "Could not open file %s\n", rundata.filename);
```

Create a new file, with name  
rundata.filename



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# 2darray-simple.c

```
#include "netcdf.h"

...
void writenetcdffile(rundata_t rundata, double **dens,
 double ***vel) {
 /* identifiers */
 int file_id;
...
 /* return status */
 int status;

 /* Create a new file - clobber anything existing */
 status = nc_create(rundata.filename, NC_CLOBBER, &file_id);
 /* netCDF routines return NC_NOERR on success */
 if (status != NC_NOERR) {
 fprintf(stderr, "Could not open file %s\n", rundata.filename);
```

Clobber anything already in  
the file



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# 2darray-simple.c

```
#include "netcdf.h"

...
void writenetcdffile(rundata_t rundata, double **dens,
 double ***vel) {
 /* identifiers */
 int file_id;
...
 /* return status */
 int status;

 /* Create a new file - clobber anything existing */
 status = nc_create(rundata.filename, NC_CLOBBER, &file_id);
 /* netCDF routines return NC_NOERR on success */
 if (status != NC_NOERR) {
 fprintf(stderr, "Could not open file %s\n", rundata.filename);
```

Test the return codes



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# f2darray-simple.f90

```
subroutine writenetcdffile(rundata, dens, vel)
 use netcdf ←
 implicit none
 type(rundata_t), intent(IN) :: rundata
 double precision, intent(IN), dimension(:,:) :: dens
 double precision, intent(IN), dimension(:,:,:,:) :: vel

 integer :: file_id
 ...
 integer :: status

 ! create the file, check return code

 status = nf90_create(path=rundata%filename, cmode=NF90_CLOBBER,
ncid=file_id)
 if (status /= NF90_NOERR) then
 print *, 'Could not open file ', rundata%filename
 return
```

Import definitions



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# f2darray-simple.f90

```
subroutine writenetcdffile(rundata, dens, vel)

 use netcdf

 implicit none

 type(rundata_t), intent(IN) :: rundata
 double precision, intent(IN), dimension(:,:) :: dens
 double precision, intent(IN), dimension(:,:,:,:) :: vel

 integer :: file_id
 ...
 integer :: status

 ! create the file, check return code
 status = nf90_create(path=rundata%filename, cmode=NF90_CLOBBER,
ncid=file_id)
 if (status /= NF90_NOERR) then
 print *, 'Could not open file ', rundata%filename
 end if
 return
```

Create file



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# f2darray-simple.f90

```
subroutine writenetcdffile(rundata, dens, vel)
 use netcdf
 implicit none
 type(rundata_t), intent(IN) :: rundata
 double precision, intent(IN), dimension(:,:) :: dens
 double precision, intent(IN), dimension(:,:,:,:) :: vel
 integer :: file_id
 ...
 integer :: status
 ! create the file, check return code
 status = nf90_create(path=rundata%filename, cmode=NF90_CLOBBER,
 ncid=file_id)
 if (status /= NF90_NOERR) then
 print *, 'Could not open file ', rundata%filename
 end if
 return
```

C definitions are NC\_,  
F90 are NF90\_



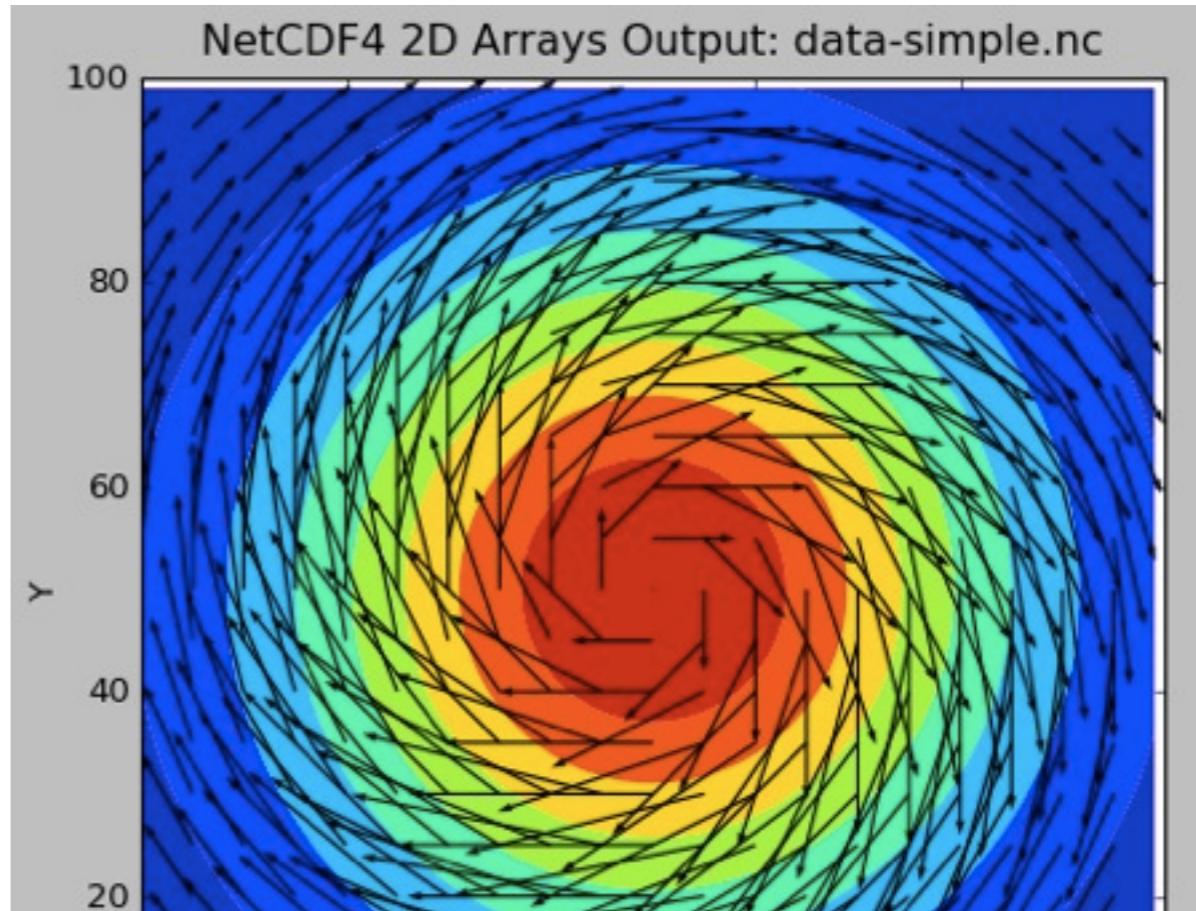
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# Writing a NetCDF File

- To write a NetCDF file, we go through the following steps:
  - **Create** the file (or open it for appending)
  - **Define dimensions** of the arrays we'll be writing
  - **Define variables** on those dimensions
  - **End definition** phase
  - **Write variables**
  - **Close file**



```
$ ncdump -h data-simple-fort.nc
netcdf data-simple-fort {
dimensions:
 X = 100 ;
 Y = 100 ;
 velocity\ components = 2 ;
variables:
 double Density(Y, X) ;
 double Velocity(Y, X, velocity\ components) ;
}
```



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# f2darray-simple.f90

```
integer :: file_id, xdim_id, ydim_id, vcomp_id
integer :: dens_id, vel_id
integer, dimension(2) :: densdims
integer, dimension(3) :: veldims
...

status = nf90_def_dim(file_id, 'x', rundata%nx, xdim_id)
status = nf90_def_dim(file_id, 'y', rundata%ny, ydim_id)
status = nf90_def_dim(file_id, 'velocity components', 2, vcomp_id)

densdims = (/ xdim_id, ydim_id /)
veldims = (/ vcomp_id, xdim_id, ydim_id /)

status = nf90_def_var(file_id, 'Density', NF90_DOUBLE, densdims, dens_id)
if (status /= NF90_NOERR) print *, trim(nf90_strerror(status)), ' Dens'
status = nf90_def_var(file_id, 'Velocity', NF90_DOUBLE, veldims, vel_id)

status = nf90_enddef(file_id)
```

Define the dimensions in the file: name, size, id



# f2darray-simple.f90

```
integer :: file_id, xdim_id, ydim_id, vcomp_id
integer :: dens_id, vel_id
integer, dimension(2) :: densdims
integer, dimension(3) :: veldims

...

status = nf90_def_dim(file_id, 'x', rundata%nx, xdim_id)
status = nf90_def_dim(file_id, 'Y', rundata%ny, ydim_id)
status = nf90_def_dim(file_id, 'velocity components', 2, vcomp_id)
```

```
densdims = (/ xdim_id, ydim_id /)
```

```
veldims = (/ vcomp_id, xdim_id, ydim_id /)
```

```
status = nf90_def_var(file_id, 'Density', NF90_DOUBLE, densdims, dens_id)
```

```
if (status /= NF90_NOERR) print *, trim(nf90_strerror(status)), ' Dens'
```

```
status = nf90_def_var(file_id, 'Velocity', NF90_DOUBLE, veldims, vel_id)
```

```
status = nf90_enddef(file_id)
```

Variables are defined in terms of these dims

# f2darray-simple.f90

```
status = nf90_enddef(file_id) ←

! Write out the values

status = nf90_put_var(file_id, dens_id, dens)
if (status /= NF90_NOERR) print *, trim(nf90_strerror(status)), ' Dens'
status = nf90_put_var(file_id, vel_id, vel)
if (status /= NF90_NOERR) print *, trim(nf90_strerror(status)), ' Vel'

status = nf90_close(file_id)
```

Once you're done defining things,

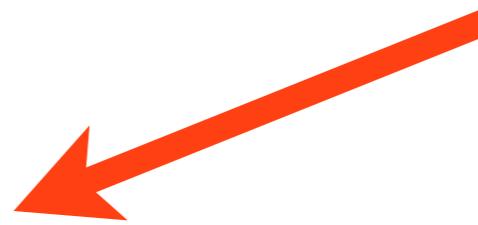
# f2darray-simple.f90

Writing data is easy.

```
status = nf90_enddef(file_id)

! Write out the values
status = nf90_put_var(file_id, dens_id, dens)
if (status /= NF90_NOERR) print *, trim(nf90_strerror(status)), ' Dens'
status = nf90_put_var(file_id, vel_id, vel)
if (status /= NF90_NOERR) print *, trim(nf90_strerror(status)), ' Vel'

status = nf90_close(file_id)
```



# f2darray-simple.f90

```
status = nf90_enddef(file_id)

! Write out the values
status = nf90_put_var(file_id, dens_id, dens)
if (status /= NF90_NOERR) print *, trim(nf90_strerror(status)), ' Dens'
status = nf90_put_var(file_id, vel_id, vel)
if (status /= NF90_NOERR) print *, trim(nf90_strerror(status)), ' Vel'

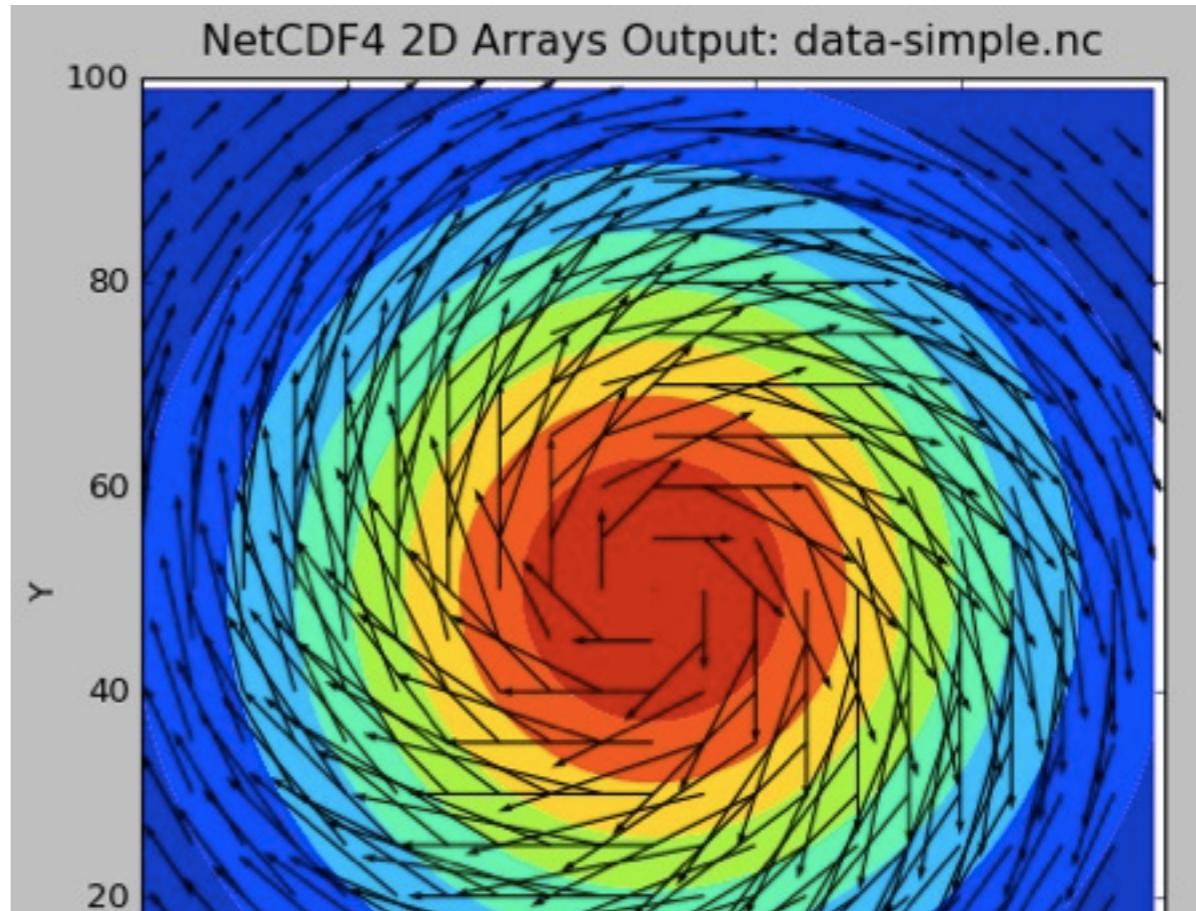
status = nf90_close(file_id)
```



**Closing the file is  
important!!**

# Reading a NetCDF File

- Flow is slightly different
  - **Open** the file for reading
  - **Get dimension ids** of the dimensions in the files
  - **Get dimension lengths** so you can allocate the files
  - **Get variable ids** so you can access the data
  - **Read variables**
  - **Close file**



```
$ ncdump -h data-simple-fort.nc
netcdf data-simple-fort {
dimensions:
 x = 100 ;
 y = 100 ;
 velocity\ components = 2 ;
variables:
 double Density(y, x) ;
 double Velocity(y, x, velocity\ components) ;
}
```



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# fread2darray-simple.f90

```
status = nf90_open(path=rundata%filename, mode=NF90_NOWRITE, ncid=file_id)
...
! find the dimensions
status = nf90_inq_dimid(file_id, 'X', xdim_id)
status = nf90_inq_dimid(file_id, 'Y', ydim_id)
status = nf90_inq_dimid(file_id, 'velocity components', vcomp_id)
! find the dimension lengths
status = nf90_inquire_dimension(file_id, xdim_id, len = rundata % nx)
status = nf90_inquire_dimension(file_id, ydim_id, len = rundata % ny)
status = nf90_inquire_dimension(file_id, vcomp_id, len = rundata % nvelcomp)
! now we can allocate variable sizes
allocate(dens(rundata%nx, rundata%ny)) !....etc...
status = nf90_inq_varid(file_id, 'Density', dens_id)
status = nf90_inq_varid(file_id, 'Velocity', vel_id)

status = nf90_get_var(file_id, dens_id, dens)
status = nf90_get_var(file_id, vel_id, vel)

status = nf90_close(file_id)
```

# fread2darray-simple.f90

```
status = nf90_open(path=rundata%filename, mode=NF90_NOWRITE, ncid=file_id)
...
! find the dimensions
status = nf90_inq_dimid(file_id, 'x', xdim_id)
status = nf90_inq_dimid(file_id, 'y', ydim_id)
status = nf90_inq_dimid(file_id, 'velocity components', vcomp_id)
! find the dimension lengths
status = nf90_inquire_dimension(file_id, xdim_id, len = rundata % nx)
status = nf90_inquire_dimension(file_id, ydim_id, len = rundata % ny)
status = nf90_inquire_dimension(file_id, vcomp_id, len = rundata % nvelcomp)
! now we can allocate variable sizes
allocate(dens(rundata%nx, rundata%ny)) !....etc...
status = nf90_inq_varid(file_id, 'Density', dens_id)
status = nf90_inq_varid(file_id, 'Velocity', vel_id)

status = nf90_get_var(file_id, dens_id, dens)
status = nf90_get_var(file_id, vel_id, vel)

status = nf90_close(file_id)
```

# fread2darray-simple.f90

```
status = nf90_open(path=rundata%filename, mode=NF90_NOWRITE, ncid=file_id)
...
! find the dimensions
status = nf90_inq_dimid(file_id, 'X', xdim_id)
status = nf90_inq_dimid(file_id, 'Y', ydim_id)
status = nf90_inq_dimid(file_id, 'velocity components', vcomp_id)
! find the dimension lengths
status = nf90_inquire_dimension(file_id, xdim_id, len = rundata % nx)
status = nf90_inquire_dimension(file_id, ydim_id, len = rundata % ny)
status = nf90_inquire_dimension(file_id, vcomp_id, len = rundata % nvelcomp)
! now we can allocate variable sizes
allocate(dens(rundata%nx, rundata%ny)) !....etc...
status = nf90_inq_varid(file_id, 'Density', dens_id)
status = nf90_inq_varid(file_id, 'Velocity', vel_id)

status = nf90_get_var(file_id, dens_id, dens)
status = nf90_get_var(file_id, vel_id, vel)

status = nf90_close(file_id)
```

# fread2darray-simple.f90

```
status = nf90_open(path=rundata%filename, mode=NF90_NOWRITE, ncid=file_id)
...
! find the dimensions
status = nf90_inq_dimid(file_id, 'X', xdim_id)
status = nf90_inq_dimid(file_id, 'Y', ydim_id)
status = nf90_inq_dimid(file_id, 'velocity components', vcomp_id)
! find the dimension lengths
status = nf90_inquire_dimension(file_id, xdim_id, len = rundata % nx)
status = nf90_inquire_dimension(file_id, ydim_id, len = rundata % ny)
status = nf90_inquire_dimension(file_id, vcomp_id, len = rundata % nvelcomp)
! now we can allocate variable sizes
allocate(dens(rundata%nx, rundata%ny)) !....etc...
status = nf90_inq_varid(file_id, 'Density', dens_id)
status = nf90_inq_varid(file_id, 'Velocity', vel_id)

status = nf90_get_var(file_id, dens_id, dens)
status = nf90_get_var(file_id, vel_id, vel)

status = nf90_close(file_id)
```

# fread2darray-simple.f90

```
status = nf90_open(path=rundata%filename, mode=NF90_NOWRITE, ncid=file_id)
...
! find the dimensions
status = nf90_inq_dimid(file_id, 'X', xdim_id)
status = nf90_inq_dimid(file_id, 'Y', ydim_id)
status = nf90_inq_dimid(file_id, 'velocity components', vcomp_id)
! find the dimension lengths
status = nf90_inquire_dimension(file_id, xdim_id, len = rundata % nx)
status = nf90_inquire_dimension(file_id, ydim_id, len = rundata % ny)
status = nf90_inquire_dimension(file_id, vcomp_id, len = rundata % nvelcomp)
! now we can allocate variable sizes
allocate(dens(rundata%nx, rundata%ny)) !....etc...
status = nf90_inq_varid(file_id, 'Density', dens_id)
status = nf90_inq_varid(file_id, 'Velocity', vel_id)

status = nf90_get_var(file_id, dens_id, dens)
status = nf90_get_var(file_id, vel_id, vel)

status = nf90_close(file_id)
```

# fread2darray-simple.f90

```
status = nf90_open(path=rundata%filename, mode=NF90_NOWRITE, ncid=file_id)
...
! find the dimensions
status = nf90_inq_dimid(file_id, 'X', xdim_id)
status = nf90_inq_dimid(file_id, 'Y', ydim_id)
status = nf90_inq_dimid(file_id, 'velocity components', vcomp_id)
! find the dimension lengths
status = nf90_inquire_dimension(file_id, xdim_id, len = rundata % nx)
status = nf90_inquire_dimension(file_id, ydim_id, len = rundata % ny)
status = nf90_inquire_dimension(file_id, vcomp_id, len = rundata % nvelcomp)
! now we can allocate variable sizes
allocate(dens(rundata%nx, rundata%ny)) !....etc...

status = nf90_inq_varid(file_id, 'Density', dens_id)
status = nf90_inq_varid(file_id, 'Velocity', vel_id)

status = nf90_get_var(file_id, dens_id, dens)
status = nf90_get_var(file_id, vel_id, vel)

status = nf90_close(file_id)
```

# read2darray-simple.c

```
status = nc_open(rundata->filename, NC_NOWRITE, &file_id);

/* Get the dimensions */

status = nc_inq_dimid(file_id, "X", &xdim_id);
if (status != NC_NOERR) fprintf(stderr, "Could not get X\n");
status = nc_inq_dimid(file_id, "Y", &ydim_id);
status = nc_inq_dimid(file_id, "velocity component", &vcomp_id);

status = nc_inq_dimlen(file_id, xdim_id, &(rundata->nx));
status = nc_inq_dimlen(file_id, ydim_id, &(rundata->ny));
status = nc_inq_dimlen(file_id, vcomp_id, &(rundata->nveldims));

...
nc_inq_varid(file_id, "Density", &dens_id);
nc_inq_varid(file_id, "Velocity", &vel_id);

nc_get_var_double(file_id, dens_id, &(*dens)[0][0]);
nc_get_var_double(file_id, vel_id, &(*vel)[0][0][0]);

nc_close(file_id);
```

# A Better example

- The above example is much more austere than a typical NetCDF file
- A more typical example is given in `2darray` (or `f2darray`)
- make this, then run it
- `./plots.py data.nc`
- (Same options as previous example)

```
$ ncdump -h data.nc
netcdf data {
dimensions:
 X = 100 ;
 Y = 100 ;
 velocity\ component = 2 ;
variables:
 float X\ coordinate(X) ;
 X\ coordinate:units = "cm" ;
 float Y\ coordinate(Y) ;
 Y\ coordinate:units = "cm" ;
 double Density(X, Y) ;
 Density:units = "g/cm^3" ;
 double Velocity(velocity\
component, X, Y) ;
 Velocity:units = "cm/s" ;
}
```



# 2darray.c

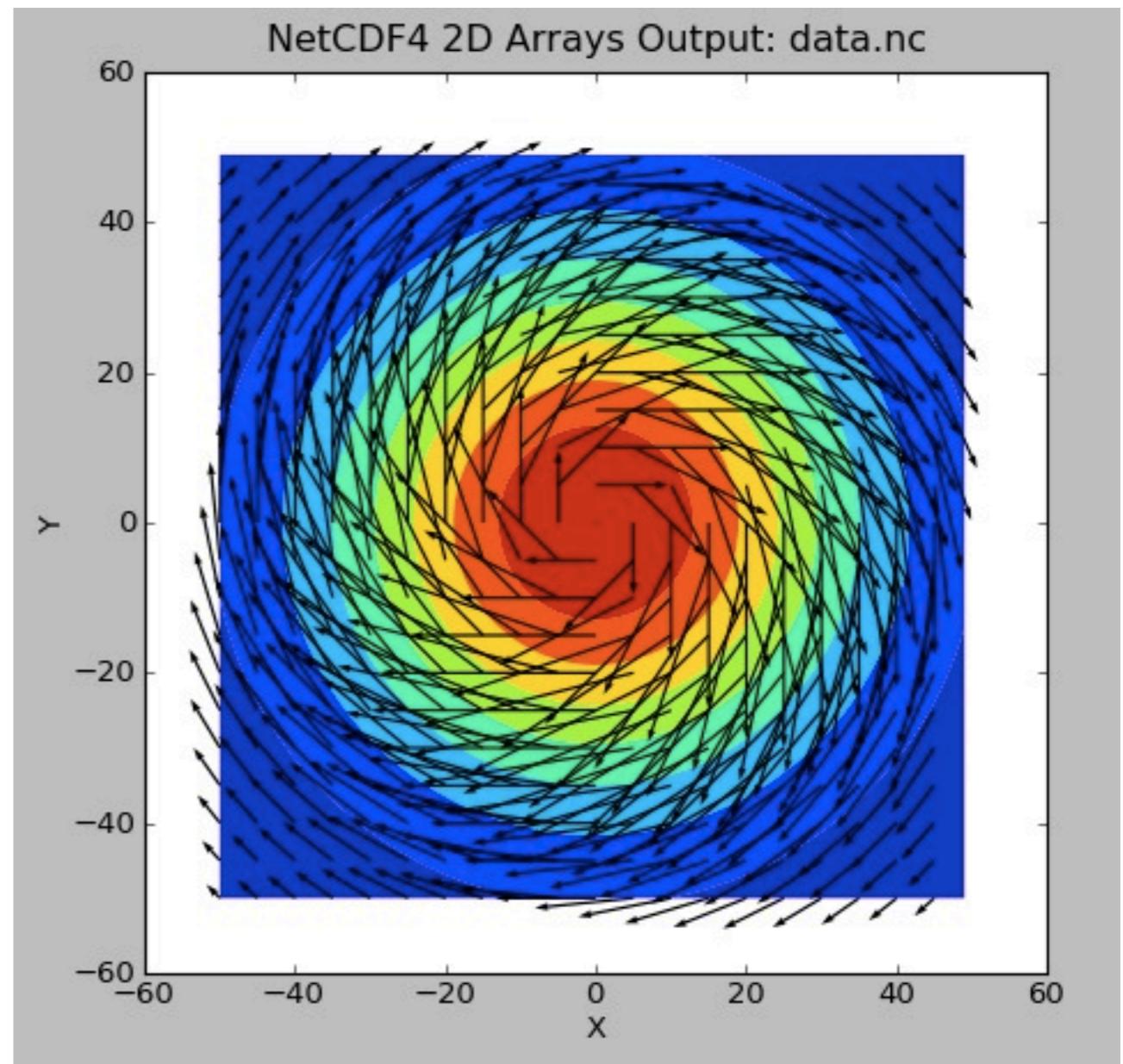
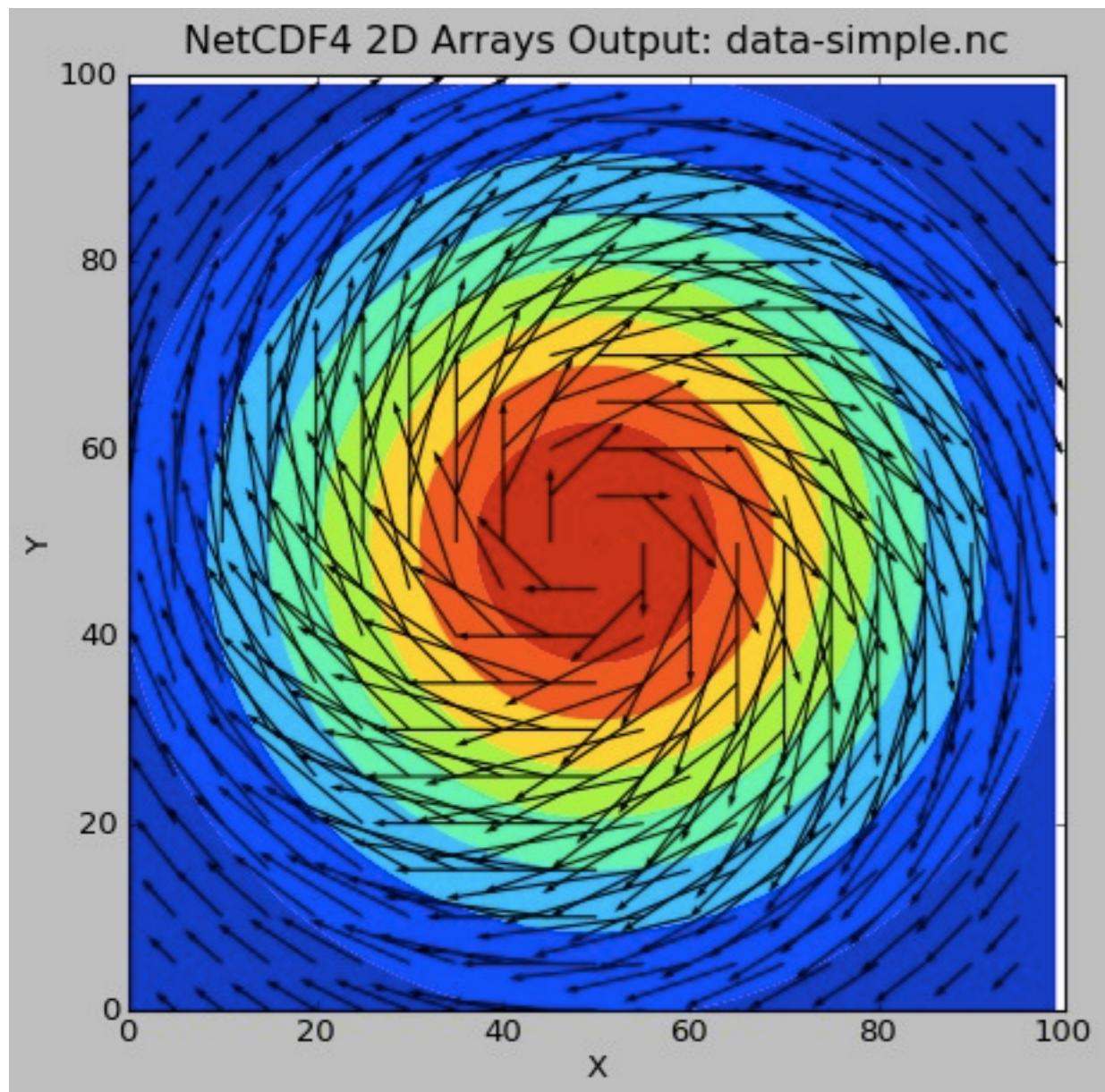
```
float *x, *y;
const char *coordunit="cm";
...
for (i=0; i<rundata.nx; i++) x[i] = (1.*i-r
for (i=0; i<rundata.ny; i++) y[i] = (1.*i-r
...
/* define the dimensions */
nc_def_dim(file_id, "X", rundata.nx, &xdim_id);
nc_def_dim(file_id, "Y", rundata.ny, &ydim_id);
nc_def_dim(file_id, "velocity component", 2, &vcomp_id);

/* define the coordinate variables,... */
nc_def_var(file_id, "X coordinate", NC_FLOAT, 1, &xdim_id, &xcoord_id);
nc_def_var(file_id, "Y coordinate", NC_FLOAT, 1, &ydim_id, &ycoord_id);

/* ...and assign units to them as an attribute */
nc_put_att_text(file_id, xcoord_id, "units", strlen(coordunit), coordunit);
nc_put_att_text(file_id, ycoord_id, "units", strlen(coordunit), coordunit);
```

Typically not only  
define dimensions but  
give coordinate values





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# 2darray.c

```
float *x, *y;
const char *coordunit="cm";
...
for (i=0; i<rundata.nx; i++) x[i] = (1.*i-r
for (i=0; i<rundata.ny; i++) y[i] = (1.*i-r
...
/* define the dimensions */
nc_def_dim(file_id, "X", rundata.nx, &xdim_id);
nc_def_dim(file_id, "Y", rundata.ny, &ydim_id);
nc_def_dim(file_id, "velocity component", 2, &vcomp_id);

/* define the coordinate variables,... */
nc_def_var(file_id, "X coordinate", NC_FLOAT, 1, &xdim_id, &xcoord_id);
nc_def_var(file_id, "Y coordinate", NC_FLOAT, 1, &ydim_id, &ycoord_id);

/* ...and assign units to them as an attribute */
nc_put_att_text(file_id, xcoord_id, "units", strlen(coordunit), coordunit);
nc_put_att_text(file_id, ycoord_id, "units", strlen(coordunit), coordunit);
```

Variables (or anything else) can have **attributes**:  
Name, and arbitrary data



# NetCDF Attributes

- Any NetCDF object (data set, dimension) can have an arbitrary number of attributes associated with it
- Name, and any type or size...
- Like a variable! (But can't access only part of it).

```
$ ncdump -h data.nc
netcdf data {
dimensions:
 X = 100 ;
 Y = 100 ;
 velocity\ component = 2 ;
variables:
 float X\ coordinate(X) ;
 X\ coordinate:units = "cm" ;
 float Y\ coordinate(Y) ;
 Y\ coordinate:units = "cm" ;
 double Density(X, Y) ;
 Density:units = "g/cm^3" ;
 double Velocity(velocity\
component, X, Y) ;
 Velocity:units = "cm/s" ;
}
```



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# NetCDF Attributes

- Attributes are assumed to be “small”, though.
- Stored in header information (not with big data)
- Don’t put large arrays in there

```
$ ncdump -h data.nc
netcdf data {
dimensions:
 X = 100 ;
 Y = 100 ;
 velocity\ component = 2 ;
variables:
 float X\ coordinate(X) ;
 X\ coordinate:units = "cm" ;
 float Y\ coordinate(Y) ;
 Y\ coordinate:units = "cm" ;
 double Density(X, Y) ;
 Density:units = "g/cm^3" ;
 double Velocity(velocity\
component, X, Y) ;
 Velocity:units = "cm/s" ;
}
```



# NetCDF Attributes

- Units are particularly useful attributes, as if a code needs data in some other units (MKS), can convert.

```
$ ncdump -h data.nc
netcdf data {
dimensions:
 X = 100 ;
 Y = 100 ;
 velocity\ component = 2 ;
variables:
 float X\ coordinate(X) ;
 X\ coordinate:units = "cm" ;
 float Y\ coordinate(Y) ;
 Y\ coordinate:units = "cm" ;
 double Density(X, Y) ;
 Density:units = "g/cm^3" ;
 double Velocity(velocity\
component, X, Y) ;
 Velocity:units = "cm/s" ;
}
```



# Limits to Self-Description

- But what if some codes expect “centimetre” and you use cm?
- Or their code uses “Dens” or “Rho” and yours uses “Density?” Or uses momentum rather than velocity?

```
$ ncdump -h data.nc
netcdf data {
dimensions:
 X = 100 ;
 Y = 100 ;
 velocity\ component = 2 ;
variables:
 float X\ coordinate(X) ;
 X\ coordinate:units = "cm" ;
 float Y\ coordinate(Y) ;
 Y\ coordinate:units = "cm" ;
 double Density(X, Y) ;
 Density:units = "g/cm^3" ;
 double Velocity(velocity\
component, X, Y) ;
 Velocity:units = "cm/s" ;
}
```



# Conventions

- There are lists of conventions that you can follow for variable names, unit names, etc.
- If you are planning for interoperability with other codes, this is the way to go
- Codes expecting data following (say) CF conventions for geophys should recognize data in that convention



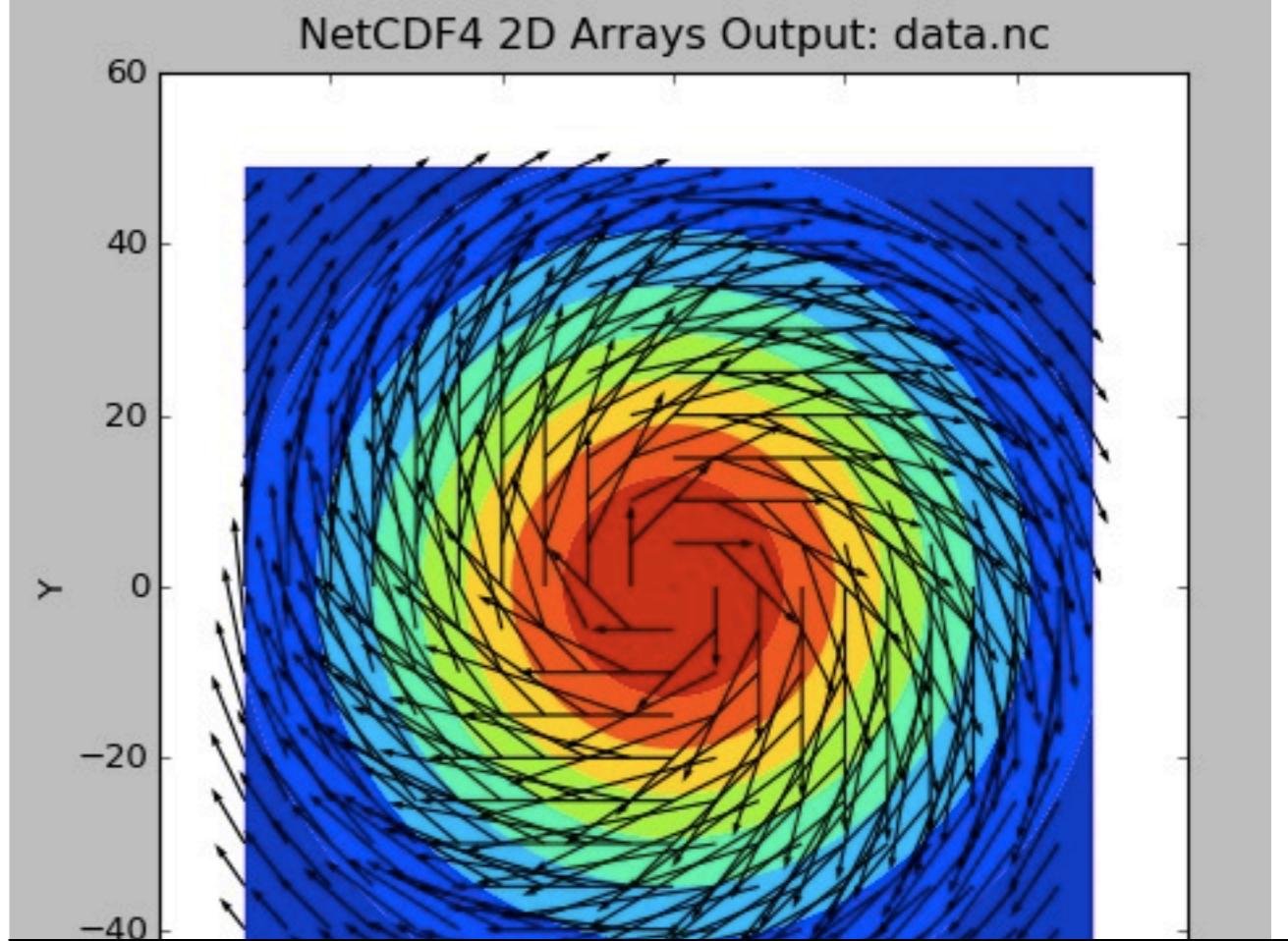
## NetCDF Conventions

Unidata offers a repository and will maintain WWW links for sets of netCDF conventions, as supported by the Conventions section of the netCDF User's Guide. The following sets of conventions are currently available:

- [CF Conventions \(Recommended, if applicable\)](#)
- [ACDD Conventions \(Attribute Convention for Dataset Discovery\)](#)
- [NCAR-RAF Conventions for Aircraft Data](#)
- [AMBER Trajectory Conventions for molecular dynamics simulations](#)
- [ARGO netCDF conventions for data centers](#)
- [National Oceanographic Data Center NetCDF Conventions](#)
- [Proposed CF Discrete Sampling Conventions \(draft CF conventions for observational and processing data\)](#)
- [Developing Conventions for NetCDF-4](#)
- [COARDS Conventions \(1995 standard that CF Conventions extends and generalizes\)](#)
- [GDT Conventions \(1999 standard that CF Conventions extends and generalizes\)](#)
- [CDC Conventions \(for gridded data, compatible with but more restrictive than COARDS\)](#)
- [NUWG Conventions \(1992-1995 effort to create some observational data conventions\)](#)

# Big advantage of self- describing:

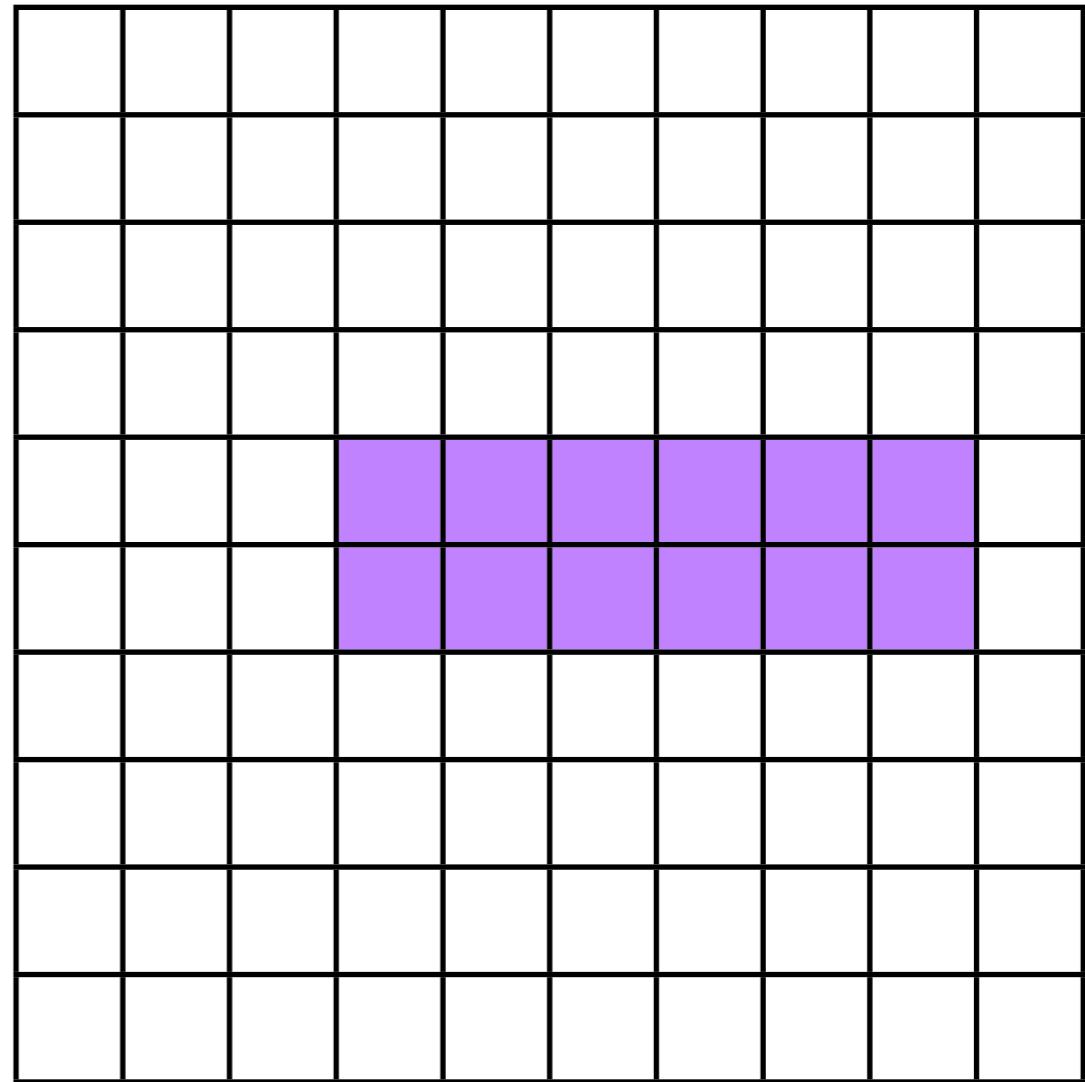
- Old program could easily read new file, even though data layout changed!
- Doesn't even need to know about attributes...
- New variables don't cause any problems - don't have to read them!
- Backwards compatibility
- But can look for them and use if available



```
$ ncdump -h data.nc
netcdf data {
dimensions:
 X = 100 ;
 Y = 100 ;
 velocity\ component = 2 ;
variables:
 float X\ coordinate(X) ;
 X\ coordinate:units = "cm" ;
 float Y\ coordinate(Y) ;
 Y\ coordinate:units = "cm" ;
 double Density(X, Y) ;
 Density:units = "g/cm^3" ;
 double Velocity(velocity\ component, X, Y)
 Velocity:units = "cm/s" ;
}
```

# Accessing subregions in file

- `nc_put_var_type` or `nf90_put_var` puts whole array (by default)
- Subarrays can be specified with starts and counts



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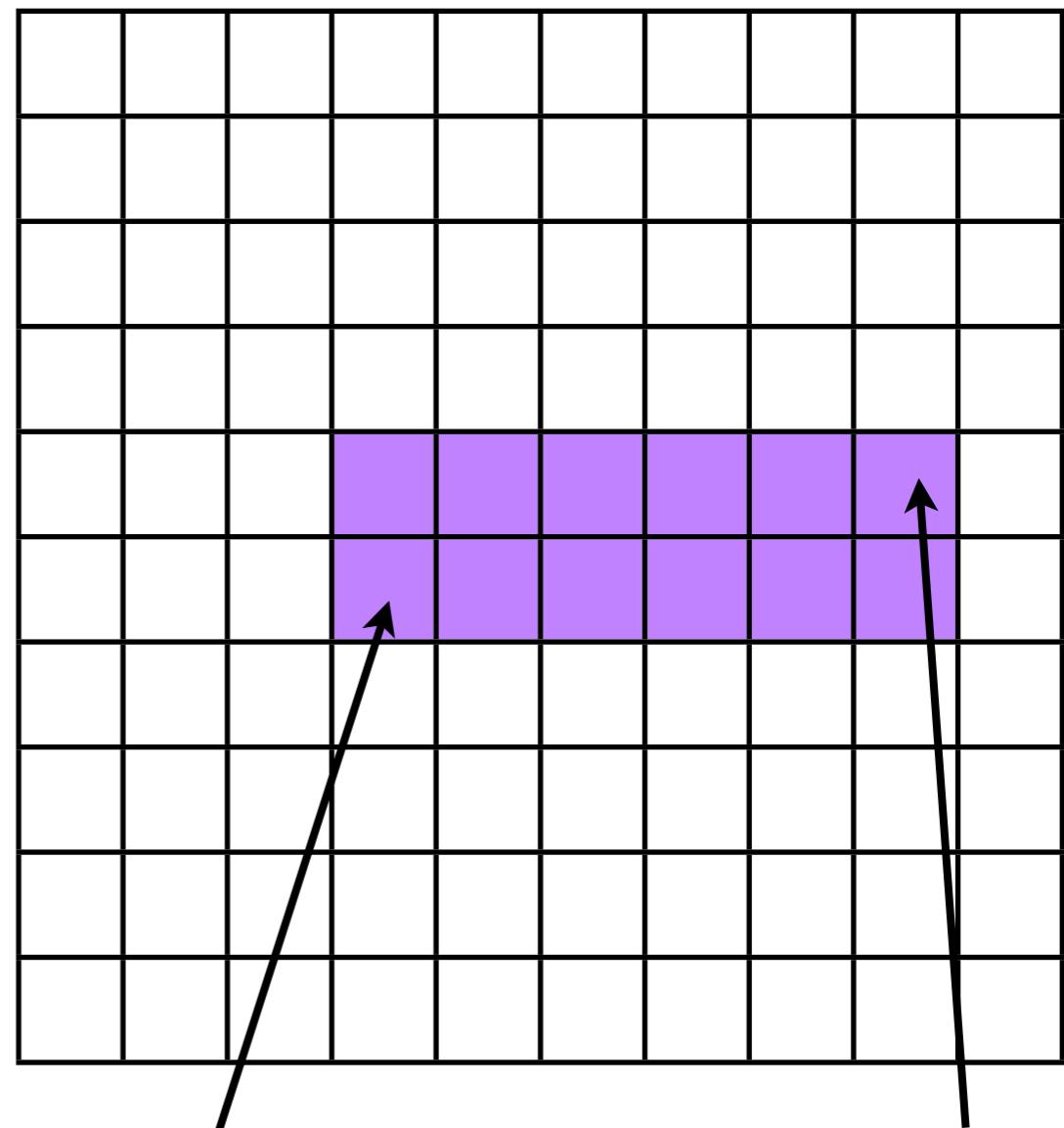
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$\text{dens}(10,10)$

```
start(1) = 4
start(2) = 5

count(1) = 6
count(2) = 2

nf90_put_var(file_id, dens_id,
data, START=start, COUNT=count)
```



$\text{dens}(4,5)$

$\text{dens}(9,6)$



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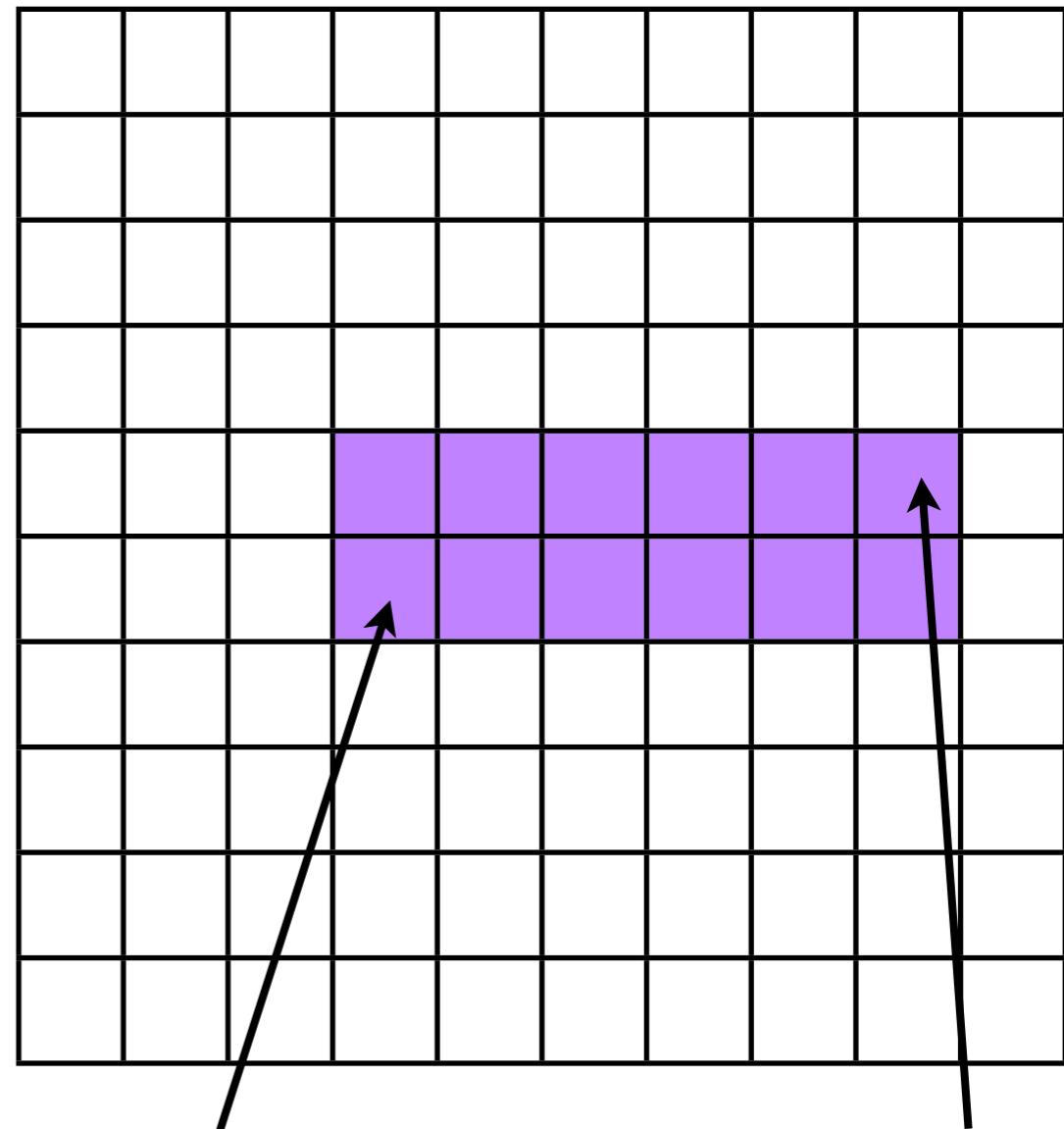
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**dens[10][10]**

```
start[0] = 3;
start[1] = 4;

count[0] = 6;
count[1] = 2;

nc_put_vara_double(file_id,
dens_id, start, count, data);
```



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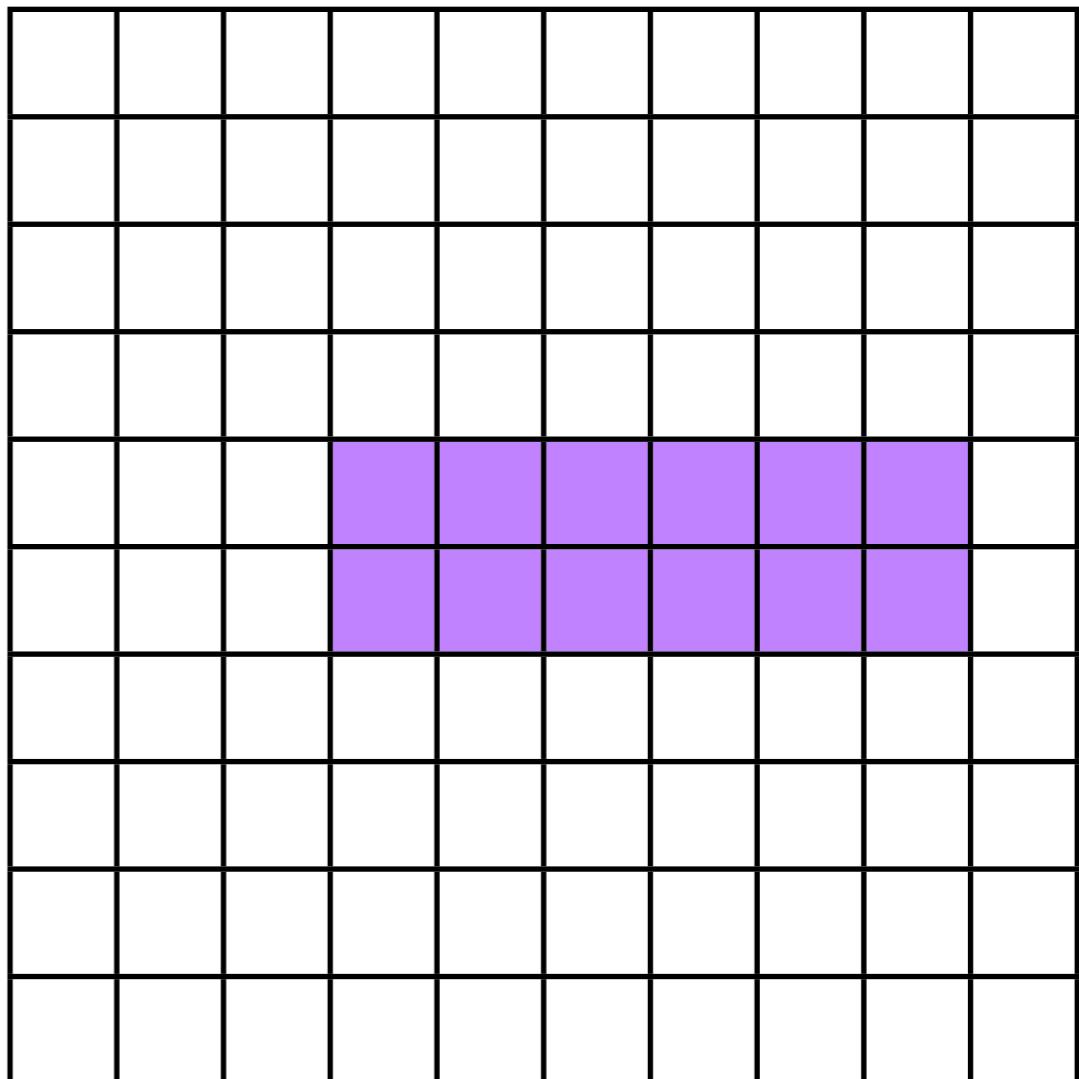
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# Accessing subregions in file

- Note that NetCDF libraries accept starting conventions of C, Fortran as appropriate.

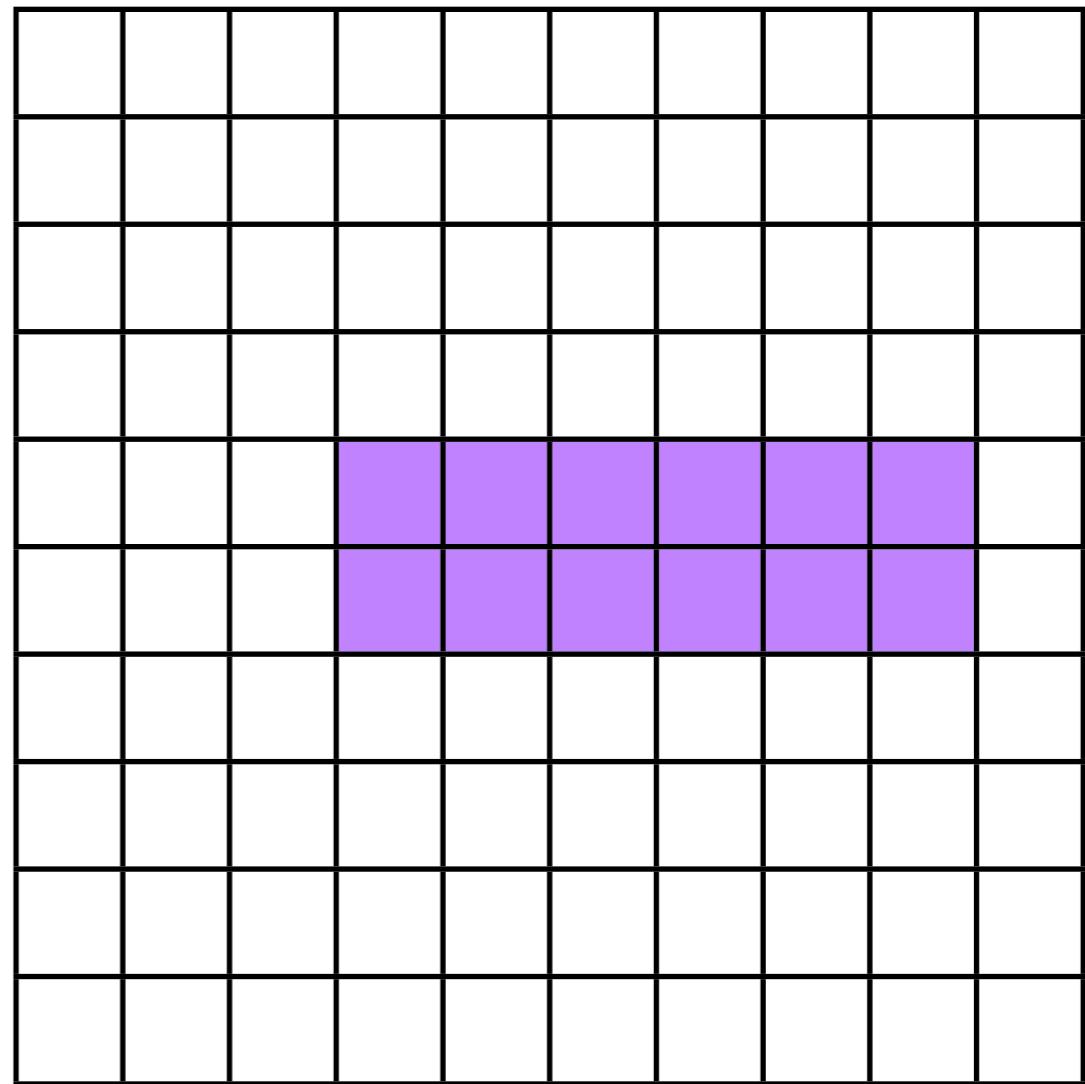


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# Accessing subregions in file

- Another thing this is good for; arrays in NetCDF can have a dimension of unlimited size (eg, can grow) - NetCDF3, only one dimension, NetCDF4, any
- Can use for timesteps, for instance.
- Any access to such a dataset is necessarily via subregions.



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# Fortran vs C array conventions

```
$ ncdump -h data.nc
netcdf data {
dimensions:
 X = 100 ;
 Y = 100 ;
 velocity\ component = 2 ;
variables:
 float X\ coordinate(X) ;
 X\ coordinate:units = "cm" ;
 float Y\ coordinate(Y) ;
 Y\ coordinate:units = "cm" ;
 double Density(X, Y) ;
 Density:units = "g/cm^3" ;
 double Velocity(velocity\
component, X, Y) ;
 Velocity:units = "cm/s" ;
}
```

```
$ ncdump -h data-fort.nc
netcdf data-fort {
dimensions:
 X = 100 ;
 Y = 100 ;
 velocity\ components = 2 ;
variables:
 float X\ coordinate(X) ;
 X\ coordinate:units = "cm" ;
 float Y\ coordinate(Y) ;
 Y\ coordinate:units = "cm" ;
 double Density(Y, X) ;
 Density:units = "g/cm^3" ;
 double Velocity(Y, X, velocity\
components) ;
 Velocity:units = "cm/s" ;
}
```



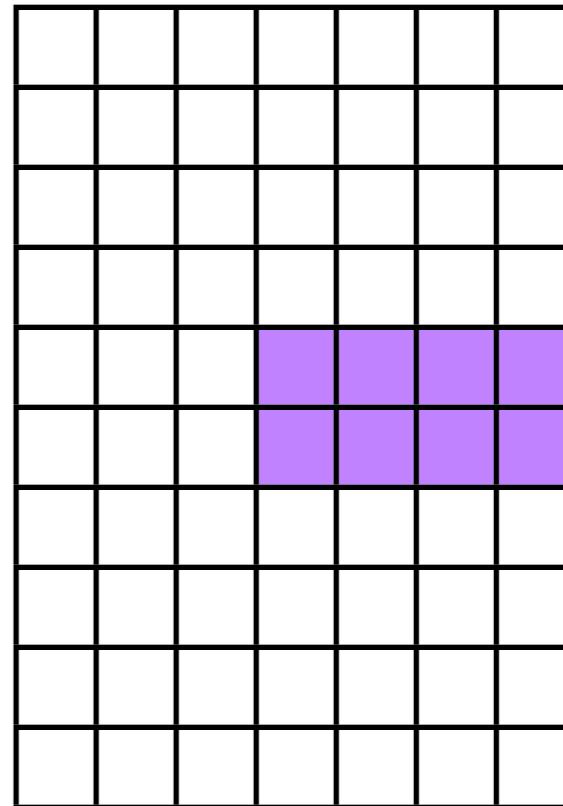
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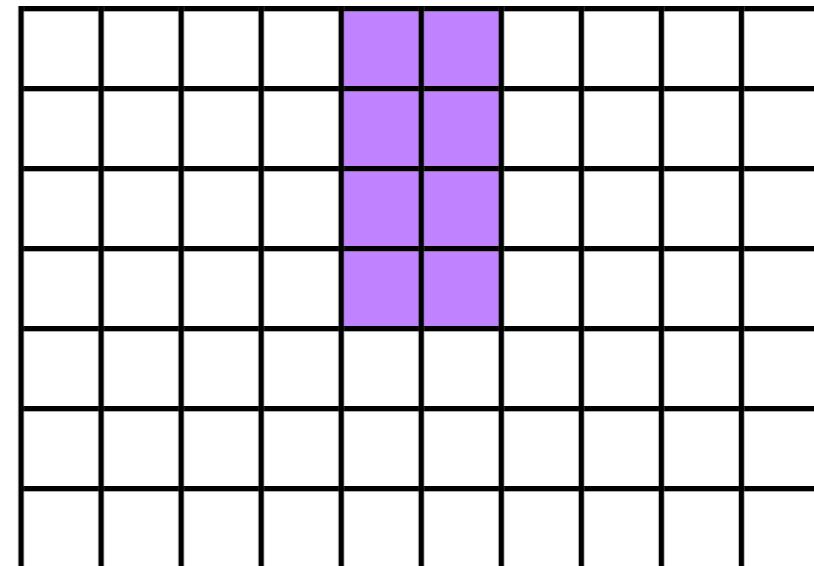
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# Mapping memory space to file

- Say in C you wanted to output in FORTRAN convention
- $(i,j)$  in your array corresponds to  $(j,i)$  in data space in file
- `nc_put_varm` allows you to do this by mapping how indicies vary in memory compared to in file.



`dens[7][10]`  
to  
`dens(10,7)`



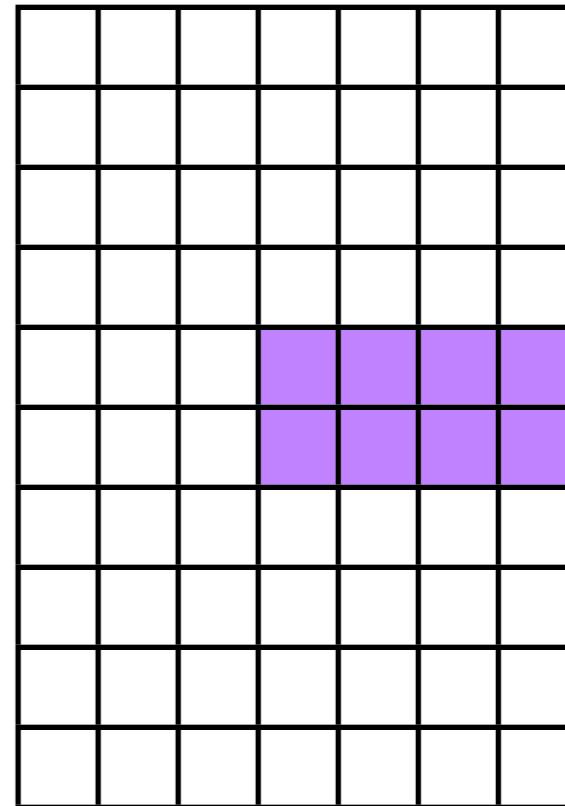
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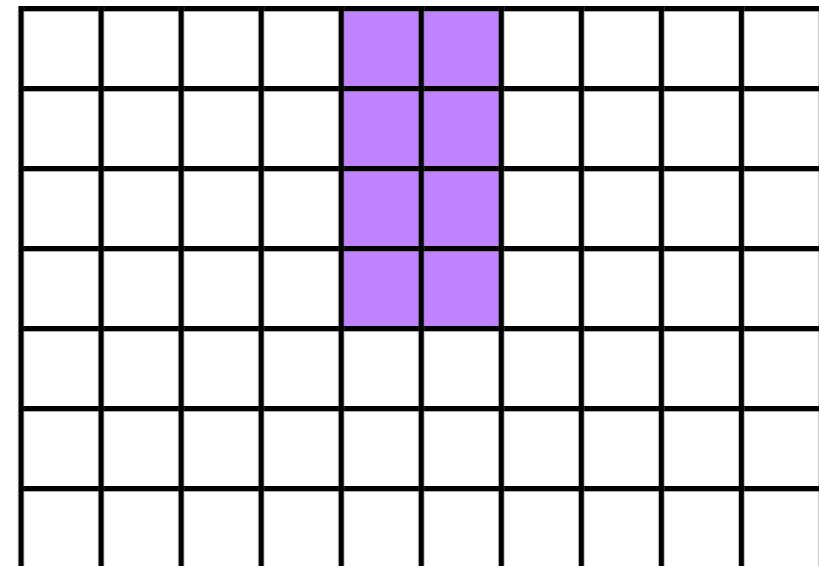
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# Mapping memory space to file

- Note - this requires understanding how memory is laid out in your data structures, as with MPI & MPI-IO
- This is *crucial* for I/O, and for HPC in general
- C has more flexibility (== potential problems) in this regard.



`dens[7][10]`  
to  
`dens(10,7)`



# Mapping memory space to file

- C: first array index most slowly varying.
- Eg, for a 3x4 array, each step in the 2nd index jumps you one position in memory,
- and in the first index, jumps you by 4.
- You could write this as (4,1)

Your picture of the array

|        |        |        |        |
|--------|--------|--------|--------|
| [2][0] | [2][1] | [2][2] | [2][3] |
| [1][0] | [1][1] | [1][2] | [1][3] |
| [0][0] | [0][1] | [0][2] | [0][3] |

In memory

|        |        |        |        |        |        |        |        |        |
|--------|--------|--------|--------|--------|--------|--------|--------|--------|
| [0][0] | [0][1] | [0][2] | [0][3] | [1][0] | [1][1] | [1][2] | [1][3] | [2][0] |
|--------|--------|--------|--------|--------|--------|--------|--------|--------|



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# Mapping memory space to file

- But if you're writing to a fortran-convention file, you want this to go the other way
- In the file, one step in the **I**st index should jump you by 1, and the second by...

Your picture of the array

|        |        |        |        |
|--------|--------|--------|--------|
| [2][0] | [2][1] | [2][2] | [2][3] |
| [1][0] | [1][1] | [1][2] | [1][3] |
| [0][0] | [0][1] | [0][2] | [0][3] |

In memory

|        |        |        |        |        |        |        |        |        |
|--------|--------|--------|--------|--------|--------|--------|--------|--------|
| [0][0] | [0][1] | [0][2] | [0][3] | [1][0] | [1][1] | [1][2] | [1][3] | [2][0] |
|--------|--------|--------|--------|--------|--------|--------|--------|--------|



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# Mapping memory space to file

- But if you're writing to a fortran-convention file, you want this to go the other way
- In the file, one step in the **I**st index should jump you by 1, and the second by 3.
- The map you want is (1,3)

Your picture of the array

|        |        |        |        |
|--------|--------|--------|--------|
| [2][0] | [2][1] | [2][2] | [2][3] |
| [1][0] | [1][1] | [1][2] | [1][3] |
| [0][0] | [0][1] | [0][2] | [0][3] |

In memory

|        |        |        |        |        |        |        |        |        |
|--------|--------|--------|--------|--------|--------|--------|--------|--------|
| [0][0] | [0][1] | [0][2] | [0][3] | [1][0] | [1][1] | [1][2] | [1][3] | [2][0] |
|--------|--------|--------|--------|--------|--------|--------|--------|--------|



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# Mapping memory space to file

```
start = count = stride = NULL;
int imap[2] = {1,3};
```

```
nc_put_varm_double(file_id,
dens_id, start, count, stride,
imap, data);
```

```
nf90_put_var(file_id, dens_id,
data, MAP=(/4,1/))
```

Your picture of the array

|        |        |        |        |
|--------|--------|--------|--------|
| [2][0] | [2][1] | [2][2] | [2][3] |
| [1][0] | [1][1] | [1][2] | [1][3] |
| [0][0] | [0][1] | [0][2] | [0][3] |

In memory

|     |        |        |        |        |        |        |        |        |
|-----|--------|--------|--------|--------|--------|--------|--------|--------|
| [0] | [0][1] | [0][2] | [0][3] | [1][0] | [1][1] | [1][2] | [1][3] | [2][0] |
|-----|--------|--------|--------|--------|--------|--------|--------|--------|



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# More on NetCDF

- [http://www.unidata.ucar.edu/  
software/netcdf/](http://www.unidata.ucar.edu/software/netcdf/)
- Docs, mailing lists, tutorials,  
sample code, API, etc.



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# Sample Code



```
$ cd parIO/hdf5

$ source /groups/h/hpc2011/bin/Parallel_IO.csh
$ make 2darray (C), or
$ make f2darray (F90)

$./{f,}2darray
$ ls *.h5

$../plots.py *.h5
```

# What is this .h5 file?

```
$ h5ls data-fort.h5
```

ArrayData

Group

OtherStuff

Group

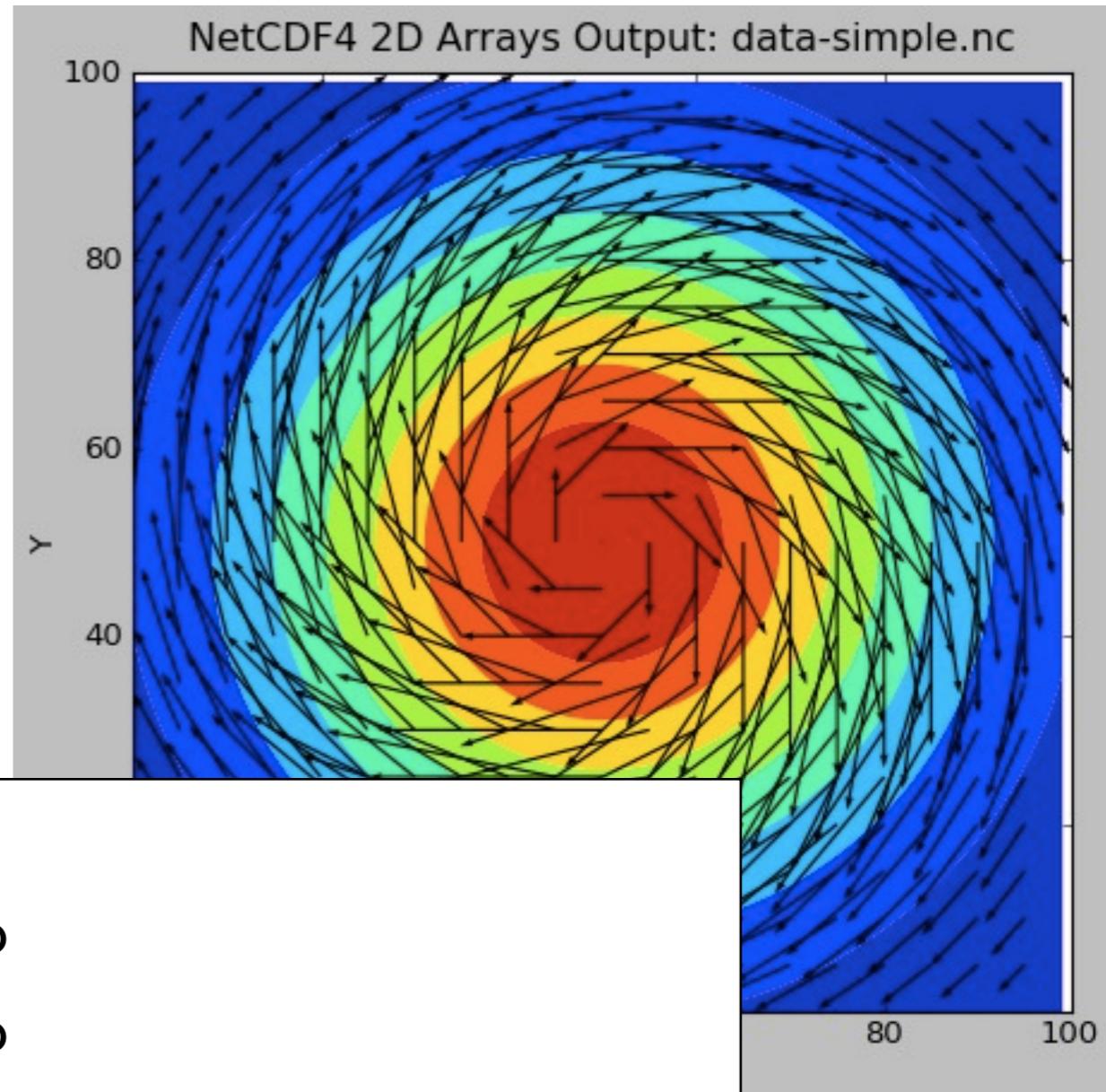
```
$ h5ls data-fort.h5/ArrayData
```

dens

Dataset {100, 100}

vel

Dataset {100, 100, 2}



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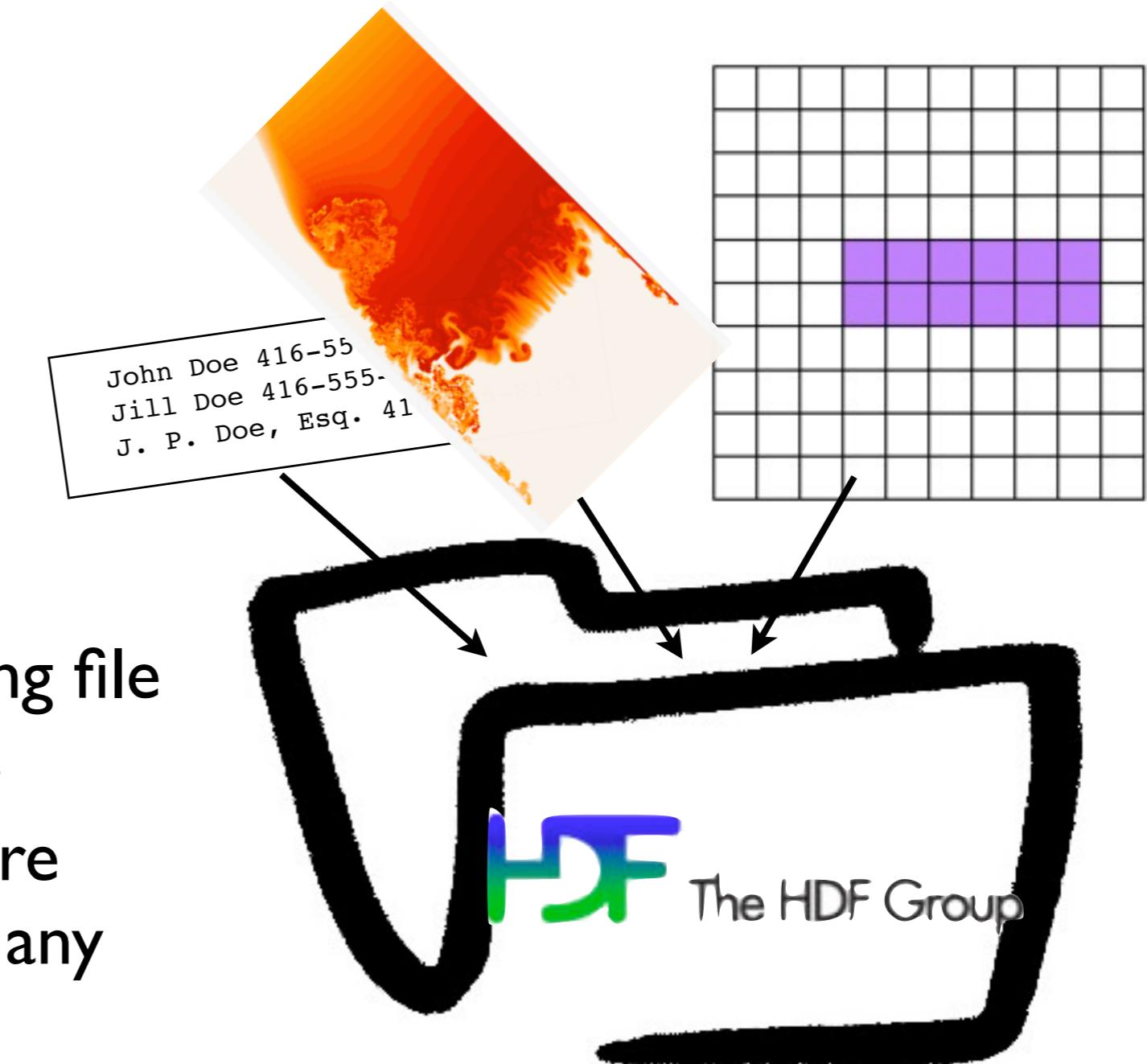
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# HDF5

- HDF5 is also self-describing file format and set of libraries
- Unlike NetCDF, much more general; can shove almost any type of data in there
- (We'll just be looking at large arrays, since that's our usual use case)

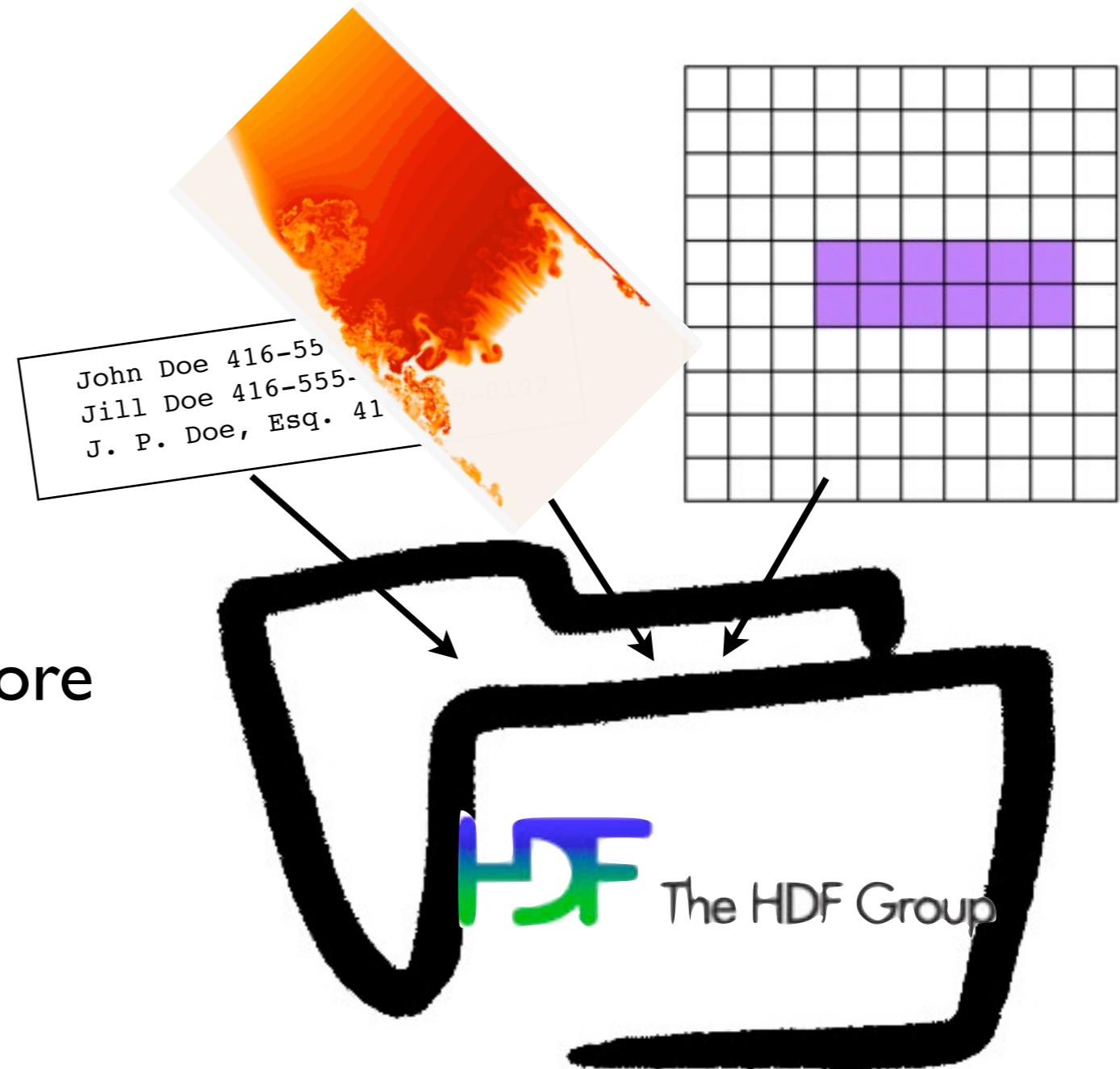


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# HDF5

- Much more general, and more low-level than NetCDF.
- (In fact, newest version of NetCDF implemented in HDF5).
- Pro: *can* do more!
- Con: **have** to do more.



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# 2darray-simple.c

```
/* identifiers */

hid_t file_id, dens_dataset_id, vel_dataset_id;
hid_t dens_dataspace_id, vel_dataspace_id;

/* sizes */

hsize_t densdims[2], veldims[3];

/* status */

herr_t status;

/* Create a new file - truncate anything existing, use default properties */
file_id = H5Fcreate(rundata.filename, H5F_ACC_TRUNC, H5P_DEFAULT,
H5P_DEFAULT);

/* HDF5 routines generally return a negative number on failure.
 * Should check return values! */

if (file_id < 0) {
 fprintf(stderr,"Could not open file %s\n", rundata.filename);
 return;
}
```

# 2darray-simple.c

```
/* identifiers */

hid_t file_id, dens_dataset_id, vel_dataset_id;

hid_t dens_dataspace_id, vel_dataspace_id;

/* sizes */
hsize_t densdims[2], veldims[3];

/* status */
herr_t status;

/* Create a new file - truncate anything
 */

file_id = H5Fcreate(rundata.filename, H5F_ACC_TRUNC, H5P_DEFAULT,
H5P_DEFAULT);

/* HDF5 routines generally return a negative number on failure.
 * Should check return values! */

if (file_id < 0) {
 fprintf(stderr, "Could not open file %s\n", rundata.filename);
 return;}
```

NetCDF used ints for everything - HDF5 distinguishes between ids, sizes, errors, uses its own types.

# 2darray-simple.c

```
/* identifiers */

hid_t file_id, dens_dataset_id, vel_dataset_id;

hid_t dens_dataspace_id, vel_dataspace_id;

/* sizes */

hsize_t densdims[2], veldims[3];

/* status */

herr_t status;

/* Create a new file - truncate anything existing, use default properties */
file_id = H5Fcreate(rundata.filename, H5F_ACC_TRUNC, H5P_DEFAULT,
H5P_DEFAULT);

/* HDF5 routines generally return a negative number on failure.
 * Should check return values! */

if (file_id < 0) {
 fprintf(stderr,"Could not open file %s\n", rundata.filename);
 return;
}
```

**H5F, H5P... ?**



# Decomposing the HDF5 API

- HDF5 API is large
  - Constants, function calls start with H5x; x tells you what part of the library
  - Table tells you (some) of those parts...
  - Fortran the same, but usually end with F
- HPC\$20T2

|     |                     |
|-----|---------------------|
| H5A | <b>Attributes</b>   |
| H5D | <b>Datasets</b>     |
| H5E | <b>Errors</b>       |
| H5F | <b>Files</b>        |
| H5G | <b>Groups</b>       |
| H5P | <b>Properties</b>   |
| H5S | Data <b>S</b> paces |
| H5T | Data <b>T</b> ypes  |



# 2darray-simple.c

```
/* Create the data space for the two datasets. */
densdims[0] = rundata.nx; densdims[1] = rundata.ny;
veldims[0] = 2; veldims[1] = rundata.nx; veldims[2] = rundata.ny;

dens_dataspace_id = H5Screate_simple(2, densdims, NULL);
vel_dataspace_id = H5Screate_simple(3, veldims, NULL);

/* Create the datasets within the file.
 * H5T_IEEE_F64LE is a standard (IEEE) double precision (64 bit)
 * floating (F) data type and will work on any machine.
 * H5T_NATIVE_DOUBLE would work too */

dens_dataset_id = H5Dcreate(file_id, "dens", H5T_IEEE_F64LE,
 dens_dataspace_id, H5P_DEFAULT,
 H5P_DEFAULT, H5P_DEFAULT);

vel_dataset_id = H5Dcreate(file_id, "vel", H5T_IEEE_F64LE,
 vel_dataspace_id, H5P_DEFAULT,
 H5P_DEFAULT, H5P_DEFAULT);
```

# 2darray-simple.c

```
/* Create the data space for the two datasets. */
densdims[0] = rundata.nx; densdims[1] = rundata.ny;
veldims[0] = 2; veldims[1] = rundata.nx; veldims[2] = rundata.ny;

dens_dataspace_id = H5Screate_simple(2, densdims, NULL);
vel_dataspace_id = H5Screate_simple(3, veldims, NULL);

/* Create the datasets within the file
 * H5T_IEEE_F64LE is a standard (IEEE)
 * floating (F) data type and will work
 * H5T_NATIVE_DOUBLE would work too */
dens_dataset_id = H5Dcreate(file_id, "density", H5T_IEEE_F64LE,
 dens_dataspace_id, H5P_DEFAULT);
H5P_SET_DEFLATE_COMPRESSION(dens_dataset_id, 5);

vel_dataset_id = H5Dcreate(file_id, "velocity", H5T_IEEE_F64LE,
 vel_dataspace_id, H5P_DEFAULT);
H5P_SET_DEFLATE_COMPRESSION(vel_dataset_id, 5);
```

All data (in file or in mem) in HDF5 has a dataspace it lives in.

In NetCDF, just cartesian product of dimensions; here more general

# 2darray-simple.c

```
/* Create the data space for the two datasets. */
densdims[0] = rundata.nx; densdims[1] =
veldims[0] = 2; veldims[1] = rundata.nx

dens_dataspace_id = H5Screate_simple(2,
vel_dataspace_id = H5Screate_simple(3,

/* Create the datasets within the file.
 * H5T_IEEE_F64LE is a standard (IEEE)
 * floating (F) data type and will work
 * H5T_NATIVE_DOUBLE would work too */

dens_dataset_id = H5Dcreate(file_id, "dens", H5T_IEEE_F64LE,
 dens_dataspace_id, H5P_DEFAULT,
 H5P_DEFAULT, H5P_DEFAULT);

vel_dataset_id = H5Dcreate(file_id, "vel", H5T_IEEE_F64LE,
 vel_dataspace_id, H5P_DEFAULT,
 H5P_DEFAULT, H5P_DEFAULT);
```

**Creating a data set like defining a variable in NetCDF.**  
**Also declare the type you want it to be on disk.**



# 2darray-simple.c

```
/* Write the data. We're writing it from memory, where it is saved
 * in NATIVE_DOUBLE format */

status = H5Dwrite(dens_dataset_id, H5T_NATIVE_DOUBLE, H5S_ALL, H5S_ALL,
H5P_DEFAULT, &(dens[0][0]));

status = H5Dwrite(vel_dataset_id, H5T_NATIVE_DOUBLE, H5S_ALL, H5S_ALL,
H5P_DEFAULT, &(vel[0][0][0]));

/* End access to groups & data sets and release resources used by them */

status = H5Sclose(dens_dataspace_id);
status = H5Dclose(dens_dataset_id);
status = H5Sclose(vel_dataspace_id);
status = H5Dclose(vel_dataset_id);

/* Close the file */
status = H5Fclose(file_id);
```

Write memory from all of memory to all of the dataset on the file.  
Values in mem are in the native double precision format.

# 2darray-simple.c

```
/* Write the data. We're writing it from memory, where it is saved
 * in NATIVE_DOUBLE format */

status = H5Dwrite(dens_dataset_id, H5T_NATIVE_DOUBLE, H5S_ALL, H5S_ALL,
H5P_DEFAULT, &(dens[0][0]));

status = H5Dwrite(vel_dataset_id, H5T_NATIVE_DOUBLE, H5S_ALL, H5S_ALL,
H5P_DEFAULT, &(vel[0][0][0]));

/* End access to groups & data sets and release resources used by them */

status = H5Sclose(dens_dataspace_id);
status = H5Dclose(dens_dataset_id);
status = H5Sclose(vel_dataspace_id);
status = H5Dclose(vel_dataset_id);

/* Close the file */
status = H5Fclose(file_id);
```

Close everything

# f2darray-simple.f90

```
integer(hid_t) :: file_id
integer(hid_t) :: dens_space_id, vel_space_id
integer(hid_t) :: dens_id, vel_id
integer(hsize_t), dimension(2) :: densdims
integer(hsize_t), dimension(3) :: veldims
integer :: status
! first we have to open the FORTRAN inter
call h5open_f(status)
```

Fortran: values are  
`integer(hid_t)` or  
`integer(hsize_t)`

```
! create the file, check return code
call h5fcreate_f(rundata%filename, H5F_ACC_TRUNC_F, file_id, status)
if (status /= 0) then
 print *, 'Could not open file ', rundata%filename
 return
endif
```

# f2darray-simple.f90

```
integer(hid_t) :: file_id
integer(hid_t) :: dens_space_id, vel_space_id
integer(hid_t) :: dens_id, vel_id
integer(hsize_t), dimension(2) :: densdims
integer(hsize_t), dimension(3) :: veldims

integer :: status

! first we have to open the FORTRAN interface.

call h5open_f(status) ←
! create the file, check return code
call h5fcreate_f(rundata%filename, H5F_ACC_TRUNC_F, file_id, status)
if (status /= 0) then
 print *, 'Could not open file ', rundata%filename
 return
endif
```

Have to start the  
FORTRAN interface

# f2darray-simple.f90

```
integer(hid_t) :: file_id
integer(hid_t) :: dens_space_id, vel_space_id
integer(hid_t) :: dens_id, vel_id
integer(hsize_t), dimension(2) :: densdims
integer(hsize_t), dimension(3) :: veldims

integer :: status

! first we have to open the FORTRAN interface.
call h5open_f(-status)
! create the file, check return value
call h5fcreate_f(:data%filename, H5F_ACC_TRUNC_F, file_id, status)
if (status /= 0) then
 print *, 'Could not open file ', rundata%filename
 return
endif
```

See what I mean about  
\_F?

# f2darray-simple.f90

```
! create the dataspaces corresponding to our variables
densdims = (/ rundata % nx, rundata % ny /)
call h5screate_simple_f(2, densdims, dens_space_id, status)

veldims = (/ 2, rundata % nx, rundata % ny /)
call h5screate_simple_f(3, veldims, vel_space_id, status)

! now that the dataspaces are defined, we can define variables on them

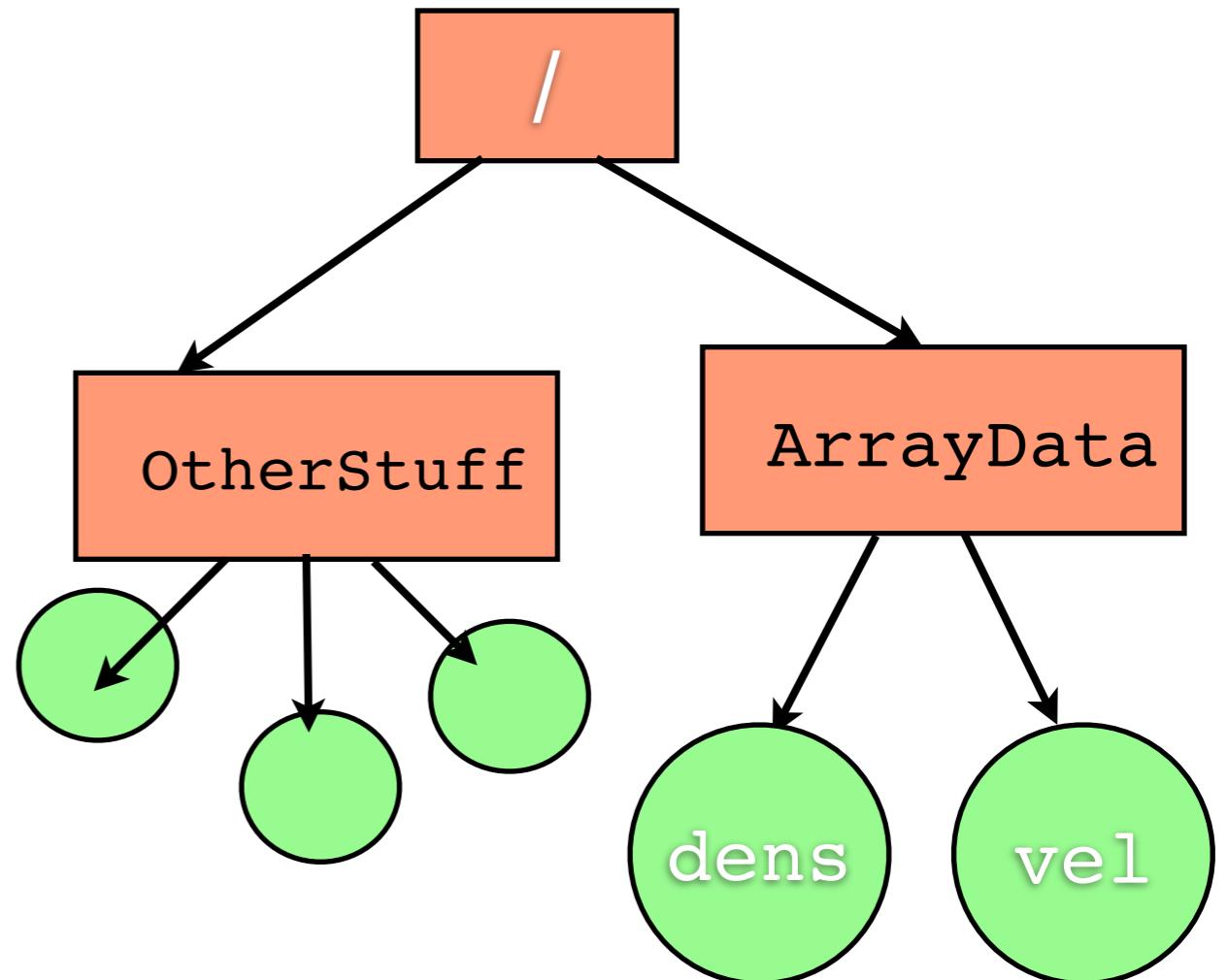
call h5dcreate_f(file_id, "dens", H5T_IEEE_F64LE, dens_space_id, dens_id,
status)
call h5dcreate_f(file_id, "vel" , H5T_IEEE_F64LE, vel_space_id, vel_id,
status)
```



In F90 interface, a lot of less-common arguments are optional; fewer H5P\_DEFAULTs kicking around

# HDF5 Groups

- HDF5 has a structure a bit like a unix filesystem:
- “Groups” - directories
- “Datasets” - files
- NetCDF4 now has these, but breaks compatibility with NetCDF3 files



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## 2darray.c

```
/* Create a new group within the new file */
arr_group_id = H5Gcreate(file_id, "/ArrayData", H5P_DEFAULT, H5P_DEFAULT,
H5P_DEFAULT);

...
dens_dataset_id = H5Dcreate(file_id, "/ArrayData/dens", H5T_IEEE_F64LE,
 dens_dataspace_id, H5P_DEFAULT,
 H5P_DEFAULT, H5P_DEFAULT);

vel_dataset_id = H5Dcreate(file_id, "/ArrayData/vel", H5T_IEEE_F64LE,
 vel_dataspace_id, H5P_DEFAULT,
 H5P_DEFAULT, H5P_DEFAULT);
```



Can specify that a dataset goes in a group by giving it an “absolute path”...

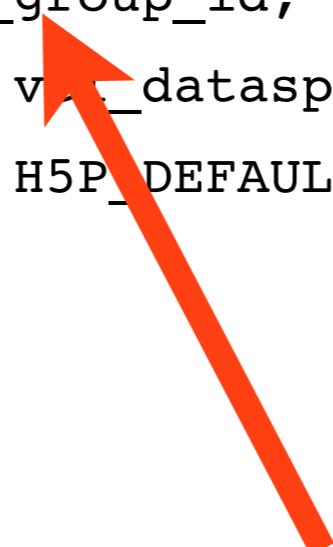
```
/* Create a new group within the new file */

arr_group_id = H5Gcreate(file_id, "/ArrayData", H5P_DEFAULT, H5P_DEFAULT,
H5P_DEFAULT);

...

dens_dataset_id = H5Dcreate(arr_group_id, "dens", H5T_IEEE_F64LE,
 dens_dataspace_id, H5P_DEFAULT,
 H5P_DEFAULT, H5P_DEFAULT);

vel_dataset_id = H5Dcreate(arr_group_id, "vel", H5T_IEEE_F64LE,
 vel_dataspace_id, H5P_DEFAULT,
 H5P_DEFAULT, H5P_DEFAULT);
```

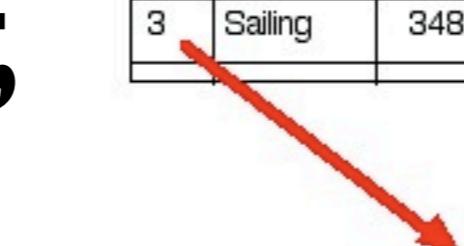


...or just by creating it *in* the group, rather than the file.

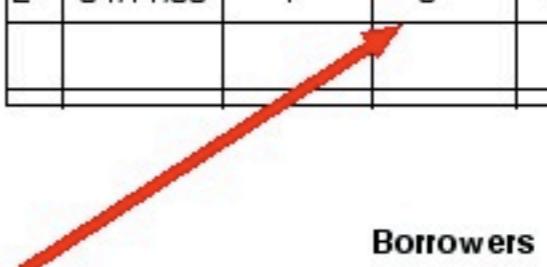
# What NetCDF, HDF aren't

- Databases
- Seem like - lots of information, in key value pairs.
- Relational databases - interrelated tables of **small** pieces of data
- Very easy/fast to query
- But can't do subarrays, etc..

| Books |          |            |             |      |        |
|-------|----------|------------|-------------|------|--------|
| bid   | title    | isbn       | author      | date | volume |
| 1     | Big Cats | 24589673-0 | Cat, Simon  | 2003 | 2      |
| 2     | Plants   | 24316759-1 | Smith, Rose | 1967 | 1      |
| 3     | Sailing  | 34817645-0 | Jones, Tom  | 1868 | 1      |



| Transactions |          |     |     |          |  |
|--------------|----------|-----|-----|----------|--|
| tid          | date     | bid | pid | duedate  |  |
| 1            | 02/11/08 | 3   | 2   | 16/11/08 |  |
| 2            | 04/11/08 | 1   | 3   | 18/11/08 |  |
|              |          |     |     |          |  |



| Borrowers |           |          |              |          |       |
|-----------|-----------|----------|--------------|----------|-------|
| pid       | firstname | lastname | address      | phone    | fines |
| 1         | Fred      | Thompson | 2 Reach Rd.  | 827-9867 | 2.25  |
| 2         | Sam       | Trunker  | 23 stone St. | 243-0955 | 0     |
| 3         | Tony      | Sandhas  | 4 two Rd.    | 123-6453 | 0     |
|           |           |          |              |          |       |



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# Databases for science

```
INSERT INTO benchmarkruns
values (newrunnum, datestr,
timestr, julianum)
...
SELECT nprocs, test, size,
transport, mpitype, runtime,
mopsperproc, run FROM
mpirundata WHERE (success=1)
```

| run# | success | size | transport | ... |
|------|---------|------|-----------|-----|
| 93   | no      | 12k  | eth       |     |
| I    | yes     | 512  | eth       |     |
| 87   | yes     | 64   | ib        |     |
| 13   | no      | 32   | eth       |     |
| ...  |         |      |           |     |



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# Parallel I/O using NetCDF4, HDF5



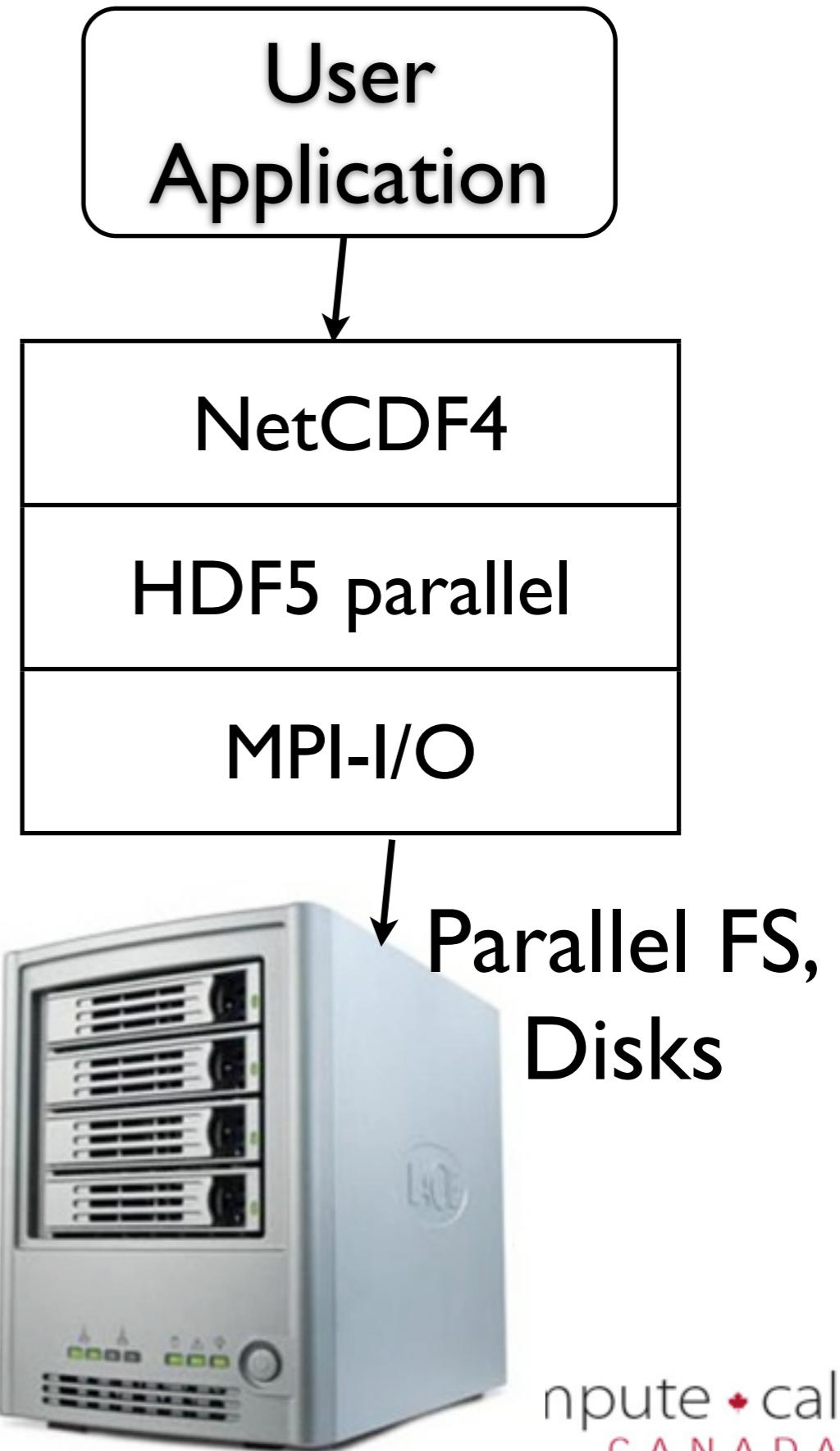
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# Parallel I/O libraries

- Can use the same NetCDF(4), HDF5 libraries to do Parallel IO on top of the MPI-I/O library
- Reading file afterwards, can't tell the difference.
- Fairly minor differences in function calls to do parallel I/O
- Hard part is figuring out what/where to write



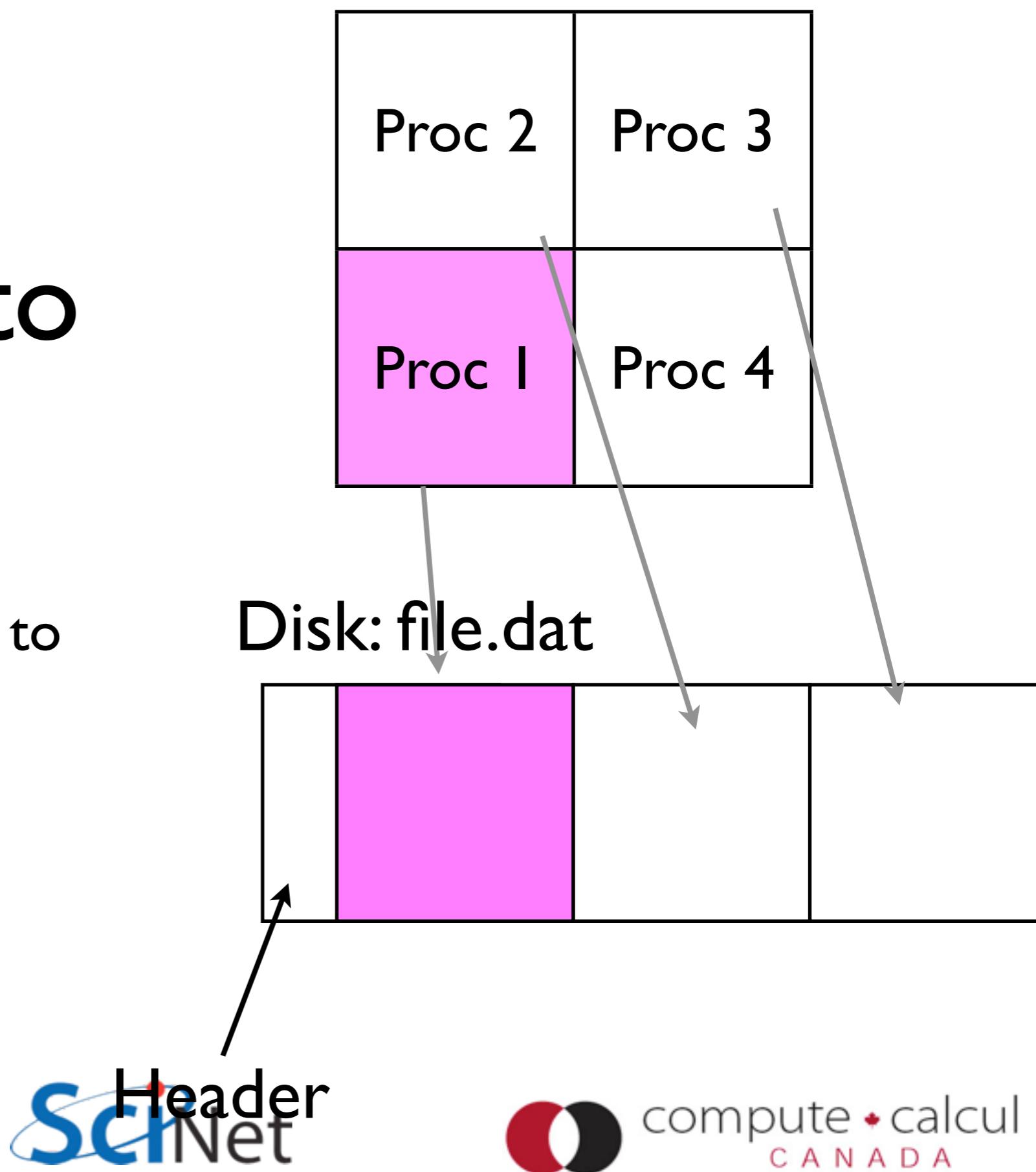
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# Parallel IO to One file

- Can be made to work efficiently, but must write to *disjoint* chunks of file
- Should write *big* disjoint chunks of file.



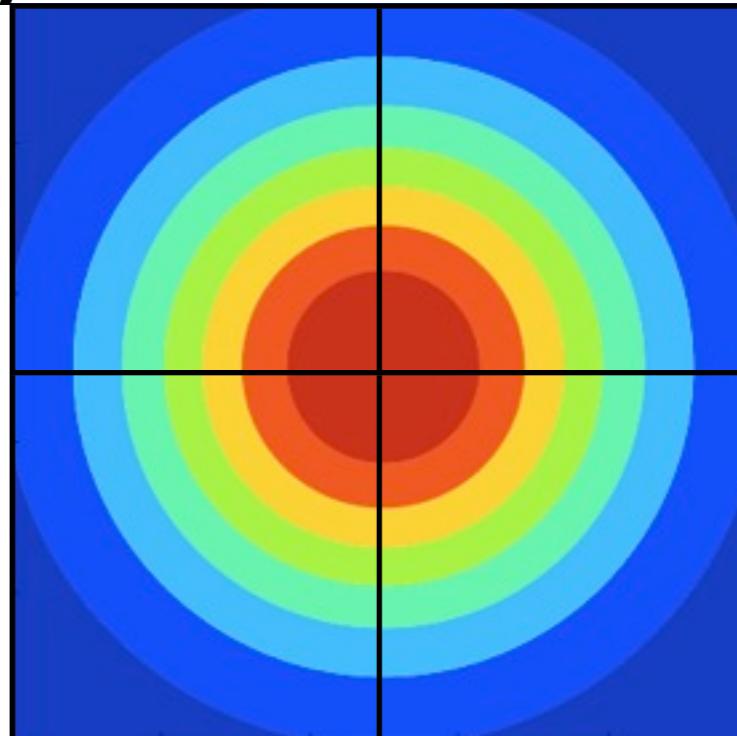
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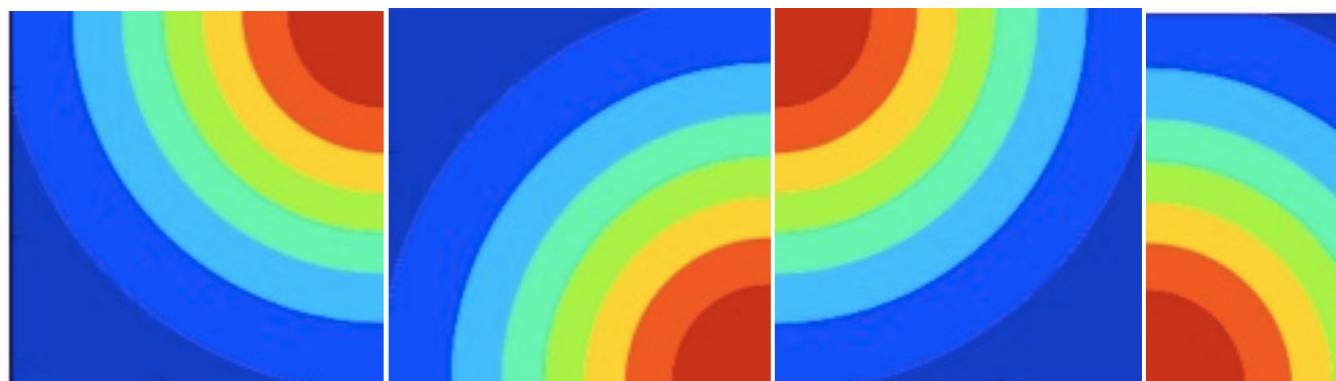
## Memory:



# How do you decide where to write?

- One possibility: each processor writes out its part of problem, in order.
- Pros - can be super fast.
- Cons - Output depends on number of processors run on. Analysis routines, restarts...

## Disk:



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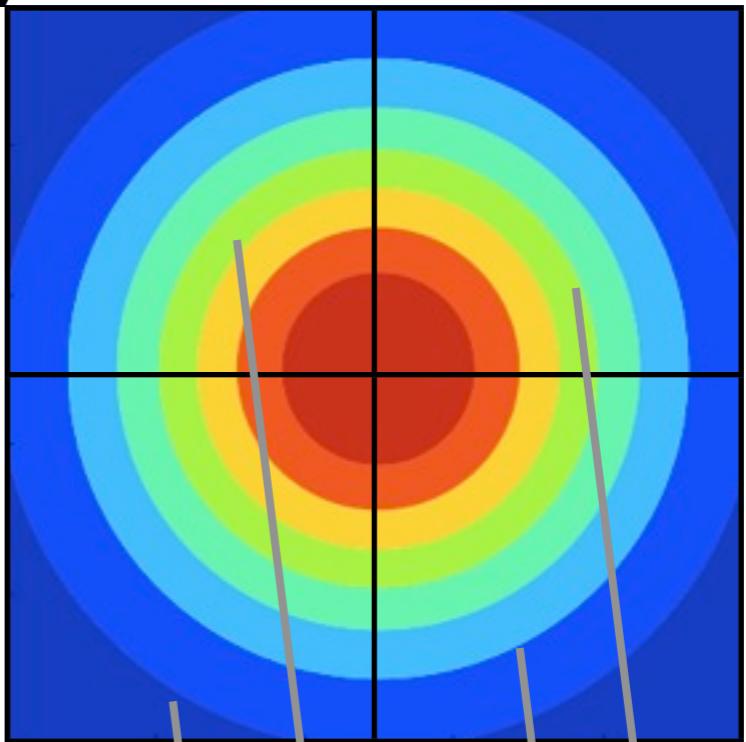


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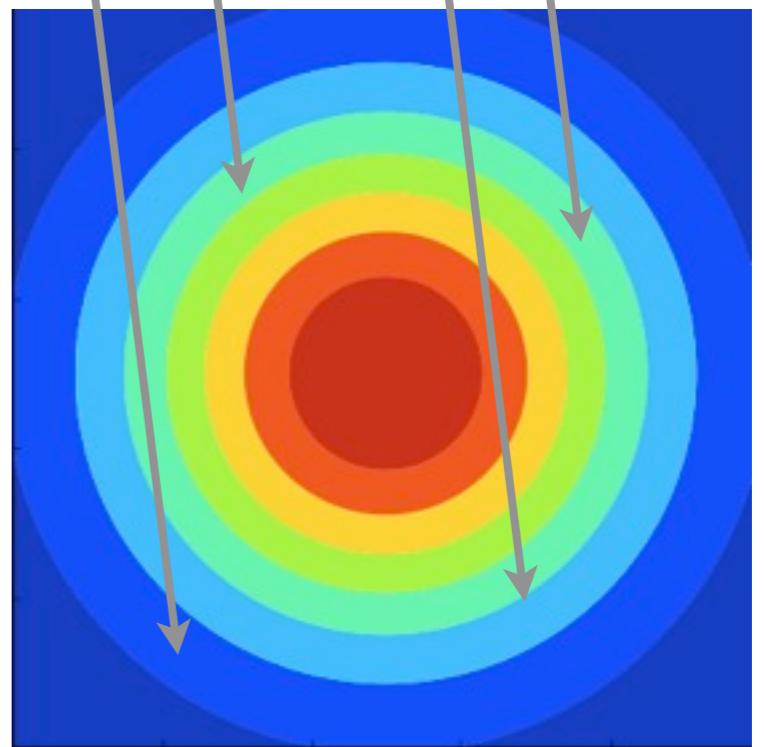
# How do you decide where to write?

- Other possibility: Write out chunks as they would be in memory on serial machine
- Pros: File looks the same no matter how many processes were used to write.
- Cons: Noncontig access; may be slower, but MPI-IO collective + good parallel FS should make competitive.

Memory:



Disk:



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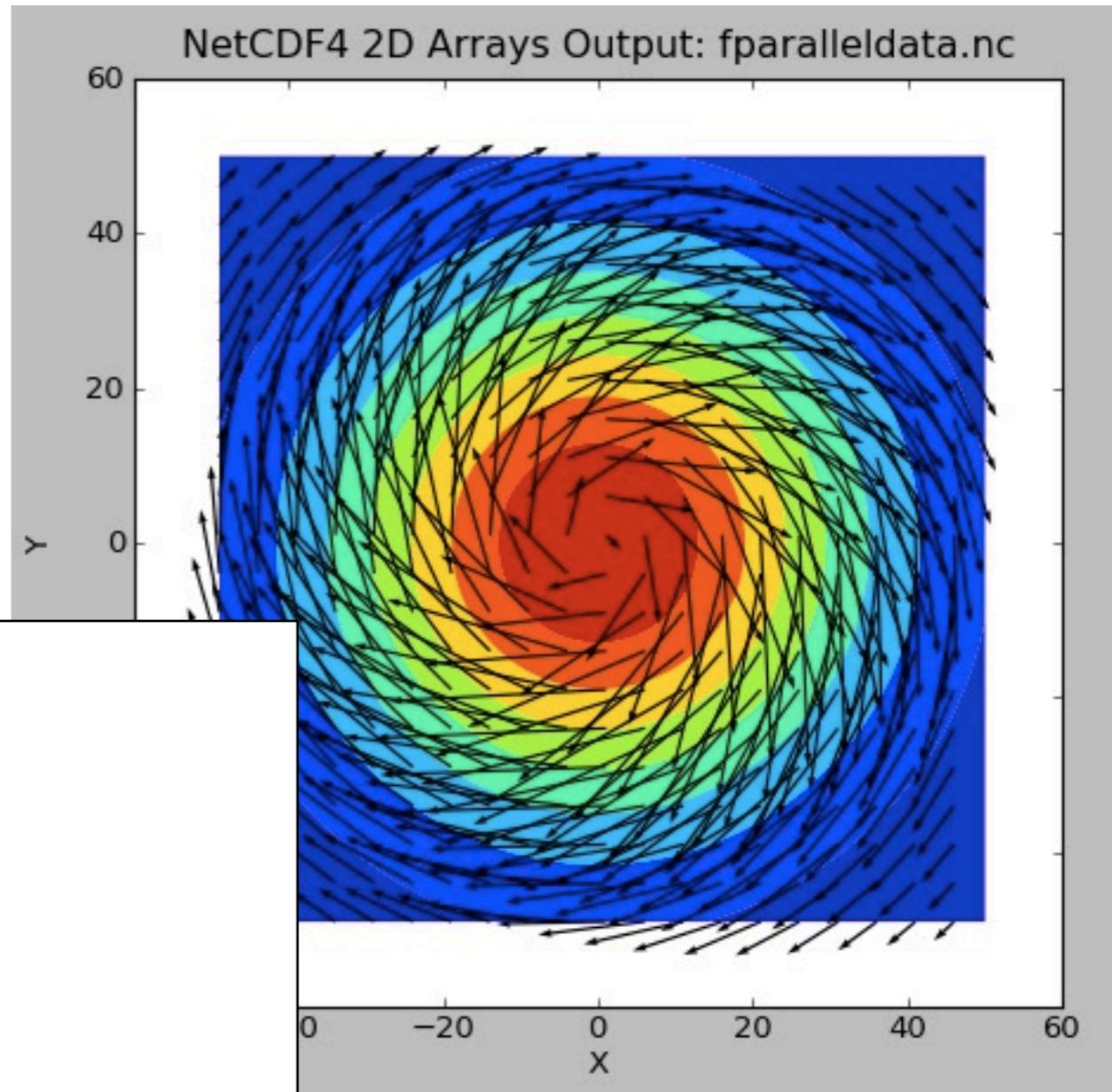
# Sample Code

```
$ cd
$ cd parIO/netcdf

$ make parallel2darray (C), or
$ make fparallel2darray (F90)

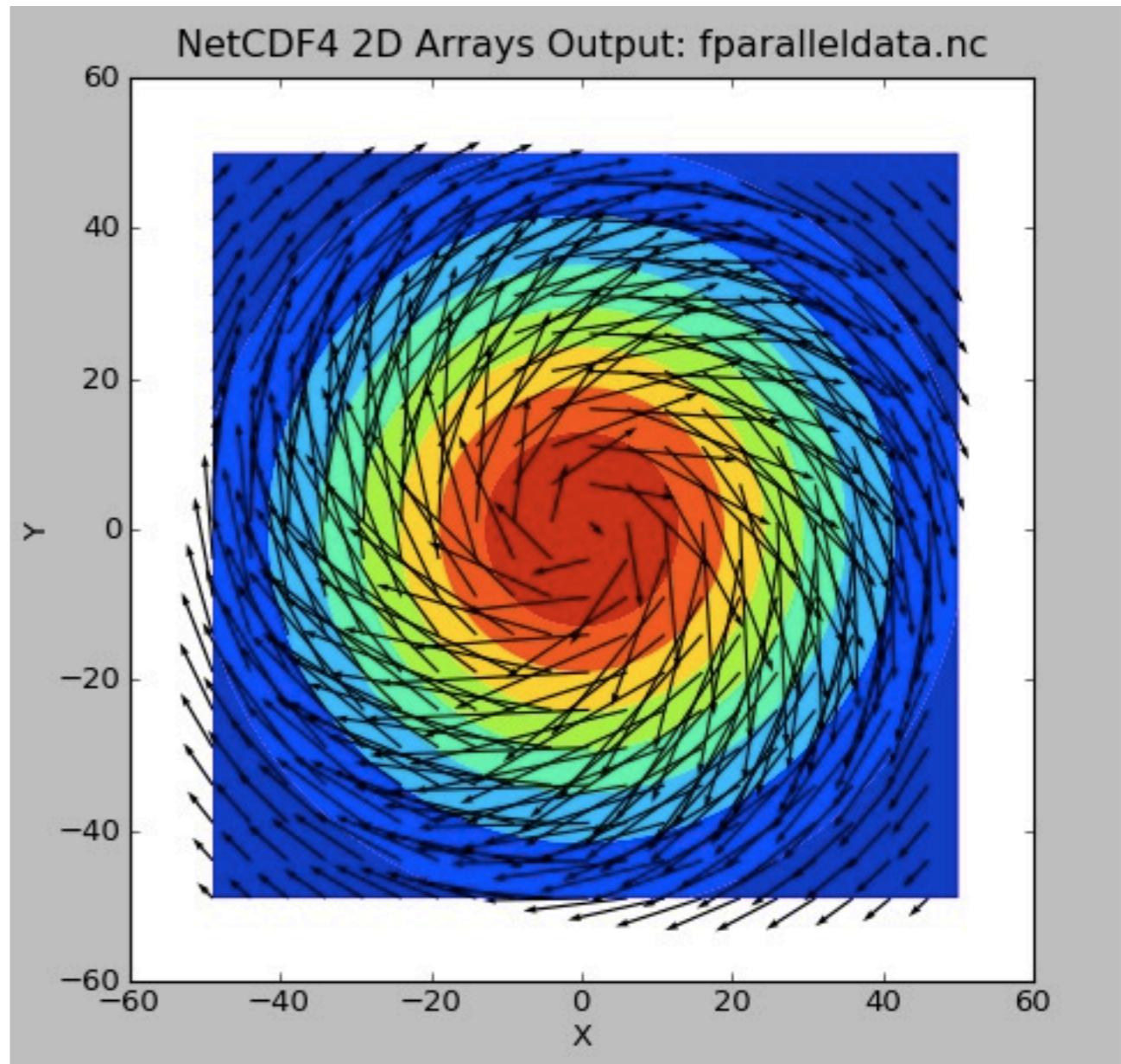
$ mpirun -np 4 parallel2darray

$ ls *.nc
$ source ../seriallibs
$../plots.py paralleldata.nc
```



# Sample Code

- Can do an ncdump -h...
- No trace of being written by different files
- Looks the same; code to read in is identical
- And not that much harder to code!
- By far the trickiest part is figuring out where in the file to write.



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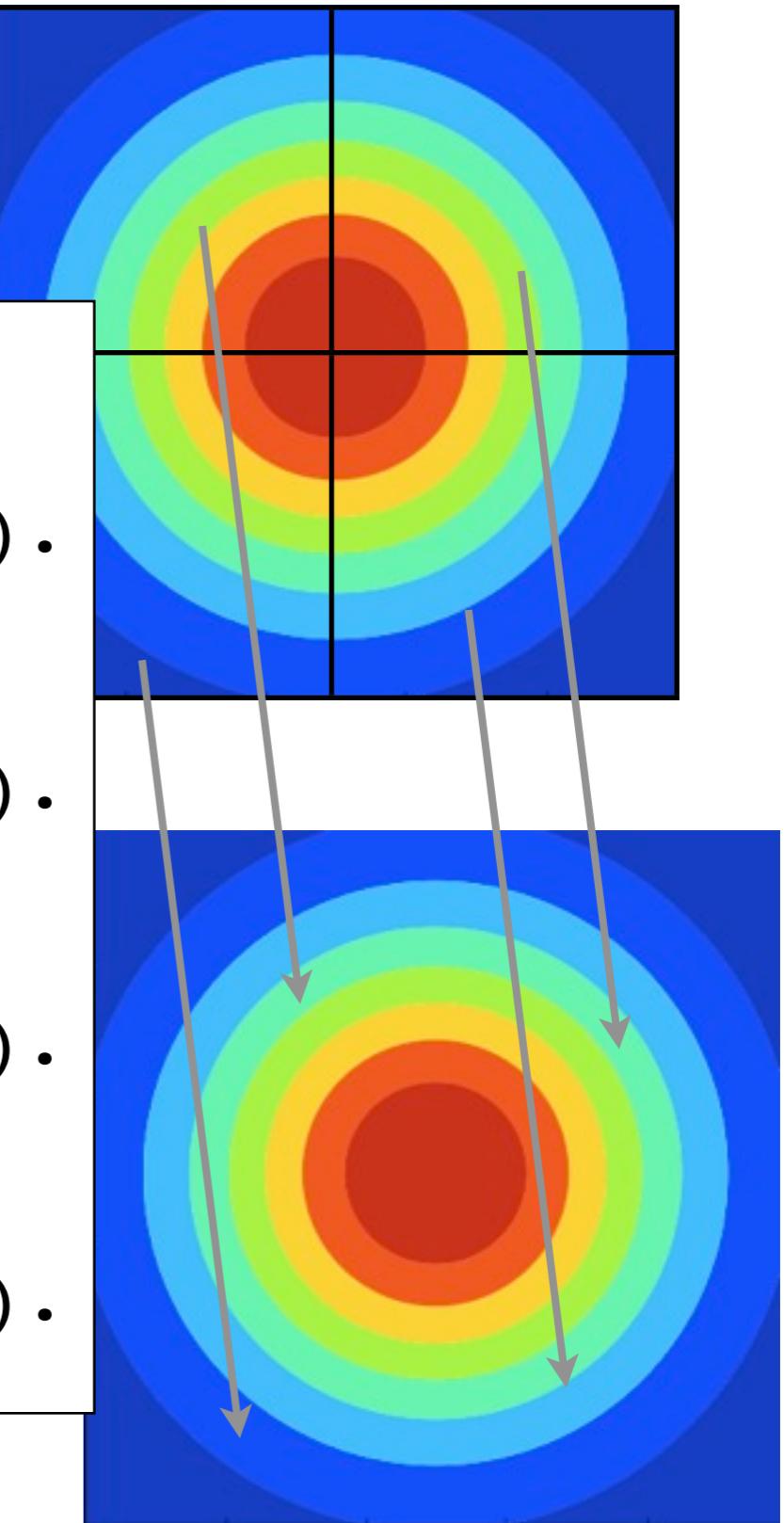
# Memory:

```
$ mpirun -np 4 ./fparallel2darray
[0] gets (0, 0): local points =
(50, 50); global points = (100,100).

[1] gets (1, 0): local points =
(50, 50); global points = (100,100).

[2] gets (0, 1): local points =
(50, 50); global points = (100,100).

[3] gets (1, 1): local points =
(50, 50); global points = (100,100).
```



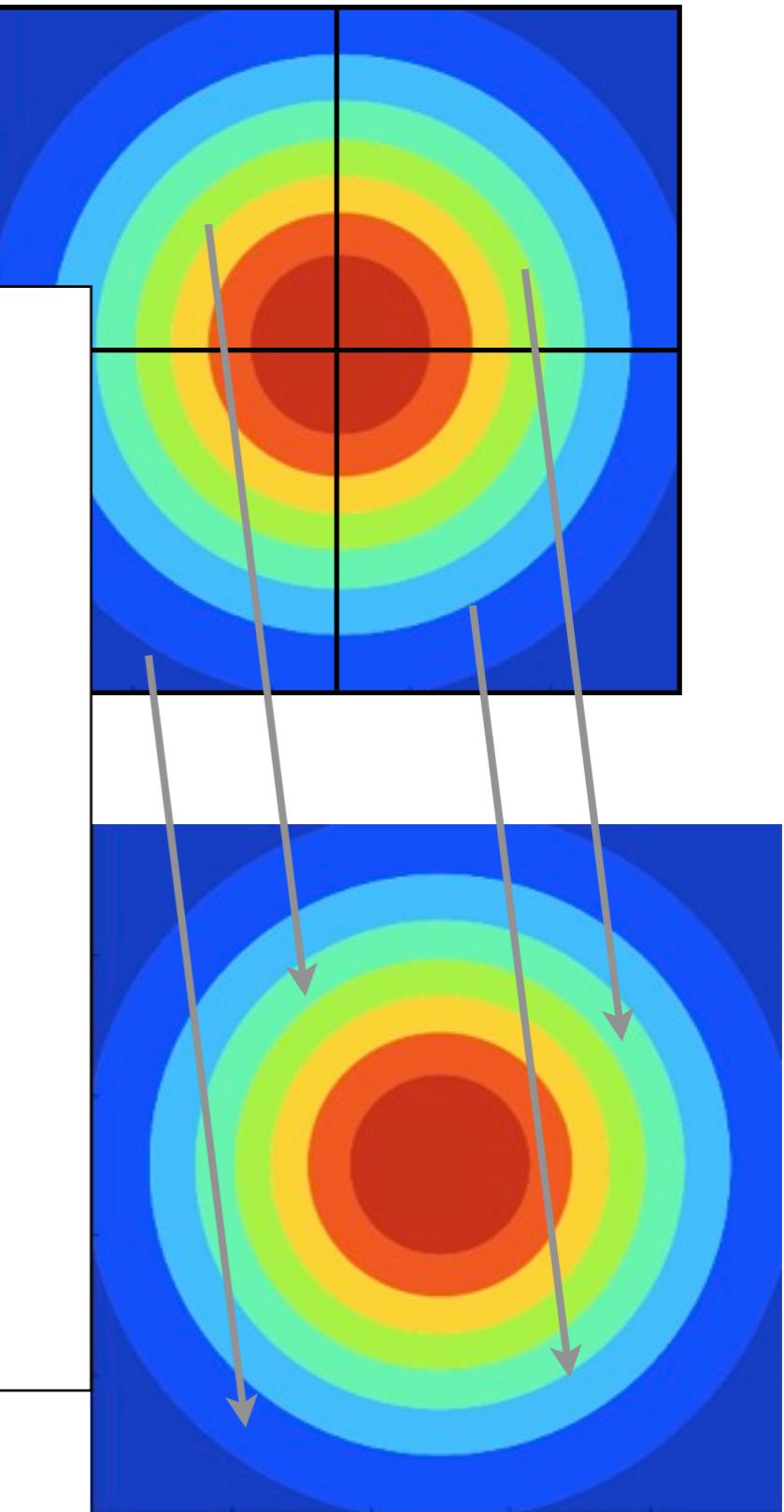
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# Memory:

```
[0]: denstarts, denscounts
 = 1 1 50 50
[1]: denstarts, denscounts
 = 51 1 50 50
[2]: denstarts, denscounts
 = 1 51 50 50
[3]: denstarts, denscounts
 = 51 51 50 50
```



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# fparallel2darray.f90

```
call MPI_Info_create(info, status)
call MPI_Info_set(info,"IBM_largeblock_io","true", status)

mode_flag = IOR(NF90_MPIIO, NF90_CLOBBER)
mode_flag = IOR(mode_flag, NF90_NETCDF4)
status = nf90_create_par(rundata%filename, mode_flag,
MPI_COMM_WORLD, info, file_id)

if (status /= NF90_NOERR) then
 print *, 'Could not open file ', rundata%filename
 return
endif
```

# fparallel2darray.f90

```
call MPI_Info_create(info, status)
call MPI_Info_set(info,"IBM_largeblock_io","true", status)

mode_flag = IOR(NF90_MPIIO, NF90_CLOBBER)
mode_flag = IOR(mode_flag, NF90_NETCDF4)
status = nf90_create_par(rundata%filename, mode_flag,
MPI_COMM_WORLD, info, file_id)

if (status /= NF90_NOERR) then
 print *, 'Could not open file ', rundata%filename
 return
endif
```



create\_par rather than create

# fparallel2darray.f90

```
call MPI_Info_create(info, status)
call MPI_Info_set(info,"IBM_largeblock_io","true", status)

mode_flag = IOR(NF90_MPIIO, NF90_CLOBBER)
mode_flag = IOR(mode_flag, NF90_NETCDF4)
status = nf90_create_par(rundata%filename, mode_flag,
MPI_COMM_WORLD, info, file_id)
if (status /= NF90_NOERR) then
 print *, 'Could not open file ', rundata%filename
 return
endif
```

**mode\_flag = CLOBBER | MPIIO | NETCDF4**

# fparallel2darray.f90

```
call MPI_Info_create(info, status)
call MPI_Info_set(info,"IBM_largeblock_io","true", status)

mode_flag = IOR(NF90_MPIIO, NF90_CLOBBER)
mode_flag = IOR(mode_flag, NF90_NETCDF4)
status = nf90_create_par(rundata%filename, mode_flag,
MPI_COMM_WORLD, info, file_id)
if (status = NF90_NOERR) then
 print *, 'Could not open file ', rundata%filename
 return
endif
```

Extra arguments: communicator that will do  
the I/O

# fparallel2darray.f90

```
call MPI_Info_create(info, status)
call MPI_Info_set(info,"IBM_largeblock_io","true", status)

mode_flag = IOR(NF90_MPIIO, NF90_CLOBBER)
mode_flag = IOR(mode_flag, NF90_NETCDF4)
status = nf90_create_par(rundata%filename, mode_flag,
MPI_COMM_WORLD, info, file_id)
if (status /= NF90_NOERR) then
 print *, 'Could not open file ', rundata%filename
 return
endif
```

Extra arguments: MPI Info; can pass MPI-I/O  
“hints”

# fparallel2darray.f90

```
status = nf90_def_dim(file_id, 'X', rundata%globalnx, xdim_id)
status = nf90_def_dim(file_id, 'Y', rundata%globalny, ydim_id)
status = nf90_def_dim(file_id, 'velocity components', 2,
vcomp_id)

! now that the dimensions are defined, define variables

densdims = (/ xdim_id, ydim_id /)
veldims = (/ vcomp_id, xdim_id, ydim_id /)

status = nf90_def_var(file_id, 'Density', NF90_DOUBLE, densdims,
dens_id)
status = nf90_def_var(file_id, 'Velocity', NF90_DOUBLE, veldims,
vel_id)
```

Defining variables identical (but global v local)



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# fparallel2darray.f90

```
status = nf90_var_par_access(file_id, dens_id, NF90_COLLECTIVE)
status = nf90_var_par_access(file_id, vel_id, NF90_COLLECTIVE)

status = nf90_put_var(file_id, dens_id, dens, start=densstarts,
count=denscounts)

status = nf90_put_var(file_id, vel_id, vel, start=velstarts,
count=velcounts)

status = nf90_close(file_id)
```



Define how we'll be accessing *variables* -  
**COLLECTIVE** vs **INDEPENDANT**.  
(eg, `Write_all` vs. `Write`).



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# fparallel2darray.f90

```
status = nf90_var_par_access(file_id, dens_id, NF90_COLLECTIVE)
status = nf90_var_par_access(file_id, vel_id, NF90_COLLECTIVE)

status = nf90_put_var(file_id, dens_id, dens, start=densstarts,
count=denscounts)

status = nf90_put_var(file_id, vel_id, vel, start=velstarts,
count=velcounts)

status = nf90_close(file_id)
```



**put\_var is exactly like serial with subsections -  
starts, counts**



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# fparallel2darray.f90

```
status = nf90_var_par_access(file_id, dens_id, NF90_COLLECTIVE)
status = nf90_var_par_access(file_id, vel_id, NF90_COLLECTIVE)

status = nf90_put_var(file_id, dens_id, dens, start=densstarts,
count=denscounts)
status = nf90_put_var(file_id, vel_id, vel, start=velstarts,
count=velcounts)

status = nf90_close(file_id)
```



close is the same as ever.



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**serial.c**

```
/* name of units for dens, vel */
const char *densunit="g/cm^3";
const char *velunit="cm/s";

/* return status */
int status;

/* set up x, y coordinates */
x = (float *)malloc(rundata.nx * sizeof(float));
y = (float *)malloc(rundata.ny * sizeof(float));
for (i=0; i<rundata.nx; i++)
 x[i] = (1.*i-rundata.nx/2.);
for (i=0; i<rundata.ny; i++)
 y[i] = (1.*i-rundata.ny/2.);

/* Create a new file - clobber anything existing */
status = nc_create(rundata.filename, NC_CLOBBER, &file_id);

/* netCDF routines return NC_NOERR on success */
if (status != NC_NOERR) {
 fprintf(stderr,"Could not open file %s\n", rundata.filename);
 return;
}

/* define the dimensions */
nc_def_dim(file_id, "X", rundata.nx, &xdim_id);
nc_def_dim(file_id, "Y", rundata.ny, &ydim_id);
nc_def_dim(file_id, "velocity component", 2, &vcomp_id);

/* define the coordinate variables,... */
```

**parallel.c**

```
/* name of units for dens, vel */
const char *densunit="g/cm^3";
const char *velunit="cm/s";

/* offsets for sub-regions of arrays */
size_t starts[3];
size_t counts[3];

/* return status */
int status;

/* MPI-IO hints for performance */
MPI_Info info;

/* set up x, y coordinates */
x = (float *)malloc(rundata.globalnx * sizeof(float));
y = (float *)malloc(rundata.globalny * sizeof(float));
for (i=0; i<rundata.globalnx; i++)
 x[i] = (1.*i-rundata.globalnx/2.);
for (i=0; i<rundata.globalny; i++)
 y[i] = (1.*i-rundata.globalny/2.);

/* set the MPI-IO hints for better performance on GPFS */
MPI_Info_create(&info);
MPI_Info_set(info, "IBM_largeblock_io", "true");

/* Create a new file - clobber anything existing */
status = nc_create_par(rundata.filename, NC_MPIIO|NC_CLOBBER|NC_COLLECTIVE,
 MPI_COMM_WORLD, info, &file_id);

/* netCDF routines return NC_NOERR on success */
if (status != NC_NOERR) {
 fprintf(stderr,"Could not open file %s\n", rundata.filename);
 return;
}

/* define the dimensions */
nc_def_dim(file_id, "X", rundata.globalnx, &xdim_id);
nc_def_dim(file_id, "Y", rundata.globalny, &ydim_id);
nc_def_dim(file_id, "velocity component", 2, &vcomp_id);

/* define the coordinate variables,... */
```

### serial.c

```
nc_def_var(file_id, "Density", NC_DOUBLE, 2, densdims, &dens_id);
nc_def_var(file_id, "Velocity", NC_DOUBLE, 3, veldims, &vel_id);

/* assign units to the variables */
nc_put_att_text(file_id, dens_id, "units", strlen(densunit), densunit);
nc_put_att_text(file_id, vel_id, "units", strlen(velunit), velunit);

/* we are now done defining variables and their attributes */
nc_enddef(file_id);

/* Write out the data to the variables we've defined */
nc_put_var_float(file_id, xcoord_id, x);
nc_put_var_float(file_id, ycoord_id, y);

nc_put_var_double(file_id, dens_id, &(dens[0][0]));
nc_put_var_double(file_id, vel_id, &(vel[0][0][0]));

nc_close(file_id);
return;
```

### parallel.c

```
nc_def_var(file_id, "Density", NC_DOUBLE, 2, densdims, &dens_id);
nc_def_var(file_id, "Velocity", NC_DOUBLE, 3, veldims, &vel_id);

/* assign units to the variables */
nc_put_att_text(file_id, dens_id, "units", strlen(densunit), densunit);
nc_put_att_text(file_id, vel_id, "units", strlen(velunit), velunit);

/* we are now done defining variables and their attributes */
nc_enddef(file_id);

/* Write out the data to the variables we've defined */
nc_put_var_float(file_id, xcoord_id, x);
nc_put_var_float(file_id, ycoord_id, y);

/* The big data will be written to collectively;
 * the alternative is NC_INDEPENDENT */
nc_var_par_access(file_id, dens_id, NC_COLLECTIVE);
nc_var_par_access(file_id, vel_id, NC_COLLECTIVE);

/* densities */
starts[0] = (rundata.globalnx/rundata.npx)*rundata.myx;
starts[1] = (rundata.globalny/rundata.npy)*rundata.myy;
counts[0] = rundata.localnx;
counts[1] = rundata.localny;

nc_put_vara_double(file_id, dens_id, starts, counts, &(dens[0][0]));

/* velocities */
starts[0] = 0;
starts[1] = (rundata.globalnx/rundata.npx)*rundata.myx;
starts[2] = (rundata.globalny/rundata.npy)*rundata.myy;
counts[0] = 2;
counts[1] = rundata.localnx;
counts[2] = rundata.localny;

nc_put_vara_double(file_id, vel_id, starts, counts, &(vel[0][0][0]));

nc_close(file_id);
return;
```



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SciNet



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# HDF5

# Hyperslabs

- Parallel HDF5 similar to parallel NetCDF - fairly modest changes to structure of code
- Different (more low-level, natch) way of dealing with sub-regions
- Offset, block, count, stride



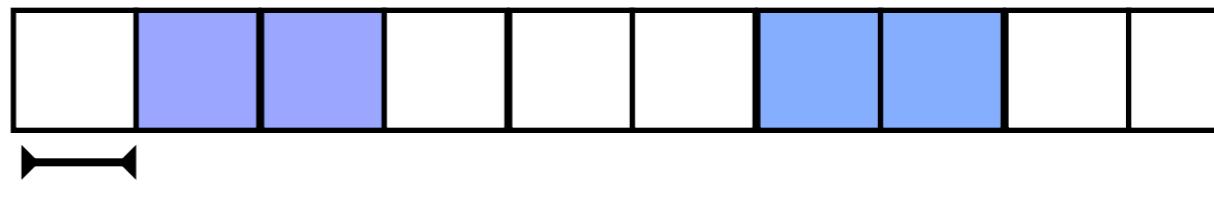
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# HDF5 Hyperslabs

- Parallel HDF5 similar to parallel NetCDF - fairly modest changes to structure of code
- Different (more low-level, natch) way of dealing with sub-regions
- Offset, block, count, stride



**Offset = 1**



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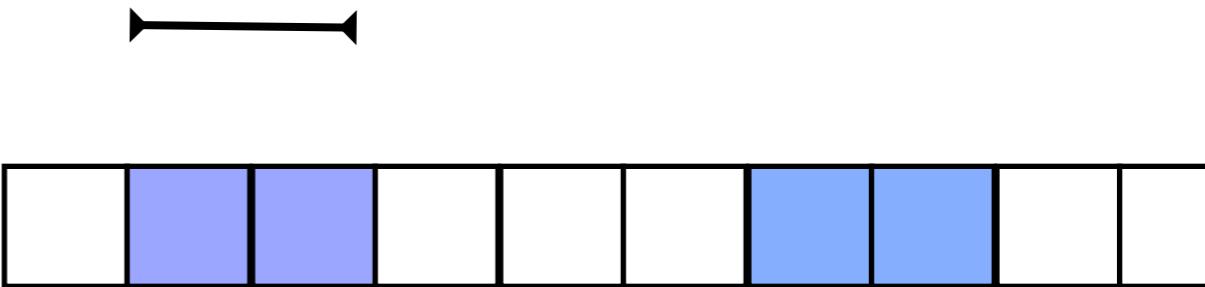


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# HDF5 Hyperslabs

- Parallel HDF5 similar to parallel NetCDF - fairly modest changes to structure of code
- Different (more low-level, natch) way of dealing with sub-regions
- Offset, block, count, stride

blocksize = 2



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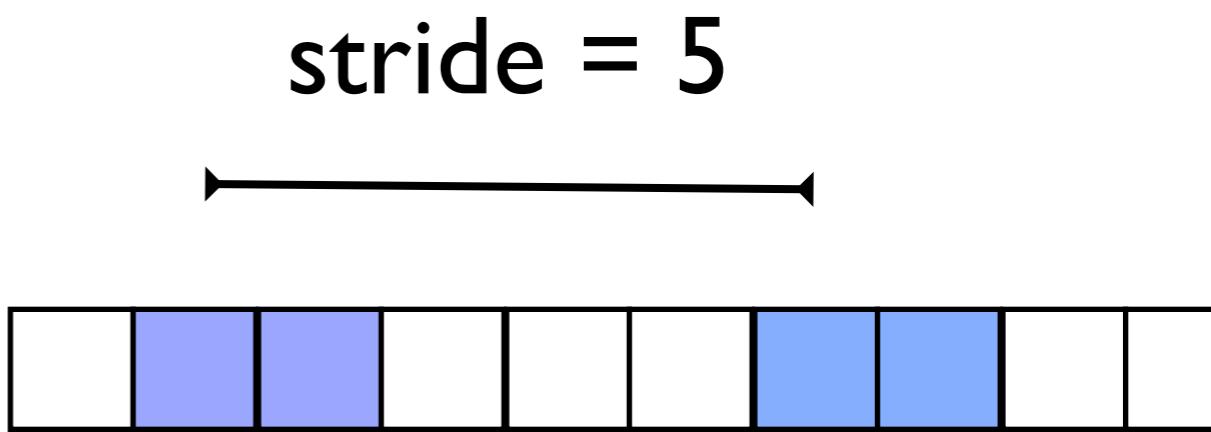


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# HDF5

## Hyperslabs

- Parallel HDF5 similar to parallel NetCDF - fairly modest changes to structure of code
- Different (more low-level, natch) way of dealing with sub-regions
- Offset, block, count, stride



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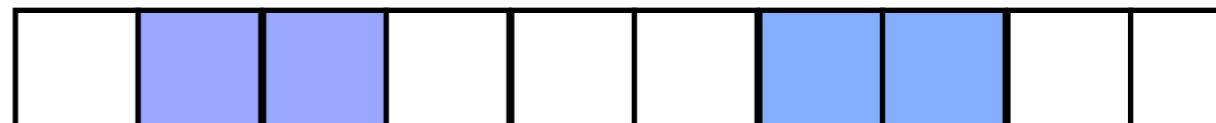


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# HDF5 Hyperslabs

count = 2

- Parallel HDF5 similar to parallel NetCDF - fairly modest changes to structure of code
- Different (more low-level, natch) way of dealing with sub-regions
- Offset, block, count, stride
- (MPI\_Type\_vector)



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# HDF5 Hyperslabs

count = 2



- Parallel HDF5 similar to parallel NetCDF - fairly modest changes to structure of code
- Different (more low-level, natch) way of dealing with sub-regions
- Offset, block, count, stride
- Hyperslab - one of these per dimensions.
- (offset,block) just like (start, counts) in netcdf.



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# parallel2darray.c

```
/* set the MPI-IO hints for better performance on GPFS */

MPI_Info_create(&info);
MPI_Info_set(info,"IBM_largeblock_io","true");

/* Set up the parallel environment for file access*/
fap_id = H5Pcreate(H5P_FILE_ACCESS);
/* Include the file access property with IBM hint */
H5Pset_fapl_mpio(fap_id, MPI_COMM_WORLD, info);

/* Set up the parallel environment */
dist_id = H5Pcreate(H5P_DATASET_XFER);
/* we'll be writing collectively */
H5Pset_dxpl_mpio(dist_id, H5FD_MPIO_COLLECTIVE);
```

# parallel2darray.c

```
/* set the MPI-IO hints for better performance on GPFS */
MPI_Info_create(&info);
MPI_Info_set(info,"IBM_largeblock_io","true");

/* Set up the parallel environment for file access*/
fap_id = H5Pcreate(H5P_FILE_ACCESS);
/* Include the file access property with IBM hint */
H5Pset_fapl_mpio(fap_id, MPI_COMM_WORLD, info);

/* Set up the parallel environment */
dist_id = H5Pcreate(H5P_DATASET_XFER);
/* we'll be writing collectively */
H5Pset_dxpl_mpio(dist_id, H5FD_MPIO_COLLECTIVE);
```



Same as NetCDF; this is a property of the *file*

# parallel2darray.c

```
/* set the MPI-IO hints for better performance on GPFS */
MPI_Info_create(&info);
MPI_Info_set(info,"IBM_largeblock_io","true");

/* Set up the parallel environment for file access*/
fap_id = H5Pcreate(H5P_FILE_ACCESS);
/* Include the file access property with IBM hint */
H5Pset_fapl_mpio(fap_id, MPI_COMM_WORLD, info);

/* Set up the parallel environment */
dist_id = H5Pcreate(H5P_DATASET_XFER);
/* we'll be writing collectively */
H5Pset_dxpl_mpio(dist_id, H5FD_MPIO_COLLECTIVE);
```

Collective/independant: this is a  
property of accessing a *variable*

## parallel2darray.c

```
offsets[0] = (rundata.globalnx/rundata.npx)*rundata.myx;
offsets[1] = (rundata.globalny/rundata.npy)*rundata.myy;
blocks[0] = rundata.localnx;
strides[0] = strides[1] = 1;
counts[0] = counts[1] = 1;

globaldensspace = H5Dget_space(dens_dataset_id);
H5Sselect_hyperslab(globaldensspace,H5S_SELECT_SET, offsets,
strides, counts, blocks);

status = H5Dwrite(dens_dataset_id, H5T_NATIVE_DOUBLE,
loc_dens_dataspace_id, globaldensspace, dist_id, &(dens[0]
[0]));
```

Select hyperslab, and write; parallelism is in  
distribution\_id



# Projects

```
$ cd parIO/hydro{c,f}
```

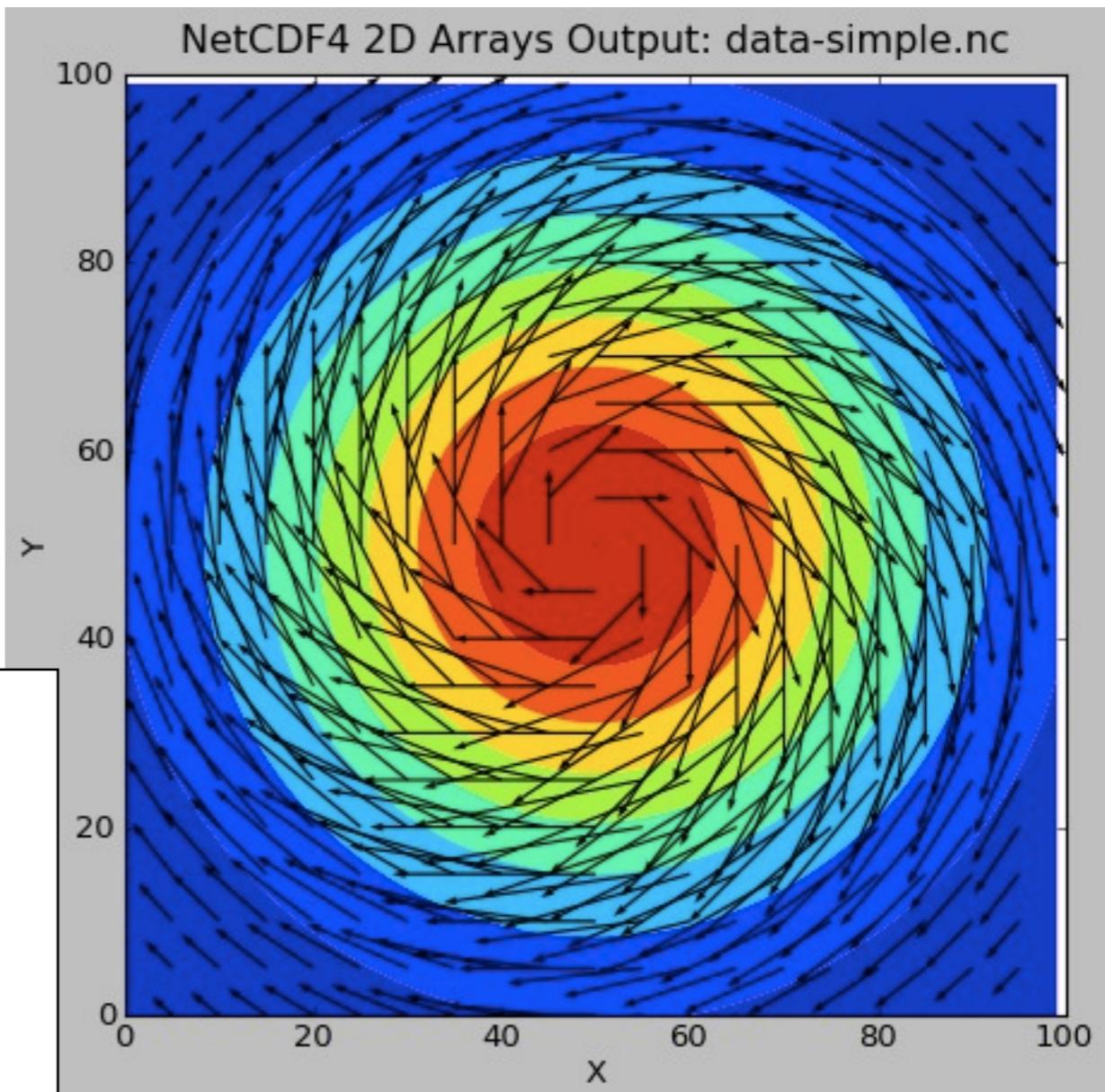
Write hdf5, netcdf outputs

```
$ cd parIO/hydro{c,f}-mpi
```

Write ppm output in MPI-IO, (started) and  
output in parallel hdf5, netcdf

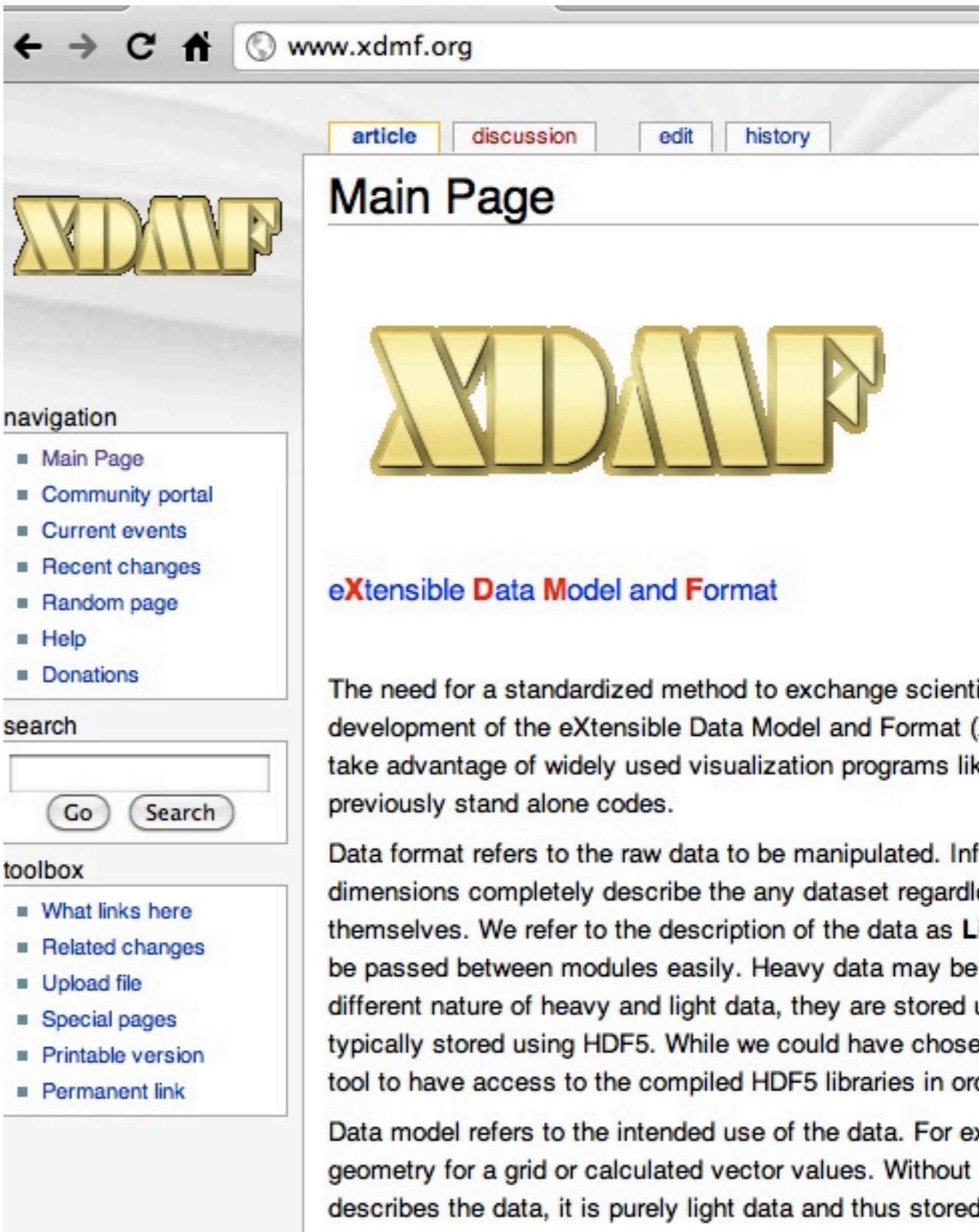
```
$ cd parIO/nbody
```

Write parallel hdf5, netcdf, MPI-IO outputs for  
gravitational particles (FORTRAN)



# Conventions for HDF5

- XDMF
- An XML description of your HDF5 files
- A way of encoding “conventions” for HDF5
- Important for interoperability (eg, w/ viz packages)



The screenshot shows a web browser displaying the XDMF.org main page. The URL in the address bar is www.xdmf.org. The page features a large, stylized XDMF logo at the top. Below it, there's a navigation menu with links to Main Page, Community portal, Current events, Recent changes, Random page, Help, and Donations. There's also a search bar with 'Go' and 'Search' buttons, and a toolbox with links to What links here, Related changes, Upload file, Special pages, Printable version, and Permanent link. The main content area contains the XDMF logo again, followed by the text "eXtensible Data Model and Format". It explains the need for a standardized method to exchange scientific data and the development of the eXtensible Data Model and Format (XDMF) to take advantage of widely used visualization programs like Paraview, which previously required stand-alone codes. It also discusses the difference between data format (raw data to be manipulated) and data model (intended use of the data), mentioning the use of HDF5 for heavy data and XML for light data.

navigation

- Main Page
- Community portal
- Current events
- Recent changes
- Random page
- Help
- Donations

search

toolbox

- What links here
- Related changes
- Upload file
- Special pages
- Printable version
- Permanent link

**Main Page**

eXtensible Data Model and Format

The need for a standardized method to exchange scientific data has led to the development of the eXtensible Data Model and Format (XDMF). This allows users to take advantage of widely used visualization programs like Paraview, which previously required stand alone codes.

Data format refers to the raw data to be manipulated. In XDMF, the data is described by its type and dimensions. These dimensions completely describe the any dataset regardless of their physical representation. We refer to the description of the data as **Light Data**. Light data can be passed between modules easily. Heavy data may be represented by a pointer to a file or a reference to a memory location. The different nature of heavy and light data, they are stored using different methods. Heavy data is typically stored using HDF5. While we could have chosen to store heavy data using XML, it would have been difficult for a tool to have access to the compiled HDF5 libraries in order to manipulate the data.

Data model refers to the intended use of the data. For example, a dataset may contain geometry for a grid or calculated vector values. Without this information, the data is purely light data and thus stored using XML.



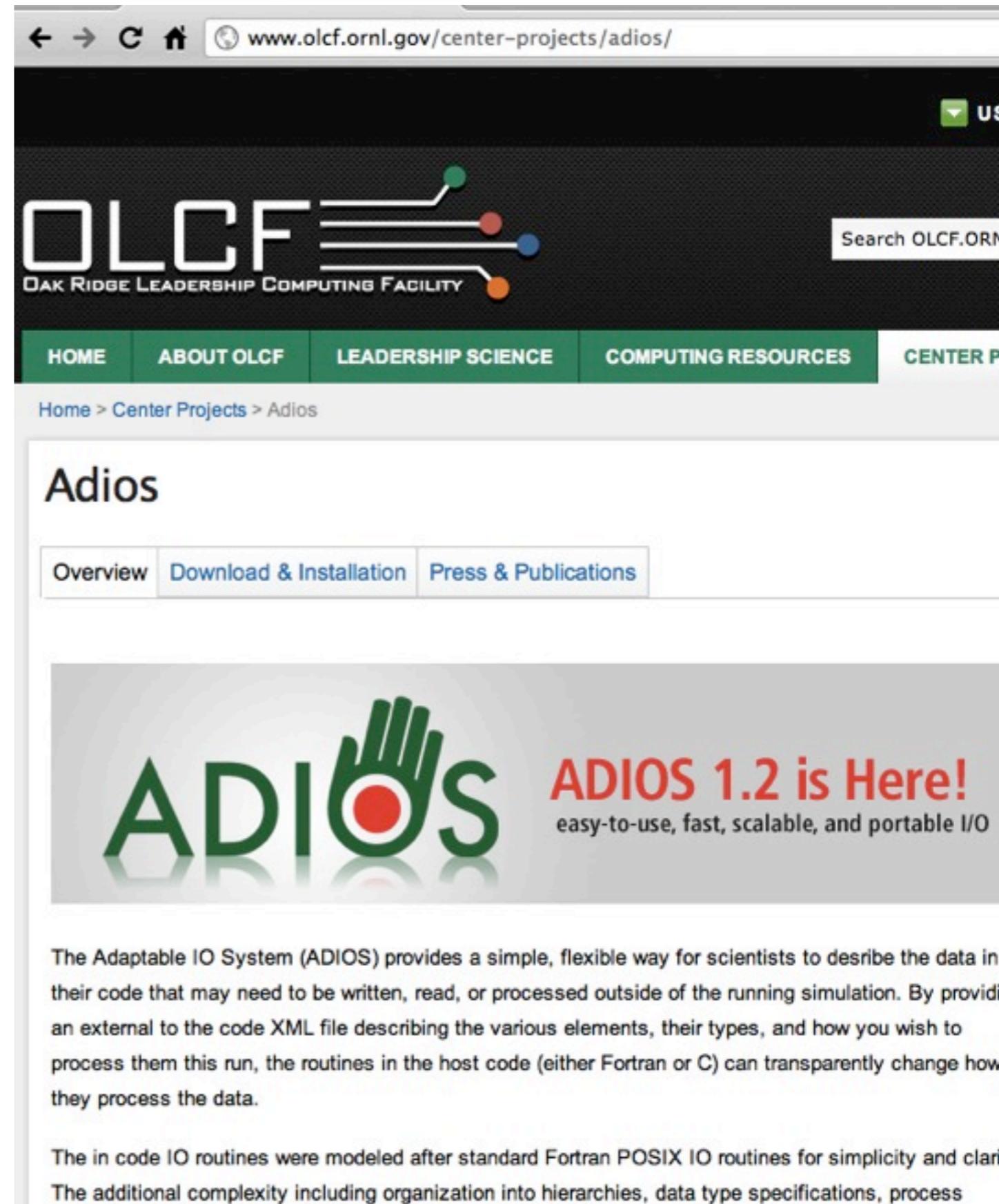
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# Adaptable IO System

- ADIOS
- A library for IO for scientific code
- Uses MPIIO, HDF5, etc... under the hood
- Allows changing of IO strategy, method; no rewriting code and maybe not even a recompile.



The screenshot shows a web browser displaying the OLCF (Oak Ridge Leadership Computing Facility) website at [www.olcf.ornl.gov/center-projects/adios/](http://www.olcf.ornl.gov/center-projects/adios/). The page features the OLCF logo with three colored dots (green, red, blue) connected by lines. The navigation menu includes links for HOME, ABOUT OLCF, LEADERSHIP SCIENCE, COMPUTING RESOURCES, and CENTER P. Below the menu, a breadcrumb trail shows Home > Center Projects > Adios. The main content area is titled "Adios" and includes tabs for Overview, Download & Installation, and Press & Publications. A large green "ADIOS" logo is prominently displayed. To the right, a red banner announces "ADIOS 1.2 is Here! easy-to-use, fast, scalable, and portable I/O". Below the banner, a paragraph describes the Adaptable IO System (ADIOS) as providing a simple, flexible way for scientists to describe data in their code, using an external XML file to define elements and processing logic. It notes that in-code IO routines are modeled after standard Fortran POSIX IO routines for simplicity and clarity.



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# parIO/adios/parallel2darray.{c,f90}

```
void writeadiosfile(rundata_t *rundata, double **dens, double ***vel) {
 int adios_err=0;
 uint64_t adios_groupsize, adios_totalsize;
 int64_t adios_handle;
 MPI_Comm comm = MPI_COMM_WORLD;
 int size;

 MPI_Comm_size(comm, &size);

 adios_init ("adios_global.xml");
 adios_open (&adios_handle, "ArrayData", rundata->filename, "w", &comm);
 #include "gwrite_ArrayData.ch"

 if (adios_err)
 fprintf(stderr,"Error doing adios write.\n");

 adios_close (adios_handle);
}
```



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# parIO/adios/{}array\_global.xml

```
<?xml version="1.0"?>
<adios-config host-language="C">

 <adios-group name="ArrayData" coordination-communicator="comm">
 <var name="rundata->localnx" type="integer" />
 <var name="rundata->localny" type="integer" />
 <var name="rundata->globalnx" type="integer" />
 <var name="rundata->globalny" type="integer" />
 <var name="rundata->startx" type="integer" />
 <var name="rundata->starty" type="integer" />
 <var name="size" type="integer" />
 <global-bounds dimensions="2,rundata->globalnx,rundata->globalny"
 offsets="0,rundata->startx,rundata->starty">
 <var name="vel" gwrite="vel[0][0]" type="double"
 dimensions="2,rundata->localnx,rundata->localny" />
 </global-bounds>
 <global-bounds dimensions="rundata->globalnx,rundata->globalny"
 offsets="rundata->startx,rundata->starty">
 <var name="dens" gwrite="dens[0]" type="double"
 dimensions="rundata->localnx,rundata->localny" />
 </global-bounds>
 </adios-group>

 <method group="ArrayData" method="PHDF5" />

 <buffer size-MB="2" allocate-time="now" />

</adios-config>
```



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# ADIOS workflow

- Write XML file describing data, layout
- gpp.py [file].xml - generates C or Fortran code: adios calls, size calculation
- Build code
- Separates data layout, code.

```
<?xml version="1.0"?>
<adios-config host-language="C">
 <adios-group name="ArrayData" coordination-communicator="MPI"
 <var name="rundata->localnx" type="integer" />
 <var name="rundata->localny" type="integer" />
 <var name="rundata->globalnx" type="integer" />
 <var name="rundata->globalny" type="integer" />
 <var name="rundata->startx" type="integer" />
 <var name="rundata->starty" type="integer" />
 <var name="size" type="integer" />
 <global-bounds dimensions="2,rundata->globalnx,rundata->globalny">
 <offsets>0,rundata->startx,rundata->starty</offsets>
 <var name="vel" gwrite="vel[0][0]" type="double" />
 <dimensions>2,rundata->localnx,rundata->localny</dimensions>
 </global-bounds>
 <global-bounds dimensions="rundata->globalnx,rundata->globalny">
 <offsets>rundata->startx,rundata->starty</offsets>
 <var name="dens" gwrite="dens[0]" type="double" />
 <dimensions>rundata->localnx,rundata->localny</dimensions>
 </global-bounds>
 </adios-group>
<method group="ArrayData" method="PHDF5" />
<buffer size-MB="2" allocate-time="now" />
</adios-config>
```

```
void writeadiosfile(rundata_t *rundata, double **dens, double *vel)
{
 int adios_err=0;
 uint64_t adios_groupsize, adios_totalsize;
 int64_t adios_handle;
 MPI_Comm comm = MPI_COMM_WORLD;
 int size;

 MPI_Comm_size(comm, &size);

 adios_init ("adios_global.xml");
 adios_open (&adios_handle, "ArrayData", rundata->filename);
 #include "gwrite_ArrayData.ch"

 if (adios_err)
 fprintf(stderr,"Error doing adios write.\n");

 adios_close (adios_handle);
}
```



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# ADIOS workflow

- Separation isn't perfect; xml file references code variables, etc.
- But allows “componentization” of I/O.
- Changes that don't result in changes to grwrite\_Array.ch don't require recompilation (eg, only changing number, size of variables in group).

```
<?xml version="1.0"?>
<adios-config host-language="C">
 <adios-group name="ArrayData" coordination-communicator="MPI"
 <var name="rundata->localnx" type="integer" />
 <var name="rundata->localny" type="integer" />
 <var name="rundata->globalnx" type="integer" />
 <var name="rundata->globalny" type="integer" />
 <var name="rundata->startx" type="integer" />
 <var name="rundata->starty" type="integer" />
 <var name="size" type="integer" />
 <global-bounds dimensions="2,rundata->globalnx,rundata->globalny"
 offsets="0,rundata->startx,rundata->starty"
 <var name="vel" gwrite="vel[0][0]" type="double"
 dimensions="2,rundata->localnx,rundata->localny" />
 </global-bounds>
 <global-bounds dimensions="rundata->globalnx,rundata->globalny"
 offsets="rundata->startx,rundata->starty"
 <var name="dens" gwrite="dens[0]" type="double"
 dimensions="rundata->localnx,rundata->localny" />
 </global-bounds>
 </adios-group>
<method group="ArrayData" method="PHDF5" />
<buffer size-MB="2" allocate-time="now" />
</adios-config>
```

```
void writeadiosfile(rundata_t *rundata, double **dens, double
 int adios_err=0;
 uint64_t adios_groupsize, adios_totalsize;
 int64_t adios_handle;
 MPI_Comm comm = MPI_COMM_WORLD;
 int size;

 MPI_Comm_size(comm, &size);

 adios_init ("adios_global.xml");
 adios_open (&adios_handle, "ArrayData", rundata->filename);
 #include "grwrite_ArrayData.ch"

 if (adios_err)
 fprintf(stderr,"Error doing adios write.\n");

 adios_close (adios_handle);
}
```



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# Variable Groups

- Multiple groups of variables possible: (eg) restart files vs. files for analysis
- Variables can appear in multiple groups
- Each group can be handled with different methods

```
<?xml version="1.0"?>
<adios-config host-language="C">
 <adios-group name="ArrayData" coordination-communicator="MPI"
 <var name="rundata->localnx" type="integer" />
 <var name="rundata->localny" type="integer" />
 <var name="rundata->globalnx" type="integer" />
 <var name="rundata->globalny" type="integer" />
 <var name="rundata->startx" type="integer" />
 <var name="rundata->starty" type="integer" />
 <var name="size" type="integer" />
 <global-bounds dimensions="2,rundata->globalnx,rundata->globalny">
 <offsets>0,rundata->startx,rundata->starty</offsets>
 <var name="vel" gwrite="vel[0][0]" type="double" dimensions="2,rundata->localnx,rundata->localny">
 </global-bounds>
 <global-bounds dimensions="rundata->globalnx,rundata->globalny">
 <offsets>rundata->startx,rundata->starty</offsets>
 <var name="dens" gwrite="dens[0]" type="double" dimensions="rundata->localnx,rundata->localny">
 </global-bounds>
 </adios-group>
<method group="ArrayData" method="PHDF5" />
<buffer size-MB="2" allocate-time="now" />
</adios-config>
```

```
void writeadiosfile(rundata_t *rundata, double **dens, double *vel)
{
 int adios_err=0;
 uint64_t adios_groupsize, adios_totalsize;
 int64_t adios_handle;
 MPI_Comm comm = MPI_COMM_WORLD;
 int size;

 MPI_Comm_size(comm, &size);

 adios_init ("adios_global.xml");
 adios_open (&adios_handle, "ArrayData", rundata->filename);
 #include "gwrite_ArrayData.ch"

 if (adios_err)
 fprintf(stderr,"Error doing adios write.\n");

 adios_close (adios_handle);
}
```



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# I/O methods

- Possible methods: parallel HDF5 (PHDF5), NetCDF (NC4), one-per-process posix files (POSIX), it's own native format (BP) using MPI-IO (MPI)
- Change between methods: edit xml file, that's it.
- P-I (PHDF5, NC4, MPI), P-P (POSIX), or even P-M possible (PHDF5, etc with multiple communicators)

```
<?xml version="1.0"?>
<adios-config host-language="C">
 <adios-group name="ArrayData" coordination-communicator="<var name="rundata->localnx" type="integer" />
 <var name="rundata->localny" type="integer" />
 <var name="rundata->globalnx" type="integer" />
 <var name="rundata->globalny" type="integer" />
 <var name="rundata->startx" type="integer" />
 <var name="rundata->starty" type="integer" />
 <var name="size" type="integer" />
 <global-bounds dimensions="2,rundata->globalnx,rundata->globalny" offsets="0,rundata->startx,rundata->starty" type="int" />
 <var name="vel" gwrite="vel[0][0]" type="double" dimensions="2,rundata->localnx,rundata->localny" />
</global-bounds>
 <global-bounds dimensions="rundata->globalnx,rundata->globalny" offsets="rundata->startx,rundata->starty" type="int" />
 <var name="dens" gwrite="dens[0]" type="double" dimensions="rundata->localnx,rundata->localny" />
</global-bounds>
</adios-group>
<method group="ArrayData" method="PHDF5" />
<buffer size-MB="2" allocate-time="now" />
</adios-config>
```

```
void writeadiosfile(rundata_t *rundata, double **dens, double *vel)
{
 int adios_err=0;
 uint64_t adios_groupsize, adios_totalsize;
 int64_t adios_handle;
 MPI_Comm comm = MPI_COMM_WORLD;
 int size;

 MPI_Comm_size(comm, &size);

 adios_init ("adios_global.xml");
 adios_open (&adios_handle, "ArrayData", rundata->filename);
 #include "gwrite_ArrayData.ch"

 if (adios_err)
 fprintf(stderr,"Error doing adios write.\n");

 adios_close (adios_handle);
}
```



# Simplifies IO code

- Even if you aren't planning to switch between IO strategies, can greatly simplify code
- Many mechanical steps (eg, pasting together rectangular multi-dimentional arrays) done for you.
- Eliminates tedious, error-prone boilerplate code

```
<?xml version="1.0"?>
<adios-config host-language="C">
 <adios-group name="ArrayData" coordination-communicator="MPI">
 <var name="rundata->localnx" type="integer" />
 <var name="rundata->localny" type="integer" />
 <var name="rundata->globalnx" type="integer" />
 <var name="rundata->globalny" type="integer" />
 <var name="rundata->startx" type="integer" />
 <var name="rundata->starty" type="integer" />
 <var name="size" type="integer" />
 <global-bounds dimensions="2,rundata->globalnx,rundata->globalny">
 <var name="vel" gwrite="vel[0][0]" type="double" dimensions="2,rundata->localnx,rundata->localny" />
 </global-bounds>
 <global-bounds dimensions="rundata->globalnx,rundata->globalny">
 <var name="dens" gwrite="dens[0]" type="double" dimensions="rundata->localnx,rundata->localny" />
 </global-bounds>
 </adios-group>
<method group="ArrayData" method="PHDF5" />
<buffer size-MB="2" allocate-time="now"/>
</adios-config>
```

```
void writeadiosfile(rundata_t *rundata, double **dens, double *vel)
{
 int adios_err=0;
 uint64_t adios_groupsize, adios_totalsize;
 int64_t adios_handle;
 MPI_Comm comm = MPI_COMM_WORLD;
 int size;

 MPI_Comm_size(comm, &size);

 adios_init ("adios_global.xml");
 adios_open (&adios_handle, "ArrayData", rundata->filename);
 #include "gwrite_ArrayData.ch"

 if (adios_err)
 fprintf(stderr,"Error doing adios write.\n");

 adios_close (adios_handle);
}
```



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```
void writeadiosfile(rundata_t *rundata, double **dens, double ***vel) {
 int adios_err=0;
 uint64_t adios_groupsize, adios_totalsize;
 int64_t adios_handle;
 MPI_Comm comm = MPI_COMM_WORLD;
 int size;

 MPI_Comm_size(comm, &size);

 adios_init ("adios_global.xml");
 adios_open (&adios_handle, "ArrayData", rundata->filename, "w", &comm);
 #include "gwrite_ArrayData.ch"

 if (adios_err)
 fprintf(stderr,"Error doing adios write.\n");

 adios_close (adios_handle);
}
```



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```
void writehdf5file(rundata_t rundata, double **dens, double ***vel) {
 /* identifiers */
 hid_t file_id, arr_group_id, dens_dataset_id, vel_dataset_id;
 hid_t dens_dataspace_id, vel_dataspace_id;
 hid_t loc_dens_dataspace_id, loc_vel_dataspace_id;
 hid_t globaldensspace,globalvelspace;
 hid_t dist_id;
 hid_t fap_id;

 /* sizes */
 hsize_t densdims[2], veldims[3];
 hsize_t locdensdims[2], locveldims[3];

 /* status */
 herr_t status;

 /* MPI-IO hints for performance */
 MPI_Info info;

 /* parameters of the hyperslab */
 hsize_t counts[3];
 hsize_t strides[3];
 hsize_t offsets[3];
 hsize_t blocks[3];
```

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```

/* set the MPI-IO hints for better performance on GPFS */
MPI_Info_create(&info);
MPI_Info_set(info,"IBM_largeblock_io","true");

/* Set up the parallel environment for file access*/
fap_id = H5Pcreate(H5P_FILE_ACCESS);
/* Include the file access property with IBM hint */
H5Pset_fapl_mpio(fap_id, MPI_COMM_WORLD, info);

/* Set up the parallel environment */
dist_id = H5Pcreate(H5P_DATASET_XFER);
/* we'll be writing collectively */
H5Pset_dxpl_mpio(dist_id, H5FD_MPIO_COLLECTIVE);

/* Create a new file - truncate anything existing, use default properties */
file_id = H5Fcreate(rundata.filename, H5F_ACC_TRUNC, H5P_DEFAULT, fap_id);

/* HDF5 routines generally return a negative number on failure.
 * Should check return values! */
if (file_id < 0) {
 fprintf(stderr,"Could not open file %s\n", rundata.filename);
 return;
}

```



```

/* Create a new group within the new file */
arr_group_id = H5Gcreate(file_id,"/ArrayData", H5P_DEFAULT, H5P_DEFAULT,
 H5P_DEFAULT);

/* Give this group an attribute listing the time of calculation */
{
 hid_t attr_id,attr_sp_id;
 struct tm *t;
 time_t now;
 int yyyyymm;
 now = time(NULL);
 t = localtime(&now);
 yyyyymm = (1900+t->tm_year)*100+t->tm_mon;

 attr_sp_id = H5Screate(H5S_SCALAR);
 attr_id = H5Acreate(arr_group_id, "Calculated on (YYYYMM)", H5T_STD_U32LE,
 attr_sp_id, H5P_DEFAULT, H5P_DEFAULT);
 printf("yyymm = %d\n",yyyyymm);
 H5Awrite(attr_id, H5T_NATIVE_INT, &yyyyymm);
 H5Aclose(attr_id);
 H5Sclose(attr_sp_id);
}

/* Create the data space for the two global datasets. */
densdims[0] = rundata.globalnx; densdims[1] = rundata.globalny;
veldims[0] = 2; veldims[1] = rundata.globalnx; veldims[2] = rundata.globalny;

```



```

dens_dataspace_id = H5Screate_simple(2, densdims, NULL);
vel_dataspace_id = H5Screate_simple(3, veldims, NULL);

/* Create the datasets within the file.
 * H5T_IEEE_F64LE is a standard (IEEE) double precision (64 bit) floating (F) data
 * and will work on any machine. H5T_NATIVE_DOUBLE would work too, but would give
 * different results on GPC and TCS */

dens_dataset_id = H5Dcreate(file_id, "/ArrayData/dens", H5T_IEEE_F64LE,
 dens_dataspace_id, H5P_DEFAULT, H5P_DEFAULT, H5P_DEFAULT);
vel_dataset_id = H5Dcreate(file_id, "/ArrayData/vel", H5T_IEEE_F64LE,
 vel_dataspace_id, H5P_DEFAULT, H5P_DEFAULT, H5P_DEFAULT);

/* Now create the data space for our sub-regions. These are the data spaces
 * of our actual local data in memory. */
locdensdims[0] = rundata.localnx; locdensdims[1] = rundata.localny;
locveldims[0] = 2; locveldims[1] = rundata.localnx; locveldims[2] = rundata.localny;

loc_dens_dataspace_id = H5Screate_simple(2, locdensdims, NULL);
loc_vel_dataspace_id = H5Screate_simple(3, locveldims, NULL);

```



```

/*
 *
 * Now we have to figure out the `hyperslab' within the global
 * data that corresponds to our local data.
 *
 * Hyperslabs are described by an array of counts, strides, offsets,
 * and block sizes.
 *
 * | -offx-- |
 * +-----+----+-----+ +-+
 * | | | |
 * | | | offy
 * | | | |
 * - +----+ - +-+
 * | | | |
 * | | | localny
 * | | | |
 * - +----+ - +-+
 * | | | |
 * +-----+----+-----+
 * localnx
 *
 * In this case the blocksizes are (localnx,localny) and the offsets are
 * (offx,offy) = ((myx)/nyp*globalnx, (myy/nyp)*globalny)
 */

```

```

offsets[0] = (rundata.globalnx/rundata.npx)*rundata.myx;
offsets[1] = (rundata.globalny/rundata.npy)*rundata.myy;
blocks[0] = rundata.localnx;
blocks[1] = rundata.localny;
strides[0] = strides[1] = 1;
counts[0] = counts[1] = 1;

/* select this subset of the density variable's space in the file */
globaldensspace = H5Dget_space(dens_dataset_id);
H5Sselect_hyperslab(globaldensspace,H5S_SELECT_SET, offsets, strides, counts, block);

/* For the velocities, it's the same thing but there's a count of two,
 * (one for each velocity component) */

offsets[1] = (rundata.globalnx/rundata.npx)*rundata.myx;
offsets[2] = (rundata.globalny/rundata.npy)*rundata.myy;
blocks[1] = rundata.localnx;
blocks[2] = rundata.localny;
strides[0] = strides[1] = strides[2] = 1;
counts[0] = 2; counts[1] = counts[2] = 1;
offsets[0] = 0;
blocks[0] = 1;

```



```
globalvinspace = H5Dget_space(vel_dataset_id);
H5Sselect_hyperslab(globalvinspace,H5S_SELECT_SET, offsets, strides, counts, blocks

/* Write the data. We're writing it from memory, where it is saved
 * in NATIVE_DOUBLE format */
status = H5Dwrite(dens_dataset_id, H5T_NATIVE_DOUBLE, loc_dens_dataspace_id, global
dist_id, &(dens[0][0]));
status = H5Dwrite(vel_dataset_id, H5T_NATIVE_DOUBLE, loc_vel_dataspace_id, globalv
dist_id, &(vel[0][0][0]));

/* End access to groups & data sets and release resources used by them */
status = H5Sclose(dens_dataspace_id);
status = H5Dclose(dens_dataset_id);
status = H5Sclose(vel_dataspace_id);
status = H5Dclose(vel_dataset_id);
status = H5Gclose(arr_group_id);
status = H5Pclose(fap_id);
status = H5Pclose(dist_id);

/* Close the file */
status = H5Fclose(file_id);
return;
```



# ADIOS hands-on:

- Modify XML file, try outputting with method of MPI (use bpls or bp2hdf on resulting file), POSIX (Netcdf won't work at this point)
- Try a different IO strategy; do contiguous parallel IO by having single file but with each process' data contiguous in file, one after another. With large file size (`--nx=10000 --ny=10000`) and 8/16 processes, what are the timings between “straight” PHDF5, MPI, POSIX, and this approach? (And how long would it have taken you to do this without ADIOS?)
- Advanced: Break up `MPI_COMM_WORLD` into 2 communicators using `MPI_Comm_split`, call the new communicator `comm`, and output 2 files from the 8/16 processes using PHDF5 or MPI.



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