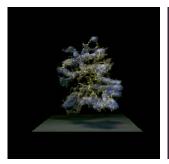
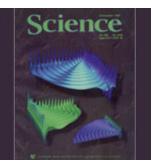
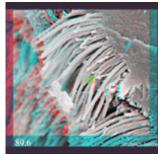
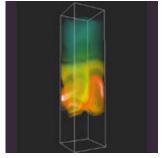


Using HDF5 for Scientific Data Analysis









NERSC Visualization Group





Before We Get Started Glossary of Terms

- Data
 - The raw information expressed in numerical form
- Metadata
 - Ancillary information about your data
- Attribute
 - Key/Value pair for accessing metadata
- Data Model / Schema
 - The choice of metadata and hierarchical organization of data to express higher-level relationships and features of a dataset.
- Self-Describing File
 - A file format that embeds explicit/queryable information about the data model
- Parallel I/O
 - Access to a single logical file image across all tasks of a parallel application that scales with the number of tasks.



- Architecture Independent self-describing binary file format
 - Well-defined APIs for specific data schemas (RIG, RIS, SDS)
 - Support for wide variety of binary FP formats (Cray Float, Convex Float, DEC Float, IEEE Float, EDBIC)
 - C and Fortran bindings
 - Very limited support for parallelism (CM-HDF, EOS-HDF/PANDA)
 - Not thread-safe
- Relationship with Unidata NetCDF
 - Similar data model for HDF Scientific Data Sets (SDS)
 - Interoperable with NetCDF

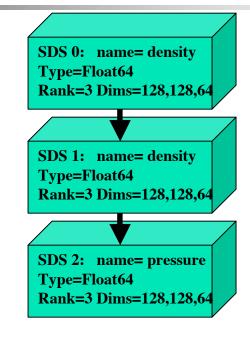






HDF4 SDS data schema

- Datasets
 - Name
 - Datatype
 - Rank,Dims



Datasets are inserted sequentially to the file

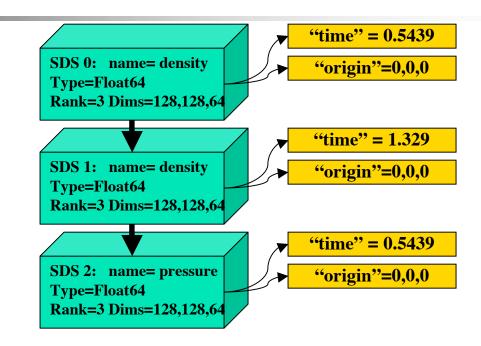




4

HDF4 SDS data schema

- Datasets
 - Name
 - Datatype
 - Rank, Dims
- Attributes
 - Key/value pair
 - DataType and length

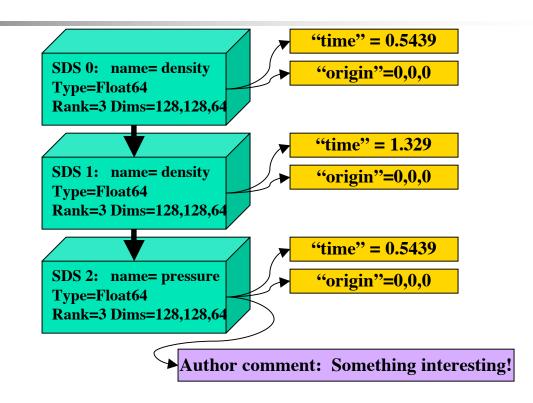






HDF4 SDS data schema

- Datasets
 - Name
 - Datatype
 - Rank, Dims
- Attributes
 - Key/value pair
 - DataType and length
- Annotations
 - Freeform text
 - String Termination

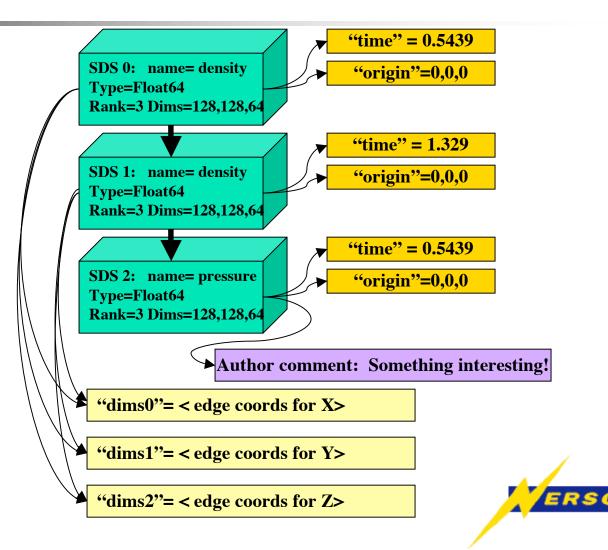






HDF4 SDS data schema

- Datasets
 - Name
 - Datatype
 - Rank, Dims
- Attributes
 - Key/value pair
 - DataType and length
- Annotations
 - Freeform text
 - String Termination
- Dimensions
 - Edge coordinates
 - Shared attribute





HDF5 Features

- Complete rewrite and API change
- Thread-safety support
- Parallel I/O support (via MPI-IO)
- Fundamental hierarchical grouping architecture
- No bundled data-schema centric APIs
- C and 100% Java implementations
- F90 and C++ API Bindings
- Pluggable compression methods
- Virtual File Driver Layer (RemoteHDF5)







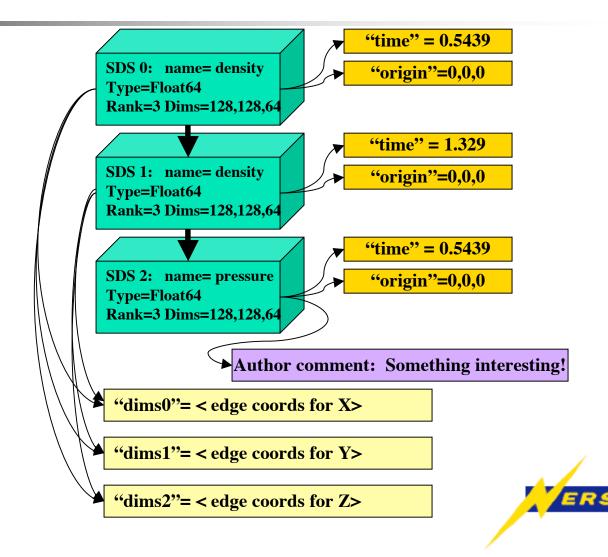
Why Use HDF5?

- Reasonably fast
 - faster than F77 binary unformatted I/O!
- Clean and Flexible Data Model
- Cross platform
 - Constant work maintaining ports to many architectures and OS revisions.
- Well documented
 - Members of the group dedicated to web docs
- Well maintained
 - Good track record for previous revs
- In general, less work for you in the long run!





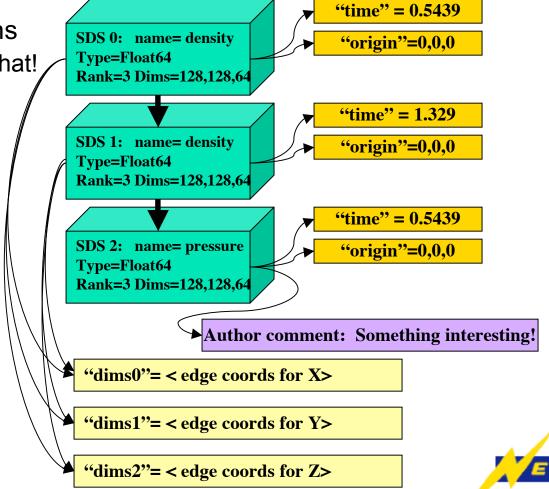








Use attributes for that!







Use attributes for that!

Eliminate Dims/Dimscales

Use attributes for that!

SDS 0: name= density
Type=Float64
Rank=3 Dims=128,128,64

"time" = 0.5439

"origin"=0,0,0

"time" = 1.329

SDS 1: name= density Type=Float64 Rank=3 Dims=128,128,64

"origin"=0,0,0

SDS 2: name= pressure
Type=Float64
Rank=3 Dims=128,128,64

"time" = 0.5439

"origin"=0,0,0

"dims0"= < edge coords for X>

"dims1"= < edge coords for Y>

"dims2"= < edge coords for Z>





Eliminate Annotations

Use attributes for that!

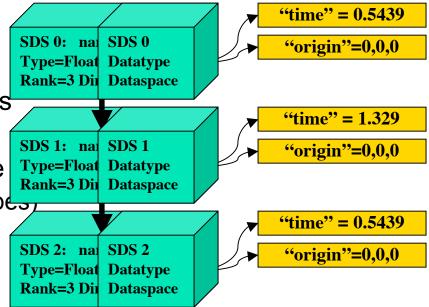
Eliminate Dims/Dimscales

Use attributes for that!

Rank, dims, type become

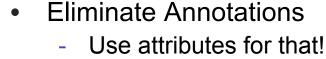
- Datatype (composite types)

- Dataspace





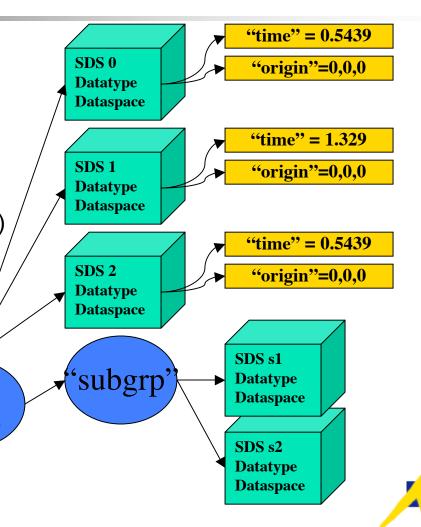




- Eliminate Dims/Dimscales
 - Use attributes for that!
- Rank, dims, type become
 - Datatype (composite types)

(root)

- Dataspace
- Groups
 - Datasets in groups
 - Groups of groups
 - Recursive nesting

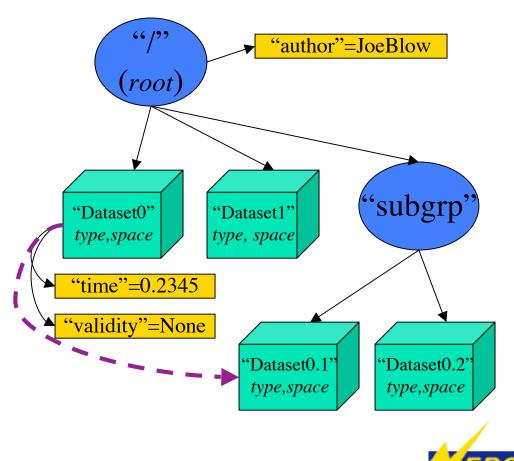






HDF5 Data Model

- Groups
 - Arranged in directory hierarchy
 - root group is always '/'
- Datasets
 - Dataspace
 - Datatype
- Attributes
 - Bind to Group & Dataset
- References – ·
 - Similar to softlinks
 - Can also be subsets of data



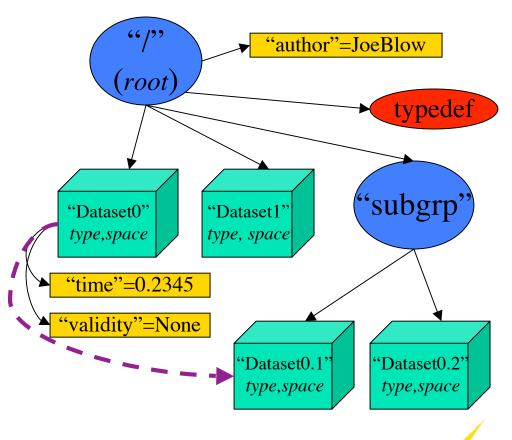




- Complex Type
 Definitions
 - Not commonly used feature of the data model.
 - Potential pitfall if you commit complex datatypes to your file
- Comments

rrrrrr

Yes, annotations actually do live on.

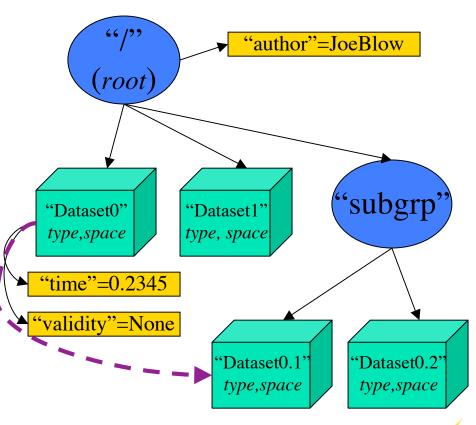


HDF5 Data Model (caveats)

- Flexible/Simple Data Model
 - You can do anything you want with it!
 - You typically define a higher level data model on top of HDF5 to describe domain-specific data relationships
 - Trivial to represent as XML!
- The perils of flexibility!

rrrrrr

- Must develop community agreement on these data models to share data effectively
- Multi-Community-standard data models required across for reusable visualization tools
- Preliminary work on Images and tables





Acquiring HDF5

- HDF5 home site
 - Information/Documentation http://hdf.ncsa.uiuc.edu/hdf5
 - Libraries (binary and source) ftp://ftp.ncsa.uiuc.edu/HDF/hdf5
- Module on NERSC and LBL systems
 - module load hdf5
 - module load hdf5_par (for parallel implementation)
- Typically build using
 - ./configure --prefix=<where you want it>
 - make
 - make install







Building With HDF5

- Build Apps using
 - #include <hdf5.h>
 - Compile options: -I\$H5HOME/include
- Link using
 - Normal Usage: -L\$H5HOME/lib -lhdf5
 - With Compression: -L\$H5HOME/lib -lhdf5 -lz
- F90/F77
 - inc "fhdf5.inc"
 - Use C linker with: -Iftn -Iftn90
 - or Use F90 linker with: -IC







The HDF5 API Organization

- API prefix denotes functional category
 - Datasets: H5D
 - Data Types: H5T
 - Data Spaces: H5S
 - Groups: H5G
 - Attributes: H5A
 - File: H5F
 - References: H5R
 - Others (compression, error management, property lists, etc...)







Comments about the API

- Every HDF5 object has a unique numerical Identifier
 - hid_t in C, INTEGER in F90
 - These IDs are explicitly allocated and deallocated (handles for objects)
 - Common pitfall: Failure to release a handle
- All objects in the file have a name
 - Objects in the same group *must* have unique names
 - API has many functions to convert between name and integer ID/handle (sometimes confusing as to when you should use ID/handle vs. when to use name string)







Open/Close A File (H5F)

- Similar to standard C file I/O
 - H5Fishdf5(): check filetype before opening
 - H5Fcreate(),H5Fopen(): create/open files
 - H5Fclose(): close file
 - *H5Fflush()*: explicitly synchronize file contents
 - H5Fmount()/H5Funmount(): create hierarchy

Typical use



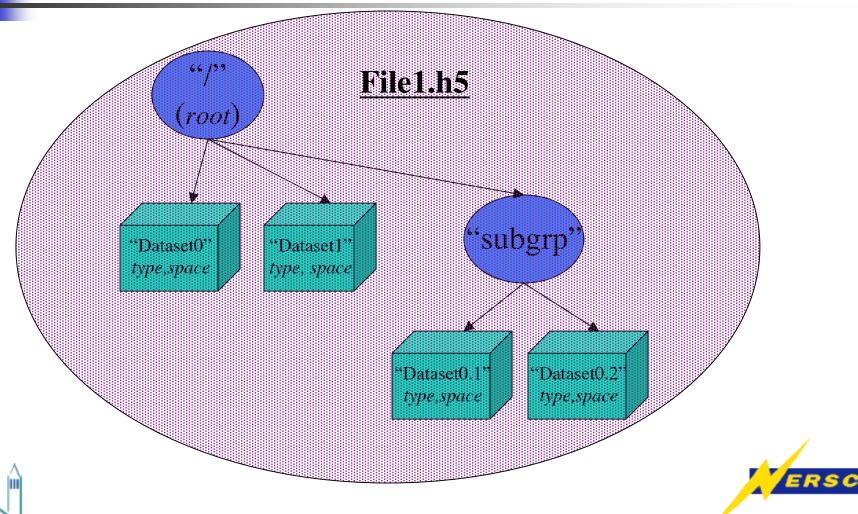


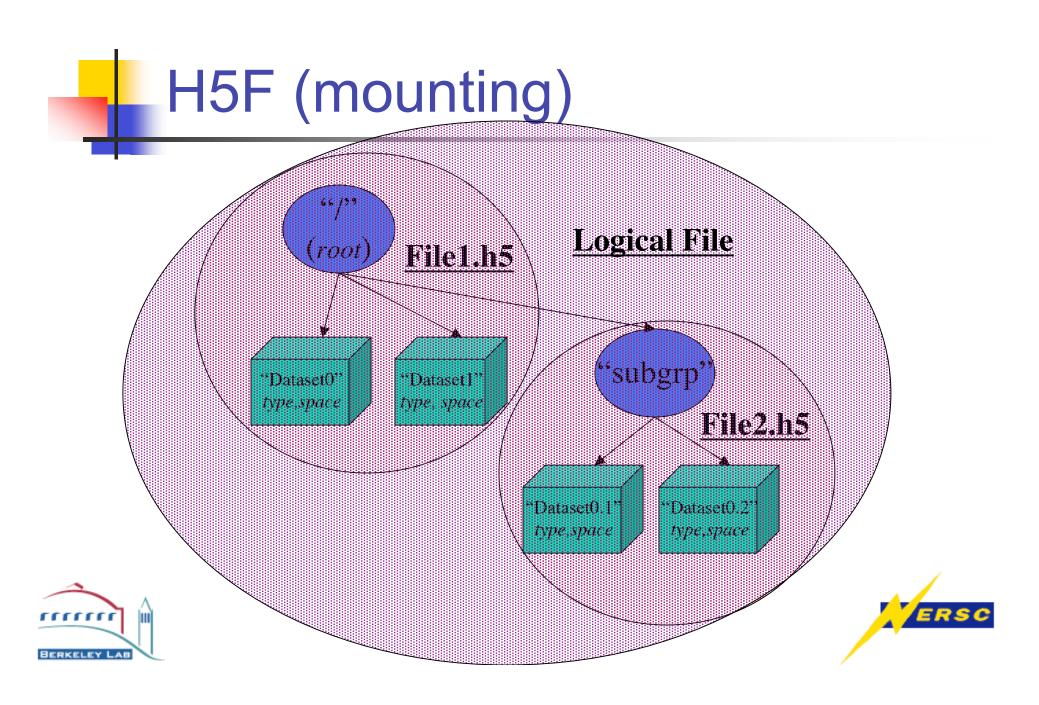


rrrrrr

BERKELEY LAB

H5F (opening)







- A serialization of a multidimensional logical structure composed of elements of same datatype
 - Each element can be complex/composite datatype
 - Logical layout (the topological meaning *you* assign to the dataset)
 - Physical layout (the actual serialization to disk or to memory.
 Can be a contiguous array or chunks that are treated as logically contiguous)
- Datasets require a "Dataspace" that defines relationship between logical and physical layout
- Dimensions are in row-major order!!!
 - ie. dims[0] is least rapidly changing and dim[n] is most rapidly changing

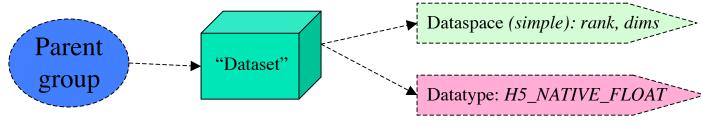


Write a Dataset (H5D)

- Required objects
 - ID for Parent Group
 - ID for DataType:

 H5T_NATIVE_FLOAT, H5T_IEEE_F32BE, H5T_IEEE_F32LE

 H5T_NATIVE_DOUBLE, H5T_NATIVE_INT, H5T_NATIVE_CHAR
 - ID for DataSpace: (logical shape of data array)
 - Simple case: rank and dimensions for logical Ndim array





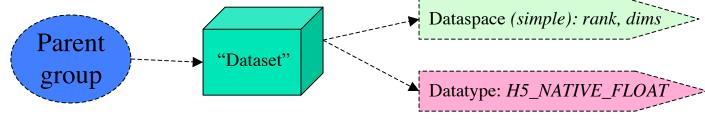




Write a Dataset (H5D)

Operations

- 1. H5Fcreate(): Open a new file for writing
- 2. H5Screate_simple(): Create data space (rank,dims)
- 3. H5Dcreate(): Create a dataset
- 4. H5Dwrite(): Commit the dataset to disk (mem-to-disk transfer)
- 5. H5Sclose(): Release data space handle
- 6. H5Fclose(): Close the file





H5D (example writing dataset)

```
hsize_t dims[3]={64,64,64};
float data[64*64*64]:
hid_t file,dataspace,datatype,dataset;
file = H5Fcreate("myfile.h5",H5F_ACC_TRUNC,
   H5P_DEFAULT,H5P_DEFAULT); /* open file (truncate if it exists) */
dataspace = H5Screate simple(3,dims,NULL); /* define 3D logical array */
datatype = H5T NATIVE FLOAT; /* use simple built-in native datatype */
   /* create simple dataset */
dataset = H5Dcreate(file,"Dataset1",datatype,dataspace,H5P_DEFAULT);
   /* and write it to disk */
H5Dwrite(dataset,datatype,H5S_ALL,H5S_ALL,H5P_DEFAULT,data);
H5Dclose(dataset);
                   /* release dataset handle */
H5Sclose(dataspace); /* release dataspace handle */
H5Fclose(file);
                    /* release file handle */
```

Read a Dataset (H5D)

- 1. H5Fopen(): Open an existing file for reading
- 2. H5Dopen(): Open a dataset in the file
- 3. H5Dget_type(): The data type stored in the dataset
 - H5Tget_class(): is it integer or float (H5T_INTEGER/FLOAT)
 - H5Tget_order(): little-endian or big-endian (H5T_LE/BE)
 - H5Tget_size(): nbytes in the data type
- H5Dget_space(): Get the data layout (the dataspace)
 - H5Dget_simple_extent_ndims(): Get number of dimensions
 - H5Dget_simple_extent_dims(): Get the dimensions
 - ... allocate memory for the dataset
- 5. H5Dread(): Read the dataset from disk
- 6. H5Tclose(): Release dataset handle
- 7. H5Sclose(): Release data space handle
- 8. H5Fclose(): Close the file





Reading a Dataset (H5D)

```
hsize t dims[3], ndims;
float *data:
hid t file, dataspace, file dataspace, mem dataspace, datatype, dataset;
file = H5Fopen("myfile.h5",H5F_ACC_RDONLY,H5P_DEFAULT); /* open file read_only */
datatype = H5Dget_type(dataset); /* get type of elements in dataset */
dataspace = H5Dget_space(dataset); /* get data layout on disk (dataspace) */
mem dataspace = file dataspace=dataspace; /* make them equivalent */
ndims = H5Sget simple extents ndims(dataspace); /* get rank */
H5Sget simple extent dims(dataspace, dims, NULL); /* get dimensions */
data = malloc(sizeof(float)*dims[0]*dims[1]*dims[2]); /* alloc memory for storage*/
H5Dread(dataset, H5T NATIVE FLOAT, mem dataspace, file dataspace, /* H5S ALL */
   H5P DEFAULT, data); /* read the data into memory buffer (H5S ALL) */
                    /* release dataset handle */
H5Dclose(dataset);
H5Tclose(datatype); /* release datatype handle */
H5Sclose(dataspace); /* release dataspace handle */
H5Fclose(file);
                           /* release file handle */
```



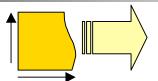
DataSpaces (H5S)

- Data Space Describes the mapping between logical and physical data layout
 - Physical is serialized array representation
 - Logical is the desired topological relationship between data elements
- Use a DataSpace to describe data layout both in memory and on disk
 - Each medium has its own DataSpace object!
 - Transfers between media involve physical-to-physical remapping but is defined as a logical-to-logical remapping operation



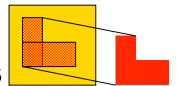
Data Storage Layout (H5S)

Elastic Arrays

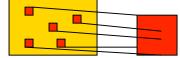


- Hyperslabs
 - Logically contiguous chunks of data

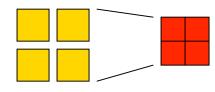
- Multidimensional Subvolumes
- Subsampling (striding, blocking)
- Union of Hyperslabs
 - Reading a non-rectangular sections



Gather/Scatter



- Chunking
 - Usually for efficient Parallel I/O







Dataset Selections (H5S)

- Array: H5Sselect_hyperslab(H5S_SELECT_SET)
 - Offset: start location of the hyperslab (default={0,0...})
 - Count: number of elements or blocks to select in each dimension (no default)
 - Stride: number of elements to separate each element or block to be selected (default={1,1...})
 - Block: size of block selected from dataspace (default={1,1...})
- Unions of Sets:
 - H5Sselect_hyperslab(H5S_SELECT_OR)
- Gather Scatter H5Sselect_elements(H5S_SELECT_SET)
 - Point lists

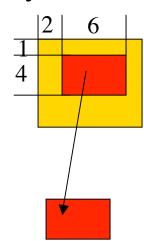


Dataspace Selections (H5S)

Transfer a subset of data from disk to fill a memory buffer

Disk Dataspace

H5Sselect_hyperslab(disk_space, H5S_SELECT_SET, offset[3]={1,2},NULL,count[2]={4,6},NULL)



Memory Dataspace

 $mem_space = H5S_ALL$

Or

 $mem_space = H5Dcreate(rank=2,dims[2]=\{4,6\});$

Transfer/Read operation

H5Dread(dataset,mem_datatype, mem_space, disk_space, H5P_DEFAULT, mem_buffer);





Dataspace Selections (H5S)

Transfer a subset of data from disk to subset in memory

Disk Dataspace

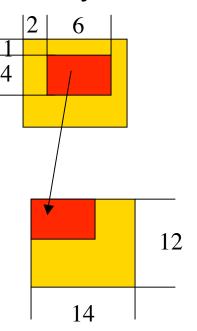
H5Sselect_hyperslab(disk_space, H5S_SELECT_SET, offset[3]={1,2},NULL,count[2]={4,6},NULL)

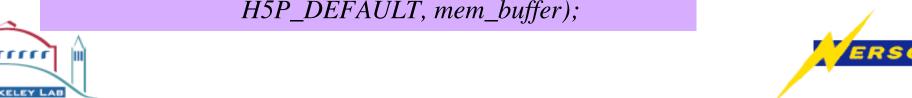
Memory Dataspace

mem_space =H5Dcreate_simple(rank=2,dims[2]={12,14}); H5Sselect_hyperslab(mem_space, H5S_SELECT_SET, offset[3]={0,0},NULL,count[2]={4,6},NULL)

Transfer/Read operation

H5Dread(dataset,mem_datatype, mem_space, disk_space, H5P_DEFAULT, mem_buffer);



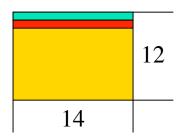


Dataspace Selections (H5S)

Row/Col Transpose

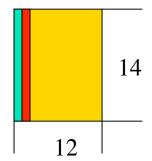
Disk Dataspace

disk_space = H5Dget_space(dataset)
rank=2 dims={12,14} canonically row-major order



Memory Dataspace

 $mem_space = H5Dcreate_simple(rank=2,dims[2]=\{14,12\});$ $H5Sselect_hyperslab(mem_space, H5S_SELECT_SET, offset[3]=\{0,0\},stride=\{14,1\},count[2]=\{1,12\},block=\{14,1\})$



Transfer/Read operation

H5Dread(dataset,mem_datatype, mem_space, disk_space, H5P_DEFAULT, mem_buffer);







DataTypes Native(H5T)

Native Types

 H5T_NATIVE_INT, H5T_NATIVE_FLOAT, H5T_NATIVE_DOUBLE, H5T_NATIVE_CHAR, ... H5T_NATIVE_<foo>

Arch Dependent Types

- Class: H5T_FLOAT, H5T_INTEGER
- Byte Order: H5T_LE, H5T_BE
- Size: 1,2,4,8,16 byte datatypes
- Composite:
 - Integer: H5T_STD_I32LE, H5T_STD_I64BE
 - Float: H5T_IEEE_F32BE, H5T_IEEE_F64LE
 - String: H5T_C_S1, H5T_FORTRAN_S1
- Arch: H5T_INTEL_I32, H5T_INTEL_F32







DataTypes (H5T)

- Type Translation for writing
 - Define Type in Memory
 - Define Type in File (native or for target machine)
 - Translates type automatically on retrieval
- Type Translation for reading
 - Query Type in file (class, size)
 - Define Type for memory buffer
 - Translates automatically on retrieval





4

DataTypes (H5T)

Writing

```
dataset=H5Dcreate(file,name,mem_datatype,dataspace,...);
H5Dwrite(dataset,file_datatype, memspace, filespace,...);
```

Reading

dataset=H5Dopen(file,name,mem_datatype,dataspace,...);







Complex DataTypes (H5T)

- Array Datatypes:
 - Vectors
 - Tensors
- Compound Objects
 - C Structures
- Variable Length Datatypes
 - Strings
 - Elastic/Ragged arrays
 - Fractal arrays
 - Polygon lists
 - Object tracking







Caveats (H5T)

- Elements of datasets that are compound/complex must be accessed in their entirety
 - It may be notationally convenient to store a tensor in a file as a dataset. However, you can select subsets of the dataset, but not sub-elements of each tensor!
 - Even if they could offer this capability, there are fundamental reasons why you would not want to use it!





Array Data Types (H5T)

- Create the Array Type
 - atype=H5Tarray_create(basetype,rank,dims,NULL)

```
hsize_t vlen=3;
flowfield = H5Tarray_create(H5T_NATIVE_DOUBLE,1,&vlen,NULL);
```

- Query the Array Type
 - H5Tget_array_ndims(atype)
 - H5Tget_array_dims(atype,dims[],NULL)







Attributes (H5A)

- Attributes are simple arrays bound to objects (datasets, groups, files) as key/value pairs
 - Name
 - Datatype
 - DataSpace: Usually a scalar H5Screate(H5S_SCALAR)
 - Data
- Retrieval
 - By name: H5Aopen_name()
 - By index: H5Aopen_idx()
 - By iterator: H5Aiterate()





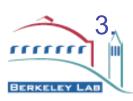


Writing/Reading Attribs (H5A)

- Write an Attribute
 - Create Scalar Dataspace: H5Screate(H5S_SCALAR)
 - 2. Create Attribute: H5Acreate()
 - 3. Write Attribute (bind to obj): H5Awrite()
 - 4. Release dataspace, attribute H5Sclose(), H5Aclose()

- Read an Attribute
 - Open Attrib: H5Aopen_name()
 - 2. Read Attribute: H5Aread()
 - Release H5Aclose()

Attrib=H5Aopen_name(object,"attribname"); H5Aread(attrib,mem_datatype,data); H5Aclose(attrib);







Caveats about Attribs (H5A)

- Do not store large or complex objects in attributes
- Do not do collective operations (parallel I/O) on attributes
- Make your life simple and make datatypes implicitly related to the attribute name
 - ie. "iorigin" vs. "origin"
 - Avoid type class discovery when unnecessary







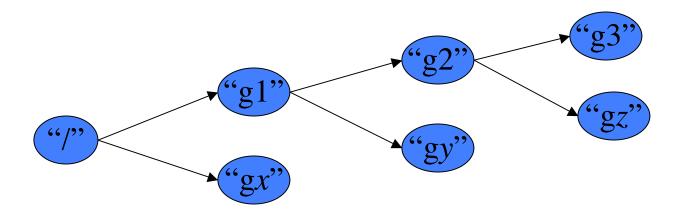
- Similar to a "directory" on a filesystem
 - Has a name
 - Parent can only be another group (except root)
- There is always a root group in a file called "/"
- Operations on Groups
 - H5Gcreate: Create a new group (ie. Unix 'mkdir')
 - H5Gopen/H5Gclose: Get/release a handle to a group
 - H5Gmove: Move a directory (ie. Unix 'mv' command)
 - H5Glink/H5Gunlink: Hardlinks or softlinks (Unix 'In') Unlink is like 'rm')
 - H5Giterate: Walk directory structures recursively







- Navigation (walking the tree)
 - Navigate groups using directory-like notation



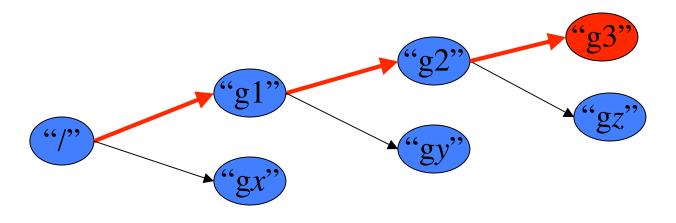


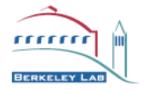




- Navigation (walking the tree)
 - Navigate groups using directory-like notation

(select group "/g1/g2/g3")







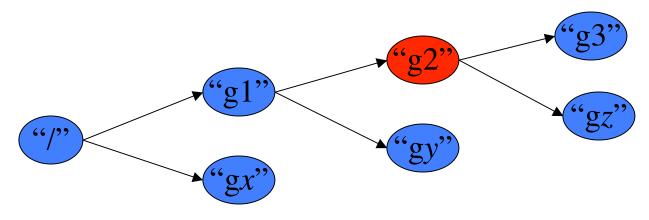


- Navigation (walking the tree)
 - Navigate groups using directory-like notation

Simple Example: Opening a particular Group "g2" hid_t gid=H5Gopen(file_ID,"/g1/g2");

Is equivalent to

hid_t gid=H5Gopen(gid_g1,"g2");



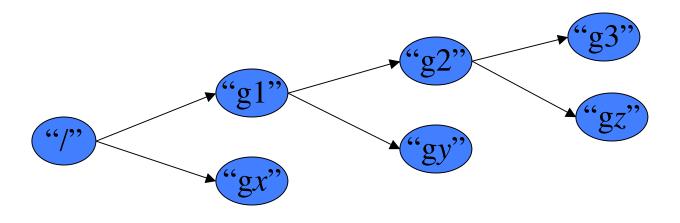






Navigation

- Navigate groups using directory-like notation
- Navigate using iterators (a recursive walk through the tree)







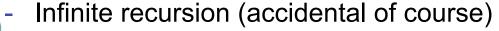
H5G (Group Iterators)

Why use iterators

- Allows discovery of directory structure without knowledge of the group names
- Simplifies implementation recursive tree walks
- Iterator Method (user-defined callbacks)
 - Iterator applies user callback to each item in group
 - Callback returns 1: Iterator stops immediately and returns current item ID to caller
 - Callback returns 0: Iterator continues to next item

Pitfalls

 No guaranteed ordering of objects in group (not by insertion order, name, or ID)



H5G (using Group Iterators)

Print names of all items in a group (pseudocode)

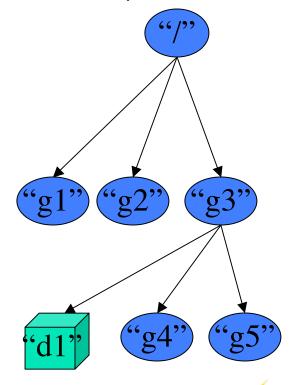
```
herr_t PrintName(objectID,objectName){
    print(objectName)
    return 0
}
```

H5Giterate(fileID, "/", PrintName, NULL(userData))

Outputs: "g1 g2 g3"

H5Giterate(fileID, "/g3", PrintName, NULL(userData))

Outputs: "d1 g4 g5"





H5G (using Group Iterators)

Depth-first walk(pseudocode)

```
herr_t DepthWalk(objectID,objectName){
    print(objectName)
    /* note: uses objectID instead of file and NULL for the name */
    H5Giterate(objectID,NULL,DepthWalk,NULL(userData));
    return 0
}

H5Giterate(fileID,"/",DepthWalk,NULL(userData))
Outputs: "g1 g2 g3 d1 g4 g5"

"g1" "g2" "g3"
```



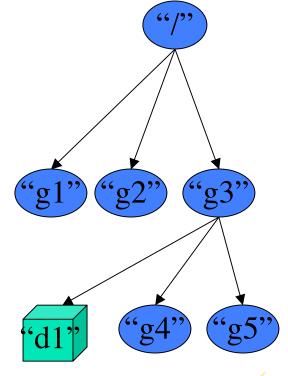


4

H5G (using Group Iterators)

Count items (pseudocode)

```
herr_t CountCallback(objectID,objectName,count){
    *userdata++;
    H5Giterate(objectID,NULL,UserCallback,count);
    return 0
}
Int count;
H5Giterate(fileID,"/",CountCallback,&count)
Print(*count)
Outputs: "6"
```









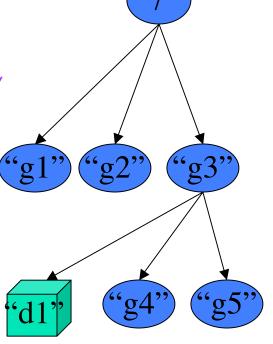
H5G (using Group Iterators)

Select item with property (pseudocode)

```
herr_t SelectCallback(objectID,objectName,property){
    if(getsomeproperty(objectID)==*property)
        return 1 /* terminate early: we found a match! */
    H5Giterate(objectID,NULL,UserCallback,property)
    return 0
}
```

Returns item that matches property

match=H5Giterate(fileID, "/", SelectCallback, & property)







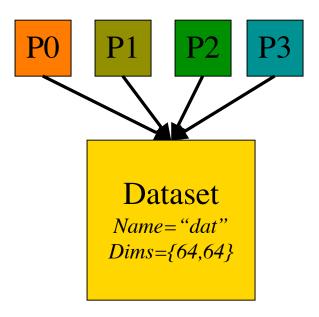


- Restricted to collective operations on datasets
 - Selecting one dataset at a time to operate on collectively or independently
 - Uses MPI/IO underneath (and hence is *not* for OpenMP threading. Use ThreadSafe HDF5 for that!)
 - Declare communicator only at file open time
 - Attributes are only actually processed by process with rank=0
- Writing to datafiles
 - Declare overall data shape (dataspace) collectively
 - Each processor then uses H5Sselect_hypeslab() to select each processor's subset of the overall dataset (the domain decomposition)
 - Write collectively or independently to those (preferably) nonoverlapping offsets in the file.

pHDF5 (example 1)

- •File open requires explicit selection of Parallel I/O layer.
- •All PE's collectively open file and declare the overall size of the dataset.

All MPI Procs!

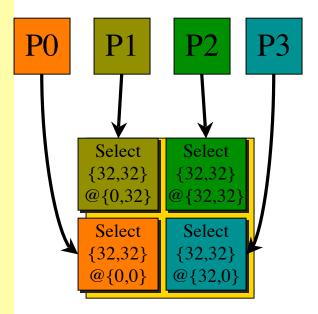




pHDF5 (example 1 cont...)

•Each proc selects a hyperslab of the dataset that represents its portion of the domain-decomposed dataset and read/write collectively or independently.

All MPI Procs!







SP2 Caveats

- Must use thread-safe compilers, or it won't even recognize the mpi/io routines.
 - mpcc_r, mpxlf90_r, mpCC_r
- Must link to -lz which isn't in default path
 - Use path to hdf4 libs to get libz
- Location of libs and includes
 - -I/usr/common/usg/hdf5/1.4.4/parallel/include
 - -L/usr/common/usg/hdf5/1.4.4/parallel/lib -lhdf5
 - -L/usr/common/usg/hdf/4.1r5 -lz -lm







- If data reorganization proves costly, then put off it off until the data analysis stage
 - The fastest writes are when layout on disk == layout in memory
 - If MPP hours are valuable, then don't waste them on massive insitu data reorganization
 - Data reorg is usually more efficient for parallel reads than for parallel writes (especially for SMPs)
 - Take maximum advantage of "chunking"
- Parallel I/O performance issues usually are direct manifestations of MPI/IO performance
- Don't store ghost zones!
- Don't store large arrays in attributes





Serial I/O Benchmarks

System	HDF4.1r5 (netCDF)	HDF5 v1.4.4	FlexIO (Custom)	F77 Unf
SGI Origin 3400 (escher.nersc.gov)	111M/s	189M/s	180M/s	140M/s
IBM SP2 (seaborg.nersc.gov)	65M/s	127M/s	110M/s	110M/s
Linux IA32 (platinum.ncsa.uiuc.edu)	34M/s	40M/s	62M/s	47M/s
Linux IA64 Teragrid node (titan.ncsa.uiuc.edu)	26M/s	83M/s	77M/s	112M/s
NEC/Cray SX-6 (rime.cray.com)				

- •Write 5-40 datasets of 128³ DP float data
- •Single CPU (multiple CPU's can improve perf. until interface saturates
- •Average of 5 trials

GPFS MPI-I/O Experiences

nTasks	I/O Rate	I/O Rate	
	16 Tasks/node	8 tasks per node	
8	-	131 Mbytes/sec	
16	7 Mbytes/sec	139 Mbytes/sec	
32	11 Mbytes/sec	217 Mbytes/sec	
64	11 Mbytes/sec	318 Mbytes/sec	
128	25 Mbytes/sec	471 Mbytes/sec	

- Block domain decomp of 512³ 3D 8-byte/element array in memory written to disk as single undecomposed 512³ logical array.
- Average throughput for 5 minutes of writes x 3 trials







Whats Next for HDF5?

- Standard Data Models with Associated APIs
 - Ad-Hoc (one model for each community)
 - Fiber Bundle (unified data model)
 - Whats in-between?
- Web Integration
 - 100% pure java implementation
 - XML/WSDL/OGSA descriptions
- Grid Integration:
 - http://www.zib.de/Visual/projects/TIKSL/HDF5-DataGrid.html
 - RemoteHDF5
 - StreamingHDF5







More Information

- HDF Website
 - Main: http://hdf.ncsa.uiuc.edu
 - HDF5: http://hdf.ncsa.uiuc.edu/HDF5
 - pHDF5: http://hdf.ncsa.uiuc.edu/Parallel_HDF
- HDF at NASA (HDF-EOS)
 - http://hdfeos.gsfc.nasa.gov/
- HDF5 at DOE/ASCI (Scalable I/O Project)
 - http://www.llnl.gov/icc/lc/siop/



