PyData SV 2013 - Tutorials

HDF5 is for Lovers



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Which you will learn today!



A Note on the Format

Intermixed, there will be:

- Slides
- Interactive Hacking
- Exercises



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Intermixed, there will be:

- Slides
- Interactive Hacking
- Exercises

Feel free to:

- Ask questions at anytime
- Explore at your own pace.



By a show of hands, how many people have used:

• HDF5 before?



- HDF5 before?
- •PyTables?



- HDF5 before?
- •PyTables?
- •h5py?



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- •h5py?
- the HDF5 C API?



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- •h5py?
- the HDF5 C API?
- •SQL?



- HDF5 before?
- •PyTables?
- •h5py?
- the HDF5 C API?
- •SQL?
- Other binary data formats?



Setup

Please clone the repo:

```
git clone git://github.com/scopatz/hdf5-is-for-lovers.git
```

Or download a tarball from:

https://github.com/scopatz/hdf5-is-for-lovers



Warm up exercise

In IPython:

```
import numpy as np
import tables as tb

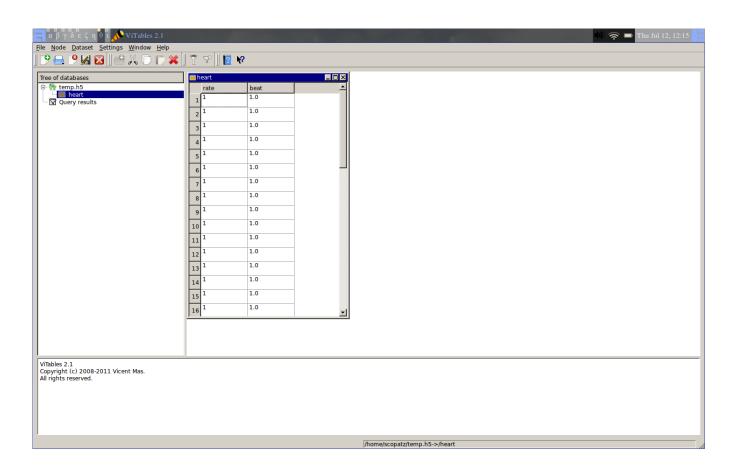
f = tb.openFile('temp.h5', 'a')
heart = np.ones(42, dtype=[('rate', int), ('beat', float)])
f.createTable('/', 'heart', heart)
f.close()
```

Or run python exer/warmup.py



Warm up exercise

You should see in ViTables:





For persisting structured numerical data, binary formats are superior to plaintext.



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For one thing, they are often smaller:



For another, binary formats are often faster for I/O because atoi() and atof() are expensive.



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However, you often want some thing more than a binary chunk of data in a file.



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Note

This is the mechanism behind numpy.save() and numpy.savez().



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In effect, HDF5 is a file system within a file.



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Unlike SQL, where every dataset lives in a flat namespace, HDF allows datasets to live in a nested tree structure.

In effect, HDF5 is a file system within a file.

(More on this later.)



Basic dataset classes include:

Array



- Array
- CArray (chunked array)



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- Table (structured array w/ named fields)



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- CArray (chunked array)
- EArray (extendable array)
- VLArray (variable length array)
- Table (structured array w/ named fields)

All of these must be composed of atomic types.



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- •complex: Complex number. 64 and 128 (default) bits.
- string: Raw string types. 8-bit positive multiples.



Other elements of the hierarchy may include:

• Groups (dirs)



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- Links



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- File Nodes



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PyTables docs may be found at http://pytables.github.com/



Opening Files

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```



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```

- 'r': Read-only; no data can be modified.
- 'w': Write; a new file is created (an existing file with the same name would be deleted).
- 'a': Append; an existing file is opened for reading and writing, and if the file does not exist it is created.
- 'r+': It is similar to 'a', but the file must already exist.

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Creating new nodes must be done on the file handle:

```
f.createGroup('/', 'a_group', "My Group")
f.root.a_group
```



Creating Datasets

The two most common datasets are Tables & Arrays.



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Appropriate create methods live on the file handle:

```
# integer array
f.createArray('/a_group', 'arthur_count', [1, 2, 5, 3])
```



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Appropriate create methods live on the file handle:

```
# integer array
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```

```
# tables, need descriptions
dt = np.dtype([('id', int), ('name', 'S10')])
knights = np.array([(42, 'Lancelot'), (12, 'Bedivere')], dtype=dt)
f.createTable('/', 'knights', dt)
f.root.knights.append(knights)
```



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```
>>> print f.root.a_group.arthur_count[:]
[1, 2, 5, 3]
>>> type(f.root.a_group.arthur_count[:])
list
>>> type(f.root.a_group.arthur_count)
tables.array.Array
```

So if they come from NumPy arrays, they may be accessed in a numpy-like fashion (slicing, fancy indexing, masking).



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```
>>> f.root.knights[1]
(12, 'Bedivere')
>>> f.root.knights[:1]
array([(42, 'Lancelot')], dtype=[('id', '<i8'), ('name', 'S10')])
>>> mask = (f.root.knights.cols.id[:] < 28)
>>> f.root.knights[mask]
array([(12, 'Bedivere')], dtype=[('id', '<i8'), ('name', 'S10')])
>>> f.root.knights[([1, 0],)]
array([(12, 'Bedivere'), (42, 'Lancelot')], dtype=[('id', '<i8'), ('name', 'S10')])</pre>
```



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```

Data accessed in this way is memory mapped.



Exercise

exer/peaks_of_kilimanjaro.py





Exercise

sol/peaks_of_kilimanjaro.py





Suppose there is a big table of like-things:



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It is tempting to throw everyone into a big people table.



However, a search over a class of people can be eliminated by splitting these tables up:



The profession column is now redundant:



Information can be embedded implicitly in the hierarchy as well:

```
root
    - England
         | - knight
          - others
      France
           - knight
           - others
```

Why bother pivoting the data like this at all?



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Ultimately, it is all about *speed*, especially for big tables.



Access Time Analogy

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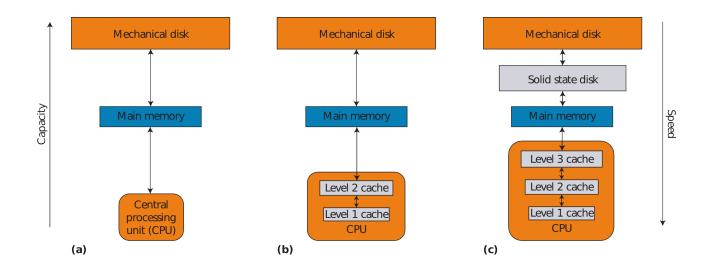
Accessing main memory is going to the break room, get a candy bar, and chatting with your co-worker (4 min).

Accessing a (mechanical) HDD is leaving your office, leaving your building, wandering the planet for a year and four months to return to your desk with the information finally made available.

Thanks K. Smith & http://duartes.org/gustavo/blog/post/what-your-computer-does-while-you-wait

Starving CPU Problem

Waiting around for access times prior to computation is known as the *Starving CPU Problem*.



Francesc Alted. 2010. Why Modern CPUs Are Starving and What Can Be Done about It. IEEE Des. Test 12, 2 (March 2010), 68-71. DOI=10.1109/MCSE.2010.51

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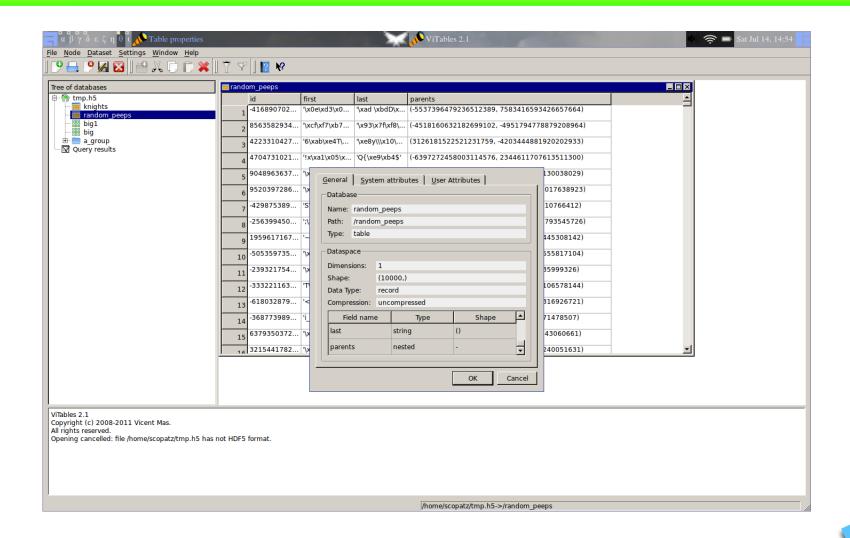
In fact, the struct / dtype / description concept is only a convenient way to assign meaning to bytes:

	,	ids		first										last										
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Data types may be nested (though they are stored in flattened way).





Python already has the ability to dynamically declare the size of descriptions.



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This is accomplished in compiled languages through normal memory allocation and careful byte counting:

```
typedef struct mat {
  double mass;
  int atoms_per_mol;
  double comp [];
} mat;
```



```
typedef struct mat {
 double mass:
 int atoms_per_mol;
 double comp [];
} mat;
size t mat size = sizeof(mat) + sizeof(double)*comp size;
hid_t desc = H5Tcreate(H5T_COMPOUND, mat_size);
hid t comptype = H5Tarray_create2(H5T_NATIVE_DOUBLE, 1, nuc_dims);
// make the data table type
H5Tinsert(desc, "mass", H0FFSET(mat, mass), H5T_NATIVE_DOUBLE);
H5Tinsert(desc, "atoms per mol", HOFFSET(mat, atoms per mol), H5T NATIVE DOUBLE);
H5Tinsert(desc, "comp", HOFFSET(mat, comp), comp type);
// make the data array for a single row, have to over-allocate
mat * mat data = new mat[mat size];
// ...fill in data array...
// Write the row
H5Dwrite(data_set, desc, mem_space, data_hyperslab, H5P_DEFAULT, mat_data);
```

Exercise

exer/boatload.py





Exercise

sol/boatload.py





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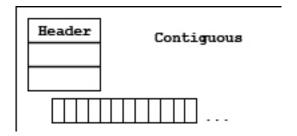
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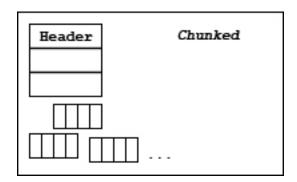
By chunking, sparse data may be stored efficiently and datasets may extend infinitely in all dimensions.

Note: Currently, PyTables only allows one extendable dim.





Contiguous Dataset





Chunked Dataset



All I/O happens by chunk. This is important for:

• edge chunks may extend beyond the dataset



- edge chunks may extend beyond the dataset
- default fill values are set in unallocated space



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- reading and writing in parallel



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- reading and writing in parallel
- small chunks are good for accessing some of data
- large chunks are good for accessing lots of data



Any chunked dataset allows you to set the chunksize.

```
f.createTable('/', 'omnomnom', data, chunkshape=(42,42))
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However, it could not have a 12x12 chunksize, since the ranks must be less than or equal to that of the array.



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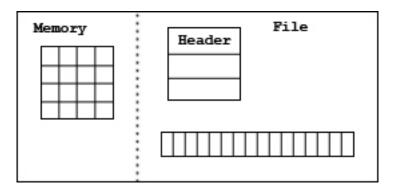
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f.createTable('/', 'omnomnom', data, chunkshape=(42,42))
```

For example, a 4x4 chunked array could have a 3x3 chunksize.

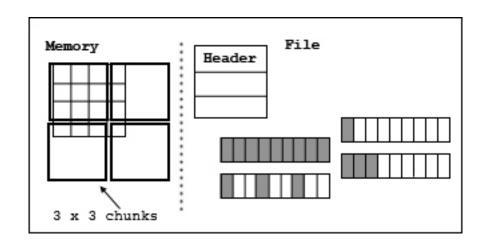
However, it could not have a 12x12 chunksize, since the ranks must be less than or equal to that of the array.

Manipulating the chunksize is a great way to fine-tune an application.





Contiguous 4x4 Dataset

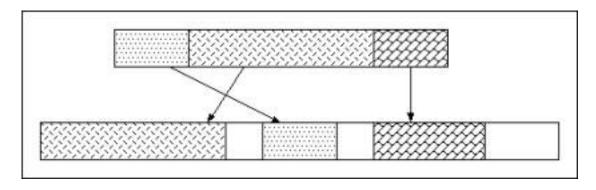




Chunked 4x4 Dataset



Note that the addresses of chunks in dataspace (memory) has no bearing on their arrangement in the actual file.



Dataspace (top) vs File (bottom) Chunk Locations



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Definitions:

- Operations which require all data to be in memory are *in-core* and may be memory bound (NumPy).
- Operations where the dataset is external to memory are *out-of-core* (or *in-kernel*) and may be CPU bound.



Say, a and b are arrays sitting in memory:

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a = np.array(...)
b = np.array(...)
c = 42 * a + 28 * b + 6
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For N operations, N-1 temporaries are made.

Wastes memory and is slow. Pulling from disk is slower.



A less memory intensive implementation would be an element-wise evaluation:

```
c = np.empty(...)
for i in range(len(c)):
    c[i] = 42 * a[i] + 28 * b[i] + 6
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But if a and b were HDF5 arrays on disk, individual element access time would kill you.

Even with in memory NumPy arrays, there are problems with gratuitous Python type checking.



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```
for i in range(0, len(a), 256):
    r0, r1 = a[i:i+256], b[i:i+256]
    multiply(r0, 42, r2)
    multiply(r1, 28, r3)
    add(r2, r3, r2); add(r2, 6, r2)
    c[i:i+256] = r2
```

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PyTables implements a tb.Expr class which backends to the numexpr VM but has additional optimizations for disk reading and writing.



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This problem lends itself nicely to parallelism.

Numexpr has low-level multithreading, avoiding the GIL.

PyTables implements a tb.Expr class which backends to the numexpr VM but has additional optimizations for disk reading and writing.

The full array need never be in memory.



Fully out-of-core expression example:

```
shape = (10, 10000)
f = tb.openFile("/tmp/expression.h5", "w")
a = f.createCArray(f.root, 'a', tb.Float32Atom(dflt=1.), shape)
b = f.createCArray(f.root, 'b', tb.Float32Atom(dflt=2.), shape)
c = f.createCArray(f.root, 'c', tb.Float32Atom(dflt=3.), shape)
out = f.createCArray(f.root, 'out', tb.Float32Atom(dflt=3.), shape)
expr = tb.Expr("a*b+c")
expr.setOutput(out)
d = expr.eval()
print "returned-->", repr(d)
f.close()
```



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Because querying is so common PyTables defines special methods on Tables.

```
tb.Table.where(cond)
tb.Table.getWhereList(cond)
tb.Table.readWhere(cond)
tb.Table.whereAppend(dest, cond)
```



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They are executed in the context of table itself combined with locals() and globals().



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The where () method itself returns an iterator over all matched (hit) rows:

```
for row in table.where('(col1 < 42) & (col2 == col3)'):
    # do something with row</pre>
```



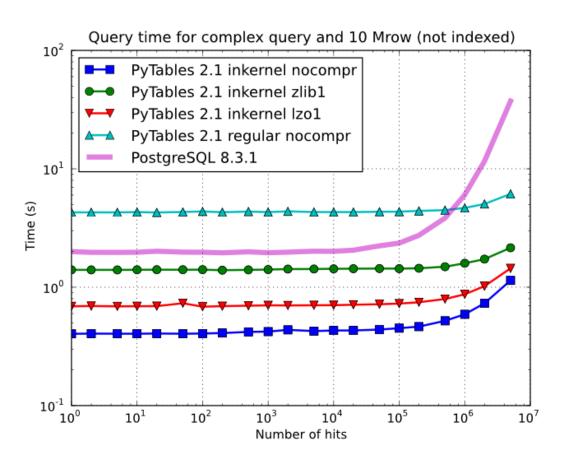
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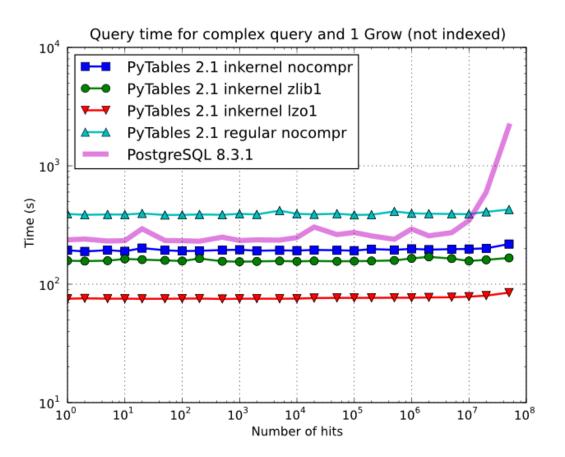
And this is the equivalent out-of-core search:





Complex query with 10 million rows. Data fits in memory.





Complex query with 1 billion rows. Too big for memory.



Exercise

exer/crono.py





Exercise

sol/crono.py





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Compression is when the dataset is piped through a zipping algorithm on write and the inverse unzipping algorithm on read.

Each chunk is compressed independently, so chunks end up with a varying number bytes.

Has some storage overhead, but may drastically reduce file sizes for very regular data.

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However, because there is *less total information* to transfer, the time spent unpacking the array can be far less than moving the array around wholesale.

This is kind of like power steering, you can either tell wheels how to turn manually or you can tell the car how you want the wheels turned.

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PyTables supports:

• zlib (default), • lzo, • bzip2, and • blosc.





```
# complevel goes from [0,9]
filters = tb.Filters(complevel=5, complib='blosc', ...)
```



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filters = tb.Filters(complevel=5, complib='blosc', ...)
# filters may be set on the whole file,
f = tb.openFile('/path/to/file', 'a', filters=filters)
f.filters = filters
```



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# filters may be set on the whole file,
f = tb.openFile('/path/to/file', 'a', filters=filters)
f.filters = filters

# filters may also be set on most other nodes
f.createTable('/', 'table', desc, filters=filters)
f.root.group._v_filters = filters
```



Compression is enabled in PyTables through *filters*.

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Filters only act on chunked datasets.



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- Use blosc all other times. It is optimized for HDF5.

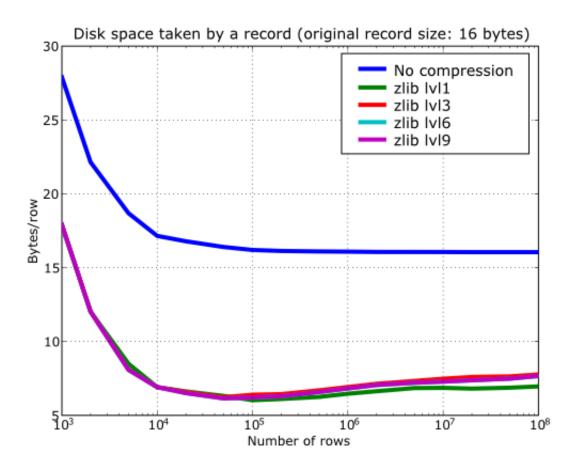


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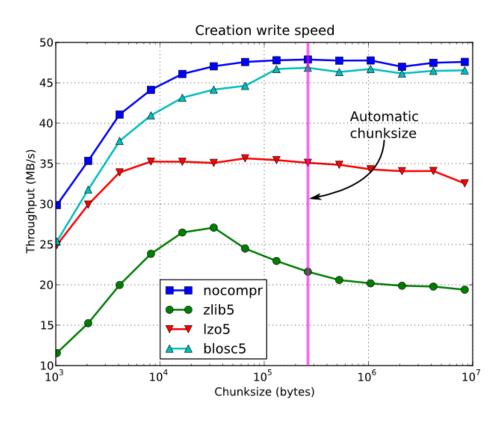
But why? (I don't have time to go into the details of blosc. However here are some justifications...)





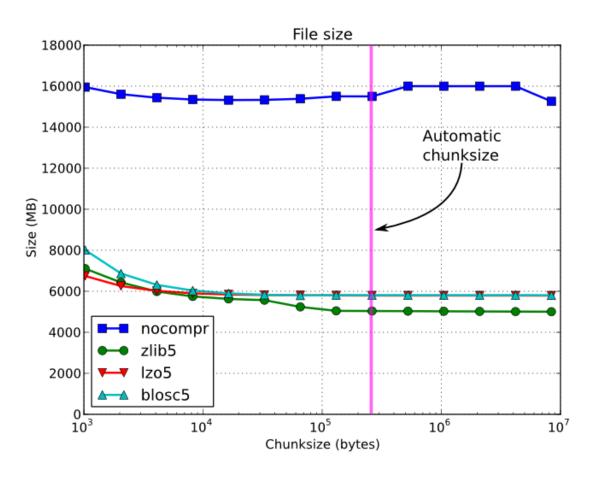
Comparison of different compression levels of zlib.





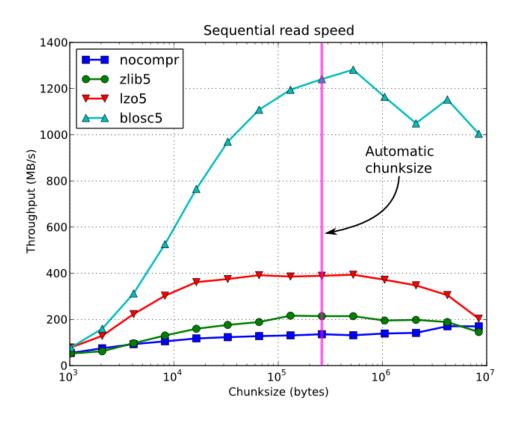
Creation time per element for a 15 GB EArray and different chunksizes.





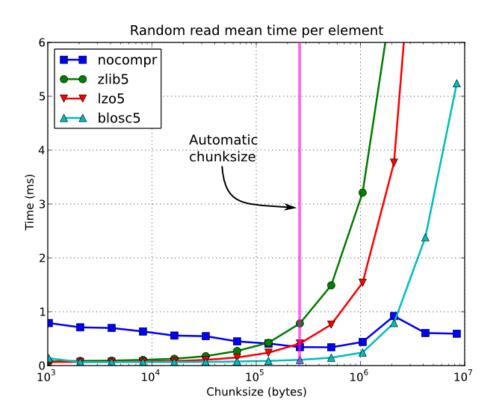
File sizes for a 15 GB EArray and different chunksizes.





Sequential access time per element for a 15 GB EArray and different chunksizes.





Random access time per element for a 15 GB EArray and different chunksizes.



Exercise

exer/spam_filter.py





Exercise

sol/spam_filter.py





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It is data structures like sets and dictionaries which do not quite map.

However, as long as all elements may be cast into the same atomic type, these structures can be stored in HDF5 with relative ease.



Sets

Example of serializing and deserializing sets:

```
>>> s = \{1.0, 42, 77.7, 6E+01, True\}
>>> f.createArray('/', 's', [float(x) for x in s])
/s (Array(4,)) ''
  atom := Float64Atom(shape=(), dflt=0.0)
  maindim := 0
  flavor := 'python'
  byteorder := 'little'
  chunkshape := None
>>> set(f.root.s)
set([1.0, 42.0, 77.7, 60.0])
```

Exercise

exer/dict_table.py





Exercise

sol/dict_table.py





What Was Missed

- Walking Nodes
- File Nodes
- Indexing
- Migrating to / from SQL
- HDF5 in other database formats
- Other Databases in HDF5
- HDF5 as a File System



Many thanks to everyone who made this possible!



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• The HDF Group



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- The HDF Group
- The PyTables Governance Team:
 - Josh Moore, Antonio Valentino, Josh Ayers



(Cont.)

• The NumPy Developers



(Cont.)

- The NumPy Developers
- •h5py, the symbiotic project



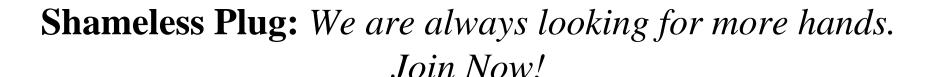
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Questions

