

# Scientific data formats

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**informatik**  
**die zukunft**

# Structure

- 1** Introduction
- 2** Requirements
- 3** Mathematical objects
- 4** Examples of scientific data formats
- 5** HDF5
- 6** Summary  
References

MP3 similar to scientific data formats?

MP3 similar to scientific data formats?

- abstraction of scientific object (MP3: wave to bits)
  - fit to your needs (MP3: reduce bandwidth of frequencies)

# Requirements

- file size
  - what is important to store?
  - meta data
  - precision
  - compression
  - fault tolerance, redundancy
- archivability, backwards compatibility
- portability
- interchangeability
- searchability
- human readable (text based)

## 1 Introduction

## 2 Requirements

## 3 Mathematical objects

- Graphs
- sparse matrices

## 4 Examples of scientific data formats

## 5 HDF5

## 6 Summary

## References

# Graphs

- traffic system (HVV map)
- state transitions (e.g. electrons)
- FIGURE 1 with mathematical Graph

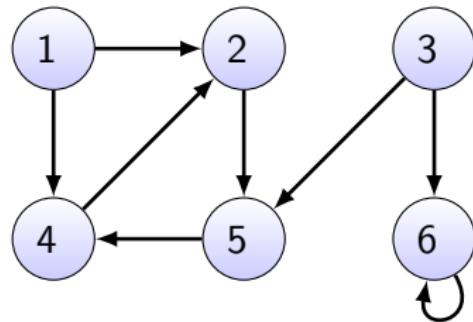
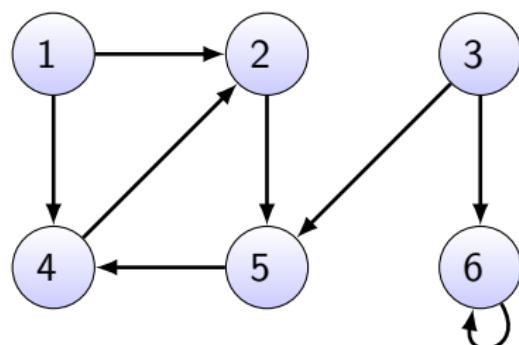


Figure: directed graph

# Adjacency lists

- each vertex has list of outgoing neighbours
- sparse graphs:  $|E| \ll |V|^2$
- usually neighbour list unsorted
- size  $\Theta(V + E)$
- reference: (Cormen u. a., 2007, 531)



1	{2, 4}
2	{5}
3	{6, 5}
4	{2}
5	{4}
6	{6}

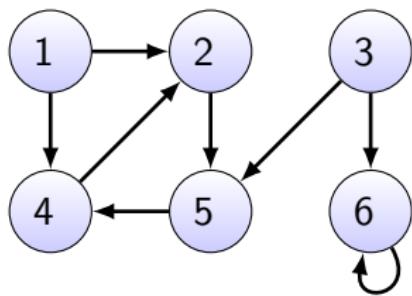
Figure: directed graph

Figure: adjacency list

## Graphs

## Adjacency matrix

- $|E| \sim |V|^2$
  - faster check if two vertices linked
  - $A^{|V| \times |V|}$  with  $a_{i,j} = \begin{cases} 1 & \text{if } (i,j) \in E \\ 0 & \text{else} \end{cases}$
  - size (const):  $\Theta(V^2)$
  - reference: (Cormen u. a., 2007, 531)



## Figure: directed graph

$$\begin{array}{cccccc} & 1 & 2 & 3 & 4 & 5 & 6 \\ \begin{matrix} 1 \\ 2 \\ 3 \\ 4 \\ 5 \\ 6 \end{matrix} & \left( \begin{array}{cccccc} 0 & 1 & 0 & 1 & 0 & 0 \\ 0 & 0 & 0 & 0 & 1 & 0 \\ 0 & 0 & 0 & 0 & 0 & 1 \\ 0 & 1 & 0 & 0 & 0 & 0 \\ 0 & 0 & 0 & 1 & 0 & 0 \\ 0 & 0 & 0 & 0 & 0 & 1 \end{array} \right) \end{array}$$

## Figure: adjacency matrix

# Sparse matrices

- often result of discretization of partial differential equation
- efficient algorithms in calculating
- save only non-null elements with their position
- special case: band matrices

$$\begin{pmatrix} a_{11} & a_{12} & 0 & \cdots & \cdots & \cdots & \cdots & 0 \\ a_{21} & a_{22} & a_{23} & \ddots & & & & \vdots \\ 0 & a_{32} & a_{33} & a_{34} & \ddots & & & \vdots \\ \vdots & \ddots & \ddots & \ddots & \ddots & \ddots & & \vdots \\ \vdots & \ddots & \ddots & \ddots & \ddots & \ddots & \ddots & \vdots \\ \vdots & & & & a_{(n-2)(n-3)} & a_{(n-2)(n-2)} & a_{(n-2)(n-1)} & 0 \\ \vdots & & & & \ddots & a_{(n-1)(n-2)} & a_{(n-1)(n-1)} & a_{(n-1)n} \\ 0 & \cdots & \cdots & \cdots & \cdots & 0 & a_{n(n-1)} & a_{nn} \end{pmatrix}$$

Figure: tridiagonal matrix

## 1 Introduction

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- Computational life science - FASTA
- Astronomy - FITS
- Climate Science - GRIB
- Climate & geo sciences - NetCDF

## 5 HDF5

## 6 Summary

## References

Computational life science - FASTA

- ASCII text based format for primary structure of nucleic acid and proteins
  - used with FASTA and BLAST algorithm (compare sequences)
  - no standard format suffix but usually .fa, .mpfa, .fna, .fsa or .fasta

## Format structure

- headline
    - begins with ">"
    - contains unique name and description of the sequence
  - comment symbol ";"
  - multiple sequences per file possible

```
1 >gi|2196610|emb|CAB09441.1| cytochrome b, partial (mitochondrion) [Acomys  
2 → spinosissimus]  
3 MTNRKTHPLLKIIINHAFIDLPPAPSNTSWWNFGSLLGICLIIQIITGLFLAMHYTSDTSTAFSSVTHIC  
4 RDVNYYGWLIRYLNHANGASMFICLFMVMGRGIYYGSYTYMETSNIGIILLFAVMATAFMGYVLPWGQMSF  
5 WGATVITNLLSAIPYIGTNLVEWIWGGFSVDKATLTRFFAHFILPFIIAALAMVHLLFLHETGSNNPTG  
6 INSDSDKIPFHPPYYTMKDLLGAFILLTLLALVLVSPDLDGDPNDYTPANPLNTPPHIKPQWYFLFAYAI  
7 LRSIPNKLGGVLALVLISLVLAILPLIHTSKQRSLMFRPISQTLFWILVANLLILT WIGGQPVEHPFIII  
8 GQLASISYFTIILILIPISGLIENKMMKWN
```

**Listing 1:** cytochrome b of Southern African spiny mouse (*Acomys spinosissimus*) (german: Zwergstachelmaus); taken from (miceFasta)

# Translation of Codes

- codes represents IUB/IUPAC standard

Code	Meaning	Code	Meaning
A	<b>A</b> denine	A	alanine
G	<b>G</b> uanine	G	glycine
T	<b>T</b> hyamine	T	threonine
K	K = GT ( <b>K</b> etone)	K	lysine
-	gap of indeterminate length	X	any
		*	translation stop
		-	gap of indeterminate length

**Table:** extract of nucleic acid codes

reference: Tao

**Table:** extract of amino acid codes, 25 acids + 3 special codes

# Compression

- very big files
- general purpose tools like gzip fall short
- many algorithms proposed to compress genomic data  $\implies$  MFCompress
- in comparison to gzip 50% additional compression but also computation time
- highly redundant datasets 8-fold of gzip possible

reference: Pinho und Pratas (2013)

# Applications

- GenomeTools & libgenometools (ZBH)
- Vmatch (ZBH)
- SeqAn
- FASTA/FASTQ parser in C<sup>1</sup>
- huge databases
  - National Center for Biotechnology Information (NCBI)
  - European Bioinformatics Institute (EBI)

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<sup>1</sup><http://lh3lh3.users.sourceforge.net/parsefastq.shtml>

# FITS

- Flexible Image Transport System
- standardized 1981, last release 2008
- designed for long-term archival storage  $\implies$  backwards compatibility - "Once FITS, always FITS"
- $\implies$  used in Vatican Apostolic Library

## Format structure

- HDU = header + data
- header: image card (80 character fixed-length ASCII strings) with key-value-pairs (size, origin, coordinates) and maybe comments
- data in multidimensional tables (variable length columns supported)
- also for non-image data e.g. spectra, data cubes
- contains several extensions e.g. x-ray and infra-red
- references supported
- internal storage sometimes in heaps

# Applications

- CFITSIO (Fortran, C)
- Aladin
- Detect the Dark Ages (LEDA) (24TB/day)<sup>2</sup>
- image viewer
  - GIMP
  - Photoshop
  - XnView
  - irfanView

references: Schwarzbürg (2005); Price u. a. (2015); Kayser (2012);  
fitsNasa

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<sup>2</sup>(see Price u. a., 2015)

GRIB

- GRIdded Binary (version 1), General Regularly-distributed Information in Binary form (version 2)
  - standardized by World Meteorological Organization's (WMO) Commission for Basic Systems (CBS)
  - store temperature, rainfall, wave height, ...

# Format

- grid discretization of room
- binary 2d-matrix
- collection of self-contained, independent records (messages)
- records contains sections
  - start and stop sequence
  - meta data e.g. origin, time, ...
  - dimension of grid, projection type
  - data
    - version 1: scaled to integer
    - version 2: compressed
- optional inventory: "table of contents" with user meta data and positions

# Tools

- *wgrib* like typical UNIX filter
- *degrib* creates indices for faster access
- *dkrz\_readgrib*<sup>3</sup>
- GUI: GRIBview<sup>4</sup>

references: Scherer (2009)

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<sup>3</sup><http://mms.dkrz.de/pdf/klimadaten/static/Pingo/post/post.dkrzgrib.html>

<sup>4</sup><http://www.theyr.com>

## Applications

- scientific applications
    - PINGO post-processing package (DKRZ)
  - common applications
    - zyGrib
    - web site: PassageWeather.com
    - Android app: Marine Weather | SailGrib Free

# Climate & geo sciences - NetCDF

- Network Common Data Format
- set of software libraries and self-describing, machine-independent data formats
- open standard maintained by University Corporation for Atmospheric Research (UCAR)
- originally based on NASA's Common Data Format but not compatible anymore
- version 4 allows use of HDF5

- self-contained, platform independent, binary
- dimensions
  - contain name and size
  - only one size unlimited (dataset dimension)
  - measurands e.g. time, length, ...
- variables
  - array of values with same type
  - contain name, datatype, shape
  - coordinate variable: one dimensional variable with same name as dimension
- attributes
  - meta data
  - used in variables and global
- conventions
  - standards for specific use case
  - compare files from different sources
  - e.g. Climate and Forecast (CF), Cooperative Ocean / Atmosphere Research Data Service (COARDS)

reference: netCdfArcGis

## Applications

- libraries for C, C++, Fortran, Java
  - third party: Perl, Python, MATLAB/Octave
  - ArcGIS

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## 5 HDF5

- Objects
- Modules
- Datatypes
- Optimization
- Tools
- Applications

# HDF5

- Hierarchical Data Format
- standardized by National Centre for Supercomputing Applications (NCSA) (1988), now developed by HDF Group
- binary file
- backward compatibility
- huge platform support
  - official: C, C++, Fortran, Java
  - third-party: Go, Python, R, MATLAB (Scilab, Octave), Mathematica, ERLANG, Perl, LabVIEW, ...
- contains datasets and groups  $\implies$  "datasystem"

# Objects

- dataset: single value or table of any dimension
- group: has name, attributes and contains groups and datasets
- attribute: any information for user e.g. simulation parameter
- meta data: information about content e.g. size of datatype (api)

# HDF5Viewer

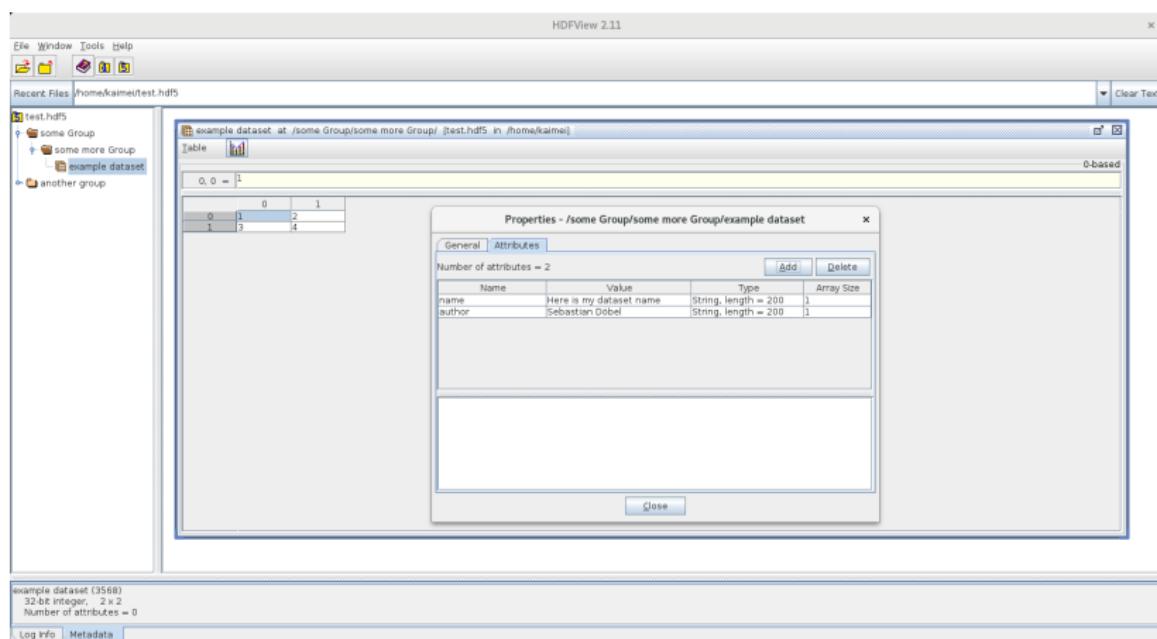


Figure: Screenshot from HDF5Viewer

# Modules

- separated in modules
- modules not independent
- e.g.
  - H5: library functions
  - H5A: annotation interface
  - H5F: file interface
  - H5G: group interface
  - H5Z: compression interface
  - ...

## Datatypes I

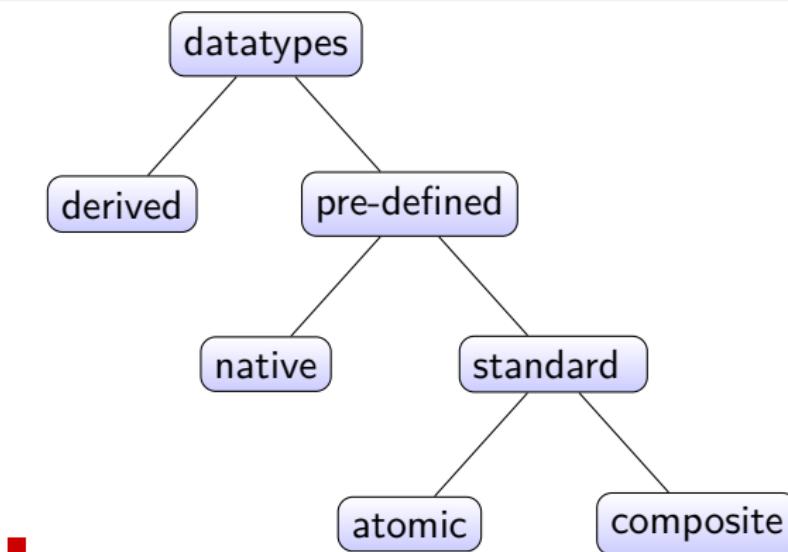
- own implementation of datatypes  $\implies$  portability
  - pre-defined and derived types
  - native: determine how standard types (atomic, composited) are stored
  - standard types: atomic and composited types

# Datatypes II

- atomic types
  - integer, float
  - string
  - reference
  - bitfield
  - date
  - time
  - opaque
- composed types
  - array
  - variable length
  - enumeration
  - compound datatypes

## Datatypes

## Datatypes III



## Figure: hierarchy of datatypes

# Datatypes IV

- pre-defined types create big bases
- possible to derive  $\implies$  change:
  - precision
  - size
  - character set
  - binary representation and interpretation
- create own application specific datatype

# Optimization I

- override system methods e.g. cache usage
- parallel IO
  - possible to read and write parallel in same datasets by multiple processes
  - file copy per process
  - uses MPI in background  $\implies$  parallel file system
  - user has to use *H5Pcreate* instead of *H5Fcreate*
  - usage only possible with C and Fortran
- compression
  - different algorithms implemented (performance vs. size)
  - possible to implement own algorithms

# Optimization II

- module *table*: any number of entries with same structure
  - unique number of types
  - unique size of types)
- module *packet table*:
  - different type sizes
  - possible to save strings with different lengths

# Tools

- post processing of already written data
- converting HD4 to HD5 (*h4toh5*, (*h5toh4*))
- import & export e.g.
  - export dataset of an image to GIF (*h52gif*)
  - import GIF file into HDF5 file (*gif2h5*)
  - create printable version (*h5dump*)
- split & merge files (*h5repart*)
- copy and compression (*h5repack*)
- compare two files (*h5diff*, *ph5diff*)
- performance tests (*h5perf*, *h5perf\_serial*)
- GUI: *HDFView* (Java-binding)

references: Kirchhart (2009)

# Applications

- FLASH (Hamburg Observatory)
- RAMS
- GNU Octave / MATLAB, Mathematica
- ParaView
- for more see ([hdf5Software](#))

reference: [hdf5Software](#); [HdfGroup](#)

# Summary

- sciences already have specialized data formats
- own format
  - what to store (data, meta data, ...)
  - text vs. binary based
- HDF5 provides fast, powerful storage interface for general problems
- HDF5 is support by many platforms
- migrating data format not easy

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