

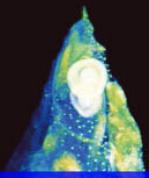
# Tomography data processing and management

P. Cloetens

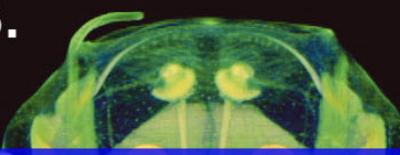
X-ray Imaging Group, ESRF

# Computed Tomography

A.



B.



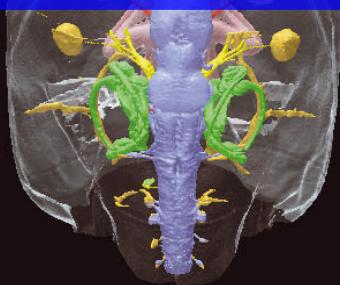
C.

*Xenopus tropicalis*  
(OsO<sub>4</sub> fixation)

**Motivation** complex materials  $\leftrightarrow$  3D microscopy  
'representative elementary volume'  
input for calculations:  $\mu$  structure  $\leftrightarrow$  properties  
**'non-destructive'**  
*in-situ* experiments ('dynamics', mechanics, ...)

D.

**Compromise spatial resolution**  $\leftrightarrow$   
micron - 100 nm

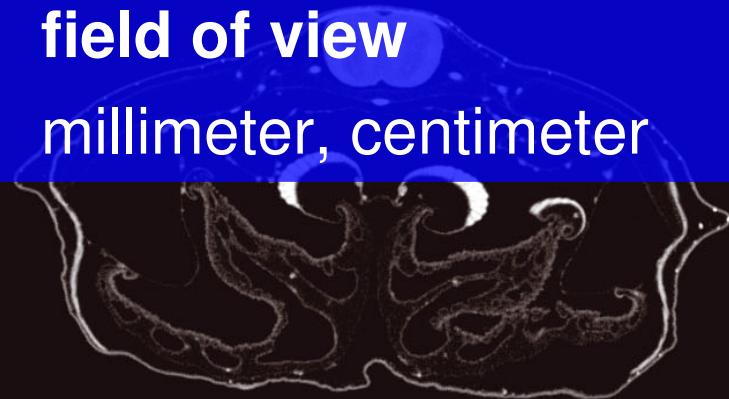


E.



F.

**field of view**  
millimeter, centimeter



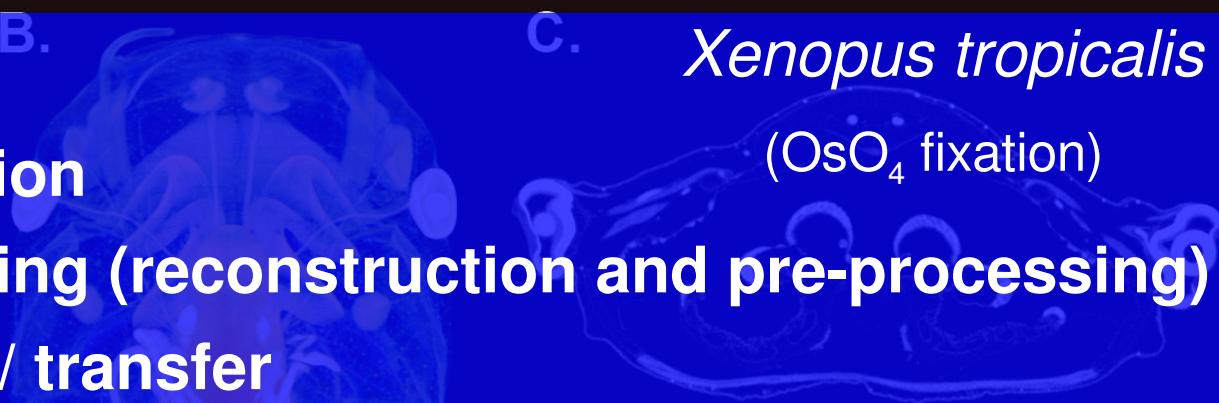
# Computed Tomography

A.  
**Current situation:**

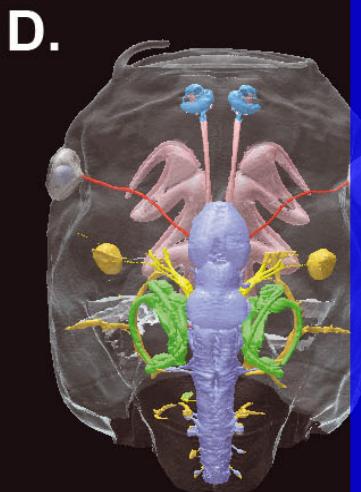
**data acquisition**

**data processing (reconstruction and pre-processing)**

**data storage / transfer**



C.  
*Xenopus tropicalis*  
( $\text{OsO}_4$  fixation)



**Grid opportunities:**

E.  
**data analysis**

**'on-line' data processing**

**large data sets**

**data transfer**

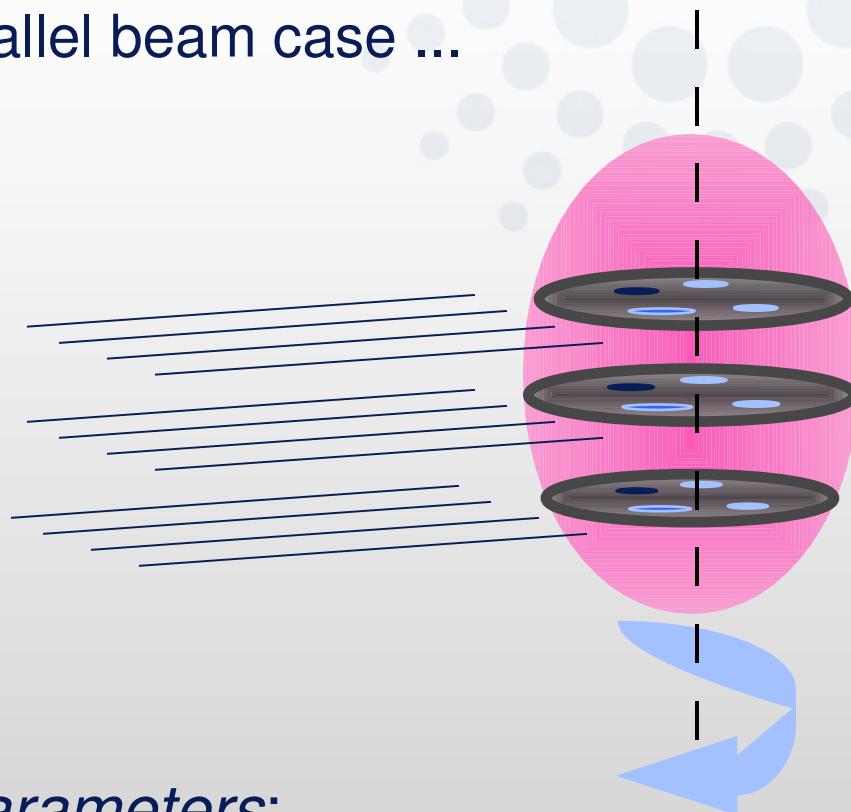
**sophisticated methods**



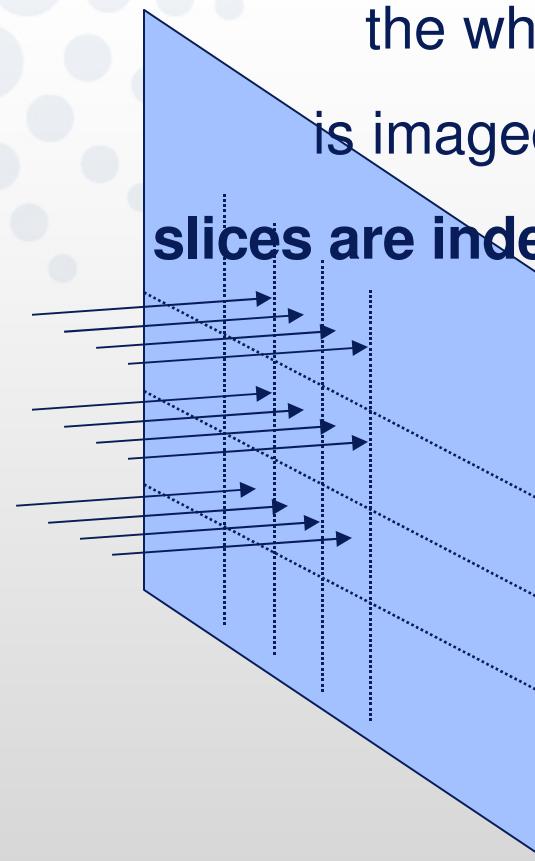
N Pollet (ibaic,u-psud), R Boistel (namc,u-psud), P Cloetens

# Synchrotron based tomography

Parallel beam case ...



the whole object  
is imaged at once,  
**slices are independent**



New parameters:

time, *in-situ* experiments

distance: holotomography, 3DXRD

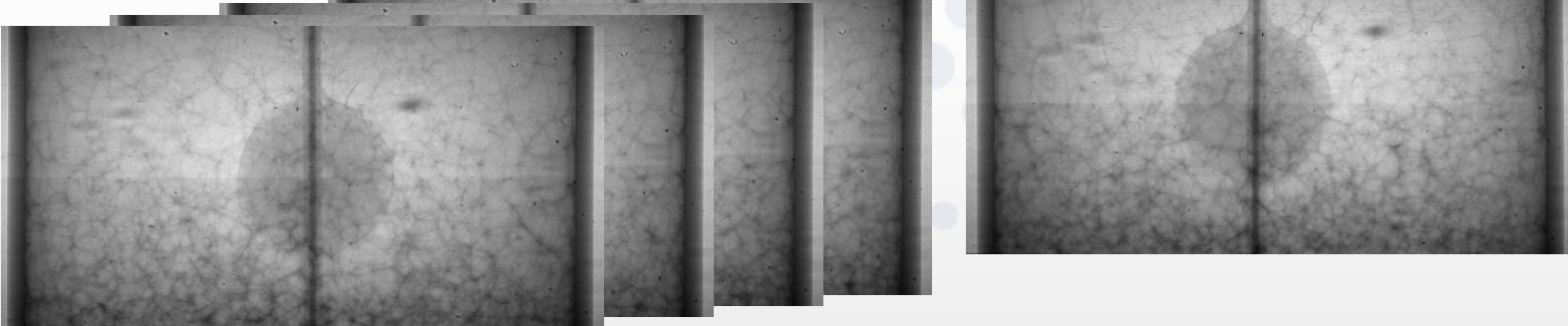
energy: edge CT, XANES, fluorescence

} **angles are  
independent**

# Micro-tomography work-flow

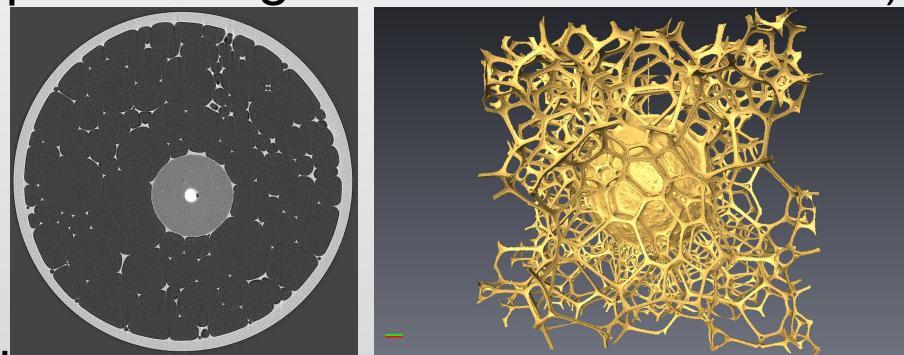
- **Data acquisition**

~ 1 second – 2 hours



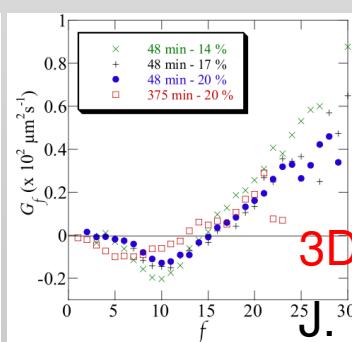
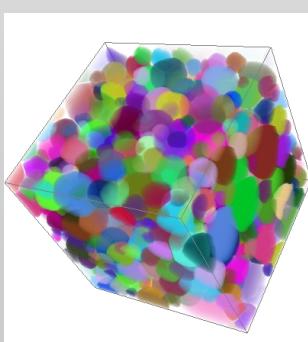
- **Data processing (pre-processing and 3D-reconstruction)**

1 hour – 6 months



- **Data analysis (extracting quantitative parameters, simulations)**

1 month –  $\infty$



J. Lambert et al, *PRL* 99, 058304 (2007)

# Data acquisition

- FReLoN (Fast Read-out Low Noise)
  - CCD (2k x 2k camera, '80 Mpixels/s', '160 MB/s')
- SPEC control (same macros on ID19, ID22NI, BM05, ...; Bliss)
  - but Device Servers for camera, ...
- Highly optimized
  - Step-by-step acquisition
  - or
  - Continuous acquisition
- Storage:
  - directly to NICE file servers over 1Gb / 10 Gb link
    - normal scans and network OK
    - local buffer on fast disks
      - fast tomography (~70 MB/s – 120 MB/s)
  - Raw data:
    - 16 GB / sample (64 GB for holo-tomography)
    - 1 TB / day in user experiment is feasible

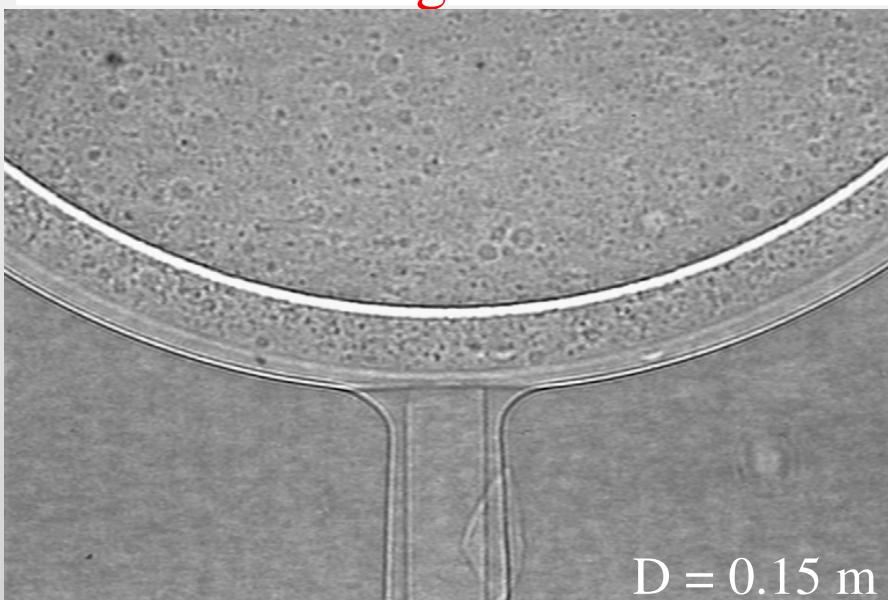
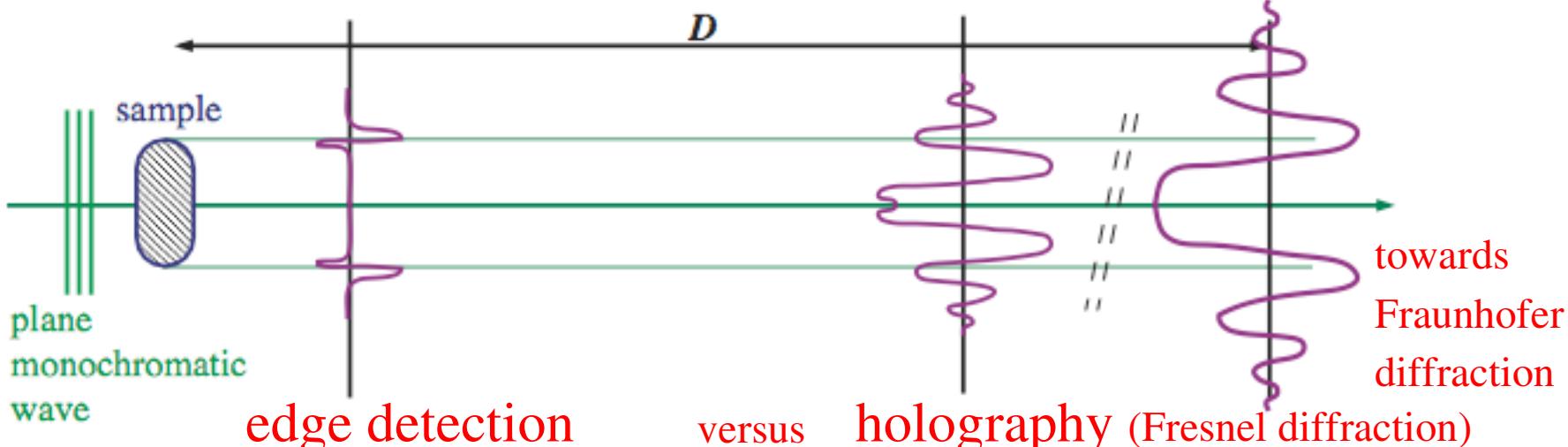
# Data processing

PyHST (SciSoft, A Mirone, C Ferrero) +



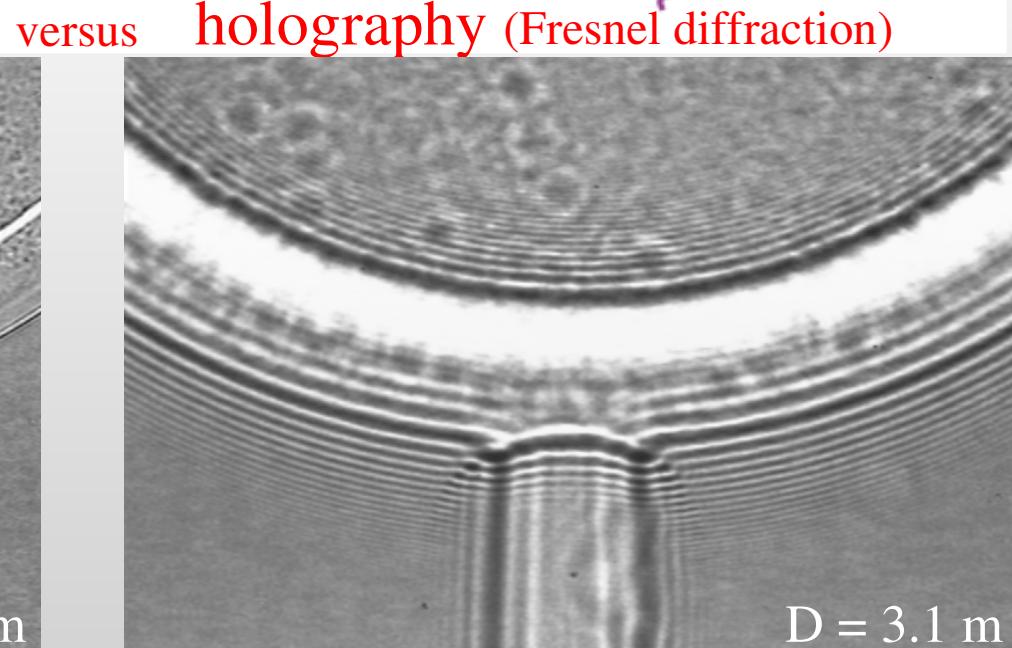
(CS, G Förstner)

- Filtered back-projection implementation
- Optimized for total through-put not for a single scan
  - parallelization in 1 GB sub-volumes not individual slices
  - 20 dedicated cpu's ( + 140 common cpu's )
- Improved through-put by including several pre-processing steps:
  - motion correction, basic ring correction, hot pixel correction, distortion correction, ...
- User interacts with Octave + ImageJ and terminal
- Special pre-processing with Octave (phase retrieval) or Matlab/IDL
- Volume reconstructed data  $\approx$  raw data ! (no data reduction)



each **edge** imaged independently

no access to **phase**, only to **border**  
 $\sqrt{\lambda} D \approx a$

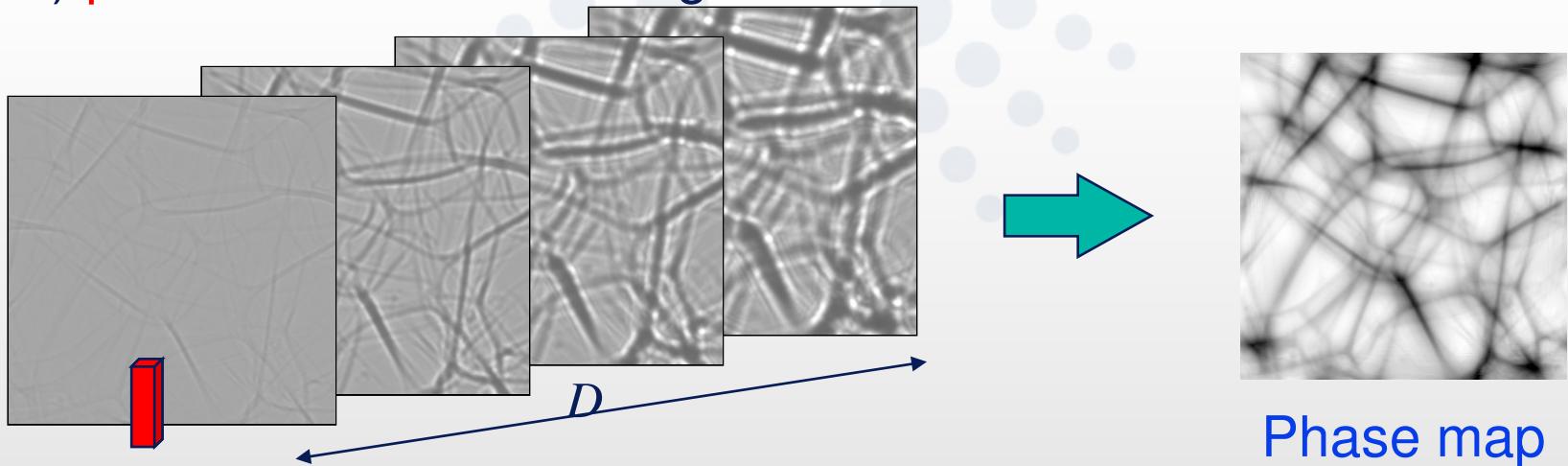


access to **phase**, if recorded at  $\neq D$ 's

$\lambda = 0.7 \text{ \AA}$  defocused **image**  
 $50 \text{ \mu m}$   
 $\sqrt{\lambda} D \ll a$

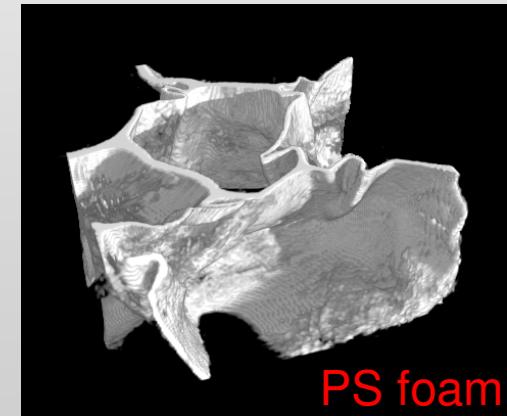
# Holo-tomography

1) phase retrieval with images at different distances



2) tomography: repeated for  $\sim 1000$  angular positions

3D distribution of  $\delta$  or the electron-density  
improved resolution  
straightforward interpretation  
processing



P.Cloetens et al., Appl. Phys. Lett. 75, 2912 (1999)

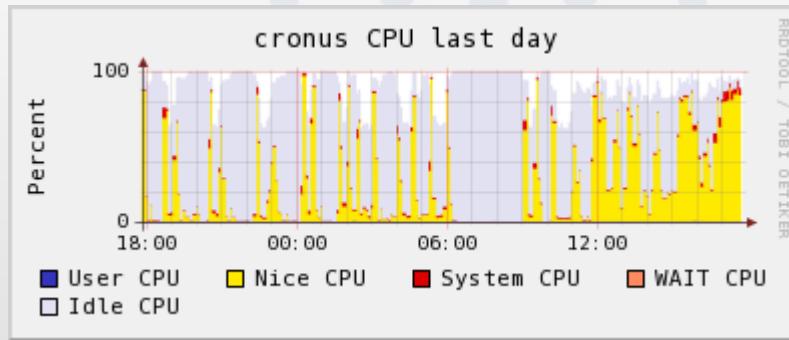
JP Guigay, M Langer, R Boistel, P Cloetens, Opt. Lett., 32, 1617 (2007)

# Data processing: limitations on parallelization

- Data Input – Output (file servers)

Too many reconstruction jobs blocks processing and acquisition (!)

Maximum 20 PyHST jobs on dedicated cluster cronus (no flocking)



- Memory intensive jobs (holo-tomography)

Only 4 or 8 GB / physical machine on NICE -> 1-2 GB / CPU

Performance collapses on swapping

In practice: only 8 GB machines used, 1 job per physical machine

# Data storage / transfer

- Tomography produces huge amounts of data (raw + reconstructed)
- Short term: NICE file servers
  - Case ID19 (~ 80 TB)

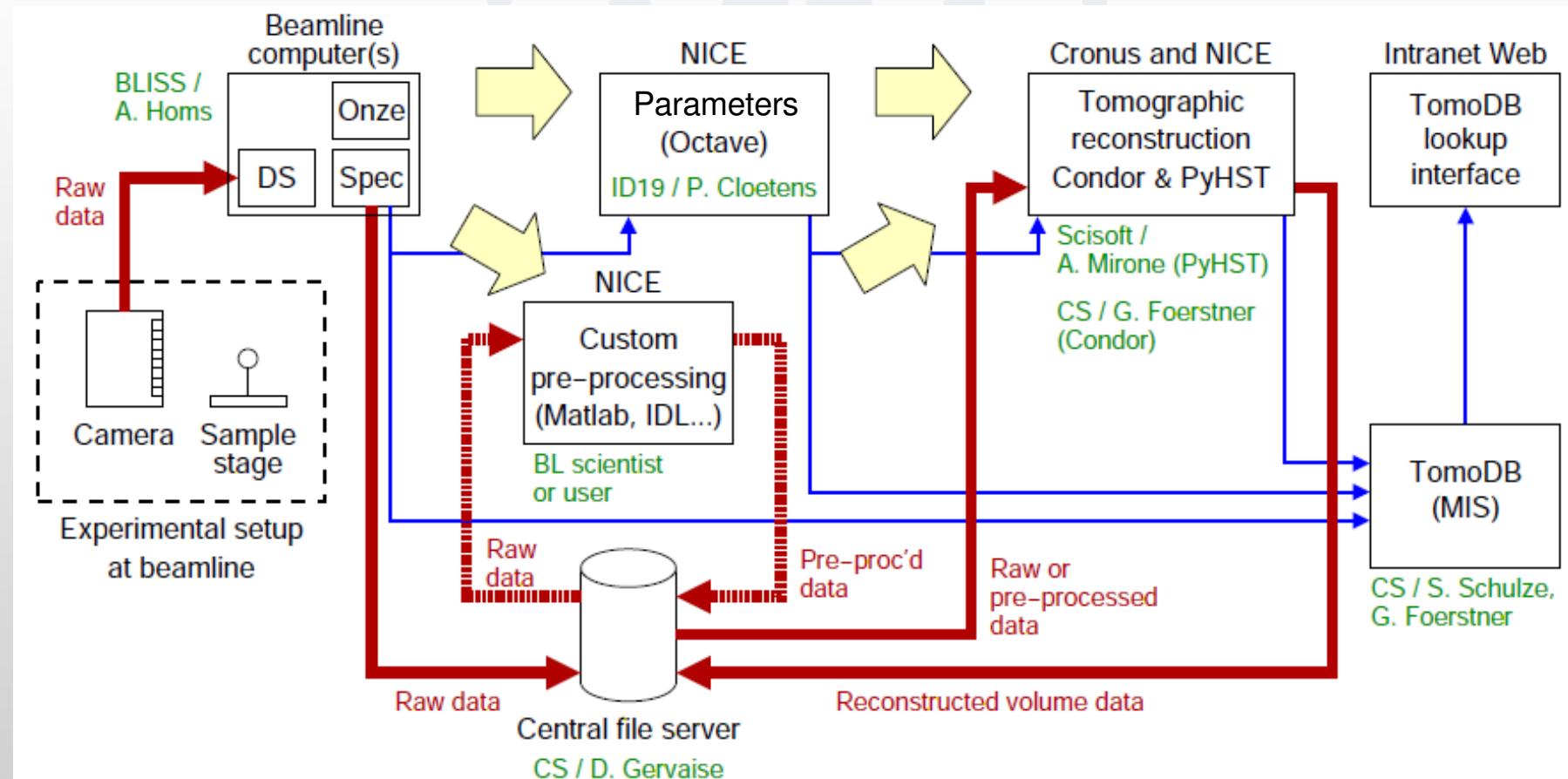
inhouse: 8TB (87% full)	paleo: 8TB (83% full)
inhouse1: 8TB (85% full)	lamino: 4TB (99% full)
external: 8.5 TB (83% full)	graintracking: 7 TB (99% full)
external1: 6 TB (89% full)	
external2: 6 TB (82% full)	
  - Medium term:
    - 6 months central backup on tape
    - many data restorations from central backup (man power)
  - Long term:
    - external hard disks / typically only reconstructed data
  - NICE also used for long term storage in practice (congestion)

# Database

- TomoDB II database (S. Schulze, MIS)
  - Most experimental conditions stored
  - Not all data processing parameters
  - Only available through intranet

The screenshot shows a Mozilla Firefox browser window with the title "TomoDB II - Consult page - Mozilla Firefox". The address bar contains the URL "http://vmis1:8080/tomodb/consult.do?id=27521". The main content area displays a tomographic reconstruction of a sample, with three views labeled "Projection 0°", "Projection 90°", and "Slice". Below the images is a section titled "Abstract". At the bottom, there is a table with columns "name", "description", and "value".

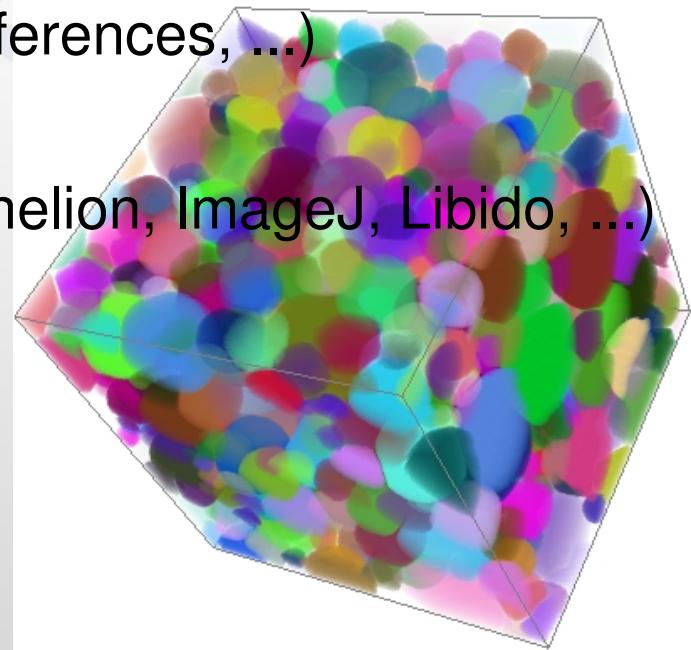
# Data Flow



courtesy: T. Weitkamp

# Grid opportunities: data analysis

- From simple segmentation over parameter extraction (local thickness, volume, ...) to simulations (finite elements, finite differences, ...)
- Presently left to user  
Several dispersed tools (VGStudio, Aphelion, ImageJ, Libido, ...)



- Faster data analysis can significantly increase publication rate
- Faster data analysis can significantly decrease data storage
- Users could perform their analysis with ESRF resources
- Users can make specialized code available

# Grid opportunities: 'on-line' data processing

- Presently: on-line reconstruction of sample slice
- On-line reconstruction of volumes
  - multi-scale approach (e.g. zoom tomography)
  - in-situ* experiments (result defines what to do next)
- Avoid processing time  $\gg$  acquisition time
  - extended stay of user
  - post-processing by user or beamline staff

# Grid opportunities: large data sets

- Nano-imaging:
    - improve spatial resolution with reasonable field of view
    - increase number of pixels, scales as  $N^3$
  - CCD's used today are only 2K x 2K
    - 2K x 2K → 16 GB
    - 4K x 2K → 64 GB
    - 4K x 4K → 128 GB
- Projection within the Upgrade Program:
- 8K x 8K → 1 TB** for a single scan !

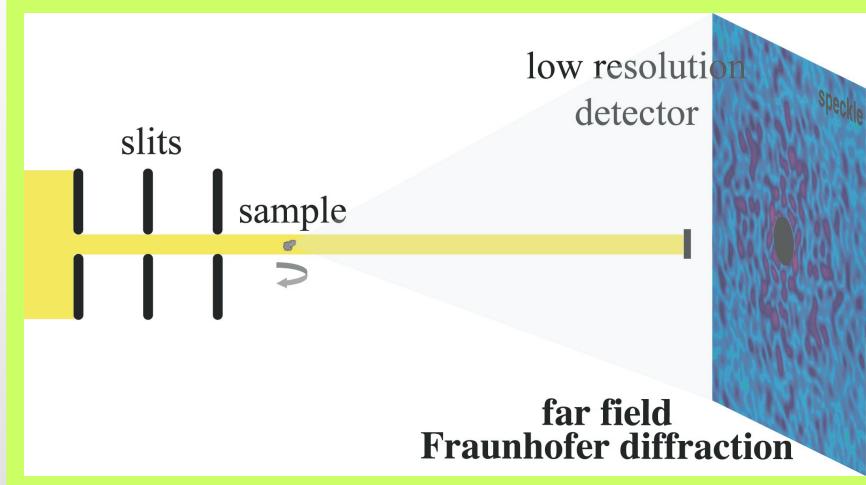
# Grid opportunities: data transfer and archiving

- External hard disk solution is impractical and not secure
- **Reliable** and fast data transfer over network
- Long term archiving for inhouse data is under discussion

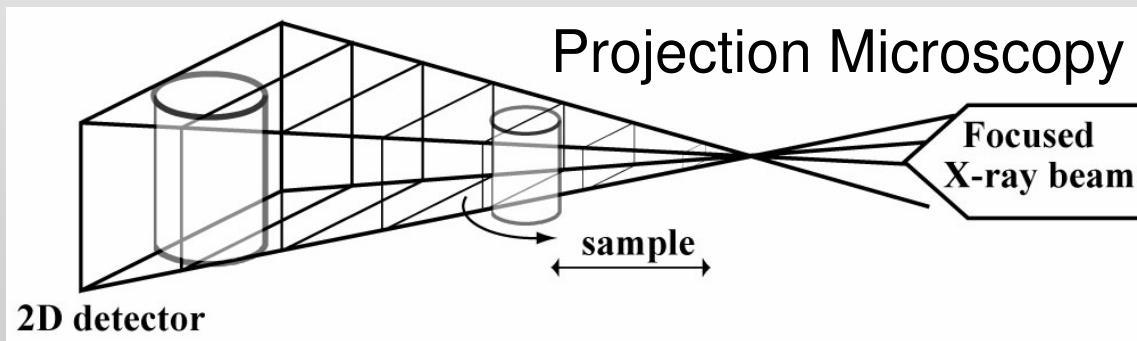
# Grid opportunities: sophisticated methods

- Nano-scale imaging:

## CDI (Coherent Diffraction Imaging)



## Projection Microscopy



- Very computer intensive methods (CPU power, memory usage)

# The Nano-Imaging end-station ID22NI



## Projection Microscopy - PXM

**electron density** Dose efficient, fast ; magnified holotomography

## Scanning Transmission Microscopy - STXM

**X-ray Fluorescence: element distribution** Slow ; trace elements

# Grid opportunities: Iterative Phase Retrieval

Fluotomography

Chemical identification

Magnified holotomography

Electron density mapping

CPU bound

Iterative minimization of cost functional

# Conclusions

- Tomography is a three steps process: data acquisition, data processing (reconstruction) and data analysis. The amount of data involved is huge. There is no data reduction before the data analysis.
- Parallelization is straightforward (independent slices or angles). It is limited in practice by data input/output.
- Possible grid opportunities:
  - data analysis
  - data transfer
  - more sophisticated methods of pre-processing, reconstruction