# HDF5/NeXus at ESRF

HDF5 Workshop ESRF - Grenoble September 2019

V. Armando SOLÉ ESRF – Data Analysis Unit

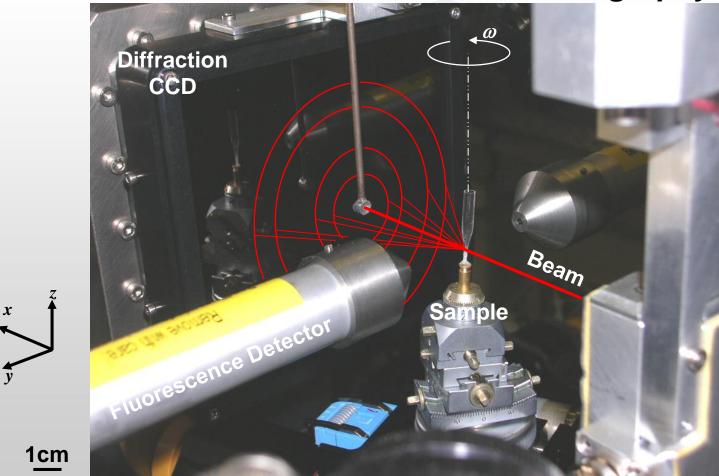


- ESRF Needs
- ESRF NeXus interpretation for raw data
- ESRF NeXus interpretation for processed data
- Other ESRF NeXus Uses: Metadata storage
- Status

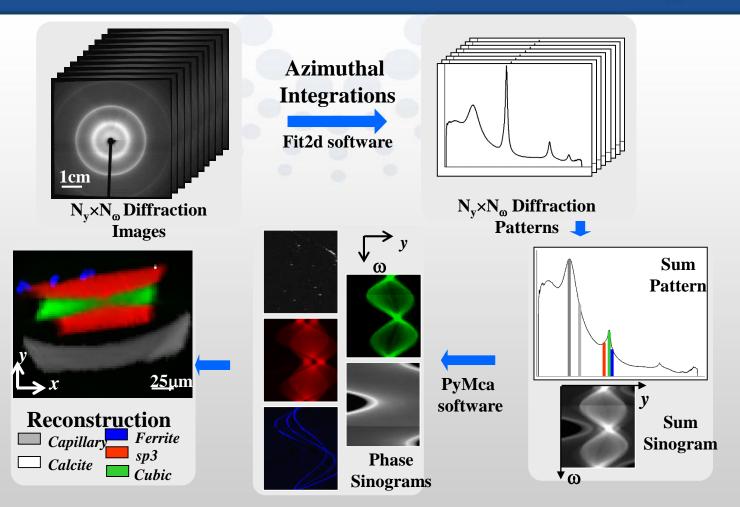




## **ID22 – Fluorescence-Diffraction Tomography**







Acknowledgements: Pierre Bleuet CEA - Grenoble



### **Data format issues**

- Currently
  - Diffraction images in EDF or MarCCD format
  - Fluorescence data in EDF or SPEC file format
  - Scan information in SPEC file format
  - Result of azimutal integration on Fit2D .chi format
- Preparing to move to HDF5

Lesson learned:

Try to avoid inventing a new data format. Use an existing one

Lesson NOT learned (yet?): Forget about ASCII just because you want to look at the file

## **ESRF** Needs

- Preparing to move to HDF5 in 2010 (previous three slides)
  - The analysis programs have changed and/or adapted
  - The ID22 beamline has moved (now ID16A and ID16B)
  - The ESRF has been dismantled and re-built
  - But we do not acquire our data in HDF5 yet

#### Still not moved to HDF5 !!!

- ESRF 2010 needs still there in 2019 (but we are prepared)
  - Single format to store different data types
    - Keep together counters, images, mca, ...
  - Compression support
  - Widespread support
  - Efficient and easy access to the data for visualization and analysis



## HDF5/NeXus – ESRF Interpretation for Raw Data

#### NXroot

Top level. One per file.

#### NXentry

One group per measurement

#### **NXinstrument**

Describe the instrument. Only one per NXentry

#### measurement (@NXcollection)

Flattened view of everything measured

Only one per NXentry

#### sample (@NXsample)

Define the physical state of the sample during the scan

#### **NXdata**

The default data to be plotted. One NXdata group per plot user (@NXuser)

Details of a user, i.e., name, affiliation, email address, etc

#### **NXsubentry**

Data or links to data for particular analysis

Exclusive Acquisition Domain

Almost exclusive Acquisition Domain

**User/Scientist** Domain

**User/Scientist** Domain

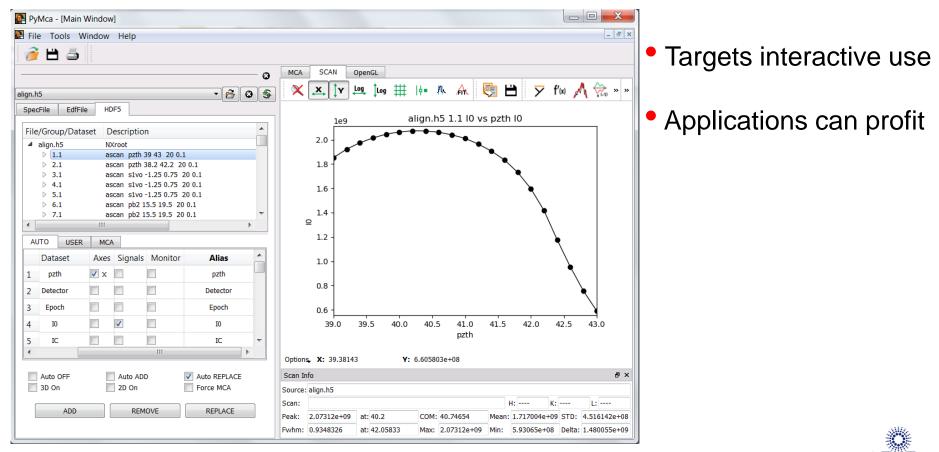
Administrative Domain (GDPR? DOI?)

Analysis Domain



## **Measurement Group Convention**

#### Name-based convention followed by ESRF and Sardana (MAX IV, ALBA...)



## **HDF5/NeXus: Requirements for Processed Data**

- NeXus conventions are fairly clear in what concerns raw data
- How to store processed data in HDF5 files?
  - Needs
    - Program used
    - Configuration parameters
    - Results
    - Minimize file creation
      - More than one data treatment step into the file
      - Describe data treatment sequence



## **NeXus: ESRF Implementation for Processed Data (v1)**

Goals can be achieved with "extended" NXprocess groups

entry

start time end time title process\_name@NXprocess program\_name version date sequence\_index configuration@NXcollection if HDF5 supported by program results@NXcollection or NXdata if plot

#### Just a name based convention added to NXprocess



## **NeXus: ESRF Implementation for Processed Data (v2)**

 A 100% pure NeXus way to specify the configuration: NXnote entry

process\_name@NXprocess program\_name version date sequence index configuration@NXnote file name type data results@NXcollection or NXdata if just a plot

The key point is that the configuration can be used back. We have to be able to feed the original program with it.



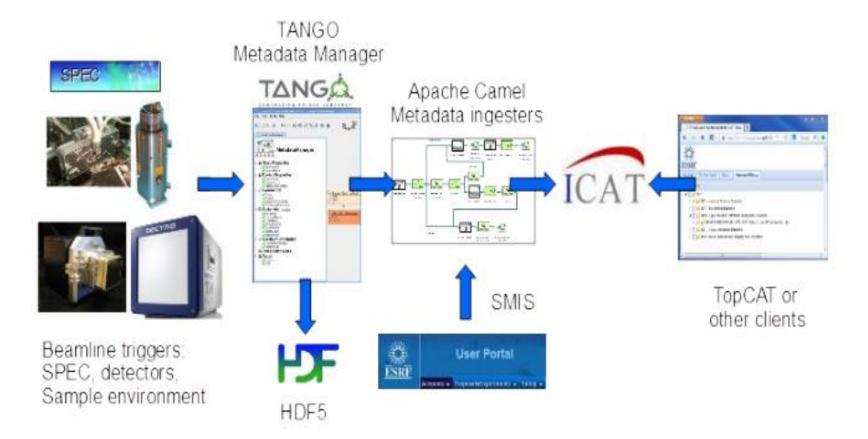
## Metada Storage: ICAT – NeXus Mapping





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## **Metadata Collection Architecture**





## ICAT – NeXus Mapping

- Clear mapping from existing NeXus conventions to ICAT
  - ICAT key = Class1Class2Class3\_dataset@attribute
  - Ex: NeXus current and mode in class Source inside class Instrument
    - InstrumentSource\_current
    - InstrumentSource\_mode

#### Technique or beamline specific information as NXsubentry based keys

#### <prove NX\_class="NXsubentry" groupName="EM"> <protein\_acronym ESRF\_description="Protein acronym" NAPItype="NX\_CHAR">\${EM\_protein\_acronym}</protein\_acronym> <voltage ESRF\_description="Voltage" NAPItype="NX\_CHAR">\${EM\_voltage} <magnification ESRF\_description="Magnification" NAPItype="NX\_CHAR">\${EM\_magnification}</magnification> <images\_count ESRF\_description="Number of images in movie" NAPItype="NX\_CHAR">\${EM\_images\_count}</magnification> <position\_x ESRF\_description="Position X" NAPItype="NX\_CHAR">\${EM\_position\_x> <position\_y ESRF\_description="Position Y" NAPItype="NX\_CHAR">\${EM\_position\_y> <dose\_initial ESRF\_description="Dose initial" NAPItype="NX\_CHAR">\${EM\_dose\_initial> <dose\_per\_frame ESRF\_description="Dose initial" NAPItype="NX\_CHAR">\${EM\_dose\_initial> <dose\_per\_frame ESRF\_description="Dose per frame" NAPItype="NX\_CHAR">\${EM\_dose\_initial> <dose\_per\_frame ESRF\_description="Spherical aberration" NAPItype="NX\_CHAR">\${EM\_dose\_initial> <dose\_per\_frame ESRF\_description="Spherical aberration" NAPItype="NX\_CHAR">\${EM\_spherical\_aberration} <spherical\_aberration ESRF\_description="Spherical aberration" NAPItype="NX\_CHAR">\${EM\_samplitude\_contrast} <spherical\_aberration="Serf\_description="Amplitude contrast" NAPItype="NX\_CHAR">\${EM\_sampling\_rate ESRF\_description="Spherical\_aberration"



## ICAT – NeXus Mapping

#### Collected files as list inside an NXcollection group named measurement

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#### Aiming at using HDF5 files and external links



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## Status of HDF5/NeXus @ ESRF – Raw Data

- Acquisition
  - SPEC (legacy control system)
    - HDF5 not worth as native output. Use *silx convert* if desired
  - Bliss (new ESRF control system)
    - HDF5 native output operational
  - LImA (ESRF Library for Image Acquisition)
    - HDF5 native output operational
    - Using Direct Chunk Write for efficient multithreaded compression

#### Why that functionality is not offered by HDF5 itself?



## Status of HDF5/NeXus @ ESRF – Data Analysis

#### Data Analysis

- Capability to read HDF5 files (preferred data analysis I/O format)
- Unified (h5py-like) API to access all data formats
- Support of NeXus NXdata I/O in viewers and analysis codes
- Provide provenance via NXprocess (pyFAI, PyMca, PyNX,...)
  - Only one NeXus application definition supported (NXcxi)



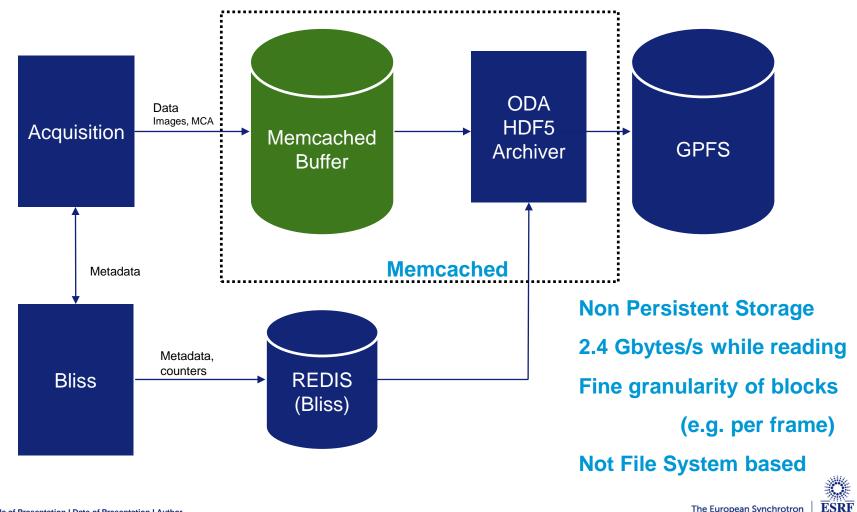
## Status of HDF5/NeXus @ ESRF – Online Data Analysis

#### HDF5 is not appropriate for online data analysis

- HDF5 Specific reasons
  - Potential concurrent access issues
    - SWMR is not enough (yet?): static file structure, flush...
      Ex. It can be OK for tomography but not for spectroscopy
- Generic reasons
  - If we have to read from a file anything will be slow
    - Basically no data format is appropriate
    - Studying to externalize via REDIS + memcached
      - Writing a file becomes a particular case of ODA
      - File writing should be the last step, not the first one!



#### **Online Data Analysis – HDF5 Writing a Particular Case of ODA**



### Status of NeXus @ ESRF - Metadata

- Data Policy and NeXus
  - Mirror ICAT and NeXus master file done
  - External links between master file and raw HDF5 files desirable

#### Ideally one should enable processing a dataset\* from its master file

(\*) A dataset in this context is not an HDF5 dataset but a collection of data



#### The Weight of Legacy

- Adoption of HDF5/NeXus has been slower at the ESRF than at other synchrotrons due to the raw data being acquired in different formats. Detector output in HDF5 and the deployment of Bliss are speeding things up.
- User experience with HDF5 files has to be better than with legacy formats HDF5 should not be the question but the answer.
- Concerning data analysis, ESRF started to provide HDF5 support in 2009. Currently making convenient **use of the NeXus formalism as output and as integral part of the ESRF data policy**.
- HDF5 and NeXus great for archival. Efficient online data analysis needs to avoid the use of files as input and HDF5 is no exception.



