

PyMca, Hyperspectral Data and HDF5

V.A. Solé - European Synchrotron Radiation Facility HDF5 as hyperspectral data format workshop



PyMca?

PyMca is set of software tools mostly known in the field of XRF analysis

It is certainly a set of programs and widgets for XRF analysis:

Spectrum modeling

Quantification

ROI imaging

Fit imaging via batch processing

But also a set of programs and generic python modules for:

Data visualization

Peak search

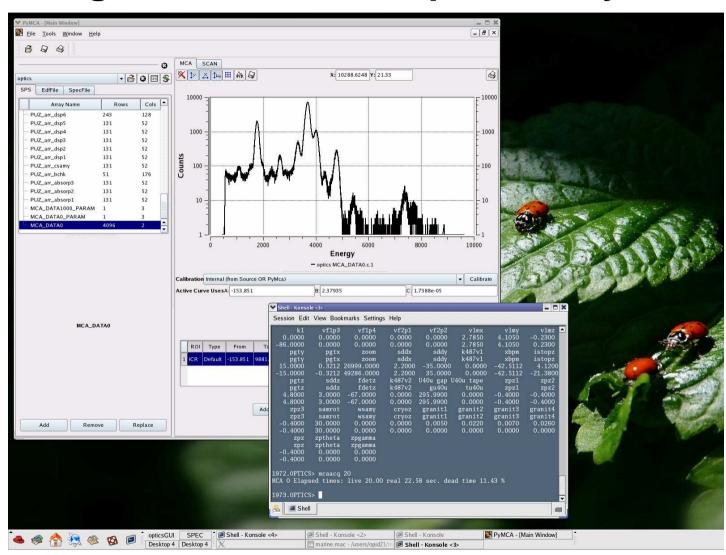
Function fitting

Imaging of stacked data

V.A. Solé, E. Papillon, M. Cotte, Ph. Walter, J. Susini, Spectrochimica Acta B 62 (2007) 63-68



Integration into our acquisition system



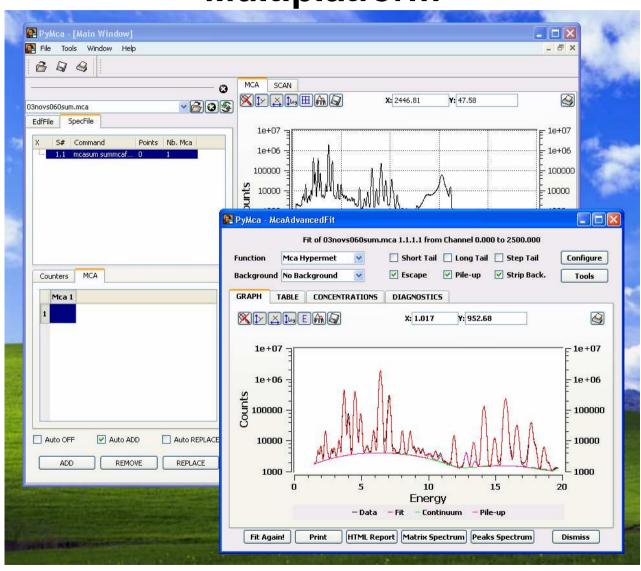


Free distribution



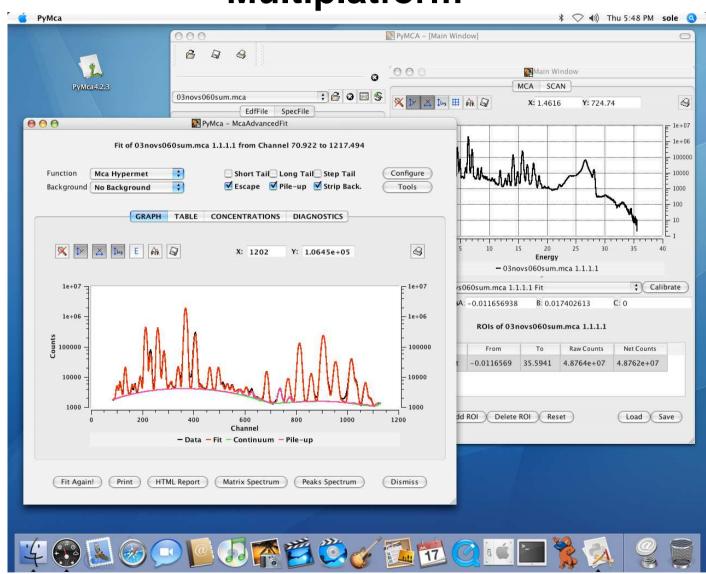


Multiplatform



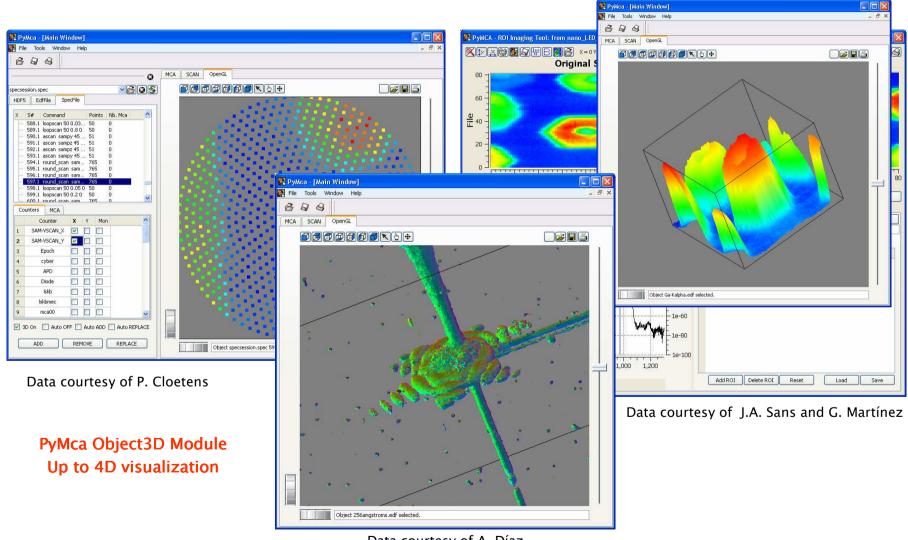


Multiplatform





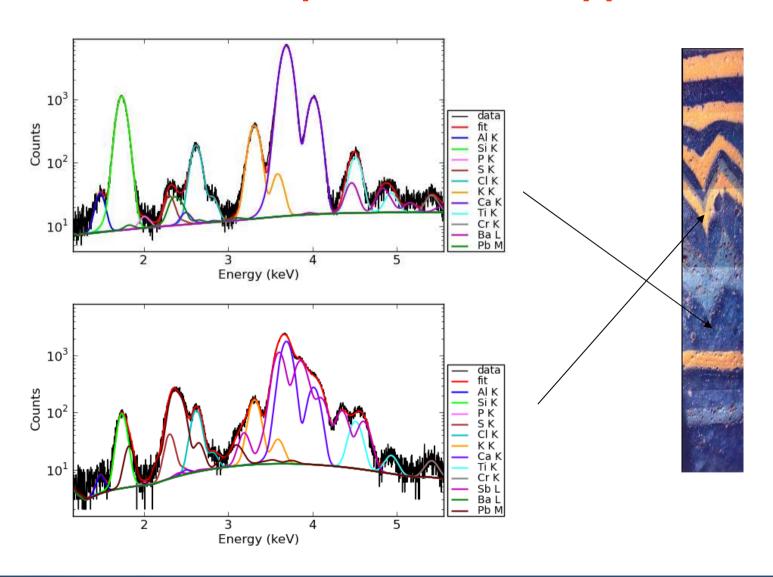
PyMca Visualization



Data courtesy of A. Díaz



Complete M-shell support





Use of (sub-)shell mass attenuation coefficients

Direct photoelectric ionization of the i (sub-)shell:

PyMca
$$P_i \propto \left(\frac{\tau_i\left(E,i\right)}{\tau(E)}\right) au(E)$$
 versus $P_i \propto \frac{J_i-1}{J_i} au(E)$

Term between parenthesis for Pb

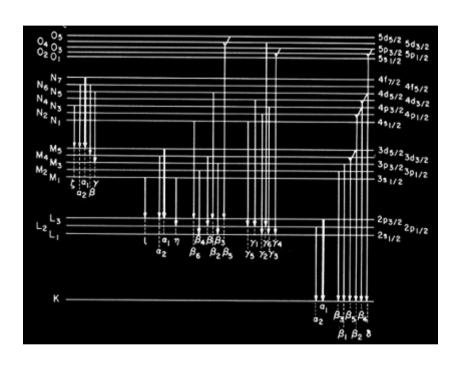
Energy	16 keV	20 keV	25 keV	30 keV
L1	0.136	0.164	0.192	0.218
L2	0.248	0.247	0.244	0.240
L3	0.367	0.345	0.323	0.303

J.H. Scofield. Theor. Photo. Cross Sections from 1 to 1500 keV, LLNL Report UCRL-51326, Livermore, Ca 1973.

A. Brunetti, M. Sánchez del Río, B. Golosio, A. Simionovici, A. Somogyi, Spectrochimica Acta B 59 (2004) 1725-1731



De-excitation cascade taken into account

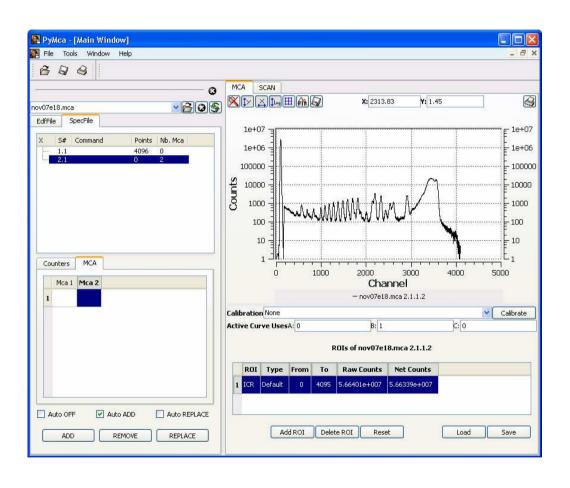


PyMca considers the vacancies produced in an atomic shell or sub-shell by the de-excitation process of an inner shell or sub-shell.

Radiative and Coster-Kronig transitions correctly considered. Auger transitions approximated.



XRF Spectrum Analysis



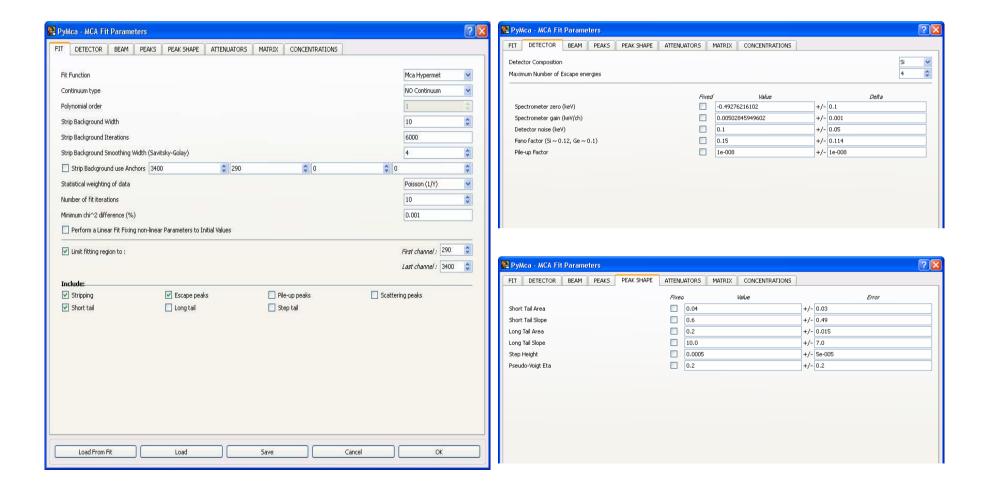
Typical procedure:

- 1. Calibration
- 2. Peak identification
- 3. Peak area extraction
 Region of interest (ROI)
 Deconvolution (FIT)
- 4. Quantification

Documentation at http://pymca.sourceforge.net/documentation.html

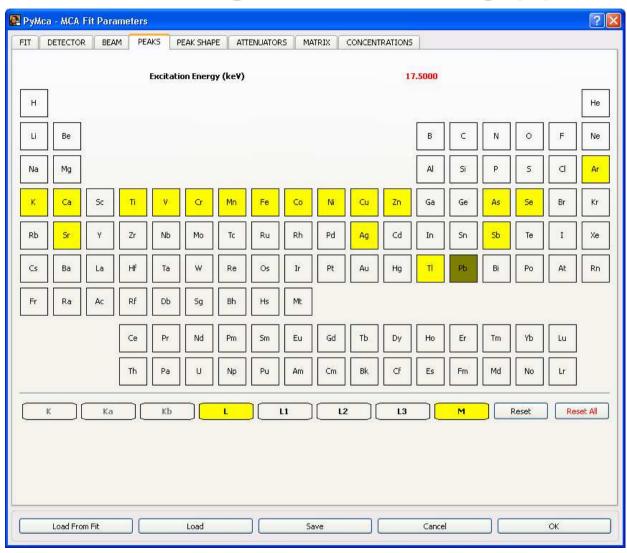


Fit configuration Dialog (I)





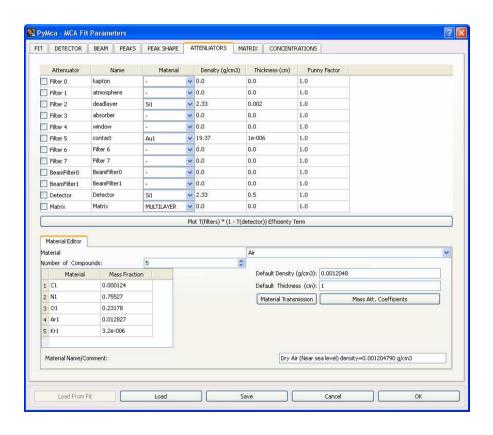
Fit Configuration Dialog (II)

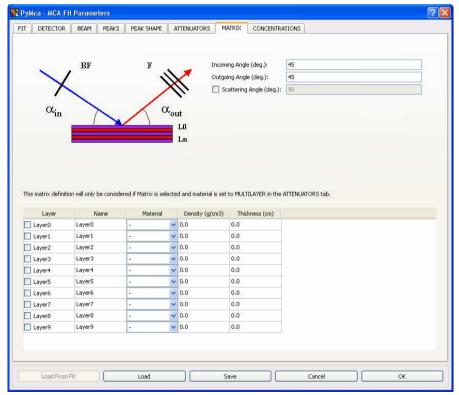




Quantification (I)

Parallel beam approximation

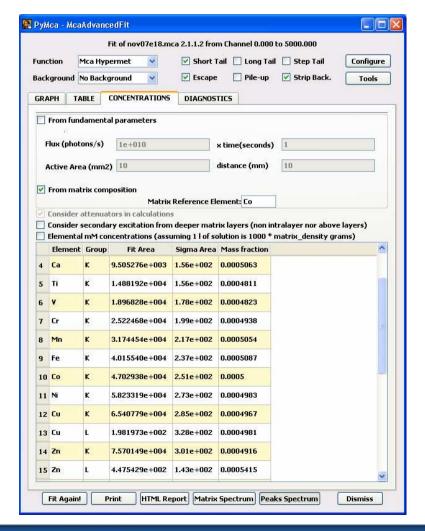


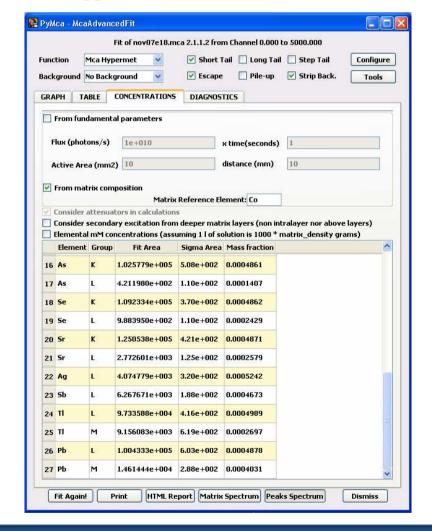




Quantification (II)

Nominal concentration 500 ppm





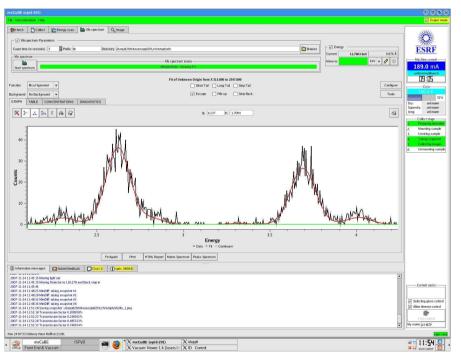


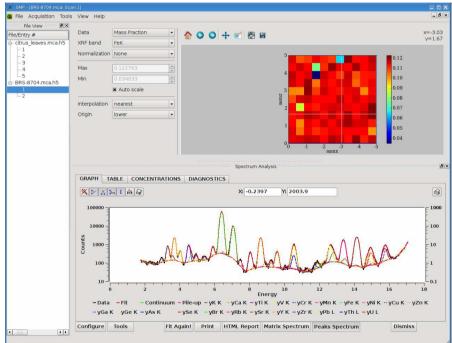
XRF Analysis Integration in other Applications

Integration in mxCuBE (ESRF)

Integration elsewhere









Is it easy to embed?

For the previous examples, basically one just needs 4 lines of code:

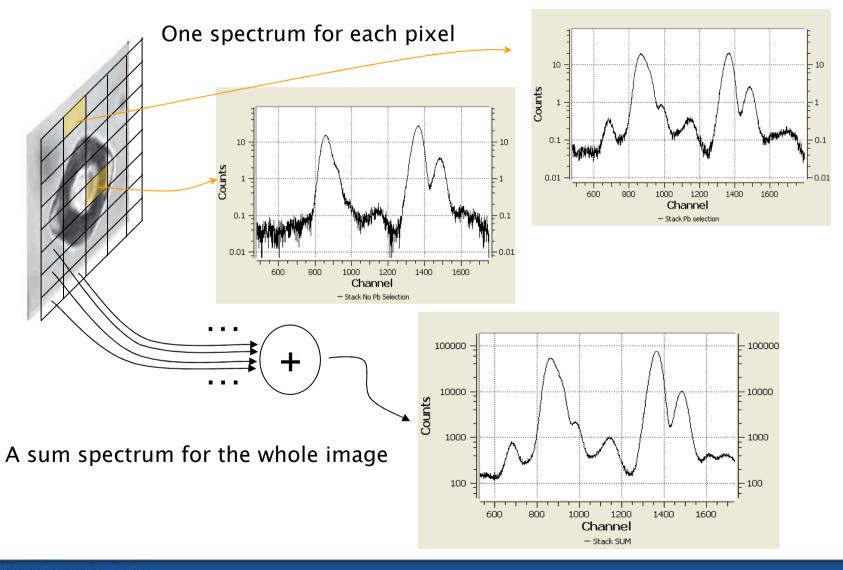
```
from PyMca import McaAdvancedFit
```

```
fitWindow = McaAdvancedFit.McaAdvancedFit()
fitWindow.setData(x, y)
fitWindow.show()
```

It can be used interactively from ipython just starting it as "ipython -q4thread"

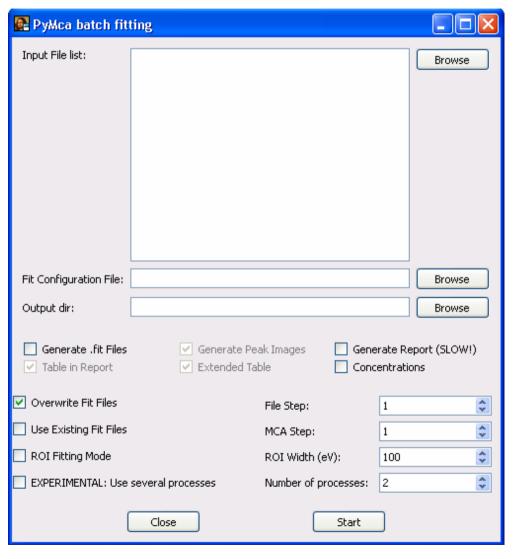


XRF Imaging





Advanced Fit Batch processing



Select the input files

Select the fit configuration Select the output directory Select the output options

Start



Output

Images in ASCII and ESRF format

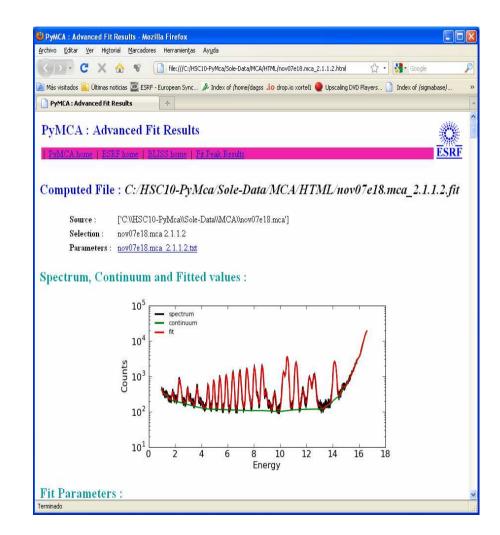
Easy to import in other programs

Individual peak contributions in ASCII

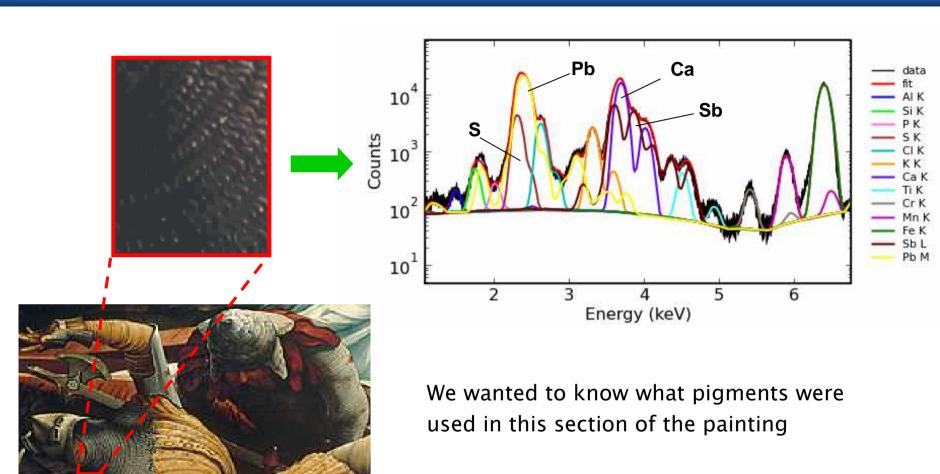
Use your own plotting program

Fully automated HTML report

Browse your results!





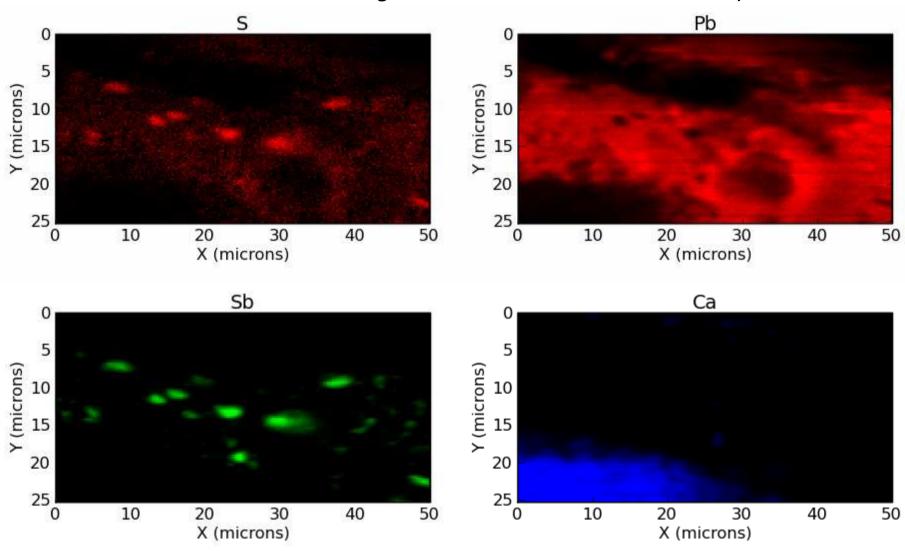


The European Light Source Slide: 21

Copyright C2RMF

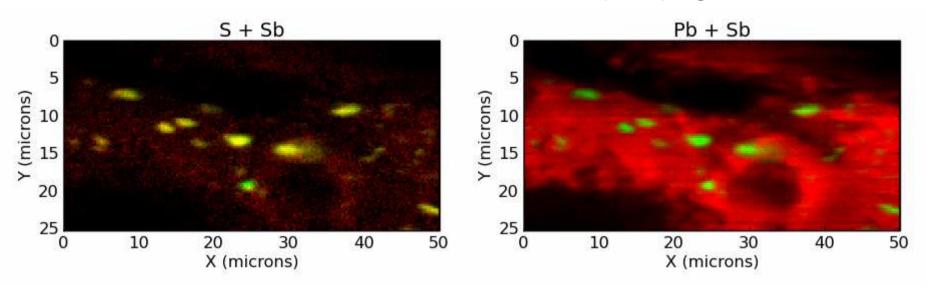


Based on the batch generated element distribution maps ...





... and their correlations as shown by the program



Sulfur and antimony correlated

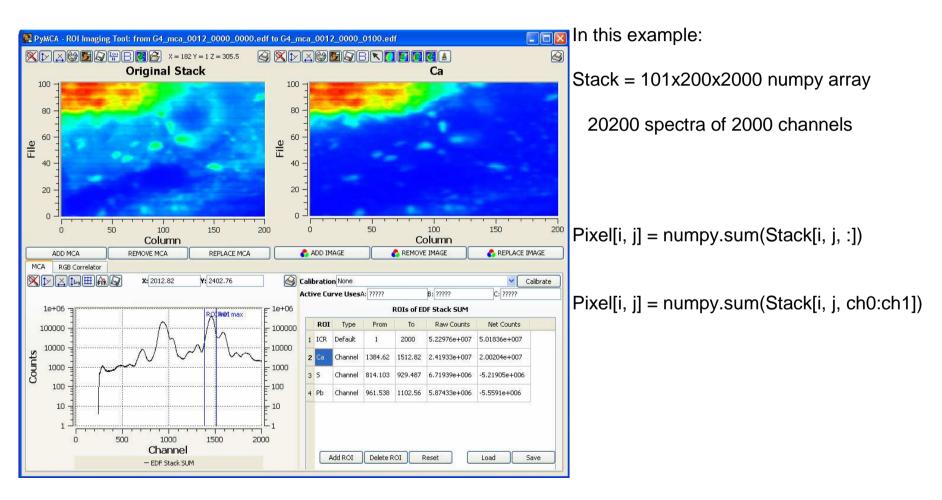
Lead and antimony not correlated

... we were able to determine the possible presence of stibnite grains (Sb_2S_3) embedded in a lead containing matrix.

M. Cotte, E. Welcomme, V.A. Solé, M. Salomé, M. Menu, Ph. Walter, J. Susini, Anal. Chem. 79 (2007) 6988-6994



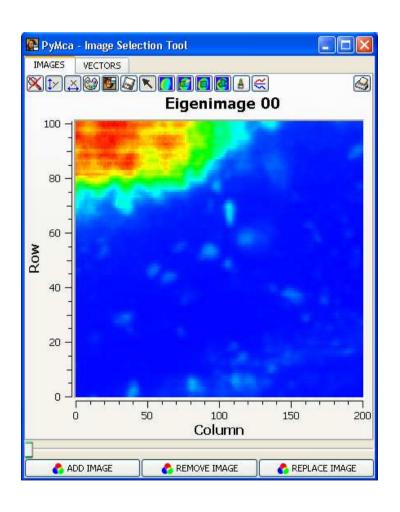
Stack ROI Imaging

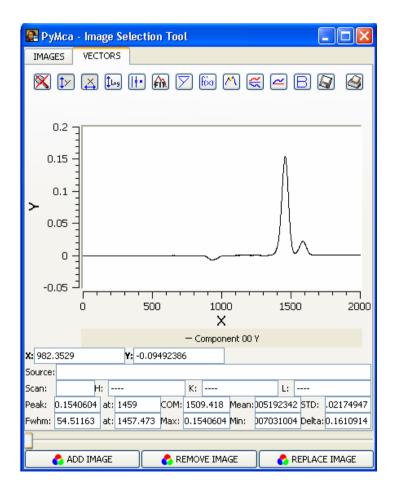


We can generate new images by moving the cursors or defining new ROIs in the table



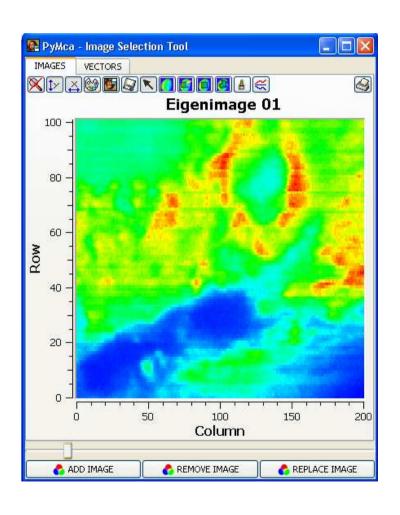
and

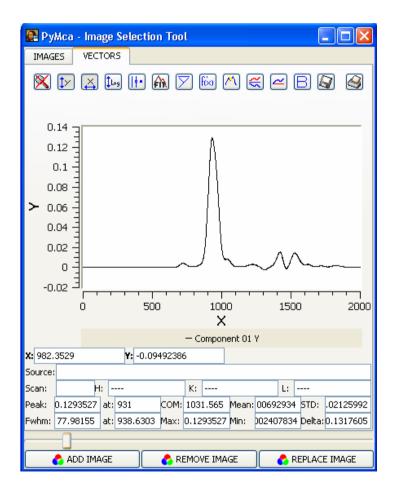






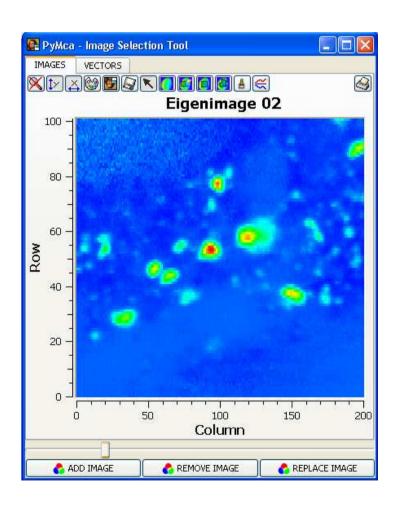
and

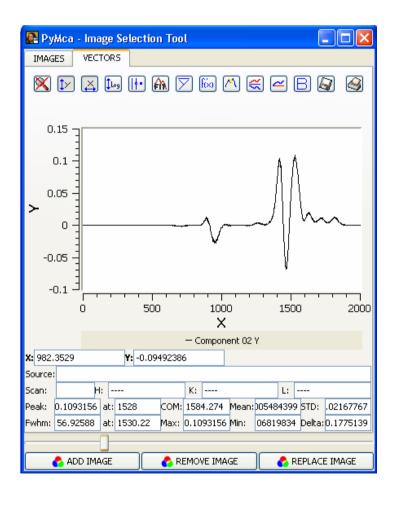






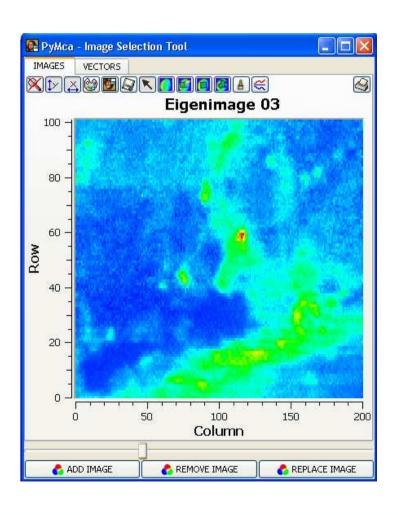
and

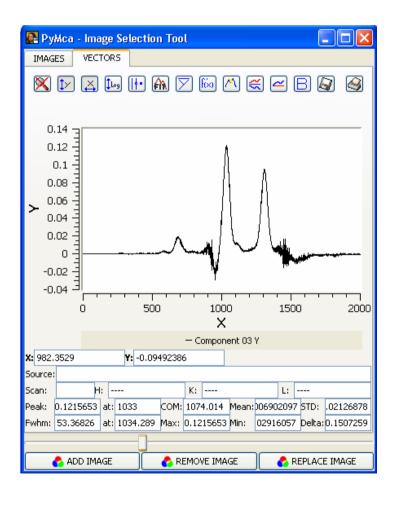






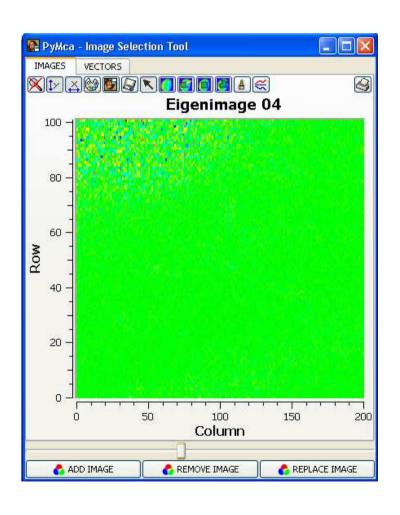
and

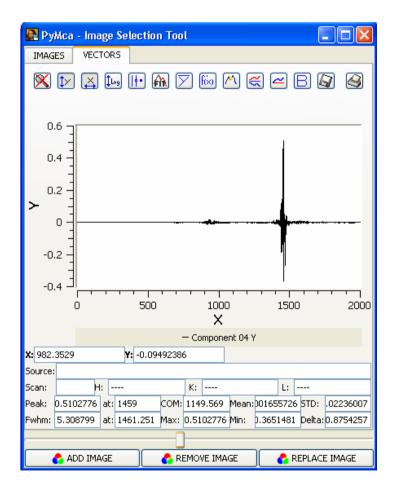






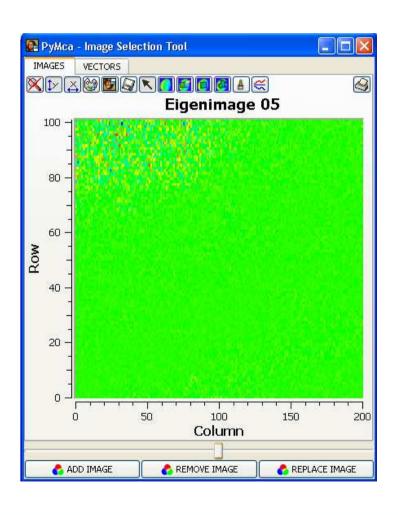
and

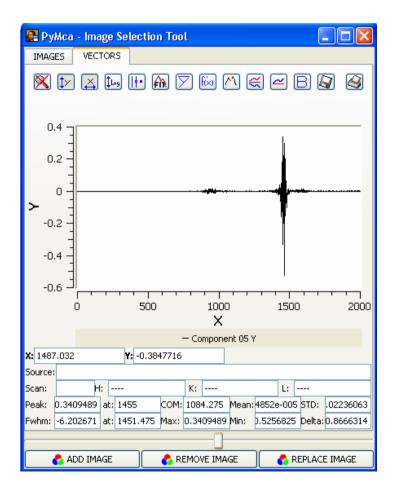






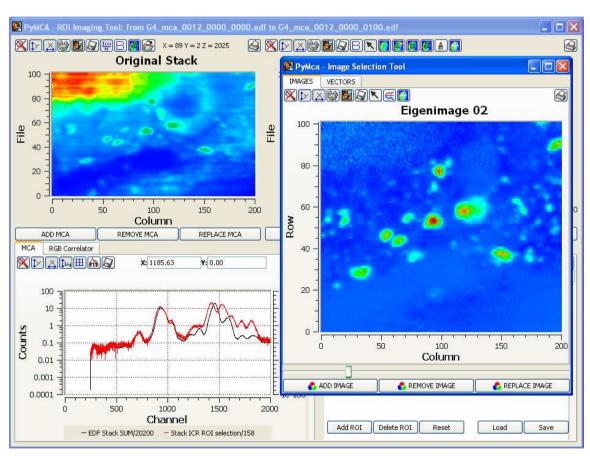
and







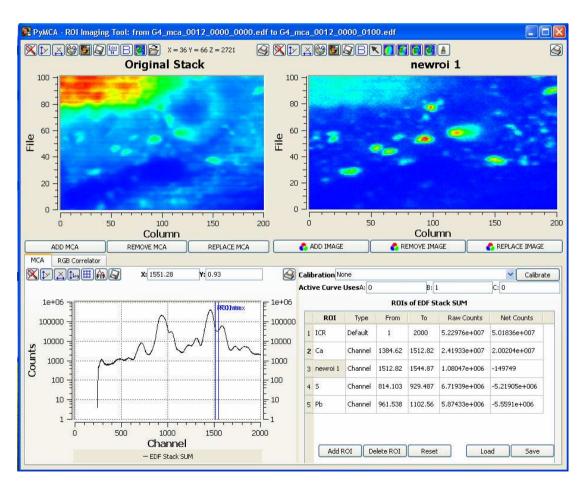
Getting the actual information



We can select a set of pixels on any of the displayed images and display the cumulative spectrum associated to those pixels.

Here we can see the average spectrum associated to the hotter pixels of the Eigenimage 02 (in red) compared to the average spectrum of the map (in black).





We could have easily missed the presence of one element if we would have just analyzed the sum spectrum via ROIs.



What have we done?

We have used multivariate analysis (PCA in this case) to know what sample regions were worth to take a closer look.

Not bad when you have a lot of data ...

This data treatment is totally generic and applicable to other techniques



Multivariate Analysis Capabilities

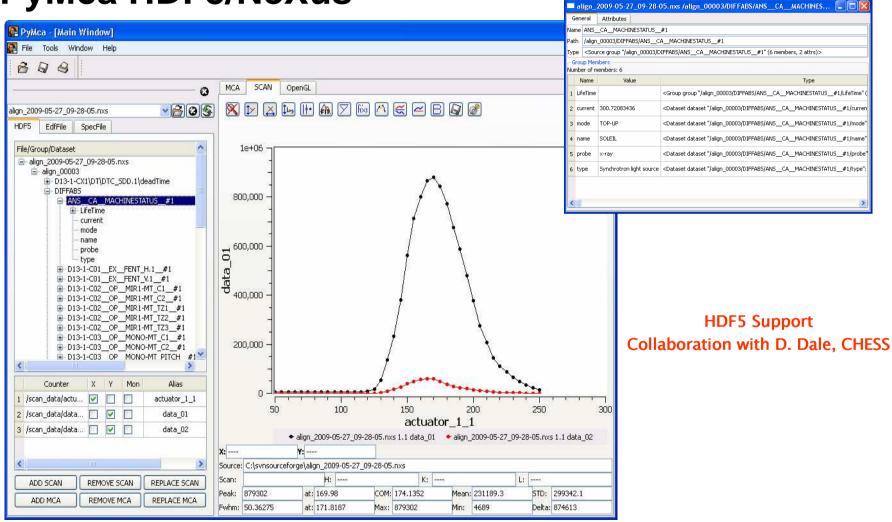
PyMca makes use of PCA, ICA (via MDP toolkit) and NNMA (via py_nnma)

For the time being, multivariate analysis is used just to identify sample regions with different properties. The associated physical spectrum gives the actual information.

It can combine information from different simultaneously acquired datasets, for instance XRD and XRF data or PIXE and RBS.



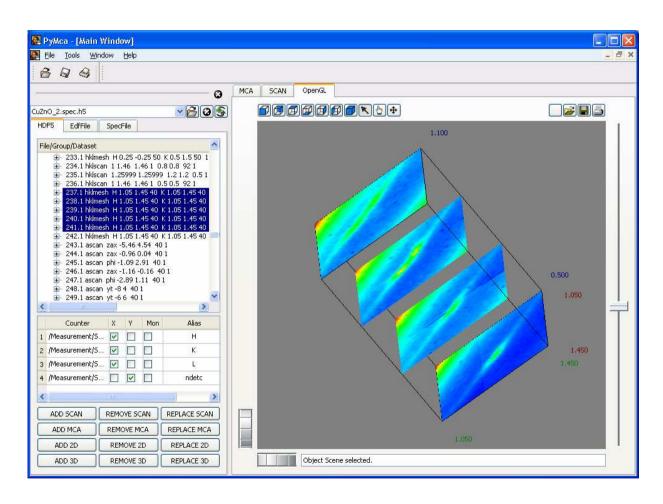
PyMca HDF5/NeXus



SOLEIL NeXus Data courtesy of J.A. Sans and G. Martínez



Generic HDF5 Visualization



Generic: Select what to plot, how to plot it, and plot it.



The PyMca HDF5 problem

PyMca can do a lot more than "just" fully analyze XRF spectra

PyMca can read HDF5 data ... but often has no clue about what to do without asking

Properly defined NXdata groups can provide default visualization and more

It would be great to automatically identify the different MCA detectors, with their counts, their energy axis, their preset time, their elapsed time, their calibration, ... NeXus NXdetector has almost everything needed to describe an MCA, but still needs user interaction to know that detector is an MCA.

It would be very simple to add two attributes to datasets. One defining the "natural" data dimensions and other one defining those dimensions are the first or the last ones of the dataset.



What can two attributes bring?

Programs would automatically provide better user choices

Programs would know what to do with datasets! Despite being able to perform 4D plots, PyMca does not know (it could just guess) what to do to visualize datasets with dimensions $1 \times 2 \times 512 \times 1024$.

```
One attribute to say "ACTUAL_DIMENSION = 2"

One attribute to say "ACTUAL_DIMENSION_IS_LAST_DIMENSION = 1"
```

And things are totally different than:

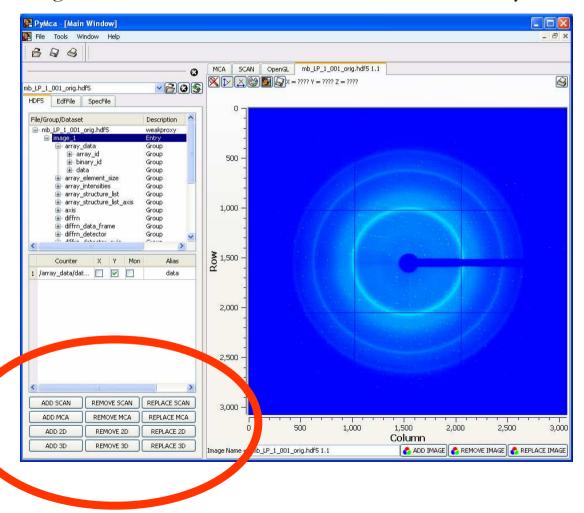
```
One attribute to say "ACTUAL_DIMENSION = 1"

One attribute to say "ACTUAL_DIMENSION_IS_LAST_DIMENSION = 1"
```



The alternative looks ugly, doesn't it?

Just imagine a wall made of buttons instead of bricks everywhere





Conclusion

PyMca is a program as well as a toolkit

- Open source and distributed under the conditions of the GPLv2+
- Supports HDF5 among other formats
- Can be used as a multivariate analysis tool
- Can be used as a fitting and visualization tool (for up to 4-dimensional data)
- Allows you to specify a physically meaningful model which can quantitatively determine element concentrations from energy dispersive X-ray spectra
- Provides high level widgets based on PyQt that can be used independently or integrated into your application
- Has an active development funded by the ESRF



Acknowledgements

My ESRF colleagues (mainly software group and microscopy beamlines ID21 and ID22)

Ph. Walter and coworkers from the Centre de Recherche et de Restauration des Musées de France

Darren Dale (Cornell High Energy Synchrotron Source)

The community of Python developers of free software

The PyMca users, for their enthusiasm and their encouragements