

**NAME**

MX-calibrate – MX-calibrate

**DESCRIPTION**

usage: MX-Calibrate **-w** 1.54 **-c** CeO2 file1.cbf file2.cbf ...

Calibrate automatically a set of frames taken at various sample–detector distance. Return the linear regression of the fit in funtion of the samplesetector distance.

**positional arguments:**

**FILE** List of files to calibrate

**optional arguments:**

**-h, --help**

show this help message and exit

**-V, --version**

show program’s version number and exit

**-v, --verbose**

switch to debug/verbose mode

**-c FILE, --calibrant FILE**

file containing d–spacing of the calibrant reference sample (MANDATORY)

**-w WAVELENGTH, --wavelength WAVELENGTH**

wavelength of the X–Ray beam in Angstrom

**-e ENERGY, --energy ENERGY**

energy of the X–Ray beam in keV ( $hc=12.398419292\text{keV}\cdot\text{A}$ )

**-P POLARIZATION\_FACTOR, --polarization POLARIZATION\_FACTOR**

polarization factor, from **-1** (vertical) to **+1** (horizontal), default is 0, synchrotrons are around 0.95

**-b BACKGROUND, --background BACKGROUND**

Automatic background subtraction if no value are provided

**-d DARK, --dark DARK**

list of dark images to average and subtract

**-f FLAT, --flat FLAT**

list of flat images to average and divide

**-s SPLINE, --spline SPLINE**

spline file describing the detector distortion

**-p PIXEL, --pixel PIXEL**

size of the pixel in micron

**-D DETECTOR\_NAME, --detector DETECTOR\_NAME**

Detector name (instead of pixel size+spline)

**-m MASK, --mask MASK**

file containing the mask (for image reconstruction)

**--filter FILTER**

select the filter, either mean(default), max or median

**--saturation SATURATION**

consider all  $\text{pixel} > \text{max} * (1 - \text{saturation})$  as saturated and reconstruct them

**-r MAX\_RINGS, --ring MAX\_RINGS**

maximum number of rings to extract

**--weighted**

weight fit by intensity

**-l** DISTANCE, **--distance** DISTANCE  
sample-detector distance in millimeter

**--tilt** Allow initially detector tilt to be refined (rot1, rot2, rot3). Default: Activated

**--no-tilt**  
Deactivated tilt refinement and set all rotation to 0

**--dist** DIST  
sample-detector distance in meter

**--poni1** PONI1  
poni1 coordinate in meter

**--poni2** PONI2  
poni2 coordinate in meter

**--rot1** ROT1  
rot1 in radians

**--rot2** ROT2  
rot2 in radians

**--rot3** ROT3  
rot3 in radians

**--fix-dist**  
fix the distance parameter

**--free-dist**  
free the distance parameter

**--fix-poni1**  
fix the poni1 parameter

**--free-poni1**  
free the poni1 parameter

**--fix-poni2**  
fix the poni2 parameter

**--free-poni2**  
free the poni2 parameter

**--fix-rot1**  
fix the rot1 parameter

**--free-rot1**  
free the rot1 parameter

**--fix-rot2**  
fix the rot2 parameter

**--free-rot2**  
free the rot2 parameter

**--fix-rot3**  
fix the rot3 parameter

**--free-rot3**  
free the rot3 parameter

**--fix-wavelength**  
fix the wavelength parameter

**--free-wavelength**  
free the wavelength parameter

- no-gui**  
force the program to run without a Graphical interface
- gui** force the program to run with a Graphical interface
- no-interactive**  
force the program to run and exit without prompting for refinements
- interactive**  
force the program to prompt for refinements
- peak-picker PEAKPICKER**  
Uses the 'massif', 'blob' or 'watershed' peak-picker algorithm (default: blob)

This tool has been developed for ESRF MX-beamlines where an acceptable calibration is usually present in the header of the image. PyFAI reads it and does a "recalib" on each of them before exporting a linear regression of all parameters versus this distance.