

# NanoLib user guide

ETH Zurich  
Laboratory of Solid State Physics  
Microstructure Research

Danilo A. Zanin - [dzanin@phys.ethz.ch](mailto:dzanin@phys.ethz.ch)  
Lorenzo G. De Pietro - [depietro@phys.ethz.ch](mailto:depietro@phys.ethz.ch)  
Quentin Peter - [qpeter@stud.phys.ethz.ch](mailto:qpeter@stud.phys.ethz.ch)

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NanoLib library allows to open and analyze data generated by the [Nanonis SPM Control System™](#) in MATLAB™. The first version was developed by Quentin Peter during its master thesis *Spin Polarized Field Emission STM and Image Processing* in the Solid State Laboratory for [Microstruture Research](#) at the ETH Zurich under the supervision of Dr. U. Ramsperger and L.G. De Pietro.

Some of the features of this library are still oriented to solve problems related to the mater thesis, e.g., *scan\_type* field in the header structure (see [2.1](#)). In next versions some of these feature may be changed and generalized.

NanoLib library is divided in package folders *+folder*. A function in a folder called *+folder* can be called as *folder.function*.

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# 1 SxM File

Images generated via the scanning interface of the Nanonis SPM Control System™ have extension *file.sxm*. Once loaded, variable, from now on *sxmFile*, is a structure divided in: *i*) **header**, a structure containing all information present in the header of the *file.sxm*, and *ii*) **channels**, an array of channel structures containing data and information about every channel. Both, **header** and **channels** can be called by

*sxmFile.header*    and    *sxmfile.channels{#}*

*#* being the number of the channel. When only one channel is loaded one refers to the channel simply as *sxmfile.channels*. More information about the substructure of header and channels is presented below.

## 1.1 *sxmFile*: header and channels tructures

The functions works with a structure that holds every relevant informations. To access the scan data on a structure named *stmFile*, one should type *stmFile.header.rec\_date*. Header and channels structure have following fields:

**header** is a structure composed of:

- scan\_file** name of the file
- rec\_date** date of the scan
- rec\_time** time of the scan
- scan\_pixels** [nx;ny], number of pixels
- scan\_range** [rx;ry], range [m]
- scan\_offset** [ox;oy], offset [m]
- scan\_angle** tilt angle of the scan
- scan\_dir** 'up' or 'down'
- bias** bias voltage [V]
- scan\_type** 'STM', 'SEMPA', 'NFESEM', etc.
- ... Others informations extracted from the file

**channels** is an array of channel structures composed of:

- Direction** 'forward' or 'backward'
- Unit** 'Z' or whatever the unit is

**Name** The name of the channel  
**data** A  $n \times m$  matrix of processed data  
**lineMedian** A  $n \times 1$  matrix of raw line median  
**lineMean** A  $n \times 1$  matrix of raw line mean  
**linePlane** A  $n \times 1$  matrix of raw line mean linear fit  
**lineResidualSlope** a  $1 \times m$  matrix of processed column mean linear fit  
**lineStd** A  $n \times 1$  matrix of processed line standard deviation

## 1.2 +load

This folder contains everything needed to load and process *.sxm* files.

### 1.2.1 loadsxm

**header = loadsxm(fn)** loads a file named *fn.sxm* and returns the Header.  
This function is called by *load.loadProcessedSxM* and **should not be called directly**.

**[header, data] = loadsxm(fn, i)** reads the channel *i* and returns its *data*.

*.sxm* files are composed of an ascii header and of single precision binary data. They are separated by 0x1A 0x04 (SUB EOT). This file is provided by NANO-NIS™ and loads a specified channel from a *.sxm* file.

### 1.2.2 processChannel

**channel = processChannel(channel, header)** Process the *channel* as described below using the informations from *header*. This function is called by *load.loadProcessedXXX* and should not be called directly.

**channel = processChannel(channel, header, corrType)** If *corrType* is set to 'Median', the median is used instead of the mean for lines corrections. If it is set to 'PlaneLineCorrection' a linear fit is used.

The processing orientate and rotate the data so that all the images are comparable. Everything that is removed is saved in the output structure to avoid losing informations.

The mean value of the measurement under the conditions of each pixel must be extracted from the data. As there is drift and other instabilities, the mean value of the data is generally not a good value. The mean of each line is used instead,

as the measurement conditions doesn't change too much during one line. Others possibility include the median or the mean plane. The mean plane along the line is also removed.

For STM, This offset is subtracted. For NFESEM and SEMPA, it is divided, as justified in the thesis.

### 1.2.3 loadProcessedSxM

**file=loadProcessedSxM(fn)** loads and process all the channels of *.sxm* file named *fn*. The structure *file* contains all the informations and is used in a large number of other functions.

**file=loadProcessedSxM(fn, chn)** only loads the channels whose numbers are in the array *chn*

**file=loadProcessedSxM(fn, corrType)** If *corrType* is set to 'MedianCorrection', the median is used instead of the mean for lines corrections. If it set to 'PlaneLineCorrection' a linear fit is used.

The loading is done with *load.loadsxm* and processing with *load.processChannel*.

## 1.3 +plot

This package contains everything needed to plot the data.

### 1.3.1 folder2png

**folder2png(folderName)** finds every *.par* and *.sxm* files in *folderName*, plot all relevant channels and saves the images in a *image* folder.

### 1.3.2 plotData

**[h, range] = plotData(data, name, unit, header)** plots the *data* using informations from the *header*. The figure title is deduced from *name* and *unit*. It returns the plot handle *h* and the chosen range *range*.

**[h, range] = plotData(data, name, unit, header, xoffset, yoffset)** adds an offset to the plot.

The range is 2 STD. If the data is STM, only the lines with low std are considered for the range.

### 1.3.3 plotChannel

**[h, range] = plotChannel(channel, header)** plots the *channel* using informations from the *header*. It returns the plot handle *h* and the chosen range *range*.

**[h, range] = plotChannel(channel, header, xoffset, yoffset)** adds an offset to the plot.

It calls *plot.plotData* on the channel data.

### 1.3.4 plotFile

**[h, range] = plotFile(file, n)** plots the  $n^{th}$  channel of *file*. It returns the plot handle *h* and the chosen range *range*.

**[h, range] = plotFile(file, n, xoffset, yoffset)** adds an offset to the plot.

It calls *plot.plotChannel*.

### 1.3.5 plotHistogram

**plotHistogram(data, range)** plots an histogram of *data* and draw lines on the limit of *range*. It removes the .1% most extreme values.

## 1.4 +op

This package contains various useful functions.

### 1.4.1 combineChannel

**channel=combineChannel(file, name, chn, chw)** combined the channels *chn* of the *file* structure with weights *chw* and return a new *channel* with name *name*.

### 1.4.2 filterData

**[filtered, removed] = filterData(data, pixSize)** filters the *data* with Fourier transform. The filtering keeps structures of approximatively *pixSize* pixels. It returns the filtered data *filtered* and the removed noise *removed*.

**[filtered, removed] = filterData(data, pixSize, 'plotFFT', zoom)** additionally plots the Fourier plane. The optional variable *zoom* has default value 8 and is used to zoom in the Fourier plane.

#### 1.4.3 getChannel

`channelNumber = getChannel(channels,channelNames)` returns all channel numbers where *channels.Name* matches the *channelNames*. *channelNames* can be either a single string or a list of strings.

`channelNumber = getChannel(channels,channelNames,direction)` returns only the channel number where *channels.Direction* matches the *direction*, too.

#### 1.4.4 getColor

`[color,colorScale] = getColor(x,xRange)` computes the ratio between a value *x* and the *xRange* (2 by 1 array). It returns the *color* – and the *colorScale* – according to a predefined color map. *Jet* is the default color map.

`[color,colorScale] = getColor(x,xRange,mapping)` allows to provide a specified *mapping* other than the default, i.e. *jet*. *mapping* should be a color map function, e.g. *hsv*, *parula*, *hot*, *summerm*, *autumn*. Note that, since *mapping* is an argument of functions *getColor*, it must be called by function handle “@” – @.

#### 1.4.5 getOffset

`[offset, XC, centerOffset] = getOffset(img1, header1, img2, header2)` compares the images matrices *img1* and *img2* using informations from the two *headeri* to find the most probable *offset*. The units of *offset* are from *header.scan\_range*. It corresponds to the maximum of the cross correlation matrix *XC*. The corresponding offset relative to the centre of the two images is returned in *centerOffset*.

`[offset, XC, centerOffset] = getOffset(img1, header1, img2, header2, 'mask')` compares masks instead of images.

The offset is from the origin of the image, which is in a corner. The offset of the center is the *centerOffset*, but is less convenient to work with.

#### 1.4.6 getRadialFFT

`[wavelength, radial_spectrum] = getRadialFFT(data)` Computes the *radial spectrum* of the image saved in *data* and the corresponding *wavelength*. The wavelength unit is pixel.

**[wavelength, radial\_spectrum] = getRadialFFT(data, pixPerUnit)** Changes the wavelength unit with the number of pixels per units, *pixPerUnit*.

This function is used to study the radial spectrum of an image computed from the FFT.

#### 1.4.7 getRadialNoise

**[noise\_fit, signal\_start, signal\_error, noise\_coeff] = getRadialNoise(wavelength, radial\_average)** tries to fit a noise from the data of *getRadialFFT*. *noise\_fit* is the detected noise. *signal\_start* is the first position where the signal is detected. *signal\_error* is the error caused by the discrete nature of the signal on *signal\_start*. *noise\_coeff* gives the power law coefficients for the first detected noise.

**[noise\_fit, signal\_start, signal\_error, noise\_coeff] = getRadialNoise(wavelength, radial\_average, maxNbrNoise)** Limits the number of noises to *maxNbrNoise*. The default value is 10.

#### 1.4.8 getRange

**[xrange, yrange] = getRange(header)** extract the ranges *xrange*, *yrange* from *header*.

#### 1.4.9 nanHighStd

**data = nanHighStd(data)** is useful for STM measurements. Usually the lines with very high std don't carry informations, and thus if a line has *std* > *3median*, it is set to nan.

#### 1.4.10 nanonisMap

**colorMap = nanonisMap(nPti)** is a color map function that generates a Nanonis color like mapping of *nPti* number of colors. *nPti* is an optional value. If not provided *nPti* = 64 per default.

#### 1.4.11 interpHighStd

**data = interpHighStd(data)** Removes the lines with high STD values and interpolates the missing values.



#### 1.4.12 interpPeaks

**data = interpPeaks(data)** Removes the data which are too far from the mean and interpolates the missing values.

### 1.5 +mask

These functions are useful to compute threshold mask and apply them.

#### 1.5.1 applyMask

**applyMask(mask)** apply the boolean mask *mask* to the current figure.

**applyMask(mask, color, alpha, xrange, yrange)** apply the boolean mask *mask* in the range *xrange*, *yrange* with color *color* and transparency *alpha*.

The ranges are vectors containing a start point and an end point. See MATLAB's *image* documentation.

#### 1.5.2 getMask

**[maskUp, maskDown, flatData] = getMask(data, pixSize, prctUp, prctDown)** flatten and filter the *data* before computing threshold masks. *flatData* is the flattened and filtered data. *maskUp* marks everything above *prctUp* and *maskDown* below *prctDown*. The filtering is done using *op.filterData*, to which *pixSize* is passed to keep features of this approximate size.

**[maskUp, maskDown, flatData] = getMask(data, pixSize, prctUp, prctDown, 'plotFFT', zoom)** Additionally passes '*plotFFT*', *zoom* to *op.filterData* to visualize the Fourier plane. *zoom* is optional.

The flattening is done using sliding mean.

### 1.6 +convolve2

This is an improved version of MATLAB's *conv2* matrix. It allows a better gestion of boundaries. It was downloaded from [MATLAB file exchange](#). See the license file.

## 2 Dat File

Images generated via the scanning interface of the Nanonis SPM Control System™ have extension *file.sxm*. Once loaded, variable, from now on *sxmFile*, is a structure divided in: *i*) **header**, a structure containing all information present in the header of the *file.sxm*, and *ii*) **channels**, an array of channel structures containing data and information about every channel. Both, **header** and **channels** can be called by

*sxmFile.header*    and    *sxmfile.channels{#}*

*#* being the number of the channel. When only one channel is loaded one refers to the channel simply as *sxmfile.channels*. More information about the substructure of header and channels is presented below.

### 2.1 *sxmFile*: header and channels tructures

The functions works with a structure that holds every relevant informations. To access the scan data on a structure named *stmFile*, one should type *stmFile.header.rec\_date*. Header and channels structure have following fields:

**header** is a structure composed of:

**scan\_file** name of the file  
... Others informations extracted from the file

**experiments** is an array of channel structures composed of:

**data** A  $n \times m$  matrix of processed data

### 2.2 +loadDat

This folder contains everything needed to load and process *.dat* files.

#### 2.2.1 loadsxm

#### 2.2.2 processChannel

#### 2.2.3 loadProcessedSxM

### 2.3 +plotDat

This package contains everything needed to plot the data.

### 2.3.1 plotData

To be written

### 2.3.2 plotChannel

To be written

### 2.3.3 plotFile

To be written

## 2.4 +opDat

### 2.4.1 getChannel

`channelNumber = getChannel(channels,channelNames)` returns all channel numbers where *channels.Name* matches the *channelNames*. *channelNames* can be either a single string or a list of strings.

`channelNumber = getChannel(channels,channelNames,direction)` returns only the channel number where *channels.Direction* matches the *direction*, too.

## 3 Examples

NanoLib library comes with few example showing the basics usage of the library. Some files are also provided in the directory *Files*. Below a list of all examples with a short explanation.

### 3.1 SxM\_Example

`example_open_SxM` shows different ways to load a *file.sxm*.

`example_process_option` shows different ways to process a file while loading a *file.sxm*.

`example_get_drift` detect XY-offset between two different *sxmFiles*.

`example_mask` generates a mask of a *sxmFile* and apply on the original image.

`example_RadialFFT` applies *op.getRadialFFT* and plots some interesting quantities.

### 3.2 dat\_Example

`example_open_Dat` shows different ways to load some *experiment.dat*.