# micropolarray

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**CHAPTER** 

ONE

# MICROPOLARRAY PACKAGE

# 1.1 Subpackages

# 1.1.1 micropolarray.processing package

# 1.1.1.1 Submodules

# 1.1.1.2 micropolarray.processing.chen\_wan\_liang\_calibration module

micropol\_phases\_prev
output\_dir,
occulter=True,
dark\_filename=None

file-

names\_list,

Performs calibration from Chen-Wang-Liang paper 2014

#### **Parameters**

- polarizer\_orientations (list[float]) List of polarizer orienataions
- **filenames\_list** (list[str]) List of filenames coupled with
- micropol\_phases\_previsions (list[float]) Previsions of the micropolarizer orientations inside a superpixel
- **output\_dir** (*str*) output path for the calibration matrices
- occulter (bool, optional) wether to exclude the occulter area. Defaults to True.
- dark\_filename (str, optional) path to the dark to be subtracted from the images. Defaults to None.
- **flat\_filename** (*str*, *optional*) path to the dark to be subtracted from the images. Defaults to None.

#### Raises

• **ValueError** – polarizer orientation list and filenames list do not have the same lenght

• ValueError – any of 0,45,90,-45 polarizations is not included in the polarizer orientation list

micropolarray.processing.chen\_wan\_liang\_calibration.chen\_wan\_liang\_calibration(data, calibration\_matrices\_dir:

str)

Calibrates the images using Chen-Wang-Liang 2014 paper calibration

#### **Parameters**

- data (np. array) data to be calibrated
- **calibration\_matrices\_dir** (*str*) path to the calibration matrices

#### Returns

calibrated data

#### **Return type**

np.array

# 1.1.1.3 micropolarray.processing.congrid module

micropolarray.processing.congrid.congrid(a, newdims, kind='linear')  $\rightarrow$  ndarray Reshapes the data into any new length and width

#### **Parameters**

- a (np.array) data to be reshaped
- **newdims** (tuple | list) new lenght and width
- **kind** (*str*, *optional*) interpolation type. Defaults to "linear".

#### Returns

numpy array of congridded image

### Return type

ndarray

micropolarray.processing.congrid.micropolarray\_jitcongrid(data, width, height, scale)

# 1.1.1.4 micropolarray.processing.convert module

micropolarray.processing.convert.average\_rawfiles\_to\_fits(filenames: list, new\_filename: str, height: int, width: int)

Saves the mean of a list of rawfiles to a new fits file.

#### **Parameters**

- **filenames** (*list*) list of raw filenames
- **new\_filename** (str) new fits filename
- height (int) image height in pix
- width (int) image width in pix

#### Raises

ValueError – trying to save in a file that does not end with .fits

micropolarray.processing.convert.convert\_rawfile\_to\_fits(filename: str, height: int, width: int, remove old: bool = False)

Converts a raw file to a fits one, using default header

#### **Parameters**

- **filename** (str) raw filename
- **height** (*int*) file height
- width (int) file width
- remove\_old (bool, optional) remove old raw file after conversion. Defaults to False.

#### Raises

**ValueError** – raised if the file does not end with ".raw"

micropolarray.processing.convert.convert\_set(filenames, new\_filename, height, width)

ANTARTICOR ONLY: Sums a set of filenames and converts them to one fits file.

#### **Parameters**

- **filenames** (*list*) list of file names to be summed before being converted
- **new\_filename** (str) new .fits file name

micropolarray.processing.convert.nparr\_from\_binary(filename)

Converts a PolarCam binary file into a numpy array. Bytes are saved like this

# • 24 bit (3 bytes)

```
1 | 3 | 2 111111111111 | 1111 | 11111111
```

#### 2 numbers

First number 12bit | Second number (little endian) 8+4=12 bit

# **Parameters**

**filename** (str) – name of the file to be converted

#### Raises

**ValueError** – file lenghts is indivisible by the number of chunks requested to parallelize operations

#### Returns

array of data from file

### Return type

np.array

micropolarray.processing.convert.three\_bytes\_to\_two\_ints(filecontent)

Needed for parallelization, this will be run by each thread for a slice of the original array.

#### Returns

array of saved data

### **Return type**

np.array

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# 1.1.1.5 micropolarray.processing.demodulation module

```
class micropolarray.processing.demodulation.Demodulator(demo_matrices_path: str)
     Bases: object
     Demodulation class needed for MicropolImage demodulation.
     flip(axis)
     rebin(binning)
          DO NOT USE THIS, calculate the tensor from the binned images
     rot90(k=1)
     show(vmin=-1, vmax=1, cmap='Greys') \rightarrow tuple
          Shows the demodulation tensor
```

#### **Parameters**

- vmin (int, optional) Minimum shown value. Defaults to -1.
- vmax (int, optional) Maximum shown value. Defaults to 1.
- cmap (str, optional) Colormap of the plot. Defaults to "Greys".

#### Returns

fig, ax tuple as returned by matplotlib.pyplot.subplots

# **Return type**

tuple

micropolarray.processing.demodulation.Malus(angle, throughput, efficiency, phase)

micropolarray.processing.demodulation.calculate\_demodulation\_tensor(polarizer orientations: list,

filenames\_list: list, micropol\_phases\_previsions: *list*, *gain*: *float*, *output\_dir*: str, binning: int = 1, occulter: list | None = *None, procs grid: list* = [4,*4]*, dark\_filename: str |  $None = None, flat_filename$ :  $str \mid None = None,$ normalizing\_S=None, DEBUG: bool = False)

Calculates the demodulation tensor images and saves them. Requires a set of images with different polarizations to fit a Malus curve model.

- polarizer\_orientations (list[float]) List containing the orientations of the incoming light for each image.
- filenames\_list (list[str]) List of input images filenames to read. Must include [0, 45, 90, -45].
- micropol\_phases\_previsions (list[float]) Previsions for the micropolarizer orientations required to initialize fit.
- **gain** (*float*) Detector [e-/DN], required to compute errors.
- **output\_dir** (*str*) output folder to save matrix to.

- **binning** (*int*, *optional*) Output matrices binning. Defaults to 1 (no binning). Be warned that binning matrices AFTER calculation is an incorrect operation.
- **occulter** (*list*, *optional*) occulter y, x center and radius to exclude from calculations. Defaults to None.
- **procs\_grid** ([int, int], optional) number of processors per side [Y, X], parallelization will be done in a Y x X grid. Defaults to [4,4] (16 procs in a 4x4 grid).
- dark\_filename (str, optional) Dark image filename to correct input images. Defaults to None.
- **flat\_filename** (*str*, *optional*) Flat image filename to correct input images. Defaults to None.
- **normalizing\_S** (*float or np.ndarray, optional*) maximum signal used to normalize single pixel signal. If not set will be estimated as the 4sigma of the signal distribution.

#### Raises

**ValueError** – Raised if any among [0, 45, 90, -45] is not included in the input polarizations.

#### **Notes**

In the binning process the sum of values is considered, which is ok because data is normalized over the maximum S before being fitted.

Utility function to parallelize calculations.

### 1.1.1.6 micropolarray.processing.demosaic module

```
micropolarray.processing.demosaic.demosaic(image_data, option='adjacent')

Returns a [4,n,m] array of polarized images, starting from a micropolarizer image array [n, m].

micropolarray.processing.demosaic.demosaicadjacent(data)

micropolarray.processing.demosaic.demosaicmean(data)

Loops over right polarization pixel location, takes 1/4 of that, stores it in the 2x2 superpixel. demo_images[0] = data[y=0, x=0] demo_images[1] = data[y=0, x=1] demo_images[2] = data[y=1, x=0] demo_images[3] = data[y=1, x=1]

micropolarray.processing.demosaic.merge_polarizations(single_pol_images: ndarray)

micropolarray.processing.demosaic.split_polarizations(data: ndarray)
```

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# 1.1.1.7 micropolarray.processing.hotpix\_cleaning module

micropolarray.processing.hotpix\_cleaning.get\_hot\_pixels(image, threshold=100)

#### 1.1.1.8 micropolarray.processing.nrgf module

micropolarray.processing.nrgf.find\_occulter\_position(data: array, method: str = 'sigmoid', threshold: float = 4.0)

Finds the occulter position of an image.

#### **Parameters**

- data (np.array) input data
- **method** (*str*, *optional*) Method to find occulter edges. If "sigmoid" it will try to fit four sigmoids at the image edges centers, inferring the occulter edges from the parameters. If "algo" it will start from the image edge center and infer the occulter position when DN[i] > threshold\*mean(DN[:i]) Defaults to "sigmoid".
- threshold (float, optional) Threshold for the algo method. Defaults to 4.0.

#### Raises

**UnboundLocalError** – couldn't converge

#### Returns

occulter y, occulter x, occulter radius

#### Return type

list

micropolarray.processing.nrgf.map\_polar\_coordinates(height, width, center)

```
micropolarray.processing.nrgf.nrgf(data: array, y\_center: int | None = None, x\_center: int | None = None, rho\_min: int | None = None, step: int = 1, phi\_to\_mean=[0.0, 360], output\_phi=[0.0, 360]) <math>\rightarrow array
```

Performs nrgf filtering on the image, starting from center and radius. Mean is performed between phi\_to\_mean, 0 is horizontal right, anti-clockwise.

- data (np. array) input array
- **y\_center** (*int*, *optional*) pixel y coordinate of the nrgf center. Defaults to None (image y center).
- **x\_center** (*int*, *optional*) pixel x coordinate of the nrgf center. Defaults to (image x center).
- **rho\_min** (*int*, *optional*) minimun radius in pixels to perform nrgf to. Defaults to None (radius 0).
- **step** (*int*, *optional*) step to which apply the nrgf from center, in pixels. Defaults to 1 pixel.
- phi\_to\_mean (list[float, float], optional) polar angle to calculate the mean value from. Defaults to [0, 360].
- output\_phi (list[float, float], optional) polar angle to include in output data. Defaults to [0, 360].

nrgf-filtered input data

### Return type

np.array

micropolarray.processing.nrgf.reject\_outliers(data, m=2.0)

micropolarray.processing.nrgf.remove\_outliers\_simple(original, neighbours=2)

EXPERIMENTAL DO NOT USE, for improving fitting on occulter position

micropolarray.processing.nrgf.roi\_from\_polar( $data: array, center: list \mid None = None, rho: list \mid None = None, theta=[0, 360], fill: float = 0.0, return boolean=False) <math>\rightarrow$  array

Returns the input array in a circular selection, otherwise an arbitrary number. If a pixel is not in the selection the ENTIRE superpixel is considered out of selection. If return\_boolean is True then a boolean array is returned instead (useful for mean/stdev operations).

#### **Parameters**

- data (np.array) input data
- **center** (*list*, *optional*) pixel coordinates of the circle center. Defaults to None (image center).
- **rho** (list, optional) radius to exclude. Defaults to None (center to image border).
- **theta** (*list*, *optional*) polar selection angle, 0 is horizonta, anti-clockwise direction. Defaults to [0, 360].
- **fill** (*float*, *optional*) number to fill the outer selection. Defaults to 0.0.
- return\_boolean (bool, optional) if set to true, function returns a boolean array of the roi. Defaults to False.

### Returns

array containing the input data inside the selection, and fill otherwise

# Return type

np.array

micropolarray.processing.nrgf.sigmoid(x, max, min, slope, intercept)

micropolarray.processing.nrgf.tile\_double(a)

#### 1.1.1.9 micropolarray.processing.rebin module

micropolarray.processing.rebin.micropolarray\_jitrebin(data, new\_height, new\_width, binning=2)

Fast rebinning function for the micropolarray image. Needs to be wrapped to print info.

 $\verb|micropolarray.processing.rebin.micropolarray_jitrebin_old| (\textit{data}, \textit{height}, \textit{width}, \textit{binning} = 2)$ 

Fast rebinning function for the micropolarray image.

micropolarray.processing.rebin.micropolarray\_rebin(data: ndarray, height: int, width: int, binning=2)
Wrapper for the faster rebinning donw with numba. First deletes last row/column until binning is possible, then calls binning on the result shape.

### **Parameters**

• data (np.ndarray) – data to rebin

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- **height** (*int*) lenght of first axis
- width (int) lenght of second axis
- binning (int, optional) Binning to be performed. Defaults to 2.

binned data, trimmed if necessary

#### Return type

ndarray

micropolarray.processing.rebin.print\_trimming\_info(height, width, new\_height, new\_width)
micropolarray.processing.rebin.standard\_jitrebin(data, height, width, binning=2)

micropolarray.processing.rebin.standard\_rebin(data, binning: int)  $\rightarrow$  array Rebins the data, binned each binningxbinning.

ms the data, binned each binningxbinn

#### **Parameters**

- image (np.array) data to be binned
- **binning** (*int*) binning to be applied. A value of 2 will result in a 2x2 binning (1 pixel is a sum of 4 neighbour pixels)

#### Raises

**KeyError** – cannot divide image height/width by the binning value

#### Returns

binned data

#### Return type

np.array

micropolarray.processing.rebin.trim\_to\_match\_2xbinning(height: int, width: int, binning: int)

Deletes the last image pixels until superpixel binning is compatible with new dimensions

#### **Parameters**

- height (int) image height
- width (int) image width
- binning (int) image binning

#### Returns

image new height and width

#### Return type

int, int

micropolarray.processing.rebin.trim\_to\_match\_binning(height, width, binning, verbose=True)

Deletes the last image pixels until simple binning is compatible with new dimensions

- height (int) image height
- width (int) image width
- binning (int) image binning
- verbose (bool, optional) warns user of trimming. Defaults to True.

image new height and width

# Return type

int, int

# 1.1.1.10 micropolarray.processing.shift module

```
micropolarray.processing.shift.shift(data: ndarray, y: int, x: int)
```

micropolarray.processing.shift.shift\_micropol(data: ndarray, y: int, x: int)

Splits the image into single polarizations, shifts each of them by y,x and then merges them back.

#### **Parameters**

- data (np.ndarray) array to shift
- **y** (*int*) vertical shift (positive inside the image)
- **x** (*int*) horizontal shift (positive inside the image)

#### Returns

shifted array

#### **Return type**

np.ndarray

#### 1.1.1.11 Module contents

# 1.2 Submodules

# 1.3 micropolarray.cameras module

```
class micropolarray.cameras.Antarticor
```

Bases: object

class micropolarray.cameras.Camera

Bases: object

**occulter\_mask**( $overoccult: int = 0, rmax: int | None = None) <math>\rightarrow$  array

Returns an array of True inside the roi, False elsewhere. Useful for mean/std operations (where=occulter\_mask).

### **Parameters**

- **overoccult** (*int*, *optional*) Pixels to overoccult. Defaults to 15.
- rmax (int, optional) Maximum r of the ROI. Defaults to image nearest border.

#### Returns

Boolean roi array

#### Return type

np.array

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```
occulter\_roi(data: array, fill: float = 0.0, overoccult: int = 0) \rightarrow array
```

Returns the array in the polar ROI, else fill

#### **Parameters**

- data (np.array) Input array
- **fill** (*float*, *optional*) Value for filling. Defaults to 0.0.
- overoccult (int, optional) Pixels to overoccult. Defaults to 0.

#### Returns

Array if in ROI, fill elsewhere

# **Return type**

np.array

class micropolarray.cameras.Kasi

Bases: Camera

class micropolarray.cameras.PolarCam

Bases: Camera

# 1.4 micropolarray.image module

class micropolarray.image.Image(initializer: str | np.ndarray | Image, averageimages: bool = True)

Bases: object

Basic image class. Can be initialized from a filename, a filenames list, a numpy array or another Image instance. If multiple filenames are provided, will perform the mean of them unless averageimages is False.

# property data: ndarray

save\_as\_fits(filename: str, fixto: str[float, float] = None)

Saves image as fits with current header

#### **Parameters**

- **filename** (str) output filename
- **fixto** (str[float, float], optional) Set a maximum and minimum range for the data. Defaults to None.

#### Raises

**ValueError** – filename does not end with ".fits"

save\_as\_raw(filename: str)

Saves the image as a raw binary file

### **Parameters**

**filename** (str) – output filename

#### Raises

ValueError - filename does not end with ".raw"

**show**( $cmap='Greys\_r'$ , vmin=None, vmax=None)  $\rightarrow$  tuple

Shows the image data

#### **Parameters**

• cmap (str, optional) - figure colorbar. Defaults to "Greys\_r".

```
• vmin (_type_, optional) – Minimum value to plot. Defaults to None.
```

• vmax (\_type\_, optional) - Maximum value to plot. Defaults to None.

#### Returns

fig, ax tuple as returned by matplotlib.pyplot.subplots

### Return type

tuple

**show\_histogram**(bins: int = 1000)  $\rightarrow$  tuple

Print the histogram of the flattened image data

#### **Parameters**

**bins** (int, optional) – Numbers of bin to compute the histogram from. Defaults to 1000.

#### Returns

fig, ax tuple as returned by matplotlib.pyplot.subplots

# Return type

tuple

# 1.5 micropolarray.micropol\_image module

Bases: Image

Micro-polarizer array image class. Can be initialized from a 2d array, a list of 1 or more file names (use the boolean keyword averageimages to select if sum or average is taken) or another MicropolImage. Dark and flat micropolarray images can also be provided to automatically correct the result.

 $congrid(newdim\_y: int, newdim\_x: int) \rightarrow MicropolImage$ 

Return type

*MicropolImage* 

```
correct_flat(flat: MicropolImage) → MicropolImage
```

Normalizes the flat and uses it to correStokes\_vecct the image.

#### **Parameters**

**flat** (MicropolImage) – flat image, does not need to be normalized.

#### Returns

copy of input image corrected by flat

#### **Return type**

MicropolImage

#### **correct\_ifov()** → *MicropolImage*

Corrects differences in single pixels fields of view inside each superpixel

#### Returns

image with data corrected for field of view differences

#### Return type

MicropolImage

**demodulate**(demodulator: Demodulator, demosaicing: bool = False)  $\rightarrow MicropolImage$ 

Returns a MicropolImage with polarization parameters calculated from the demodulation tensor provided.

#### **Parameters**

- **demodulator** (Demodulator) Demodulator object containing the demodulation tensor components (see processing.new\_demodulation)
- **demosaicing** (*bool*, *optional*) wether to apply demosaicing to the image or not. Set it to False if demodulation matrices have half the dimension of the image. Defaults to True.

#### Raises

**ValueError** – raised if image and demodulator do not have the same dimension, for example in case of different binning

#### Returns

copy of the input imagreturn e with I, Q, U, pB, DoLP, AoLP calculated from the demodulation tensor.

#### Return type

MicropolImage

 $demosaic(demosaic\_mode='adjacent') \rightarrow MicropolImage$ 

Returns a demosaiced copy of the image with updated polarization parameters. Demoisacing is done IN PLACE.

#### **Parameters**

 ${\tt demosaic\_mode}\ (str,\ optional) - {\tt demosaicing}\ mode\ (see\ processing.demosaic).$  Defaults to "adjacent".

#### Returns

demosaiced image

### **Return type**

MicropolImage

```
first call = True
```

```
mask_occulter(y: int = 919, x: int = 950, r: int = 531, overoccult: int = 0) \rightarrow None
```

Masks occulter for all image parameters

- y (int, optional) Occulter y position. Defaults to PolarCam().occulter\_pos\_last[0].
- **x** (int, optional) Occulter x position. Defaults to PolarCam().occulter\_pos\_last[1].
- **r** (int, optional) Occulter radius. Defaults to PolarCam().occulter\_pos\_last[2].
- **overoccult** (*int*, *optional*) Pixels to overoccult. Defaults to 0.
- camera (\_type\_, optional) Camera image type. Defaults to PolarCam().

None

property pB: PolParam

property pol0: PolParam

property pol45: PolParam

property pol90: PolParam

property pol\_45: PolParam

property polparam\_list: list

**rebin**(binning: int)  $\rightarrow MicropolImage$ 

Rebins the micropolarizer array image, binned each binningxbinning. Sum bins by default.

#### **Parameters**

**binning** (int) – binning to perform. A value of n will be translated in a nxn binning.

#### **Raises**

**ValueError** – negative binning provided

#### Returns

copy of the input image, rebinned.

#### Return type

MicropolImage

 $rotate(angle: float) \rightarrow MicropolImage$ 

Rotates an image of angle degrees, counter-clockwise.

```
save\_all\_pol\_params\_as\_fits(filename: str) \rightarrow None
```

Saves the image and all polarization parameters as fits file with the same name

#### **Parameters**

**filename** (str) – filename of the output image. Will be saved as filename\_[I, Q, U, pB, AoLP, DoLP].fits

### Raises

**ValueError** – filename is not a valid .fits file

 $\textbf{save\_demosaiced\_images\_as\_fits}(\textit{filename: str, fixto: list[float, float]} = \textit{None}) \rightarrow \textit{None}$ 

Saves the four demosaiced images as fits files

- **filename** (*str*) filename of the output image. The four images will be saved as filename\_POLXX.fits
- **fixto** (list[float, float], optional) set the minimum and maximum value for the output images. Defaults to None.

#### Raises

**ValueError** – an invalid file name is provided

 $save\_param\_as\_fits(polparam: str, filename: str, fixto: list[float, float] = None) \rightarrow None$ 

Saves chosen polarization parameter as a fits file

#### **Parameters**

- **polparam** (*str*) polarization parameter to save. Can be one among [I, Q, U, pB, AoLP, DoLP]
- **filename** (str) filename of the output image.
- **fixto** (list[float, float], optional) set the minimum and maximum value for the output images. Defaults to None.

#### Raises

**ValueError** – filename is not a valid .fits file

**save\_single\_pol\_images**( $filename: str, fixto: list[float, float] = None) <math>\rightarrow$  None

Saves the four polarized images as fits files

#### **Parameters**

- **filename** (*str*) filename of the output image. The four images will be saved as filename\_POLXX.fits
- **fixto** (list[float, float], optional) set the minimum and maximum value for the output images. Defaults to None.

#### Raises

**ValueError** – an invalid file name is provided

```
shift(y: int, x: int) \rightarrow MicropolImage
```

Shifts image by y, x pixels and fills with 0 the remaining space. Positive numbers for up/right shift and negative for down/left shift. Image is split into polarizations, each one is shifted, then they are merged again.

#### **Parameters**

- **y** (*int*) vertical shift in pix
- **x** (*int*) horizontal shift in pix

#### Returns

shifted image copied from the original

#### **Return type**

MicropolImage

show\_demo\_images(cmap='Greys\_r', vmin=None, vmax=None, \*\*kwargs)

Plots the four demosaiced images.

#### **Parameters**

- **cmap** (str, optional) colormap for the plot. Defaults to "Greys\_r".
- \*\*kwargs arguments passed to matplotlib.pyplot.imshow.

# Returns

a (figure, axis) couple same as matplotlib.pyplot.subplots

#### Return type

tuple

```
show_pol_param(polparam: str, cmap='Greys_r', vmin=None, vmax=None, **kwargs)
```

Plots a single polarization parameter given as input

#### **Parameters**

- **polparam** (*str*) image PolParam containing the parameter to plot. Can be one among [I, Q, U, pB, AoLP, DoLP]
- cmap (str, optional) colormap for the plot. Defaults to "Greys\_r".
- **\*\*kwargs** arguments passed to matplotlib.pyplot.imshow.

#### Returns

a (figure, axis) couple same as matplotlib.pyplot.subplots

### Return type

tuple

### show\_single\_pol\_images(cmap='Greys\_r', \*\*kwargs)

Plots the four polarizations images.

#### **Parameters**

- **cmap** (str, optional) colormap for the plot. Defaults to "Greys\_r".
- **\*\*kwargs** arguments passed to matplotlib.pyplot.imshow.

#### Returns

a (figure, axis) couple same as matplotlib.pyplot.subplots

### Return type

tuple

# **show\_with\_pol\_params**( $cmap='Greys\_r'$ ) $\rightarrow$ tuple

Returns a tuple containing figure and axis of the plotted data, and figure and axis of polarization parameters (3x2 subplots). User must callplt.show after this is called.

#### **Parameters**

```
cmap (str, optional) – colormap string. Defaults to "Greys_r".
```

#### Returns

a (figure, axis, figure, axis) couple same as matplotlib.pyplot.subplots for the image data and another for the six polarization parameters

# Return type

tuple

#### property single\_pol\_subimages

#### **subtract\_dark**(*dark*: MicropolImage) → *MicropolImage*

Correctly subtracts the input dark image from the image

#### **Parameters**

dark (MicropolImage) - dark to subtract

#### Returns

copy of input image with dark subtracted

#### **Return type**

MicropolImage

```
class micropolarray.micropol_image.PolParam(ID: str, data: ndarray, title: str, measure_unit: str, fix data: bool = False)
```

Bases: object

Auxiliary class for polarization parameters.

#### Members:

ID (str): parameter identifier data (np.array): parameter image as numpy 2D array title (str): brief title of the parameter, useful for plotting measure\_unit (str): initial measure units of the parameter fix\_data (bool): controls whether data has to be constrained to [0, 4096] interval (not implemented yet)

ID: str

data: ndarray

fix\_data: bool = False

measure\_unit: str

title: str

micropolarray.micropol\_image.set\_default\_angles(angles\_dic: dict)

Sets the default micropolarizer orientations for images.

#### **Parameters**

**angles\_dic** (dict) – dictionary {value : pos} where value is the angle in degrees from -90 to 90 and pos is the pixel position in superpixel, from 0 to 3 (position [y, x], fast index x)

# 1.6 micropolarray.polarization\_functions module

```
micropolarray.polarization_functions.AoLP(Stokes_vec_components)
Angle of linear polarization in [rad]
micropolarray.polarization_functions.DoLP(Stokes_vec_components)
Degree of linear polarization in [%]
micropolarray.polarization_functions.normalize2pi(angles_list)
Normalizes the input angle list in the -90,90 range

Parameters
angles_list(_type_) - list of input angles in degrees

Returns
normalized angles

Return type
__type_

micropolarray.polarization_functions.pB(Stokes_vec_components)
Polarized brighness in [%]
```

# 1.7 micropolarray.utils module

micropolarray.utils.align\_keywords\_and\_data(header, data, sun\_center, platescale, binning=1)

Fixes antarticor keywords and data to reflect each other.

#### **Parameters**

- header (dict) fits file header
- data (ndarray) data as np array
- platescale (float) plate scale in arcsec/pixel
- **binning** (*int*, *optional*) binning applied to image. Defaults to 1 (no binning).

#### Returns

new fixed header and data

#### Return type

header, data

micropolarray.utils.fix\_data(data: array, min, max)

micropolarray.utils.get\_Bsun\_units( $diffuser_I$ : float,  $texp\_image$ : float = 1.0,  $texp\_diffuser$ : float = 1.0)  $\rightarrow$  float

Returns the conversion unit for expressing brightness in units of sun brightness. Usage is data [units of B\_sun] = data[DN] \* get\_Bsun\_units(mean\_Bsun\_brightness, texp\_image, texp\_diffuser)

#### **Parameters**

- mean\_sun\_brightness (float) diffuser mean in DN.
- **texp\_image** (*float*, *optional*) image exposure time. Defaults to 1.0.
- texp\_diffuser (float, optional) diffuser exposure time. Defaults to 1.0.

#### Returns

Bsun units conversion factor

# Return type

float

micropolarray.utils.get\_malus\_normalization(four\_peaks\_images, show\_hist=False)

micropolarray.utils.mean\_minus\_std( $data: array, stds_n: int = 1$ )  $\rightarrow$  float

Returns the value at the mean - standard deviation for the input data

#### **Parameters**

- data (np.array) input data
- **stds\_n** (*int*, *optional*) number of standard deviations. Defaults to 1.

#### Returns

mean value - n\*stdevs

#### Return type

float

micropolarray.utils.mean\_plus\_std( $data: array, stds_n: int = 1$ )  $\rightarrow$  float

Returns the value at the mean + standard deviation for the input data

- data (np.array) input data
- **stds\_n** (*int*, *optional*) number of standard deviations. Defaults to 1.

mean value + n\*stdevs

#### **Return type**

float

micropolarray.utils.median\_minus\_std( $data: array, stds_n: int = 1$ )  $\rightarrow$  float

Returns the value at the median - median deviation for the input data

#### **Parameters**

- data (np.array) input data
- **stds\_n** (*int*, *optional*) number of standard deviations. Defaults to 1.

#### **Returns**

median value - n\*mediandevs

# Return type

float

micropolarray.utils.median\_plus\_std( $data: array, stds_n: int = 1$ )  $\rightarrow$  float

Returns the value at the median + median deviation for the input data

#### **Parameters**

- data (np.array) input data
- **stds\_n** (*int*, *optional*) number of standard deviations. Defaults to 1.

### Returns

median value + n\*mediandevs

# Return type

float

micropolarray.utils.normalize2pi(angles\_list)

micropolarray.utils.sigma\_DN(pix\_DN)

micropolarray.utils.timer(func)

Use this to time function execution

#### **Parameters**

**func** (function) – function of which to measure execution time

# 1.8 Module contents

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