# 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\_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.merge\_rawfiles\_to\_fits(filenames: list, new\_filename: str, height: int, width: int, mode='sum')

Saves the average or sum 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
- mode (str) wether to "average" or "sum" the images. Defaults to "sum".

### Raises

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

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

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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
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.
     property Cij: ndarray
     property angle_dic: dict
          Dictionary representing the correlation between pix family and fitted angle
               Returns
                  key[value] where key is the angle and value is the pixel family index (y, x) with fast index x
               Return type
                  dict
     property eff: ndarray
     flip(axis)
     property phi: ndarray
     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
     property tk: ndarray
```

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

 $\label{lem:micropolarray.processing.demodulation.calculate\_demodulation\_tensor(\textit{polarizer\_orientations: list}, \\$ 

filenames\_list: list, micropol\_phases\_previsions:
list, gain: float, output\_dir:
str, binning: int = 1,
occulter: list = None,
procs\_grid: list = [4, 4],
dark\_filename: str = None,
flat\_filename: str = None,
normalizing\_S=None,
tk\_boundary: list = None,
eff\_boundary: list = 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.

### **Parameters**

- **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.
- **tk\_boundary** (*list*) if provided, sets the transmittancy [initial guess, boundary\_inf, boundary\_sup] of the Malus curve (max value). Defaults to [0.5, 0.1, 1.-1.e-6].
- **eff\_boundary** (*list*) if provided, sets the efficiency [initial guess, boundary\_inf, boundary\_sup] of the Malus curve (max value). Defaults to [0.5, 0.1, 1.-1.e-6].

### Raises

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

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### **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.

```
micropolarray.processing.demodulation.compute_demodulation_by_chunk(splitted_normalized_dara_arr, splitted_pixel_erorrs, splitted_occulter_flag, splitted_occulter_flag, polarizer_orientations, rad_micropol_phases_previsions, tk_boundary, eff_boundary, DEBUG)
```

Utility function to parallelize calculations.

# 1.1.1.6 micropolarray.processing.demodulation\_errors module

Bases: object

```
micropolarray.processing.demodulation_errors.get_error_on_AoLP(S: ndarray, sigma\_S: ndarray) \rightarrow ndarray
```

 $\label{eq:micropolarray.processing.demodulation\_errors. \textbf{get\_error\_on\_DoLP}(\textit{S: ndarray}, \textit{sigma\_S: ndarray}) \rightarrow \\ \text{ndarray}$ 

```
micropolarray.processing.demodulation_errors.get\_error\_on\_Stokes(image\_error: ndarray, demodulator: Demodulator) <math>\rightarrow ndarray
```

Returns the error on the image, propagated through the demodulation matrix. If M[i, j] is the demodulation matrix, sigma\_I[k] are the four pixel values in a superpixel, and S[i, j] is the Stokes vector, returns the matrix product  $sqrt(M^2 @ I^2)$ 

### **Parameters**

- **image\_error** (*np.ndarray*) array containing the pixel by pixel error to propagate.
- **demodulator** (Demodulator) demodulator containing the demodulation matrix.

# Returns

errors of the computed Stokes vector as a [3, y, x] array.

# Return type

np.ndarray

 $\label{eq:micropolarray.processing.demodulation\_errors. \textbf{get\_error\_on\_pB}(S: ndarray, sigma\_S: ndarray) \rightarrow \\ \text{ndarray}$ 

# 1.1.1.7 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)
```

# 1.1.1.8 micropolarray.processing.image\_cleaning module

```
micropolarray.processing.image_cleaning.auto_threshold(data: ndarray) \rightarrow float
```

Get the threshold following Otsu's algorithm. This assumes that there are two populations (noise + signal) and minimizes the intra- class variance

#### **Parameters**

data (np. ndarray) – array on which to perform the treshold

### Returns

Otsu's threshold

### Return type

float

```
micropolarray.processing.image_cleaning.get_hot_pixels(image, threshold=100)
micropolarray.processing.image_cleaning.reject_outliers(data, m=2.0)
micropolarray.processing.image_cleaning.remove_outliers_simple(original, neighbours=2)
EXPERIMENTAL DO NOT USE, for improving fitting on occulter position
```

### 1.1.1.9 micropolarray.processing.linear roi module

```
micropolarray.processing.linear_roi.DDA(start: list, end: list) \rightarrow ndarray
```

Digital\_differential\_analyzer algorithm for line rasterizing. Unlike bresenham, works in every quadrant. NOTE: even if the distance between start and end coordinates is the same, a different number of points is selected depending on the line slope, so the ratio between distance and number of points is also returned.

### **Parameters**

- start (list) starting point coordinates
- end (list) ending point coordinates

# Returns

interpolated points locations float: ratio between the distance from start to end point and the number of returned locations

### Return type

np.ndarray

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micropolarray.processing.linear\_roi.bresenham(start: list, end: list) → ndarray

Bresenham algorithm for generating integers on a line. Efficient BUT works ONLY in the first octant

### **Parameters**

- **start** (*list*) starting point coordinates
- end (list) ending point coordinates

### **Returns**

coordinates of the points under the line from start to end

### **Return type**

np.ndarray

micropolarray.processing.linear\_roi.linear\_roi(data: ndarray, start: list, end: list) → ndarray
Get values

### **Parameters**

- data (np.ndarray) \_description\_
- start (list) \_description\_
- end (list) \_description\_

### Returns

\_description\_

# Return type

np.ndarray

micropolarray.processing.linear\_roi\_linear\_roi\_from\_polar( $data: ndarray, center: list, theta: float, r: list = None) <math>\rightarrow$  list

Returns a linear roi starting from the center and extending to r or to the edge of the input data array. Angles start horizontally and rotate anti-clockwise (0deg corresponds to fixed y and increasing x).

### **Parameters**

- data (np.ndarray) input array from which to select a roi
- center (list) center coordinates
- theta (float) angle of the linear roi
- **r** (list, optional) Maximum radius for the roi. Defaults to
- None.

### Returns

1-dimensional array containing the selected values from data np.ndarray: roi indexes along the first (y) dimension of data np.ndarray: roi indexes along the second (x) dimension of data float: ratio between the pixel length and the length of the returned roi (see linear\_roi.DDA)

# **Return type**

np.ndarray

# 1.1.1.10 micropolarray.processing.nrgf module

micropolarray.processing.nrgf.find\_occulter\_hough( $data: array, minr: float = 1, maxr: float = None, **kwargs) <math>\rightarrow$  tuple

Uses Hough Gradient from cv2 and computes the coronagraph occulter coordinates. Returns Y, X, Radius of the occulter.

### **Parameters**

**data** (*np.array*) – input data

#### Returns

occulter y, x, r

### **Return type**

tuple

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, x_center: int = None, rho_min: int = None, step: int = 1, phi_to_mean=[0.0, 360], output_phi=[0.0, 360]) \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.

### **Parameters**

- 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).

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- **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].

### Returns

nrgf-filtered input data

### Return type

np.array

```
micropolarray.processing.nrgf.roi_from_polar(data: array, center: list = None, rho: list = None, theta=[0, 360], fill: float = 0.0, return_boolean: bool = False, include_superpixels: bool = True) <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.
- include\_superpixels (bool, optional) if set to true then exclude entire superpixel if one pixel is in the occulter

# 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)

# 1.1.1.11 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.

micropolarray.processing.rebin.micropolarray\_jitrebin\_old(data, height, width, binning=2)

Fast rebinning function for the micropolarray image.

micropolarray.processing.rebin.micropolarray\_rebin(data: ndarray, 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
- **height** (*int*) lenght of first axis
- width (int) lenght of second axis
- binning (int, optional) Binning to be performed. Defaults to 2.

### Returns

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, new\_height, new\_width, binning=2)

micropolarray.processing.rebin.standard\_rebin(data, binning: int)  $\rightarrow$  array

Rebins the data, binned each binningxbinning.

### **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

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# 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

### **Parameters**

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

### **Returns**

image new height and width

### Return type

int, int

# 1.1.1.12 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)
- $\boldsymbol{x}$  (int) horizontal shift (positive inside the image)

# Returns

shifted array

# Return type

np.ndarray

# 1.1.1.13 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) \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

**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 | 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"

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

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.

### **Parameters**

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

### Returns

shifted image copied from the original

# **Return type**

*Image* 

**show**( $cmap='Greys\_r'$ , vmin=None, vmax=None, \*\*kwargs)  $\rightarrow$  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(split\_pols=False, **kwargs) \rightarrow tuple
```

Print the histogram of the flattened image data

# **Parameters**

\*\*kwargs (int, optional) – arguments to pass to numpy.histogram(), like bins and range.

### **Returns**

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

# Return type

tuple

# 1.5 micropolarray.micropol\_image module

```
class micropolarray.micropol_image.MicropolImage(initializer: str | ndarray | list | MicropolImage,
                                                           angle dic: dict = None, dark: MicropolImage =
                                                           None, flat: MicropolImage = None, averageimages:
                                                           bool = True)
     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.
     property AoLP: PolParam
     property DoLP: PolParam
     property I: PolParam
     property Q: PolParam
     property U: PolParam
     clean_hot_pixels(flagged_hot_pix_map: MicropolImage)
           Returns a copy of the image with gaussian smeared pixels where flagged_hot_pix_map == 1.
               Parameters
                   flagged_hot_pix_map (MicropolImage) – hot pixels map.
                   copy of the original image, gaussian smeared where flagged_hot_pix_map == 1
               Return type
                   MicropolImage
     congrid(newdim\_y: int, newdim\_x: int) \rightarrow MicropolImage
           Reshapes a MicropolImage into any new length and width. This is done separately for each pixel family.
               Parameters
                   • newdim_y (int) – new height
                   • newdim_x (int) – new width
               Returns
                   image with reshaped data.
               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.
                   copy of input image corrected by flat
```

MicropolImage

Return type

# $correct_ifov() \rightarrow \textit{MicropolImage}$

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

 ${\it MicropolImage}$ 

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

Returns a demosaiced copy of the image with updated polarization parameters. Demoisacing is done IN PLACE and using the THEORETICAL MATRIX. If demodulation and demosaicing are required, please use demodulate(demosaic=True)

### **Parameters**

**demosaic\_mode** (*str*, *optional*) – demosaicing mode (see processing.demosaic). Defaults to "adjacent".

# Returns

demosaiced image

# **Return type**

MicropolImage

first\_call = True

 $mask_occulter(y: int = 919, x: int = 941, r: int = 536, overoccult: int = 0) \rightarrow None$ 

Masks occulter for all image parameters

### **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().

### Returns

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

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

Saves the four demosaiced 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

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

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

 $\mathbf{shift}(y: int, x: int) \rightarrow MicropolImage$ 

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\_histogram(split\_pols: bool = True, \*\*kwargs)

Print the histogram of the flattened image data

# **Parameters**

• **split\_pols** (*bool*, *optional*) – Whether to overplot histograms of same family pixels separately. Defaults to False.

• \*\*kwargs (int, optional) — arguments to pass to numpy.histogram(), like bins and range.

### Returns

fig, ax tuple as returned by 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_pol_params(cmap='Greys_r', figsize=None, **kwargs) \rightarrow tuple
```

Returns a tuple containing figure and axis of polarization parameters (3x2 subplots). User must call plt.show after this is called.

### **Parameters**

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

### Returns

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

### 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\_

 $\verb|micropolarray.polarization_functions.pB| (Stokes\_vec\_components)|$ 

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

### **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.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)

Returns the list of angles (in degrees) normalized between -90 and 90 degrees.

### **Parameters**

angles\_list (list) - list of angles to normalize

### Returns

list of normalized angles

# Return type

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

1.8. Module contents

# CHAPTER

# TWO

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