



stistools Documentation

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This package provides data processing tools for working with STIS data.

Contents:

BASIC2D

```
stistools.basic2d.basic2d(input, output='', outblev='', dqicorr=True, blevcorr=True, doppcorr=True, lorscorr=True, glincorr=True, lflgcorr=True, biascorr=True, darkcorr=True, flatcorr=True, photcorr=True, statflag=True, darkscale='', verbose=False, timestamps=False, trailer='', print_version=False, print_revision=False)
```

Perform basic 2-D calibration of STIS raw data.

Some calibration steps are relevant only for CCD or only for MAMA, and since an output file of calstis or basic2d may be used as the input, some steps may have already been done. Most calibration steps will not be done if they are not relevant or if they have already been done, regardless of the value of the calibration switch (e.g. flatcorr).

Parameters**input: str :**

Name of the input raw file.

output: str :

Name of the output file, or "" (the default). If no name was specified, the output name will be constructed from the input name.

outblev: str :

Name of the output text file for blev info, or "" (the default).

dqicorr: bool :

If True, update the DQ array.

blevcorr: bool :

If True, subtract a bias level based on the overscan values. (CCD only.)

doppcorr: bool :

If True, convolve reference files (bpixtab, darkfile, flatfile) as needed with the Doppler shift offset throughout the exposure, if Doppler correction was done on-board. (MAMA only, because for the CCD Doppler correction is not done on-board.)

lorscorr: bool :

If True, bin high-res data to lo-res. (MAMA only.)

glincorr: bool :

If True, correct for global non-linearity. (MAMA only.)

lflgcorr: bool :

If True, flag local non-linearity. (MAMA only.)

biascorr: bool :

If True, subtract the bias image. (CCD only.)

darkcorr: bool :

If True, subtract the dark image, scaled by the exposure time and possibly also a temperature-dependent factor.

flatcorr: bool :

If True, divide by the flat field image.

photcorr: bool :

If True, determine the photometric parameters and populate keywords PHOTFLAM, PHOTZPT, PHOTPLAM and PHOTBW. (Imaging data only.)

statflag: bool :

If True, compute statistics for image arrays and update keywords.

darkscale: str :

This may be used to override the time and/or temperature dependent scale factor that would normally be applied to the dark image before subtracting from the raw data. It's a string rather than a float in order to accept a different scale factor for each image set in the input data. calstis reads the value or values (separated by blanks) from the string, and if the value is greater than zero, it will be used instead of the value determined from the temperature and time. (CCD or NUV-MAMA only.)

verbose: bool :

If True, calstis will print more info.

timestamps: bool :

If True, calstis will print the date and time at various points during processing.

trailer: str :

If specified, the standard output and standard error will be written to this file instead of to the terminal. Note, however, that if `print_version` or `print_revision` is specified, the value will be printed to the terminal, and any name given for the trailer will be ignored.

print_version: bool :

If True, calstis will print the version number (a string) and then return 0.

print_revision: bool :

If True, calstis will print the full version string and then return 0.

Returns

status: int :

0 is OK. 1 is returned if `cs1.e` (the calstis host executable) returned a non-zero status. If `verbose` is True, the value returned by `cs1.e` will be printed. 2 is returned if the specified input file or files were not found, or if there is a mismatch between the number of input, output, and/or outblev files specified.

`stistools.basic2d.getHelpAsString` (*fulldoc=True*)

Return documentation on the `basic2d` function.

`stistools.basic2d.main` (*args*)

stistools.basic2d.**prtOptions**()

Print a list of command-line options and arguments.

stistools.basic2d.**run**(*configobj=None*)

TEAL interface for the basic2d function.

CALSTIS

`stistools.calstis.calstis` (*input*, *wavecal*='', *outroot*='', *savetmp*=False, *verbose*=False, *timesteps*=False, *trailer*='', *print_version*=False, *print_revision*=False)

Calibrate STIS data.

Parameters

input: str :

Name of the input file.

wavecal: str :

Name of the input wavecal file, or "" (the default). This is only needed if the name is not the "normal" name (`rootname_wav.fits`).

outroot: str :

Root name for the output files, or "" (the default). This can be a directory name, in which case the string must end in '/

savetmp: bool :

True if `calstis` should not delete temporary files.

verbose: bool :

If True, `calstis` will print more info.

timesteps: bool :

If True, `calstis` will print the date and time at various points during processing.

trailer: str :

If specified, the standard output and standard error will be written to this file instead of to the terminal. Note, however, that if `print_version` or `print_revision` is specified, the value will be printed to the terminal, and any name given for the trailer will be ignored.

print_version: bool :

If True, `calstis` will print the version number (a string) and then return 0.

print_revision: bool :

If True, `calstis` will print the full version string and then return 0.

Returns

status: int :

0 is OK. 1 is returned if `cs0.e` (the `calstis` host executable) returned a non-zero status. If `verbose` is True, the value returned by `cs0.e` will be printed. 2 is returned if the specified input file or files were not found.

`stistools.calstis.getHelpAsString (fulldoc=True)`

Return documentation on the calstis function.

`stistools.calstis.main (args)`

`stistools.calstis.prtOptions ()`

Print a list of command-line options and arguments.

`stistools.calstis.run (configobj=None)`

TEAL interface for the calstis function.

SSHIFT

A Python module for aligning the spectra in different flat-fielded images of an IMSET. These files can then be combined with along-the-slit dithering to reject hot pixels and cosmic rays. The POSTARG2 keyword is used to determine the number of rows to be shifted.

`stistools.sshift.shift_image` (*infile, outfile, shift=0*)

Shift each image extension of an input file by N rows and write the new image extension to the output file.

`stistools.sshift.sshift` (*input, output=None, shifts=None, platescale=None, tolerance=None*)

Align spectra from different images of an imset.

Parameters

input : list

A list of input filenames. These must be STIS flat- fielded (`_flt`) image FITS files. This argument will accept a single filename or a list of filenames.

shifts : list, optional

A list of integers indicating the number of rows to shift each image of each file in the cross-dispersion (Y-) direction.

platescale : float, optional

The size of a pixel in arcseconds. Used to convert the value of the POSTARG2 keyword to pixels.

tolerance : float, optional

The allowed difference between calculated shifts and integer pixel shifts (fraction of pixel).

Returns

output : list, optional

A list of output filenames. The number of output filenames must match the number of input filenames. If no output is given, then the `_flt` substring of the input file is replaced by the `_sfl` substring to create an output file. This option will accept a single filename or a list of filenames.

Notes

Author:

•Paul Barrett (STScI)

STISNOISE

`stistools.stisnoise.gauss` (*x, x0, dx, ymax*)

`stistools.stisnoise.medianfilter` (*time_series, width*)

`stistools.stisnoise.stisnoise` (*infile, exten=1, outfile=None, dc=1, verbose=1, boxcar=0, wipe=None, window=None*)

Computes an FFT on STIS CCD frames to evaluate fixed pattern noise.

Fixed pattern noise is most obvious in a FFT of bias frames. Optional filtering to correct the fixed pattern noise is provided through keywords `boxcar`, `wipe`, and `window`. Filtered data can be saved as an output file.

Parameters

infile : string

STIS FITS file

exten : int, optional

fits extension to be read

dc : int, optional

the power in the first freq bin is set to zero for better plotting of the power spectrum.

verbose : int, optional [Default: 1]

set to 0 if you do not want brief information about each image.

boxcar : int

width of boxcar smoothing to be applied.

wipe : ndarray

a 3-element array, specifying how to modify the data in frequency space. If set, the image is converted to a 1-D time series, fourier transformed to frequency space, modified, inverse transformed back to time space, and converted back to a 2-D image. The first and second elements specify the range in frequencies to be scaled (in hz), and the third element specifies the scaling factor (should be 0-1).

window : ndarray

a 3 element array, specifying how to modify the data in frequency space. The first element is the center of the window (in hz). The second element is the width of the window (in hz). The third element controls the tapering of the window - it is the scale (in hz) of the tapering width. Specifically, a square bandstop is convolved with a gaussian having the FWHM given by the third parameter.

outfile : string,optional
name of filtered image file

Returns

noise_terms : tuple of arrays

A tuple containing the arrays; namely, the arrays:

```
freq = frequency in power spectrum (hz)
magn = magnitude in power spectrum
```

Notes

Authors:

- Original algorithm: Thomas M. Brown (STScI)
- Python version: Paul Barrett (STScI)

`stistools.stisnoise.windowfilter` (*time_series, image_type, sst, freqpeak, width, taper*)

`stistools.stisnoise.wipefilter` (*time_series, image_type, sst, freqmin, freqmax, scale*)

MKTRACE

Refine a STIS trace table.

- A trace is generated from the science file and a trace center is computed.
- The two traces bracketing the trace center are extracted from the trace table and interpolated
- The correction is computed as the difference between the linear fit to the science and interpolated traces
- The correction is applied to all traces in the trace file for that particular OPT_ELEM and CENWAVE
- A new trace table is written to the current directory and the relevant keywords are updates in the header of the input file.

Usage

Simple example of running mktrace on a STIS file named 'file.fits':

```
>>>import mktrace >>>mktrace.mktrace('file.fits', [tracecen=509.4], [weights=[(x1,x2),(x3,x4)])
```

Authors

- Author (IDL): Linda Dressel
- Python version: Nadia Dencheva

class `stistools.mktrace.Trace` (*file*, *kwinfo*)
Trace class for a crj or fit file.

Notes

`tr=Trace(file)` file is a crj or fit file.

`opt_elem`, `cenwave`, `sporder` are read from the header of the science file `a2center` is a2center of the trace generated from the science file

`tr_ind= tr.getTraceInd(a2center)`

`tr_ind` is the index of the row in the trace file which brackets from below `a2center` as computed fro the generated trace

`tr.readTrace(tr_ind)`

`a2center = tr.generateTrace(...)`

gFitTrace (*specimage*, *y1*, *y2*)

Fit a gaussian to each column of an image.

generateTrace (*data*, *kwinfo*, *tracecen=0.0*, *wind=None*)

Generates a trace from a science file.

getTraceInd (*a2center*)

Finds the first trace in the trace table whose A2CENTER is larger than the specified a2center

openTraceFile (*filename*)

Returns a spectrum trace table

readTrace (*tr_ind*)

reads the specified row from the 1dttab.fits

writeTrace (*fname, sciline, refline, interp_trace, trace1024, tr_ind, a2disp_ind*)

The 'writeTrace' method performs the following steps:

- Adds sciline-refline to all traces with the relevent OPT_ELEM, CENWAVE and SPORDER.
- Writes the new trace table to the current directory.
- Updates the SPTRCTAB keyword in the header to point to the new table.
- Writes out fits files with the
 - science trace - '_sci'
 - the fit to the science trace - '_scifit'
 - the interpolated trace - '_interp'
 - the linear fit to the interpolated trace - '_interpfit'

stistools.mktrace.**getKWInfo** (*hdr0, hdr1*)

stistools.mktrace.**interp** (*y, n*)

Given a 1D array of size m, interpolates it to a size n (m<n).

stistools.mktrace.**iterable** (*v*)

stistools.mktrace.**mktrace** (*fname, tracecen=0.0, weights=None*)

Refine a stis spectroscopic trace.

stistools.mktrace.**trace_interp** (*tr1, tr2, cen*)

EVALDISP

`stistools.evaldisp.evalDisp` (*coeff*, *wl*)

Return the pixel corresponding to wavelength *wl*.

Parameters

coeff : array_like object

a list of eight elements containing the dispersion coefficients as read from a STIS `_dsp.fits` table

wl : float or ndarray

a single wavelength or an array (numarray) of wavelengths, in Angstroms

Returns

pix_number : float or ndarray

the pixel number (or array of pixel numbers) corresponding to the input wavelength(s); note that these are zero indexed

Notes

The expression in the calstis code is:

```
x = coeff[0] +  
    coeff[1] * m * wl +  
    coeff[2] * m**2 * wl**2 +  
    coeff[3] * m +  
    coeff[4] * wl +  
    coeff[5] * m**2 * wl +  
    coeff[6] * m * wl**2 +  
    coeff[7] * m**3 * wl**3
```

This version of the function to evaluate the dispersion relation assumes that the grating is first order, i.e. $m = 1$. The dispersion coefficients give one-indexed pixel coordinates (reference pixels), but this function converts to zero-indexed pixels.

`stistools.evaldisp.newton` (*x*, *coeff*, *cenwave*, *niter=4*)

Return the wavelength corresponding to pixel *x*.

The dispersion solution is evaluated iteratively, and the slope (dispersion) for Newton's method is determined numerically, using a difference in wavelength of one Angstrom. Note that the `evalDisp` in this file assumes that the grating is first order.

Parameters

x : float or ndarray

a single pixel number or an array of pixel numbers

coeff : array_like object

a list of eight elements containing the dispersion coefficients as read from a STIS
_dsp.fits table

cenwave : int or float

central wavelength, in Angstroms

niter : int

number of iterations

Returns

wavelength : float or ndarray

a single wavelength or an array (numarray) of wavelengths, in Angstroms

WAVELEN

`stistools.wavelen.adjust_disp`(*ncoeff*, *coeff*, *delta_offset1*, *shifta1*, *inang_info*, *delta_tan*,
delta_row, *binaxis1*)

Adjust the dispersion coefficients.

The changes to the coefficients are for the incidence angle correction, the offset from the SHIFTA1 keyword, and the tilt of the slit. The coefficients will be modified in-place.

Parameters

ncoeff : int

number of dispersion coefficients

coeff : ndarray of float64

array of dispersion coefficients, modified in-place

delta_offset1 : float

incidence angle offset in degrees

shifta1 : float

MSM offset (ref. pixels) in the dispersion direction

delta_tan : float

difference in tangents of slit angle and ref angle

delta_row : float

difference between current row number and CRPIX2

binaxis1 : float

binning factor in dispersion direction

inang_info : rec_array

rows from the incidence-angle table

`stistools.wavelen.compute_wavelengths`(*shape*, *phdr*, *hdr*, *helcorr*)

Compute a 2-D array of wavelengths, one value for each image pixel.

Parameters

shape : tuple of two ints

the number of rows and columns in the output image

phdr : pyfits Header object

primary header

hdr : pyfits Header object

extension header

helcorr : string

“PERFORM” if heliocentric correction should be done

Returns

wavelengths : ndarray of float64

an array of wavelengths, of the same shape (nrows, ncols) as the output image

`stistools.wavelen.get_delta_offset1` (*apdestab*, *aperture*, *ref_aper*)

Get the incidence angle offset.

Parameters

apdestab : string

name of the aperture description table

aperture : string

aperture (slit) name

ref_aper : string

name of the reference aperture, the one that was used to calculate the dispersion relation

Returns

angle : float

incidence angle offset in degrees

```
stistools.wx2d.apply_trace(image, a2center, a2displ, subdiv, offset=0.0, shiffta2=0.0,  
                           extname='SCI')
```

Add together 'subdiv' rows of 'image', following the trace.

Parameters

image : ndarray

input 2-D image array, oversampled by 'subdiv' in axis 0

a2center : ndarray

1-D array of Y locations

a2displ : ndarray

array of traces, one for each a2center; the length of each trace must be the same as the number of columns in the input image

subdiv : int

number of rows to add together

offset : float

offset of the first row in 'image' from the beginning of the data block in the original file, needed for trace

shiffta2 : float

offset of the row from nominal (from shiffta2 keyword)

extname : string

which type of extension (SCI, ERR, DQ)?

Returns

x2d : ndarray

resampled 2-D image array

Notes

The function value is a 2-D array containing the resampled image. This is binned by subdiv in Y (axis 0), after shifting by trace (multiplied by subdiv).

For extname = "ERR" the result differs in these ways:

- 1.fractions of pixels at the endpoints of the extraction region are not included
- 2.the values are combined as the average of the sum of the squares

For extname = "DQ" the result differs in these ways:

- 1.the output is type int16
- 2.the output values are nominally the same as the input, while for SCI the output are subdiv times larger than the input
- 3.fractions of pixels at the endpoints of the extraction region are not included
- 4.the values are combined via bitwise OR rather than an average or sum

`stistools.wx2d.bin_traces` (*a2displ*, *binaxis1*, *ltv*)

bin the traces by the factor *binaxis1*

Parameters

a2displ : ndarray

an array of one or more arrays of Y displacements (traces)

binaxis1 : int

binning factor in the dispersion axis

ltv : float

offset in the dispersion axis (one indexing)

Returns

a2displ : ndarray

an array of traces (*a2displ*), but with the trace arrays binned and shorter by the factor *binaxis1*

`stistools.wx2d.extract` (*image*, *locn*, *subdiv*)

Add together 'subdiv' rows of 'image', centered on 'locn'.

Parameters

image : ndarray

input array, oversampled by 'subdiv' in axis 0

locn : ndarray

a 1-D array giving the location at which to extract; an integer value corresponds to the center of the pixel. The length must be the same as the number of columns in the input image.

subdiv : int

number of rows to add together

Returns

spec : ndarray

a 1-D array containing the extracted row

`stistools.wx2d.extract_err` (*image*, *locn*, *subdiv*)

Average 'subdiv' rows of 'image', centered on 'locn'.

Parameters

image : ndarray

input array, oversampled by 'subdiv' in axis 0

locn : ndarray

a 1-D array giving the location at which to extract; an integer value corresponds to the center of the pixel

subdiv : int

number of rows to add together

Returns

spec : ndarray

a 1-D array containing the extracted row

Notes

This takes the square root of the average of the squares, intended to be used for interpolating the ERR array. Fractions of pixels at the upper and lower edges are excluded.

`stistools.wx2d.extract_i16` (*image*, *locn*, *subdiv*)

Bitwise OR 'subdiv' rows of 'image', centered on 'locn'.

Parameters

image : ndarray

input array, oversampled by 'subdiv' in axis 0

locn : ndarray

a 1-D array giving the location at which to extract; an integer value corresponds to the center of the pixel

subdiv : int

number of rows to add together

Returns

spec : ndarray

a 1-D array containing the extracted row

`stistools.wx2d.get_trace` (*tracefile*, *phdr*, *hdr*)

Read 1-D traces from the 1dt table (sptctab).

Parameters

tracefile : string or array

either a trace array or the name of a FITS 1dt table

phdr : pyfits Header object

primary header of input file

hdr : pyfits Header object

extension header of input image (for binning info and time of exposure)

Returns

trace_arrays : tuple of 2 arrays

a pair of arrays, one is the Y location at the middle column, and the other is an array of trace arrays

Notes

If 'tracefile' is already a trace array, it will just be returned, together with an arbitrary Y location of 0 (because that will always be within the image).

`opt_elem` and `cenwave` are criteria for selecting the relevant rows from the 1dt table. There will normally be several rows that match, and they should have different values of the Y location; the output list will be sorted on Y location.

`stistools.wx2d.interpolate_trace` (*a2center, a2displ, y, length*)

Interpolate within the array of traces, and return a trace.

Parameters

a2center : ndarray

array of Y locations

a2displ : ndarray

array of traces, one trace for each element of a2center

y : float

Y location on the detector

length : int

length of a trace; needed only if traces is empty

`stistools.wx2d.inv_avg_interp` (*order, image*)

`stistools.wx2d.inv_haar` (*image*)

`stistools.wx2d.kd_apply_trace` (*image, a2center, a2displ, offset=0.0, shiffta2=0.0*)

Kris Davidson's resampling algorithm, following the trace.

Parameters

image : ndarray

input 2-D image array

a2center : ndarray

array of Y locations

a2displ : ndarray

array of traces, one for each a2center; the length of each trace must be the same as the number of columns in 'image'

offset : float

offset of the first row in 'image' from the beginning of the data block in the original file, needed for trace

shiffta2 : float

offset of the row from nominal (from shiffta2 keyword)

Returns

x2d : ndarray

2-D array containing the resampled image

`stistools.wx2d.kd_resampling` (*img, errimg, original_nrows, nrows, ncols, rows, a2center, a2displ, offset, shiffta2*)

Apply Kris Davidson's resampling method.

Parameters

img : ndarray

SCI image array (could be a subset of full image)

errimg : ndarray

ERR image array (could be a subset of full image)

original_nrows : int

number of image lines (NAXIS2) in input image

nrows : int

number of image lines in subset

ncols : int

number of image columns (NAXIS1)

rows : tuple

tuple giving the slice of rows to process

a2center : ndarray

1-D array of Y locations

a2displ : ndarray

array of traces, one for each a2center; the length of each trace must be the same as the number of columns in the input image

offset : float

offset of the first row in 'image' from the beginning of the data block in the original file, needed for trace

shiffta2 : float

offset of the row from nominal (from shiffta2 keyword)

Returns

img_arr : tuple

the image and error arrays (to replace the input img and errimg)

`stistools.wx2d.polynomial(x, y, z, n)`

used for interpolation

Parameters

x : ndarray

the integer values from 0 through n-1 inclusive (but float64)

y : ndarray

a 2-D array, axis 0 of length n

z : float

n / 2.

n : int

1 + order of polynomial fit

`stistools.wx2d.stis_psf(x, a)`

Evaluate the cross-dispersion PSF at x.

Parameters

x : float

offset in pixels from the center of the profile

a : float

a measure of the width of the PSF

Returns

val : float

the PSF evaluated at x

`stistools.wx2d.trace_name` (*trace, phdr*)

Return the 1dt table name or array.

Parameters

trace : string or array or None

if trace is None the header keyword SPTRCTAB will be gotten from phdr; else if this is a string it should be the name of a trace file (possibly using an environment variable); otherwise, it should be a trace, in which case it will be returned unchanged

phdr : pyfits Header object

primary header, used only if trace is None

Returns

tracefile : string or array

name of a trace file (with environment variable expanded), or an actual trace array

`stistools.wx2d.wavelet_resampling` (*hdu, img, errimg, original_nrows, nrows, ncols, rows, a2center, a2displ, offset, shifta2, imset, order, subdiv, psf_width, subsampled, convolved*)

Resample img and errimg using wavelets.

Parameters

hdu : pyfits header/data unit object

header/data unit for a SCI extension

img : ndarray

SCI image array (could be a subset of full image)

errimg : ndarray

ERR image array (could be a subset of full image)

original_nrows : int

number of image lines (NAXIS2) in input image

nrows : int

number of image lines in subset

ncols : int

number of image columns (NAXIS1)

rows : tuple

tuple giving the slice of rows to process

a2center : ndarray

1-D array of Y locations

a2displ : ndarray

array of traces, one for each a2center; the length of each trace must be the same as the number of columns in the input image

offset : float

offset of the first row in ‘image’ from the beginning of the data block in the original file, needed for trace

shiffta2 : float

offset of the row from nominal (from shiffta2 keyword)

imset : int

number of the current image set (keyword EXTVER)

order : int

polynomial order

subdiv : int

number of subpixels per input pixel

psf_width : float

width of PSF for convolution (e.g. 1.3);

subsamped : string or None

name of the output file with the subsampled image

convolved : string or None

name of the output file with the convolved image

Returns

img_arr: tuple of ndarrays :

the image and error arrays (to replace the input img and errimg)

`stistools.wx2d.wx2d(input, output, wavelengths=None, helcorr='', algorithm='wavelet', trace=None, order=7, subdiv=8, psf_width=0.0, rows=None, subsamped=None, convolved=None)`

Resample the input, correcting for geometric distortion.

Parameters

input : string

name of input file containing an image set

output : string

name of the output file

wavelengths : string, optional [Default: None]

name of the output file for wavelengths

helcorr : string

specify “perform” or “omit” to override header keyword

algorithm : { ‘wavelet’, ‘kd’ }

algorithm to use in resampling the input

trace : string or array, or None

trace array, or name of FITS table containing trace(s)

order : int [Default: 7]

polynomial order (an odd number, e.g. 5 or 7)

subdiv : int [Default: 8]

number of subpixels (a power of 2, e.g. 8 or 16)

psf_width : float [Default: 0.]

width of PSF for convolution (e.g. 1.3); 0 means no convolution

rows : tuple, optional [Default: None]

a tuple giving the slice of rows to process; output values in all other rows will be set to zero. The default of None means all rows, same as (0, 1024)

subsamped : string, optional [Default: None]

name of the output file with the subsampled image

convolved : string, optional [Default: None]

name of the output file with the convolved image

`stistools.wx2d.wx2d_imset` (*ft, imset, output, wavelengths, helcorr, algorithm, tracefile, order, subdiv, psf_width, rows, subsampled, convolved*)

Resample one image set, and append to output file(s).

Parameters

ft : HDUList

pyfits HDUList object for the input file

imset : int

one-indexed image set number

output : string

name of the output file

wavelengths : string or None

name of the output file for wavelengths

helcorr : { 'perform', 'omit' }

specify "perform" or "omit" to override header keyword

algorithm : { "wavelet", "kd" }

algorithm to use to process input

tracefile : string or array

trace array, or name of FITS table containing trace(s)

order : int

polynomial order

subdiv : int

number of subpixels

psf_width : float

width of PSF for convolution

rows : tuple

a tuple giving the slice of rows to process

subsamped : string, or None

name of the output file with the subsampled image

convolved : string, or None

name of the output file with the convolved image

RADIALVEL

`stistools.radialvel.earthVel` (*mjd*)

Compute and return the velocity of the Earth at the specified time.

This function computes the Earth's orbital velocity around the Sun in celestial rectangular coordinates. The expressions are from the *Astronomical Almanac*, p C24, which gives low precision formulas for the Sun's coordinates. We'll apply these formulas directly to get the velocity of the Sun relative to the Earth, then we'll convert to km per sec and change the sign to get the velocity of the Earth.

Parameters

mjd : float

time, Modified Julian Date

Returns

vel : ndarray

the velocity vector of the Earth around the Sun, in celestial coordinates
(shape=(3,),ndtype=float64)

Notes

We get the velocity of the Sun relative to the Earth as follows:

The velocity in the ecliptic plane with the X-axis aligned with the radius vector is:

- $V_x = \text{radius_dot}$,
- $V_y = \text{radius} * \text{elong_dot}$,
- $V_z = 0$

where:

- **radius** is the radial distance from Earth to Sun
- **elong** is the ecliptic longitude of the Sun
- **eps** is the obliquity of the ecliptic
- **_dot** means the time derivative

Rotate in the XY-plane by **elong** to get the velocity in ecliptic coordinates:

```
radius_dot * cos (elong) - radius * elong_dot * sin (elong)
radius_dot * sin (elong) + radius * elong_dot * cos (elong)
0
```

Rotate in the YZ-plane by **eps** to get the velocity in equatorial coordinates:

```
radius_dot * cos (elong) - radius * elong_dot * sin (elong)
(radius_dot * sin (elong) + radius * elong_dot * cos (elong)) * cos (eps)
(radius_dot * sin (elong) + radius * elong_dot * cos (elong)) * sin (eps)
```

`stistools.radialvel.precess` (*mjd*, *target*)

Precess target coordinates from J2000 to the date *mjd*.

Parameters

mjd : float

time, Modified Julian Date

target : array_like object

unit vector pointing toward the target, J2000 coordinates

Returns

vector : ndarray

the target vector (or matrix) precessed to *mjd* as an array object of type float64 and the same shape as *target*, i.e. either (3,) or (n,3)

Notes

target can be a single vector, e.g. [x0, y0, z0], or it can be a 2-D array; in the latter case, the shape should be (n,3):

```
target = [[x0, x1, x2, x3, x4],
          [y0, y1, y2, y3, y4],
          [z0, z1, z2, z3, z4]]
```

The algorithm used in this function was based on [R1] and [R2].

References

[R1], [R2]

`stistools.radialvel.radialVel` (*ra_targ*, *dec_targ*, *mjd*)

Compute the heliocentric velocity of the Earth.

This function computes the radial velocity of a target based on the Earth's orbital velocity around the Sun. The space motion of the target is not taken into account. That is, the radial velocity is just the negative of the component of the Earth's orbital velocity in the direction toward the target.

Parameters

ra_targ : float

right ascension of the target (degrees)

dec_targ : float

declination of the target (degrees)

mjd : float

Modified Julian Date at the time of observation

Returns

radial_vel : float

the radial velocity in km/s

`stistools.r_util.expandFileName` (*filename*)

Expand environment variable in a file name.

If the input file name begins with either a Unix-style or IRAF-style environment variable (e.g. `$lref/name_dqi.fits` or `lref$name_dqi.fits` respectively), this routine expands the variable and returns a complete path name for the file.

Parameters

filename : string

a file name, possibly including an environment variable

Returns

fullname : string

the file name with environment variable expanded

`stistools.r_util.interpolate` (*x, values, xp*)

Interpolate.

Linear interpolation is used. If the specified independent variable value *xp* is outside the range of the array *x*, the first (or last) value in *values* will be returned.

Parameters

x : a sequence object, e.g. an array, int or float

array of independent variable values

values : a sequence object, e.g. an array (not character)

array of dependent variable values

xp : int or float

independent variable value at which to interpolate

Returns

interp_vals : the same type as one element of *values*

linearly interpolated value

GETTABLE

`stistools.gettable.getTable` (*table*, *filter*, *sortcol=None*, *exactly_one=False*, *at_least_one=False*)

Return row(s) of a table that match the filter.

Rows that match every item in the filter (a dictionary of column_name=value) will be returned. If the value in the table is `STRING_WILDCARD` or `INT_WILDCARD` (depending on the data type of the column), that value is considered to match the filter for that column. Also, for a given filter key, if the corresponding value in the filter is `STRING_WILDCARD`, the test on filter will be skipped for that key (i.e. a wildcard filter element matches any row).

If more than one row matches the filter, there is an option to sort these rows based on the values of one of the table columns.

It is an error if `exactly_one` or `at_least_one` is `True` but no row matches the filter. A warning will be printed if `exactly_one` is `True` but more than one row matches the filter.

Parameters

table : string

name of the reference table

filter : dict

each key is a column name, and the corresponding value is a possible table value in that column

sortcol : string

the name of a column on which to sort the table rows (if there is more than one matching row), or `None` to disable sorting

exactly_one : bool

set this to `True` if there must be one and only one matching row

at_least_one : bool

set this to `True` if there must be at least one matching row

Returns

match_rows : rec_array

an array of the rows of the table that match the filter; note that if only one row matches the filter, the function value will still be an array

`stistools.gettable.rotateTrace` (*trace_info*, *expstart*)

Rotate `a2displ`, if `MJD` and `DEGPERYR` are in the trace table.

Parameters

trace_info : rec_array

an array of the relevant rows of the table; the A2DISPL column will be modified in-place if the MJD and DEGPERYR columns are present

expstart : float

exposure start time (MJD)

`stistools.gettable.sortrows` (*rowdata*, *sortcol*, *ascend=True*)

Return a copy of rowdata, sorted on sortcol.

INDICES AND TABLES

- *genindex*
- *modindex*
- *search*

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- [R1] Lieske, et al. 1976, *Astron & Astrophys* vol 58, p 1.
- [R2] J.H. Lieske, 1979, *Astron & Astrophys* vol 73, 282-284.

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