
stingray Documentation

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Stingray Developers

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Part I

Stingray API

Library of Time Series Methods For Astronomical X-ray Data.

class stingray.AveragedPowerspectrum(*lc*, *segment_size*, *norm*='rms')

Make an averaged periodogram from a light curve by segmenting the light curve, Fourier-transforming each segment and then averaging the resulting periodograms.

Parameters

lc: lightcurve.Lightcurve object OR

iterable of lightcurve.Lightcurve objects The light curve data to be Fourier-transformed.

segment_size: float

The size of each segment to average. Note that if the total duration of each Lightcurve object in *lc* is not an integer multiple of the *segment_size*, then any fraction left-over at the end of the time series will be lost.

norm: {"leahy" | "rms"}, optional, default "rms"

The normalization of the periodogram to be used. Options are "leahy" or "rms", default is "rms".

Attributes

norm: {"leahy" "rms"}	the normalization of the periodogram
freq: numpy.ndarray	The array of mid-bin frequencies that the Fourier transform samples
ps: numpy.ndarray	The array of normalized squared absolute values of Fourier amplitudes
df: float	The frequency resolution
m: int	The number of averaged periodograms
n: int	The number of data points in the light curve
nphots: float	The total number of photons in the light curve

class stingray.Lightcurve(*time*, *counts*, *input_counts*=True)

Make a light curve object from an array of time stamps and an array of counts.

Parameters

time: iterable

A list or array of time stamps for a light curve

counts: iterable, optional, default None

A list or array of the counts in each bin corresponding to the bins defined in *time* (note: **not** the count rate, i.e. counts/second, but the counts/bin).

input_counts: bool, optional, default True

If True, the code assumes that the input data in 'counts' is in units of counts/bin. If False, it assumes the data in 'counts' is in counts/second.

Attributes

time: numpy.ndarray	The array of midpoints of time bins
counts: numpy.ndarray	The counts per bin corresponding to the bins in <i>time</i> .
countrate: numpy.ndarray	The counts per second in each of the bins defined in <i>time</i> .
ncounts: int	The number of data points in the light curve.
dt: float	The time resolution of the light curve.
tseg: float	The total duration of the light curve.
tstart: float	The start time of the light curve.

static `make_lightcurve(toa, dt, tseg=None, tstart=None)`

Make a light curve out of photon arrival times.

Parameters

toa: iterable

list of photon arrival times

dt: float

time resolution of the light curve (the bin width)

tseg: float, optional, default None

The total duration of the light curve. If this is `None`, then the total duration of the light curve will be the interval between the arrival between the first and the last photon in `toa`.

Note: If `tseg` is not divisible by `dt` (i.e. if `tseg/dt` is not an integer number), then the last fractional bin will be dropped!

tstart: float, optional, default None

The start time of the light curve. If this is `None`, the arrival time of the first photon will be used as the start time of the light curve.

Returns

lc: `Lightcurve` object

A light curve object with the binned light curve

rebin_lightcurve(dt_new, method='sum')

Rebin the light curve to a new time resolution. While the new resolution need not be an integer multiple of the previous time resolution, be aware that if it is not, the last bin will be cut off by the fraction left over by the integer division.

Parameters

dt_new: float

The new time resolution of the light curve. Must be larger than the time resolution of the old light curve!

method: {"sum" | "mean" | "average"}, optional, default "sum"

This keyword argument sets whether the counts in the new bins should be summed or averaged.

Returns

lc_new: `Lightcurve` object

The `Lightcurve` object with the new, binned light curve.

class `stingray.Powerspectrum(lc=None, norm='rms')`

Make a Periodogram (power spectrum) from a (binned) light curve. Periodograms can be Leahy normalized or fractional rms normalized. You can also make an empty Periodogram object to populate with your own fourier-transformed data (this can sometimes be useful when making binned periodograms).

Parameters

lc: lightcurve.Lightcurve object, optional, default None

The light curve data to be Fourier-transformed.

norm: {"leahy" | "rms"}, optional, default "rms"

The normaliation of the periodogram to be used. Options are “leahy” or “rms”, default is “rms”.

Attributes

norm: {“leahy” “rms”}	the normalization of the periodogram
freq: numpy.ndarray	The array of mid-bin frequencies that the Fourier transform samples
ps: numpy.ndarray	The array of normalized squared absolute values of Fourier amplitudes
df: float	The frequency resolution
m: int	The number of averaged powers in each bin
n: int	The number of data points in the light curve
nphotos: float	The total number of photons in the light curve

classical_significances(*threshold=1, trial_correction=False*)

Compute the classical significances for the powers in the power spectrum, assuming an underlying noise distribution that follows a chi-square distributions with 2M degrees of freedom, where M is the number of powers averaged in each bin.

Note that this function will *only* produce correct results when the following underlying assumptions are fulfilled: (1) The power spectrum is Leahy-normalized (2) There is no source of variability in the data other than the periodic signal to be determined with this method. This is important! If there are other sources of (aperiodic) variability in the data, this method will *not* produce correct results, but instead produce a large number of spurious false positive detections! (3) There are no significant instrumental effects changing the statistical distribution of the powers (e.g. pile-up or dead time)

By default, the method produces (index,p-values) for all powers in the power spectrum, where index is the numerical index of the power in question. If a threshold is set, then only powers with p-values *below* that threshold with their respective indices. If trial_correction is set to True, then the threshold will be corrected for the number of trials (frequencies) in the power spectrum before being used.

Parameters

threshold : float

The threshold to be used when reporting p-values of potentially significant powers. Must be between 0 and 1. Default is 1 (all p-values will be reported).

trial_correction : bool

A Boolean flag that sets whether the threshold will be correted by the number of frequencies before being applied. This decreases the threshold (p-values need to be lower to count as significant). Default is False (report all powers) though for any application where threshold is set to something meaningful, this should also be applied!

Returns

pvals : iterable

A list of (index, p-value) tuples for all powers that have p-values lower than the threshold specified in threshold.

compute_rms(*min_freq, max_freq*)

Compute the fractional rms amplitude in the periodgram between two frequencies.

Parameters

min_freq: float

The lower frequency bound for the calculation

max_freq: float

The upper frequency bound for the calculation

Returns

rms: float

The fractional rms amplitude contained between min_freq and max_freq

rebin(df, method='mean')

Rebin the periodogram to a new frequency resolution df.

Parameters

df: float

The new frequency resolution

Returns

bin_ps = Periodogram object

The newly binned periodogram

rebin_log(f=0.01)

Logarithmic rebin of the periodogram. The new frequency depends on the previous frequency modified by a factor f:

$$\text{dnu}_j = \text{dnu}_{j-1} * (1+f)$$

Parameters

f: float, optional, default 0.01

parameter that steers the frequency resolution

Returns

binfreq: numpy.ndarray

the binned frequencies

bins: numpy.ndarray

the binned powers

nsamples: numpy.ndarray

the samples of the original periodogram included in each frequency bin

stingray.is_iterable(stuff)

Test if stuff is an iterable.

stingray.is_string(s)

Portable function to answer this question.

stingray.jit(fun)

stingray.optimal_bin_time(ffilen, tbin)

Vary slightly the bin time to have a power of two number of bins.

Given an FFT length and a proposed bin time, return a bin time slightly shorter than the original, that will produce a power-of-two number of FFT bins.

stingray.rebin_data(x, y, dx_new, method='sum')

Rebin some data to an arbitrary new data resolution. Either sum the data points in the new bins or average them.

Parameters

x: iterable

The dependent variable with some resolution $\text{dx_old} = x[1] - x[0]$

y: iterable

The independent variable to be binned

dx_new: float

The new resolution of the dependent variable x

method: {"sum" | "average" | "mean"}, optional, default "sum"

The method to be used in binning. Either sum the samples y in each new bin of x, or take the arithmetic mean.

Returns

xbin: numpy.ndarray

The midpoints of the new bins in x

ybin: numpy.ndarray

The binned quantity y

stingray.**simon**(message, **kwargs)

The Statistical Interpretation MONitor.

A warning system designed to always remind the user that Simon is watching him/her.

Parameters

message : string

The message that is thrown

kwargs : dict

The rest of the arguments that are passed to warnings.warn

stingray.**test**(package=None, test_path=None, args=None, plugins=None, verbose=False, pastebin=None, remote_data=False, pep8=False, pdb=False, coverage=False, open_files=False, **kwargs)
Run the tests using [py.test](#). A proper set of arguments is constructed and passed to [pytest.main](#).

Parameters

package : str, optional

The name of a specific package to test, e.g. 'io.fits' or 'utils'. If nothing is specified all default tests are run.

test_path : str, optional

Specify location to test by path. May be a single file or directory. Must be specified absolutely or relative to the calling directory.

args : str, optional

Additional arguments to be passed to [pytest.main](#) in the args keyword argument.

plugins : list, optional

Plugins to be passed to [pytest.main](#) in the plugins keyword argument.

verbose : bool, optional

Convenience option to turn on verbose output from [py.test](#). Passing True is the same as specifying '-v' in args.

pastebin : {'failed', 'all', None}, optional

Convenience option for turning on [py.test](#) pastebin output. Set to 'failed' to upload info for failed tests, or 'all' to upload info for all tests.

remote_data : bool, optional

Controls whether to run tests marked with `@remote_data`. These tests use online data and are not run by default. Set to `True` to run these tests.

pep8 : bool, optional

Turn on PEP8 checking via the [pytest-pep8 plugin](#) and disable normal tests. Same as specifying `'--pep8 -k pep8'` in args.

pdb : bool, optional

Turn on PDB post-mortem analysis for failing tests. Same as specifying `'--pdb'` in args.

coverage : bool, optional

Generate a test coverage report. The result will be placed in the directory `htmlcov`.

open_files : bool, optional

Fail when any tests leave files open. Off by default, because this adds extra run time to the test suite. Requires the [psutil](#) package.

parallel : int, optional

When provided, run the tests in parallel on the specified number of CPUs. If `parallel` is negative, it will use all the cores on the machine. Requires the [pytest-xdist](#) plugin installed. Only available when using Astropy 0.3 or later.

kwargs

Any additional keywords passed into this function will be passed on to the astropy test runner. This allows use of test-related functionality implemented in later versions of astropy without explicitly updating the package template.

Part II

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