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# **DrizzlePac Documentation**

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This package supports the use of `AstroDrizzle` as an integrated set of modules that can be run in an automated manner to combine images, along with other tasks to support image alignment and coordinate transformations with distortion included. The version of `DrizzlePac` described here implements a single task to run the entire `AstroDrizzle` processing pipeline, while also providing the framework for users to create their own custom pipeline based on the modules in this package merged with their own custom code if desired. These pages document what functions and classes are available for use under Python while providing the syntax for calling those functions from Python tasks.

Full documentation of how to run the primary `AstroDrizzle` and `TweakReg` tasks, along with fully worked examples, can be found in the [DrizzlePac Handbook](#).

This package relies on the `STWCS` package in order to provide the support for the WCS-based distortion models and alignment of the input images.

Contents:



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## Primary User Interface: AstroDrizzle()

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`AstroDrizzle` - Python implementation of `MultiDrizzle`

`AstroDrizzle` automates the process of aligning images in an output frame, identifying cosmic-rays, removing distortion, and then combining the images after removing the identified cosmic-rays.

**This process involves a number of steps, such as:**

- Processing the input images and input parameters
- Creating a static mask
- Performing sky subtraction
- Drizzling onto separate output images
- Creating the median image
- Blotting the median image
- Identifying and flagging cosmic-rays
- Final combination
- Cleaning-up of temporary files (when applicable)

A full description of this process can be found in the [DrizzlePac Handbook](#).

The primary output from this task is the distortion-corrected, cosmic-ray cleaned, and combined image as a FITS file.

This task requires numerous user-settable parameters to control the primary aspects of each of the processing steps.

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`drizzlepac.astrodrizzle.AstroDrizzle` (*input=None*, *mdriztab=False*, *editpars=False*, *configobj=None*, *wcsmmap=None*, *\*\*input\_dict*)

## Parameters

**input** [str or list of str (Default = `'*flt.fits'`)] The name or names of the input files to be processed, which can be provided in any of the following forms:

- filename of a single image
- filename of an association (ASN) table
- wild-card specification for files in directory
- comma-separated list of filenames
- `@file` filelist containing list of desired input filenames. The file list needs to be provided as an ASCII text file containing a list of filenames for all input images with one filename on each line of the file. If inverse variance maps (IVM maps) have also been created by the user and are to be used (by specifying `'IVM'` to the parameter `final_wht_type`), then these are simply provided as a second column in the filelist, with each IVM filename listed on the same line as a second entry, after its corresponding exposure filename.

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**Note:** If the user specifies IVM for the `final_wht_type`, but does not provide the names of IVM files, `AstroDrizzle` will automatically generate the IVM files itself for each input exposure.

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**editpars** [bool (Default = False)] A parameter that allows user to edit input parameters by hand in the GUI. `True` to use the GUI to edit parameters.

**configobj** [ConfigObjPars, ConfigObj, dict (Default = None)] An instance of `stsci.tools.cfgpars.ConfigObjPars` or `stsci.tools.configobj.ConfigObj` which overrides default parameter settings. When `configobj` is `defaults`, default parameter values are loaded from the user local configuration file usually located in `~/.teal/astrodrizzle.cfg` or a matching configuration file in the current directory. This configuration file stores most recent settings that an user used when running `AstroDrizzle` through the TEAL interface. When `configobj` is `None`, `AstroDrizzle` parameters not provided explicitly will be initialized with their default values as described in the “Other Parameters” section.

**input\_dict** [dict, optional] An optional list of parameters specified by the user, which can also be used to override the defaults.

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**Note:** This list of parameters **can** include the `updatewcs` parameter, even though this parameter no longer can be set through the TEAL GUI.

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**Note:** This list of parameters **can** contain parameters specific to the `AstroDrizzle` task itself described here in the “Other Parameters” section.

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## Other Parameters

**output** [str (Default = ‘’)] The rootname for the output drizzled products. This step can result in the creation of several files, including:

- copies of each input image as a FITS image, if `workinplace='Yes'` and/or input images are in GEIS format.
- mask files and coeffs files created by `PyDrizzle` for use by `drizzle`.

If an association file has been given as input, the specified filename will be used instead of the product name specified in the ASN file. Similarly, if a single exposure is provided, the rootname of the single exposure will be used for the output product instead of relying on the input rootname. If no value is provided when a filelist or wild-card specification is given as input, then a rootname of `'final'` will be used for the output file name.

**mdriztab** [bool (Default = False)] This button will immediately update the parameter values in the TEAL GUI based on those provided by the MDRIZTAB reference table referenced in the first input image. This requires that the MDRIZTAB reference file be available locally.

**runfile** [str (Default = ‘astrodrizzle.log’)] This log file will contain all the output messages generated during processing, including full details of any errors/exceptions. These messages will be a super-set of those reported to the screen during processing.

**wcskey** [str (Default = ‘’)] This parameter corresponds to the *key* for the WCS being selected by the user. It allows the user to select which WCS solution should be used for processing the images when multiple WCS’s have been updated in each input image header using the Paper I Multiple WCS FITS standard.

**Warning:** Use of this parameter should be done only when all input images have been updated using the Paper I FITS standard for specifying Multiple WCS’s in each image header. This parameter assumes that the same WCS letter corresponds to WCS’s that have been updated in a consistent manner. For example, all input images have been updated to be consistent with their distortion model in the WCS’s with key of *A*.

**proc\_unit** [str (Default = ‘native’)] The units to be used for the final output drizzled product. Valid values and definitions are:

- `'native'`: Output DRZ product and input ‘values’ given in the native units of the input image.

- `'electrons'`: Output DRZ product and input 'values' given in units of electrons.

**coeffs** [bool (Default = Yes)] This parameter determines whether or not to use the coefficients stored in the each input image header. If turned off, no distortion coefficients will be applied during the coordinate transformations.

**context** [bool (Default = Yes)] This parameter specifies whether or not to create a context image during the final drizzle combination. The context image contains the information regarding which image(s) contributed to each pixel encoded as a bit-mask. More information on context images can be obtained from the ACS Data Handbook.

**group** [int (Default = None)] This parameter establishes whether or not a single FITS extension, or group will be drizzled. If an extension is provided, then only that chip will be drizzled onto the output frame. Either a FITS extension number, a GEIS group number (such as '1'), or a FITS extension name (such as 'sci, 1') may be specified.

**build** [bool (Default = No)] When this parameter is set to 'Yes' (`True`), AstroDrizzle will combine the separate 'drizzle' output files into a single multi-extension format FITS file. This combined output file will contain separate SCI (science), WHT (weight), and CTX (context) extensions. If this parameter is set to 'No' (`False`), a separate simple FITS file will be created for each aforementioned extension.

**crbit** [int (Default = 4096)] This parameter sets the bit value for CR identification in the DQ array.

**stepsize** [int (Default = 10)] This parameter controls the internal grid of points used in the coordinate transformation from the input image to the output frame. The default value of 10 indicates that every 10th pixel will be transformed using the full WCS-based transformation. All remaining pixels will then be transformed using bilinear interpolation based on those pixels (i.e. every 10th pixel in the case of the default parameter setting) that were fully transformed.

**resetbits** [int (Default = 4096)] This parameter allows the user to specify which DQ bits of each input image DQ array should be reset to a value of 0. This operation is performed on the copy of the input data after updating the headers based on the 'updatewcs' parameter, and prior to starting any of the AstroDrizzle processing steps (static mask, sky subtraction, and so on).

**num\_cores** [int (Default = None)] This specifies the number of CPU cores to use during processing. Any value less than 2 will disable all use of parallel processing.

**in\_memory** [bool (Default = False)] This parameter sets whether or not to keep all intermediate products in memory when processing. This includes all single drizzle products (`*single_sci` and `*single_wht`), median image, blot images, and crmask images. The use of this option will therefore require significantly more memory than usual to process the data while reducing the overall processing time by eliminating most of the disk activity. *Only* the products of the final drizzle step will get written out when this parameter gets specified as `True`.

**rules\_file** [str (Default = "")] Rules for how to blend the header keyword values for all the input exposures into a single header for the drizzle products are specified using this `rules_file`. The `fitsblender` package uses this file to determine what keywords should be written out to the drizzle product headers. If no file is specified (default), the rules file for the instrument as included with the `fitsblender` package will be used for defining the product headers.

#### **\*\*STATE OF INPUT FILES\*\***

**restore: bool (Default = No)** Setting this to 'Yes' (True) directs `AstroDrizzle` to copy the input images from the 'OrIg\_files' sub-directory and use them for processing, if they had been archived by `AstroDrizzle` using the `preserve` or `overwrite` parameters already. If set to 'Yes' and the input files had not been archived already, it will simply ignore this and work with the current input images.

**preserve** [bool (Default = Yes)] Copy input files to archive directory, if not already archived. This parameter determines whether or not `AstroDrizzle` creates a copy of the input file in a sub-directory called 'OrIg\_files'. If a copy already exists in this directory, then the previously existing version will NOT be overwritten.

**overwrite** [bool (Default = No)] Copy input files into archive, overwriting older files if required? This parameter will cause `AstroDrizzle` to make a copy of each input file in the 'OrIg\_files' directory regardless of whether a previous copy existed or not, and will overwrite any previous copy should it be present.

**clean** [bool (Default = No)] The temporary files created by `AstroDrizzle` can be automatically removed by setting this parameter to 'Yes' (True). The affected files include the coefficient and static mask files created by `PyDrizzle`, in addition to other intermediate files created by `AstroDrizzle`. It is often useful to retain the intermediate files and examine them when first learning how to run `AstroDrizzle`. However, when running `AstroDrizzle` routinely, or on a small disk drive, these files can be removed to conserve space.

#### **\*STEP 1: STATIC MASK\***

**static** [bool (Default = Yes)] Create a static bad-pixel mask from the data? This mask flags all pixels that deviate by more than a value of 'static\_sig' sigma below the image median, since these pixels are typically the result of bad pixel oversubtraction in the dark image during calibration.

**static\_sig** [float (Default = 4.0)] The number of sigma below the RMS to use as the clipping limit for creating the static mask.

#### **\*\*STEP 2: SKY SUBTRACTION\*\***

**skysub** [bool (Default = Yes)] Turn on or off sky subtraction on the input data. When `skysub` is set to `no`, then `skyuser` field will be enabled and if user specifies a header keyword showing the sky value in the image, then that value will be used for CR-rejection but it will not be subtracted from the (drizzled) image data. If user sets `skysub` to `yes` then `skyuser` field will be disabled (and if it is not empty - it will be ignored) and user can use one of the methods available through

the `skymethod` parameter to compute the sky or provide a file (see `skyfile` parameter) with values that should be subtracted from (single) drizzled images.

**skymethod** [{‘localmin’, ‘globalmin+match’, ‘globalmin’, ‘match’}] (Default = ‘localmin’)] Select the algorithm for sky computation:

- **‘localmin’**: compute a common sky for all members of *an exposure*. For a typical use, it will compute sky values for each chip/image extension (marked for sky subtraction in the `input` parameter) in an input image, and it will subtract the previously found minimum sky value from all chips (marked for sky subtraction) in that image. This process is repeated for each input image.

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**Note:** This setting is recommended when regions of overlap between images are dominated by “pure” sky (as opposite to extended, diffuse sources).

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**Note:** This is similar to the “skysub” algorithm used in previous versions of `AstroDrizzle`.

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- **‘globalmin’**: compute a common sky value for all members of **all** “sky-lines”. It will compute sky values for each chip/image extension (marked for sky subtraction in the `input` parameter) in **all** input images, find the minimum sky value, and then it will subtract the **same** minimum sky value from **all** chips (marked for sky subtraction) in **all** images. This method *may* useful when input images already have matched background values.
- **‘match’**: compute differences in sky values between images in common (pair-wise) sky regions. In this case computed sky values will be relative (delta) to the sky computed in one of the input images whose sky value will be set to (reported to be) 0. This setting will “equalize” sky values between the images in large mosaics. However, this method is not recommended when used in conjunction with `AstroDrizzle` because it computes relative sky values while `AstroDrizzle` needs “measured” sky values for median image generation and CR rejection.
- **‘globalmin+match’**: first find a minimum “global” sky value in all input images and then use `‘match’` method to equalize sky values between images.

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**Note:** This is the *recommended* setting for images containing diffuse sources (e.g., galaxies, nebulae) covering significant parts of the image.

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**skywidth** [float (Default = 0.3)] Bin width, in sigma, used to sample the distribution of pixel flux values in order to compute the sky background statistics.

**skystat** [{‘median’, ‘mode’, ‘mean’}] (Default = ‘median’)] Statistical method for determining the sky value from the image pixel values.

**skylower** [float (Default = None)] Lower limit of usable pixel values for computing the sky. This value should be specified in the units of the input image(s).

**skyupper** [float (Default = None)] Upper limit of usable pixel values for computing the sky. This value should be specified in the units of the input image(s).

**skyclip** [int (Default = 5)] Number of clipping iterations to use when computing the sky value.

**skylsigma** [float (Default = 4.0)] Lower clipping limit, in sigma, used when computing the sky value.

**skyusigma** [float (Default = 4.0)] Upper clipping limit, in sigma, used when computing the sky value.

**skymask\_cat** [str (Default = '')] File name of a catalog file listing user masks to be used with images.

**use\_static** [bool (Default = True)] Specifies whether or not to use static mask to exclude masked image pixels from sky computations.

**sky\_bits** [int, str, None (Default = 0)] Integer sum of all the DQ bit values from the input image's DQ array that should be considered "good" when building masks for sky computations. For example, if pixels in the DQ array can be combinations of 1, 2, 4, and 8 flags and one wants to consider DQ "defects" having flags 2 and 4 as being acceptable for sky computations, then `sky_bits` should be set to  $2+4=6$ . Then a DQ pixel having values 2,4, or 6 will be considered a good pixel, while a DQ pixel with a value, e.g.,  $1+2=3$ ,  $4+8=12$ , etc. will be flagged as a "bad" pixel.

Alternatively, one can enter a comma- or '+'-separated list of integer bit flags that should be added to obtain the final "good" bits. For example, both 4, 8 and 4+8 are equivalent to setting `sky_bits` to 12.

Default value (0) will make *all* non-zero pixels in the DQ mask to be considered "bad" pixels, and the corresponding image pixels will not be used for sky computations.

Set `sky_bits` to `None` to turn off the use of image's DQ array for sky computations.

In order to reverse the meaning of the `sky_bits` parameter from indicating values of the "good" DQ flags to indicating the "bad" DQ flags, prepend '~' to the string value. For example, in order not to use pixels with DQ flags 4 and 8 for sky computations and to consider as "good" all other pixels (regardless of their DQ flag), set `sky_bits` to `~4+8`, or `~4, 8`. To obtain the same effect with an `int` input value (except for 0), enter  $-(4+8+1)=-9$ . Following this convention, a `sky_bits` string value of '~0' would be equivalent to setting `sky_bits=None`.

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**Note:** DQ masks (if used), *will be* combined with user masks specified in the input @-file.

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**Note:** To summarize, below are provided allowable syntaxes for `sky_bits`:

- Specify that bits 4,8, and 512 be considered “good” bits and that all other bits be considered “bad” bits:
    - Integer: 524 [numerically: 4+8+512=524]
    - String: 4+8+512
    - String: 4,8,512
  - Specify that only bits 4,8, and 512 be considered “bad” bits and that all other bits be considered “good” bits:
    - Integer: -525 [numerically: ~(4+8+512)=~524=-(524+1)=-525]
    - String: ~4+8+512 or ~(4+8+512)
    - String: ~4,8,512 or ~(4,8,512)
- 

**skyfile** [str (Default = ‘’)] Name of file containing user-computed sky values to be used with each input image. This ASCII file should only contain 2 columns: image filename in column 1 and sky value in column 2 (and higher for multi-chip images). The sky value should be provided in units that match the units of the input image and for multi-chip images, if only one sky value was provided in column 2, the same value will be applied to all chips. If more than one sky value are provided (in columns 2, 3, ...) then the number of sky values should match the number of SCI extensions in the images.

**skyuser** [str (Default = ‘’)] Name of header keyword which records the sky value already subtracted from the image by the user. The `skyuser` parameter is ignored when `skysub` is set to `yes`.

Alternatively, user can enter the name of a file that contains user-computed sky values. To distinguish a file name from a header keyword, prepend '@' to the file name. For example '@my\_sky\_values.txt'. The format of the file with user-supplied sky values is the same as that of a `skyfile`.

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**Note:** When `skysub='no'` and `skyuser` field is empty, then `AstroDrizzle` will assume that sky background is 0.0 for the purpose of cosmic-ray rejection.

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### **\*\*STEP 3: DRIZZLE SEPARATE IMAGES\*\***

**driz\_separate** [bool (Default = Yes)] This parameter specifies whether or not to drizzle each input image onto separate output images. The separate output images

will all have the same WCS as the final combined output frame. These images are used to create the median image, needed for cosmic ray rejection.

**driz\_sep\_kernel** [str {'square', 'point', 'gaussian', 'turbo', 'tophat', 'lanczos3'} (Default = 'turbo')] Used for the initial separate drizzling operation only, this parameter specifies the form of the kernel function used to distribute flux onto the separate output images. The current options are:

- 'square': original classic drizzling kernel
- 'point': this kernel is a point so each input pixel can only contribute to the single pixel that is closest to the output position. It is equivalent to the limit as  $\text{pixfrac} \rightarrow 0$ , and is very fast.
- 'gaussian': this kernel is a circular gaussian with a FWHM equal to the value of  $\text{pixfrac}$ , measured in input pixels.
- 'turbo': this is similar to  $\text{kernel} = \text{'square'}$  but the box is always the same shape and size on the output grid, and is always aligned with the X and Y axes. This may result in a significant speed increase.
- 'tophat': this kernel is a circular “top hat” shape of width  $\text{pixfrac}$ . It effects only output pixels within a radius of  $\text{pixfrac}/2$  from the output position.
- 'lanczos3': a Lanczos style kernel, extending a radius of 3 pixels from the center of the detection. The Lanczos kernel is a damped and bounded form of the “sinc” interpolator, and is very effective for resampling single images when  $\text{scale} = \text{pixfrac} = 1$ . It leads to less resolution loss than other kernels, and typically results in reduced correlated noise in outputs.

**Warning:** The 'lanczos3' kernel tends to result in much slower processing as compared to other kernel options. This option should never be used for  $\text{pixfrac} \neq 1.0$ , and is not recommended for  $\text{scale} \neq 1.0$ .

The default for this step is 'turbo' since it is much faster than 'square', and it is quite satisfactory for the purposes of generating the median image. More information about the different kernels can be found in the help file for the drizzle task.

**driz\_sep\_wt\_scl** [float (Default = exptime)] This parameter specifies the weighting factor for input image. If  $\text{driz\_sep\_wt\_scl} = \text{exptime}$ , then the scaling value will be set equal to the exposure time found in the image header. The use of the default value is recommended for producing optimal behavior for most scenarios. It is possible to set  $\text{wt\_scl} = \text{expsq}$  for weighting by the square of the exposure time, which is optimal for read-noise dominated images.

**driz\_sep\_pixfrac** [float (Default = 1.0)] Fraction by which input pixels are “shrunk” before being drizzled onto the output image grid, given as a real number between 0 and 1. This specifies the size of the footprint, or “dropsizes”, of a pixel in units of the input pixel size. If  $\text{pixfrac}$  is set to less than 0.001, the kernel parameter will

be reset to 'point' for more efficient processing. In the step of drizzling each input image onto a separate output image, the default value of 1.0 is best in order to ensure that each output drizzled image is fully populated with pixels from the input image. For more information, see the help for the `drizzle` task.

**driz\_sep\_fillval** [int (Default = None)] Value to be assigned to output pixels that have zero weight, or that receive flux from any input pixels during drizzling. This parameter corresponds to the `fillval` parameter of the 'drizzle' task. If the default of `None` is used, and if the weight in both the input and output images for a given pixel are zero, then the output pixel will be set to the value it would have had if the input had a non-zero weight. Otherwise, if a numerical value is provided (e.g. 0), then these pixels will be set to that value.

**driz\_sep\_bits** [int, str, or None (Default = 0)] Integer sum of all the DQ bit values from the input image's DQ array that should be considered 'good' when building the weighting mask. This can also be used to reset pixels to good if they had been flagged as cosmic rays during a previous run of `AstroDrizzle`, by adding the value 4096 for ACS and WFPC2 data. For possible input formats, see the description for `sky_bits` parameter.

**\*\*STEP 3a: CUSTOM WCS FOR SEPARATE OUTPUTS\*\***

**driz\_sep\_wcs** [bool (Default = No)] Define custom WCS for separate output images?

**driz\_sep\_refimage** [str (Default = '')] Reference image from which a WCS solution can be obtained.

**driz\_sep\_rot** [float (Default = None)] Position Angle of output image's Y-axis relative to North. A value of 0.0 would orient the final output image to be North up. The default of `None` specifies that the images will not be rotated, but will instead be drizzled in the default orientation for the camera with the x and y axes of the drizzled image corresponding approximately to the detector axes. This conserves disk space, as these single drizzled images are only used in the intermediate step of creating a median image.

**driz\_sep\_scale** [float (Default = None)] Linear size of the output pixels in arcseconds/pixel for each separate drizzled image (used in creating the median for cosmic ray rejection). The default value of `None` specifies that the undistorted pixel scale for the first input image will be used as the pixel scale for all the output images.

**driz\_sep\_outnx** [int (Default = None)] Size, in pixels, of the X axis in the output images that each input will be drizzled onto. If no value is specified, the smallest size that can accommodate the full dithered field will be used.

**driz\_sep\_outny** [int (Default = None)] Size, in pixels, of the Y axis in the output images that each input will be drizzled onto. If no value is specified, the smallest size that can accommodate the full dithered field will be used.

**driz\_sep\_ra** [float (Default = None)] Right ascension (in decimal degrees) specifying the center of the output image. If this value is not designated, the center will automatically be calculated based on the distribution of image dither positions.



**driz\_sep\_dec** [float (Default = None)] Declination (in decimal degrees) specifying the center of the output image. If this value is not designated, the center will automatically be calculated based on the distribution of image dither positions.

**\*\*STEP 4: CREATE MEDIAN IMAGE\*\***

**median** [bool (Default = Yes)] This parameter specifies whether or not to create a median image. This median image will be used as the comparison ‘truth’ image in the cosmic ray rejection step.

**median\_newmasks** [bool (Default = Yes)] This parameter specifies whether or not new mask files will be created when the median image is created. These masks are generated from weight files previously produced by the “driz\_separate” step, and contain all bad pixel information used to exclude pixels when calculating the median. Generally this step should be set to ‘Yes’ (True), unless for some reason, it is desirable to include bad pixel information when generating the median.

**combine\_maskpt** [float (Default = 0.3)] Percentage of weight image values, below which the are flagged.

**combine\_type** [str {‘median’, ‘mean’, ‘minmed’, ‘imedian’, ‘imean’, ‘iminmed’}] (Default = ‘minmed’) This parameter defines the method that will be used to create the median image. The ‘mean’ and ‘median’ options set the calculation type when running ‘numcombine’, a numpy method for median-combining arrays to create the median image. The ‘minmed’ option will produce an image that is generally the same as the median, except in cases where the median is significantly higher than the minimum good pixel value. In this case, ‘minmed’ will choose the minimum value. The sigma thresholds for this decision are provided by the ‘combine\_nsigma’ parameter. However, as the ‘combine\_nsigma’ parameter does not adjust for the larger probability of a single “nsigma” event with a greater number of images, ‘minmed’ will bias the comparison image low for a large number of images. The value of sigma is computed as  $\sigma = \sqrt{(M + S + R^2)}$ , where  $M$  is the median image data (in electrons),  $S$  is the value of the subtracted sky (in electrons), and  $R$  is the value of the readout noise (in electrons). ‘minmed’ is highly recommended for three images, and is good for four to six images, but should be avoided for ten or more images.

A value of ‘median’ is the recommended method for a large number of images, and works equally well as minmed down to approximately four images. However, the user should set the `combine_nhigh` parameter to a value of 1 when using “median” with four images, and consider raising this parameter’s value for larger numbers of images. As a median averages the two inner values when the number of values being considered is even, the user may want to keep the total number of images minus `combine_nhigh` odd when using median.

The options starting with ‘i’, such as ‘imedian’, works just like the normal median operation except when dealing with a pixel were all the values are flagged as ‘bad’. In this case, the ‘i’ functions return the last pixel in the stack as if it were good. This will prevent saturated pixels in the image from leaving holes in the middle of the stars, for example.

**combine\_nsigma** [float (Default = ‘4 3’)] This parameter defines the sigmas used

for accepting minimum values, rather than median values, when using the 'minmed' combination method. If two values are specified the first value will be used in the initial choice between median and minimum, while the second value will be used in the “growing” step to reject additional pixels around those identified in the first step. If only one value is specified, then it is used in both steps.

**combine\_nlow** [int (Default = 0)] This parameter sets the number of low value pixels to reject automatically during image combination.

**combine\_nhigh** [int (Default = 0)] This parameter sets the number of high value pixels to reject automatically during image combination.

**combine\_lthresh** [float (Default = None)] Sets the lower threshold for clipping input pixel values during image combination. This value gets passed directly to `imcombine` for use in creating the median image. If the parameter is set to `None`, no thresholds will be imposed.

**combine\_hthresh** [float (Default = None)] This parameter sets the upper threshold for clipping input pixel values during image combination. The value for this parameter is passed directly to `imcombine` for use in creating the median image. If the parameter is set to `None`, no thresholds will be imposed.

**combine\_grow** [int (Default = 1)] Width, in pixels, beyond the limit set by the rejection algorithm being used, for additional pixels to be rejected in an image. This parameter is used to set the `grow` parameter in `imcombine` for use in creating the median image **only when** `combine_type` is '(i)minmed'. When `combine_type` is anything other than '(i)minmed', this parameter is ignored (set to 0).

**combine\_bufsize** [float (Default = None)] Size of buffer, in MB (MiB), to use when reading in each section of each input image. The default buffer size is 1MB. The larger the buffer size, the fewer times the code needs to open each input image and the more memory will be required to create the median image. A larger buffer can be helpful when using compression, since slower copies need to be made of each set of rows from each input image instead of using memory-mapping.

#### **\*\*STEP 5: BLOT BACK THE MEDIAN IMAGE\*\***

**blot** [bool (Default = Yes)] Perform the blot operation on the median image? If set to 'Yes' (`True`), the output will be median smoothed images that match each input chips location, and will be used in the cosmic ray rejection step.

**blot\_interp** [str{'nearest', 'linear', 'poly3', 'poly5', 'sinc'} (Default = 'poly5')] This parameter defines the method of interpolation to be used when blotting drizzled images back to their original WCS solution. Valid options include:

- 'nearest': Nearest neighbor
- 'linear': Bilinear interpolation in x and y
- 'poly3': Third order interior polynomial in x and y
- 'poly5': Fifth order interior polynomial in x and y

- 'sinc': Sinc interpolation (accurate but slow)

The 'poly5' interpolation method has been chosen as the default because it is relatively fast and accurate.

If 'sinc' interpolation is selected, then the value of the parameter for `blot_sinscl` will be used to specify the size of the sinc interpolation kernel.

**blot\_sinscl** [float (Default = 1.0)] Size of the sinc interpolation kernel in pixels.

**blot\_addsky** [bool (Default = Yes)] Add back a sky value using the MDRIZSKY value from the header. If 'Yes' (`True`), the `blot_skyval` parameter is ignored.

**blot\_skyval** [float (Default = 0.0)] This is a user-specified custom sky value to be added to the blot image. This is only used if `blot_addsky` is 'No' (`False`).

#### **\*\*STEP 6: REMOVE COSMIC RAYS WITH DERIV, DRIZ\_CR\*\***

**driz\_cr** [bool (Default = Yes)] Perform cosmic-ray detection? If set to 'Yes' (`True`), cosmic-rays will be detected and used to create cosmic-ray masks based on the algorithms from 'deriv' and `driz_cr`.

**driz\_cr\_corr** [bool (Default = No)] Create a cosmic-ray cleaned input image? If set to 'Yes' (`True`), a cosmic-ray cleaned `_crclean` image will be generated directly from the input image, and a corresponding `_crmask` file will be written to document detected pixels affected by cosmic-rays.

**driz\_cr\_snr** [list of floats (Default = '3.5 3.0')] The values for this parameter specify the signal-to-noise ratios for the `driz_cr` task to be used in detecting cosmic rays. See the help file for `driz_cr` for further discussion of this parameter.

**driz\_cr\_grow** [int (Default = 1)] The radius, in pixels, around each detected cosmic-ray, in which more stringent detection criteria for additional cosmic rays will be used.

**driz\_cr\_ctegrow** [int (Default = 0)] Length, in pixels, of the CTE tail that should be masked in the drizzled output.

**driz\_cr\_scale** [str (Default = '1.2 0.7')] Scaling factor applied to the derivative in `driz_cr` when detecting cosmic-rays. See the help file for `driz_cr` for further discussion of this parameter.

#### **\*\*STEP 7: DRIZZLE FINAL COMBINED IMAGE\*\***

**driz\_combine** [bool (Default = Yes)] This parameter specifies whether or not to drizzle each input image onto the final output image. This applies the generated cosmic-ray masks to the input images and creates a final, cleaned, distortion-corrected image.

**final\_wht\_type** [{'EXP', 'ERR', 'IVM'}] (Default = 'EXP') Specify the type of weighting image to apply with the bad pixel mask for the final drizzle step. The options for this parameter include:

- 'EXP': The default of 'EXP' indicates that the images will be weighted according to their exposure time, which is the standard behavior for drizzle. This weighting is a good approximation in the regime where the noise is dominated

by photon counts from the sources, while contributions from sky background, read-noise and dark current are negligible. This option is provided as the default since it produces reliable weighting for all types of data, including older instruments (eg., WFPC2), where more sophisticated options may not be available.

- **'ERR'**: Specifying **'ERR'** is an alternative for ACS and STIS data. In these cases, the final drizzled images will be weighted according to the inverse variance of each pixel in the input exposure files, calculated from the error array data extension that is in each calibrated input exposure file. This array is exposure time dependent, and encapsulates all of the noise sources in each exposure including read-noise, dark current, sky background, and Poisson noise from the sources themselves. For WFPC2, the ERR array is not produced during the calibration process, and therefore is not a viable option. We advise extreme caution when selecting the **'ERR'** option, since the nature of this weighting scheme can introduce photometric discrepancies in sharp unresolved sources, although these effects are minimized for sources with gradual variations between pixels. The “EXP” weighting option does not suffer from these effects, and is therefore the recommended option.
- **'IVM'**: Specifying **'IVM'** allows the user to either supply their own inverse-variance weighting map, or allow `AstroDrizzle` to generate one automatically on-the-fly during the final drizzle step. This parameter option may be necessary for specific purposes. For example, to create a drizzled weight file for software such as `SExtractor`, it is expected that a weight image containing all of the background noise sources (sky level, read-noise, dark current, etc), but not the Poisson noise from the objects themselves will be available. The user can create the inverse variance images and then specify their names using the `input` parameter for `AstroDrizzle` to specify an '@file'. This would be a single ASCII file containing the list of input calibrated exposure filenames (one per line), with a second column containing the name of the IVM file corresponding to each calibrated exposure. Each IVM file must have the same file format as the input file, and if provided as multi-extension FITS files (e.g., ACS or STIS data) then the IVM extension must have the `EXTNAME` of **'IVM'**. If no IVM files are specified on input, then `AstroDrizzle` will rely on the flat-field reference file and computed dark value from the image header to automatically generate an IVM file specific to each exposure.

**final\_kernel** [{'square', 'point', 'gaussian', 'turbo', 'tophat', 'lanczos3'}] (Default = 'square') This parameter specifies the form of the kernel function used to distribute flux onto the separate output images, for the initial separate drizzling operation only. The value options for this parameter include:

- **'square'**: original classic drizzling kernel
- **'point'**: this kernel is a point so each input pixel can only contribute to the single pixel that is closest to the output position. It is equivalent to the limit as `pixfrac`  $\rightarrow 0$ , and is very fast.
- **'gaussian'**: this kernel is a circular gaussian, measured in input pixels, with a FWHM value equal to the value of `pixfrac`.

- 'turbo': this is similar to kernel="square", except that the box is always the same shape and size on the output grid, and is always aligned with the X and Y axes. This may result in a significant speed increase.
- 'tophat': this kernel is a circular "top hat" shape of width `pixfrac`. It effects only output pixels within a radius of `pixfrac/2` from the output position.
- 'lanczos3': a Lanczos style kernel, extending a radius of 3 pixels from the center of the detection. The Lanczos kernel is a damped and bounded form of the "sinc" interpolator, and is very effective for resampling single images when `scale=pixfrac=1`. It leads to less resolution loss than other kernels, and typically results in reduced correlated noise in outputs.

**Warning:** The 'lanczos3' kernel tends to result in much slower processing as compared to other kernel options. This option should never be used for `pixfrac != 1.0`, and is not recommended for `scale != 1.0`.

The default for this step is 'turbo' since it is much faster than 'square', and it is quite satisfactory for the purposes of generating the median image. More information about the different kernels can be found in the help file for the drizzle task.

**final\_wt\_scl** [float (Default = exptime)] This parameter specifies the weighting factor for input image. If `final_wt_scl=exptime`, then the scaling value will be set equal to the exposure time found in the image header. The use of the default value is recommended for producing optimal behavior for most scenarios. It is possible to set `wt_scl='expsq'` for weighting by the square of the exposure time, which is optimal for read-noise dominated images.

**final\_pixfrac** [float (Default = 1.0)] Fraction by which input pixels are "shrunk" before being drizzled onto the output image grid, given as a real number between 0 and 1. This specifies the size of the footprint, or "dropsize", of a pixel in units of the input pixel size. If `pixfrac` is set to less than 0.001, the kernel parameter will be reset to 'point' for more efficient processing. In the step of drizzling each input image onto a separate output image, the default value of 1.0 is best in order to ensure that each output drizzled image is fully populated with pixels from the input image. For more information, see the help for the 'drizzle' task.

**final\_fillval** [float (Default = None)] The value for this parameter is to be assigned to the output pixels that have zero weight or which do not receive flux from any input pixels during drizzling. This parameter corresponds to the `fillval` parameter of the `drizzle` task. If the default of `None` is used, and if the weight in both the input and output images for a given pixel are zero, then the output pixel will be set to the value it would have had if the input had a non-zero weight. Otherwise, if a numerical value is provided (e.g. 0), then these pixels will be set to that numerical value.

**final\_bits** [int, str, or None (Default = 0)] Integer sum for all of the DQ bit values

from the input image's DQ array that should be considered 'good' when building the weight mask. This can also be used to reset pixels to good if they had been flagged as cosmic rays during a previous run of `AstroDrizzle`, by adding the value 4096 for ACS and WFPC2 data. For possible input formats, see the description for `sky_bits` parameter.

**final\_units** [str (Default = 'cps')] This parameter determines the units of the final drizzle-combined image, and can either be 'counts' or 'cps'. It is passed through to `drizzle` in the final drizzle step.

**\*\*STEP 7a: CUSTOM WCS FOR FINAL OUTPUT\*\***

**final\_wcs** [bool (Default = No)] Obtain the WCS solution from a user-designated reference image?

**final\_refimage** [str (Default = '')] Reference image from which a WCS solution can be obtained. If no extension is specified (such as 'sci,1' or '4'), then `AstroDrizzle` will automatically look for the **first** extension which contains a valid HSTWCS object to read in as the WCS. Otherwise, the user can explicitly provide the extension name for multi-extension FITS files.

**final\_rot** [float (Default = None)] Position Angle of output image's Y-axis relative to North. A value of 0.0 would orient the final output image to be North up. The default of `None` specifies that the images will not be rotated, but will instead be drizzled in the default orientation for the camera with the x and y axes of the drizzled image corresponding approximately to the detector axes. This conserves disk space, as these single drizzled images are only used in the intermediate step of creating a median image.

**final\_scale** [float (Default = None)] Linear size of the output pixels in arcseconds/pixel for each separate drizzled image (used in creating the median for cosmic ray rejection). The default value of `None` specifies that the undistorted pixel scale for the first input image will be used as the pixel scale for all the output images.

**final\_outnx** [int (Default = None)] Size, in pixels, of the X axis in the output images that each input will be drizzled onto. If no value is specified, the smallest size that can accommodate the full dithered field will be used.

**final\_outny** [int (Default = None)] Size, in pixels, of the Y axis in the output images that each input will be drizzled onto. If no value is specified, the smallest size that can accommodate the full dithered field will be used.

**final\_ra** [float (Default = None)] Right ascension (in decimal degrees) specifying the center of the output image. If this value is not designated, the center will automatically be calculated based on the distribution of image dither positions.

**final\_dec** [float (Default = None)] Declination (in decimal degrees) specifying the center of the output image. If this value is not designated, the center will automatically be calculated based on the distribution of image dither positions.

**\*\*INSTRUMENT PARAMETERS\*\***

**gain** [float (Default = None)] Value used to override instrument specific default gain values. The value is assumed to be in units of electrons/count. This parameter should not be populated if the `gainkeyword` parameter is in use.

**gainkeyword** [str (Default = '')] Keyword used to specify a value, which is used to override the instrument specific default gain values. The value is assumed to be in units of electrons/count. This parameter should not be populated if the `gain` parameter is in use.

**rdnoise** [float (Default = None)] Value used to override instrument specific default readnoise values. The value is assumed to be in units of electrons. This parameter should not be populated if the `rnkeyword` parameter is in use.

**rnkeyword** [str (Default = '')] Keyword used to specify a value, which is used to override the instrument specific default readnoise values. The value is assumed to be in units of electrons. This parameter should not be populated if the `rdnoise` parameter is in use.

**exptime** [float (Default = None)] Value used to override default exposure time image header values. The value is assumed to be in units of seconds. This parameter should not be populated if the `expkeyword` parameter is in use.

**expkeyword** [str (Default = '')] Keyword used to specify a value, which is used to override the default exposure time image header values. The value is assumed to be in units of seconds. This parameter should not be populated if the `exptime` parameter is in use.

**\*\*ADVANCED PARAMETERS AVAILABLE FROM COMMAND LINE\*\***

**updatewcs** [bool (Default = No)] This parameter specifies whether the WCS keywords are to be updated by running `updatewcs` on the input data, or left alone. The update performed by `updatewcs` not only recomputes the WCS based on the currently used `IDCTAB`, but also populates the header with the `SIP` coefficients. For ACS/WFC images, the time-dependence correction will also be applied to the WCS and `SIP` keywords. This parameter should be set to 'No' (`False`) when the WCS keywords have been carefully set by some other method, and need to be passed through to `drizzle` 'as is', otherwise those updates will be over-written by this update.

---

**Note:** This parameter was preserved in the API for compatibility purposes with existing user processing pipe-lines. However, it has been removed from the `TEAL` interface because it is easy to have it set to 'Yes' (especially between consecutive runs of `AstroDrizzle`) with potentially disastrous effects on input image WCS (for example it could wipe-out previously aligned WCS).

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**See also:**

`drizzlepac.adrizzle` Apply the 'drizzle' algorithm to the images

`drizzlepac.ablot` Apply the 'blot' algorithm to drizzled images

`drizzlepac.sky` Perform sky subtraction

`stsci.skypac.skymatch` Sky computation and equalization

`drizzlepac.createMedian` Create a median combined image from a set of drizzled images

`drizzlepac.drizCR` Identify cosmic-rays by comparing blotted, median images to the original input images.

## Notes

Something to keep in mind is that the full `AstroDrizzle` interface will make backup copies of your original files and place them in the `'Orig_files'` directory of your current working directory.

All calibrated input images must have been updated using `updatewcs` from the `STWCS` package, to include the full distortion model in the header. Alternatively, one can set `updatewcs` parameter to `True` when running either `TweakReg` or `AstroDrizzle` from command line (Python interpreter) **the first time** on such images.

## Examples

The `AstroDrizzle` task can be run from either the `TEAL` GUI or from the command-line using `PyRAF` or `Python`. These examples illustrate the various syntax options available.

**Example 1:** Drizzle a set of calibrated (`_flt.fits`) images using mostly default parameters. Select the ‘World Coordinate System’ keyword to the updated solution computed from `TweakReg`. When combining images, ignore pixel flags of 64 and 32 in the `DQ` array of the (`_flt.fits`) images. Align the final product such that North is up, and set the final pixel scale to 0.05 arcseconds/pixel.

1. Run the task from `PyRAF` using the `TEAL` GUI:

```
>>> import drizzlepac
>>> epar astrodrizzle
```

2. Run the task from `PyRAF` using the command line.

```
>>> import drizzlepac
>>> from drizzlepac import astrodrizzle
>>> astrodrizzle.AstroDrizzle('*flt.fits', output='final',
...     wcskey='TWEAK', driz_sep_bits='64,32', final_wcs=True,
...     final_scale=0.05, final_rot=0)
```

Or, run the same task from the `PyRAF` command line, but specify all parameters in a config file named `myparam.cfg`:

```
>>> astrodrizzle.AstroDrizzle('*flt.fits', configobj='myparam.cfg')
```

3. Run the task directly from `Python`:



```
>>> from drizzlepac import astrodrizzle
>>> astrodrizzle.AstroDrizzle('*flt.fits', output='final',
...     wcskey='TWEAK', driz_sep_bits='64,32', final_wcs=True,
...     final_scale=0.05, final_rot=0)
```

4. Help can be accessed via the “Help” pulldown menu in the TEAL GUI. It can also be accessed from the PyRAF command-line and saved to a text file:

```
>>> from drizzlepac import astrodrizzle
>>> astrodrizzle.help()
```

or

```
>>> astrodrizzle.help(file='help.txt')
>>> page help.txt
```

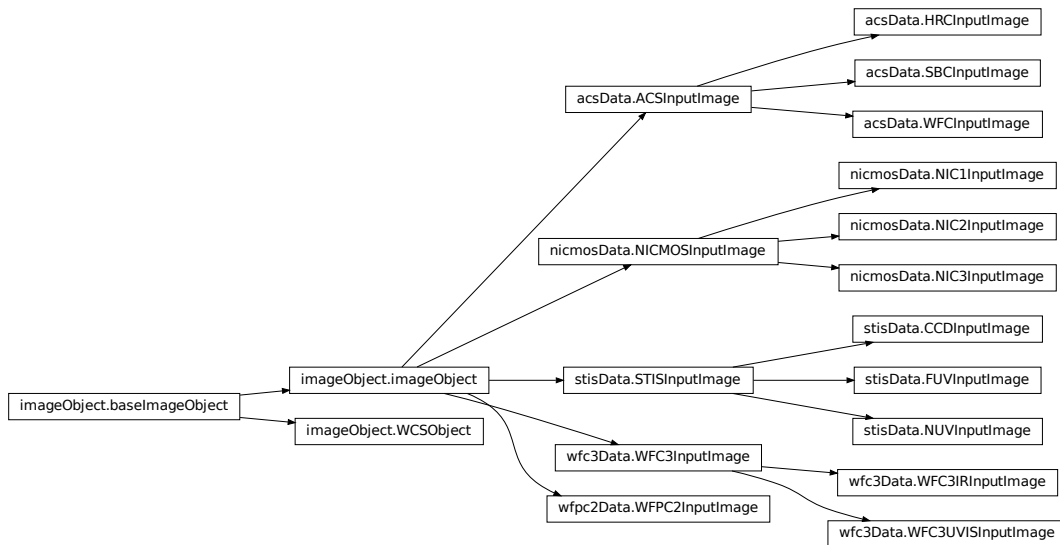


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## imageObject Classes

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This class and related sub-classes manage all the instrument-specific images for processing by AstroDrizzle.



### 2.1 Base ImageObject Classes

A class which makes image objects for each input filename.

A class which makes image objects for each input filename.

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**License** *LICENSE*

**class** drizzlepac.imageObject.**baseImageObject** (*filename*)

Bases: `object`

Base ImageObject which defines the primary set of methods.

**buildERRmask** (*chip, dqarr, scale*)

Builds a weight mask from an input DQ array and an ERR array associated with the input image.

**buildEXPmask** (*chip, dqarr*)

Builds a weight mask from an input DQ array and the exposure time per pixel for this chip.

**buildIVMmask** (*chip, dqarr, scale*)

Builds a weight mask from an input DQ array and either an IVM array provided by the user or a self-generated IVM array derived from the flat-field reference file associated with the input image.

**buildMask** (*chip, bits=0, write=False*)

Build masks as specified in the user parameters found in the configObj object.

We should overload this function in the instrument specific implementations so that we can add other stuff to the badpixel mask? Like vignetting areas and chip boundries in nicmos which are camera dependent? these are not defined in the DQ masks, but should be masked out to get the best results in multidrizzle.

**clean** ()

Deletes intermediate products generated for this imageObject.

**close** ()

Close the object nicely and release all the data arrays from memory YOU CANT GET IT BACK, the pointers and data are gone so use the getData method to get the data array returned for future use. You can use putData to reattach a new data array to the imageObject.

**findExtNum** (*extname=None, extver=1*)

Find the extension number of the give extname and extver.

**find\_DQ\_extension** ()

Return the suffix for the data quality extension and the name of the file which that DQ extension should be read from.

**getAllData** (*extname=None, exclude=None*)

This function is meant to make it easier to attach ALL the data extensions of the image object so that we can write out copies of the original image nicer.

If no extname is given, the it retrieves all data from the original file and attaches it. Otherwise, give the name of the extensions you want and all of those will be restored.

Ok, I added another option. If you want to get all the data extensions EXCEPT a particular one, leave extname=NONE and set exclude=EXTNAME. This is helpfull cause you might not know all the extnames the image has, this will find out and exclude the one you do not want overwritten.

**getData** (*exten=None*)

Return just the data array from the specified extension fileutil is used instead of fits to account for non-FITS input images. openImage returns a fits object.

**getExtensions** (*extname='SCI', section=None*)

Return the list of EXTVER values for extensions with name specified in extname.

**getGain** (*exten*)

**getHeader** (*exten=None*)

Return just the specified header extension fileutil is used instead of fits to account for non-FITS input images. openImage returns a fits object.

**getInstrParameter** (*value, header, keyword*)

This method gets a instrument parameter from a pair of task parameters: a value, and a header keyword.

**The default behavior is:**

- if the value and header keyword are given, raise an exception.
- if the value is given, use it.
- if the value is blank and the header keyword is given, use the header keyword.
- if both are blank, or if the header keyword is not found, return None.

**getKeywordList** (*kw*)

Return lists of all attribute values for all active chips in the `imageObject`.

**getNumpyType** (*irafType*)

Return the corresponding numpy data type.

**getOutputName** (*name*)

Return the name of the file or PyFITS object associated with that name, depending on the setting of `self.inmemory`.

**getReadNoiseImage** (*chip*)

## Notes

Method for returning the readnoise image of a detector (in electrons).

The method will return an array of the same shape as the image.

**Units** electrons

**getdarkcurrent** ()

## Notes

Return the dark current for the detector. This value will be contained within an instrument specific keyword. The value in the image header will be converted to units of electrons.

**Units** electrons

`getdarkimg` (*chip*)

### Notes

Return an array representing the dark image for the detector.

The method will return an array of the same shape as the image.

**Units** electrons

`getexptimeimg` (*chip*)

### Returns

**exptimeimg** [numpy array] The method will return an array of the same shape as the image.

### Notes

Return an array representing the exposure time per pixel for the detector. This method will be overloaded for IR detectors which have their own EXP arrays, namely, WFC3/IR and NICMOS images.

**Units** None

`getflat` (*chip*)

Method for retrieving a detector's flat field.

### Returns

**flat: array** This method will return an array the same shape as the image in **units of electrons**.

`getskyimg` (*chip*)

### Notes

Return an array representing the sky image for the detector. The value of the sky is what would actually be subtracted from the exposure by the skysub step.

**Units** electrons

`info` ()

Return fits information on the `_image`.

`putData` (*data=None, exten=None*)

Now that we are removing the data from the object to save memory, we need something that cleanly puts the data array back into the object so that we can write out everything together using something like `fits.writeto`. . . this method is an attempt to make sure that when you add an array back to the `.data` section of the `hdu` it still matches the header information for that section ( ie. update the `bitpix` to reflect the datatype of the array you are adding). The other header stuff is up to you to verify.

Data should be the data array `exten` is where you want to stick it, either extension number or a string like `'sci,1'`

**returnAllChips** (*extname=None, exclude=None*)

Returns a list containing all the chips which match the `extname` given minus those specified for exclusion (if any).

**saveVirtualOutputs** (*outdict*)

Assign in-memory versions of generated products for this `imageObject` based on dictionary `'outdict'`.

**set\_mt\_wcs** (*image*)

Reset the WCS for this image based on the WCS information from another `imageObject`.

**set\_units** ()

Record the units for this image, both BUNITS from header and `in_units` as needed internally. This method will be defined specifically for each instrument.

**set\_wtscl** (*chip, wtscl\_par*)

Sets the value of the `wt_scl` parameter as needed for drizzling.

**updateContextImage** (*contextpar*)

Reset the name of the context image to `None` if parameter `context` is `False`.

**updateData** (*exten, data*)

Write out updated data and header to the original input file for this object.

**updateIVMName** (*ivmname*)

Update `outputNames` for image with user-supplied IVM filename.

**updateOutputValues** (*output\_wcs*)

Copy info from output `WCSObject` into `outputnames` for each chip for use in creating `outputimage` object.

**class** `drizzlepac.imageObject.imageObject` (*filename, group=None, inmemory=False*)

Bases: `drizzlepac.imageObject.baseImageObject`

This returns an `imageObject` that contains all the necessary information to run the image file through any `multidrizzle` function. It is essentially a `PyFits` object with extra attributes.

There will be generic keywords which are good for the entire image file, and some that might pertain only to the specific chip.

**compute\_wcslin** (*undistort=True*)

Compute the undistorted WCS based solely on the known distortion model information associated with the WCS.

**setInstrumentParameters** (*instrpars*)

Define instrument-specific parameters for use in the code. By definition, this definition will need to be overridden by methods defined in each instrument's sub-class.

**set\_units** (*chip*)

Define units for this image.

```
class drizzlepac.imageObject.WCSObject (filename, suffix='_drz')
    Bases: drizzlepac.imageObject.baseImageObject

    restore_wcs ()
```

## 2.2 ACS ImageObjects

Class used to model ACS specific instrument data.

**Authors** Christopher Hanley, Warren Hack, Ivo Busko, David Grumm

**License** *LICENSE*

```
class drizzlepac.acsData.ACSInputImage (filename=None, group=None)
    Bases: drizzlepac.imageObject.imageObject
```

```
SEPARATOR = '_'
```

```
doUnitConversions ()
```

```
getdarkcurrent (extver)
```

Return the dark current for the ACS detector. This value will be contained within an instrument specific keyword. The value in the image header will be converted to units of electrons.

### Returns

**darkcurrent: float** Dark current value for the ACS detector in **units of electrons**.

```
class drizzlepac.acsData.WFCInputImage (filename=None, group=None)
    Bases: drizzlepac.acsData.ACSInputImage
```

```
setInstrumentParameters (instrpars)
```

This method overrides the superclass to set default values into the parameter dictionary, in case empty entries are provided.

This method gets called from processInput.

```
class drizzlepac.acsData.HRCInputImage (filename=None, group=None)
    Bases: drizzlepac.acsData.ACSInputImage
```

```
setInstrumentParameters (instrpars)
```

This method overrides the superclass to set default values into the parameter dictionary, in case empty entries are provided.

This method gets called from processInput.

```
class drizzlepac.acsData.SBCInputImage (filename=None, group=None)
    Bases: drizzlepac.acsData.ACSInputImage
```

```
setInstrumentParameters (instrpars)
```

Sets the instrument parameters.



## 2.3 WFC3 ImageObjects

wfc3Data module provides classes used to import WFC3 specific instrument data.

**Authors** Megan Sosey, Christopher Hanley

**License** *LICENSE*

**class** drizzlepac.wfc3Data.**WFC3InputImage** (*filename=None, group=None*)

Bases: *drizzlepac.imageObject.imageObject*

**SEPARATOR** = '\_'

**class** drizzlepac.wfc3Data.**WFC3UVISInputImage** (*filename=None, group=None*)

Bases: *drizzlepac.wfc3Data.WFC3InputImage*

**doUnitConversions** ()

**getdarkcurrent** (*chip*)

Return the dark current for the WFC3 UVIS detector. This value will be contained within an instrument specific keyword.

**Returns**

**darkcurrent: float** The dark current value with **units of electrons**.

**setInstrumentParameters** (*instrpars*)

This method overrides the superclass to set default values into the parameter dictionary, in case empty entries are provided.

**class** drizzlepac.wfc3Data.**WFC3IRInputImage** (*filename=None, group=None*)

Bases: *drizzlepac.wfc3Data.WFC3InputImage*

**doUnitConversions** ()

WF3 IR data come out in electrons, and I imagine the photometry keywords will be calculated as such, so no image manipulation needs be done between native and electrons

**getdarkcurrent** ()

Return the dark current for the WFC3/IR detector. This value will be contained within an instrument specific keyword.

**Returns**

**darkcurrent: float** The dark current value in **units of electrons**.

**getdarkimg** (*chip*)

Return an array representing the dark image for the detector.

**Returns**

**dark: array** Dark image array in the same shape as the input image with **units of cps**

**getexptimeimg** (*chip*)

Return an array representing the exposure time per pixel for the detector.

**Returns**

**dark:** array Exposure time array in the same shape as the input image

**getskyimg** (*chip*)

### Notes

Return an array representing the sky image for the detector. The value of the sky is what would actually be subtracted from the exposure by the skysub step.

**Units** electrons

**setInstrumentParameters** (*instrpars*)

This method overrides the superclass to set default values into the parameter dictionary, in case empty entries are provided.

## 2.4 STIS ImageObjects

stisData module provides classes used to import STIS specific instrument data.

**Authors** Megan Sosey, Christopher Hanley, Mihai Cara

**License** *LICENSE*

**class** drizzlepac.stisData.**STISInputImage** (*filename=None, group=None*)

Bases: *drizzlepac.imageObject.imageObject*

**SEPARATOR** = '\_'

**doUnitConversions** ()

Convert the data to electrons.

This converts all science data extensions and saves the results back to disk. We need to make sure the data inside the chips already in memory is altered as well.

**getflat** (*chip*)

Method for retrieving a detector's flat field. For STIS there are three. This method will return an array the same shape as the image.

**class** drizzlepac.stisData.**CCDInputImage** (*filename=None, group=None*)

Bases: *drizzlepac.stisData.STISInputImage*

**getReadNoise** ()

Method for returning the readnoise of a detector (in DN).

**Units** DN

This should work on a chip, since different chips to be consistent with other detector classes where different chips have different gains.

**getdarkcurrent** ()

Returns the dark current for the STIS CCD chip.

**Returns**

**darkcurrent** [float] Dark current value in **units of electrons** (or counts, if `proc_unit=='native'`).

**setInstrumentParameters** (*instrpars*)

This method overrides the superclass to set default values into the parameter dictionary, in case empty entries are provided.

**class** `drizzlepac.stisData.NUVInputImage` (*filename, group=None*)

Bases: `drizzlepac.stisData.STISInputImage`

**doUnitConversions** ()

Convert the data to electrons.

This converts all science data extensions and saves the results back to disk. We need to make sure the data inside the chips already in memory is altered as well.

**getdarkcurrent** ()

Returns the dark current for the STIS NUV detector.

#### Returns

**darkcurrent** [float] Dark current value in **units of electrons** (or counts, if `proc_unit=='native'`).

**setInstrumentParameters** (*instrpars*)

This method overrides the superclass to set default values into the parameter dictionary, in case empty entries are provided.

**class** `drizzlepac.stisData.FUVInputImage` (*filename=None, group=None*)

Bases: `drizzlepac.stisData.STISInputImage`

**doUnitConversions** ()

Convert the data to electrons.

This converts all science data extensions and saves the results back to disk. We need to make sure the data inside the chips already in memory is altered as well.

**getdarkcurrent** ()

Returns the dark current for the STIS FUV detector.

#### Returns

**darkcurrent** [float] Dark current value in **units of electrons** (or counts, if `proc_unit=='native'`).

**setInstrumentParameters** (*instrpars*)

This method overrides the superclass to set default values into the parameter dictionary, in case empty entries are provided.

## 2.5 NICMOS ImageObjects

Class used to model NICMOS specific instrument data.

**Authors** Christopher Hanley, David Grumm, Megan Sosey

License *LICENSE*

**class** drizzlepac.nicmosData.NICMOSInputImage (*filename=None*)

Bases: *drizzlepac.imageObject.imageObject*

**SEPARATOR** = '\_'

**doUnitConversions** ()

Convert the data to electrons

This converts all science data extensions and saves the results back to disk. We need to make sure the data inside the chips already in memory is altered as well.

**getdarkcurrent** ()

Return the dark current for the NICMOS detectors.

**Returns**

**darkcurrent** [float] Dark current value with **units of cps**.

**getdarkimg** (*chip*)

Return an array representing the dark image for the detector.

**Returns**

**dark** [array] The dark array in the same shape as the image with **units of cps**.

**getexptimeimg** (*chip*)

Return an array representing the exposure time per pixel for the detector.

**Returns**

**dark: array** Exposure time array in the same shape as the input image

**getflat** (*chip*)

Method for retrieving a detector's flat field.

**Returns**

**flat** [array] The flat field array in the same shape as the input image with **units of cps**.

**isCountRate** ()

isCountRate: Method or IRInputObject used to indicate if the science data is in units of counts or count rate. This method assumes that the keyword 'BUNIT' is in the header of the input FITS file.

**class** drizzlepac.nicmosData.NIC1InputImage (*filename=None*)

Bases: *drizzlepac.nicmosData.NICMOSInputImage*

**setInstrumentParameters** (*instrpars*)

This method overrides the superclass to set default values into the parameter dictionary, in case empty entries are provided.

**class** drizzlepac.nicmosData.NIC2InputImage (*filename=None*)

Bases: *drizzlepac.nicmosData.NICMOSInputImage*

**createHoleMask** ()

Add in a mask for the coronographic hole to the general static pixel mask.

**setInstrumentParameters** (*instrpars*)

This method overrides the superclass to set default values into the parameter dictionary, in case empty entries are provided.

**class** drizzlepac.nicmosData.NIC3InputImage (*filename=None*)

Bases: *drizzlepac.nicmosData.NICMOSInputImage*

**setInstrumentParameters** (*instrpars*)

This method overrides the superclass to set default values into the parameter dictionary, in case empty entries are provided.

## 2.6 WFPC2 ImageObjects

wfpc2Data module provides classes used to import WFPC2 specific instrument data.

**Authors** Warren Hack, Ivo Busko, Christopher Hanley

**License** *LICENSE*

**class** drizzlepac.wfpc2Data.WFPC2InputImage (*filename, group=None*)

Bases: *drizzlepac.imageObject.imageObject*

**SEPARATOR** = '\_'

**buildMask** (*chip, bits=0, write=False*)

Build masks as specified in the user parameters found in the configObj object.

**doUnitConversions** ()

Apply unit conversions to all the chips, ignoring the group parameter. This insures that all the chips get the same conversions when this gets done, even if only 1 chip was specified to be processed.

**find\_DQ\_extension** ()

Return the suffix for the data quality extension and the name of the file which that DQ extension should be read from.

**getEffGain** ()

Method used to return the effective gain of a instrument's detector.

**Returns**

**gain** [float] The effective gain.

**getReadNoise** (*exten*)

Method for returning the readnoise of a detector (in counts).

**Returns**

**readnoise** [float] The readnoise of the detector in **units of counts/electrons**.

**getdarkcurrent** (*exten*)

Return the dark current for the WFPC2 detector. This value will be contained within an instrument specific keyword. The value in the image header will be converted to units of electrons.

**Returns**

**darkcurrent** [float] Dark current for the WFPC3 detector in **units of counts/electrons**.

**getflat** (*chip*)

Method for retrieving a detector's flat field.

**Returns**

**flat** [array] The flat-field array in the same shape as the input image.

**setInstrumentParameters** (*instrpars*)

This method overrides the superclass to set default values into the parameter dictionary, in case empty entries are provided.

This module supports the interpretation and initial verification of all the input files specified by the user. These functions:

- reads in parameter values from MDRIZTAB reference file and merges those values in with the rest of the parameters from the GUI/configObj, if use of MDRIZTAB was specified
- insure that all input files are multi-extension FITS files and converts them if they are not
- updates all input WCS's to be consistent with IDCTAB, if specified
- generates the `ImageObject` instances for each input file
- resets the DQ bits if specified by the user
- adds info about any user-provided IVM files to the `ImageObjects`
- generates the output WCS based on user inputs

Process input to `MultiDrizzle/PyDrizzle`.

**Authors** Warren Hack

**License** *LICENSE*

The input can be one of:

- a python list of files
- a comma separated string of filenames (including wild card characters)
- an association table
- an @file (can have a second column with names of ivm files)

No mixture of instruments is allowed. No mixture of association tables, @files and regular fits files is allowed. Files can be in GEIS or MEF format (but not waiver fits).

Runs some sanity checks on the input files. If necessary converts files to MEF format (this should not be left to `makewcs` because `updatewcs` may be `False`). Runs `makewcs`. The function `process_input` returns an association table, `ivm` list, output name

The common interface interpreter for MultiDrizzle tasks, `processCommonInput()`, not only runs `process_input()` but `createImageObject()` and `defineOutput()` as well to fully setup all inputs for use with the rest of the MultiDrizzle steps either as stand-alone tasks or internally to MultiDrizzle itself.

`drizzlepac.processInput.addIVMInputs` (*imageObjectList*, *ivm*list)

Add IVM filenames provided by user to `outputNames` dictionary for each input `imageObject`.

`drizzlepac.processInput.applyContextPar` (*imageObjectList*, *contextpar*)

Apply the value of the parameter `context` to the input list, setting the name of the output context image to `None` if `context` is `False`

`drizzlepac.processInput.buildASNList` (*rootnames*, *asnname*,  
*check\_for\_duplicates=True*)

Return the list of filenames for a given set of rootnames

`drizzlepac.processInput.buildEmptyDRZ` (*input*, *output*)

Create an empty DRZ file.

This module creates an empty DRZ file in a valid FITS format so that the HST pipeline can handle the Multidrizzle zero exposure time exception where all data has been excluded from processing.

### Parameters

**input** [str] filename of the initial input to `process_input`

**output** [str] filename of the default empty `_drz.fits` file to be generated

`drizzlepac.processInput.buildFileList` (*input*, *output=None*, *ivm*list=*None*,  
*wcskey=None*, *updatewcs=True*, *\*\*workinplace*)

Builds a file list which has undergone various instrument-specific checks for input to MultiDrizzle, including splitting STIS associations.

`drizzlepac.processInput.buildFileListOrig` (*input*, *output=None*, *ivm*list=*None*,  
*wcskey=None*, *updatewcs=True*, *\*\*workinplace*)

Builds a file list which has undergone various instrument-specific checks for input to MultiDrizzle, including splitting STIS associations. Compared to `buildFileList`, this version returns the list of the original file names as specified by the user (e.g., before GEIS->MEF, or WAIVER FITS->MEF conversion).

`drizzlepac.processInput.changeSuffixinASN` (*asnfile*, *suffix*)

Create a copy of the original `asn` file and change the name of all members to include the suffix.

`drizzlepac.processInput.checkDGEOfile` (*filenames*)

Verify that input file has been updated with NPOLFILE

This function checks for the presence of 'NPOLFILE' kw in the primary header when 'DGEOfile' kw is present and valid (i.e. 'DGEOfile' is not blank or 'N/A'). It handles the case of science files downloaded from the archive before the new software was installed there. If 'DGEOfile' is present and 'NPOLFILE' is missing, print a message and let the user choose whether to (q)uit and update the



headers or (c)ontinue and run astrodrizzle without the non-polynomial correction. 'NPOLFILE' will be populated in the pipeline before astrodrizzle is run.

In the case of WFPC2 the old style dgeo files are used to create detector to image correction at runtime.

### Parameters

**filenames** [list of str] file names of all images to be checked

`drizzlepac.processInput.checkForDuplicateInputs` (*rootnames*)

Check input files specified in ASN table for duplicate versions with multiple valid suffixes (`_flt` and `_flc`, for example).

`drizzlepac.processInput.checkMultipleFiles` (*input*)

Evaluates the input to determine whether there is 1 or more than 1 valid input file.

`drizzlepac.processInput.createImageObjectList` (*files*, *instrpars*, *group=None*,  
*undistort=True*, *inmemory=False*)

Returns a list of `imageObject` instances, 1 for each input image in the list of input filenames.

`drizzlepac.processInput.getMdriztabPars` (*input*)

High-level function for getting the parameters from MDRIZTAB

Used primarily for TEAL interface.

`drizzlepac.processInput.manageInputCopies` (*filelist*, *\*\*workinplace*)

Creates copies of all input images in a sub-directory.

The copies are made prior to any processing being done to the images at all, including updating the WCS keywords. If there are already copies present, they will NOT be overwritten, but instead will be used to over-write the current working copies.

`drizzlepac.processInput.processFilenames` (*input=None*, *output=None*, *infilesOnly=False*)

Process the input string which contains the input file information and return a `filelist`, `output`

`drizzlepac.processInput.process_input` (*input*, *output=None*, *ivmlist=None*,  
*updatewcs=True*, *prodonly=False*,  
*wcskey=None*, *\*\*workinplace*)

Create the full input list of filenames after verifying and converting files as needed.

`drizzlepac.processInput.reportResourceUsage` (*imageObjectList*, *outwcs*,  
*num\_cores*, *interactive=False*)

Provide some information to the user on the estimated resource usage (primarily memory) for this run.

`drizzlepac.processInput.resetDQBits` (*imageObjectList*, *cr\_bits\_value=4096*)

Reset the CR bit in each input image's DQ array

`drizzlepac.processInput.runmakewcs` (*input*)

Runs make wcs and recomputes the WCS keywords

### Parameters

**input** [str or list of str] a list of files

## Returns

**output** [list of str] returns a list of names of the modified files (For GEIS files returns the translated names.)

`drizzlepac.processInput.setCommonInput (configObj, createOutwcs=True)`

The common interface interpreter for MultiDrizzle tasks which not only runs ‘process\_input()’ but ‘createImageObject()’ and ‘defineOutput()’ as well to fully setup all inputs for use with the rest of the MultiDrizzle steps either as stand-alone tasks or internally to MultiDrizzle itself.

## Parameters

**configObj** [object] configObj instance or simple dictionary of input parameters

**imageObjectList** [list of imageObject objects] list of imageObject instances, 1 for each input exposure

**outwcs** [object] imageObject instance defining the final output frame

## Notes

**At a minimum, the configObj instance (dictionary) should contain:** `configObj = {‘input’:None,‘output’:None }`

If provided, the configObj should contain the values of all the multidrizzle parameters as set by the user with TEAL. If no configObj is given, it will retrieve the default values automatically. In either case, the values from the input\_dict will be merged in with the configObj before being used by the rest of the code.

## Examples

You can set `createOutwcs=False` for the cases where you only want the images processed and no output wcs information in necessary; as in:

```
>>> imageObjectList, outwcs = processInput.processCommonInput (configObj)
```

`drizzlepac.processInput.update_member_names (oldasndict, pydr_input)`

Update names in a member dictionary.

Given an association dictionary with rootnames and a list of full file names, it will update the names in the member dictionary to contain ‘\_\*’ extension. For example a rootname of ‘u9600201m’ will be replaced by ‘u9600201m\_c0h’ making sure that a Mef file is passed as an input and not the corresponding GEIS file.

`drizzlepac.processInput.userStop (message)`

## 3.1 ResetBits Update of Input

This module provides the capability to set a specific set of bit values in the input DQ arrays to zero. This allows a user to reset pixels previously erroneously flagged as cosmic-rays to good for reprocessing with

improved alignment or detection parameters.

resetbits - A module to set the value of specified array pixels to zero

This module allows a user to reset the pixel values of any integer array, such as the DQ array from an HST image, to zero.

**Authors** Warren Hack

**License** *LICENSE*

### 3.1.1 PARAMETERS

**filename** [str] full filename with path

**bits** [str] sum or list of integers corresponding to all the bits to be reset

### 3.1.2 Optional Parameters

**extver** [int, optional] List of version numbers of the arrays to be corrected (default: None, will reset all matching arrays)

**extname** [str, optional] EXTNAME of the arrays in the FITS files to be reset (default: 'dq')

### 3.1.3 NOTES

This module performs a simple bitwise-and on all the pixels in the specified array and the integer value provided as input using the operation (array & ~bits).

### 3.1.4 Usage

It can be called not only from within Python, but also from the host-level operating system command line using the syntax:

```
resetbits filename bits [extver [extname]]
```

### 3.1.5 EXAMPLES

1. The following command will reset the 4096 bits in all the DQ arrays of the file 'input\_flt.fits':

```
resetbits input_flt.fits 4096
```

or from the Python command line:

```
>>> import resetbits
>>> resetbits.reset_dq_bits("input_file_flt.fits", 4096)
```

- To reset the 2,32,64 and 4096 (sum of 4194) bits in the second DQ array, specified as ‘dq,2’, in the file ‘input\_ft.fits’:

```
resetbits input_ft.fits 4194 2
```

or from the Python command line:

```
>>> import resetbits
>>> resetbits.reset_dq_bits("input_file_ft.fits", 2+32+64+4096,
↳extver=2)
```

`drizzlepac.resetbits.getHelpAsString` (*docstring=False, show\_ver=True*)

return useful help from a file in the script directory called `__taskname__.help`

`drizzlepac.resetbits.help` (*file=None*)

Print out syntax help for running `astrodrizzle`

### Parameters

**file** [str (Default = None)] If given, write out help to the filename specified by this parameter Any previously existing file with this name will be deleted before writing out the help.

`drizzlepac.resetbits.reset_dq_bits` (*input, bits, extver=None, extname='dq'*)

This function resets bits in the integer array(s) of a FITS file.

### Parameters

**filename** [str] full filename with path

**bits** [str] sum or list of integers corresponding to all the bits to be reset

**extver** [int, optional] List of version numbers of the DQ arrays to be corrected [Default Value: None, will do all]

**extname** [str, optional] EXTNAME of the DQ arrays in the FITS file [Default Value: ‘dq’]

### Notes

The default value of None for the ‘extver’ parameter specifies that all extensions with EXTNAME matching ‘dq’ (as specified by the ‘extname’ parameter) will have their bits reset.

### Examples

- The following command will reset the 4096 bits in all the DQ arrays of the file `input_file_ft.fits`:

```
reset_dq_bits("input_file_ft.fits", 4096)
```

- To reset the 2,32,64 and 4096 bits in the second DQ array, specified as ‘dq,2’, in the file `input_file_ft.fits`:

```
reset_dq_bits("input_fileflt.fits", "2,32,64,4096", extver=2)
```

`drizzlepac.resetbits.run` (*configobj=None*)  
Teal interface for running this code.



---

## Static Mask Step

---

This step focuses entirely on creating a static mask for each detector used in the input images of all negative (presumably bad) pixel values.

This module provides functions and classes that manage the creation of the global static masks.

For *staticMask*, the user interface function is *createMask()*.

**Authors** Ivo Busko, Christopher Hanley, Warren Hack, Megan Sosey

**License** *LICENSE*

`drizzlepac.staticMask.buildSignatureKey(signature)`

Build static file filename suffix used by mkstemp()

`drizzlepac.staticMask.constructFilename(signature)`

Construct an output filename for the given signature:

```
signature=[instr+detector, (nx,ny), detnum]
```

The signature is in the image object.

`drizzlepac.staticMask.createMask(input=None, static_sig=4.0, group=None, edit_pars=False, configObj=None, **inputDict)`

Create a static mask for all input images. The mask contains pixels that fall more than `static_sig` RMS below the mode for a given chip or extension. Those severely negative, or low pixels, might result from oversubtraction of bad pixels in the dark image, or high sky levels during calibration. For example, each ACS WFC image contains a separate image for each of 2 CCDs, and separate masks will be generated for each chip accordingly.

The final static mask for each chip contains all of the bad pixels that meet this criteria from all of the input images along with any bad pixels that satisfy the `final_bits` value specified by the user, and found in the images DQ mask.

Users should consider the details of their science image and decide whether or not creating this mask is appropriate for their resulting science. For example, if your field is very crowded, or contains mostly nebulous or extended objects, then the statistics could be heavily skewed and the mask could end up containing sources.

The generated static masks are saved to disk for use in later steps with the following naming convention:

```
[Instrument][Detector]_[xsize]x[ysize]_[detector number]_staticMask.fits
```

so an ACS image would produce a static mask with the name:

```
ACSWFC_2048x4096_1_staticMask.fits
```

and this would be the only file saved to disk, storing the logic and of all the badpixel masks created for each acs image in the set.

For more information on the science applications of the static mask task, see the [DrizzlePac Handbook](#)

### Parameters

**input** [str, None (Default = None)] A list of images or associations you would like to use to compute the mask.

**static** [bool (Default = True)] Create a static bad-pixel mask from the data? This mask flags all pixels that deviate by more than a value of `static_sig` sigma below the image median, since these pixels are typically the result of bad pixel oversubtraction in the dark image during calibration.

**static\_sig** [float (Default = 4.0)] The number of sigma below the RMS to use as the clipping limit for creating the static mask.

**editpars** [bool (Default = False)] Set to `True` if you would like to edit the parameters using the GUI interface.

### Examples

These tasks are designed to work together seamlessly when run in the full `AstroDrizzle` interface. More advanced users may wish to create specialized scripts for their own datasets, making use of only a subset of the predefined `AstroDrizzle` tasks, or add additional processing, which may be useful for their particular data. In these cases, individual access to the tasks is important.

Something to keep in mind is that the full `AstroDrizzle` interface will make backup copies of your original files and place them in the `Orig/` directory of your current working directory. If you are working with the stand alone interfaces, it is assumed that the user has already taken care of backing up their original datafiles as the input file with be directly altered.

Basic example of how to call static yourself from a python command line, using the default parameters for the task.

```
>>> from drizzlepac import staticMask
>>> staticMask.createMask('*flt.fits')
```



`drizzlepac.staticMask.createStaticMask` (*imageObjectList=[]*, *configObj=None*,  
*procSteps=None*)

`drizzlepac.staticMask.getHelpAsString` (*docstring=False*, *show\_ver=True*)  
return useful help from a file in the script directory called `__taskname__.help`

`drizzlepac.staticMask.help` (*file=None*)  
Print out syntax help for running astrodrizzle

### Parameters

**file** [str (Default = None)] If given, write out help to the filename specified by this parameter Any previously existing file with this name will be deleted before writing out the help.

`drizzlepac.staticMask.run` (*configObj*)

**class** `drizzlepac.staticMask.staticMask` (*configObj=None*)

This class manages the creation of the global static mask which masks pixels that are unwanted in the SCI array. A static mask object gets created for each global mask needed, one for each chip from each instrument/detector. Each static mask array has type Int16, and resides in memory.

**Notes** Class that manages the creation of a global static mask which is used to mask pixels that are some sigma BELOW the mode computed for the image.

**addMember** (*imagePtr=None*)

Combines the input image with the static mask that has the same signature.

### Parameters

**imagePtr** [object] An imageObject reference

### Notes

The signature parameter consists of the tuple:

```
(instrument/detector, (nx,ny), chip_id)
```

The signature is defined in the image object for each chip

**close** ()

Deletes all static mask objects.

**deleteMask** (*signature*)

Delete just the mask that matches the signature given.

**getFilename** (*signature*)

Returns the name of the output mask file that should reside on disk for the given signature.

**getMaskArray** (*signature*)

Returns the appropriate StaticMask array for the image.

**getMaskname** (*chipid*)

Construct an output filename for the given signature:

```
signature=[instr+detector, (nx,ny), detnum]
```

The signature is in the image object and the name of the static mask file is saved as `sci_chip.outputNames["staticMask"]`.

**saveToFile** (*imageObjectList*)

Saves the static mask to a file it uses the signatures associated with each mask to construct the filename for the output mask image.

---

## Sky-Subtraction Step

---

This step measures, subtracts and/or equalizes the sky from each input image while recording the subtracted value in the image header.

**Authors** Christopher Hanley, Megan Sosey, Mihai Cara

**License** *LICENSE*

`drizzlepac.sky.sky` (*input=None, outExt=None, configObj=None, group=None, edit-*  
*pars=False, \*\*inputDict*)

Function for computing and subtracting (or equalizing/matching) the background in input images. The algorithm for sky subtraction can be selected through the `skymethod` parameter. This function will update the `MDRIZSKY` keyword in the headers of the input files.

Sky subtraction is generally recommended for optimal flagging and removal of CR's when the sky background is more than a few electrons. However, some science applications may require the sky to not be removed, allowing for the final drizzle step to be performed with no sky subtraction. If you turn off sky subtraction, you should also set `drizzle.pixfrac` to 1, otherwise variations in sky between images will add noise to your data.

In addition to the “pure” sky computation, this task can be used for sky “equalization”, that is, it can match sky values in the images that are part of a mosaic.

For cameras with multiple detectors (such as ACS/WFC, WFPC2, or WFC3), the sky values in each exposure are first measured separately for the different detectors. These different values are then compared, and the lowest measured sky value is used as the estimate for all of the detectors for that exposure. This is based on the premise that for large extended or bright targets, the pixel intensity distribution in one or more of the detectors may be significantly skewed toward the bright end by the target itself, thereby overestimating the sky on that detector. If the other detector is less affected by such a target, then its sky value will be lower, and can therefore also be substituted as the sky value for the detector with the bright source.

For more information on the science applications of the sky task, see the [DrizzlePac Handbook](#).

## Parameters

**input** [str or list of str (Default = None)] A python list of image filenames, or just a single filename.

**outExt** [str (Default = None)] The extension of the output image. If the output already exists then the input image is overwritten.

**configObj** [configObject (Default = None)] An instance of `configObject`

**group** [int (Default = None)] The group of the input image.

**editpars** [bool (Default = False)] A parameter that allows user to edit input parameters by hand in the GUI.

**inputDict** [dict, optional] An optional list of parameters specified by the user.

---

**Note:** These are parameters that `configObj` should contain by default. These parameters can be altered on the fly using the `inputDict`. If `configObj` is set to None and there is no `inputDict` information, then the values for the parameters will be pulled from the default configuration files for the task. These are the same configuration files that are referenced in the TEAL parameter interface GUI. If you wish to edit these parameters by hand in the GUI, simply set `editpars=True`.

---

Table of optional parameters that should be in `configobj` and can also be specified in `inputDict`.

Name	Definition
<code>skyuser</code>	'KEYWORD in header which indicates a sky subtraction value to use'
<code>skymethod</code>	'Sky computation method'
<code>skysub</code>	'Perform sky subtraction?'
<code>skywidth</code>	'Bin width of histogram for sampling sky statistics (in sigma)'
<code>skystat</code>	'Sky correction statistics parameter'
<code>skylower</code>	'Lower limit of usable data for sky (always in electrons)'
<code>skyupper</code>	'Upper limit of usable data for sky (always in electrons)'
<code>skyclip</code>	'Number of clipping iterations'
<code>skylsigma</code>	'Lower side clipping factor (in sigma)'
<code>skyusigma</code>	'Upper side clipping factor (in sigma)'
<code>skymask_cat</code>	'Catalog file listing image masks'
<code>use_static</code>	'Use static mask for skymatch computations?'
<code>sky_bits</code>	'Bit flags for identifying bad pixels in DQ array'
<code>skyuser</code>	'KEYWORD indicating a sky subtraction value if done by user'
<code>skyfile</code>	'Name of file with user-computed sky values'
<code>in_memory</code>	'Optimize for speed or for memory use'

These optional parameters are described in more detail below in the “Other Parameters” section.

## Returns

**None** [The input file's primary headers is updated with the computed sky value.]

## Other Parameters

**skysub** [bool (Default = Yes)] Turn on or off sky subtraction on the input data. When `skysub` is set to `no`, then `skyuser` field will be enabled and if user specifies a header keyword showing the sky value in the image, then that value will be used for CR-rejection but it will not be subtracted from the (drizzled) image data. If user sets `skysub` to `yes` then `skyuser` field will be disabled (and if it is not empty - it will be ignored) and user can use one of the methods available through the `skymethod` parameter to compute the sky or provide a file (see `skyfile` parameter) with values that should be subtracted from (single) drizzled images.

**skymethod** [{ 'localmin', 'globalmin+match', 'globalmin', 'match' }, optional (Default = 'localmin')] Select the algorithm for sky computation:

- **'localmin'**: compute a common sky for all members of *an exposure* (see NOTES below). For a typical use, it will compute sky values for each chip/image extension (marked for sky subtraction in the `input` parameter) in an input image, and it will subtract the previously found minimum sky value from all chips (marked for sky subtraction) in that image. This process is repeated for each input image.

---

**Note:** This setting is recommended when regions of overlap between images are dominated by “pure” sky (as opposite to extended, diffuse sources).

---



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**Note:** This is similar to the “skysub” algorithm used in previous versions of `astrodrizzle`.

---

- **'globalmin'**: compute a common sky value for all members of *all exposures* (see NOTES below). It will compute sky values for each chip/image extension (marked for sky subtraction in the `input` parameter) in **all** input images, find the minimum sky value, and then it will subtract the **same** minimum sky value from **all** chips (marked for sky subtraction) in **all** images. This method *may* useful when input images already have matched background values.
- **'match'**: compute differences in sky values between images in common (pair-wise) sky regions. In this case computed sky values will be relative (delta) to the sky computed in one of the input images whose sky value will be set to (reported to be) 0. This setting will “equalize” sky values between the images in large mosaics. However, this method is not recommended when used in conjunction with `AstroDrizzle` because it computes relative sky values while `AstroDrizzle` needs “measured” sky values for median image generation and CR rejection.
- **'globalmin+match'**: first find a minimum “global” sky value in all input images and then use **'match'** method to equalize sky values between images.

---

**Note:** This is the *recommended* setting for images containing diffuse sources (e.g., galaxies, nebulae) covering significant parts of the image.

---

**skywidth** [float, optional (Default Value = 0.1)] Bin width, in sigma, used to sample the distribution of pixel flux values in order to compute the sky background statistics.

**skystat** [{ 'median', 'mode', 'mean' }, optional (Default Value = 'median')] Statistical method for determining the sky value from the image pixel values.

**skylower** [float, optional (Default Value = INDEF)] Lower limit of usable pixel values for computing the sky. This value should be specified in the units of the input image.

**skyyupper** [float, optional (Default Value = INDEF)] Upper limit of usable pixel values for computing the sky. This value should be specified in the units of the input image.

**skyclip** [int, optional (Default Value = 5)] Number of clipping iterations to use when computing the sky value.

**skylsigma** [float, optional (Default Value = 4.0)] Lower clipping limit, in sigma, used when computing the sky value.

**skyusigma** [float, optional (Default Value = 4.0)] Upper clipping limit, in sigma, used when computing the sky value.

**skymask\_cat** [str, optional (Default Value = '')] File name of a catalog file listing user masks to be used with images.

**use\_static** [bool, optional (Default Value = True)] Specifies whether or not to use static mask to exclude masked image pixels from sky computations.

**sky\_bits** [int, None, optional (Default = 0)] Integer sum of all the DQ bit values from the input image's DQ array that should be considered "good" when building masks for sky computations. For example, if pixels in the DQ array can be combinations of 1, 2, 4, and 8 flags and one wants to consider DQ "defects" having flags 2 and 4 as being acceptable for sky computations, then `sky_bits` should be set to  $2+4=6$ . Then a DQ pixel having values 2,4, or 6 will be considered a good pixel, while a DQ pixel with a value, e.g.,  $1+2=3$ ,  $4+8=12$ , etc. will be flagged as a "bad" pixel.

Default value (0) will make *all* non-zero pixels in the DQ mask to be considered "bad" pixels, and the corresponding image pixels will not be used for sky computations.

Set `sky_bits` to `None` to turn off the use of image's DQ array for sky computations.

---

**Note:** DQ masks (if used), *will be* combined with user masks specified in the input @-file.

---

**skyfile** [str, optional (Default Value = '')] Name of file containing user-computed sky values to be used with each input image. This ASCII file should only contain 2 columns: image filename in column 1 and sky value in column 2. The sky value should be provided in units that match the units of the input image and for multi-chip images, the same value will be applied to all chips.

**skyuser** [str (Default = '')] Name of header keyword which records the sky value already subtracted from the image by the user. The `skyuser` parameter is ignored when `skysub` is set to `yes`.

---

**Note:** When `skysub`=`no` and `skyuser` field is empty, then `AstroDrizzle` will assume that sky background is 0.0 for the purpose of cosmic-ray rejection.

---

**in\_memory** [bool, optional (Default Value = False)] Specifies whether to optimize execution for speed (maximum memory usage) or use a balanced approach in which a minimal amount of image data is kept in memory and retrieved from disk as needed. The default setting is recommended for most systems.

## Notes

`sky()` provides new algorithms for sky value computations and enhances previously available algorithms used by, e.g., `Astrodrizzle`.

First, the standard sky computation algorithm (see `skymethod = 'localmin'`) was upgraded to be able to use DQ flags and user supplied masks to remove “bad” pixels from being used for sky statistics computations.

Second, two new methods have been introduced: `'globalmin'` and `'match'`, as well as a combination of the two – `'globalmin+match'`.

- The `'globalmin'` method computes the minimum sky value across *all* chips in *all* input images. That sky value is then considered to be the background in all input images.
- The `'match'` algorithm is somewhat similar to the traditional sky subtraction method (`skymethod='localmin'`) in the sense that it measures the sky independently in input images (or detector chips). The major differences are that, unlike the traditional method,
  1. `'match'` algorithm computes *relative* sky values with regard to the sky in a reference image chosen from the input list of images; *and*
  2. Sky statistics is computed only in the part of the image that intersects other images.

This makes `'match'` sky computation algorithm particularly useful for “equalizing” sky values in large mosaics in which one may have only (at least) pair-wise intersection of images without

having a common intersection region (on the sky) in all images.

The 'match' method works in the following way: for each pair of intersecting images, an equation is written that requires that average surface brightness in the overlapping part of the sky be equal in both images. The final system of equations is then solved for unknown background levels.

**Warning:** Current algorithm is not capable of detecting cases when some groups of intersecting images (from the input list of images) do not intersect at all other groups of intersecting images (except for the simple case when *single* images do not intersect any other images). In these cases the algorithm will find equalizing sky values for each group. However since these groups of images do not intersect each other, sky will be matched only within each group and the “inter-group” sky mismatch could be significant.

Users are responsible for detecting such cases and adjusting processing accordingly.

**Warning:** Because this method computes *relative sky values* compared to a reference image (which will have its sky value set to 0), the sky values computed with this method usually are smaller than the “absolute” sky values computed, e.g., with the 'localmin' algorithm. Since *AstroDrizzle* expects “true” (as opposite to *relative*) sky values in order to correctly compute the median image or to perform cosmic-ray detection, this algorithm is not recommended to be used *alone* for sky computations to be used with *AstroDrizzle*.

For the same reason, IVM weighting in *AstroDrizzle* should **not** be used with 'match' method: sky values reported in MDRIZSKY header keyword will be relative sky values (sky offsets) and derived weights will be incorrect.

- The 'globalmin+match' algorithm combines 'match' and 'globalmin' methods in order to overcome the limitation of the 'match' method described in the note above: it uses 'globalmin' algorithm to find a baseline sky value common to all input images and the 'match' algorithm to “equalize” sky values in the mosaic. Thus, the sky value of the “reference” image will be equal to the baseline sky value (instead of 0 in 'match' algorithm alone) making this method acceptable for use in conjunction with *AstroDrizzle*.

**Glossary: Exposure** – a *subset* of FITS image extensions in an input image that correspond to different chips in the detector used to acquire the image. The subset of image extensions that form an exposure is defined by specifying extensions to be used with input images (see parameter *input*).

See help for `skypac.parseat.parse_at_line()` for details on how to specify image extensions.

**Footprint** – the outline (edge) of the projection of a chip or of an exposure on the celestial sphere.

---

**Note:**



- Footprints are managed by the `SphericalPolygon` class.
  - Both footprints *and* associated exposures (image data, WCS information, and other header information) are managed by the `SkyLine` class.
  - Each `SkyLine` object contains one or more `SkyLineMember` objects that manage both footprints *and* associated *chip* data that form an exposure.
- 

**Remarks:**

- `sky()` works directly on *geometrically distorted* flat-fielded images thus avoiding the need to perform an additional drizzle step to perform distortion correction of input images.

Initially, the footprint of a chip in an image is approximated by a 2D planar rectangle representing the borders of chip's distorted image. After applying distortion model to this rectangle and projecting it onto the celestial sphere, it is approximated by spherical polygons. Footprints of exposures and mosaics are computed as unions of such spherical polygons while overlaps of image pairs are found by intersecting these spherical polygons.

**Limitations and Discussions:** Primary reason for introducing “sky match” algorithm was to try to equalize the sky in large mosaics in which computation of the “absolute” sky is difficult due to the presence of large diffuse sources in the image. As discussed above, `sky()` accomplishes this by comparing “sky values” in a pair of images in the overlap region (that is common to both images). Quite obviously the quality of sky “matching” will depend on how well these “sky values” can be estimated. We use quotation marks around *sky values* because for some image “true” background may not be present at all and the measured sky may be the surface brightness of large galaxy, nebula, etc.

Here is a brief list of possible limitations/factors that can affect the outcome of the matching (sky subtraction in general) algorithm:

- Since sky subtraction is performed on *flat-fielded* but *not distortion corrected* images, it is important to keep in mind that flat-fielding is performed to obtain uniform surface brightness and not flux. This distinction is important for images that have not been distortion corrected. As a consequence, it is advisable that point-like sources be masked through the user-supplied mask files. Alternatively, one can use `upper` parameter to limit the use of bright objects in sky computations.
- Normally, distorted flat-fielded images contain cosmic rays. This algorithm does not perform CR cleaning. A possible way of minimizing the effect of the cosmic rays on sky computations is to use `clipping (nclip > 0)` and/or set `upper` parameter to a value larger than most of the sky background (or extended source) but lower than the values of most CR pixels.
- In general, clipping is a good way of eliminating “bad” pixels: pixels affected by CR, hot/dead pixels, etc. However, for images with complicated backgrounds (extended galaxies, nebulae, etc.), affected by CR and noise, clipping process may mask different pixels in different images. If variations in the background are too strong, clipping may converge to different sky values in different images even when factoring in the “true” difference in the sky background between the two images.
- In general images can have different “true” background values (we could measure it if

images were not affected by large diffuse sources). However, arguments such as `lower` and `upper` will apply to all images regardless of the intrinsic differences in sky levels.

**How to use the tasks stand alone interface in your own scripts:** These tasks are designed to work together seamlessly when run in the full `AstroDrizzle` interface. More advanced users may wish to create specialized scripts for their own datasets, making use of only a subset of the predefined `AstroDrizzle` tasks, or add additional processing, which may be useful for their particular data. In these cases, individual access to the tasks is important.

Something to keep in mind is that the full `AstroDrizzle` interface will make backup copies of your original files and place them in the `Orig/` directory of your current working directory. If you are working with the stand alone interfaces, it is assumed that the user has already taken care of backing up their original datafiles as the input file will be directly altered.

## Examples

Basic example of how to call sky yourself from a python command line, this example will use the default parameter settings and subtract a sky value from each `*flt.fits` image in the current directory, saving the output file with the extension of “mysky”:

```
>>> from drizzlepac import sky
>>> sky.sky('*flt.fits', outExt='mysky')
```

`drizzlepac.sky.help` (*file=None*)

Print out syntax help for running `astrodrizzle`

### Parameters

**file** [str (Default = None)] If given, write out help to the filename specified by this parameter Any previously existing file with this name will be deleted before writing out the help.

---

## Image Drizzling Step

---

The operation of drizzling each input image needs to be performed twice during processing:

- single drizzle step: this initial step drizzles each image onto the final output WCS as separate images
- final drizzle step: this step produces the final combined image based on the cosmic-ray masks determined by `AstroDrizzle`

Interfaces to main drizzle functions.

**Authors** Warren Hack

**License** *LICENSE*

`drizzlepac.adrizzle.drizzle` (*input*, *outdata*, *wscmap=None*, *editpars=False*, *configObj=None*, *\*\*input\_dict*)

Each input image gets drizzled onto a separate copy of the output frame. When stacked, these copies would correspond to the final combined product. As separate images, they allow for treatment of each input image separately in the undistorted, final WCS system. These images provide the information necessary for refining image registration for each of the input images. They also provide the images that will be succeedingly combined into a median image and then used for the subsequent blot and cosmic ray detection steps.

Aside from the input parameters, this step requires:

- valid input images with SCI extensions
- valid distortion coefficients tables
- any optional secondary distortion correction images
- numpy object (in memory) for static mask

This step produces:

- singly drizzled science image (simple FITS format)

- singly drizzled weight images (simple FITS format)

These images all have the same WCS based on the original input parameters and those provided for this step; specifically, output shape, pixel size, and orientation, if any have been specified at all.

For more information on the science applications of the drizzle task, see the [AstroDrizzle Handbook](#)

### Other Parameters

**driz\_separate** [bool (Default = No)] This parameter specifies whether or not to drizzle each input image onto separate output images. The separate output images will all have the same WCS as the final combined output frame. These images are used to create the median image, needed for cosmic ray rejection.

**driz\_sep\_kernel** [{'square', 'point', 'gaussian', 'turbo', 'tophat', 'lanczos3'}] (Default = 'turbo') Used for the initial separate drizzling operation only, this parameter specifies the form of the kernel function used to distribute flux onto the separate output images. The current options are:

- **square**: original classic drizzling kernel
- **point**: this kernel is a point so each input pixel can only contribute to the single pixel that is closest to the output position. It is equivalent to the limit as `pixfrac`  $\rightarrow 0$ , and is very fast.
- **gaussian**: this kernel is a circular gaussian with a FWHM equal to the value of `pixfrac`, measured in input pixels.
- **turbo**: this is similar to kernel="square" but the box is always the same shape and size on the output grid, and is always aligned with the X and Y axes. This may result in a significant speed increase.
- **tophat**: this kernel is a circular "top hat" shape of width `pixfrac`. It effects only output pixels within a radius of `pixfrac/2` from the output position.
- **lanczos3**: a Lanczos style kernel, extending a radius of 3 pixels from the center of the detection. The Lanczos kernel is a damped and bounded form of the "sinc" interpolator, and is very effective for resampling single images when `scale=pixfrac=1`. It leads to less resolution loss than other kernels, and typically results in reduced correlated noise in outputs.

**Warning:** The 'lanczos3' kernel tends to result in much slower processing as compared to other kernel options. This option should never be used for `pixfrac!=1.0`, and is not recommended for `scale != 1.0`.

The default for this step is "**turbo**" since it is much faster than "**square**", and it is quite satisfactory for the purposes of generating the median image. More information about the different kernels can be found in the help file for the drizzle task.

**driz\_sep\_wt\_scl** [float (Default = exptime)] This parameter specifies the weighting factor for input image. If `driz_sep_wt_scl=exptime`, then the scaling

value will be set equal to the exposure time found in the image header. The use of the default value is recommended for producing optimal behavior for most scenarios. It is possible to set `wt_scl='expsq'` for weighting by the square of the exposure time, which is optimal for read-noise dominated images.

**driz\_sep\_pixfrac** [float (Default = 1.0)] Fraction by which input pixels are “shrunk” before being drizzled onto the output image grid, given as a real number between 0 and 1. This specifies the size of the footprint, or “dropsize”, of a pixel in units of the input pixel size. If `pixfrac` is set to less than 0.001, the kernel parameter will be reset to ‘point’ for more efficient processing. In the step of drizzling each input image onto a separate output image, the default value of 1.0 is best in order to ensure that each output drizzled image is fully populated with pixels from the input image. For more information, see the help for the *drizzle* task.

**driz\_sep\_fillval** [int or INDEF (Default = INDEF)] Value to be assigned to output pixels that have zero weight, or that receive flux from any input pixels during drizzling. This parameter corresponds to the `fillval` parameter of the *drizzle* task. If the default of `INDEF` is used, and if the weight in both the input and output images for a given pixel are zero, then the output pixel will be set to the value it would have had if the input had a non-zero weight. Otherwise, if a numerical value is provided (e.g. 0), then these pixels will be set to that value.

**driz\_sep\_bits** [int (Default = 0)] Integer sum of all the DQ bit values from the input image’s DQ array that should be considered ‘good’ when building the weighting mask. This can also be used to reset pixels to good if they had been flagged as cosmic rays during a previous run of *AstroDrizzle*, by adding the value 4096 for ACS and WFPC2 data. Please see the section on Selecting the Bits Parameter for a more detailed discussion.

**driz\_sep\_wcs** [bool (Default = No)] Define custom WCS for separate output images?

**driz\_sep\_refimage** [str (Default = ‘’)] Reference image from which a WCS solution can be obtained.

**driz\_sep\_rot** [float (Default = INDEF)] Position Angle of output image’s Y-axis relative to North. A value of 0.0 would orient the final output image to be North up. The default of `INDEF` specifies that the images will not be rotated, but will instead be drizzled in the default orientation for the camera with the x and y axes of the drizzled image corresponding approximately to the detector axes. This conserves disk space, as these single drizzled images are only used in the intermediate step of creating a median image.

**driz\_sep\_scale** [float (Default = INDEF)] Linear size of the output pixels in arcseconds/pixel for each separate drizzled image (used in creating the median for cosmic ray rejection). The default value of `INDEF` specifies that the undistorted pixel scale for the first input image will be used as the pixel scale for all the output images.

**driz\_sep\_outnx** [int (Default = INDEF)] Size, in pixels, of the X axis in the output images that each input will be drizzled onto. If no value is specified, the smallest size that can accommodate the full dithered field will be used.

**driz\_sep\_outny** [int (Default = INDEF)] Size, in pixels, of the Y axis in the output images that each input will be drizzled onto. If no value is specified, the smallest size that can accommodate the full dithered field will be used.

**driz\_sep\_ra** [float (Default = INDEF)] Right ascension (in decimal degrees) specifying the center of the output image. If this value is not designated, the center will automatically be calculated based on the distribution of image dither positions.

**driz\_sep\_dec** [float (Default = INDEF)] Declination (in decimal degrees) specifying the center of the output image. If this value is not designated, the center will automatically be calculated based on the distribution of image dither positions.

## Notes

These tasks are designed to work together seamlessly when run in the full `AstroDrizzle` interface. More advanced users may wish to create specialized scripts for their own datasets, making use of only a subset of the predefined `AstroDrizzle` tasks, or add additional processing, which may be useful for their particular data. In these cases, individual access to the tasks is important.

Something to keep in mind is that the full `AstroDrizzle` interface will make backup copies of your original files and place them in the `Orig/` directory of your current working directory. If you are working with the stand alone interfaces, it is assumed that the user has already taken care of backing up their original datafiles as the input file will be directly altered.

There are two user interface function for this task, one to allow you to create separately drizzled images of each image in your list and the other to create one single output drizzled image, which is the combination of all of them:

```
def drizSeparate(imageObjectList, output_wcs, configObj, wcsmap=wcs_
    ↪functions.WCSMap)
def drizFinal(imageObjectList, output_wcs, configObj, build=None,
    ↪wcsmap=wcs_functions.WCSMap)
if configObj[single_step]['driz_separate']:
    drizSeparate(imgObjList, outwcs, configObj, wcsmap=wcsmap)
else:
    drizFinal(imgObjList, outwcs, configObj, wcsmap=wcsmap)
```

## Examples

Basic example of how to call static yourself from a python command line, using the default parameters for the task.

```
>>> from drizzlepac import adrizzle
```

```
drizzlepac.adrizzle.run(configObj, wcsmap=None)
```

Interface for running `wdrizzle` from TEAL or Python command-line.

This code performs all file I/O to set up the use of the drizzle code for a single exposure to replicate the functionality of the original `wdrizzle`.

`drizzlepac.adrizzle.drizSeparate` (*imageObjectList*, *output\_wcs*, *configObj*, *wcsmmap=None*, *procSteps=None*)

`drizzlepac.adrizzle.drizFinal` (*imageObjectList*, *output\_wcs*, *configObj*, *build=None*, *wcsmmap=None*, *procSteps=None*)

`drizzlepac.adrizzle.mergeDQarray` (*maskname*, *dqarr*)

Merge static or CR mask with mask created from DQ array on-the-fly here.

`drizzlepac.adrizzle.updateInputDQArray` (*dqfile*, *dq\_extn*, *chip*, *crmaskname*, *cr\_bits\_value*)

`drizzlepac.adrizzle.buildDrizParamDict` (*configObj*, *single=True*)

`drizzlepac.adrizzle.interpret_maskval` (*paramDict*)

Apply logic for interpreting final\_maskval value...

`drizzlepac.adrizzle.run_driz` (*imageObjectList*, *output\_wcs*, *paramDict*, *single*, *build*, *wcsmmap=None*)

Perform drizzle operation on input to create output. The input parameters originally was a list of dictionaries, one for each input, that matches the primary parameters for an IRAF *drizzle* task.

This method would then loop over all the entries in the list and run *drizzle* for each entry.

**Parameters required for input in paramDict:** build, single, units, wt\_scl, pixfrac, kernel, fillval, rot, scale, xsh, ysh, blotnx, blotny, outnx, outny, data

`drizzlepac.adrizzle.run_driz_img` (*img*, *chiplist*, *output\_wcs*, *outwcs*, *template*, *paramDict*, *single*, *num\_in\_prod*, *build*, *\_versions*, *\_numctx*, *\_nplanes*, *chipIdxCopy*, *\_outsci*, *\_outwht*, *\_outctx*, *\_hdrlist*, *wcsmmap*)

Perform the drizzle operation on a single image. This is separated out from *run\_driz()* so as to keep together the entirety of the code which is inside the loop over images. See the *run\_driz()* code for more documentation.

`drizzlepac.adrizzle.run_driz_chip` (*img*, *chip*, *output\_wcs*, *outwcs*, *template*, *paramDict*, *single*, *doWrite*, *build*, *\_versions*, *\_numctx*, *\_nplanes*, *\_numchips*, *\_outsci*, *\_outwht*, *\_outctx*, *\_hdrlist*, *wcsmmap*)

Perform the drizzle operation on a single chip. This is separated out from *run\_driz\_img* so as to keep together the entirety of the code which is inside the loop over chips. See the *run\_driz* code for more documentation.

`drizzlepac.adrizzle.do_driz` (*inisci*, *input\_wcs*, *inwht*, *output\_wcs*, *outsci*, *outwht*, *outcon*, *expin*, *in\_units*, *wt\_scl*, *wcslin\_pscale=1.0*, *uniqid=1*, *pixfrac=1.0*, *kernel='square'*, *fillval='INDEF'*, *stepsize=10*, *wcsmmap=None*)

Core routine for performing 'drizzle' operation on a single input image All input values will be Python objects such as ndarrays, instead of filenames. File handling (input and output) will be performed by calling routine.

`drizzlepac.adrizzle.get_data` (*filename*)

`drizzlepac.adrizzle.create_output` (*filename*)

`drizzlepac.adrizzle.help` (*file=None*)  
Print out syntax help for running astrodrizzle

**Parameters**

**file** [str (Default = None)] If given, write out help to the filename specified by this parameter Any previously existing file with this name will be deleted before writing out the help.

`drizzlepac.adrizzle.getHelpAsString` (*docstring=False, show\_ver=True*)  
return useful help from a file in the script directory called `__taskname__.help`



---

## Median Image Computation Step

---

A median image gets generated from the stack of undistorted single drizzle images.

Create a median image from the singly drizzled images.

**Authors** Warren Hack, Mihai Cara

**License** *LICENSE*

`drizzlepac.createMedian.createMedian` (*imgObjList, configObj, procSteps=None*)

Top-level interface to createMedian step called from top-level AstroDrizzle.

This function parses the input parameters then calls the `_median()` function to median-combine the input images into a single image.

`drizzlepac.createMedian.getHelpAsString` (*docstring=False, show\_ver=True*)

return useful help from a file in the script directory called `__taskname__.help`

`drizzlepac.createMedian.help` (*file=None*)

Print out syntax help for running astrodrizzle

### Parameters

**file** [str (Default = None)] If given, write out help to the filename specified by this parameter Any previously existing file with this name will be deleted before writing out the help.

`drizzlepac.createMedian.median` (*input=None, configObj=None, editpars=False, \*\*inputDict*)

The singly drizzled science images are combined to create a single median image. This median combination gets performed section-by-section from the input single drizzled images. Each section corresponds to a contiguous set of lines from each image taking up no more than 1Mb in memory, such that combining 10 input images would only require 10Mb for these sections. The goal of this step is to establish an estimate for what the fully cleaned image should look like in order to enable better

bad pixel detection, in addition to improving the alignment of the image stack. Creating a median image from the aligned and undistorted input images allows for a statistical rejection of bad pixels.

The final median image serves as the only output from this step.

For more information on the science applications of the *createMedian* task, see the [DrizzlePac Handbook](#).

### Parameters

**input** [str or list of str (Default = None)] A python list of drizzled image filenames, or just a single filename.

**configObj** [configObject (Default = None)] An instance of `configObject` which overrides default parameter settings.

**editpars** [bool (Default = False)] A parameter that allows user to edit input parameters by hand in the GUI. True to use the GUI to edit parameters.

**inputDict** [dict, optional] An optional list of parameters specified by the user, which can also be used to override the defaults.

### Other Parameters

**median** [bool (Default = No)] This parameter specifies whether or not to create a median image. This median image will be used as the comparison ‘truth’ image in the cosmic ray rejection step.

**median\_newmasks** [bool (Default = Yes)] This parameter specifies whether or not new mask files will be created when the median image is created. These masks are generated from weight files previously produced by the “driz\_separate” step, and contain all bad pixel information used to exclude pixels when calculating the median. Generally this step should be set to “Yes”, unless for some reason, it is desirable to include bad pixel information when generating the median.

**combine\_maskpt** [float (Default = 0.7)] Percentage of weight image values, below which the are flagged.

**combine\_type** [str {‘average’, ‘median’, ‘sum’, ‘minmed’} (Default = ‘minmed’)] This parameter defines the method that will be used to create the median image. The ‘average’, ‘median’, and ‘sum’ options set the calculation type when running ‘numcombine’, a numpy method for median-combining arrays to create the median image. The “minmed” option will produce an image that is generally the same as the median, except in cases where the median is significantly higher than the minimum good pixel value. In this case, “minmed” will choose the minimum value. The sigma thresholds for this decision are provided by the “combine\_nsigma” parameter. However, as the “combine\_nsigma” parameter does not adjust for the larger probability of a single “nsigma” event with a greater number of images, “minmed” will bias the comparison image low for a large number of images. The value of sigma is computed as  $\sigma = \sqrt{(M + S + R^2)}$ , where  $M$  is the median image data (in electrons),  $S$  is the value of the subtracted sky (in electrons), and  $R$  is the value of the readout noise (in electrons). “minmed” is highly recommended for three images, and is good for four to six images, but should be avoided for ten or more images.

A value of ‘median’ is the recommended method for a large number of images, and works equally well as minmed down to approximately four images. However, the user should set the “combine\_nhigh” parameter to a value of 1 when using “median” with four images, and consider raising this parameter’s value for larger numbers of images. As a median averages the two inner values when the number of values being considered is even, the user may want to keep the total number of images minus “combine\_nhigh” odd when using “median”.

**combine\_nsigma** [float (Default = ‘4 3’)] This parameter defines the sigmas used for accepting minimum values, rather than median values, when using the ‘minmed’ combination method. If two values are specified the first value will be used in the initial choice between median and minimum, while the second value will be used in the “growing” step to reject additional pixels around those identified in the first step. If only one value is specified, then it is used in both steps.

**combine\_nlow** [int (Default = 0)] This parameter sets the number of low value pixels to reject automatically during image combination.

**combine\_nhigh** [int (Default = 0)] This parameter sets the number of high value pixels to reject automatically during image combination.

**combine\_lthresh** [float (Default = INDEF)] Sets the lower threshold for clipping input pixel values during image combination. This value gets passed directly to ‘imcombine’ for use in creating the median image. If the parameter is set to “None”, no thresholds will be imposed.

**combine\_hthresh** [float (Default = INDEF)] This parameter sets the upper threshold for clipping input pixel values during image combination. The value for this parameter is passed directly to ‘imcombine’ for use in creating the median image. If the parameter is set to “None”, no thresholds will be imposed.

**combine\_grow** [int (Default = 1)] Width, in pixels, beyond the limit set by the rejection algorithm being used, for additional pixels to be rejected in an image. This parameter is used to set the ‘grow’ parameter in ‘imcombine’ for use in creating the median image **only when** combine\_type is ‘(i)minmed’. When combine\_type is anything other than ‘(i)minmed’, this parameter is ignored (set to 0).

**combine\_bufsize** [float (Default = None)] Size of buffer, in MB (MiB), to use when reading in each section of each input image. The default buffer size is 1MB. The larger the buffer size, the fewer times the code needs to open each input image and the more memory will be required to create the median image. A larger buffer can be helpful when using compression, since slower copies need to be made of each set of rows from each input image instead of using memory-mapping.

## Examples

For `createMedian`, the user interface function is `median`:

```
>>> from drizzlepac import createMedian
>>> createMedian.median('*flt.fits')
```

`drizzlepac.createMedian.run` (*configObj*)

---

## Median Image Blotting Step

---

In this step the median image now gets blotted back to create median-cleaned images which can be compared directly with each input image to identify cosmic-rays.

**Authors** Warren Hack

**License** *LICENSE*

`drizzlepac.ablot.blot` (*data, outdata, configObj=None, wcsmap=wcs\_functions.WCSMap, editpars=False, \*\*input\_dict*)

The median image is the combination of the WCS aligned input images that have already had the distortion model applied. Taking the median of the aligned images allows for a statistical rejection of bad pixels from the image stack. The resulting median image can then be input for the blot task with the goal of creating ‘cleaned’ versions of the input images at each of their respective dither locations. These “blotted” images can then be directly compared to the original distorted input images for detection of image artifacts (i.e. bad-pixels, hot pixels, and cosmic-rays) whose locations will be saved to the output badpixel masks.

Aside from the input parameters, this step only requires opening the single median image created from all the input images. A distorted version of the median image corresponding to each input ‘chip’ (extension) is written as output from this step as separate simple FITS images.

For more information on the science applications of the blot task, see the [DrizzlePac Handbook](#)

### Parameters

**data** [str] Input distortion-corrected (median or drizzled) image to be used as the source for creating blotted images.

**reference** [str] Filename of image to read to define the blotted WCS; image with distortion to be matched by output blotted image.

**outdata** [str] Filename for output blotted image.

**coeffs** [bool (Default Value = True)] This parameter specifies whether or not to use the header-based distortion coefficients when creating the blotted, distorted image. If False, no distortion will be applied at all, effectively working as a cut-out operation.

**interp** [str{‘nearest’, ‘linear’, ‘poly3’, ‘poly5’, ‘sinc’} (Default = ‘poly5’)] This parameter defines the method of interpolation to be used when blotting drizzled images back to their original WCS solution. Valid options include:

- **nearest**: Nearest neighbor
- **linear**: Bilinear interpolation in x and y
- **poly3**: Third order interior polynomial in x and y
- **poly5**: Fifth order interior polynomial in x and y
- **sinc**: Sinc interpolation (accurate but slow)

The ‘poly5’ interpolation method has been chosen as the default because it is relatively fast and accurate.

If ‘sinc’ interpolation is selected, then the value of the parameter for `blot_sinscl` will be used to specify the size of the sinc interpolation kernel.

**sinscl** [float (Default Value = 1.0)] Size of the sinc interpolation kernel in pixels.

**stepsize** [int (Default Value = 10)] Number of pixels for WCS interpolation. The distortion model will be sampled exactly and completely every `stepsize` pixel with bi-linear interpolation being used to compute the distortion for intermediate pixels. This optimization speeds up the computation significantly when `stepsize`  $\gg$  1 at the expense of interpolation errors for intermediate pixels.

**addsky** [bool (Default Value = Yes)] Add back a sky value using the `MDRIZSKY` value from the header. If ‘Yes’ (True), the `blot_skyval` parameter is ignored.

**skyval** [float (Default Value = 0.0)] This is a user-specified custom sky value to be added to the blot image. This is only used if `blot_addsky` is ‘No’ (False).

**in\_units** [str{‘cps’, ‘counts’} (Default Value= ‘cps’)] Units of input (drizzled) image. Valid options are ‘cps’ and ‘counts’.

**out\_units** [str{‘cps’, ‘counts’} (Default Value = ‘counts’)] Units of the output (blotted) image. Valid options are ‘cps’ and ‘counts’.

**expkey** [str (Default Value = ‘exptime’)] Name of keyword to use to extract exposure time value, which will be used to scale the blotted image to the final output flux values when `out_units` is set to **counts**.

**expout** [str or float (Default Value = ‘input’)] Value of exposure time to use in scaling the output blotted image when `out_units` is set to **counts**. If set to ‘input’, the value will be read in from the input image header keyword specified by `expkey`.

---

**Note:** The following parameters, when set, will override any value determined

from `refimage` if a reference image was specified.

---

**outscale** [float,optional] Absolute size of output pixels in arcsec/pixel  
**orient** [float] Orientation of output (PA of Y axis, N through E)  
**raref** [float] RA of reference point on output image(CRVAL1,degrees)  
**decref** [float] Dec of reference point on output image (CRVAL2, degrees)  
**xrefpix** [float] Reference pixel X position on output (CRPIX1)  
**yrefpix** [float] Reference pixel Y position on output (CRPIX2)  
**outnx** [float] Size of output image's X-axis (pixels)  
**outny** [float] Size of output image's Y-axis (pixels)

## Notes

These tasks are designed to work together seamlessly when run in the full `AstroDrizzle` interface. More advanced users may wish to create specialized scripts for their own datasets, making use of only a subset of the predefined `AstroDrizzle` tasks, or add additional processing, which may be useful for their particular data. In these cases, individual access to the tasks is important.

Something to keep in mind is that the full `AstroDrizzle` interface will make backup copies of your original files and place them in the `Orig/` directory of your current working directory. If you are working with the stand alone interfaces, it is assumed that the user has already taken care of backing up their original datafiles as the input file will be directly altered.

## Examples

1. Basic example of how to call `blot()` yourself from a python command line, using the default parameter settings:

```
>>> from drizzlepac import ablot
>>> ablot.blot()
```

2. Creation of a blotted image from the products generated by running the `AstroDrizzle` task can be done for the median image “`adriz_aligned_wcs_f814w_med.fits`” to re-create the (SCI,1) chip from “`j8c0d1bwq_flc.fits`” using:

```
>>> from drizzlepac import ablot
>>> from stsci.tools import teal
>>> blotobj = teal.load('ablot') # get default values
>>> ablot.blot('adriz_aligned_wcs_f814w_med.fits', 'j8c0d1bwq_flc.
↳fits[sci,1]',
'aligned_f814w_sci1_blot.fits', configObj=blotobj)
```

or

```
>>> a = teal.teal('ablot')
# set data = adriz_aligned_wcs_f814w_med.fits
# set reference = j8c0d1bwq_flc.fits[sci,1]
# set outdata = aligned_f814w_sci1_blot.fits
```

`drizzlepac.ablot.runBlot` (*imageObjectList*, *output\_wcs*, *configObj={}*, *wcsm*  
*smap=wcs\_functions.WCSMap*, *procSteps=None*)

`drizzlepac.ablot.help` (*file=None*)

Print out syntax help for running astrodrizzle

#### Parameters

**file** [str (Default = None)] If given, write out help to the filename specified by this parameter Any previously existing file with this name will be deleted before writing out the help.

`drizzlepac.ablot.getHelpAsString` (*docstring=False*, *show\_ver=True*)

return useful help from a file in the script directory called `__taskname__.help`



---

## Cosmic-ray Identification Step

---

The cosmic rays and bad pixels are now identified by comparing the input images with the associated blotted, median-cleaned images created.

Mask blemishes in dithered data by comparison of an image with a model image and the derivative of the model image.

**Authors** Warren Hack

**License** *LICENSE*

`drizzlepac.drizCR.createCorrFile` (*outfile, arrlist, template*)

Create a `_cor` file with the same format as the original input image.

The DQ array will be replaced with the mask array used to create the `_cor` file.

`drizzlepac.drizCR.drizCR` (*input=None, configObj=None, editpars=False, \*\*inputDict*)

The blotted median images that are now transformed back into the original reference frame, get compared to the original input images to detect any spurious pixels (which may include pixels impacted by cosmic rays) in each input. Those spurious pixels then get flagged as ‘bad’ in the output cosmic ray mask files, which get used as input for the final combination so that they do not show up in the final product. The identified bad pixels get flagged by updating the input mask files. Optionally, copies of the original images with the bad pixels removed can be created through the use of the `driz_cr_corr` parameter.

### Parameters

**input** [str or list of str (Default = None)] A python list of blotted median image filenames, or just a single filename.

**configObj** [configObject (Default = None)] An instance of `configObject` which overrides default parameter settings.

**editpars** [bool (Default = False)] A parameter that allows user to edit input parameters by hand in the GUI. True to use the GUI to edit parameters.

**inputDict** [dict, optional] An optional list of parameters specified by the user, which can also be used to override the defaults.

### Other Parameters

**driz\_cr** [bool (Default = False)] Perform cosmic-ray detection? If set to `True`, cosmic-rays will be detected and used to create cosmic-ray masks based on the algorithms from `deriv` and `driz_cr`.

**driz\_cr\_corr** [bool (Default = False)] Create a cosmic-ray cleaned input image? If set to `True`, a cosmic-ray cleaned `_cor` image will be generated directly from the input image, and a corresponding `_crmask` file will be written to document detected pixels affected by cosmic-rays.

**driz\_cr\_snr** [list of floats (Default = '3.5 3.0')] The values for this parameter specify the signal-to-noise ratios for the `driz_cr` task to be used in detecting cosmic rays. See the help file for `driz_cr` for further discussion of this parameter.

**driz\_cr\_grow** [int (Default = 1)] The radius, in pixels, around each detected cosmic-ray, in which more stringent detection criteria for additional cosmic rays will be used.

**driz\_cr\_ctegrow** [int (Default = 0)] Length, in pixels, of the CTE tail that should be masked in the drizzled output.

**driz\_cr\_scale** [str (Default = '1.2 0.7')] Scaling factor applied to the derivative in `driz_cr` when detecting cosmic-rays. See the help file for `driz_cr` for further discussion of this parameter.

### Notes

These tasks are designed to work together seamlessly when run in the full `AstroDrizzle` interface. More advanced users may wish to create specialized scripts for their own datasets, making use of only a subset of the predefined `AstroDrizzle` tasks, or add additional processing, which may be useful for their particular data. In these cases, individual access to the tasks is important.

Something to keep in mind is that the full `AstroDrizzle` interface will make backup copies of your original files and place them in the `Orig/` directory of your current working directory. If you are working with the stand alone interfaces, it is assumed that the user has already taken care of backing up their original datafiles as the input file will be directly altered.

### Examples

Basic example of how to call `drizCR` yourself from a python command line using the default parameters for the task.

```
>>> from drizzlepac import drizCR
>>> drizCR.drizCR('*flt.fits')
```

`drizzlepac.drizCR.getHelpAsString` (*docstring=False, show\_ver=True*)  
Return useful help from a file in the script directory called `__taskname__.help`

`drizzlepac.drizCR.help` (*file=None*)  
Print out syntax help for running `astrodrizzle`

#### **Parameters**

**file** [str (Default = None)] If given, write out help to the filename specified by this parameter Any previously existing file with this name will be deleted before writing out the help.

`drizzlepac.drizCR.run` (*configObj*)

`drizzlepac.drizCR.rundrizCR` (*imgObjList, configObj, procSteps=None*)

`drizzlepac.drizCR.setDefault`s (*configObj={}*)  
Return a dictionary of the default parameters which also been updated with the user overrides.



Various utilities get used by `MultiDrizzle`, including some to handle WCS interpretation, trailer file generation, output file generation and interpretation of the `MDRIZTAB` reference file.

### 10.1 Utility Functions

These functions perform various small operations within `MultiDrizzle`.

A library of utility functions

**Authors** Warren Hack

**License** *LICENSE*

**class** `drizzlepac.util.ProcSteps`

This class allows `MultiDrizzle` to keep track of the start and end times of each processing step that gets run as well as computing/reporting the elapsed time for each step.

The code for each processing step must call the `'addStep()'` method to initialize the information for that step, then the `'endStep()'` method to record the end and elapsed times.

The `'reportTimes()'` method can then be used to provide a summary of all the elapsed times and total run time.

**addStep** (*key*)

Add information about a new step to the dict of steps The value `'ptime'` is the output from `'_ptime()'` containing both the formatted and unformatted time for the start of the step.

**endStep** (*key*)

Record the end time for the step.

If `key==None`, simply record `ptime` as end time for class to represent the overall runtime since the initialization of the class.

**reportTimes ()**

Print out a formatted summary of the elapsed times for all the performed steps.

**class** `drizzlepac.util.WithLogging`

`drizzlepac.util.applyUserPars_steps (configObj, input_dict, step='3a')`

Apply logic to turn on use of user-specified output WCS if user provides any parameter on command-line regardless of how `final_wcs` was set.

`drizzlepac.util.atfile_ivm (filename)`

Return the filename of the IVM file which is assumed to be the second word in the atfile the user gave.

`drizzlepac.util.atfile_sci (filename)`

Return the filename of the science image which is assumed to be the first word in the atfile the user gave.

`drizzlepac.util.base_taskname (taskname, packagename=None)`

Extract the base name of the task.

Many tasks in the `drizzlepac` have “compound” names such as ‘`drizzlepac.sky`’. This function will search for the presence of a dot in the input `taskname` and if found, it will return the string to the right of the right-most dot. If a dot is not found, it will return the input string.

### Parameters

**taskname** [str, None] Full task name. If it is `None`, `base_taskname ()` will return `None`.

**packagename** [str, None (Default = None)] Package name. It is assumed that a compound task name is formed by concatenating `packagename + '.' + taskname`. If `packagename` is not `None`, `base_taskname ()` will check that the string to the left of the right-most dot matches `packagename` and will raise an `AssertionError` if the package name derived from the input `taskname` does not match the supplied `packagename`. This is intended as a check for discrepancies that may arise during the development of the tasks. If `packagename` is `None`, no such check will be performed.

### Raises

**AssertionError** Raised when package name derived from the input `taskname` does not match the supplied `packagename`

`drizzlepac.util.check_blank (cvar)`

Converts blank value (from `configObj`?) into a value of `None`.

`drizzlepac.util.computeRange (corners)`

Determine the range spanned by an array of pixel positions.

`drizzlepac.util.compute_texptime (imageObjectList)`

Add up the exposure time for all the members in the pattern, since ‘`drizzle`’ doesn’t have the necessary information to correctly set this itself.

`drizzlepac.util.countImages (imageObjectList)`

`drizzlepac.util.count_sci_extensions (filename)`

Return the number of SCI extensions and the EXTNAME from a input MEF file.

`drizzlepac.util.createFile (dataArray=None, outfile=None, header=None)`

Create a simple fits file for the given data array and header. Returns either the FITS object in-membory when outfile==None or None when the FITS file was written out to a file.

`drizzlepac.util.displayBadRefimageWarningBox (display=True, parent=None)`

Displays a warning box for the 'input' parameter.

`drizzlepac.util.displayEmptyInputWarningBox (display=True, parent=None)`

Displays a warning box for the 'input' parameter.

`drizzlepac.util.displayMakewcsWarningBox (display=True, parent=None)`

Displays a warning box for the 'makewcs' parameter.

`drizzlepac.util.end_logging (filename=None)`

Close log file and restore system defaults.

`drizzlepac.util.findWCSExtn (filename)`

Return new filename with extension that points to an extension with a valid WCS.

#### Returns

**extnum** [str, None] Value of extension name as a string either as provided by the user or based on the extension number for the first extension which contains a valid HSTWCS object. Returns None if no extension can be found with a valid WCS.

#### Notes

The return value from this function can be used as input to create another HSTWCS with the syntax:

```
`HSTWCS ('{}[{}]' .format (filename, extnum) )
```

`drizzlepac.util.findrootname (filename)`

Return the rootname of the given file.

`drizzlepac.util.getConfigObjPar (configObj, parname)`

Return parameter value without having to specify which section holds the parameter.

`drizzlepac.util.getDefaultConfigObj (taskname, configObj, input_dict={}, load-Only=True)`

Return default configObj instance for task updated with user-specified values from input\_dict.

#### Parameters

**taskname** [string] Name of task to load into TEAL

**configObj** [string] The valid values for 'configObj' would be:

<b>None</b>	- loads last saved user .cfg file
'defaults'	- loads task default .cfg file
name of .cfg file (string)	- loads user-specified .cfg file

**input\_dict** [dict] Set of parameters and values specified by user to be different from what gets loaded in from the .cfg file for the task

**loadOnly** [bool] Setting 'loadOnly' to False causes the TEAL GUI to start allowing the user to edit the values further and then run the task if desired.

`drizzlepac.util.getFullParList (configObj)`

Return a single list of all parameter names included in the configObj regardless of which section the parameter was stored

`drizzlepac.util.getRotatedSize (corners, angle)`

Determine the size of a rotated (meta)image.

`drizzlepac.util.getSectionName (configObj, stepnum)`

Return section label based on step number.

`drizzlepac.util.get_detnum (hstwcs, filename, extnum)`

`drizzlepac.util.get_expstart (header, primary_hdr)`

shouldn't this just be defined in the instrument subclass of imageobject?

`drizzlepac.util.get_pool_size (usr_config_value, num_tasks)`

Determine size of thread/process-pool for parallel processing. This examines the `cpu_count` to decide and return the right pool size to use. Also take into account the user's wishes via the config object value, if specified. On top of that, don't allow the pool size returned to be any higher than the number of parallel tasks, if specified. Only use what we need (`mp.Pool` starts `pool_size` processes, needed or not). If number of tasks is unknown, call this with "num\_tasks" set to `None`. Returns 1 when indicating that parallel processing should not be used. Consolidate all such logic here, not in the caller.

`drizzlepac.util.init_logging (logfile='astrodrizzle.log', default=None, level=20)`

Set up logger for capturing stdout/stderr messages.

Must be called prior to writing any messages that you want to log.

`drizzlepac.util.isASNTable (inputFilelist)`

Return TRUE if inputFilelist is a fits ASN file.

`drizzlepac.util.isCommaList (inputFilelist)`

Return True if the input is a comma separated list of names.

`drizzlepac.util.is_blank (val)`

Determines whether or not a value is considered 'blank'.

`drizzlepac.util.loadFileList (inputFilelist)`

Open up the '@ file' and read in the science and possible ivm filenames from the first two columns.

`drizzlepac.util.parse_colnames (colnames, coords=None)`

Convert colnames input into list of column numbers.

`drizzlepac.util.printParams (paramDictionary, all=False, log=None)`

Print nicely the parameters from the dictionary.

`drizzlepac.util.print_cfg (cfg, logfn=None)`

`drizzlepac.util.print_key (key, val, lev=0, logfn=<built-in function print>)`



`drizzlepac.util.print_pkg_versions` (*packages=None, git=False, svn=False, log=None*)

`drizzlepac.util.readCommaList` (*fileList*)  
Return a list of the files with the commas removed.

`drizzlepac.util.readcols` (*infile, cols=[0, 1, 2, 3], hms=False*)  
Read the columns from an ASCII file as numpy arrays.

#### Parameters

**infile** [str] Filename of ASCII file with array data as columns.

**cols** [list of int] List of 0-indexed column numbers for columns to be turned into numpy arrays (DEFAULT- [0,1,2,3]).

#### Returns

**outarr** [list of numpy arrays] Simple list of numpy arrays in the order as specified in the 'cols' parameter.

`drizzlepac.util.removeFileSafely` (*filename, clobber=True*)  
Delete the file specified, but only if it exists and clobber is True.

`drizzlepac.util.runmakewcs` (*input*)  
Runs 'updatewcs' to recompute the WCS keywords for the input image.

#### Parameters

**input** [list of str] A list of file names.

#### Returns

**output** [list of str] Returns a list of names of the modified files (For GEIS files returns the translated names).

`drizzlepac.util.updateNEXTENDKw` (*fobj*)  
Update NEXTEND keyword in PRIMARY header (if present) to accurately reflect the number of extensions in the MEF file.

`drizzlepac.util.update_input` (*filelist, ivmlist=None, removed\_files=None*)  
Removes files flagged to be removed from the input filelist. Removes the corresponding ivm files if present.

`drizzlepac.util.validateUserPars` (*configObj, input\_dict*)  
Compares input parameter names specified by user with those already recognized by the task.

Any parameters provided by the user that does not match a known task parameter will be reported and a ValueError exception will be raised.

`drizzlepac.util.verifyFilePermissions` (*filelist, chmod=True*)  
Verify that images specified in 'filelist' can be updated.

A message will be printed reporting the names of any images which do not have write-permission, then quit.

`drizzlepac.util.verifyRefimage` (*refimage*)  
Verify that the value of refimage specified by the user points to an extension with a proper WCS

defined. It starts by making sure an extension gets specified by the user when using a MEF file. The final check comes by looking for a CD matrix in the WCS object itself. If either test fails, it returns a value of False.

`drizzlepac.util.verifyUniqueWcsname (fname, wcsname, include_primary=True)`  
Report whether or not the specified WCSNAME already exists in the file

`drizzlepac.util.verifyUpdatewcs (fname)`  
Verify the existence of WCSNAME in the file. If it is not present, report this to the user and raise an exception. Returns True if WCSNAME was found in all SCI extensions.

## 10.2 WCS Utilities

These functions read and interpret the WCS information from input images and create the output WCS objects based on STWCS routines.

**Authors** Warren Hack

**License** *LICENSE*

**class** `drizzlepac.wcs_functions.IdentityMap (input, output)`

**forward** (*pixx, pixy*)

**class** `drizzlepac.wcs_functions.LinearMap (xsh=0.0, ysh=0.0, rot=0.0, scale=1.0)`

**forward** (*pixx, pixy*)

**class** `drizzlepac.wcs_functions.WCSMap (input, output, origin=1)`

Sample class to demonstrate how to define a coordinate transformation

**backward** (*pixx, pixy*)

Transform *pixx, pixy* positions from the output frame back onto their original positions in the input frame.

**checkWCS** (*obj, name*)

**forward** (*pixx, pixy*)

Transform the input *pixx, pixy* positions in the input frame to pixel positions in the output frame.

This method gets passed to the drizzle algorithm.

**get\_pix\_ratio** ()

Return the ratio of plate scales between the input and output WCS. This is used to properly distribute the flux in each pixel in ‘tdriz’.

**rd2xy** (*wcs, ra, dec*)

Transform input sky positions into pixel positions in the WCS provided.

**xy2rd** (*wcs, pixx, pixy*)

Transform input pixel positions into sky positions in the WCS provided.

`drizzlepac.wcs_functions.apply_fitlin (data, P, Q)`

`drizzlepac.wcs_functions.build_hstwcs` (*crval1, crval2, crpix1, crpix2, naxis1, naxis2, pscale, orientat*)

Create an HSTWCS object for a default instrument without distortion based on user provided parameter values.

`drizzlepac.wcs_functions.build_pixel_transform` (*chip, output\_wcs*)

`drizzlepac.wcs_functions.calcNewEdges` (*wcs, shape*)

This method will compute sky coordinates for all the pixels around the edge of an image AFTER applying the geometry model.

#### Parameters

**wcs** [obj] HSTWCS object for image

**shape** [tuple] numpy shape tuple for size of image

#### Returns

**border** [arr] array which contains the new positions for all pixels around the border of the edges in alpha,dec

`drizzlepac.wcs_functions.computeEdgesCenter` (*edges*)

`drizzlepac.wcs_functions.convertWCS` (*inwcs, drizwcs*)

Copy WCSObject WCS into Drizzle compatible array.

`drizzlepac.wcs_functions.createWCSObject` (*output, default\_wcs, imageObjectList*)

Converts a astropy.wcs WCS object into a WCSObject(baseImageObject) instance.

`drizzlepac.wcs_functions.create_CD` (*orient, scale, cx=None, cy=None*)

Create a (un?)distorted CD matrix from the basic inputs.

The 'cx' and 'cy' parameters, if given, provide the X and Y coefficients of the distortion as returned by reading the IDCTAB. Only the first 2 elements are used and should correspond to the 'OC[X/Y]10' and 'OC[X/Y]11' terms in that order as read from the expanded SIP headers.

The units of 'scale' should be 'arcseconds/pixel' of the reference pixel. The value of 'orient' should be the absolute orientation on the sky of the reference pixel.

`drizzlepac.wcs_functions.create_mosaic_pars` (*mosaic\_wcs*)

Return dict of values to use with AstroDrizzle to specify output mosaic

#### Parameters

**mosaic\_wcs** [object] WCS as generated by make\_mosaic\_wcs

#### Returns

**mosaic\_pars** [dict] This dictionary can be used as input to astrodrizzle.AstroDrizzle with the syntax 'AstroDrizzle(filename, \*\*mosaic\_pars)'

`drizzlepac.wcs_functions.ddtohms` (*xsky, ysky, verbose=False, precision=6*)

Convert sky position(s) from decimal degrees to HMS format.

`drizzlepac.wcs_functions.fitlin` (*imgarr, refarr*)

Compute the least-squares fit between two arrays. A Python translation of 'FITLIN' from 'drutil.f' (Drizzle V2.9).

`drizzlepac.wcs_functions.fitlin_clipped(xy, uv, verbose=False, mode='rscale',  
nclip=3, reject=3)`

Perform a clipped fit based on the number of iterations and rejection limit (in sigma) specified by the user. This will more closely replicate the results obtained by 'geomap' using 'maxiter' and 'reject' parameters.

`drizzlepac.wcs_functions.fitlin_rscales(xy, uv, verbose=False)`

Performs a linear, orthogonal fit between matched lists of positions 'xy' (input) and 'uv' (output).

Output: (same as for `fit_arrays_general`)

`drizzlepac.wcs_functions.get_extns(fimg, extname='SCI')`

**Examines the specified fits image and returns the extension number(s) whose name(s) match the specified by input value.**

#### Parameters

**fimg** [string] Name of the image to probe

**extname** [string] Extension name to search for in specified image. Default value is 'SCI'.

#### Returns

**extns** [list] List of matching extension numbers

`drizzlepac.wcs_functions.get_hstwcs(filename, hdulist, extnum)`

Return the HSTWCS object for a given chip.

`drizzlepac.wcs_functions.get_pix_ratio_from_WCS(input, output)`

[Functional form of `.get_pix_ratio()` method of `WCSTMap`]

`drizzlepac.wcs_functions.make_mosaic_wcs(filenamees, rot=None, scale=None)`

Combine WCSs from all input files into a single meta-WCS

#### Parameters

**filenamees** [list] list of filenames to process

**rot** [float] desired rotation of meta-WCS. Default value is None.

**scale** [float] desired scale of meta-WCS. Default value is None.

#### Returns

**mosaic\_wcs** [HSTWCS object] the merged composite WCS

`drizzlepac.wcs_functions.make_outputwcs(imageObjectList, output, con-  
figObj=None, perfect=False)`

Computes the full output WCS based on the set of input imageObjects provided as input, along with the pre-determined output name from `process_input`. The user specified output parameters are then used to modify the default WCS to produce the final desired output frame. The input `imageObjectList` has the `outputValues` dictionary updated with the information from the computed output WCS. It then returns this WCS as a `WCSTObject(imageObject)` instance.

`drizzlepac.wcs_functions.make_perfect_cd(wcs)`

Create a perfect (square, orthogonal, undistorted) CD matrix from the input WCS.

`drizzlepac.wcs_functions.mergeWCS(default_wcs, user_pars)`

Merges the user specified WCS values given as dictionary derived from the input configObj object with the output astropy.wcs object computed using `distortion.output_wcs()`.

The `user_pars` dictionary needs to have the following set of keys:

```
user_pars = {'ra':None, 'dec':None, 'scale':None, 'rot':None,
            'outnx':None, 'outny':None, 'crpix1':None, 'crpix2':None}
```

`drizzlepac.wcs_functions.readAltWCS(fobj, ext, wcskey=' ', verbose=False)`

Reads in alternate primary WCS from specified extension.

#### Parameters

**fobj** [str, `astropy.io.fits.HDUList`] fits filename or fits file object containing alternate/primary WCS(s) to be converted

**wcskey** [str] [” “,A-Z] alternate/primary WCS key that will be replaced by the new key

**ext** [int] fits extension number

#### Returns

\_\_\_\_\_

**hdr:** `fits.Header` header object with ONLY the keywords for specified alternate WCS

`drizzlepac.wcs_functions.removeAllAltWCS(hdulist, extlist)`

Removes all alternate WCS solutions from the header

`drizzlepac.wcs_functions.restoreDefaultWCS(imageObjectList, output_wcs)`

Restore WCS information to default values, and update imageObject accordingly.

`drizzlepac.wcs_functions.updateImageWCS(imageObjectList, output_wcs)`

`drizzlepac.wcs_functions.updateWCS(drizwcs, inwcs)`

Copy output WCS array from Drizzle into WCSObject.

`drizzlepac.wcs_functions.update_linCD(cdmat, delta_rot=0.0, delta_scale=1.0, cx=[0.0, 1.0], cy=[1.0, 0.0])`

Modify an existing linear CD matrix with rotation and/or scale changes and return a new CD matrix. If ‘cx’ and ‘cy’ are specified, it will return a distorted CD matrix.

Only those terms which are varying need to be specified on input.

`drizzlepac.wcs_functions.wcsfit(img_wcs, ref_wcs)`

Perform a linear fit between 2 WCS for shift, rotation and scale. Based on the WCSLIN function from ‘drutil.f’(Drizzle V2.9) and modified to allow for differences in reference positions assumed by PyDrizzle’s distortion model and the coeffs used by ‘drizzle’.

#### Parameters

**img** [obj] `ObsGeometry` instance for input image

`ref_wcs` [obj] Undistorted WCSObject instance for output frame

## 10.3 Output Image Generation

This module manages the creation of the output image FITS file.

**Authors** Warren Hack

**License** *LICENSE*

**class** `drizzlepac.outputimage.OutputImage` (*plist*, *input\_pars*, *build=True*,  
*wcs=None*, *single=False*, *blot=False*)

This class manages the creation of the array objects which will be used by Drizzle. The three arrays, SCI/WHT/CTX, will be setup either as extensions in a single multi-extension FITS file, or as separate FITS files.

The object 'plist' must contain at least the following members:

```
plist['output'] - name of output FITS image (for SCI)
plist['outnx']  - size of X axis for output array
plist['outny']  - size of Y axis for output array
```

If 'single=yes', then 'plist' also needs to contain:

```
plist['outsingle']
plist['outsweight']
plist['outscontext']
```

If 'blot=yes', then 'plist' also needs:

```
plist['data']
plist['blotImage']
plist['blotnx'],plist['blotny']
```

If 'build' is set to 'no', then each extension/array must be in separate FITS objects. This would also require:

```
plist['outdata'] - name of output SCI FITS image
plist['outweight'] - name of output WHT FITS image
plist['outcontext'] - name of output CTX FITS image
```

Optionally, the overall exposure time information can be passed as:

```
plist['texptime'] - total exptime for output
plist['expstart'] - start time of combined exposure
plist['expend']   - end time of combined exposure
```

**addDrizKeywords** (*hdr*, *versions*)

Add drizzle parameter keywords to header.

**find\_kwupdate\_location** (*hdr*, *keyword*)

Find the last keyword in the output header that comes before the new keyword in the original, full

input headers. This will rely on the original ordering of keywords from the original input files in order to place the updated keyword in the correct location in case the keyword was removed from the output header prior to calling this method.

**set\_bunit** (*bunit*)

Method used to update the value of the bunit attribute.

**set\_units** (*units*)

Method used to record what units were specified by the user for the output product.

**writeFITS** (*template*, *sciarr*, *wharr*, *ctxarr=None*, *versions=None*, *overwrite=True*,  
*blend=True*, *virtual=False*, *rules\_file=None*)

Generate PyFITS objects for each output extension using the file given by ‘template’ for populating headers.

The arrays will have the size specified by ‘shape’.

## 10.4 MultiDrizzle Reference Table

This module supports the interpretation of the MDRIZTAB for processing as used in the pipeline.

This module supports the interpretation of the MDRIZTAB for processing as used in the pipeline.

**Authors** Warren Hack, Ivo Busko, Christopher Hanley

**License** *LICENSE*

`drizzlepac.mdzhandler.cleanBlank` (*value*)

`drizzlepac.mdzhandler.cleanInt` (*value*)

`drizzlepac.mdzhandler.cleanNaN` (*value*)

`drizzlepac.mdzhandler.findFormat` (*format*)

`drizzlepac.mdzhandler.getMdriztabParameters` (*files*)

Gets entry in MDRIZTAB where task parameters live. This method returns a record array mapping the selected row.

`drizzlepac.mdzhandler.toBoolean` (*flag*)





*tweakback* - propagate the “tweaked” solutions back to the original input files.

Version 0.4.0 - replaced previous algorithm that used fitting of WCS footprints to reconstruct the transformation that was applied to the old drizzled image (to align it with another image) to obtain the new drizzled image WCS with an algorithm that is based on linearization of the exact compound operator that transforms current image coordinates to the “aligned” (to the new drizzled WCS) image coordinates.

**Authors** Warren Hack, Mihai Cara

**License** *LICENSE*

`drizzlepac.tweakback.determine_extnum` (*drzfile*, *extname='SCI'*)

`drizzlepac.tweakback.determine_orig_wcsname` (*header*, *wnames*, *wkeys*)  
Determine the name of the original, unmodified WCS solution

`drizzlepac.tweakback.extract_input_filenames` (*drzfile*)  
Generate a list of filenames from a drizzled image’s header

`drizzlepac.tweakback.getHelpAsString` (*docstring=False*, *show\_ver=True*)  
return useful help from a file in the script directory called `__taskname__.help`

`drizzlepac.tweakback.help` (*file=None*)  
Print out syntax help for running `astrodrizzle`

### Parameters

**file** [str (Default = None)] If given, write out help to the filename specified by this parameter Any previously existing file with this name will be deleted before writing out the help.

`drizzlepac.tweakback.linearize` (*wcsim*, *wcsima*, *wcs\_olddrz*, *wcs\_newdrz*, *imcrpix*,  
*hx=1.0*, *hy=1.0*)

`drizzlepac.tweakback.run` (*configobj*)

`drizzlepac.tweakback.tweakback` (*drzfile, input=None, origwcs=None, newname=None, wcsname=None, extname='SCI', force=False, verbose=False*)

Apply WCS solution recorded in drizzled file to distorted input images (`_flt.fits` files) used to create the drizzled file. This task relies on the original WCS and updated WCS to be recorded in the drizzled image's header as the last 2 alternate WCSs.

### Parameters

**drzfile** [str (Default = '')] filename of undistorted image which contains the new WCS and WCS prior to being updated

**newname** [str (Default = None)] Value of WCSNAME to be used to label the updated solution in the output (eq., `_flt.fits`) files. If left blank or None, it will default to using the current WCSNAME value from the input drzfile.

**input** [str (Default = '')] filenames of distorted images to be updated using new WCS from 'drzfile'. These can be provided either as an `@-file`, a comma-separated list of filenames or using wildcards.

---

**Note:** A blank value will indicate that the task should derive the filenames from the 'drzfile' itself, if possible. The filenames will be derived from the `D*DATA` keywords written out by `AstroDrizzle`. If they can not be found, the task will quit.

---

**origwcs** [str (Default = None)] Value of WCSNAME keyword prior to the drzfile image being updated by `TweakReg`. If left blank or None, it will default to using the second to last `WCSNAME*` keyword value found in the header.

**wcsname** [str (Default = None)] Value of WCSNAME for updated solution written out by `TweakReg` as specified by the `wcsname` parameter from `TweakReg`. If this is left blank or `None`, it will default to the current WCSNAME value from the input drzfile.

**extname** [str (Default = 'SCI')] Name of extension in `input` files to be updated with new WCS

**force** [bool (Default = False)] This parameters specified whether or not to force an update of the WCS even though WCS already exists with this solution or `wcsname`?

**verbose** [bool (Default = False)] This parameter specifies whether or not to print out additional messages during processing.

See also:

`stwcs.wcsutil.altwcs` Alternate WCS implementation

## Notes

The algorithm used by this function is based on linearization of the exact compound operator that converts input image coordinates to the coordinates (in the input image) that would result in alignment with the new drizzled image WCS.

If no input distorted files are specified as input, this task will attempt to generate the list of filenames from the drizzled input file's own header.

## Examples

An image named `acswfc_mos2_drz.fits` was created from 4 images using `astrodrizzle`. This drizzled image was then aligned to another image using `tweakreg` and the header was updated using the `WCSNAME = TWEAK_DRZ`. The new WCS can then be used to update each of the 4 images that were combined to make up this drizzled image using:

```
>>> from drizzlepac import tweakback
>>> tweakback.tweakback('acswfc_mos2_drz.fits')
```

If the same WCS should be applied to a specific set of images, those images can be updated using:

```
>>> tweakback.tweakback('acswfc_mos2_drz.fits',
...                       input='img_mos2aflt.fits,img_mos2eflt.fits')
```

```
drizzlepac.tweakback.update_chip_wcs(chip_wcs, drz_old_wcs, drz_new_wcs,
                                     xrms=None, yrms=None)
```

```
drizzlepac.tweakback.tweakback(drzfile, input=None, origwcs=None, newname=None,
                               wcsname=None, extname='SCI', force=False, ver-
                              bose=False)
```

Apply WCS solution recorded in drizzled file to distorted input images (`_flt.fits` files) used to create the drizzled file. This task relies on the original WCS and updated WCS to be recorded in the drizzled image's header as the last 2 alternate WCSs.

### Parameters

**drzfile** [str (Default = "")] filename of undistorted image which contains the new WCS and WCS prior to being updated

**newname** [str (Default = None)] Value of `WCSNAME` to be used to label the updated solution in the output (eq., `_flt.fits`) files. If left blank or `None`, it will default to using the current `WCSNAME` value from the input `drzfile`.

**input** [str (Default = "")] filenames of distorted images to be updated using new WCS from 'drzfile'. These can be provided either as an `@-file`, a comma-separated list of filenames or using wildcards.

---

**Note:** A blank value will indicate that the task should derive the filenames from the 'drzfile' itself, if possible. The filenames will be derived from the `D*DATA`

keywords written out by `AstroDrizzle`. If they can not be found, the task will quit.

**origwcs** [str (Default = None)] Value of `WCSNAME` keyword prior to the drzfile image being updated by `TweakReg`. If left blank or `None`, it will default to using the second to last `WCSNAME*` keyword value found in the header.

**wcsname** [str (Default = None)] Value of `WCSNAME` for updated solution written out by `TweakReg` as specified by the `wcsname` parameter from `TweakReg`. If this is left blank or `None`, it will default to the current `WCSNAME` value from the input drzfile.

**extname** [str (Default = 'SCI')] Name of extension in `input` files to be updated with new WCS

**force** [bool (Default = False)] This parameters specified whether or not to force an update of the WCS even though WCS already exists with this solution or `wcsname`?

**verbose** [bool (Default = False)] This parameter specifies whether or not to print out additional messages during processing.

See also:

**stwcs.wcsutil.altwcs** Alternate WCS implementation

## Notes

The algorithm used by this function is based on linearization of the exact compound operator that converts input image coordinates to the coordinates (in the input image) that would result in alignment with the new drizzled image WCS.

If no input distorted files are specified as input, this task will attempt to generate the list of filenames from the drizzled input file's own header.

## Examples

An image named `acswfc_mos2_drz.fits` was created from 4 images using `astrodrizzle`. This drizzled image was then aligned to another image using `tweakreg` and the header was updated using the `WCSNAME = TWEAK_DRZ`. The new WCS can then be used to update each of the 4 images that were combined to make up this drizzled image using:

```
>>> from drizzlepac import tweakback
>>> tweakback.tweakback('acswfc_mos2_drz.fits')
```

If the same WCS should be applied to a specific set of images, those images can be updated using:

```
>>> tweakback.tweakback('acswfc_mos2_drz.fits',
...                       input='img_mos2aflt.fits,img_mos2eflt.fits')
```

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## DrizzlePac Release Notes

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The code for this package gets released through a number of methods: namely, the use of the package for pipeline and archive processing of ACS and WFC3 data, SSB's semi-annual public release of the stsci\_python package, and a weekly beta release of the development version. The following notes provide some details on what has been revised for each version.

### 13.1 DrizzlePac Release Notes

The version of DrizzlePac can be identified using

```
> python
>>> import drizzlepac
>>> drizzlepac.__version__
```

The following notes provide some details on what has been revised for each version in reverse chronological order (most recent version at the top of the list).

#### 13.1.1 3.1.8 (Aug-2020)

A number of changes have been implemented to either correct problems or improve the processed results. The most significant of the changes are:

- rscale only used for alignment.
- a minimum of 6 sources now gets used for alignment
- no proper motions used in astrometric (GAIA) catalog when attempting a posteriori fitting
- chip-to-chip alignment errors were corrected

In addition to a few dozen bug fixes, the following updates to the algorithms were also implemented.

- Simplified the logic in `tweakreg` for deciding how to archive primary WCS resulting in a reduction of duplicate WCSes in image headers. [#715]
- Added polynomial look-up table distortion keywords to the list of distortion keywords used by `outputimage.deleteDistortionKeywords` so that distortions can be removed from ACS images that use `NPOLFILE`. This now allows removal of alternate WCS from blotted image headers. [#709]
- Added `rules_file` parameter to `AstroDrizzle` to enable use of custom files in pipeline processing. [#674]
- Only apply solutions from the astrometry database which were non-aposteriori WCS solutions as the PRIMARY WCS. This allows the pipeline to compare the true apriori WCS solutions (e.g., GSC or HSC WCSs) to aposteriori solutions computed using the latest distortion-models and alignment algorithms being used at the time of processing. [#669]
- Verification using a similarity index gets reported in the trailer file and does not get used as a Pass/Fail criteria for alignment. [#619]
- If verification fails for either pipeline-default or apriori solution, reset cosmic-ray(CR) flag (4096) in DQ arrays. This will allow subsequent attempt to align the images to not be impacted by potentially mis-identified CRs that most likely blanked out real sources in the field. As a result, the image alignment process became more robust when computing the aposteriori alignment. [#614]
- Fix a crash in `tweakreg` when finding sources in very large images due to a bug in `scipy.signal.convolve2d`. [#670]
- Fix a bug in `tweakreg` due to which the number of matched sources needed to be *strictly* greater than `minobj`. Now the minimum number of matched sources must be *at least* equal or greater than `minobj`. [#604]
- Fix a crash in `tweakreg` when `2dhist` is enabled and `numpy` version is `1.18.1` and later. [#583, #587]
- Update calibrated (FLC/FLT) files with RMS and NMATCH keywords when it successfully aligns the data to GAIA using the a posteriori fit. Headerlet files for this fit which already have these keywords are now retained and provided as the final output headerlets as well. [#555]
- Insure HDRNAME keyword gets added to successfully aligned FLC/FLT files. [#580]
- Fix problem with ‘tweakback’ task when trying to work with updated WCS names. [#551]
- Fix problems found in processing data with `NGOODPIX==0`, DRC files not getting generated for singletons, alignment trying to use a source too near the chip edge, catch the case were all inputs have zero exposure time, lazily remove alignment sub-directories, fixed a bug in overlap computation that showed up in oblong mosaics, recast an input to `histogram2d` as `int`, defined default values for tables when no sources were found. [#593]
- Updated to be compatible with `tweakwcs v0.6.0` to correct chip-to-chip alignment issues in aposteriori WCS solutions. [#596]
- Correctly define output drizzle product filename during pipeline processing for exposures with ‘drz’ in the rootname. [#523]



- Implement multiple levels of verification for the drizzle products generated during pipeline processing (using `runastrodriz`); including overlapp difference computations [#520], and magnitude correlation [#512].
- Replace `alignimages` module with O-O based `align` [#512]
- Fix problem with NaNs when looking for sources to use for aligning images [#512]
- Fixed code that selected the brightest sources to use for alignment allowing alignment to work (more often) for images with saturated sources. [#512]
- Use logic for defining the PSF extracted from the images to shrink it in each axis by one-half for images of crowded fields to allow for more sources to be extracted by `daofind`-like algorithm. This enables source finding and alignment to work more reliably on crowded field images. [#512]
- Insure all input files, especially those with zero exposure time or grism images, get updated with the latest pipeline calibration for the distortion. [ #495]

This version also relies on updates in the following packages to get correctly aligned and combined images with correctly specified WCS keywords:

- TWEAKWCS 0.6.4: This version corrects problems with the chip-to-chip separation that arose when applying a single fit solution to the entire observation.
- STWCS 1.5.4: This version implements a couple of fixes to insure that use of headerlets defines the full correct set of keywords from the headerlet for the PRIMARY WCS in the science exposure without introducing multiple copies of some keywords.
- Numpy 1.18: Changes in numpy data type definitions affected some of the code used for computing the offset between images when performing a posteriori alignment during pipeline processing and when running the ‘`tweakreg`’ task.

### **13.1.2 3.1.3 (5-Dec-2019)**

- Fixed a bug in the `updatehdr.update_from_shiftfile()` function that would crash while reading shift files. [#448]
- Migration of the HAP portion of the package to an object-oriented implementation. [#427]
- Added support for providing HSTWCS object as input to ‘`final_refimage`’ or ‘`single_refimage`’ parameter. [#426]
- Implementation of grid definition interface to support returning `SkyCell` objects that overlap a mosaic footprint. [#425]
- Complete rewrite of `runastrodriz` for pipeline processing to include multi-level verification of alignment. [#440]

### **13.1.3 3.0.2 (15-Jul-2019)**

- Removed deprecated parameter `coords` from the parameter list of `pixtopix.tran()` function. [#406]

- Modified the behavior of the `verbose` parameter in `pixtopix.tran()` to not print coordinates when not run as a script and when `output` is `None`. [#406]
- Fixed a compatibility issue in `tweakutils` that would result in crash in `skypix` when converting coordinates in `hms` format. [#385]
- Fixed a bug in the `astrodrizzle.sky` module due to which sky matching fails with “Keyword ‘MDRIZSKY’ not found” error when some of the input images do not overlap at all with the other images. [#380]
- Fixed a bug in the `util.WithLogging` decorator due to which incorrect log file was reported when user-supplied log file name does not have `.log` extension. [#365]
- Fixed a bug introduced in #364 returning in `finally` block. [#365]
- Improved `util.WithLogging` decorator to handle functions that return values. [#364]
- Fixed a bug in the automatic computation of the IVM weights when IVM was not provided by the user. [#320]
- Fixed a bug in the 2D histogram code used for estimating shifts for catalog pre-matching. This may result in better matching. [#286]
- Now `tolerance` (in `tweakreg`) is no longer ignored when `use2dhist` is enabled. [#286]
- Fixed VS compiler errors with pointer arithmetic on void pointers. [#273]
- Fix logic so that code no longer tries to update headers when no valid fit could be determined. [#241]
- Fixed a bug in the computation of interpolated large scale flat field for STIS data. The bug was inconsequential in practice. Removed the dependency on `stsci.imagemanip` package. [#227]
- Removed the dependency on `stsci.ndimage` (using `scipy` routines instead). [#225]
- Added 'Advanced Pipeline Products' alignment code to `drizzlepac` package. Enhance `runastrodriz` to compute and apply absolute astrometric corrections to GAIA (or related) frame to images where possible. [#200, #213, #216, #223, #234, #235, #244, #248, #249, #250, #251, #259, #260, #268, #271, #283, #294, #302]
- Add computation and reporting of the fit’s **Root-Mean-Square Error (RMSE)** and **Mean Absolute Error (MAE)**. [#210]
- Replaced the use of `WCS._naxis1` and `WCS._naxis2` with `WCS.pixel_shape` [#207]
- Removed support for Python 2. Only versions `>= 3.5` are supported. [#207]
- Use a more numerically stable `numpy.linalg.inv` instead of own matrix inversion. [#205]
- The intermediate fit match catalog, with the name `_catalog_fit.match` generated by `tweakreg` now has correct RA and DEC values for the sources after applying the fit. [#200, #202]
- Simplify logic for determining the chip ID for each source. [#200]

### 13.1.4 2.2.6 (02-Nov-2018)

- Fix a bug that results in `tweakreg` crashing when no sources are found with user-specified source-finding parameters and when `tweakreg` then attempts to find sources using default parameters. [#181]
- Updated `unit_tests` to use original inputs, rather than updated inputs used by nightly regression tests.
- Fix `numpy` “floating” deprecation warnings. [#175]
- Fix incorrect units in CR-cleaned images created by `astrodrizzle`. Now CR-cleaned images should have the same units as input images. [#190]

### 13.1.5 2.2.5 (14-Aug-2018)

- Changed the color scheme of the `hist2d` plots to `viridis`. [#167]
- Refactored test suite
- `sdist` now packages C extension source code

### 13.1.6 2.2.4 (28-June-2018)

- Replace `pyregion` with `stregion`

### 13.1.7 2.2.3 (13-June-2018)

- Updated links in the documentation to point to latest `drizzlepac` website and online API documentation.
- Code cleanup.
- Updated C code to be more compatible with latest `numpy` releases in order to reduce numerous compile warnings.
- Updated documentation to eliminate (at this moment) all `sphinx` documentation generation warnings.
- Moved `'release_notes.rst'` to `'CHANGELOG.rst'` in the top-level directory.
- Improved setup to allow documentation build. See [drizzlepac PR #142](#) and [Issue #129](#) for more details.
- Fixed a bug in a print statement in the create median step due to which background values for input images used in this step were not printed.
- Fixed a bug due to which `TweakReg` may have effectively ignored `verbose` setting.
- Fixed a bug in `drizzlepac.util.WithLogging` due to which `astrodrizzle` would throw an error trying when to raise another error. See [Issue #157](#) for more details.

### 13.1.8 2.2.2 (18-April-2018)

- Fixed a bug in `TweakReg` introduced in `v2.2.0` due to which, when `TweakReg` is run from the interpreter, the code may crash when trying to interpret input files.

### 13.1.9 2.2.1 (12-April-2018)

- Fixed problems with processing WFPC2 data provided by the archive. User will need to make sure they run `updatewcs` on all input WFPC2 data before combining them with `astrodrizzle`.

### 13.1.10 2.2.0 (11-April-2018)

- Implemented a major refactor of the project directory structure. Building no longer requires `d2to1` or `stsci.distutils`. Drizzlepac's release information (i.e. version, build date, etc) is now handled by `relic`. See <https://github.com/spacetelescope/relic>
- Added basic support for compiling Drizzlepac's C extensions under Windows.
- Documentation is now generated during the build process. This ensures the end-user always has access to documentation that applies to the version of `drizzlepac` being used.
- Swapped the effect of setting `configobj` to `None` or `'defaults'` in `AstroDrizzle` and `TweakReg`. When calling one of these tasks with `configobj` parameter set to `None`, values for the not-explicitly-specified parameters should be set to the default values for the task. When `configobj` is set to `'defaults'` not-explicitly-specified parameters will be loaded from the `~/.teal/astrodrizzle.cfg` or `~/.teal/tweakreg.cfg` files that store latest used settings (or from matching configuration files in the current directory). See <https://github.com/spacetelescope/drizzlepac/pull/115> for more details.

### 13.1.11 2.1.22 (15-March-2018)

- Changed the definition of Megabyte used to describe the size of the buffer for `create median` step (`combine_bufsize`). Previously a mixed (base-2 and base-10) definition was used with  $1\text{MB} = 1000 \times 1024\text{B} = 1024000\text{B}$ . Now `1MB` is defined in base-2 (MiB) as  $1\text{MB} = 1024 \times 1024\text{B} = 1048576\text{B}$ .
- Redesigned the logic in `createMedian` step used to split large `single_sci` images into smaller chunks: new logic is more straightforward and fixes errors in the old algorithm that resulted in crashes or unnecessarily small chunk sizes that slowed down `createMedian` step.
- Due to the above mentioned redesign in the logic for splitting large images into smaller chunks, now `overlap` can be set to 0 if so desired in the `minmed` combine type. Also, it is automatically ignored (set to 0) for all non-`minmed` combine types. This will result in additional speed-up in the `Create Median` step.
- Both `AstroDrizzle()` and `TweakReg()` now can be called with `configobj` parameter set to `'defaults'` in order to indicate that values for the not-explicitly-specified parameters should be set to the default values for the task instead of being loaded from the `~/.teal/astrodrizzle.cfg` or `~/.teal/tweakreg.cfg` files that store latest used settings.

- Updated documentation.

### 13.1.12 2.1.21 (12-January-2018)

- Restore recording of correct EXPTIME value in the headers of single drizzled (“single\_sci”) images. See <https://github.com/spacetelescope/drizzlepac/issues/93> for more details.
- Fixed a bug in drizzlepac due to which user provided `combine_lthresh` or `combine_hthresh` in the CREATE MEDIAN IMAGE step were not converted correctly to electrons (processing unit). This bug affected processing of WFPC2, STIS, NICMOS, and WFC3 data. See <https://github.com/spacetelescope/drizzlepac/issues/94> for more details.
- Modified print format so that scales, skew and rotations are printed with 10 significant digits while shifts are printed with 4 digits after the decimal point.

### 13.1.13 2.1.20 (07-October-2017)

- Fixed a bug in expanding reference catalog in `TweakReg` that would result in the code crashing. See <https://github.com/spacetelescope/drizzlepac/pull/87> for more details.
- Fixed a bug due to which user catalog fluxes would be interpreted as magnitudes when `fluxunits` was set to 'cps'. See <https://github.com/spacetelescope/drizzlepac/pull/88> for more details.
- Fixed a bug due to which user-supplied flux limits were ignored for the reference catalog. See <https://github.com/spacetelescope/drizzlepac/pull/89> for more details.

### 13.1.14 2.1.19 (29-September-2017)

- Fixed a bug in computing optimal order of expanding reference catalog that resulted in code crashes. See <https://github.com/spacetelescope/drizzlepac/pull/86> for more details.

### 13.1.15 2.1.18 (05-September-2017)

- Fixed `astrodrizzle` lowers the case of the path of output images issue. See <https://github.com/spacetelescope/drizzlepac/issues/79> for more details.
- Fixed `tweakreg` ignores user-specified units of image catalogs (provided through the `refcat` parameter) issue. See <https://github.com/spacetelescope/drizzlepac/issues/81> for more details.
- Corrected a message printed by `tweakreg` about used WCS for alignment. Also improved documentation for the `refimage` parameter.

### 13.1.16 2.1.17 (13-June-2017)

- `drizzlepac.adrizzle` updated to work with numpy  $\geq 1.12$  when they implemented more strict array conversion rules for math. Any input which still has INT format will be converted to a float

before any operations are performed, explicitly implementing what was an automatic operation prior to numpy 1.12.

### **13.1.17 2.1.16 (05-June-2017)**

- Fixed a bug introduced in release v2.1.15 in the logic for merging WCS due to which custom WCS scale was being ignored.

### **13.1.18 2.1.15 (26-May-2017)**

- `fits.io` operations will no longer use memory mapping in order to reduce the number of file handles used when running either `astrodrizzle` or `tweakreg`. See [issue #39](#) for more details.
- Fixed bugs and improved the logic for merging WCS that is used to define `astrodrizzle`'s output WCS.
- Added `crpix1` and `crpix2` parameters to custom WCS.

### **13.1.19 2.1.14 (28-Apr-2017)**

- Supressed info messages related inconsistent WCS - see [issue #60](#) and [stwcs issue #25](#) for more details.

### **13.1.20 2.1.13 (11-Apr-2017)**

- Fixed a bug due to which sky background was subtracted by `astrodrizzle` from the images even though `skysub` was set to `False` when `MDRIZSKY` was already present in input images' headers.

### **13.1.21 2.1.12 (04-Apr-2017)**

- `astrodrizzle` now will run `updatewcs()` on newly created images when necessary, e.g., after converting `WAVORED FITS` to `MEF` format (`*c0f.fits` to `*_c0h.fits`) or after unpacking multi-imset `STIS_flt` files. See [PR #56](#) for more details.
- Fixed a bug that was preventing processing `STIS` image data.
- Fixed a bug in reading user input (see [issue #51](#)).

### **13.1.22 2.1.11 (24-Mar-2017)**

Bug fix release (a bug was introduced in v2.1.10).

### 13.1.23 2.1.10 (23-Mar-2017)

Some of the changes introduced in release v2.1.9 were not backward compatible. This release makes those changes backward compatible.

### 13.1.24 2.1.9 (22-Mar-2017)

Compatibility improvements with Python 3 and other STScI software packages.

### 13.1.25 2.1.8 (08-Feb-2017)

- Drizzlepac code will no longer attempt to delete “original” (WCS key ‘O’) resulting in a decreased number of warnings (see [issue #35](#)).
- Negative values are now zeroed in the ‘minmed’ step before attempting to estimate Poisson errors (see [issue #22](#)).
- Fixed a bug in `tweakreg` due to incorrect matrix inversion.
- Improved compatibility with `astropy.io.fits` (‘clobber’ parameter) and `numpy` which has reduced the number of deprecation warnings).
- Existing static masks in the working directory are now overwritten and not simply re-used (see [issue #23](#)).
- Corrected formula for  $\sigma$  computation in the “create median” step to convert background to electrons before computations. This bug was producing incorrect  $\sigma$  for instruments whose gain was different from one.
- Improved `astrodrizzle` documentation for `combine_type` parameter which now also documents the formula for  $\sigma$  computation when `combine_type` parameter is set to ‘minmed’.

### 13.1.26 2.1.6 and 2.1.7rc (15-Aug-2016)

Package maintenance release.

### 13.1.27 2.1.5 (09-Aug-2016)

Technical re-release of v2.1.4.

### 13.1.28 2.1.4 (01-Jul-2016)

The following bug fixes have been implemented:

- `tweakreg` crashes when run with a single input image and a reference catalog.
- Fixes an issue due to which `tweakreg`, when updating image headers, would not add ‘-SIP’ suffix to CTYPE

### 13.1.29 2.1.3 (16-Mar-2016)

- Improved ASN input file handling.
- `astrodrizzle` does not delete `d2imfile` anylonger allowing multiple runs of `updatewcs` on the same WFPC2 image, see [Ticket 1244](#) for more details.
- Allow exclusion regions in `tweakreg` to be in a different directory and allow relative path in exclusion region file name.
- Improved handling of empty input image lists.
- `tweakreg` bug fix: use absolute value of polygon area.

### 13.1.30 2.1.2 (12-Jan-2016)

- `runastrodriz` moved to `drizzlepac` from `acstools` and `wfc3tools` packages.
- Improved logic for duplicate input detection.
- Improved logic for handling custom WCS parameters in `astrodrizzle`.
- Compatibility improvements with Python 3.

### 13.1.31 2.1.1

**Available under SSBX/IRAFX starting:** Nov 17, 2015

This release includes the following bug fixes:

- Resolved order of operation problems when processing WFPC2 data with DGEOFILES.
- The conversion of the WFPC2 DGEOFILE into D2IMFILE is now incorporated into STWCS v1.2.3 (r47112, r47113, r47114) rather than a part of `astrodrizzle`. This requires users to run `updatewcs` first, then `astrodrizzle/tweakreg` will work with that WFPC2 data seamlessly (as if they were ACS or WFC3 data).
- Compatibility improvements with Python 3.

### 13.1.32 2.1.0

**Available under SSBX/IRAFX starting:** Nov 2, 2015

This version builds upon the major set of changes implemented in v2.0.0 by not only fixing some bugs, but also cleaning up/changing/revising some APIs and docstrings. The complete list of changes includes:

- [API Change] The ‘`updatewcs`’ parameter was removed from both the `astrodrizzle` and `tweakreg` interactive TEAL interfaces. The ‘`updatewcs`’ parameter can still be used with the Python interface for both the `astrodrizzle.astrodrizzle``()` and ```tweakreg`. Call the `stwcs.updatewcs.updatewcs()` function separately before running `astrodrizzle` or `tweakreg`.



- [API Change] The stand-alone interface for the blot routine (`ablot.blot()`) has been revised to work seamlessly with astrodrizzle-generated products while being more obvious how to call it correctly. The help file for this task was also heavily revised to document all the input parameters and to provide an example of how to use the task.
- [API Change] Coordinate transformation task (`pixtopix/pixtosky/skytopix`) interfaces changed to be more consistent, yet remain backward-compatible for now.
- Both `astrodrizzle` and `tweakreg` now return an output CD matrix which has identical cross-terms indicating the same scale and orientation in each axis (an orthogonal CD matrix). This relies on a revision to the `stwcs.distortion.utils.output_wcs()` function.
- The user interfaces to all 3 coordinate transformation tasks now use ‘coordfile’ as the input file of coordinates to transform. The use of ‘coords’ has been deprecated, but still can be used if needed. However, use of ‘coordfile’ will always override any input provided simultaneously with ‘coords’ parameter. Help files have been updated to document this as clearly as possible for users.
- User-provided list of input catalogs no longer needs to be matched exactly with input files. As long as all input images are included in input catalog list in any order, `tweakreg` will apply the correct catalog to the correct file.
- `tweakreg` has been updated to correctly and fully apply source selection criteria for both input source catalogs and reference source catalogs based on `fluxmin`, `fluxmax` and `nbright` for each.
- All use of keyword deletion has been updated in `drizzlepac` (and `fitsblender`) to avoid warnings from `astropy`.
- All 3 coordinate transformation tasks rely on the input of valid WCS information for the calculations. These tasks now warn the user when it could not find a valid WCS and instead defaulted to using a unity WCS, so that the user can understand what input needs to be checked/revised to get the correct results.
- Exclusion/inclusion region files that can be used with `tweakreg` can now be specified in image coordinates and sky coordinates and will only support files written out using DS9-compatible format.
- The filename for ‘final\_refimage’ in `astrodrizzle` and ‘refimage’ in `tweakreg` can now be specified with OR without an extension, such as ‘[sci,1]’ or ‘[0]’. If no extension is specified, it will automatically look for the first extension with a valid HSTWCS and use that. This makes the use of this parameter in both place consistent and more general than before.
- The reported fit as written out to a file has been slightly modified to report more appropriate numbers of significant digits for the results.
- Use of `astrolib.coords` was removed from `drizzlepac` and replaced by use of `astropy` functions instead. This eliminated one more obsolete dependency in our software.
- Code was revised to rely entirely on `astropy.wcs` instead of stand-alone `pywcs`.
- Code was revised to rely entirely on `astropy.io.fits` instead of stand-alone `pyfits`.
- Added `photeq` task to account for inverse sensitivity variations across detector chips and/or epochs.
- WFPC2 data from the archive with `DGEOFILE` reference files will now need to be processed using `stwcs.updatewcs` before running them through `astrodrizzle` or `tweakreg`. This update

converts the obsolete, unsupported `DGEOFILE` correction for the WFPC2 data into a `D2IMFILE` specific for each WFPC2 observation, then uses that to convert the WCS based on the new conventions used for ACS and WFC3.

This set of changes represents the last major development effort for `DrizzlePac` in support of HST. Support of this code will continue throughout the lifetime of HST, but will be limited primarily to bug fixes to keep the code viable as Python libraries used by `DrizzlePac` continue to develop and evolve with the language.

### 13.1.33 2.0.0

**\*\* Available under SSBX/IRAFX starting:\*\* Aug 4, 2014**

This version encompasses a large number of updates and revisions to the `DrizzlePac` code, including the addition of new tasks and several parameter name changes. The scope of these changes indicates the level of effort that went into improving the `DrizzlePac` code to make it easier and more productive for users. The most significant updates to the `DrizzlePac` code include:

- The Python code has been updated to work identically (without change) under both Python 2.7 and Python 3.x.
- Implementing sky matching, a new algorithm for matching the sky across a set of images being combined by `astrodrizzle`.
- Updating `tweakreg` to now align full mosaics where some images may not overlap others in the mosaic.
- Added the option to write out single drizzle step images as compressed images (to save disk space for large mosaics, and I/O time for single drizzle step).
- Improved `tweakreg` residual plots visually while allowing them to be written out automatically when `tweakreg` gets run in non-interactive mode.
- Renamed parameters in `tweakreg` and `imagefind` to eliminate name clashes.
- Added option to select sources based on sharpness/roundness when `tweakreg` searches for sources.
- Added support for exclusion and inclusion regions arbitrary shape/size when `tweakreg` searches for sources.
- Added a full set of source detection parameters for reference image to support multi-instrument alignment in `tweakreg`.
- Added support for new (simpler, more robust) ACS calibration of time-dependent distortion.
- A full 6-parameter general linear fit can now be performed using `tweakreg`, in addition to shift and `rscale`.
- Cleaned up logic for sky-subtraction: user can now turn off sky-subtraction with `skysub=no`, and still specify a user-defined sky value as the `skyuser` keyword. This will reduce(eliminate?) the need to manually set `MDRIZSKY=0`.

In addition to these major updates/changes, numerous smaller bugs were fixed and other revisions were implemented which affected a small portion of the use cases, such as:

- headerlet code now accepts lists of files to be updated.
- source sky positions (RA and Dec) now included in match file.
- DQ flags can now be taken into account when performing source finding in `tweakreg`.
- all intermediate files generated by `astrodrizzle` will now be removed when using `'clean'='yes'`.
- a problem was fixed that caused `createMedian` to crash where there were no good pixels in one of the images (when they did not overlap).
- interpretation of `shiftfile` now improved to handle arbitrarily-long filenames, rather than being limited to 24 character filenames.
- documentation has been updated, sometimes with a lot more extensive descriptions.

This version of `DrizzlePac` also requires use of the latest release version of `astropy` primarily for WCS and FITS I/O support.

### 13.1.34 1.1.16

**Publicly Released through PyPI:** Mar 27, 2014

**Available under SSBX/IRAFX starting:** Mar 13, 2014

- Support for WFPC2 GEIS input images improved to correctly find the associated DQ images.
- Static mask files created for all chips in an image now get deleted when using the `'group'` parameter to only drizzle a single chip or subset of chips.
- Fixed problem caused by changes to `stsci.tools` code so that `drizzlepac` will reference the correct extensions in input images.

### 13.1.35 1.1.15 (30-Dec-2013)

**Publicly Released through PyPI:** Jan 14, 2014

**Available under SSBX/IRAFX starting:** Jan 6, 2014

#### Bug fixes

- Files created or updated by `drizzlepac`, `fitsblender`, or STWCS tasks, e.g. `tweakreg` or `apply_headerlet`, will now ensure that the `NEXTEND` keyword value correctly reflects the number of extensions in the FITS file upon completion.

### 13.1.36 1.1.14dev (21-Oct-2013)

**Installed in OPUS:** Dec 11, 2013

**Available starting:** Oct 28, 2013

## Bug fixes

- DQ arrays in input images now get updated with cosmic-ray masks computed by `astrodrizzle` when run with the parameter `in_memory=True`. This restored the cosmic-ray masks detected during pipeline processing.

### 13.1.37 v1.1.13dev (11-Oct-2013)

**available starting:** Oct 21, 2013

- `tweakreg` can now be run in ‘batch’ mode. This allows the user to generate plots and have them saved to disk automatically without stopping processing and requiring any user input.

### 13.1.38 1.1.12dev (05-Sep-2013)

**available starting:** Sept 9, 2013

This version fixed a couple of bugs in `astrodrizzle`; namely,

- Logic was updated to support `pixfrac = 0.0` without crashing. This code will now automatically reset the kernel to ‘point’ in that case.
- `astrodrizzle` now forcibly removes all OPUS WCS keywords from drizzle product headers.
- Default rules for generating drizzle product headers (as used in the archive) were modified to add definitions for ‘float\_one’, ‘int\_one’, ‘zero’ that generate output values of 1.0, 1, and 0 (zero) respectively for use as keyword values. This allows the LTM\* rules to replace ‘first’ with ‘float\_one’ so that the physical and image coordinates for drizzle products are consistent.

Additionally, changes were made to `STWCS` for reprocessing use:

- Problems with using `apply_headerlet_as_primary()` from the `STWCS` package on WFPC2 data have been corrected in this revision.

### 13.1.39 1.1.11dev (05-Jul-2013)

**Available starting:** July 15, 2013

- `AstroDrizzle` now can process all STIS data without crashing.

### 13.1.40 1.1.10dev (06-Feb-2013)

**available starting:** May 6, 2013

- The output drizzle image header no longer contains references to D2IM arrays. This allows `tweakreg` to work with drizzled images as input where 2-D D2IM corrections were needed.
- Deprecated references to PyFITS `.has_key()` methods were also removed from the entire package, making it compatible with PyFITS 3.2.x and later.

### 13.1.41 1.1.8dev (06-Feb-2013)

**available starting:** Feb 11, 2013

- Fixed a bug in `astrodrizzle` which caused `blot` to raise an exception when using ‘`sinc`’ interpolation.
- Cleaned up the logic for writing out the results from the `pixtopix`, `pixtosky`, and `skytopix` tasks to avoid an Exception when a list of inputs are provided and no output file is specified.
- A new parameter was added to the `tweakback` task to allow a user to specify the value of `WCSNAME` when updating the FLT images with a new solution from a DRZ image header.
- Code in `tweakback` for updating the header with a new WCS will now automatically generate a unique `WCSNAME` if there is a WCS solution in the FLT headers with the default or user-defined value of `WCSNAME`.

### 13.1.42 1.1.7dev (18-Dec-2012)

**available starting:** Feb 4, 2013

- Updated `astrodrizzle` to work with input images which do not have `WCSNAME` defined. This should make it easier to support non-HST input images in the future.
- cleared up confusion between flux parameters in `imagefindpars` and catalog inputs in `tweakreg`.
- turned off use of fluxes for trimming input source catalogs when no flux column can be found in input source catalogs.

### 13.1.43 1.1.7dev (18-Dec-2012)

**available starting:** Dec 10, 2012

- Update `tweakreg` 2d histogram building mode to correctly find the peak when all the inputs match with the same offset (no spurious sources in either source catalog).
- Fixed a bug so that Ctrl-C does not cause an exception when used while `tweakreg` is running.
- revised the source finding logic to ignore sources near the image edge, a change from how `daofind` works (`daofind` expands the image with blanks then fits anyway).
- created a new function to apply the `nsigma` separation criteria to (try to) eliminate duplicate entries for the same source from the source list. It turns out `daofind` does have problems with reporting some duplicate sources as well. This function does not work perfectly, but works to remove nearly all (if not all) duplicates in most cases.

### 13.1.44 1.1.7dev (8-Jan-2012)

**available starting:** Jan 14, 2013

- Bug fixed in `updatehdr` module to allow shiftfiles without RMS columns to work as inputs to manually apply shifts to headers of input images.

- Revised `astrodrizzle` to update WCS of all input images BEFORE checking whether or not they are valid. This ensures that all files provided as input to `astrodrizzle` in the pipeline have the headers updated with the distortion model and new WCS.
- Images with `NGOODPIX=0` now identified for WFC3 and WFPC2 inputs, so they can be ignored during `astrodrizzle` processing.
- Replaced 2d histogram building code originally written in Python with a C function that run about 4x faster.

### **13.1.45 1.1.6dev (5-Dec-2012)**

**available starting:** Dec 10, 2012

- `tweakreg v1.1.0` source finding algorithm now runs many times faster (no algorithmic changes). No changes have been made yet to speed up the 2d histogram source matching code.
- The ‘`pixtopix`’ task was updated to make the ‘`outimage`’ parameter optional by using the input image as the default. This required no API changes, but the help files were updated.
- Very minor update to guard against `MDRIZTAB` being specified without any explicit path.
- Update `astrodrizzle` to correctly report the exposure time, exposure start, and exposure end for the single drizzle products, in addition to insuring the final drizzle values remain correct.
- `astrodrizzle` also includes initial changes to safeguard the C code from getting improperly cast values from the `configObj(TEAL)` input.

### **13.1.46 1.1.5dev (23-Oct-2012)**

**available starting:** Oct 29, 2012

- Scaling of sky array for WFC3/IR IVM generation now correct.
- template mask files for WFPC2 no longer generated so that WFPC2 data can now be processed using `num_cores > 1` (parallel processing).
- interpretation of the ‘`group`’ parameter fixed to support a single integer, a comma-separated list of integers or a single ‘`sci,<n>`’ value. The values correspond to the FITS extension number of the extensions that should be combined. This fix may also speed up the initialization step as more direct use of `pyfits` was implemented for the interpretation of the ‘`group`’ parameter.

### **13.1.47 1.1.1 (31-Aug-2012)**

**available starting:** Sept 26, 2012

The HST Archive and operational calibration pipeline started using this version on Sept 26, 2012.

### 13.1.48 1.1.4dev (20-Sep-2012)

**available starting:** Sept 24, 2012

- Bug fixed to allow use of `final_wht_type=IVM` for processing WFPC2 data.
- Revised Initialization processing to speed it up by using more up-to-date, direct pyfits calls.

### 13.1.49 1.1.3 (7-Sep-2012)

**available starting:** Sept 17, 2012

- Fixed the logic so that `crclean` images always get created regardless of the value of the ‘clean’ parameter.

### 13.1.50 1.1.2 (5-Sep-2012)

**available starting:** Sept 10, 2012

- Remove the restriction of only being able to process images which have `WCSNAME` keyword as imposed by r15631. The removal of this restriction will now allow for processing of non-updated input files with `updatewcs=False` for cases where no distortion model exists for the data (as required by CADCE).
- Added log statements reporting what sky value was actually used in the drizzle and blot steps

### 13.1.51 1.1.1 (30-Aug-2012)

**available starting:** Sept 3, 2012

- Major revision to `astrodrizzle` allowing the option to process without writing out any intermediate products to disk. The intermediate products remain in memory requiring significantly more memory than usual. This improves the overall processing time by eliminating as much disk activity as possible as long as the OS does not start disk swapping due to lack of RAM.
- revised to turn off ‘updatewcs’ when `coeffs=False(no)` so that exposures with filter combinations not found in the IDCTAB will not cause an error.

### 13.1.52 1.0.7 (21-Aug-2012)

**available starting:** Aug 27, 2012

- Fixes problems with missing `single_sci` images.
- Static mask step revised to skip updates to static mask if all pixel data falls within a single histogram bin. This avoids problems with masking out entire images, which happens if low S/N SBC data is processed with `static_mask=yes`.

### 13.1.53 1.0.6 (14-Aug-2012)

**available starting:** Aug 20, 2012

Use of IVM for final\_wht now correct, as previous code used wrong inputs when IVM weighting was automatically generated by `astrodrizzle`.

### 13.1.54 1.0.5 (8-Aug-2012)

**available starting:** Aug 13, 2012

- Completely removed the use of the TIME arrays for weighting IR drizzle products so that the photometry for saturated sources in drizzled products now comes out correct.
- Corrected a problem with `astrodrizzle` which affected processing of WFPC2 data where CR-PIX2 was not found when creating the output single sci image.

### 13.1.55 1.0.2 (13-July-2012)

**available starting:** Aug 3, 2012

The complete version of `stsci_python` can be downloaded from our [download page](#)

- [stsci\\_python v2.13 Release Notes](#)
- [Old stsci\\_python release notes](#)

### 13.1.56 1.0.1 (20-June-2012)

**Used in archive/pipeline starting:** July 10, 2012

Pipeline and archive started processing ACS data with this version.

### 13.1.57 1.0.0 (25-May-2012)

**Used in archive/pipeline starting:** June 6, 2012

Pipeline and archive first started using `astrodrizzle` by processing WFC3 images.



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## Image Registration Tasks

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Documentation for the replacement task for IRAF's `tweakshifts`, currently named `TweakReg`, has been added to this package. These new modules describe how to run the new TEAL-enabled task, as well as use the classes in the task to generate catalogs interactively for any chip and work with that catalog. The current implementation of this code relies on a very basic source finding algorithm loosely patterned after the DAOFIND algorithm and does not provide all the same features or outputs found in DAOFIND. The fitting algorithm also reproduces the fitting performed by IRAF's `geomap` in a limited fashion; primarily, it only performs fits equivalent to `geomap`'s 'shift' and 'rscale' solutions. These algorithms will be upgraded as soon as replacements are available.

### 14.1 TWEAKREG: Image Alignment

Combining images using `AstroDrizzle` requires that the WCS information in the headers of each input image align to within sub-pixel accuracy. The `TweakReg` task allows the user to align sets of images to each other and/or to an external astrometric reference frame or image.

`TweakReg` - A replacement for IRAF-based `tweakshifts`

**Authors** Warren Hack, Mihai Cara

**License** *LICENSE*

`drizzlepac.tweakreg.TweakReg` (*files=None, editpars=False, configobj=None, imagefind-  
cfg=None, refimagefindcfg=None, \*\*input\_dict*)

`Tweakreg` provides an automated interface for computing residual shifts between input exposures being combined using `AstroDrizzle`. The offsets computed by `Tweakreg` correspond to pointing differences after applying the WCS information from the input image's headers. Such errors would, for example, be due to errors in guide-star positions when combining observations from different observing visits or from slight offsets introduced upon re-acquiring the guide stars in a slightly different position.

## Parameters

**file** [str or list of str (Default = `*flt.fits`)] Input files (passed in from *files* parameter) This parameter can be provided in any of several forms:

- filename of a single image
- filename of an association (ASN)table
- wild-card specification for files in directory (using `\*`, `?` etc.)
- comma-separated list of filenames
- `@file` filelist containing list of desired input filenames with one filename on each line of the file.

**editpars** [bool (Default = False)] A parameter that allows user to edit input parameters by hand in the GUI. `True` to use the GUI to edit parameters.

**configobj** [ConfigObjPars, ConfigObj, dict (Default = None)] An instance of `stsci.tools.cfgpars.ConfigObjPars` or `stsci.tools.configobj.ConfigObj` which overrides default parameter settings. When `configobj` is defaults, default parameter values are loaded from the user local configuration file usually located in `~/.teal/tweakreg.cfg` or a matching configuration file in the current directory. This configuration file stores most recent settings that an user used when running `TweakReg` through the TEAL interface. When `configobj` is `None`, `TweakReg` parameters not provided explicitly will be initialized with their default values as described in the “Other Parameters” section.

**imagefindcfg** [dict, configObject (Default = None)] An instance of `dict` or `configObject` which overrides default source finding (for input images) parameter settings. See help for `imagefindpars` PSET for a list of available parameters. **Only** the parameters that are different from default values **need** to be specified here.

**refimagefindcfg** [dict, configObject (Default = None)] An instance of `dict` or `configObject` which overrides default source finding (for input reference image) parameter settings. See help for `refimagefindpars` PSET for a list of available parameters. **Only** the parameters that are different from default values **need** to be specified here.

**input\_dict** [dict, optional] An optional list of parameters specified by the user, which can also be used to override the defaults.

---

**Note:** This list of parameters **can** include the `updatewcs` parameter, even though this parameter no longer can be set through the TEAL GUI.

---



---

**Note:** This list of parameters **can** contain parameters specific to the `TweakReg` task itself described here in the “Other Parameters” section and **may not** contain

parameters from the `refimagefindpars` PSET.

---

**Note:** For compatibility purpose with previous `TweakReg` versions, `input_dict` may contain parameters from the `imagefindpars` PSET. However, if `imagefindcfg` is not `None`, then `imagefindpars` parameters specified through `input_dict` may not duplicate parameters specified through `imagefindcfg`.

---

### Other Parameters

**refimage** [str (Default = ‘’)] Filename of reference image. Sources derived from this image will be used as the reference for matching with sources from all input images unless a separate catalog is provided through the `refcat` parameter. In addition, this image file must contain a valid not distorted WCS that will define the projection plane in which image alignment is performed (“reference WCS”). When `refimage` is not provided, a reference WCS will be derived from input images.

**expand\_refcat** [bool (Default = False)] Specifies whether to add new sources from just matched images to the reference catalog to allow next image to be matched against an expanded reference catalog.

**enforce\_user\_order** [bool (Default = True)] Specifies whether images should be aligned in the order specified in the `file` input parameter or `TweakReg` should optimize the order of alignment by intersection area of the images. Default value (`True`) will align images in the user specified order, except when some images cannot be aligned in which case `TweakReg` will optimize the image alignment order. Alignment order optimization is available *only* when `expand_refcat = True`.

**exclusions: string (Default = ‘’)** This parameter allows the user to specify a set of files which contain regions in the image to ignore (or to include) when finding sources. This file **MUST** have 1 line for each input image with the name of the input file in the first column. Subsequent columns would be used to specify an inclusion and/or exclusion file for each chip in 'SCI, <n>' index order. If a chip does not require an exclusion file, the string `None` or `INDEF` can be used as a placeholder for that chip. Each exclusion file can be either a mask provided as a simple FITS file or a region file in DS9-format.

When a mask file is provided, `TweakReg` will look for the first image-like extension with image data of the same dimensions as the input image. Zeros in the mask will be interpreted as “bad” (excluded from search) pixels while non-zero pixels will be interpreted as “good” pixels. It is recommended that mask files be FITS files without extensions and mask data (preferably of integer type) reside in the primary HDU.

If a region file is provided then it should conform to the ‘region’ file format generated by DS9. The region files can contain both regular (“include”) regions as well as “exclude” regions. Regular (“include”) regions indicate the regions of the

image that should be searched for sources while “exclude” regions indicate parts of the image that should not be used for source detection. The “ruler”, “compass”, and “projection” regions are not supported (ignored). When **all regions** in a region file are “exclude” regions, then it will be assumed that the entire image is “good” before the exclude regions are processed. In other words, an “include” region corresponding to the entire image will be *prepended* to the list of exclude regions.

---

**Note:** Regions in a region file are processed in the order they appear in the region file. Thus, when region files contain *both* “include” and “exclude” regions, the order in which these regions appear may affect the results.

---

**Warning:** `TweakReg` relies on `pyregion` package for work with region files. At the time of writing, `pyregion` uses a different algorithm from DS9 for converting regions from sky coordinates to image coordinate (this conversion is performed before regions are converted to masks). For these reasons, regions provided in sky coordinates may not produce the expected (from DS9) results. While in most instances these discrepancies should be tolerable, it is important to keep this in mind.

During testing it was observed that conversion to image coordinates is most accurate for polygonal regions and less accurate for other regions. Therefore, if one must provide regions in sky coordinates, it is recommended to use polygonal and circular regions and to avoid elliptical and rectangular regions as their conversion to image coordinates is less accurate. One may use `mapreg` task in the `drizzlepac` package to convert region files from sky coordinates to image coordinates. This will allow one to see the actual regions that will be used by source finding routine in `TweakReg`.

**updatewcs** [bool (Default = No)] **NOT available through TEAL GUI interface.**

This parameter can only be set through the Python interface to `Tweakreg` by passing it in as part of the `input_dict` in order to insure that running `updatewcs` **does not overwrite** a previously determined solution written out to the input file headers.

**writecat** [bool (Default = Yes)] Specify whether or not to write out the source catalogs generated for each input image by the built-in source extraction algorithm.

**clean** [bool (Default = No)] Specify whether or not to remove the temporary files created by `TweakReg`, including any catalog files generated for the shift determination.

**interactive** [bool (Default = Yes)] This switch controls whether the program stops and waits for the user to examine any generated plots before continuing on to the next image. If turned off, plots will still be displayed, but they will also be saved to disk automatically as a PNG image with an autogenerated name without requiring any user input.

**verbose** [bool (Default = No)] Specify whether or not to print extra messages during processing.

**runfile** [string (Default = 'tweakreg.log')] Specify the filename of the processing log.

**\*UPDATE HEADER\***

**updatehdr** [bool (Default = No)] Specify whether or not to update the headers of each input image directly with the shifts that were determined. This will allow the input images to be combined by `AstroDrizzle` without having to provide the shiftfile as well.

**wcsname** [str (Default = 'TWEAK')] Name of updated primary WCS.

**reusename** [bool (Default = False)] Allows overwriting of an existing primary WCS with the same name as specified by `wcsname` parameter.

**\*HEADERLET CREATION\***

**headerlet: bool (Default = No)** Specify whether or not to generate a headerlet from the images at the end of the task? If turned on, this will create a headerlet from the images regardless of the value of the `updatehdr` parameter.

**attach: bool (Default = Yes)** If creating a headerlet, choose whether or not to attach the new headerlet to the input image as a new extension.

**hdrfile: string (Default = '')** Filename to use for writing out headerlet to a separate file. If the name does not contain `.fits`, it will create a filename from the rootname of the input image, the value of this string, and it will end in `'_hlet.fits'`. For example, if only `'hdrlet1'` is given, the full filename created will be `'j99da1f2q_hdrlet1_hlet.fits'` when creating a headerlet for image `'j99da1f2qflt.fits'`.

**lobber: bool (Default = No)** If a headerlet with `'hdrfile'` already exists on disk, specify whether or not to overwrite that previous file.

**hdrname: string (Default = '')** Unique name to give to headerlet solution. This name will be used to identify this specific WCS alignment solution contained in the headerlet.

**author: string, optional (Default = '')** Name of the creator of the headerlet.

**descrip: string, optional (Default = '')** Short (1-line) description to be included in headerlet as `DESCRIP` keyword. This can be used to provide a quick look description of the WCS alignment contained in the headerlet.

**catalog: string, optional (Default = '')** Name of reference catalog used as the basis for the image alignment.

**history: string, optional (Default = '')** Filename of a file containing detailed information regarding the history of the WCS solution contained in the headerlet. This can include information on the catalog used for the alignment, or notes on processing that went into finalizing the WCS alignment stored in this headerlet. This information will be reformatted as 70-character wide FITS `HISTORY` keyword section.

**\*OPTIONAL SHIFTFILE OUTPUT\***

**shiftfile** [bool (Default = No)] Create output shiftfile?

**outshifts** [str (Default = 'shifts.txt')] The name for the output shift file created by `TweakReg`. This shiftfile will be formatted for use as direct input to `AstroDrizzle`.

**outwcs** [str (Default = 'shifts\_wcs.fits')] Filename to be given to the OUTPUT reference WCS file created by `TweakReg`. This reference WCS defines the WCS from which the shifts get measured, and will be used by `AstroDrizzle` to interpret those shifts. This reference WCS file will be a FITS file that only contains the WCS keywords in a Primary header with no image data itself. The values will be derived from the FIRST input image specified.

**\*COORDINATE FILE DESCRIPTION\***

**catfile** [str (Default = '')] Name of file that contains a list of input images and associated catalog files generated by the user. Each line of this file will contain the name of an input image in the first column. The remaining columns will provide the names of the source catalogs for each chip in order of the science extension numbers ((SCI,1), (SCI,2), ...).

A sample catfile, with one line per image would look like:

```
image1_flt.fits  cat1_sci1.coo  cat1_sci2.coo
image2_flt.fits  cat2_sci1.coo  cat2_sci2.coo
```

---

**Note:** Catalog files themselves must be text files containing “white space”-separated list of values (`xcol`, `ycol`, etc.)

---

**xcol** [int (Default = 1)] Column number of X position from the user-generated catalog files specified in the catfile.

**ycol** [int (Default = 2)] Column number of Y position from the user-generated catalog files specified in the catfile.

**fluxcol** [int (Default = None)] Column number for the flux values from the user-generated catalog files specified in the catfile. These values will only be used if a flux limit has been specified by the user using the `maxflux` or `minflux` parameters.

**maxflux** [float (Default = None)] Limiting flux value for selecting valid objects in the input image’s catalog. If specified, this flux will serve as the upper limit of a range for selecting objects to be used in matching with objects identified in the reference image. If the value is set to `None`, all objects with fluxes brighter than the minimum specified in `minflux` will be used. If both values are set to `None`, all objects will be used.

**minflux** [float (Default = None)] Limiting flux value for selecting valid objects in the input image’s catalog. If specified, this flux value will serve as the lower limit of a range for selecting objects to be used in matching with objects identified in

the reference image. If the value is set to `None`, all objects fainter than the limit specified by `maxflux` will be used. If both values are set to `None`, all objects will be used.

**fluxunits** [str {'counts', 'cps', 'mag'} (Default = 'counts')] This allows the task to correctly interpret the flux limits specified by `maxflux` and `minflux` when sorting the object list for trimming of fainter objects.

**xyunits** [str {'pixels', 'degrees'} (Default = 'pixels')] Specifies whether the positions in this catalog are already sky pixel positions, or whether they need to be transformed to the sky.

**nbright** [int (Default = None)] The number of brightest objects to keep after sorting the full object list. If `nbright` is set equal to `None`, all objects will be used.

#### \*REFERENCE CATALOG DESCRIPTION\*

**refcat** [str (Default = '')] Name of the external reference catalog file to be used in place of the catalog extracted from one of the input images. When `refimage` is not specified, reference WCS to be used with reference catalog will be derived from input images.

---

**Note:** Reference catalog must be text file containing “white space”-separated list of values (`xcol`, `ycol`, etc.)

---

**refxcol** [int (Default = 1)] Column number of RA in the external catalog file specified by the `refcat`.

**refycol** [int (Default = 2)] Column number of Dec in the external catalog file specified by the `refcat`.

**refxyunits** [str {'pixels', 'degrees'} (Default = 'degrees')] Units of sky positions.

**rfluxcol** [int (Default = None)] Column number of flux/magnitude values in the external catalog file specified by the `refcat`.

**rmaxflux** [float (Default = None)] Limiting flux value used to select valid objects in the external catalog. If specified, the flux value will serve as the upper limit of a range for selecting objects to be used in matching with objects identified in the reference image. If the value is set to `None`, all objects with fluxes brighter than the minimum specified in `rminflux` will be used. If both values are set to `None`, all objects will be used.

**rminflux** [float (Default = None)] Limiting flux value used to select valid objects in the external catalog. If specified, the flux will serve as the lower limit of a range for selecting objects to be used in matching with objects identified in the reference image. If the value is set to `None`, all objects fainter than the limit specified by `rmaxflux` will be used. If both values are set to `None`, all objects will be used.

**rfluxunits** [{ 'counts', 'cps', 'mag' } (Default = 'mag')] This allows the task to correctly interpret the flux limits specified by `rmaxflux` and `rminflux` when sorting the object list for trimming of fainter objects.

**refnbright** [int (Default = None)] Number of brightest objects to keep after sorting the full object list. If refnbright is set to `None`, all objects will be used. Used in conjunction with refcat.

**\*OBJECT MATCHING PARAMETERS\***

**minobj** [int (Default = 15)] Minimum number of identified objects from each input image to use in matching objects from other images.

**searchrad** [float (Default = 1.0)] The search radius for a match.

**searchunits** [str (Default = 'arcseconds')] Units for search radius.

**use2dhist** [bool (Default = Yes)] Use 2d histogram to find initial offset?

**see2dplot** [bool (Default = Yes)] See 2d histogram for initial offset?

**tolerance** [float (Default = 1.0)] The matching tolerance in pixels after applying an initial solution derived from the 'triangles' algorithm. This parameter gets passed directly to `xyxymatch` for use in matching the object lists from each image with the reference image's object list.

**separation** [float (Default = 0.0)] The minimum separation for objects in the input and reference coordinate lists. Objects closer together than 'separation' pixels are removed from the input and reference coordinate lists prior to matching. This parameter gets passed directly to `xyxymatch` for use in matching the object lists from each image with the reference image's object list.

**xoffset** [float (Default = 0.0)] Initial estimate for the offset in X between the images and the reference frame. This offset will be used for all input images provided. If the parameter value is set to `None`, no offset will be assumed in matching sources in `xyxymatch`.

**yoffset** [float (Default = 0.0)] Initial estimate for the offset in Y between the images and the reference frame. This offset will be used for all input images provided. If the parameter value is set to `None`, no offset will be assumed in matching sources in `xyxymatch`.

**\*CATALOG FITTING PARAMETERS\***

**fitgeometry** [str {'shift', 'rscale', 'general'} (Default = 'rscale')] The fitting geometry to be used in fitting the matched object lists. This parameter is used in fitting the offsets, rotations and/or scale changes from the matched object lists. The 'general' fit geometry allows for independent scale and rotation for each axis.

**residplot** [str {'No plot', 'vector', 'residuals', 'both'} (Default = 'both')] Plot residuals from fit? If 'both' is selected, the 'vector' and 'residuals' plots will be displayed in separate plotting windows at the same time.

**nclip** [int (Default = 3)] Number of clipping iterations in fit.

**sigma** [float (Default = 3.0)] Clipping limit in sigma units.

**\*ADVANCED PARAMETERS AVAILABLE FROM COMMAND LINE\***



**updatewcs** [bool (Default = No)] This parameter specifies whether the WCS keywords are to be updated by running `updatewcs` on the input data, or left alone. The update performed by `updatewcs` not only recomputes the WCS based on the currently used `IDCTAB`, but also populates the header with the `SIP` coefficients. For `ACS/WFC` images, the time-dependence correction will also be applied to the WCS and `SIP` keywords. This parameter should be set to 'No' (`False`) when the WCS keywords have been carefully set by some other method, and need to be passed through to `drizzle` 'as is', otherwise those updates will be over-written by this update.

---

**Note:** This parameter was preserved in the API for compatibility purposes with existing user processing pipe-lines. However, it has been removed from the `TEAL` interface because it is easy to have it set to 'yes' (especially between consecutive runs of `AstroDrizzle`) with potentially disastrous effects on input image WCS (for example it could wipe-out previously aligned WCS).

---

**See also:**

**astrodrizzle**

## Notes

`Tweakreg` supports the use of calibrated, distorted images (such as `FLT` images for `ACS` and `WFC3`, or `_c0m.fits` images for `WFPC2`) as input images. All coordinates for sources derived from these images (either by this task or as provided by the user directly) will be corrected for distortion using the distortion model information specified in each image's header. This eliminates the need to run `AstroDrizzle` on the input images prior to running `TweakReg`.

---

**Note:** All calibrated input images must have been updated using `updatewcs` from the `STWCS` package, to include the full distortion model in the header. Alternatively, one can set `updatewcs` parameter to `True` when running either `TweakReg` or `AstroDrizzle` from command line (Python interpreter) **the first time** on such images.

---

This task will use catalogs, and catalog-matching, based on the `xyxymatch` algorithm to determine the offset between the input images. The primary mode of operation will be to extract a catalog of source positions from each input image using either a 'DAOFIND-like' algorithm or `SExtractor` (if the user has `SExtractor` installed). Alternatively, the user can provide their catalogs of source positions derived from **each input chip**.

---

**Note:** Catalog files must be text files containing "white space"-separated list of values (`xcol`, `ycol`, etc.)

---

The reference frame will be defined either by:

- the image with the largest overlap with another input image AND with the largest total overlap with the rest of the input images,
- a catalog derived from a reference image specified by the user, or
- a catalog of undistorted sky positions (RA/Dec) and fluxes provided by the user.

For a given observation, the distortion model is applied to all distorted input positions, and the sources from each chip are then combined into a single catalog of undistorted positions.

The undistorted positions for each observation then get passed to `xyxymatch` for matching to objects from the reference catalog.

The source lists from each image will generally include cosmic-rays as detected sources, which can at times significantly confuse object identification between images. Observations that include long exposures often have more cosmic-ray events than source objects. As such, isolating the cosmic-ray events in those cases would significantly improve the efficiency of common source identification between images. One such method for trimming potential false detections from each source list would be to set a flux limit to exclude detections below that limit. As the fluxes reported in the default source object lists are provided as magnitude values, setting the `maxflux` or `minflux` parameter value to a magnitude-based limit, and then setting the `ascend` parameter to `True`, will allow for the creation of catalogs trimmed of all sources fainter than the provided limit. The trimmed source list can then be used in matching sources between images and in establishing the final fitting for the shifts.

A fit can then be performed on the matched set of positions between the input and the reference to produce the ‘shiftfile’. If the user is confident that the solution will be correct, the header of each input image can be updated directly with the fit derived for that image. Otherwise, the ‘shiftfile’ can be passed to `AstroDrizzle` for aligning the images.

---

**Note:** Because of the nature of the used algorithm it may be necessary to run this task multiple times until new shifts, rotations, and/or scales are small enough for the required precision.

---

New sources (that are not in the reference catalog) from the matched images are added to the reference catalog in order to allow next image to be matched to a larger reference catalog. This allows alignment of images that do not overlap directly with the reference image and/or catalog and it is particularly useful in image registration of large mosaics. Addition of new sources to the reference catalog can be turned off by setting `expand_refcat` to `False` when using an external reference catalog. When an external catalog is not provided (`refcat=''`) or when using an external reference catalog with `expand_refcat` set to `True` (assuming `writecat = True` and `clean = False`), the list of all sources in the expanded reference catalog is saved in a catalog file named `cumulative_sky_refcat_###.coo` where `###` is the base file name derived from either the external catalog (if provided) or the name of the image used as the reference image.

When `enforce_user_order` is `False`, image catalogs are matched to the reference catalog in order of decreasing overlap area with the reference catalog, otherwise user order of files specified in the `file` parameter is used.

### Format of Exclusion Catalog

The format for the exclusions catalog requires 1 line in the file for every input image, regardless of whether or not that image has any defined exclusion regions. A sample file would look like:

```
j99dalemq_flt.fits
j99dalf2q_flt.fits test_exclusion.reg
```

This file specifies no exclusion files for the first image, and only an regions file for SCI,1 of the second image. NOTE: The first file can be dropped completely from the exclusion catalog file.

In the above example, should an exclusion regions file only be needed for the second chip in the second image, the file would need to look like:

```
j99dalemq_flt.fits
j99dalf2q_flt.fits None test_sci2_exclusion.reg
```

The value `None` could also be replaced by `INDEF` if desired, but either string needs to be present to signify no regions file for that chip while the code continues parsing the line to find a file for the second chip.

### Format of Region Files

The format of the exclusions catalogs referenced in the ‘exclusions’ file defaults to the format written out by DS9 using the ‘DS9/Funtools’ region file format. A sample file with circle() regions will look like:

```
# Region file format: DS9 version 4.1
# Filename: j99dalf2q_flt.fits[SCI]
global color=green dashlist=8 3 width=1 font="helvetica 10 normal roman"
select=1 highlite=1 dash=0 fixed=0 edit=1 move=1 delete=1 include=1
↔source=1
image
circle(3170,198,20)
ellipse(3269,428,30,10,45) # a rotated ellipse
box(3241.1146,219.78132,20,20,15) # a rotated box
circle(200,200,50) # outer circle
-circle(200,200,30) # inner circle
```

This region file will be interpreted as “find all sources in the image that **are inside** the four regions above but **not inside** the region `-circle(200,200,30)`”. Effectively we will instruct `TweakReg` to find all the sources *inside* the following regions:

```
circle(3170,198,20)
ellipse(3269,428,30,10,45) # a rotated ellipse
box(3241.1146,219.78132,20,20,15) # a rotated box
annulus(200,200,30,50) # outer circle(r=50) - inner circle(r=30)
```

### Examples

The `tweakreg` task can be run from either the TEAL GUI or from the command-line using PyRAF or Python. These examples illustrate the various syntax options available.

**Example 1:** Align a set of calibrated (`_flt.fits`) images using `IMAGEFIND`, a built-in source finding algorithm based on DAOPHOT. Auto-detect the sky sigma value and select sources >

200 sigma. (Auto-sigma is computed from the first input exposure as: `1.5*imstat(image, nclip=3, fields='stddev')`.) Set the convolution kernel width to  $\sim 2\times$  the value of the PSF FWHM. Save the residual offsets (`dx`, `dy`, `rot`, `scale`, `xfit_rms`, `yfit_rms`) to a text file.

1. Run the task from PyRAF using the TEAL GUI:

```
>>> import drizzlepac
>>> epar tweakreg
```

2. Run the task from PyRAF using the command line while individually specifying source finding parameters for the reference image and input images:

```
>>> import drizzlepac
>>> from drizzlepac import tweakreg
>>> tweakreg.TweakReg('*flt.fits',
...     imagefindcfg={'threshold' : 200, 'conv_width' : 3.5},
...     refimagefindcfg={'threshold' : 400, 'conv_width' : 2.5},
...     updatehdr=False, shiftfile=True, outshifts='shift.txt')
```

or, using `dict` constructor,

```
>>> import drizzlepac
>>> from drizzlepac import tweakreg
>>> tweakreg.TweakReg('*flt.fits',
...     imagefindcfg=dict(threshold=200, conv_width=3.5),
...     refimagefindcfg=dict(threshold=400, conv_width=2.5),
...     updatehdr=False, shiftfile=True, outshifts='shift.txt')
```

Or, run the same task from the PyRAF command line, but specify all parameters in a config file named “myparam.cfg”:

```
>>> tweakreg.TweakReg('*flt.fits', configobj='myparam.cfg')
```

Alternately, edit the imagefind parameters in a TEAL GUI window prior to running the task:

```
>>> tweakreg.edit_imagefindpars()
```

3. Help can be accessed via the “Help” pulldown menu in the TEAL GUI. It can also be accessed from the PyRAF command-line and saved to a text file:

```
>>> from drizzlepac import tweakreg
>>> tweakreg.help()
```

or

```
>>> tweakreg.help(file='help.txt')
>>> page help.txt
```

`drizzlepac.tweakreg.help` (*file=None*)

Print out syntax help for running `astrodrizzle`

### Parameters

**file** [str (Default = None)] If given, write out help to the filename specified by this parameter. Any previously existing file with this name will be deleted before writing out the help.

## 14.2 Refmagefindpars: Source finding parameters for the reference image

This interface provides a mechanism for setting the parameters used by the built-in source finding algorithm based on the published algorithms used by DAOFIND. These settings apply only to source finding in the reference images.

These parameters control the operation of the algorithm that extracts sources from the reference image (if specified) as called by *TWEAKREG: Image Alignment*. The algorithm implemented follows the concepts defined by DAOFIND (without actually using any DAOFIND code).

### Attributes

**computesig: bool (Default = True)** This parameter controls whether or not to automatically compute a sigma value to be used for object identification. If set to `True`, then the value computed will override any user input for the parameter `skysigma`. The automatic sigma value gets computed from each input exposure as:

$$\sigma = \sqrt{2|mode|}$$

This single value will then be used for object identification for all input exposures.

**skysigma: float (Default = 0.0)** The standard deviation of the sky pixels. This value will only be used if `computesig` is `False`.

**conv\_width: float (Default = 3.5)** The convolution kernel width in pixels. Recommended values (~2x the PSF FWHM): ACS/WFC & WFC3/UVIS ~3.5 pix and WFC3/IR ~2.5 pix.

**peakmin: float, None (Default = None)** This parameter allows the user to select only those sources whose peak value (in the units of the input image) is greater than this value.

**peakmax: float, None (Default = None)** This parameter allows the user to select only those sources whose peak value (in the units of the input image) is less than this value.

**threshold: float (Default = 4.0)** The object detection threshold above the local background in units of sigma.

**nsigma: float (Default = 1.5)** The semi-major axis of the Gaussian convolution kernel used to compute the density enhancement and mean density images in Gaussian sigma.

**ratio: float (Default = 1.0)** The ratio of the sigma of the Gaussian convolution kernel along the minor axis direction to the sigma along the major axis direction. For a circularly-symmetric kernel use `ratio = 1.0`.

**theta: float (Default = 0.0)** The position angle (degrees) of the major axis of the Gaussian convolution kernel. Theta is measured counter-clockwise from the x axis.

**fluxmin: float, None (Default = None)** This parameter allows the user to select only those sources whose total flux (in the units of the input image) is greater than this value.

**fluxmax: float, None (Default = None)** This parameter allows the user to select only those sources whose total flux (in the units of the input image) is less than this value.

**dqbits: int, str, None, optional (Default = None)** Integer sum of all the DQ bit values from the input image's DQ array that should be considered "good" when building masks for source finding. For example, if pixels in the DQ array can be combinations of 1, 2, 4, and 8 flags and one wants to consider DQ "defects" having flags 2 and 4 as being acceptable for source finding, then `dqbits` should be set to  $2+4=6$ . Then a DQ pixel having values 2,4, or 6 will be considered a good pixel, while a DQ pixel with a value, e.g.,  $1+2=3$ ,  $4+8=12$ , etc. will be flagged as a "bad" pixel.

Alternatively, one can enter a comma- or '+'-separated list of integer bit flags that should be added to obtain the final "good" bits. For example, both `4, 8` and `4+8` are equivalent to setting `dqbits` to 12.

Setting `dqbits` to 0 will make *all* non-zero pixels in the DQ mask to be considered "bad" pixels, and the corresponding image pixels will not be used for source finding.

The default value of `None` will turn off the use of image's DQ array for source finding.

In order to reverse the meaning of the `dqbits` parameter from indicating values of the "good" DQ flags to indicating the "bad" DQ flags, prepend '~' to the string value. For example, in order not to use pixels with DQ flags 4 and 8 for source finding and to consider as "good" all other pixels (regardless of their DQ flag), set `dqbits` to `~4+8`, or `~4, 8`. To obtain the same effect with an `int` input value (except for 0), enter  $-(4+8+1)=-9$ . Following this convention, a `dqbits` string value of `'~0'` would be equivalent to setting `dqbits=None`.

**use\_sharp\_round: bool (Default = False)** This parameter controls whether or not to enable selection of sources based on their sharpness and roundness statistics.

**sharplo: float (Default = 0.2)** `sharplo` and `sharphi` are designed to eliminate brightness maxima that are due to bad pixels rather than to astronomical objects. Only sources with sharpness above the `sharplo` value will be selected.

**sharphi: float (Default = 1.0)** `sharplo` and `sharphi` are designed to eliminate brightness maxima that are due to bad pixels rather than to astronomical objects. Only sources with sharpness below the `sharphi` value will be selected.

**roundlo: float (Default = -1.0)** `roundlo` and `roundhi` are designed to eliminate brightness maxima that are due to bad rows or columns, rather than to astronomical objects. Only sources with roundness above the `roundlo` value will be selected.

**roundhi: float (Default = 1.0)** `roundlo` and `roundhi` are designed to eliminate brightness maxima that are due to bad rows or columns, rather than to astronomical objects. Only sources with roundness below the `roundhi` value will be selected.

`drizzlepac.refimagefindpars.getHelpAsString (docstring=False, show_ver=True)`  
return useful help from a file in the script directory called `__taskname__.help`

`drizzlepac.refimagefindpars.help (file=None)`  
Print out syntax help for running `astrodrizzle`

### Parameters

**file** [str (Default = None)] If given, write out help to the filename specified by this parameter. Any previously existing file with this name will be deleted before writing out the help.

## 14.3 Imagefindpars: Source finding parameters

This interface provides a mechanism for setting the parameters used by the built-in source finding algorithm based on the published algorithms used by DAOFIND.

These parameters control the operation of the algorithm that extracts sources from the image as called by *TWEAKREG: Image Alignment*. The algorithm implemented follows the concepts defined by DAOFIND (without actually using any DAOFIND code).

### Attributes

**computesig: bool (Default = True)** This parameter controls whether or not to automatically compute a sigma value to be used for object identification. If set to `True`, then the value computed will override any user input for the parameter `skysigma`. The automatic sigma value gets computed from each input exposure as:

$$\sigma = \sqrt{2 |mode|}$$

This single value will then be used for object identification for all input exposures.

**skysigma: float (Default = 0.0)** The standard deviation of the sky pixels. This value will only be used if `computesig` is `False`.

**conv\_width: float (Default = 3.5)** The convolution kernel width in pixels. Recommended values (~2x the PSF FWHM): ACS/WFC & WFC3/UVIS ~3.5 pix and WFC3/IR ~2.5 pix.

**peakmin: float, None (Default = None)** This parameter allows the user to select only those sources whose peak value (in the units of the input image) is greater than this value.

**peakmax:** float, None (Default = None) This parameter allows the user to select only those sources whose peak value (in the units of the input image) is less than this value.

**threshold:** float (Default = 4.0) The object detection threshold above the local background in units of sigma.

**nsigma:** float (Default = 1.5) The semi-major axis of the Gaussian convolution kernel used to compute the density enhancement and mean density images in Gaussian sigma.

**ratio:** float (Default = 1.0) The ratio of the sigma of the Gaussian convolution kernel along the minor axis direction to the sigma along the major axis direction. For a circularly-symmetric kernel use ratio = 1.0.

**theta:** float (Default = 0.0) The position angle (degrees) of the major axis of the Gaussian convolution kernel. Theta is measured counter-clockwise from the x axis.

**fluxmin:** float, None (Default = None) This parameter allows the user to select only those sources whose total flux (in the units of the input image) is greater than this value.

**fluxmax:** float, None (Default = None) This parameter allows the user to select only those sources whose total flux (in the units of the input image) is less than this value.

**dqbits:** int, str, None, optional (Default = None) Integer sum of all the DQ bit values from the input image's DQ array that should be considered "good" when building masks for source finding. For example, if pixels in the DQ array can be combinations of 1, 2, 4, and 8 flags and one wants to consider DQ "defects" having flags 2 and 4 as being acceptable for source finding, then `dqbits` should be set to `2+4=6`. Then a DQ pixel having values 2,4, or 6 will be considered a good pixel, while a DQ pixel with a value, e.g., `1+2=3`, `4+8=12`, etc. will be flagged as a "bad" pixel.

Alternatively, one can enter a comma- or '+'-separated list of integer bit flags that should be added to obtain the final "good" bits. For example, both `4, 8` and `4+8` are equivalent to setting `dqbits` to `12`.

Setting `dqbits` to 0 will make *all* non-zero pixels in the DQ mask to be considered "bad" pixels, and the corresponding image pixels will not be used for source finding.

The default value of `None` will turn off the use of image's DQ array for source finding.

In order to reverse the meaning of the `dqbits` parameter from indicating values of the "good" DQ flags to indicating the "bad" DQ flags, prepend '~' to the string value. For example, in order not to use pixels with DQ flags 4 and 8 for source finding and to consider as "good" all other pixels (regardless of their DQ flag), set `dqbits` to `~4+8`, or `~4, 8`. To obtain the same effect with an `int` input value



(except for 0), enter  $-(4+8+1)=-9$ . Following this convention, a `dqbits` string value of `'~0'` would be equivalent to setting `dqbits=None`.

**use\_sharp\_round: bool (Default = False)** This parameter controls whether or not to enable selection of sources based on their sharpness and roundness statistics.

**sharplo: float (Default = 0.2)** `sharplo` and `sharphi` are designed to eliminate brightness maxima that are due to bad pixels rather than to astronomical objects. Only sources with sharpness above the `sharplo` value will be selected.

**sharphi: float (Default = 1.0)** `sharplo` and `sharphi` are designed to eliminate brightness maxima that are due to bad pixels rather than to astronomical objects. Only sources with sharpness below the `sharphi` value will be selected.

**roundlo: float (Default = -1.0)** `roundlo` and `roundhi` are designed to eliminate brightness maxima that are due to bad rows or columns, rather than to astronomical objects. Only sources with roundness above the `roundlo` value will be selected.

**roundhi: float (Default = 1.0)** `roundlo` and `roundhi` are designed to eliminate brightness maxima that are due to bad rows or columns, rather than to astronomical objects. Only sources with roundness below the `roundhi` value will be selected.

```
drizzlepac.imagefindpars.getHelpAsString (docstring=False, show_ver=True)
    return useful help from a file in the script directory called __taskname__.help
```

```
drizzlepac.imagefindpars.help (file=None)
    Print out syntax help for running astrodrizzle
```

#### Parameters

**file** [str (Default = None)] If given, write out help to the filename specified by this parameter. Any previously existing file with this name will be deleted before writing out the help.

## 14.4 Image Class

imgclasses.ReflImage

imgclasses.Image

Classes to keep track of all WCS and catalog information.

Used by `TweakReg`.

**Authors** Warren Hack, Mihai Cara

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```
class drizzlepac.imgclasses.Image (filename, input_catalogs=None, exclu-  
sions=None, **kwargs)
```

Bases: `object`

Primary class to keep track of all WCS and catalog information for a single input image. This class also performs all matching and fitting.

#### Parameters

**filename** [str] Filename for image.

**input\_catalogs** [list of str or None] Filename of catalog files for each chip, if specified by user.

**kwargs** [dict] Parameters necessary for processing derived from input configObj object.

**buildDefaultRefWCS** ()

Generate a default reference WCS for this image.

**buildSkyCatalog** ()

Convert sky catalog for all chips into a single catalog for the entire field-of-view of this image.

**clean** ()

Remove intermediate files created.

**close** ()

Close any open file handles and flush updates to disk

**compute\_fit\_rms** ()

**get\_shiftfile\_row** ()

Return the information for a shiftfile for this image to provide compatibility with the IRAF-based MultiDrizzle.

**get\_wcs** ()

Helper method to return a list of all the input WCS objects associated with this image.

**get\_xy\_catnames** ()

Return a string with the names of input\_xy catalog names

**match** (*refimage*, *quiet\_identity*, *\*\*kwargs*)

Uses xyxymatch to cross-match sources between this catalog and a reference catalog (`refCatalog`).

**openFile** (*openDQ=False*)

Open file and set up filehandle for image file

**performFit** (*\*\*kwargs*)

Perform a fit between the matched sources.

#### Parameters

**kwargs** [dict] Parameter necessary to perform the fit; namely, *fitgeometry*.

## Notes

This task still needs to implement (eventually) interactive iteration of the fit to remove outliers.

### `sortSkyCatalog()`

Sort and clip the source catalog based on the flux range specified by the user. It keeps a copy of the original full list in order to support iteration.

### `transformToRef(ref_wcs, force=False)`

Transform sky coords from ALL chips into X,Y coords in reference WCS.

### `updateHeader(wcsname='TWEAK', reusename=False)`

Update header of image with shifts computed by `perform_fit()`.

### `writeHeaderlet(**kwargs)`

Write and/or attach a headerlet based on update to PRIMARY WCS

### `write_fit_catalog()`

Write out the catalog of all sources and resids used in the final fit.

### `write_outxy(filename)`

Write out the output(transformed) XY catalog for this image to a file.

### `write_skycatalog(filename)`

Write out the all\_radec catalog for this image to a file.

```
class drizzlepac.imgclasses.RefImage(wcs_list, catalog, xycatalog=None,
                                     cat_origin=None, **kwargs)
```

Bases: `object`

This class provides all the information needed by to define a reference tangent plane and list of source positions on the sky.

**Warning:** When `wcs_list` is a Python list of WCS objects, each element must be an instance of `stwcs.wcsutil.HSTWCS`.

### `append_not_matched_sources(image)`

### `clean()`

Remove intermediate files created

### `clear_dirty_flag()`

### `close()`

### `get_shiftfile_row()`

Return the information for a shiftfile for this image to provide compatibility with the IRAF-based MultiDrizzle.

### `set_dirty()`

**transformToRef** ()

Transform reference catalog sky positions (`self.all_radec`) to reference tangent plane (`self.wcs`) to create output X,Y positions.

**write\_skycatalog** (*filename*, *show\_flux=False*, *show\_id=False*)

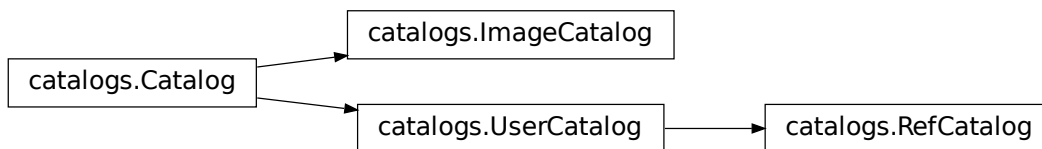
Write out the all\_radec catalog for this image to a file.

## 14.5 Classes to manage Catalogs and WCS's

This module provides the classes used to generate and manage source catalogs for each input chip. Those positions can be transformed to undistorted sky positions, written out to files, or plotted using various methods defined for these classes.

**Authors** Warren Hack

**License** *LICENSE*



`drizzlepac.catalogs.generateCatalog` (*wcs*, *mode='automatic'*, *catalog=None*, *src\_find\_filters=None*, *\*\*kwargs*)

Function which determines what type of catalog object needs to be instantiated based on what type of source selection algorithm the user specified.

### Parameters

**wcs** [obj] WCS object generated by STWCS or PyWCS

**catalog** [str or ndarray] Filename of existing catalog or ndarray of image for generation of source catalog.

**kwargs** [dict] Parameters needed to interpret source catalog from input catalog with `findmode` being required.

### Returns

**catalog** [obj] A Catalog-based class instance for keeping track of WCS and associated source catalog

**class** `drizzlepac.catalogs.ImageCatalog` (*wcs*, *catalog\_source*, *src\_find\_filters=None*, *\*\*kwargs*)

Bases: `drizzlepac.catalogs.Catalog`

Class which generates a source catalog from an image using Python-based, daofind-like algorithms

Required input `kwargs` parameters:

```
computesig, skysigma, threshold, peakmin, peakmax,
hmin, conv_width, [roundlim, sharplim]
```

**generateXY** (\*\*kwargs)

Generate source catalog from input image using DAOFIND-style algorithm

**class** `drizzlepac.catalogs.UserCatalog` (`wcs`, `catalog_source`, \*\*kwargs)

Bases: `drizzlepac.catalogs.Catalog`

Class to manage user-supplied catalogs as inputs.

Required input `kwargs` parameters:

```
xyunits, xcol, ycol[, fluxcol, [idcol]]
```

**COLNAMES** = ['xcol', 'ycol', 'fluxcol']

**IN\_UNITS** = None

**generateXY** (\*\*kwargs)

Method to interpret input catalog file as columns of positions and fluxes.

**plotXYCatalog** (\*\*kwargs)

Plots the source catalog positions using matplotlib's `pyplot.plot()`

Plotting `kwargs` that can also be passed include any keywords understood by matplotlib's `pyplot.plot()` function such as:

```
vmin, vmax, cmap, marker
```

**set\_colnames** ()

Method to define how to interpret a catalog file Only needed when provided a source catalog as input

**class** `drizzlepac.catalogs.RefCatalog` (`wcs`, `catalog_source`, \*\*kwargs)

Bases: `drizzlepac.catalogs.UserCatalog`

Class which manages a reference catalog.

## Notes

A *reference catalog* is defined as a catalog of undistorted source positions given in RA/Dec which would be used as the master list for subsequent matching and fitting.

**COLNAMES** = ['refxcol', 'refycol', 'rfluxcol']

**IN\_UNITS** = 'degrees'

**PAR\_NBRIGHT\_PREFIX** = 'ref'

**PAR\_PREFIX** = 'r'

**buildXY** (catalogs)

**generateRaDec** ()

Convert XY positions into sky coordinates using STWCS methods.

**generateXY** (\*\*kwargs)

Method to interpret input catalog file as columns of positions and fluxes.

**class** drizzlepac.catalogs.**Catalog** (wcs, catalog\_source, \*\*kwargs)

Bases: `object`

Base class for keeping track of a source catalog for an input WCS

**Warning:** This class should never be instantiated by itself, as necessary methods are not defined yet.

This class requires the input of a WCS and a source for the catalog, along with any arguments necessary for interpreting the catalog.

### Parameters

**wcs** [obj] Input WCS object generated using STWCS or HSTWCS

**catalog\_source** [str] Name of the file from which to read the catalog.

**kwargs** [dict] Parameters for interpreting the catalog file or for performing the source extraction from the image. These will be set differently depending on the type of catalog being instantiated.

**PAR\_NBRIGHT\_PREFIX** = ''

**PAR\_PREFIX** = ''

**apply\_exclusions** (exclusions)

Trim sky catalog to remove any sources within regions specified by exclusions file.

**apply\_flux\_limits** ()

Apply any user-specified limits on source selection Limits based on fluxes.

**buildCatalogs** (exclusions=None, \*\*kwargs)

Primary interface to build catalogs based on user inputs.

**generateRaDec** ()

Convert XY positions into sky coordinates using STWCS methods.

**generateXY** (\*\*kwargs)

Method to generate source catalog in XY positions Implemented by each subclass

**plotXYCatalog** (\*\*kwargs)

Method which displays the original image and overlays the positions of the detected sources from this image's catalog.

Plotting `kwargs` that can be provided are:

`vmin`, `vmax`, `cmap`, `marker`

Default colormap is `summer`.

`set_colnames()`

Method to define how to interpret a catalog file Only needed when provided a source catalog as input

`writeXYCatalog(filename)`

Write out the X,Y catalog to a file

## 14.6 Catalog Generation

### 14.6.1 Point (Aperture) Photometric Catalog Generation

#### 1: Source Detection

##### 1.1: Important Clarifications

As previously discussed in *Single-visit Mosaic Processing*, AstroDrizzle creates a single multi-filter detector-level drizzle-combined image for source identification and one or more detector/filter-level drizzle-combined images (depending on which filters were used in the dataset) for photometry. The same set of sources identified in the multi-filter detection image is used to measure photometry for each filter. We use this method to maximize the signal across all available wavelengths at the source detection stage, thus providing photometry with the best quality source list across all available input filters.

It should also be stressed here that the point and segment photometry source list generation algorithms identify source catalogs independently of each other and DO NOT use a shared common source catalog for photometry.

##### 1.2: Preliminaries

###### 1.2.1: Generation of the Bad Pixel Mask

Before any source identification takes place, a bad pixel mask is created to identify regions of the detection image where signal quality is known to be degraded. These are areas near the edge of the image, areas with little to no input image contribution, and areas that contain saturated pixels. To minimize the impact of these regions on source identification and subsequent photometric measurements, the regions flagged in this bad pixel mask are iteratively “grown” for 10 steps using the `ndimage.binary_dilation` scipy tool. Pixels in the immediate vicinity of a given masked region may also be impacted to some degree. As we cannot be fully certain that these pixels are or are not impacted, or to the degree of the impact, they are all flagged.

###### 1.2.2: Detection Image Background Subtraction

To ensure optimal source detection, the multi-filter detection image is background-subtracted. We computed a 2-dimensional background image using the `photutils.background.Background2d` Astropy tool. This algorithm uses sigma-clipped statistics to determine background and RMS values across

the image. An initial low-resolution estimate of the background is performed by computing sigma-clipped median values in 27x27 pixel boxes across the image. This low-resolution background image is then median-filtered using a 3x3 pixel sample window to correct for local small-scale overestimates and/or underestimates. It should be noted these are configurable values. Our catalogs use these values deeming them to be the best for the general situation, but users can tune these values to optimize for their own data. To this end, users can adjust parameter values “bkg\_box\_size” and/or “bkg\_filter\_size” in the <instrument>\_<detector>\_catalog\_generation\_all.json files in the following path: /drizzlepac/pars/hap\_pars/default\_parameters/<instrument>/<detector>/.

### 1.3: Source Identification with DAOSTarFinder

We use the `photutils.detection.DAOSTarFinder` Astropy tool to identify sources in the background-subtracted multi-filter detection image. Regions flagged in the previously created bad pixel mask are ignored by DAOSTarFinder. This algorithm works by identifying local brightness maxima with roughly gaussian distributions whose peak values are above a predefined minimum threshold. Full details of the process are described in Stetson 1987; PASP 99, 191. The exact set of input parameters fed into DAOSTarFinder is detector-dependent. The parameters can be found in the <instrument>\_<detector>\_catalog\_generation\_all.json files mentioned in the previous section.

## 2: Aperture Photometry Measurement

### 2.1: Flux determination

Aperture photometry is then performed on the previously identified sources using a pair of concentric photometric apertures. The sizes of these apertures depend on the specific detector being used, and are listed below in table 1:

Table 1: Table 1: Aperture photometry aperture sizes

Instrument/Detector	Inner aperture size (arcsec)	Outer aperture size (arcsec)
ACS/HRC	0.03	0.125
ACS/SBC	0.07	0.125
ACS/WFC	0.05	0.15
WFC3/IR	0.15	0.45
WFC3/UVIS	0.05	0.15

Raw (non-background-subtracted) flux values are computed by summing up the enclosed flux within the two specified apertures using the `photutils.aperture.aperture_photometry` tool. Input values are detector-dependent, and can be found in the \*\_catalog\_generation\_all.json files described above in section 1.3.

Local background values are computed based on the 3-sigma-clipped mode of pixel values present in a circular annulus with an inner radius of 0.25 arcseconds and an outer radius of 0.50 arcseconds surrounding each identified source. This local background value is then subtracted from the raw inner and outer aperture flux values to compute the background-subtracted inner and outer aperture flux values found in the output .ecsv catalog file by the formula

$$f_{bgs} = f_{raw} - f_{bg} \cdot a$$



where

- $f_{bgs}$  is the background-subtracted flux, in electrons per second
- $f_{raw}$  is the raw, non-background-subtracted flux, in electrons per second
- $f_{bg}$  is the per-pixel background flux, in electrons per second per pixel
- $a$  is the area of the photometric aperture, in pixels

The overall standard deviation and mode values of pixels in the background annulus are also reported for each identified source in the output .ecsv catalog file in the “STDEV” and “MSKY” columns respectively (see Section 3 for more details).

## 2.2: Calculation of photometric errors

### 2.2.1: Calculation of flux uncertainties

For every identified source, the `photutils.aperture_photometry()` tool calculates standard deviation values for each aperture based on a 2-dimensional RMS array computed using the `photutils.background.Background2d()` tool that we previously utilized to compute the 2-dimensional background array in order to background-subtract the detection image for source identification. We then compute the final flux errors as seen in the output .ecsv catalog file using the following formula:

$$\Delta f = \sqrt{\frac{\sigma^2}{g} + (a \cdot \sigma_{bg}^2) \cdot \left(1 + \frac{a}{n_{sky}}\right)}$$

where

- $\Delta f$  is the flux uncertainty, in electrons per second
- $\sigma$  is the standard deviation of photometric aperture signal, in counts per second
- $g$  is effective gain in electrons per count
- $a$  is the photometric aperture area, in pixels
- $\sigma_{bg}$  is standard deviation of the background
- $n_{sky}$  is the sky annulus area, in pixels

### 2.2.2: Calculation of ABmag uncertainties

Magnitude error calculation comes from computing  $\frac{d(ABMAG)}{d(flux)}$ . We use the following formula:

$$\Delta mag_{AB} = 1.0857 \cdot \frac{\Delta f}{f}$$

where

- $\Delta mag_{AB}$  is the uncertainty in AB magnitude
- $\Delta f$  is the flux uncertainty, in electrons per second

- $f$  is the flux, in electrons per second

## 2.3: Calculation of concentration index (CI) values and flag values

### 2.3.1: Calculation of concentration index (CI) values

The Concentration index is a measure of the “sharpness” of a given source’s PSF, and computed with the following formula:

$$CI = m_{inner} - m_{outer}$$

where

- $CI$  is the concentration index, in AB magnitude
- $m_{inner}$  is the inner aperture AB magnitude
- $m_{outer}$  is the outer aperture AB magnitude

We use the concentration index to automatically classify each identified photometric source as either a point source (i.e. stars), an extended source (i.e. galaxies, nebosity, etc.), or as an “anomalous” source (i.e. saturation, hot pixels, cosmic ray hits, etc.). This designation is described by the value in the “flags” column

### 2.3.2: Determination of flag values

The flag value associated with each source provides users with a means to distinguish between legitimate point sources, legitimate extended sources, and scientifically dubious sources (those likely impacted by low signal to noise, detector artifacts, saturation, cosmic rays, etc.). The values in the “flags” column of the catalog are a sum of a one or more of these values. Specific flag values are defined below in table 2:

Table 2: Table 2: Flag definitions

Flag value	Meaning
0	Point source ( $CI_{lower} < CI < CI_{upper}$ )
1	Extended source ( $CI > CI_{upper}$ )
2	Bit value 2 not used in ACS or WFC3 sourcelists
4	Saturated Source
8	Faint Detection Limit
16	Hot pixels ( $CI < CI_{lower}$ )
32	False Detection: Swarm Around Saturated Source
64	False detection due proximity of source to image edge or other region with a low number of input images

#### 2.3.2.1: Assignment of flag values 0 (point source), 1 (extended source), and 16 (hot pixels)

Assignment of flag values 0 (point source), 1 (extended source), and 16 (hot pixels) are determined purely based on the concentration index (CI) value. The majority of commonly used filters for all ACS and WFC3

detectors have filter-specific CI threshold values that are automatically set at run-time. However, if filter-specific CI threshold values cannot be found, default instrument/detector-specific CI limits are used instead. Instrument/detector/filter combinations that do not have filter-specific CI threshold values are listed below in table 3 and the default CI values are listed below in table 4.

Table 3: Table 3: Instrument/detector/filter combinations that **do not** have filter-specific CI threshold values

Instrument/Detector	Filters without specifically defined CI limits
ACS/HRC	F344N
ACS/SBC	All ACS/SBC filters
ACS/WFC	F892N
WFC3/IR	None
WFC3/UVIS	None

---

**Note:** As photometry is not performed on observations that utilized grisms, prisms, polarizers, ramp filters, or quad filters, these elements were omitted from the above list.

---

Table 4: Table 4: Default concentration index threshold values

Instrument/Detector	$CI_{lower}$	$CI_{upper}$
ACS/HRC	0.9	1.6
ACS/SBC	0.15	0.45
ACS/WFC	0.9	1.23
WFC3/IR	0.25	0.55
WFC3/UVIS	0.75	1.0

### 2.3.2.2: Assignment of flag value 4 (Saturated Source)

A flag value of 4 is assigned to sources that are saturated. The process of identifying saturated sources starts by first transforming the input image XY coordinates of all pixels flagged as saturated in the data quality arrays of each input flc/ft.fits images (the images drizzled together to produce the drizzle-combined filter image being used to measure photometry) from non-rectified, non-distortion-corrected coordinates to the rectified, distortion-corrected frame of reference of the filter-combined image. We then identify impacted sources by cross-matching this list of saturated pixel coordinates against the positions of sources in the newly created source catalog and assign flag values where necessary.

### 2.3.2.3: Assignment of flag value 8 (faint detection limit)

A flag value of 8 is assigned to sources whose signal to noise ratio is below a predefined value. We define sources as being above the faint object limit if the following is true:

$$\Delta ABmag_{outer} \leq \frac{2.5}{snr \cdot \log(10)}$$

## Where

- $\Delta ABmag_{outer}$  is the outer aperture AB magnitude uncertainty
- *snr* is the signal to noise ratio, which is 1.5 for ACS/WFC and 5.0 for all other detectors.

### 2.3.2.4: Assignment of flag value 32 (false detection: swarm around saturated source)

The source identification routine has been shown to identify false sources in regions near bright or saturated sources, and in image artifacts associated with bright or saturated sources, such as diffraction spikes, and in the pixels surrounding saturated PSF where the brightness level “plateaus” at saturation. We identify impacted sources by locating all sources within a predefined radius of a given source and checking if the brightness of each of these surrounding sources is less than a radially-dependent minimum brightness value defined by a pre-defined stepped encircled energy curve. The parameters used to determine assignment of this flag are instrument-dependent, can be found in the “swarm filter” section of the \*\_quality\_control\_all.json files in the path described above in section 1.3.

### 2.3.2.5: Assignment of flag value 64 (False detection due proximity of source to image edge or other region with a low number of input images)

Sources flagged with a value of 64 are flagged as “bad” because they are inside of or in close proximity to regions characterized by low or null input image contribution. These are areas where for some reason or another, very few or no input images contributed to the pixel value(s) in the drizzle-combined image. We identify sources impacted with this effect by creating a two-dimensional weight image that maps the number of contributing exposures for every pixel. We then check each source against this map to ensure that all sources are flagged appropriately.

## 3: The output catalog file

### 3.1: Filename format

Source positions and photometric information are written to a .ecsv (Enhanced Character Separated Values) file. The naming of this file is fully automatic and follows the following format: <TELESCOPE>\_<PROPOSAL ID>\_<OBSERVATION SET ID>\_<INSTRUMENT>\_<DETECTOR>\_<FILTER>\_<DATASET NAME>\_<CATALOG TYPE>.ecsv

So, for example if we have the following information:

- Telescope = HST
- Proposal ID = 98765
- Observation set ID = 43
- Instrument = acs
- Detector = wfc
- Filter name = f606w

- Dataset name = j65c43
- Catalog type = point\_cat

**The resulting auto-generated catalog filename will be:**

- hst\_98765\_43\_acs\_wfc\_f606w\_j65c43\_point-cat.ecsv

### **3.2: File format**

The .ecsv file format is quite flexible and allows for the storage of not only character-separated datasets, but also metadata. The first section (lines 4-17) contains a mapping that defines the datatype, units, and formatting information for each data table column. The second section (lines 19-27) contains information explaining STScI's use policy for HAP data in refereed publications. The third section (lines 28-48) contains relevant image metadata. This includes the following items:

- WCS (world coordinate system) name
- WCS (world coordinate system) type
- Proposal ID
- Image filename
- Target name
- Observation date
- Observation time
- Instrument
- Detector
- Target right ascension
- Target declination
- Orientation
- Aperture right ascension
- Aperture declination
- Aperture position angle
- Exposure start (MJD)
- Total exposure duration in seconds
- CCD Gain
- Filter name
- Total Number of sources in catalog

The next section (lines 50-66) contains important notes regarding the coordinate systems used, magnitude system used, apertures used, concentration index definition and flag value definitions:

- X, Y coordinates listed below use are zero-indexed (origin = 0,0)
- RA and Dec values in this table are in sky coordinates (i.e. coordinates at the epoch of observation and fit to GAIADR1 (2015.0) or GAIADR2 (2015.5)).
- Magnitude values in this table are in the ABMAG system.
- Inner aperture radius in pixels and arcseconds (based on detector platescale)
- Outer aperture radius in pixels and arcseconds (based on detector platescale)
- Concentration index (CI) formulaic definition
- Flag value definitions

Finally, the last section contains the catalog of source locations and photometry values. It should be noted that the specific columns and their ordering were deliberately chosen to facilitate a 1:1 exact mapping to the `_daophot.txt` catalogs produced by Hubble Legacy Archive. As this code was designed to be the HLA's replacement, we sought to minimize any issues caused by the transition. The column names are as follows (Note that this is the same left-to-right ordering in the `.ecsv` file as well):

- X-Center: 0-indexed X-coordinate position
- Y-Center: 0-indexed Y-coordinate position
- RA: Right ascension (sky coordinates), in degrees
- DEC: Declination (sky coordinates), in degrees
- ID: Object catalog index number
- MagAp1: Inner aperture brightness, in AB magnitude
- MagErrAp1: Inner aperture brightness uncertainty, in AB magnitude
- MagAp2: Outer aperture brightness, in AB magnitude
- MagErrAp2: Outer aperture brightness uncertainty, in AB magnitude
- MSkyAp2: Outer aperture background brightness, in AB magnitude
- StdevAp2: Standard deviation of the outer aperture background brightness, in AB magnitude
- FluxAp2: Outer aperture flux, in electrons/sec
- CI: Concentration index ( $\text{MagAp1} - \text{MagAp2}$ ), in AB magnitude
- Flags: See Section 2.3.2 for flag value definitions

### 14.6.2 Segment Photometric Catalog Generation

Michele's documentation goes here!

## 14.7 Functions to Manage WCS Table Extension

These functions provide the basic support for initializing, creating and updating the WCS table extension which serves as the archive of updates made to the WCS information in the image headers.

`stwcs.wcsutil.wcscorr.archive_wcs_file` (*image*, *wcs\_id=None*)

Update WCSCORR table with rows for each SCI extension to record the newly updated WCS key-word values.

`stwcs.wcsutil.wcscorr.create_wcscorr` (*descrip=False*, *numrows=1*, *padding=0*)

Return the basic definitions for a WCSCORR table. The dtype definitions for the string columns are set to the maximum allowed so that all new elements will have the same max size which will be automatically truncated to this limit upon updating (if needed).

The table is initialized with rows corresponding to the OPUS solution for all the ‘SCI’ extensions.

`stwcs.wcsutil.wcscorr.delete_wcscorr_row` (*wcstab*, *selections=None*, *rows=None*)

Sets all values in a specified row or set of rows to default values

This function will essentially erase the specified row from the table without actually removing the row from the table. This avoids the problems with trying to resize the number of rows in the table while preserving the ability to update the table with new rows again without resizing the table.

### Parameters

**wcstab: object** PyFITS binTable object for WCSCORR table

**selections: dict** Dictionary of wcscorr column names and values to be used to select the row or set of rows to erase

**rows: int, list** If specified, will specify what rows from the table to erase regardless of the value of ‘selections’

`stwcs.wcsutil.wcscorr.find_wcscorr_row` (*wcstab*, *selections*)

Return an array of indices from the table (NOT HDU) ‘wcstab’ that matches the selections specified by the user.

The row selection criteria must be specified as a dictionary with column name as key and value(s) representing the valid desired row values. For example, {‘wcs\_id’:‘OPUS’,‘extver’:2}.

`stwcs.wcsutil.wcscorr.init_wcscorr` (*input*, *force=False*)

This function will initialize the WCSCORR table if it is not already present, and look for WCS keywords with a prefix of ‘O’ as the original OPUS generated WCS as the initial row for the table or use the current WCS keywords as initial row if no ‘O’ prefix keywords are found.

This function will NOT overwrite any rows already present.

This function works on all SCI extensions at one time.

`stwcs.wcsutil.wcscorr.restore_file_from_wcscorr` (*image*, *id=‘OPUS’,*  
*wcskey=’’*)

Copies the values of the WCS from the WCSCORR based on ID specified by user. The default will be to restore the original OPUS-derived values to the Primary WCS. If *wcskey* is specified, the WCS with that key will be updated instead.

```
stwcs.wcsutil.wcscorr.update_wcscorr (dest, source=None, extname='SCI',
                                       wcs_id=None, active=True)
```

Update WCSCORR table with a new row or rows for this extension header. It copies the current set of WCS keywords as a new row of the table based on keyed WCSs as per Paper I Multiple WCS standard).

#### Parameters

**dest** [HDUList] The HDU list whose WCSCORR table should be appended to (the WCSCORR HDU must already exist)

**source** [HDUList, optional] The HDU list containing the extension from which to extract the WCS keywords to add to the WCSCORR table. If None, the dest is also used as the source.

**extname** [str, optional] The extension name from which to take new WCS keywords. If there are multiple extensions with that name, rows are added for each extension version.

**wcs\_id** [str, optional] The name of the WCS to add, as in the WCSNAMEa keyword. If unspecified, all the WCSs in the specified extensions are added.

**active: bool, optional** When True, indicates that the update should reflect an update of the active WCS information, not just appending the WCS to the file as a headerlet

```
stwcs.wcsutil.wcscorr.update_wcscorr_column (wcstab, column, values, selections=None, rows=None)
```

Update the values in 'column' with 'values' for selected rows

#### Parameters

**wcstab: object** PyFITS binTable object for WCSCORR table

**column: string** Name of table column with values that need to be updated

**values: string, int, or list** Value or set of values to copy into the selected rows for the column

**selections: dict** Dictionary of wcscorr column names and values to be used to select the row or set of rows to erase

**rows: int, list** If specified, will specify what rows from the table to erase regardless of the value of 'selections'

## 14.8 Functions to Manage Legacy OPUS WCS Keywords in the WCS Table

The previously released versions of `makewcs` provided with `MultiDrizzle archives` the original OPUS generated WCS keywords using header keywords which have a prefix of "O", such as "OCRPIX1". In order to avoid overwriting or ignoring these original values, these functions can be used to convert the prefixed OPUS WCS keywords into WCS table entries compatible with the new code.



Strictly to provide complete support for these OPUS keywords, the code will also create, if the user desires, prefix “O” WCS keywords from the alternate WCS FITS conventions OPUS keywords. This would allow images processed using the new code only can then be used with older versions of `MultiDrizzle`, if the user needs such compatibility.

`stwcs.wcsutil.convertwcs.archive_prefix_OPUS_WCS (fobj, extname='SCI')`

Identifies WCS keywords which were generated by OPUS and archived using a prefix of ‘O’ for all ‘SCI’ extensions in the file

#### Parameters

**fobj** [str or `astropy.io.fits.HDUList`] Filename or fits object of a file

`stwcs.wcsutil.convertwcs.create_prefix_OPUS_WCS (fobj, extname='SCI')`

Creates alternate WCS with a prefix of ‘O’ for OPUS generated WCS values to work with old `MultiDrizzle`.

#### Parameters

**fobj** [str or `astropy.io.fits.HDUList`] Filename or fits object of a file

#### Raises

**IOError:** if input FITS object was not opened in ‘update’ mode

## 14.9 TWEAKUTILS: Utility Functions for `Tweakreg`

The functions in this module support the various aspects of `TweakReg`, including finding the objects in the images and plotting the residuals.

**Authors** Warren Hack

**License** *LICENSE*

`drizzlepac.tweakutils.parse_atfile_cat (input)`

Return the list of catalog filenames specified as part of the input @-file

`drizzlepac.tweakutils.isfloat (value)`

Return True if all characters are part of a floating point value

`drizzlepac.tweakutils.parse_skypos (ra, dec)`

Function to parse RA and Dec input values and turn them into decimal degrees

**Input formats could be:** [“nn”,”nn”,”nn.nn”] “nn nn nn.nnn” “nn:nn:nn.nn” “nnH nnM nn.nnS” or “nnD nnM nn.nnS” nn.nnnnnnnn “nn.nnnnnnnn”

`drizzlepac.tweakutils.radec_hmstodd (ra, dec)`

Function to convert HMS values into decimal degrees.

This function relies on the `astropy.coordinates` package to perform the conversion to decimal degrees.

#### Parameters

**ra** [list or array] List or array of input RA positions

**dec** [list or array] List or array of input Dec positions

**Returns**

**pos** [arr] Array of RA,Dec positions in decimal degrees

**See also:**

`astropy.coordinates`

**Notes**

This function supports any specification of RA and Dec as HMS or DMS; specifically, the formats:

```
["nn", "nn", "nn.nn"]
"nn nn nn.nnn"
"nn:nn:nn.nn"
"nnH nnM nn.nnS" or "nnD nnM nn.nnS"
```

`drizzlepac.tweakutils.parse_exclusions` (*exclusions*)

Read in exclusion definitions from file named by 'exclusions' and return a list of positions and distances

`drizzlepac.tweakutils.parse_colname` (*colname*)

Common function to interpret input column names provided by the user.

This function translates column specification provided by the user into a column number.

**Parameters**

**colname** : Column name or names to be interpreted

**Returns**

**cols** [list] The return value will be a list of strings.

**Notes**

This function will understand the following inputs:

```
'1,2,3' or 'c1,c2,c3' or ['c1','c2','c3']
'1-3' or 'c1-c3'
'1:3' or 'c1:c3'
'1 2 3' or 'c1 c2 c3'
'1' or 'c1'
1
```

`drizzlepac.tweakutils.readcols` (*infile, cols=None*)

Function which reads specified columns from either FITS tables or ASCII files

This function reads in the columns specified by the user into numpy arrays regardless of the format of the input table (ASCII or FITS table).

**Parameters**

**infile** [string] Filename of the input file

**cols** [string or list of strings] Columns to be read into arrays

### Returns

**outarr** [array] Numpy array or arrays of columns from the table

`drizzlepac.tweakutils.read_FITS_cols(infile, cols=None)`

Read columns from FITS table

`drizzlepac.tweakutils.read_ASCII_cols(infile, cols=[1, 2, 3])`

Interpret input ASCII file to return arrays for specified columns.

### Returns

**outarr** [list of arrays] The return value will be a list of numpy arrays, one for each 'column'.

### Notes

The specification of the columns should be expected to have lists for each 'column', with all columns in each list combined into a single entry.

For example:

```
cols = ['1,2,3', '4,5,6', 7]
```

where '1,2,3' represent the X/RA values, '4,5,6' represent the Y/Dec values and 7 represents the flux value for a total of 3 requested columns of data to be returned.

`drizzlepac.tweakutils.write_shiftfile(image_list, filename, out-wcs='tweak_wcs.fits')`

Write out a shiftfile for a given list of input Image class objects

`drizzlepac.tweakutils.createWcsHDU(wcs)`

Generate a WCS header object that can be used to populate a reference WCS HDU.

For most applications, `stwcs.wcsutil.HSTWCS.wcs2header()` will work just as well.

`drizzlepac.tweakutils.idlgauss_convolve(image, fwhm)`

Deprecated since version 3.0.0: The `idlgauss_convolve` function is deprecated and may be removed in a future version.

`drizzlepac.tweakutils.gauss_array(nx, ny=None, fwhm=1.0, sigma_x=None, sigma_y=None, zero_norm=False)`

Computes the 2D Gaussian with size `nx*ny`.

### Parameters

**nx** [int]

**ny** [int [Default: None]] Size of output array for the generated Gaussian. If `ny == None`, output will be an array `nx X nx` pixels.

**fwhm** [float [Default: 1.0]] Full-width, half-maximum of the Gaussian to be generated

**sigma\_x** [float [Default: None]]

**sigma\_y** [float [Default: None]] Sigma\_x and sigma\_y are the stddev of the Gaussian functions.

**zero\_norm** [bool [Default: False]] The kernel will be normalized to a sum of 1 when True.

### Returns

**gauss\_arr** [array] A numpy array with the generated gaussian function

`drizzlepac.tweakutils.gauss(x, sigma)`

Compute 1-D value of gaussian at position x relative to center.

`drizzlepac.tweakutils.make_vector_plot(coordfile, columns=[1, 2, 3, 4], data=None, figure_id=None, title=None, axes=None, every=1, labelsize=8, ylimit=None, limit=None, xlower=None, ylower=None, output=None, headl=4, headw=3, xsh=0.0, ysh=0.0, fit=None, scale=1.0, vector=True, textscale=5, append=False, linfit=False, rms=True, plotname=None)`

Convert a XYXYMATCH file into a vector plot or set of residuals plots.

This function provides a single interface for generating either a vector plot of residuals or a set of 4 plots showing residuals. The data being plotted can also be adjusted for a linear fit on-the-fly.

### Parameters

**coordfile** [string] Name of file with matched sets of coordinates. This input file can be a file compatible for use with IRAF's geomap.

**columns** [list [Default: [0,1,2,3]]] Column numbers for the X,Y positions from each image

**data** [list of arrays] If specified, this can be used to input matched data directly

**title** [string] Title to be used for the generated plot

**axes** [list] List of X and Y min/max values to customize the plot axes

**every** [int [Default: 1]] Slice value for the data to be plotted

**limit** [float] Radial offset limit for selecting which sources are included in the plot

**labelsiz** [int [Default: 8] or str] Font size to use for tick labels, either in font points or as a string understood by tick\_params().

**ylimit** [float] Limit to use for Y range of plots.

**xlower** [float]

**ylower** [float] Limit in X and/or Y offset for selecting which sources are included in the plot

**output** [string] Filename of output file for generated plot

**headl** [int [Default: 4]] Length of arrow head to be used in vector plot

**headw** [int [Default: 3]] Width of arrow head to be used in vector plot

**xsh** [float]

**ysh** [float] Shift in X and Y from linear fit to be applied to source positions from the first image

**scale** [float] Scale from linear fit to be applied to source positions from the first image

**fit** [array] Array of linear coefficients for rotation (and scale?) in X and Y from a linear fit to be applied to source positions from the first image

**vector** [bool [Default: True]] Specifies whether or not to generate a vector plot. If False, task will generate a set of 4 residuals plots instead

**textscales** [int [Default: 5]] Scale factor for text used for labelling the generated plot

**append** [bool [Default: False]] If True, will overplot new plot on any pre-existing plot

**linfit** [bool [Default: False]] If True, a linear fit to the residuals will be generated and added to the generated residuals plots

**rms** [bool [Default: True]] Specifies whether or not to report the RMS of the residuals as a label on the generated plot(s).

**plotname** [str [Default: None]] Write out plot to a file with this name if specified.

`drizzlepac.tweakutils.find_xy_peak` (*img*, *center=None*, *sigma=3.0*)

Deprecated since version 3.0.0: The `find_xy_peak` function is deprecated and may be removed in a future version.

Find the center of the peak of offsets

`drizzlepac.tweakutils.plot_zeropoint` (*pars*)

Plot 2d histogram.

**Pars will be a dictionary containing:** *data*, *figure\_id*, *vmax*, *title\_str*, *xp,yp*, *searchrad*

`drizzlepac.tweakutils.build_xy_zeropoint` (*imgxy*, *refxy*, *searchrad=3.0*, *hist-plot=False*, *figure\_id=1*, *plot-name=None*, *interactive=True*)

Deprecated since version 3.0.0: The `build_xy_zeropoint` function is deprecated and may be removed in a future version.

**Create a matrix which contains the delta between each XY position and** each UV position.

`drizzlepac.tweakutils.build_pos_grid` (*start*, *end*, *nstep*, *mesh=False*)

Deprecated since version 3.0.0: The `build_pos_grid` function is deprecated and may be removed in a future version.

Return a grid of positions starting at X,Y given by ‘start’, and ending at X,Y given by ‘end’. The grid will be completely filled in X and Y by every ‘step’ interval.

## 14.10 UPDATEHDR: Functions for Updating WCS with New Solutions

The functions in this module support updating the WCS information in distorted images with the alignment solution determined by `TweakReg` or saved in a shiftfile.

**Authors** Warren Hack, Mihai Cara

**License** *LICENSE*

`drizzlepac.updatehdr.create_unique_wcsname` (*fimg*, *extnum*, *wcsname*)

This function evaluates whether the specified `wcsname` value has already been used in this image. If so, it automatically modifies the name with a simple version ID using `wcsname_NNN` format.

### Parameters

**fimg** [obj] PyFITS object of image with WCS information to be updated

**extnum** [int] Index of extension with WCS information to be updated

**wcsname** [str] Value of WCSNAME specified by user for labelling the new WCS

### Returns

**uniqname** [str] Unique WCSNAME value

`drizzlepac.updatehdr.interpret_wcsname_type` (*wcsname*)

Interpret WCSNAME as a standardized human-understandable description

`drizzlepac.updatehdr.linearize` (*wcsim*, *wcsima*, *wcsref*, *imcrpix*, *f*, *shift*, *hx=1.0*,  
*hy=1.0*)

linearization using 5-point formula for first order derivative

`drizzlepac.updatehdr.update_from_shiftfile` (*shiftfile*, *wcsname=None*,  
*force=False*)

Update headers of all images specified in `shiftfile` with shifts from `shiftfile`.

### Parameters

**shiftfile** [str] Filename of `shiftfile`.

**wcsname** [str] Label to give to new WCS solution being created by this fit. If a value of `None` is given, it will automatically use ‘TWEAK’ as the label. [Default=`None`]

**force** [bool] Update header even though WCS already exists with this solution or `wcsname`? [Default=`False`]

`drizzlepac.updatehdr.update_refchip_with_shift` (*chip\_wcs*, *wcslin*, *fit-geom='rscale'*, *rot=0.0*,  
*scale=1.0*, *xsh=0.0*, *ysh=0.0*,  
*fit=None*, *xrms=None*,  
*yrms=None*)

Compute the matrix for the scale and rotation correction

### Parameters

- chip\_wcs:** **wcs object** HST of the input image
- wcslin:** **wcs object** Reference WCS from which the offsets/rotations are determined
- fitgeom:** **str** NOT USED
- rot** [float] Amount of rotation measured in fit to be applied. [Default=0.0]
- scale** [float] Amount of scale change measured in fit to be applied. [Default=1.0]
- xsh** [float] Offset in X pixels from defined tangent plane to be applied to image. [Default=0.0]
- ysh** [float] Offset in Y pixels from defined tangent plane to be applied to image. [Default=0.0]
- fit** [arr] Linear coefficients for fit [Default = None]
- xrms** [float] RMS of fit in RA (in decimal degrees) that will be recorded as CRDER1 in WCS and header [Default = None]
- yrms** [float] RMS of fit in Dec (in decimal degrees) that will be recorded as CRDER2 in WCS and header [Default = None]

```
drizzlepac.updatehdr.update_wcs(image, extnum, new_wcs, wcsname="", reuse-
                                name=False, verbose=False)
```

Updates the WCS of the specified extension number with the new WCS after archiving the original WCS.

The value of 'new\_wcs' needs to be the full HSTWCS object.

### Parameters

- image** [str] Filename of image with WCS that needs to be updated
- extnum** [int] Extension number for extension with WCS to be updated/replaced
- new\_wcs** [object] Full HSTWCS object which will replace/update the existing WCS
- wcsname** [str] Label to give newly updated WCS
- reusename** [bool] User can choose whether to over-write WCS with same name or not. [Default: False]
- verbose** [bool, int] Print extra messages during processing? [Default: False]

```
drizzlepac.updatehdr.updatewcs_with_shift(image, reference, wc-
                                           sname='TWEAK', reusename=False,
                                           fitgeom='rscale', rot=0.0, scale=1.0,
                                           xsh=0.0, ysh=0.0, fit=None,
                                           xrms=None, yrms=None, ver-
                                           bose=False, force=False, sci-
                                           ext='SCI')
```

Update the SCI headers in 'image' based on the fit provided as determined in the WCS specified by 'reference'. The fit should be a 2-D matrix as generated for use with 'make\_vector\_plot()'.

## Parameters

- image** [str or PyFITS.HDUList object] Filename, or PyFITS object, of image with WCS to be updated. All extensions with EXTNAME matches the value of the ‘sciext’ parameter value (by default, all ‘SCI’ extensions) will be updated.
- reference** [str] Filename of image/headerlet (FITS file) which contains the WCS used to define the tangent plane in which all the fit parameters (shift, rot, scale) were measured.
- wcsname** [str, None, optional] Label to give to new WCS solution being created by this fit. If a value of None is given, it will automatically use ‘TWEAK’ as the label. [Default =None]
- reusename** [bool] User can specify whether or not to over-write WCS with same name. [Default: False]
- rot** [float] Amount of rotation measured in fit to be applied. [Default=0.0]
- scale** [float] Amount of scale change measured in fit to be applied. [Default=1.0]
- xsh** [float] Offset in X pixels from defined tangent plane to be applied to image. [Default=0.0]
- ysh** [float] Offset in Y pixels from defined tangent plane to be applied to image. [Default=0.0]
- fit** [arr] Linear coefficients for fit [Default = None]
- xrms** [float] RMS of fit in RA (in decimal degrees) that will be recorded as CRDER1 in WCS and header [Default = None]
- yrms** [float] RMS of fit in Dec (in decimal degrees) that will be recorded as CRDER2 in WCS and header [Default = None]
- verbose** [bool] Print extra messages during processing? [Default=False]
- force** [bool] Update header even though WCS already exists with this solution or wcsname? [Default=False]
- sciext** [string] Value of FITS EXTNAME keyword for extensions with WCS headers to be updated with the fit values. [Default=‘SCI’]

## Notes

The algorithm used to apply the provided fit solution to the image involves applying the following steps to the WCS of each of the input image’s chips:

1. **compute RA/Dec with full distortion correction for** reference point as (Rc\_i,Dc\_i)
2. **find the Xc,Yc for each Rc\_i,Dc\_i and get the difference from the** CRPIX position for the reference WCS as (dXc\_i,dYc\_i)
3. **apply fit (rot&scale) to (dXc\_i,dYc\_i) then apply shift, then add** CRPIX back to get new (Xcs\_i,Ycs\_i) position



4. compute (Rcs\_i,Dcs\_i) as the sky coordinates for (Xcs\_i,Ycs\_i)
5. compute delta of (Rcs\_i-Rc\_i, Dcs\_i-Dcs\_i) as (dRcs\_i,dDcs\_i)
6. **apply the fit to the chip's undistorted CD matrix, the apply linear** distortion terms back in to create a new CD matrix
7. add (dRcs\_i,dDcs\_i) to CRVAL of the reference chip's WCS
8. update header with new WCS values

## 14.11 Region mapping for TweakReg

This module provides functions for mapping DS9 region files given in sky coordinates to DS9 region files specified in image coordinates of multiple images using the WCS information from the images.

**Authors** Mihai Cara

**License** *LICENSE*

```
drizzlepac.mapreg.MapReg (input_reg, images, img_wcs_ext='sci', refimg="",
                           ref_wcs_ext='sci', chip_reg="", outpath='./regions', filter="",
                           catfname="", interactive=False, append=False, verbose=True)
```

`MapReg()` provides an automated interface for converting a region file to the image coordinate system (CS) of multiple images (and their extensions) using WCS information from the image(s) header(s). This conversion does not take into account pointing errors and, therefore, an examination and adjustment (if required) of output region files is highly recommended. This task is designed to simplify the creation of the exclusions and/or inclusions region files used with `TweakReg()` task for sources finding.

### Parameters

**input\_reg** [string or list of strings (Default = '')] Input region files that need to be mapped to image CS using WCS information from `images` (see below). Only region files saved in sky CS are allowed in this release. Regions specified in image-like coordinates (e.g., image, physical) will be ignored.

This parameter can be provided in any of several forms:

- filename of a single image
- comma-separated list of filenames
- `@-file` filelist containing list of desired input region filenames

The `@-file` filelist needs to be provided as an ASCII text file containing a list of filenames for all input region files with one filename on each line of the file.

**images** [string or list of strings (Default = \*.fits)] FITS images onto which the region files `input_reg` will be mapped. These image files must contain WCS information in their headers in order to convert `input_reg` from sky coordinates to correct image coordinates. This parameter can be provided in any of several forms:

- filename of a single image
- filename of an association (ASN)table
- wild-card specification for files in directory (using \*, ? etc.)
- comma-separated list of filenames
- @-file filelist containing list of desired input filenames (and optional inverse variance map filenames)

The @-file filelist needs to be provided as an ASCII text file containing a list of filenames for all input images (to which `input_reg` regions should be mapped) with one filename on each line of the file.

**img\_wcs\_ext** [string or list of strings (Default = SCI)] Extension name, extension name and version, or extension number of FITS extensions in the `images` to which the input regions `input_reg` should be mapped. The header of each extension must contain WCS information that will be used to convert `input_reg` from sky CS to image-like CS. Multiple extensions must be separated by semi-colon while extension name and version (if present) must be separated by comma, e.g., 'SCI;DQ,1;0'. When specifying the extension name only, internally it will be expanded into a list of extension names and versions for each version of that extension name present in the input `images`. For example, if a FITS file has four SCI and four DQ extensions, then 'SCI;DQ,1;0' will be expanded into 'SCI,1;SCI,2;SCI,3;SCI,4;DQ,1;0'.

**refimg** [string (Default = '')] **Reserved for future use.** Filename of the reference image. May contain extension specifier: [extname,extver], [extname], or [extnumber].

---

**Note:** This parameter is reserved for future use and it is not available through TEAL interface.

---

**ref\_wcs\_ext** [string (Default = SCI)] **Reserved for future use.** Extension name and/or version of FITS extensions in the `refimg` that contain WCS information that will be used to convert `input_reg` from image-like CS to sky CS. NOTE: Only extension name is allowed when `input_reg` is a list of region files that contain regions in image-like CS. In this case, the number of regions in `input_reg` must agree with the number of extensions with name specified by `ref_wcs_ext` present in the `refimg` FITS image.

---

**Note:** This parameter is reserved for future use and it is not available through TEAL interface.

---

**chip\_reg** [string or list of strings (Default = '')] Input region files in image CS associated with each extension specified by the `img_wcs_ext` parameter above. These regions will be added directly (without any transformation) to the

`input_reg` regions mapped to each extension of the input images. These regions must be specified in image-like coordinates. Typically, these regions should contain “exclude” regions to exclude parts of the image specific to the detector **chip** (e.g., vignetted regions due to used filters, or occulting finger in ACS/HRC images) from being used for source finding. This parameter can be provided in one of the following forms:

- filename of a single image (if `img_wcs_ext` specifies a single FITS extension);
- comma-separated list of filenames (if `img_wcs_ext` specifies more than one extension) or `None` for extensions that do not need any chip-specific regions to be excluded/included;
- “” (empty string) or `None` if no chip-specific region files are provided.

The number of regions ideally must be equal to the number of extensions specified by the `img_wcs_ext` parameter. If the number of chip-specific regions is less than the number of `img_wcs_ext` extensions then ‘chip\_reg’ regions will be assigned to the first extensions from `img_wcs_ext` (after internal expansion described in help for the `img_wcs_ext` parameter above). If the number of ‘chip\_reg’ is larger than the number of `img_wcs_ext` extensions then extra regions will be ignored.

**output** [string (Default = `./regions`)] The directory to which the transformed regions should be saved.

**filter** [string {‘None’, ‘fast’, or ‘precise’ } (Default = ‘None’)] Specify whether or not to remove the regions in the transformed region files that are outside the image array. With the ‘fast’ method only intersection of the bounding boxes is being checked.

---

**Note:** The ‘precise’ method is not implemented in this release and, if specified, defaults to ‘fast’. The ‘precise’ option is not available through the TEAL interface.

---

**catfname** [string (Default = `exclusions_cat.txt`)] The file name of the output exclusions catalog file to be created from the supplied image and region file names. This file can be passed as an input to TweakReg task. Verify that the created file is correct!

**append** [bool (Default = False)] Specify whether or not to append the transformed regions to the existing region files with the same name.

**interactive** [bool (Default = False)] **Reserved for future use.** (This switch controls whether the program stops and waits for the user to examine any generated region files before continuing on to the next image.)

---

**Note:** This parameter is reserved for future use and it is not available through

TEAL interface.

---

**verbose** [bool (Default = False)] Specify whether or not to print extra messages during processing.

## Notes

**NOTE 1:** This task takes a region file (or multiple files) that describe(s) what regions of sky should be used for source finding (*include* regions) and what regions should be avoided (*exclude* regions) and transforms/maps this region file onto a number of image files that need to be aligned.

The idea behind this task is automate the creation of region files that then can be passed to *exclusions* parameter of the `TweakReg` task.

The same thing can be achieved manually using, for example, external FITS viewers, e.g., SAO DS9. For example, based on some image `refimg.fits` we can select a few small regions of sky that contain several good (bright, not saturated) point-like sources that could be used for image alignment of other images (say `img1.fits`, `img2.fits`, etc.). We can save this region file in sky coordinates (e.g., `fk5`), e.g., under the name `input_reg.reg`. We can then load a specific extension of each of the images `img1.fits`, `img2.fits`, etc. one by one into DS9 and then load onto those images the previously created include/exclude region file `input_reg.reg`. Now we can save the regions using *image* coordinates. To do conversion from the sky coordinates to image coordinates, DS9 will use the WCS info from the image onto which the region file was loaded. The `MapReg()` task tries to automate this process.

**NOTE 2:** `MapReg()` relies on the `pyregion` package for region file parsing and coordinate transformation. Unfortunately, as of writing, **pyregion** does not consider distortion corrections when performing coordinate transformations. Therefore, there might be a slight discrepancy between the regions produced by `MapReg()` and the DS9 regions obtained as described in the NOTE 1 above.

**NOTE 3:** `MapReg()` does not take into account pointing errors and thus the produced region files can be somewhat misaligned compared to their intended position around the sources identified in the “reference” image. Therefore, it is highly recommended that the produced region files be loaded into DS9 and their position be adjusted manually to include the sources of interest (or to avoid the regions that need to be avoided). If possible, the *include* or *exclude* regions should be large enough as to allow for most pointing errors.

## Examples

Let’s say that based on some image `refimg.fits` we have produced a “master” reference image (`master.reg`) that includes regions around sources that we want to use for image alignment in task `TweakReg()` and excludes regions that we want to avoid being used for image alignment (e.g, diffraction spikes, saturated quasars, stars, etc.). We save the file `master.reg` in sky CS (e.g., `fk5`).

Also, let’s assume that we have a set of images `img1.fits`, `img2.fits`, etc. with four FITS extensions named ‘SCI’ and ‘DQ’. For some of the extensions, after analyzing the `img*.fits` images we have identified parts of the chips that cannot be used for image alignment. We create region

files for those extensions and save the files in image CS as, e.g., `img1_chip_sci2.reg` (in our example this will be the only chip that needs “special” treatment).

Finally, let’s say we want to “replicate” the “master” region file to all SCI extensions of the `img*.fits` images as well as to the 2nd DQ extension and to the 8th extension of the `img*.fits` images.

To do this we run:

```
>>> mapreg(input_reg = 'master.reg', images='img*.fits',
...        img_wcs_ext='sci;dq,2;8', chip_reg='None',
...        img1_chip_sci2.reg, None, None, None, None)
```

This will produce six region files in the `.regions` subdirectory for *each* input image:

```
`img1_sci1_twreg.reg`,    `img1_sci2_twreg.reg`,    `img1_sci3_
↪twreg.reg`,
`img1_sci4_twreg.reg`,    `img1_dq2_twreg.reg`,    `img1_extn8_
↪twreg.reg`
...
```

```
`img2_sci1_twreg.reg`,    `img2_sci2_twreg.reg`,    `img2_sci3_
↪twreg.reg`,
`img2_sci4_twreg.reg`,    `img2_dq2_twreg.reg`,    `img2_extn8_
↪twreg.reg`
...
```

```
drizzlepac.mapreg.map_region_files(input_reg, images, img_wcs_ext='sci', re-
fimg=None, ref_wcs_ext='sci', chip_reg=None,
outpath='./regions', filter=None, catf-
name=None, interactive=False, append=False,
verbose=True)
```

```
drizzlepac.mapreg.help(file=None)
Print out syntax help for running astrodrizzle
```

### Parameters

**file** [str (Default = None)] If given, write out help to the filename specified by this parameter Any previously existing file with this name will be deleted before writing out the help.

## 14.12 Photometric equalization for AstroDrizzle

A tool to adjust data values of images by equalizing each chip’s PHOTFLAM value to a single common value so that all chips can be treated equally by `AstroDrizzle`.

**Authors** Mihai Cara

**License** *LICENSE*

```
drizzlepac.photeq.photeq (files='*_flt.fits', sciext='SCI', errex='ERR', ref_phot=None,
                           ref_phot_ext=None, phot_kwd='PHOTFLAM',
                           aux_phot_kwd='PHOTFNU', search_primary=True, read-
                           only=True, clobber=False, logfile='photeq.log')
```

Adjust data values of images by equalizing each chip's PHOTFLAM value to a single common value so that all chips can be treated equally by AstroDrizzle.

### Parameters

**files** [str (Default = '\*\_flt.fits')] A string containing one of the following:

- a comma-separated list of valid science image file names, e.g.: 'j1234567q\_flt.fits, j1234568q\_flt.fits';
- an @-file name, e.g., '@files\_to\_match.txt'. See notes section for details on the format of the @-files.

---

**Note: Valid science image file names** are:

- file names of existing FITS, GEIS, or WAIVER FITS files;
  - partial file names containing wildcard characters, e.g., '\*\_flt.fits';
  - Association (ASN) tables (must have `_asn`, or `_asc` suffix), e.g., 'j12345670\_asn.fits'.
- 

**sciext** [str (Default = 'SCI')] Extension *name* of extensions whose data and/or headers should be corrected.

**errex** [str (Default = 'ERR')] Extension *name* of the extensions containing corresponding error arrays. Error arrays are corrected in the same way as science data.

**ref\_phot** [float, None (Default = None)] A number indicating the new value of PHOTFLAM or PHOTFNU (set by 'phot\_kwd') to which the data should be adjusted.

**ref\_phot\_ext** [int, str, tuple, None (Default = None)] Extension from which the *photeq* should get the reference photometric value specified by the `phot_kwd` parameter. This parameter is ignored if `ref_phot` is **not** None. When `ref_phot_ext` is None, then the reference inverse sensitivity value will be picked from the first `sciext` of the first input image containing `phot_kwd`.

**phot\_kwd** [str (Default = 'PHOTFLAM')] Specifies the primary keyword which contains inverse sensitivity (e.g., PHOTFLAM). It is used to compute conversion factors by which data should be rescaled.

**aux\_phot\_kwd** [str, None, list of str (Default = 'PHOTFNU')] Same as `phot_kwd` but describes *other* photometric keyword(s) that should be corrected by inverse of the scale factor used to correct data. These keywords are *not* used to compute conversion factors. Multiple keywords can be specified as a Python list of strings: ['PHOTFNU', 'PHOTOHMY'].

---

**Note:** If specifying multiple secondary photometric keywords in the TEAL interface, use a comma-separated list of keywords.

---

**search\_primary** [bool (Default = True)] Specifies whether to first search the primary header for the presence of `phot_kwd` keyword and compute conversion factor based on that value. This is (partially) ignored when `ref_phot` is not `None` in the sense that the value specified by `ref_phot` will be used as the reference *but* in all images primary will be searched for `phot_kwd` and `aux_phot_kwd` and those values will be corrected (if `search_primary=True`).

**readonly** [bool (Default = True)] If `True`, `photeq` will not modify input files (nevertheless, it will convert input GEIS or WAVED FITS files to MEF and could overwrite existing MEF files if `clobber` is set to `True`). The (console or log file) output however will be identical to the case when `readonly=False` and it can be examined before applying these changes to input files.

**clobber** [bool (Default = False)] Overwrite existing MEF files when converting input WAVED FITS or GEIS to MEF.

**logfile** [str, None (Default = 'photeq.log')] File name of the log file.

## Notes

By default, `photeq` will search for the first inverse sensitivity value (given by the header keyword specified by the `phot_kwd` parameter, e.g., PHOTFLAM or PHOTFNU) found in the input images and it will equalize all other images to this reference value.

It is possible to tell `photeq` to look for the reference inverse sensitivity value only in a specific extension of input images, e.g.: 3, ('sci',3), etc. This can be done by setting `ref_phot_ext` to a specific extension. This may be useful, for example, for WFPC2 images: WF3 chip was one of the better calibrated chips, and so, if one prefers to have inverse sensitivities equalized to the inverse sensitivity of the WF3 chip, one can set `ref_phot_ext=3`.

Alternatively, one can provide their own reference inverse sensitivity value to which all other images should be “equalized” through the parameter `ref_phot`.

---

**Note:** Default parameter values (except for `files`, `readonly`, and `clobber`) should be acceptable for most HST images.

---

**Warning:** If images are intended to be used with `AstroDrizzle`, it is recommended that sky background measurement be performed on “equalized” images as the `photeq` is not aware of sky user keyword in the image headers and thus it cannot correct sky values already recorded in the headers.

## Examples

1. In most cases the default parameters should suffice:

```
>>> from drizzlepac import photeq
>>> photeq.photeq(files='*_flt.fits', readonly=False)
```

2. If the re-calibration needs to be done on PHOTFNU rather than PHOTFLAM, then:

```
>>> photeq.photeq(files='*_flt.fits', ref_phot='PHOTFNU',
... aux_phot_kwd='PHOTFLAM')
```

3. If for WFPC2 data one desires that PHOTFLAM from WF3 be used as the reference in WFPC2 images, then:

```
>>> photeq.photeq(files='*_flt.fits', ref_phot_ext=3) # or ('sci',3)
```

`drizzlepac.photeq.help` (*file=None*)

Print out syntax help for running skymatch

### Parameters

**file** [str (Default = None)] If given, write out help to the filename specified by this parameter Any previously existing file with this name will be deleted before writing out the help.

## 14.13 pixreplace: Replace pixels which have one value with another value

This task allows a user to replace pixels in an image or set of images which have one value with a new value; for example, replace all NaNs with a value of -999.9.

Pixreplace – Replace pixels which have one value with another value

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### 14.13.1 PARAMETERS

**input** [str, @-file, list of filenames] Filename(s) of image to be processed.

**pixvalue** [float] Pixel value from `input` file to be replaced. [Default: np.nan]

**newvalue** [float] New pixel value to use to replace `pixvalue`. [Default: 0.0]

**ext** [int, list of ints, None] Extensions from `input` file to process with new pixel values. If None (default), all image extensions (and only image extensions) will be processed.



### 14.13.2 Usage

It can be called from within Python using the syntax:

```
>>> from drizzlepac import pixreplace
>>> pixreplace.replace('adriz_nanSCI_drz.fits')
or
>>> epar pixreplace
```

### 14.13.3 EXAMPLES

1. Replace all pixels in all extensions which have a value of NaN in 'adriz\_nanSCI\_drz.fits' with a constant value of 0.0.

```
>>> from drizzlepac import pixreplace
>>> pixreplace.replace('adriz_nanSCI_drz.fits')
```

2. Replace pixels in first extension only which have a value of NaN in both 'j8c061vnq\_drc.fits' and 'j8c061nyq\_drc.fits' with a constant value of 0.0.

```
>>> pixreplace.replace('j8c061vnq_drc.fits,j8c061nyq_drc.fits', ext=1)
```

3. Replace pixels in first and fourth extensions which have a value of 0.0 in 'adriz\_nanSCI\_drz.fits' with a value of -999.

```
>>> pixreplace.replace('adriz_nanSCI_drz.fits',pixvalue=0.0, newvalue=-999,
↳ext=[1,4])
```

`drizzlepac.pixreplace.replace(input, **pars)`

Replace pixels in `input` that have a value of `pixvalue` with a value given by `newvalue`.



---

## Coordinate Transformation Tasks

---

These tasks support transformations of source positions to and from distorted and drizzled images.

### 15.1 pixtopix: Coordinate transformation to/from drizzled images

This task allows a user to perform coordinate transformations with the full WCS and distortion model to and from drizzled image positions. This task serves as a replacement for the STSDAS.dither task ‘tran’.

`pixtopix` transforms pixel positions given as X,Y values into positions in another WCS frame based on the WCS information and any recognized distortion keywords from the input image header.

**Authors** Warren Hack

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#### 15.1.1 Parameters

**inimage** [str] full filename with path of input image, an extension name ['sci', 1] should be provided if input is a multi-extension FITS file

**outimage** [str, optional] full filename with path of output image, an extension name ['sci', 1] should be provided if output is a multi-extension FITS file. If no image gets specified, the input image will be used to generate a default output WCS using `stwcs.distortion.util.output_wcs()`.

**direction** [str] Direction of transform (forward or backward). The ‘forward’ transform takes the pixel positions (assumed to be from the ‘input’ image) and determines their position in the ‘output’ image. The ‘backward’ transform converts the pixel positions (assumed to be from the ‘output’ image) into pixel positions in the ‘input’ image.

### 15.1.2 Optional Parameters

**x** [float, optional] X position from image

**y** [float, optional] Y position from image

**coords** [str, deprecated] [DEPRECATED] full filename with path of file with x,y coordinates  
Filename given here will be *ignored* if a file has been specified in `coordfile` parameter.

**coordfile** [str, optional] full filename with path of file with starting x,y coordinates

**colnames** [str, optional] comma separated list of column names from 'coords' files containing x,y coordinates, respectively. Will default to first two columns if None are specified. Column names for ASCII files will use 'c1','c2',... convention.

**separator** [str, optional] non-blank separator used as the column delimiter in the coords file

**precision** [int, optional] Number of floating-point digits in output values

**output** [str, optional] Name of output file with results, if desired

**verbose** [bool] Print out full list of transformation results (default: False)

### 15.1.3 RETURNS

**outx** [float] X position of transformed pixel. If more than 1 input value, then it will be a numpy array.

**outy** [float] Y position of transformed pixel. If more than 1 input value, then it will be a numpy array.

### 15.1.4 Notes

This task performs a full distortion-correction coordinate transformation based on all WCS keywords and any recognized distortion keywords from the input image header. The transformation recognizes the conventions for describing distortion implemented as part of the SIP and Paper IV conventions used with Astro-Drizzle. Input images can be updated to use these conventions through the use of the 'updatewcs' module the STWCS package.

### 15.1.5 See Also

stwcs

### 15.1.6 Examples

This task can be run from either the TEAL GUI or from the Python command-line. These examples illustrate the syntax that can be used to run the task in a couple of common modes.

1. Run the task using the TEAL GUI under PyRAF:

```
>>> import drizzlepac
>>> epar pixtopix
```

- Convert the position 256,256 from 'input\_ft.fits[sci,1]' into a position on the output image 'output\_drz.fits[sci,1]' using:

```
>>> from drizzlepac import pixtopix
>>> outx,outy = pixtopix.tran("input_file_ft.fits[sci,1]",
...                          "output_drz.fits[sci,1]", "forward", 256,256)
```

or if you the default direction of 'forward' is already appropriate and you don't want to explicitly set that value:

```
>>> outx,outy = pixtopix.tran("input_file_ft.fits[sci,1]",
...                          "output_drz.fits[sci,1]", x=256,y=256)
```

- The set of X,Y positions from 'output\_drz.fits[sci,1]' stored as the 3rd and 4th columns from the ASCII file xy\_sci1.dat will be transformed into pixel positions from 'input\_ft.fits[sci,1]' and written out to xy\_flt1.dat using:

```
>>> from drizzlepac import pixtopix
>>> x,y = pixtopix.tran("input_ft.fits[sci,1]",
...                    "output_drz.fits[sci,1]", "backward",
...                    coordfile='xy_sci1.dat', colnames=['c3','c4'],
...                    output="xy_flt1.dat")
```

`drizzlepac.pixtopix.tran` (*inimage*, *outimage*, *direction='forward'*, *x=None*, *y=None*, *coordfile=None*, *colnames=None*, *separator=None*, *precision=6*, *output=None*, *verbose=True*)

Primary interface to perform coordinate transformations in pixel coordinates between 2 images using STWCS and full distortion models read from each image's header.

## 15.2 pixtosky: Coordinate transformation to sky coordinates

This task allows a user to perform coordinate transformations with the full WCS and distortion model on source positions from an input image to sky coordinates. This task serves as a replacement for the IRAF.STSDAS task `xy2rd`, albeit with the added capability of understanding the full distortion model provided in the headers of images updated for use with `astrodrizzle` and `tweakreg`.

`pixtosky` transforms pixel positions given as X,Y values into positions on the sky based on the WCS information and any recognized distortion keywords from the input image header.

**Authors** Warren Hack

**License** [http://www.stsci.edu/resources/software\\_hardware/pyraf/LICENSE](http://www.stsci.edu/resources/software_hardware/pyraf/LICENSE)

### 15.2.1 Parameters

**input** [str] full filename with path of input image, an extension name ['sci',1] should be provided if input is a multi-extension FITS file.

### 15.2.2 Optional Parameters

**x** [float, optional] X position from input image

**y** [float, optional] Y position from input image

**coordfile** [str, optional] full filename with path of file with x, y coordinates

**colnames** [str, optional] comma separated list of column names from 'coordfile' file containing x, y coordinates. Will default to first two columns if None are specified. Column names for ASCII files will use 'c1','c2',... convention.

**separator** [str, optional] non-blank separator used as the column delimiter in the `coordfile`

**hms** [bool, optional (Default: False)] Produce output in HH:MM:SS.S format instead of decimal degrees?

**precision** [int, optional] Number of floating-point digits in output values

**output** [str, optional] Name of output file with results, if desired

**verbose** [bool (Default: False)] Print out full list of transformation results

### 15.2.3 Returns

**ra** [float] Right Ascension of pixel. If more than 1 input value, then it will be a numpy array.

**dec** [float] Declination of pixel. If more than 1 input value, then it will be a numpy array.

### 15.2.4 Notes

This task performs a full distortion-correction coordinate transformation based on all WCS keywords and any recognized distortion keywords from the input image header. The transformation recognizes the conventions for describing distortion implemented as part of the SIP and Paper IV conventions used with `AstroDrizzle`. Input images can be updated to use these conventions through the use of the `updatewcs` module the `STWCS` package.

### 15.2.5 See Also

`stwcs`

## 15.2.6 Examples

This task can be run from either the TEAL GUI or from the Python command-line. These examples illustrate the syntax that can be used to run the task in a couple of common modes.

1. Run the task using the TEAL GUI under PyRAF:

```
>>> import drizzlepac
>>> epar pixtosky
```

2. Convert a single X,Y position (100,100) from an calibrated ACS image (`j8bt06nyq_flt.fits`) into an undistorted sky position (RA,Dec) without using the TEAL GUI:

```
>>> from drizzlepac import pixtosky
>>> r,d = pixtosky.xy2rd("j8bt06nyq_flt.fits[sci,1]", 100, 100)
```

3. Convert a list of X,Y positions from the file 'xyfile.dat' for a calibrated ACS image (`j8bt06nyq_flt.fits`) into undistorted sky positions and write out the result to the file 'radec.dat' without using the TEAL GUI:

```
>>> r,d = pixtosky.xy2rd("j8bt06nyq_flt.fits[sci,1]", coordfile="xyfile.
↳ dat",
...     output="radec.dat")
```

4. The set of X,Y positions from 'input\_flt.fits[sci,1]' stored as the 3rd and 4th columns from the ASCII file 'xy\_sci1.dat' will be transformed and written out to 'radec\_sci1.dat' using:

```
>>> from drizzlepac import pixtosky
>>> r,d = pixtosky.xy2rd("input_flt.fits[sci,1]", coordfile='xy_sci1.dat
↳ ',
...     colnames=['c3','c4'], output="radec_sci1.dat")
```

`drizzlepac.pixtosky.xy2rd` (*input*, *x=None*, *y=None*, *coords=None*, *coordfile=None*, *colnames=None*, *separator=None*, *hms=True*, *precision=6*, *output=None*, *verbose=True*)

Primary interface to perform coordinate transformations from pixel to sky coordinates using STWCS and full distortion models read from the input image header.

## 15.3 skytopix: Coordinate transformation from sky coordinates

This task allows a user to perform coordinate transformations with the full WCS and distortion model on source positions from sky coordinates to the WCS defined by an image. This task serves as a replacement for the IRAF.STSDAS task `rd2xy`, albeit with the added capability of understanding the full distortion model provided in the headers of images updated for use with `astrodrizzle` and `tweakreg`.

`skytopix` transforms source positions given as RA, Dec values into pixel positions on the image based on the WCS information and any recognized distortion keywords contained in the input image header.

**Authors** Warren Hack

License [http://www.stsci.edu/resources/software\\_hardware/pyraf/LICENSE](http://www.stsci.edu/resources/software_hardware/pyraf/LICENSE)

### 15.3.1 Parameters

**input** [str] full filename with path of input image, an extension name ['sci',1] should be provided if input is a multi-extension FITS file.

### 15.3.2 Optional Parameters

**ra** [string, optional] RA position in either decimal degrees or HMS format (with or without ':') like '19:10:50.337406303' or '19 10 50.337406303'

**dec** [string, optional] Dec position in either decimal degrees or HMS format (with or without ':') like '-60:2:22.186557409' or '-60 2 22.186557409'

**coordfile** [str, optional] full filename with path of file with sky coordinates

**colnames** [str, optional] comma separated list of column names from 'coordfile' file containing sky coordinates, respectively. Will default to first two columns if None are specified. Column names for ASCII files will use 'c1','c2',... convention.

**precision** [int, optional] Number of floating-point digits in output values

**output** [str, optional] Name of output file with results, if desired

**verbose** [bool] Print out full list of transformation results (default: `False`)

### 15.3.3 Returns

**x** [float] X position of pixel. If more than 1 input value, then it will be a numpy array.

**y** [float] Y position of pixel. If more than 1 input value, then it will be a numpy array.

### 15.3.4 Notes

This task performs a full distortion-correction coordinate transformation based on all WCS keywords and any recognized distortion keywords from the input image header. The transformation recognizes the conventions for describing distortion implemented as part of the SIP and Paper IV conventions used with `AstroDrizzle`. Input images can be updated to use these conventions through the use of the `updatewcs` module the `STWCS` package.

### 15.3.5 See Also

`stwcs`



### 15.3.6 Examples

This task can be run from either the TEAL GUI, or from the Python command-line. These examples illustrate the syntax that can be used to run the task in a couple of common modes.

1. Run the task using the TEAL GUI under PyRAF:

```
>>> import drizzlepac
>>> epar skytopix
```

2. Convert a single sky position (0:22:07.0088,-72:03:05.429) from a calibrated ACS image (j94f05bgq\_fit.fits) into a pixel position (X,Y) without using the TEAL GUI:

```
>>> from drizzlepac import skytopix
>>> x,y = skytopix.rd2xy("j8bt06nyq_fit.fits[sci,1]",
...                     '0:22:07.0088', '-72:03:05.429')
```

3. Convert a list of (undistorted) sky positions from the file,

**‘radec.dat’ for a calibrated ACS image (j8bt06nyq\_fit.fits) into distorted pixel positions, and write out the result to the file ‘xypos.dat’ without using the TEAL GUI:**

```
>>> x,y = skytopix.rd2xy("j8bt06nyq_fit.fits[sci,1]",
...                     coordfile="radec.dat", output="xypos.dat")
```

`drizzlepac.skytopix.rd2xy` (*input, ra=None, dec=None, coordfile=None, colnames=None, precision=6, output=None, verbose=True*)

Primary interface to perform coordinate transformations from pixel to sky coordinates using STWCS and full distortion models read from the input image header.



---

## ACS Header Update Task

---

A task, ‘updatenpol’, has been written to automate the updating of ACS image headers with the filename of the appropriate NPOLFILE based on the DGEOFILE specified in the image header. This task should be used to update all ACS images prior to processing them with ‘astrodrizzle’.

### 16.1 Updatenpol

`updatenpol`: Update the header of ACS file(s) with the names of new NPOLFILE and D2IMFILE reference files for use with the C version of MultiDrizzle (astrodrizzle).

**Authors** Warren Hack

**License** *LICENSE*

**Usage** This task can be run from the operating system command line with:

```
updatenpol [options] input [refdir]
```

#### Command-line Options

**input** The specification of the files to be updated, either as a single filename, an ASN table name, or wild-card specification of a list of files.

**refdir** The name of the directory containing all the new reference files (`*_npl.fits` and `*_d2i.fits` files). If no directory is given, it will look in `jref$` by default.

**-h** Print the help (this text).

**-l** If specified, copy NPOLFILES and D2IMFILES to local directory for use with the input files.

- i If specified, the program will interactively request the exact names of the NPOLFILE and D2IMFILE reference files to be used for updating the header of each file. The value of 'refdir' will be ignored in interactive mode.

**Warning:** It will ask for the names of the NPOLFILE and D2IMFILE for EACH separate INPUT file when the option -i has been specified.

**Example 1.** This command will update all the FLT files in the current directory with the new NPOLFILE and D2IMFILE reference files found in the 'myjref' directory as defined in the environment:

```
updatenpol *flt.fits myjref$
```

**Compatibility with MultiDrizzle** The new version of MultiDrizzle (AstroDrizzle) and updatewcs only work with the new NPOLFILE reference file for the DGEO correction (to replace the use of DGEOFILE). In fact, AstroDrizzle has been extensively modified to prompt the user with a very lengthy explanation on whether it should stop and allow the user to update the header or continue without applying the DGEO correction under circumstances when the NPOLFILE keyword can not be found for ACS.

drizzlepac.updatenpol.**find\_d2ifile** (*flist, detector*)

Search a list of files for one that matches the detector specified.

drizzlepac.updatenpol.**find\_npolfile** (*flist, detector, filters*)

Search a list of files for one that matches the configuration of detector and filters used.

drizzlepac.updatenpol.**getHelpAsString** (*docstring=False, show\_ver=True*)

return useful help from a file in the script directory called `__taskname__.help`

drizzlepac.updatenpol.**help** (*file=None*)

Print out syntax help for running astrodrizzle

### Parameters

**file** [str (Default = None)] If given, write out help to the filename specified by this parameter Any previously existing file with this name will be deleted before writing out the help.

drizzlepac.updatenpol.**run** (*configobj=None, editpars=False*)

Teal interface for running this code.

drizzlepac.updatenpol.**update** (*input, refdir='jref\$', local=None, interactive=False, wcsupdate=True*)

Updates headers of files given as input to point to the new reference files NPOLFILE and D2IMFILE required with the new C version of MultiDrizzle.

### Parameters

**input** [string or list]

**Name of input file or files acceptable forms:**

- single filename with or without directory

- @-file
- association table
- python list of filenames
- wildcard specification of filenames

**refdir** [string] Path to directory containing new reference files, either environment variable or full path.

**local** [boolean] Specifies whether or not to copy new reference files to local directory for use with the input files.

**interactive** [boolean] Specifies whether or not to interactively ask the user for the exact names of the new reference files instead of automatically searching a directory for them.

**updatewcs** [boolean] Specifies whether or not to update the WCS information in this file to use the new reference files.

## Notes

**Warning:** This program requires access to the `jref$` directory in order to evaluate the DGE-OFI FILE specified in the input image header. This evaluation allows the program to get the information it needs to identify the correct NPOLFILE.

The use of this program now requires that a directory be set up with all the new NPOLFILE and D2IMFILE reference files for ACS (a single directory for all files for all ACS detectors will be fine, much like `jref`). Currently, all the files generated by the ACS team has initially been made available at:

```
/grp/hst/acs/lucas/new-npl/
```

The one known limitation to how this program works comes from confusion if more than 1 file could possibly be used as the new reference file. This would only happen when NPOLFILE reference files have been checked into CDBS multiple times, and there are several versions that apply to the same detector/filter combination. However, that can be sorted out later if we get into that situation at all.

## Examples

1. A set of associated images specified by an ASN file can be updated to use the NPOLFILES and D2IMFILE found in the local directory defined using the `myjref$` environment variable under PyRAF using:

```
>>> import updatenpol
>>> updatenpol.update('j8bt06010_asn.fits', 'myref$')
```

2. Another use under Python would be to feed it a specific list of files to be updated using:

```
>>> updatenpol.update(['file1_flt.fits', 'file2_flt.fits'], 'myjref$')
```

3. Files in another directory can also be processed using:

```
>>> updatenpol.update('data$*flt.fits', '../new/ref/')
```

---

## Reproducing Pipeline Processing

---

The task ‘runastrodriz’ can be used to reproduce the same Drizzle processing that gets performed on HST data when retrieving data from the HST archive.

### 17.1 Running Astrodrizzle

`runastrodriz` is a module to control operation of `astrodrizzle` which removes distortion and combines HST images in the pipeline.

#### 17.1.1 Typical Usage

```
>>> runastrodriz.py [-fhibn] inputFilename [newpath]
```

#### 17.1.2 Alternative Usage

```
>>> python
>>> from wfc3tools import runastrodriz
>>> runastrodriz.process(inputFilename, force=False, newpath=None,
↳ inmemory=False)
```

#### 17.1.3 GUI Usage under Python

```
>>> python
>>> from stsci.tools import teal
>>> import wfc3tools
>>> cfg = teal.teal('runastrodriz')
```

#### 17.1.4 PyRAF Usage

```
>>> epar runastrodriz
```

#### 17.1.5 Options

If the ‘-i’ option gets specified, no intermediate products will be written out to disk. These products, instead, will be kept in memory. This includes all single drizzle products (*single\_sci* and *single\_wht*), median image, blot images, and crmask images. The use of this option will therefore require significantly more memory than usual to process the data.

If a value has been provided for the `newpath` parameter, all processing will be performed in that directory/ramdisk. The steps involved are:

- create a temporary directory under that directory named after the input file
- copy all files related to the input to that new directory
- change to that new directory and run `astrodrizzle`
- change back to original directory
- move (not copy) ALL files from temp directory to original directory
- delete temp sub-directory

The ‘-b’ option will run this task in BASIC mode without creating headerlets for each input image.

The ‘-n’ option allows the user to specify the number of cores to be used in running `AstroDrizzle`.

#### 17.1.6 Pipeline Astrometric Calibration Description

A lot of effort goes into trying to determine the WCS solution which provides closest agreement with the GAIA astrometry frame. Combining the images successfully requires the most accurate alignment possible between all the input exposures taking into account the best available distortion model for the input exposures. Being able to compare the combined image to other exposures taken of the same field at other times, or even with other telescopes, requires that the WCS be defined to come as close to the GAIA frame as possible. The processing done by `runastrodriz` attempts to not only apply the most current distortion model, but also the best available pre-computed GAIA-based WCS solutions while proceeding to determine it’s own solution to the available GAIA reference stars for the field-of-view. It also performs numerous checks to see which of these solutions results in the most accurately aligned images to each other. The processing continued to attempt an alignment to GAIA using sources it identifies from the observations and checks to see if any fit was successful. Finally, it selects the WCS solution most closely aligned to GAIA



that as the basis for creating the final, distorted-corrected, combined drizzle products for the set of exposures being processed.

## Overview

The overall logic implemented by `runastrodriz` to generate the final set of drizzle products involves creating multiple sets of drizzle products and ultimately selecting one as the ‘best’. The basic steps are outlined here, with following sections providing additional detail on each step.

1. Initialization
  - a) Get any defined environment variables to control processing
  - b) Interpret input file
  - c) Make sure all input files are in a local directory
  - d) Check for DRIZCORR calibration switch in input files
  - e) Create name for output log files
  - f) Define lists of individual input files to be processed
1. Run `updatewcs` without astrometry database update on all input exposures (FLCs? and FLTs)
2. Generate initial default products and perform verification
  - a) perform cosmic-ray identification and generate drizzle products using `astrodrizzle` for all sets of inputs
  - b) verify relative alignment with focus index after masking out CRs
  - c) if alignment fails, update trailer file with failure information
3. If alignment is verified,
  - a) copy inputs to separate sub-directory for processing
  - b) run `updatewcs` to get a priori updates
    - apply ‘best’ a priori (not a posteriori) solution
  - c) generate drizzle products for all sets of inputs (FLC and/or FLT)
  - d) verify alignment using focus index on FLC or, if no FLC, FLT products
  - e) if alignment fails, update trailer file with info on failure
  - f) if product alignment verified:
    - copy all drizzle products to parent directory
    - copy updated input exposures to parent directory
4. If a posteriori correction enabled,
  - a) copy all inputs to separate sub-directory for processing

- b) run align to align the images
  - c) generate drizzle products for all sets of inputs (FLC and/or FLT) without CR identification
  - d) verify alignment using focus index on FLC or, if no FLC, FLT products
  - e) determine similarity index relative to pipeline default product
  - f) if either focus or similarity indicates a problem, update trailer file with info on failure
  - g) if product alignment verified:
    - copy all drizzle products to parent directory
    - copy updated input exposures to parent directory
5. Remove all processing sub-directories

## Initialization

### Environment Variables

The pipeline processing code starts out by looking to see whether the user has defined any processing behavior through the use of these environment variables:

- **‘ASTROMETRY\_COMPUTE\_APOSTERIORI’**: This environment variable specifies whether or not to attempt an *a posteriori* alignment where the code looks for sources in each of the images and uses those positions to perform relative alignment between the images and then fit those images to the GAIA frame.
- **‘ASTROMETRY\_APPLY\_APRIORI’**: This environment variable turns on/off application of any pre-defined(*a priori*) WCS solution found in the astrometry database.
- **‘ASTROMETRY\_STEP\_CONTROL’ [DEPRECATED, do not use]**: Old variable replaced by **‘ASTROMETRY\_APPLY\_APRIORI’**.

Values that can be provided for setting these variables are:

- ‘on’, ‘yes’, ‘true’: Any of these values will turn **on** the processing controlled by the variable
- ‘off’, ‘no’, ‘false’: Any of these values will turn **off** the processing controlled by the variable

By default, all the processing steps are turned **on** during pipeline processing in order to maximize the chances of aligning the data as closely as possible to the absolute astrometry standard coordinate system defined through the use of the GAIA catalogs. However, these controls are provided to support those observations which would not be suitable for such alignment, including observations of single sources.

## Input Data

The processing code needs to be told what data to process, and for `runastrodriz`, a single input filename is all that **can** be provided. This single input will be either:

- the name of an association table for a whole set of input exposures with a filename that looks like ‘<rootname>\_asn.fits’, where <rootname> is the designation for the association, such as ‘ie6d07030\_asn.fits’.
- the name of a single (uncalibrated) exposure with a filename that looks like ‘<rootname>\_raw.fits’.

This one input filename, though, will simply provide the code with the information it needs to find all the calibrated input exposures which need to have their distortion-models updated and applied. The whole set of input files required for processing includes:

- ASN (\*\_asn.fits) files: These small FITS tables provide the relationship between the input exposures and the output products with the output filenames defined in the table. There will NOT be an ASN table for exposures which were taken by themselves (called ‘singletons’).
- RAW (\*\_raw.fits) files: Not processed directly, but required in order to get the intended value of the DRIZCORR calibration switch. The ASN files also only give the rootname, and with the possibility of multiple suffixes (\_flt, \_flc, ...) for calibrated products, the code starts with the \_raw files to insure that what is specified in the ASN table is actually present and has been calibrated before processing.
- FLT/FLC (\*\_flt.fits or \*\_flc.fits) files: These are the non-CTE-corrected (\_flt) and CTE-corrected (\_flc) calibrated exposures to be processed.

The FLT/FLC files will be the ones that actually get processed and updated with the new distortion models and WCSs, while the others allow the code to know what FLT/FLC files should be included in the processing. This allows for multiple associations of data to live in the same directory and not interfere with each other as they are re-processed. That can be useful when interested in combining data from multiple visits, for example.

**Warning:** Should any of these files not be available (found in the local directory), the code will raise an Exception when trying to run ‘drizzlepac.astrodrizzle.AstroDrizzle’ on the data. The message will indicate what file was missing with something like: “**Exception: File ie6d07ujq\_flt not found.**”

## Calibration Switches

This processing serves as an official calibration step defined for HST data through the use of the **DRIZCORR** header keyword. This keyword can be found along with all the other calibration switches in the PRIMARY header (extension 0) of the exposures FITS file. A quick way to view this (or any keyword) value would be with:

```
from astropy.io import fits
val = fits.getval('ie6d07ujq_flt.fits', 'drizcorr')
```

This switch must be set to ‘PERFORM’ in order to allow the processing to be done. Processing will be completely skipped should the value of this switch in the ‘\_raw.fits’ file be set to ‘OMIT’.

## Log Files

A number of log files, or ‘trailer’ files, get generated during processing, and their filenames get defined as early as possible in the processing. The primary file will be a file with a ‘.tra’ extension and should have the same ‘<rootname>’ as the input file used to start the code. For example, if you were to reprocess ‘ie6d07030\_asn.fits’, you would end up with a trailer file with the name ‘ie6d07030.tra’.

This log file contains the messages generated from performing all the updates to the distortion model, updates from the astrometry database (if any), and all the image combinations performed by ‘AstroDrizzle()’ to create the final set of calibrated, drizzled exposures. Should any problems arise when during the processing, the log can provide the error messages and tracebacks to determine what went wrong.

## Data to be Processed

Once the code has performed all the initialization, it prepares the processing by defining what files need to be combined together from the input files it can find. This includes looking for CTE-corrected versions of the calibrated exposures (FLC files) as well as all the non-CTE-corrected files (FLT files) and creating a separate list of each type. Many types of data do not get CTE-corrected by the instruments calibration software, such as calacs.e or calwf3.e, and so no list of FLC files will be made. This will tell the code that it only needs to process the FLT files by themselves. If FLC files are found, all updates to the astrometry and WCS will be performed on those files and the results then get copied into the FLT file headers upon completion of the processing.

## Update the WCS

The first operation on the calibrated input files focuses on applying the calibrations for the distortion model to the WCS. This operation gets performed using the `updatewcs` task using the syntax:

```
from stwcs.updatewcs import updatewcs
updatewcs(calfiles_flg, use_db=False)
```

where `calfiles_flg` is the list of CTE-corrected FLC files or in the case there are no CTE-corrected files, the list of calibrated FLT files. Crucially, the use of `use_db=False` forces `updatewcs` to only apply the distortion model to the default WCS to create what is referred to as the **pipeline-default WCS**. This WCS has a WCSNAME associated with it that has the format `IDC_<rootname>` where `<rootname>` is the rootname of the IDCTAB reference files applied to the WCS.

This default WCS serves as the basis for all subsequent processing as the code tries to determine the WCS which is aligned most closely to the GAIA astrometric coordinate system.

## Generate the initial default products

The instrument teams have calibrated the distortion models extremely well for nearly all imaging modes with the latest calibration model being applied to the WCS keywords when the observations were updated in the previous step. The observations at this point represent what the best calibration of the pointings as observed

by the telescope. The accuracy of the guiding allows for sub-pixel alignment of the observations for most of the data and this step applies the distortion model to generate the ‘pipeline-default’ drizzle products.

The default products get generated using the `astrodrizzle` task. This initial run relies on a couple of default settings to generate the default drizzle products; namely,

- reads and applies default parameter settings from MDRIZTAB specified in observation header
- uses `resetbits=4096`
- runs with `crbit=4096` to define cosmic-rays/bad-pixels with DQ flag of 4096

## Identify Cosmic-Rays

Generating these drizzle products serves as the initial attempt to identify and to flag bad-pixels or cosmic-rays in each of the observations. Assuming the relative alignment of the initial pointing by the telescope is good (aligned to  $<0.1$  pixels), most of the cosmic-rays will be successfully identified at this point by flagging those pixels with a value of 4096 in the DQ array for each chip. This will make it easier to find sources and confirm alignment without having to weed through so many false sources. However, there are times when the default alignment by the telescope was not maintained which can result in all sources (real and cosmic-rays alike) to be flagged, so subsequent steps can reset the DQ bits from 4096 to 0 while processing the data again with `astrodrizzle` using different WCS solutions.

These initial products will only be generated for the CTE-corrected versions of the observations (`*_flc.fits` or FLC files) if they are present, and the standard calibrated versions of the observations (`*flt.fits` or FLT files) otherwise.

## Verifying Alignment

The relative alignment of these pipeline-default products relies entirely on the guiding accuracy of the telescope. Unfortunately, there are times when guiding problems impact the observations. These guiding errors can occur due to any of several reasons, including but not limited to:

- re-acquisition of a different guide star from one orbit to another, usually as a result of using a close binary that was not previously identified in the guide star catalog
- high slew rate due to only guiding on gyros due to problems with acquiring guide stars
- spurious guiding problems due to the aging telescope and guiding systems

## Computing the Focus Index

Verifying whether or not we can identify any problems with the relative alignment for these products starts by measuring the focus index for the drizzled products. The focus index was based on using the properties of the Laplacian of Gaussian (LoG) operator as an edge detector. See [http://alumni.media.mit.edu/~maov/classes/vision09/lect/09\\_Image\\_Filtering\\_Edge\\_Detection\\_09.pdf](http://alumni.media.mit.edu/~maov/classes/vision09/lect/09_Image_Filtering_Edge_Detection_09.pdf) for background on the Laplacian of Gaussian operator and its use in image filtering. The index that has been implemented is based on the maximum value of the LoG operation on each drizzled product.

The process for computing this index is:

- use the drizzled product, with as many cosmic-rays removed as possible, as the input

- mask out all the saturated sources as well as possible
- apply the LoG operator to the image
- pick out the pixel with the maximum value to serve as the value of the focus index

This measurement process gets applied to the total drizzle product for an association, as well as the drizzle product for each input exposure as well, known as ‘single drizzled’ products. The single drizzled products represent the optimal focus since there is only a single exposure with only telescope focus changes affecting the image focus value. The range of values from the single drizzled products establishes the distribution of ‘good’ focus values that gets used to evaluate whether the total drizzle product passes focus verification. This range of values comes as a result of the changing focus of the telescope from one exposure to another and to a lesser extent the effect of noise in low-S/N observations.

A Z-score then gets computed for the focus index value of each single drizzle product. In simplest terms, the Z-score is a measure of how many sigma above or below the population mean a measured value is. The actual computation is:

```
from scipy.stats as st

p = st.norm.cdf(x=val, loc=mean, scale=sigma)
z_score = st.norm.ppf(p)
```

A Z-score then gets computed for the focus index value derived from the total drizzle product. If this score falls within the range of values defined by the single drizzle focus index Z-score values, this WCS solution is considered to have passed the ‘focus verification’ check.

### Computing the Similarity Index

In addition to the focus index, a similarity index can also be computed between the single drizzle products (again treated as ‘truth’) and the total drizzle product. The function used to compute this is the `max_overlap_diff` function in `astrometric_utils`. The similarity index gets computed only for the region of maximum overlap of all the input exposures. This region of overlap gets determined using the `SkyFootprint` class from the `cell_utils` module. Should an input exposure not overlap the regions where most of the exposures overlap, then the region which overlaps at least 1 other exposure will be used for computing the index.

Point sources are detected in the selected region of overlap with a mask being generated for each source containing a value of 1 for the point source and 0 for the background. The sources are identified in the single drizzle image overlap region and the total drizzle product overlap region. These single drizzle mask then gets subtracted from the total drizzle mask, then scaled by the number of non-zero pixels in the single drizzle mask resulting in a Hamming distance between the two images. The Hamming distance, simply put, provides the percentage of differences pixel-by-pixel between two arrays as described in the `scipy` package `spatial.distance`. This distance then gets scaled by the relative exposure time of the single drizzle image to account for uncertainties introduced by readout noise, low S/N detection of sources and other variances due to exposure time.

We then compute a variant of the Mean Squared Error (MSE) algorithm used in the `AmphiIndex` image comparison code used for comparing images taken of amphibians. One description of how the MSE measures the similarity between images can be found at <https://www.pyimagesearch.com/2014/09/15/python-compare-two-images/>. This similarity index is sensitive to small offsets between exposures, as well as differences in noise, overall S/N, and even presence of cosmic-rays. In contrast, the Hamming-

distance is not as sensitive to noise. Therefore, we compare the MSE similarity with the Hamming distance and take the minimum of the two values as a more robust measure of the similarity of the images. Both values share one key characteristic: values  $> 1.0$  indicate more pixels are different than similar. The code takes the maximum value of the similarity indices computed for the total drizzle product compared to all the single drizzle products as the final measure of the similarity. If this value is less than 1.0, then this WCS is considered to have passed the similarity check.

## Updating the Trailer File

Associations where there are problems with the alignment will cause this verification to fail since the sources will not be ‘as sharp’ based on the LoG operator. As a result, it can flag situations where even sub-pixel offsets down less than 0.5 pixels are identified. For the default pipeline alignment, failure at this point is only noted in the log with the hope that later alignment efforts will resolve the problem affecting the original input data as noted in this check.

## Applying A Priori WCS Solutions

A priori WCS solutions defined for use with HST data refer to improvements to the WCS solutions that were pre-computed. As of 2020, there were 2 primary sources of a priori WCS solutions:

- GSC240: correcting the previous guide star coordinates to the GAIA frame
- HSC30: corrections derived using the Hubble Source Catalog(HSC) coordinates cross-matched to the GAIA catalog

The updated a priori solutions are stored as `headerlets` in the database. The headerlet format allows them to be applied directly to the exposure using the STWCS package while requiring very little storage space (typically,  $< 120\text{Kb}$  per headerlet). More details on the `headerlet` can be found at <https://stwcs.readthedocs.io/en/latest/headerlet.html>.

## GSC240: GAIA and the HST Guide Stars

Observations taken prior to October 2017 used guide star coordinates which were based on guide star coordinates derived primarily from ground-based observations. This resulted in an uncertainty of 1 arcsecond in the absolute pointing of the telescope for any given observation. The development and availability of the space-based GAIA astrometric catalog finally allowed for the guide star coordinates to be known to better than 10 milli-arcseconds in 2015 with proper motion uncertainties increasing by 5 milli-arcseconds per year on average. The GAIA astrometric catalog was then cross-matched to the HST guide star catalog used for pointing the telescope, and corrections were determined. These corrections were then applied to every HST observation taken before Oct 2017 as if the telescope used the GAIA coordinates originally to generate updated WCS solutions to describe the GAIA-based pointing. These updated WCS solutions were labelled with ‘GSC240’ in the WCSNAME and stored in an astrometry database to be applied on-demand to all observations taken before Oct 2017.

These solutions will not result in perfect alignment to the GAIA catalog, due to temporal uncertainties in the calibration of the instrument’s field of view relative to the FGS’s used to point and to guid the telescope

during the observations. This uncertainty can be up to 0.5 arcseconds, but it still represents a significant improvement in the absolute astrometry from the 1-sigma of 1 arcsecond for previous WCS solutions.

All observations taken after Oct 2017 already used guide-star coordinates based on GAIA, so no new WCS was needed as it would simply be the same as the pipeline default WCS.

## HSC30: Hubble Source Catalog WCSs

The Hubble Source Catalog(HSC) (<https://archive.stsci.edu/hst/hsc/>) developed a comprehensive catalog of a majority of the sources observed in Hubble data. This catalog was then cross-matched to the GAIA catalog to determine improved positions for those sources. By using the updated positions from Version 3.0 of the HSC and comparing them to the original positions based on the pipeline default WCS solutions, updates were derived for all observations with sources from the HSC. The updates were then used to recompute the WCS solutions for those observations which were labelled as 'HSC30' in the WCSNAME and stored in the astrometry database.

## Separate Directories

One mechanism used to enable comparisons of various WCS solutions is to keep copies of the observations with different types of WCS solutions in separate directories. Up until this point in the processing, the data has been processed in the directory where the processing was started. In order to keep the `a priori` solutions separate, a sub-directory gets created with name based on the association table rootname or the rootname of the single exposure being processed using the convention: `<rootname>_apriori`. All the FLC (or FLT, if no FLC files are present), and ASN file (if processing an association) are copied from the main directory into the new sub-directory and the process moves to the sub-directory to continue its processing.

## Applying the A Priori Solutions

Application of these `a priori` WCS solutions simply involves running the `updatewcs` task with `use_db=True` (the default setting). This queries the astrometry database and retrieves the headerlets for all the `a priori` solutions. The database also reports what solution is flagged as the `best` solution, which will typically result in the closest alignment to GAIA. All the headerlets get appended as new extensions to the observations FITS file, then the `a priori` solution flagged as `best` gets applied to replace the active or primary WCS in the observation after saving a copy of the original primary WCS. Other solutions could be provided by the database that were derived directly from the observation itself, perhaps in previous pipeline processing runs. These solutions are retained, but not applied at this point since it is not clear whether the distortion model has changed from those saved in the database, or whether the pipeline software has been improved to provide a more accurate or more robust solution. Finally, we are only interested in seeing whether there are any issues in applying the pre-defined `a priori` corrections.

## Generating A Priori Products

The FLC images updated with the `a priori` WCS solutions now get combined using `astrodrizzle`. If the pipeline default focus verification succeeded, then `resetbits` will be set to 0 so that the previous



DQ flags can be used. If the verification failed, though, `resetbits` gets set to 4096 so that the cosmic-rays can be identified and flagged fresh based on the alignment provided by the `a priori` WCS solutions.

This processing will result in a total combined drizzle product based on the `a priori` solution.

## Evaluating Alignment

Confirming that the relative alignment between the images in the association was maintained with the `a priori` WCS now can be done. Although the `a priori` WCS solutions are vetted for accuracy, HST has taken a few hundred thousand different exposures in dozens of configurations and not all of those exposures were taken exactly as planned. Therefore, considerable effort goes into trying to verify that the alignment between the images has been maintained.

This verification starts by computing the focus index and similarity values for the total drizzle product and the single drizzle products using the same code used to verify the pipeline default WCS drizzle product. It then extends to include computing the similarity index between the `a priori` drizzle products and the pipeline default drizzle products. This will attempt to measure whether or not the `a priori` alignment is significantly different than the presumably good pipeline default alignment. Once again, if the similarity index is less than 1, the `a priori` alignment is considered to be successful.

## Keeping the A Priori Alignment

Should all the verification steps indicate a successful alignment, the `a priori` WCS solution should be retained as an improved WCS solution over the pipeline default WCS. This gets done by simply copying the calibrated images which have been updated with the WCS solution (both the FLC and FLT images) from the `<rootname>_apriori` sub-directory to the main processing directory. This will replace the FLC and FLT files with the pipeline default solutions so that should no other WCS prove to be better, the `a priori` WCS solution will end up being used to generate the final drizzle products which get archived and provided to the end-user.

## Performing An A Posteriori Alignment

The ultimate goal of this processing would be to have the input observations aligned as closely to an astrometric standard coordinate system as much as possible. The highest quality, highest precision astrometric catalog available would be the GAIA astrometric catalog and this processing seeks to align HST observations as closely to that catalog's coordinate system.

The `a priori` solutions provide an update to the astrometry based on either the guide stars used (the GSC240 and related solutions) or manually verified alignment of sources from the observations field-of-view performed using the Hubble Source Catalog (the HSC30 solution). Unfortunately, both of these types of solutions fail to account for sources of astrometric error which can still affect the observations and result in offsets from the GAIA system due to updates in the distortion calibration for the instruments or uncertainties in the position of the detectors field-of-view relative to the Fine Guidance Sensors (FGS) and the guide stars used for taking the observations.

The only way to correct for those effects remains to identify sources from the observations and perform a fit to the GAIA catalog directly. This is called an `a posteriori` solution when it can be done successfully.

However, this can only be performed for observations which contain enough detectable sources, specifically sources found in the GAIA catalog. Not all observations meet this criteria either due to exposure time (too long or too short), wavelength of observation, filter bandpass (narrowband vs wide-band) and even number of sources in the field. This processing code makes no assumptions about the possibility of success and tries to perform this `a posteriori` fit on all observations.

## Copying the Observations

Copies of the observations are made in a sub-directory named after the input file used to start the processing with the convention:

```
<rootname>_aposteriori
```

For example, if the association `icw402010_asn.fits` was being processed, this directory would be named `icw402010_aposteriori`.

All the calibrated FLC and/or FLT images along with the ASN file are copied into this sub-directory. These files, at this point, have the best available WCS at this time which is most likely an `a priori` solution. This improves the chance that the `a posteriori` fit will work by minimizing the offset from GAIA which needs to be searched to find a cross-match with the GAIA sources in the field-of-view.

## Aligning the Observations

The alignment process gets performed using the `perform_align()` function from the `drizzlepac/align` module. This function performs the following steps in an attempt to perform an `a posteriori` fit to GAIA:

- Evaluates all the input observations to identify any which can not be aligned, such as GRISM or SCAN mode observations. For a full description of all the type of observations that can be filtered out, see *haputils.analyze*.
- Compute a 2D background for all the observations using `photutils`
- Determine a PSF kernel from the detectable sources in the image, if possible.
- Segments the image after applying the 2D background to identify as many sources as possible above a threshold using `photutils.segmentation`
- Performs source centering using `photutils.DAOStarFinder`
- Keeps the position of the single brightest source nearest the center of the segment as the catalog position for each segment's object.
- Checks whether there are enough sources to potentially get a viable linear fit.
  - If not, the attempt at an `a posteriori` fit quits without updating the WCS of the input files.
- Queries the GAIA DR2 catalog through the STScI web service to obtain a catalog of GAIA sources that overlap the field-of-view of the combined set of observations. This catalog will serve as the **reference catalog** for the fitting process.

- If there are not enough GAIA sources overlapping these observations, then the fit attempt quits without updating the WCS of the input files.
- Provide the source catalogs for each input image, each input images’s WCS, and the GAIA reference catalog to function `align_wcs()` in the `tweakwcs` package.
  - This function cross-matches the source catalog from each image with the GAIA catalog and performs an **r**scale linear fit (as defined by `runastrodriz`), then updates the input WCS with the results of the fit upon success. See the [tweakwcs readthedocs pages](#) for more details.
  - The function `align_wcs` is first called without using the GAIA reference catalog in order to perform a relative alignment between the observations.
  - The function `align_wcs` is then called with the GAIA catalog as the reference in order to finally perform a single fit to the GAIA catalog for all the observations at the same time.
- Evaluate the success/failure state of the fit and the quality of any successful fit.
- Repeat the fit with `tweakwcs.align_wcs` with other GAIA catalogs; including GAIA DR1 or any others specified for use in `runastrodriz` itself.
- Select the fit to the GAIA catalog which results in the lowest RMS.
  - Some fields are dominated by external galaxies with no proper motion for which GAIA DR1 without proper motions provides the best fit (lowest RMS).
  - Other fields are dominated by local galactic stars with appreciable proper motions best accounted for (still with some error) by the GAIA DR2 catalog with its proper motions.
- Keep the WCS’s updated with the **best** solution and update the **WCSNAME** keyword for those WCSs to reflect the type of fit that was successful and the catalog that was used.
  - The naming convention is more fully described on the [Drizzlepac Astrometry description](#).

The result of this lengthy process is a set of WCS objects which have been updated with a fit to a GAIA catalog representing an `a posteriori` solution.

## Generate the Aligned Drizzle Products

Successful alignment of the WCSs to a GAIA catalog means that these `a posteriori` updated exposures can be combined to create a drizzled product using `AstroDrizzle`.

## Verify A Posteriori Alignment

These newly updated drizzle products still need to be evaluated to insure that the fit performed to GAIA maintained relative alignment between the images as well. Mis-alignment of the images to each other can result from too few sources being used for the fit imprinting the errors in those source positions on the relative alignment. The verification used is the same focus and similarity checks that were performed on the `a priori` updated drizzle products and even the pipeline default drizzle products.

## A Posteriori Failure

At any number of points throughout this computation and verification, it could end up quitting and flagging this attempt as a failure. If this happens, no updated WCS solutions get created or saved and processing returns to the parent directory while deleting the entire `<dataset>_aposteriori` directory along with all the mis-aligned or un-alignable files. This allows the processing to revert to using the previously verified WCS solutions as the `best` WCS solution available for these observations.

## A Posteriori Success

Successfully fitting to GAIA can only be declared after the verification process returned values indicating good alignment in the drizzle product. The processing would then copy these `a posteriori`-updated input exposures from the sub-directory these computations were being performed in based up to the parent directory to replace the previously updated versions of the input files. This entire sub-directory then gets deleted, unless the processing was being run in debug mode.

## Creation of Final Aligned Products

The starting directory now contains updated input FLC/FLT files based on WCSs which have been verified to have maintained relative alignment and with alignment as close to the GAIA astrometric coordinate system as possible. These exposures get processed by `AstroDrizzle` to create the final, combined drizzle products for the user and for archiving at STScI in the Mikulski Archive for Space Telescopes (MAST). These products include the calibrated drizzle(DRZ) products as well as any CTE-corrected drizzle(DRC) products depending on what input exposures are available.

## 17.2 Hubble Advanced Products API

These modules provide the basic functionality used to process automatically data using this package to apply the distortion models to the WCS of HST observations and to verify the alignment of the observations.

### 17.2.1 haputils.astrometric\_utils

Utilities to support creation of astrometrically accurate reference catalogs

The function, `create_astrometric_catalog`, allows the user to query an astrometric catalog online to generate a catalog of astrometric sources that should fall within the field-of-view of all the input images.

This module relies on the definition of an environment variable to specify the URL of the astrometric catalog to use for generating this reference catalog.

```
ASTROMETRIC_CATALOG_URL -- URL of web service that can be queried to
                        obtain listing of astrometric sources,
                        sky coordinates, and magnitudes.
```

```
drizzlepac.haputils.astrometric_utils.create_astrometric_catalog(inputs,
                                                                cat-
                                                                a-
                                                                log='GAIADR2',
                                                                out-
                                                                put='ref_cat.ecsv',
                                                                gaia_only=False,
                                                                ta-
                                                                ble_format='ascii.ecsv',
                                                                ex-
                                                                ist-
                                                                ing_wcs=None,
                                                                num_sources=None,
                                                                use_footprint=False,
                                                                full_catalog=False)
```

Create an astrometric catalog that covers the inputs' field-of-view.

### Parameters

- input** [str, list] Filenames of images to be aligned to astrometric catalog
- catalog** [str, optional] Name of catalog to extract astrometric positions for sources in the input images' field-of-view. Default: GAIADR2. Options available are documented on the catalog web page.
- output** [str, optional] Filename to give to the astrometric catalog read in from the master catalog web service. If None, no file will be written out.
- gaia\_only** [bool, optional] Specify whether or not to only use sources from GAIA in output catalog
- existing\_wcs** [HSTWCS] existing WCS object specified by the user
- num\_sources** [int] Maximum number of brightest/faintest sources to return in catalog. If `num_sources` is negative, return that number of the faintest sources. By default, all sources are returned.
- use\_footprint** [bool, optional] Use the image footprint as the source identification bounds insted of using a circle centered on a given RA and Dec? By default, `use_footprint = False`.
- full\_catalog** [bool, optional] Return the full set of columns provided by the web service.

### Returns

- ref\_table** [Table] Astropy Table object of the catalog

### Notes

This function will point to astrometric catalog web service defined through the use of the `ASTROMETRIC_CATALOG_URL` environment variable.

```
drizzlepac.haputils.astrometric_utils.get_catalog(ra, dec, sr=0.1,  
                                                  epoch=None, catalog='GSC241')
```

Extract catalog from VO web service.

#### Parameters

- ra** [float] Right Ascension (RA) of center of field-of-view (in decimal degrees)
- dec** [float] Declination (Dec) of center of field-of-view (in decimal degrees)
- sr** [float, optional] Search radius (in decimal degrees) from field-of-view center to use for sources from catalog. Default: 0.1 degrees
- epoch** [float, optional] Catalog positions returned for this field-of-view will have their proper motions applied to represent their positions at this date, if a value is specified at all, for catalogs with proper motions.
- catalog** [str, optional] Name of catalog to query, as defined by web-service. Default: 'GSC241'

#### Returns

- csv** [CSV object] CSV object of returned sources with all columns as provided by catalog

```
drizzlepac.haputils.astrometric_utils.find_gsc_offset(image, input_catalog='GSC1',  
                                                    output_catalog='GAIA')
```

Find the GSC to GAIA offset based on guide star coordinates

#### Parameters

- image** [str] Filename of image to be processed.

#### Returns

- delta\_ra, delta\_dec** [tuple of floats] Offset in decimal degrees of image based on correction to guide star coordinates relative to GAIA.

```
drizzlepac.haputils.astrometric_utils.extract_sources (img,          dq-
                                                    mask=None,
                                                    fwhm=3.0,    ker-
                                                    nel=None,    phot-
                                                    mode=None,   seg-
                                                    ment_threshold=None,
                                                    dao_threshold=None,
                                                    source_box=7,
                                                    classify=True,
                                                    center-
                                                    ing_mode='starfind',
                                                    nlargest=None,
                                                    outroot=None,
                                                    plot=False,
                                                    vmax=None,
                                                    deblend=False)
```

Use photutils to find sources in image based on segmentation.

### Parameters

**img** [ndarray] Numpy array of the science extension from the observations FITS file.

**dqmask** [ndarray] Bitmask which identifies whether a pixel should be used (1) in source identification or not(0). If provided, this mask will be applied to the input array prior to source identification.

**fwhm** [float] Full-width half-maximum (fwhm) of the PSF in pixels.

**threshold** [float or None] Value from the image which serves as the limit for determining sources. If None, compute a default value of  $(\text{background} + 5 * \text{rms}(\text{background}))$ . If  $\text{threshold} < 0.0$ , use absolute value as scaling factor for default value.

**source\_box** [int] Size of box (in pixels) which defines the minimum size of a valid source.

**classify** [bool] Specify whether or not to apply classification based on invariant moments of each source to determine whether or not a source is likely to be a cosmic-ray, and not include those sources in the final catalog.

**centering\_mode** [str] “segmentation” or “starfind” Algorithm to use when computing the positions of the detected sources. Centering will only take place after `threshold` has been determined, and sources are identified using segmentation. Centering using segmentation will rely on `photutils.segmentation.source_properties` to generate the properties for the source catalog. Centering using `starfind` will use `photutils.IRAFStarFinder` to characterize each source in the catalog.

**nlargest** [int, None] Number of largest (brightest) sources in each chip/array to measure when using ‘starfind’ mode.

**outroot** [str, optional] If specified, write out the catalog of sources to the file with this name rootname.

**plot** [bool, optional] Specify whether or not to create a plot of the sources on a view of the image.

**vmax** [float, optional] If plotting the sources, scale the image to this maximum value.

**deblend** [bool, optional] Specify whether or not to apply photutils deblending algorithm when evaluating each of the identified segments (sources) from the chip.

`drizzlepac.haputils.astrometric_utils.classify_sources` (*catalog*,  
*sources=None*)

Convert `moments_central` attribute for source catalog into star/cr flag.

This algorithm interprets the `central_moments` from the `source_properties` generated for the sources as more-likely a star or a cosmic-ray. It is not intended or expected to be precise, merely a means of making a first cut at removing likely cosmic-rays or other artifacts.

### Parameters

**catalog** [`SourceCatalog`] The photutils catalog for the image/chip.

**sources** [tuple] Range of objects from catalog to process as a tuple of (min, max). If None (default) all sources are processed.

### Returns

**srctype** [ndarray] An ndarray where a value of 1 indicates a likely valid, non-cosmic-ray source, and a value of 0 indicates a likely cosmic-ray.

`drizzlepac.haputils.astrometric_utils.generate_source_catalog` (*image*,  
*dq-*  
*name='DQ'*,  
*out-*  
*put=False*,  
*fwhm=3.0*,  
*\*\*de-*  
*tec-*  
*tor\_pars*)

Build source catalogs for each chip using photutils.

The catalog returned by this function includes sources found in all chips of the input image with the positions translated to the coordinate frame defined by the reference WCS `refwcs`. The sources will be - identified using photutils segmentation-based source finding code - ignore any input pixel which has been flagged as 'bad' in the DQ array, should a DQ array be found in the input HDUList. - classified as probable cosmic-rays (if enabled) using `central_moments` properties of each source, with these sources being removed from the catalog.

### Parameters

**image** [`HDUList`] Input image as an `astropy.io.fits` HDUList.

**dqname** [str] EXTNAME for the DQ array, if present, in the input image HDUList.

**output** [bool] Specify whether or not to write out a separate catalog file for all the sources found in each chip.

**fwhm** [float] Full-width half-maximum (fwhm) of the PSF in pixels.



**Returns**

**source\_cats** [dict] Dict of astropy Tables identified by chip number with each table containing sources from image extension ('sci', chip).

```
drizzlepac.haputils.astrometric_utils.generate_sky_catalog(image,
                                                            refwcs, dq-
                                                            name='DQ',
                                                            out-
                                                            put=False)
```

Build source catalog from input image using photutils.

This script borrows heavily from build\_source\_catalog.

The catalog returned by this function includes sources found in all chips of the input image with the positions translated to the coordinate frame defined by the reference WCS `refwcs`. The sources will be - identified using photutils segmentation-based source finding code - ignore any input pixel which has been flagged as 'bad' in the DQ array, should a DQ array be found in the input HDUList. - classified as probable cosmic-rays (if enabled) using `central_moments` properties of each source, with these sources being removed from the catalog.

**Parameters**

**image** [HDUList] Input image.

**refwcs** [HSTWCS] Definition of the reference frame WCS.

**dqname** [str, optional] EXTNAME for the DQ array, if present, in the input image.

**output** [bool, optional] Specify whether or not to write out a separate catalog file for all the sources found in each chip.

**Returns**

**master\_cat** [Table] Source catalog for all 'valid' sources identified from all chips of the input image with positions translated to the reference WCS coordinate frame.

```
drizzlepac.haputils.astrometric_utils.compute_photometry(catalog, phot-
                                                            vals)
```

Compute magnitudes for sources from catalog based on observations photmode.

Magnitudes will be AB mag values.

**Parameters**

**catalog** [Table] Astropy Table with 'source\_sum' column for the measured flux for each source.

**photmode** [str] Specification of the observation filter configuration used for the exposure as reported by the 'PHOTMODE' keyword from the PRIMARY header.

**Returns**

**phot\_cat** [Table] Astropy Table object of input source catalog with added column for ABMAG photometry (in magnitudes).

```
drizzlepac.haputils.astrometric_utils.filter_catalog(catalog,
                                                    bright_limit=1.0,
                                                    max_bright=None,
                                                    min_bright=20, col-
                                                    name='vegamag')
```

Create a new catalog selected from input based on photometry.

### Parameters

**catalog** [Table] Table containing the full set of identified sources.

**bright\_limit** [float] Fraction of catalog based on brightness that should be retained.  
Value of 1.00 means full catalog.

**max\_bright** [int] Maximum number of sources to keep regardless of  
`bright_limit`.

**min\_bright** [int] Minimum number of sources to keep regardless of  
`bright_limit`.

**colname** [str] Name of column to use for selection/sorting.

### Returns

**new\_catalog** [Table] New table which only has the sources that meet the selection  
criteria.

```
drizzlepac.haputils.astrometric_utils.build_self_reference(filename,
                                                            clean_wcs=False)
```

This function creates a reference, undistorted WCS that can be used to apply a correction to the WCS of the input file.

### Parameters

**filename** [str] Filename of image which will be corrected, and which will form the  
basis of the undistorted WCS.

**clean\_wcs** [bool] Specify whether or not to return the WCS object without any dis-  
tortion information, or any history of the original input image. This converts the  
output from `utils.output_wcs()` into a pristine HSTWCS object.

### Returns

**customwcs** [HSTWCS] HSTWCS object which contains the undistorted WCS repre-  
senting the entire field-of-view for the input image.

## Examples

This function can be used with the following syntax to apply a shift/rot/scale change to the same image:

```
>>> import buildref
>>> from drizzlepac import updatehdr
>>> filename = "jce501erq_flg.fits"
```

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```
>>> wcslin = buildref.build_self_reference(filename)
>>> updatehdr.updatewcs_with_shift(filename, wcslin, xsh=49.5694,
... ysh=19.2203, rot = 359.998, scale = 0.9999964)
```

`drizzlepac.haputils.astrometric_utils.within_footprint` (*img*, *wcsobj*, *x*, *y*)  
Determine whether input *x*, *y* fall in the science area of the image.

**Parameters**

**img** [ndarray] ndarray of image where non-science areas are marked with value of NaN.

**wcsobj** [HSTWCS] HSTWCS or WCS object with naxis terms defined.

**x, y** [ndarray] arrays of *x*, *y* positions for sources to be checked.

**Returns**

**mask** [ndarray] Boolean array of same length as *x*, *y* arrays where sources that fall within the footprint are True.

`drizzlepac.haputils.astrometric_utils.find_hist2d_offset` (*filename*,  
*reference*,  
*refwcs=None*,  
*ref-*  
*names=['ra'*,  
*'dec']*,  
*match\_tolerance=5.0*,  
*chip\_catalog=True*,  
*search\_radius=15.0*,  
*min\_match=10*,  
*classify=True*)

Iteratively look for the best cross-match between the catalog and ref.

**Parameters**

**filename** [HDUList or str] Single image to extract sources for matching to the external astrometric catalog.

**reference** [str or Table] Reference catalog, either as a filename or `astropy.Table` containing astrometrically accurate sky coordinates for astrometric standard sources.

**refwcs** [HSTWCS] This WCS will define the coordinate frame which will be used to determine the offset. If None is specified, use the WCS from the input image *filename* to build this WCS using `build_self_reference()`.

**refnames** [list] List of table column names for sky coordinates of astrometric standard sources from reference catalog.

**match\_tolerance** [float] Tolerance (in pixels) for recognizing that a source position matches an astrometric catalog position. Larger values allow for lower accuracy source positions to be compared to astrometric catalog

**chip\_catalog** [bool] Specify whether or not to write out individual source catalog for each chip in the image.

**search\_radius** [float] Maximum separation (in arcseconds) from source positions to look for valid cross-matches with reference source positions.

**min\_match** [int] Minimum number of cross-matches for an acceptable determination of the offset.

**classify** [bool] Specify whether or not to use `central_moments` classification to ignore likely cosmic-rays/bad-pixels when generating the source catalog.

### Returns

**best\_offset** [tuple]

Offset in input image pixels between image source positions and astrometric catalog positions that results in largest number of matches of astrometric sources with image sources

**seg\_xy, ref\_xy** [Table] Source catalog and reference catalog, respectively, used for determining the offset. Each catalog includes sources for the entire field-of-view, not just a single chip.

```
drizzlepac.haputils.astrometric_utils.build_wscat(image, group_id,
                                                    source_catalog)
```

Return a list of FITSWCS objects for all chips in an image.

### Parameters

**image** [str, HDUList] Either filename or HDUList of a single HST observation.

**group\_id** [int] Integer ID for group this image should be associated with; primarily used when separate chips are in separate files to treat them all as one exposure.

**source\_catalog** [dict] If provided, these catalogs will be attached as `catalog` entries in each chip's FITSWCS object. It should be provided as a dict of astropy Tables identified by chip number with each table containing sources from image extension ('sci', chip) as generated by `generate_source_catalog()`.

### Returns

**wcs\_catalogs** [list of FITSWCS] List of FITSWCS objects defined for all chips in input image.

```
drizzlepac.haputils.astrometric_utils.compute_similarity(image, reference)
```

Compute a similarity index for an image compared to a reference image.

Similarity index is based on a the general algorithm used in the AmphiIndex algorithm.

- identify slice of image that is a factor of 256 in size
- rebin image slice down to a (256,256) image
- rebin same slice from reference down to a (256,256) image

- sum the differences of the rebinned slices
- divide absolute value of difference scaled by reference slice sum

This index will typically return values  $< 0.1$  for similar images, and values  $> 1$  for dis-similar images.

### Parameters

**image** [ndarray] Image (as ndarray) to measure

**reference** [ndarray] Image which serves as the ‘truth’ or comparison image.

### Returns

**similarity\_index** [float] Value of similarity index for image

```
drizzlepac.haputils.astrometric_utils.determine_focus_index(img,
                                                         sigma=1.5)
```

Determine blurriness indicator for an image

This returns a single value that serves as an indication of the sharpness of the image based on the max pixel value from the image after applying a Laplacian-of-Gaussian filter with sigma.

This index needs to be based on ‘max’ value in order to avoid field-dependent biases, since the ‘max’ value will correspond to point-source-like sources regardless of the field (nebula, galaxy, ...). Care must be taken, though, to ignore cosmic-rays as much as possible as they will mimic real sources without providing information on the actual focus through the optics. Similarly, saturation regions must also be ignored as they also only indicate a detector feature, not the focus through the optics or alignment of actual sources.

```
drizzlepac.haputils.astrometric_utils.max_overlap_diff(total_mask, singlefiles, prodfile,
                                                         sigma=2.0,
                                                         scale=1,
                                                         lsigma=3.0)
```

Determines the difference in the region of max overlap for all drizzled products

### Parameters

**total\_mask** [ndarray] Mask (array) showing where each input exposure contributes to the final drizzle product `prodfile`. This could be created using `cell_utils.SkyFootprint`.

**singlefiles** [list] List of filenames for each single input exposure drizzled onto the same WCS as the final drizzle product `prodfile`

**prodfile** [str] Filename for the final drizzle product

**scale** [int, optional] Factor to use in downsizing (resizing smaller) the images to be evaluated. The larger the value, the less sensitive this measurement becomes.

**sigma** [float, optional] Size of default kernel (in pixels) to use for determining the focus.

### Returns

**diff\_dict** [dictionary] Dictionary of difference scores for each input exposure drizzle product (from `singlefiles`) calculated for the region of maximum overlap with the final drizzle product `prodfile`. Entries for each singlefile includes:

- `distance` : Hamming distance of singlefile from `prodfile`
- `focus` : focus index of singlefile
- `focus_pos` : position for best focus in singlefile
- `product_focus` : focus index for `prodfile`
- `product_focus_pos` : position for best focus in `prodfile`

```
drizzlepac.haputils.astrometric_utils.detect_point_sources (arr, back-
                                                             ground=None,
                                                             nsigma=4,
                                                             log_sigma=3.0,
                                                             scale=1,
                                                             sigma=3.0,
                                                             exp_weight=None)
```

## 17.2.2 haputils.analyze

Utility to analyze an input dataset and determine whether the dataset can be aligned

The function `analyze_data` opens an input list containing FLT and/or FLC FITS filenames in order to access the primary header data. Based upon the values of specific FITS keywords, the function determines whether or not each file within this dataset can or should be reconciled against an astrometric catalog and, for multiple images, used to create a mosaic.

```
drizzlepac.haputils.analyze.analyze_data (input_file_list, log_level=0)
```

Determine if images within the dataset can be aligned

### Parameters

**input\_file\_list** [list] List containing FLT and/or FLC filenames for all input images which comprise an associated dataset where ‘associated dataset’ may be a single image, multiple images, an HST association, or a number of HST associations

**log\_level** [int, optional] The desired level of verbosity in the log statements displayed on the screen and written to the `.log` file. Default value is 20, or ‘info’.

### Returns

**output\_table** [object] Astropy Table object containing data pertaining to the associated dataset, including the `do_process` bool. It is intended this table is updated by subsequent functions for bookkeeping purposes.

### Notes

The keyword/value pairs below define the “cannot process categories”. `OBSTYPE` : is not IMAGING  
`MTFLAG` : T `SCAN-TYP` : C or D (or !N) `FILTER` : G\*, PR\*, where G=Grism and PR=Prism `FIL-`

TER1 : G\*, PR\*, where G=Grism and PR=Prism  
 FILTER2 : G\*, PR\*, where G=Grism and PR=Prism  
 TARGNAME : DARK, TUNGSTEN, BIAS, FLAT, EARTH-CALIB, DEUTERIUM  
 EXPTIME : 0  
 CHINJECT : is not NONE

The keyword/value pairs below define the category which the data can be processed, but the results may be compromised  
 FGSLOCK : FINE/GYRO, FINE/GY, COARSE, GYROS

FITS Keywords only for WFC3 data: SCAN\_TYP, FILTER, and CHINJECT (UVIS)  
 FITS Keywords only for ACS data: FILTER1 and FILTER2

Please be aware of the FITS keyword value NONE vs the Python None.

### 17.2.3 haputils.align\_utils

```
class drizzlepac.haputils.align_utils.AlignmentTable (input_list, clobber=False, dq_name='DQ', log_level=0, **alignment_pars)
```

**\*\*alignment\_pars** needs to contain the following entries: # kernel defining, source finding par fwhmpsf=0.12, # background computing pars box\_size=BKG\_BOX\_SIZE, win\_size=BKG\_FILTER\_SIZE, bkg\_estimator=SExtractorBackground, rms\_estimator=StdBackgroundRMS, nsigma=5., threshold\_flag=None, # object finding pars source\_box=7, classify=True, centering\_mode="starfind", nlargest=None, plot=False, vmax=None, deblend=False

```
drizzlepac.haputils.align_utils.match_relative_fit (imglist, reference_catalog, **fit_pars)
```

Perform cross-matching and final fit using relative matching algorithm

#### Parameters

**imglist** [list] List of input image FITSWCS objects with metadata and source catalogs  
**reference\_catalog** [Table] Astropy Table of reference sources for this field

#### Returns

**imglist** [list] List of input image FITSWCS objects with metadata and source catalogs

```
drizzlepac.haputils.align_utils.match_default_fit (imglist, reference_catalog, **fit_pars)
```

Perform cross-matching and final fit using default tolerance matching

#### Parameters

**imglist** [list] List of input image FITSWCS objects with metadata and source catalogs  
**reference\_catalog** [Table] Astropy Table of reference sources for this field

#### Returns

**imglist** [list] List of input image FITSWCS objects with metadata and source catalogs

`drizzlepac.haputils.align_utils.match_2dhist_fit` (*imglist*, *reference\_catalog*,  
\*\**fit\_pars*)

Perform cross-matching and final fit using 2dHistogram matching

#### Parameters

**imglist** [list] List of input image FITSWCS objects with metadata and source catalogs

**reference\_catalog** [Table] Astropy Table of reference sources for this field

#### Returns

**imglist** [list] List of input image FITSWCS objects with metadata and source catalogs

## 17.3 Single-visit Mosaic Processing

Standard calibration pipeline processing focuses on aligning data taken as associations or as single exposures as closely to a standard astrometric coordinate frame as possible (see *Pipeline Astrometric Calibration Description* for full details). This involves alignment of very few images all taken under as identical conditions as possible; namely, detector, filter, guide stars, guiding mode, and so on. These observations were intended by the user to represent a single view of the object of interest in a way that allows for removal of as many calibration effects as possible.

Observations taken within a single visit, on the other hand, represent data intended to produce a multi-wavelength view of the objects in the desired field-of-view. Most of the time, those observations are taken using pairs of guide stars to provide very stable field-of-view throughout the entire visit resulting in images which overlap almost perfectly. Unfortunately, this is not always possible due to increasing limitations of the aging telescope systems. The result is that an increasing number of visits are taken where observations drift and/or roll during the course of the visit or re-acquire at slightly different pointings from one orbit to the next. Whatever the reasons, data across each visit cannot automatically be assumed to align. Single-visit mosaic (SVM) processing attempts to correct these relative alignment errors and to align the data to an absolute astrometric frame so that all the data across all the filters used can be drizzled onto the same pixel grid.

Understanding the quality of the SVM products requires knowing what processing took place to generate that product, and more importantly, what limitations that processing may have had. The following sections provide the description of the single-visit processing. Definitions of a number of processing-specific terms used in this description can be found in the hap-glossary.

### 17.3.1 Primary User-Interface

One task has been written to perform the single-visit processing: `runsinglehap`. It gets used by STScI to generate the single-visit products which can be found in the Mikulsk Archive for Space Telescopes (MAST) archive. This task can also be run from the operating system command-line or from within a Python session to reproduce those results, or with modification of the input parameters, perhaps improve on the standard archived results. Full details on how to run this task can be found in the description of the task at *API for runsinglehap*.



## 17.3.2 Processing Steps

Single-visit processing performed by `runsinglehap` relies on the results of the standard astrometric processing of the individual exposures and associations as the starting point for alignment. This processing then follows these steps to create the final products:

- interpret the list of filenames for all exposures taken as part of a single visit
- copy the pipeline-calibrated (FLT/FLC) files to the current directory for processing
- rename the input files to conform to the single-visit naming conventions
  - This step insures that the original pipeline results remain available in the archive unchanged
- Define what output products can be generated
- Align all exposures in a relative sense (all to each other)
- Create a composite source catalog from all aligned input exposures
- Cross-match and fit this composite catalog to GAIA to determine new WCS solution
- Update renamed input exposures with results of alignment to GAIA
- Create each of the output products using the updated WCS solutions

## 17.3.3 Single Visit Naming Convention

All files processed as part of a single visit get renamed from the standard pipeline filenames into something which describes the data more clearly. The convention used for the names of these input and output files uses these components:

- **<propid>** : the proposal ID for this visit
- **<obsetid>** : the 2-digit visit ID from this proposal
- **<instr>** : 3 letter designation of the instrument used for the observations
- **<detector>** : name of the detector used for the observations
- **<filter>** : hyphen-separated list of filter names used for the observations
- **<ippssoo>** : standard 8 character rootname for a single exposure defined by the pipeline
- **<ippsss>** : standard 6 character designation of the **<instr>/<propid>/<obsetid>** for this visit
- **dr[*cz*].fits** : suffix for drizzled products
- **fl[*ct*].fits** : suffix for pipeline-calibrated files
- **asn.fits** : suffix for the association table
- **hlet.fits** : suffix for headerlet files containing the WCS solution used to create the final drizzle products/mosaics

These components get combined to create filenames specific to each type of file being processed. The following table provides a complete list of all the products created as a result of single-visit processing.

Table 1: Single-visit product filenames

Product	File Type	Filename for Files Produced	
Exposure	drizzle product	hst_<propid>_<obsetid>_<instr>_<detector>_<filter>_<ippssoo>_dr[cz].fits	
	flat-field product	hst_<propid>_<obsetid>_<instr>_<detector>_<filter>_<ippssoo>_fl[ct].fits	
	headerlet file	hst_<propid>_<obsetid>_<instr>_<detector>_<filter>_<ippssoo>_hlet.fits	
	trailer file	hst_<propid>_<obsetid>_<instr>_<detector>_<filter>_<ippssoo>_trl.txt	
	preview (full size)	hst_<propid>_<obsetid>_<instr>_<detector>_<filter>_<ippssoo>_dr[cz].jpg	
Filter	drizzle product	hst_<propid>_<obsetid>_<instr>_<detector>_<filter>_<ippss>_dr[cz].fits	
	point-source catalog	hst_<propid>_<obsetid>_<instr>_<detector>_<filter>_<ippss>_point-cat.ecsv	
	segment-source catalog	hst_<propid>_<obsetid>_<instr>_<detector>_<filter>_<ippss>_segment-cat.ecsv	
	trailer file	hst_<propid>_<obsetid>_<instr>_<detector>_<filter>_<ippss>_trl.txt	
	preview (full size)	hst_<propid>_<obsetid>_<instr>_<detector>_<filter>_<ippss>_dr[cz].jpg	
	preview (thumbnail)	hst_<propid>_<obsetid>_<instr>_<detector>_<filter>_<ippss>_dr[cz]_thumb.jpg	
	Total	drizzle product	hst_<propid>_<obsetid>_<instr>_<detector>_total_<ippss>_dr[cz].fits
		point-source catalog	hst_<propid>_<obsetid>_<instr>_<detector>_total_<ippss>_point-cat.ecsv
		segment-source catalog	hst_<propid>_<obsetid>_<instr>_<detector>_total_<ippss>_segment-cat.ecsv
		trailer file	hst_<propid>_<obsetid>_<instr>_<detector>_total_<ippss>_trl.txt
	preview (full size)	hst_<propid>_<obsetid>_<instr>_<detector>_total_<ippss>_dr[cz].jpg	
	preview (thumbnail)	hst_<propid>_<obsetid>_<instr>_<detector>_total_<ippss>_dr[cz]_thumb.jpg	
	color preview (full size)	hst_<propid>_<obsetid>_<instr>_<detector>_total_<ippss>_<filters>_dr[cz].jpg	
	color preview (thumbnail)	hst_<propid>_<obsetid>_<instr>_<detector>_total_<ippss>_<filters>_dr[cz]_thumb.jpg	

### 17.3.4 Processing the Input Data

SVM processing starts with a list of all the single exposures which were taken as part of a visit. Any associations which were defined by the proposal are ignored, since the visit itself gets treated, in essence, as a new association. The input files can be specified either using the **poller** file format used by the STScI automated processing or a file with a simple list of filenames (one filename per line).

#### Automated poller input file format

The automated processing performed to populate the MAST archive at STScI provides a file with the following format:

```
ic0s17h4q_flt.fits,12861,C0S,17,602.937317,F160W,IR,ic0s/ic0s17h4q/ic0s17h4q_
↳flt.fits
ic0s17h5q_flt.fits,12861,C0S,17,602.937317,F160W,IR,ic0s/ic0s17h5q/ic0s17h5q_
↳flt.fits
```

(continues on next page)

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```
ic0s17h7q_flt.fits,12861,C0S,17,602.937317,F160W,IR,ic0s/ic0s17h7q/ic0s17h7q_
↳flt.fits
ic0s17hhq_flt.fits,12861,C0S,17,602.937317,F160W,IR,ic0s/ic0s17hhq/ic0s17hhq_
↳flt.fits
```

This example comes from the ‘ic0s1’ visit where the columns are:

1. exposure filename
2. proposal ID (numeric value)
3. program ID - ppp value from exposure filename
4. obset\_id - visit number from proposal
5. exposure time of the exposure
6. filters used for the exposure
7. detector used to take the exposure
8. location of the exposure in a local cache

## Status of Input Data

The list of filenames which should be processed as a single-visit provides the raw science data for creating the new combined output products. However, these files need to be properly calibrated prior to SVM processing. Specifically, the exposures need to be:

- fully calibrated using the instruments calibration software, such as `calacs.e` for ACS and `calwf3.e` for WFC3 data. This should also include CTE-correction for the images whenever possible.
- processed using `runastrodriz` in order to apply the latest distortion model calibrations to the astrometry and to align the exposures as closely as possible to an external astrometric reference when possible.

These steps insure that the latest calibrations get applied to the data making it easier for the SVM processing to cross-match the data with minimal interference from artifacts in the data. In addition, the CTE-corrected versions of the data get used during pipeline processing in order to allow for better alignment of the exposures and to improve the photometry of the data as much as possible.

These processing steps can be verified in the input data using header keywords from the exposures

Table 2: Processing keywords

Header Keyword	Valid Values	Notes
FLATCORR	COMPLETED	Completion of basic calibration
DRIZCORR	COMPLETED	Completion of distortion calibration
WCSNAME	-FIT	Successful <b>a posteriori</b> alignment
	-HSC30	Successful <b>a priori</b> alignment
	-GSC240	Successful <b>a priori</b> alignment

The full set of possibilities for updated WCSs as reported using the `WCSNAME` keyword can be found in the description of the *Interpreting WCS names*.

As long as the input data meets these requirements, then SVM processing will have the best chance of success. Data which has not been able to be aligned successfully with an **a priori** or **a posteriori** solution can still be processed as part of a single-visit, however, the alignment may be more difficult to determine due to the larger uncertainties for HST pointing prior to October 2017.

## Filtering the input data

Not all HST imaging observations can be aligned using SVM processing. Observations taken with the GRISM or in SPATIAL SCAN mode result in sources which can not be aligned, for example. The *haputils.analyze* module evaluates all input exposures using these header keywords for the stated rejection criteria.

Table 3: Single-visit product filenames

Header Keyword	Values Which Trigger Rejection	Explanation
OBSTYPE	(not IMAGING)	Only Imaging mode data processed
MTFLAG	T	No moving targets, WCS and background sources vary
SCAN_TYP	C or D (or not N)	Can not align streaked sources
FILTER or FILTER1, FILTER2	G*, PR*, BLOCK	G=Grism and PR=Prism, Can not align streaked sources
EXPTIME	0	no exposure time, no data to align
TARGNAME	DARK, TUNGSTEN, BIAS, FLAT,	No alignable external sources in these calibration modes
	EARTH-CALIB, DEUTERIUM	No alignable external sources in these calibration modes
CHINJECT	not NONE	No alignable external sources in these calibration modes

Any observation which meets any of these criteria are flagged to be ignored (not processed). In addition, any data taken where the FGSLOCK keyword contains ‘COARSE’ or ‘GY’ will be flagged as potentially compromised in the comments generated during processing.

All observations which are alignable based on these criteria are then passed along as a table to create the SVM products. Those inputs which can be processed are then copied and renamed using the *Single Visit Naming Convention*. This insures that no SVM processing will affect or otherwise modify the original pipeline-processed input files. Only the SVM named input files will be updated with new SVM-aligned WCS solutions and then used to produce the drizzle products.

### 17.3.5 Defining the Output Products

The table with the set of observations which can be processed now gets interpreted. The goal is to identify what exposures can be combined to create unique products. This grouping will be used to create the **product list**. The **product list** is a Python list of `drizzlepac/haputils/product/HAPProduct` objects,

described in *drizzlepac.haputils.product* API docs, which represent each and every output product to be created for the visit. Each **Product** instance contains:

- list of filenames for all input exposures that will contribute to the output drizzle product
- WCS for output drizzle product
- pre-defined names for all output files associated with this **Product** including:
  - drizzle-combined image
  - point-source catalog determined from the drizzle-combined image
  - segmentation-based catalog determined from the drizzle-combined image
  - astrometric catalog used to align the input exposures
- methods for:
  - determining average number of images per pixel
  - defining the final WCS
  - aligning the exposures to an astrometric reference (GAIA)
  - applying the selected parameters to `AstroDrizzle`
  - drizzling the inputs to create the output drizzle product
  - determining the source catalogs from the drizzle product

This interpretation of the list of input filenames gets performed using the code in *drizzlepac.haputils.poller\_utils* by grouping similar observations. The rules used for grouping the inputs into output products result in outputs which have the same detector and filter. These output products are referred to as **filter products** defined as a `product/FilterProduct` instance.

All exposures for a single detector are also identified and grouped to define a **total product** using the `product/TotalProduct` class. This **total product** drizzle image provides the deepest available view of the field-of-view from this visit which will be used to produce the master catalog of sources for this visit. The master catalog of source positions will be used to perform photometry on each exposure, whether the source can be identified in the exposure at that position or not. This **forced photometry** results in limits for the photometry in cases where the sources are not bright enough to be identified in a given filter.

Two separate source catalogs for each filter are also pre-defined; namely,

- a point-source catalog derived using `photutils DAOSTarFinder`
- a segmentation-based catalog derived using `photutils segmentation` code

These two catalogs provide complimentary views of each field-of-view to try to highlight all types of compact sources found in the exposures.

## Example Visit

For example, a relatively simple visit of a fairly bright and crowded field with 6 F555W exposures (two 15-second and four 30-second exposures) and 6 F814W exposures (two 5-second and four 15-second exposures) would result in the definition of these output products:

- a drizzled image for each separate exposure
- a single F555W product
- a single F814W product, and
- a single **total product**
- a point-source catalog for the F555W product
- a segmentation-based source catalog for the F555W product
- a point-source catalog for the F814W product
- a segmentation-based source catalog for the F814W product
- a point-source catalog for the total product
- a segmentation-based catalog for the total product

The function `haputils.poller_utils.interpret_obset_input` serves as the sole interface for this interpretation. A basic tree gets defined (as a dictionary of dictionaries) by this function where the output exposures are identified along with all the names of the input exposures. This tree then serves as the basis for organizing the rest of the SVM processing.

In addition to defining what output products need to be generated, all the SVM products names are defined using the *Single Visit Naming Convention*. This insures that all the output products have filenames which are not only unique but also understandable (if a bit long) that are easily grouped on disk.

### 17.3.6 Aligning the Input Data

All input exposures should have already been aligned either individually or by association table as close to GAIA as possible during standard pipeline calibration processing. However, each exposure or association (of exposures) can be aligned to slightly different fits or catalogs due to differences in the source objects which can be identified in each separate exposure. The primary goal of SVM processing is to refine this alignment so that all exposures in the visit for the same detector (those exposures which contribute to each **total product**) share the same WCS (pixels on the sky).

Alignment of all the exposures for a **total product** uses the same alignment code as the standard calibration pipeline. The basic steps it follows is:

- generate a source catalog for each exposure (using `haputils.astrometric_utils`)
- obtain the WCS from each exposure
- perform a relative fit between the exposures using `tweakwcs`
- obtain an astrometric catalog for the field-of-view
- perform a final fit of all the exposures at once to the astrometric catalog
- update each WCS with the final corrected WCS generated by `tweakwcs`

The limits for performing the relative alignment and absolute fit to the astrometric catalog (defaults to **GAIADR2**) are lower under the expectation that large offsets (> 0.5 arcseconds) have already been removed in the pipeline processing. This makes the SVM alignment more robust across a wider range of types of

fields-of-view. The final updated WCS will be provided with a name that reflects this cross-filter alignment using **-FIT-SVM-<catalog name>** as the final half of the **WCSNAME** keyword. More details on the WCS naming conventions can be found in the *Interpreting WCS names* section.

### 17.3.7 Creating the Output Products

Successful alignment of the exposures allows them to be combined into the pre-defined output products; primarily, the **filter products** and the **total product**. These products get created using `drizzlepac.astrodrizzle.AstroDrizzle`.

#### Selecting Drizzle Parameters

Optimal parameters for creating every possible type of output product or mosaic would require knowledge of not only the input exposures, but also expert knowledge of the science. Parameters optimized for one science goal may not be optimal for another science goal. Therefore, automated pipeline processing has defined a basic set of parameters which will result in a reasonably consistent set of products as opposed to trying to optimize for any specific science case.

The default parameters have been included as part of the `drizzlepac` package in the `drizzlepac/pars/hap_pars` directory. Index JSON files provide the options that have been developed for selecting the best available default parameter set for processing. The INDEX JSON files point to different parameter files (also in JSON format) that are also stored in sub-directories the code organized by instrument and detector.

Selection criteria are also listed in these Index JSON files for each step in the SVM processing pipeline; namely,

- alignment
- astrodrizzle
- catalog generation
- quality control

Initially, only the **astrodrizzle** step defines any selection criteria for use in processing. The criteria is based on the number of images being combined for the specific instrument and detector of the exposures.

The SVM processing interprets the input data and verifies what input data can be processed. At that point, the code determines what selection criteria apply to the data and uses that to obtain the appropriate parameter settings for the processing steps. Applying the selection to select the appropriate parameter file simply requires matching up the key in the JSON file with the selection information. For example, a **filter product** would end up using the **filter\_basic** criteria, while an 8 exposure ACS/WFC association would end up selecting the **acs\_wfc\_any\_n6** entry.

#### User-customization of Parameters

The parameter configuration file now included in the `drizzlepac` package are designed to be easily customized for manual processing with both `runastrodriz` (pipeline astrometry processing) and

`runsinglehap` (SVM processing). These ASCII JSON files can be edited prior to manual reprocessing to include whatever custom settings would best suit the science needs of the research being performed with the data.

## Defining the Output WCS

The SVM processing steps through the **product list** to generate each of the pre-defined products one at a time after the input exposures have all been aligned. One of the primary goals of SVM processing is to produce combined images which share the same WCS for all the data from the same detector. This simply requires defining a common WCS which can be used to define the output for all the **filter products** from the visit.

The common WCS, or **metawcs**, gets defined by reading in all the WCS definitions as `stwcs.wcsutil.HSTWCS` objects for all the input exposures taken with the same **instrument** in the visit. This list of **HST-WCS** objects then gets fed to `stwcs.distortion.utils.output_wcs`, the same function used by `AstroDrizzle` to define the default output WCS when the user does not specify one before-hand. This results in the definition of a WCS which spans the entire field-of-view for all the input exposures with the same plate scale and orientation as the first **HSTWCS** in the input list. This **metawcs** then gets used to define the shape, size and WCS pointing for all drizzle products taken with the same detector in the visit.

## Drizzling

Each output product gets created using `AstroDrizzle`. This step:

- combines all the input exposures associated with the product
- uses the parameters read in from the configuration files
- defines the output image using the **metawcs** WCS definition
- writes out a multi-extension FITS (MEF) file for the drizzled image using the pre-defined name

This drizzled output image has the same structure as the standard pipeline drizzle products; namely,

- PRIMARY extension: all information common to the product such as instrument and detector.
- SCI extension: the drizzled science image along with header keywords describing the combined array such as total exposure time.
- WHT extension: an array reporting the drizzled weight for each pixel
- CON extension: an array reporting what input exposures contributed to each output pixel

The headers of each extension gets defined as using the `fitsblender` software with much the same rules used to create the standard pipeline drizzle product headers. In short, it uses simple rules files to determine what keywords should be kept in the output headers from all the input exposures, and how to select or compute the value from all the input headers for each keyword.



## Unique SVM Keywords

A small set of keywords have been added to the standard drizzle headers to reflect the unique characteristics of the SVM products. These keywords are:

**NPIXFRAC** Fraction of pixels with data

**MEANEXPT** Mean exposure time per pixel with data

**MEDEXPT** Median exposure time per pixel with data

**MEANNEXP** Mean number of exposures per pixel with data

**MEDNEXP** Median number of exposures per pixel with data

## Catalog Generation

SVM processing does not stop with the creation of the output drizzled images like the standard calibration pipeline. Instead, it derives 2 separate source catalogs from each drizzled **filter product** to provide a standardized measure of each visit. For more details on how the catalogs are produced, please refer to the *Catalog Generation* documentation page.

## 17.4 API for runsinglehap

The task `runsinglehap` serves as the primary interface for processing data from a single-visit into a uniform set of images.

`runsinglehap.py` - Module to control processing of single-visit mosaics

**License** *LICENSE*

USAGE: `runsinglehap [-d] inputFilename`

The ‘-d’ option will run this task in DEBUG mode producing additional outputs.

**Python USAGE:** `python from drizzlepac import runsinglehap runsinglehap.perform(inputFilename,debug=True)`

This script defines the HST Advanced Products (HAP) generation portion of the calibration pipeline. This portion of the pipeline produces mosaic and catalog products. This script provides similar functionality as compared to the Hubble Legacy Archive (HLA) pipeline in that it acts as the controller for the overall sequence of processing.

Note regarding logging. . . During instantiation of the log, the logging level is set to NOTSET which essentially means all message levels (debug, info, etc.) will be passed from the logger to the underlying handlers, and the handlers will dispatch all messages to the associated streams. When the command line option of setting the logging level is invoked, the logger basically filters which messages are passed on to the handlers according the level chosen. The logger is acting as a gate on the messages which are allowed to be passed to the handlers.

The output products can be evaluated to determine the quality of the alignment and output data through the use of the environment variable:

- `SVM_QUALITY_TESTING` : Turn on quality assessment processing. This environment variable, if found with an affirmative value, will turn on processing to generate a JSON file which contains the results of evaluating the quality of the generated products.

NOTE: Step 9 compares the output HAP products to the Hubble Legacy Archive (HLA) products. In order for step 9 (`run_sourcelist_comparison()`) to run, the following environment variables need to be set: - `HLA_CLASSIC_BASEPATH` - `HLA_BUILD_VER`

Alternatively, if the HLA classic path is unavailable, The comparison can be run using locally stored HLA classic files. The relevant HLA classic imagery and sourcelist files must be placed in a subdirectory of the current working directory called 'hla\_classic'.

```
drizzlepac.hapsequencer.run_hap_processing (input_filename,          diag-
                                           nostic_mode=False,
                                           use_defaults_configs=True,    in-
                                           put_custom_pars_file=None,
                                           output_custom_pars_file=None,
                                           phot_mode='both', log_level=20)
```

Run the HST Advanced Products (HAP) generation code. This routine is the sequencer or controller which invokes the high-level functionality to process the single visit data.

### Parameters

**input\_filename: string** The 'poller file' where each line contains information regarding an exposures taken during a single visit.

**diagnostic\_mode** [bool, optional] Allows printing of additional diagnostic information to the log. Also, can turn on creation and use of pickled information.

**use\_defaults\_configs: bool, optional** If True, use the configuration parameters in the 'default' portion of the configuration JSON files. If False, use the configuration parameters in the "parameters" portion of the file. The default is True.

**input\_custom\_pars\_file: string, optional** Represents a fully specified input filename of a configuration JSON file which has been customized for specialized processing. This file should contain ALL the input parameters necessary for processing. If there is a filename present for this parameter, the 'use\_defaults\_configs' parameter is ignored. The default is None.

**output\_custom\_pars\_file: string, optional** Fully specified output filename which contains all the configuration parameters available during the processing session. The default is None.

**phot\_mode** [str, optional] Which algorithm should be used to generate the sourcelists? 'aperture' for aperture photometry; 'segment' for segment map photometry; 'both' for both 'segment' and 'aperture'. Default value is 'both'.

**log\_level** [int, optional] The desired level of verbosity in the log statements displayed on the screen and written to the .log file. Default value is 20, or 'info'.

### Returns

**return\_value: integer** A return exit code used by the calling Condor/OWL workflow code: 0 (zero) for success, 1 for error

### 17.4.1 Supporting code

These modules and functions provide the core functionality for the single-visit processing.

#### drizzlepac.haputils.product

Definition of Super and Subclasses for the mosaic output image\_list

Classes which define the total (“white light” image), filter, and exposure drizzle products.

```
class drizzlepac.haputils.product.HAPPProduct (prop_id, obset_id, instrument,  
detector, filename, filetype,  
log_level)
```

HAPPProduct is the base class for the various products generated during the astrometry update and mosaicing of image data.

```
class drizzlepac.haputils.product.TotalProduct (prop_id, obset_id, instrument,  
detector, filename, filetype,  
log_level)
```

A Total Detection Product is a ‘white’ light mosaic comprised of images acquired with one instrument, one detector, all filters, and all exposure times. The exposure time characteristic may change - TBD.

The “tdp” is short hand for TotalProduct.

```
class drizzlepac.haputils.product.FilterProduct (prop_id, obset_id, instru-  
ment, detector, filename, fil-  
ters, filetype, log_level)
```

A Filter Detection Product is a mosaic comprised of images acquired during a single visit with one instrument, one detector, a single filter, and all exposure times. The exposure time characteristic may change - TBD.

The “fdp” is short hand for FilterProduct.

```
class drizzlepac.haputils.product.ExposureProduct (prop_id, obset_id, instru-  
ment, detector, filename,  
filters, filetype, log_level)
```

An Exposure Product is an individual exposure/image (flt/flc).

The “edp” is short hand for ExposureProduct.

#### drizzlepac.haputils.poller\_utils

Utilities to interpret the pipeline poller obset information and generate product filenames

The function, interpret\_obset\_input, parses the file generated by the pipeline poller, and produces a tree listing of the output products. The function, parse\_obset\_tree, converts the tree into product categories.

#### drizzlepac.haputils.catalog\_utils

This script contains code to support creation of photometric sourcelists using two techniques: aperture photometry and segmentation-map based photometry.

## drizzlepac.haputils.sourcelist\_generation

This script contains code to support creation of source extractor-like and daophot-like sourcelists.

## drizzlepac.haputils.photometry\_tools

Tools for aperture photometry with non native bg/error methods

This function serves to ease the computation of photometric magnitudes and errors using PhotUtils by replicating DAOPHOT's photometry and error methods. The formula for DAOPHOT's error is:

$$\text{err} = \sqrt{(\text{Poisson\_noise} / \text{epadu} + \text{area} * \text{stdev}^2 + \text{area}^2 * \text{stdev}^2 / \text{nsky})}$$

Which gives a magnitude error:

$$\text{mag\_err} = 1.0857 * \text{err} / \text{flux}$$

Where epadu is electrons per ADU (gain), area is the photometric aperture area, stdev is the uncertainty in the sky measurement and nsky is the sky annulus area. To get the uncertainty in the sky we must use a custom background tool, which also enables computation of the mean and median of the sky as well (more robust statistics). All the stats are sigma clipped. These are calculated by the functions in aperture\_stats\_tbl.

---

**Note:** Currently, the background computations will fully include a pixel that has ANY overlap with the background aperture (the annulus). This is to simplify the computation of the median, as a weighted median is nontrivial, and slower. Copied from [https://grit.stsci.edu/HLA/software/blob/master/HLApipeline/HLApipe/scripts/photometry\\_tools.py](https://grit.stsci.edu/HLA/software/blob/master/HLApipeline/HLApipe/scripts/photometry_tools.py)

---

## Authors

- Varun Bajaj, January 2018

## Use

```
from photometry_tools import iraf_style_photometry
phot_aps = CircularAperture((sources['xcentroid'], sources['ycentroid']), r=10.
↪)
bg_aps = CircularAnnulus((sources['xcentroid'], sources['ycentroid']), r_
↪in=13., r_out=15.)
```

Simplest call:

```
photometry_tbl = iraf_style_photometry(phot_aps, bg_aps, data)
```

Pass in pixelwise error and set background to mean

```
photometry_tbl = iraf_style_photometry(phot_aps, bg_aps, data, error_
↪array=data_err, bg_method='mean')
```

Can also set the gain (if image units are DN)

```
photometry_tbl = iraf_style_photometry(phot_aps, bg_aps, data, epadu=2.5)
```

## **Classes and Functions**

### **drizzlepac.haputils.processing\_utils**

Utilities to support creation of Hubble Advanced Pipeline(HAP) products.



---

## Astrometry and Advanced Pipeline Products

---

The `drizzlepac` package can be used for many purposes, all related to aligning and combining images to create products which can provide the deepest available views of the data. Combining the data with `drizzlepac` relies on the WCS solution specified in the input image headers. These WCS solutions are expected to align the images to each other (relative astrometry) as well as align the image to the correct position on the sky (absolute astrometry). The telemetry from HST allows the relative astrometry to be known extremely accurately (sub-milli-arcsecond level) when all images use the same guide stars and when the images were taken in the same visit. However, data taken at different times using different guide stars have historically had errors in the alignment with a sigma of 1 arc-second (or more). As a result, corrections to the alignment need to be made in order to successfully combine the images.

The code being used in the automated HST calibration pipeline to generate the products served by the HST Archive through the MAST Portal relies on multiple methods to do the best job possible in aligning and combining data regardless of whether they came from the same visit (or instrument) or not.

### 18.1 Headerlets and You: Astrometry in Drizzle Products

The astrometry for any given observation relies upon accurate pointing information from the telescope. However, HST has evolved over time since it was put into orbit, with the focus changing over time as the telescope desorbs. The instruments have also changed positions over time relative to the FGS guides, and the coordinates for the guide stars were originally determined using ground-based information. All this has limited the calculation of the pointing of any given observation on the sky (absolute astrometry) to no better than 1-2 arc-seconds.

Calibration of the distortion model for each instrument has been done well enough to allow observations to be combined (relative astrometry) with an accuracy of better than 5 milli-arcseconds. This has made the absolute astrometry inaccuracy stand out even more, making it more difficult to compare observations taken at different times or to compare HST data with observations from other telescopes (like Chandra).

Therefore, multiple efforts have been undertaken to improve the absolute astrometry to match the accuracy of the relative astrometry. These efforts have resulted in multiple new world coordinate systems (WCS) solutions to be developed for HST observations, starting with ACS, WFC3, and WFPC2. Complete details of the results of calibrating the distortion models and astrometry for each instrument can be found in each of the instruments web pages; namely,

- **ACS:** [ACS Distortion](#)
- **WFC3:** [WFC3 Data Handbook Chapter 4: WFC3 Images: Distortion Correction and AstroDrizzle](#)

Each solution has its own advantages and errors, making some good for one use but inadequate for others. As a result, **all new WCS solutions which are approved by STScI** are being offered with HST data provided by MAST with headerlets serving as the mechanism for providing and applying all WCS solutions.

### 18.1.1 Where are all the WCS solutions?

All calibrated HST products delivered by the Barbara A. Mikulski Archive for Space Telescopes (MAST) contain the best available WCS solutions available at the time the data was last processed. Unfortunately, only 1 WCS can be used at a time to transform the position in the image to an undistorted position on the sky. The FITS Standard, however, describes how multiple WCS solutions can be defined for an image in FITS Paper I (Greisen, E. W., and Calabretta, M. R., *Astronomy & Astrophysics*, 395, 1061-1075, 2002).

This standard defines a keyword, WCSNAME, which serves as the label for the WCS specified in the header of the observation. It also defines additional keywords, WCSNAME[A-Z], that represent additional WCS solutions which are specified as alternate WCS solutions in the same header as the active WCS associated with WCSNAME. Headerlets rely on the use of the WCSNAME keyword to manage the multiple solutions that may be defined for a given observation, with each WCS having it's own unique WCSNAME value. Headerlets also build the HDRNAME keyword which specifies the exposure name as well WCS to insure that this WCS will only ever be applied to this exposure and no other.

The use of headerlets extends this standard by allowing each new WCS, specified using the FITS Paper I standards along with the SIP convention, to be stored separately as a new extension at the end of the image's FITS file. This can be seen in the extensions listed by `astropy.fits` for a sample ACS/WFC image which includes 6 additional WCS solutions from the astrometry database.:

No.	Name	Ver	Type	Cards	Dimensions	Format
0	PRIMARY	1	PrimaryHDU	279	()	
1	SCI	1	ImageHDU	201	(4096, 2048)	float32
2	ERR	1	ImageHDU	57	(4096, 2048)	float32
3	DQ	1	ImageHDU	49	(4096, 2048)	int16
4	SCI	2	ImageHDU	199	(4096, 2048)	float32
5	ERR	2	ImageHDU	57	(4096, 2048)	float32
6	DQ	2	ImageHDU	49	(4096, 2048)	int16
7	D2IMARR	1	ImageHDU	15	(64, 32)	float32
8	D2IMARR	2	ImageHDU	15	(64, 32)	float32
9	D2IMARR	3	ImageHDU	15	(64, 32)	float32
10	D2IMARR	4	ImageHDU	15	(64, 32)	float32
11	WCSDVARR	1	ImageHDU	15	(64, 32)	float32
12	WCSDVARR	2	ImageHDU	15	(64, 32)	float32
13	WCSDVARR	3	ImageHDU	15	(64, 32)	float32
14	WCSDVARR	4	ImageHDU	15	(64, 32)	float32

(continues on next page)



(continued from previous page)

15	WCSCORR	1	BinTableHDU	59	14R x 24C	[40A, I, A, 24A, 24A, ↵ ↪24A, 24A, D, D, D, D, D, D, D, D, 24A, 24A, D, D, D, D, J, 40A, 128A]
16	HDRLET	1	HeaderletHDU	22	( )	
17	HDRLET	2	HeaderletHDU	22	( )	
18	HDRLET	3	HeaderletHDU	26	( )	
19	HDRLET	4	HeaderletHDU	26	( )	
20	HDRLET	5	HeaderletHDU	26	( )	
21	HDRLET	6	HeaderletHDU	26	( )	

In this example, extensions 16 through 21 contain headerlets each of which specifies different astrometric alignment (each may potentially be based on a different distortion model) and each has been assigned a unique name given by the **WCSNAME** keyword. A quick listing of the names of all these WCSs can be obtained using the Python library **STWCS**:

```
from stwcs.wcsutil import headerlet
headerlet.get_headerlet_kw_names("j95y04hq_flg.fits", kw='WCSNAME')
['OPUS',
 'IDC_0461802ej',
 'OPUS-GSC240',
 'IDC_0461802ej-GSC240',
 'OPUS-HSC30',
 'IDC_0461802ej-HSC30']
```

## Interpreting WCS names

The first WCS solution listed, **OPUS**, corresponds to the default WCS defined by the HST calibration pipeline based on original telemetry and only a first-order distortion model. Solutions which contain **IDC\_** are based on the distortion model based on the reference data associated with the **IDCTAB** reference file whose rootname is specified in the **WCSNAME**. For example, **IDC\_0461802ej** refers to a WCS solution based on the **IDCTAB** reference file `0461802ej_idc.fits` along with all the associated secondary distortion reference files given by **DGEOFILE**, **NPOLFILE** and, for WFPC2, **OFFTAB** and **DXYFILE**.

Additional WCS solutions will then be based on either the original **OPUS** WCS or the distortion-corrected **IDC\_** WCS solutions. Two types of solutions can be defined for images; namely, *a priori* and *a posteriori* solutions.

### *a priori* solutions

The *a priori* solutions have been determined for **ALL HST data** by correcting the coordinates of the guide stars that were used from the originally specified coordinates to the coordinates of those guide stars as determined by **GAIA**. The naming convention for these *a priori* solutions are:

```
<Starting WCS>-<Astrometric Catalog>

For example,
'IDC_0461802ej-GSC240'
```

where the **Astrometric Catalog** refers the exact astrometric catalog used to correct the guide star positions. A number of **Astrometric Catalogs** are available through MAST for aligning images. Solutions generated for the database were initially based on catalogs which were based on the GAIA catalog. These GAIA-based catalogs include:

- **GSC240**

- This catalog contains version 2.4.0 of the *Guide Star Coordinates* (GSC) catalog,
- All guide stars in the catalog were cross-matched with the GAIA DR1 catalog and corrected to the coordinates reported in GAIA DR1.
- **APPLIES TO:** All HST datasets which had a successful guide star acquisition, which is nearly all data in the archive.
- **NOTE:** HST Observations taken after September 2017.

- **HSC30**

- This catalog contains version 3.0 of the *Hubble Source Catalog* (HSC)
- Technically, this is an *a posteriori* solution, but it is applied blindly without further verification that the correction fully aligns the image to GAIA; hence, it is included as an *a priori* solution.
- Sources in the HSC were cross-matched with the GAIA DR1 catalog.
- Those cross-matched sources were then used to determine a fit to the GAIA catalog.
- The fit to GAIA was then applied to all remaining sources in the catalog.
- **APPLIES TO:** Only datasets which had a sufficient number of sources in the exposure to be aligned to GAIA by the *Hubble Legacy Archive* (HLA) project.

- **GAIADR1**

- A MAST-provided version of the first data release (DR1) version of the official GAIA astrometric catalog.
- This version does not have proper motions for a majority of the sources in the catalog.

- **GAIADR2**

- A MAST-provided version of the second data release version of the official GAIA astrometric catalog.
- This catalog contains initial proper motion measurements (and errors) for most sources in the catalog.

Although all solutions are appended to each FITS file, only 1 WCS (referred to as the **‘active’ WCS**) can be used at a time to represent the transformation from pixel coordinates to world coordinates. The active WCS is defined by the standard WCS keywords found in the header of the science extension for each chip in the exposure; *e.g.*, CRVAL1, CRVAL2, CRPIX1, CRPIX2, and so on.

The *a priori* solution which gets selected to replace the active WCS solution represents the most accurate solution available in the astrometry database at the time, and will be chosen based on the following hierarchy (as of Summer 2019):

1. HSC30

2. GSC240
3. IDC\_<rootname> (distortion-corrected pipeline default WCS)

If the first type of solution is not available, the next solution in the list is selected. As new solutions are added to the astrometry database, these rules will be modified to always try to return the WCS correction which best aligns the data to the GAIA catalog.

## a posteriori solutions

The *a posteriori* solutions, on the other hand, get determined from measuring sources in each image, finding overlapping sources from an astrometric catalog, identifying and cross-matching image sources with sources from the astrometric catalog and performing a fit to correct the WCS. These type of solutions can not be determined for all datasets due to a number of reasons, such as lack of sources in the image and/or lack of overlapping sources from an astrometric catalog. When these solutions can be determined for an observation, they are given a value for the WCSNAME keyword which follows the convention:

**<Starting WCS>-FIT\_<REL|IMG|EVM|SVM>\_<Astrometric Catalog>**

For example,

`'IDC_0461802ej-FIT_REL_GAIADR2'`

The terms are defined as:

- **<Starting WCS>**
  - Value of WCSNAME for the exposure prior to applying any astrometric Solutions
  - IDC\_<rootname> (like IDC\_041802ej) refers to a distortion-corrected model based on the IDCTAB reference file 0461802ej\_idc.fits.
- **'FIT'**
  - This term refers to the fact that sources from the image were identified, cross-matched and fit to sources from an astrometric catalog to create an *a posteriori* WCS solution.
- **'<REL|IMG|EVM|SVM>'**
  - REL : This term denotes the fact that all images were aligned relative (REL) to each other and then aligned to an astrometric catalog. This attempts to maintain the original relative alignment between the images in a given visit.
  - IMG : This term denotes the fact the the images were fit individually to the astrometric catalog. These solutions are applied only when relative alignment does not yield a viable fit to the astrometric catalog.
  - EVM : The cross-match and fit to an astrometric catalog was performed on a single exposure by itself as part of processing the exposures of an entire visit. This will typically only apply to those rare visits which do not have enough valid exposures in the visit for alignment.
  - SVM : This term refers to alignment of all the exposures in a single-visit to an astrometric catalog. The exposures of a visit are aligned to each other (relative alignment), then, as a group, all the exposures are cross-matched and fit to the astrometric catalog specified in the next term in the WCSNAME.

- **<Astrometric Catalog>**

- This term describes the astrometric catalog, as listed for use with the *a priori* solutions, which was used for the cross-matching and fitting sources identified in the image(s). If a value of **NONE** is specified here, it indicates that although the image appears (according to the code) to have been successfully relatively aligned one exposure to another, there were indications that the alignment to an astrometric catalog like **GAIADR2** failed. The user will need to carefully review the state of alignment of this data when **NONE** is listed in the output WCS.

These separate terms provide as succinct a description of the solution determined for and applied to the exposure as possible. Additional keywords have been written out to the headerlet extension for the *a posteriori* fit which further describe the solution, including:

- number of sources used in the fit
- RMS in RA and Dec of the fit
- parameters determined for the fit
- and more...

---

**Note:** A successfully determined *a posteriori* solution will **always** be used to replace the active WCS (after insuring the previous WCS has been saved as a headerlet extension already) regardless of the original solution.

---

## Pipeline Processing

All HST observations get processed in an automated environment using standard parameters for the calibration code, including the alignment and combination of individual exposures into undistorted products. The standard pipeline processing to create the undistorted drizzled images (drc.fits or drz.fits) gets performed using the ‘runastrodriz’ task in this package. This same processing can be run at any time using:

```
runastrodriz j8cw03010_asn.fits
runastrodriz j8cw03f6q_raw.fits
```

The files which need to be present are:

- RAW files (\*raw.fits)
- FLT files (\*flt.fits)
- FLC files (\*flc.fits, if any were created by the pipeline)
- ASN file (\*asn.fits, if applicable)

This processing includes a lot of logic intended to not only apply pre-defined (apriori) WCS solutions, but also to try and determine a new aposteriori solution then verify which solution (default pipeline, apriori or aposteriori) actually provides the WCS which comes closest to the GAIA astrometric frame. The [Pipeline Astrometric Calibration Description](#) of the runastrodriz task provides the full discussion of the logic used to define the defined ‘active’ WCS that gets used to create the products which get archived.

## Choosing a WCS

The **only** WCS solution that gets used to perform coordinate transformations on the pixel values will be the ‘active’ or ‘primary’ WCS associated with the WCSNAME keyword. The pipeline generated products will include an active WCS which the pipeline specifies as the *best* available WCS given the information used at the time of processing. However, this default ‘active’ WCS may not be appropriate for all science, so this WCS may need to be replaced by one of the other WCSs instead to best support the analysis necessary for the research.

## Dependent Packages

Working with the WCS solutions and headerlets gets performed using [STWCS package](#). Examples of how to work with this package will assume that the user has already installed this package into their working Python environment and has started a python shell. In addition, the following example relies on the Astropy IO package to work with the FITS headers and extensions.

Finally, the example described here will rely on additional functionality included in the V3.2.0 or later of the Drizzlepac package. These new functions support the generation of drizzle combined products which have been aligned to an astrometric standard catalog such as GAIA DR2.

## Sample Session

This example will work with 4 exposures taken using ACS/WFC as the association *j95y04010*, with most of the example being performed on the single exposure *j95y04hpq\_flc.fits*.

All the necessary libraries for working on this example can be imported using:

```
from drizzlepac.haputils import astroquery_utils as aquils
from drizzlepac.haputils import astrometric_utils as amutils
from astropy.io import fits
from stwcs.wcsutil import headerlet
```

The data can be obtained from MAST with `astroquery` using a simplified interface developed in `drizzlepac` using the commands:

```
filenames = aquils.retrieve_observation('j95y04010')
# filenames = ['j95y04hpq_flc.fits', 'j95y04hqq_flc.fits',
#             'j95y04hsq_flc.fits', 'j95y04huq_flc.fits']
```

The default ‘active’ WCS can be determined using:

```
default_wcsname = fits.getval(filenames[0], 'wcsname', ext=1)
```

For this example, we find that `default_wcsname='IDC_0461802ej'`, or as noted earlier, the default distortion correction based WCS provided by the pipeline with no correction for any astrometric catalogs, has been designated by the pipeline as the ‘active’ WCS.

All available WCSs provided by the astrometry database and attached as headerlet extensions can be queried to find the WCSNAMEs for all the new WCSs using:

```
new_wcsnames = headerlet.get_headerlet_kw_names(filenamees[0], kw='WCSNAME')
```

These will be the same ones listed earlier. For this example, we decide we would like to have this observation aligned using the guide stars corrected to GAIA DR1 through the use of the GSC240-based WCS; specifically, **IDC\_0461802ej-GSC240**. We can replace the ‘active’ WCS with this new one using:

```
new_hdrnames = headerlet.get_headerlet_kw_names(filenamees[0])
# identify hdrname that corresponds with desired WCS with name of IDC_
  ↳ 041802ej-GSC240
new_wcs = new_hdrnames[new_wcsnames.index('IDC_0416802ej-GSC240')]
headerlet.restore_from_headerlet(filenamees[0], hdrname=new_wcs, force=True)
# confirm new WCS is now 'active'
fits.getval(filenamees[0], 'wcsname', ext=1)
```

At this point, the exposure has been updated to perform all coordinate transformations with the new GAIA DR1-based WCS as if the guide stars used for taking the observation has GAIA DR1 coordinates in the first place. This will not mean it will align perfectly with GAIA, but should be within 0.5 arcseconds due to drift in the telescope field-of-view for each of the instruments relative to the FGSs.

## 18.1.2 Headerlet Primer

The headerlet file itself conforms to FITS standards with the PRIMARY header containing global information about the WCS solution and how it was determined. Separate extensions in the headerlet then contain the header keywords for specifying the WCS for each chip in the exposure or for the distortion information necessary to correct the pixel positions from the image to the un-distorted position on the sky. These solutions rely on calibration reference data that describe the distortion observed in each instrument to better than 0.1 pixels in each detector. Instead of having to retrieve separate files with this distortion information, that distortion information has been folded into the header of each WFC3, ACS and WFPC2 dataset.

### Headerlet File Structure

This new object complete with the NPOLFILE and the D2IMFILE extensions derived from the full FITS file fully describes the WCS of each chip and serves without further modification as the definition of the headerlet. The listing of the FITS extensions for a headerlet for the sample ACS/WFC exposure after writing it out to a file would then be:

EXT#	FITSNAME	FILENAME	EXTVE	DIMENS	BITPI	OBJECT
0	j8hw27c4q	j8hw27c4q_hdr.fits			16	
1	IMAGE	D2IMARR	1	4096	-32	
2	IMAGE	WCSDVARR	1	64x32	-32	
3	IMAGE	WCSDVARR	2	64x32	-32	
4	IMAGE	WCSDVARR	3	64x32	-32	
5	IMAGE	WCSDVARR	4	64x32	-32	
6	IMAGE	SIPWCS	1		8	
7	IMAGE	SIPWCS	2		8	

## **Detailed Description of headerlet**

The full details on the headerlet, it's required set of keywords, and how the distortion models get described in the headerlet can be found in the [Technical Report on headerlets](#).

## **Code Interface to headerlets**

The [STWCS package](#) provides the code used to work with headerlets and WCS solutions.





## CHAPTER 19

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