



reftools Documentation

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MAKING IMPHTTAB TABLES

Making IMPHTTAB tables requires a connection to CDBS and that the `PYSYN_CDBS` environmental keyword point to CDBS.

Use `reftools.mkimphttab.create_table()` and provide the name of your output file and the base observation mode from which to build a set observation modes for which photometry keywords will be calculated.

Examples of a base observation mode include "acs,hrc", "wfc3,uvis1", or "cos".

1.1 Examples

```
>>> from reftools import mkimphttab
>>> mkimphttab.create_table('acs_wfc1', 'acs,wfc1', 'WFC1',
...                        'Mar 01 2002 00:00:00', clobber=True, verbose=False)
>>> mkimphttab.create_table('acs_sbc', 'acs,sbc', 'SBC',
...                        'Mar 01 2002 00:00:00', clobber=True)
```

```
reftools.mkimphttab.create_table(output, basemode, detector, useafter, tmgtab=None, tmctab=None, tmttab=None, mode_list=[], nmodes=None, clobber=False, verbose=True)
```

Create an IMPHTTAB reference file for a specified base configuration, `basemode`.

Parameters

output : str

Output IMPHTTAB filename. (`_imp.fits` will be appended if only prefix is given.)

basemode : str

Base observation mode for which to generate IMPHTTAB (e.g., `acs,hrc`). This is ignored if `mode_list` is given.

detector : str

Detector name.

useafter : str

Useafter date in the format of `MMM DD YYYY HH:MM:SS`.

tmgtab, tmctab, tmttab : str, optional

Graph (TMG), component (TMC), and thermal component (TMT) tables to use. If `None`, the most recent version in CDBS is used.

mode_list : list of str, optional

A list of observation modes which should be used to make an IMPHTTAB reference file. If given, `basemode` is ignored.

nmodes : int, optional

Set to limit the number of modes to calculate. This is for testing only, otherwise set to `None`.

clobber : bool, optional

Overwrite existing IMPHTTAB?

verbose : bool, optional

Display extra information.

`reftools.mkimphttab.create_table_from_table(output, detector, useafter, imphttab, **kwargs)`

Use a previously created IMPHTTAB reference file to generate a new IMPHTTAB reference file.

Parameters

output, detector, useafter : str

See `create_table()`.

imphttab : str

File name of `_imp.fits` IMPHTTAB table from which to take observation modes.

kwargs : dict

Keywords accepted by `create_table()`, except `mode_list`.

`reftools.mkimphttab.create_nicmos_table(output, detector, useafter, pht_table, **kwargs)`

Use a NICMOS `_pht.fits` table to generate an IMPHTTAB table for observation modes listed in the given table.

Parameters

output, detector, useafter : str

See `create_table()`.

pht_table : str

File name of `_pht.fits` table from which to take observation modes.

kwargs : dict

Keywords accepted by `create_table()`, except `mode_list`.

`reftools.mkimphttab.compute_values(obsmode, component_dict)`

Compute the 3 photometric values needed for a given observation mode string using `pysynphot`.

Values calculated:

- PHOTFLAM - Unit response in FLAM.
- PHOTPLAM - Pivot wavelength.
- PHOTBW - Bandwidth.

Parameters

obsmode : str

Observation mode for which to calculate values.

component_dict : dict

A dictionary in which to cache opened component objects. May be empty.

Returns

valdict : dict

Dictionary with photometry keywords as keys.

`reftools.mkimphttab.compute_synphot_values` (*obsmode*)

Calculate the same values as `compute_values()` but using IRAF SYNPHOT.

Note: This is replaced by `compute_values()` but kept for debugging.

Parameters

obsmode : str

Observation mode for which to calculate values.

Returns

valdict : dict

Dictionary with photometry keywords as keys.

COMPARE IMPHTTAB TABLES

Tools for comparing two IMPHTTAB tables from the same instrument and detector.

`class reftools.imphtcomp.ImphttabComp (tab1, tab2)`

Class for comparing two IMPHTTAB tables from the same instrument and detector.

Parameters

tab1 : str

Filename of first IMPHTTAB for comparison.

tab2 : str

Filename of second IMPHTTAB for comparison.

Attributes

tab1_name	str	Filename of first IMPHTTAB.
tab2_name	str	Filename of second IMPHTTAB.
modes	ndarray	Obsmodes present in both input files.
flams1	ndarray	PHOTFLAM values from <code>tab1</code> for obsmodes in <code>modes</code> .
plams1	ndarray	PHOTPLAM values from <code>tab1</code> for obsmodes in <code>modes</code> .
bws1	ndarray	PHOTBW values from <code>tab1</code> for obsmodes in <code>modes</code> .
flams2	ndarray	PHOTFLAM values from <code>tab2</code> for obsmodes in <code>modes</code> .
plams2	ndarray	PHOTPLAM values from <code>tab2</code> for obsmodes in <code>modes</code> .
bws2	ndarray	PHOTBW values from <code>tab2</code> for obsmodes in <code>modes</code> .
flamdiff	ndarray	Percent differences between <code>flams1</code> and <code>flams2</code> calculated as $(flams1 - flams2) / flams1$.
plamdiff	ndarray	Percent differences between <code>plams1</code> and <code>plams2</code> calculated as $(plams1 - plams2) / plams1$.
bwdiff	ndarray	Percent differences between <code>bws1</code> and <code>bws2</code> calculated as $(bws1 - bws2) / bws1$.

`make_plot (outname='imphttab_comp.pdf')`

Make a plot with histograms of the percent differences between PHOTFLAM, PHOTPLAM, and PHOTBW for the IMPHTTAB tables.

Differences plotted are $100 * (table1 - table2) / table1$.

Parameters

outname : str, optional

Filename of output plot, including extension. Defaults to 'imphttab_comp.pdf'.

print_diffs (*orderby='photflam', lines=25*)

Print obsmodes and parameters ordered by orderby parameter, with the largest absolute differences in that parameter at the top. This is for seeing which obsmodes have the largest difference in the specified parameter. Prints the number of modes given in the lines parameter.

Differences shown are calculated as $100 * (table1 - table2)/table1$.

Parameters

orderby : str, optional

The parameter by which to order the printed results, with modes having the largest absolute difference in this parameter printed at the top.

May be one of: 'photflam', 'photplam', or 'photbw'. An ImphtcompError is raised if the input does not match one of these.

Defaults to 'photflam'.

lines : int, optional

The number of lines to print. Defaults to 25.

Raises

ImphtcompError :

If orderby does not match a valid option.

reftools.imphtcomp.**plot_table_diffs** (*table1, table2, outname='imphttab_comp.pdf'*)

Make a plot with histograms of the percent differences between PHOTFLAM, PHOTPLAM, and PHOTBW for the IMPHTTAB tables.

Differences plotted are $100 * (table1 - table2) / table1$.

Parameters

table1 : str

Filename of first IMPHTTAB for comparison.

table2 : str

Filename of the second IMPHTTAB for comparison.

outname : str, optional

Filename of output plot, including extension. Defaults to 'imphttab_comp.pdf'.

reftools.imphtcomp.**print_table_diffs** (*table1, table2, orderby='photflam', lines=25*)

Compare two IMPHTTAB tables and print their differences to the terminal. Prints any obsmodes which are in either table but not in both.

Also prints the obsmodes and parameters for the modes which most differ in the parameter given in orderby. This is for seeing which obsmodes have the largest percent difference in the specified parameter. Prints the number of modes given in the lines parameter.

Differences shown are calculated as $100 * (table1 - table2)/table1$.

Parameters

table1 : str

Filename of first IMPHTTAB for comparison.

table2 : str

Filename of the second IMPHTTAB for comparison.

orderby : str, optional

This specifies one of 'photflam', 'photplam', 'photbw', or 'all'. The printed results are ordered according to the absolute difference in the specified parameter, with the mode with the largest absolute difference at the top of the list.

Specifying 'all' will print 3 tables, one ordered by each of the parameters.

Defaults to 'photflam'.

lines : int, optional

Number of lines of differences to print. Defaults to 25.

Raises

ImphtcompError :

If `orderby` does not match a valid option.

COMPARE SYNPHOT AND PYSYNPHOT

Tools for comparing pysynphot and synphot photometry calculations.

class `reftools.synpysyncomp.SynPysynComp` (*imphttab*)

Compare synphot and pysynphot values for obsmodes listed in the input IMPHTTAB. Only obsmodes with three or fewer listed components are compared. For example, `acs,wfc1,f555w` would be processed but not `acs,wfc1,f555w,MJD#`.

This can be a long process for large numbers of obsmodes so it is recommended that users use the CSV functionality of this class to save the results to a .csv file and then use the supporting functions `print_synpysyn_diffs` and `plot_synpysyn_diffs` in this module to investigate them.

Parameters

imphttab : str

Filename of IMPHTTAB from which to take obsmodes for comparison.

calculate_diffs (*verbose=False*)

Calculate diffs for all obsmodes and return.

Parameters

verbose: bool, optional :

If True, print obsmodes as they are processed. Defaults to False.

Returns

res : dict

Dictionary containing lists of all pysynphot and synphot calculated values and their differences calculated as $(\text{pysynphot} - \text{synphot})/\text{synphot}$.

comp_synpysyn (*mode*)

Returns a dictionary of pysynphot and synphot values and their percent differences calculated as $(\text{pysynphot} - \text{synphot})/\text{synphot}$.

Parameters

mode : str

Obsmode string.

Returns

comp : dict

Dictionary containing calculated values.

get_pysyn_vals (*mode*)

Get comparison values from pysynphot.

Parameters**mode** : str

Obsmode string.

Returns**ret** : dict

Dictionary containing calculated values.

get_syn_vals (*mode*)

Get comparison values from synphot.

Parameters**mode** : str

Obsmode string.

Returns**ret** : dict

Dictionary containing calculated values.

write_csv (*outfile*, *verbose=False*)Write a CSV file containing the pysynphot and synphot values for all obsmodes and their percent differences calculated as $(\text{pysynphot} - \text{synphot})/\text{synphot}$.**Parameters****outfile** : str

Name of file to write to.

verbose : bool, optional

If True, print obsmodes as they are processed. Defaults to False.

class reftools.synpysyncomp.**SynPysynPlot** (*csvfile*)Make a plot from a CSV file created by `SynPysynComp.write_csv` illustrating differences between synphot and pysynphot calculated products.The plots show differences between 7 parameters calculated by both synphot and pysynphot. The differences are shown as histograms. The difference plotted is calculated as $100 * (\text{pysynphot} - \text{synphot}) / \text{synphot}$.**Parameters****csvfile** : strFilename of input CSV file containing comparison results. Should be a file created by `SynPysynComp.write_csv`.**Attributes :****fig** : matplotlib.figure.FigureUseful for setting manipulating figure properties such as the title, which otherwise defaults to `csvfile`.**save_plot** (*outname='synpysyn_comp.pdf'*)

Save plots to a file.

Parameters**outname** : str, optional

Name of file to save to, including extension. Defaults to 'synpysyn_comp.pdf'.

`reftools.synpysyncomp.plot_synpysyn_diffs(csvfile, outname='synpysyn_comp.pdf')`

Make and save a plot illustrating differences between parameters calculated by both `synphot` and `pysynphot`. Data are taken from a CSV file made by `SynPysynComp.write_csv`.

Differences are shown as histograms. The difference plotted is calculated as $100 * (\text{pysynphot} - \text{synphot}) / \text{synphot}$.

Parameters

csvfile : str

Filename of input CSV file containing comparison results. Should be a file created by `SynPysynComp.write_csv`.

outname : str, optional

Name of file to save to, including extension. Defaults to 'synpysyn_comp.pdf'.

`reftools.synpysyncomp.print_synpysyn_diffs(csvfile, orderby='photflam', lines=25)`

Print `synphot/pysynphot` comparison results from a CSV file produced by `SynPysynComp.write_csv` to the terminal.

Prints the obsmodes and parameters for the modes which most differ in the parameter given in `orderby`. This is for seeing which obsmodes have the largest percent difference in the specified parameter. Prints the number of modes given in the `lines` parameter.

Only prints data for the parameters PHOTFLAM, PHOTPLAM (PIVOT WV), and PHOTBW (RMSWIDTH). Differences are given as $100 * (\text{pysynphot} - \text{synphot}) / \text{synphot}$.

Parameters

csvfile : str

Filename of input CSV file containing comparison results. Should be a file created by `SynPysynComp.write_csv`.

orderby : str, optional

This specifies one of 'photflam', 'photplam', 'photbw', or 'all'. The printed results are ordered according to the absolute difference in the specified parameter, with the mode with the largest absolute difference at the top of the list.

Specifying 'all' will print 3 tables, one ordered by each of the parameters.

Defaults to 'photflam'.

lines : int, optional

Number of lines of differences to print. Defaults to 25.

Raises

SynPysynCompError :

If `orderby` does not match a valid option.

`reftools.synpysyncomp.read_synpysyn(csvfile)`

Read a CSV file created by `SynPysynComp.write_csv` into numpy arrays.

Parameters

csvfile : str

Filename of CSV to read. Should be a file created by `SynPysynComp.write_csv`.

Returns

res : dict

Contains one field for each column of the CSV file with a numpy array in that field.

CALCULATE PHOTOMETRY KEYWORDS FROM IMPHTTAB

getphotpars includes utilities for calculating the photometry keywords PHOTZPT, PHOTFLAM, PHOTPLAM, and PHOTBW for a given obsmode and IMPHTTAB. The calculations are performed in the same way here as they are in hstcal pipelines.

To calculate a single set of keywords use the function `get_phot_pars`.

If you are calculating for several obsmodes from a single IMPHTTAB file it's best to use the `GetPhotPars` class. For example:

```
get_phot = GetPhotPars(imphttab)
for obs in obsmodes:
    photzpt, photflam, photplam, photbw = get_phot.get_phot_pars(obs)
    ...
get_phot.close()
```

exception `reftools.getphotpars.ImphttabError`

Class for errors associated with the imphttab file.

class `reftools.getphotpars.GetPhotPars(imphttab)`

This object can be used to get photometry parameters from a given IMPHTTAB reference file. Initialize with the name of an IMPHTTAB reference file and then call this class or the `get_phot_pars` method with a complete obsmode to get the photometry parameters.

Example obsmodes are:

'acs,wfc1,f625w,f660n' 'acs,wfc1,f625w,f814w,MJD#55000.0' 'acs,wfc1,f625w,fr505n#5000.0,MJD#55000.0'

Parameters

imphttab : str

Filename and path of IMPHTTAB reference file.

Attributes

<code>impht-tab_name</code>	str	Filename and path of IMPHTTAB reference file. Same as input <code>imphttab</code> .
<code>imphttab_fits</code>	<code>pyfits.HDUList</code>	Open <code>pyfits.HDUList</code> object from <code>imphttab</code> .

close()

Close `imphttab_fits` attribute.

get_phot_pars(obsmode)

Return PHOTZPT, PHOTFLAM, PHOTPLAM, and PHOTBW for specified obsmode.

Parameters

obsmode : str

obsmode string

Returns

photzpt : float

PHOTZPT from `imphttab_fits` header.

photflam : float

Interpolated PHOTFLAM for `obsmode`.

photplam : float

Interpolated PHOTPLAM for `obsmode`.

photbw : float

Interpolated PHOTBW for `obsmode`.

`reftools.getphotpars.get_phot_pars(obsmode, imphttab)`

Return PHOTZPT, PHOTFLAM, PHOTPLAM, and PHOTBW for specified `obsmode` and `imphttab`.

Parameters

obsmode : str

Complete `obsmode` string including any parameterized values.

Example obsmodes are:

'acs,wfc1,f625w,f660n'

'acs,wfc1,f625w,f814w,MJD#55000.0'

'acs,wfc1,f625w,fr505n#5000.0,MJD#55000.0'

imphttab : str

Path and filename of IMPHTTAB reference file.

Returns

photzpt : float

PHOTZPT from IMPHTTAB header.

photflam : float

Interpolated PHOTFLAM for `obsmode`.

photplam : float

Interpolated PHOTPLAM for `obsmode`.

photbw : float

Interpolated PHOTBW for `obsmode`.

MAKE PCTETAB REFERENCE FILE

The PCTETAB reference file contains data in both its primary header and in several table extensions. The parameters that go into the primary header are single numbers that must be specified in the call to `MakePCTETab`.

The data that goes into the table extensions is kept in text files. The names of these text files are given to `MakePCTETab`, which reads them to populate the table extensions. See the documentation for `MakePCTETab` for more detailed descriptions, argument names, and default values.

5.1 Primary Header Parameters

- Number of times the readout is simulated to arrive at the corrected image.
- Number of times the pixels are shifted per readout simulation.
- The read noise, in electrons, of the image. This is technically different for each amp but here we pick a single representative value.
- The default value selecting a model for how read noise is handled before CTE correction.
- Threshold for re-correcting over-subtracted pixels. Sometimes the CTE correction removes too much flux from a trail leaving a large divot.

5.2 Table Extensions

Functions for ACS PCTETAB reference file.

Authors

Pey Lian Lim, Matt Davis

Organization

Space Telescope Science Institute

History

- 2010-08-31 PLL created this module.
- 2010-11-09 PLL added RN2_NIT keyword and updated documentation.
- 2011-04-25 MRD updated for new CTE algorithm parameters
- 2011-07-18 MRD updated to handle time dependence
- 2011-11-29 MRD updated with column-by-column CTE scaling

- 2013-08-13 PLL removed deprecated PyFITS calls and cleaned up codes.

5.2.1 Examples

```
>>> from reftools import pctetab
>>> pctetab.MakePCTETab(
...     'pctetab_pcte.fits', 'pctetab_dtde.txt',
...     ['pctetab_chgleak-1.txt', 'pctetab_chgleak-2.txt'],
...     'pctetab_levels.txt', 'pctetab_scaling.txt',
...     'pctetab_column_scaling.txt', history_file='pctetab_history.txt')
```

exception reftools.pctetab.PCTEFileError

Generic exception for errors in this module.

```
reftools.pctetab.MakePCTETab(out_name, dtde_file, chg_leak_file, levels_file, scale_file, col-
                             umn_file, sim_nit=7, shft_nit=7, read_noise=5.0, noise_model=1,
                             oversub_thresh=-10, useafter='Mar 01 2002 00:00:00', pedi-
                             gree='INFLIGHT 01/03/2002 22/07/2010', creatorName='ACS
                             Team', history_file='', detector='WFC')
```

Make the CTE parameters reference file.

Parameters

out_name : str

Name of pcte fits file being created. May include path.

dtde_file : str

Path to text file containing dtde data.

The file should have 2 columns with the following format:

```
DTDE  Q
float int
... ..
```

Lines beginning with # are ignored.

chg_leak_file : str or list of str

Path to text file(s) containing charge leak data. If passed as a string the string may contain wild cards so that multiple files are specified.

The input file should contain 5 columns with following format:

```
NODE LOG_Q_1 LOG_Q_2 LOG_Q_3 LOG_Q_4
int  float  float  float  float
... ..   ...   ...   ...
```

Lines beginning with # are ignored.

levels_file : str

Text file containing charge levels at which to do CTE evaluation.

The input file should have a single column with the following format:

```
LEVELS
int
...
```

Lines beginning with # are ignored.

scale_file : str

Text file containing CTE scaling parameters

The input file should have two columns with the following format:

```
MJD      SCALE
float    float
...      ...
```

Lines beginning with # are ignored.

column_file : str

Text file containing CTE column-by-column scaling.

The input file should have 5 columns with the following format:

```
COLUMN  AMPA    AMPB    AMPC    AMPD
int      float  float  float  float
...      ...    ...    ...    ...
```

Lines beginning with # are ignored.

sim_nit : int, optional

Number of iterations of readout simulation per column.

shft_nit : int, optional

Number of shifts each readout simulation is broken up into. A large number means pixels are shifted a smaller number of rows before the CTE is evaluated again.

read_noise : float

Value for RN_CLIP keyword in PCTEFILE EXT 0. This is the maximum amplitude of read noise used in the read noise mitigation. Unit is in electrons.

noise_model : {0, 1, 2}

Select the method to be used for readnoise removal:

```
0: no read noise smoothing
1: standard smoothing
2: strong smoothing
```

oversub_thresh : float

Value for SUBTHRSH keyword in PCTEFILE header. CTE corrected pixels taken below this value are re-corrected. Unit is in electrons.

useafter : str, optional

Value for USEAFTER keyword. Defaults to 'Mar 01 2002 00:00:00'

pedigree : str, optional

Value for PEDIGREE keyword. Defaults to 'INFLIGHT 01/03/2002 22/07/2010'

creatorName : str, optional

Name of the person generating this fitsFile. Defaults to 'ACS Team'

historyFile : str, optional

ASCII file containing HISTORY lines for EXT 0. Include path. Each row will produce one HISTORY line. Defaults to ‘

detector : str, optional

Supported detector. Defaults to ‘WFC’

Examples

Saving file `pctetab_pcte.fits` with the command:

```
>>> MakePCTETab(  
...     'pctetab_pcte.fits', 'pctetab_dtdel.txt',  
...     ['pctetab_chgleak-1.txt', 'pctetab_chgleak-2.txt'],  
...     'pctetab_levels.txt', 'pctetab_scaling.txt',  
...     'pctetab_column_scaling.txt', history_file='pctetab_history.txt')
```


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