



# **nictools Documentation**

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

Contents:



## SAACLEAN

**saaclean: Module for estimating and removing persistent CR signal due to a prior**

SAA passage.

**Usage**

Normally used via the STSDAS task saaclean in the nicmos package. To use as pure python, create a params object to override any of the default parameters if desired, then invoke clean:

```
>>> mypars=saaclean.params(thresh=0.23)
>>> saaclean.clean('inputfile.fits','outputfile.fits',pars=mypars)
```

**For more information**

Additional user information, including parameter definitions and more examples, can be found in the help file for the STSDAS saaclean task, located in `nicmos$doc/saaclean.hlp`.

The algorithm and IDL prototype are described in the NICMOS ISR 2003-009, by Bergeron and Dickinson, available through the NICMOS webpage.

**Dependencies**

- numpy 1.0.2.dev3534 or higher
- pyfits v1.1b4 or higher
- imagestats v1.3 or higher

```
exception nictools.saaclean.AlreadyDone
```

```
exception nictools.saaclean.BadThreshError
```

```
exception nictools.saaclean.InsuffImprovement
```

```
exception nictools.saaclean.NegScaleError
```

```
exception nictools.saaclean.NoPersistError
```

```
class nictools.saaclean.Domain(name, pixellist, range)
```

Stores a list of pixels for a (typically high or low) signal domain

```
getmin ()
```

```
stripowerthan (factor)
```

`self.pp` is defined in `Exposure.getscales` It contains the (bin, stddev, mode) for the statistical analysis.

`striplowerthan(factor)` examines the `stddev` column only, and replaces all values of the `stddev` that are less than `factor*the zeroth bin`, with the maximum `stddev`.

`writeto(filename, clobber=False)`

**class** `nictools.saaclean.Exposure` (*imgfile, nickname=None*)

Stores a collection of keywords and the image data for an exposure.

`apply_domains` (*saaper, badmask, noisethresh, appimage=None*)

`apply_mask` (*mask*)

`dark_subtract` (*dark*)

`getmask` (*dim=256, border=3, writename='mask.dat', clobber=False*)

Computes a mask to use for pixels to omit

`getscales` (*saaper, mask, pars*)

`pedskyish` ()

Performs something like the IRAF `pedsky` task, but with a bit more sophistication in handling the central row and column

`update_header` (*pars, tag='default', header=None*)

Update the FITS header with all this good stuff we've done

`writeto(outname, clobber=False)`

**class** `nictools.saaclean.params` (*scale=0.54, wf1=0.7, wf2=0.3, stepsize=0.008, thresh=None, hirange=0.4, lorange=0.25, dofit=1, crthresh=0.3, noisethresh=1.0, binsigfrac=0.3, readsaaper=False, writesaaper=True, saaperfile='saaper.fits', fitthresh=True, histbinwidth=0.01, nclip=10, clobber=False, flatsaaper=True, flatsaaperfile=None, maskfile=None, darkpath=None, diagfile=None*)

`nictools.saaclean.clean` (*usr\_calfile, usr\_targfile, usr\_outfile, pars=None*)

`nictools.saaclean.create_saaper_header` (*im1, im2, saaper*)

`nictools.saaclean.flat_saaper` (*saaper, img*)

`nictools.saaclean.gausspoly_eval` (*coeffs, t*)

`nictools.saaclean.gausspoly_fit` (*thedata, guesscoeff*)

`nictools.saaclean.gausspoly_model` (*coeffs, t*)

`nictools.saaclean.get_dark_data` (*imgfile, darkpath*)



`nictools.saaclean.get_postsaa_darks` (*imgfile*)

Return the filenames containing the post-saa dark exposures, if present. Otherwise raise an exception and exit.

`nictools.saaclean.getdark` (*camera, tdkfile, darkpath*)

Get the right dark file for a given NICMOS camera. This is definitely not the right way to do this.

`nictools.saaclean.make_saaper` (*imgfile, pars, crthresh=1*)

`nictools.saaclean.median` (*a*)

`nictools.saaclean.osfn` (*filename*)

Return a filename with iraf syntax and os environment names substituted out

`nictools.saaclean.parabola_min` (*thedata, startguess*)

`nictools.saaclean.parabola_model` (*coeffs, t*)

`nictools.saaclean.smartopen` (*fname, mode, clobber=True*)

Allows specifying a clobber behavior

`nictools.saaclean.thresh_from_gausspoly_fit` (*saa, parbinwidth=0.5, nclip=3, diag-  
file=None, clobber=False*)

Some massaging of the SAApersistence image histogram is performed in order to obtain an optimal fit. Unfortunately this involves some magic numbers taken from the IDL code.

`nictools.saaclean.writeimage` (*image, filename, header=None, comment=None, clobber=False*)



## NIC\_REM\_PERSIST

```
class nictools.nic_rem_persist.NicRemPersist (calcfile, targfile, verbosity=1, per-
                                             sist_lo=None, used_lo=None, per-
                                             sist_model=None, persist_mask=None,
                                             run_stdout=None)
```

Remove bright earth persistence persistence from CAL file of NICMOS data (on which pedsub has been run) using a method based on correction calculated from PED file.

### Notes

The syntax for using this class under Python or PyRAF:

```
--> nrp = nic_rem_persist.NicRemPersist('n9r7b2bjq_ped.fits', persist_model = 'persistentstring.fits',
                                         used_lo = .3, persist_lo = 0.8)
--> nrp.persist()
```

The linux command line syntax for calling this module:

```
hal> ./nic_rem_persist.py 'n9r7b2bjq_ped.fits', 'n9r7b2bjq_cal.fits' -d 'same_as_persistentstring.fits'
```

The full description of the parameters required by this class is as follows.

constructor

#### Parameters

**calcfile** : string

name of ped file

**targfile** : string

name of cal file

**persist\_lo** : float

minimum allowed value of the persistence

**used\_lo** : float

minimum allowed value of the fraction of pixels used

**persist\_model** : string

filename containing persistence frame (ring median of)

**persist\_mask** : string

filename containing pixel mask

**run\_stdout** : file handle  
open trailer file (pipeline use only)

**persist** ()  
remove persistence due to the full bright Earth.

**print\_pars** ()  
Print input parameters and calculated values.

`nictools.nic_rem_persist.check_cl_pars` (*calcfile*, *targfile*, *persist\_lo*, *used\_lo*)  
When run from linux command line, verify that each parameter is valid.

#### Parameters

**calcfile** : string  
name of ped file

**targfile** : string  
name of cal file

**persist\_lo** : float  
minimum allowed value of the persistence

**used\_lo** : float  
minimum allowed value of the fraction of pixels used

#### Returns

**persist\_lo**, **used\_lo** : float, float

`nictools.nic_rem_persist.check_py_pars` (*self*, *calcfile*, *targfile*, *persist\_lo*, *used\_lo*, *persist\_model*, *persist\_mask*)

When run under python, check validity of input parameters. For unspecified \*\_lo parameters, the user will be given the option of typing in a value or accepting the default value. For an unspecified model(mask) file, will try to get file name from PMODFILE(PMSKFILE) from input file header, otherwise will get default value from persutil.

#### Parameters

**calcfile** : string  
name of PED file

**targfile** : string  
name of CAL file

**persist\_lo** : float  
minimum allowed value of the persistence

**used\_lo** : float  
minimum allowed value of the fraction of pixels used

**persist\_model** : string  
filename containing persistence frame (ring median of)

**persist\_mask** : string  
filename containing pixel mask

**Returns**

**persist\_lo, used\_lo** : float

**persist\_model, persist\_mask** : string

`nictools.nic_rem_persist.iterstatc(clip, d)`

**version of nicmos iterstat: calculate sigma-clipped mean and standart deviation**

**Parameters**

**clip** : float

number of std to use in sigma clipping

**d** : float

array of values to sigma clip

**Returns**

**clipped mean, clipped std** : float, float

`nictools.nic_rem_persist.make_footprint(rin, rout)`

**Make an annular mask footprint for use in ndimage.median\_filter to create a ring median image.**  
Sets all pixels between *rin* and *rout* to 1.

**Parameters**

**rin** : int

inner radius

**rout** : int

outer radius

**Returns**

**mask** : ndarray

`nictools.nic_rem_persist.median(y, mask)`

Return the median of the array *y*, ignoring masked elements.

**Parameters**

**y** : float

array of values [2d]

**mask** : int

array of ones or zeros (0 indicates a good value) [2d]

**Returns**

**median\_y** : float

median of *y*, ignoring masked elements



## CALTEMPFROMBIAS

```
class nictools.CalTempFromBias.CalTempFromBias (input_file, edit_type=None, hdr_key=None,  
err_key=None, nref_par=None,  
force=None, noclean=False, dry_run=1,  
verbosity=0)
```

Calculate the temperature from the bias for a given filename.

### Notes

Basic syntax for using this class is:

```
tfb = CalTempFromBias( filename, edit_type=edit_type, hdr_key=hdr_key, err_key=err_key,  
    nref_par=nref_par, force=force, noclean=noclean, dry_run=dry_run, verbosity=verbosity)  
[temp, sigma, winner, in_flag, dry_run ]= tfb.calctemp()  
tfb.print_pars()  
stat = tfb.update_header( temp, sigma, winner)
```

The full set of parameters for the methods:

constructor

#### Parameters

**input\_file** : string

name of the file or filelist to be processed

**edit\_type** : string type of file to update

**hdr\_key** : string

name of keyword to update in file

**err\_key** : string

name of keyword for error estimate

**nref\_par** : string

name of the directory containing the nonlinearity file

**force** : string

name of algorithm whose value is to be returned

**noclean** : {'True', 'False'}

flag to force use of UNCLEANed 0th read.

**dry\_run** : {0,1} [Default: 1]

flag to force not writing to header

**verbosity** : {0,1,2}

verbosity level (0 for quiet, 1 verbose, 2 very verbose)

**calctemp** ()

Calculate the temperature from the bias for the given input file

**Returns**

**temp** : float

**sig** : float

**winner** : int

**in\_flag** : str

**print\_pars** ()

Print parameters used.

**update\_header** (*temp, sig, winner, edit\_type=None, hdr\_key=None, err\_key=None*)

Update header method

**Parameters**

**temp** : float

calculated temperature

**sig** : float

standard deviation of calculated temperature

**winner** : int

algorithm used

**edit\_type** : string

type of file to be updated

**hdr\_key** : string

name of keyword to update in file

**err\_key** : string

name of keyword for error estimate

**Returns**

**status** : int (not None for failure due to key not being specified)

`nictools.CalTempFromBias.do_blind` (*camera, quads, verbosity*)

Calculate temperature using the blind correction

**Parameters**

**camera** : int

number of camera

**quads** : float

value of quad from quadmean

**verbosity** : int

level of verbosity

**Returns**

**temp, sig** : float, float



`nictools.CalTempFromBias.do_quietest` (*camera, quads, verbosity*)

Calculate temperature using the quietest quad correction

#### Parameters

**camera** : int  
number of camera

**quads** : float  
value of quad from quadmean

**verbosity** : int  
level of veborsity

#### Returns

**temp, sig** : float

`nictools.CalTempFromBias.poly` (*var\_x, coeffs*)

Return linear polynomial with given coefficients.

#### **var\_x**

[scalar, vector, or array] input argument

#### **coeffs**

[float] vector of polynomial coefficients

`nictools.CalTempFromBias.quadmean` (*im, border*)

This function computes the mean in the 4 quadrants of an input NxM array

#### Parameters

**im** : ndarray  
input rectangular array

**border** : int  
border size (in pixels) around the perimeter of each quad to be excluded from the mean

#### Returns

**quads** : float

#### Notes

Following Eddie's convention, the quads are numbered as follows:

----- -----	
Q4   Q3	
----- -----	as seen in the standard NICMOS/HST data format.
Q1   Q2	
----- -----	

optionally, you can specify a border, in pixels, around the perimeter of EACH QUAD to be excluded from the mean



## FINESKY

**class** `nictools.finesky.Makemedmask` (*thresh=None, medfile=None, callist=None, verbosity=0*)  
Create and output a median mask from cal files and blt files

### Notes

Syntax for using this class:

```
m_mask = finesky.Makemedmask( medfile='medout2.fits', callist='/hal/data2/dev/nicmos_ped/inlist1
    thresh = 0.7, verbosity = 1)
m_mask.makemask()
```

Full set of parameters for class methods are as follows.

constructor

#### Parameters

**thresh** : real

threshold used in making mask

**medfile** : string

name of output masked median image

**callist** : string

name of text file containing cal file names

**verbosity** : {0,1,2}

verbosity level (0 for quiet, 1 verbose, 2 very verbose)

**makemask** ()

Make and output mask

**print\_pars** ()

Print parameters.

`nictools.finesky.write_to_file` (*data, filename, hdr, verbosity*)

Write data to specified filename with specified header

#### Parameters

**data** : ndarray

numpy array

**filename** : string

name of output file

**hdr** : pyfits Header object

header for output file

**verbosity** : {0,1,2}

verbosity level (0 for quiet, 1 verbose, 2 very verbose)

## PUFTCORR

**puftcorr: Module for estimating and removing “Mr. Staypuft” signal from**  
a NICMOS exposure.

### Usage

Normally used via the STSDAS task `puftcorr` in the `nicmos` package. To use as pure python, just invoke the `clean` method:

```
>>> puftcorr.clean('inputfile.fits', 'outputfile.fits')
```

### For more information

Additional user information, including parameter definitions and more examples, can be found in the help file for the STSDAS `puftcorr` task, located in `nicmos$doc/puftcorr.hlp`.

The algorithm and IDL prototype were developed by L.Bergeron, but never made publicly available.

### Dependencies

- numpy v1.0.2dev3534 or higher
- astropy
- scipy

**exception** `nictools.puftcorr.NoPuftError`

**class** `nictools.puftcorr.InputFile` (*imgfile*)

Stores a collection of keywords and the header for an exposure.

**class** `nictools.puftcorr.Readout` (*input, sampnum*)

**class** `nictools.puftcorr.params` (*camera*)

`nictools.puftcorr.clean` (*usr\_imgfile, usr\_outfile*)

`nictools.puftcorr.get_corr` (*im, pars*)

`nictools.puftcorr.get_totSIG` (*im, la*)

`nictools.puftcorr.osfn` (*filename*)

Return a filename with iraf syntax and os environment names substituted out



## RNLINCORR

**rnlincor: Module to correct for the countrate-dependent nonlinearity in a NICMOS image.**

### Usage

Normally used via the STSDAS task rnlincor in the nicmos package. To use as pure python, just invoke the run method:

```
>>> rnlincor.run('inputfile.fits', 'outputfile.fits')
```

It may also be run from the shell:

```
% rnlincor.py infile.fits [outfile.fits] [--nozpcorr]
```

### For more information

Additional user information, including parameter definitions and more examples, can be found in the help file for the STSDAS rnlincor task, located in `nicmos$doc/rnlincor.hlp`.

This task is based on prototype code developed by R. de Jong. The algorithm is described in more detail in ISR NICMOS 2006-003 by de Jong.

### Dependencies

- numpy v1.0.2dev3534 or higher
- pyfits v1.1b4 or higher

**class** `nictools.rnlincor.FitsRowObject` (*fitsrecord*)

Class to facilitate working with single table rows.

`nictools.rnlincor.check_infile` (*infile*)

Open the input file and check all the things that can go wrong. If we pass all the tests, return the handle to the open file to pass to the main routine.

`nictools.rnlincor.expandname` (*pattern*)

Select the latest file that matches the pattern in the directory specified in the pattern

`nictools.rnlincor.getcurve` (*fname*, *col1='wavelength'*, *col2='correction'*, *pad=0*)

Gets a 2-column table from the first extension of a FITS file. Defaults to “wavelength” and “correction” for column names, but others can be specified. If pad keyword is nonzero, the wavelength table will be extended by the pad amount in each direction.

`nictools.rnlincor.getrow` (*photmode*, *corrname*)

Pick out the correction parameters from the proper row of the table located in `corrname`, based on `photmode`. Return an object that contains the row fields as attributes.

`nictools.rnlincor.parrun` (*parfile*)

`nictools.rnlincor.rnlincor` (*infile, outfile, \*\*opt*)

The main routine

`nictools.rnlincor.run` (*\*args, \*\*inopt*)

`nictools.rnlincor.set_default_options` (*inopt*)

`nictools.rnlincor.update_data` (*f, imgext, img, mul*)

`nictools.rnlincor.update_header` (*f, alpha, zpcorr, zpratio=None*)

Update all the header keywords



## READTDD

The `readTDD.py` is a helper module used to extract the linear dark and amp glow components from a NICMOS time dependent dark file.

**author**

Christopher Hanley

**dependencies**

`stsci.tools.fileutil`

**class** `nictools.readTDD.darkobject` (*hdulist*)

`darkobject`: This class takes as input a `pyfits hdulist` object. The linear dark and amp glow noise components are then extracted from the `hdulist`.

**getampglow** ()

`getampglow`: `darkobject` method which is used to return the amp glow component from a NICMOS temperature dependent dark file.

**getampglowheader** ()

`getampglowheader`: `darkobject` method used to return the header information of the amp glow extension of a TDD file.

**getlindark** ()

`getlindark`: `darkobject` method which is used to return the linear dark component from a NICMOS temperature dependent dark file.

**getlindarkheader** ()

`getlindarkheader`: `darkobject` method used to return the header information of the linear dark extension of a TDD file.

`nictools.readTDD.fromcalfile` (*filename*)

`fromcalfile`: function that returns a `darkobject` instance given the name of a `cal.fits` file as input. If there is no `TEMPFILE` keyword in the primary header of the `cal.fits` file or if the file specified by `TEMPFILE` cannot be found, a `None` object is returned.



## MAKEMEDMASK

**class** `nictools.makemedmask.Makemedmask` (*thresh=None, medfile=None, callist=None, verbosity=0*)

Create and output a median mask from cal files and blt files

### Notes

Syntax for using this class:

```
m_mask = makemedmask.Makemedmask( medfile='medout2.fits', callist='/hal/data2/dev/nicmos_ped/inl
    thresh = 0.7, verbosity = 1)
m_mask.makemask()
```

Full signatures for methods are as follows:

constructor

#### Parameters

**thresh** : float

threshold used in making mask

**medfile** : string

name of output masked median image

**callist** : string

name of text file containing cal file names

**verbosity** : {0,1,2}

verbosity level (0 for quiet, 1 verbose, 2 very verbose)

**makemask** ()

Make and output mask

**print\_pars** ()

Print parameters.

`nictools.makemedmask.write_to_file` (*data, filename, hdr, verbosity*)

Write data to specified filename with specified header

#### Parameters

**data** : ndarray

numpy array

**filename** : string

name of output file

**hdr** : pyfits Header object

header for output file

**verbosity** : {0,1,2}

verbosity level (0 for quiet, 1 verbose, 2 very verbose)

## UTILITIES

There are a number of modules which provide utility functions for use in this package.

### 9.1 fsutil

`nictools.fsutil.all_printMsg` (*message*, *level=1*)

**Parameters**

**message** : string  
message to print  
**level** : int  
verbosity level

`nictools.fsutil.checkVerbosity` (*level*)

**Parameters**

**level** : int  
level of verbosity

**Returns**

**level** : bool  
true if verbosity is at least as great as level.

`nictools.fsutil.printMsg` (*message*, *level=0*)

**Parameters**

**message** : string  
message to print  
**level** : int  
verbosity level

`nictools.fsutil.setCallist` (*callist\_value*)

Copy callist to a variable that is global for this file.

**Parameters**

**callist\_value** : string  
name of file listing cal files

`nictools.fsutil.setMedfile (medfile_value)`

Copy medfile to a variable that is global for this file.

**Parameters**

**medfile\_value** : string

name of output file for masked median image

`nictools.fsutil.setThresh (thresh_value)`

Copy thresh to a variable that is global for this file.

**Parameters**

**thresh\_value** : float

level of threshold

`nictools.fsutil.setVerbosity (verbosity_level)`

Copy verbosity to a variable that is global for this file.

**Parameters**

**verbosity\_level** : int

level of verbosity

## 9.2 opusutil

`nictools.opusutil.CloseTrl ()`

`nictools.opusutil.FileToList (filename)`

`nictools.opusutil.OpenTrl (filespec)`

`nictools.opusutil.PrintMsg (level, msg, module_name='')`

`nictools.opusutil.RemoveIfThere (filename)`

`nictools.opusutil.ResourceToMap (filename)`

`nictools.opusutil.StretchFile (stretched_filename)`

`nictools.opusutil.UsingLvl (level)`

## 9.3 persutil

`nictools.persutil.all_printMsg (message, level=1)`

`nictools.persutil.checkVerbosity (level)`

Return true if verbosity is at least as great as level.

`nictools.persutil.getOptions()`

`nictools.persutil.getPersist_lo(calcfile)`

Get value of `persist_lo` from BEPVALLO in the persistence model file PMODFILE or from a default specified by this module. This will only be called if user did not specify a value on the command-line.

**Parameters**

**calcfile** : string

input ped file

**Returns**

**persist\_lo** : float

`nictools.persutil.getPersist_mask(calcfile)`

Get name of persistence mask from PMSKFILE in the input file This will only be called if user did not specify a model on the command-line.

**Parameters**

**calcfile** : string

input ped file

**Returns**

**persist\_mask** : string

`nictools.persutil.getPersist_model(calcfile)`

Get name of persistence model from PMODFILE in the input file. This will only be called if user did not specify a model on the command-line.

**Parameters**

**calcfile** : string

input ped file

**Returns**

**persist\_model** : string

`nictools.persutil.getUsed_lo(calcfile)`

Get value of `used_lo` from BEPUSELO in the persistence model file PMODFILE or from a default specified by this module. This will only be called if user did not specify a value on the command-line.

**Parameters**

**calcfile** : string

input ped file

**Returns**

**used\_lo** : float

`nictools.persutil.printMsg(message, level=0)`

`nictools.persutil.setVerbosity(verbosity_level)`

Copy verbosity to a variable that is global for this file. argument: `verbosity_level` - an integer value indicating the level of verbosity

## 9.4 SP\_FirstDerivatives

Automatic differentiation for functions with any number of variables

Instances of the class `DerivVar` represent the values of a function and its partial X{derivatives} with respect to a list of variables. All common mathematical operations and functions are available for these numbers. There is no restriction on the type of the numbers fed into the code; it works for real and complex numbers as well as for any Python type that implements the necessary operations.

This module is as far as possible compatible with the n-th order derivatives module `Derivatives`. If only first-order derivatives are required, this module is faster than the general one.

Example:

```
print sin(DerivVar(2))
```

produces the output:

```
(0.909297426826, [-0.416146836547])
```

The first number is the value of  $\sin(2)$ ; the number in the following list is the value of the derivative of  $\sin(x)$  at  $x=2$ , i.e.  $\cos(2)$ .

When there is more than one variable, `DerivVar` must be called with an integer second argument that specifies the number of the variable.

Example:

```
>>>x = DerivVar(7., 0)
>>>y = DerivVar(42., 1)
>>>z = DerivVar(pi, 2)
>>>print (sqrt(pow(x,2)+pow(y,2)+pow(z,2)))
```

produces the output

```
>>>(42.6950770511, [0.163953328662, 0.98371997197, 0.0735820818365])
```

The numbers in the list are the partial derivatives with respect to  $x$ ,  $y$ , and  $z$ , respectively.

Note: It doesn't make sense to use `DerivVar` with different values for the same variable index in one calculation, but there is no check for this. I.e.:

```
>>>print DerivVar(3, 0)+DerivVar(5, 0)
```

produces

```
>>>(8, [2])
```

but this result is meaningless.

**class** `nictools.SP_FirstDerivatives.DerivVar` (*value*, *index=0*, *order=1*)  
Numerical variable with automatic derivatives of first order

**Parameters**

**value** : int or float

the numerical value of the variable

**index** : int

the variable index, which serves to distinguish between variables and as an index for the derivative lists. Each explicitly created instance of `DerivVar` must have a unique index.

**order** : int

the derivative order, must be zero or one



**Raises**

---

**ValueError: if order < 0 or order > 1****arccos** ()**arcsin** ()**arctan** ()**arctan2** (*other*)**cos** ()**cosh** ()**exp** ()**gamma** ()**log** ()**log10** ()**sign** ()**sin** ()**sinh** ()**sqrt** ()**tan** ()**tanh** ()`nictools.SP_FirstDerivatives.DerivVector` (*x*, *y*, *z*, *index=0*)**Parameters****x** : float or int

x component of the vector

**y** : float or int

y component of the vector

**z** : float or int

z component of the vector

**index** : int

the DerivVar index for the x component. The y and z components receive consecutive indices.

**Returns**

**vector** : Scientific.Geometry.VectorModule.Vector

a vector whose components are DerivVar objects

nictools.SP\_FirstDerivatives.**isDerivVar**(x)

**Parameters**

**x** :

an arbitrary object

**Returns**

**result** : bool

True if x is a DerivVar object, False otherwise

## 9.5 SP\_LeastSquares

Non-linear least squares fitting

### 9.5.1 Examples

Usage example:

```
from Scientific.N import exp

def f(param, t):
    return param[0]*exp(-param[1]/t)

data_quantum = [(100, 3.445e+6), (200, 2.744e+7),
                (300, 2.592e+8), (400, 1.600e+9)]
data_classical = [(100, 4.999e-8), (200, 5.307e+2),
                  (300, 1.289e+6), (400, 6.559e+7)]

print leastSquaresFit(f, (1e13,4700), data_classical)

def f2(param, t):
    return 1e13*exp(-param[0]/t)

print leastSquaresFit(f2, (3000.,), data_quantum)
```

**exception** nictools.SP\_LeastSquares.**IterationCountExceededError**

nictools.SP\_LeastSquares.**leastSquaresFit**(*model*, *parameters*, *data*, *max\_iterations=None*,  
*stopping\_limit=0.005*)

General non-linear least-squares fit using the X{Levenberg-Marquardt} algorithm and X{automatic differentiation}.

**model**

[function] the function to be fitted. It will be called with two parameters: the first is a tuple containing all fit parameters, and the second is the first element of a data point (see below). The return value must be a number. Since automatic differentiation is used to obtain the derivatives with respect to the parameters, the function may only use the mathematical functions known to the module FirstDerivatives.

**parameters**

[tuple of numbers] a tuple of initial values for the fit parameters

**data**

[list] a list of data points to which the model is to be fitted. Each data point is a tuple of length two or three. Its first element specifies the independent variables of the model. It is passed to the model function as its first parameter, but not used in any other way. The second element of each data point tuple is the number that the return value of the model function is supposed to match as well as possible. The third element (which defaults to 1.) is the statistical variance of the data point, i.e. the inverse of its statistical weight in the fitting procedure.

**Returns**

**fitlist** : list

a list containing the optimal parameter values

**chisq** : float

chi-squared value describing the quality of the fit

`nictools.SP_LeastSquares.polynomialLeastSquaresFit` (*parameters*, *data*)  
Least-squares fit to a polynomial whose order is defined by the number of parameter values.

---

**Note:** This could also be done with a linear least squares fit from `L{LinearAlgebra}`

---

**Parameters**

**parameters** : tuple

a tuple of initial values for the polynomial coefficients

**data** : list

the data points, as for `L{leastSquaresFit}`

## 9.6 tfbutil

`nictools.tfbutil.all_printMsg` (*message*, *level=1*)

`nictools.tfbutil.checkVerbosity` (*level*)

Return true if verbosity is at least as great as level.

`nictools.tfbutil.printMsg` (*message*, *level=0*)

`nictools.tfbutil.setDry_run` (*dry\_run\_value*)

Copy `dry_run` to a variable that is global for this file.

**Parameters**

**dry\_run** : string

string that is either True or False

`nictools.tfbutil.setEdit_type_key` (*edit\_type\_value*)

Copy `edit_type_key` to a variable that is global for this file.

**Parameters**

**edit\_type\_key** : string

a string for the keyword name to write

`nictools.tfbutil.setErr_key` (*err\_key\_value*)

Copy `err_key` to a variable that is global for this file.

**Parameters**

**err\_key** : string

a string for the keyword for the error estimate

`nictools.tfbutil.setForce` (*force\_value*)

Copy `force` to a variable that is global for this file.

**Parameters**

**force** : string

string that is either None, Q, B, or A

`nictools.tfbutil.setHdr_key` (*hdr\_key\_value*)

Copy `hdr_key` to a variable that is global for this file.

**Parameters**

**hdr\_key** : string

a string for the keyword name to write

`nictools.tfbutil.setNoclean` (*noclean\_value*)

Copy `no_clean` to a variable that is global for this file.

**Parameters**

**no\_clean** : string

string that is either True or False

`nictools.tfbutil.setNref` (*nref\_value*)

Copy `nref` to a variable that is global for this file.

**Parameters**

**nref** : string

string for name of directory containing nonlinearity file

`nictools.tfbutil.setVerbosity` (*verbosity\_level*)

Copy `verbosity` to a variable that is global for this file.

**Parameters**

**verbosity\_level** **L** int

an integer value indicating the level of verbosity

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