

---

# healpy Documentation

*Release 1.12.9*

March 21, 2019



---

## Contents

---

<b>1 Tutorial</b>	<b>3</b>
<b>2 Installation</b>	<b>7</b>
<b>3 Reference</b>	<b>11</b>
<b>4 Indices and tables</b>	<b>75</b>



*healpy* is a Python package to handle pixelated data on the sphere. It is based on the [Hierarchical Equal Area isoLatitude Pixelization \(HEALPix\)](#) scheme and bundles the *HEALPix* C++ library.

*HEALPix* was developed to efficiently process Cosmic Microwave Background data from Cosmology experiments like BOOMERANG and WMAP but it is now used in other branches of Astrophysics to store data from all-sky surveys. The target audience used to be primarily the Cosmology scientific community but currently anyone interested in handling pixelated data on the sphere is very welcome to propose new features.

*healpy* provides utilities to:

- convert between sky coordinates and pixel indices in HEALPix nested and ring schemes
- find pixels within a disk, a polygon or a strip in the sky
- apply coordinate transformations between Galactic, Ecliptic and Equatorial reference frames
- apply custom rotations either to vectors or full maps
- read and write HEALPix maps to disk in FITS format
- upgrade and downgrade the resolution of existing HEALPix maps
- visualize maps in Mollweide, Gnomonic and Cartographic projections
- transform maps to Spherical Harmonics space and back using multi-threaded C++ routines
- compute Auto and Cross Power Spectra from maps and create map realizations from spectra



# CHAPTER 1

---

## Tutorial

---

### 1.1 healpy tutorial

See the Jupyter Notebook version of this tutorial at [https://github.com/healpy/healpy/blob/master/doc/healpy\\_tutorial.ipynb](https://github.com/healpy/healpy/blob/master/doc/healpy_tutorial.ipynb)

See a executed version of the notebook with embedded plots at <https://gist.github.com/zonca/9c14608e0903a3b8ea0bfe41c96f255>

Choose the `inline` backend of `matplotlib` to display the plots inside the Jupyter Notebook

```
import matplotlib.pyplot as plt  
%matplotlib inline
```

```
import numpy as np  
import healpy as hp
```

#### 1.1.1 NSIDE and ordering

Maps are simply numpy arrays, where each array element refers to a location in the sky as defined by the Healpix pixelization schemes (see the [healpix website](#)).

Note: Running the code-block below in a regular Python session will not display the maps; it's recommended to use an IPython shell or a Jupyter notebook.

The resolution of the map is defined by the `NSIDE` parameter, which is generally a power of 2.

```
NSIDE = 32  
print(  
    "Approximate resolution at NSIDE {} is {:.2} deg".format(  
        NSIDE, hp.nside2resol(NSIDE, arcmin=True) / 60  
    )  
)
```

The function `healpy.pixelfunc.nside2npix` gives the number of pixels  $NPIX$  of the map:

```
NPIX = hp.nside2npix(NSIDE)
print(NPIX)
```

The same pixels in the map can be ordered in 2 ways, either RING, where they are numbered in the array in horizontal rings starting from the North pole:

```
m = np.arange(NPIX)
hp.mollview(m, title="Mollview image RING")
hp.graticule()
```

The standard coordinates are the **colatitude**  $\theta$ , 0 at the North Pole,  $\pi/2$  at the equator and  $\pi$  at the South Pole and the **longitude**  $\phi$  between 0 and  $2\pi$  eastward, in a Mollview projection,  $\phi = 0$  is at the center and increases eastward toward the left of the map.

We can also use vectors to represent coordinates, for example `vec` is the normalized vector that points to  $\theta = \pi/2, \phi = 3/4\pi$ :

```
vec = hp.ang2vec(np.pi / 2, np.pi * 3 / 4)
print(vec)
```

We can find the indices of all the pixels within 10 degrees of that point and then change the value of the map at those indices:

```
ipix_disc = hp.query_disc(nside=32, vec=vec, radius=np.radians(10))
```

```
m = np.arange(NPIX)
m[ipix_disc] = m.max()
hp.mollview(m, title="Mollview image RING")
```

We can retrieve colatitude and longitude of each pixel using `pix2ang`, in this case we notice that the first 4 pixels cover the North Pole with pixel centers just  $\sim 1.5$  degrees South of the Pole all at the same latitude. The fifth pixel is already part of another ring of pixels.

```
theta, phi = np.degrees(hp.pix2ang(nside=32, ipix=[0, 1, 2, 3, 4]))
```

```
theta
```

```
phi
```

The RING ordering is necessary for the Spherical Harmonics transforms, the other option is NESTED ordering which is very efficient for map domain operations because scaling up and down maps is achieved just multiplying and rounding pixel indices. See below how pixel are ordered in the NESTED scheme, notice the structure of the 12 HEALPix base pixels (NSIDE 1):

```
hp.mollview(m, nest=True, title="Mollview image NESTED")
```

All `healpy` routines assume RING ordering, in fact as soon as you read a map with `read_map`, even if it was stored as NESTED, it is transformed to RING. However, you can work in NESTED ordering passing the `nest=True` argument to most `healpy` routines.

### 1.1.2 Reading and writing maps to file

For the following section, it is required to download larger maps by executing from the terminal the bash script `healpy_get_wmap_maps.sh` which should be available in your path.

This will download the higher resolution WMAP data into the current directory.

```
!healpy_get_wmap_maps.sh

wmap_map_I = hp.read_map("wmap_band_iqumap_r9_7yr_W_v4.fits")
```

By default, input maps are converted to *RING* ordering, if they are in *NESTED* ordering. You can otherwise specify `nest=True` to retrieve a map in *NESTED* ordering, or `nest=None` to keep the ordering unchanged.

By default, `read_map` loads the first column, for reading other columns you can specify the `field` keyword.

`write_map` writes a map to disk in FITS format, if the input map is a list of 3 maps, they are written to a single file as I,Q,U polarization components:

```
hp.write_map("my_map.fits", wmap_map_I, overwrite=True)
```

### 1.1.3 Visualization

As shown above, mollweide projection with `mollview` is the most common visualization tool for HEALPIX maps. It also supports coordinate transformation, `coord` does Galactic to ecliptic coordinate transformation, `norm='hist'` sets a histogram equalized color scale and `xsize` increases the size of the image. `graticule` adds meridians and parallels.

```
hp.mollview(
    wmap_map_I,
    coord=["G", "E"],
    title="Histogram equalized Ecliptic",
    unit="mK",
    norm="hist",
    min=-1,
    max=1,
)
hp.graticule()
```

`gnomview` instead provides gnomonic projection around a position specified by `rot`, for example you can plot a projection of the galactic center, `xsize` and `ysize` change the dimension of the sky patch.

```
hp.gnomview(wmap_map_I, rot=[0, 0.3], title="GnomView", unit="mK", format=".2g")
```

`mollzoom` is a powerful tool for interactive inspection of a map, it provides a mollweide projection where you can click to set the center of the adjacent gnomview panel. ## Masked map, partial maps

By convention, HEALPIX uses  $-1.6375 * 10^{30}$  to mark invalid or unseen pixels. This is stored in `healpy` as the constant `UNSEEN`.

All `healpy` functions automatically deal with maps with `UNSEEN` pixels, for example `mollview` marks in grey those sections of a map.

There is an alternative way of dealing with `UNSEEN` pixel based on the `numpyMaskedArray` class, `hp.ma` loads a map as a masked array, by convention the mask is 0 where the data are masked, while `numpy` defines data masked when the mask is True, so it is necessary to flip the mask.

```
mask = hp.read_map("wmap_temperature_analysis_mask_r9_7yr_v4.fits").astype(np.bool)
wmap_map_I_masked = hp.ma(wmap_map_I)
wmap_map_I_masked.mask = np.logical_not(mask)
```

Filling a masked array fills in the UNSEEN value and return a standard array that can be used by `mollview`. `compressed()` instead removes all the masked pixels and returns a standard array that can be used for examples by the `matplotlib hist()` function:

```
hp.mollview(wmap_map_I_masked.filled())
```

```
plt.hist(wmap_map_I_masked.compressed(), bins=1000)
```

### 1.1.4 Spherical Harmonics transforms

healpy provides bindings to the C++ HEALPIX library for performing spherical harmonic transforms. `hp.anafast` computes the angular power spectrum of a map:

```
LMAX = 1024
cl = hp.anafast(wmap_map_I_masked.filled(), lmax=LMAX)
ell = np.arange(len(cl))
```

therefore we can plot a normalized CMB spectrum and write it to disk:

```
plt.figure(figsize=(10, 5))
plt.plot(ell, ell * (ell + 1) * cl)
plt.xlabel("$\ell$")
plt.ylabel("$\ell(\ell+1)C_\ell$")
plt.grid()
hp.write_cl("cl.fits", cl, overwrite=True)
```

Gaussian beam map smoothing is provided by `hp.smoothing`:

```
wmap_map_I_smoothed = hp.smoothing(wmap_map_I, fwhm=np.radians(1.))
hp.mollview(wmap_map_I_smoothed, min=-1, max=1, title="Map smoothed 1 deg")
```

For more information see the [HEALPix primer](#)

# CHAPTER 2

---

## Installation

---

### 2.1 Installation procedure for Healpy

#### 2.1.1 Requirements

Healpy depends on the HEALPix C++ and cfitsio C libraries. Source code for both is included with Healpy and is built automatically, so you do not need to install them yourself. Only Linux and MAC OS X are supported, not Windows.

#### 2.1.2 Binary installation with conda (RECOMMENDED)

Conda forge provides a [conda channel](#) with a pre-compiled version of healpy for linux 64bit and MAC OS X platforms, you can install it in Anaconda with:

```
conda config --add channels conda-forge  
conda install healpy
```

#### 2.1.3 Source installation with Pip

It is possible to build the latest healpy with pip

```
pip install --user healpy
```

If you have installed with pip, you can keep your installation up to date by upgrading from time to time:

```
pip install --user --upgrade healpy
```

On Linux with newer compilers many users reported compilation errors like `configure: error: cannot run C compiled programs`, the solution is to specify the flags for the C and CXX compiler:

```
CC=gcc CXX=g++ CFLAGS='-fPIC' CXXFLAGS='-fPIC' pip install --user healpy
```

## 2.1.4 Installation on Mac OS with MacPorts

If you are using a Mac and have the [MacPorts](#) package manager, it's even easier to install Healpy with:

```
sudo port install py27-healpy
```

Binary *apt-get* style packages are also available in the development versions of [Debian \(sid\)](#) and [Ubuntu \(utopic\)](#).

## 2.1.5 Almost-as-quick installation from official source release

Healpy is also available in the [Python Package Index \(PyPI\)](#). You can download it with:

```
curl -O https://pypi.python.org/packages/source/h/healpy/healpy-1.7.4.tar.gz
```

and build it with:

```
tar -xzf healpy-1.7.4.tar.gz  
pushd healpy-1.7.4  
python setup.py install --user  
popd
```

If everything goes fine, you can test it:

```
python
```

```
>>> import matplotlib.pyplot as plt  
>>> import numpy as np  
>>> import healpy as hp  
>>> hp.mollview(np.arange(12))  
>>> plt.show()
```

or run the test suite with nose:

```
cd healpy-1.7.4 && python setup.py test
```

## 2.1.6 Building against external Healpix and cfitsio

Healpy uses `pkg-config` to detect the presence of the Healpix and cfitsio libraries. `pkg-config` is available on most systems. If you do not have `pkg-config` installed, then Healpy will download and use (but not install) a Python clone called `pykg-config`.

If you want to provide your own external builds of Healpix and cfitsio, then download the following packages:

- [pkg-config](#)
- [HEALPix autotools-style C++ package](#)
- [cfitsio](#)

If you are going to install the packages in a nonstandard location (say, `--prefix=/path/to/local`), then you should set the environment variable `PKG_CONFIG_PATH=/path/to/local/lib/pkgconfig` when building. No other environment variable settings are necessary, and you do not need to set `PKG_CONFIG_PATH` to use Healpy after you have built it.

Then, unpack each of the above packages and build them with the usual `configure; make; make install` recipe.

## 2.1.7 Development install

Developers building from a snapshot of the github repository need:

- autoconf and libtool (in Debian or Ubuntu: `sudo apt-get install autoconf automake libtool pkg-config`)
- *cython > 0.16*
- run `git submodule init` and `git submodule update` to get the bundled HEALPix sources

the best way to install healpy if you plan to develop is to build the C++ extensions in place with:

```
python setup.py build_ext --inplace
```

then add the `healpy/healpy` folder to your `PYTHONPATH`.

In case of compilation errors, see the note above in the `pip` section.

## 2.1.8 Clean

When you run “`python setup.py`”, temporary build products are placed in the “build” directory. If you want to clean out and remove the `build` directory, then run:

```
python setup.py clean --all
```



# CHAPTER 3

---

## Reference

---

### 3.1 pixelfunc – Pixelisation related functions

#### 3.1.1 conversion from/to sky coordinates

<code>pix2ang(nside, ipix[, nest, lonlat])</code>	<code>pix2ang : nside,ipix,nest=False,lonlat=False -&gt; theta[rad],phi[rad] (default RING)</code>
<code>pix2vec(nside, ipix[, nest])</code>	<code>pix2vec : nside,ipix,nest=False -&gt; x,y,z (default RING)</code>
<code>ang2pix(nside, theta, phi[, nest, lonlat])</code>	<code>ang2pix : nside,theta[rad],phi[rad],nest=False,lonlat=False -&gt; ipix (default:RING)</code>
<code>vec2pix(nside, x, y, z[, nest])</code>	<code>vec2pix : nside,x,y,z,nest=False -&gt; ipix (default:RING)</code>
<code>vec2ang(vectors[, lonlat])</code>	<code>vec2ang: vectors [x, y, z] -&gt; theta[rad], phi[rad]</code>
<code>ang2vec(theta, phi[, lonlat])</code>	<code>ang2vec : convert angles to 3D position vector</code>
<code>get_all_neighbours(nside, theta[, phi, ...])</code>	<code>Return the 8 nearest pixels.</code>
<code>get_interp_weights(nside, theta[, phi, ...])</code>	<code>Return the 4 closest pixels on the two rings above and below the location and corresponding weights.</code>
<code>get_interp_val(m, theta, phi[, nest, lonlat])</code>	<code>Return the bi-linear interpolation value of a map using 4 nearest neighbours.</code>

#### healpy.pixelfunc.pix2ang

```
healpy.pixelfunc.pix2ang(nside, ipix, nest=False, lonlat=False)
pix2ang : nside,ipix,nest=False,lonlat=False -> theta[rad],phi[rad] (default RING)
```

##### Parameters

**nside** [int or array-like] The healpix nside parameter, must be a power of 2, less than  $2^{**}30$

**ipix** [int or array-like] Pixel indices

**nest** [bool, optional] if True, assume NESTED pixel ordering, otherwise, RING pixel ordering

**lonlat** [bool, optional] If True, return angles will be longitude and latitude in degree, otherwise, angles will be co-latitude and longitude in radians (default)

#### Returns

**theta, phi** [float, scalar or array-like] The angular coordinates corresponding to ipix. Scalar if all input are scalar, array otherwise. Usual numpy broadcasting rules apply.

See also:

[ang2pix](#), [vec2pix](#), [pix2vec](#)

#### Examples

```
>>> import healpy as hp
>>> hp.pix2ang(16, 1440)
(1.5291175943723188, 0.0)
```

```
>>> hp.pix2ang(16, [1440, 427, 1520, 0, 3068])
(array([ 1.52911759,  0.78550497,  1.57079633,  0.05103658,  3.09055608]),
 array([ 0.,  0.78539816,  1.61988371,  0.78539816,  0.78539816]))
```

```
>>> hp.pix2ang([1, 2, 4, 8], 11)
(array([ 2.30052398,  0.84106867,  0.41113786,  0.2044802 ]), array([ 5.49778714,
 5.89048623,  5.89048623,  5.89048623]))
```

```
>>> hp.pix2ang([1, 2, 4, 8], 11, lonlat=True)
(array([ 315.,  337.5,  337.5,  337.5]), array([-41.8103149,  41.8103149,  66.
-44353569,  78.28414761]))
```

## healpy.pixelfunc.pix2vec

healpy.pixelfunc.**pix2vec**(nside, ipix, nest=False)  
pix2vec : nside,ipix,nest=False -> x,y,z (default RING)

#### Parameters

**nside** [int, scalar or array-like] The healpix nside parameter, must be a power of 2, less than  $2^{30}$

**ipix** [int, scalar or array-like] Healpix pixel number

**nest** [bool, optional] if True, assume NESTED pixel ordering, otherwise, RING pixel ordering

#### Returns

**x, y, z** [floats, scalar or array-like] The coordinates of vector corresponding to input pixels. Scalar if all input are scalar, array otherwise. Usual numpy broadcasting rules apply.

See also:

[ang2pix](#), [pix2ang](#), [vec2pix](#)

#### Examples

```
>>> import healpy as hp
>>> hp.pix2vec(16, 1504)
(0.99879545620517241, 0.049067674327418015, 0.0)
```

```
>>> hp.pix2vec(16, [1440, 427])
(array([ 0.99913157,  0.5000534]), array([ 0.          ,  0.5000534]), array([ 0.
-04166667,  0.70703125]))
```

```
>>> hp.pix2vec([1, 2], 11)
(array([ 0.52704628,  0.68861915]), array([-0.52704628, -0.28523539]), array([-0.
-66666667,  0.66666667]))
```

## healpy.pixelfunc.ang2pix

`healpy.pixelfunc.ang2pix(nside, theta, phi, nest=False, lonlat=False)`  
`ang2pix : nside,theta[rad],phi[rad],nest=False,lonlat=False -> ipix (default:RING)`

### Parameters

**nside** [int, scalar or array-like] The healpix nside parameter, must be a power of 2, less than  $2^{**30}$

**theta, phi** [float, scalars or array-like] Angular coordinates of a point on the sphere

**nest** [bool, optional] if True, assume NESTED pixel ordering, otherwise, RING pixel ordering

**lonlat** [bool] If True, input angles are assumed to be longitude and latitude in degree, otherwise, they are co-latitude and longitude in radians.

### Returns

**pix** [int or array of int] The healpix pixel numbers. Scalar if all input are scalar, array otherwise. Usual numpy broadcasting rules apply.

### See also:

`pix2ang`, `pix2vec`, `vec2pix`

## Examples

Note that some of the test inputs below that are on pixel boundaries such as  $\theta=\pi/2$ ,  $\phi=\pi/2$ , have a tiny value of  $1e-15$  added to them to make them reproducible on i386 machines using x87 floating point instruction set (see <https://github.com/healpy/healpy/issues/528>).

```
>>> import healpy as hp
>>> hp.ang2pix(16, np.pi/2, 0)
1440
```

```
>>> print(hp.ang2pix(16, [np.pi/2, np.pi/4, np.pi/2, 0, np.pi], [0., np.pi/4, np.
->pi/2 + 1e-15, 0, 0]))
[1440 427 1520 0 3068]
```

```
>>> print(hp.ang2pix(16, np.pi/2, [0, np.pi/2 + 1e-15]))
[1440 1520]
```

```
>>> print(hp.ang2pix([1, 2, 4, 8, 16], np.pi/2, 0))
[ 4 12 72 336 1440]
```

```
>>> print(hp.ang2pix([1, 2, 4, 8, 16], 0, 0, lonlat=True))
[ 4 12 72 336 1440]
```

## healpy.pixelfunc.vec2pix

healpy.pixelfunc.**vec2pix**(*nside*, *x*, *y*, *z*, *nest=False*)  
vec2pix : nside,x,y,z,nest=False -> ipix (default:RING)

### Parameters

**nside** [int or array-like] The healpix nside parameter, must be a power of 2, less than  $2^{**30}$

**x,y,z** [floats or array-like] vector coordinates defining point on the sphere

**nest** [bool, optional] if True, assume NESTED pixel ordering, otherwise, RING pixel ordering

### Returns

**ipix** [int, scalar or array-like] The healpix pixel number corresponding to input vector. Scalar if all input are scalar, array otherwise. Usual numpy broadcasting rules apply.

### See also:

*ang2pix*, *pix2ang*, *pix2vec*

## Examples

```
>>> import healpy as hp
>>> hp.vec2pix(16, 1, 0, 0)
1504
```

```
>>> print(hp.vec2pix(16, [1, 0], [0, 1], [0, 0]))
[1504 1520]
```

```
>>> print(hp.vec2pix([1, 2, 4, 8], 1, 0, 0))
[ 4 20 88 368]
```

## healpy.pixelfunc.vec2ang

healpy.pixelfunc.**vec2ang**(*vectors*, *lonlat=False*)  
vec2ang: vectors [x, y, z] -> theta[rad], phi[rad]

### Parameters

**vectors** [float, array-like] the vector(s) to convert, shape is (3,) or (N, 3)

**lonlat** [bool, optional] If True, return angles will be longitude and latitude in degree, otherwise, angles will be co-latitude and longitude in radians (default)

### Returns

**theta, phi** [float, tuple of two arrays] the colatitude and longitude in radians

**See also:**

`ang2vec`, `rotator.vec2dir`, `rotator.dir2vec`

## healpy.pixelfunc.ang2vec

`healpy.pixelfunc.ang2vec(theta, phi, lonlat=False)`  
`ang2vec` : convert angles to 3D position vector

### Parameters

**theta** [float, scalar or arry-like] colatitude in radians measured southward from north pole (in [0,pi]).  
**phi** [float, scalar or array-like] longitude in radians measured eastward (in [0, 2\*pi]).  
**lonlat** [bool] If True, input angles are assumed to be longitude and latitude in degree, otherwise, they are co-latitude and longitude in radians.

### Returns

**vec** [float, array] if theta and phi are vectors, the result is a 2D array with a vector per row otherwise, it is a 1D array of shape (3,)

**See also:**

`vec2ang`, `rotator.dir2vec`, `rotator.vec2dir`

## healpy.pixelfunc.get\_all\_neighbours

`healpy.pixelfunc.get_all_neighbours(nside, theta, phi=None, nest=False, lonlat=False)`  
 Return the 8 nearest pixels.

### Parameters

**nside** [int] the nside to work with  
**theta, phi** [scalar or array-like] if phi is not given or None, theta is interpreted as pixel number, otherwise, theta[rad],phi[rad] are angular coordinates  
**nest** [bool] if True, pixel number will be NESTED ordering, otherwise RING ordering.  
**lonlat** [bool] If True, input angles are assumed to be longitude and latitude in degree, otherwise, they are co-latitude and longitude in radians.

### Returns

**ipix** [int, array] pixel number of the SW, W, NW, N, NE, E, SE and S neighbours, shape is (8,) if input is scalar, otherwise shape is (8, N) if input is of length N. If a neighbor does not exist (it can be the case for W, N, E and S) the corresponding pixel number will be -1.

**See also:**

`get_interp_weights`, `get_interp_val`

## Examples

```
>>> import healpy as hp
>>> print(hp.get_all_neighbours(1, 4))
[11  7  3 -1  0  5  8 -1]
```

```
>>> print(hp.get_all_neighbours(1, np.pi/2, np.pi/2))
[ 8  4  0 -1  1  6  9 -1]
```

```
>>> print(hp.get_all_neighbours(1, 90, 0, lonlat=True))
[ 8  4  0 -1  1  6  9 -1]
```

## healpy.pixelfunc.get\_interp\_weights

healpy.pixelfunc.**get\_interp\_weights**(*nside*, *theta*, *phi=None*, *nest=False*, *lonlat=False*)

Return the 4 closest pixels on the two rings above and below the location and corresponding weights. Weights are provided for bilinear interpolation along latitude and longitude

### Parameters

**nside** [int] the healpix nside

**theta, phi** [float, scalar or array-like] if phi is not given, theta is interpreted as pixel number, otherwise theta[rad],phi[rad] are angular coordinates

**nest** [bool] if True, NESTED ordering, otherwise RING ordering.

**lonlat** [bool] If True, input angles are assumed to be longitude and latitude in degree, otherwise, they are co-latitude and longitude in radians.

### Returns

**res** [tuple of length 2] contains pixel numbers in res[0] and weights in res[1]. Usual numpy broadcasting rules apply.

### See also:

*get\_interp\_val*, *get\_all\_neighbours*

## Examples

Note that some of the test inputs below that are on pixel boundaries such as theta=pi/2, phi=pi/2, have a tiny value of 1e-15 added to them to make them reproducible on i386 machines using x87 floating point instruction set (see <https://github.com/healpy/healpy/issues/528>).

```
>>> import healpy as hp
>>> pix, weights = hp.get_interp_weights(1, 0)
>>> print(pix)
[0 1 4 5]
>>> weights
array([ 1.,  0.,  0.,  0.])
```

```
>>> pix, weights = hp.get_interp_weights(1, 0, 0)
>>> print(pix)
[1 2 3 0]
>>> weights
array([ 0.25,  0.25,  0.25,  0.25])
```

```
>>> pix, weights = hp.get_interp_weights(1, 0, 90, lonlat=True)
>>> print(pix)
[1 2 3 0]
>>> weights
array([ 0.25,  0.25,  0.25,  0.25])
```

```
>>> pix, weights = hp.get_interp_weights(1, [0, np.pi/2 + 1e-15], 0)
>>> print(pix)
[[ 1  4]
 [ 2  5]
 [ 3 11]
 [ 0  8]]
>>> np.testing.assert_allclose(
...     weights,
...     np.array([[ 0.25,  1.  ],
...               [ 0.25,  0.  ],
...               [ 0.25,  0.  ],
...               [ 0.25,  0.  ]]), rtol=0, atol=1e-14)
```

## healpy.pixelfunc.get\_interp\_val

healpy.pixelfunc.**get\_interp\_val** (*m, theta, phi, nest=False, lonlat=False*)

Return the bi-linear interpolation value of a map using 4 nearest neighbours.

### Parameters

**m** [array-like] a healpix map, accepts masked arrays

**theta, phi** [float, scalar or array-like] angular coordinates of point at which to interpolate the map

**nest** [bool] if True, the is assumed to be in NESTED ordering.

**lonlat** [bool] If True, input angles are assumed to be longitude and latitude in degree, otherwise, they are co-latitude and longitude in radians.

### Returns

**val** [float, scalar or arry-like] the interpolated value(s), usual numpy broadcasting rules apply.

### See also:

[get\\_interp\\_weights](#), [get\\_all\\_neighbours](#)

## Examples

```
>>> import healpy as hp
>>> hp.get_interp_val(np.arange(12.), np.pi/2, 0)
4.0
>>> hp.get_interp_val(np.arange(12.), np.pi/2, np.pi/2)
5.0
>>> hp.get_interp_val(np.arange(12.), np.pi/2, np.pi/2 + 2*np.pi)
5.0
>>> hp.get_interp_val(np.arange(12.), np.linspace(0, np.pi, 10), 0)
array([ 1.5        ,  1.5        ,  1.5        ,  2.20618428,  3.40206143,
       5.31546486,  7.94639458,  9.5        ,  9.5        ,  9.5        ])
>>> hp.get_interp_val(np.arange(12.), 0, np.linspace(90, -90, 10), lonlat=True)
array([ 1.5        ,  1.5        ,  1.5        ,  2.20618428,  3.40206143,
       5.31546486,  7.94639458,  9.5        ,  9.5        ,  9.5        ])
```

## 3.1.2 conversion between NESTED and RING schemes

<code>nest2ring(nside, ipix)</code>	Convert pixel number from NESTED ordering to RING ordering.
<code>ring2nest(nside, ipix)</code>	Convert pixel number from RING ordering to NESTED ordering.
<code>reorder(map_in[, inp, out, r2n, n2r])</code>	Reorder a healpix map from RING/NESTED ordering to NESTED/RING

## healpy.pixelfunc.nest2ring

`healpy.pixelfunc.nest2ring(nside, ipix)`  
Convert pixel number from NESTED ordering to RING ordering.

### Parameters

**nside** [int, scalar or array-like] the healpix nside parameter  
**ipix** [int, scalar or array-like] the pixel number in NESTED scheme

### Returns

**ipix** [int, scalar or array-like] the pixel number in RING scheme

### See also:

`ring2nest, reorder`

## Examples

```
>>> import healpy as hp
>>> hp.nest2ring(16, 1130)
1504
```

```
>>> print(hp.nest2ring(2, np.arange(10)))
[13  5  4  0 15  7  6  1 17  9]
```

```
>>> print(hp.nest2ring([1, 2, 4, 8], 11))
[ 11    2   12 211]
```

## healpy.pixelfunc.ring2nest

`healpy.pixelfunc.ring2nest(nside, ipix)`  
Convert pixel number from RING ordering to NESTED ordering.

### Parameters

**nside** [int, scalar or array-like] the healpix nside parameter  
**ipix** [int, scalar or array-like] the pixel number in RING scheme

### Returns

**ipix** [int, scalar or array-like] the pixel number in NESTED scheme

### See also:

`nest2ring, reorder`

## Examples

```
>>> import healpy as hp
>>> hp.ring2nest(16, 1504)
1130
```

```
>>> print(hp.ring2nest(2, np.arange(10)))
[ 3  7 11 15  2  1  6  5 10  9]
```

```
>>> print(hp.ring2nest([1, 2, 4, 8], 11))
[ 11  13  61 253]
```

## healpy.pixelfunc.reorder

`healpy.pixelfunc.reorder(map_in, inp=None, out=None, r2n=None, n2r=None)`  
Reorder a healpix map from RING/NESTED ordering to NESTED/RING

### Parameters

`map_in` [array-like] the input map to reorder, accepts masked arrays

`inp, out` ['RING' or 'NESTED'] define the input and output ordering

`r2n` [bool] if True, reorder from RING to NESTED

`n2r` [bool] if True, reorder from NESTED to RING

### Returns

`map_out` [array-like] the reordered map, as masked array if the input was a masked array

### See also:

`nest2ring`, `ring2nest`

## Notes

if `r2n` or `n2r` is defined, override `inp` and `out`.

## Examples

```
>>> import healpy as hp
>>> hp.reorder(np.arange(48), r2n = True)
array([13,  5,  4,  0, 15,  7,  6,  1, 17,  9,  8,  2, 19, 11, 10,  3, 28,
       20, 27, 12, 30, 22, 21, 14, 32, 24, 23, 16, 34, 26, 25, 18, 44, 37,
       36, 29, 45, 39, 38, 31, 46, 41, 40, 33, 47, 43, 42, 35])
>>> hp.reorder(np.arange(12), n2r = True)
array([ 0,  1,  2,  3,  4,  5,  6,  7,  8,  9, 10, 11])
>>> hp.reorder(hp.ma(np.arange(12.)), n2r = True)
masked_array(data = [ 0.  1.  2.  3.  4.  5.  6.  7.  8.  9. 10. 11.],
             mask = False,
             fill_value = -1.6375e+30)
<BLANKLINE>
>>> m = [np.arange(12.), np.arange(12.), np.arange(12.)]
>>> m[0][2] = hp.UNSEEN
>>> m[1][2] = hp.UNSEEN
```

(continues on next page)

(continued from previous page)

```
>>> m[2][2] = hp.UNSEEN
>>> m = hp.ma(m)
>>> hp.reorder(m, n2r = True)
masked_array(data =
 [[0.0 1.0 -- 3.0 4.0 5.0 6.0 7.0 8.0 9.0 10.0 11.0]
 [0.0 1.0 -- 3.0 4.0 5.0 6.0 7.0 8.0 9.0 10.0 11.0]
 [0.0 1.0 -- 3.0 4.0 5.0 6.0 7.0 8.0 9.0 10.0 11.0]],
 mask =
 [[False False True False False False False False False]
 [False False True False False False False False False]
 [False False True False False False False False False]], fill_value = -1.6375e+30)
<BLANKLINE>
```

### 3.1.3 nside/npix/resolution

<code>nside2npix(nside)</code>	Give the number of pixels for the given nside.
<code>npix2nside(npix)</code>	Give the nside parameter for the given number of pixels.
<code>nside2order(nside)</code>	Give the resolution order for a given nside.
<code>order2nside(order)</code>	Give the nside parameter for the given resolution order.
<code>nside2resol(nside[, arcmin])</code>	Give approximate resolution (pixel size in radian or arcmin) for nside.
<code>nside2pixarea(nside[, degrees])</code>	Give pixel area given nside in square radians or square degrees.
<code>max_pixrad(nside[, degrees])</code>	Maximum angular distance between any pixel center and its corners
<code>isnsideok(nside[, nest])</code>	Returns True if nside is a valid nside parameter, False otherwise.
<code>isnpixok(npix)</code>	Return True if npix is a valid value for healpix map size, False otherwise.
<code>get_map_size(m)</code>	Returns the npix of a given map (implicit or explicit pixellation).
<code>get_min_valid_nside(npix)</code>	Returns the minimum acceptable nside so that npix <= nside2npix(nside).
<code>get_nside(m)</code>	Return the nside of the given map.
<code>maptype(m)</code>	Describe the type of the map (valid, single, sequence of maps).
<code>ud_grade(map_in, nside_out[, pess, ...])</code>	Upgrade or degrade resolution of a map (or list of maps).

#### healpy.pixelfunc.nside2npix

`healpy.pixelfunc.nside2npix(nside)`

Give the number of pixels for the given nside.

##### Parameters

`nside` [int] healpix nside parameter; an exception is raised if nside is not valid (nside must be a power of 2, less than  $2^{**30}$ )

##### Returns

`npix` [int] corresponding number of pixels

## Notes

Raise a ValueError exception if nside is not valid.

## Examples

```
>>> import healpy as hp
>>> import numpy as np
>>> hp.nside2npix(8)
768
```

```
>>> np.all([hp.nside2npix(nside) == 12 * nside**2 for nside in [2**n for n in range(12)]])
True
```

```
>>> hp.nside2npix(7)
588
```

## healpy.pixelfunc.npix2nside

`healpy.pixelfunc.npix2nside(npix)`

Give the nside parameter for the given number of pixels.

### Parameters

`npix` [int] the number of pixels

### Returns

`nside` [int] the nside parameter corresponding to npix

## Notes

Raise a ValueError exception if number of pixel does not correspond to the number of pixel of a healpix map.

## Examples

```
>>> import healpy as hp
>>> hp.npix2nside(768)
8
```

```
>>> np.all([hp.npix2nside(12 * nside**2) == nside for nside in [2**n for n in range(12)]])
True
```

```
>>> hp.npix2nside(1000)
Traceback (most recent call last):
...
ValueError: Wrong pixel number (it is not 12*nside**2)
```

## healpy.pixelfunc.nside2order

healpy.pixelfunc.**nside2order**(*nside*)  
Give the resolution order for a given nside.

### Parameters

**nside** [int] healpix nside parameter; an exception is raised if nside is not valid (nside must be a power of 2, less than  $2^{**30}$ )

### Returns

**order** [int] corresponding order where  $\text{nside} = 2^{**(\text{order})}$

### Notes

Raise a ValueError exception if nside is not valid.

### Examples

```
>>> import healpy as hp
>>> import numpy as np
>>> hp.nside2order(128)
7
```

```
>>> np.all([hp.nside2order(2**o) == o for o in range(30)])
True
```

```
>>> hp.nside2order(7)
Traceback (most recent call last):
...
ValueError: 7 is not a valid nside parameter (must be a power of 2, less than
2**30)
```

## healpy.pixelfunc.order2nside

healpy.pixelfunc.**order2nside**(*order*)  
Give the nside parameter for the given resolution order.

### Parameters

**order** [int] the resolution order

### Returns

**nside** [int] the nside parameter corresponding to order

### Notes

Raise a ValueError exception if order produces an nside out of range.

## Examples

```
>>> import healpy as hp
>>> hp.order2nside(7)
128
```

```
>>> print(hp.order2nside(np.arange(8)))
[ 1   2   4   8  16  32  64 128]
```

```
>>> hp.order2nside(31)
Traceback (most recent call last):
...
ValueError: 2147483648 is not a valid nside parameter (must be a power of 2, less than 2**30)
```

## healpy.pixelfunc.nside2resol

`healpy.pixelfunc.nside2resol(nside, arcmin=False)`

Give approximate resolution (pixel size in radian or arcmin) for nside.

Resolution is just the square root of the pixel area, which is a gross approximation given the different pixel shapes

### Parameters

`nside` [int] healpix nside parameter, must be a power of 2, less than  $2^{**30}$

`arcmin` [bool] if True, return resolution in arcmin, otherwise in radian

### Returns

`resol` [float] approximate pixel size in radians or arcmin

## Notes

Raise a ValueError exception if nside is not valid.

## Examples

```
>>> import healpy as hp
>>> hp.nside2resol(128, arcmin = True)
27.483891294539248
```

```
>>> hp.nside2resol(256)
0.0039973699529159707
```

```
>>> hp.nside2resol(7)
0.1461895297066412
```

## healpy.pixelfunc.nside2pixarea

`healpy.pixelfunc.nside2pixarea(nside, degrees=False)`

Give pixel area given nside in square radians or square degrees.

### Parameters

**nside** [int] healpix nside parameter, must be a power of 2, less than  $2^{**30}$   
**degrees** [bool] if True, returns pixel area in square degrees, in square radians otherwise

### Returns

**pixarea** [float] pixel area in square radian or square degree

### Notes

Raise a ValueError exception if nside is not valid.

### Examples

```
>>> import healpy as hp
>>> hp.nside2pixarea(128, degrees = True)
0.2098234113027917
```

```
>>> hp.nside2pixarea(256)
1.5978966540475428e-05
```

```
>>> hp.nside2pixarea(7)
0.021371378595848933
```

## healpy.pixelfunc.max\_pixrad

healpy.pixelfunc.**max\_pixrad**(nside, degrees=False)

Maximum angular distance between any pixel center and its corners

### Parameters

**nside** [int] the nside to work with  
**degrees** [bool] if True, returns pixel radius in degrees, in radians otherwise

### Returns

**rads: double** angular distance (in radians or degrees)

### Examples

```
>>> '%.14f' % max_pixrad(1)
'0.84106867056793'
>>> '%.14f' % max_pixrad(16)
'0.06601476143251'
```

## healpy.pixelfunc.isnsideok

healpy.pixelfunc.**isnsideok**(nside, nest=False)

Returns True if nside is a valid nside parameter, False otherwise.

NSIDE needs to be a power of 2 only for nested ordering

**Parameters**

**nside** [int, scalar or array-like] integer value to be tested

**Returns**

**ok** [bool, scalar or array-like] True if given value is a valid nside, False otherwise.

**Examples**

```
>>> import healpy as hp
>>> hp.isnsideok(13, nest=True)
False
```

```
>>> hp.isnsideok(13, nest=False)
True
```

```
>>> hp.isnsideok(32)
True
```

```
>>> hp.isnsideok([1, 2, 3, 4, 8, 16], nest=True)
array([ True,  True, False,  True,  True,  True], dtype=bool)
```

**healpy.pixelfunc.isnpixok**

healpy.pixelfunc.**isnpixok**(*npix*)

Return True if npix is a valid value for healpix map size, False otherwise.

**Parameters**

**npix** [int, scalar or array-like] integer value to be tested

**Returns**

**ok** [bool, scalar or array-like] True if given value is a valid number of pixel, False otherwise

**Examples**

```
>>> import healpy as hp
>>> hp.isnpixok(12)
True
```

```
>>> hp.isnpixok(768)
True
```

```
>>> hp.isnpixok([12, 768, 1002])
array([ True,  True, False], dtype=bool)
```

**healpy.pixelfunc.get\_map\_size**

healpy.pixelfunc.**get\_map\_size**(*m*)

Returns the npix of a given map (implicit or explicit pixelization).

If map is a dict type, assumes explicit pixelization: use nside key if present, or use nside attribute if present, otherwise use the smallest valid npix given the maximum key value. otherwise assumes implicit pixelization and returns len(m).

#### Parameters

**m** [array-like or dict-like] a map with implicit (array-like) or explicit (dict-like) pixellization

#### Returns

**npix** [int] a valid number of pixel

#### Notes

In implicit pixellization, raise a ValueError exception if the size of the input is not a valid pixel number.

#### Examples

```
>>> import healpy as hp
>>> m = {0: 1, 1: 1, 2: 1, 'nside': 1}
>>> print(hp.get_map_size(m))
12
```

```
>>> m = {0: 1, 767: 1}
>>> print(hp.get_map_size(m))
768
```

```
>>> print(hp.get_map_size(np.zeros(12 * 8 ** 2)))
768
```

## healpy.pixelfunc.get\_min\_valid\_nside

healpy.pixelfunc.**get\_min\_valid\_nside**(npix)

Returns the minimum acceptable nside so that npix <= nside2npix(nside).

#### Parameters

**npix** [int] a minimal number of pixel

#### Returns

**nside** [int] a valid healpix nside so that  $12 * \text{nside}^{** 2} \geq \text{npix}$

#### Examples

```
>>> import healpy as hp
>>> hp.pixelfunc.get_min_valid_nside(355)
8
>>> hp.pixelfunc.get_min_valid_nside(768)
8
```

## healpy.pixelfunc.get\_nside

`healpy.pixelfunc.get_nside(m)`

Return the nside of the given map.

### Parameters

**m** [sequence] the map to get the nside from.

### Returns

**nside** [int] the healpix nside parameter of the map (or sequence of maps)

## Notes

If the input is a sequence of maps, all of them must have same size. If the input is not a valid map (not a sequence, invalid number of pixels), a `TypeError` exception is raised.

## healpy.pixelfunc.maptyle

`healpy.pixelfunc.maptyle(m)`

Describe the type of the map (valid, single, sequence of maps). Checks : the number of maps, that all maps have same length and that this length is a valid map size (using `isnpxok()`).

### Parameters

**m** [sequence] the map to get info from

### Returns

**info** [int] -1 if the given object is not a valid map, 0 if it is a single map,  $info > 0$  if it is a sequence of maps ( $info$  is then the number of maps)

## Examples

```
>>> import healpy as hp
>>> hp.pixelfunc.maptyle(np.arange(12))
0
>>> hp.pixelfunc.maptyle([np.arange(12), np.arange(12)])
2
```

## healpy.pixelfunc.ud\_grade

`healpy.pixelfunc.ud_grade(map_in, nside_out, pess=False, order_in='RING', order_out=None, power=None, dtype=None)`

Upgrade or degrade resolution of a map (or list of maps).

in degrading the resolution, ud\_grade sets the value of the superpixel as the mean of the children pixels.

### Parameters

**map\_in** [array-like or sequence of array-like] the input map(s) (if a sequence of maps, all must have same size)

**nside\_out** [int] the desired nside of the output map(s)

**pess** [bool] if True, in degrading, reject pixels which contains a bad sub\_pixel. Otherwise, estimate average with good pixels

**order\_in, order\_out** [str] pixel ordering of input and output ('RING' or 'NESTED')

**power** [float] if non-zero, divide the result by (nside\_in/nside\_out)\*\*power Examples: power=-2 keeps the sum of the map invariant (useful for hitmaps), power=2 divides the mean by another factor of (nside\_in/nside\_out)\*\*2 (useful for variance maps)

**dtype** [type] the type of the output map

#### Returns

**map\_out** [array-like or sequence of array-like] the upgraded or degraded map(s)

### Examples

```
>>> import healpy as hp
>>> hp.ud_grade(np.arange(48.), 1)
array([ 5.5 ,  7.25,  9. ,  10.75, 21.75, 21.75, 23.75, 25.75,
       36.5 , 38.25, 40. , 41.75])
```

## 3.1.4 Masking pixels

<i>UNSEEN</i>	Special value used for masked pixels
<code>mask_bad(m[, badval, rtol, atol])</code>	Returns a bool array with True where m is close to badval.
<code>mask_good(m[, badval, rtol, atol])</code>	Returns a bool array with False where m is close to badval.
<code>ma(m[, badval, rtol, atol, copy])</code>	Return map as a masked array, with badval pixels masked.

### healpy.pixelfunc.UNSEEN

`healpy.pixelfunc.UNSEEN = -1.6375e+30`

Special value used for masked pixels

### healpy.pixelfunc.mask\_bad

`healpy.pixelfunc.mask_bad(m, badval=-1.6375e+30, rtol=1e-05, atol=1e-08)`

Returns a bool array with True where m is close to badval.

#### Parameters

**m** [a map (may be a sequence of maps)]

**badval** [float, optional] The value of the pixel considered as bad (*UNSEEN* by default)

**rtol** [float, optional] The relative tolerance

**atol** [float, optional] The absolute tolerance

#### Returns

**mask** a bool array with the same shape as the input map, True where input map is close to badval, and False elsewhere.

**See also:**

`mask_good, ma`

## Examples

```
>>> import healpy as hp
>>> import numpy as np
>>> m = np.arange(12.)
>>> m[3] = hp.UNSEEN
>>> hp.mask_bad(m)
array([False, False, False, True, False, False, False, False,
       False, False], dtype=bool)
```

## healpy.pixelfunc.mask\_good

`healpy.pixelfunc.mask_good(m, badval=-1.6375e+30, rtol=1e-05, atol=1e-08)`

Returns a bool array with `False` where `m` is close to `badval`.

### Parameters

`m` [a map (may be a sequence of maps)]

`badval` [float, optional] The value of the pixel considered as bad (`UNSEEN` by default)

`rtol` [float, optional] The relative tolerance

`atol` [float, optional] The absolute tolerance

### Returns

a bool array with the same shape as the input map, “`False`“ where input map is close to `badval`, and “`True`“ elsewhere.

**See also:**

`mask_bad, ma`

## Examples

```
>>> import healpy as hp
>>> m = np.arange(12.)
>>> m[3] = hp.UNSEEN
>>> hp.mask_good(m)
array([ True,  True,  True, False,  True,  True,  True,  True,
       True,  True,  True], dtype=bool)
```

## healpy.pixelfunc.ma

`healpy.pixelfunc.ma(m, badval=-1.6375e+30, rtol=1e-05, atol=1e-08, copy=True)`

Return map as a masked array, with `badval` pixels masked.

### Parameters

`m` [a map (may be a sequence of maps)]

`badval` [float, optional] The value of the pixel considered as bad (`UNSEEN` by default)

**rtol** [float, optional] The relative tolerance  
**atol** [float, optional] The absolute tolerance  
**copy** [bool, optional] If `True`, a copy of the input map is made.

#### Returns

a masked array with the same shape as the input map,  
masked where input map is close to `badval`.

#### See also:

`mask_good`, `mask_bad`, `numpy.ma.masked_values`

#### Examples

```
>>> import healpy as hp
>>> m = np.arange(12.)
>>> m[3] = hp.UNSEEN
>>> hp.ma(m)
masked_array(data = [0.0 1.0 2.0 -- 4.0 5.0 6.0 7.0 8.0 9.0 10.0 11.0],
             mask = [False False False True False False False False False,
                     False False],
             fill_value = -1.6375e+30)
<BLANKLINE>
```

### 3.1.5 Map data manipulation

<code>fit_dipole(m[, nest, bad, gal_cut])</code>	Fit a dipole and a monopole to the map, excluding bad pixels.
<code>fit_monopole(m[, nest, bad, gal_cut])</code>	Fit a monopole to the map, excluding unseen pixels.
<code>remove_dipole(m[, nest, bad, gal_cut, ...])</code>	Fit and subtract the dipole and the monopole from the given map <code>m</code> .
<code>remove_monopole(m[, nest, bad, gal_cut, ...])</code>	Fit and subtract the monopole from the given map <code>m</code> .
<code>get_interp_val(m, theta, phi[, nest, lonlat])</code>	Return the bi-linear interpolation value of a map using 4 nearest neighbours.

#### healpy.pixelfunc.fit\_dipole

`healpy.pixelfunc.fit_dipole(m, nest=False, bad=-1.6375e+30, gal_cut=0)`

Fit a dipole and a monopole to the map, excluding bad pixels.

#### Parameters

**m** [float, array-like] the map to which a dipole is fitted and subtracted, accepts masked maps  
**nest** [bool] if `False` `m` is assumed in RING scheme, otherwise map is NESTED  
**bad** [float] bad values of pixel, default to `UNSEEN`.  
**gal\_cut** [float [degrees]] pixels at latitude in  $[-\text{gal\_cut}; +\text{gal\_cut}]$  degrees are not taken into account

#### Returns

**res** [tuple of length 2] the monopole value in `res[0]` and the dipole vector (as array) in `res[1]`

See also:

`remove_dipole`, `fit_monopole`, `remove_monopole`

## healpy.pixelfunc.fit\_monopole

`healpy.pixelfunc.fit_monopole(m, nest=False, bad=-1.6375e+30, gal_cut=0)`

Fit a monopole to the map, excluding unseen pixels.

### Parameters

**m** [float, array-like] the map to which a dipole is fitted and subtracted, accepts masked arrays

**nest** [bool] if `False` m is assumed in RING scheme, otherwise map is NESTED

**bad** [float] bad values of pixel, default to `UNSEEN`.

**gal\_cut** [float [degrees]] pixels at latitude in [-gal\_cut;+gal\_cut] degrees are not taken into account

### Returns

**res: float** fitted monopole value

See also:

`fit_dipole`, `remove_monopole`, `remove_monopole`

## healpy.pixelfunc.remove\_dipole

`healpy.pixelfunc.remove_dipole(m, nest=False, bad=-1.6375e+30, gal_cut=0, fitval=False, copy=True, verbose=True)`

Fit and subtract the dipole and the monopole from the given map m.

### Parameters

**m** [float, array-like] the map to which a dipole is fitted and subtracted, accepts masked arrays

**nest** [bool] if `False` m is assumed in RING scheme, otherwise map is NESTED

**bad** [float] bad values of pixel, default to `UNSEEN`.

**gal\_cut** [float [degrees]] pixels at latitude in [-gal\_cut;+gal\_cut] are not taken into account

**fitval** [bool] whether to return or not the fitted values of monopole and dipole

**copy** [bool] whether to modify input map or not (by default, make a copy)

**verbose** [bool] print values of monopole and dipole

### Returns

**res** [array or tuple of length 3] if `fitval` is `False`, returns map with monopole and dipole subtracted, otherwise, returns map (array, in `res[0]`), monopole (float, in `res[1]`), dipole\_vector (array, in `res[2]`)

See also:

`fit_dipole`, `fit_monopole`, `remove_monopole`

**healpy.pixelfunc.remove\_monopole**

```
healpy.pixelfunc.remove_monopole(m, nest=False, bad=-1.6375e+30, gal_cut=0, fitval=False,  
    copy=True, verbose=True)
```

Fit and subtract the monopole from the given map *m*.

**Parameters**

- m*** [float, array-like] the map to which a monopole is fitted and subtracted
- nest*** [bool] if *False* *m* is assumed in RING scheme, otherwise map is NESTED
- bad*** [float] bad values of pixel, default to *UNSEEN*.
- gal\_cut*** [float [degrees]] pixels at latitude in [-*gal\_cut*;+*gal\_cut*] are not taken into account
- fitval*** [bool] whether to return or not the fitted value of monopole
- copy*** [bool] whether to modify input map or not (by default, make a copy)
- verbose: bool*** whether to print values of monopole

**Returns**

- res*** [array or tuple of length 3] if *fitval* is False, returns map with monopole subtracted, otherwise, returns map (array, in *res[0]*) and monopole (float, in *res[1]*)

**See also:**

*fit\_dipole*, *fit\_monopole*, *remove\_dipole*

## 3.2 sphtfunc – Spherical harmonic transforms

### 3.2.1 From map to spherical harmonics

<i>anafast</i> ( <i>map1</i> [, <i>map2</i> , <i>nspec</i> , <i>lmax</i> , <i>mmax</i> , ...])	Computes the power spectrum of a Healpix map, or the cross-spectrum between two maps if <i>map2</i> is given.
<i>map2alm</i> ( <i>maps</i> [, <i>lmax</i> , <i>mmax</i> , <i>iter</i> , <i>pol</i> , ...])	Computes the alm of a Healpix map.

**healpy.sphtfunc.anafast**

```
healpy.sphtfunc.anafast(map1, map2=None, nspec=None, lmax=None, mmax=None, iter=3,  
    alm=False, pol=True, use_weights=False, datapath=None, gal_cut=0,  
    use_pixel_weights=False)
```

Computes the power spectrum of a Healpix map, or the cross-spectrum between two maps if *map2* is given. No removal of monopole or dipole is performed. The input maps must be in ring-ordering.

**Parameters**

- map1*** [float, array-like shape (Npix,) or (3, Npix)] Either an array representing a map, or a sequence of 3 arrays representing I, Q, U maps. Must be in ring ordering.
- map2*** [float, array-like shape (Npix,) or (3, Npix)] Either an array representing a map, or a sequence of 3 arrays representing I, Q, U maps. Must be in ring ordering.
- nspec*** [None or int, optional] The number of spectra to return. If None, returns all, otherwise returns *cls[:nspec]*
- lmax*** [int, scalar, optional] Maximum l of the power spectrum (default: 3\*nside-1)

**mmax** [int, scalar, optional] Maximum m of the alm (default: lmax)  
**iter** [int, scalar, optional] Number of iteration (default: 3)  
**alm** [bool, scalar, optional] If True, returns both cl and alm, otherwise only cl is returned  
**pol** [bool, optional] If True, assumes input maps are TQU. Output will be TEB cl's and correlations (input must be 1 or 3 maps). If False, maps are assumed to be described by spin 0 spherical harmonics. (input can be any number of maps) If there is only one input map, it has no effect. Default: True.  
**datapath** [None or str, optional] If given, the directory where to find the weights data.  
**gal\_cut** [float [degrees]] pixels at latitude in [-gal\_cut;+gal\_cut] are not taken into account  
**use\_pixel\_weights: bool, optional** If True, use pixel by pixel weighting, healpy will automatically download the weights, if needed

#### Returns

**res** [array or sequence of arrays] If *alm* is False, returns cl or a list of cl's (TT, EE, BB, TE, EB, TB for polarized input map) Otherwise, returns a tuple (cl, alm), where cl is as above and alm is the spherical harmonic transform or a list of almT, almE, almB for polarized input

## healpy.sphtfunc.map2alm

```
healpy.sphtfunc.map2alm(maps, lmax=None, mmax=None, iter=3, pol=True, use_weights=False,
                        datapath=None, gal_cut=0, use_pixel_weights=False)
```

Computes the alm of a Healpix map. The input maps must all be in ring ordering.

#### Parameters

**maps** [array-like, shape (Npix,) or (n, Npix)] The input map or a list of n input maps. Must be in ring ordering.  
**lmax** [int, scalar, optional] Maximum l of the power spectrum. Default: 3\*nside-1  
**mmax** [int, scalar, optional] Maximum m of the alm. Default: lmax  
**iter** [int, scalar, optional] Number of iteration (default: 3)  
**pol** [bool, optional] If True, assumes input maps are TQU. Output will be TEB alm's. (input must be 1 or 3 maps) If False, apply spin 0 harmonic transform to each map. (input can be any number of maps) If there is only one input map, it has no effect. Default: True.  
**use\_weights: bool, scalar, optional** If True, use the ring weighting. Default: False.  
**datapath** [None or str, optional] If given, the directory where to find the weights data.  
**gal\_cut** [float [degrees]] pixels at latitude in [-gal\_cut;+gal\_cut] are not taken into account  
**use\_pixel\_weights: bool, optional** If True, use pixel by pixel weighting, healpy will automatically download the weights, if needed

#### Returns

**alms** [array or tuple of array] alm or a tuple of 3 alm (almT, almE, almB) if polarized input.

#### Notes

The pixels which have the special *UNSEEN* value are replaced by zeros before spherical harmonic transform. They are converted back to *UNSEEN* value, so that the input maps are not modified. Each map have its own, independent mask.

### 3.2.2 From spherical harmonics to map

<code>synfast</code> (cls, nside[, lmax, mmax, alm, pol, ...])	Create a map(s) from cl(s).
<code>alm2map</code> (alms, nside[, lmax, mmax, pixwin, ...])	Computes a Healpix map given the alm.
<code>alm2map_der1</code> (alm, nside[, lmax, mmax])	Computes a Healpix map and its first derivatives given the alm.

#### healpy.sphtfunc.synfast

```
healpy.sphtfunc.synfast (cls, nside, lmax=None, mmax=None, alm=False, pol=True, pixwin=False,  
                      fwhm=0.0, sigma=None, new=False, verbose=True)
```

Create a map(s) from cl(s).

##### Parameters

- cls** [array or tuple of array] A cl or a list of cl (either 4 or 6, see [synalm\(\)](#))
- nside** [int, scalar] The nside of the output map(s)
- lmax** [int, scalar, optional] Maximum l for alm. Default: min of 3\*nside-1 or length of the cls - 1
- mmax** [int, scalar, optional] Maximum m for alm.
- alm** [bool, scalar, optional] If True, return also alm(s). Default: False.
- pol** [bool, optional] If True, assumes input cls are TEB and correlation. Output will be TQU maps. (input must be 1, 4 or 6 cl's) If False, fields are assumed to be described by spin 0 spherical harmonics. (input can be any number of cl's) If there is only one input cl, it has no effect. Default: True.
- pixwin** [bool, scalar, optional] If True, convolve the alm by the pixel window function. Default: False.
- fwhm** [float, scalar, optional] The fwhm of the Gaussian used to smooth the map (applied on alm) [in radians]
- sigma** [float, scalar, optional] The sigma of the Gaussian used to smooth the map (applied on alm) [in radians]

##### Returns

- maps** [array or tuple of arrays] The output map (possibly list of maps if polarized input). or, if alm is True, a tuple of (map,alm) (alm possibly a list of alm if polarized input)

#### Notes

The order of the spectra will change in a future release. The new= parameter help to make the transition smoother. You can start using the new order by setting new=True. In the next version of healpy, the default will be new=True. This change is done for consistency between the different tools (alm2cl, synfast, anafast). In the new order, the spectra are ordered by diagonal of the correlation matrix. Eg, if fields are T, E, B, the spectra are TT, EE, BB, TE, EB, TB with new=True, and TT, TE, TB, EE, EB, BB if new=False.

#### healpy.sphtfunc.alm2map

```
healpy.sphtfunc.alm2map (alms, nside, lmax=None, mmax=None, pixwin=False, fwhm=0.0,  
                      sigma=None, pol=True, inplace=False, verbose=True)
```

Computes a Healpix map given the alm.

The alm are given as a complex array. You can specify lmax and mmax, or they will be computed from array size (assuming lmax==mmax).

### Parameters

**alms** [complex, array or sequence of arrays] A complex array or a sequence of complex arrays.

Each array must have a size of the form:  $mmax * (2 * lmax + 1 - mmax) / 2 + lmax + 1$

**nside** [int, scalar] The nside of the output map.

**lmax** [None or int, scalar, optional] Explicitly define lmax (needed if mmax!=lmax)

**mmax** [None or int, scalar, optional] Explicitly define mmax (needed if mmax!=lmax)

**pixwin** [bool, optional] Smooth the alm using the pixel window functions. Default: False.

**fwhm** [float, scalar, optional] The fwhm of the Gaussian used to smooth the map (applied on alm) [in radians]

**sigma** [float, scalar, optional] The sigma of the Gaussian used to smooth the map (applied on alm) [in radians]

**pol** [bool, optional] If True, assumes input alms are TEB. Output will be TQU maps. (input must be 1 or 3 alms) If False, apply spin 0 harmonic transform to each alm. (input can be any number of alms) If there is only one input alm, it has no effect. Default: True.

**inplace** [bool, optional] If True, input alms may be modified by pixel window function and beam smoothing (if alm(s) are complex128 contiguous arrays). Otherwise, input alms are not modified. A copy is made if needed to apply beam smoothing or pixel window.

### Returns

**maps** [array or list of arrays] A Healpix map in RING scheme at nside or a list of T,Q,U maps (if polarized input)

### Notes

Running map2alm then alm2map will not return exactly the same map if the discretized field you construct on the sphere is not band-limited (for example, if you have a map containing pixel-based noise rather than beam-smoothed noise). If you need a band-limited map, you have to start with random numbers in lm space and transform these via alm2map. With such an input, the accuracy of map2alm->alm2map should be quite good, depending on your choices of lmax, mmax and nside (for some typical values, see e.g., section 5.1 of <https://arxiv.org/pdf/1010.2084.pdf>).

## healpy.sphtfunc.alm2map\_der1

`healpy.sphtfunc.alm2map_der1(alm, nside, lmax=None, mmax=None)`

Computes a Healpix map and its first derivatives given the alm.

The alm are given as a complex array. You can specify lmax and mmax, or they will be computed from array size (assuming lmax==mmax).

### Parameters

**alm** [array, complex] A complex array of alm. Size must be of the form  $mmax(lmax-mmax+1)/2+lmax$

**nside** [int] The nside of the output map.

**lmax** [None or int, optional] Explicitly define lmax (needed if mmax!=lmax)

**mmax** [None or int, optional] Explicitly define mmax (needed if mmax!=lmax)

**Returns**

**m, d\_theta, d\_phi** [tuple of arrays] The maps correponding to alm, and its derivatives with respect to theta and phi. d\_phi is already divided by sin(theta)

### 3.2.3 Spherical harmonic transform tools

<code>smoothing(map_in[, fwhm, sigma, ...])</code>	Smooth a map with a Gaussian symmetric beam.
<code>smoothalm(alms[, fwhm, sigma, beam_window, ...])</code>	Smooth alm with a Gaussian symmetric beam function.
<code>alm2cl(alms1[, alms2, lmax, mmax, lmax_out, ...])</code>	Computes (cross-)spectra from alm(s).
<code>synalm(cls[, lmax, mmax, new, verbose])</code>	Generate a set of alm given cl.
<code>almxfl(alm, fl[, mmax, inplace])</code>	Multiply alm by a function of l.
<code>pixwin(nside[, pol, lmax])</code>	Return the pixel window function for the given nside.
<code>Alm()</code>	This class provides some static methods for alm index computation.

#### healpy.sphtfunc.smoothing

```
healpy.sphtfunc.smoothing(map_in,      fwhm=0.0,      sigma=None,      beam_window=None,
                           pol=True,     iter=3,       lmax=None,     mmax=None,     use_weights=False,
                           use_pixel_weights=False, datapath=None, verbose=True)
```

Smooth a map with a Gaussian symmetric beam.

No removal of monopole or dipole is performed.

**Parameters**

**map\_in** [array or sequence of 3 arrays] Either an array representing one map, or a sequence of 3 arrays representing 3 maps, accepts masked arrays

**fwhm** [float, optional] The full width half max parameter of the Gaussian [in radians]. Default: 0.0

**sigma** [float, optional] The sigma of the Gaussian [in radians]. Override fwhm.

**beam\_window: array, optional** Custom beam window function. Override fwhm and sigma.

**pol** [bool, optional] If True, assumes input maps are TQU. Output will be TQU maps. (input must be 1 or 3 alms) If False, each map is assumed to be a spin 0 map and is treated independently (input can be any number of alms). If there is only one input map, it has no effect. Default: True.

**iter** [int, scalar, optional] Number of iteration (default: 3)

**lmax** [int, scalar, optional] Maximum l of the power spectrum. Default: 3\*nside-1

**mmax** [int, scalar, optional] Maximum m of the alm. Default: lmax

**use\_weights: bool, scalar, optional** If True, use the ring weighting. Default: False.

**use\_pixel\_weights: bool, optional** If True, use pixel by pixel weighting, healpy will automatically download the weights, if needed

**datapath** [None or str, optional] If given, the directory where to find the weights data.

**verbose** [bool, optional] If True prints diagnostic information. Default: True

**Returns**

**maps** [array or list of 3 arrays] The smoothed map(s)

## healpy.sphtfunc.smoothalm

```
healpy.sphtfunc.smoothalm(alms, fwhm=0.0, sigma=None, beam_window=None, pol=True,
                           mmax=None, verbose=True, inplace=True)
```

Smooth alm with a Gaussian symmetric beam function.

### Parameters

**alms** [array or sequence of 3 arrays] Either an array representing one alm, or a sequence of arrays. See *pol* parameter.

**fwhm** [float, optional] The full width half max parameter of the Gaussian. Default:0.0 [in radians]

**sigma** [float, optional] The sigma of the Gaussian. Override fwhm. [in radians]

**beam\_window: array, optional** Custom beam window function. Override fwhm and sigma.

**pol** [bool, optional] If True, assumes input alms are TEB. Output will be TQU maps. (input must be 1 or 3 alms) If False, apply spin 0 harmonic transform to each alm. (input can be any number of alms) If there is only one input alm, it has no effect. Default: True.

**mmax** [None or int, optional] The maximum m for alm. Default: mmax=lmax

**inplace** [bool, optional] If True, the alm's are modified inplace if they are contiguous arrays of type complex128. Otherwise, a copy of alm is made. Default: True.

**verbose** [bool, optional] If True prints diagnostic information. Default: True

### Returns

**alms** [array or sequence of 3 arrays] The smoothed alm. If alm[i] is a contiguous array of type complex128, and *inplace* is True the smoothing is applied inplace. Otherwise, a copy is made.

## healpy.sphtfunc.alm2cl

```
healpy.sphtfunc.alm2cl(alms1, alms2=None, lmax=None, mmax=None, lmax_out=None,
                       nspec=None)
```

Computes (cross-)spectra from alm(s). If alm2 is given, cross-spectra between alm and alm2 are computed. If alm (and alm2 if provided) contains n alm, then n(n+1)/2 auto and cross-spectra are returned.

### Parameters

**alm** [complex, array or sequence of arrays] The alm from which to compute the power spectrum. If n $\geq$ 2 arrays are given, computes both auto- and cross-spectra.

**alms2** [complex, array or sequence of 3 arrays, optional] If provided, computes cross-spectra between alm and alm2. Default: alm2=alm, so auto-spectra are computed.

**lmax** [None or int, optional] The maximum l of the input alm. Default: computed from size of alm and mmax\_in

**mmax** [None or int, optional] The maximum m of the input alm. Default: assume mmax\_in = lmax\_in

**lmax\_out** [None or int, optional] The maximum l of the returned spectra. By default: the lmax of the given alm(s).

**nspec** [None or int, optional] The number of spectra to return. None means all, otherwise returns cl[:nspec]

### Returns

**cl** [array or tuple of n(n+1)/2 arrays] the spectrum  $\langle alm \times alm2 \rangle$  if *alm* (and *alm2*) is one *alm*, or the auto- and cross-spectra  $\langle alm^*[i] \times *alm2^*[j] \rangle$  if *alm* (and *alm2*) contains more than one spectra. If more than one spectrum is returned, they are ordered by diagonal. For example, if  $*alm$  is *almT*, *almE*, *almB*, then the returned spectra are: TT, EE, BB, TE, EB, TB.

## healpy.sphtfunc.synalm

`healpy.sphtfunc.synalm(cls, lmax=None, mmax=None, new=False, verbose=True)`

Generate a set of *alm* given *cl*. The *cl* are given as a float array. Corresponding *alm* are generated. If *lmax* is None, it is assumed *lmax*=*cl.size-1*. If *mmax* is None, it is assumed *mmax*=*lmax*.

### Parameters

**cls** [float, array or tuple of arrays] Either one *cl* (1D array) or a tuple of either 4 *cl* or of n\*(n+1)/2 *cl*. Some of the *cl* may be None, implying no cross-correlation. See *new* parameter.

**lmax** [int, scalar, optional] The *lmax* (if None or <0, the largest size-1 of *cls*)

**mmax** [int, scalar, optional] The *mmax* (if None or <0, =*lmax*)

**new** [bool, optional] If True, use the new ordering of *cl*'s, ie by diagonal (e.g. TT, EE, BB, TE, EB, TB or TT, EE, BB, TE if 4 *cl* as input). If False, use the old ordering, ie by row (e.g. TT, TE, TB, EE, EB, BB or TT, TE, EE, BB if 4 *cl* as input).

### Returns

**alms** [array or list of arrays] the generated *alm* if one spectrum is given, or a list of *n* *alms* (with n(n+1)/2 the number of input *cl*, or n=3 if there are 4 input *cl*).

## Notes

The order of the spectra will change in a future release. The *new*= parameter help to make the transition smoother. You can start using the new order by setting *new*=True. In the next version of healpy, the default will be *new*=True. This change is done for consistency between the different tools (*alm2cl*, *synfast*, *anafast*). In the new order, the spectra are ordered by diagonal of the correlation matrix. Eg, if fields are T, E, B, the spectra are TT, EE, BB, TE, EB, TB with *new*=True, and TT, TE, TB, EE, EB, BB if *new*=False.

## healpy.sphtfunc.almxfl

`healpy.sphtfunc.almxfl(alm, fl, mmax=None, inplace=False)`

Multiply *alm* by a function of *l*. The function is assumed to be zero where not defined.

### Parameters

**alm** [array] The *alm* to multiply

**fl** [array] The function (at  $l=0..fl.size-1$ ) by which *alm* must be multiplied.

**mmax** [None or int, optional] The maximum *m* defining the *alm* layout. Default: *lmax*.

**inplace** [bool, optional] If True, modify the given *alm*, otherwise make a copy before multiplying.

### Returns

**alm** [array] The modified *alm*, either a new array or a reference to input *alm*, if *inplace* is True.

## healpy.sphtfunc.pixwin

`healpy.sphtfunc.pixwin(nside, pol=False, lmax=None)`

Return the pixel window function for the given nside.

### Parameters

**nside** [int] The nside for which to return the pixel window function

**pol** [bool, optional] If True, return also the polar pixel window. Default: False

**lmax** [int, optional] Maximum l of the power spectrum (default: 3\*nside-1)

### Returns

**pw or pwT,pwP** [array or tuple of 2 arrays] The temperature pixel window function, or a tuple with both temperature and polarisation pixel window functions.

## healpy.sphtfunc.Alm

**class** `healpy.sphtfunc.Alm`

This class provides some static methods for alm index computation.

### Methods

<code>getlm(lmax[, i])</code>	Get the l and m from index and lmax.
<code>getidx(lmax, l, m)</code>	Returns index corresponding to (l,m) in an array describing alm up to lmax.
<code>getsize(lmax[, mmax])</code>	Returns the size of the array needed to store alm up to <i>lmax</i> and <i>mmax</i>
<code>getlmax(s[, mmax])</code>	Returns the lmax corresponding to a given array size.

### healpy.sphtfunc.Alm.getlm

**static** `Alm.getlm(lmax, i=None)`

Get the l and m from index and lmax.

### Parameters

**lmax** [int] The maximum l defining the alm layout

**i** [int or None] The index for which to compute the l and m. If None, the function return l and m for i=0..Alm.getsize(lmax)

### healpy.sphtfunc.Alm.getidx

**static** `Alm.getidx(lmax, l, m)`

Returns index corresponding to (l,m) in an array describing alm up to lmax.

### Parameters

**lmax** [int] The maximum l, defines the alm layout

**l** [int] The l for which to get the index

**m** [int] The m for which to get the index

**Returns**

**idx** [int] The index corresponding to (l,m)

**healpy.sphtfunc.Alm.getsize**

**static Alm.getsize(lmax, mmax=None)**

Returns the size of the array needed to store alm up to *lmax* and *mmax*

**Parameters**

**lmax** [int] The maximum l, defines the alm layout

**mmax** [int, optional] The maximum m, defines the alm layout. Default: lmax.

**Returns**

**size** [int] The size of the array needed to store alm up to lmax, mmax.

**healpy.sphtfunc.Alm.getlmax**

**static Alm.getlmax(s, mmax=None)**

Returns the lmax corresponding to a given array size.

**Parameters**

**s** [int] Size of the array

**mmax** [None or int, optional] The maximum m, defines the alm layout. Default: lmax.

**Returns**

**lmax** [int] The maximum l of the array, or -1 if it is not a valid size.

### 3.2.4 Other tools

<code>gauss_beam(fwhm[, lmax, pol])</code>	Gaussian beam window function
<code>beam2bl(beam, theta, lmax)</code>	Computes a transfer (or window) function b(l) in spherical harmonic space from its circular beam profile b(theta) in real space.
<code>bl2beam(bl, theta)</code>	Computes a circular beam profile b(theta) in real space from its transfer (or window) function b(l) in spherical harmonic space.

**healpy.sphtfunc.gauss\_beam**

`healpy.sphtfunc.gauss_beam(fwhm, lmax=512, pol=False)`

Gaussian beam window function

Computes the spherical transform of an axisymmetric gaussian beam

For a sky of underlying power spectrum  $C(l)$  observed with beam of given FWHM, the measured power spectrum will be  $C(l)_{\text{meas}} = C(l) B(l)^2$  where  $B(l)$  is given by `gaussbeam(Fwhm,Lmax)`. The polarization beam is also provided (when `pol = True`) assuming a perfectly co-polarized beam (e.g., Challinor et al 2000, astro-ph/0008228)

**Parameters**

**fw hm** [float] full width half max in radians

**lmax** [integer] ell max

**pol** [bool] if False, output has size (lmax+1) and is temperature beam if True output has size (lmax+1, 4) with components: \* temperature beam \* grad/electric polarization beam \* curl/magnetic polarization beam \* temperature \* grad beam

#### Returns

**beam** [array] beam window function [0, lmax] if dim not specified otherwise (lmax+1, 4) contains polarized beam

## healpy.sphtfunc.beam2bl

`healpy.sphtfunc.beam2bl(beam, theta, lmax)`

Computes a transfer (or window) function  $b(l)$  in spherical harmonic space from its circular beam profile  $b(\theta)$  in real space.

#### Parameters

**beam** [array] Circular beam profile  $b(\theta)$ .

**theta** [array] Radius at which the beam profile is given. Has to be given in radians with same size as beam.

**lmax** [integer] Maximum multipole moment at which to compute  $b(l)$ .

#### Returns

**bl** [array] Beam window function  $b(l)$ .

## healpy.sphtfunc.bl2beam

`healpy.sphtfunc.bl2beam(bl, theta)`

Computes a circular beam profile  $b(\theta)$  in real space from its transfer (or window) function  $b(l)$  in spherical harmonic space.

#### Parameters

**bl** [array] Window function  $b(l)$  of the beam.

**theta** [array] Radius at which the beam profile will be computed. Has to be given in radians.

#### Returns

**beam** [array] (Circular) beam profile  $b(\theta)$ .

## 3.3 visufunc – Visualisation

### 3.3.1 Map projections

---

`mollview([map, fig, rot, coord, unit, ...])`

Plot a healpix map (given as an array) in Mollweide projection.

---

`gnomview([map, fig, rot, coord, unit, ...])`

Plot a healpix map (given as an array) in Gnomonic projection.

Continued on next page

Table 11 – continued from previous page

<code>cartview([map, fig, rot, zat, coord, unit, ...])</code>	Plot a healpix map (given as an array) in Cartesian projection.
<code>orthview([map, fig, rot, coord, unit, ...])</code>	Plot a healpix map (given as an array) in Orthographic projection.

## healpy.visufunc.mollview

```
healpy.visufunc.mollview(map=None, fig=None, rot=None, coord=None, unit='', xsize=800,  
                      title='Mollweide view', nest=False, min=None, max=None,  
                      flip='astro', remove_dip=False, remove_mono=False, gal_cut=0,  
                      format='%g', format2='%g', cbar=True, cmap=None, notext=False,  
                      norm=None, hold=False, margins=None, sub=None, nlocs=2, re-  
                      turn_projected_map=False)
```

Plot a healpix map (given as an array) in Mollweide projection.

### Parameters

- map** [float, array-like or None] An array containing the map, supports masked maps, see the `ma` function. If None, will display a blank map, useful for overplotting.
- fig** [int or None, optional] The figure number to use. Default: create a new figure
- rot** [scalar or sequence, optional] Describe the rotation to apply. In the form (lon, lat, psi) (unit: degrees) : the point at longitude *lon* and latitude *lat* will be at the center. An additional rotation of angle *psi* around this direction is applied.
- coord** [sequence of character, optional] Either one of ‘G’, ‘E’ or ‘C’ to describe the coordinate system of the map, or a sequence of 2 of these to rotate the map from the first to the second coordinate system.
- unit** [str, optional] A text describing the unit of the data. Default: “”
- xsize** [int, optional] The size of the image. Default: 800
- title** [str, optional] The title of the plot. Default: ‘Mollweide view’
- nest** [bool, optional] If True, ordering scheme is NESTED. Default: False (RING)
- min** [float, optional] The minimum range value
- max** [float, optional] The maximum range value
- flip** [{‘astro’, ‘geo’}, optional] Defines the convention of projection : ‘astro’ (default, east towards left, west towards right) or ‘geo’ (east towards right, west towards left)
- remove\_dip** [bool, optional] If True, remove the dipole+monopole
- remove\_mono** [bool, optional] If True, remove the monopole
- gal\_cut** [float, scalar, optional] Symmetric galactic cut for the dipole/monopole fit. Removes points in latitude range [-gal\_cut, +gal\_cut]
- format** [str, optional] The format of the scale label. Default: ‘%g’
- format2** [str, optional] Format of the pixel value under mouse. Default: ‘%g’
- cbar** [bool, optional] Display the colorbar. Default: True
- notext** [bool, optional] If True, no text is printed around the map
- norm** [{‘hist’, ‘log’, None}] Color normalization, hist= histogram equalized color mapping, log= logarithmic color mapping, default: None (linear color mapping)

**hold** [bool, optional] If True, replace the current Axes by a MollweideAxes. use this if you want to have multiple maps on the same figure. Default: False

**sub** [int, scalar or sequence, optional] Use only a zone of the current figure (same syntax as subplot). Default: None

**margins** [None or sequence, optional] Either None, or a sequence (left,bottom,right,top) giving the margins on left,bottom,right and top of the axes. Values are relative to figure (0-1). Default: None

**return\_projected\_map** [bool] if True returns the projected map in a 2d numpy array

See also:

[gnomview](#), [cartview](#), [orthview](#), [azeqview](#)

## healpy.visufunc.gnomview

```
healpy.visufunc.gnomview(map=None, fig=None, rot=None, coord=None, unit='', xsize=200,
                         ysize=None, reso=1.5, title='Gnomonic view', nest=False, remove_dip=False,
                         remove_mono=False, gal_cut=0, min=None, max=None, flip='astro',
                         format='%.3g', cbar=True, cmap=None, norm=None, hold=False,
                         sub=None, margins=None, notext=False, return_projected_map=False,
                         no_plot=False)
```

Plot a healpix map (given as an array) in Gnomonic projection.

### Parameters

**map** [array-like] The map to project, supports masked maps, see the *ma* function. If None, use a blank map, useful for overplotting.

**fig** [None or int, optional] A figure number. Default: None= create a new figure

**rot** [scalar or sequence, optional] Describe the rotation to apply. In the form (lon, lat, psi) (unit: degrees) : the point at longitude *lon* and latitude *lat* will be at the center. An additional rotation of angle *psi* around this direction is applied.

**coord** [sequence of character, optional] Either one of ‘G’, ‘E’ or ‘C’ to describe the coordinate system of the map, or a sequence of 2 of these to rotate the map from the first to the second coordinate system.

**unit** [str, optional] A text describing the unit of the data. Default: ‘’

**xsize** [int, optional] The size of the image. Default: 200

**ysize** [None or int, optional] The size of the image. Default: None= xsize

**reso** [float, optional] Resolution (in arcmin). Default: 1.5 arcmin

**title** [str, optional] The title of the plot. Default: ‘Gnomonic view’

**nest** [bool, optional] If True, ordering scheme is NESTED. Default: False (RING)

**min** [float, scalar, optional] The minimum range value

**max** [float, scalar, optional] The maximum range value

**flip** [{‘astro’, ‘geo’}, optional] Defines the convention of projection : ‘astro’ (default, east towards left, west towards right) or ‘geo’ (east towards right, west towards left)

**remove\_dip** [bool, optional] If True, remove the dipole+monopole

**remove\_mono** [bool, optional] If True, remove the monopole

**gal\_cut** [float, scalar, optional] Symmetric galactic cut for the dipole/monopole fit. Removes points in latitude range [-gal\_cut, +gal\_cut]

**format** [str, optional] The format of the scale label. Default: ‘%g’

**hold** [bool, optional] If True, replace the current Axes by a MollweideAxes. use this if you want to have multiple maps on the same figure. Default: False

**sub** [int or sequence, optional] Use only a zone of the current figure (same syntax as subplot). Default: None

**margins** [None or sequence, optional] Either None, or a sequence (left,bottom,right,top) giving the margins on left,bottom,right and top of the axes. Values are relative to figure (0-1). Default: None

**notext: bool, optional** If True: do not add resolution info text. Default=False

**return\_projected\_map** [bool, optional] if True returns the projected map in a 2d numpy array

**no\_plot** [bool, optional] if True no figure will be created

**See also:**

*mollview, cartview, orthview, azeqview*

## healpy.visufunc.cartview

```
healpy.visufunc.cartview(map=None, fig=None, rot=None, zat=None, coord=None, unit="",
                         xsize=800, ysize=None, lonra=None, latra=None, title='Cartesian
view', nest=False, remove_dip=False, remove_mono=False, gal_cut=0,
min=None, max=None, flip='astro', format='%.3g', cbar=True,
cmap=None, norm=None, aspect=None, hold=False, sub=None,
margins=None, notext=False, return_projected_map=False)
```

Plot a healpix map (given as an array) in Cartesian projection.

### Parameters

**map** [float, array-like or None] An array containing the map, supports masked maps, see the *ma* function. If None, will display a blank map, useful for overplotting.

**fig** [int or None, optional] The figure number to use. Default: create a new figure

**rot** [scalar or sequence, optional] Describe the rotation to apply. In the form (lon, lat, psi) (unit: degrees) : the point at longitude *lon* and latitude *lat* will be at the center. An additional rotation of angle *psi* around this direction is applied.

**coord** [sequence of character, optional] Either one of ‘G’, ‘E’ or ‘C’ to describe the coordinate system of the map, or a sequence of 2 of these to rotate the map from the first to the second coordinate system.

**unit** [str, optional] A text describing the unit of the data. Default: “”

**xsize** [int, optional] The size of the image. Default: 800

**lonra** [sequence, optional] Range in longitude. Default: [-180,180]

**latra** [sequence, optional] Range in latitude. Default: [-90,90]

**title** [str, optional] The title of the plot. Default: ‘Mollweide view’

**nest** [bool, optional] If True, ordering scheme is NESTED. Default: False (RING)

**min** [float, optional] The minimum range value

**max** [float, optional] The maximum range value

**flip** [{‘astro’, ‘geo’}, optional] Defines the convention of projection : ‘astro’ (default, east towards left, west towards right) or ‘geo’ (east towards right, west towards left)

**remove\_dip** [bool, optional] If True, remove the dipole+monopole

**remove\_mono** [bool, optional] If True, remove the monopole

**gal\_cut** [float, scalar, optional] Symmetric galactic cut for the dipole/monopole fit. Removes points in latitude range [-gal\_cut, +gal\_cut]

**format** [str, optional] The format of the scale label. Default: ‘%g’

**cbar** [bool, optional] Display the colorbar. Default: True

**notext** [bool, optional] If True, no text is printed around the map

**norm** [{‘hist’, ‘log’, None}, optional] Color normalization, hist= histogram equalized color mapping, log= logarithmic color mapping, default: None (linear color mapping)

**hold** [bool, optional] If True, replace the current Axes by a CartesianAxes. use this if you want to have multiple maps on the same figure. Default: False

**sub** [int, scalar or sequence, optional] Use only a zone of the current figure (same syntax as subplot). Default: None

**margins** [None or sequence, optional] Either None, or a sequence (left,bottom,right,top) giving the margins on left,bottom,right and top of the axes. Values are relative to figure (0-1). Default: None

**return\_projected\_map** [bool] if True returns the projected map in a 2d numpy array

#### See also:

[mollview](#), [gnomview](#), [orthview](#), [azeqview](#)

### healpy.visufunc.orthview

```
healpy.visufunc.orthview(map=None, fig=None, rot=None, coord=None, unit="", xsize=800,
                         half_sky=False, title='Orthographic view', nest=False, min=None,
                         max=None, flip='astro', remove_dip=False, remove_mono=False,
                         gal_cut=0, format='%.g', format2='%.g', cbar=True, cmap=None,
                         notext=False, norm=None, hold=False, margins=None, sub=None,
                         return_projected_map=False)
```

Plot a healpix map (given as an array) in Orthographic projection.

#### Parameters

**map** [float, array-like or None] An array containing the map. If None, will display a blank map, useful for overplotting.

**fig** [int or None, optional] The figure number to use. Default: create a new figure

**rot** [scalar or sequence, optional] Describe the rotation to apply. In the form (lon, lat, psi) (unit: degrees) : the point at longitude *lon* and latitude *lat* will be at the center. An additional rotation of angle *psi* around this direction is applied.

**coord** [sequence of character, optional] Either one of ‘G’, ‘E’ or ‘C’ to describe the coordinate system of the map, or a sequence of 2 of these to rotate the map from the first to the second coordinate system.

**half\_sky** [bool, optional] Plot only one side of the sphere. Default: False

**unit** [str, optional] A text describing the unit of the data. Default: “”

**xsize** [int, optional] The size of the image. Default: 800

**title** [str, optional] The title of the plot. Default: ‘Orthographic view’

**nest** [bool, optional] If True, ordering scheme is NESTED. Default: False (RING)

**min** [float, optional] The minimum range value

**max** [float, optional] The maximum range value

**flip** [{‘astro’, ‘geo’}, optional] Defines the convention of projection : ‘astro’ (default, east towards left, west towards right) or ‘geo’ (east towards right, west towards left)

**remove\_dip** [bool, optional] If True, remove the dipole+monopole

**remove\_mono** [bool, optional] If True, remove the monopole

**gal\_cut** [float, scalar, optional] Symmetric galactic cut for the dipole/monopole fit. Removes points in latitude range [-gal\_cut, +gal\_cut]

**format** [str, optional] The format of the scale label. Default: ‘%g’

**format2** [str, optional] Format of the pixel value under mouse. Default: ‘%g’

**cbar** [bool, optional] Display the colorbar. Default: True

**notext** [bool, optional] If True, no text is printed around the map

**norm** [{‘hist’, ‘log’, None}] Color normalization, hist= histogram equalized color mapping, log= logarithmic color mapping, default: None (linear color mapping)

**hold** [bool, optional] If True, replace the current Axes by an OrthographicAxes. use this if you want to have multiple maps on the same figure. Default: False

**sub** [int, scalar or sequence, optional] Use only a zone of the current figure (same syntax as subplot). Default: None

**margins** [None or sequence, optional] Either None, or a sequence (left,bottom,right,top) giving the margins on left,bottom,right and top of the axes. Values are relative to figure (0-1). Default: None

**return\_projected\_map** [bool] if True returns the projected map in a 2d numpy array

See also:

[mollview](#), [gnomview](#), [cartview](#), [azeqview](#)

### 3.3.2 Graticules

---

<code>graticule([dpar, dmer, coord, local])</code>	Draw a graticule on the current Axes.
<code>delgraticules()</code>	Delete all graticules previously created on the Axes.

---

#### healpy.visufunc.graticule

`healpy.visufunc.graticule(dpar=None, dmer=None, coord=None, local=None, **kwds)`  
Draw a graticule on the current Axes.

##### Parameters

**dpar, dmer** [float, scalars] Interval in degrees between meridians and between parallels

**coord** [{‘E’, ‘G’, ‘C’}] The coordinate system of the graticule (make rotation if needed, using coordinate system of the map if it is defined).

**local** [bool] If True, draw a local graticule (no rotation is performed, useful for a gnomonic view, for example)

See also:

[delgraticules](#)

## Notes

Other keyword parameters will be transmitted to the projplot function.

### healpy.visufunc.delgraticules

healpy.visufunc.**delgraticules**()

Delete all graticules previously created on the Axes.

See also:

[graticule](#)

### 3.3.3 Tracing lines or points

<code>projplot(*args, **kwds)</code>	projplot is a wrapper around <code>matplotlib.Axes.plot()</code> to take into account the spherical projection.
<code>projscatter(*args, **kwds)</code>	Projscatter is a wrapper around <code>matplotlib.Axes.scatter()</code> to take into account the spherical projection.
<code>projtext(*args, **kwds)</code>	Projtext is a wrapper around <code>matplotlib.Axes.text()</code> to take into account the spherical projection.

### healpy.visufunc.projplot

healpy.visufunc.**projplot**(\*args, \*\*kwds)

projplot is a wrapper around `matplotlib.Axes.plot()` to take into account the spherical projection.

You can call this function as:

```
projplot(theta, phi)          # plot a line going through points at coord (theta, φ)
projplot(theta, phi, 'bo')    # plot 'o' in blue at coord (theta, phi)
projplot(theta, phi)          # plot a line going through points at coord
                             # (theta, φ)
projplot(theta, phi, 'bx')    # idem but with blue 'x'
```

#### Parameters

**theta, phi** [float, array-like] Coordinates of point to plot. Can be put into one 2-d array, first line is then *theta* and second line is *phi*. See *lonlat* parameter for unit.

**fmt** [str] A format string (see `matplotlib.Axes.plot()` for details)

**lonlat** [bool, optional] If True, theta and phi are interpreted as longitude and latitude in degree, otherwise, as colatitude and longitude in radian

**coord** [{‘E’, ‘G’, ‘C’, None}] The coordinate system of the points, only used if the coordinate coordinate system of the Axes has been defined and in this case, a rotation is performed

**rot** [None or sequence] rotation to be applied =(lon, lat, psi) : lon, lat will be position of the new Z axis, and psi is rotation around this axis, all in degree. if None, no rotation is performed

**direct** [bool] if True, the rotation to center the projection is not taken into account

#### See also:

`projscatter, projtext`

#### Notes

Other keywords are passed to `matplotlib.Axes.plot()`.

## healpy.visufunc.projscatter

`healpy.visufunc.projscatter(*args, **kwds)`

`Projscatter` is a wrapper around `matplotlib.Axes.scatter()` to take into account the spherical projection.

You can call this function as:

```
projscatter(theta, phi)      # plot points at coord (theta, phi)
projplot(thetaphi)          # plot points at coord (thetaphi[0], thetaphi[1])
```

#### Parameters

**theta, phi** [float, array-like] Coordinates of point to plot. Can be put into one 2-d array, first line is then *theta* and second line is *phi*. See *lonlat* parameter for unit.

**lonlat** [bool, optional] If True, theta and phi are interpreted as longitude and latitude in degree, otherwise, as colatitude and longitude in radian

**coord** [{‘E’, ‘G’, ‘C’, None}, optional] The coordinate system of the points, only used if the coordinate coordinate system of the axes has been defined and in this case, a rotation is performed

**rot** [None or sequence, optional] rotation to be applied =(lon, lat, psi) : lon, lat will be position of the new Z axis, and psi is rotation around this axis, all in degree. if None, no rotation is performed

**direct** [bool, optional] if True, the rotation to center the projection is not taken into account

#### See also:

`projplot, projtext`

#### Notes

Other keywords are passed to `matplotlib.Axes.plot()`.

## healpy.visufunc.projtext

`healpy.visufunc.projtext(*args, **kwds)`

Projtext is a wrapper around `matplotlib.Axes.text()` to take into account the spherical projection.

### Parameters

**theta, phi** [float, array-like] Coordinates of point to plot. Can be put into one 2-d array, first line is then *theta* and second line is *phi*. See *lonlat* parameter for unit.

**text** [str] The text to be displayed.

**lonlat** [bool, optional] If True, theta and phi are interpreted as longitude and latitude in degree, otherwise, as colatitude and longitude in radian

**coord** [{‘E’, ‘G’, ‘C’, None}, optional] The coordinate system of the points, only used if the coordinate coordinate system of the axes has been defined and in this case, a rotation is performed

**rot** [None or sequence, optional] rotation to be applied =(lon, lat, psi) : lon, lat will be position of the new Z axis, and psi is rotation around this axis, all in degree. if None, no rotation is performed

**direct** [bool, optional] if True, the rotation to center the projection is not taken into account

### See also:

`projplot`, `projscatter`

### Notes

Other keywords are passed to `matplotlib.Axes.text()`.

## 3.4 fitsfunc – FITS file related functions

### 3.4.1 Reading/writing maps

<code>read_map(filename[, field, dtype, nest, ...])</code>	Read a healpix map from a fits file.
<code>write_map(filename, m[, nest, dtype, ...])</code>	Writes a healpix map into a healpix file.

## healpy.fitsfunc.read\_map

`healpy.fitsfunc.read_map(filename, field=0, dtype=<class ‘numpy.float64’>, nest=False, partial=False, hdu=1, h=False, verbose=True, memmap=False)`

Read a healpix map from a fits file. Partial-sky files, if properly identified, are expanded to full size and filled with UNSEEN.

### Parameters

**filename** [str or HDU or HDUList or pathlib.Path instance] the fits file name

**field** [int or tuple of int, or None, optional] The column to read. Default: 0. By convention 0 is temperature, 1 is Q, 2 is U. Field can be a tuple to read multiple columns (0,1,2) If the fits file is a partial-sky file, field=0 corresponds to the first column after the pixel index column. If None, all columns are read in.

**dtype** [data type or list of data types, optional] Force the conversion to some type. Passing a

list allows different types for each field. In that case, the length of the list must correspond to the length of the field parameter. Default: np.float64

**nest** [bool, optional] If True return the map in NEST ordering, otherwise in RING ordering; use fits keyword ORDERING to decide whether conversion is needed or not If None, no conversion is performed.

**partial** [bool, optional] If True, fits file is assumed to be a partial-sky file with explicit indexing, if the indexing scheme cannot be determined from the header. If False, implicit indexing is assumed. Default: False. A partial sky file is one in which OBJECT=PARTIAL and INDXSCHM=EXPLICIT, and the first column is then assumed to contain pixel indices. A full sky file is one in which OBJECT=FULLSKY and INDXSCHM=IMPLICIT. At least one of these keywords must be set for the indexing scheme to be properly identified.

**hdu** [int, optional] the header number to look at (start at 0)

**h** [bool, optional] If True, return also the header. Default: False.

**verbose** [bool, optional] If True, print a number of diagnostic messages

**memmap** [bool, optional] Argument passed to astropy.io.fits.open, if True, the map is not read into memory, but only the required pixels are read when needed. Default: False.

### Returns

**m | (m0, m1, ... ) [, header]** [array or a tuple of arrays, optionally with header appended] The map(s) read from the file, and the header if *h* is True.

## healpy.fitsfunc.write\_map

```
healpy.fitsfunc.write_map(filename, m, nest=False, dtype=<class 'numpy.float32'>,  
                         fits_IDL=True, coord=None, partial=False, column_names=None,  
                         column_units=None, extra_header=(), overwrite=False)
```

Writes a healpix map into a healpix file.

### Parameters

**filename** [str] the fits file name

**m** [array or sequence of 3 arrays] the map to write. Possibly a sequence of 3 maps of same size. They will be considered as I, Q, U maps. Supports masked maps, see the *ma* function.

**nest** [bool, optional] If True, ordering scheme is assumed to be NESTED, otherwise, RING. Default: RING. The map ordering is not modified by this function, the input map array should already be in the desired ordering (run *ud\_grade* beforehand).

**fits\_IDL** [bool, optional] If True, reshapes columns in rows of 1024, otherwise all the data will go in one column. Default: True

**coord** [str] The coordinate system, typically ‘E’ for Ecliptic, ‘G’ for Galactic or ‘C’ for Celestial (equatorial)

**partial** [bool, optional] If True, fits file is written as a partial-sky file with explicit indexing. Otherwise, implicit indexing is used. Default: False.

**column\_names** [str or list] Column name or list of column names, if None here the default column names based on the number of columns: 1 : “TEMPERATURE”, 2 : [“Q\_POLARISATION”, “U\_POLARISATION”], 3 : [“TEMPERATURE”, “Q\_POLARISATION”, “U\_POLARISATION”], 6 : [“II”, “IQ”, “IU”, “QQ”, “QU”, “UU”] COLUMN\_1, COLUMN\_2... otherwise (FITS is 1-based)

**column\_units** [str or list] Units for each column, or same units for all columns.

**extra\_header** [list] Extra records to add to FITS header.

**dtype: np.dtype or list of np.dtypes, optional** The datatype in which the columns will be stored. Will be converted internally from the numpy datatype to the fits convention. If a list, the length must correspond to the number of map arrays. Default: np.float32.

**overwrite** [bool, optional] If True, existing file is silently overwritten. Otherwise trying to write an existing file raises an OSError (IOError for Python 2).

### 3.4.2 Reading/writing alm

---

<code>read_alm(filename[, hdu, return_mmax])</code>	Read alm from a fits file.
<code>write_alm(filename, alms[, out_dtype, lmax, ...])</code>	Write alms to a fits file.

---

#### healpy.fitsfunc.read\_alm

`healpy.fitsfunc.read_alm(filename, hdu=1, return_mmax=False)`

Read alm from a fits file.

In the fits file, the alm are written with explicit index scheme,  $\text{index} = l^{**}2+l+m+1$ , while healpix cxx uses  $\text{index} = m*(2*lmax+1-m)/2+l$ . The conversion is done in this function.

##### Parameters

**filename** [str or HDUList or HDU or pathlib.Path instance] The name of the fits file to read

**hdu** [int, or tuple of int, optional] The header to read. Start at 0. Default: hdu=1

**return\_mmax** [bool, optional] If true, both the alms and mmax is returned in a tuple. Default: return\_mmax=False

##### Returns

**alms[, mmax]** [complex array or tuple of a complex array and an int] The alms read from the file and optionally mmax read from the file

#### healpy.fitsfunc.write\_alm

`healpy.fitsfunc.write_alm(filename, alms, out_dtype=None, lmax=-1, mmax=-1, mmax_in=-1, overwrite=False)`

Write alms to a fits file.

In the fits file the alms are written with explicit index scheme,  $\text{index} = l^{**}l + 1 + m + 1$ , possibly out of order. By default write\_alm makes a table with the same precision as the alms. If specified, the lmax and mmax parameters truncate the input data to include only alms for which  $l \leq lmax$  and  $m \leq mmax$ .

##### Parameters

**filename** [str] The filename of the output fits file

**alms** [array, complex or list of arrays] A complex ndarray holding the alms,  $\text{index} = m*(2*lmax+1-m)/2+l$ , see Alm.getidx

**lmax** [int, optional] The maximum l in the output file

**mmax** [int, optional] The maximum m in the output file

**out\_dtype** [data type, optional] data type in the output file (must be a numpy dtype). Default: `alms.real.dtype`

**mmax\_in** [int, optional] maximum m in the input array

### 3.4.3 Reading/writing cl

<code>read_cl(filename[, dtype, h])</code>	Reads Cl from a healpix file, as IDL fits2cl.
<code>write_cl(filename, cl[, dtype, overwrite])</code>	Writes Cl into a healpix file, as IDL cl2fits.

#### healpy.fitsfunc.read\_cl

`healpy.fitsfunc.read_cl(filename, dtype=<class 'numpy.float64'>, h=False)`

Reads Cl from a healpix file, as IDL fits2cl.

##### Parameters

**filename** [str or HDUList or HDU or pathlib.Path instance] the fits file name

**dtype** [data type, optional] the data type of the returned array

##### Returns

**cl** [array] the cl array

#### healpy.fitsfunc.write\_cl

`healpy.fitsfunc.write_cl(filename, cl, dtype=<class 'numpy.float64'>, overwrite=False)`

Writes Cl into a healpix file, as IDL cl2fits.

##### Parameters

**filename** [str] the fits file name

**cl** [array] the cl array to write to file

**overwrite** [bool, optional] If True, existing file is silently overwritten. Otherwise trying to write an existing file raises an OSError (IOError for Python 2).

### 3.4.4 Reading/writing column in fits file

---

`mrdfits`

---

`mwrdfits`

---

### 3.4.5 Helper

---

`getformat(t)`

---

Get the FITS convention format string of data type t.

---

#### healpy.fitsfunc.getformat

`healpy.fitsfunc.getformat(t)`

Get the FITS convention format string of data type t.

##### Parameters

**t** [data type] The data type for which the FITS type is requested

**Returns**

**fits\_type** [str or None] The FITS string code describing the data type, or None if unknown type.

## 3.5 Pixel querying routines

<code>query_disc</code> (nside, vec, radius[, inclusive, ...])	Returns pixels whose centers lie within the disk defined by <i>vec</i> and <i>radius</i> (in radians) (if <i>inclusive</i> is False), or which overlap with this disk (if <i>inclusive</i> is True).
<code>query_polygon</code> (nside, vertices[, inclusive, ...])	Returns the pixels whose centers lie within the convex polygon defined by the <i>vertices</i> array (if <i>inclusive</i> is False), or which overlap with this polygon (if <i>inclusive</i> is True).
<code>query_strip</code> (nside, theta1, theta2[, ...])	Returns pixels whose centers lie within the colatitude range defined by <i>theta1</i> and <i>theta2</i> (if <i>inclusive</i> is False), or which overlap with this region (if <i>inclusive</i> is True).
<code>boundaries</code> (nside, pix[, step, nest])	Returns an array containing vectors to the boundary of the nominated pixel.

### 3.5.1 healpy.query\_disc

`healpy.query_disc(nside, vec, radius, inclusive=False, fact=4, nest=False, ndarray buff=None)`

Returns pixels whose centers lie within the disk defined by *vec* and *radius* (in radians) (if *inclusive* is False), or which overlap with this disk (if *inclusive* is True).

**Parameters**

**nside** [int] The nside of the Healpix map.

**vec** [float, sequence of 3 elements] The coordinates of unit vector defining the disk center.

**radius** [float] The radius (in radians) of the disk

**inclusive** [bool, optional] If False, return the exact set of pixels whose pixel centers lie within the disk; if True, return all pixels that overlap with the disk, and maybe a few more. Default: False

**fact** [int, optional] Only used when inclusive=True. The overlapping test will be done at the resolution fact\*nside. For NESTED ordering, fact must be a power of 2, less than 2\*\*30, else it can be any positive integer. Default: 4.

**nest: bool, optional** if True, assume NESTED pixel ordering, otherwise, RING pixel ordering

**buff: int array, optional** if provided, this numpy array is used to contain the return values and must be at least long enough to do so

**Returns**

**ipix** [int, array] The pixels which lie within the given disk.

### 3.5.2 healpy.query\_polygon

`healpy.query_polygon(nside, vertices, inclusive=False, fact=4, nest=False, ndarray buff=None)`

Returns the pixels whose centers lie within the convex polygon defined by the *vertices* array (if *inclusive* is False), or which overlap with this polygon (if *inclusive* is True).

### Parameters

**nside** [int] The nside of the Healpix map.

**vertices** [float, array-like] Vertex array containing the vertices of the polygon, shape (N, 3).

**inclusive** [bool, optional] If False, return the exact set of pixels whose pixel centers lie within the polygon; if True, return all pixels that overlap with the polygon, and maybe a few more. Default: False.

**fact** [int, optional] Only used when inclusive=True. The overlapping test will be done at the resolution fact\*nside. For NESTED ordering, fact must be a power of 2, less than  $2^{**30}$ , else it can be any positive integer. Default: 4.

**nest: bool, optional** if True, assume NESTED pixel ordering, otherwise, RING pixel ordering

**buff: int array, optional** if provided, this numpy array is used to contain the return values and must be at least long enough to do so

### Returns

**ipix** [int, array] The pixels which lie within the given polygon.

## 3.5.3 healpy.query\_strip

`healpy.query_strip(nside, theta1, theta2, inclusive=False, nest=False, ndarray buff=None)`

Returns pixels whose centers lie within the colatitude range defined by *theta1* and *theta2* (if *inclusive* is False), or which overlap with this region (if *inclusive* is True). If  $\text{theta1} < \text{theta2}$ , the region between both angles is considered, otherwise the regions  $0 < \text{theta} < \text{theta2}$  and  $\text{theta1} < \text{theta} < \pi$ .

### Parameters

**nside** [int] The nside of the Healpix map.

**theta1** [float] First colatitude (radians)

**theta2** [float] Second colatitude (radians)

**inclusive ; bool** If False, return the exact set of pixels whose pixels centers lie within the region; if True, return all pixels that overlap with the region.

**nest: bool, optional** if True, assume NESTED pixel ordering, otherwise, RING pixel ordering

**buff: int array, optional** if provided, this numpy array is used to contain the return values and must be at least long enough to do so

### Returns

**ipix** [int, array] The pixels which lie within the given strip.

## 3.5.4 healpy.boundaries

`healpy.boundaries(nside, pix, step=1, nest=False)`

Returns an array containing vectors to the boundary of the nominated pixel.

The returned array has shape (3, 4\*step), the elements of which are the x,y,z positions on the unit sphere of the pixel boundary. In order to get vector positions for just the corners, specify step=1.

### Parameters

**nside** [int] The nside of the Healpix map.

**pix** [int] Pixel identifier

**step** [int, optional] Number of elements for each side of the pixel.

**nest** [bool, optional] if True, assume NESTED pixel ordering, otherwise, RING pixel ordering

#### Returns

**boundary** [float, array] x,y,z for positions on the boundary of the pixel

## 3.6 rotator – Rotation and geometry functions

### 3.6.1 Rotation

<code>Rotator([rot, coord, inv, deg, eulertype])</code>	Rotation operator, including astronomical coordinate systems.
<code>rotateVector(rotmat, vec[, vy, vz, do_rot])</code>	Rotate a vector (or a list of vectors) using the rotation matrix given as first argument.
<code>rotateDirection(rotmat, theta[, phi, ...])</code>	Rotate the vector described by angles theta,phi using the rotation matrix given as first argument.

#### healpy.rotator.Rotator

```
class healpy.rotator.Rotator(rot=None, coord=None, inv=None, deg=True, eulertype='ZYX')
```

Rotation operator, including astronomical coordinate systems.

This class provides tools for spherical rotations. It is meant to be used in the healpy library for plotting, and for this reason reflects the convention used in the Healpix IDL library.

#### Parameters

**rot** [None or sequence] Describe the rotation by its euler angle. See `euler_matrix_new()`.

**coord** [None or sequence of str] Describe the coordinate system transform. If `rot` is also given, the coordinate transform is applied first, and then the rotation.

**inv** [bool] If True, the inverse rotation is defined. (Default: False)

**deg** [bool] If True, angles are assumed to be in degree. (Default: True)

**eulertype** [str] The Euler angle convention used. See `euler_matrix_new()`.

#### Examples

```
>>> r = Rotator(coord=['G', 'E']) # Transforms galactic to ecliptic coordinates
>>> theta_gal, phi_gal = np.pi/2., 0.
>>> theta_ecl, phi_ecl = r(theta_gal, phi_gal) # Apply the conversion
>>> print(theta_ecl)
1.66742286715
>>> print(phi_ecl)
-1.62596400306
>>> theta_ecl, phi_ecl = Rotator(coord='ge')(theta_gal, phi_gal) # In one line
>>> print(theta_ecl)
1.66742286715
>>> print(phi_ecl)
-1.62596400306
>>> vec_gal = np.array([1, 0, 0]) #Using vectors
```

(continues on next page)

(continued from previous page)

```
>>> vec_ecl = r(vec_gal)
>>> print(vec_ecl)
[-0.05488249 -0.99382103 -0.09647625]
```

## Attributes

- mat** The matrix representing the rotation.
- coordin** The input coordinate system.
- coordout** The output coordinate system.
- coordinstr** The input coordinate system in str.
- coordoutstr** The output coordinate system in str.
- rots** The sequence of rots defining the rotation.
- coords** The sequence of coords defining the rotation.

## Methods

<code>I(*args, **kwds)</code>	Rotate the given vector or direction using the inverse matrix.
<code>__call__(*args, **kwds)</code>	Use the rotator to rotate either spherical coordinates (theta, phi) or a vector (x,y,z).
<code>angle_ref(*args, **kwds)</code>	Compute the angle between transverse reference direction of initial and final frames
<code>do_rot(i)</code>	Returns True if rotation is not (close to) identity.
<code>rotate_alm(alm[, lmax, mmax])</code>	Rotate Alms with the transform defined in the Rotator object
<code>rotate_map_alm(m[, use_pixel_weights, ...])</code>	Rotate a HEALPix map to a new reference frame in spherical harmonics space
<code>rotate_map_pixel(m)</code>	Rotate a HEALPix map to a new reference frame in pixel space

### healpy.rotator.Rotator.I

`Rotator.I (*args, **kwds)`Rotate the given vector or direction using the inverse matrix. `rot.I(vec) <==> rot(vec,inv=True)`

### healpy.rotator.Rotator.\_\_call\_\_

`Rotator.__call__(*args, **kwds)`

Use the rotator to rotate either spherical coordinates (theta, phi) or a vector (x,y,z). You can use lonlat keyword to use longitude, latitude (in degree) instead of theta, phi (in radian). In this case, returns longitude, latitude in degree.

Accepted forms:

`r(x,y,z)` # x,y,z either scalars or arrays  
`r(theta,phi)` # theta, phi scalars or arrays  
`r(lon,lat,lonlat=True)` # lon, lat scalars or arrays  
`r(vec)` # vec 1-D array with 3 elements, or 2-D array 3xN  
`r(direction)` # direction 1-D array with 2 elements, or 2xN array

**Parameters**

**vec\_or\_dir** [array or multiple arrays] The direction to rotate. See above for accepted formats.

**lonlat** [bool, optional] If True, assumes the input direction is longitude/latitude in degrees. Otherwise, assumes co-latitude/longitude in radians. Default: False

**inv** [bool, optional] If True, applies the inverse rotation. Default: False.

**healpy.rotator.Rotator.angle\_ref**

`Rotator.angle_ref(*args, **kwds)`

Compute the angle between transverse reference direction of initial and final frames

For example, if angle of polarisation is psi in initial frame, it will be psi+angle\_ref in final frame.

**Parameters**

**dir\_or\_vec** [array] Direction or vector (see `Rotator.__call__`)

**lonlat: bool, optional** If True, assume input is longitude,latitude in degrees. Otherwise, theta,phi in radian. Default: False

**inv** [bool, optional] If True, use the inverse transforms. Default: False

**Returns**

**angle** [float, scalar or array] Angle in radian (a scalar or an array if input is a sequence of direction/vector)

**healpy.rotator.Rotator.do\_rot**

`Rotator.do_rot(i)`

Returns True if rotation is not (close to) identity.

**healpy.rotator.Rotator.rotate\_alm**

`Rotator.rotate_alm(alm, lmax=None, mmax=None)`

Rotate Alms with the transform defined in the Rotator object

see the docstring of the `rotate_alm` function defined in the healpy package, this function **returns** the rotated alms, does not rotate in place

**healpy.rotator.Rotator.rotate\_map\_alm**

`Rotator.rotate_map_alm(m, use_pixel_weights=True, lmax=None, mmax=None)`

Rotate a HEALPix map to a new reference frame in spherical harmonics space

This is generally the best strategy to rotate/change reference frame of maps. If the input map is band-limited, i.e. it can be represented exactly by a spherical harmonics transform under a specific lmax, the map rotation will be invertible.

**Parameters**

**m** [np.ndarray] Input map, 1 map is considered I, 2 maps:[Q,U], 3 maps:[I,Q,U]

**use\_pixel\_weights** [bool, optional] Use pixel weights in map2alm

**Returns**

**m\_rotated** [np.ndarray] Map in the new reference frame

## healpy.rotator.Rotator.rotate\_map\_pixel

`Rotator.rotate_map_pixel(m)`

Rotate a HEALPix map to a new reference frame in pixel space

It is generally better to rotate in spherical harmonics space, see the `rotate_map_alm`s method. A case where pixel space rotation is better is for heavily masked maps where the spherical harmonics transform is not well defined. This function first rotates the pixels centers of the new reference frame to the original reference frame, then uses `hp.get_interp_val` to interpolate bilinearly the pixel values, finally fixes Q and U polarization by the modification to the psi angle caused by the Rotator using `Rotator.angle_ref`. Due to interpolation, this function generally suppresses the signal at high angular scales.

**Parameters**

**m** [np.ndarray] Input map, 1 map is considered I, 2 maps:[Q,U], 3 maps:[I,Q,U]

**Returns**

**m\_rotated** [np.ndarray] Map in the new reference frame

**get\_inverse**

## healpy.rotator.rotateVector

`healpy.rotator.rotateVector(rotmat, vec, vy=None, vz=None, do_rot=True)`

Rotate a vector (or a list of vectors) using the rotation matrix given as first argument.

**Parameters**

**rotmat** [float, array-like shape (3,3)] The rotation matrix

**vec** [float, scalar or array-like] The vector to transform (shape (3,) or (3,N)), or x component (scalar or shape (N,)) if vy and vz are given

**vy** [float, scalar or array-like, optional] The y component of the vector (scalar or shape (N,))

**vz** [float, scalar or array-like, optional] The z component of the vector (scalar or shape (N,))

**do\_rot** [bool, optional] if True, really perform the operation, if False do nothing.

**Returns**

**vec** [float, array] The component of the rotated vector(s).

**See also:**

`Rotator`

## healpy.rotator.rotateDirection

`healpy.rotator.rotateDirection(rotmat, theta, phi=None, do_rot=True, lonlat=False)`

Rotate the vector described by angles theta,phi using the rotation matrix given as first argument.

**Parameters**

**rotmat** [float, array-like shape (3,3)] The rotation matrix

**theta** [float, scalar or array-like] The angle theta (scalar or shape (N,)) or both angles (scalar or shape (2, N)) if phi is not given.

**phi** [float, scalar or array-like, optionnal] The angle phi (scalar or shape (N,)).

**do\_rot** [bool, optional] if True, really perform the operation, if False do nothing.

**lonlat** [bool] If True, input angles are assumed to be longitude and latitude in degree, otherwise, they are co-latitude and longitude in radians.

#### Returns

**angles** [float, array] The angles of describing the rotated vector(s).

#### See also:

*Rotator*

## 3.6.2 Geometrical helpers

<code>vec2dir(vec[, vy, vz, lonlat])</code>	Transform a vector to angle given by theta,phi.
<code>dir2vec(theta[, phi, lonlat])</code>	Transform a direction theta,phi to a unit vector.
<code>angdist(dir1, dir2[, lonlat])</code>	Returns the angular distance between dir1 and dir2.

### healpy.rotator.vec2dir

`healpy.rotator.vec2dir(vec, vy=None, vz=None, lonlat=False)`

Transform a vector to angle given by theta,phi.

#### Parameters

**vec** [float, scalar or array-like] The vector to transform (shape (3,) or (3,N)), or x component (scalar or shape (N,)) if vy and vz are given

**vy** [float, scalar or array-like, optional] The y component of the vector (scalar or shape (N,))

**vz** [float, scalar or array-like, optional] The z component of the vector (scalar or shape (N,))

**lonlat** [bool, optional] If True, return angles will be longitude and latitude in degree, otherwise, angles will be longitude and co-latitude in radians (default)

#### Returns

**angles** [float, array] The angles (unit depending on *lonlat*) in an array of shape (2,) (if scalar input) or (2, N)

#### See also:

*dir2vec()*, *pixelfunc.ang2vec()*, *pixelfunc.vec2ang()*

### healpy.rotator.dir2vec

`healpy.rotator.dir2vec(theta, phi=None, lonlat=False)`

Transform a direction theta,phi to a unit vector.

#### Parameters

**theta** [float, scalar or array-like] The angle theta (scalar or shape (N,)) or both angles (scalar or shape (2, N)) if phi is not given.

**phi** [float, scalar or array-like, optionnal] The angle phi (scalar or shape (N,)).

**lonlat** [bool] If True, input angles are assumed to be longitude and latitude in degree, otherwise, they are co-latitude and longitude in radians.

#### Returns

**vec** [array] The vector(s) corresponding to given angles, shape is (3,) or (3, N).

#### See also:

`vec2dir()`, `pixelfunc.ang2vec()`, `pixelfunc.vec2ang()`

## healpy.rotator.angdist

`healpy.rotator.angdist(dir1, dir2, lonlat=False)`

Returns the angular distance between dir1 and dir2.

#### Parameters

**dir1, dir2** [float, array-like] The directions between which computing the angular distance. Angular if `len(dir) == 2` or vector if `len(dir) == 3`. See `lonlat` for unit

**lonlat** [bool, scalar or sequence] If True, angles are assumed to be longitude and latitude in degree, otherwise they are interpreted as colatitude and longitude in radian. If a sequence, `lonlat[0]` applies to `dir1` and `lonlat[1]` applies to `dir2`.

#### Returns

**angles** [float, scalar or array-like] The angle(s) between `dir1` and `dir2` in radian.

#### Examples

```
>>> import healpy as hp
>>> hp.rotator.angdist([.2, 0], [.2, 1e-6])
array([-1.98669331e-07])
```

## 3.7 projector – Spherical projections

### 3.7.1 Basic classes

<code>SphericalProj</code> ([rot, coord, flipconv])	This class defines functions for spherical projection.
<code>GnomonicProj</code> ([rot, coord, xsize, ysize, reso])	This class provides class methods for Gnomonic projection.
<code>MollweideProj</code> ([rot, coord, xsize])	This class provides class methods for Mollweide projection.
<code>CartesianProj</code> ([rot, coord, xsize, ysize, ...])	This class provides class methods for Cartesian projection.

## healpy.projector.SphericalProj

`class healpy.projector.SphericalProj(rot=None, coord=None, flipconv=None, **kwds)`

This class defines functions for spherical projection.

This class contains class method for spherical projection computation. It should not be instantiated. It should be inherited from and methods should be overloaded for desired projection.

### Attributes

**arrayinfo** Dictionary with information on the projection array

### Methods

<code>ang2xy(theta[, phi, lonlat, direct])</code>	From angular direction to position in the projection plane (%s).
<code>get_center([lonlat])</code>	Get the center of the projection.
<code>get_extent()</code>	Get the extension of the projection plane.
<code>get_fov()</code>	Get the field of view in degree of the plane of projection
<code>ij2xy([i, j])</code>	From image array indices to position in projection plane (%s).
<code>projmap(map, vec2pix_func[, rot, coord])</code>	Create an array containing the projection of the map.
<code>set_flip(flipconv)</code>	flipconv is either ‘astro’ or ‘geo’.
<code>vec2xy(vx[, vy, vz, direct])</code>	From unit vector direction to position in the projection plane (%s).
<code>xy2ang(x[, y, lonlat, direct])</code>	From position in the projection plane to angular direction (%s).
<code>xy2ij(x[, y])</code>	From position in the projection plane to image array index (%s).
<code>xy2vec(x[, y, direct])</code>	From position in the projection plane to unit vector direction (%s).

### healpy.projector.SphericalProj.ang2xy

SphericalProj. **ang2xy** (*theta, phi=None, lonlat=False, direct=False*)

From angular direction to position in the projection plane (%s).

#### Input:

- theta: if phi is None, theta[0] contains theta, theta[1] contains phi
- phi : if phi is not None, theta,phi are direction
- lonlat: if True, angle are assumed in degree, and longitude, latitude
- flipconv is either ‘astro’ or ‘geo’. None will be default.

#### Return:

- x, y: position in %s plane.

### healpy.projector.SphericalProj.get\_center

SphericalProj. **get\_center** (*lonlat=False*)

Get the center of the projection.

#### Input:

- lonlat [if True, will return longitude and latitude in degree,] otherwise, theta and phi in radian

#### Return:

- theta,phi or lonlat depending on lonlat keyword

### `healpy.projector.SphericalProj.get_extent`

`SphericalProj.get_extent()`

Get the extension of the projection plane.

**Return:** extent = (left,right,bottom,top)

### `healpy.projector.SphericalProj.get_fov`

`SphericalProj.get_fov()`

Get the field of view in degree of the plane of projection

**Return:** fov: the diameter in radian of the field of view

### `healpy.projector.SphericalProj.ij2xy`

`SphericalProj.ij2xy(i=None, j=None)`

From image array indices to position in projection plane (%s).

**Input:**

- if i and j are None, generate arrays of i and j as input
- i : if j is None, i[0], j[1] define array indices in %s image.
- j : if defined, i,j define array indices in image.
- projinfo : additional projection information.

**Return:**

- x,y : position in projection plane.

### `healpy.projector.SphericalProj.projmap`

`SphericalProj.projmap(map, vec2pix_func, rot=None, coord=None)`

Create an array containing the projection of the map.

**Input:**

- vec2pix\_func: a function taking theta,phi and returning pixel number
- **map: an array containing the spherical map to project**, the pixelisation is described by vec2pix\_func

**Return:**

- a 2D array with the projection of the map.

Note: the Projector must contain information on the array.

## healpy.projector.SphericalProj.set\_flip

SphericalProj.**set\_flip** (*flipconv*)

*flipconv* is either ‘astro’ or ‘geo’. None will be default.

With ‘astro’, east is toward left and west toward right. It is the opposite for ‘geo’

## healpy.projector.SphericalProj.vec2xy

SphericalProj.**vec2xy** (*vx*, *vy*=*None*, *vz*=*None*, *direct=False*)

From unit vector direction to position in the projection plane (%s).

### Input:

- *vx*: if *vy* and *vz* are *None*, *vx[0]*,*vx[1]*,*vx[2]* defines the unit vector.
- *vy*,*vz*: if defined, *vx*,*vy*,*vz* define the unit vector
- *lonlat*: if True, angle are assumed in degree, and longitude, latitude
- *flipconv* is either ‘astro’ or ‘geo’. None will be default.

### Return:

- *x*, *y*: position in %s plane.

## healpy.projector.SphericalProj.xy2ang

SphericalProj.**xy2ang** (*x*, *y*=*None*, *lonlat=False*, *direct=False*)

From position in the projection plane to angular direction (%s).

### Input:

- *x* : if *y* is *None*, *x[0]*, *x[1]* define the position in %s plane.
- *y* : if defined, *x*,*y* define the position in projection plane.
- *lonlat*: if True, angle are assumed in degree, and longitude, latitude
- *flipconv* is either ‘astro’ or ‘geo’. None will be default.

### Return:

- *theta*, *phi* : angular direction.

## healpy.projector.SphericalProj.xy2ij

SphericalProj.**xy2ij** (*x*, *y*=*None*)

From position in the projection plane to image array index (%s).

### Input:

- *x* : if *y* is *None*, *x[0]*, *x[1]* define the position in %s plane.
- *y* : if defined, *x*,*y* define the position in projection plane.
- *projinfo* : additional projection information.

### Return:

- *i,j* : image array indices.

## healpy.projector.SphericalProj.xy2vec

SphericalProj.**xy2vec**(*x*, *y*=None, *direct*=False)

From position in the projection plane to unit vector direction (%s).

### Input:

- *x* : if *y* is None, *x*[0], *x*[1] define the position in %s plane.
- *y* : if defined, *x*,*y* define the position in projection plane.
- lonlat: if True, angle are assumed in degree, and longitude, latitude
- flipconv is either ‘astro’ or ‘geo’. None will be default.

### Return:

- theta, phi : angular direction.

<code>get_proj_plane_info</code>	
<code>mkcoord</code>	
<code>set_proj_plane_info</code>	

## healpy.projector.GnomonicProj

**class** healpy.projector.**GnomonicProj**(*rot*=None, *coord*=None, *xsize*=None, *ysize*=None, *reso*=None, \*\**kwds*)

This class provides class methods for Gnomonic projection.

### Attributes

**arrayinfo** Dictionary with information on the projection array

### Methods

<code>ang2xy(theta[, phi, lonlat, direct])</code>	From angular direction to position in the projection plane (Gnomonic).
<code>get_center([lonlat])</code>	Get the center of the projection.
<code>get_extent()</code>	Get the extension of the projection plane.
<code>get_fov()</code>	Get the field of view in degree of the plane of projection
<code>ij2xy([i, j])</code>	From image array indices to position in projection plane (Gnomonic).
<code>projmap(map, vec2pix_func[, rot, coord])</code>	Create an array containing the projection of the map.
<code>set_flip(flipconv)</code>	flipconv is either ‘astro’ or ‘geo’.
<code>vec2xy(vx[, vy, vz, direct])</code>	From angular direction to position in the projection plane (Gnomonic).
<code>xy2ang(x[, y, lonlat, direct])</code>	From position in the projection plane to angular direction (Gnomonic).
<code>xy2ij(x[, y])</code>	From position in the projection plane to image array index (Gnomonic).
<code>xy2vec(x[, y, direct])</code>	From position in the projection plane to unit vector direction (Gnomonic).

## healpy.projector.GnomonicProj.ang2xy

GnomonicProj.ang2xy (*theta, phi=None, lonlat=False, direct=False*)

From angular direction to position in the projection plane (Gnomonic).

### Input:

- theta: if phi is None, theta[0] contains theta, theta[1] contains phi
- phi : if phi is not None, theta,phi are direction
- lonlat: if True, angle are assumed in degree, and longitude, latitude
- flipconv is either ‘astro’ or ‘geo’. None will be default.

### Return:

- x, y: position in Gnomonic plane.

## healpy.projector.GnomonicProj.get\_center

GnomonicProj.get\_center (*lonlat=False*)

Get the center of the projection.

### Input:

- lonlat [if True, will return longitude and latitude in degree,] otherwise, theta and phi in radian

### Return:

- theta,phi or lonlat depending on lonlat keyword

## healpy.projector.GnomonicProj.get\_extent

GnomonicProj.get\_extent ()

Get the extension of the projection plane.

**Return:** extent = (left,right,bottom,top)

## healpy.projector.GnomonicProj.get\_fov

GnomonicProj.get\_fov ()

Get the field of view in degree of the plane of projection

**Return:** fov: the diameter in radian of the field of view

## healpy.projector.GnomonicProj.ij2xy

GnomonicProj.ij2xy (*i=None, j=None*)

From image array indices to position in projection plane (Gnomonic).

### Input:

- if i and j are None, generate arrays of i and j as input
- i : if j is None, i[0], j[1] define array indices in Gnomonic image.
- j : if defined, i,j define array indices in image.

- projinfo : additional projection information.

**Return:**

- x,y : position in projection plane.

### healpy.projector.GnomonicProj.projmap

GnomonicProj.**projmap** (*map*, *vec2pix\_func*, *rot=None*, *coord=None*)

Create an array containing the projection of the map.

**Input:**

- vec2pix\_func: a function taking theta,phi and returning pixel number
- **map:** an array containing the spherical map to project, the pixelisation is described by vec2pix\_func

**Return:**

- a 2D array with the projection of the map.

Note: the Projector must contain information on the array.

### healpy.projector.GnomonicProj.set\_flip

GnomonicProj.**set\_flip** (*flipconv*)

flipconv is either ‘astro’ or ‘geo’. None will be default.

With ‘astro’, east is toward left and west toward right. It is the opposite for ‘geo’

### healpy.projector.GnomonicProj.vec2xy

GnomonicProj.**vec2xy** (*vx*, *vy=None*, *vz=None*, *direct=False*)

From angular direction to position in the projection plane (Gnomonic).

**Input:**

- theta: if phi is None, theta[0] contains theta, theta[1] contains phi
- phi : if phi is not None, theta,phi are direction
- lonlat: if True, angle are assumed in degree, and longitude, latitude
- flipconv is either ‘astro’ or ‘geo’. None will be default.

**Return:**

- x, y: position in Gnomonic plane.

### healpy.projector.GnomonicProj.xy2ang

GnomonicProj.**xy2ang** (*x*, *y=None*, *lonlat=False*, *direct=False*)

From position in the projection plane to angular direction (Gnomonic).

**Input:**

- x : if y is None, x[0], x[1] define the position in Gnomonic plane.

- y : if defined, x,y define the position in projection plane.
- lonlat: if True, angle are assumed in degree, and longitude, latitude
- flipconv is either ‘astro’ or ‘geo’. None will be default.

**Return:**

- theta, phi : angular direction.

**healpy.projector.GnomonicProj.xy2ij****GnomonicProj.xy2ij**(x, y=None)

From position in the projection plane to image array index (Gnomonic).

**Input:**

- x : if y is None, x[0], x[1] define the position in Gnomonic plane.
- y : if defined, x,y define the position in projection plane.
- projinfo : additional projection information.

**Return:**

- i,j : image array indices.

**healpy.projector.GnomonicProj.xy2vec****GnomonicProj.xy2vec**(x, y=None, direct=False)

From position in the projection plane to unit vector direction (Gnomonic).

**Input:**

- x : if y is None, x[0], x[1] define the position in Gnomonic plane.
- y : if defined, x,y define the position in projection plane.
- lonlat: if True, angle are assumed in degree, and longitude, latitude
- flipconv is either ‘astro’ or ‘geo’. None will be default.

**Return:**

- theta, phi : angular direction.

get_proj_plane_info	
mkcoord	
set_proj_plane_info	

**healpy.projector.MollweideProj****class** `healpy.projector.MollweideProj`(rot=None, coord=None, xsize=800, \*\*kwds)

This class provides class methods for Mollweide projection.

**Attributes****arrayinfo** Dictionary with information on the projection array

## Methods

<code>ang2xy(theta[, phi, lonlat, direct])</code>	From angular direction to position in the projection plane (Mollweide).
<code>get_center([lonlat])</code>	Get the center of the projection.
<code>get_extent()</code>	Get the extension of the projection plane.
<code>get_fov()</code>	Get the field of view in degree of the plane of projection
<code>ij2xy([i, j])</code>	From image array indices to position in projection plane (Mollweide).
<code>projmap(map, vec2pix_func[, rot, coord])</code>	Create an array containing the projection of the map.
<code>set_flip(flipconv)</code>	flipconv is either ‘astro’ or ‘geo’.
<code>vec2xy(vx[, vy, vz, direct])</code>	From unit vector direction to position in the projection plane (Mollweide).
<code>xy2ang(x[, y, lonlat, direct])</code>	From position in the projection plane to angular direction (Mollweide).
<code>xy2ij(x[, y])</code>	From position in the projection plane to image array index (Mollweide).
<code>xy2vec(x[, y, direct])</code>	From position in the projection plane to unit vector direction (Mollweide).

### healpy.projector.MollweideProj.ang2xy

`MollweideProj.ang2xy(theta, phi=None, lonlat=False, direct=False)`

From angular direction to position in the projection plane (Mollweide).

#### Input:

- theta: if phi is None, theta[0] contains theta, theta[1] contains phi
- phi : if phi is not None, theta,phi are direction
- lonlat: if True, angle are assumed in degree, and longitude, latitude
- flipconv is either ‘astro’ or ‘geo’. None will be default.

#### Return:

- x, y: position in Mollweide plane.

### healpy.projector.MollweideProj.get\_center

`MollweideProj.get_center(lonlat=False)`

Get the center of the projection.

#### Input:

- lonlat [if True, will return longitude and latitude in degree,] otherwise, theta and phi in radian

#### Return:

- theta,phi or lonlat depending on lonlat keyword

### healpy.projector.MollweideProj.get\_extent

MollweideProj.**get\_extent**()  
Get the extension of the projection plane.  
**Return:** extent = (left,right,bottom,top)

### healpy.projector.MollweideProj.get\_fov

MollweideProj.**get\_fov**()  
Get the field of view in degree of the plane of projection  
**Return:** fov: the diameter in radian of the field of view

### healpy.projector.MollweideProj.ij2xy

MollweideProj.**ij2xy** (*i=None, j=None*)  
From image array indices to position in projection plane (Mollweide).

#### Input:

- if i and j are None, generate arrays of i and j as input
- i : if j is None, i[0], j[1] define array indices in Mollweide image.
- j : if defined, i,j define array indices in image.
- projinfo : additional projection information.

#### Return:

- x,y : position in projection plane.

### healpy.projector.MollweideProj.projmap

MollweideProj.**projmap** (*map, vec2pix\_func, rot=None, coord=None*)  
Create an array containing the projection of the map.

#### Input:

- vec2pix\_func: a function taking theta,phi and returning pixel number
- **map: an array containing the spherical map to project**, the pixelisation is described by vec2pix\_func

#### Return:

- a 2D array with the projection of the map.

Note: the Projector must contain information on the array.

### healpy.projector.MollweideProj.set\_flip

MollweideProj.**set\_flip** (*flipconv*)  
flipconv is either ‘astro’ or ‘geo’. None will be default.  
With ‘astro’, east is toward left and west toward right. It is the opposite for ‘geo’

## healpy.projector.MollweideProj.vec2xy

MollweideProj.**vec2xy** (*vx*, *vy*=*None*, *vz*=*None*, *direct*=*False*)

From unit vector direction to position in the projection plane (Mollweide).

### Input:

- *vx*: if *vy* and *vz* are *None*, *vx*[0],*vx*[1],*vx*[2] defines the unit vector.
- *vy*,*vz*: if defined, *vx*,*vy*,*vz* define the unit vector
- *lonlat*: if True, angle are assumed in degree, and longitude, latitude
- *flipconv* is either ‘astro’ or ‘geo’. *None* will be default.

### Return:

- *x*, *y*: position in Mollweide plane.

## healpy.projector.MollweideProj.xy2ang

MollweideProj.**xy2ang** (*x*, *y*=*None*, *lonlat*=*False*, *direct*=*False*)

From position in the projection plane to angular direction (Mollweide).

### Input:

- *x* : if *y* is *None*, *x*[0], *x*[1] define the position in Mollweide plane.
- *y* : if defined, *x*,*y* define the position in projection plane.
- *lonlat*: if True, angle are assumed in degree, and longitude, latitude
- *flipconv* is either ‘astro’ or ‘geo’. *None* will be default.

### Return:

- *theta*, *phi* : angular direction.

## healpy.projector.MollweideProj.xy2ij

MollweideProj.**xy2ij** (*x*, *y*=*None*)

From position in the projection plane to image array index (Mollweide).

### Input:

- *x* : if *y* is *None*, *x*[0], *x*[1] define the position in Mollweide plane.
- *y* : if defined, *x*,*y* define the position in projection plane.
- *projinfo* : additional projection information.

### Return:

- *i*,*j* : image array indices.

## healpy.projector.MollweideProj.xy2vec

MollweideProj.**xy2vec** (*x*, *y*=*None*, *direct*=*False*)

From position in the projection plane to unit vector direction (Mollweide).

### Input:

- x : if y is None, x[0], x[1] define the position in Mollweide plane.
- y : if defined, x,y define the position in projection plane.
- lonlat: if True, angle are assumed in degree, and longitude, latitude
- flipconv is either ‘astro’ or ‘geo’. None will be default.

**Return:**

- theta, phi : angular direction.

<code>get_proj_plane_info</code>	
<code>mkcoord</code>	
<code>set_proj_plane_info</code>	

**healpy.projector.CartesianProj**

```
class healpy.projector.CartesianProj(rot=None, coord=None, xsize=800, ysize=None,
                                      lonra=None, latra=None, **kwds)
```

This class provides class methods for Cartesian projection.

**Attributes**

`arrayinfo` Dictionary with information on the projection array

**Methods**

<code>ang2xy(theta[, phi, lonlat, direct])</code>	From angular direction to position in the projection plane (Cartesian).
<code>get_center([lonlat])</code>	Get the center of the projection.
<code>get_extent()</code>	Get the extension of the projection plane.
<code>get_fov()</code>	Get the field of view in degree of the plane of projection
<code>ij2xy([i, j])</code>	From image array indices to position in projection plane (Cartesian).
<code>projmap(map, vec2pix_func[, rot, coord])</code>	Create an array containing the projection of the map.
<code>set_flip(flipconv)</code>	flipconv is either ‘astro’ or ‘geo’.
<code>vec2xy(vx[, vy, vz, direct])</code>	From unit vector direction to position in the projection plane (Cartesian).
<code>xy2ang(x[, y, lonlat, direct])</code>	From position in the projection plane to angular direction (Cartesian).
<code>xy2ij(x[, y])</code>	From position in the projection plane to image array index (Cartesian).
<code>xy2vec(x[, y, direct])</code>	From position in the projection plane to unit vector direction (Cartesian).

**healpy.projector.CartesianProj.ang2xy**

`CartesianProj.ang2xy(theta, phi=None, lonlat=False, direct=False)`

From angular direction to position in the projection plane (Cartesian).

**Input:**

- theta: if phi is None, theta[0] contains theta, theta[1] contains phi

- phi : if phi is not None, theta,phi are direction
- lonlat: if True, angle are assumed in degree, and longitude, latitude
- flipconv is either ‘astro’ or ‘geo’. None will be default.

**Return:**

- x, y: position in Cartesian plane.

## healpy.projector.CartesianProj.get\_center

CartesianProj.**get\_center** (*lonlat=False*)

Get the center of the projection.

**Input:**

- **lonlat** [if True, will return longitude and latitude in degree,] otherwise, theta and phi in radian

**Return:**

- theta,phi or lonlat depending on lonlat keyword

## healpy.projector.CartesianProj.get\_extent

CartesianProj.**get\_extent** ()

Get the extension of the projection plane.

**Return:** extent = (left,right,bottom,top)

## healpy.projector.CartesianProj.get\_fov

CartesianProj.**get\_fov** ()

Get the field of view in degree of the plane of projection

**Return:** fov: the diameter in radian of the field of view

## healpy.projector.CartesianProj.ij2xy

CartesianProj.**ij2xy** (*i=None, j=None*)

From image array indices to position in projection plane (Cartesian).

**Input:**

- if i and j are None, generate arrays of i and j as input
- i : if j is None, i[0], j[1] define array indices in Cartesian image.
- j : if defined, i,j define array indices in image.
- projinfo : additional projection information.

**Return:**

- x,y : position in projection plane.

## healpy.projector.CartesianProj.projmap

CartesianProj.**projmap** (*map*, *vec2pix\_func*, *rot=None*, *coord=None*)

Create an array containing the projection of the map.

### Input:

- *vec2pix\_func*: a function taking theta,phi and returning pixel number
- **map**: an array containing the spherical map to project, the pixelisation is described by *vec2pix\_func*

### Return:

- a 2D array with the projection of the map.

Note: the Projector must contain information on the array.

## healpy.projector.CartesianProj.set\_flip

CartesianProj.**set\_flip** (*flipconv*)

*flipconv* is either ‘astro’ or ‘geo’. None will be default.

With ‘astro’, east is toward left and west toward right. It is the opposite for ‘geo’

## healpy.projector.CartesianProj.vec2xy

CartesianProj.**vec2xy** (*vx*, *vy=None*, *vz=None*, *direct=False*)

From unit vector direction to position in the projection plane (Cartesian).

### Input:

- *vx*: if *vy* and *vz* are None, *vx[0]*,*vx[1]*,*vx[2]* defines the unit vector.
- *vy*,*vz*: if defined, *vx*,*vy*,*vz* define the unit vector
- *lonlat*: if True, angle are assumed in degree, and longitude, latitude
- *flipconv* is either ‘astro’ or ‘geo’. None will be default.

### Return:

- *x*, *y*: position in Cartesian plane.

## healpy.projector.CartesianProj.xy2ang

CartesianProj.**xy2ang** (*x*, *y=None*, *lonlat=False*, *direct=False*)

From position in the projection plane to angular direction (Cartesian).

### Input:

- *x* : if *y* is None, *x[0]*, *x[1]* define the position in Cartesian plane.
- *y* : if defined, *x*,*y* define the position in projection plane.
- *lonlat*: if True, angle are assumed in degree, and longitude, latitude
- *flipconv* is either ‘astro’ or ‘geo’. None will be default.

### Return:

- theta, phi : angular direction.

### healpy.projector.CartesianProj.xy2ij

CartesianProj.**xy2ij**(*x, y=None*)

From position in the projection plane to image array index (Cartesian).

#### Input:

- *x* : if *y* is None, *x*[0], *x*[1] define the position in Cartesian plane.
- *y* : if defined, *x,y* define the position in projection plane.
- *projinfo* : additional projection information.

#### Return:

- *i,j* : image array indices.

### healpy.projector.CartesianProj.xy2vec

CartesianProj.**xy2vec**(*x, y=None, direct=False*)

From position in the projection plane to unit vector direction (Cartesian).

#### Input:

- *x* : if *y* is None, *x*[0], *x*[1] define the position in Cartesian plane.
- *y* : if defined, *x,y* define the position in projection plane.
- lonlat: if True, angle are assumed in degree, and longitude, latitude
- flipconv is either ‘astro’ or ‘geo’. None will be default.

#### Return:

- theta, phi : angular direction.

<code>get_proj_plane_info</code>	
<code>mkcoord</code>	
<code>set_proj_plane_info</code>	

## 3.8 zoomtool – Interactive visualisation

### 3.8.1 Interactive map visualization

---

`mollzoom([map, fig, rot, coord, unit, ...])`

Interactive mollweide plot with zoomed gnomview.

---

#### healpy.zoomtool.mollzoom

healpy.zoomtool.**mollzoom**(*map=None, fig=None, rot=None, coord=None, unit=”, xscale=800, title='Mollweide view', nest=False, min=None, max=None, flip='astro', remove\_dip=False, remove\_mono=False, gal\_cut=0, format='%g', cmap=None, norm=None, hold=False, margins=None, sub=None*)

Interactive mollweide plot with zoomed gnomview.

# CHAPTER 4

---

## Indices and tables

---

- genindex
- modindex
- search

## 4.1 License

### 4.1.1 Licenses

#### Healpy License

Healpy is licensed under the GNU General Public License.

**GNU GENERAL PUBLIC LICENSE** Version 2, June 1991

**Copyright (C) 1989, 1991 Free Software Foundation, Inc.** 51 Franklin St, Fifth Floor, Boston, MA 02110-1301 USA

Everyone is permitted to copy and distribute verbatim copies of this license document, but changing it is not allowed.

#### Preamble

The licenses for most software are designed to take away your

freedom to share and change it. By contrast, the GNU General Public License is intended to guarantee your freedom to share and change free software—to make sure the software is free for all its users. This General Public License applies to most of the Free Software Foundation's software and to any other program whose authors commit to using it. (Some other Free Software Foundation software is covered by the GNU Library General Public License instead.) You can apply it to your programs, too.

When we speak of free software, we are referring to freedom, not

price. Our General Public Licenses are designed to make sure that you have the freedom to distribute copies of free software (and charge for this service if you wish), that you receive source code or can get it if you want it, that you can change the software or use pieces of it in new free programs; and that you know you can do these things.

To protect your rights, we need to make restrictions that forbid anyone to deny you these rights or to ask you to surrender the rights. These restrictions translate to certain responsibilities for you if you distribute copies of the software, or if you modify it.

For example, if you distribute copies of such a program, whether gratis or for a fee, you must give the recipients all the rights that you have. You must make sure that they, too, receive or can get the source code. And you must show them these terms so they know their rights.

We protect your rights with two steps: (1) copyright the software, and  
(2) offer you this license which gives you legal permission to copy, distribute and/or modify the software.

Also, for each author's protection and ours, we want to make certain that everyone understands that there is no warranty for this free software. If the software is modified by someone else and passed on, we want its recipients to know that what they have is not the original, so that any problems introduced by others will not reflect on the original authors' reputations.

Finally, any free program is threatened constantly by software patents. We wish to avoid the danger that redistributors of a free program will individually obtain patent licenses, in effect making the program proprietary. To prevent this, we have made it clear that any patent must be licensed for everyone's free use or not licensed at all.

The precise terms and conditions for copying, distribution and modification follow.

#### GNU GENERAL PUBLIC LICENSE

#### TERMS AND CONDITIONS FOR COPYING, DISTRIBUTION AND MODIFICATION

0. This License applies to any program or other work which contains a notice placed by the copyright holder saying it may be distributed under the terms of this General Public License. The "Program", below, refers to any such program or work, and a "work based on the Program" means either the Program or any derivative work under copyright law: that is to say, a work containing the Program or a portion of it, either verbatim or with modifications and/or translated into another language. (Hereinafter, translation is included without limitation in the term "modification".) Each licensee is addressed as "you".

Activities other than copying, distribution and modification are not covered by this License; they are outside its scope. The act of running the Program is not restricted, and the output from the Program is covered only if its contents constitute a work based on the Program (independent of having been made by running the Program). Whether that is true depends on what the Program does.

1. You may copy and distribute verbatim copies of the Program's source code as you receive it, in any medium, provided that you conspicuously and appropriately publish on each copy an appropriate copyright notice and disclaimer of warranty; keep intact all the notices that refer to this License and to the absence of any warranty; and give any other recipients of the Program a copy of this License along with the Program.

You may charge a fee for the physical act of transferring a copy, and you may at your option offer warranty protection in exchange for a fee.

2. You may modify your copy or copies of the Program or any portion of it, thus forming a work based on the Program, and copy and distribute such modifications or work under the terms of Section 1 above, provided that you also meet all of these conditions:

- a) You must cause the modified files to carry prominent notices stating that you changed the files and the date of any change.
- b) You must cause any work that you distribute or publish, that in whole or in part contains or is derived from the Program or any part thereof, to be licensed as a whole at no charge to all third parties under the terms of this License.
- c) If the modified program normally reads commands interactively when run, you must cause it, when started running for such interactive use in the most ordinary way, to print or display an announcement including an appropriate copyright notice and a notice that there is no warranty (or else, saying that you provide a warranty) and that users may redistribute the program under these conditions, and telling the user how to view a copy of this License. (Exception: if the Program itself is interactive but does not normally print such an announcement, your work based on the Program is not required to print an announcement.)

These requirements apply to the modified work as a whole. If identifiable sections of that work are not derived from the Program, and can be reasonably considered independent and separate works in themselves, then this License, and its terms, do not apply to those sections when you distribute them as separate works. But when you distribute the same sections as part of a whole which is a work based on the Program, the distribution of the whole must be on the terms of this License, whose permissions for other licensees extend to the entire whole, and thus to each and every part regardless of who wrote it.

Thus, it is not the intent of this section to claim rights or contest your rights to work written entirely by you; rather, the intent is to exercise the right to control the distribution of derivative or collective works based on the Program.

In addition, mere aggregation of another work not based on the Program with the Program (or with a work based on the Program) on a volume of a storage or distribution medium does not bring the other work under the scope of this License.

3. You may copy and distribute the Program (or a work based on it,

under Section 2) in object code or executable form under the terms of Sections 1 and 2 above provided that you also do one of the following:

- a) Accompany it with the complete corresponding machine-readable source code, which must be distributed under the terms of Sections 1 and 2 above on a medium customarily used for software interchange; or,
- b) Accompany it with a written offer, valid for at least three years, to give any third party, for a charge no more than your cost of physically performing source distribution, a complete machine-readable copy of the corresponding source code, to be distributed under the terms of Sections 1 and 2 above on a medium customarily used for software interchange; or,
- c) Accompany it with the information you received as to the offer to distribute corresponding source code. (This alternative is allowed only for noncommercial distribution and only if you received the program in object code or executable form with such an offer, in accord with Subsection b above.)

The source code for a work means the preferred form of the work for making modifications to it. For an executable work, complete source code means all the source code for all modules it contains, plus any associated interface definition files, plus the scripts used to control compilation and installation of the executable. However, as a special exception, the source code distributed need not include anything that is normally distributed (in either source or binary form) with the major components (compiler, kernel, and so on) of the operating system on which the executable runs, unless that component itself accompanies the executable.

If distribution of executable or object code is made by offering access to copy from a designated place, then offering equivalent access to copy the source code from the same place counts as distribution of the source code, even though third parties are not compelled to copy the source along with the object code.

4. You may not copy, modify, sublicense, or distribute the Program

except as expressly provided under this License. Any attempt otherwise to copy, modify, sublicense or distribute the Program is void, and will automatically terminate your rights under this License. However, parties who have received

copies, or rights, from you under this License will not have their licenses terminated so long as such parties remain in full compliance.

5. You are not required to accept this License, since you have not

signed it. However, nothing else grants you permission to modify or distribute the Program or its derivative works. These actions are prohibited by law if you do not accept this License. Therefore, by modifying or distributing the Program (or any work based on the Program), you indicate your acceptance of this License to do so, and all its terms and conditions for copying, distributing or modifying the Program or works based on it.

6. Each time you redistribute the Program (or any work based on the

Program), the recipient automatically receives a license from the original licensor to copy, distribute or modify the Program subject to these terms and conditions. You may not impose any further restrictions on the recipients' exercise of the rights granted herein. You are not responsible for enforcing compliance by third parties to this License.

7. If, as a consequence of a court judgment or allegation of patent

infringement or for any other reason (not limited to patent issues), conditions are imposed on you (whether by court order, agreement or otherwise) that contradict the conditions of this License, they do not excuse you from the conditions of this License. If you cannot distribute so as to satisfy simultaneously your obligations under this License and any other pertinent obligations, then as a consequence you may not distribute the Program at all. For example, if a patent license would not permit royalty-free redistribution of the Program by all those who receive copies directly or indirectly through you, then the only way you could satisfy both it and this License would be to refrain entirely from distribution of the Program.

If any portion of this section is held invalid or unenforceable under any particular circumstance, the balance of the section is intended to apply and the section as a whole is intended to apply in other circumstances.

It is not the purpose of this section to induce you to infringe any patents or other property right claims or to contest validity of any such claims; this section has the sole purpose of protecting the integrity of the free software distribution system, which is implemented by public license practices. Many people have made generous contributions to the wide range of software distributed through that system in reliance on consistent application of that system; it is up to the author/donor to decide if he or she is willing to distribute software through any other system and a licensee cannot impose that choice.

This section is intended to make thoroughly clear what is believed to be a consequence of the rest of this License.

8. If the distribution and/or use of the Program is restricted in

certain countries either by patents or by copyrighted interfaces, the original copyright holder who places the Program under this License may add an explicit geographical distribution limitation excluding those countries, so that distribution is permitted only in or among countries not thus excluded. In such case, this License incorporates the limitation as if written in the body of this License.

9. The Free Software Foundation may publish revised and/or new versions

of the General Public License from time to time. Such new versions will be similar in spirit to the present version, but may differ in detail to address new problems or concerns.

Each version is given a distinguishing version number. If the Program specifies a version number of this License which applies to it and "any later version", you have the option of following the terms and conditions either of that version or of any later version published by the Free Software Foundation. If the Program does not specify a version number of this License, you may choose any version ever published by the Free Software Foundation.

10. If you wish to incorporate parts of the Program into other free

programs whose distribution conditions are different, write to the author to ask for permission. For software which is copyrighted by the Free Software Foundation, write to the Free Software Foundation; we sometimes make exceptions for this. Our decision will be guided by the two goals of preserving the free status of all derivatives of our free software and of promoting the sharing and reuse of software generally.

## NO WARRANTY

### 11. BECAUSE THE PROGRAM IS LICENSED FREE OF CHARGE, THERE IS NO WARRANTY

FOR THE PROGRAM, TO THE EXTENT PERMITTED BY APPLICABLE LAW. EXCEPT WHEN OTHERWISE STATED IN WRITING THE COPYRIGHT HOLDERS AND/OR OTHER PARTIES PROVIDE THE PROGRAM "AS IS" WITHOUT WARRANTY OF ANY KIND, EITHER EXPRESSED OR IMPLIED, INCLUDING, BUT NOT LIMITED TO, THE IMPLIED WARRANTIES OF MERCHANTABILITY AND FITNESS FOR A PARTICULAR PURPOSE. THE ENTIRE RISK AS TO THE QUALITY AND PERFORMANCE OF THE PROGRAM IS WITH YOU. SHOULD THE PROGRAM PROVE DEFECTIVE, YOU ASSUME THE COST OF ALL NECESSARY SERVICING, REPAIR OR CORRECTION.

### 12. IN NO EVENT UNLESS REQUIRED BY APPLICABLE LAW OR AGREED TO IN WRITING

WILL ANY COPYRIGHT HOLDER, OR ANY OTHER PARTY WHO MAY MODIFY AND/OR REDISTRIBUTE THE PROGRAM AS PERMITTED ABOVE, BE LIABLE TO YOU FOR DAMAGES, INCLUDING ANY GENERAL, SPECIAL, INCIDENTAL OR CONSEQUENTIAL DAMAGES ARISING OUT OF THE USE OR INABILITY TO USE THE PROGRAM (INCLUDING BUT NOT LIMITED TO LOSS OF DATA OR DATA BEING RENDERED INACCURATE OR LOSSES SUSTAINED BY YOU OR THIRD PARTIES OR A FAILURE OF THE PROGRAM TO OPERATE WITH ANY OTHER PROGRAMS), EVEN IF SUCH HOLDER OR OTHER PARTY HAS BEEN ADVISED OF THE POSSIBILITY OF SUCH DAMAGES.

## END OF TERMS AND CONDITIONS

### How to Apply These Terms to Your New Programs

If you develop a new program, and you want it to be of the greatest possible use to the public, the best way to achieve this is to make it free software which everyone can redistribute and change under these terms.

To do so, attach the following notices to the program. It is safest

to attach them to the start of each source file to most effectively convey the exclusion of warranty; and each file should have at least the "copyright" line and a pointer to where the full notice is found.

<one line to give the program's name and a brief idea of what it does.> Copyright (C) <year> <name of author>

This program is free software; you can redistribute it and/or modify it under the terms of the GNU General Public License as published by the Free Software Foundation; either version 2 of the License, or (at your option) any later version.

This program is distributed in the hope that it will be useful, but WITHOUT ANY WARRANTY; without even the implied warranty of MERCHANTABILITY or FITNESS FOR A PARTICULAR PURPOSE. See the GNU General Public License for more details.

You should have received a copy of the GNU General Public License along with this program; if not, write to the Free Software Foundation, Inc., 51 Franklin St, Fifth Floor, Boston, MA 02110-1301 USA

Also add information on how to contact you by electronic and paper mail.

If the program is interactive, make it output a short notice like this when it starts in an interactive mode:

Gnomovision version 69, Copyright (C) year name of author Gnomovision comes with ABSOLUTELY NO WARRANTY; for details type 'show w'. This is free software, and you are welcome to redistribute it under certain conditions; type 'show c' for details.

The hypothetical commands 'show w' and 'show c' should show the appropriate parts of the General Public License. Of course, the commands you use may be called something other than 'show w' and 'show c'; they could even be mouse-clicks or menu items—whatever suits your program.

You should also get your employer (if you work as a programmer) or your school, if any, to sign a “copyright disclaimer” for the program, if necessary. Here is a sample; alter the names:

Yoyodyne, Inc., hereby disclaims all copyright interest in the program ‘Gnomovision’ (which makes  
passes at compilers) written by James Hacker.

<signature of Ty Coon>, 1 April 1989 Ty Coon, President of Vice

This General Public License does not permit incorporating your program into proprietary programs. If your program is a subroutine library, you may consider it more useful to permit linking proprietary applications with the library. If this is what you want to do, use the GNU Library General Public License instead of this License.

### Symbols

`__call__()` (*healpy.rotator.Rotator method*), 56

### A

`Alm` (*class in healpy.sphfunc*), 39  
`alm2cl()` (*in module healpy.sphfunc*), 37  
`alm2map()` (*in module healpy.sphfunc*), 34  
`alm2map_der1()` (*in module healpy.sphfunc*), 35  
`almxfl()` (*in module healpy.sphfunc*), 38  
`anafast()` (*in module healpy.sphfunc*), 32  
`ang2pix()` (*in module healpy.pixelfunc*), 13  
`ang2vec()` (*in module healpy.pixelfunc*), 15  
`ang2xy()` (*healpy.projector.CartesianProj method*), 71  
`ang2xy()` (*healpy.projector.GnomonicProj method*), 65  
`ang2xy()` (*healpy.projector.MollweideProj method*), 68  
`ang2xy()` (*healpy.projector.SphericalProj method*), 61  
`angdist()` (*in module healpy.rotator*), 60  
`angle_ref()` (*healpy.rotator.Rotator method*), 57

### B

`beam2bl()` (*in module healpy.sphfunc*), 41  
`bl2beam()` (*in module healpy.sphfunc*), 41  
`boundaries()` (*in module healpy*), 54

### C

`CartesianProj` (*class in healpy.projector*), 71  
`cartview()` (*in module healpy.visufunc*), 44

### D

`delgraticules()` (*in module healpy.visufunc*), 47  
`dir2vec()` (*in module healpy.rotator*), 59  
`do_rot()` (*healpy.rotator.Rotator method*), 57

### F

`fit_dipole()` (*in module healpy.pixelfunc*), 30  
`fit_monopole()` (*in module healpy.pixelfunc*), 31

### G

`gauss_beam()` (*in module healpy.sphfunc*), 40

`get_all_neighbours()` (*in module healpy.pixelfunc*), 15  
`get_center()` (*healpy.projector.CartesianProj method*), 72  
`get_center()` (*healpy.projector.GnomonicProj method*), 65  
`get_center()` (*healpy.projector.MollweideProj method*), 68  
`get_center()` (*healpy.projector.SphericalProj method*), 61  
`get_extent()` (*healpy.projector.CartesianProj method*), 72  
`get_extent()` (*healpy.projector.GnomonicProj method*), 65  
`get_extent()` (*healpy.projector.MollweideProj method*), 69  
`get_extent()` (*healpy.projector.SphericalProj method*), 62  
`get_fov()` (*healpy.projector.CartesianProj method*), 72  
`get_fov()` (*healpy.projector.GnomonicProj method*), 65  
`get_fov()` (*healpy.projector.MollweideProj method*), 69  
`get_fov()` (*healpy.projector.SphericalProj method*), 62  
`get_interp_val()` (*in module healpy.pixelfunc*), 17  
`get_interp_weights()` (*in module healpy.pixelfunc*), 16  
`get_map_size()` (*in module healpy.pixelfunc*), 25  
`get_min_valid_nside()` (*in module healpy.pixelfunc*), 26  
`get_nside()` (*in module healpy.pixelfunc*), 27  
`getformat()` (*in module healpy.fitsfunc*), 52  
`getidx()` (*healpy.sphfunc.Alm static method*), 39  
`getlm()` (*healpy.sphfunc.Alm static method*), 39  
`getlmax()` (*healpy.sphfunc.Alm static method*), 40  
`getsize()` (*healpy.sphfunc.Alm static method*), 40  
`GnomonicProj` (*class in healpy.projector*), 64  
`gnomview()` (*in module healpy.visufunc*), 43

graticule() (*in module healpy.visufunc*), 46

## I

I() (*healpy.rotator.Rotator method*), 56  
ij2xy() (*healpy.projector.CartesianProj method*), 72  
ij2xy() (*healpy.projector.GnomonicProj method*), 65  
ij2xy() (*healpy.projector.MollweideProj method*), 69  
ij2xy() (*healpy.projector.SphericalProj method*), 62  
isnpixok() (*in module healpy.pixelfunc*), 25  
isinsideok() (*in module healpy.pixelfunc*), 24

## M

ma() (*in module healpy.pixelfunc*), 29  
map2alm() (*in module healpy.sphfunc*), 33  
maptype() (*in module healpy.pixelfunc*), 27  
mask\_bad() (*in module healpy.pixelfunc*), 28  
mask\_good() (*in module healpy.pixelfunc*), 29  
max\_pixrad() (*in module healpy.pixelfunc*), 24  
mollview() (*in module healpy.visufunc*), 42  
MollweideProj (*class in healpy.projector*), 67  
mollzoom() (*in module healpy.zoomtool*), 74

## N

nest2ring() (*in module healpy.pixelfunc*), 18  
npix2nside() (*in module healpy.pixelfunc*), 21  
nside2npix() (*in module healpy.pixelfunc*), 20  
nside2order() (*in module healpy.pixelfunc*), 22  
nside2pixarea() (*in module healpy.pixelfunc*), 23  
nside2resol() (*in module healpy.pixelfunc*), 23

## O

order2nside() (*in module healpy.pixelfunc*), 22  
orthview() (*in module healpy.visufunc*), 45

## P

pix2ang() (*in module healpy.pixelfunc*), 11  
pix2vec() (*in module healpy.pixelfunc*), 12  
pixwin() (*in module healpy.sphfunc*), 39  
projmap() (*healpy.projector.CartesianProj method*), 73  
projmap() (*healpy.projector.GnomonicProj method*), 66  
projmap() (*healpy.projector.MollweideProj method*), 69  
projmap() (*healpy.projector.SphericalProj method*), 62  
projplot() (*in module healpy.visufunc*), 47  
projscatter() (*in module healpy.visufunc*), 48  
projtext() (*in module healpy.visufunc*), 49

## Q

query\_disc() (*in module healpy*), 53  
query\_polygon() (*in module healpy*), 53

query\_strip() (*in module healpy*), 54

## R

read\_alm() (*in module healpy.fitsfunc*), 51  
read\_cl() (*in module healpy.fitsfunc*), 52  
read\_map() (*in module healpy.fitsfunc*), 49  
remove\_dipole() (*in module healpy.pixelfunc*), 31  
remove\_monopole() (*in module healpy.pixelfunc*), 32  
reorder() (*in module healpy.pixelfunc*), 19  
ring2nest() (*in module healpy.pixelfunc*), 18  
rotate\_alm() (*healpy.rotator.Rotator method*), 57  
rotate\_map\_alsms() (*healpy.rotator.Rotator method*), 57  
rotate\_map\_pixel() (*healpy.rotator.Rotator method*), 58  
rotateDirection() (*in module healpy.rotator*), 58  
rotateVector() (*in module healpy.rotator*), 58  
Rotator (*class in healpy.rotator*), 55

## S

set\_flip() (*healpy.projector.CartesianProj method*), 73  
set\_flip() (*healpy.projector.GnomonicProj method*), 66  
set\_flip() (*healpy.projector.MollweideProj method*), 69  
set\_flip() (*healpy.projector.SphericalProj method*), 63  
smoothalm() (*in module healpy.sphfunc*), 37  
smoothing() (*in module healpy.sphfunc*), 36  
SphericalProj (*class in healpy.projector*), 60  
synalm() (*in module healpy.sphfunc*), 38  
synfast() (*in module healpy.sphfunc*), 34

## U

ud\_grade() (*in module healpy.pixelfunc*), 27  
UNSEEN (*in module healpy.pixelfunc*), 28

## V

vec2ang() (*in module healpy.pixelfunc*), 14  
vec2dir() (*in module healpy.rotator*), 59  
vec2pix() (*in module healpy.pixelfunc*), 14  
vec2xy() (*healpy.projector.CartesianProj method*), 73  
vec2xy() (*healpy.projector.GnomonicProj method*), 66  
vec2xy() (*healpy.projector.MollweideProj method*), 70  
vec2xy() (*healpy.projector.SphericalProj method*), 63

## W

write\_alm() (*in module healpy.fitsfunc*), 51  
write\_cl() (*in module healpy.fitsfunc*), 52  
write\_map() (*in module healpy.fitsfunc*), 50

## X

`xy2ang()` (*healpy.projector.CartesianProj method*), 73  
`xy2ang()` (*healpy.projector.GnomonicProj method*), 66  
`xy2ang()` (*healpy.projector.MollweideProj method*), 70  
`xy2ang()` (*healpy.projector.SphericalProj method*), 63  
`xy2ij()` (*healpy.projector.CartesianProj method*), 74  
`xy2ij()` (*healpy.projector.GnomonicProj method*), 67  
`xy2ij()` (*healpy.projector.MollweideProj method*), 70  
`xy2ij()` (*healpy.projector.SphericalProj method*), 63  
`xy2vec()` (*healpy.projector.CartesianProj method*), 74  
`xy2vec()` (*healpy.projector.GnomonicProj method*), 67  
`xy2vec()` (*healpy.projector.MollweideProj method*), 70  
`xy2vec()` (*healpy.projector.SphericalProj method*), 64