

# Tools for data analyses in Cosmology

- Aula 5 -

Camila Novaes

Observatório Nacional

May 23, 2017

Healpy

Previously on `healpy` class ...

- Writing fits files -

## `healpy.fitsfunc.write_map`

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

## `healpy.fitsfunc.mwrfits`

```
healpy.fitsfunc.mwrfits(filename, data, hdu=1, colnames=None, keys=None)
```

ERROR -

Previously on `healpy` class ...

- Writing fits files -  
solving partially (and properly) the  
problem

In `fitsfunc.py` substitute:

<code>clobber</code>	→	<code>overwrite</code>
<code>pf.new_table</code>	→	<code>pf.BinTableHDU.from_columns</code>
<code>header.update</code>	→	<code>header.set</code>

## `healpy.fitsfunc.write_map`

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

## `healpy.fitsfunc.mwrfits`

```
healpy.fitsfunc.mwrfits(filename, data, hdu=1, colnames=None, keys=None)
```



Previously on `healpy` class ...

How to use:

```
In [95]: import pyfits
...:
...: x = data[10]
...: hdu = pyfits.PrimaryHDU(x)
...: hdulist = pyfits.HDUList([hdu])
...: hdulist.writeto('new.fits')
...: hdulist.close()
...:
...: hdulist = pyfits.open('new.fits')
...: data2 = hdulist[0].data
...: |
```

Still from VISUALIZATION tools

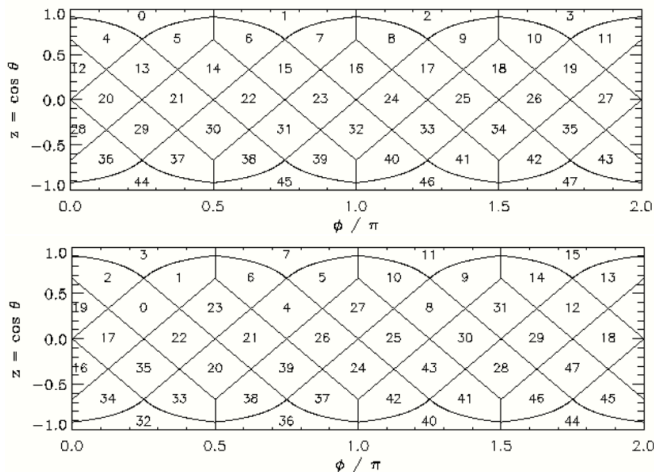
Tracing lines or points

**projplot** (\*args, \*\*kwargs)

**projscatter** (\*args, \*\*kwargs)

**projtext** (\*args, \*\*kwargs)

## Conversion between NESTED and RING schemes



**healpy.pixelfunc.ring2nest**    **healpy.pixelfunc.nest2ring**

`healpy.pixelfunc.ring2nest(nside, ipix)`

`healpy.pixelfunc.nest2ring(nside, ipix)`

## Conversion between NESTED and RING schemes

How to use:

RING  $\Rightarrow$  NEST

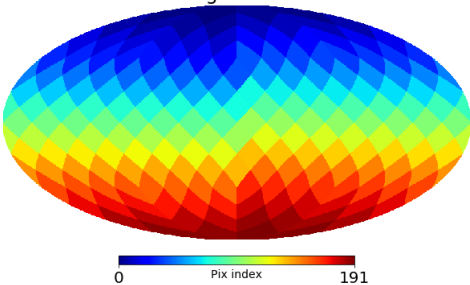
```
In [34]: Nside = 4
...: pix_ring = 25
...: pix_nest = hp.ring2nest(Nside, pix_ring)
...: print('pix_nest =', pix_nest)
...:
pix_nest = 9

In [35]: all_pix_ring = np.arange(192)
...: all_pix_nest = hp.ring2nest(Nside, all_pix_ring)
...: hp.mollview(all_pix_nest, title='Nest scheme',
unit='Pix index')
...: pyplot.savefig('mollweide_view_nest.png')
...:
```

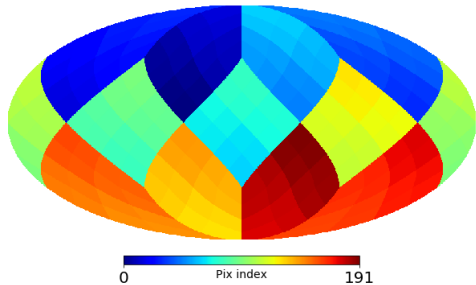


## Conversion between NESTED and RING schemes

Ring scheme



Nest scheme



## Conversion between NESTED and RING schemes

How to use:

NEST  $\Rightarrow$  RING

```
In [36]: # Nside = 4
...: print('pix_nest =', pix_nest)
...: pix_ring = hp.nest2ring(Nside, pix_nest)
...: print('pix_ring =', pix_ring)
...:
pix_nest = 9
pix_ring = 25

In [37]: all_pix_ring = hp.nest2ring(Nside, all_pix_nest)
...: hp.mollview(all_pix_ring, title='Ring scheme',
unit='Pix index')
...: pyplot.savefig('mollweide_view_ring.png')
...:
```

## Conversion between NESTED and RING schemes

**healpy.pixelfunc.reorder**

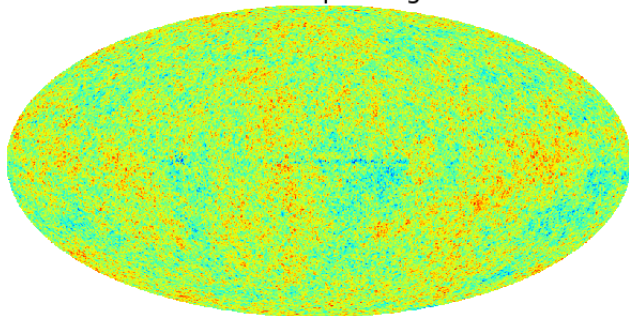
**healpy.pixelfunc.reorder**(*map\_in*, \*args, \*\*kwargs)

NEST  $\rightleftharpoons$  RING

## Conversion between NESTED and RING schemes

```
In [50]: mapa_ring = hp.read_map('COM_CMB_IQU-smica_1024_R2.02_full.fits')
...: hp.mollview(mapa_ring, title='CMB map - Ring', unit='K')
...:
NSIDE = 1024
ORDERING = NESTED in fits file <---
INDXSCHM = IMPLICIT
/home/camila/anaconda3_4p3p1/lib/python3.6/site-packages/healpy/fitsfunc.py:
339: UserWarning: No INDXSCHM keyword in header file : assume IMPLICIT
      "assume {}".format(schm))
Ordering converted to RING <---
```

CMB map - Ring



## Conversion between NESTED and RING schemes

How to use:

NEST  $\rightleftharpoons$  RING

```
In [52]: mapa_nest = hp.reorder(mapa_ring, inp='RING', out='NEST')
```

```
In [53]: # OR:::
```

```
....:
....: mapa_nest = hp.reorder(mapa_ring, r2n=True)
```

Exercise:

- Read the map in ring ordering.
- Reorder it according to the nested scheme.
- Visualize it.
- Reorder back to ring, and visualize it.

## $N_{\text{side}}$ / $N_{\text{pix}}$ / Resolution

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

$N_{\text{side}}$  /  $N_{\text{pix}}$  / Resolution

$$N_{\text{side}} \Rightarrow N_{\text{pix}}$$

```
In [72]: Nside1 = 64 # = 1, 2, 4, 8, 16, 64, ...
...: Npix1 = 12*Nside**2 # 12 x Nside^2
...: print('(1)', Nside1, '-->', Npix1)
...:
...: # OR:::
...:
...: Npix2 = hp.nside2npix(Nside1) # 12 x Nside^2
...: print('(2)', Nside1, '-->', Npix2)
...:
...: Nside2 = hp.npix2nside(Npix2) # 12 x Nside^2
...: print('(3)', Nside2, '<--', Npix2)
...:
(1) 64 --> 49152
(2) 64 --> 49152
(3) 64 <-- 49152
```

Exercise:

- $N_{\text{side}} = 4, 129, 1024$ .

$N_{\text{side}} / N_{\text{pix}} / \text{Resolution}$

$N_{\text{side}} \rightarrow \text{Resolution}$

```
In [81]: Nside = 4 # = 1, 2, 4, 8, 16, 64, ...
...: Resolution = hp.nside2resol(Nside)
...: print('(1)', Nside, '-->', Resolution, 'radians')
...:
(1) 4 --> 0.255831676987 radians

In [82]: Resolution = hp.nside2resol(Nside, arcmin=True)
...: print('(2)', Nside, '-->', Resolution, 'arcmin =', Resolution/60., 'deg')
...:
(2) 4 --> 879.484521425 arcmin = 14.6580753571 deg
```

$N_{\text{side}} \rightarrow \text{Pix Area}$

```
In [85]: Area = hp.nside2pixarea(Nside, degrees=True)
...: print('(3)', Nside, '-->', Area, 'square deg')
...:
(3) 4 --> 214.859173174 square deg
```

Exercise:

- $N_{\text{side}} = 2048$  (Planck).



## $N_{\text{side}}$ / $N_{\text{pix}}$ / Resolution

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

$N_{\text{side}}$  /  $N_{\text{pix}}$  / Resolution

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

**healpy.pixelfunc.ud\_grade**

**healpy.pixelfunc.ud\_grade**(*map\_in*, \*args, \*\*kws)

args = nside\_out, order\_in, order\_out, ...

$N_{\text{side}} / N_{\text{pix}} / \text{Resolution}$

How to use:

$N_{\text{side}} = 1024 \rightarrow N_{\text{side}} = 64$

```
In [103]: mapa = hp.read_map('COM_CMB_IQU-smica_1024_R2.02_full.fits')
NSIDE = 1024
ORDERING = NESTED in fits file
INDXSCHM = IMPLICIT
/home/camila/anaconda3_4p3p1/lib/python3.6/site-packages/healpy/fitsfunc.py:339:
UserWarning: No INDXSCHM keyword in header file : assume IMPLICIT
    "assume {}".format(schm))
Ordering converted to RING

In [104]: mapa_deg = hp.ud_grade(mapa, 64, order_in = 'RING', order_out='NEST')
...:
...: hp.mollview(mapa_deg, title='CMB map - Nside=64', unit='K', nest=True)
...:
```

$N_{\text{side}}$  /  $N_{\text{pix}}$  / Resolution

### Exercise:

- Repeat the degrading process using  $N_{\text{side\_out}} = 16$  and  $\text{order\_out} = \text{'RING'}$ .
- Degrade the mask<sup>1</sup> to  $N_{\text{side}}=128$  and to  $N_{\text{side}}=2048$ .
- Visualize the 3 cases.

---

<sup>1</sup>The fits file with the CMB map has one of its column corresponding to the mask. Use  $h=\text{True}$  to find it.