Tools for data analyses in Cosmology

- Aula 11 -

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June 13, 2017

Healpy

Previously: you could define ℓ_{\max} , m_{\max} .

Now you can select them ...

healpy.sphtfunc.Alm

class healpy.sphtfunc.Alm

getlmax (s[, mmax])

This class provides some static methods for alm index computation.

Methods

	getlm (Imax[, i])	Get the I and m from index and Imax.
	getidx (Imax, I, m)	Returns index corresponding to (I,m) in an array describing alm up to Imax.
	getsize (Imax[, mmax])	Returns the size of the array needed to store alm up to Imax and Imax

Returns the Imax corresponding to a given array size.

healpy.sphtfunc.Alm.getlm

static Alm.getlm(Imax, i=None)

Get the I and m from index and Imax.

healpy.sphtfunc.Alm.getidx

static Alm.getidx(Imax, I, m)

Returns index corresponding to (I,m) in an array describing alm up to Imax.

healpy.sphtfunc.Alm.getsize

static Alm.getsize(Imax, mmax=None)

Returns the size of the array needed to store alm up to Imax and Imax

healpy.sphtfunc.Alm.getlmax

static Alm.getlmax(s, mmax=None)

Returns the Imax corresponding to a given array size.

```
In [6]: map_in = hp.read_map('COM_CMB_IQU-smica_1024_R2.02_full.fits')
   ...: alm = hp.map2alm(map in)
   ...: l_max = hp.Alm.getlmax(len(alm))
   ...: size = hp.Alm.getsize(l_max)
   ...: print(len(alm), '/', size)
   ...: print('lmax =', l max)
NSIDE = 1024
ORDERING = NESTED in fits file
TNDXSCHM = TMPLTCTT
/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
4720128 / 4720128
lmax = 3071
```

```
Spherical harmonic transforms: tools
  How to use:
```

```
In [17]: l, m = hp.Alm.getlm(l_max) # getlm(l_max, index)
In [18]: l,m
Out[18]:
(array([ 0, 1, 2, ..., 3070, 3071, 3071]),
```

```
array([ 0, 0, 0, ..., 3070, 3070, 3071]))
In [19]: l[0:5], m[0:5]
```

Out[24]: 4720128

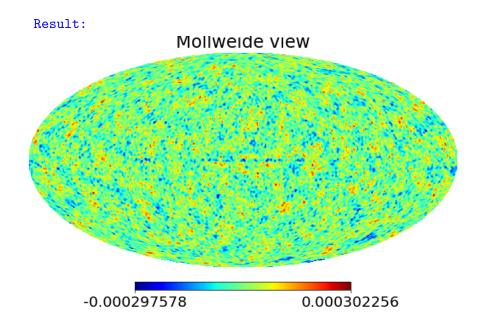
```
Out[19]: (array([0, 1, 2, 3, 4]), array([0, 0, 0, 0, 0]))
In [20]: l[3071:3076],m[3071:3076]
Out[20]: (array([3071, 1, 2, 3, 4]), array([0, 1, 1, 1, 1]))
```

```
In [22]: index = hp.Alm.getidx(l max, l, m)
In [23]: index
Out[23]: array([ 0, 1, 2, ..., 4720125, 4720126, 4720127])
In [24]: len(index)
```

Exercise:

o Reconstruct the CMB map ['COM_CMB_IQU-smica_1024_R2.02_full.fits'] corresponding to the range of multipole $50 \le \ell \le 200$. Steps:

- o Read the map.
- Calculate the $a_{\ell m}$'s [map2alm].
- Calculate the ℓ_{\max} [Alm.getlmax].
- \circ Calculate the ℓ and m for the whole $a_{\ell m}$ array [Alm.getlm].
- \circ Set to zero the $a_{\ell m}$ components for multipoles out of the chosen range [alm[1 < l_min] = (0+0j), alm[1 > l_max] = (0+0j)]
- Rebuild the map [alm2map].
- Visualize it.



healpy.sphtfunc.alm2cl

healpy.sphtfunc.alm2cl(alms1, alms2=None, Imax=None, Imax_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.

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 Imax is None, it is assumed Imax=cl.size-1 If mmax is None, it is assumed mmax=lmax.

healpy.sphtfunc.almxfl

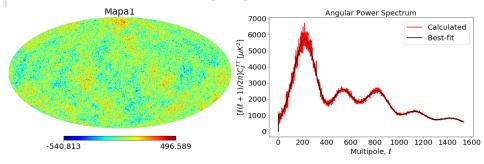
healpy.sphtfunc.almxfl(alm, fl, mmax=None, inplace=False)

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

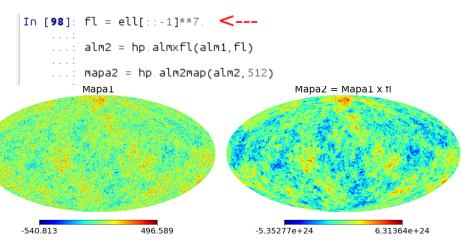
How to use:

```
In [93]: Cls = hp.read_cl('Cls_bestfitLCDM_PLA2_TT_lmax2508.fits')
In [94]: alm1 = hp.synalm(Cls,lmax=1535) # = synfast, but generates alm's
    ...: mapa1 = hp.alm2map(alm1,512)
```

In [97]: Cls_calc = hp.alm2cl(alm1) # = anafast, but upon alm's.



How to use:



—> Also to select a multipole range.

healpy.sphtfunc.smoothing

healpy.sphtfunc.smoothing(map_in, *args, **kwds)

Smooth a map with a Gaussian symmetric beam.

No removal of monopole or dipole is performed.

healpy.sphtfunc.smoothalm

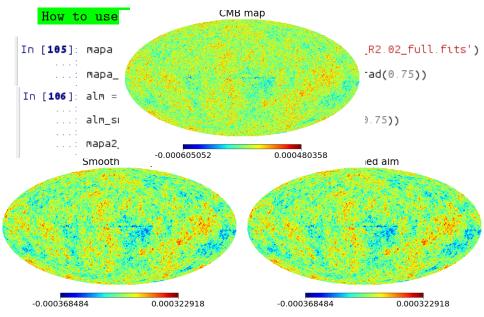
 $\label{lem:healpy.sphtfunc.smoothalm} \textbf{(alms, fwhm=0.0, sigma=None, pol=True, mmax=None, verbose=True, inplace=True)}$

Smooth alm with a Gaussian symmetric beam function.

...: $mapa2_alm_smo = hp.alm2map(alm_smo, 1024)$

```
In [105]: mapa = hp.read_map('COM_CMB_IQU-smica_1024_R2.02_full.fits')
...:
mapa_smo = hp.smoothing(mapa, fwhm=np.deg2rad(0.75))
In [106]: alm = hp.map2alm(mapa)
...:
alm_smo = hp.smoothalm(alm, fwhm=np.deg2rad(0.75))
```

```
In [105]: mapa = hp.read_map('COM_CMB_IQU-smica_1024_R2.02_full.fits')
     ...: mapa_smo = hp.smoothing(mapa, fwhm=np.deg2rad(0.75))
In [106]: alm = hp.map2alm(mapa)
      ...: alm_smo = hp.smoothalm(alm, fwhm=np.deg2rad(0.75))
      ...: mapa2_alm_smo = hp.alm2map(alm_smo, 1024)
          Smoothed map
                                                   Smoothed alm
  -0.000368484
                     0.000322918
                                          -0.000368484
                                                              0.000322918
```



$$C_{\ell}^{\text{obs}} = B_{\ell}^2 C_{\ell}$$

healpy.sphtfunc.pixwin

$$C_{\ell}^{\text{obs}} = B_{\ell}^2 C_{\ell}$$

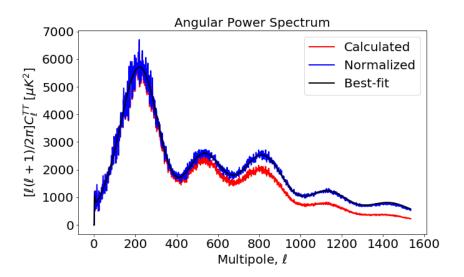
healpy.sphtfunc.pixwin

healpy.sphtfunc.pixwin(nside, pol=False) %

Exercise:

- Use the pixwin function to correct the angular power spectrum calculated from a map generated considering the pixel window function.
 Steps:
 - Read the Planck best-fit C_{ℓ} 's.
 - Generate a map (Nside = 512) including the pixel window effect [pixwin = True].
 - Calculate its C_{ℓ} 's [anafast].
 - Calculate the pixel window function [pixwin].
 - Use the result to correct the calculated C_ℓ 's $[C_\ell^{\rm obs} = B_\ell^2 C_\ell]$.
 - Compare them in a Plot.

Result:



Result: We can also correct the effect of degrading a map. The example shows the C_{ℓ} 's of a map degraded from Nside = 2048 to 512.

