# tutorial\_projection

December 29, 2019

#### 0.1 Introduction

This is a tutorial of generation of simulations and projection. We first specify two templates, one in equatorian coordinates with CAR pixellisation and one in equatorian coordinates with HEALPIX pixellisation. We generate alms from a CAMB lensed power spectrum file and use them to generate a random CMB realisation in both template. We then project the HEALPIX simulation and plot both the native CAR simulation and the projected HEALPIX simulation. We chose a low resolution nside to emphasize the effect of resolution

#### 0.2 Preamble

matplotlib magic

```
[1]: %matplotlib inline
```

Print versions used

```
[2]: import numpy as np
  import matplotlib as mpl
  import matplotlib.pyplot as plt
  import pspy, pixell
  print(" Numpy:", np.__version__)
  print("Matplotlib:", mpl.__version__)
  print(" pixell:", pixell.__version__)
  print(" pspy:", pspy.__version__)
```

```
Numpy: 1.18.0
Matplotlib: 3.1.2
pixell: 0.6.0+34.g23be32d
pspy: 0+untagged.88.g1ef44db
```

Get default data dir from pspy and set Planck colormap as default

```
[3]: from pspy.so_config import DEFAULT_DATA_DIR pixell.colorize.mpl_setdefault("planck")
```

#### 0.3 Generation of the templates

The CAR template will go from right ascension ra0 to ra1 and from declination dec0 to dec1 (all in degrees). It will have a resolution of 1 arcminute and it allows 3 components (stokes parameter

in the case of CMB anisotropies).

```
[4]: ra0, ra1 = -5, 5
  dec0, dec1 = -5, 5
  res = 1
  ncomp = 3
  from pspy import so_map
  template_car = so_map.car_template(ncomp, ra0, ra1, dec0, dec1, res)
```

We also generate an HEALPIX template for which we choose nside=256 so that the resolution of HEALPIX is much smaller

```
[5]: template_healpix = so_map.healpix_template(ncomp, nside=256, coordinate="equ")
```

### 0.4 Read power spectrum and alm generation

We first have to compute  $C_{\ell}$  data using a cosmology code such as CAMB and we need to install it since this is a prerequisite of pspy. We can do it within this notebook by executing the following command

```
[6]: %pip install camb
```

```
Requirement already satisfied: camb in
```

/home/garrido/Workdir/CMB/development/pspy/pyenv/lib/python3.8/site-packages (1.1.0)

Requirement already satisfied: scipy>=1.0 in

/home/garrido/Workdir/CMB/development/pspy/pyenv/lib/python3.8/site-packages (from camb) (1.4.1)

Requirement already satisfied: six in

/home/garrido/Workdir/CMB/development/pspy/pyenv/lib/python3.8/site-packages (from camb) (1.13.0)

Requirement already satisfied: sympy>=1.0 in

/home/garrido/Workdir/CMB/development/pspy/pyenv/lib/python3.8/site-packages (from camb) (1.5)

Requirement already satisfied: numpy>=1.13.3 in

/home/garrido/Workdir/CMB/development/pspy/pyenv/lib/python3.8/site-packages (from scipy>=1.0->camb) (1.18.0)

Requirement already satisfied: mpmath>=0.19 in

/home/garrido/Workdir/CMB/development/pspy/pyenv/lib/python3.8/site-packages (from sympy>=1.0->camb) (1.1.0)

Note: you may need to restart the kernel to use updated packages.

To make sure everything goes well, we can import CAMB and check its version

```
[7]: import camb print("CAMB version:", camb.__version__)
```

CAMB version: 1.1.0

Now that CAMB is properly installed, we will produce  $C_{\ell}$  data from  $\ell \text{min}=2$  to  $\ell \text{max}=104$  for the following set of  $\Lambda \text{CDM}$  parameters

We finally have to write  $C_{\ell}$  into a file to read back using the pixell.powspec function

and generate alms from the power spectrum up to lmax = 5000

```
[10]: from pixell import curvedsky lmax = 5000 alms = curvedsky.rand_alm(ps, lmax=lmax)
```

## 0.5 Computation of stokes parameters

We compute the stokes parameters from the alms in both templates

```
[11]: from pspy import sph_tools
map_healpix = sph_tools.alm2map(alms, template_healpix)
map_car = sph_tools.alm2map(alms, template_car)
```

and we project the HEALPIX map into the CAR template

```
[12]: map_healpix_proj = so_map.healpix2car(map_healpix, map_car, lmax=lmax)
```

```
WARNING: your lmax is too large, setting it to 3*nside-1 now Preparing SHT T -> alm float64 complex128 P -> alm Projecting
```

## 0.6 Showing maps

We plot both the native CAR map and the HEALPIX projected to CAR map. They contain the same CMB but have different resolutions.

```
fig, axes = plt.subplots(2, 3, figsize=(9, 6), sharex=True, sharey=True)
fields = ["T", "Q", "U"]
kwargs = dict(extent=[ra1, ra0, dec0, dec1], origin="lower")
for i, field in enumerate(fields):
    kwargs["vmin"] = np.min([map_car.data[i], map_healpix_proj.data[i]])
    kwargs["vmax"] = np.max([map_car.data[i], map_healpix_proj.data[i]])
    axes[0, i].imshow(map_car.data[i], **kwargs)
    axes[1, i].imshow(map_healpix_proj.data[i], **kwargs)
    axes[0, i].set_title(fields[i])

axes[0, 0].set_ylabel("CAR")
axes[1, 0].set_ylabel("HEALPIX")
plt.tight_layout()
```

We can also use the plot function from pspy.so\_map and set the output path to get individual images for each component T, Q, U.

```
[14]: map_car.plot(file_name=output_dir + "/map_car")
map_healpix_proj.plot(file_name=output_dir + "/map_healpix")
```