PSCF demo

1 Preliminaries

There are two steps for PSCF analysis:

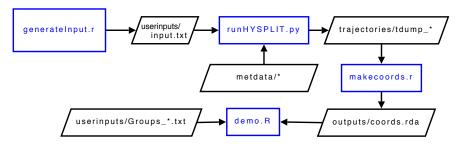
- 1. Generate back-trajectories using HYSPLIT
- 2. Create a grid over your domain and overlay HYSPLIT backtrajectories for the PSCF analysis.

Two important things to consider:

- 1. Regarding trajectory generation: how often to generate, time resolution, how far back in time
- 2. For PSCF or density calculation: spacing of grid points, whether to count each point in trajectory or the trajectory itself which passes over a given grid cell.

2 Part 1: Running HYSPLIT

Code diagram (note, demo.R may now be called PSCFdemo.R):



- 1. Download and place meteorological files in metdata/.
- 2. Edit the desired conditions in generateInput.r

This will generate a file called input.txt which is a table containing the inputs (by row) for HYSPLIT. input.txt is a tab-delimited file with the following format:

```
Latitude Longitude Altitude Time Run_hours Vert_coord Model_top 32.1200 -116.9700 1000.0000 04/25/09 00:00:00 -5 0 10000000.0000 32.1200 -116.9700 1200.0000 04/25/09 00:00:00 -5 0 1000000.0000 32.1200 -116.9700 1000.0000 04/25/09 02:00:00 -5 0 1000000.0000 32.1200 -116.9700 1200.0000 04/25/09 02:00:00 -5 0 1000000.0000 32.1200 -116.9700 1000.0000 04/25/09 02:00:00 -5 0 1000000.0000 32.1200 -116.9700 1000.0000 04/25/09 04:00:00 -5 0 1000000.0000
```

If you generate input.txt with your own script or program, please make certain the the "Time" format is exactly as shown (each field is 2-digits).

3. Execute runHYSPLIT.py, which will read input.txt and [create a CONTROL file and] run HYS-PLIT for each specified location/time. Trajectories are saved in a folder called trajectories/. Each trajectory is saved in a file with the format of tdump-%y_%m_%d_%H-Alt.

You may have to specify the location of directories containing the met files and name of executable file if they are different from the following (hymoldelt.exe was from HYSPLIT Version 4.6):

```
Input_file = 'input.txt'
Exec_file = './hymodelt.exe'
Meteo_path = './metdata/'
Output_path = './trajectories/'
Output_base = 'tdump'
Control = 'CONTROL'
```

3 Part 2: PSCF analysis

- 1. Run makecoords.r, which will read files in the trajectories directory and create a binary file called coords.rda in a folder called outputs/.
- 2. Use the script, PSCFdemo.R to load coords.rda and run the PSCF functions. Note: PSCFdemo.R is extracted from PSCFdemo.Rnw with Stangle("demo.Rnw") in R, so the code contained in the PSCFdemo.R file is exactly the same as that which produced the output and graphics in the PSCFdemo.pdf file.