## IPART <br> Release 2.0

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## The automated AR detect/tracking workflow:

1 Introduction ..... 3
2 Installation ..... 5
2.1 Install from conda-forge ..... 5
2.2 Create a conda environment using the environment file ..... 5
3 Dependencies ..... 7
4 Main functionalities ..... 9
4.1 Data preparation ..... 9
4.2 Perform the THR computation on IVT data ..... 12
4.3 Detect AR appearances from THR output ..... 14
4.4 Find the axis from detected AR ..... 20
4.5 Track ARs at individual time steps to form tracks ..... 23
5 ipart module contents ..... 29
5.1 Documentation page for thr.py ..... 29
5.2 Documentation page for AR_detector.py ..... 31
5.3 Documentation page for AR_tracer.py ..... 42
5.4 Documentation page for utils/funcs.py ..... 46
5.5 Documentation page for utils/plot.py ..... 48
6 Github and Contact ..... 53
7 Contributing and getting help ..... 55
8 License ..... 57
9 Indices and tables ..... 59
Bibliography ..... 61
Python Module Index ..... 63
Index ..... 65

Table of Contents

- Introduction
- Installation
- Install from conda-forge
- Create a conda environment using the environment file
- Dependencies
- Main functionalities
- ipart module contents
- Github and Contact
- Contributing and getting help
- License


## CHAPTER 1

## Introduction

IPART (Image-Processing based Atmospheric River Tracking) is a Python package for automated Atmospheric River (AR) detection, axis finding and AR tracking from gridded Integrated Vapor Transport (IVT) data, for instance Reanalysis datasets.

An overview of what ARs are can be found in this review paper: Atmospheric rivers: a mini-review.
IPART is intended for researchers and students who are interested in the field of atmospheric river studies in the present day climate or future projections. Different from commonly used AR detection methods that rely on thresholding on the IVT magnitude, this package includes a method inspired by an image processing technique - Top-hap by reconstruction (THR).

Below is an example output figure:
(a) 1984-01-04 06:00 IVT

(b) 1984-01-04 06:00 Reconstruction

(c) 1984-01-04 06:00 THR


Fig. 1.1: (a) The IVT field in $\mathrm{kg} /(\mathrm{m} * \mathrm{~s})$ at 1984-01-04 06:00 UTC over the North Hemisphere. (b) the IVT reconstruction field at the same time point. (c) the IVT anomaly field from the THR process at the same time point. In all three subplots, the detected ARs are outlined in black contour. The AR axes are drawn in green dashed lines.

Installation

### 2.1 Install from conda-forge

ipart can be installed in an existing conda environment:
conda install - conda-forge ipart
will install ipart and its dependencies for Python 3.

### 2.2 Create a conda environment using the environment file

Alternatively, users can obtain the code of this pacakge from the github page, and create a new conda environment using the environment files provided. This way will install the optional cartopy package and allow you to run the notebook examples.

```
git clone https://github.com/ihesp/IPART
cd IPART
conda env create -f environment_py3.yml
```

This creates a new environment named ipartpy3. Activate the environment using:

```
conda activate ipartpy3
```

After that, you can check the list of packages installed by:

```
conda list
```

Similarly for Python 2, use:

```
conda env create -f environment_py2.yml
```

- OS: Linux or MacOS. Windows is not tested.
- Python2.7 or Python3.7.
- netCDF4 (tested 1.4.2, 1.5.3 in py2, tested 1.5 .3 in py3)
- numpy (developed in 1.16 .5 in py2, tested $1.18 .1,1.19 .0$ in py3)
- scipy (developed in 1.2.1 in py2, tested 1.4.1, 1.5.1 in py3)
- matplotlib (tested 2.2.5 in py2, tested 3.3.1 in py3)
- pandas (developed in 0.23.4, 0.24.2 in py2, tested 1.0.3, 1.0.5 in py3)
- networkx (developed in 1.11 and 2.2 in py2, tested 2.4 in py 3 )
- scikit-image (developed in $0.14 .2,0.14 .3$ in py2, tested $0.16 .2,0.17 .2$ in py3)
- cartopy (optional, only used for plotting. Tested 0.17 .0 in py2, tested 1.18 .0 in py3)


## chapter 4

## Main functionalities

There are four main functionalities provided by the package that collectively constitute a specific workflow for the automated AR detection/tracking task:

1. Perform THR computation on input data.
2. Detect ARs from the outputs from the previous step, and at the same time,
3. Identify the AR axis.
4. Track ARs detected at individual time steps to form tracks.

More details regarding these steps are provided in separate pages below.
Applications on example data can be found in a series of example notebooks at github repository.

### 4.1 Data preparation

### 4.1.1 netcdf data

Source data are the $u$ - and $v$ - components of the vertically integrated vapor fluxes, in a rectangular grid.
The u-component of the vertically integrated vapor fluxes have a standard_name of eastward_atmosphere_water_transport_across_unit_distance.
The v-component of the vertically integrated vapor fluxes have a standard_name of northward_atmosphere_water_transport_across_unit_distance.

These are usually computed as:

$$
\left\{\begin{array}{l}
F_{u}=\int_{T O A}^{P_{s}} \frac{u q}{g} d P \\
F_{v}=\int_{T O A}^{P_{s}} \frac{v q}{g} d P
\end{array}\right.
$$

where

- $F_{u}\left(F_{v}\right)$ is the zonal (meridional) component of the integrated flux, both in $\mathrm{kg} /(\mathrm{m} * s)$.
- $u(v)$ : is the zonal (meridional) wind speed (in $\mathrm{m} / \mathrm{s}$ ) at a given level, and $q$ is the specific humidity (in $\mathrm{kg} / \mathrm{kg}$ ) at the same level.
- $d P$ is the pressure increment, in $P a$, and $g$ is acceleration by gravity.
- TOA can be substituted with a sensibly high level, e.g. 300hPa.

No strict requirement for the spatial or temporal resolution of input data is imposed, however, for better results, one should use something greater than $\sim 1.5^{\circ}$ latitude/longitude. 6-hourly temporal resolution is a standard for Reanalysis datasets, daily would probably work, but one should do some parameter adjustments in such a case.

Data are supposed to saved in netcdf files.

### 4.1.2 Metadata

Note: the user is responsible for making sure that the data are saved in the following rank order:

```
(time, level, latitude, longitude)
```

or:

```
(time, latitude, longitude)
```

The level dimension is optional. As data are vertical integrals, the length of the level dimension, if present, should be 1 .

The user also needs to provide the time, latitude and longitude axes values. This is because these temporal and geographical information is used in the computation.

To test the sanity of your input data, run this script against the netcdf data file:

```
cd /path/to/IPART/folder/you/cloned
python test_data.py /path/to/your/netcdf/file 'id_of_variable'
```

For instance

```
cd ~/Downloads/IPART
python test_data.py ~/datasets/erai/uflux_file.nc 'uflux'
```

The expected outputs would look like this:

```
### <test_data>: Read in file:
/home/guangzhi/scripts/IPART/notebooks/uflux_s_6_1984_Jan.nc
##################################################
Variable time axis:
##################################################
### Description of slab ###
    id: time
    shape: (124,)
    filename: None
    missing_value: None
    comments: None
    grid_name: None
    grid_type: None
    long_name: time
    units: hours since 1900-01-01 00:00:00.0
```

```
    standard_name: None
    Order: []
### End of description ###
None
Time axis values:
[datetime.datetime(1984, 1, 1, 0, 0) datetime.datetime(1984, 1, 1, 6, 0)
    datetime.datetime(1984, 1, 1, 12, 0) datetime.datetime(1984, 1, 1, 18, 0)
    datetime.datetime(1984, 1, 2, 0, 0) datetime.datetime(1984, 1, 2, 6, 0)
    datetime.datetime(1984, 1, 2, 12, 0) datetime.datetime(1984, 1, 2, 18, 0)
    datetime.datetime(1984, 1, 29, 0, 0) datetime.datetime(1984, 1, 29, 6, 0)
    datetime.datetime(1984, 1, 31, 12, 0)
    datetime.datetime(1984, 1, 31, 18, 0)]
##################################################
Variable latitude axis:
##################################################
### Description of slab ###
    id: latitude
    shape: (94,)
    filename: None
    missing_value: None
    comments: None
    grid_name: None
    grid_type: None
    long_name: latitude
    units: degrees_north
    standard_name: None
    Order: []
### End of description ###
None
Latitude axis values:
\begin{tabular}{clllllllllll} 
[10. & 10.75 & 11.5 & 12.25 & 13. & 13.75 & 14.5 & 15.25 & 16. & 16.75 & 17.5 & 18.25 \\
19. & 19.75 & 20.5 & 21.25 & 22. & 22.75 & 23.5 & 24.25 & 25. & 25.75 & 26.5 & 27.25 \\
28. & 28.75 & 29.5 & 30.25 & 31. & 31.75 & 32.5 & 33.25 & 34. & 34.75 & 35.5 & 36.25 \\
37. & 37.75 & 38.5 & 39.25 & 40. & 40.75 & 41.5 & 42.25 & 43. & 43.75 & 44.5 & 45.25 \\
46. & 46.75 & 47.5 & 48.25 & 49. & 49.75 & 50.5 & 51.25 & 52. & 52.75 & 53.5 & 54.25 \\
55. & 55.75 & 56.5 & 57.25 & 58. & 58.75 & 59.5 & 60.25 & 61. & 61.75 & 62.5 & 63.25 \\
64. & 64.75 & 65.5 & 66.25 & 67. & 67.75 & 68.5 & 69.25 & 70. & 70.75 & 71.5 & 72.25 \\
73. & 73.75 & 74.5 & 75.25 & 76. & 76.75 & 77.5 & 78.25 & 79. & \(79.75]\) &
\end{tabular}
##################################################
Variable longitude axis:
##################################################
### Description of slab ###
    id: longitude
    shape: (480,)
    filename: None
    missing_value: None
    comments: None
    grid_name: None
    grid_type: None
    long_name: longitude
```

```
    units: degrees_east
    standard_name: None
    Order: []
### End of description ###
None
Longitude axis values:
[-180. -179.25-178.5 -177.75 -177. -176.25 -175.5 -174.75 -174.
-173.25 -172.5 -171.75 -171. -170.25 -169.5 -168.75 -168. -167.25
-166.5 -165.75 -165. -164.25 -163.5 -162.75 -162. -161.25 -160.5
...
    164.25 165. 165.75 166.5 167.25 168. 168.75 169.5 170.25
    171. 171.75 172.5 173.25 174. 174.75 175.5 176.25 177.
    177.75 178.5 179.25]
Data have unit of "kg m**-1 s**-1"
```

Pay some attention to the values listed in the latitude and longitude axes blocks, to make sure the values make physical sense. For high resolution data, the input variable may have a fairly large size, e.g. a longitude axis of length 720 (if your data have a resolution of $0.5 \times 0.5^{\circ}$ ). If the longitude axis reports a largest value of 720 , it is probably reporting the size of the longitude dimension, rather than the actual longitude label (as the maximum possible longitude label should be 360 ). In such cases, the user should take some extra steps to make sure that the data have proper metadata associated with them.

### 4.1.3 Compute IVT

With $F_{u}$ and $F_{v}$, compute the IVT as

$$
I V T=\sqrt{F_{u}^{2}+F_{v}^{2}}
$$

This is trivial to achieve, you can use the compute_ivt. py script provided in the package for this computation.

### 4.2 Perform the THR computation on IVT data

### 4.2.1 The Top-hat by Reconstruction (THR) algorithm

The AR detection method is inspired by the image processing technique top-hat by reconstruction (THR), which consists of subtracting from the original image a greyscale reconstruction by dilation image. Some more details of the THR algorithm and its applications can be found in this work of [Vincent1993].

In the context of AR detection, the greyscale image in question is the non-negative IVT distribution, denoted as $I$.
The greyscale reconstruction by dilation component (hereafter reconstruction) corresponds to the background IVT component, denoted as $\delta(I)$.
The difference between $I$ and $\delta(I)$ gives the transient IVT component, from which AR candidates are searched.

Note: we made a modification based on the THR algorithm as descripted in [Vincent1993]. The marker image used in this package is obtained by a grey scale erosion ${ }^{1}$ with a structuring element $E$, while in a standard THR process

[^0]as in [Vincent1993], the marker image is obtained by a global substraction $I-\delta h$, where $\delta h$ is the pixel intensity subtracted globally from the original input image $I$.

The introduction of the grey scale erosion process allows us to have a control over on what spatio-temporal scales ARs are to be detected. An important parameter in this erosion process is the size of the structuring element $E$.

We then extend the processes of erosion and reconstruction to 3D (i.e. time, $x$ - and $y$ - dimensions), measuring "spatiotemporal spikiness". The structuring element used for 3D erosion is a 3D ellipsoid:

$$
E=\left\{(z, x, y) \in \mathbb{Z}^{3} \mid(z / t)^{2}+(x / s)^{2}+(y / s)^{2} \leq 1\right\}
$$

with the axis length along the time dimension being $t$, and the axes for the x - and y -dimensions sharing the same length $s$. Both $t$ and $s$ are measured in pixels/grids.

Note: the axis length of an ellipsoid is half the size of the ellipsoid in that dimension. For relatively large sized $E$, the difference in the THR results using an ellipsoid structuring element and a 3D cube with size $(2 t+1,2 s+1,2 s+1)$ is fairly small.

Considering the close physical correspondences between ARs and extra-tropical storm systems, the "correct" THR parameter choices of $t$ and $s$ should be centered around the spatio-temporal scale of ARs.
Let's assume the data we are working with is 6 -hourly in time, and $0.75 \times 0.75^{\circ}$ in space.
The typical synoptic time scale is about a week, giving $t=4$ days (recall that $t$ is only half the size of the time dimension).

The typical width of ARs is within 1000 km , therefore $s=6 \mathrm{grids}$ is chosen. Given the $0.75^{\circ}$ resolution of data, this corresponds to a distance of about $80 \mathrm{~km} / \mathrm{grid} \times(6 \times 2+1) \mathrm{grids}=1040 \mathrm{~km}$. An extra grid is added to ensure an odd numbered grid length, same for the $t$ parameter: the number of time steps is 4 steps /day $\times 4$ days $\times 2+1$ step $=$ 33 steps.

### 4.2.2 Compute THR

Using the above setup, the THR process is computed using following code:

```
from ipart import thr
ivt, ivtrec, ivtano = thr.THR(ivt_input, [16, 6, 6])
```

where ivt_input is the input IVT data, ivtrec is the reconstruction component, and ivtano is the anomalous component.

Note: the thr. $\operatorname{THR()}$ function accepts an optional argument oro, which is to provide the algorithm with some surface elevation information, with the help of which detection sensitivity of landfalling ARs can be enhanced.

## See also:

```
thr.THR().
```


### 4.2.3 Dedicated Python script

The package provides two script to help doing this computation:

- compute_thr_singlefile: when your IVT data are saved in a single file.
- compute_thr_multifile: when your IVT data are too large to fit in a single file, e.g. data spanning multiple decades and saved into one-file-per year. Like in the case of a simple moving average, discontinuity at the end of one year and the beginning of the next may introduce some errors. When the data are too large to fit into RAM, one possible solution is to read in 2 years at a time, concatenate them then perform the filtering/THR process to achieve a smooth year-to-year transition. Then read in the 3rd year to form another 2-year concatenation with the 2 nd year. Then the process rotates on untill all years are processed.


### 4.2.4 Example output

### 4.2.5 Notebook example

An example of this process is given in this notebook.

### 4.2.6 References

### 4.3 Detect AR appearances from THR output

### 4.3.1 Definition of AR occurrence

An AR occurrence at a given time point is defined using these following rules:

1. A connected region in the IVT anomaly field $(I-\delta(I)$, computed in the section "The Top-hat by Reconstruction (THR) algorithm") where its values is greater than 0 .
2. The centroid (weighted by IVT values of the grid cells) of the region is north of $20^{\circ} \mathrm{N}$ (or south of $20^{\circ} S$ for the Southern Hemisphere), and south of $80^{\circ} N$ (or north of $80^{\circ} S$ for the Southern Hemisphere), i.e. we are only interested in mid-latitude systems.
3. The region's area has to be within $50-1800 \times 10^{4} \mathrm{~km}^{2}$.
4. After the computation of this AR candidates axis (see Find the axis from detected $A R$ ) and the effective width (defined as area/length ratio), the length has to be $\geq 1500 \mathrm{~km}$, and length/width ratio has to be $\geq 2$ if length is below 2000 km .

Note: API is provided to control all these parameters.

### 4.3.2 Input data

These are the input data required for AR occurrence detection:

- u- and v-components of the integrated vapor fluxes ( $F_{u}$ and $F_{v}$ ).
- IVT (as $\sqrt{F_{u}^{2}+F_{v}^{2}}$ ), see Compute IVT for more.
- The output from the THR process: the reconstruction component $(\delta(I))$ and the anomaly component $(I-\delta(I))$. See The Top-hat by Reconstruction (THR) algorithm for more.

Additional inputs:

- latitude, longitude and time axis. See Metadata for more.
- detection parameters, see below.
(a) 1984-1-26 0:0:0.0, IVT (=IVT_rec + IVT_ano)

(b) 1984-1-26 0:0:0.0, IVT_rec



Fig. 4.1: (a) The IVT field in $\mathrm{kg} /\left(\mathrm{m}^{*} \mathrm{~s}\right)$ at 1984-01-26 00:00 UTC over the North Hemisphere. (b) the IVT reconstruction field $(\delta(J)$ ) at the same time noint (c) the IVT anomaly field $(I-\delta(I))$ from the THR nrocess at the same time

```
PARAM_DICT={
    # kg/(m*s), define AR candidates as regions >= than this anomalous ivt.
    'thres_low' : 1,
    # km^2, drop AR candidates smaller than this area.
    'min_area': 50*1e4,
    # km^2, drop AR candidates larger than this area.
    'max_area': 1800*1e4,
    # float, min length/width ratio, applied only when length<min_length.
    'min_LW': 2,
    # degree, exclude systems whose centroids are lower than this latitude.
    'min_lat': 20,
    # degree, exclude systems whose centroids are higher than this latitude.
    'max_lat': 80,
    # km, ARs shorter than this length is treated as relaxed.
    'min_length': 2000,
    # km, ARs shorter than this length is discarded.
    'min_length_hard': 1500,
    # degree lat/lon, error when simplifying axis using rdp algorithm.
    'rdp_thres': 2,
    # grids. Remove small holes in AR contour.
    'fill_radius': None,
    # do peak partition or not, used to separate systems that are merged
    # together with an outer contour.
    'single_dome': False,
    # max prominence/height ratio of a local peak. Only used when SINGLE_DOME=True
    'max_ph_ratio': 0.6,
    # minimal proportion of flux component in a direction to total flux to
    # allow edge building in that direction
    'edge_eps': 0.4
    }
```


### 4.3.3 Usage in Python scripts

The following snippet shows the detection function calls:

```
from ipart.AR_detector import findARs
time_idx, labelsNV, anglesNV, crossfluxesNV, result_df = findARs(ivtNV.data,
    ivtrecNV.data, ivtanoNV.data, quNV.data, qvNV.data, latax, lonax,
    times=timeax, **PARAM_DICT)
```

where these input arguments are:

- ivtNV is a ipart. utils.NCVAR data object, which is a rudimentary wrapper object designed to achieve a tighter bound between data values and metadata. Same for the other variables with an NV suffix.
- ivtNV.data is the IVT data values in numpy. ndarray format, with dimensions of (time, level, latitude, longitude) or (time, latitude, longitude).
- ivtrecNV is $\delta(I)$, and ivtanoNV is $I-\delta(I)$, see The Top-hat by Reconstruction (THR) algorithm for more details.
- quNV: is $F_{u}$, and qvNV is $F_{v}$.
- latax: is an 1d array storing the latitude coordinates of ivtNV and others.
- lonax: is an 1d array storing the longitude coordinates of ivtNV and others.
- timeax is a list of python datetime objects storing time stamps of the data in ivtNV and others.
- PARAM_DICT is the parameter dictionary as defined above.

The return values are:

- time_idx is a list of indices of the time dimension when any AR is found.
- labelsNV is a ipart.utils. NCVAR object, whose data attribute is an ndarray variable saving the numerical labels of all found ARs in each time step. It has shape of (time, lat, lon).
- anglesNV is a ipart.utils.NCVAR object storing an ndarray variable saving the difference in the orientation of IVT vectors in all found ARs, wrt the AR axis.
- crossfluxesNV is a ipart.utils.NCVAR object storing an ndarray variable saving the cross-sectional IVT flux, computed as the projection of IVT vectors onto the AR axis, using angles in angles.
- The result_df return value is a pandas. DataFrame object saving in a table the various attributes of all detected ARs at this time point.


## See also:

AR_detector.findARs(), AR_detector.findARsGen(), AR_detector.getARData().

## AR records DataFrame

The rows of ardf are different AR records, the columns of ardf are listed below:

- id: integer numeric id for this AR at this particular time point. ARs at different time points can share the same id, and an AR can be uniquely identified with the combination of time stamp + id.
- time : time stamp in the YYYY-MM-DD HH:mm:ss format.
- contour_y : list of floats, the y-coordinates (latitudes) of the AR contour in degrees North.
- contour_x : list of floats, the x-coordinates (longitude) of the AR contour in degrees North.
- centroid_y : float, latitude of the AR centroid, weighted by the IVT value.
- centroid_x: float, longitude of the AR centroid, weighted by the IVT value.
- axis_y : list of floats, latitudes of the AR axis.
- axis_x: list of floats, longitude of the AR axis.
- axis_rdp_y : list of floats, latitude of the simplified AR axis.
- axis_rdp_x: list of floats, longitude of the simplified AR axis.
- area: float, area of the AR in $\mathrm{km}^{2}$.
- length : float, length of the AR in km .
- width : float, effective width in $k m$, as area/length.
- LW_ratio: float, length/width ratio.
- strength: float, spatially averaged IVT value within the $\operatorname{AR}$ region, in $\mathrm{kg} /(\mathrm{m} \cdot \mathrm{s})$.
- strength_ano: float, spatially averaged anomalous IVT value within the AR region, in $k g /(m \cdot s)$.
- strength_std: float, standard deviation of IVT within the AR region, in $k g /(m \cdot s)$.
- max_strength : float, maximum IVT value within the AR region, in $k g /(m \cdot s)$.
- mean_angle : float, spatially averaged angle between the IVT vector and the AR axis, in degrees.
- is_relaxed : True or False, whether the AR is flagged as "relaxed".
- qv_mean : float, spatially averaged meridional integrated vapor flux, in $\mathrm{kg} /(\mathrm{m} \cdot \mathrm{s})$.


### 4.3.4 Dedicated Python script

You can use the scripts/detect_ARs.py or scripts/detect_ARs_generator_version.py script (check them out in the github repo). for AR detection process in production. The former does the computation and returns all outpus in one go, and the latter yields results at each time step separately, so the outputs can be saved to disk rather than accumulating in RAM. Note that this process is essentially time-independent, i.e. the computation of one time point does not rely on another, therefore you can potentially parallelize this process to achieve greater efficiency.

### 4.3.5 Example output

The resultant detected ARs can be visualized using the following snippet:

```
import matplotlib.pyplot as plt
from ipart.utils import plot
import cartopy.crs as ccrs
plot_vars=[slab,slabrec,slabano]
titles=['IVT', 'Reconstruction', 'THR']
iso=plot.Isofill(plot_vars,12,1,1,min_level=0,max_level=800)
figure=plt.figure(figsize=(12,10),dpi=100)
for jj in range(len(plot_vars)):
    ax=figure.add_subplot(3,1,jj+1,projection=ccrs.PlateCarree())
    pobj=plot.plot2(plot_vars[jj],iso,ax,
            xarray=lonax, yarray=latax,
            title='%s %s' %(timett_str, titles[jj]),
            fix_aspect=False)
plot.plotAR(ardf,ax,lonax)
figure.show()
```


## See also:

```
utils.plot.Isofill,utils.plot.plot2().
```

One example output figure is shown below:

### 4.3.6 Notebook example

An example of this process is given in this notebook.
(a) 1984-01-04 06:00 IVT

(b) 1984-01-04 06:00 Reconstruction

(c) 1984-01-04 06:00 THR


$\mathrm{kg} / \mathrm{m} / \mathrm{s}$

Fig. 4.2: (a) The IVT field in $\mathrm{kg} /(\mathrm{m} * \mathrm{~s})$ at 1984-01-04 06:00 UTC over the North Hemisphere. (b) the IVT reconstruction field at the same time point. (c) the IVT anomaly field from the THR process at the same time point. In all three subplots, the detected ARs are outlined in black contour. The AR axes are drawn in green dashed lines.

### 4.4 Find the axis from detected AR

### 4.4.1 Axis-finding in a planar graph framework

After Detect AR appearances from THR output, a binary mask $I_{k}$ representing the spatial region of each candidate AR is obtained. An axis is sought from this region that summarizes the shape and orientation of the AR. A solution in a planar graph framework is proposed here.
A directed planar graph is build using the coordinate pairs $\left(\lambda_{k}, \phi_{k}\right)$ as nodes (see Figure Fig. 4.3 below). At each node, directed edges to its eight neighbors are created, so long as the moisture flux component along the direction of the edge exceeds a user-defined fraction ( $\epsilon$, see the PARAM_DICT in Input data) to the total flux. The along-edge flux is defined as:
$f_{i j}=u_{i} \sin (\alpha)+v_{i} \cos (\alpha)(4.1)$
where $f_{i j}$ is the flux along the edge $e_{i j}$ that points from node $n_{i}$ to node $n_{j}$, and $\alpha$ is the azimuth angle of $e_{i j}$.
Therefore an edge can be created if $f_{i j} / \sqrt{u_{i}^{2}+v_{i}^{2}} \geq \epsilon$. It is advised to use a relatively small $\epsilon=0.4$ is used, as the orientation of an AR can deviate considerably from its moisture fluxes, and denser edges in the graph allows the axis to capture the full extent of the AR.
The boundary pixels of the AR region are then found, labeled $L_{k}$. The trans-boundary moisture fluxes are compute as the dot product of the gradients of $I_{k}$ and $\left(u_{k}, v_{k}\right): \nabla I_{k} \cdot\left(u_{k}, v_{k}\right)$.
Then the boundary pixels with net input moisture fluxes can be defined as:

$$
L_{k, i n}=\left\{p \in L_{k} \mid\left(\nabla I_{k} \cdot\left(u_{k}, v_{k}\right)\right)(p)>0\right\}
$$

Similarly, boundary pixels with net output moisture fluxes is the set

$$
L_{k, \text { out }}=\left\{p \in L_{k} \mid\left(\nabla I_{k} \cdot\left(u_{k}, v_{k}\right)\right)(p)<0\right\}
$$

These boundary pixels are colored in green and black, respectively, in Fig. 4.4.
For each pair of boundary nodes $\left\{\left(n_{i}, n_{j}\right) \mid n_{i} \in L_{k, i n}, n_{j} \in L_{k, \text { out }}\right\}$, a simple path (a path with no repeated nodes) is sought that, among all possible paths that connect the entry node $n_{i}$ and the exit node $n_{j}$, is the shortest in the sense that its path-integral of weights is the lowest.

The weight for edge $e_{i j}$ is defined as:

$$
w_{i j}=e^{-f_{i j} / A}
$$

where $f_{i, j}$ is the projected moisture flux along edge $e_{i, j}$ and $A=100 \mathrm{~kg} /(\mathrm{m} \cdot \mathrm{s})$ is a scaling factor.
This formulation ensures a non-negative weight for each edge, and penalizes the inclusion of weak edges when a weighted shortest path search is performed.

The Dijkstra path-finding algorithm is used to find this shortest path $p_{i j}^{*}$.
Then among all $p_{i j}^{*}$ that connect all entry-exit pairs, the one with the largest path-integral of along-edge fluxes is chosen as the AR axis, as highlighted in yellow in Fig. 4.4.
It could be seen that various aspects of the physical processes of ARs are encoded. The shortest path design gives a natural looking axis that is free from discontinuities and redundant curvatures, and never shoots out of the AR boundary. The weight formulation assigns smaller weights to edges with larger moisture fluxes, "urging: the shortest path to pass through nodes with greater intensity. The found axis is by design directed, which in certain applications can provide the necessary information to orient the AR with respect to its ambiance, such as the horizontal temperature gradient, which relates to the low level jet by the thermal wind relation.


Fig. 4.3: Schematic diagram illustrating the planar graph build from the AR pixels and horizontal moisture fluxes. Nodes are taken from pixels within region $I_{k}$, and are represented as circles. Red vectors denote $I V T$ vectors. The one at node $n_{i}$ forms an angle $\theta$ with the x -axis, and has components $(u, v)$. Black arrows denote directed edges between nodes, using an 8 -connectivity neighborhood scheme. The edge between node $n_{i}$ and $n_{j}$ is $e_{i j}$, and forms an azimuth angle $\alpha$ with the y-axis. $w_{i j}$ is the weight attribute assigned to edge $e_{i j}$, and $f_{i j}$ is along-edge moisture flux.


Fig. 4.4: Application of the axis finding algorithm on the AR in the North Pacific, 2007-Dec-1, 00 UTC. IVT within the AR is shown as colors, in $\mathrm{kg} /(\mathrm{m} \cdot \mathrm{s})$. The region of the AR $\left(I_{k}\right)$ is shown as a collection of gray dots, which constitute nodes of the directed graph. Edges among neighboring nodes are created. A square marker is drawn at each boundary node, and is filled with green if the boundary node has net input moisture fluxes ( $n_{i} \in L_{k, i n}$ ), and black if it has net output moisture fluxes $\left(n_{i} \in L_{k, o u t}\right)$. The found axis is highlighted in yellow.

### 4.4.2 Usage in Python scripts

The following snippet shows the axis finding process:

```
from ipart.AR_detector import findARAxis
axis_list, axismask=findARAxis(quslab, qvslab, mask_list, costhetas,
    sinthetas, param_dict)
```

where:

- quslab, qvslab are the $u$ - and v- component of integrated vapor fluxes at a given time point.
- mask_list is a list of binary masks denoting the region of an each found AR.
- sinthetas and costhetas are used to compute the azimuthal angles for each grid cell.
- param_dict is the parameter dictionary as defined in Input data.


## See also:

```
AR_detector.findARAxis(),AR_detector.maskToGraph(),AR_detector.getARAxis().
```


### 4.4.3 Dedicated Python script

No detected Python script is offered for this process, as it is performed in the AR_detector.findARsGen() function.

### 4.4.4 Notebook example

An example of this process is given in this notebook.

### 4.5 Track ARs at individual time steps to form tracks

### 4.5.1 The modified Hausdorff distance definition

After the Detect AR appearances from THR output process, assume we have detected

- $n$ ARs at time $t$, and
- $m$ ARs at time $t+1$.

There are theoretically $n \times m$ possible associations to link these two groups of ARs. Of cause not all of them are meaningful. The rules that are applied in the association process are:

1. The nearest neighbor link method: for any AR at time $t$, the nearest AR at time $t+1$ "wins" and is associated with it, subject to that:
2. the inter-AR distance $(\mathbf{H})$ is $\leq 1200 \mathrm{~km}$.
3. no merging or splitting is allowed, any AR at time $t$ can only be linked to one AR at time $t+1$, similarly, any AR at time $t+1$ can only be linked to one AR at time $t$.
4. after all associations at any give time point have been created, any left-over AR at time $t+1$ forms a track on their own, and waits to be associated in the next iteration between $t+1$ and $t+2$.
5. any track that does not get updated during the $t-(t+1)$ process terminates. This assumes that no gap in the track is allowed.

The remaining important question is how to define that inter-AR distance (H). Here we adopt a modified Hausdorff distance definition:

$$
H(A, B) \equiv \min \left\{h_{f}(A, B), h_{b}(A, B)\right\}
$$

where $H(A, B)$ is the modified Hausdorff distance from track $\mathbf{A}$ to track $\mathbf{B}, h_{f}(A, B)$ is the forward Hausdorff distance from $\mathbf{A}$ to $\mathbf{B}$, and $h_{b}(A, B)$ the backward Hausdorff distance from $\mathbf{A}$ to $\mathbf{B}$. They are defined, respectively, as:

$$
h_{f}(A, B) \equiv \max _{a \in A}\left\{\min _{b \in B}\left\{d_{g}(a, b)\right\}\right\}
$$

namely, the largest great circle distance of all distances from a point in $\mathbf{A}$ to the closest point in $\mathbf{B}$. And the backward Hausdorff distance is:

$$
h_{b}(A, B) \equiv \max _{b \in B}\left\{\min _{a \in A}\left\{d_{g}(a, b)\right\}\right\}
$$

Note that in general $h_{f} \neq h_{b}$. Unlike the standard definition of Hausdorff distance that takes the maximum of $h_{f}$ and $h_{b}$, we take the minimum of the two.

The rationale behind this modification is that merging/splitting of ARs mostly happen in an end-to-end manner, during which a sudden increase/decrease in the length of the AR induce misalignment among the anchor points. Specifically, merging (splitting) tends to induce large backward (forward) Hausdorff distance. Therefore $\min \left\{h_{f}(A, B), h_{b}(A, B)\right\}$ offers a more faithful description of the spatial closeness of ARs. For merging/splitting events in a side-to-side manner, this definition works just as well.

### 4.5.2 The nearest neighbor link method

To link AR records to form a track, a nearest neighbor method is used that the two AR axes found in consecutive time steps with a Hausdorff distance $\leq 1200 \mathrm{~km}$ are linked, with an exclusive preference to the smallest Hausdorff distance.

Formally, suppose $n$ tracks have been found at $t=t: A=\left\{a_{1}, a_{2}, \cdots, a_{n}\right\}$, and $t=t+1$ has $m$ new records: $B=\left\{b_{1}, b_{2}, \cdots, b_{m}\right\}$. The Hausdorff distances between all pairs of possible associations form a distance matrix:

$$
M=\left[\begin{array}{cccc}
H\left(a_{1}, b_{1}\right) & H\left(a_{1}, b_{2}\right) & \cdots & H\left(a_{1}, b_{m}\right) \\
H\left(a_{2}, b_{1}\right) & H\left(a_{2}, b_{2}\right) & \cdots & H\left(a_{2}, b_{m}\right) \\
\vdots & \vdots & \vdots & \vdots \\
H\left(a_{n}, b_{1}\right) & H\left(a_{n}, b_{2}\right) & \cdots & H\left(a_{n}, b_{m}\right)
\end{array}\right]
$$

Then Algorithm shown in Fig. 4.5 is called with these arguments:

$$
\left(A=A, B=B, M=M, H^{*}=1200 \mathrm{~km}, R^{-}=[], C^{-}=[]\right)
$$

The algorithm iteratively links two AR records with the smallest distance, so long as the distance does not exceed a given threshold $H^{*}$.
It ensures that no existing track connects to more than one new records, and no new record connects to more than one existing tracks. After this, any left-over records in $B$ form a new track on their own. Then the same procedure repeats with updated time $t:=t+1$. Tracks that do not get any new record can be removed from the stack list, which only maintains a few active tracks at any given time. Therefore the complexity does not scale with time.

Note: One can use 3 consecutive calls of the above algorithm, with different input arguments, to achieve merging and splitting in the tracks.

```
Algorithm 1: Nearest neighbor algorithm
    Data: \(A\) : list of existing tracks at \(t, B\) : list of new records at \(t+1, M\) : distance
        matrix, \(H^{*}\) : maximum allowed distance, \(R^{-}\): list of indices of tracks not
        allowed to link, \(C^{-}\): list of indices of records not allowed to link.
    Result: \(A\) : updated list of tracks. \(R^{+}\): list of indices of tracks get linked. \(C^{+}\):
            list of indices of records get linked.
    create empty lists \(R^{+}=[]\)and \(C^{+}=[]\);
    while \(\min (M)<H^{*}\) and \(\operatorname{len}\left(R^{+}\right)<\min (n, m)\) do
            for each \((i, j)\) that \(M(i, j)=\min (M)\) do
                if \(i \notin R^{-}\)and \(j \notin C^{-}\)then
            let \(a_{i}=A[i] ;\)
            let \(b_{j}=B[j]\);
            if time stamp of \(b_{j}=\) latest time stamp of \(a_{i}\) then
                /* relevant for network scheme only;
                    the track has got a new record, splitting happening;
                    retrieve the original \(a_{i} \quad * /\)
                        let \(a_{i}=A^{\prime}[i]\);
                /* make a new track */
                    append \(a_{i}\) to A;
            end
            add \(i\) to \(R^{+}\);
            add \(j\) to \(C^{+}\);
            add \(i\) to \(R^{-}\);
            add \(j\) to \(C^{-}\);
            append \(b_{j}\) to \(a_{i}\);
            let \(M(i, j):=\infty\);
            else
            let \(M(i, j):=\infty\);
            end
        end
    end
```

Fig. 4.5: Algorithm for the nearest neighbor link method.

### 4.5.3 Input data

This step takes as inputs the AR records detected at individual time steps as computed in Detect AR appearances from THR output.

The tracker parameters used:

```
# Int, hours, gap allowed to link 2 records. Should be the time resolution of
# the data.
TIME_GAP_ALLOW=6
# tracking scheme. 'simple': all tracks are simple paths.
# 'full': use the network scheme, tracks are connected by their joint points.
TRACK_SCHEME='simple' # 'simple' / 'full'
# int, max Hausdorff distance in km to define a neighborhood relationship
MAX_DIST_ALLOW=1200 # km
# int, min duration in hrs to keep a track.
MIN_DURATION=24
# int, min number of non-relaxed records in a track to keep a track.
MIN_NONRELAX=1
# whether to plot linkage schematic plots or not
SCHEMATIC=True
```


### 4.5.4 Usage in Python scripts

The tracking process is handled by the $A R \_t r a c e r . t r a c k A R s() f u n c t i o n: ~$

```
from ipart.AR_tracer import trackARs
from ipart.AR_tracer import readCSVRecord
ardf=readCSVRecord(RECORD_FILE)
track_list=trackARs(ardf, TIME_GAP_ALLOW, MAX_DIST_ALLOW,
    track_scheme=TRACK_SCHEME, isplot=SCHEMATIC, plot_dir=plot_dir)
```

where

- RECORD_FILE is the path to the CSv file saving the individual AR records. Refer to this notebook for more information on the creation of this file.
- ardf is a pandas. DataFrame object containing the AR records at individual time points.
- track_list is a list of AR objects, each stores a sequence of AR records that form a single track. The data attribute of the AR object is a pandas. DataFrame object, with the same columns as shown in $A R$ records DataFrame.

After this, one can optionally perform a filtering on the obtained tracks, using AR_tracer.filterTracks(), to remove, for instance, tracks that do not last for more than 24 hours:

```
from ipart.AR_tracer import filterTracks
track_list=filterTracks(track_list, MIN_DURATION, MIN_NONRELAX)
```


### 4.5.5 Example output

The resultant AR track can be visualized using the following snippet:

```
from ipart.utils import plot
import cartopy.crs as ccrs
latax=np.arange (0, 90)
lonax=np.arange(80, 440) # degree east, shifted by 80 to ensure monotonically
\hookrightarrowincreasing axis
plot_ar=track_list[6] # plot the 7th track in list
figure=plt.figure(figsize=(12,6),dpi=100)
ax=figure.add_subplot(111, projection=ccrs.Platecarree())
plotplotARTrack(plot_ar,latax,lonax,ax, full=True)
```


## See also:

```
utils.plot.plotARTrack().
```

The output figure looks like Fig. 4.6 below.


Fig. 4.6: Locations of a track labelled "198424" found in year 1984. Black to yellow color scheme indicates the evolution.

### 4.5.6 Dedicated Python script

You can use the scripts/trace_ARs.py script for AR tracking process in production.

Note: Unlike the AR occurrence detection process, this tracking process is time-dependent and therefore can not be
paralleized. Also, if you divide the detection process into batches, e.g. one for each year, you may want to combine
the output csv records into one big data file, and perform the tracking on this combined data file. This would prevent a track lasting from the end of one year into the next from being broken into 2 .

### 4.5.7 Notebook example

An example of this process is given in this notebook.

## chapter 5

## ipart module contents

### 5.1 Documentation page for thr.py

## Perform THR computation on IVT data

Author: guangzhi XU (xugzhi1987@gmail.com) Update time: 2020-07-22 09:58:36.
thr.THR (ivtNV, kernel, oroNV=None, high_terrain=600, verbose=True)
Perform THR filtering process on 3d data

## Parameters

- ivtNV (NCVAR) - 3D or 4D input IVT data, with dimensions (time, lat, lon) or (time, level, lat, lon).
- kernel (list or tuple) - list/tuple of integers specifying the shape of the kernel/structuring element used in the gray erosion process.


## Keyword Arguments

- oroNV (NCVAR or None) - 2D array, surface orographic data in meters. This optional surface height info is used to perform a separate reconstruction computation for areas with high elevations, and the results can be used to enhance the continent-penetration ability of landfalling ARs. Sensitivity in landfalling ARs is enhanced, other areas are not affected. Needs to have compatible (lat, lon) shape as <ivt>. If None, omit this process and treat areas with different heights all equally.
New in v2.0.
- high_terrain (float) - minimum orographic height (in m) to define as high terrain area, within which a separate reconstruction is performed. Only used if <oroNV> is not None.

New in v2.0.

- verbose (bool) - print some messages or not.


## Returns

ivtNV (NCVAR) - 3D or 4D array, input <ivt>. ivtrecNV (NCVAR): 3D or 4D array, the reconstruction component from the

THR process.
ivtanoNV (NCVAR): 3D or 4D array, the difference between input <ivt> and <ivtrecNV>.
thr.getAttrDict (ref_var, component)
Prepare a metadata dict from a reference variable

## Parameters

- ref_var(ipart.utils.NCVAR) - NCVAR obj from which meta is obtained and modified.
- component (str) - 'reconstruction' or 'anomaly'. Strings as modifiers to modify attributes in <ref_var>.
Returns attdict (dict) - attribute dictionary.
thr.rotatingTHR (filelist, varin, kernel, outputdir, oroNV=None, selector=None, high_terrain=600, verbose=True)
Compute time filtering on data in different files.


## Parameters

- filelist (list) - list of abs paths to data files. User is responsible to make sure files in list have correct chronological order. Note that time axis in data files should be the 1st axis.
- varin (str) - variable id in files.
- kernel (list or tuple) - list/tuple of integers specifying the shape of the kernel/structuring element used in the gray erosion process.
- selector (utils.funcs.Selector) - selector obj to select subset of data.
- outputdir (str) - path to folder to save outputs.


## Keyword Arguments

- oroNV (NCVAR) - 2D array, surface orographic data in meters. This additional surface height info is used to perform a separate reconstruction computation for areas with high elevations, and the results can be used to enhance the continent-penetration ability of landfalling ARs. Sensitivity in landfalling ARs is enhanced, other areas are not affected. Needs to have compatible shape as <ivt>. If None, omit this process and treat areas with different heights all equally.
New in v2.0.
- high_terrain (float) - minimum orographic height to define as high terrain area, within which a separate reconstruction is performed. Only used if <oroNV> is not None.
New in v2.0.
- verbose (bool) - print some messages or not.

Designed to perform temporal filtering on data that are too large to fit into memory, e.g. high-resolution data across multiple decades.

Function will read in 2 files at once, call the filtering function on the concatenated data, and shift 1 step to the next 2 files. If at the begining, pad 0 s to the left end. If in the mid, pad filtered data in the mid of the concatenated data in the previous step. If at the end, pad 0 s to the right end.

The filtering function <func> is assumed to apply a filtering window with odd length $n$, and truncates ( $\mathrm{n}-1$ )/2 points from both ends. If the function doesn't truncate data, will raise an exception.

### 5.2 Documentation page for AR_detector.py

$A R$ detection functions.
Author: guangzhi XU (xugzhi1987@gmail.com) Update time: 2020-07-22 09:27:22.
AR_detector.applyCropIdx (slab, cropidx)
Cut out a bounding box from given 2 d slab given corner indices

## Parameters

- slab (NCVAR or ndarray) - 2D array to cut a box from.
- cropidx (tuple) - (y, x) coordinate indices, output from $\operatorname{cropMask}()$.


## Returns

cropslab (NCVAR or ndarray) -
2D sub array cut from <slab> using <cropidx> as boundary indices.
AR_detector.areaFilt (mask, area, min_area=None, max_area=None, zonal_cyclic=False)
Filter AR binary masks by region areas

## Parameters

- mask (ndarray) - 2D binary mask with detected objects shown as 1s.
- area (ndarray) - 2D map showing grid cell areas in $\mathrm{km}^{\wedge} 2$.


## Keyword Arguments

- min_area (float or None) - if not None, minimum area to filter objects in <mask>.
- max_area (float or None) - if not None, maximum area to filter objects in <mask>.
- zonal_cyclic (bool) - if True, treat zonal boundary as cyclic.

Returns result (ndarray) - 2D binary mask with objects area-filtered.
AR_detector.cart2Spherical ( $x, y, z$, shift_lon)
Convert Cartesian to spherical coordiantes :param $\mathrm{x}, \mathrm{y}, \mathrm{z}: \mathrm{x}-, \mathrm{y}$ - and z - coordiantes. :type $\mathrm{x}, \mathrm{y}, \mathrm{z}$ : float :param shift_lon: longitude to shift so the longitude dimension
starts from this number.

## Returns

result (ndrray) -
$1 \times 3$ array, the columns are the lat-, lon- and r-coordinates. r-coordinates are all 1 s (unit sphere).

AR_detector.cart2Wind (vs, lats, lons)
Convert winds in Cartesian coordinates to $u, v$, inverse to wind2Cart. :param vs: Cartesian representation of the horizontal winds.

Parameters lats,lons (float or ndarray) - latitude and longitude coordinates corresponding to the wind vectors given by $<u>$ and $<v>$.
Returns $u, v$ (float or ndarray) - u- and v-components of horizontal winds.
AR_detector.checkCyclic (mask)
Check binary mask is zonally cyclic or not
Parameters mask (ndarray) - 2d binary array, with 1s denoting existance of a feature.

## Returns

result (bool) -
True if feature in <mask> is zonally cyclic, False otherwise.
AR_detector.computeTheta ( $p 1, p 2$ )
Tangent line to the arc |p1-p2|
Parameters p1, p2 (float) - (lat,lon) coordinates

## Returns

theta (float) -
unit vector tangent at point $<\mathbf{p} 1>$ pointing at $<\mathbf{p} 2>$ on unit sphere.
AR_detector.cropMask (mask, edge=4)
Cut out a bounding box around mask==1 areas

## Parameters

- mask (ndarray) - 2D binary map showing the location of an AR with 1s.
- edge (int) - number of pixels as edge at 4 sides.


## Returns

mask[yl -
$\mathbf{y 2}, \mathbf{x 1}: \mathbf{x 2}$ ] (ndarray): a sub region cut from <mask> surrouding regions with value=1.
$(\mathbf{y} y, \mathbf{x x}): \mathbf{y}$-, $\mathbf{x}$ - indices of the box of the cut region. Can later by used in applyCropIdx(new_slab, (yy,xx)) to crop out the same region from a new array <new_slab>.

AR_detector.crossSectionFlux (mask, quslabNV, qvslabNV, axis_rdp)
Compute setion-wise orientation differences and cross-section fluxes in an AR

## Parameters

- mask (ndarray) - CROPPED (see cropMask and applyCropIdx) 2D binary map showing the location of an AR with 1 s .
- quslab (NCVAR) - CROPPED ( $\mathrm{n} * \mathrm{~m}$ ) 2 D array of u-flux, in $\mathrm{kg} / \mathrm{m} / \mathrm{s}$.
- quslab (NCVAR) - CROPPED ( n * m) 2D array of v-flux, in $\mathrm{kg} / \mathrm{m} / \mathrm{s}$.
- axis_rdp (ndarray) - Nx2 array storing the (lat, lon) coordinates of rdp-simplified AR axis.


## Returns

angles (ndarray) -
2D map with the same shape as <mask>, showing section-wise orientation differences between horizontal flux (as in <quslab>, <qvslab>) and the AR axis of that section. In degrees. Regions outside of AR ( 0 s in <mask>) are masked.
anglesmean (float): area-weighted averaged of <angles> inside <mask>. crossflux (ndarray): 2D map with the same shape as <mask>,
the section-wise cross-section fluxes in the AR, defined as the projection of fluxes onto the AR axis, i.e. flux multiplied by the cos of <angles>.
seg_thetas (list): list of $(\mathbf{x}, \mathbf{y}, \mathbf{z})$ Cartesian coordinates of the tangent vectors along section boundaries.

AR_detector.cyclicLabel (mask, connectivity=1, iszonalcyclic=False)
Label connected region, zonally cyclic version
Parameters mask (ndarray) - 2d binary mask with 1s denoting feature regions.

## Keyword Arguments

- connectivity (int) - 1 or 2 connectivity. 2 probaby won't work for zonally cyclic labelling.
- iszonalcyclic (bool) - doing zonally cyclic labelling or not. If False, call skimage.measure.label(). If True, call skimage.measure.label() for initial labelling, then shift the map zonally by half the zonal length, do another measure.label() to find zonally linked regions. Then use this to update the original labels.


## Returns

result (ndarray) -
2d array with same shape as <mask>. Each connected region in <mask> is given an int label.
AR_detector.determineThresLow (anoslab, sill=0.8)
Determine the threshold for anomalous IVT field, experimental
Parameters anoslab (ndarray) - $(\mathrm{n} * \mathrm{~m}) 2 \mathrm{D}$ anomalous IVT slab, in $\mathrm{kg} / \mathrm{m} / \mathrm{s}$.
Keyword Arguments sill (float) - float in ( 0,1 ). Fraction of max score to define as the 1 st time the score is regarded as reaching stable level.

Returns result (float) - determined lower threshold.
Method of determining the threshold: 1. make a loop through an array of thresholds from 1 to the 99th percentile of <ivtano>. 2. at each level, record the number of pixels $>$ threshold. 3. after looping, pixel number counts will have a curve with a
lower-left elbow. Compute the product of number counts and thresholds P. P will have a peak value around the elbow.
4. choose the 1 st time $P$ reaches the $80 \%$ of $\max (\mathrm{P})$, and pick the corresponding threshold as result.

Largely empirical, but seems to work good on MERRA2 IVT results.
AR_detector.findARAxis (quslab, qvslab, armask_list, costhetas, sinthetas, param_dict, verbose=True)
Find AR axis

## Parameters

- quslab (ndarray) - ( $\mathrm{n} * \mathrm{~m}$ ) 2D u-flux slab, in $\mathrm{kg} / \mathrm{m} / \mathrm{s}$.
- qvslab (ndarray) - ( n * m) 2D v-flux slab, in $\mathrm{kg} / \mathrm{m} / \mathrm{s}$.
- armask_list (list) - list of 2D binary masks, each with the same shape as <quslab> etc., and with 1 s denoting the location of a found AR.
- costhetas (ndarray) - ( $\mathrm{n}^{*} \mathrm{~m}$ ) 2D slab of grid cell shape: $\cos =\mathrm{dx} / \mathrm{sqrt}\left(\mathrm{dx} \wedge 2+\mathrm{dy}{ }^{\wedge} 2\right)$.
- sinthetas (ndarray) - ( $\left.\mathrm{n}^{*} \mathrm{~m}\right)$ 2D slab of grid cell shape: $\sin =d y / s q r t\left(d x^{\wedge} 2+d y^{\wedge} 2\right)$.
- param_dict $(d i c t)$ - a dict containing parameters controlling the detection process. Keys of the dict: 'thres_low', 'min_area', 'max_area', 'max_isoq', 'max_isoq_hard', 'min_lat', 'max_lat', 'min_length', 'min_length_hard', 'rdp_thres', 'fill_radius', 'single_dome', 'max_ph_ratio', 'edge_eps'. See the doc string of findARs() for more.
Keyword Arguments verbose (bool) - print some messages or not.


## Returns

axes (list) -
list of AR axis coordinates. Each coordinate is defined as a $N x 2$ ndarray storing ( $\mathrm{y}, \mathrm{x}$ ) indices of the axis (indices defined in the matrix of corresponding mask in <armask_list>.)
axismask (ndarray): 2D binary mask showing all axes in <axes> merged into one map.
New in v2.0.
AR_detector.findARs (ivt, ivtrec, ivtano, qu, qv, lats, lons, times=None, ref_time='days since 1900-01-01', thres_low $=1$, min_area $=500000.0$, max_area $=18000000.0$, min_LW=2, min_lat=20, max_lat=80, min_length=2000, min_length_hard $=1500$, rdp_thres $=2$, fill_radius=None, single_dome=False, max_ph_ratio=0.6, edge_eps=0.4, zonal_cyclic=False, verbose=True)
Find ARs from THR results, get all results in one go.

## Parameters

- ivt (ndarray) - 3D or 4D input IVT data, with dimensions (time, lat, lon) or (time, level, lat, lon).
- ivtrec (ndarray) - 3D or 4D array, the reconstruction component from the THR process.
- ivtano (ndarray) - 3D or 4D array, the difference between input <ivt> and <ivtrec>.
- qu (ndarray) - 3D or 4D array, zonal component of integrated moisture flux.
- qv (ndarray) - 3D or 4D array, meridional component of integrated moisture flux.
- lats (ndarray) - 1D, latitude coordinates, the length needs to be the same as the lat dimension of <ivt>.
- lons (ndarray) - 1D, longitude coordinates, the length needs to be the same as the lon dimension of <ivt>.


## Keyword Arguments

- times (list or array) - time stamps of the input data as a list of strings, e.g. ['2007-01-01 06:00:00', '2007-01-01 12:00', ...]. Needs to have the same length as the time dimension of <ivt>. If None, default to create a dummy 6-hourly time axis, using <ref_time> as start, with a length as the time dimension of <ivt>.
- ref_time (str) - reference time point to create dummy time axis, if no time stamps are given in <times>.
- thres_low (float or None) - $\mathrm{kg} / \mathrm{m} / \mathrm{s}$, define AR candidates as regions >= this anomalous ivt level. If None is given, compute a threshold based on anomalous ivt data in <ivtano> using an empirical method:

1. make a loop through an array of thresholds from 1 to the 99th percentile of <ivtano>. 2. at each level, record the number of pixels $>$ threshold. 3. after looping, pixel number counts will have a curve with a
lower-left elbow. Compute the product of number counts and thresholds P. P will have a peak value around the elbow.
2. choose the 1 st time $P$ reaches the $80 \%$ of $\max (\mathrm{P})$, and pick the corresponding threshold as result.

- min_area $(f l o a t)-\mathrm{km}^{\wedge} 2$, drop AR candidates smaller than this area.
- max_area $(f l o a t)-\mathrm{km}^{\wedge} 2$, drop AR candidates larger than this area.
- min_LW (float) - minimum length/width ratio.
- min_lat (float) - degree, exclude systems whose centroids are lower than this latitude. NOTE this is the absolute latitude for both NH and SH. For SH, systems with centroid latitude north of - min_lat will be excluded.
- max_lat (float) - degree, exclude systems whose centroids are higher than this latitude. NOTE this is the absolute latitude for both NH and SH. For SH, systems with centroid latitude south of - max_lat will be excluded.
- min_length (float) - km, ARs shorter than this length is treated as relaxed.
- min_length_hard (float) - km, ARs shorter than this length is discarded.
- rdp_thres (float) - degree lat/lon, error when simplifying axis using rdp algorithm.
- fill_radius (int or None) - number of grids as radius to fill small holes in AR contour. If None, computed as
$\max (1, \operatorname{int}(4 * 0.75 /$ RESO $))$
where RESO is the approximate resolution in degrees of lat/lon, estimated from <lat>, <lon>.
- single_dome (bool) - do peak partition or not, used to separate systems that are merged together with an outer contour.
- max_ph_ratio (float) - max prominence/height ratio of a local peak. Only used when single_dome=True
- edge_eps (float) - minimal proportion of flux component in a direction to total flux to allow edge building in that direction.
- zonal_cyclic (bool) - if True, treat the data as zonally cyclic (e.g. entire hemisphere or global). ARs covering regions across the longitude bounds will be correctly treated as one. If your data is not zonally cyclic, or a zonal shift of the data can put the domain of interest to the center, consider doing the shift and setting this to False, as it will save computations.
- verbose (bool) - print some messages or not.


## Returns

time_idx (list) - indices of the time dimension when any AR is found. labels_allNV (NCVAR): 3D array, with dimension
(time, lat, lon). At each time slice, a unique int label is assign to each detected AR at that time, and the AR region is filled out with the label value in the (lat, lon) map.
angles_allNV (NCVAR): 3D array showing orientation differences between AR axes and fluxes, for all ARs. In degrees.
crossfluxes_allNV (NCVAR): 3D array showing cross- sectional fluxes in all ARs. In $\mathrm{kg} / \mathrm{m} / \mathrm{s}$.
result_df (DataFrame): AR record table. Each row is an AR, see code in getARData() for columns.

## See also:

findARsGen(): generator version, yields results at time points separately.
AR_detector.findARsGen (ivt, ivtrec, ivtano, qu, qv, lats, lons, times=None, ref_time='days since 1900-01-01', thres_low $=1$, min_area $=500000.0$, max_area $=18000000.0$, min_LW=2, min_lat=20, max_lat=80, min_length $=2000$, min_length_hard $=1500, \quad$ rdp_thres $=2, \quad$ fil_radius $=$ None, $\quad$ single_dome $=$ False, max_ph_ratio=0.6, edge_eps $=0.4, \quad$ zonal_cyclic $=$ False, verbose=True)
Find ARs from THR results, generator version

## Parameters

- ivt (ndarray) - 3D or 4D input IVT data, with dimensions (time, lat, lon) or (time, level, lat, lon).
- ivtrec (ndarray) - 3D or 4D array, the reconstruction component from the THR process.
- ivtano (ndarray) - 3D or 4D array, the difference between input <ivt> and <ivtrec>.
- qu (ndarray) - 3D or 4D array, zonal component of integrated moisture flux.
- qv (ndarray) - 3D or 4D array, meridional component of integrated moisture flux.
- lats (ndarray) - 1D, latitude coordinates, the length needs to be the same as the lat dimension of <ivt>.
- lons (ndarray) - 1D, longitude coordinates, the length needs to be the same as the lon dimension of <ivt>.


## Keyword Arguments

- times (list or array) - time stamps of the input data as a list of strings, e.g. ['2007-01-01 06:00:00', '2007-01-01 12:00', ...]. Needs to have the same length as the time dimension of <ivt>. If None, default to create a dummy 6-hourly time axis, using <ref_time> as start, with a length as the time dimension of <ivt>.
- ref_time (str) - reference time point to create dummy time axis, if no time stamps are given in <times>.
- thres_low (float or None) - $\mathrm{kg} / \mathrm{m} / \mathrm{s}$, define AR candidates as regions >= this anomalous ivt level. If None is given, compute a threshold based on anomalous ivt data in <ivtano> using an empirical method:

1. make a loop through an array of thresholds from 1 to the 99th percentile of <ivtano>.
2. at each level, record the number of pixels $>$ threshold. 3. after looping, pixel number counts will have a curve with a
lower-left elbow. Compute the product of number counts and thresholds P. P will have a peak value around the elbow.
3. choose the 1st time $P$ reaches the $80 \%$ of $\max (\mathrm{P})$, and pick the corresponding threshold as result.

- min_area $(f l o a t)-\mathrm{km}^{\wedge} 2$, drop AR candidates smaller than this area.
- max_area $($ float $)-\mathrm{km} \wedge 2$, drop AR candidates larger than this area.
- min_LW (float) - minimum length/width ratio.
- min_lat (float) - degree, exclude systems whose centroids are lower than this latitude. NOTE this is the absolute latitude for both NH and SH. For SH, systems with centroid latitude north of - min_lat will be excluded.
- max_lat (float) - degree, exclude systems whose centroids are higher than this latitude. NOTE this is the absolute latitude for both NH and SH. For SH, systems with centroid latitude south of - max_lat will be excluded.
- min_length (float) - km, ARs shorter than this length is treated as relaxed.
- min_length_hard (float) - km, ARs shorter than this length is discarded.
- rdp_thres (float) - degree lat/lon, error when simplifying axis using rdp algorithm.
- fill_radius (int or None) - number of grids as radius to fill small holes in AR contour. If None, computed as $\max (1, \operatorname{int}(4 * 0.75 / \mathrm{RESO}))$
where RESO is the approximate resolution in degrees of lat/lon, estimated from <lat>, <lon>.
- single_dome (bool) - do peak partition or not, used to separate systems that are merged together with an outer contour.
- max_ph_ratio (float) - max prominence/height ratio of a local peak. Only used when single_dome=True
- edge_eps (float) - minimal proportion of flux component in a direction to total flux to allow edge building in that direction.
- zonal_cyclic (bool) - if True, treat the data as zonally cyclic (e.g. entire hemisphere or global). ARs covering regions across the longitude bounds will be correctly treated as one. If your data is not zonally cyclic, or a zonal shift of the data can put the domain of interest to the center, consider doing the shift and setting this to False, as it will save computations.
- verbose (bool) - print some messages or not.


## Returns

ii (int) - index of the time dimension when any AR is found. timett_str (str): time when any AR is found, in string format. labelsNV (NCVAR): 2D array, with dimension
(lat, lon). A unique int label is assign to each detected AR at the time, and the AR region is filled out with the label value in the (lat, lon) map.
anglesNV (NCVAR): 2D array showing orientation differences between AR axes and fluxes, for all ARs. In degrees.
crossfluxesNV (NCVAR): 2D array showing cross- sectional fluxes in all ARs. In $\mathrm{kg} / \mathrm{m} / \mathrm{s}$. ardf (DataFrame): AR record table. Each row is an AR, see code in getARData() for columns.

## See also:

findARs(): collect and return all results in one go.
New in v2.0.
AR_detector. getARAxis ( $g$, quslab, quslab, mask)
Find AR axis from AR region mask

## Parameters

- $\mathbf{g}$ (networkx.DiGraph) - directed planar graph constructed from AR mask and flows. See maskToGraph().
- quslab (ndarray) - 2D map of u-flux.
- qvslab (ndarray) - 2D map of v-flux.
- mask (ndarray) - 2D binary map showing the location of an AR with 1s.


## Returns

## path (ndarray) -

$\mathbf{N x} 2$ array storing the AR axis coordinate indices in ( $y, x$ ) format.
axismask (ndarray): 2D binary map with same shape as <mask>, with grid cells corresponding to coordinates in <path> set to 1 s .

AR_detector.getARData (slab, quslab, qvslab, anoslab, quano, qvano, areas, lats, lons, mask_list, axis_list, timestr, param_dict)
Fetch AR related data

## Parameters

- slab (ndarray) - (n * m) 2D array of IVT, in $\mathrm{kg} / \mathrm{m} / \mathrm{s}$.
- quslab (ndarray) - ( $\mathrm{n} * \mathrm{~m}$ ) 2D array of u-flux, in $\mathrm{kg} / \mathrm{m} / \mathrm{s}$.
- quslab (ndarray) - (n*m) 2D array of v-flux, in $\mathrm{kg} / \mathrm{m} / \mathrm{s}$.
- anoslab (ndarray) - ( $\mathrm{n} * \mathrm{~m}$ ) 2D array of IVT anomalies, in $\mathrm{kg} / \mathrm{m} / \mathrm{s}$.
- quano (ndarray) - (n * m) 2D array of u-flux anomalies, in $\mathrm{kg} / \mathrm{m} / \mathrm{s}$.
- quano (ndarray) - ( $\mathrm{n} * \mathrm{~m}$ ) 2D array of v-flux anomalies, in $\mathrm{kg} / \mathrm{m} / \mathrm{s}$.
- areas (ndarray) - $(\mathrm{n} * \mathrm{~m})$ 2D grid cell area slab, in $\mathrm{km}{ }^{\wedge} 2$.
- lats (ndarray) - 1d array, latitude coordinates.
- lons (ndarray) - 1d array, longitude coordinates.
- mask_list (list) - list of 2D binary masks, each with the same shape as <anoslab> etc., and with 1 s denoting the location of a found AR.
- axis_list (list) - list of AR axis coordinates. Each coordinate is defined as a Nx2 ndarray storing ( $\mathrm{y}, \mathrm{x}$ ) indices of the axis (indices defined in the matrix of corresponding mask in <masks>.)
- timestr (str) - string of time snap.
- param_dict $(d i c t)$ - a dict containing parameters controlling the detection process. Keys of the dict: 'thres_low', 'min_area', 'max_area', 'max_isoq', 'max_isoq_hard', 'min_lat', 'max_lat', 'min_length', 'min_length_hard', 'rdp_thres', 'fill_radius', 'single_dome', 'max_ph_ratio', 'edge_eps'. See the doc string of findARs() for more.


## Returns

labelsNV (NCVAR) -
$(\mathbf{n} * \mathbf{m}) 2 \mathrm{D}$ int map showing all ARs at current time. Each AR is labeled by an int label, starting from 1. Background is filled with 0s.
anglesNV (NCVAR): ( $\mathbf{n} * \mathbf{m}$ ) 2D map showing orientation differences between AR axes and fluxes, for all ARs. In degrees.
crossfluxesNV (NCVAR): ( $\mathbf{n} * \mathbf{m}$ ) 2D map showing cross- sectional fluxes in all ARs. In $\mathrm{kg} / \mathrm{m} / \mathrm{s}$.
df (pandas.DataFrame): AR record table. Each row is an AR, see code below for columns.
AR_detector.getNormalVectors (point_list, idx)
Get the normal vector and the tagent vector to the plane dividing 2 sections along the AR axis.

## Parameters

- point_list (list) - list of (lat, lon) coordinates.
- idx (int) - index of the point in <point_list>, denoting the point in question.


## Returns

normi (tuple) -
the ( $\mathbf{x}, \mathbf{y}, \mathbf{z}$ ) Cartesian coordinate of the unit normal vector, at the point denoted by <idx>, on the Earth surface. This is the normal vector to the plane spanned by the vector Theta and $P$. Where $P$ is the vector pointing to the point in question (point_list[idx]), and Theta is the tangent vector evenly dividing the angle formed by $\langle\mathrm{P}, \mathrm{P} 1\rangle$, and $\langle\mathrm{P}, \mathrm{P} 2\rangle$. Where $\mathrm{P} 1, \mathrm{P} 2$ are 2 points on both side of $P$.
thetai (tuple): the ( $\mathbf{x}, \mathbf{y}, \mathbf{z}$ ) Cartesian coordinate of the tangent vector Theta above.
AR_detector.insertCropSlab (shape, cropslab, cropidx)
Insert the cropped sub-array back to a larger empty slab

## Parameters

- shape (tuple) - ( $\mathrm{n}, \mathrm{m}$ ) size of the larger slab.
- cropslab (ndarray) - 2D array to insert.
- cropidx (tuple) - (y, x) coordinate indices, output from cropMask(), defines where <cropslab> will be inserted into.


## Returns

result (ndarray) -
2D slab with shape ( $\mathbf{n}, \mathrm{m}$ ), an empty array with a box at <cropidx> replaced with data from <cropslab>.
AR_detector.maskToGraph (mask, quslab, qvslab, costhetas, sinthetas, edge_eps, connectivity=2)
Create graph from AR mask

## Parameters

- mask (ndarray) - 2D binary map showing the location of an AR with 1 s .
- quslab (ndarray) - 2D map of u-flux.
- qvslab (ndarray) - 2D map of v-flux.
- costhetas (ndarray) - ( $\left.\mathrm{n}^{*} \mathrm{~m}\right) 2 \mathrm{D}$ slab of grid cell shape: $\cos =\mathrm{dx} / \mathrm{sqrt}\left(\mathrm{dx}^{\wedge} 2+\mathrm{dy}{ }^{\wedge} 2\right)$.
- sinthetas (ndarray) - (n* m) 2D slab of grid cell shape: $\sin =d y / s q r t\left(d x \wedge 2+d y^{\wedge} 2\right)$.
- edge_eps (float) - float in $(0,1)$, minimal proportion of flux component in a direction to total flux to allow edge building in that direction. Defined in Global preamble.
- connectivity (int) - 1 or 2 . 4 - or 8 - connectivity in defining neighbor- hood relationship in a 2D square grid.


## Returns

g (networkx.DiGraph) -
directed planar graph constructed from AR mask and flows.
AR_detector.partPeaks (cropmask, cropidx, orislab, area, min_area, max_ph_ratio, fill_radius)
Separate local maxima by topographical prominence, watershed version

## Parameters

- cropmask (ndarray) - 2D binary array, defines regions of local maxima.
- cropidx (tuple) - (y, x) coordinate indices, output from cropMask().
- orislab (ndarray) - 2D array, giving magnitude/height/intensity values defining the topography.
- area (ndarray) - ( $\mathrm{n} * \mathrm{~m}$ ) 2D grid cell area slab, in $\mathrm{km}{ }^{\wedge} 2$.
- min_area (float) - $\mathrm{km}^{\wedge} 2$, drop AR candidates smaller than this area.
- max_ph_ratio (float) - maximum peak/height ratio. Local peaks with a peak/height ratio larger than this value is treated as an independent peak.
- fill_radius (int) - number of grids as radius to further separate peaks.


## Returns

result (ndarray) -
2D binary array, similar as the input <cropmask> but with connected peaks (if any) separated so that each connected region (with 1 s ) denotes an independent local maximum.

AR_detector.partPeaksOld (cropmask, cropidx, orislab, max_ph_ratio)
Separate local maxima by topographical prominence

## Parameters

- cropmask (ndarray) - 2D binary array, defines regions of local maxima.
- cropidx (tuple) - (y, x) coordinate indices, output from cropMask().
- orislab (ndarray) - 2D array, giving magnitude/height/intensity values defining the topography.
- max_ph_ratio (float) - maximum peak/height ratio. Local peaks with a peak/height ratio larger than this value is treated as an independent peak.


## Returns

result (ndarray) -
2D binary array, similar as the input <cropmask> but with connected peaks (if any) separated so that each connected region (with 1s) denotes an independent local maximum.
AR_detector.plotGraph (graph, ax=None, show=True)
Helper func to plot the graph of an AR coordinates. For debugging. :param graph: networkx Graph obj to visualize. :type graph: networkx.Graph

## Keyword Arguments

- $\mathbf{a x}$ (matplotlib axis obj)-axis to plot on. If None, create a new one.
- show (bool) - whether to show the plot or not.

AR_detector.prepareMeta (lats, lons, times, ntime, nlat, nlon, ref_time='days since 1900-01-01', verbose=True)
Prepare metadata for AR detection function calls

## Parameters

- lats (ndarray) - 1D, latitude coordinates, the length needs to equal <nlat>.
- lons (ndarray) - 1D, longitude coordinates, the length needs to equal <nlon>.
- times (list or array) - time stamps of the input data as a list of strings, e.g. ['2007-01-01 06:00:00', '2007-01-01 12:00', ...]. Needs to have the a length of <ntime>.
- ntime (int) - length of the time axis, should equal the length of <times>.
- nlat (int) - length of the latitude axis, should equal the length of <lats>.
- nlon (int) - length of the longitude axis, should equal the length of <lons>.


## Keyword Arguments

- ref_time (str) - reference time point to create time axis.
- verbose (bool) - print some messages or not.


## Returns

timeax (list) - a list of datetime objs. areamap (ndarray): grid cell areas in $\mathrm{km}^{\wedge} 2$, with shape (<nlat> $x<n l o n>$ ). costhetas (ndarray): ratios of $d x / s q r t\left(d x \wedge 2+d y^{\wedge} 2\right.$ ) for all grid cells. with shape (<nlat> $\mathrm{x}<$ nlon>).
sinthetas (ndarray): ratios of $\mathbf{d y} / \mathbf{s q r t}\left(\mathrm{dx}^{\wedge} \mathbf{2}+\mathrm{dy}^{\wedge} \mathbf{2}\right)$ for all grid cells. with shape (<nlat> x <nlon>).
reso (float): (approximate) horizontal resolution in degrees of lat/lon estimate from <lats> and <lons>.

New in v2.0.

## AR_detector.save2DF (result_dict)

Save AR records to a pandas DataFrame
Parameters result_dict (dict) - key: time str in 'yyyy-mm-dd hh:00' value: pandas dataframe. See getARData().

## Returns

result_df(pandas.DataFrame) -
AR record table containing records from multiple time steps sorted by time.
AR_detector.spherical2Cart (lat, lon)
Convert spherical to Cartesian coordiantes :param lat,lon: latitude and longitude coordinates. :type lat,lon: float or ndarray

## Returns

result (ndarray) -
Nx3 array, the columns are the $\mathbf{x}-, \mathbf{y}$ - and $\mathbf{z}$ - coordinates.

AR_detector.uvDecomp ( $u 0, v 0, i 1, i 2$ )
Decompose background-transient components of $u-, v$ - fluxes

## Parameters

- u0 (ndarray) - nd array of total u-flux.
- v0 (ndarray) - nd array of total v-flux.
- i1 (ndarray) - nd array of the reconstruction component of IVT.
- i2 (ndarray) - nd array of the anomalous component of IVT (i2 = IVT - i1).


## Returns

ul (ndarray) -
nd array of the u-flux component corresponding to <i1>, i.e. the background component.
$\mathbf{v 1}$ (ndarray): nd array of the $\mathbf{v}$-flux component corresponding to <i1>, i.e. the background component.
$\mathbf{u} 2$ (ndarray): nd array of the $u$-flux component corresponding to <i2>, i.e. the transient component.
$\mathbf{v} 2$ (ndarray): nd array of the $\mathbf{v}$-flux component corresponding to <i2>, i.e. the transient component.

AR_detector.wind2Cart ( $u$, $v$, lats, lons)
Convert $\mathrm{u}, \mathrm{v}$ winds to Cartesian, consistent with spherical2Cart.

## Parameters

- u,v(float or ndarray) - u- and v-components of horizontal winds.
- lats,lons (float or ndarray) - latitude and longitude coordinates corresponding to the wind vectors given by $\langle u\rangle$ and $\langle v\rangle$.


## Returns

vs (float or ndarray) -
Cartesian representation of the horizontal winds.

### 5.3 Documentation page for AR_tracer.py

Functions to compute Hausdorff distance between AR axes pairs and link ARs across time steps to form tracks. Author: guangzhi XU (xugzhi1987@gmail.com) Update time: 2020-06-05 22:46:19.

```
class AR_tracer.AR(id,data)
```

Ojbect representing an AR entity
Hausdorff (lats, lons)
Compute modified Hausdorff distance from the lastest record to given axis

## Parameters

- lats (ndarray) - 1d array, the target axis's latitude coordinates.
- lons (ndarray) - 1d array, the target axis's longitude coordinates.

Returns float - modified Hausdorff distance from this AR to the given axis.
__init (id, data)

## Parameters

- id (int) - a numeric id for each AR.
- data (pandas. DataFrame) - DataFrame storing an AR's records.


## anchor_lats

$1 d$ array, get the latitude coordinates from roughly evenly spaced points from the AR axis.
Type ndarray

## anchor_lons

1d array, get the longitude coordinates from roughly evenly spaced points from the AR axis.
Type ndarray

## append (ar)

Add new records to the AR track
backwardHausdorff (lats, lons)
Compute backward Hausdorff distance from the lastest record to given axis

## Parameters

- lats (ndarray) - 1d array, the target axis's latitude coordinates.
- lons (ndarray) - 1d array, the target axis's longitude coordinates.

Returns float - backward Hausdorff distance from this AR to the given axis.
coor
(Nx3) ndarray, (time, lat_centroid, lon_centroid) coordinates of an AR track.
Type ndarray

## duration

track duration in hours.
Type int

## forwardHausdorff(lats, lons)

Compute forward Hausdorff distance from the lastest record to given axis

## Parameters

- lats (ndarray) - 1d array, the target axis's latitude coordinates.
- lons (ndarray) - 1d array, the target axis's longitude coordinates.

Returns float - forward Hausdorff distance from this AR to the given axis.

## latest

the AR record of the latest time point in an AR track.
Type Series

## lats

1d array, the latitude coordinates of the AR axis in the latest record in an AR's track.
Type ndarray
lons
1 d array, the longitude coordinates of the AR axis in the latest record in an AR's track.
Type ndarray
rdp_lats
1d array, the latitude coordinates of the simplified AR axis in the latest record in an AR's track.

Type ndarray
rdp_lons
1d array, the longitude coordinates of the simplified AR axis in the latest record in an AR's track.
Type ndarray

## times

sorted time stamps of an AR track.
Type Series
AR_tracer.filterTracks (tr_list, min_duration, min_nonrelax, verbose=True)
Filter tracks

## Parameters

- tr_list (list) - list of AR objects, found tracks.
- min_duration $(i n t)$ - min duration in hrs to keep a track.
- min_nonrelax (int) - min number of non-relaxed records in a track to keep a track.

Keyword Arguments verbose (b○○l) - print some messages or not.
Returns tr_list (list) - list of AR objects, filtered tracks.
Tracks that are filtered:

- tracks that are too short, controlled by 'min_duration'
- tracks that consist of solely relaxed records.

AR_tracer.forwardHausdorff(lats1, lons1, lats2, lons2)
Compute forward Hausdorff distance betweem 2 tracks

## Parameters

- lats1 (list or 1D array)- latitudes of track1.
- lons1 (list or 1D array)- longitudes of track1.
- lats2 (list or 1D array)- latitudes of track2.
- lons2 (list or 1D array)-longitudes of track2.

Returns forward Hausdorff distance in km.
AR_tracer.getAnchors (arr, num_anchors=7)
Get anchor points along from an axis.
Parameters arr (ndarray) - 1D array from which to sample the anchor points.
Returns (ndarray) - 1D array of the sampled anchor points from <arr>.
AR_tracer.getDistMatrix(tr_list, newlats, newlons)
Compute distance matrix among track axis anchors

## Parameters

- tr_list (list) - list of AR objs, existing systems at time $\mathrm{t}=\mathrm{t}$.
- newlats (list or 1D array)- latitudes at $\mathrm{t}=\mathrm{t}+1$.
- newlons (list or $1 D$ array) - longitudes at $t=t+1$.


## Returns

dists (ndarray) -
n*m matrix consisting distances between existing and new tracks. Rows as new records at tnow, columns as existing tracks.
AR_tracer.matchCenters (tr_list, newrec, time_gap_allow, max_dist_allow, track_scheme='simple', isplot=False, plot_dir=None, verbose=True)
Match and link nearby centers at 2 consecutive time steps

## Parameters

- tr_list (list) - list of AR objs, existing systems at time $\mathrm{t}=\mathrm{t}$.
- newrec (DataFrame) - new center data at time $t=t+1$.
- time_gap_allow (int) - max allowed gap between 2 records, in number of hours.
- max_dist_allow (float) - max allowed Hausdorff distance allowed between 2 records, in km.


## Keyword Arguments

- track_scheme (str) - tracking scheme. 'simple': all tracks are simple
- 'full' (paths.) - use the network scheme, tracks are connected by their
- points. (joint)-
- isplot (bool) - create schematic plot or not.
- plot_dir (str) - folder to save schematic plot. Only used if isplot=True.
- verbose (bool) - print some messages or not.


## Returns

tr_list (list) -
list of AR objs, ARs with new matching records appended at the end.
allocated_recs (list): list of ints, ids of new records that are attributed to existing systems during the process.
Matching is based on geo-distances and uses nearest neighbour strategy.
AR_tracer.plotHD ( $y 1, x 1, y 2, x 2$, timelabel $=$ None, linkflag $="$, a $x=$ None, show=True $)$
Plot Hausdorff links

## Parameters

- $\mathbf{y} \mathbf{1}, \mathbf{x 1}$ (ndarray) - 1d array, $\mathrm{y}, \mathrm{x}$ coordinates of AR axis A.
- $\mathbf{y} \mathbf{2 , \mathbf { x } 2 ( n d a r r a y ) - 1 d ~ a r r a y , ~} \mathbf{y}, \mathbf{x}$ coordinates of AR axis B.


## Keyword Arguments

- timelabel (str or None) - string of the time stamp. If given, plot as subplot title.
- linkflag (str) - a single char to denote the type of linking, used in generated plot. '" for initial linking, ' $M$ ' for a merging, ' $S$ ' for a splitting.
- $\mathbf{a x}(p l t$ axis obj) - if not give, create a new axis to plot with.
- show (bool) - whether to show the figure or not.

AR_tracer.readCSVRecord (abpath_in)
Read in individual AR records from .csv file

Parameters abpath_in (str) - absolute file path to AR record file.
Returns $\operatorname{ardf}$ (pandas.DataFrame) - record saved in DataFrame.
New in v2.0.
AR_tracer.trackARs (record, time_gap_allow, max_dist_allow, track_scheme='simple', isplot=False, plot_dir=None, verbose=True)
Track ARs at consecutive time points to form tracks

## Parameters

- record (DataFrame) - AR records at different time slices.
- time_gap_allow (int) - max allowed gap between 2 records, in number of hours.
- max_dist_allow (float) - max allowed Hausdorff distance allowed between 2 records, in km.


## Keyword Arguments

- track_scheme (str) - tracking scheme. 'simple': all tracks are simple
- 'full' (paths.) - use the network scheme, tracks are connected by their
- points. (joint)-
- isplot (bool) - whether to create schematic plots of linking.
- plot_dir (str) - folder to save schematic plot.
- verbose (bool) - print some messages or not.

Returns finished_list (list) - list of AR objs. Found tracks.

### 5.4 Documentation page for utils/funcs.py

Only some functions from this module are documented. The other undocumented functions are intended as private functions, not to be exposed to the user.

Utility functions
Author: guangzhi XU (xugzhi1987@gmail.com) Update time: 2020-07-22 09:27:36.
utils.funcs.get3DEllipse ( $t, y, x$ )
Get a binary 3D ellipse structuring element

## Parameters

- $t$ (int) - ellipse axis length in the $t(1 s t)$ dimension.
- y (int) - ellipse axis length in the y (2nd) dimension.
- $\mathbf{x}(i n t)$ - ellipse axis length in the $\mathrm{x}(3 \mathrm{rd})$ dimension.
- that the axis length is half the size of the ellipse (Note)-
- that dimension. (in)-


## Returns

result (ndarray) -
3D binary array, with 1 s side the ellipse defined as $(\mathrm{T} / \mathrm{t})^{\wedge} 2+(\mathrm{Y} / \mathrm{y})^{\wedge} 2+(\mathrm{X} / \mathrm{x})^{\wedge} 2<=1$.
utils.funcs.dLatitude (lats, lons, $R=6371000$, verbose $=$ True)
Return a slab of latitudinal increment (meter) delta_y.

## Parameters

- lats (ndarray) - 1d array, latitude coordinates in degrees.
- lons (ndarray) - 1d array, longitude coordinates in degrees.

Keyword Arguments R (float) - radius of Earth;

## Returns

delta_x (ndarray) -
2d array, latitudinal increments. <var>.
utils.funcs.dLongitude (lats, lons, side $=$ ' $c$ ', $R=6371000$ )
Return a slab of longitudinal increment (meter) delta_x.

## Parameters

- lats (ndarray) - 1d array, latitude coordinates in degrees.
- lons (ndarray) - 1d array, longitude coordinates in degrees.


## Keyword Arguments

- side (str)-
' $\mathbf{n}$ ': northern boundary of each latitudinal band;
's': southern boundary of each latitudinal band; 'c': central line of latitudinal band;
$\qquad$
' n '
$\qquad$
1 $\qquad$ 's'
- $\mathbf{R}$ (float) - radius of Earth.

Returns delta_x (ndarray) - 2d array, longitudinal increments.
utils.funcs.readVar (abpath_in, varid)
Read in netcdf variable

## Parameters

- abpath_in (str) - absolute file path to nc file.
- varid (str) - id of variable to read.

Returns var (TransientVariable) - 4d TransientVariable.
NOTE: deprecated, use netCDF4 instead of CDAT.
utils.funcs.getTimeAxis (times, ntime, ref_time='days since 1900-01-01')
Create a time axis

## Parameters

- times (list or tuple or array) - array of datetime objs, or strings, giving the time stamps in the format of 'yyyy-mm-dd HH:MM'. It is assumed to be in chronological order. If None, default to create a dummy time axis, with 6-hourly time step, starting from <ref_time>, with a length of <ntime>.
- ntime (int) - length of the time axis. If <times> is not None, it is checked to insure that length of <times> equals <ntime>. If <times> is None, <ntime> is used to create a dummy time axis with a length of <ntime>.
Keyword Arguments ref_time (str) - reference time point. If <times> is not None, used to create the numerical values of the resulant time axis with this reference time. If <times> is None, used to create a dummy time axis with this reference time.


## Returns

result (list) -
a list of datetime objs. If <times> in not None, the datetime objs are using the provided time stamps. Otherwise, it is a new time series, with 6-hourly time step, starting from <ref_time>, with a length of <ntime>.
New in v2.0.

### 5.5 Documentation page for utils/plot.py

Only some functions from this module are documented. The other undocumented functions are intended as private functions, not to be exposed to the user.

Plotting Functions.
Author: guangzhi XU (xugzhi1987@gmail.com) Update time: 2020-07-22 09:27:30.
utils.plot.plot2 (var, method, ax, legend='global', xarray=None, yarray=None, title=None, latlon=True, latlongrid=False, fill_color='0.8', legend_ori='horizontal', clean=False, iscartopy=True, fix_aspect $=$ True, verbose $=$ True )
A helper function for quickly create $2 \bar{D}$ plots

## Parameters

- $\operatorname{var}(N C V A R$ or ndarray) - variable to plot. At least 2D.
- method - plotting method, could be an instance of Boxfill, Isofill.
- ax - matplotlib axis obj.


## Keyword Arguments

- legend (str) - location of colorbar. Could be: 'global': all subplots share the colorbar of the 1 st subplot in figure. or 'local': each subplot in figure uses its own colorbar.
- xarray (ndarray) - 1d array, the array values for the x-axis. If None, use the int indices for the x -dimension.
- yarray (ndarray) - 1d array, the array values for the y-axis. If None, use the int indices for the $y$-dimension.
- title (str) - title to plot at subtitle. If None, plot only an alphabetical index.
- latlon (bool) - plot lat/lon axis labels or not.
- latlongrid (bool) - plot lat/lon grid lines or not.
- fill_color - color to fill continent or masked regions.
- legend_ori (str) - 'horizontal' or 'vertical', colorbar orientation.
- clean (bool) - if True, omit axis labels, colorbar, subtitle, continents, boundaries etc.. Useful to overlay plots.
- iscartopy (bool) - plot using cartopy or not. Usually used to force plot as a normal 2 d plot instead of geographical plot using cartopy.
- fix_aspect (bool) - passed to the cartopy plotting function (e.g. contourf()) for control of aspect ratio. NOTE: needs to be deprecated.
utils.plot.plotAR (ardf, ax, lonax)
Helper function to plot the regions and axes of ARs


## Parameters

- ardf (pandas. DataFrame) - table containing AR records.
- $\mathbf{a x}$ (matplotlib axis) - axis to plot onto.
- lonax (ndarray) - 1d array of the longitude axis the plot is using.
utils.plot.plotARTrack (arlist, latax, lonax, ax, full=False, label=None, linestyle='solid', marker=None)
Plot AR tracks


## Parameters

- arlist (list) - list of AR objects to plot.
- latax, lonax (ndarray) - 1darrays giving latitude- and longitude- coordinates of the plotting domain.
- $\mathbf{a x}$ (matplotlib.axis) - axis to plot onto.


## Keyword Arguments

- full (bool) - if True, plot tracks of an AR from its entire lifecycle. if False, plot only the track of the last time step.
- label (str or None) - type of label to label tracks. 'id': label with AR id. 'time': label with time stamp. None: don't put label.
- linestyle (str) - line style to plot the tracks.
- marker (str) - marker to plot track axes.
class utils.plot.Boxfill(vars, zero=1, split=2, max_level=None, min_level=None, ql=None, $q r=$ None, cmap $=$ None, verbose $=$ True )
__init__(vars, zero=1, split=2, max_level=None, min_level=None, ql=None, qr=None, cmap=None, verbose=True)
Return an isofill object with specified color scheme.
Parameters vars - one or a list of variables, from which the <minlevel> and <maxlevel> is obtained. If <vars> has more than 1 variables, then function calculates the minimum and maximum of all variables, thus the color legend would be unified for all subplots. Note that if <max_level> and/or <min_level> are given, they will override the maximum/minimal levels derived from <vars>.


## Keyword Arguments

- zero (int) - controls 0 in created <levels>: $-1: 0$ is NOT allowed to be a level; 1: 0 is permitted to be a level; 2: 0 forced to be a level.
- split (int) - int, control behavior of negative and positive values 0: Do not split negatives and positives, map onto entire range of $[0,1] ; 1$ : split only when vmin $<0$ and vmax $>0$, otherwise map onto entire range of $[0,1]$;

If split is to be applied, negative values are mapped onto first half [0,0.5], and postives onto second half $(0.5,1]$.

2: force split, if vmin<0 and vmax>0, same as $<$ split>==1; If vmin<0 and vmax $<0$, map onto 1 st half $[0,0.5]$; If vmin $>0$ and $v \max >0$, map onto 2nd half $(0.5,1]$.

- max_level, min_level (float) - the max/min limits to be plotted out, values outside the range will be grouped into the last level intervals on both ends.
- ql, $q \mathbf{r}(f l o a t)$ - extreme percentiles for the lower and upper boundaries. Could be one of the values in the list:

```
percents=[0.001,0.005,0.01,0.025,0.05,0.1]
```

E.g. $\mathrm{ql}=0.001 ; \mathrm{qr}=0.005$ means that the $0.1 \%$ percentile will be set to the minimum level; $0.005 \%$ (from the right most, or $99.95 \%$ from the left most) percentil will be set to the maximum level.

If both <ql> and <min_level> are given, use <min_level>. If both <qr> and <max_level> are given, use <max_level>.

- cmap - specify a color map. Could be: 1) None, a default blue-white-red (bwr) color map will be created. 2) a matplotlib cmap obj. 3) a string name of a matplotlib cmap obj, to list of few:
'bwr': blue-white-red 'RdBu': darkred-white-blue 'RdY1Bu': red-yellow-white-blue 'RdYlGn': red-yellow-white-green 'spectral': purple-yellow-cyanblue 'seismic': darkblue-white-darkred 'jet': rainbow darkblue-darkred 'rainbow': rainbow purple-red

Append '_r' to get the reversed colormap.

> Note: <minlevel> and <maxlevel> are better derived using numpy.min() and numpy.max(). MV.min() and MV.max() may have problems. Iso levels are computed using vcs function (mkscale()), and a matplotlib colormap is created (if not given), and the colormap will be changed so positive/negative splits (if required) is achieved.

Update time: 2015-04-27 14:55:33
class utils.plot.Isofill(vars, num=15, zero=1, split=2, max_level=None, min_level=None, $q l=$ None, $q r=$ None, cmap $=$ None, verbose=True)
__init__(vars, num=15, zero=1, split=2, max_level=None, min_level=None, ql=None, qr=None, cmap $=$ None, verbose $=$ True)
Return an isofill object with specified color scheme.
Parameters vars - one or a list of variables, from which the <minlevel> and <maxlevel> is obtained. If <vars> has more than 1 variables, then function calculates the minimum and maximum of all variables, thus the color legend would be unified for all subplots. Note that if <max_level> and/or <min_level> are given, they will override the maximum/minimal levels derived from <vars>.

## Keyword Arguments

- Num (int) - is the (maximum) number of isoline levels;
- zero (int) - controls 0 in created <levels>: $-1: 0$ is NOT allowed to be a level; 1: 0 is permitted to be a level; 2: 0 forced to be a level.
- split (int) - int, control behavior of negative and positive values 0: Do not split negatives and positives, map onto entire range of $[0,1] ; 1$ : split only when vmin $<0$ and vmax $>0$, otherwise map onto entire range of $[0,1]$;

If split is to be applied, negative values are mapped onto first half $[0,0.5]$, and postives onto second half $(0.5,1]$.

2: force split, if vmin<0 and vmax $>\mathbf{0}$, same as $<$ split>==1; If vmin<0 and vmax $<0$, map onto 1 st half $[0,0.5]$; If vmin $>0$ and $v m a x>0$, map onto 2nd half $(0.5,1]$.

- max_level, min_level (float) - the max/min limits to be plotted out, values outside the range will be grouped into the last level intervals on both ends.
- $\mathbf{q l}_{1}$ qr (float) - extreme percentiles for the lower and upper boundaries. Could be one of the values in the list:

```
percents=[0.001,0.005,0.01,0.025,0.05,0.1]
```

E.g. $\mathrm{ql}=0.001 ; \mathrm{qr}=0.005$ means that the $0.1 \%$ percentile will be set to the minimum level; $0.005 \%$ (from the right most, or $99.95 \%$ from the left most) percentil will be set to the maximum level.

If both <ql> and <min_level> are given, use <min_level>. If both <qr> and <max_level> are given, use <max_level>.

- cmap - specify a color map. Could be: 1) None, a default blue-white-red (bwr) color map will be created. 2) a matplotlib cmap obj. 3) a string name of a matplotlib cmap obj, to list a few:
'bwr': blue-white-red 'RdBu': darkred-white-blue 'RdY1Bu': red-yellow-white-blue 'RdYlGn': red-yellow-white-green 'spectral': purple-yellow-cyanblue 'seismic': darkblue-white-darkred 'jet': rainbow darkblue-darkred 'rainbow': rainbow purple-red
Append '_r' to get the reversed colormap.


#### Abstract

Note: <minlevel> and <maxlevel> are better derived using numpy.min() and numpy.max(). MV.min() and MV.max() may have problems. Iso levels are computed using vcs function (mkscale()), and a matplotlib colormap is created (if not given), and the colormap will be changed so positive/negative splits (if required) is achieved.


Update time: 2015-04-27 14:55:33
class utils.plot.Plot2D(var, method, ax=None, xarray=None, yarray=None, title=None, latlon=True, latlongrid=False, legend='global', legend_ori='horizontal', clean=False)
__init__(var, method, ax=None, xarray=None, yarray=None, title=None, latlon=True, latlongrid=False, legend='global', legend_ori='horizontal', clean=False )
Utility class for 2D plots

## Parameters

- $\operatorname{var}($ NCVAR or ndarray) - variable to plot. At least 2D.
- method - plotting method, could be an instance of Boxfill, Isofill.


## Keyword Arguments

- ax - matplotlib axis obj. If None, create a new.
- xarray (ndarray) - 1d array, the array values for the $x$-axis. If None, use the int indices for the x -dimension.
- yarray (ndarray) - 1d array, the array values for the $y$-axis. If None, use the int indices for the $y$-dimension.
- title (str) - title to plot at subtitle. If None, plot only an alphabetical index.
- latlon (bool) - plot lat/lon axis labels or not.
- latlongrid (bool) - plot lat/lon grid lines or not.
- legend (str) - location of colorbar. Could be: 'global': all subplots share the colorbar of the 1st subplot in figure. or 'local': each subplot in figure uses its own colorbar.
- legend_ori (str) - 'horizontal' or 'vertical', colorbar orientation. clean (bool): if True, omit axis labels, colorbar, subtitle, continents, boundaries etc.. Useful to overlay plots.
classmethod checkBasemap (var, xarray, yarray)
Check variable should be plotted using basemap or not
getGrid()
Get lat/lon grid info from data


## classmethod getSlab (var)

Get a 2D slab from variable

## Parameters

- $\operatorname{var}(N C V A R$ or ndarray) - nd variable. If is transient variable, try
- slice its 1st time point. If numpy.ndarray, try to take a slab (to) -
- its last 2 dimensions. (from)-

Returns result (ndarray) - a 2d slab from input <var> to plot.
class utils.plot.Plot2Cartopy (var, method, ax, legend='global', title=None, xarray=None, yarray =None, latlon=True, latlongrid=False, fill_color='0.8', legend_ori='horizontal', clean=False, fix_aspect=False )
classmethod checkBasemap (var, xarray, yarray)
Check variable should be plotted using basemap or not
getGrid()
Get lat/lon grid info from data

## classmethod getSlab (var)

Get a 2D slab from variable

## Parameters

```
- var (NCVAR or ndarray) - nd variable. If is transient variable, try
- slice its 1st time point. If numpy.ndarray, try to take
a slab(to)-
- its last 2 dimensions. (from)-
```

Returns result (ndarray) - a 2d slab from input <var> to plot.

The code of this package is hosted at https://github.com/ihesp/IPART.
For any queries, please contact xugzhi1987@gmail.com.

## CHAPTER 7

## Contributing and getting help

We welcome contributions from the community. Please create a fork of the project on GitHub and use a pull request to propose your changes. We strongly encourage creating an issue before starting to work on major changes, to discuss these changes first.

For help using the package, please post issues on the project GitHub page.

# CHAPTER 8 

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## Bibliography

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## Python Module Index

```
a
AR_detector,31
AR_tracer,42
t
thr, 29
U
utils.funcs,46
utils.plot,48
```


## Symbols

$\qquad$ init__ () (AR_tracer.AR method), 42
_init__() (utils.plot.Boxfill method), 49
__init__() (utils.plot.Isofill method), 50
__init__() (utils.plot.Plot2D method), 51

## A

anchor_lats (AR_tracer.AR attribute), 43
anchor_lons (AR_tracer.AR attribute), 43
append () (AR_tracer. $A R$ method), 43
applyCropIdx () (in module AR_detector), 31
AR (class in AR_tracer), 42
AR_detector (module), 31
AR_tracer (module), 42
areaFilt() (in module AR_detector), 31

## B

backwardHausdorff() (AR_tracer.AR method), 43 Boxfill (class in utils.plot), 49

## C

cart2Spherical() (in module AR_detector), 31
cart2Wind() (in module AR_detector), 31
checkBasemap () (utils.plot.Plot2Cartopy class method), 52
checkBasemap () (utils.plot.Plot 2 D class method), 52 checkCyclic() (in module AR_detector), 32 computeTheta () (in module $A R$ _detector), 32 coor (AR_tracer.AR attribute), 43
cropMask () (in module AR_detector), 32
crossSectionFlux() (in module AR_detector), 32
cyclicLabel () (in module AR_detector), 33

## D

determineThresLow () (in module AR_detector), 33 dLatitude () (in module utils.funcs), 46 dLongitude () (in module utils.funcs), 47 duration (AR_tracer.AR attribute), 43

## F

filterTracks() (in module AR_tracer), 44 findARAxis () (in module AR_detector), 33
findARs () (in module AR_detector), 34
findARsGen () (in module AR_detector), 36
forwardHausdorff () (AR_tracer.AR method), 43
forwardHausdorff() (in module AR_tracer), 44

## G

get3DEllipse () (in module utils.funcs), 46 getAnchors () (in module AR_tracer), 44 getARAxis() (in module AR_detector), 38 getARData () (in module AR_detector), 38 getAttrDict() (in module thr), 30 getDistMatrix() (in module AR_tracer), 44 getGrid() (utils.plot.Plot2Cartopy method), 52 getGrid() (utils.plot.Plot2D method), 52 getNormalVectors() (in module AR_detector), 39 getSlab() (utils.plot.Plot2Cartopy class method), 52
getSlab() (utils.plot.Plot2D class method), 52
get TimeAxis() (in module utils.funcs), 47

## H

Hausdorff()(AR_tracer.AR method), 42
I
insertCropSlab() (in module AR_detector), 39
Isofill (class in utils.plot), 50

## L

latest (AR_tracer.AR attribute), 43
lats (AR_tracer. $A R$ attribute), 43
lons (AR_tracer.AR attribute), 43

## M

maskToGraph () (in module AR_detector), 39
matchCenters () (in module $A R$ _tracer), 45

## P

```
partPeaks() (in module AR_detector), 40
partPeaksOld() (in module AR_detector), 40
plot2() (in module utils.plot),48
Plot2Cartopy (class in utils.plot),52
Plot2D (class in utils.plot),51
plotAR() (in module utils.plot),49
plotARTrack()(in module utils.plot),49
plotGraph() (in module AR_detector), 40
plotHD () (in module AR_tracer), 45
prepareMeta() (in module AR_detector), 41
```


## R

```
rdp_lats (AR_tracer.AR attribute), 43
rdp_lons (AR_tracer.AR attribute), 44
readCSVRecord() (in module AR_tracer), 45
readVar() (in module utils.funcs), 47
rotatingTHR() (in module thr), 30
```


## S

save2DF () (in module AR_detector), 41
spherical2Cart() (in module AR_detector), 41

## T

thr (module), 29
THR () (in module thr), 29
times (AR_tracer.AR attribute), 44
trackARs () (in module AR_tracer), 46

## U

utils.funcs (module), 46
utils.plot (module), 48
uvDecomp () (in module AR_detector), 41
W
wind2Cart() (in module AR_detector), 42


[^0]:    ${ }^{1}$ Greyscale erosion (also known as minimum filtering) can be understood by analogy with a moving average. Instead of the average within a neighborhood, erosion replaces the central value with the neighborhood minimum. Similarly, dilation replaces with the maximum. And the neighborhood is defined by the structuring element $E$.

