

Jupyter notebook demonstrating the use of additional PmagPy functions

This Jupyter notebook demonstrates a number of PmagPy functions within a notebook environment running a Python 2.7 kernel. The benefits of working within these notebooks include: reproducibility, interactive code development, convenient workspace for projects, version control (when integrated with GitHub or other version control software) and ease of sharing.

The notebook can be viewed as html at the following link where the code and tables are better rendered than in this PDF: http://pmagpy.github.io/Additional_PmagPy_Examples.html

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Note: This notebook makes use of pandas for reading, displaying, and using data with a dataframe structure. More information about the pandas module and its use within PmagPy can be found [here](#) within the documentation of the [PmagPy Cookbook](#).

```
In [1]: # With the PmagPy folder in the PYTHONPATH,
        # the function modules from PmagPy can be imported
import pmagpy.ipmag as ipmag
import pmagpy.pmagplotlib as pmagplotlib
import pmagpy.pmag as pmag

from mpl_toolkits.basemap import Basemap
import numpy as np
import pandas as pd
```

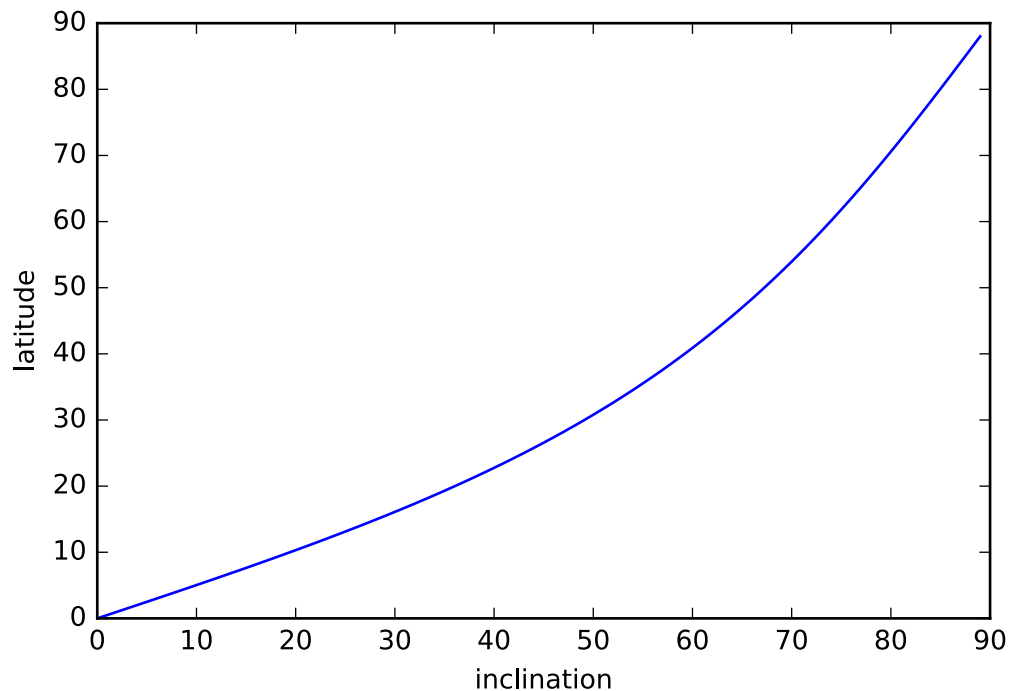
```
import matplotlib.pyplot as plt
import os
%matplotlib inline
%config InlineBackend.figure_formats = {'svg',}
```

2 The dipole equation

The following demonstrates the use of a simple function (**ipmag.lat_from_inc**) which uses the dipole equation to return expected latitude from inclination data as predicted by a pure geocentric axial dipole. The expected inclination for the geomagnetic field can be calculated from a specified latitude using **ipmag.inc_from_lat**.

```
In [2]: inclination = range(0,90,1)
        latitude = []
        for inc in inclination:
            lat = ipmag.lat_from_inc(inc)
            latitude.append(lat)
```

```
In [3]: plt.plot(inclination,latitude)
        plt.ylabel('latitude')
        plt.xlabel('inclination')
        plt.show()
```



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3 Get local geomagnetic field estimate from IGRF

The function **ipmag.igrf** uses the International Geomagnetic Reference Field (IGRF) model to estimate the geomagnetic field direction at a particular location and time. Let's find the direction of the geomagnetic field in Berkeley, California (37.87° N, 122.27° W, elevation of 52 m) on August 27, 2013 (in decimal format, 2013.6544).

```
In [4]: berk_igrf = ipmag.igrf([2013.6544, .052, 37.871667, -122.272778])
        ipmag.igrf_print(berk_igrf)
```

```
Declination: 13.950
Inclination: 61.354
Intensity: 13.950 nT
```

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4 Plotting Directions

We can plot this direction using **matplotlib (plt)** in conjunction with a few **ipmag** functions. To do this, we first initiate a figure (numbered as Fig. 0, with a size of 6x6) with the following syntax:

```
plt.figure(num=0,figsize=(6,6))
```

We then draw an equal area stereonet within the figure, specifying the figure number:

```
ipmag.plot_net(0)
```

Now we can plot the direction we just pulled from IGRF using **ipmag.plot_di()**:

```
ipmag.plot_di(berk_igrf[0],berk_igrf[1])
```

To label or color the plotted points, we would pass the same code as above but with a few extra arguments and one additional line of code:

```
ipmag.plot_di(berk_igrf[0],berk_igrf[1], color='r', label="Berkeley, CA -- August 27, 2013")
plt.legend()
```

We may wish to save the figure we just created. To do so, we would pass the following *save* function, specifying 1) the relative path to the folder where we want the figure to be saved and 2) the name of the file with the desired extension (.pdf in this example):

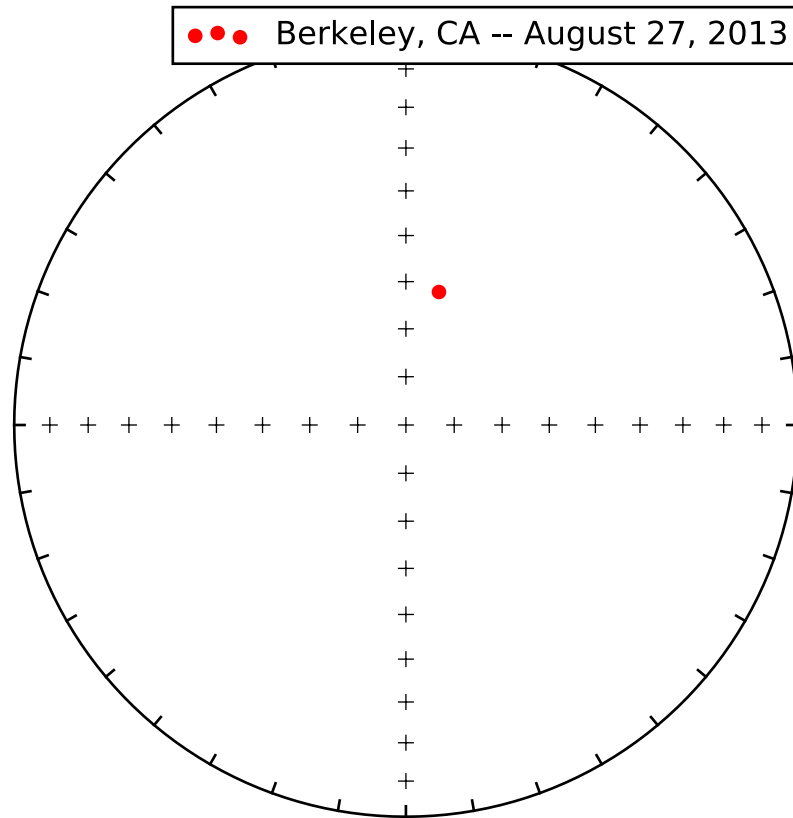
```
plt.savefig("../Additional_Notebook_Output/Berkeley_IGRF.pdf")
```

To ensure the figure is displayed properly and then cleared from the namespace, it is good practice to end such a code block with the following:

```
plt.show()
```

Now let's run the code we just developed.

```
In [5]: plt.figure(num=0,figsize=(5,5))
        ipmag.plot_net(0)
        ipmag.plot_di(berk_igrf[0],berk_igrf[1], color='r', label="Berkeley, CA -- August 27, 2013")
        plt.legend()
        plt.savefig("./Additional_Notebook_Output/Berkeley_IGRF.pdf")
        plt.show()
```



Let's see how this magnetic direction compares to the Geocentric Axial Dipole (GAD) model of the geomagnetic field. We can estimate the expected GAD inclination by passing Berkeley's latitude to the function `ipmag.inc_from_lat`.

We also demonstrate below how to manipulate the placement of the figure legend to ensure no data points are obscured. `plt.legend` uses the "best" location by default, but this can be changed with the following:

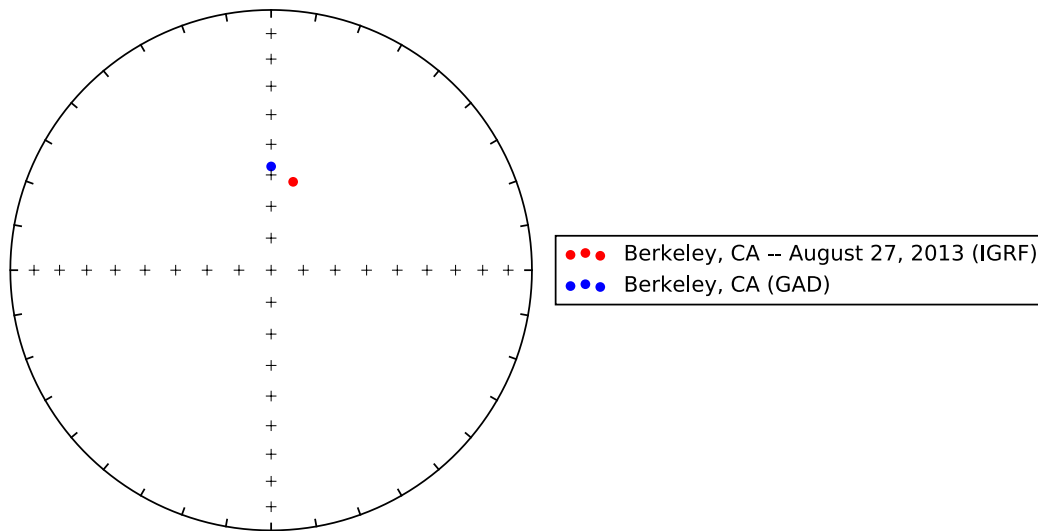
```
plt.legend(loc="upper right")
```

or

```
plt.legend(loc="lower center")
```

See the **plt.legend** documentation for the complete list of placement options. Alternatively, you can give (x,y) coordinates to the **loc=** keyword argument (with the origin (0,0) at the lower left of the figure). To manipulate placement even more precisely, use the keyword **bbox_to_anchor** in conjunction with **loc**. If this is done, **loc** becomes the anchor point on the legend, and **bbox_to_anchor** places this anchor point at the specified coordinates. The latter method is demonstrated below. Play around with the **plt.legend** arguments to see how this changes things.

```
In [6]: GAD_inc = ipmag.inc_from_lat(37.87)
plt.figure(num=0,figsize=(5,5))
ipmag.plot_net(0)
ipmag.plot_di(berk_igrf[0],berk_igrf[1], color='r', label="Berkeley, CA -- August 27, 2013 (IGRF)"
ipmag.plot_di(0,GAD_inc, color='b', label="Berkeley, CA (GAD)")
plt.legend(loc='center left', bbox_to_anchor=(1.0, 0.5))
plt.show()
```



Below, we calculate the angular difference between these two directions.

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5 Calculate the Angle Between Directions

While **ipmag** functions have been optimized to perform tasks within an interactive computing environment such as the Jupyter notebook, the **pmag** functions which are used extensively within **ipmag** can also be directly called. Here is a demonstration of the function **pmag.angle**, which calculates the angle between two directions and outputs a **numpy** array. Continuing our comparison from the last section, let's calculate the angle between the IGRF and GAD-estimated magnetic directions calculated and plotted above.

```
In [7]: direction1 = [berk_igrf[0],berk_igrf[1]]
        direction2 = [0,GAD_inc]
        print pmag.angle(direction1,direction2)[0]
```

8.18973048085

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6 Generate and plot Fisher distributed unit vectors from a specified distribution

Let's use the function `ipmag.fishrot` to generate a set of 50 Fisher-distributed directions at a declination of 200° and inclination of 45° . These directions will serve as an example paleomagnetic dataset that will be used for the next several examples. The output from `ipmag.fishrot` is a nested list of lists of vectors (declination, inclination, intensity). Generally these vectors are unit vectors with an intensity of 1.0. We refer to this data structure as a `di_block`. In the code below the first two vectors are shown.

```
In [8]: fisher_directions = ipmag.fishrot(k=40, n=50, dec=200, inc=50)
        fisher_directions[0:2]
```

```
Out[8]: [[183.99588954660211, 45.029054467016621, 1.0],
         [200.58837258558225, 41.811402826036407, 1.0]]
```

This `di_block` can be unpacked in separate lists of declination and inclination using the `ipmag.unpack_di_block` function.

```
In [9]: fisher_decs, fisher_incs = ipmag.unpack_di_block(fisher_directions)
        print fisher_decs[0]
        print fisher_incs[0]
```

183.995889547
45.029054467

Another way to deal with the `di_block` is to make it into a pandas dataframe which allows for the direction to be nicely displayed and analyzed. In the code below, a dataframe is made from the *fisher_directions* `di_block` and then the first 5 rows are displayed with `.head()`.

```
In [10]: directions = pd.DataFrame(fisher_directions,columns=['dec','inc','length'])
        directions.head()
```

```
Out[10]:
```

	dec	inc	length
0	183.995890	45.029054	1
1	200.588373	41.811403	1
2	199.546318	36.865786	1
3	182.021565	57.114399	1
4	221.438217	38.160275	1

Now let's calculate the Fisher and Bingham means of these data.

```
In [11]: fisher_mean = ipmag.fisher_mean(directions.dec,directions.inc)
        bingham_mean = ipmag.bingham_mean(directions.dec,directions.inc)
```

Here's the raw output of the Fisher mean which is a dictionary containing the mean direction and associated statistics:

```
In [12]: fisher_mean
```

```
Out[12]: {'alpha95': 3.159239986865467,
          'csd': 12.553483567656313,
          'dec': 198.86339034639676,
          'inc': 49.196847025953076,
          'k': 41.633365663104868,
          'n': 50,
          'r': 48.823059360693883}
```

The function **ipmag.print_direction_mean** prints formatted output from this Fisher mean dictionary:

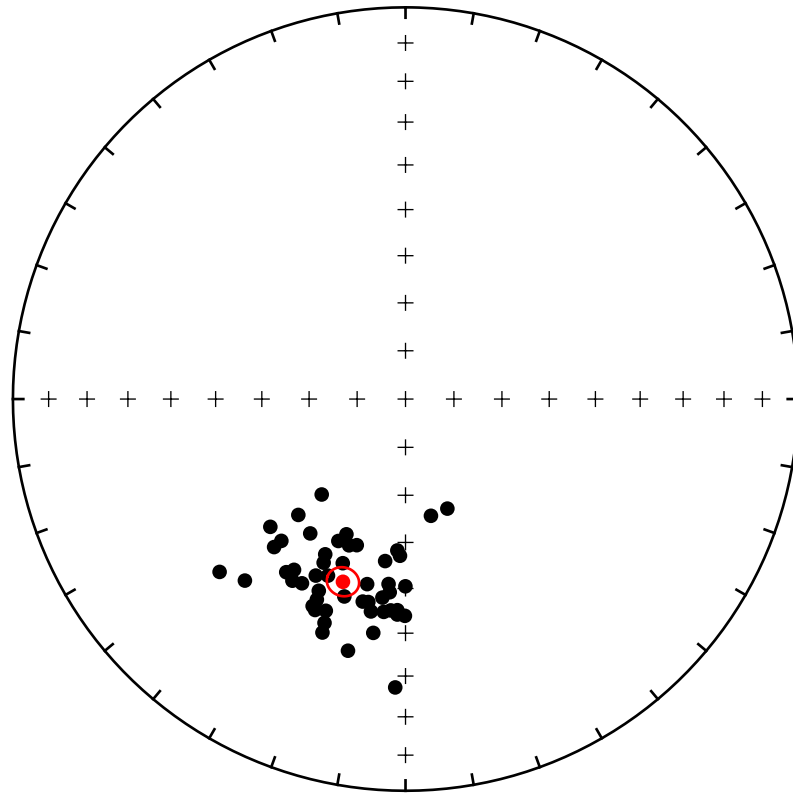
```
In [13]: ipmag.print_direction_mean(fisher_mean)
```

```
Dec: 198.9  Inc: 49.2
Number of directions in mean (n): 50
Angular radius of 95% confidence (a_95): 3.2
Precision parameter (k) estimate: 41.6
```

Now we can plot all of our data using the function **ipmag.plot_di**. We can also plot the Fisher mean with its angular radius of 95% confidence (α_{95}) using **ipmag.plot_di_mean**.

```
In [14]: declinations = directions.dec.tolist()
        inclinations = directions.inc.tolist()

        plt.figure(num=1,figsize=(5,5))
        ipmag.plot_net(1)
        ipmag.plot_di(declinations,inclinations)
        ipmag.plot_di_mean(fisher_mean['dec'],fisher_mean['inc'],fisher_mean['alpha95'],color='r')
```



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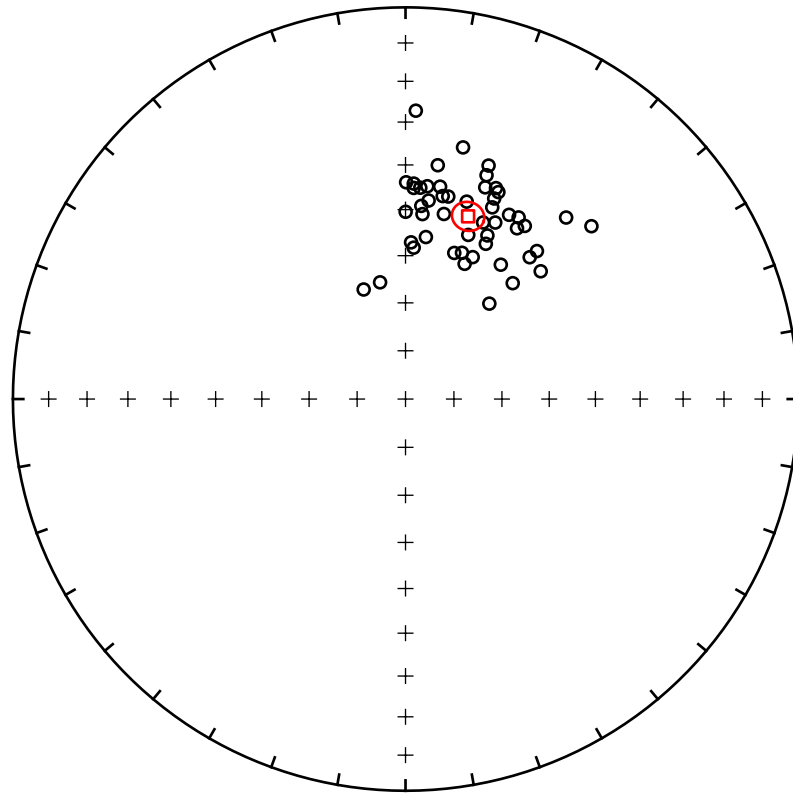
7 Flip polarity of directional data

Let's flip all the directions (find their antipodes) of the Fisher-distributed population using the function `ipmag.do_flip()` function and plot the resulting directions.

```
In [15]: # get reversed directions
         dec_reversed, inc_reversed = ipmag.do_flip(declinations, inclinations)

         # take the Fisher mean of these reversed directions
         rev_mean = ipmag.fisher_mean(dec_reversed, inc_reversed)

         # plot the flipped directions
         plt.figure(num=1, figsize=(5,5))
         ipmag.plot_net(1)
         ipmag.plot_di(dec_reversed, inc_reversed)
         ipmag.plot_di_mean(rev_mean['dec'], rev_mean['inc'], rev_mean['alpha95'], color='r', marker='s')
```

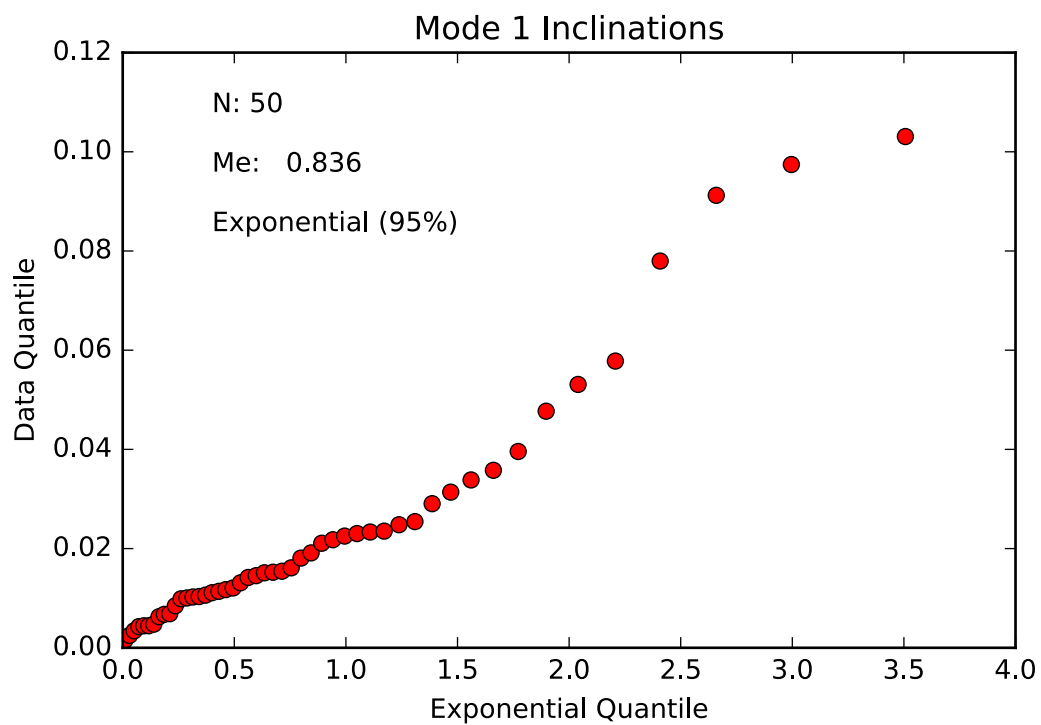
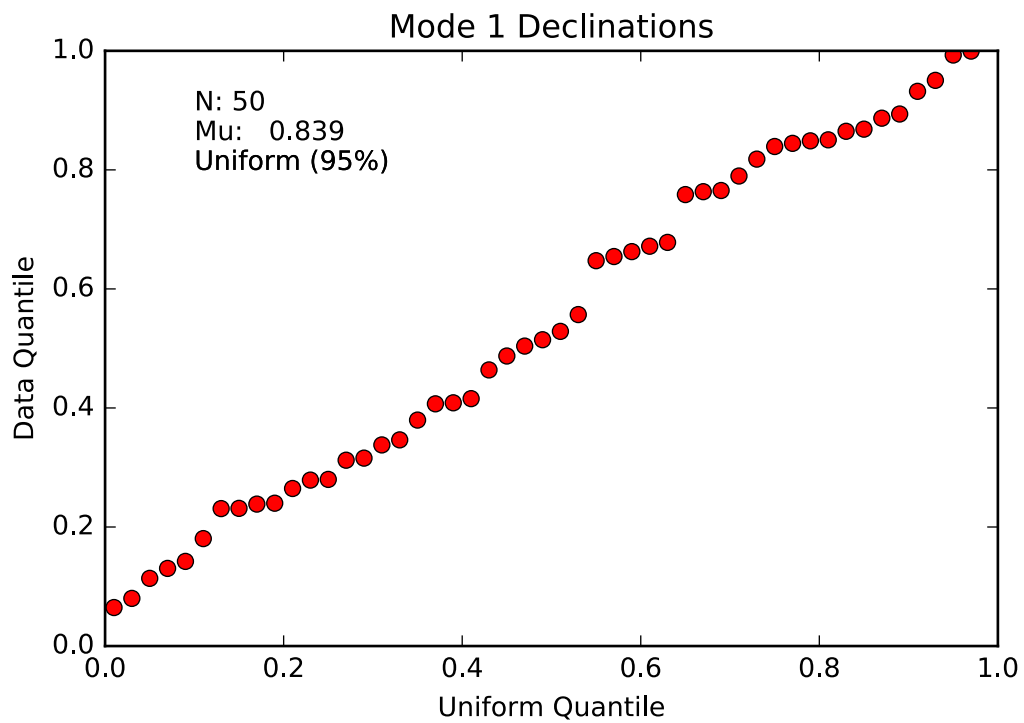
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8 Test directional data for Fisher distribution

The function `ipmag.fishqq` tests whether directional data are Fisher-distributed. Let's use this test on the random Fisher-distributed directions we just created (it should pass!).

```
In [16]: ipmag.fishqq(declinations, inclinations)
```

```
Out[16]: {'Dec': 198.86305338122148,
          'Inc': 49.174480281318843,
          'Me': 0.83647984581601553,
          'Me_critical': 1.094,
          'Mode': 'Mode 1',
          'Mu': 0.83924082417308898,
          'Mu_critical': 1.207,
          'N': 50,
          'Test_result': 'consistent with Fisherian model'}
```



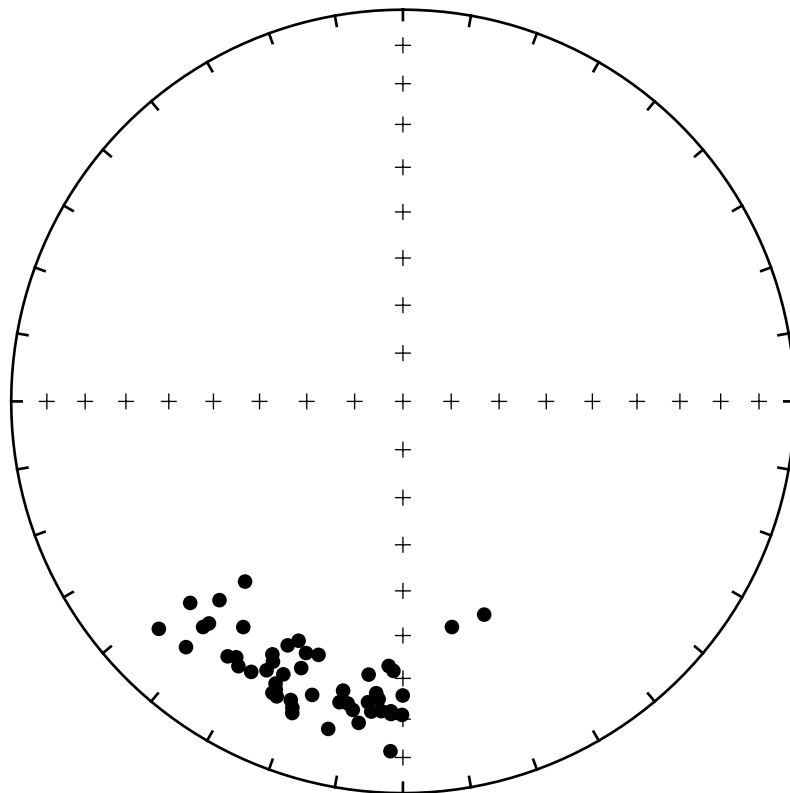
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9 Squish directional data

Inclination flattening can occur for magnetizations in sedimentary rocks. We can simulate inclination error of a specified “flattening factor” with the function **ipmag.squish**. Flattening factors range from 0 (completely flattened) to 1 (no flattening). Let’s squish our directions with a 0.4 flattening factor.

```
In [17]: # squish all inclinations
squished_incs = []
for inclination in inclinations:
    squished_incs.append(ipmag.squish(inclination, 0.4))

# plot the squished directional data
plt.figure(num=1,figsize=(5,5))
ipmag.plot_net(1)
ipmag.plot_di(declinations,squished_incs)
squished_DIs = np.array(zip(declinations,squished_incs))
```



```
In [18]: ipmag.fisher_mean(di_block=squished_DIs)
```

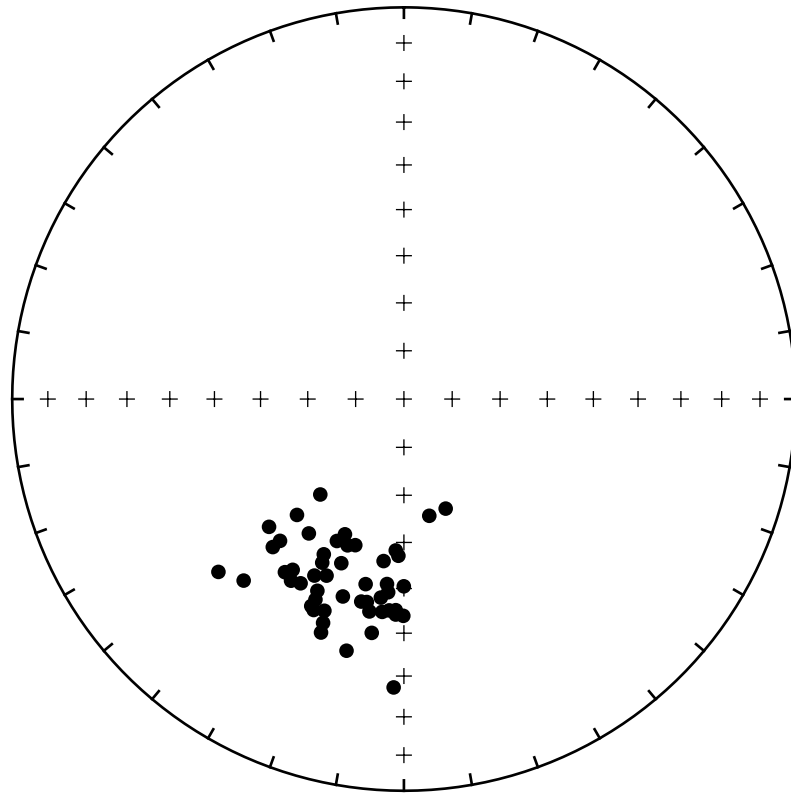
```
Out[18]: {'alpha95': 3.7940228548357475,  
          'csd': 14.997170953500827,  
          'dec': 198.72106646496212,  
          'inc': 25.567588455989114,  
          'k': 29.171002445333873,  
          'n': 50,  
          'r': 48.320249703731456}
```

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10 Unsquish directional data

We can also “unsquish” data by a specified flattening factor. Let’s unsquish the data we squished above with the function `ipmag.unsquish`. Using a flattening factor of 0.4 will restore the data to its original state.

```
In [19]: unsquished_incs = []  
         for squished_inc in squished_incs:  
             unsquished_incs.append(ipmag.unsquish(squished_inc, 0.4))  
  
         # plot the squished directional data  
         plt.figure(num=1,figsize=(5,5))  
         ipmag.plot_net(1)  
         ipmag.plot_di(declinations,unsquished_incs)
```



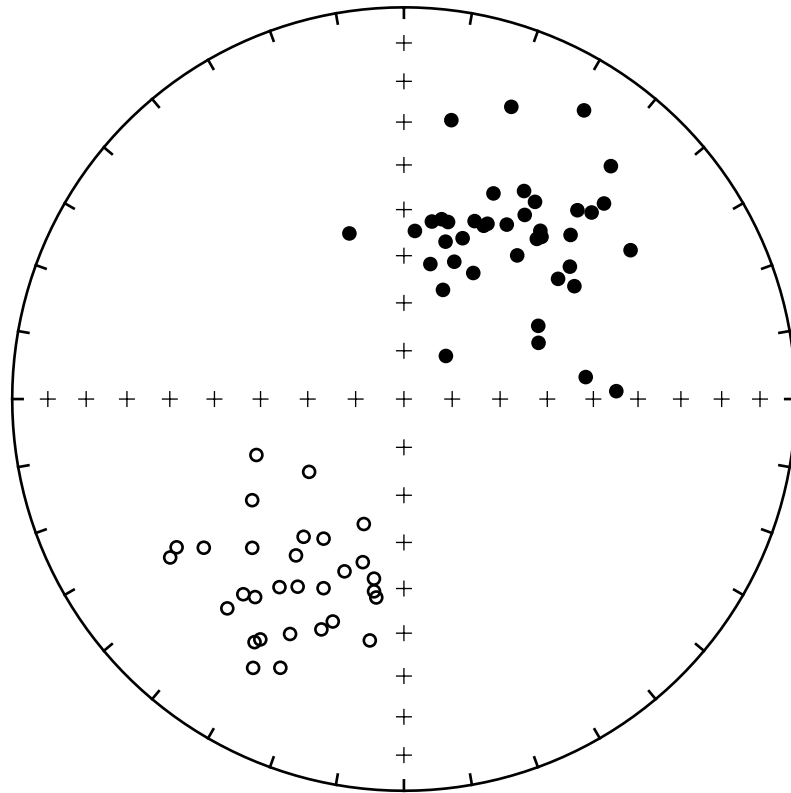
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11 Bootstrap Reversal Test

Here we carry out two types of reversal tests with **ipmag** to test if two populations are antipodal to one another: the bootstrap reversal test (Tauxe, 2010; **ipmag.reversal_test_bootstrap**) and the McFadden and McElhinny (1990) reversal test, which is an adaptation of the Watson V test for a common mean (**ipmag.reversal_test_MM1990**). The code below uses **ipmag.fishrot** to simulate normal directions and reversed directions from antipodal Fisher distributions.

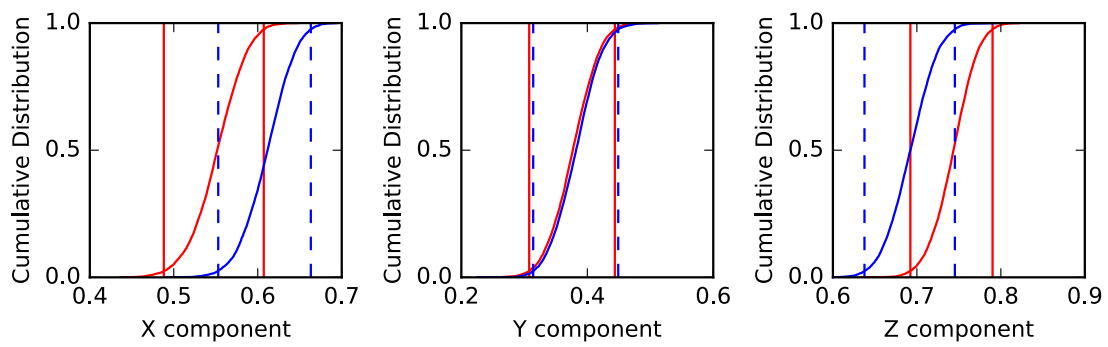
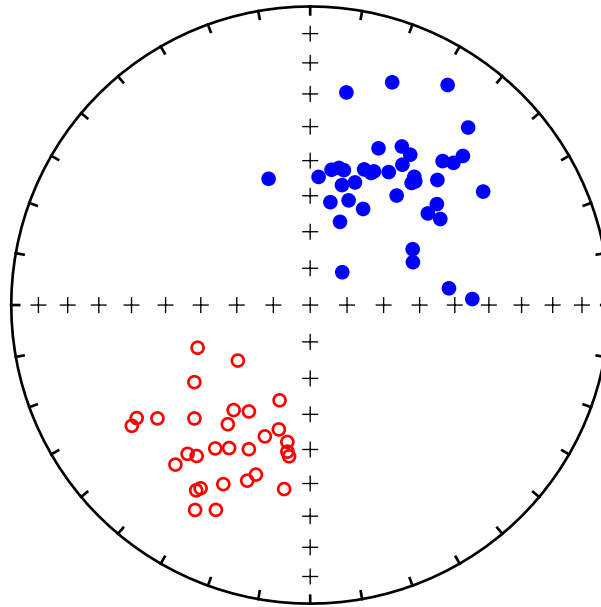
```
In [20]: normal_directions = ipmag.fishrot(k=20,n=40,dec=30,inc=45)
         reversed_directions = ipmag.fishrot(k=20,n=30,dec=210,inc=-45)
         combined_directions = normal_directions + reversed_directions

         plt.figure(num=1,figsize=(5,5))
         ipmag.plot_net(1)
         ipmag.plot_di(di_block=combined_directions)
```



```
In [21]: ipmag.reversal_test_bootstrap(di_block=combined_directions,
    plot_stereo=True, save=True,
    save_folder='./Additional_Notebook_Output/')
```

Here are the results of the bootstrap test for a common mean:



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McFadden and McElhinny (1990) Reversal Test

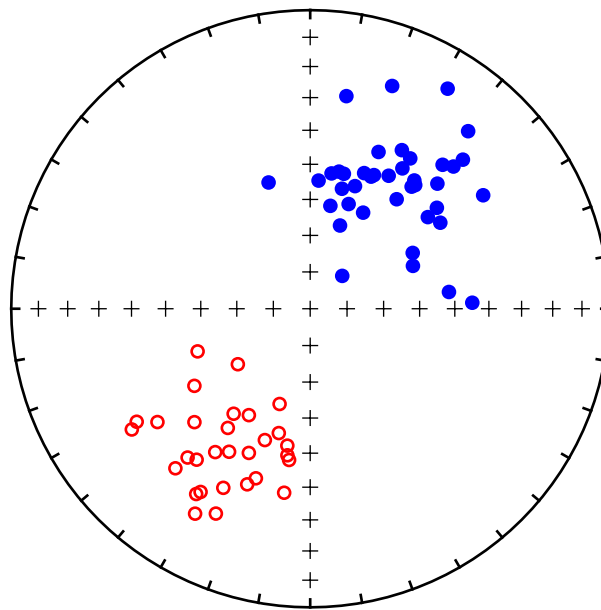
```
In [22]: ipmag.reversal_test_MM1990(di_block=combined_directions,
    plot_CDF=True, plot_stereo=True,
    save=True, save_folder= './Additional_Notebook_Output/')
```

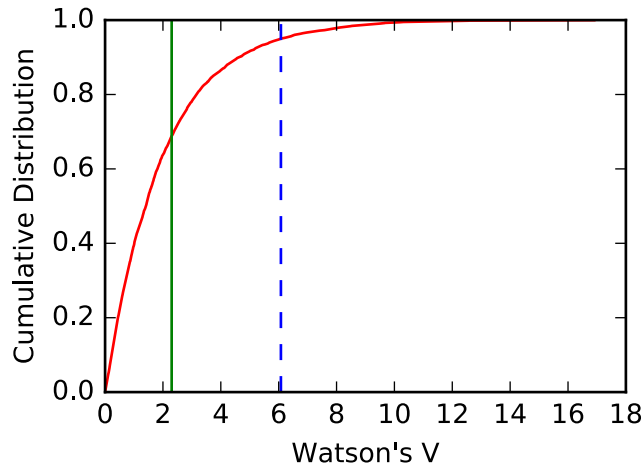
Results of Watson V test:

Watson's V: 2.3
Critical value of V: 6.1
"Pass": Since V is less than Vcrit, the null hypothesis
that the two populations are drawn from distributions
that share a common mean direction can not be rejected.

M&M1990 classification:

Angle between data set means: 4.6
Critical angle for M&M1990: 7.4
The McFadden and McElhinny (1990) classification for
this test is: 'B'





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12 Working with Poles

A variety of plotting functions within PmagPy, together with the Basemap package of matplotlib, provide a great way to work with paleomagnetic poles, virtual geomagnetic poles, and polar wander paths.

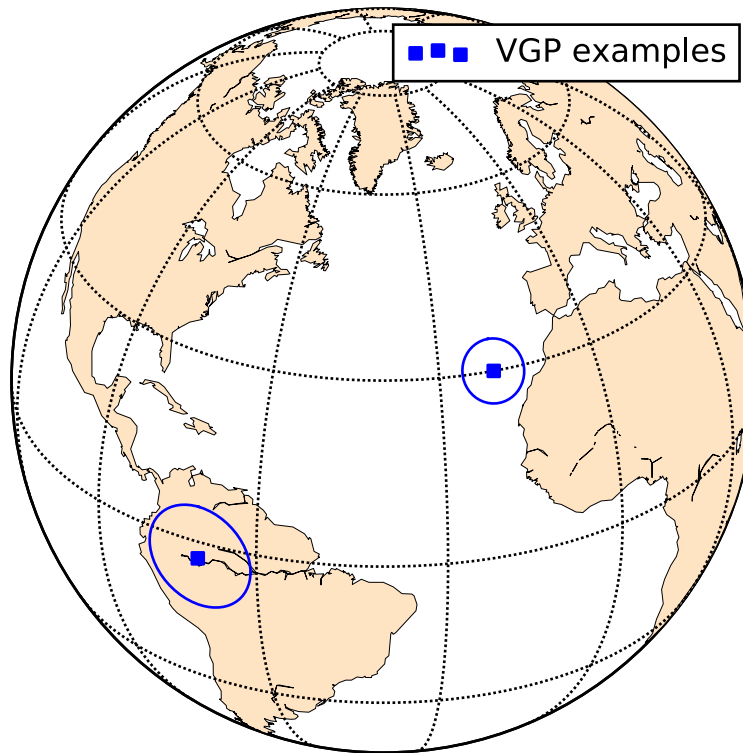
```
In [23]: # initiate figure and specify figure size
plt.figure(figsize=(5, 5))

# initiate a Basemap projection, specifying the latitude and
# longitude (lat_0 and lon_0) at which our figure is centered.
pmap = Basemap(projection='ortho',lat_0=30,lon_0=320,
               resolution='c',area_thresh=50000)
# other optional modifications to the globe figure
pmap.drawcoastlines(linewidth=0.25)
pmap.fillcontinents(color='bisque',lake_color='white',zorder=1)
pmap.drawmapboundary(fill_color='white')
pmap.drawmeridians(np.arange(0,360,30))
pmap.drawparallels(np.arange(-90,90,30))

# Here we plot a pole at 340 E longitude, 30 N latitude with an
# alpha 95 error angle of 5 degrees. Keyword arguments allow us
# to specify the label, shape, and color of this data.
ipmag.plot_pole(pmap,340,30,5,label='VGP examples',
               marker='s',color='Blue')

# We can plot multiple poles sequentially on the same globe using
# the same plot_pole function.
ipmag.plot_pole(pmap,290,-3,9,marker='s',color='Blue')
```

```
plt.legend()
# Optional save (uncomment to save the figure)
# plt.savefig('Code_output/VGP_example.pdf')
plt.show()
```



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13 Calculate and Plot VGPs

Using the function `ipmag.vgp_calc`, we can calculate virtual geomagnetic poles (VGPs) of our fFisher-distributed directions. We'll need to first assign a location to these magnetic directions - let's assume they are from Berkeley, CA (37.87° N, 122.27° W).

```
In [24]: # plug in site latitude and longitude to the "directions" dataframe
directions['site_lat'] = 37.97
directions['site_lon'] = -122.27

# calculate VGPs (this automatically adds VGP data to the dataframe)
ipmag.vgp_calc(directions, dec_tc = 'dec', inc_tc = 'inc')
directions.head()
```

```
Out[24]:
```

	dec	inc	length	site_lat	site_lon	paleolatitude \
0	183.995890	45.029054	1	37.97	-122.27	26.588302

1	200.588373	41.811403	1	37.97	-122.27	24.095659
2	199.546318	36.865786	1	37.97	-122.27	20.553229
3	182.021565	57.114399	1	37.97	-122.27	37.715082
4	221.438217	38.160275	1	37.97	-122.27	21.449872

	vgp_lat	vgp_lon	vgp_lat_rev	vgp_lon_rev
0	-25.333013	233.776578	25.333013	53.776578
1	-24.992308	216.987184	24.992308	36.987184
2	-28.660199	216.813085	28.660199	36.813085
3	-14.291968	236.079855	14.291968	56.079855
4	-18.969679	197.086706	18.969679	17.086706

We have already calculated the Fisher mean of this data, so let's translate it to a VGP too. For a one-line dataset, we plug the Fisher mean data into a **pandas** *Series* instead of a *DataFrame* (a *DataFrame* can be considered a sequence of concatenated *Series*).

```
In [25]: mean_pole = pd.Series(fisher_mean)
mean_pole['site_lat'] = 37.97
mean_pole['site_lon'] = -122.27
ipmag.vgp_calc(mean_pole, dec_tc = 'dec', inc_tc = 'inc')
mean_pole
```

```
Out[25]: alpha95          3.15924
csd                    12.5535
dec                   198.863
inc                   49.1968
k                     41.6334
n                      50
r                   48.8231
site_lat              37.97
site_lon             -122.27
paleolatitude         30.079
vgp_lat              -19.7048
vgp_lon             220.44194226
vgp_lat_rev          19.7048
vgp_lon_rev          40.4419
dtype: object
```

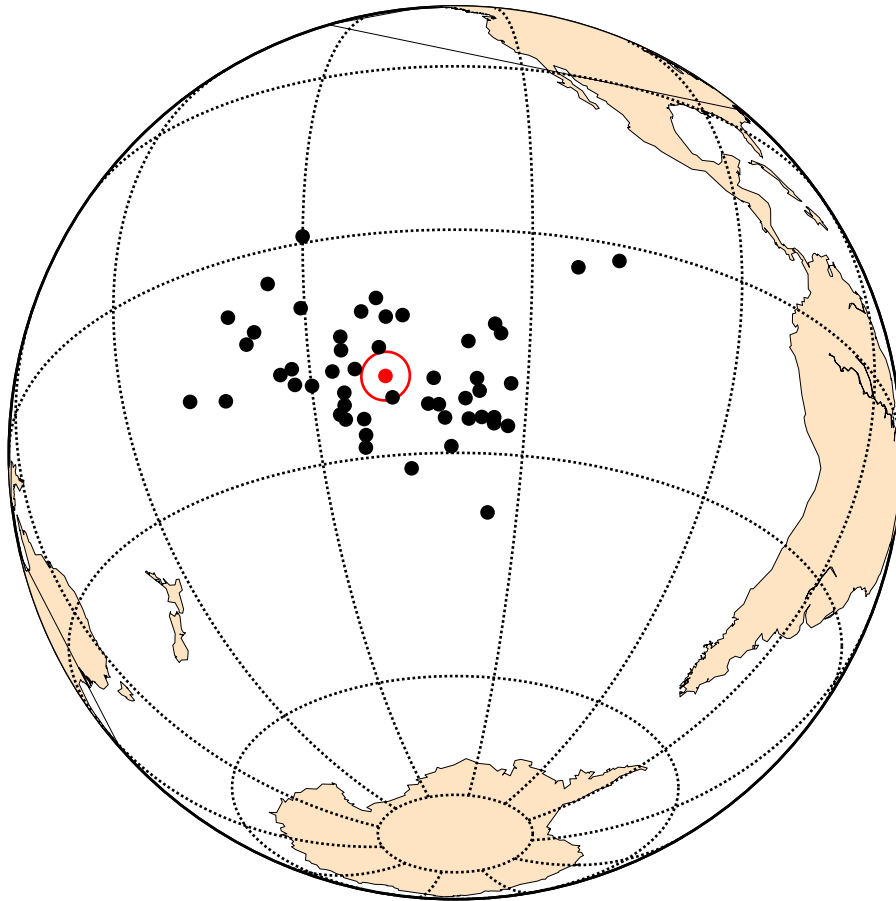
```
In [26]: plt.figure(figsize=(6, 6))
pmap = Basemap(projection='ortho',lat_0=-30,lon_0=-130,
               resolution='c',area_thresh=50000)
pmap.drawcoastlines(linewidth=0.25)
pmap.fillcontinents(color='bisque',lake_color='white',zorder=1)
pmap.drawmapboundary(fill_color='white')
pmap.drawmeridians(np.arange(0,360,30))
pmap.drawparallels(np.arange(-90,90,30))

# use the print_pole_mean function to print the mean data above the globe
ipmag.print_pole_mean(mean_pole)
for n in range(len(directions)):
    ipmag.plot_vgp(pmap, directions['vgp_lon'][n],
                  directions['vgp_lat'][n])
ipmag.plot_pole(pmap, mean_pole['vgp_lon'], mean_pole['vgp_lat'],
               mean_pole['alpha95'], color='r')
```

```

Plong: 198.9  Plat: 49.2
Number of directions in mean (n): 50.0
Angular radius of 95% confidence (A_95): 3.2
Precision parameter (k) estimate: 41.6

```



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14 Plotting APWPs

The capability to plot multiple poles in sequence provides a good way to visualize polar wander paths. Here we use the Phanerozoic APWP of Laurentia (*Torsvik, 2012*) to demonstrate the `plot_pole_colorbar` function.

We first upload the Torsvik (2012) data using the pandas function `read_csv`.

```

In [27]: Laurentia_Pole_Compilation = pd.read_csv('./Additional_Data/Torsvik2012/Laurentia_Pole_Compilation.csv')
Laurentia_Pole_Compilation.head()

```

```

Out[27]:   Q  A95  Com      Formation  Lat   Lon  CLat  CLon  RLat  \
0  5  3.9  NaN      Dunkard Formation -44.1  301.5 -41.5  300.4 -38.0

```

1	5	2.1	NaN	Laborcita	Formation	-42.1	312.1	-43.0	313.4	-32.7
2	5	3.4	#	Wescogame	Formation	-44.1	303.9	-46.3	306.8	-38.2
3	6	3.1	I	Glenshaw	Formation	-28.6	299.9	-28.6	299.9	-28.6
4	5	1.8	NaN	Lower Casper	Formation	-45.7	308.6	-50.5	314.6	-37.6

	RLon		EULER	Age	GPDB	RefNo/Reference
0	43.0	(63.2/_	13.9/79.9)	300		302, T
1	52.9	(63.2/_	13.9/79.9)	301		1311, T
2	51.4	(63.2/_	13.9/79.9)	301		1311, T
3	32.4	(63.2/_	13.9/79.9)	303		Kodama (2009)
4	59.8	(63.2/_	13.9/79.9)	303		1455, T

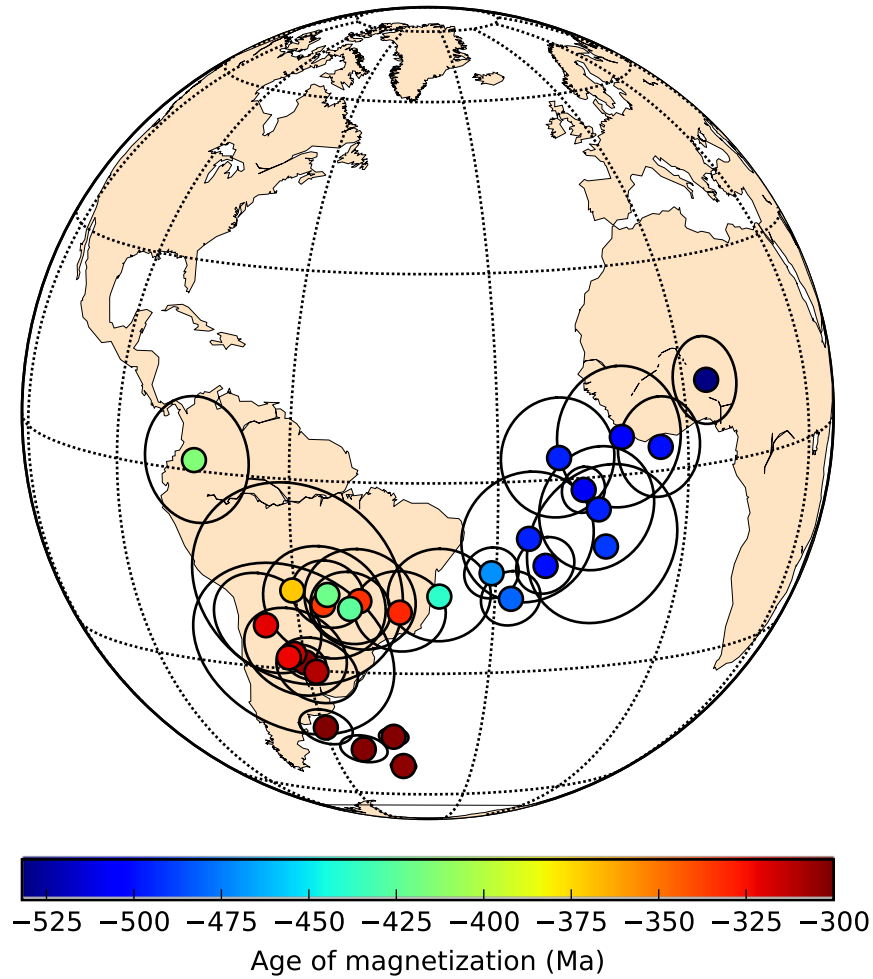
```
In [28]: # initiate the figure as in the plot_pole example
plt.figure(figsize=(6, 6))
pmap = Basemap(projection='ortho',lat_0=10,lon_0=320,
                resolution='c',area_thresh=50000)
pmap.drawcoastlines(linewidth=0.25)
pmap.fillcontinents(color='bisque',lake_color='white',zorder=1)
pmap.drawmapboundary(fill_color='white')
pmap.drawmeridians(np.arange(0,360,30))
pmap.drawparallels(np.arange(-90,90,30))

# Loop through the uploaded data and use the plot_pole_colorbar function
# (instead of plot_pole) to plot the individual poles. The input of this
# function is very similar to that of plot_pole but has the additional
# arguments of (1)AGE, (2)MINIMUM AND (3)MAXIMUM AGES OF PLOTTED POLES.
# Note that the ages are treated as negative numbers -- this just determines
# the direction of the colorbar.
for n in xrange(0, len(Laurentia_Pole_Compilation)):
    m = ipmag.plot_pole_colorbar(pmap, Laurentia_Pole_Compilation['CLon'][n],
                                Laurentia_Pole_Compilation['CLat'][n],
                                Laurentia_Pole_Compilation['A95'][n],
                                -Laurentia_Pole_Compilation['Age'][n],
                                -532,
                                -300,
                                markersize=80, color="k", alpha=1)

pmap.colorbar(m,location='bottom',pad="5%",label='Age of magnetization (Ma)')

# Optional save (uncomment to save the figure)
#plt.savefig('Additional_Notebook_Output/plot_pole_colorbar_example.pdf')

plt.show()
```



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15 Working with anisotropy data

The following code demonstrates reading magnetic anisotropy data into a pandas DataFrame.

```
In [29]: aniso_data = pd.read_csv('./Additional_Data/ani_depthplot/rmag_anisotropy.txt',
                                   delimiter='\\t', skiprows=1)
        aniso_data.head()
```

```
Out[29]:
```

	anisotropy_n	anisotropy_s1	anisotropy_s2	anisotropy_s3	anisotropy_s4	\
0	192	0.332294	0.332862	0.334844	-0.000048	
1	192	0.333086	0.332999	0.333916	-0.000262	
2	192	0.333750	0.332208	0.334041	-0.000699	
3	192	0.330565	0.333928	0.335507	0.000603	
4	192	0.332747	0.332939	0.334314	-0.001516	

	anisotropy_s5	anisotropy_s6	anisotropy_sigma	anisotropy_tilt_correction \
0	0.000027	-0.000263	0.000122	-1
1	-0.000322	0.000440	0.000259	-1
2	0.000663	0.002888	0.000093	-1
3	0.000212	-0.000932	0.000198	-1
4	-0.000311	-0.000099	0.000162	-1

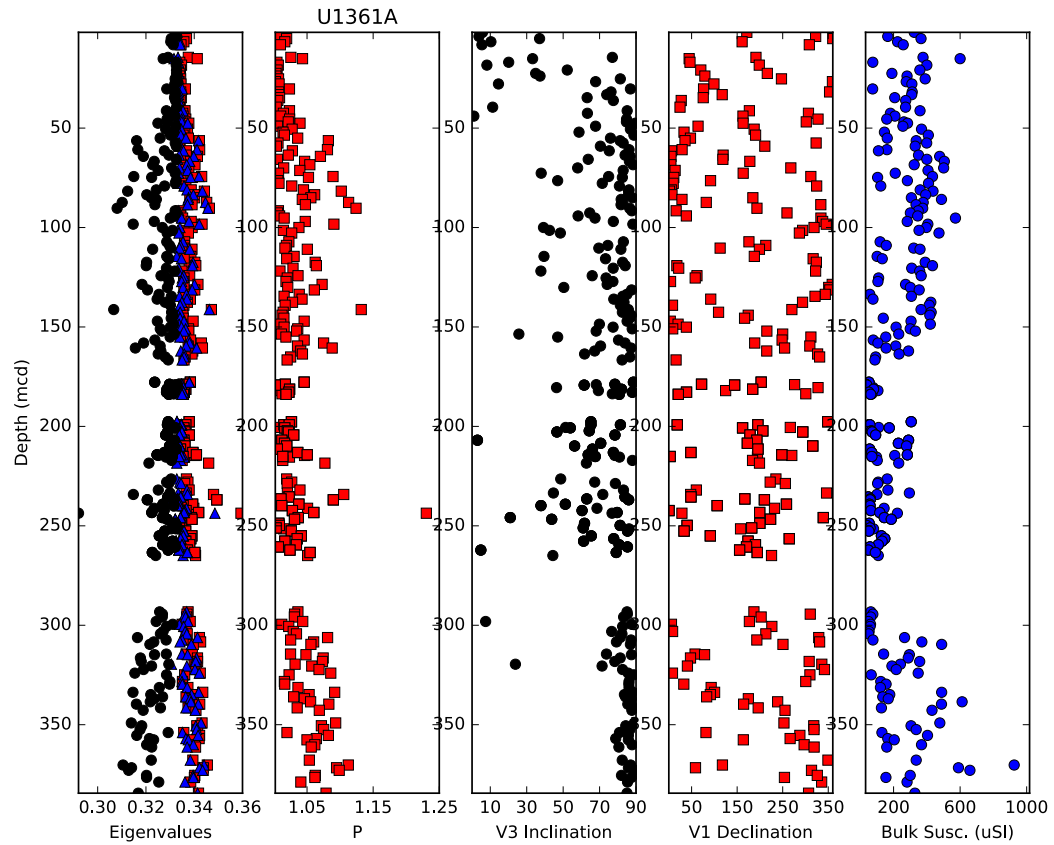
	anisotropy_type	anisotropy_unit	er_analyst_mail_names \
0	AMS	Normalized by trace	NaN
1	AMS	Normalized by trace	NaN
2	AMS	Normalized by trace	NaN
3	AMS	Normalized by trace	NaN
4	AMS	Normalized by trace	NaN

	er_citation_names	er_location_name	er_sample_name \
0	This study	U1361A	318-U1361A-001H-2-W-35
1	This study	U1361A	318-U1361A-001H-3-W-90
2	This study	U1361A	318-U1361A-001H-4-W-50
3	This study	U1361A	318-U1361A-001H-5-W-59
4	This study	U1361A	318-U1361A-001H-6-W-60

	er_site_name	er_specimen_name	magic_method_codes
0	318-U1361A-001H-2-W-35	318-U1361A-001H-2-W-35	LP-X:AE-H:LP-AN-MS:SO-V
1	318-U1361A-001H-3-W-90	318-U1361A-001H-3-W-90	LP-X:AE-H:LP-AN-MS:SO-V
2	318-U1361A-001H-4-W-50	318-U1361A-001H-4-W-50	LP-X:AE-H:LP-AN-MS:SO-V
3	318-U1361A-001H-5-W-59	318-U1361A-001H-5-W-59	LP-X:AE-H:LP-AN-MS:SO-V
4	318-U1361A-001H-6-W-60	318-U1361A-001H-6-W-60	LP-X:AE-H:LP-AN-MS:SO-V

The function `ipmag.aniso_depthplot` is one example of how PmagPy works with such data to generate plots.

```
In [30]: ipmag.aniso_depthplot(dir_path='./Additional_Data/ani_depthplot/');
```



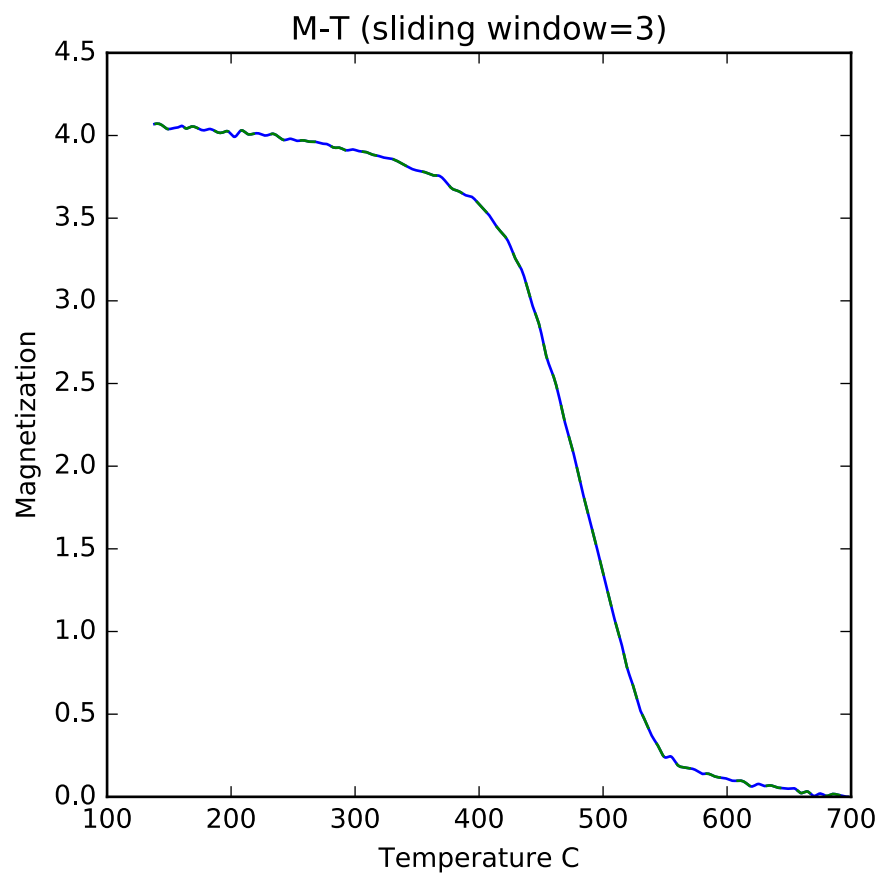
pmagpy-3.4.0

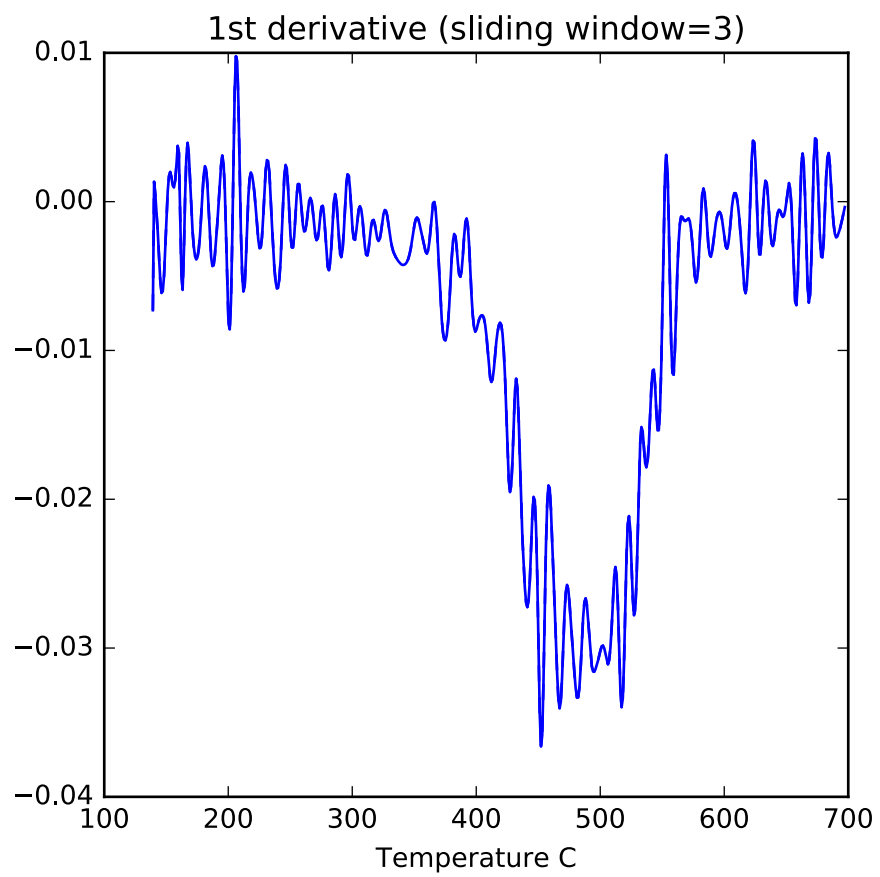
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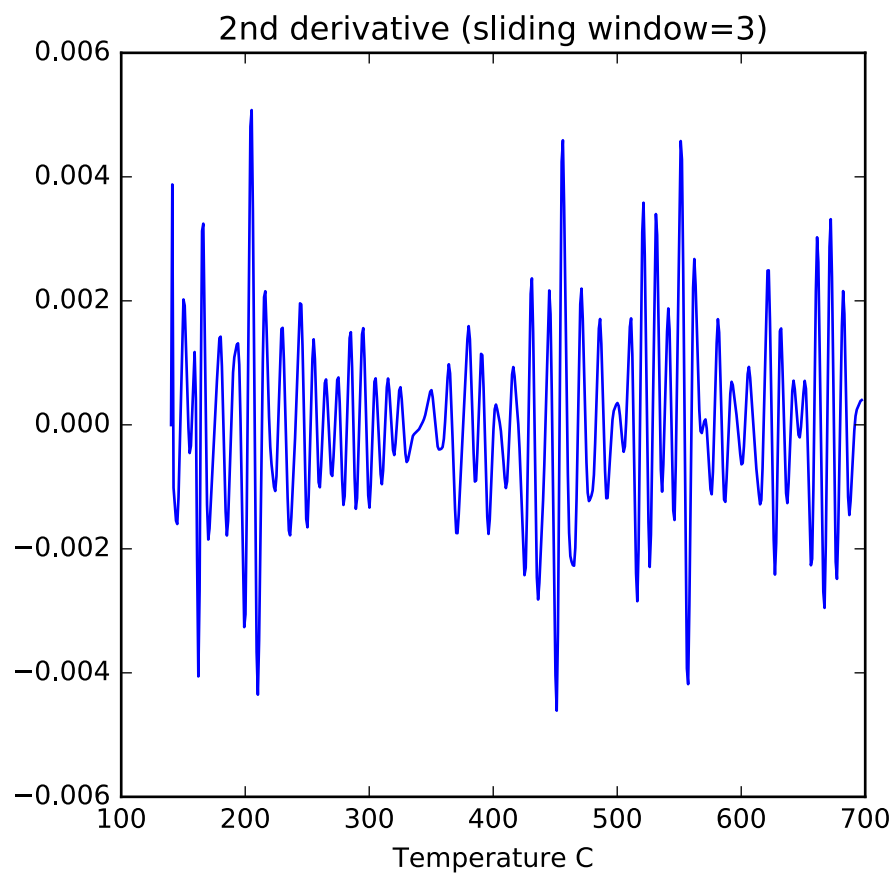
16 Curie temperature data

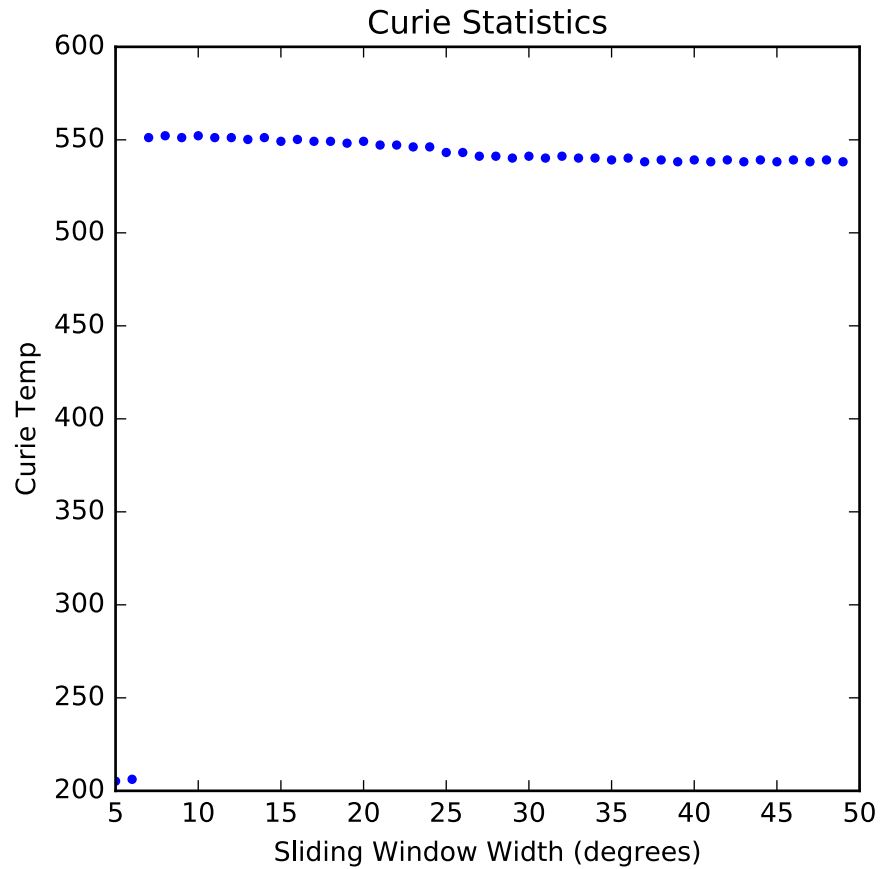
```
In [31]: ipmag.curie(path_to_file='./Additional_Data/curie/',
                    file_name='curie_example.dat', save=True,
                    save_folder='Additional_Notebook_Output/curie/')
```

second deriative maximum is at T=205







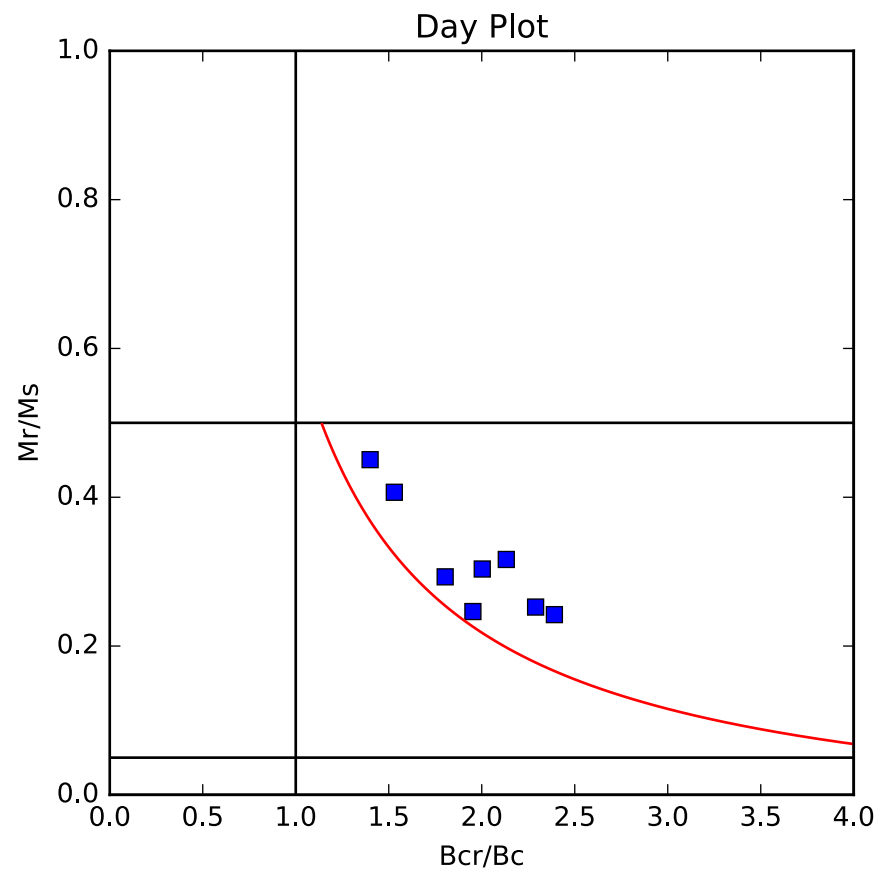


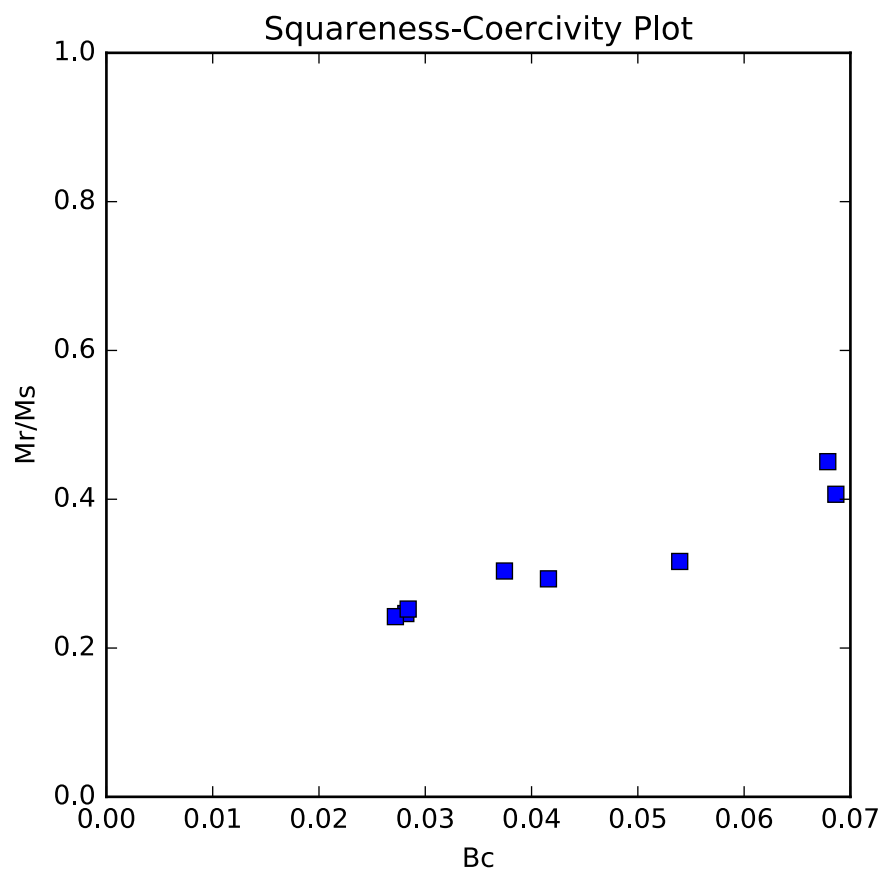
[Go to Top](#)

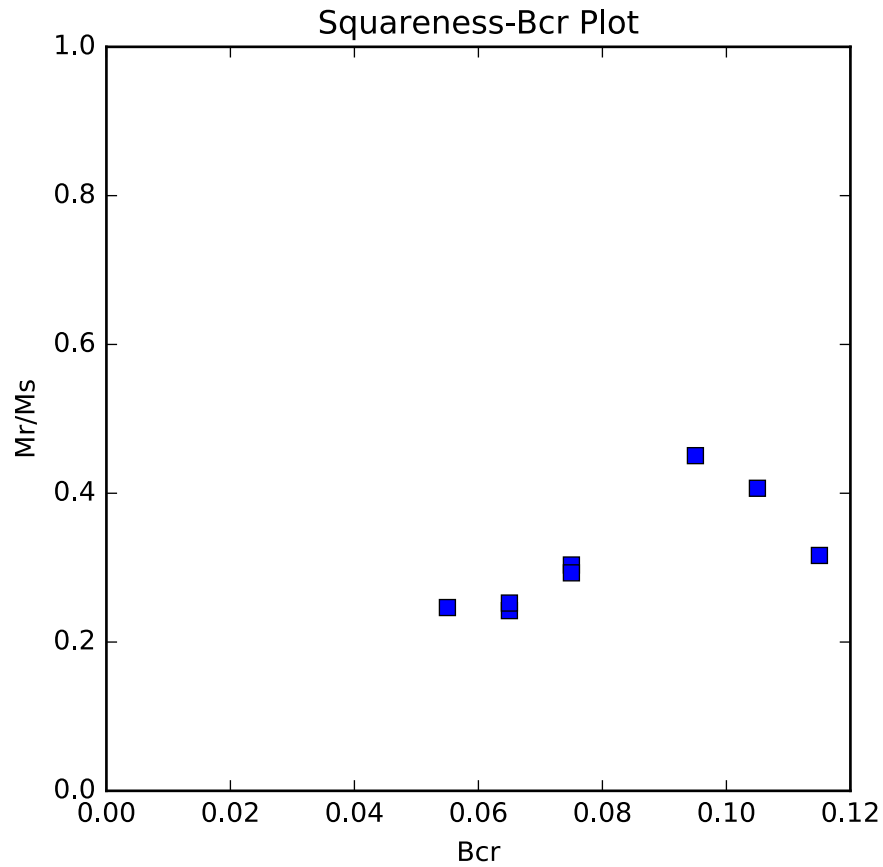
17 Day plots

Here we demonstrate the function `ipmag.dayplot`, which creates Day plots, squareness/coercivity and squareness/coercivity of remanence diagrams using hysteresis data.

```
In [32]: ipmag.dayplot(path_to_file='./Additional_Data/dayplot_magic/',  
                        hyst_file='dayplot_magic_example.dat',  
                        save=True, save_folder='Additional_Notebook_Output/day_plots/');
```







<matplotlib.figure.Figure at 0x114da9290>

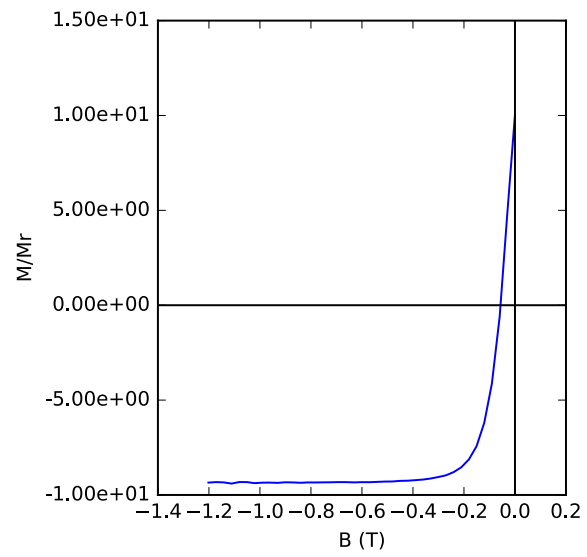
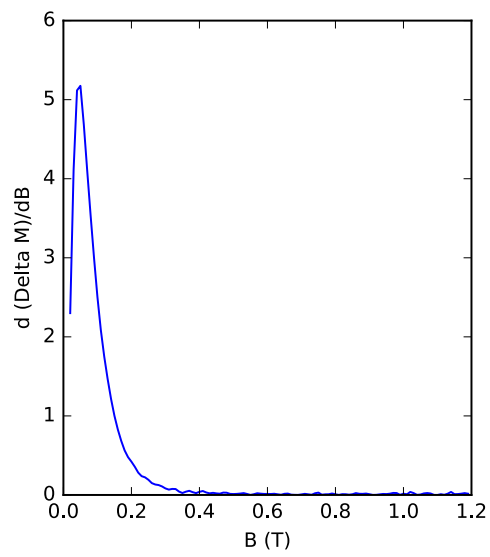
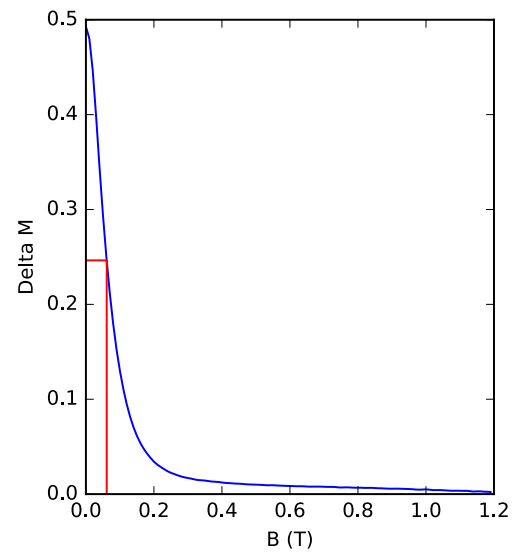
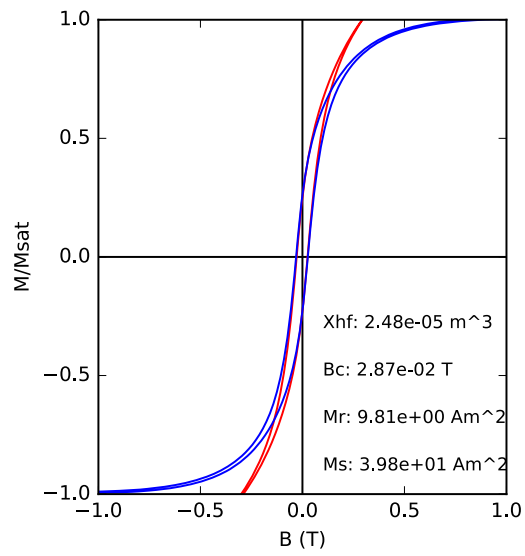
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18 Hysteresis Loops

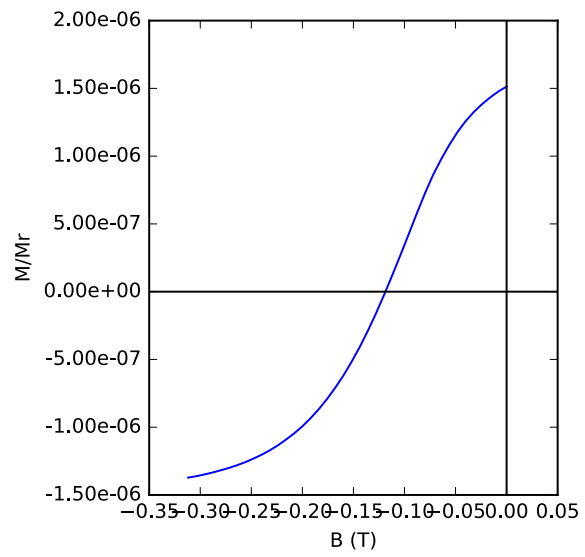
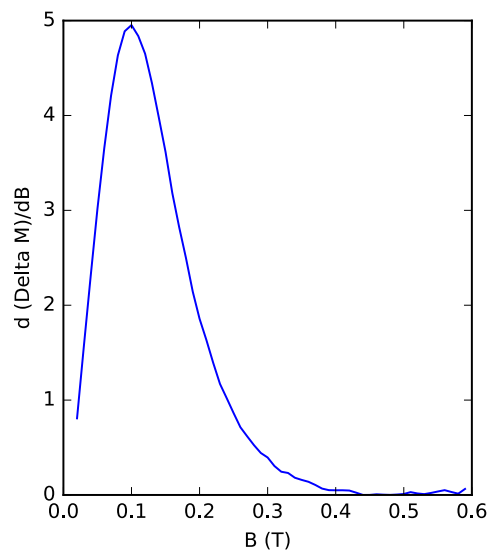
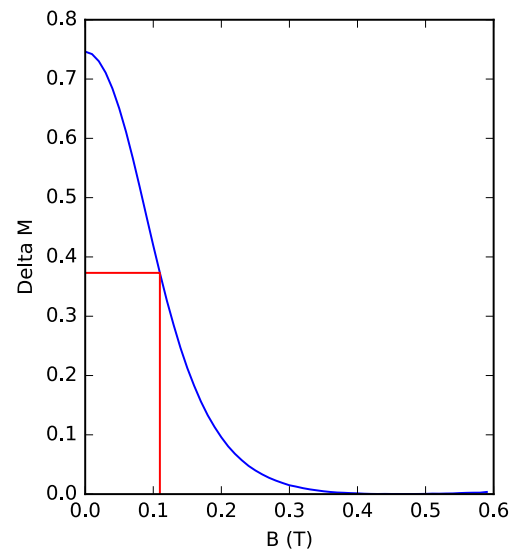
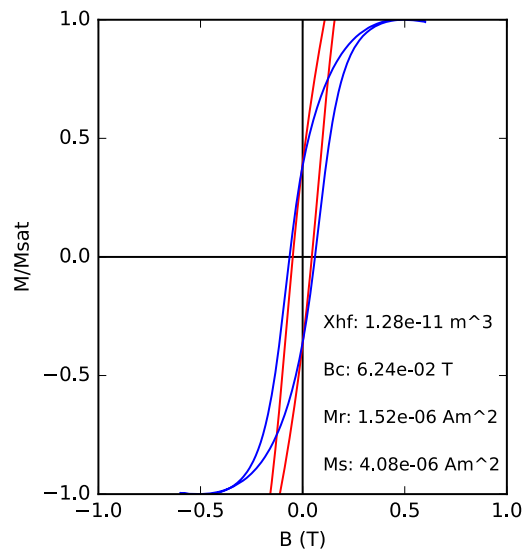
The function `ipmag.hysteresis_magic` also generates a set of hysteresis plots with data from a *magic_measurements* file.

```
In [33]: ipmag.hysteresis_magic(path_to_file='./Additional_Data/hysteresis_magic/',  
                                hyst_file='hysteresis_magic_example.dat', save=True,  
                                save_folder='./Additional_Notebook_Output/hysteresis')
```

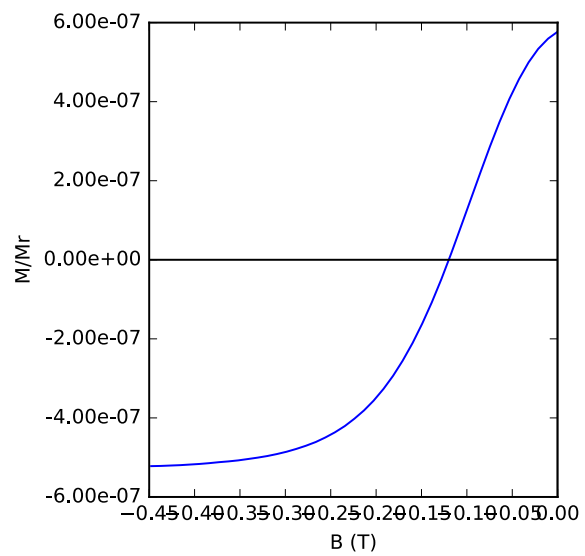
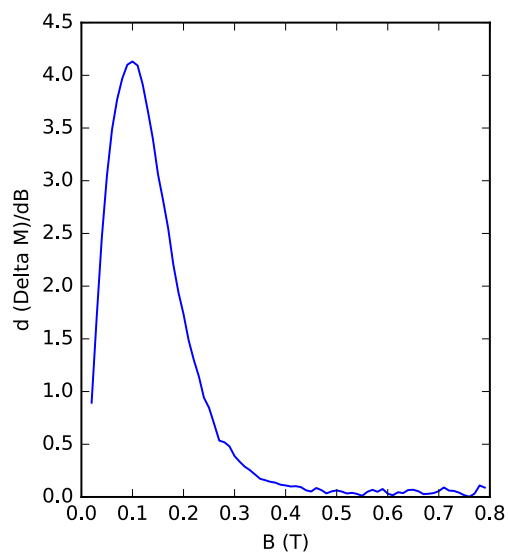
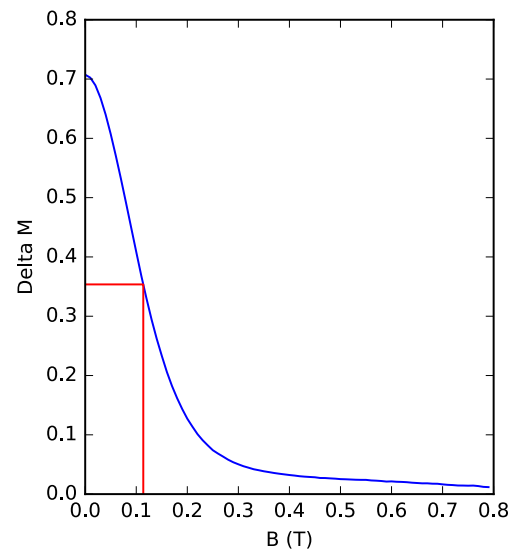
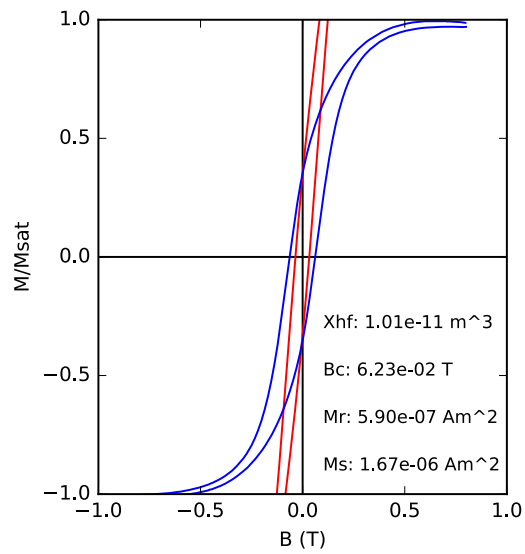
IS06a-1 1 out of 8



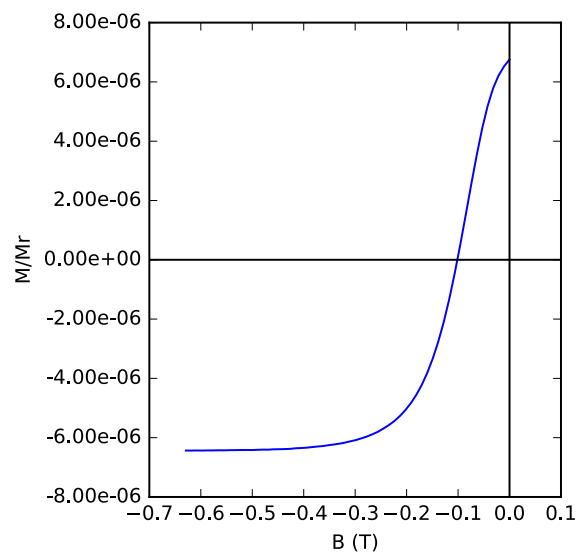
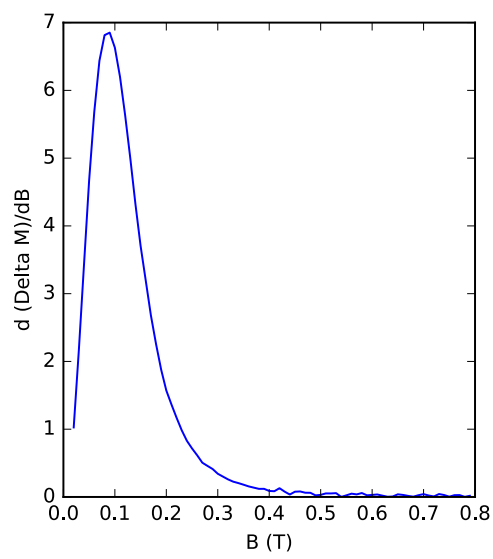
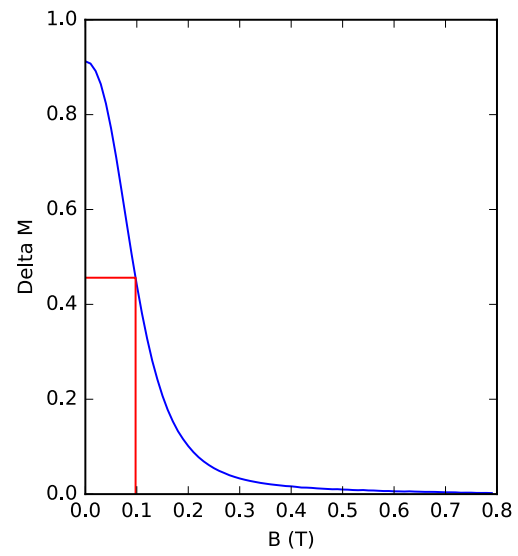
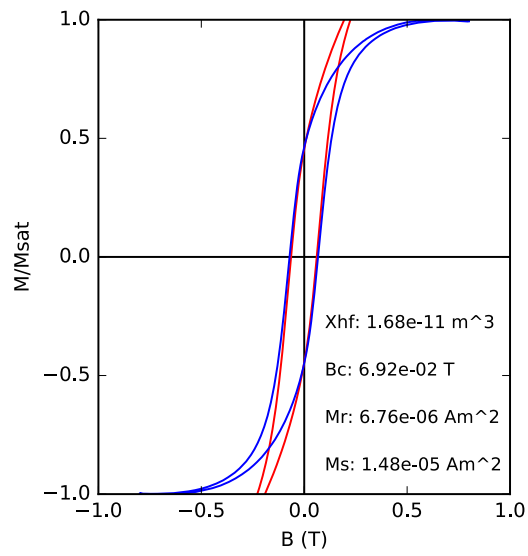
IS06a-2 2 out of 8



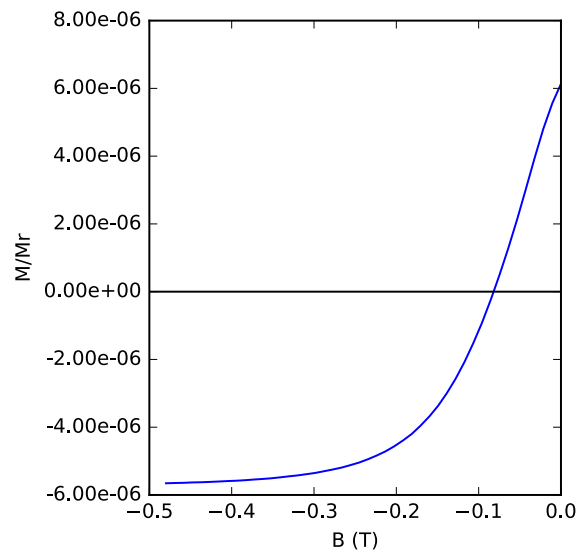
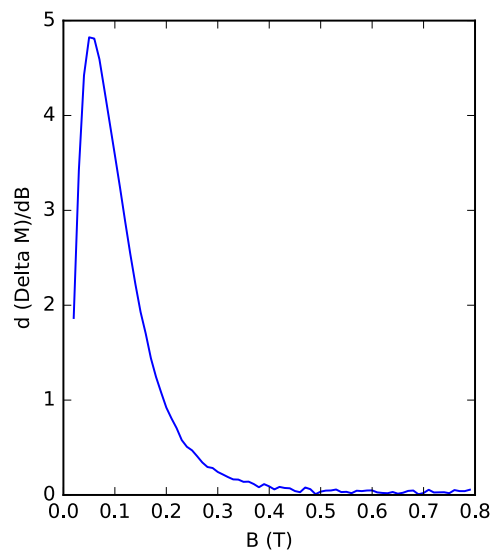
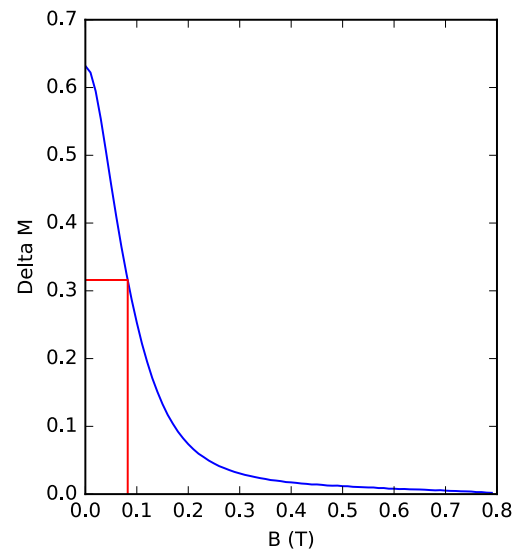
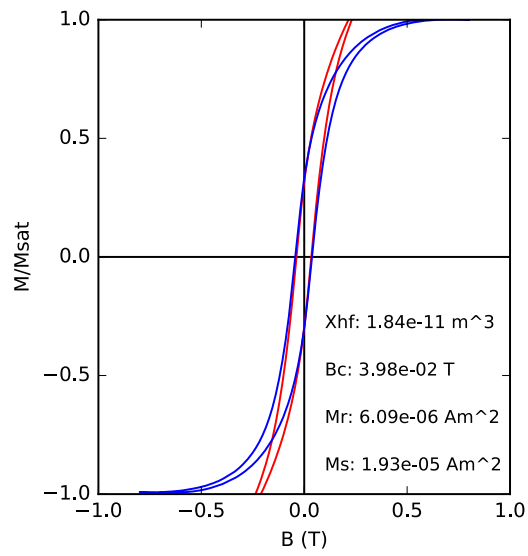
IS06a-3 3 out of 8



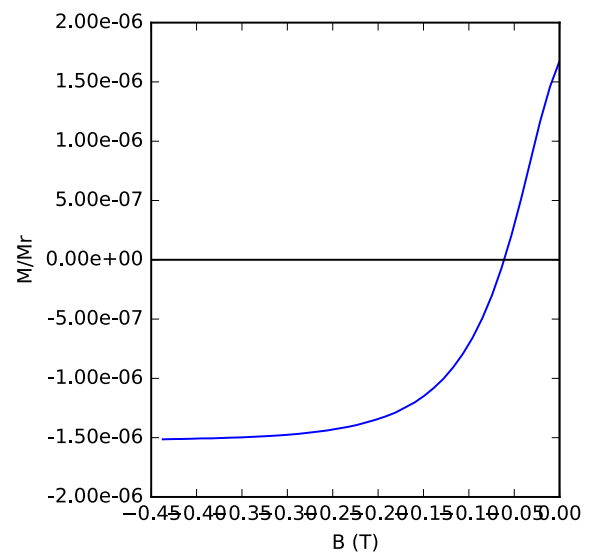
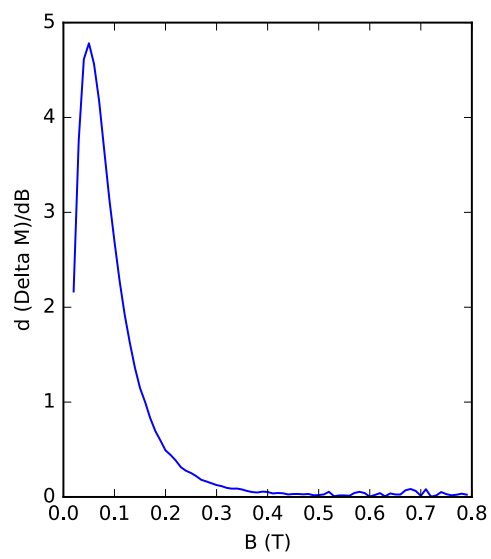
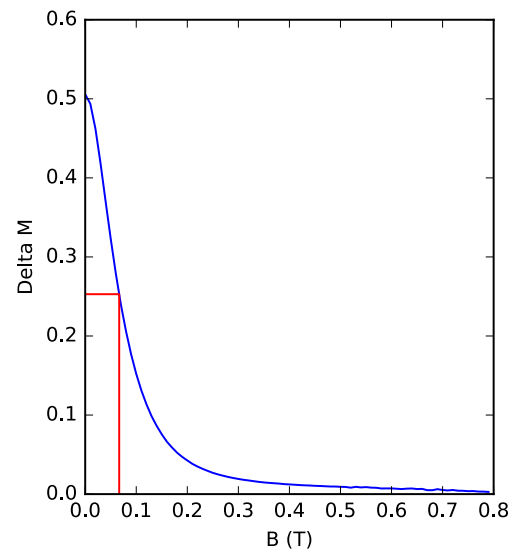
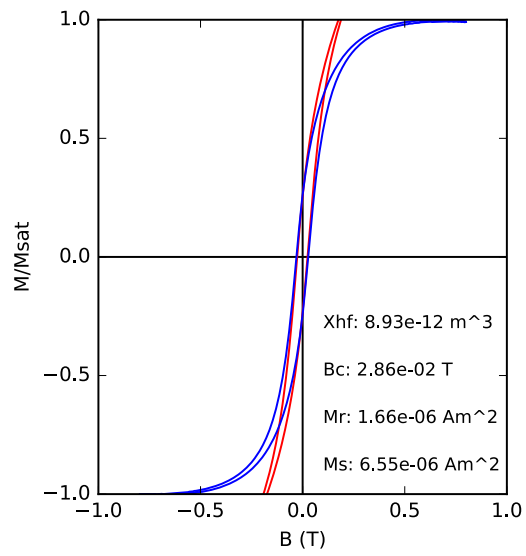
IS06a-4 4 out of 8



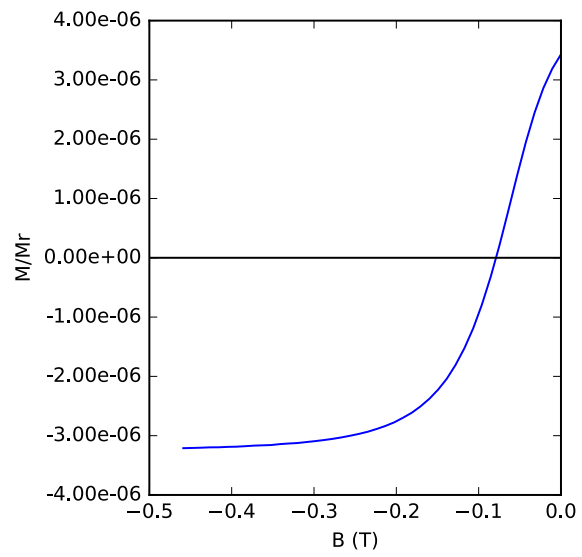
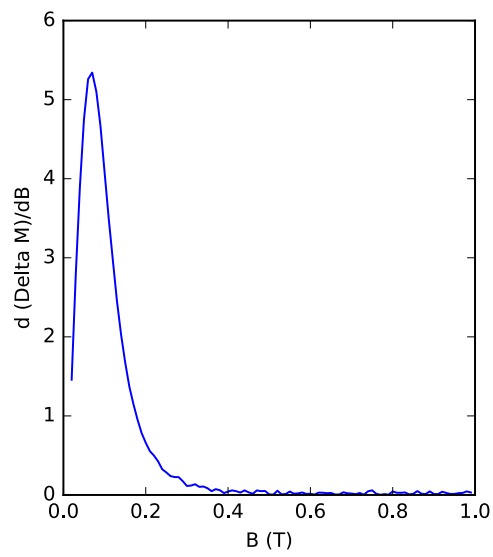
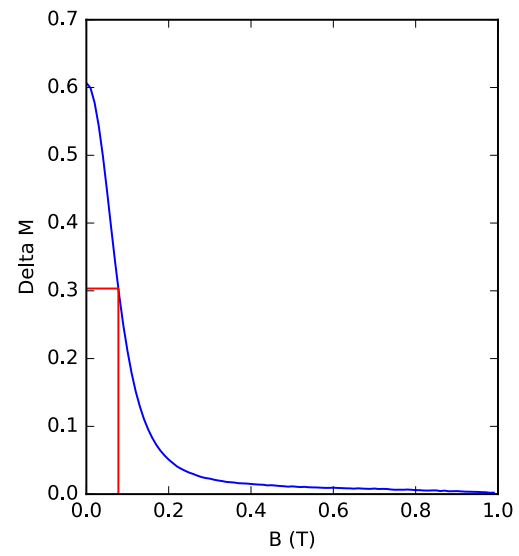
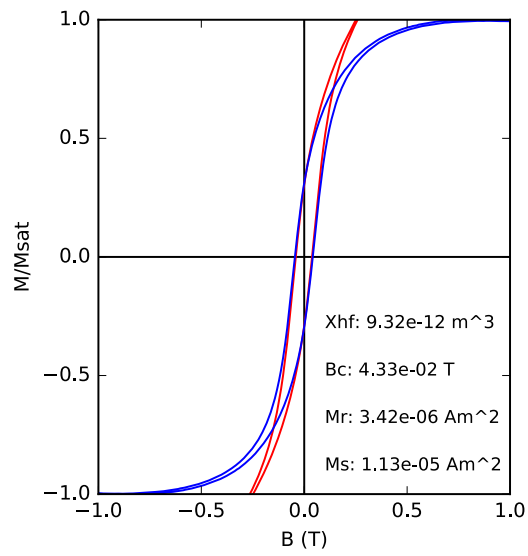
IS06a-5 5 out of 8



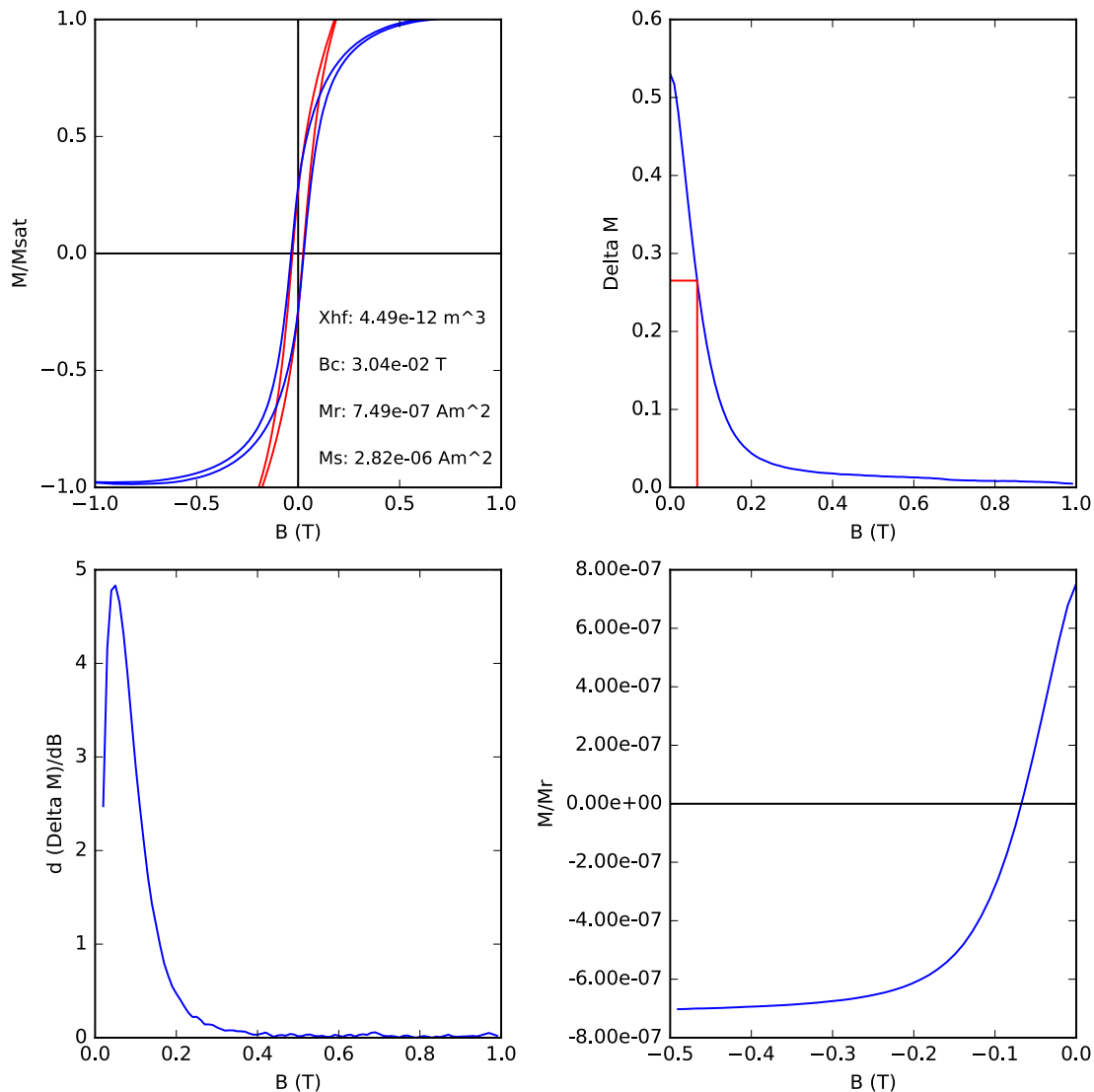
IS06a-6 6 out of 8



IS06a-8 7 out of 8



IS06a-9 8 out of 8



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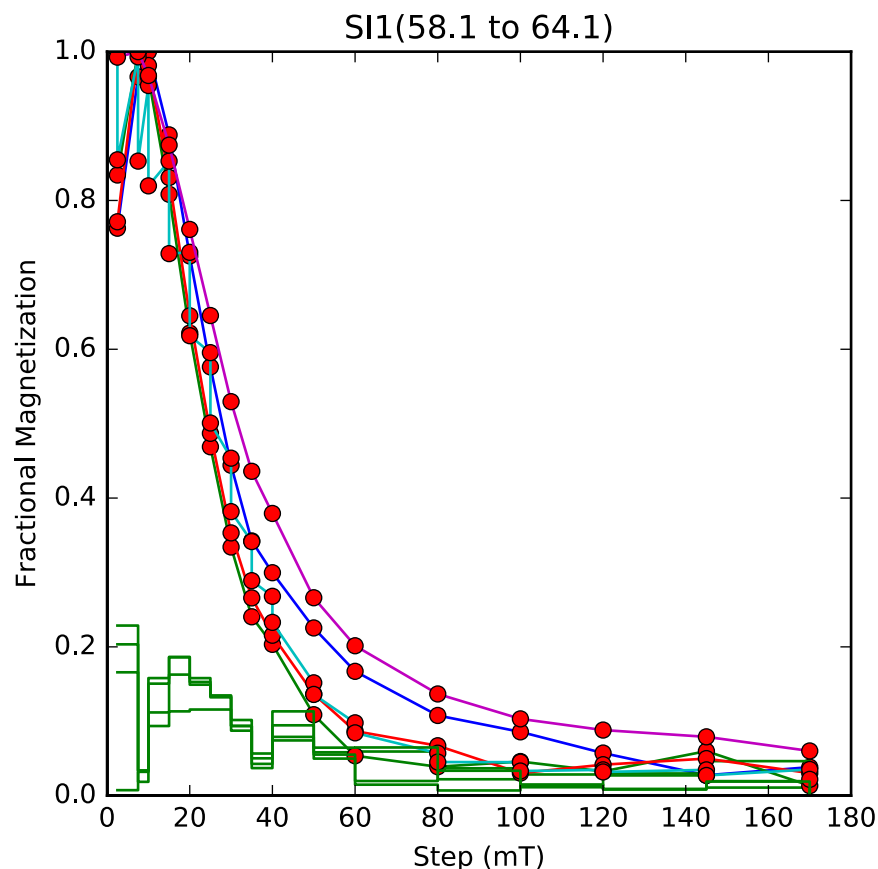
19 Demagnetization Curves

The function `ipmag.demag_magic` filters and plots demagnetization data. These data will be read and combined by expedition, location, site or sample according to the *plot_by* keyword argument. Alternatively, you can choose to plot each specimen measurement individually. By default, all plots generated by this function will be shown. If you only wish to plot a single subset of data, you can use the keyword argument *individual* to specify the name of the one site, location, sample, etc. that you would like to see.

Below, we use the *magic_measurements.txt* file of Swanson-Hysell et al., 2014 to plot demagnetization data by site. We then specify an individual site ('SI1(58.1 to 64.1)') that will plot alone. Like other functions, these plots can be optionally saved out of the notebook.

```
In [34]: ipmag.demag_magic(path_to_file='./Example_Data/Swanson-Hysell2014/',
                             plot_by='site', treat='AF', individual= 'SI1(58.1 to 64.1)')
```

13395 records read from ./Example_Data/Swanson-Hysell2014/magic_measurements.txt
 SI1(58.1 to 64.1) plotting by: er_site_name



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20 Interactive plotting

IPy Widgets are part of what makes the Jupyter notebook environment so powerful – these widgets allow user interaction with figures. We first demonstrate the use of the **interact** widget, imported below.

Note: If you do not have the ipywidgets package installed, you may choose to either install it through Anaconda or Enthought (depending on your Python distribution), manually install it (a bit more difficult), or simply skip the next few blocks of code. Below are quick installation instructions for those with either an Anaconda or Enthought Canopy distribution.

Installation on Anaconda

On the command line, enter


```
conda install ipywidgets
```

Make sure this installs within the Python 2 environment (if you have Python 3 as your default environment).

Installation on Enthought Canopy

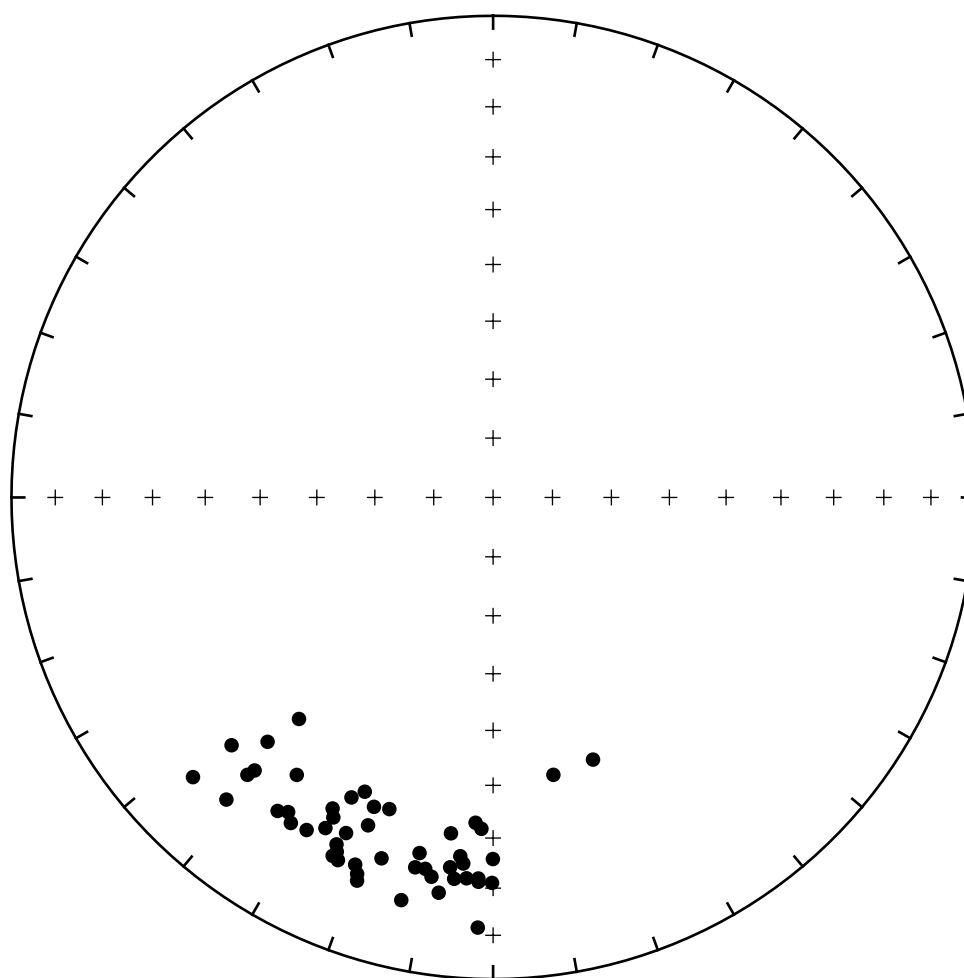
Open the Canopy application and navigate to the Package Manager. Search for and install ipywidgets.

```
In [35]: from ipywidgets import interact
```

The **interact** widget allows adjustable values (within specified bounds) to all keyword arguments of a function. It can be used as a wrapper function, as seen below. Here we create a new function, **squish_interactive**, which streamlines the **ipmag.squish** function and automatically inputs the fisher-distributed directions created at the beginning of the notebook. This new function also allows us to reduce the keyword arguments to the *factor* variable, which is the only value we want to be actively adjustable. Finally, to make the **squish_interactive** function interactive in the notebook, we “wrap” this function with **@interact** placed directly above our new function.

```
In [36]: @interact
def squish_interactive(flattening_factor=(0.,1.,.1)):
    squished_incs = []
    for inclination in inclinations:
        squished_incs.append(ipmag.squish(inclination, flattening_factor))

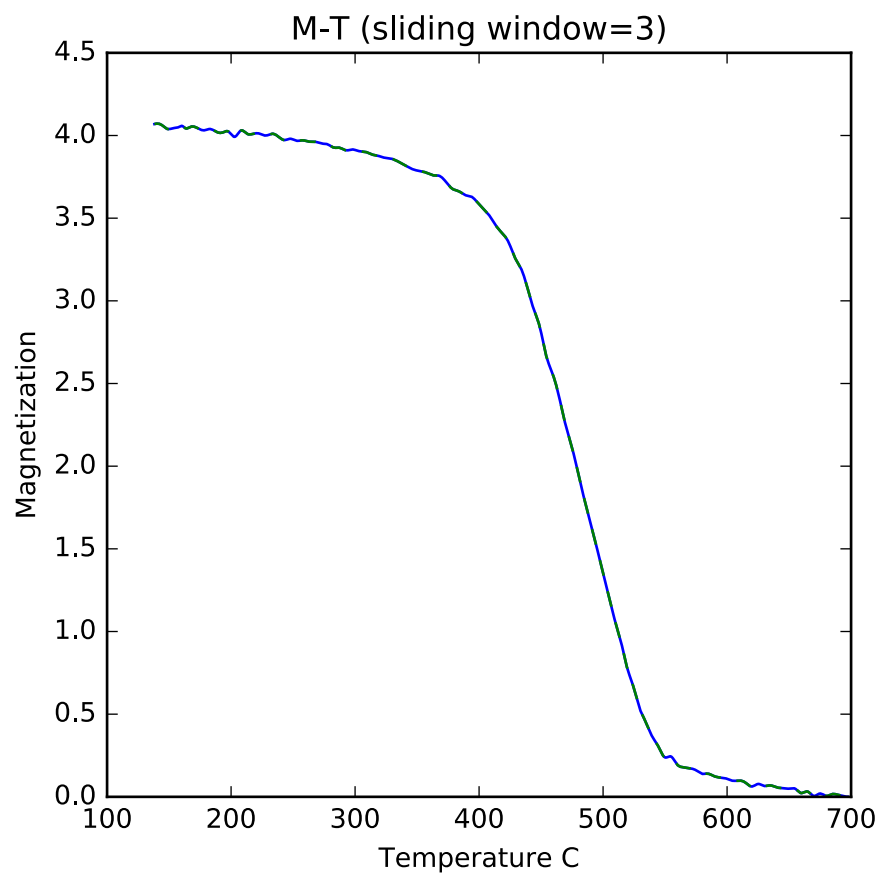
    # plot the squished directional data
    plt.figure(num=1,figsize=(6,6))
    ipmag.plot_net(1)
    ipmag.plot_di(declinations,squished_incs)
```

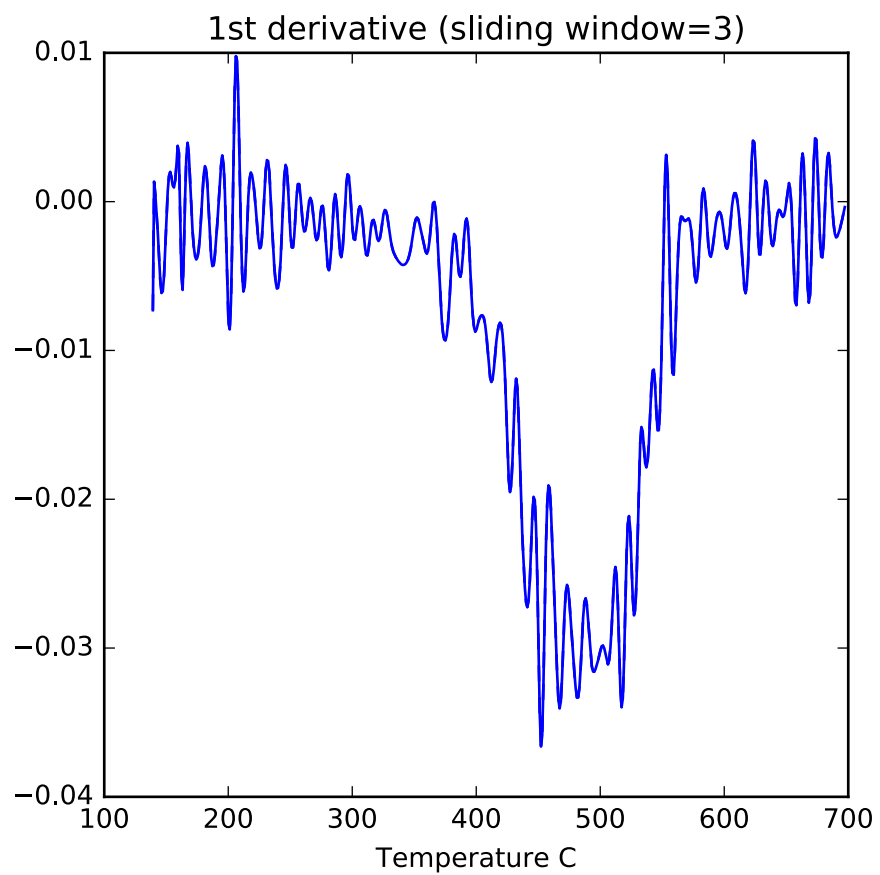


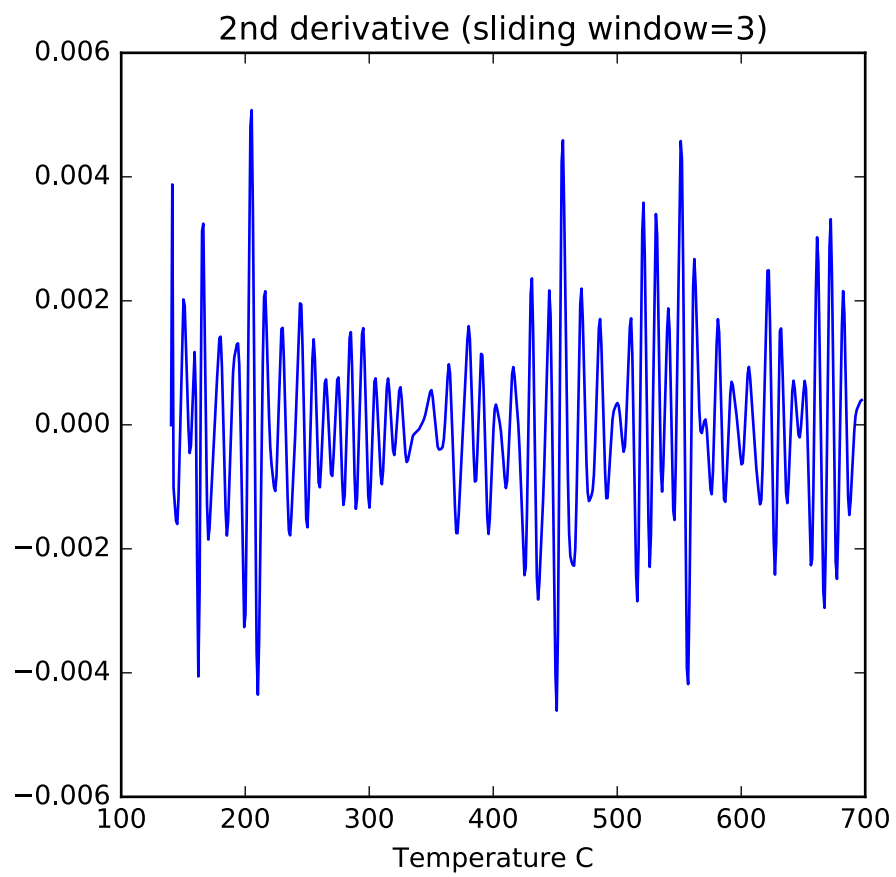
interact can also be used as a regular function call – the name of the interactive function is passed as the first argument, followed by the adjustable keyword arguments. Below, we demonstrate passing the *curie* function's parameters to **interact**.

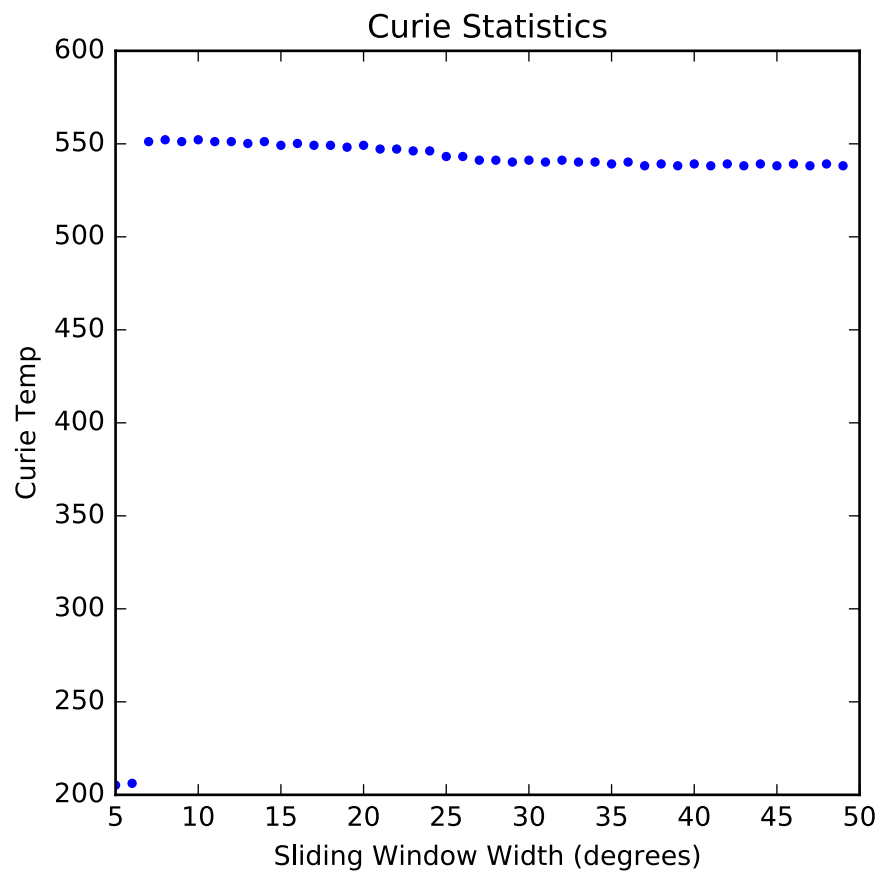
```
In [37]: interact(ipmag.curie, path_to_file='./Additional_Data/curie/',file_name='curie_example.dat',wi
```

```
second deriative maximum is at T=205
```









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In []: