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 other example PmagPy notebook in this repository (Example_PmagPy_Notebook.ipynb in https://github.com/PmagPy/2016_Tauxe-et-al_PmagPy_Notebooks (https://github.com/PmagPy/2016_Tauxe-et-al_PmagPy_Notebooks) which can also be seen here: http://pmagpy.github.io/Example_PmagPy_Notebook.html (http://pmagpy.github.io/Example_PmagPy_Notebook.html)) includes additional instructions on how to get started using PmagPy in the notebook. That example notebook contains a more extended work flow on two data sets while this notebook contains numerous code vignettes to illustrate additional availible functionality within the PmagPy module.

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Paleomagnetic Data Analysis

Basic Functions

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Additional Features of the Jupyter Notebook

Note: This notebook makes use of additional scientific Python modules: ***pandas*** *for reading, displaying, and using data with a dataframe structure,* ***numpy*** *array computations and *matplotlib* for plotting. These modules come standard with scientific computing distributions of Python or can be installed separately as needed.

In [1]:

```
# With the PmagPy folder in the PYTHONPATH,
# the function modules from PmagPy can be imported
import pmagpy.ipmag as ipmag
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',}
```

The dipole equation

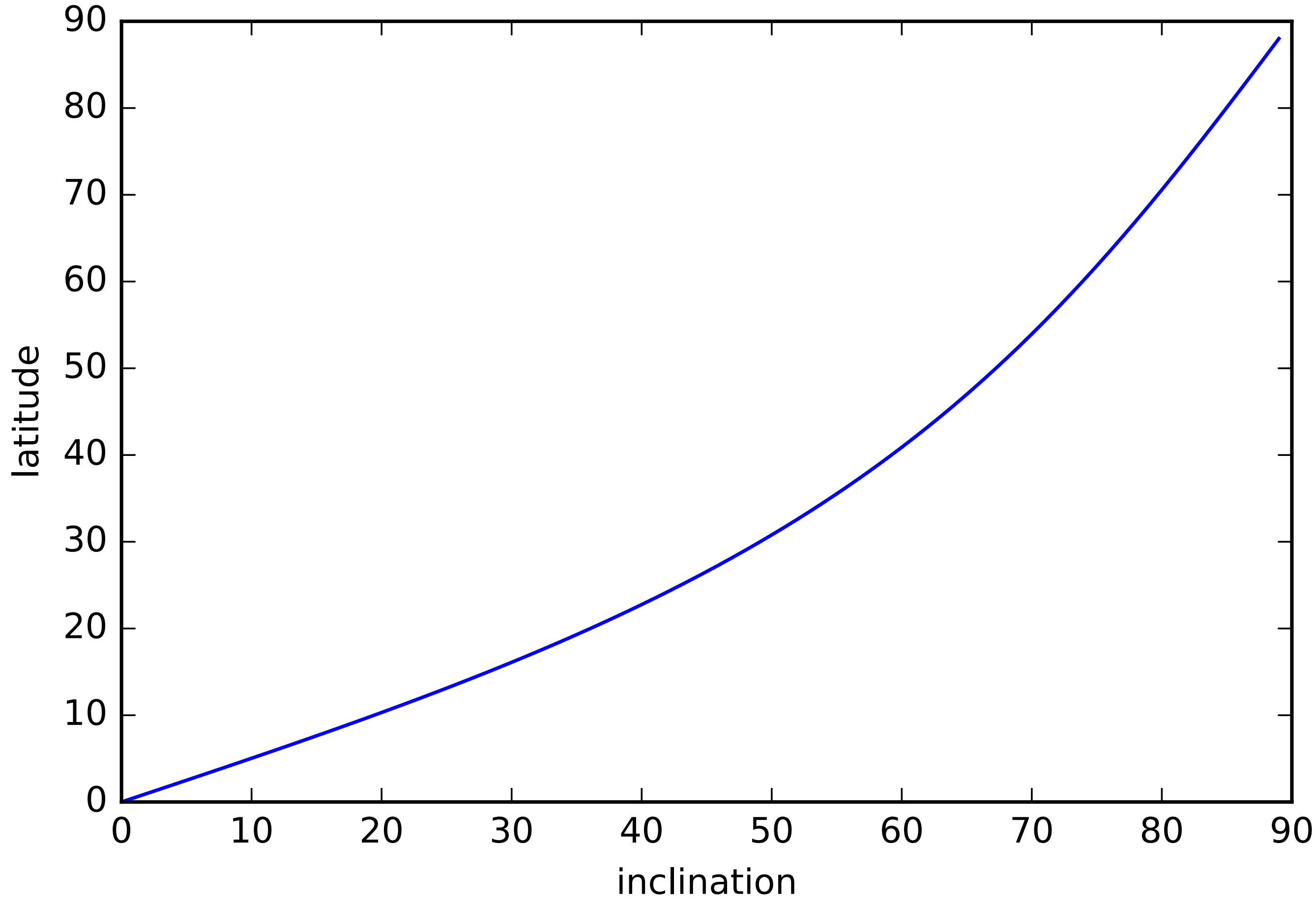
The following demonstrates the use of a simple function (**ipmag.lat_from_inc**) which uses the dipole equation to return expected latitude from inclination data as predicted by a pure geocentric axial dipole. The expected inclination for the geomagnetic field can be calculated from a specified latitude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

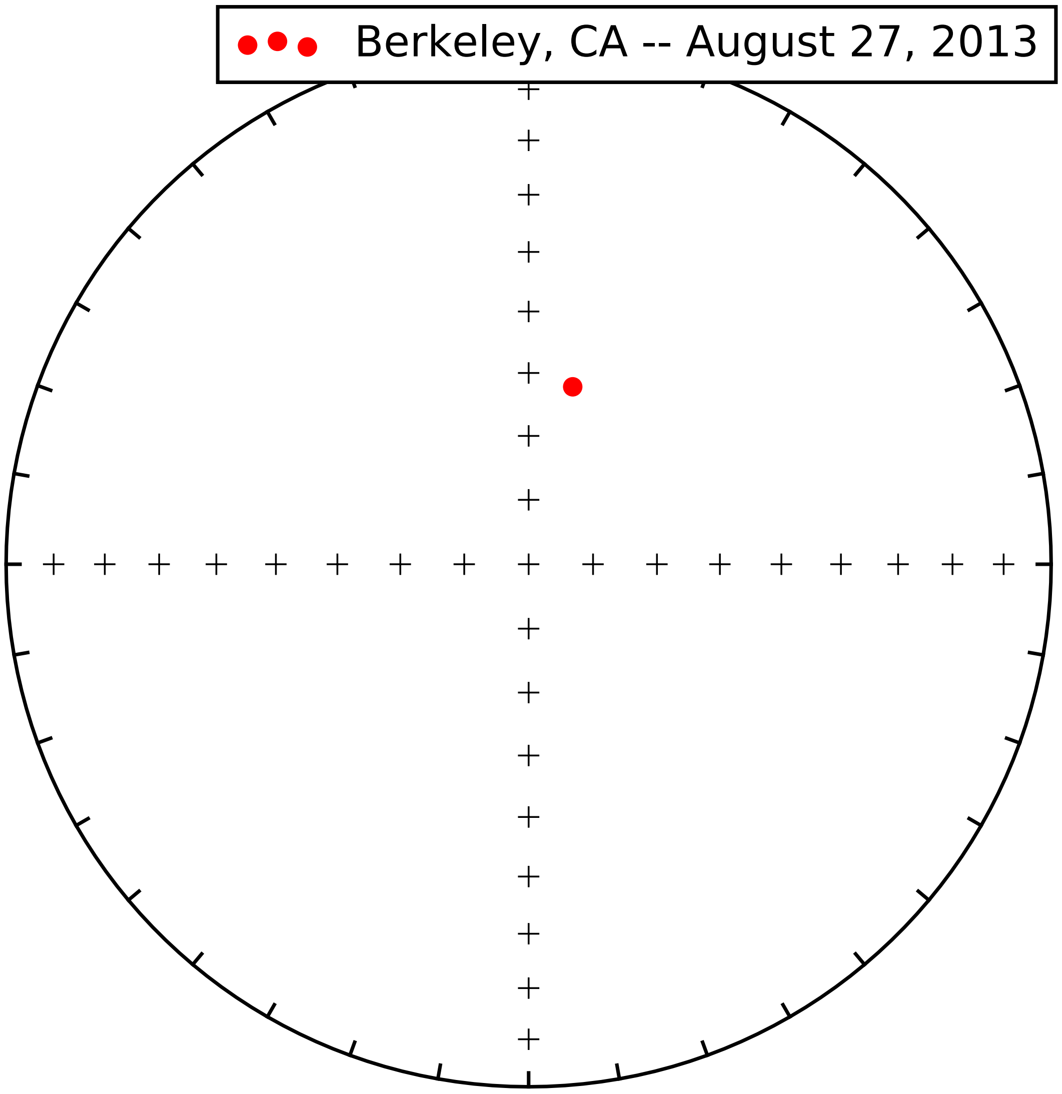
To ensure the figure is displayed properly and then cleared from the namespace, it is good practice to end such a code block with the following:

```
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 -- Augus
t 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")
```

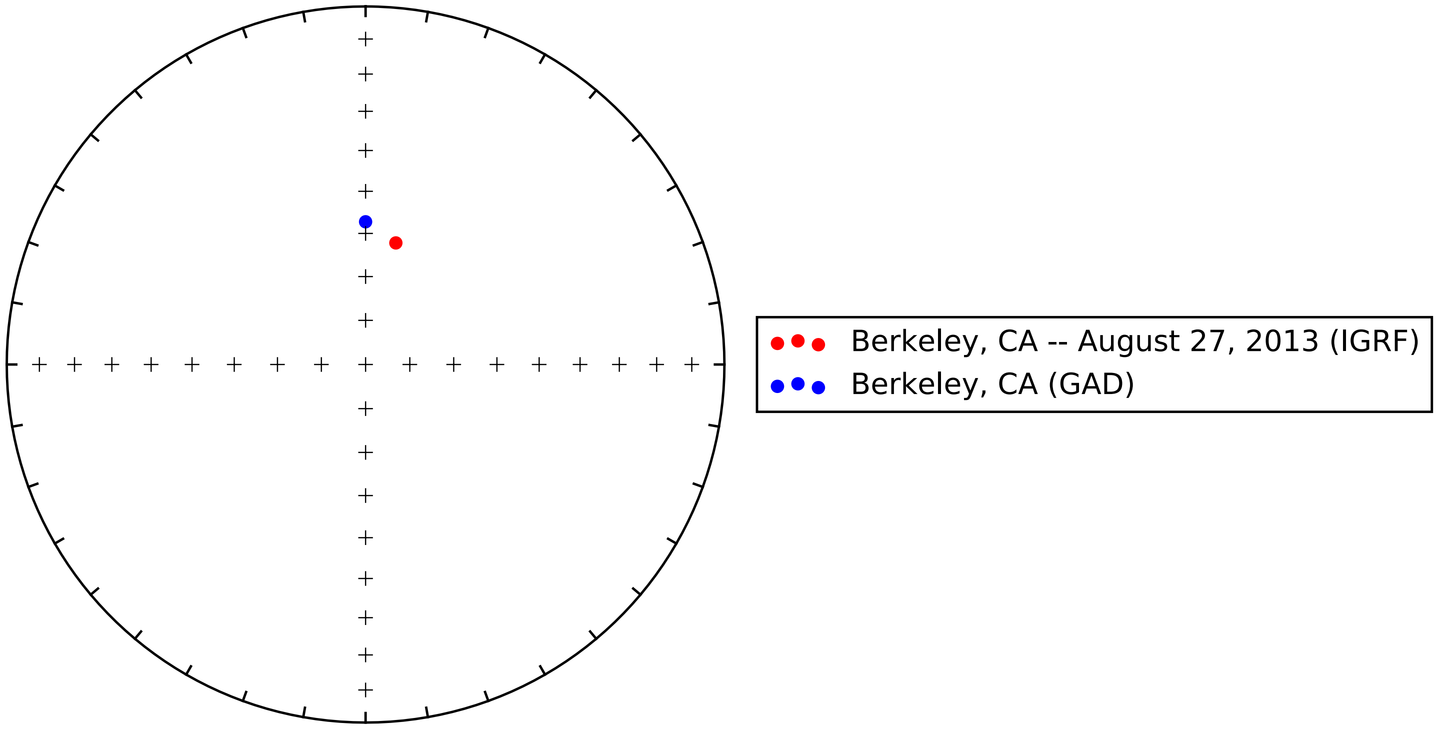
or

```
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 -- Augus
t 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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Calculate the Angle Between Directions

While **ipmag** functions have been optimized to preform 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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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]:

```
[[190.45811580955265, 50.061261452480771, 1.0],
 [199.33544934875593, 55.403087445353542, 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]
```

```
190.45811581
50.0612614525
```

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	190.458116	50.061261	1
1	199.335449	55.403087	1
2	172.790533	54.044842	1
3	186.632529	48.049753	1
4	215.501766	45.890581	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': 3.3503228585732363,
 'csd': 13.293077796925324,
 'dec': 202.06246797501123,
 'inc': 51.041290971640322,
 'k': 37.129486661729402,
 'n': 50,
 'r': 48.680294170333738}
```

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

In [13]:

```
ipmag.print_direction_mean(fisher_mean)
```

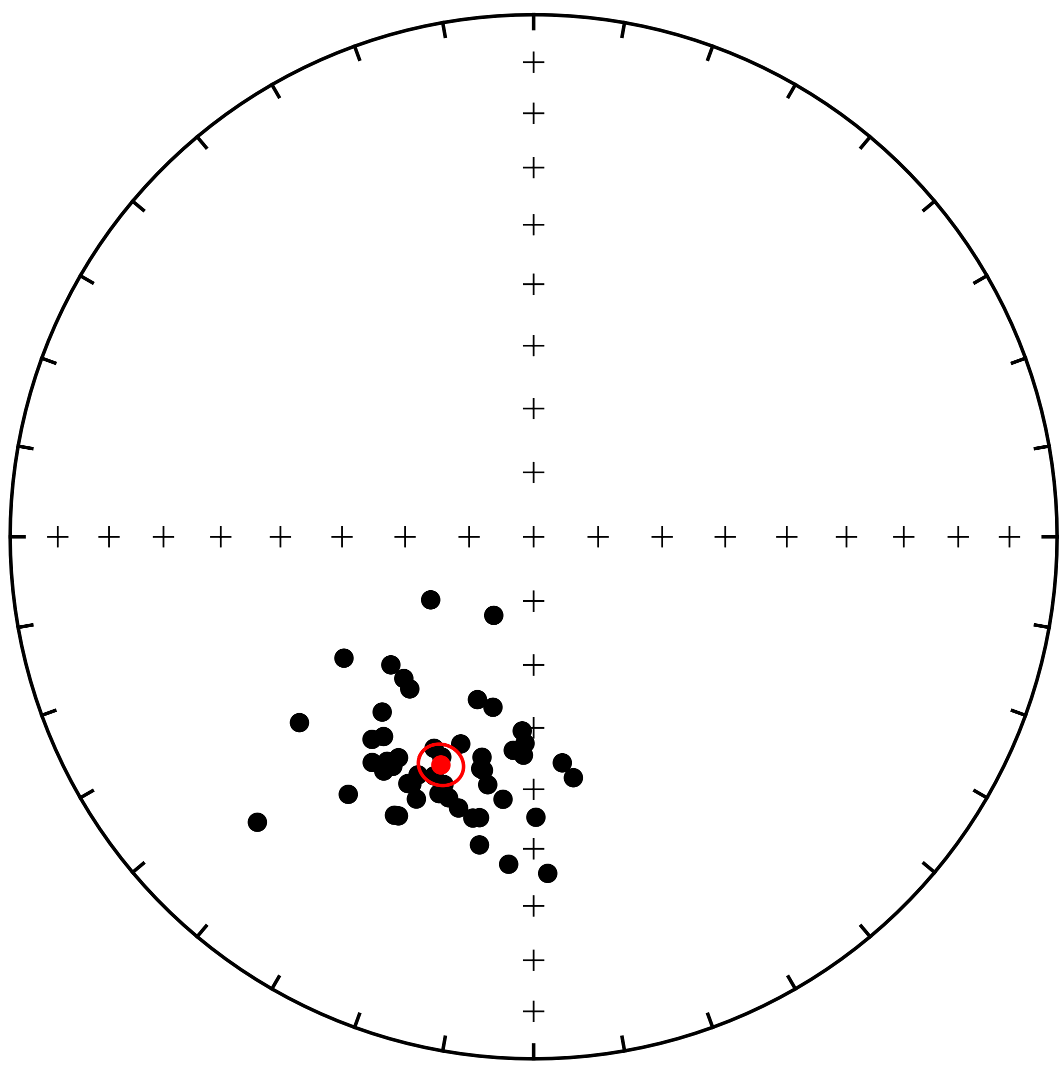
```
Dec: 202.1  Inc: 51.0
Number of directions in mean (n): 50
Angular radius of 95% confidence (a_95): 3.4
Precision parameter (k) estimate: 37.1
```

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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Flip polarity of directional data

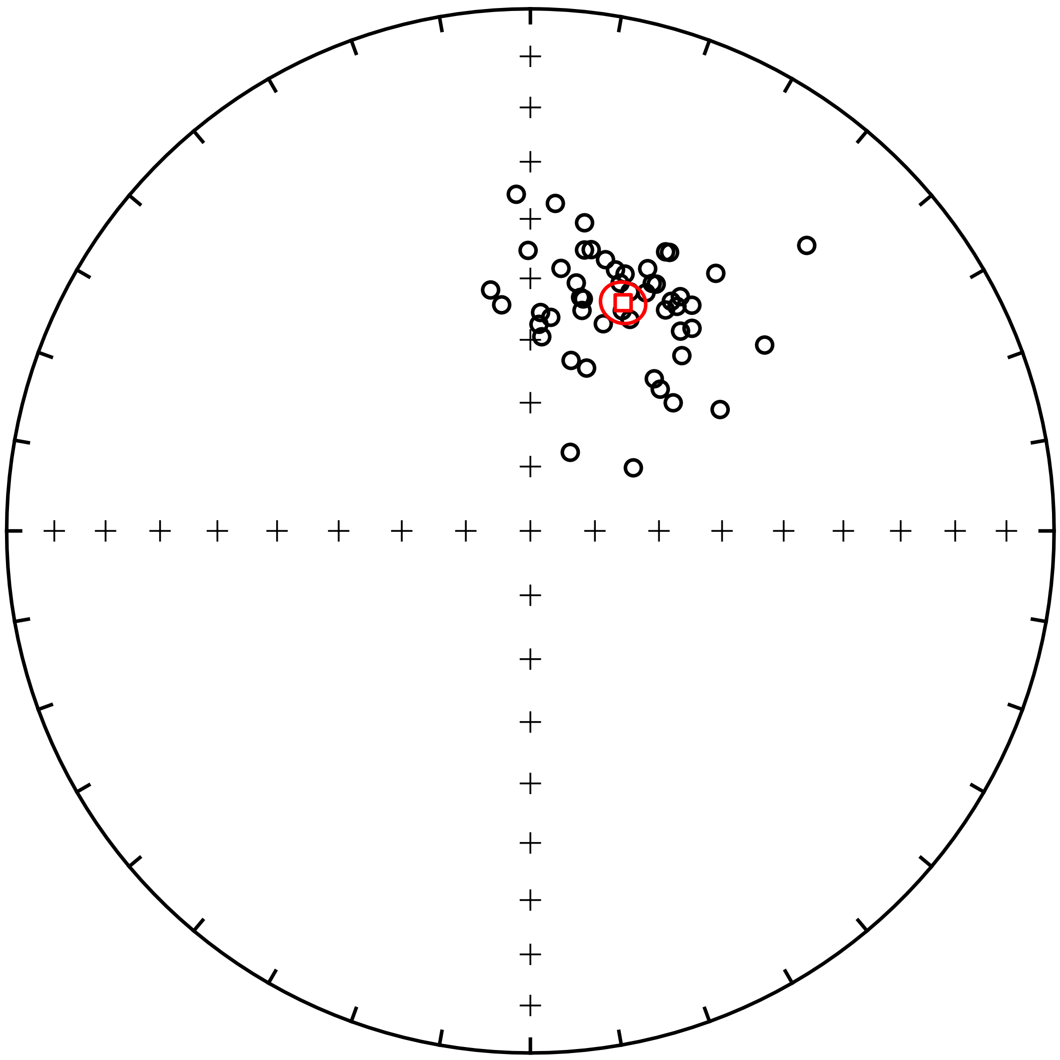
Let's flip all the directions (find their antipodes) of the Fisher-distributed population using the function **ipmag.do_flip()** function and plot the resulting directions.

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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Test directional data for Fisher distribution

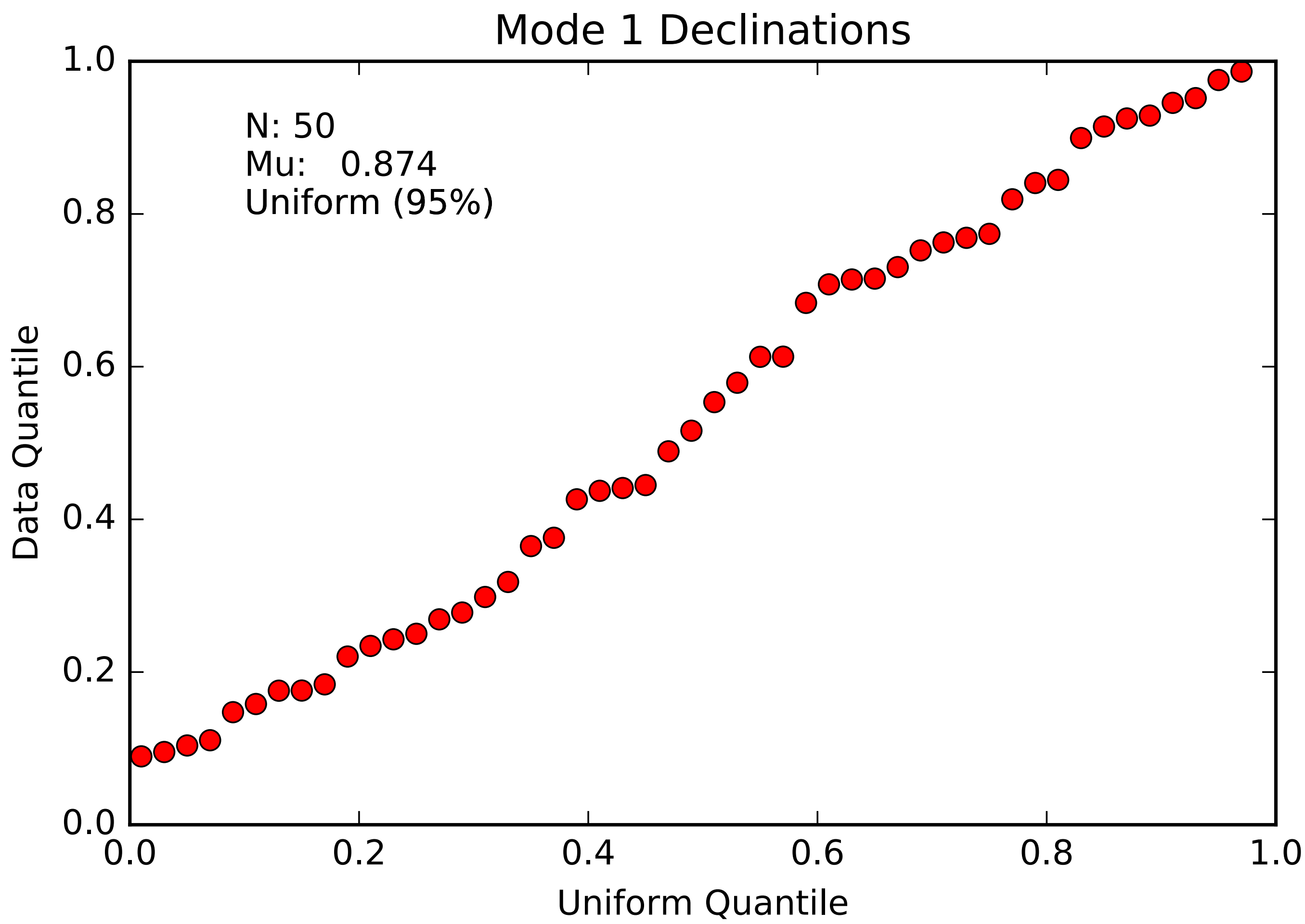
The function **ipmag.fishqq** tests whether directional data are Fisher-distributed. Let's use this test on the random Fisher-distributed directions we just created (it should pass!).

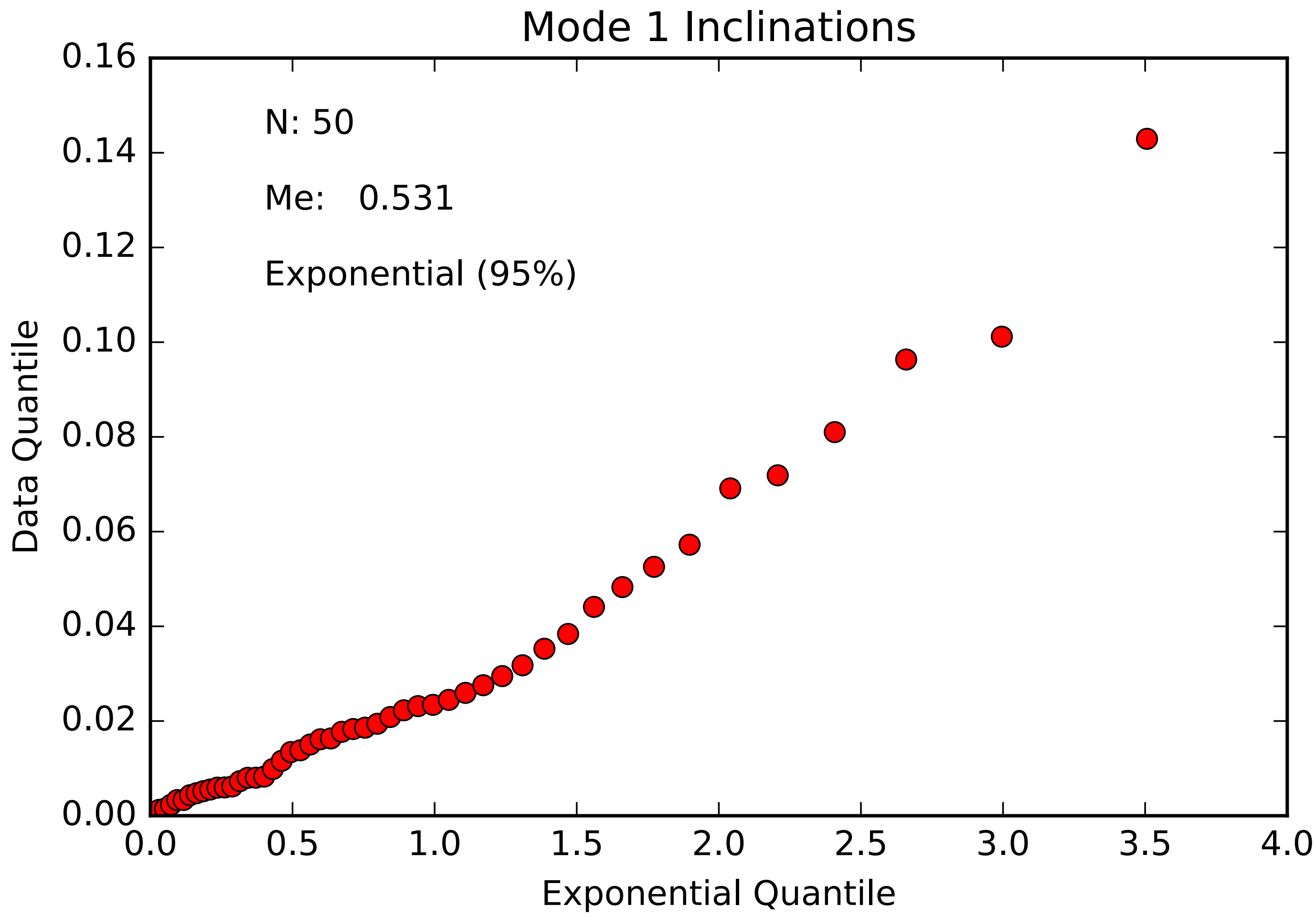
In [16]:

```
ipmag.fishqq(declinations, inclinations)
```

Out[16]:

```
{'Dec': 201.97276914475819,
 'Inc': 51.014325707259353,
 'Me': 0.53066629539385524,
 'Me_critical': 1.094,
 'Mode': 'Mode 1',
 'Mu': 0.87432565831513398,
 'Mu_critical': 1.207,
 'N': 50,
 'Test_result': 'consistent with Fisherian model'}
```





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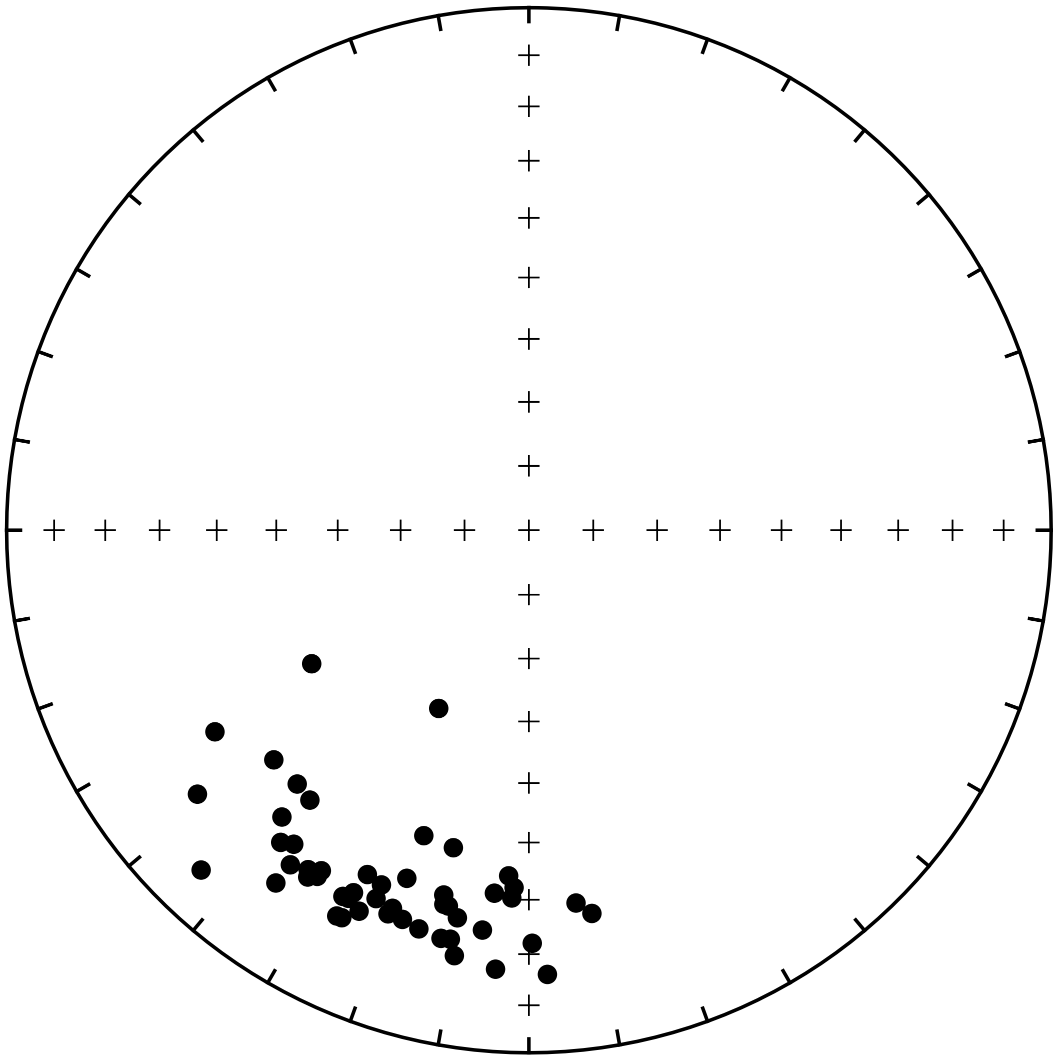
Squish directional data

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': 4.1230242552381018,
 'csd': 16.247832872173813,
 'dec': 202.15768542577069,
 'inc': 27.285568627714493,
 'k': 24.853018972860934,
 'n': 50,
 'r': 48.02840853847546}
```

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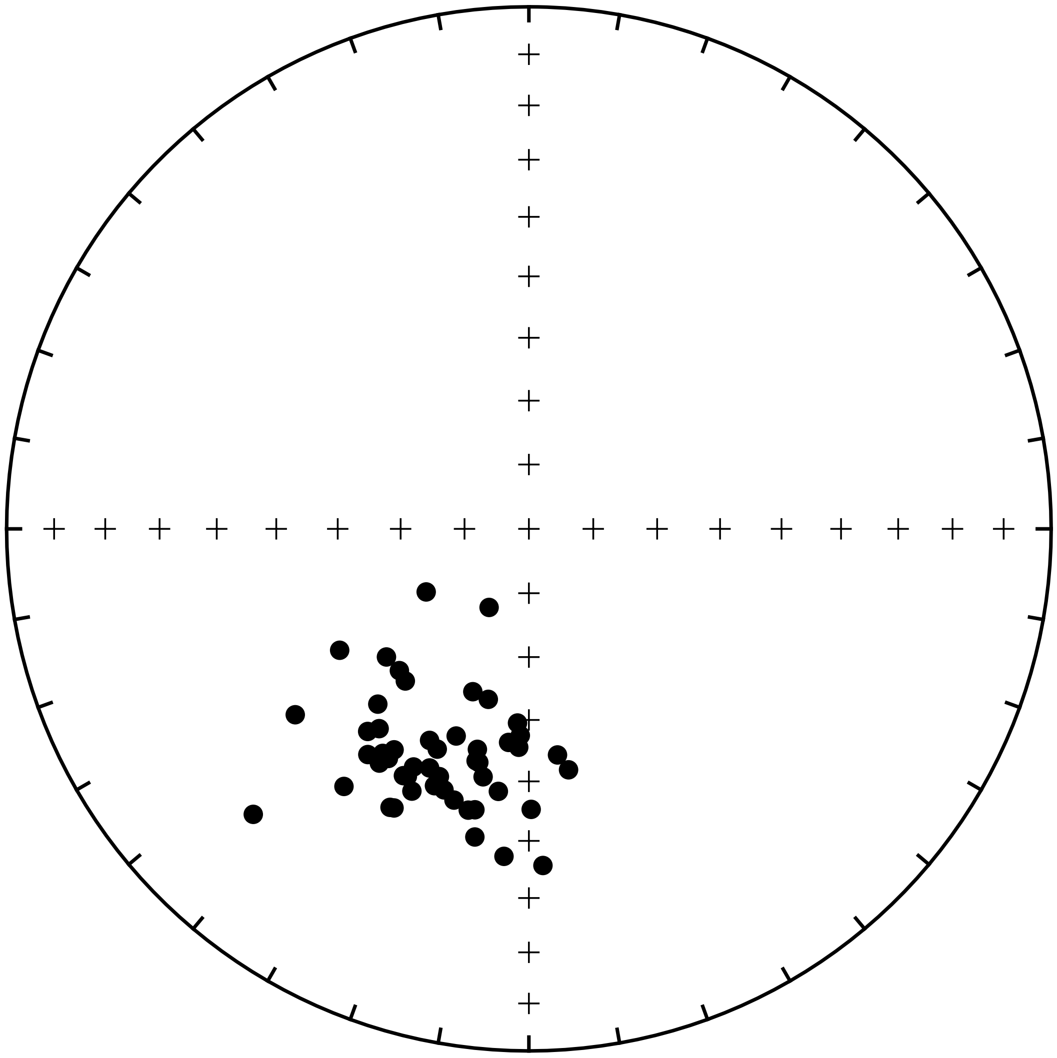
Unsquish directional data

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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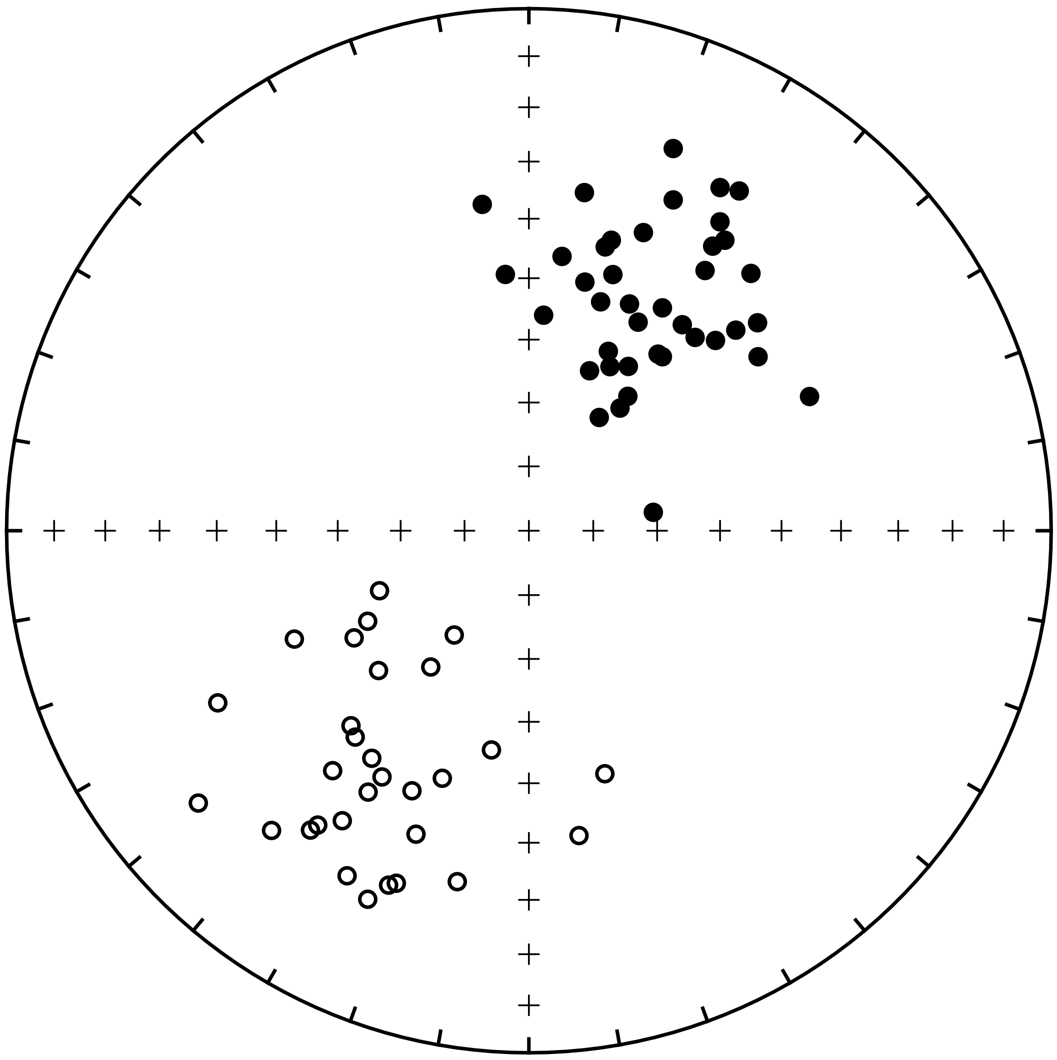
Bootstrap Reversal Test

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]:

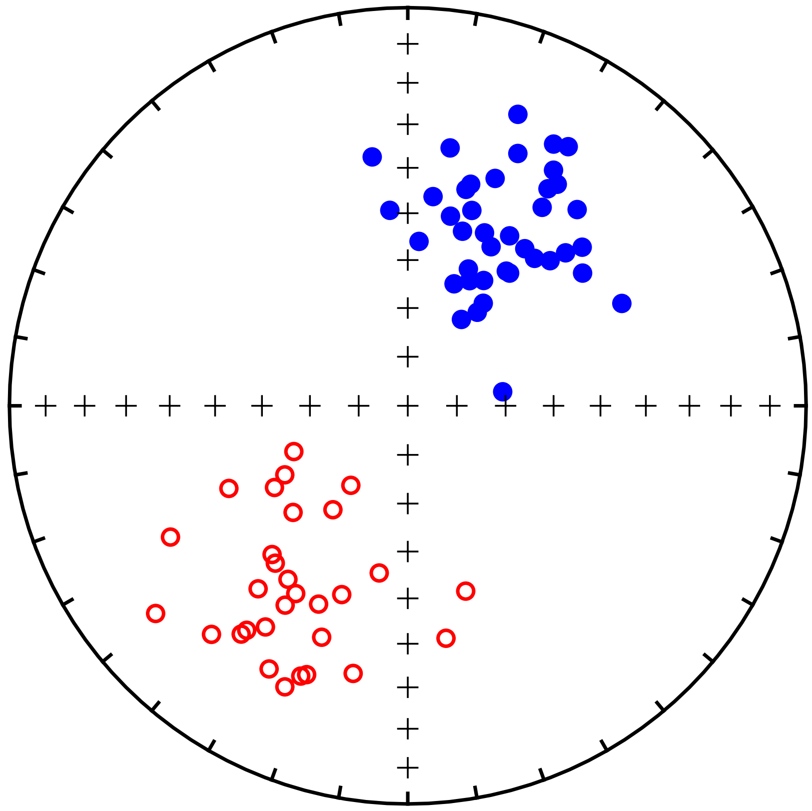
```
normal_directions = ipmag.fishrot(k=20,n=40,dec=30,inc=45)
reversed_directions = ipmag.fishrot(k=20,n=30,dec=210,inc=-45)
combined_directions = normal_directions + reversed_directions

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

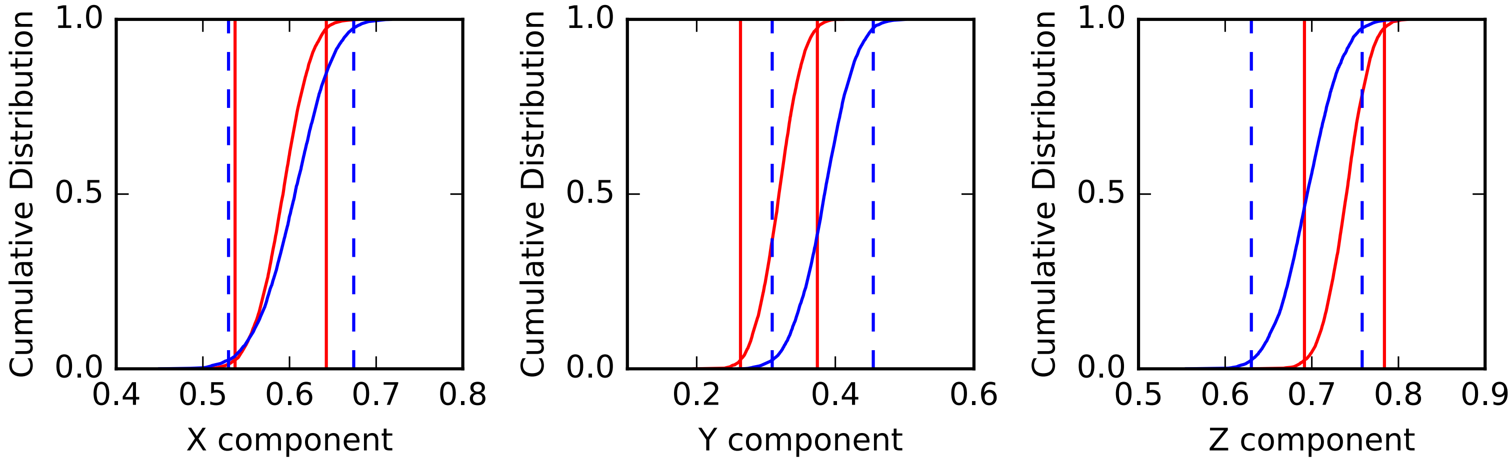


In [21]:

```
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]:

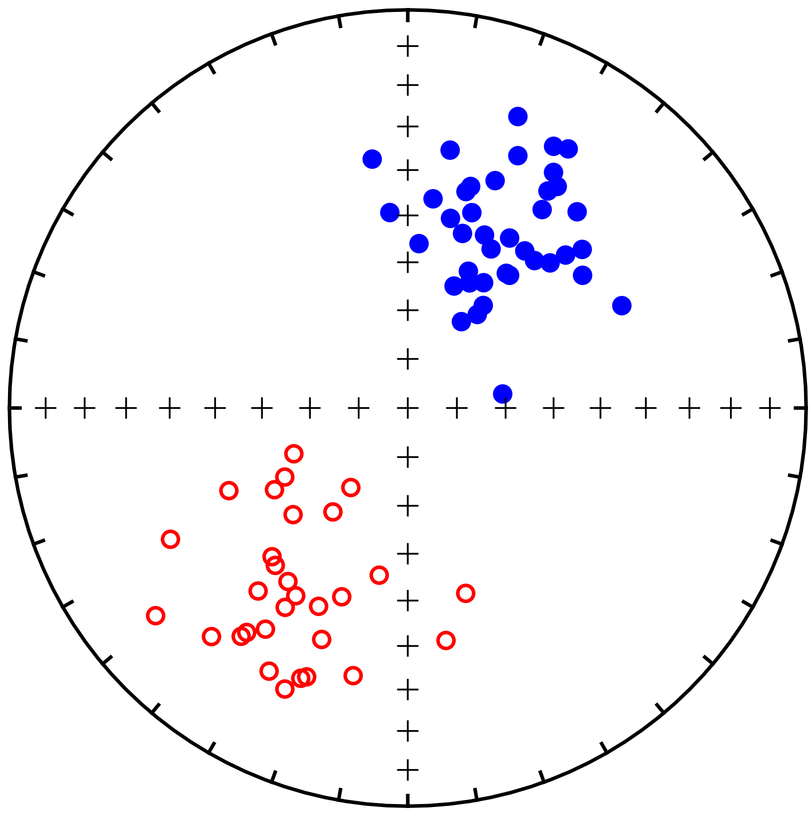
```
ipmag.reversal_test_MM1990(di_block=combined_directions,
                           plot_CDF=True, plot_stereo=True,
                           save=True, save_folder= './Additional_Notebook_Output
/')
```

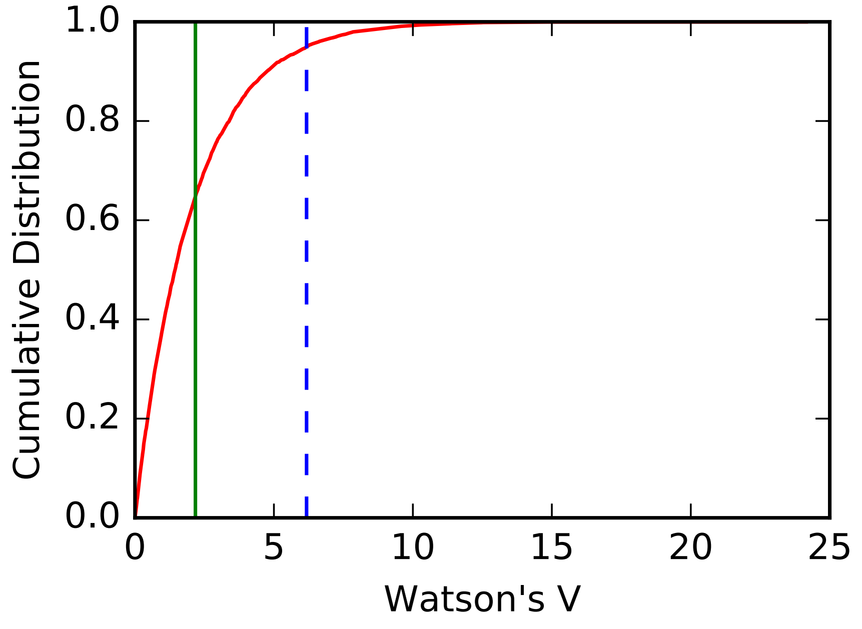
```
Results of Watson V test:

Watson's V:           2.2
Critical value of V:  6.2
"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:   7.8
The McFadden and McElhinny (1990) classification for
this test is: 'B'
```





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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]:

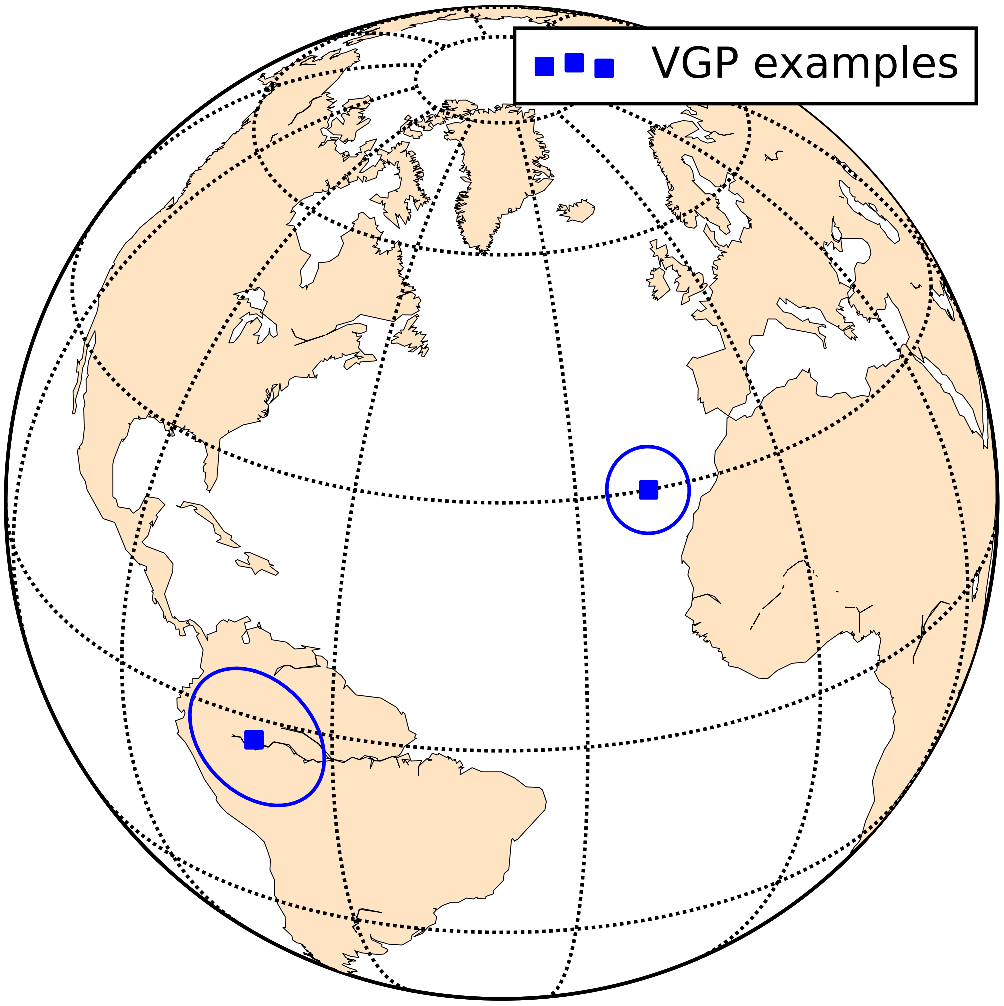
```
# initiate figure and specify figure size
plt.figure(figsize=(5, 5))

# 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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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	vgp_lat	vgp_lon
0	190.458116	50.061261	1	37.97	-122.27	30.844481	-20.496202	228.1528
1	199.335449	55.403087	1	37.97	-122.27	35.937411	-13.956726	221.6945
2	172.790533	54.044842	1	37.97	-122.27	34.579460	-17.142599	243.9375
3	186.632529	48.049753	1	37.97	-122.27	29.086177	-22.657258	231.4506
4	215.501766	45.890581	1	37.97	-122.27	27.284244	-16.758869	205.1136

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]:

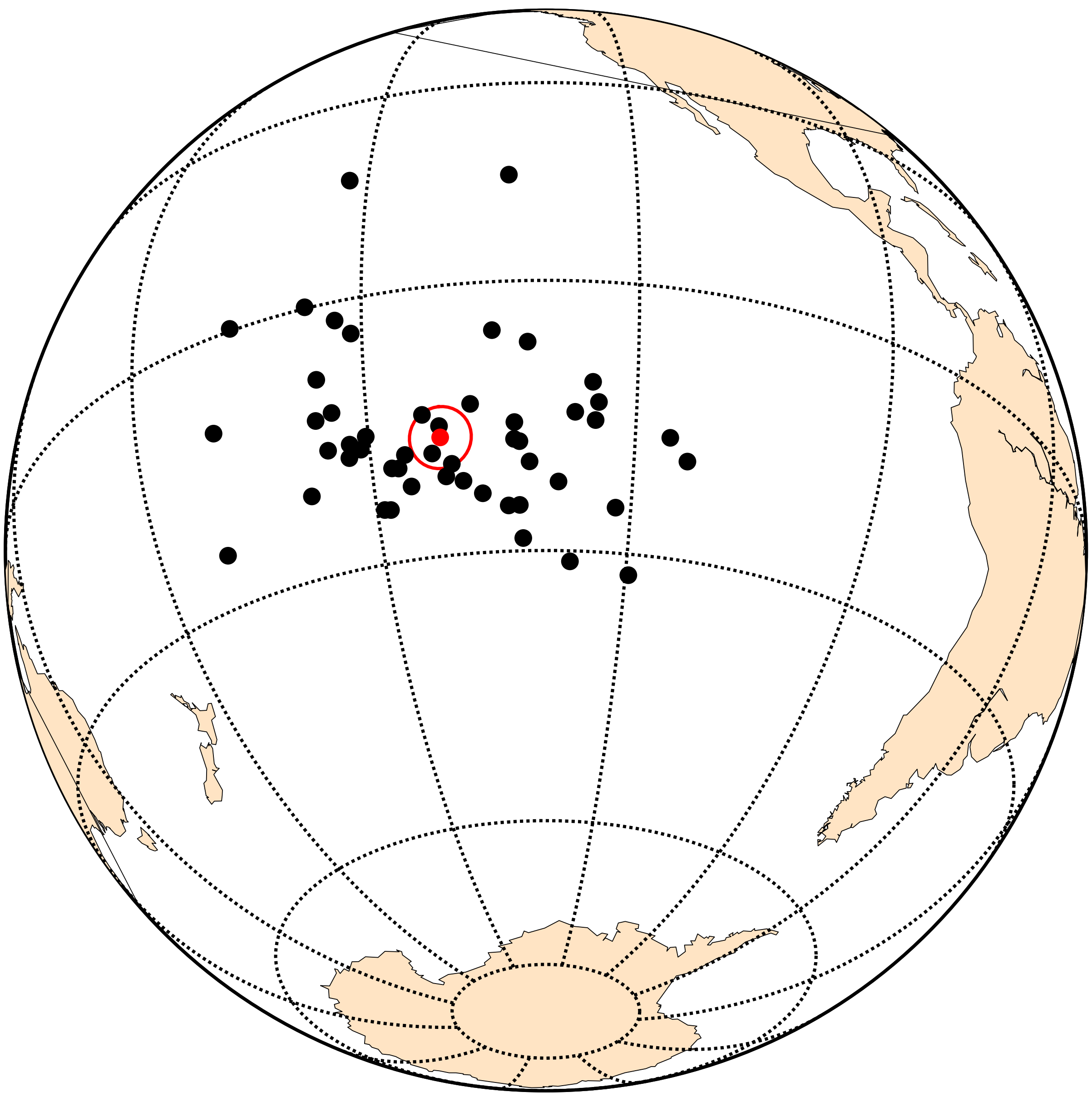
```
alpha95              3.350323
csd                  13.29308
dec                  202.0625
inc                  51.04129
k                    37.12949
n                          50
r                    48.68029
site_lat                37.97
site_lon              -122.27
paleolatitude        31.73096
vgp_lat             -17.32687
vgp_lon         218.178192667
vgp_lat_rev          17.32687
vgp_lon_rev          38.17819
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: 202.1  Plat: 51.0
Number of directions in mean (n): 50.0
Angular radius of 95% confidence (A_95): 3.4
Precision parameter (k) estimate: 37.1
```



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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/Laurenti
a_Pole_Compilation.csv')
Laurentia_Pole_Compilation.head()
```

Out[27]:

	Q	A95	Com	Formation	Lat	Lon	CLat	CLon	RLat	RLon	EULER	Age	G R
0	5	3.9	NaN	Dunkard Formation	-44.1	301.5	-41.5	300.4	-38.0	43.0	(63.2/_ 13.9/79.9)	300	3(
1	5	2.1	NaN	Laborcita Formation	-42.1	312.1	-43.0	313.4	-32.7	52.9	(63.2/_ 13.9/79.9)	301	1:
2	5	3.4	#	Wescogame Formation	-44.1	303.9	-46.3	306.8	-38.2	51.4	(63.2/_ 13.9/79.9)	301	1:
3	6	3.1	I	Glenshaw Formation	-28.6	299.9	-28.6	299.9	-28.6	32.4	(63.2/_ 13.9/79.9)	303	K
4	5	1.8	NaN	Lower Casper Formation	-45.7	308.6	-50.5	314.6	-37.6	59.8	(63.2/_ 13.9/79.9)	303	1

In [28]:

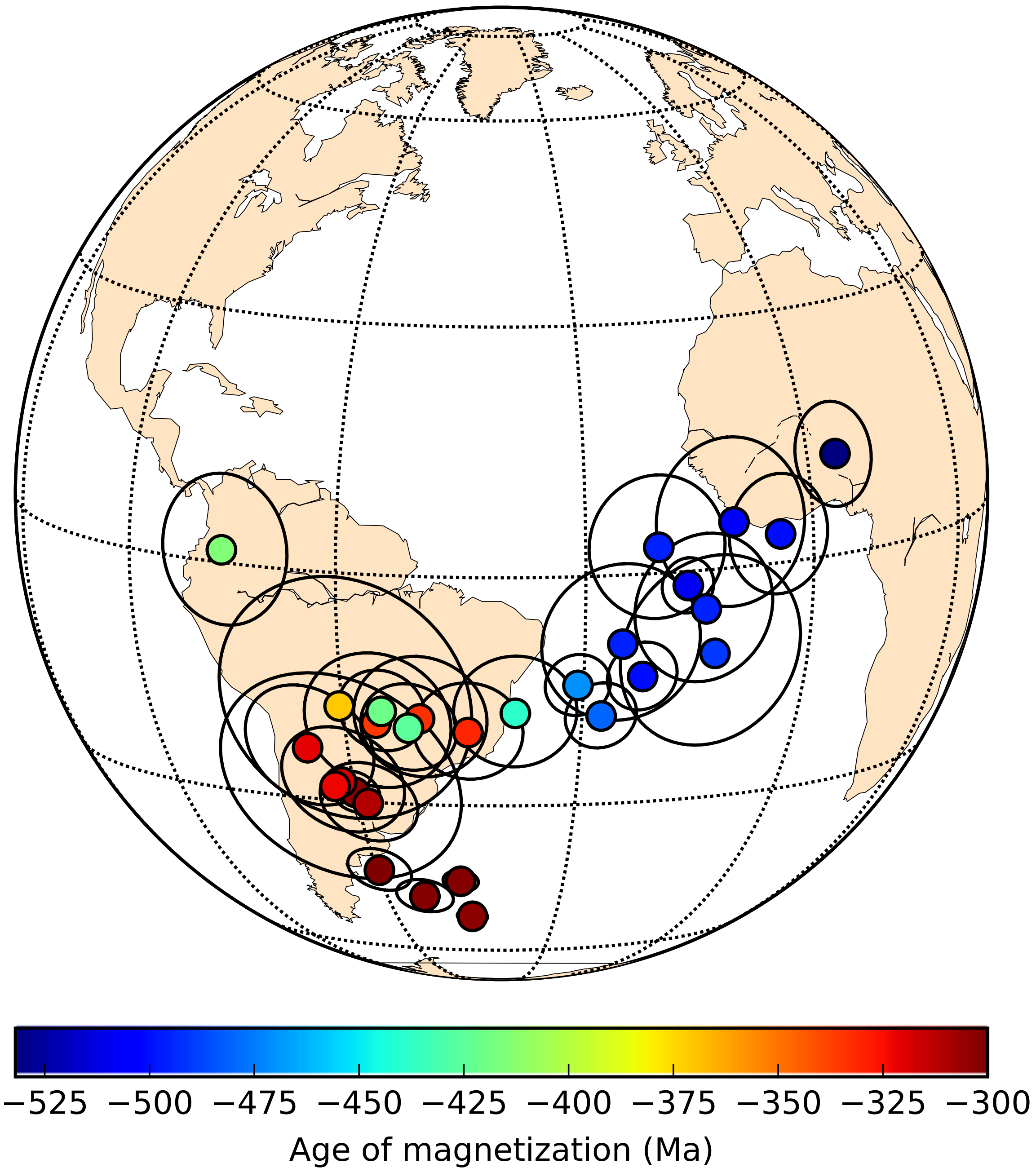
```
# initiate the figure as in the plot_pole example
plt.figure(figsize=(6, 6))
pmap = Basemap(projection='ortho',lat_0=10,lon_0=320,
               resolution='c',area_thresh=50000)
pmap.drawcoastlines(linewidth=0.25)
pmap.fillcontinents(color='bisque',lake_color='white',zorder=1)
pmap.drawmapboundary(fill_color='white')
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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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()
```

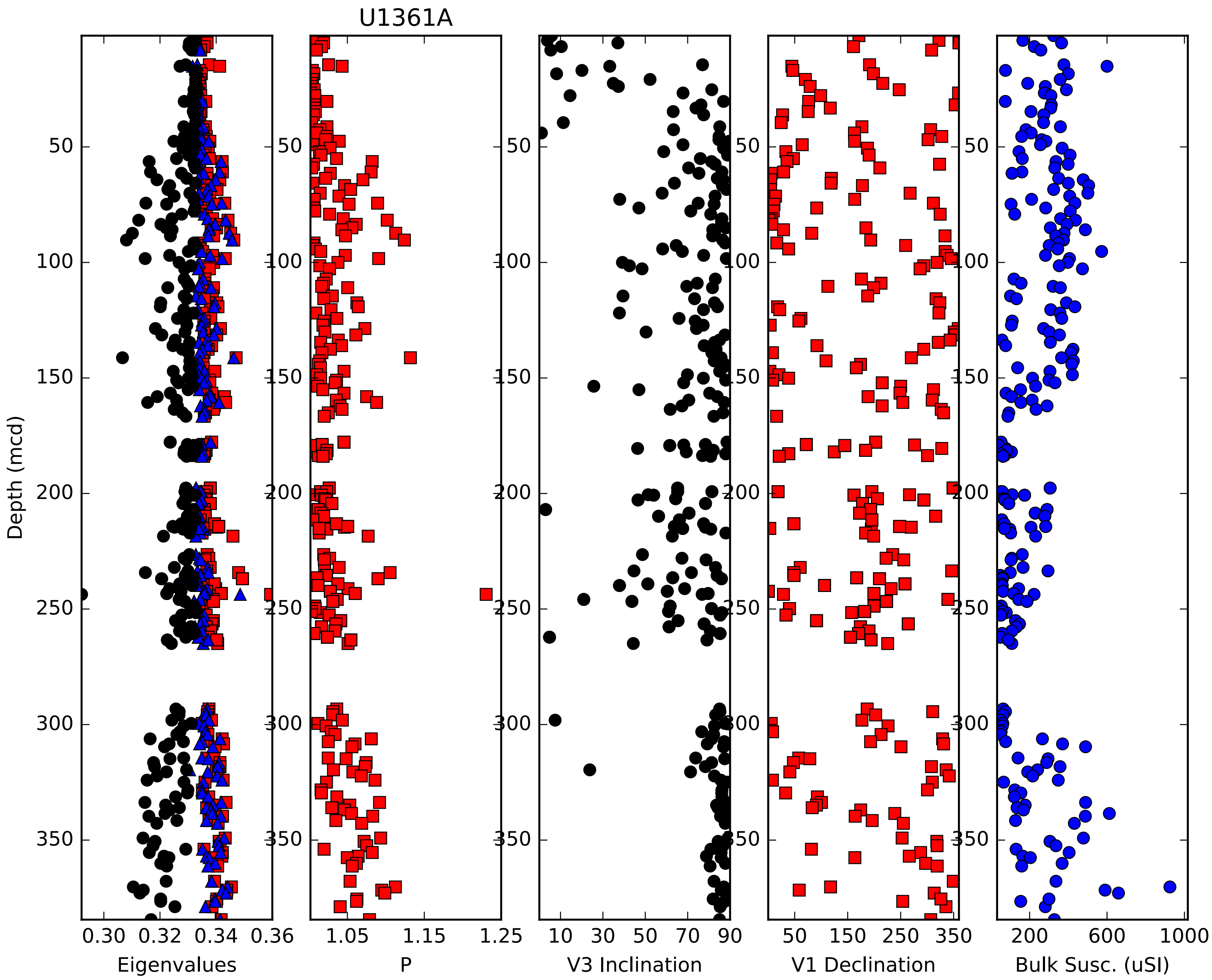
Out[29]:

	anisotropy_n	anisotropy_s1	anisotropy_s2	anisotropy_s3	anisotropy_s4	anisotrop
0	192	0.332294	0.332862	0.334844	-0.000048	0.000027
1	192	0.333086	0.332999	0.333916	-0.000262	-0.00032
2	192	0.333750	0.332208	0.334041	-0.000699	0.000663
3	192	0.330565	0.333928	0.335507	0.000603	0.000212
4	192	0.332747	0.332939	0.334314	-0.001516	-0.00031

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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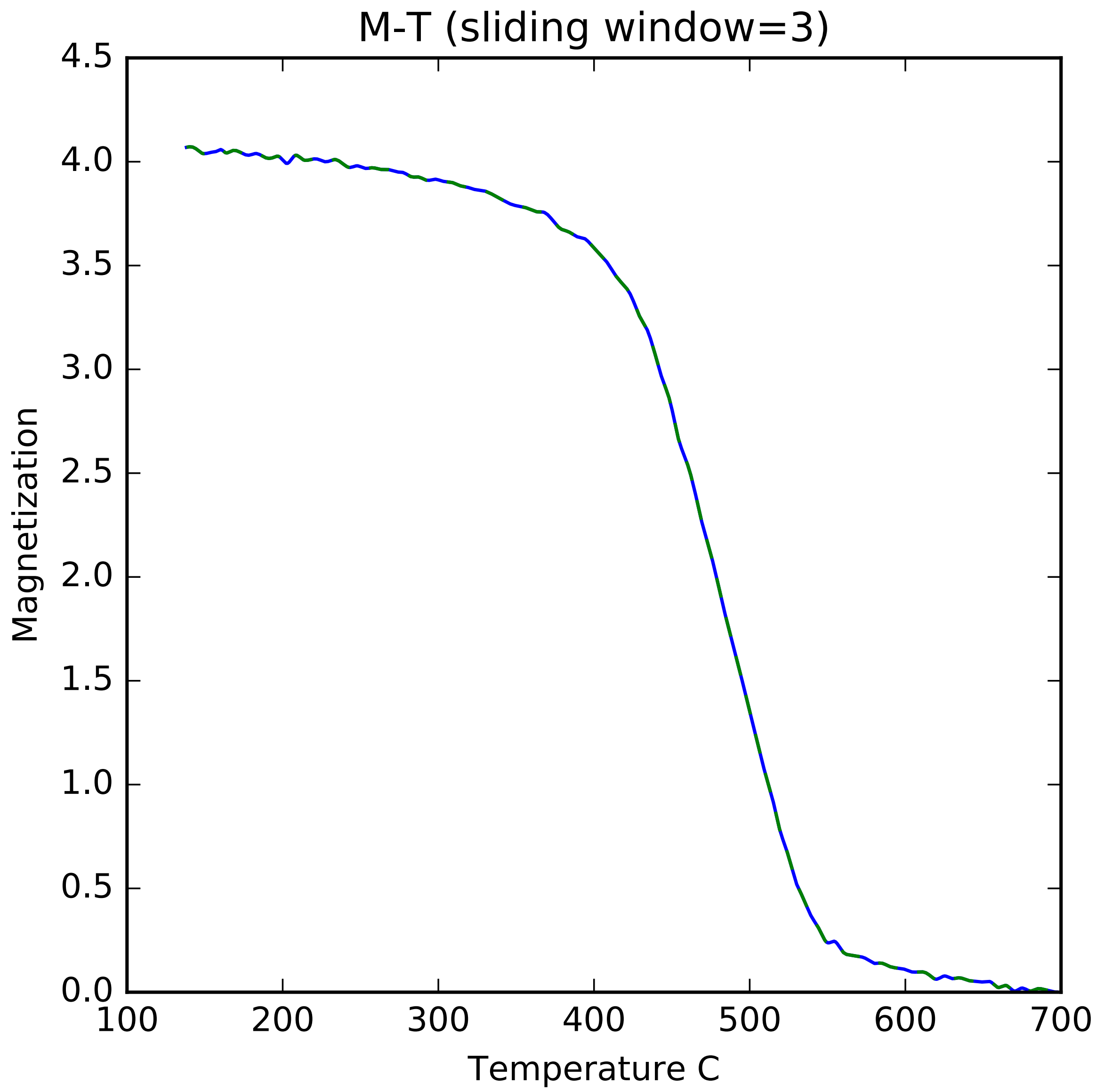
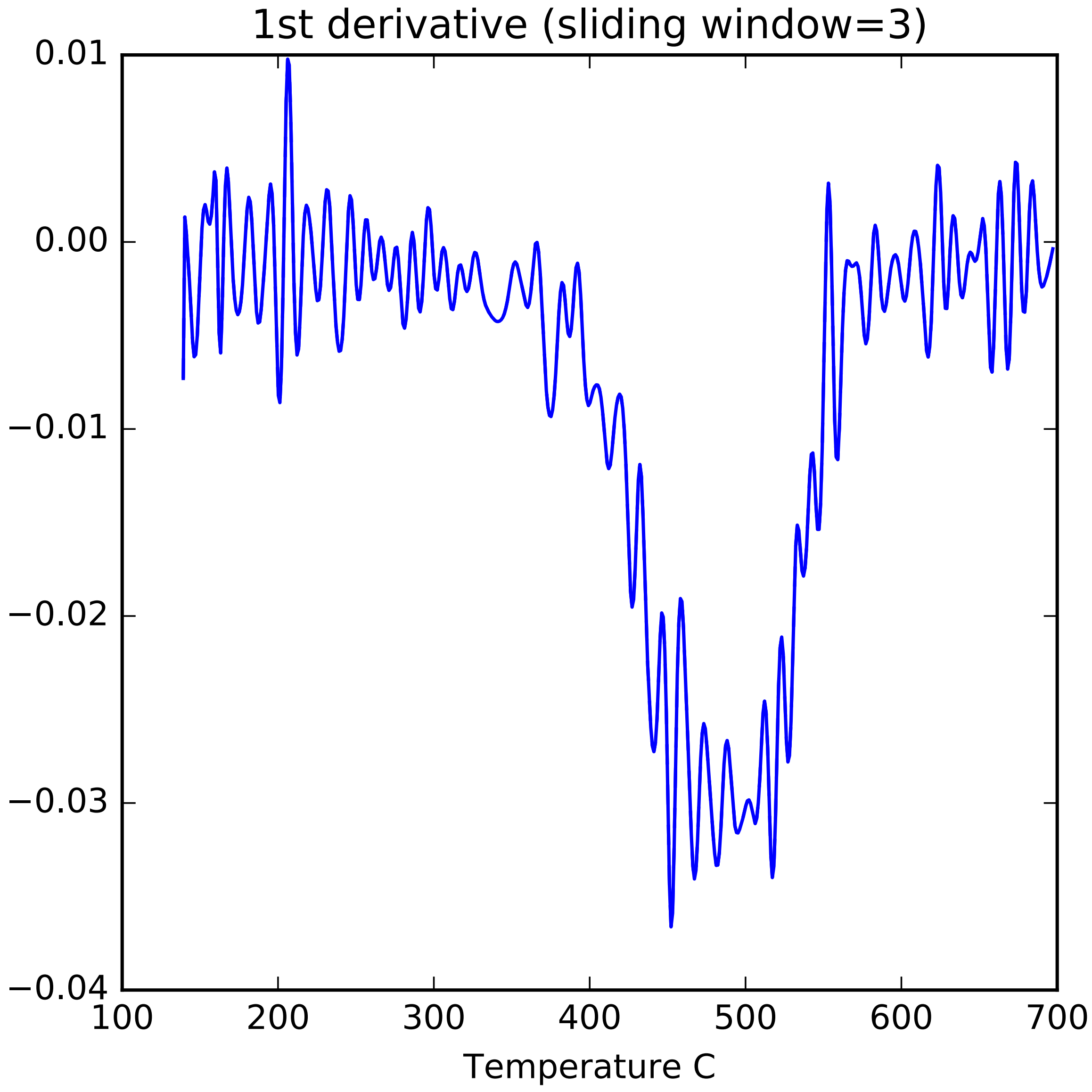
Curie temperature data

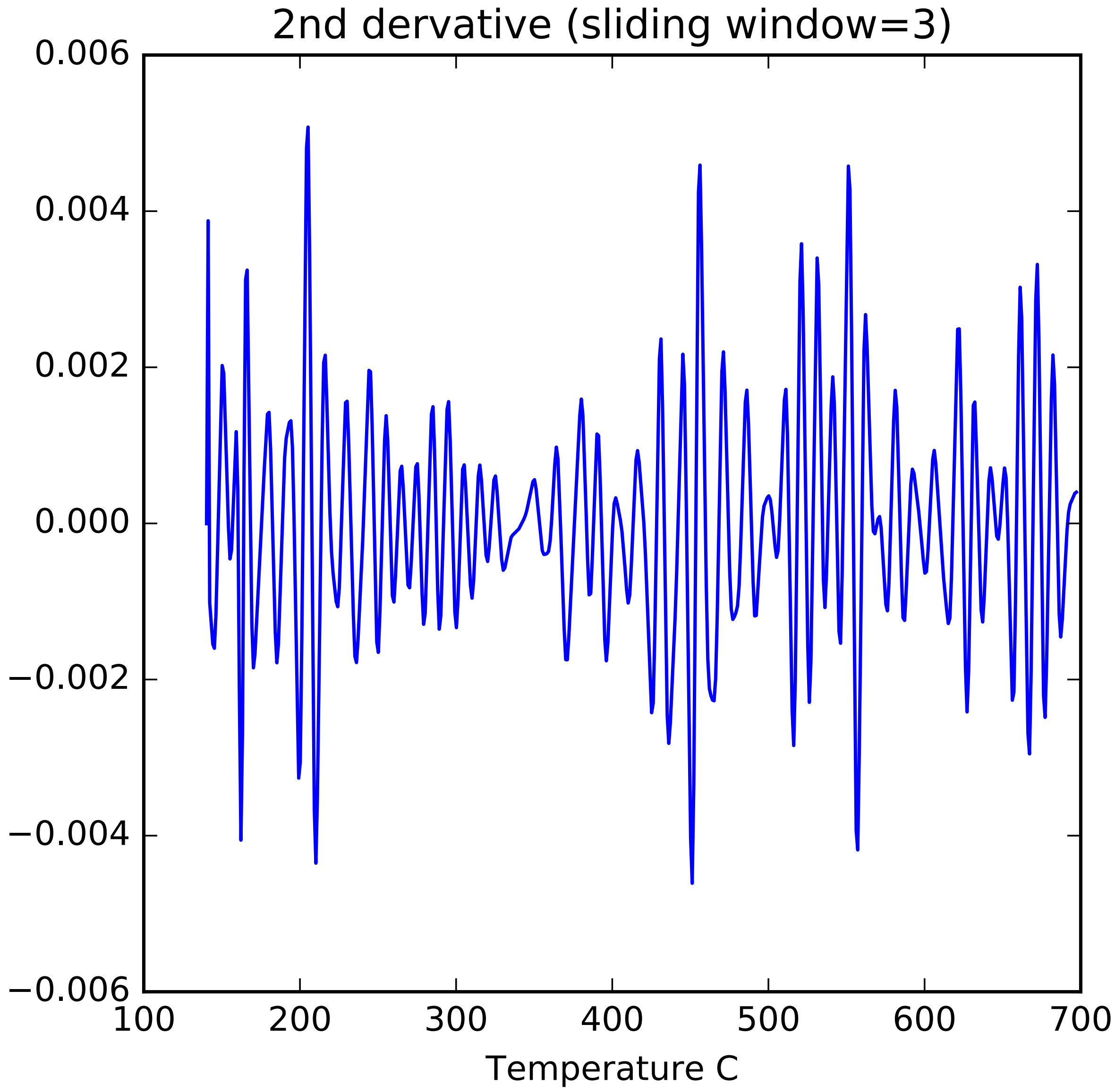
In [31]:

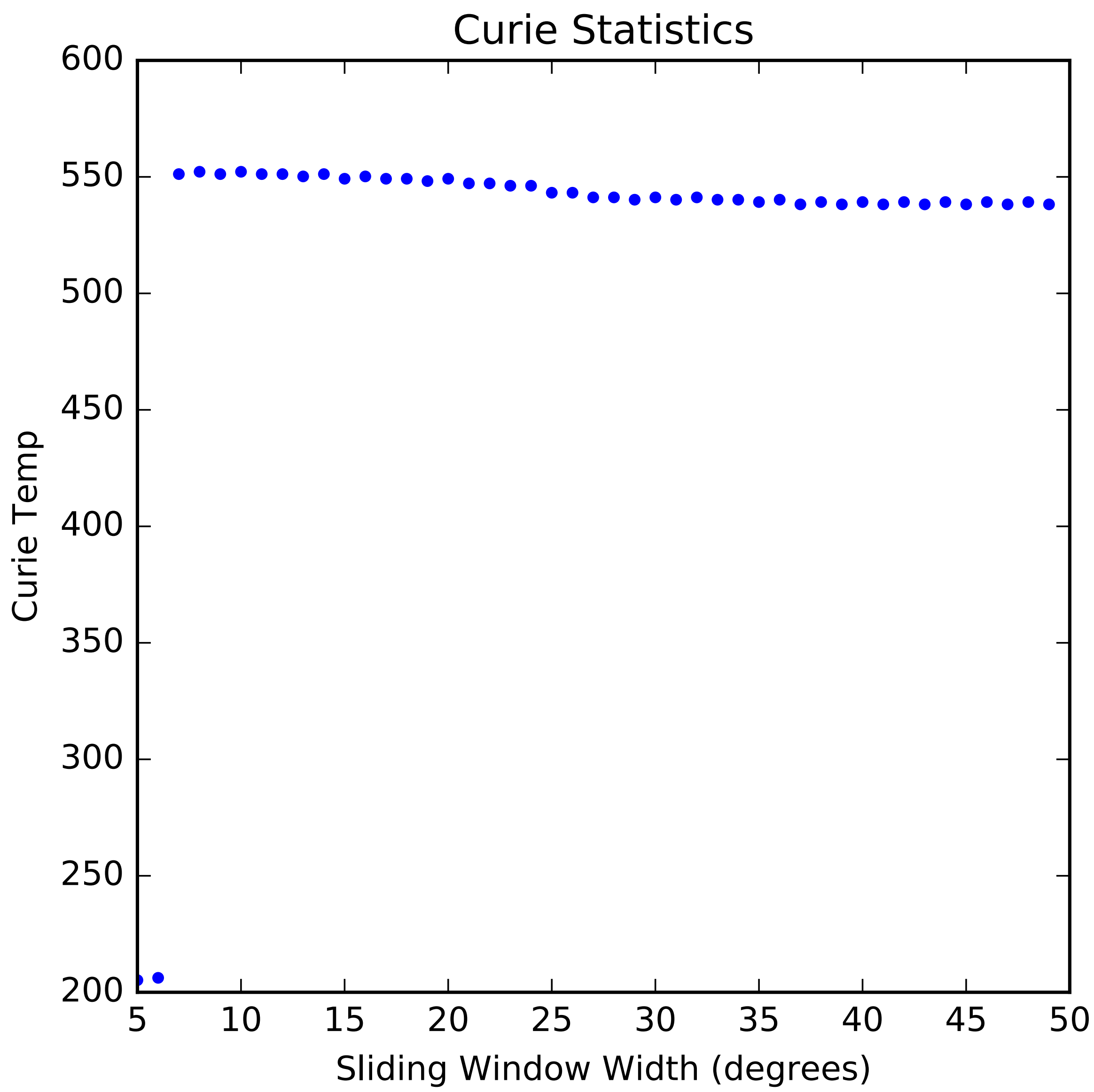
```
ipmag.curie(path_to_file='./Additional_Data/curie/',
            file_name='curie_example.dat', save=True,
            save_folder='Additional_Notebook_Output/curie/')
```

second deriative maximum is at T=205




1st derivative (sliding window=3)
0.01
0.00
−0.01
−0.02
−0.03
−0.04
100
200
300
400
500
600
700
Temperature C


2nd dervative (sliding window=3)
0.006
0.004
0.002
0.000
−0.002
−0.004
−0.006
100
200
300
400
500
600
700
Temperature C



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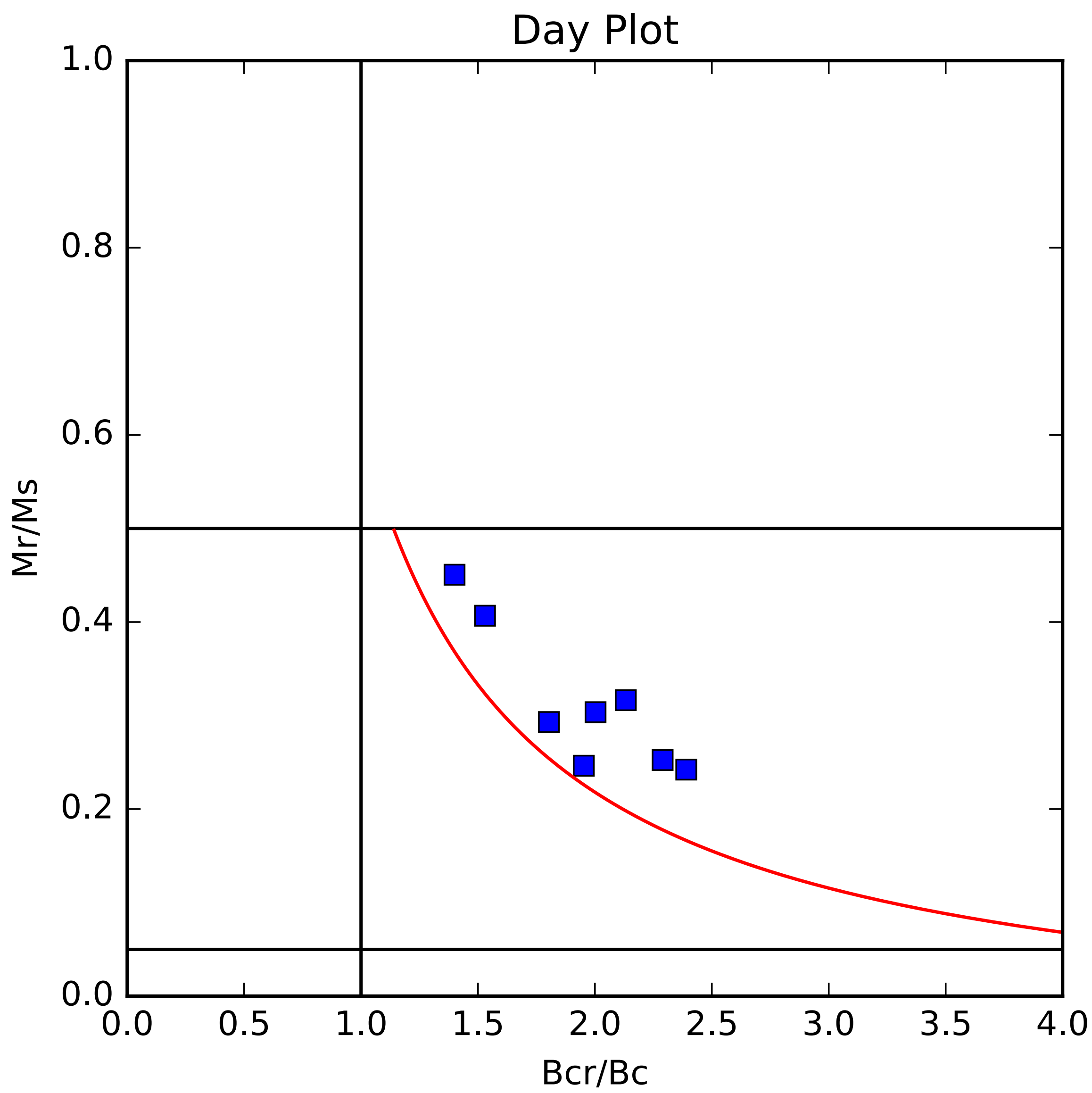
Day plots

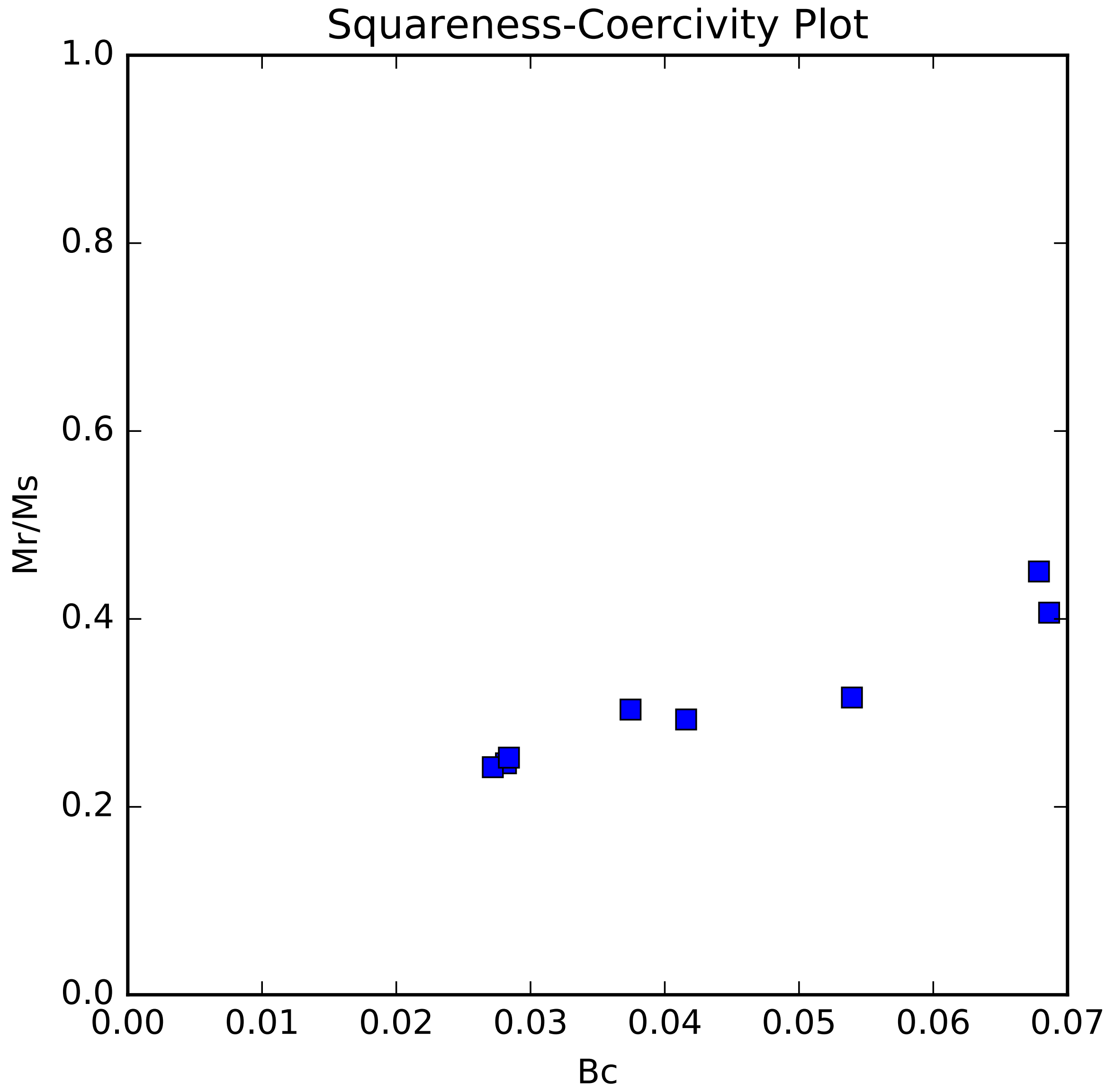
Here we demonstrate the function **ipmag.dayplot**, which creates Day plots, squareness/coercivity and squareness/coercivity of remanence diagrams using hysteresis data.

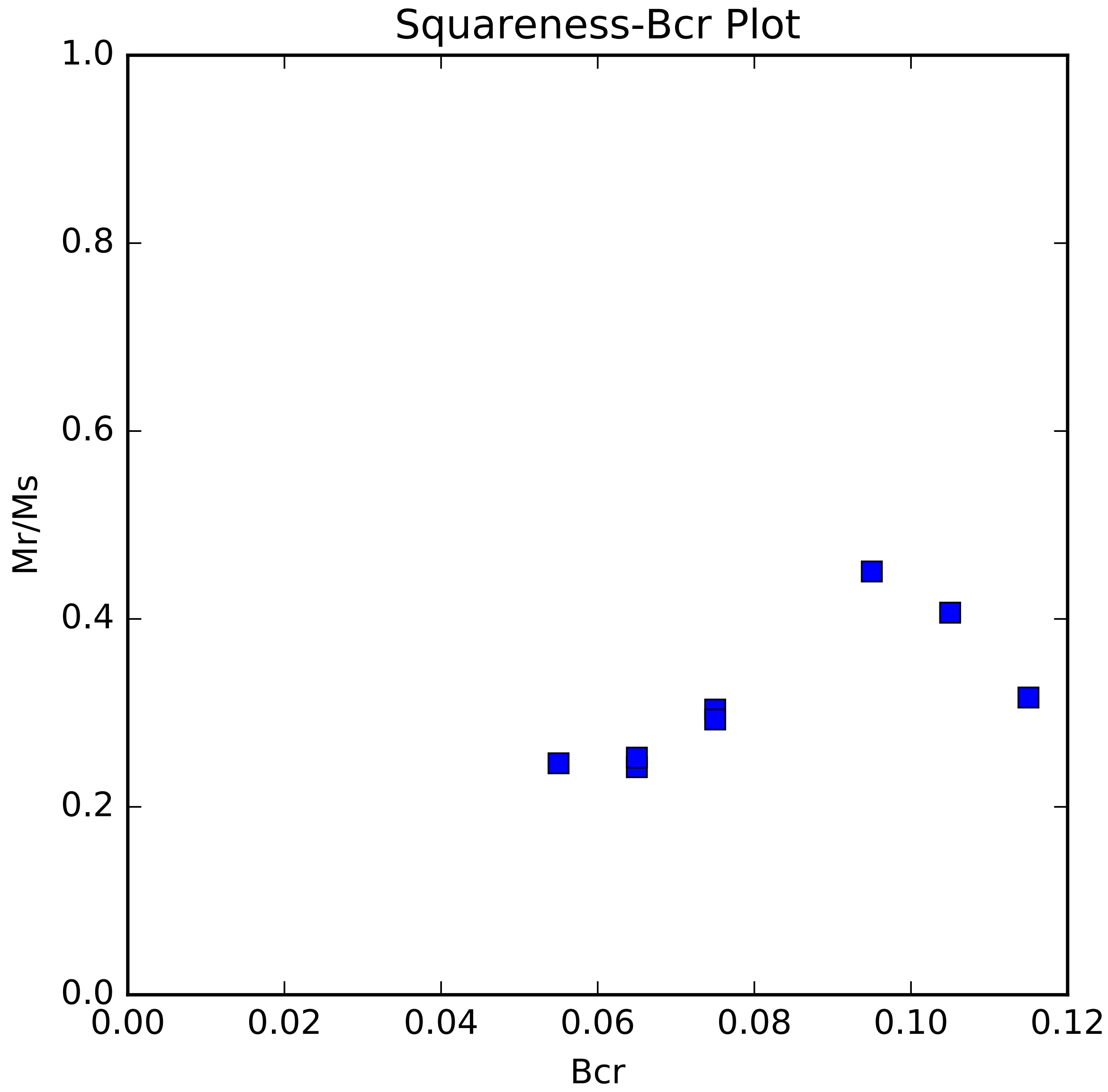
In [32]:

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



Day Plot
1.0
0.8
0.6
0.4
0.2
0.0
Mr/Ms
0.0
0.5
1.0
1.5
2.0
2.5
3.0
3.5
4.0
Bcr/Bc


Squareness-Coercivity Plot
Mr/Ms
Bc
1.0
0.8
0.6
0.4
0.2
0.0
0.00
0.01
0.02
0.03
0.04
0.05
0.06
0.07




```
<matplotlib.figure.Figure at 0x10aca2e90>
```

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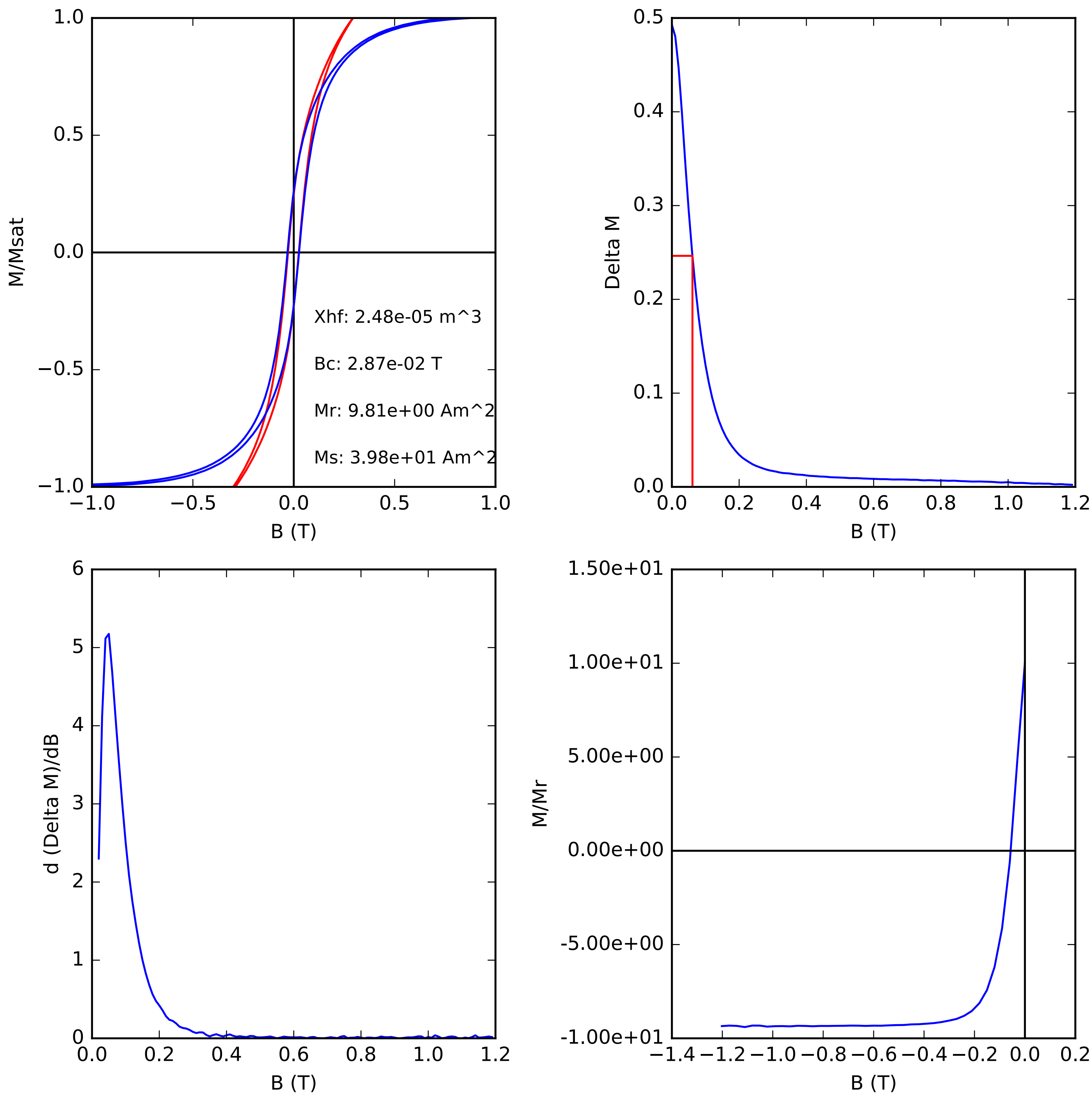
Hysteresis Loops

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

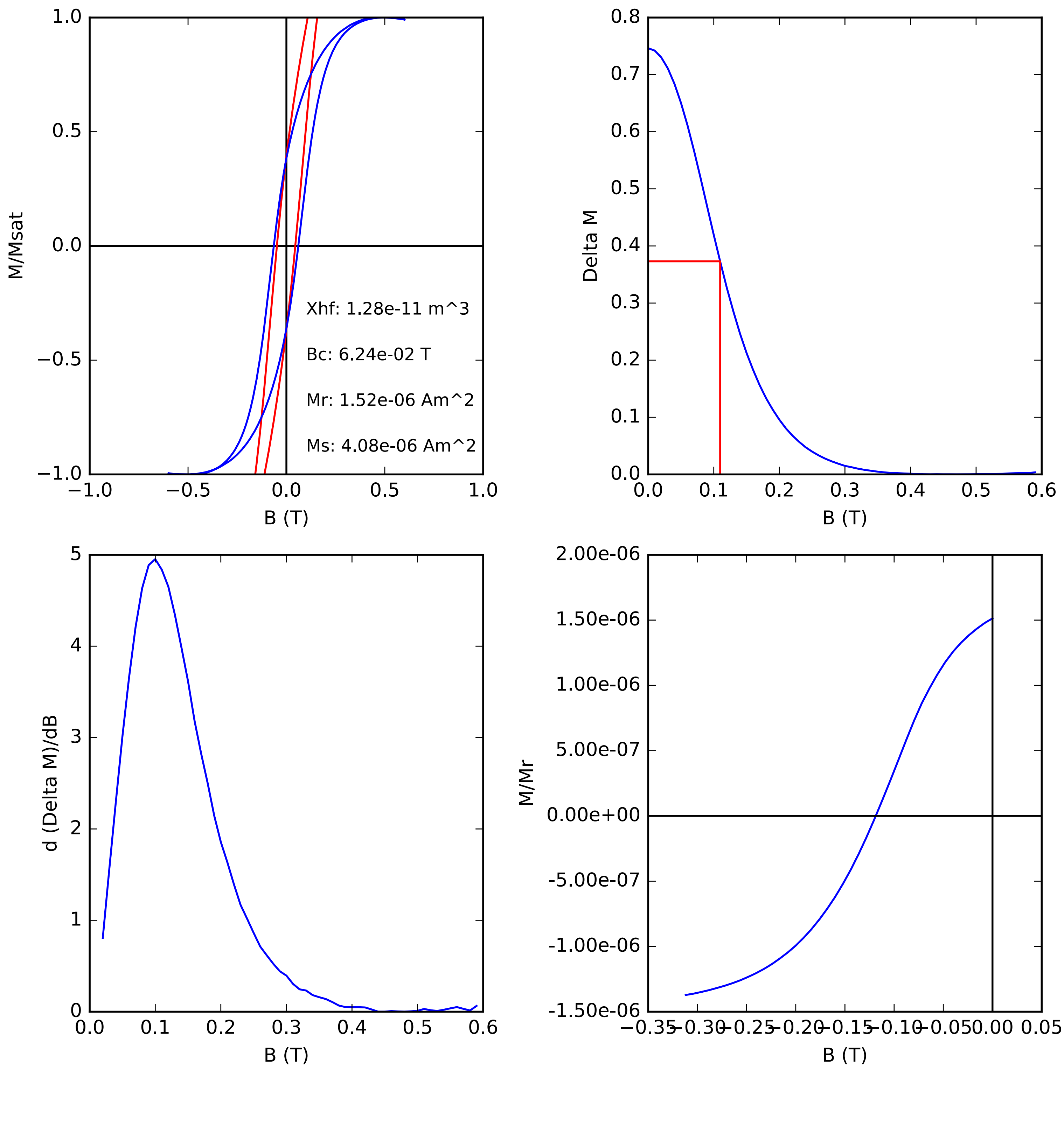
In [33]:

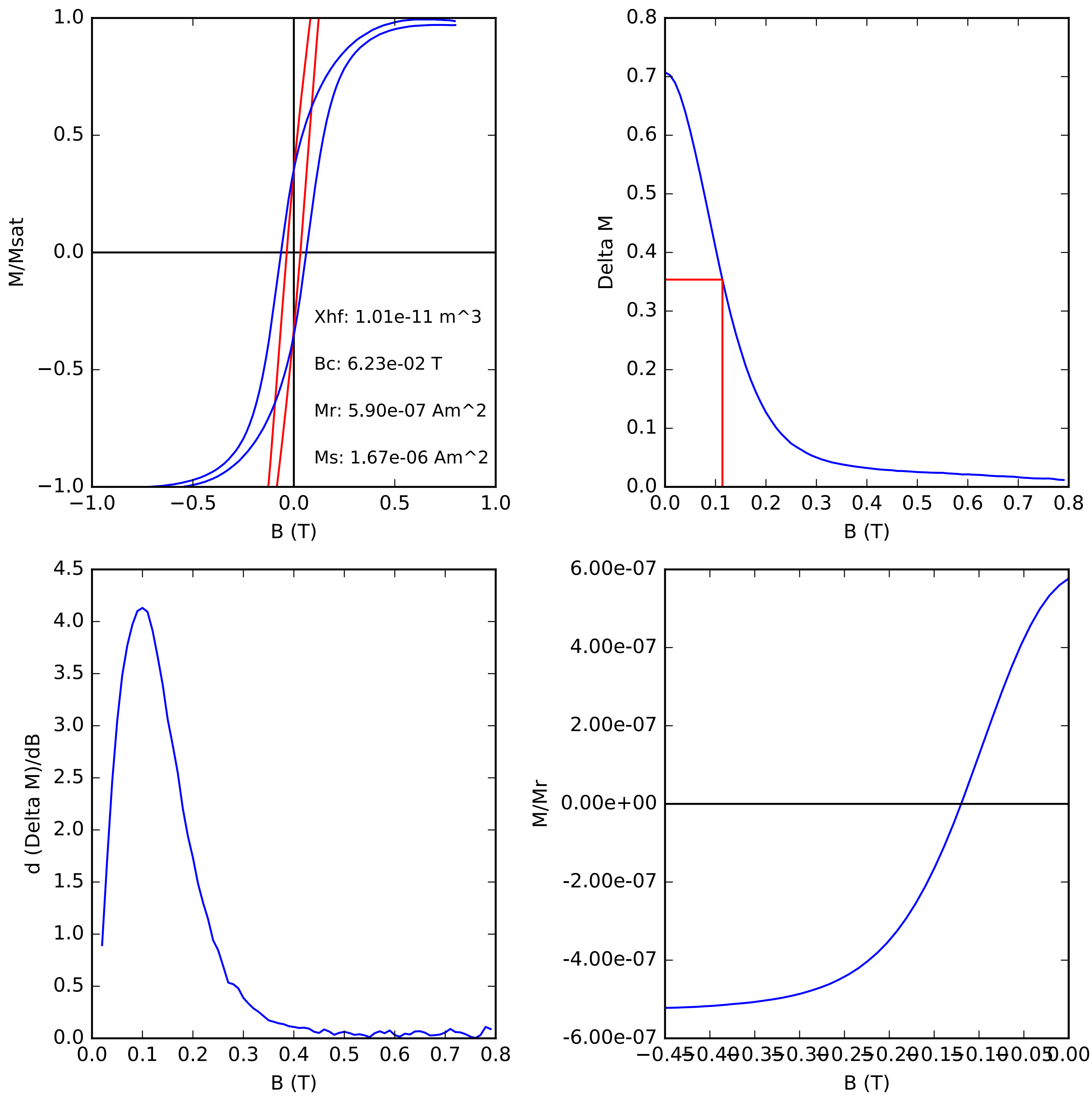
```
ipmag.hysteresis_magic(path_to_file='./Additional_Data/hysteresis_magic/',
                 hyst_file='hysteresis_magic_example.dat', save=True,
                save_folder='./Additional_Notebook_Output/hysteresis')
```

```
IS06a-1 1 out of  8
```

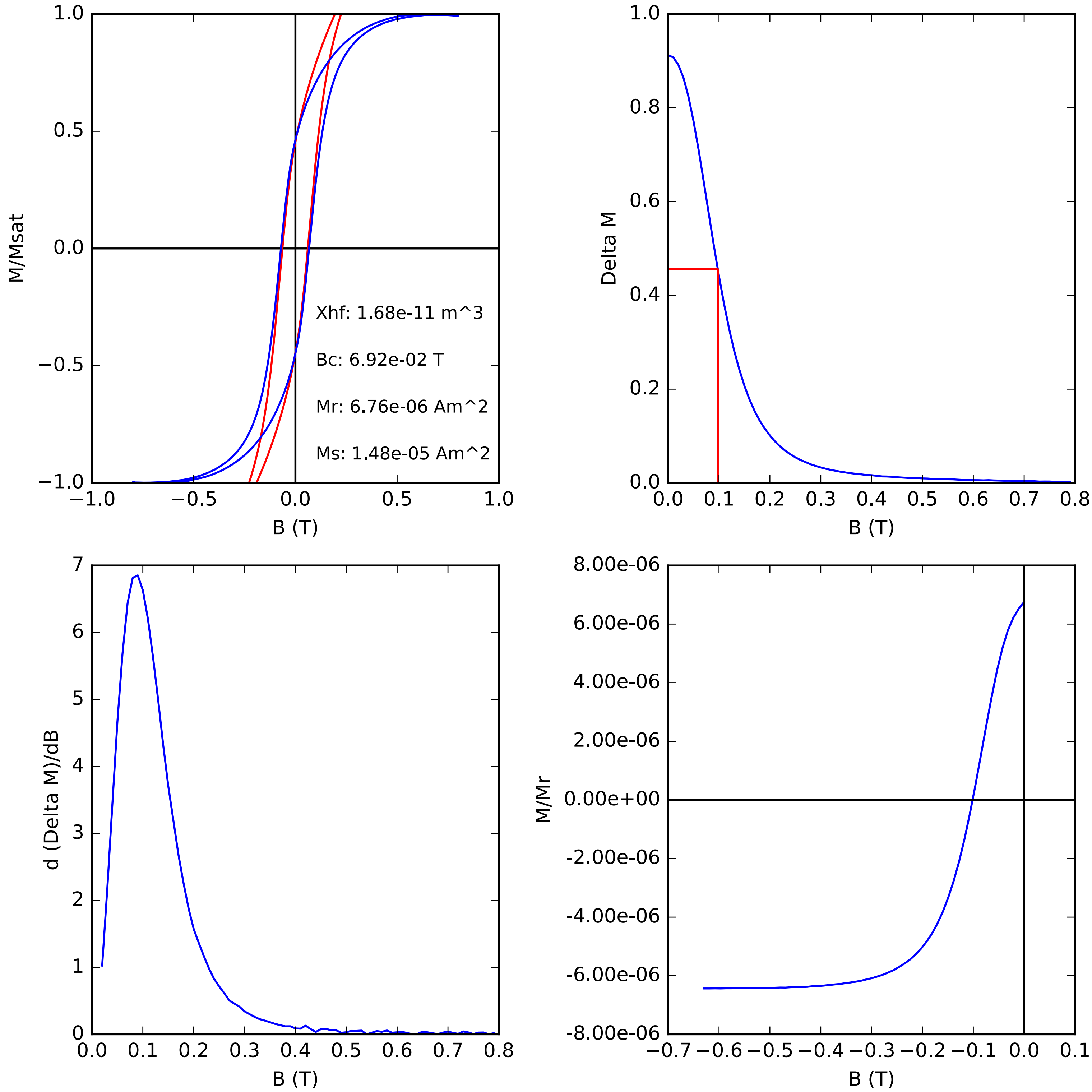


IS06a-2 2 out of 8

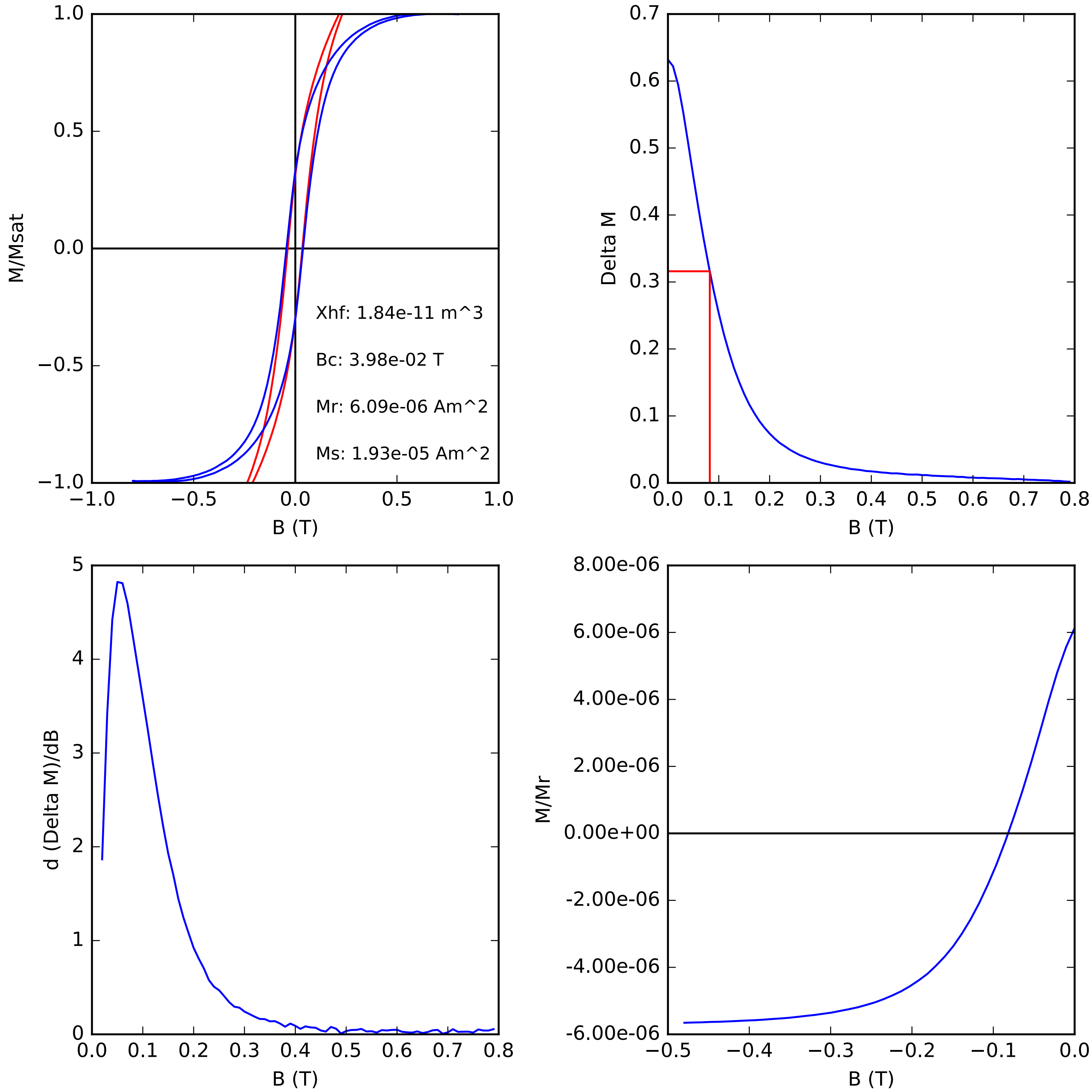

Xhf: 1.28e-11 m^3
Bc: 6.24e-02 T
Mr: 1.52e-06 Am^2
Ms: 4.08e-06 Am^2
M/Msat
B (T)
Delta M
B (T)
d (Delta M)/dB
B (T)
M/Mr
B (T)



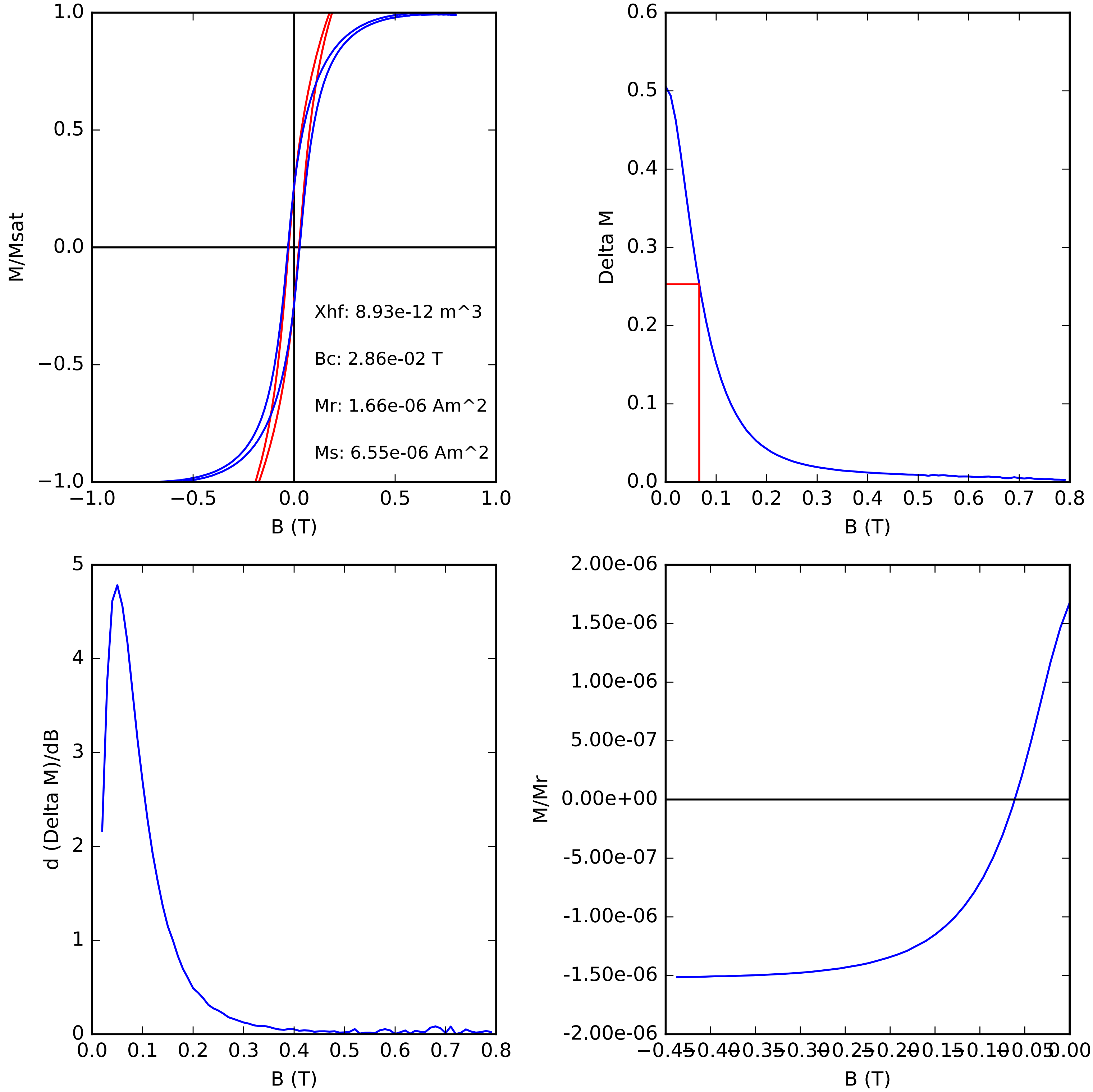
IS06a-4 4 out of 8



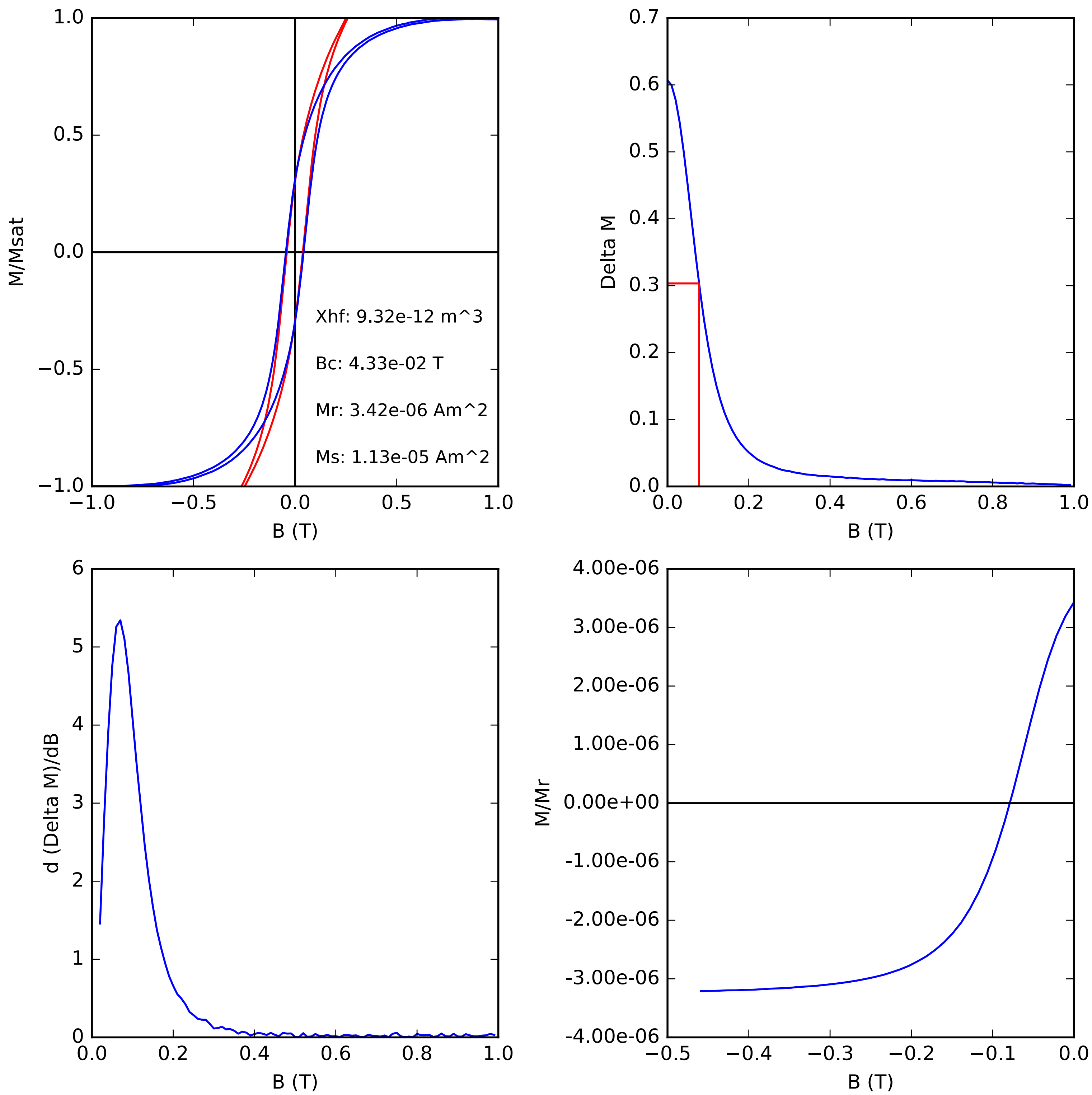
M/Msat
B (T)
Xhf: 1.68e-11 m^3
Bc: 6.92e-02 T
Mr: 6.76e-06 Am^2
Ms: 1.48e-05 Am^2
Delta M
B (T)
d (Delta M)/dB
B (T)
M/Mr
B (T)



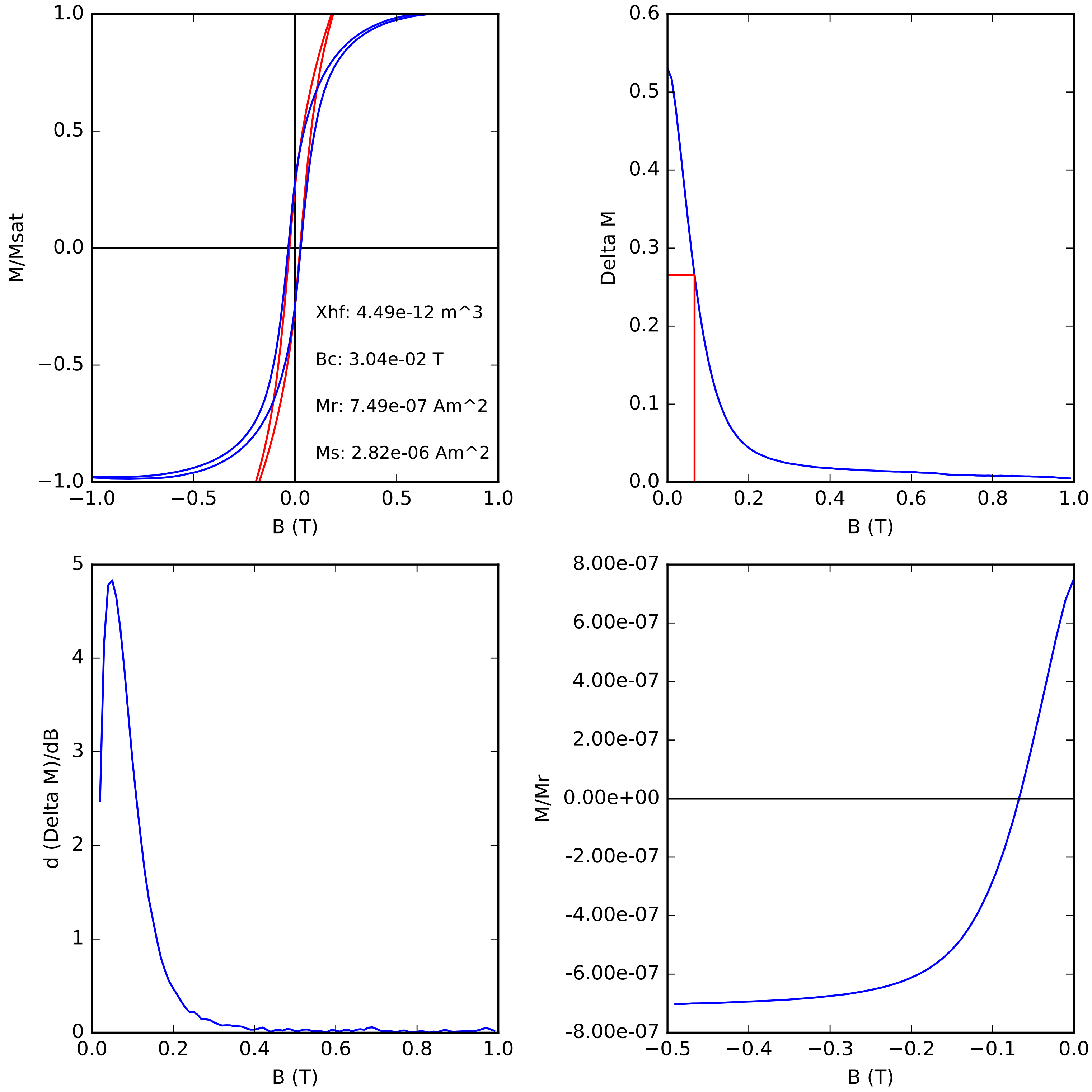
IS06a-6 6 out of 8



IS06a-8 7 out of 8



IS06a-9 8 out of 8



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Demagnetization Curves

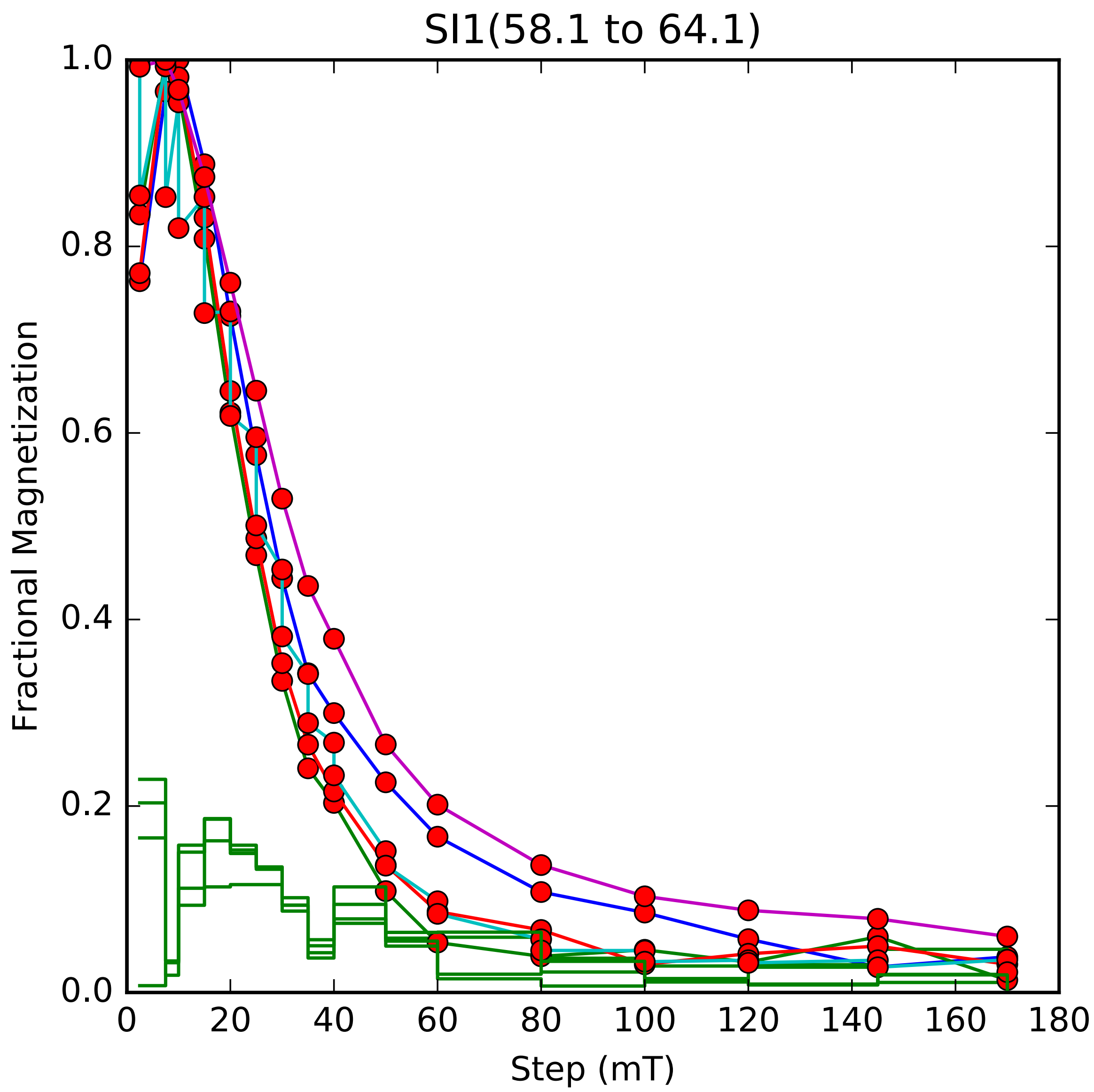
The function **ipmag.demag_magic** filters and plots demagnetization data. These data will be read and combined by expedition, location, site or sample according to the *plot_by* keyword argument. Alternatively, you can choose to plot each specimen measurement individually. By default, all plots generated 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_me
asurements.txt
SI1(58.1 to 64.1) plotting by:  er_site_name
```



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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]:

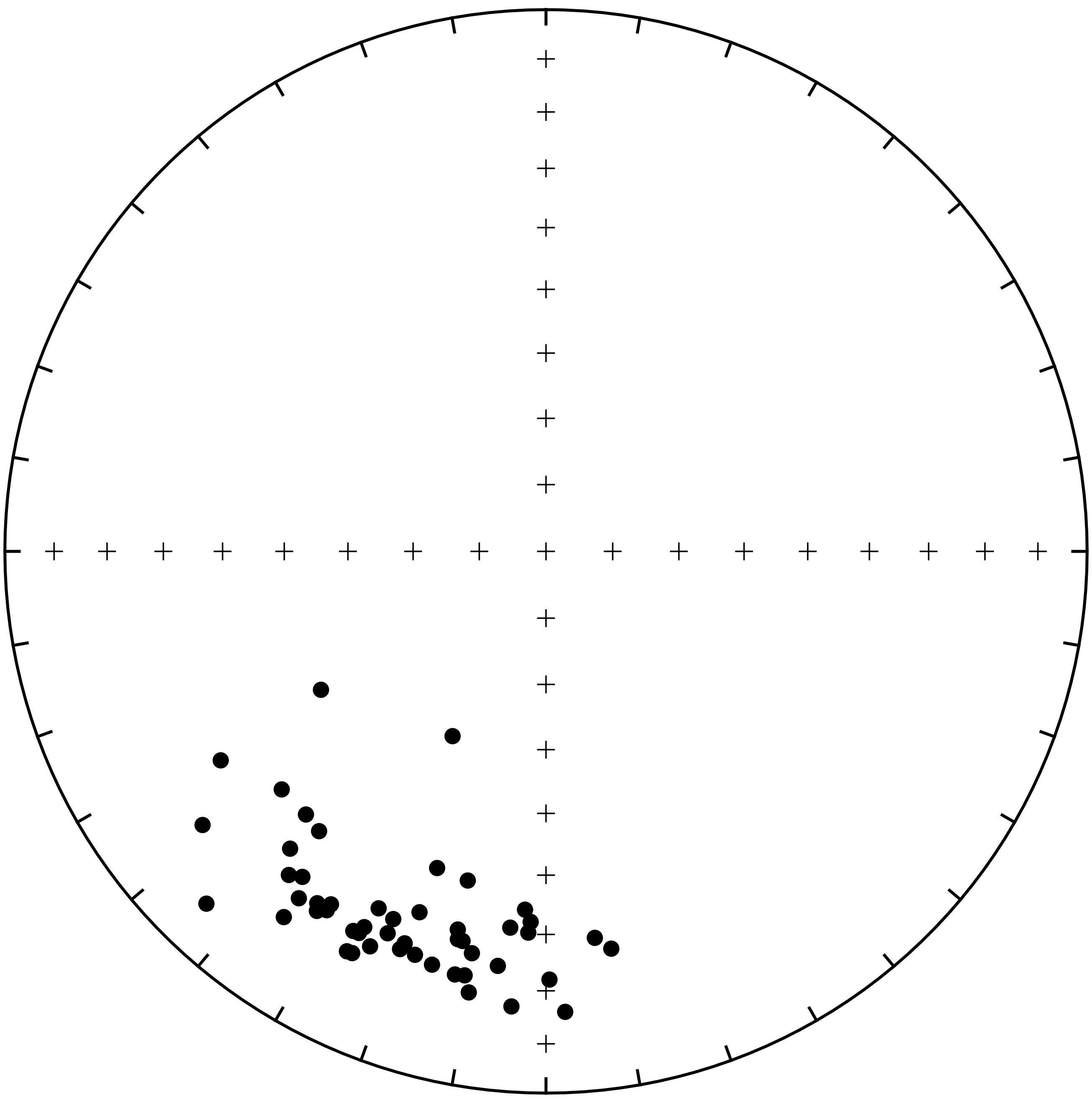
```
from ipywidgets import interact
```

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)
```

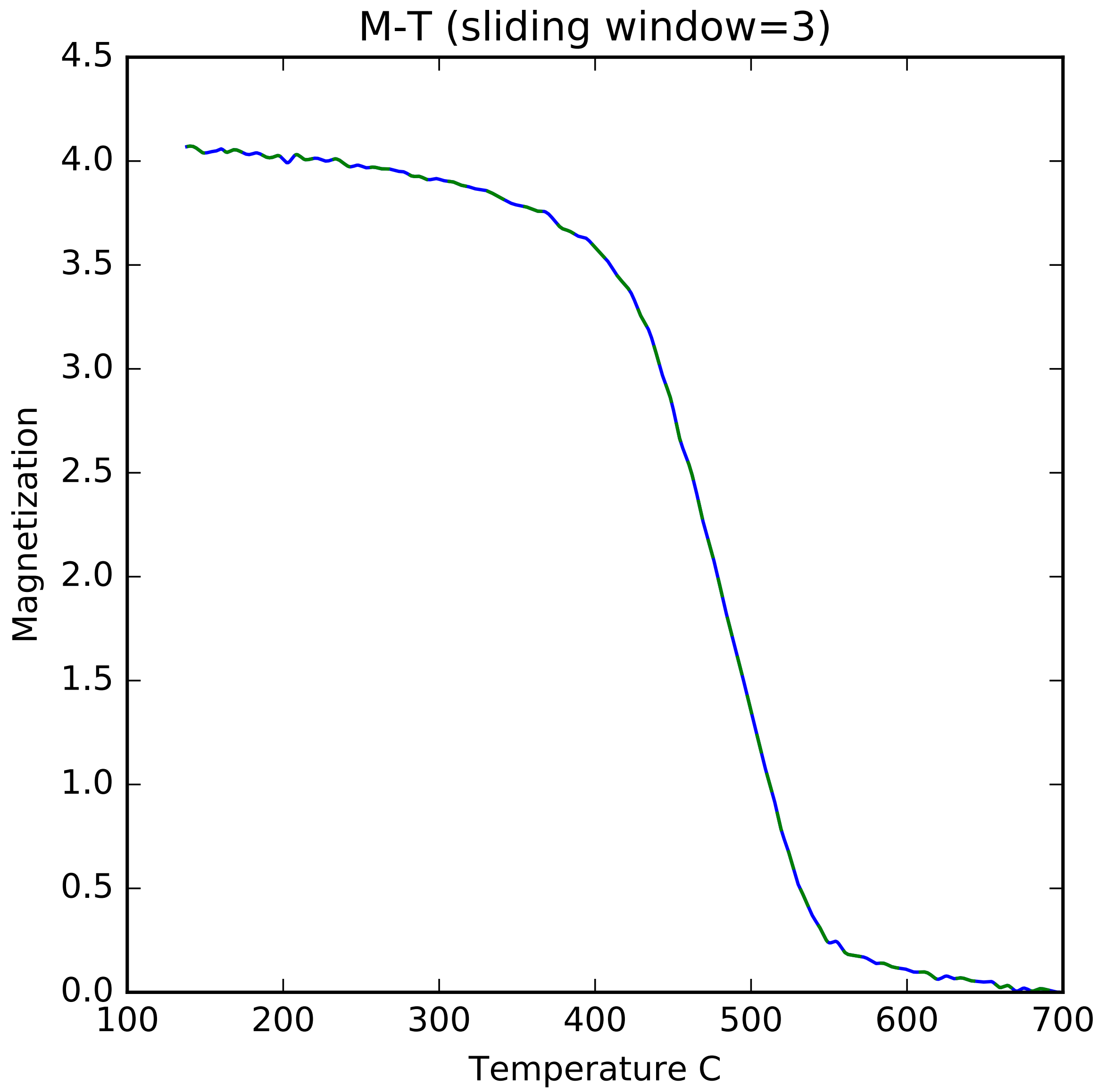


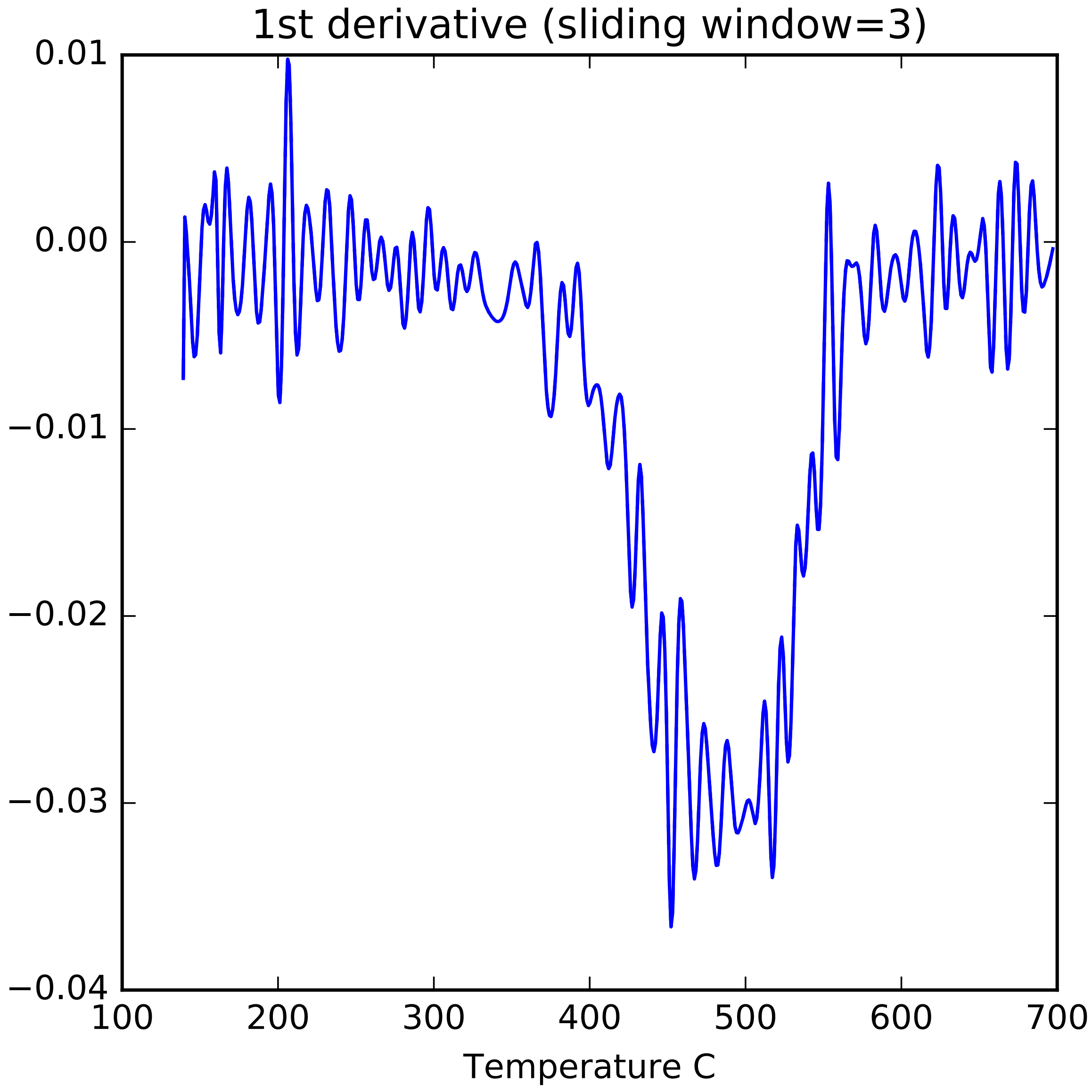
interact can also be used as a regular function call -- the name of the interactive function is passed as the first argument, followed by the adjustable keyword arguments. Below, we demonstrate passing the *curie* function's parameters to **interact**.

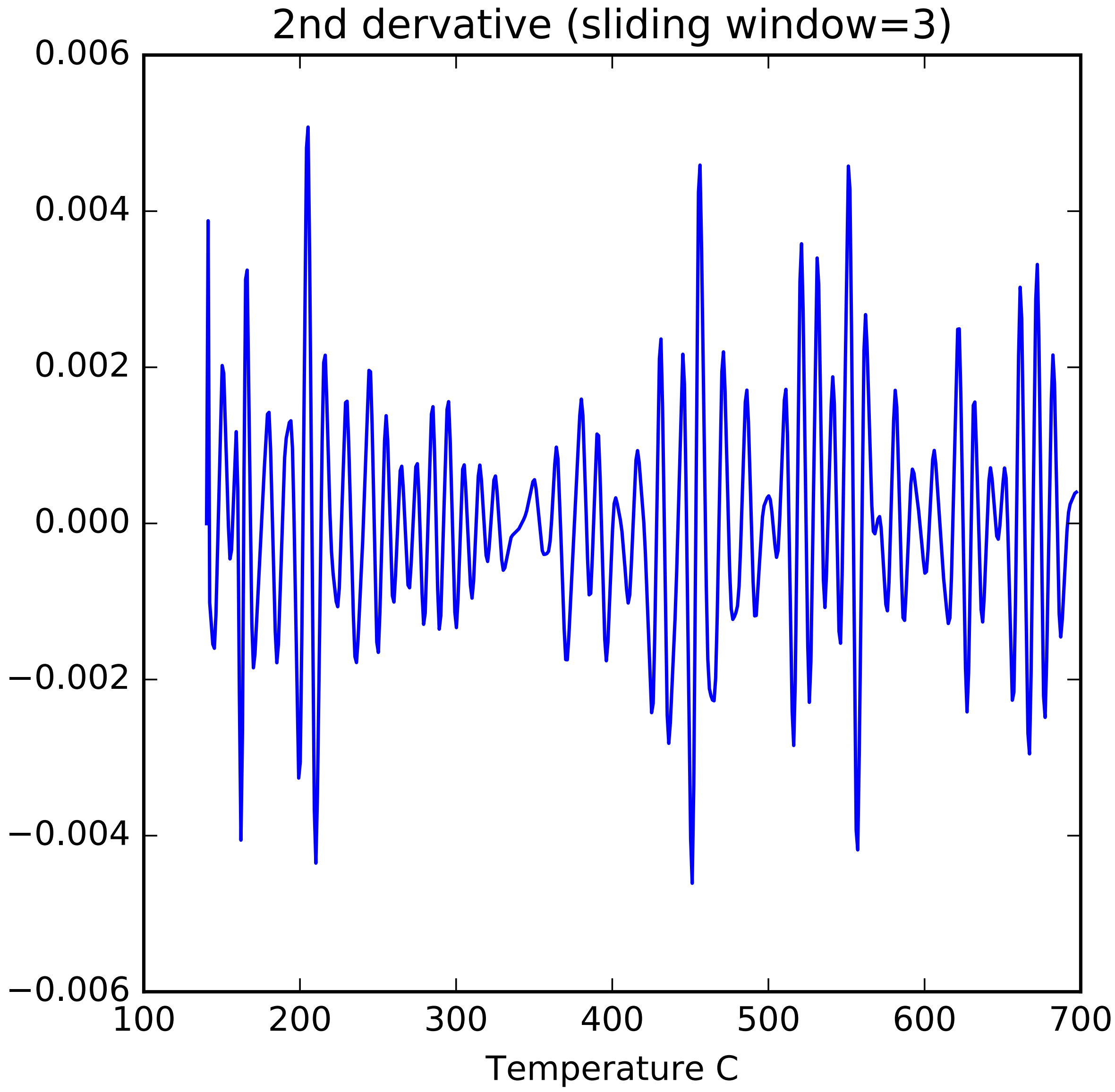
In [37]:

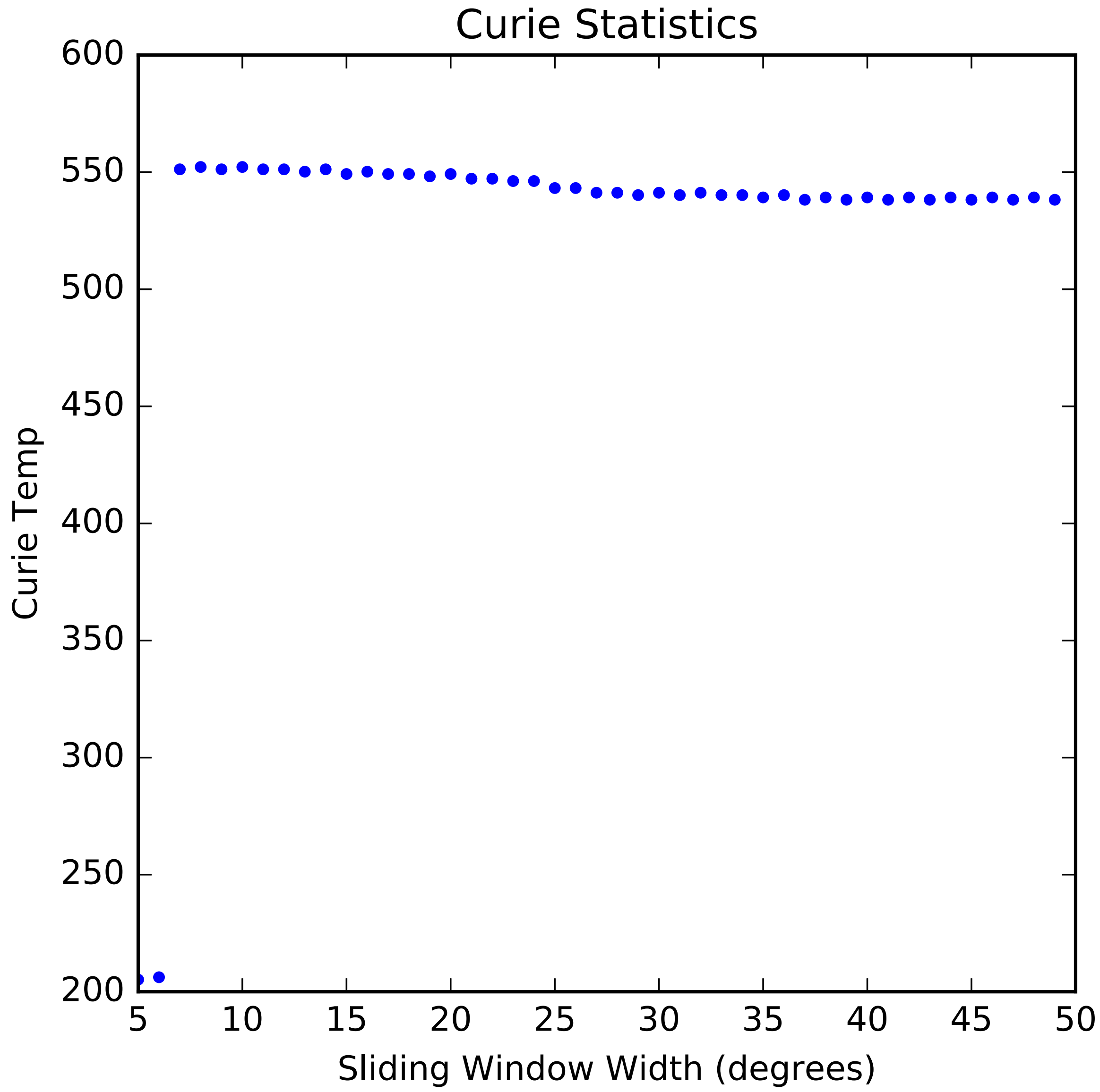
```
interact(ipmag.curie, path_to_file='./Additional_Data/curie/',file_name='curie_e
xample.dat',window_length = (1,60));
```

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




1st derivative (sliding window=3)
0.01
0.00
−0.01
−0.02
−0.03
−0.04
100
200
300
400
500
600
700
Temperature C


2nd dervative (sliding window=3)
0.006
0.004
0.002
0.000
−0.002
−0.004
−0.006
100
200
300
400
500
600
700
Temperature C


Curie Statistics
Curie Temp
Sliding Window Width (degrees)
600
550
500
450
400
350
300
250
200
5
10
15
20
25
30
35
40
45
50

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```
In [ ]:
```