# Pantheon Parameter guide

## Contents

NOTE: Please set the paths at the beginning and ending of the code!p	2
Magnet	
CSA and Aysm	
Spinning_Speed	
Number_of_points	
Crystal_file	
Gamma_Angle	
DCcor	2
Weighting_type	2
Weighting_value	2
Searchedppm	2
Additional_Sim_Scaling	3
Spectral_Width	3
SimType	3
MathType3DWeighting	3
Experiment_Type	3
PlotMode	3
ExpFile	3
ExcelProtonData	3
Artifact_Removal Artifact_mid_point Artifact_Removal_width	3
Bonus_RFinhomogeneity	3
Mirror_f1	4
Exp_2D_Hz_width	4
Processing R16_3_2	5
Example: Performing a 2D simulation of SC212 at 20ppm for CSA and 0.5 for asymmetry and comparing it the an experimetnal proton at 12ppm	11
Removing the artifact for help calculating the CSA	12
Example: Comparing a 3D simulation of data to experiment data in an R16_3_2 experiment	15
Easy_Pantheon	17

## NOTE: Please set the paths at the beginning and ending of the code!

#### Magnet

This is the strength of the magnet in MHz, so typically this can be left at 400.

#### CSA and Aysm

These are the values of the CSA and asymmetry that you wish to simulate in a 2D experiment. If you are sim'ing a 3D experiment please see ExcelProtonData

#### Spinning Speed

This is the spinning speed of the experiment.

#### Number\_of\_points

This is the number of points in the f1 fid of the simulated data. Note this does not have to match the experimental and in general can be left at either 96 or 120.

#### Crystal file

This is the size of the crystal file. The larger the value you input the longer the experiment will take. There are more sizes that are listed in the comment in the code. If you enter an incorrect number the code will not run but it will show a list of possible numbers you can run. Tho it only works for the zcw crystal type.

## Gamma\_Angle

This is the number of gamma angles used in the experiment. In general the larger the better quality of the simulation but ~5 is suitable for most purposes.

#### **DCcor**

This is the number of points in the f1 sim fid that are used to lower the centre of the R16\_3\_2 sequence. The lower this value then the higher the middle of the spectrum becomes and vise versa. This should be set to a value so that it matches the experimental. Note this only effects R16\_3\_2 sequences.

## Weighting\_type

This is the type of line broadening that is applied. This should never be changed.

## Weighting\_value

This is the value of the line broadening that is applied. This simply needs to be changed to match the simulated to the experimental data.

## Searchedppm

This is the ppm value that the 2D simulated data will be compared to. (if the experimental cross section does not look correct then plot a 3D version to check the ppm values of the experimental data)

#### Additional Sim Scaling

This simply scales the height of the simulated data in a 2D plot right at the end.

#### Spectral Width

This is the sw of an experiment, note this only needs to be set for R16\_3\_2 experiments.

#### SimType

This can be 2D or 3D and simply decides what type of simulation is run.

#### MathType3DWeighting

This can be 'Gauss' or 'Matt.' This is the technique that is used for line broadening in the f2 axis for the simulated data. 'Gauss' uses a back and fourth fid weighting technique while 'Matt' uses a normal distribution technique. The difference is that 'Gauss' could be see as being slightly more 'NMR'ish but produces a artifact along the baseline. 'Matt' is done though pure maths and does not produce this baseline error.

#### **Experiment Type**

This lets you chose which type of experiment you are going to simulate: R16\_3\_2 SC212 C313

#### PlotMode

This is whether you want to go 'Solo' and plot just the simulated data or to 'Compare' it to experimental data.

## ExpFile

This is the file name of your experimental data. For exporting to a txt file you can type totxt in topspin. This works for SC212 C313. For R16\_3\_2 you need to first process the experimental data in matlab before you can compare it. Please see the 'Processing R16\_3\_2' section for how to do that.

#### ExcelProtonData

This is a Excel file containing the Iso ppm, csa value and asymmetry value (in that order) for the protons you wish to simulate in a 3D spectrum. (note there must be a minimum of 3 protons)

## Artifact\_Removal Artifact\_mid\_point Artifact\_Removal\_width

Artifact removal smooths a window of the spectrum, where the middle of the window is the 'mid\_point' and the width is the 'width.' A good tip if you wish to do this is to the run code with this off. Then you can click on the tip of the artifact to be able to obtain its X value. You then input this as the mid point and set the width to between 1-15 (depending on the experimental type) and It will remove the artifact. NOTE the width is not the width in ppm or hz but ar arbitrary scale. Between 1-15 should be used. NOTE this must be turned off for R16\_3\_2 experiments.

## Bonus\_RFinhomogeneity

This simulated additional RF inhomogeneity

## Mirror\_f1

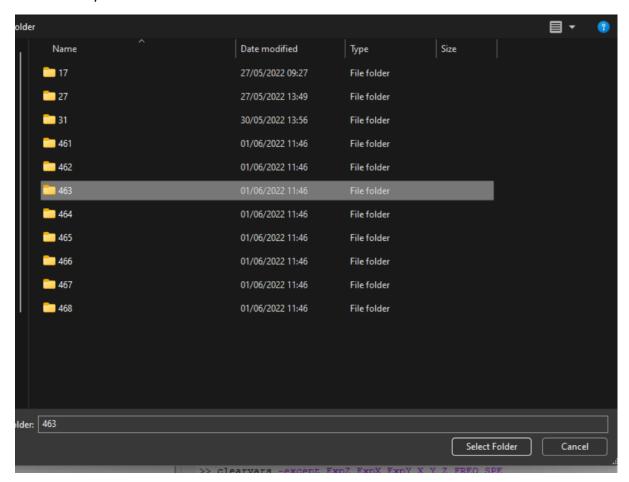
This mirrors the f1 axis in 2D simulations, does not change values simply mirrors the plotting.

## Exp\_2D\_Hz\_width

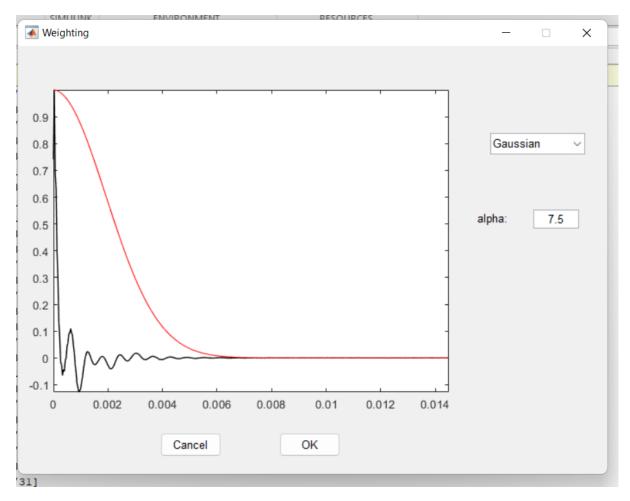
This is the width of the experiment f1 axis. Note this should be set to be ever so slightly under(could be exactly the width but if mistakenly set to big then the code will use this and mess up the plotting)

## Processing R16\_3\_2

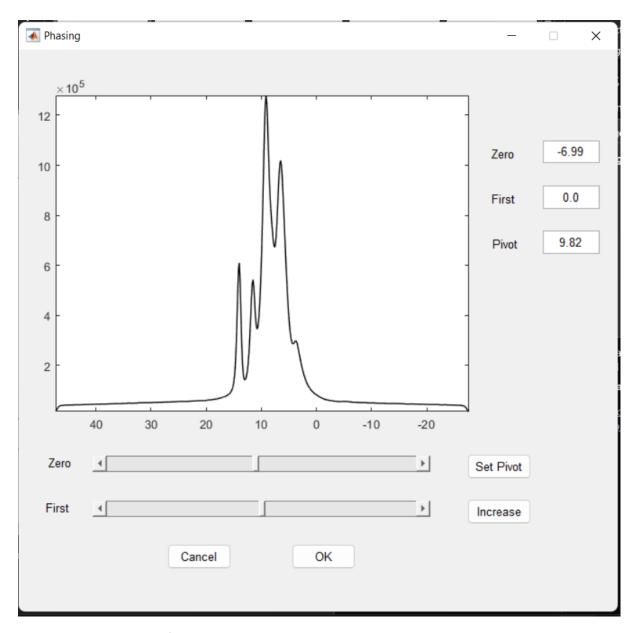
For comparing Experimental data for this sequence you should first run the code titled CSA. Initially this will lead you to a selection window.



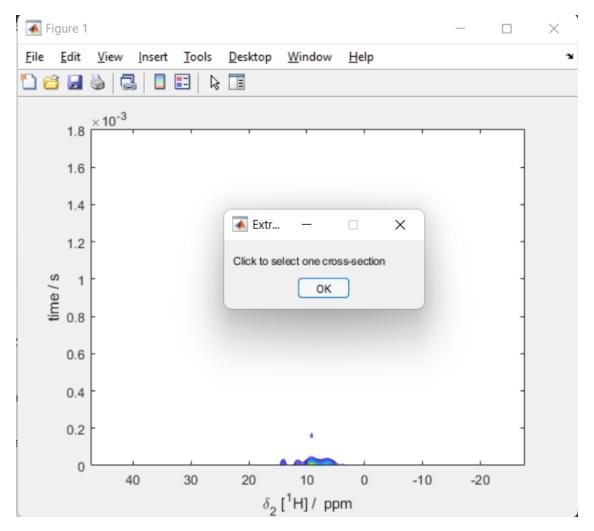
You select the folder that contains the experiment you are interested in.



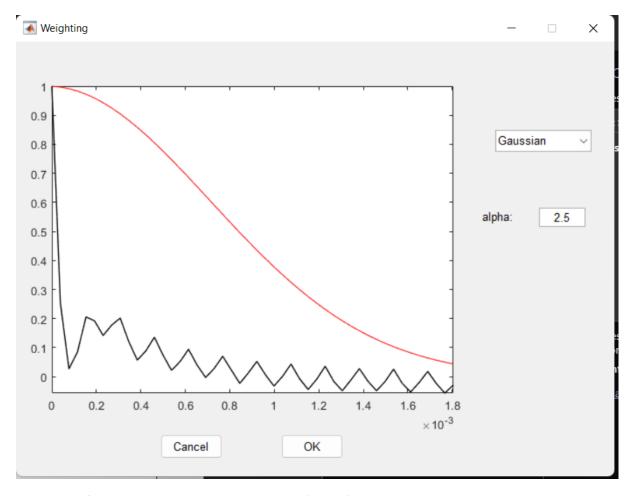
You then weight the f2 axis, a line like above is ideal as to minimise the noise but not the signal.



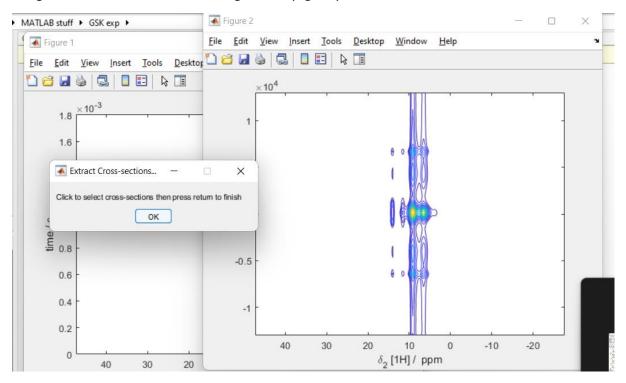
Phase the experiment in f2.



If this window pops up next click ok and click anywhere on the plot, if not ignore this sentence.

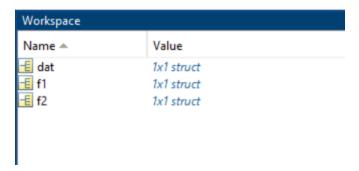


Weight the f1 axis, this should be weighted very 'gently' as above.



Depending on your version it will either as you to select a cross section or one will appear.

If it asks you to select the ok, click a point on the graph and hit enter. This will pop up a cross section. After this you can close down all of the graphs

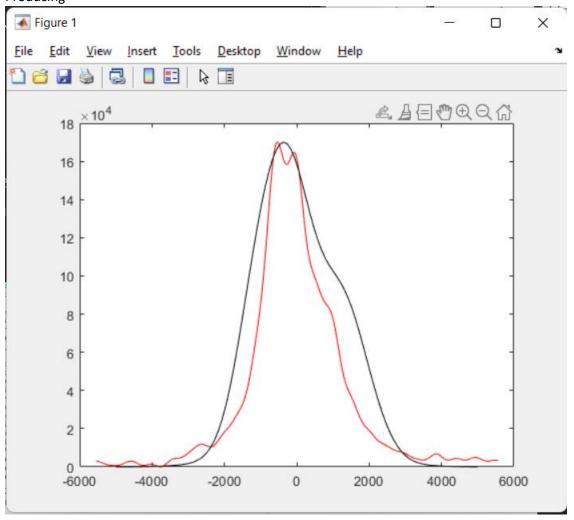


In your work space you will be left with these three structures. Simply select them all and save them together as something '.mat'

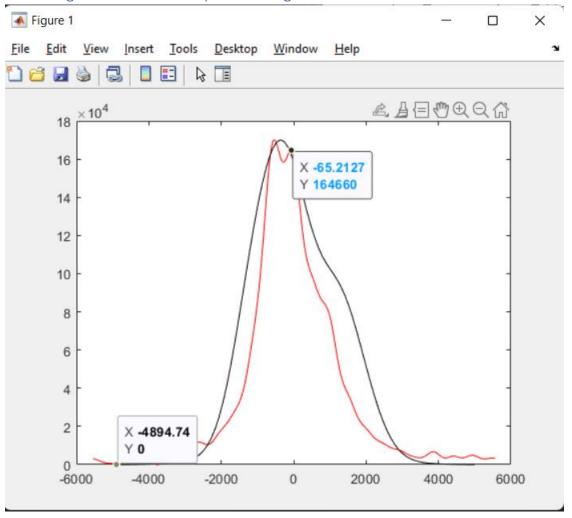
Then simply copy this to the folder containing Pantheon and past it in there. This can then be used instead of a .txt file for ExpFile while doing R16\_3\_2 experiments.

Example: Performing a 2D simulation of SC212 at 20ppm for CSA and 0.5 for asymmetry and comparing it the an experimetral proton at 12ppm.

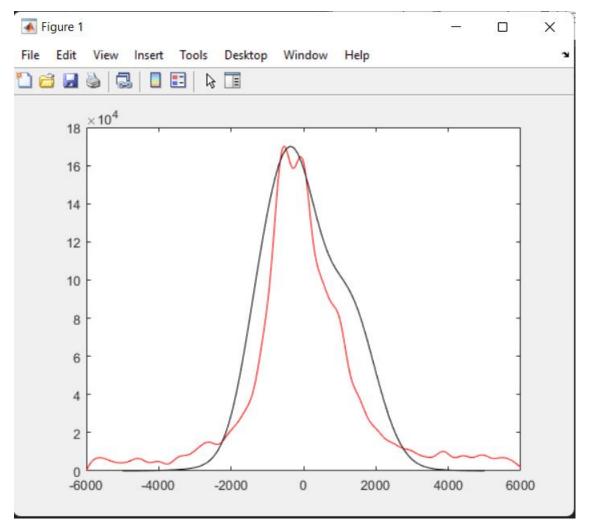
#### **Producing**



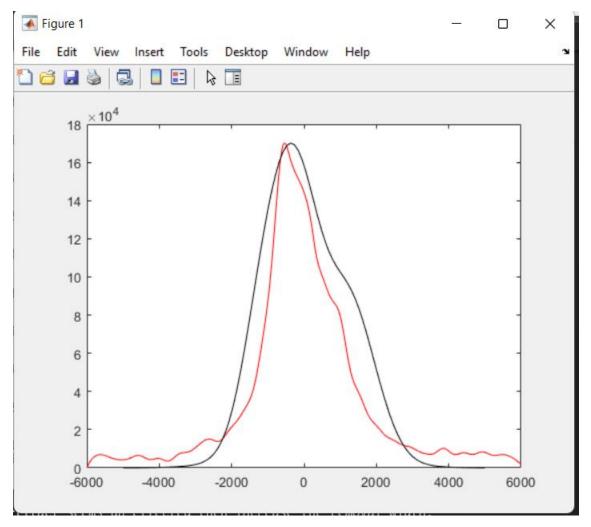
## Removing the artifact for help calculating the CSA.



Find the X value of the artifact by clicking it. Then set Artifact\_mid\_point = -65;



If the artifact seems un effected then increase the removal width.



In this case going to 15 was enough.

# Example: Comparing a 3D simulation of data to experiment data in an R16\_3\_2 experiment.

The csatext.xlsx is used for this. This contains 3 protons that are evenly spaced throughout the typical proton ppm range.

```
%% Settable Parameters
Magnet = 400;
CSA = 20;
                                                                  % Magnet strength in MHz
                                                                  % 2D simulated Chemical Shift Anisotropy value
                                                                  % 2D simulated Asymmetry
% 2D or 3D simulation spinning speed
% Number of points in the f1 dimension of simulation
 Avsm = 0.5:
 Spinning_Speed = 58600;
Spinning_Speed = 58600;
Number_of_points = 96;
Crystal_file = 20;
Gamma_Angle = 5;
DCcor = 46;
Weighting_type = 'Gaussian';
Weighting_value = 2.5;
Searchedppm = 12;
Additional_Sim_Scaling = 1;
Spectral_Width = 26041;
SimType = '30';
MathType3Nowighting = 'Gauss':
                                                               % Number of points in the f1 dimension of simulation

% Crystal file number. Typically use 20, 143, 232, 615 NOTE takes longer

% Number of angles tested. ~5-10 for quick analysis, ~32 for high quality NOTE takes longer

% Number points of the FID that are averaged to bring the dipole-dipole coulping down.

% Type of weighting parameter

% 'Power' of the above weighting type

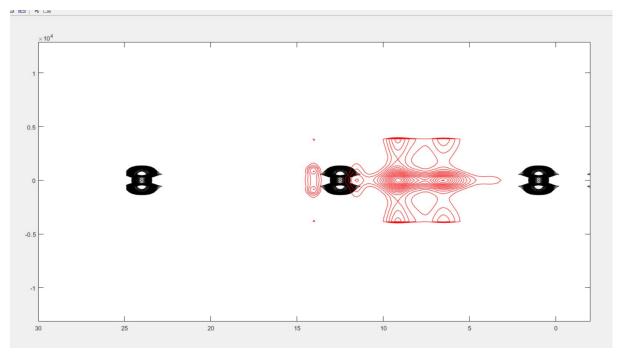
% The ppm value that the '2D' data will compare itself to.

% Used to bring down peak heights to combat artifact

% Only needs to be set for R16_3_2

% 2D or 3D

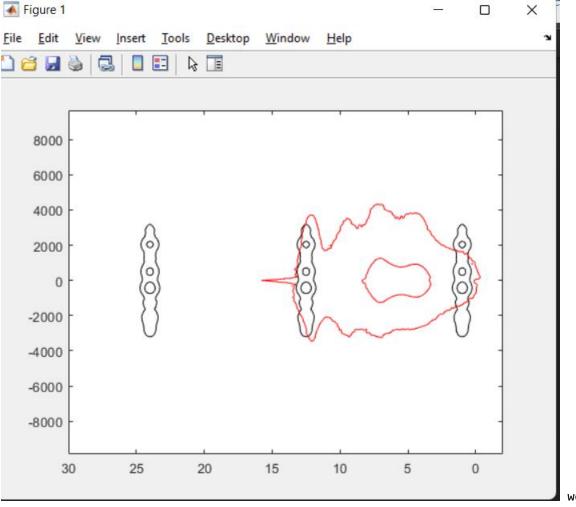
% Gauss or Matt
MathType3DWeighting = 'Gauss';
                                                               % Gauss or Matt
% Currently Available: R16_3_2 SC212 C313
Artifact_Removal_width = 15;
Bonus_RFinhomogeneity = 'Off';
                                                                    "Sets the size of the removal window
                                                               % Uses a larger profile for rf inhomogeneity. NOTE takes longer % Reverses the fl axis
Mirror_f1 = 'Off';
Exp_2D_Hz_width = 12000;
                                                                       % Width of F1 in Hz (only affected when artifact removal is on)
```



If we wanted to decrease the amount of noise that shows we can decrease the number of contours that are seen we can quickly change this using the 'QuickPlot' code.

The pl are the contour levels that show for the Sim and experiment data and these can be changed to whatever you want.

For example setting both to pl=[1.0,0.5,0.1];



get this ugly monster.

## Easy\_Pantheon

