

Preparation for 2D

1. Run and process a normal ^1H experiment running any 2D experiment.

2. Expand the region of interest in the ^1H NMR spectrum. Then click .

3. The window pictured below will open. Write down the values for “SW” and “O1”.



4. After referencing the ^1H spectrum, type “sr” and note down the value.

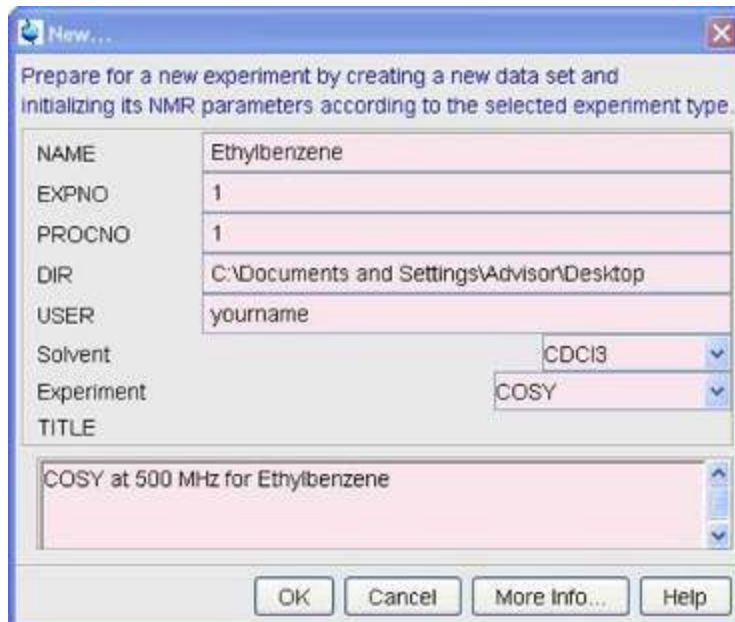
5. Type “**ro off**” to stop sample spinning.

6. Type “**topshim gui**” to open the TOPSHIM menu. From the “Before” drop-down list, choose “Z-X-Y-XZ-YZ-Z”. Click “Start” to start TOPSHIM.

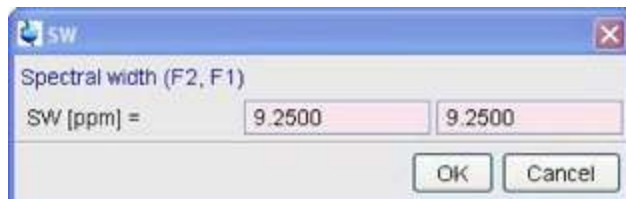
1H-1H COSY

Set-Up:

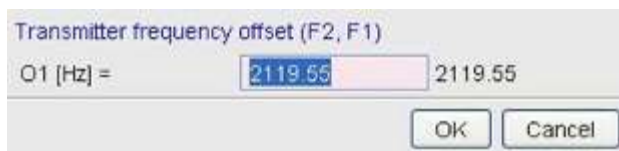
1. Type “**new**” on the TOPSPIN command line to create a new data set.



2. Type “**sw**” and enter the value that you had noted down from the ^1H NMR spectrum into both F1 and F2 dimension

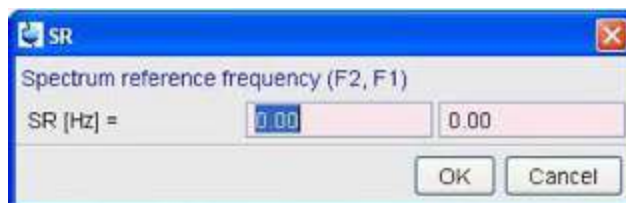


3. Type “**o1**” and enter the value that you had noted down from the ^1H NMR spectrum.



4. Set the required number of scans by typing “**ns**”
5. You can check the time required for finishing the experiment by typing “**expt**”.
6. Type “**rga**” to set receiver gain.
7. Type “**zg**” to start the acquisition.

1. Type “**sr**” and enter the value noted down from the 1H spectrum.

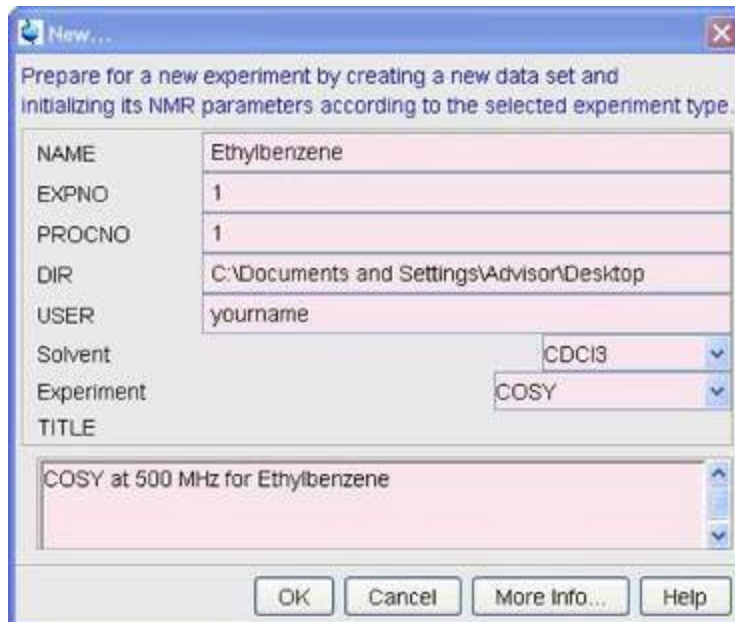


2. Type “**xfb**” - Fourier Transform in both the dimensions.
3. Type “**abs2**” - automatic baseline correction in the f2 dimension.
4. Type “**abs1**” - for automatic baseline correction in the f1 dimension.
5. Type “**sym**” – makes the 2D spectrum symmetric.

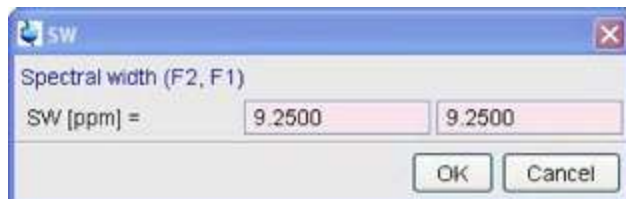
1H-13C HMBC

Set-Up:

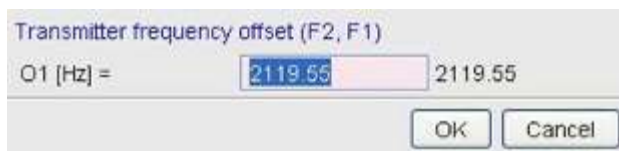
8. Type “**new**” on the TOPSPIN command line to create a new data set.



9. Type “**sw**” and enter the value that you had noted down from the ^1H NMR spectrum into the F2 dimension only.

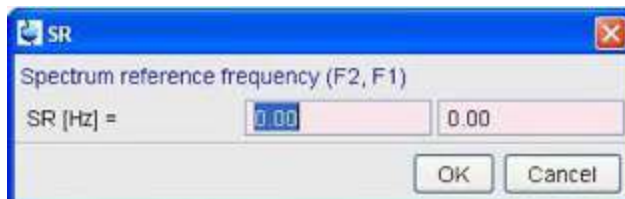


10. Type “**o1**” and enter the value that you had noted down from the ^1H NMR spectrum.



11. Set the required number of scans by typing “**ns**”
12. You can check the time required for finishing the experiment by typing “**expt**”.
13. Type “**rga**” to set receiver gain.
14. Type “**zg**” to start the acquisition.

2. Type “**sr**” and enter the value noted down from the ¹H spectrum for the F2 dimension only.



2. Type “**xfb**” - Fourier Transform in both the dimensions.
3. Type “**abs2**” - automatic baseline correction in the f2 dimension.
4. Type “**abs1**” - for automatic baseline correction in the f1 dimension.