Standard NMR Procedures

Prepping the sample:

- -Wipe NMR tube with a chem wipe
- -Place NMR tube in blue spinner using depth gauge to set the tube to the correct depth
- -Only touch the blue spinner with a chem wipe
- -Place sample into a numbered slot

Take note of number

Taking NMR:

- -Open Top Spin 3.0
 - 1. $sx \# \rightarrow changes sample to the correct numbered slot (from above)$
 - 2. ro on →turns spinner on
 - 3. edc \rightarrow creates new data set
 - -Name
 - -Change directory to your own
 - -Exp/proc are typically 1
 - -Change title to represent sample
 - -Click OK
 - 4. lock \rightarrow find solvent being used
 - 5. rpar \rightarrow read parameter file
 - -Click what NMR is being run
 - -Click READ
 - -Parameter window (acquisition, processing, output data)
 - -Click OK
 - 6. atma →automatic tune and match
 - 7. rsh \rightarrow read shim file
 - -Read most recent date (lastbest)
 - -Click READ
 - 8. topshim 1d \rightarrow shims sample
 - -Lock signal disappears and sample still spins
 - -The magnet shifts along z axis
 - -Already averaged along x and y
 - 9. eda → edit acquisition parameters
 - -Click blue test tube (will not highlight)
 - -NS: number of scans → left at 16
 - -DS: dummy scans → changed to 0 for fast scans
 - -Changed to 4 for carbon
 - 10. rga → pulses to change gain
 - -If the sample is concentrated then the number will be low

- 11. zg →zero go: zeros out waiting commands
 - -Wait for all scans to finish
 - -Y axis is relative intensity
- 12. fp \rightarrow Fourier processes the .fid file
- 13. apk → automatic phase correction

Editing Data:

- 1. Find and reference solvent
 - -Click Process → Calibrate axis
 - -Find and reference middle of peak
- 2. Integrate peaks
 - -Integrate Tab
 - -Yellow highlight button (goal post button)
 - -Click and drag to integrate
 - -Click floppy disk button to save
- 3. Pick peaks
 - -Click and highlight with box
- 4. Plot
 - -Plot tab
 - -Click double folder button
 - -Select H+pp+info
 - -Click paper sun icon to edit graph
 - -Print

LAST:

-sx 1

-Places the standard "ethyl benzene in chloroform" back in instrument

TRACS Site: Dept. of Chemistry and Biochemistry NMR

Use the calendar feature to reserve time.