

NMR New User Workshop

First session

Rui Huang

Notes reproduced from Dr. Weiguo Hu

Link:

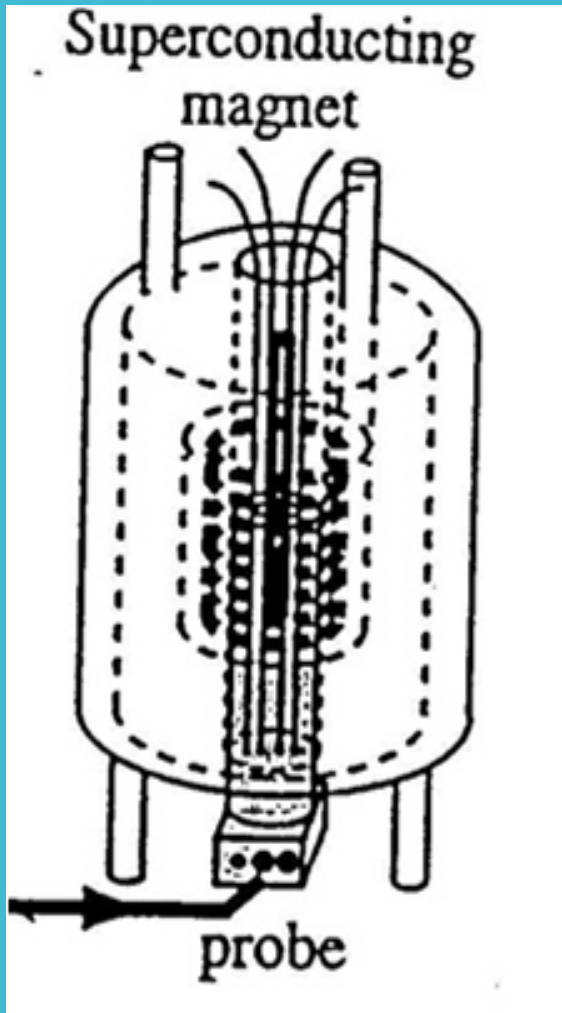
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Booklet:

<https://udrive.oit.umass.edu/weiguoh/nmrhandouts/Introduction-NMR.pdf>

Magnet Safety



- Very strong magnetic field!
- ca. 100,000 times stronger than earth magnetic field
- **Pregnant women** should stay away
- **People with pacemakers** should stay away

Magnet Quench

Quenching is the process whereby there is a sudden loss of superconductivity in the magnet coils, so that all the electric energy in the coils becomes heat and suddenly boils the liquid helium, causing a loud “boom” and people in the room will experience difficulty breathing. You should leave the room immediately when this happens!

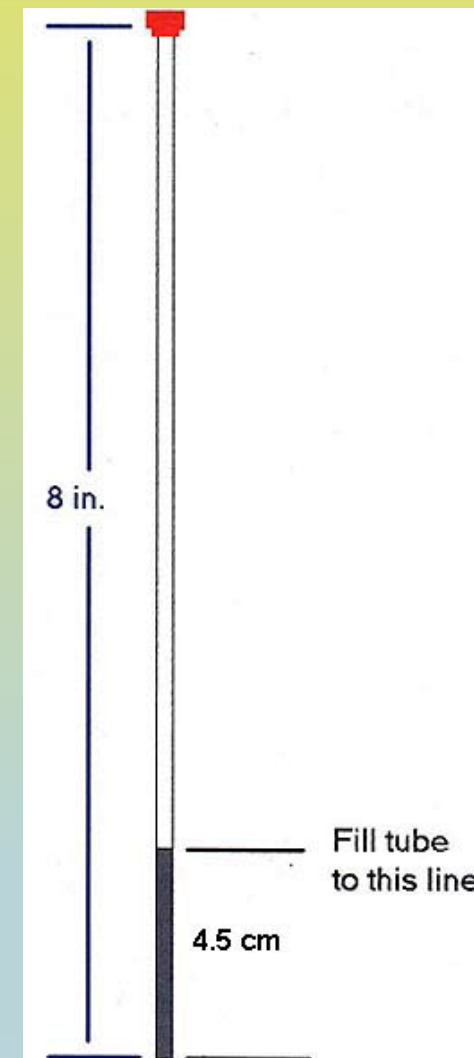


- Never get iron object close to magnet
 - Wrenches, heat gun, etc could fly to the magnet
 - The “bang” would **quench** the magnet

Keep credit cards, cell phone, watches, etc. away

Sample Handling Safety

- **NMR tubes are brittle; handle with care**
 - Breaking sample tubes are frequent events!
 - Select a spinner that fits the tube properly
 - Stuffing kimwipes to fit the tube in spinner is NOT Okay !
- **Common mistake**
 - Insert tube without spinner
- **Measure the correct sample depth**
 - Incorrect depth makes lock and shim difficult
 - too much depth will get the sample stuck
 - If you have less than 4cm sample, center your sample around the thin black line rather than push it to the bottom



What to do if you have insufficient solvent?

<https://blogs.umass.edu/weiguoh/>

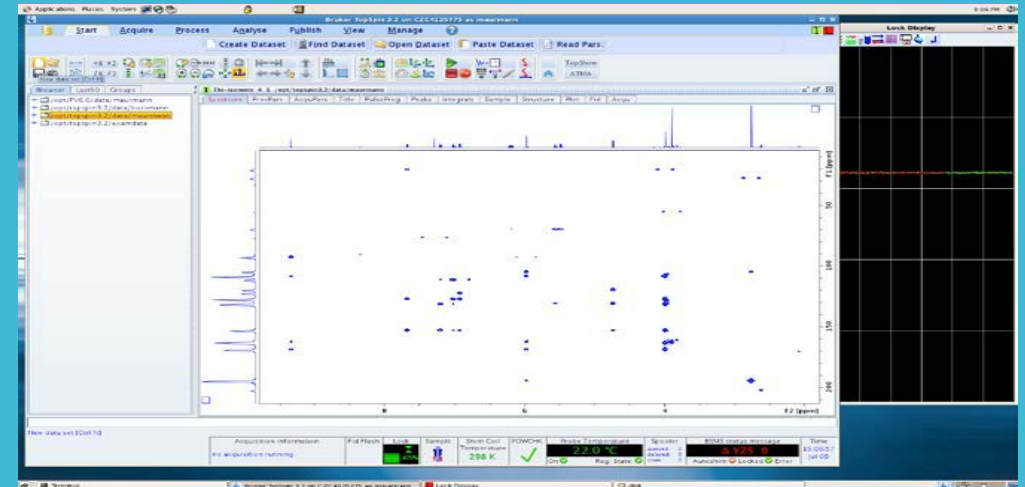
Choices that you have

- Icon NMR



- Multiple samples
- Multiple spectrum
- Undergraduates

- Manual mode



- More choices and flexible
- Faster

Preparations

1. Inject your samples :

- `sx #`

2. Create new file: you must find the correct template file

- `edc`

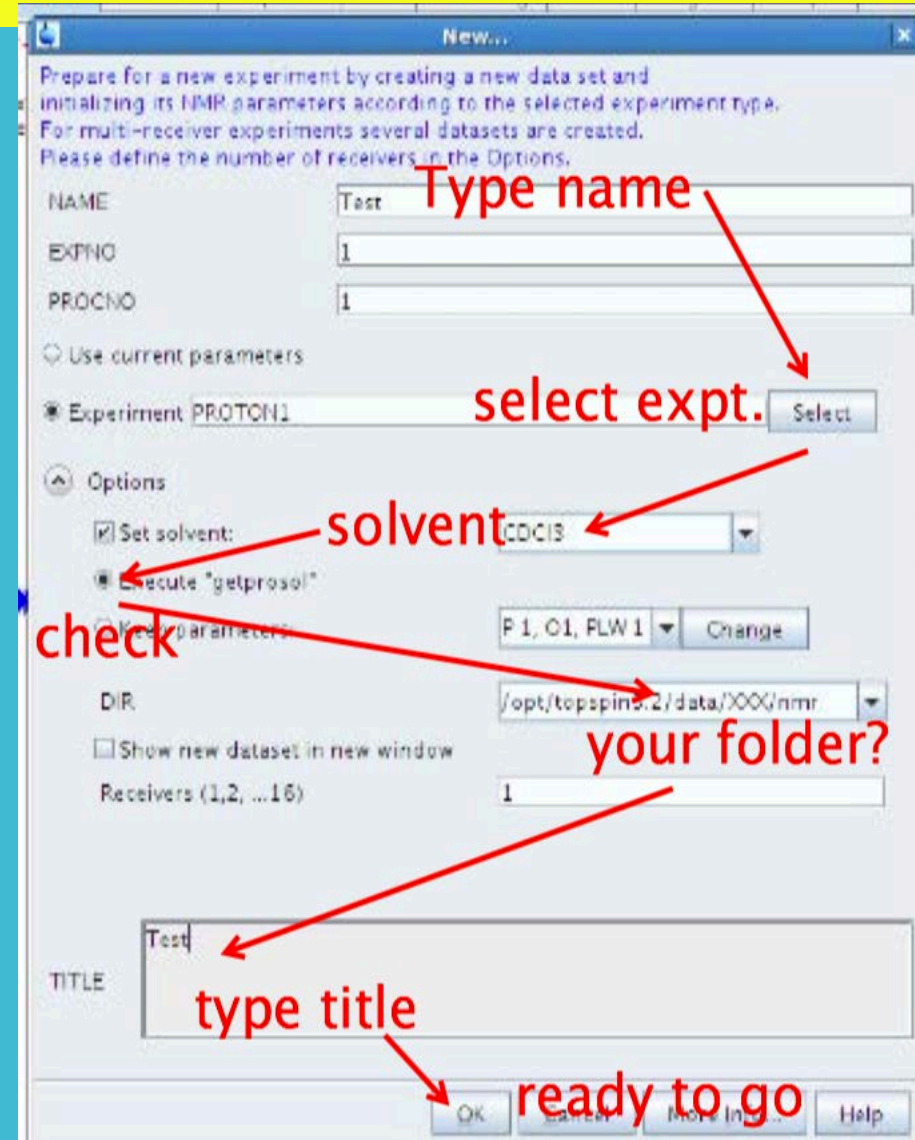
- check get prosol, and see if the DIR is directed to your folder

3. Tuning the probe frequency to what you are going to detect

- `atma`

4. Load a generally good shim set.

- `rsh shims.best`



Shimming and Locking

Shimming: to eliminate inhomogeneities in magnetic field

- the magnetic field is far from homogeneous initially
- the sample become slightly magnetized and create additional inhomogeneous fields.
- the process of correcting for these inhomogeneities is called shimming the magnet

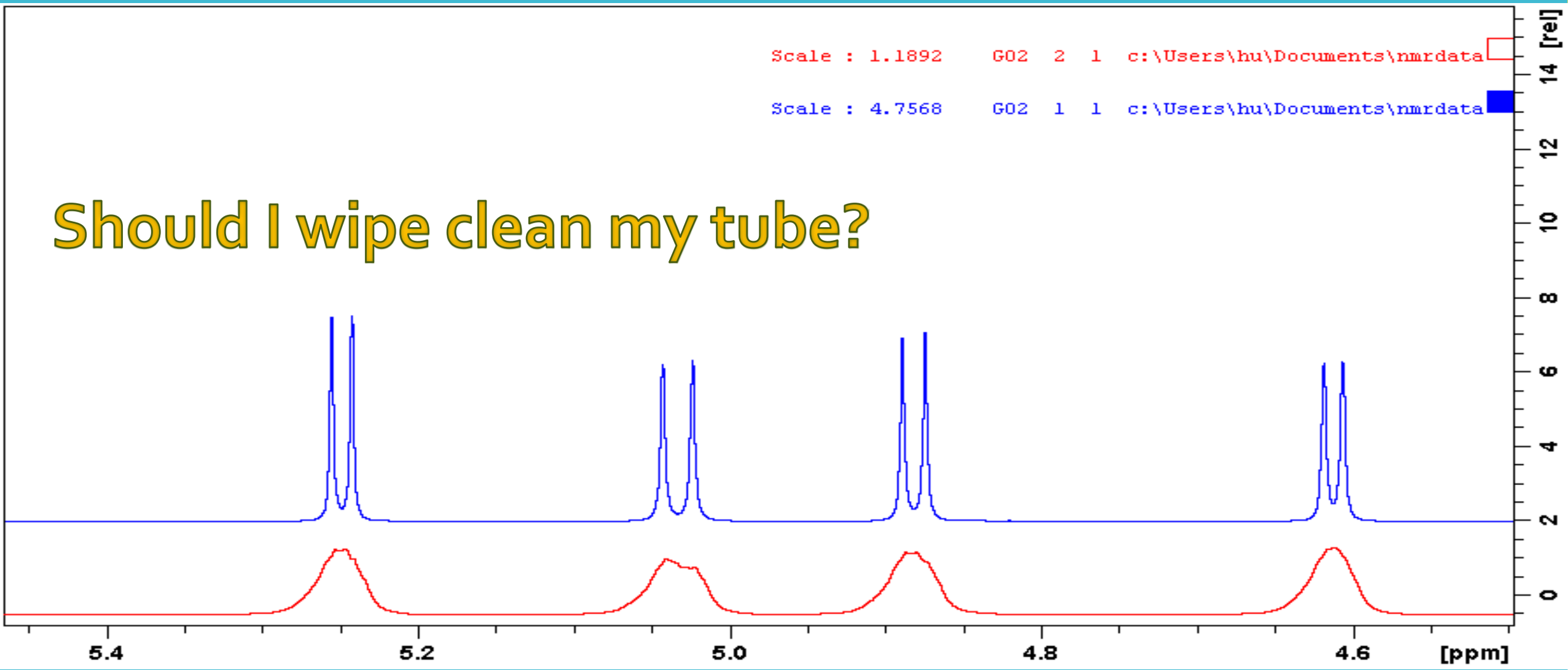
Locking: to compensate for the slight drift of magnetic field

- If the magnetic field strength changes during the course of the acquisition, the signals will appear broadened and the spectrum will have poor resolution.
- use ^2H signal
- This is one reason why we need deuterated solvent to do NMR!
- “**lock [solvent name]**” also calibrates chemical shift

Stable and homogeneous

Shimming and Locking

Should I wipe clean my tube?



Stable and homogeneous

Important Factors That Affect Shimming

rsh shims.best reads in the most recent good shimming parameters

– However, best shimming condition is sample dependent, so you still need to touch-up shim for each sample yourself.

Spinning is optional. Gives better resolution for simple 1D.

– Turn off spin when running 2D

Magnetic susceptibility – everything becomes magnetic in a big magnet!

– Sample homogeneity: insoluble particles, dirt stains on tube wall

– Quality of tube

– Shim would change significantly if you switch between non-polar and polar solvents

Gradient shimming (topshim): directly detect coefficients

How to manual shim?

<https://blogs.umass.edu/weiguoh/>

Running Experiments

- rga** sets receiver gain (rg – just like the volume knob in your radio).
- zg** will start the experiment
- ns** number of scans.
- tr** transfer data to hard disc without stopping expt
- halt** stop expt and save data
- stop** will stop expt immediately and discard data.
- efp** Fourier transformation (with optional line broadening and phase correction)

Processing functions:
feel free to play with all the buttons

<http://www.youtube.com/channel/UCZlxixehEgJONeGTVqMjEjw>

Phasing; Baseline Correction; Integration

Phasing is the “p” in efp

– apk and manual phasing (0th and 1st order)

Baseline must be corrected before integration

– abs is usually adequate for routine uses

– To integrate small peaks on the shoulders of big peaks, manual baseline correction (bas) is necessary

Integrations of ¹H spectra are crudely reliable

Peak heights are not a good measure of population

Phase Correction

Phase distortions cannot be corrected by baseline correction

apk often does not do an ideal job

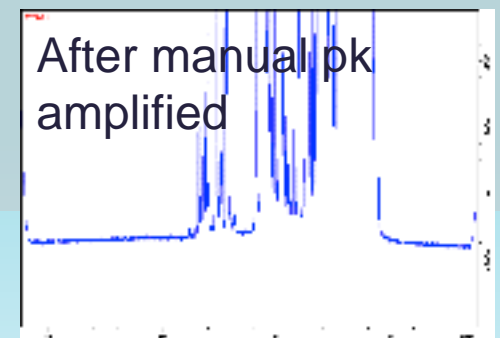
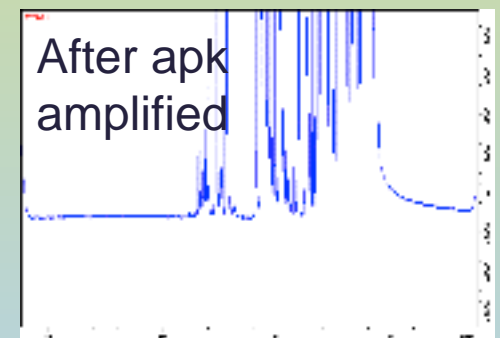
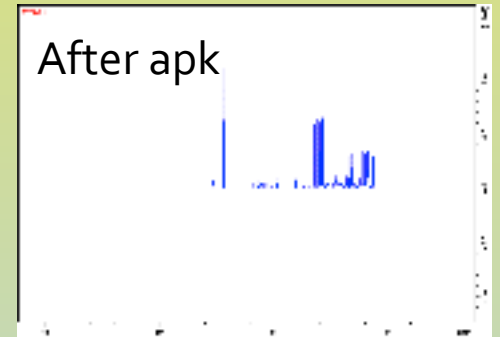
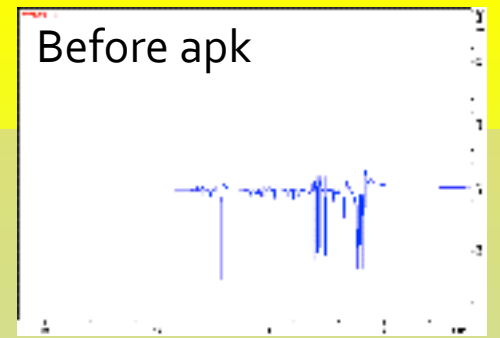
Manual phase correction:

0th order: correct phase of all peaks to the same degree

1st order : correction amount \propto distance to the pivot line

Objective of phase correction:

to make baseline visually continuous so that baseline correction can work (baseline correction requires a continuous baseline)

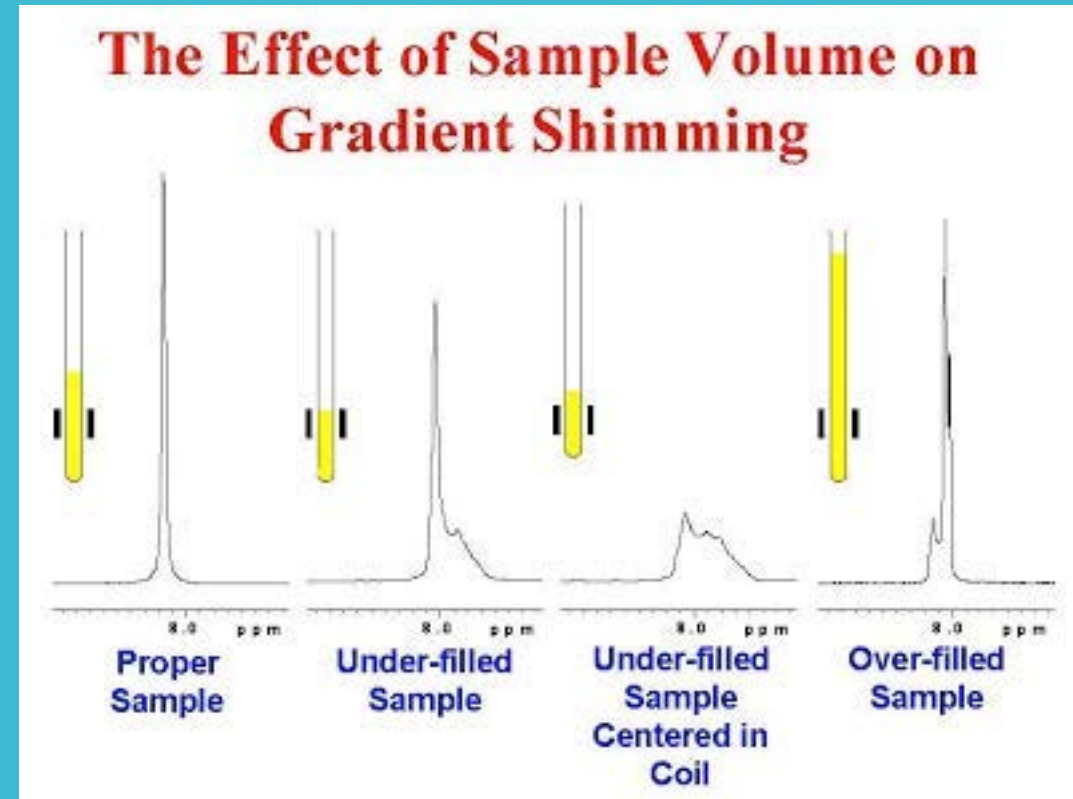


Outline of Procedures

- Remove wallets, watches, keys. Keep steel objects far from the magnet
- Check the blue LED light on the cryo-plateform (Must be **solid** on)
 - Insert your sample: **sx #**
 - Create new file: **edc**
 - **atma**; **rsh shims.best**; lockdisp
 - **lock**; select solvent
 - Shim (manual (bsmsdisp; several iterations of z and z2) or **topshim**)
 - **rga**; **ns**; **zg**; **efp**
 - Phase correction: **apk**
 - Baseline correction: **abs**
 - Integration; transfer data out
 - Get your sample back : **ej**
 - Logout (not logged out promptly will result in excessive charges)

Common Experimental Problems

- Difficulty locking:
 - Bad shimming (2H signal is broad and low)
 - Did not do rsh shims.best
 - Sample depth not correct
 - If less than full sample, did not center sample around black line
 - Instrument may be “Fatigued”... try typing the command again
- Spectrum has a poor resolution:
 - Indications of a shimming problem:
 - Every peak has the same peak shape
 - Peak shapes are often asymmetric



A brief introduction to NMR

<https://blogs.umass.edu/weiguoh/>

How would you explain NMR to a non-scientist?

<https://blogs.umass.edu/weiguoh/>