

# 600MHz NMR

## NMR Training

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# Agilent Hardware

# Basic Operation

## 600 MHz NMR

*“In the absence of a magnetic field the period of all these oscillations is the same. But as soon as the electron is exposed to the effect of a magnetic field, its motion changes.” – Zeeman, P.*

## FIRST PROCEDURES

### Content:

- 1. Sample preparation**
- 2. Logging in**
- 3. NMR instrument**
- 4. Some important line commands**
- 5. Data acquisition**
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## 1. Sample preparation

*“There are no good results using bad samples.”*

CAUTION! Clean the sample of precipitates, cells debris, bubbles, foam, any solid and heterogeneous phases at all in your sample, because different phases will cause distortion on the field and will impact the result. To deal with this issue, use ultracentrifuge in a highest speed as possible to deposit any solid, and/or filter the sample using at least 0,22  $\mu\text{m}$  pore size filter (attention to the filter composition to avoid dissolution of the material by the solvent) right before fill the NMR tube. This step is crucial to obtain a well-adjusted shimming applied to your experiment.

ALWAYS! Clean and dry the sample tube. A tiny scratch, humidity, oily, or dust can destroy the high quality properties of the tube making it useless for high resolution experiments, and also leaving dirty to the inside of the probe, reducing the sensitivity and harming the magnet.

Make your sample using a deuterated solvent (often deuterated water is used for polar samples at 10-20% v/v) until the total volume reaches a volume of  $\sim 500$  to  $600\ \mu\text{L}$  (or  $0.5 - 0.6\text{ mL}$ ) in a 5-mm tube,  $3.1\text{ mL}$  in a 10-mm tube, and  $150\ \mu\text{L}$  (or  $0.15\text{ mL}$ ) for a Shigemi® tube are adequate for removing the end effects. In general, the NMR samples are compatible with: Acetone- $\text{d}_6$ , chloroform- $\text{d}$ , methylene chloride- $\text{d}_2$ , and DMSO- $\text{d}_6$ . Make sure that the solvent will not react or degrade the sample and has a high purity for NMR, free of any impurity. Remember, the peaks of the impurity will appear in the same spectrum of your sample, which can cause a misinterpretation of the result and loss of the sensibility (gain).

If the impurity is part of the solvent, is possible to dry the contaminated solvent through  $\text{N}_2$  stream and dissolving the sample in an appropriate solvent again (repeat the process if you observe any improvement on the result, usually monitored by 1D NMR screening is enough).

The liquid column length must be at least three times the length of the observe coil window to minimize end effects (fig. 1). A typical sample length is  $5\text{ cm}$  regardless of the diameter of the NMR tube used, positioned properly. Make every sample up to the same height in order to obtain similar shim values using samples of that height.

ATTENTION! Some probes may require particular volume and height adjustments. Reduction of sample volume to attain higher concentration usually fails, at this case some special plugs for low volume samples are available and will help with line shape. As well the temperature. Leave the sample become in thermal equilibrium with the probe surrounds, the suggested procedure is to wait for at least 5 minutes or the VT panel/monitor shows a stable behavior with the sample tube inside of the probe before to start any intervention on the equipment, is known that a difference of  $1^\circ\text{C}$  can affect the chemical shift in  $0.01\text{ ppm}$ .

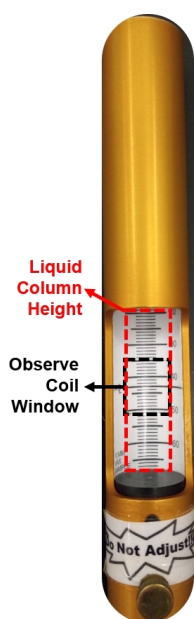


Figure 1. Sample depth gauge:

As a final point, there are few questions to do just before you perform the NMR experiment, as follows:

1. Do I used the depth gauge properly?
2. Do I filled the NMR tube with sufficient volume? The tube is under filled or overfilled, which might cause shim problems?
3. Is the sample homogenous?
4. Are there some particles suspended or precipitated?
6. Are there some paramagnetic substance or high ionic concentration?
7. Is there any possible intrinsic chemical dynamic or exchange in some of the components of my sample?

If your answer was yes for any of these questions, the standard parameters will be modified caused by this specific environment, consider go back to fix any inconvenient, if not possible, consider this specificity during the whole procedure.

## **2. Logging in**

*"Last night I slept like a log. I woke up in the fire place." – Cooper T.*

### **2.1. Lock**

### **2.2. Shim**

## **3. NMR instrument**

*"It is the photographer, not the camera, which is the instrument." – Ardold, E.*

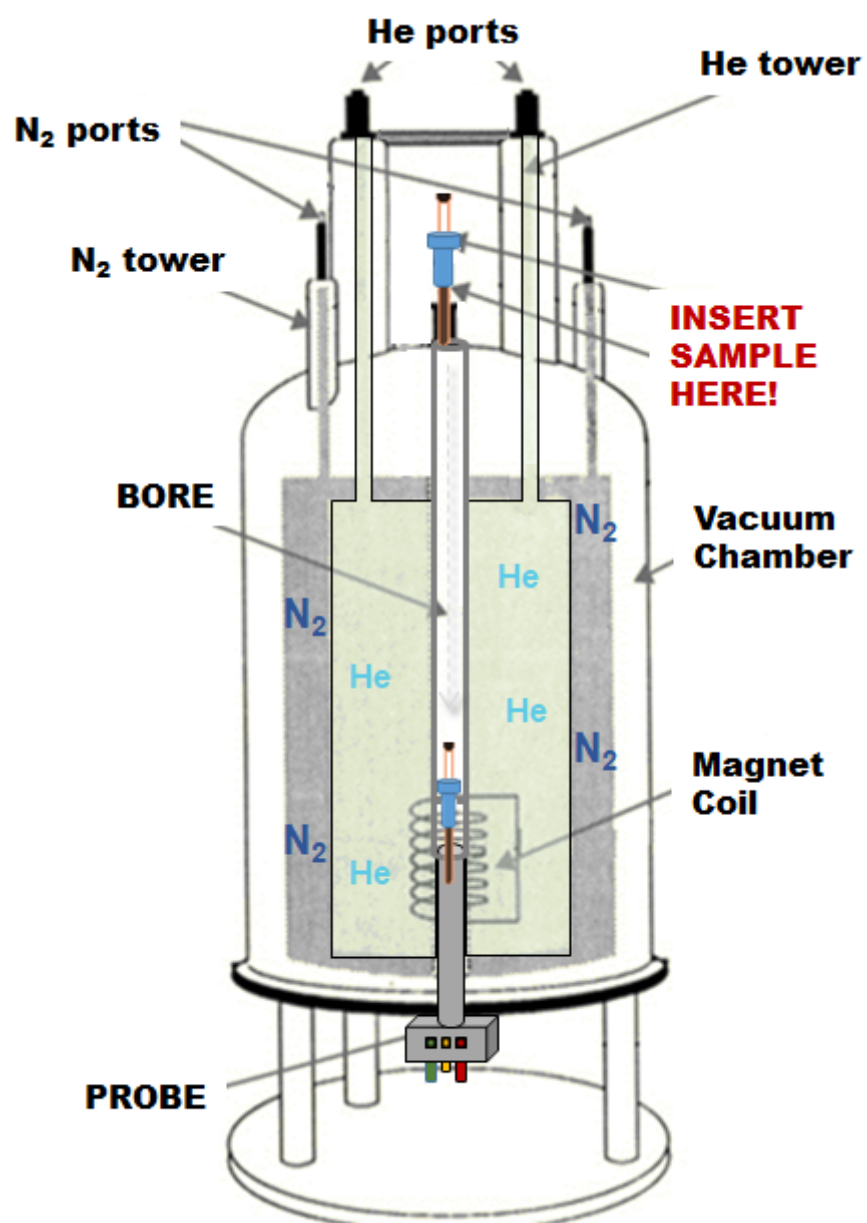


Figure XX:

### 3.1. Locking

### 3.2. Shim

#### **4. Some important line commands**

*"No command, no work."*

#### **5. Data acquisition**

"Experts often possess more data than judgment." - Powel, C.

Always remember because these experiments involve a fourier transform that to double your signal-to-noise ratio requires four times as many scans. So if you can't see anything after 1h you are not going to have much to look at after 4h, likewise if you can not see anything after 2h it is not going to improve a lot after 8h. So on long experiments check your data after an hour or so and if can not see any correlations you might as well stop the experiment.

##### **5.1. Locking**

##### **5.2. Shim**

When the magnet shims are not optimized, the lines in your NMR spectra can take on distinct shapes. The shape of the peak will give you a hint as to which shims need adjustment. Note that lineshapes can also be much more complicated than those below and can even show splittings.

#### **6. Annexes**

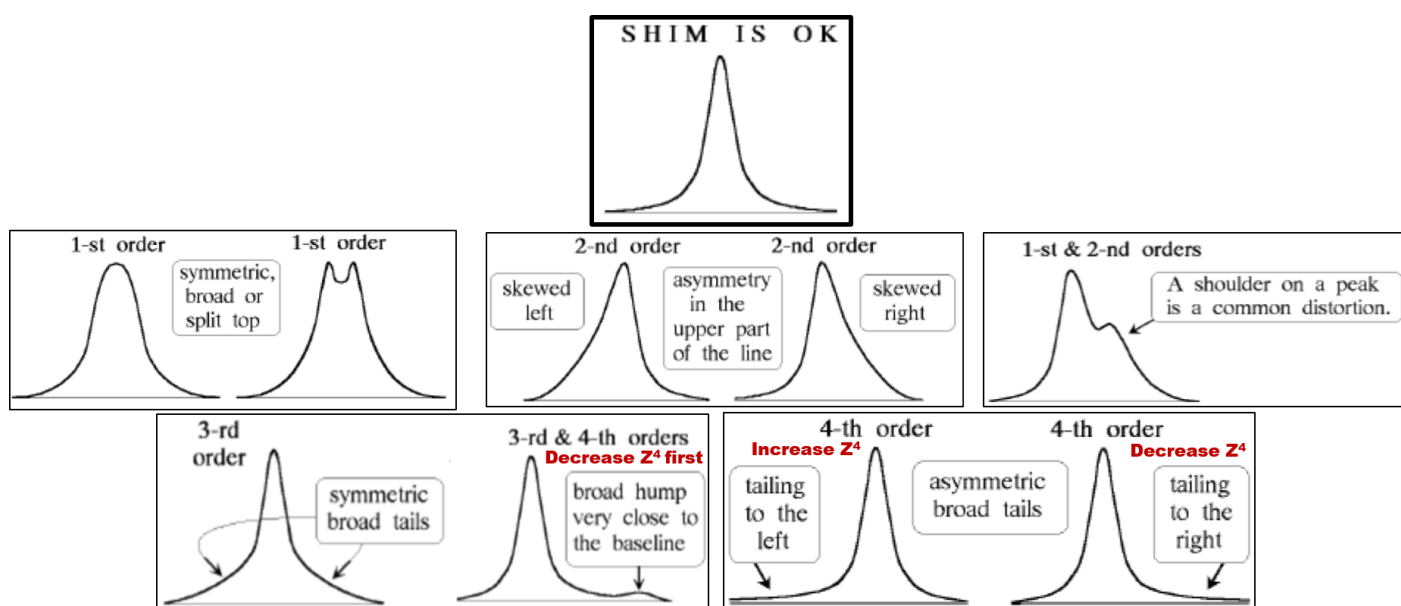
"The reward for work well done is the opportunity to do more" - Salk, J.

#### **7. References**

"Everything I do references something that influenced me" - Abloh, V.

## 8. Troubleshooting

"It's so much easier to suggest solutions when you don't know too much about the problem." – Forbes, M.





# NMR Theory