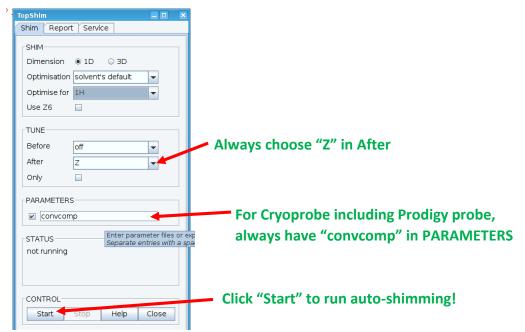
NMR Acquisition Guide

20190819 lzhu

Topspin has a command line to type command. **Each version of Topspin uses the same command.** But the graphic button may change depending on the Topspin version. So I use commands in the following steps:

- 1. Create new dataset: type new, →; Then fill all the information; See page2 for more information
- 2. Insert your sample if you haven't done so. Read "NMR Sample Preparation_UCR" for more information
- Read Shim file(optional): type rsh↓; Then choose the most up-to-date shim: e.g. For Avance600, choose BBFO_AV600.shim; For NEO400, choose Prodigy_NEO400.shim.
- 4. Lock solvent: type lock, ; Then choose the solvent in your sample
- 5. Tune probe: type atma, ; If the probe needs tuned mannually, type wobb,
- 6. Shim via gui panel: type topshim gui, ;



- 7. Revew parameters: type ased, ; check d1, ns, sw, o1p and so on...
- 8. Adjust receiver gain: type rga, ; this could take 30 seconds or so
- 9. Acquire data: type zg, ;

Topspin 2.1 Create New Dataset Window

Please define th	ne number of r	eceivers in the box below.		
NAME	20190816_1	9F 4	<u> </u>	me folder under USER folder. You can change it.
EXPNO	1		Numb	er folder under Name folder. It contains your data.
PROCNO	1		Use di	fferent EXPNO for each dataset.
DIR	/home/opt/nmrsu<			ur group data folder path. Don't change it.
USER	Izhu 🕂			Sub USER(e.g. Izhu)folder under group data folder.
Solvent Choo	se your solv	cDCI3	-	
Experiment Di	rs.	/opt/topspin/exp/stan/nmr/par	user s	— Don't change it.
Experiment		ACIF_13C	-	¹ Usually Expt starting ACIF is good ones, ready to
TITLE				go. Choose expt by using the drop down menu.
Your da	ta title. Add	l all the notes you want.		
1 Receivers	(1.28)			

You can use different NAME folders for each dataset. Or you can use different EXPNO number folders under the same NAME folder.

Topspin 4 Create New Dataset Window

NAME 20190819_test	NAME folder under Directory. You can change it. Number folder under Name folder. It contains your dat		
EXPNO 1			
Directory /opt/nmrsu/CRL	Your main data folder path. Please make su		
🗹 Open in new window	that is the right folder.		
> Parameters			
○ Use current parameters			
Read parameterset ACIF_1H	Select		
	Usually Parameterset starting ACIF is good ones,		
Do nothing Choose your solvent	ready to go. Choose your preferred parameterset		
 Execute getprosol 			
○ Keep parameters P 1, O1, PLW 1 Change	by clicking the "Select".		
Advanced			
Number of datasets (receivers)	1		
Title			

You can use different NAME folders for each dataset. Or you can use different EXPNO number folders under the same NAME folder.