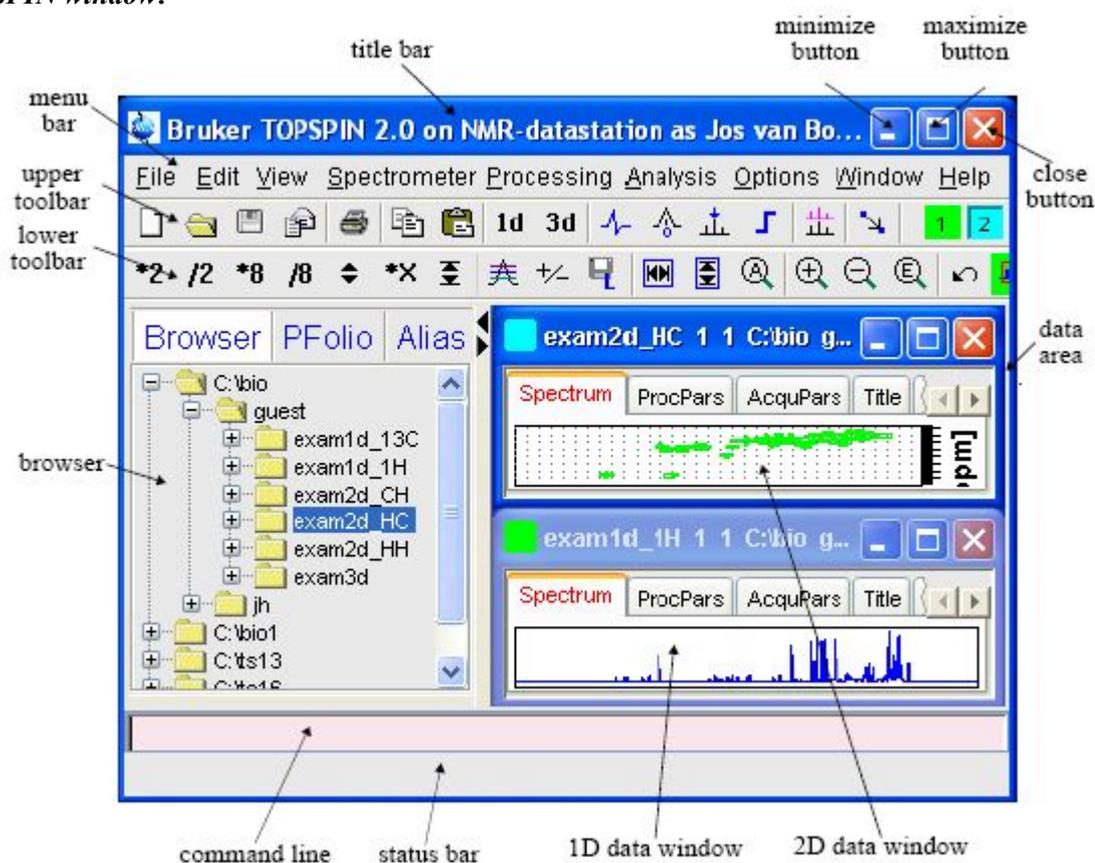


## 1. Start TOPSPIN

- Login using your group ID and password
- Double click TOPSPIN icon to start TOPSPIN software

*TOPSPIN window:*



## 2. Insert the sample

**blue spinner** (temperature 15-35 °C), **white ceramic spinner** (other temperature). Replace **ONLY** with the dummy standby sample ( $\text{CDCl}_3$ ) into the magnet when you are done.

- Type **ej** in the command line or Press the “lift on/off” (green = on) on BSMS Window, take the dummy sample out, wipe your tube, then insert the sample to Spinner into the depth measure then slide the tube down to the bottom.
- Type **ej** in the command line or Press the “lift on/off” (green = on) on BSMS Window



**Make sure your NMR tube is supported by compressed air before releasing it.**

- Type **ij** in the command line or Press the “lift off” button (no light) on BSMS Window

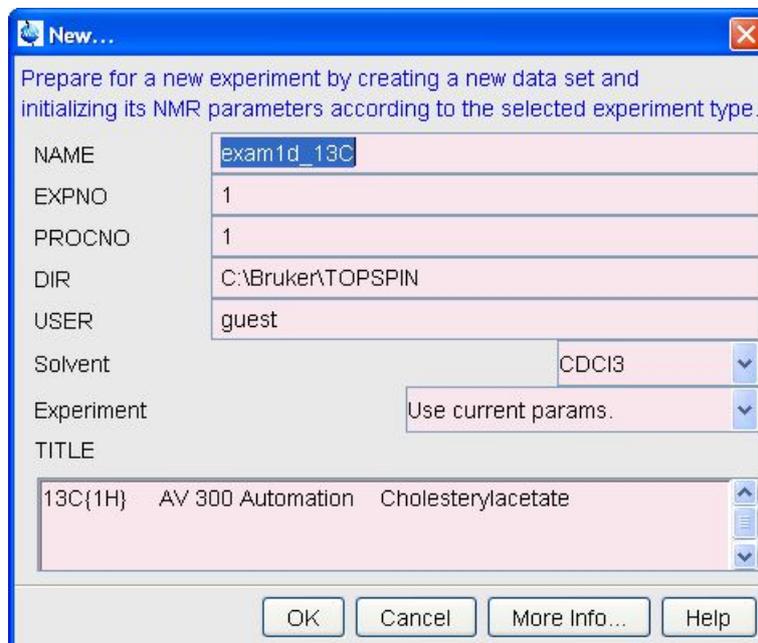
### 3. Open an old dataset

In the Browser window, locate your data, right-click a dataset name, and choose **Display** in the popup menu

### 4. Create a new dataset

a. Type **edc** in the command line

In the popup dialog box:



The screenshot shows a 'New...' dialog box with the following fields and values:

NAME	exam1d_13C
EXPNO	1
PROCNO	1
DIR	C:\Bruker\TOPSPIN
USER	guest
Solvent	CDC13
Experiment	Use current params.
TITLE	13C{1H} AV 300 Automation Cholesterylacetate

b. Specify name, expno, procno, dir and user

c. Click the down-arrow of the **Solvent** box to choose a solvent from the list

d. In **Experiment** box, select Use current params

e. Type the dataset title in the **TITLE** box

f. Type **rpar** in the command line to choose a parameter set from the list

For example:

Parameter set name: proton experiment = 1\_protonstd

Carbon experiment = 1\_carbonstd

### 5. Tune, lock and shim

a. Type **atma** in the command line

b. Type **lock** in the command line and select a solvent from the popup list

c. Type **topshim** in the command line

\*type **ro** in the command line, then type **20** at rate to the popup box to turn the spin on (not recommended; definitely do not spin for 2D experiments)

d. If shim is messed up, type **rsh** in the command line to select most current shim file from the list, then type **topshim**.

### 6. Acquire FID signal and modify acquisition parameters

a. Type **rga** in the command line, then

b. Type **zg** in the command line

c. Sometimes it is necessary to modify acquisition parameters

- Modify acquisition parameters

- Clicking **AcquPars** tab in the tab bar of the data window

•or type **eda** in the command line

## 7. Process 1D spectrum

a. Type **efp** in the command line

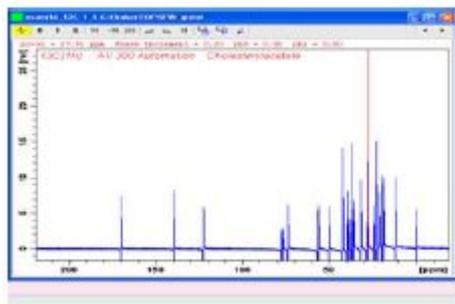
b. phase correction

Automatic method:

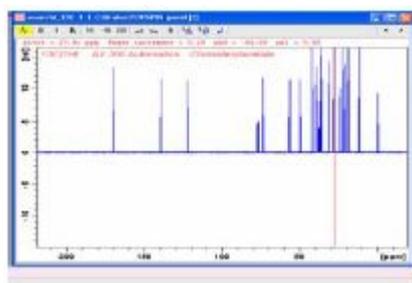
Type **apk** in the command line to execute automatic phase correction

Manual method:

- Click phase correction  button in the upper toolbar
- The Tab bar of the active data window will be replaced by the following toolbar

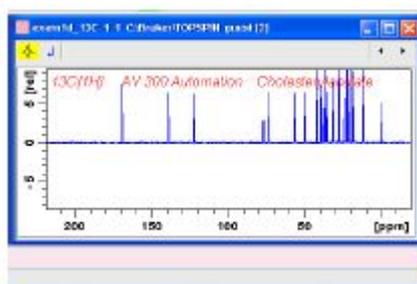


- Left-click-hold the  button and move the mouse until the reference peak is exactly in absorption mode
- Left-click-hold the  button and move the mouse until the entire spectrum is exactly in absorption mode
- Click the  button to save and execute the phase correction



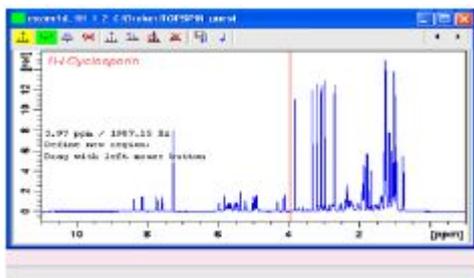
c. Chemical shift calibration

- ◆ Click the  button in the upper toolbar, and the Tab bar of the active data window will be replaced by the following toolbar



- ◆ Position the red cursor line at the reference peak
- ◆ Left-click at that position and enter the chemical shift of the reference peak at the popup dialog box





✧ Define peak picking regions: Note: the active button is highlighted in green



: Define peak picking range



: Change peak picking range



: delete all peak picking regions

When the  button is green, put the cursor at the upper-left corner of a peak picking range, then left-click-hold and drag the cursor to the low-right corner of the range. You can use this  button to modify the peak picking range.

•Other buttons in the toolbar



: Define peaks manually



: Define peaks semi-automatically



: Delete all peaks



: Save the peak region and peak list and return



: Return, discarding any changes

## 8. Plot 1D spectrum

Click **file** in the menu bar, then select **print**, click **OK** in the popup dialog box, then click **print** in the popup dialog box.

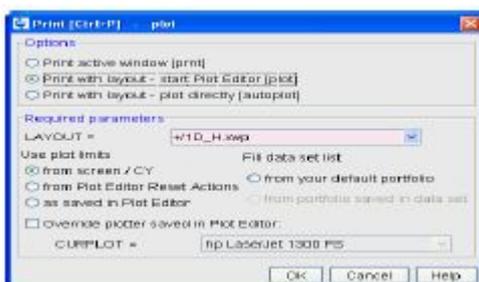
## TOPSPIN PLOT EDITOR INSTRUCTION

### 1. Use TOPSPIN Plot Editor to plot a spectrum

- a. Type **Layout** in the command line to select the desired layout by clicking down-arrow button of LAYOUT box , then type **plot** in the command line and TOPSPIN Plot Editor will start



- b. OR **File**→**Print**, and select **Print with layout-start Plot Editor** in the popup window

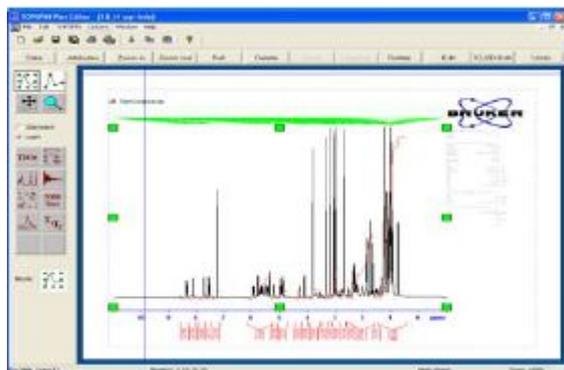


In the required parameters, select the desired layout by clicking down-arrow button in **LAYOUT** box. After clicking **OK** button, the TOPSPIN Plot Editor will start

The layout can be specified by using one of the following abbreviations:

- + : the standard layout directory: ../topspin/plot/layout
- ~ : the user home directory
- # : current processed data directory

- c. Preview the current plot layout and plot (Click **File**→ **Print**)



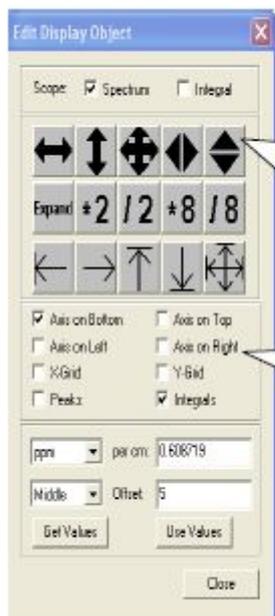
- d. Modify the plot layout

Move, resize and delete an object (spectrum, title, parameter or logo):

- Mark an object by clicking the  button and then clicking the object
- Click-hold the object and move the mouse to move the object
- Click-hold one of the green markers and move the mouse to resize the object
- Click the  button in command bar to delete the object

### Modify the spectrum

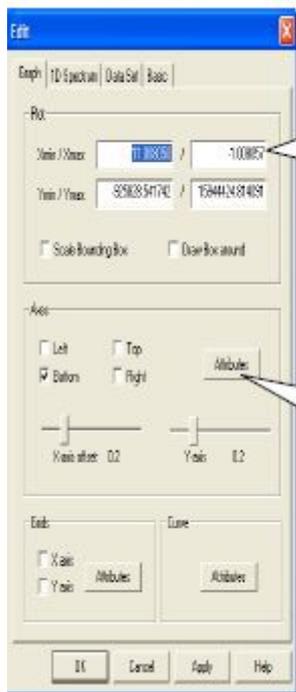
- Mark the spectrum object and click the **1D/2D-Edit** button in the command bar



All buttons, which have same functions as in TOPSPIN, are active for spectrum or integral or both by setting the scope

Checking the various buttons in this part to change the spectrum object

- Mark the spectrum object and click the **Edit** button in the command bar



Under **Graph** tab, you can expand a spectrum by entering the exact limits of a certain spectral region

You can modify the attributes of Axes and Curve



Under **1D Spectrum** tab, you can modify the units of axes and peak (ppm or Hz), attributes of peak labels and integral labels

Under Linux, all parts (Graph, 1D spectrum. DataSet and Basic) are shown simultaneously

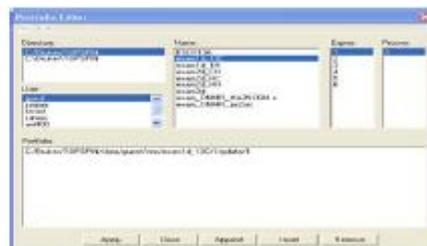


Modify parameters and title

- Right-click the object and choose corresponding buttons in the popup menu to modify the object

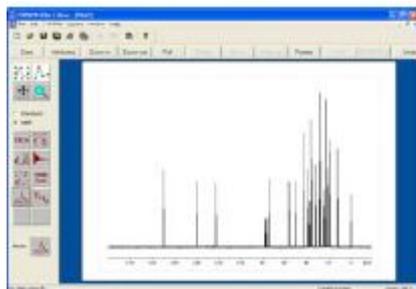
**2. Plot several 1D spectra in stack mode in Topspin Plot Editor**

- Click the **Data** button in the command bar, click **Edit** button in the popup **Data Set Selector** window. The **Portfolio Editor** window will pop up



- In **Portfolio Editor** window, choose right **Directory** and **User**, all datasets will show up.
- Choose the first spectrum by clicking the respective entries in the sections **Name**, **Expno** and **Procno**. Then click the **Append** button.

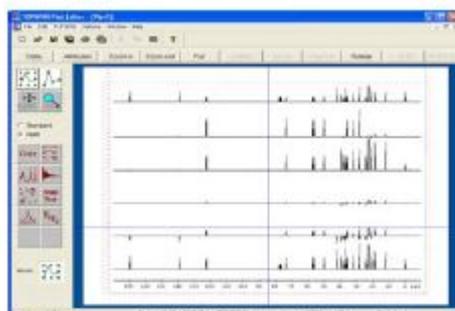
- d. Repeat step c for the rest of spectra, then click **Apply** back to **Data Set Selector** window, click **OK**
- e. In **TOPSPIN Plot Editor**, click **File** → **New** to open a new layout
- f. Click the  button, click-hold left mouse button and drag in the layout area



- g. Mark the spectrum by click the  button, then click **Edit** button in command bar. The popup window is



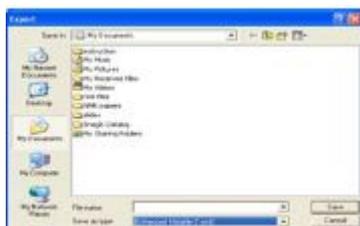
- h. Click **Stacked** menu bar, fill the box. In Spectra Offset box, the first number is offset of X-axes, and the second number is offset of Y-axes. By adjusting these two offsets, you will get the desired layout



### 3. Export a spectrum as PDF or PNG or EMF format file so you can insert it to your report / thesis

#### a. From Topspin Plot Editor

- A spectrum is appeared in the Plot Editor with desired layout
- Click **File** → **Export**



- In the box of **Save as type**, you can choose the type you want, and put

