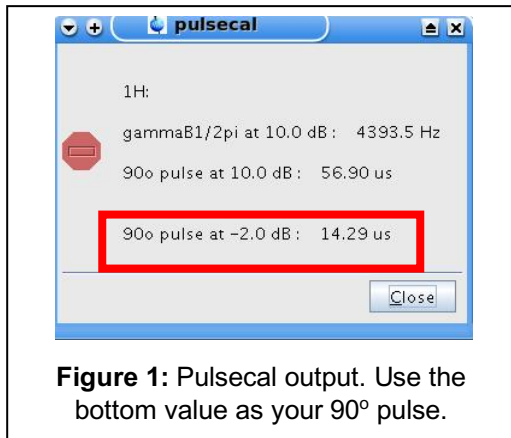


## Calibration of 90° Pulse Lengths

<p>Do all of the usual set-up stuff</p>	<p><b>PART I – INTRODUCTION</b></p> <ol style="list-style-type: none"> <li>Occasionally, some experiments will require more careful pulse length calibration.</li> <li>This handout will teach you how to calibrate the 90° pulse for <sup>1</sup>H (or other nuclei). It assumes basic familiarity with running a routine 1D NMR experiment.             <ol style="list-style-type: none"> <li>You will need to do the usual things: log on, insert your sample, create a new experiment, read in the parameters, lock, shim, tune, etc.</li> </ol> </li> </ol>
<p>new</p> <p>rpar</p> <p>pulsecal</p>	<p><b>PART II – CALIBRATE <sup>1</sup>H 90° PULSE</b></p> <p><b>1. Automatic 90° pulse Calibration</b></p> <ol style="list-style-type: none"> <li>Type <b>new</b> to set up a new experiment. Select the proton experiment from the panel on the right side of Topspin or type <b>rpar</b> to read in the proton parameters.</li> <li>Execute the automatic 90° pulse calculation by running the command <b>pulsecal</b>. Pulsecal returns three values. Take note of the bottom one, your 90° pulse (μs) and the power level (dB) (Fig 1.) If pulsecal fails ("too many entries in peaklist"), proceed to manual calibration.</li> </ol>
<p>pulprog zg</p> <p>getprosol</p> <p>rga</p> <p>ns 1 ds 0 zg ef apk</p> <p>change p1 to p1*4</p>	<p><b>2. Quick Manual 90° Pulse Calibration</b></p> <ol style="list-style-type: none"> <li>Follow steps 1 and 2 of the Automatic Pulse Calibration procedure above.</li> <li>Set the pulse program <b>pulprog</b> to zg (should have been set as zg30 before).</li> <li>Type <b>getprosol</b> to ensure that the most recently determined values for pulse times and power levels for all nuclei in the particular pulse program will be used.</li> <li>Type <b>rga</b> to start an automatic receiver gain set routine. When this is finished, the message <b>rga:finished</b> is output above the pink command line. Record the number that is set as <b>rg</b>.</li> <li>Set <b>ns 1, ds 0</b>.</li> <li>Execute the experiment by running the command <b>zg</b>. After the experiment is done, process the data by running the commands <b>ef</b> (exponential multiplication and fourier transform) and <b>apk</b> (automatic phase correction).</li> <li>Change <b>p1</b> to a value that is four times the initial value, e.g., from p1 = 10,</li> </ol>



change to  $p1 = 40$ . You can do the math in the box by entering  $*4$  after the current value. Rerun the experiment with **zg**.

a. After the experiment is done, process the data with **efp**. Do NOT run apk! If you do, you will need to restart from step 3!

b. Inspect the spectrum, then change **p1** and acquire again until you get approximately a null signal (equal parts positive and negative).

c. Make sure you wait at least 5 seconds between each acquisition to allow for sufficient relaxation.

d. The null signal is your  $360^\circ$  pulse. Divide the **p1** value corresponding to your null signal by 4 to get the  $90^\circ$  pulse. To save keystrokes, you can use the command **zefp**, which combines **zg + efp** and **overwrites your data without asking**.

7) Make note of the **p1** value for your 2D or other experiments.

or you can use  
paropt

### 3. $90^\circ$ Pulse Calibration with paropt

1) Alternatively, you can use the *paropt* command to calibrate your  $90^\circ$  pulse length. This will give you a full sinusoidal pulse calibration profile.

2) Follow steps 1-4 of the Quick Manual Calibration step above

3) Set  $p1$  to  $p1/2$  (e.g., if  $p1 = 14$  after getprosol, set  $p1 = 7$ ).

4) Acquire a spectrum with **zg**, then **ef; apk**.

5) Zoom in on the peak you wish to calculate the  $90^\circ$  pulse for.

6) Type **dpl1**. This will set the left and right x-coordinates (F1/F2 or F1P/F2P) and the vertical scaling of the result.

7) Type **paropt** and enter  $p1$ , the first value, the increment, and the number of measurements in the series. The first scan should be set to a few microseconds less than  $p1*4$ . Increments of a few microseconds is usually good enough.

8) Note: Each time *paropt* runs, you are left in *procno 999*. You can return to the *expno* and *procno* from which you started by using the *re* command (i.e., if you started in *expno 1 procno 1*, type *re 1 1*).

9) Expand the vertical scale as necessary to judge the null point as the point with symmetrically distributed residual signal. You can either use the

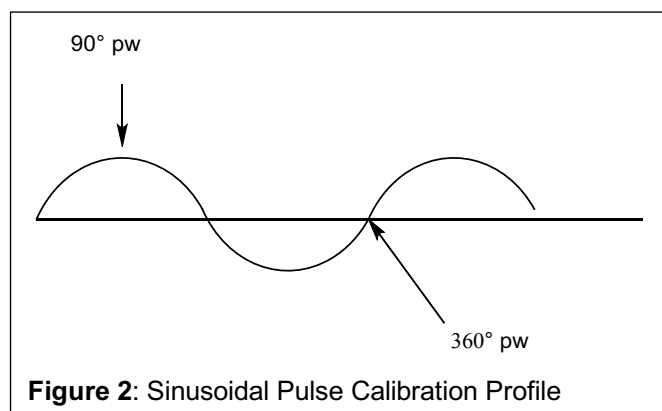


Figure 2: Sinusoidal Pulse Calibration Profile

set  $p1 = p1/2$

zg; ef; apk

dpl1

paropt  
and follow the  
prompts

Look for the null  
point. That is your  
 $360^\circ$  pulse; the  $90^\circ$   
is the  $p1$  value  
corresponding to  
the null point  
divided by 4.

kill if you need to

scaling buttons on the left or the equivalent command line commands to rescale the display while *paropt* is running. You'll have to mentally count the number of increments over to the target peak, so choose start values and increments that make the arithmetic easy.

- a. If you do not want to wait for the full array to run, you can **kill** the experiment. Type **kill**, and a window of all of the running processes will pop-up. Select *paropt* and hit the "Kill" button. Exit out of the pop-up window.
- 9) Consult one of the NMR staff or the Bruker manual if you have any further questions on *paropt*.

### **PART III – CALIBRATE A 90° PULSE FOR ANOTHER NUCLEUS**

- 1) You can use any of the above methods to calibrate pulse lengths for other nuclei.
- 2) However, if you do not have enough SNR to do a single scan, you must use labeled compounds for low abundance nuclei and signal average. This means you will need to change your *d1* value appropriately to account for the  $T_1$  relaxation of your compound! Contact facility staff to setup pulse parameters for currently unused nuclei.