

Basic Acquisition:

- new (edc) Make new data set using all the current parameters
- atma automatic probe tuning
- rsh <name> Read shim file, e.g., rsh LAST
- lock Lock on deuterated solvent (or run lock <solventname>)
- topshim, topshim gui Automated gradient shimming Z₁₋₅ (use GUI for X, Y, etc. as well)
- rpar <name> Read parameter set
- rga, rg Automatic receiver gain adjust/manual adjustment
- ns, td0 Set number of scans, number of times to acquire ns scans
- zg, go Zero existing data (if any) and acquire/add to existing data
- halt, stop Halt acquisition and save fid/stop without (!) saving fid
- ft,ef Fourier transform (ft), exponential multiply + Fourier transform (ef)
- efp Combines em, ft, pk (exponential multiply, ft, phase correct)
- apk/.ph Automatically/manually determine phase correction
- abs n/abs Automatic baseline correction without/with automatic integration
- pps, pp Peak pick to screen, print peak pick
- int Integrate peaks
- plot Plot program

More Advanced:

- ased, eda Edit current pulse program parameters/edit all parameters
- sw, swh Spectral width in ppm/Hz
- o1p, o1 Center of spectrum in ppm/Hz
- getprosol Read probe/solvent dependent pulse width and power
- tr Write accumulated scans to disk
- gs Go setup mode, observe fid/spectrum (Hz/ppm) in acqu window
- xppdir Display, load pulse program
- edte Variable temperature control
- pulprog Pulse program
- zefp Combines zg and efp AND OVERWRITES CURRENT DATA WITHOUT A PROMPT!
- p1 90° observe channel pulse (or run p1 <new value>)
- d1 Recycle delay (in most pulse programs) (or run d1 <new value>)
- <command1>;<command2> Execute multiple commands in sequence (can't interrupt)
- (up arrow), (down arrow) Scroll through history of previous commands
- re <expno>, rew <expno> Open a different experiment number in the current/new window
- wrpa <expno> Copy current data set to a new experiment number

Expert Level:

- edasp Set up rf channels and nuclei

Useful Features:

- CTRL + \ (backslash) Restart display
- CTRL + ALT + BACKSPACE Log off your (or someone else's) NMR account