

Commonly Used VNMR Commands and Parameters:

The following is a **short** list of VNMR commands and parameters. User fluent in Bruker-speak will recognize the equivalent Bruker commands appearing in parenthesis.

- su **(ii)** Setup the system hardware to match the current parameters. *Does not* start the acquisition.
- acqi **(bsmsdisp)** Opens the acquisition interface window for interactive locking and shimming.
- ds Display spectrum in the Graphics Window
- dps Display pulse sequence in the Graphics Window
- dg Display the 'group' processing/ acquisition parameters
- flip Switches the Graphics Window or the Text Window to the front of the screen
- rt **(re)** Retrieve FID's from a file into the current experiment
(eg: rt('/directory path/filename'))
- svf **(wr)** Saves current parameters, text, and FID in the current experiment to a file. You will be prompted for a path and a filename. Alternatively you could specify both a path and a filename on the command line.
(eg: svf('/directory path/filename'))
- rtp **(rj)** Retrieves parameters from a file or FID into the current experiment. By default rtp will prompt you for a filename or you may add the filename and/ or path. Example: rtp('/directory/mydatafile')
- svp **(wj)** Saves parameters from the current experiment to a file. By default svp will prompt you for a filename or you may add the filename and/ or path. Example: svp('/directory/mydatafile')
- rts **(rsh)** Retrieves a shim file and loads the shim values into the current experiment

- **svs (wsh)** Save the current shim values to a shim file. By default you will be prompted for a filename or you may add the filename and/ or path as options.
- **e (ej)** Eject sample
- **i (ij)** Insert sample
- **tn** Changes the value of the transmitter nucleus. Examples: `tn='H1'` or `tn='C13'` etc. Can also be used to query the current status. Example: `tn?`
- **dn** Changes the value of the decoupler nucleus. Examples: `dn='H1'` or `dn='C13'` etc. Can also be used as a query.
- **tof (o1)** Controls the exact positioning of the transmitter. As the value of `tof` increases the transmitter moves to higher frequencies (moves toward the left).
- **dof (o2)** Controls the decoupler offset in the same manner as `tof`.
- **tpwr** Controls the transmitter power level (in dB). [Can be used as a command, i.e., `tpwr=30` or as a query, i.e., `tpwr?`](#)
- **dpwr** Work identical to `tpwr` for the decoupler channel.
- **dm** Determines state of the decoupler (decoupler mode) at different times during the pulse sequence. Typical options are 'y' and 'n'.
- **pw** Transmitter pulse width in ms. If you are receiving an ADC overflow error type `pw=pw/2 ga`. Repeat as necessary.
- **at (aq)** Length of each individual FID acquisition time.
- **sw (sw)** Sets the total width of the spectrum. To enter a value 200ppm type `sw=200p`
- **np (td)** sets number of data points to be acquired.
- **fn (si)** Fourier number.
- **nt (ns)** Set the number of tranients (or scans) the instrument will perform of make up a FID. Range is 1 to 1e9.

- ct Number of completed transients in the current experiment.
- bs Block size - determines how often the data in the current experiment is saved.
- lb Lorentzian line broadening
- time (expt) Calculate the time required for the experiment to complete based on the current parameters.
- array Array an experimental parameter. Array will prompt you for the parameter to array, the number of steps, the starting point, and the step size.
- da Displays arrayed acquisition parameters in the Text Window.