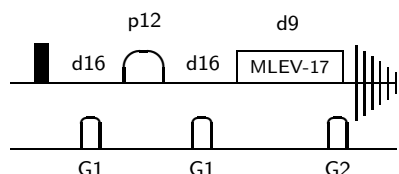


## Obtaining 1D-TOCSY Spectra

A 1D-TOCSY spectrum will show the peaks within the same spin-group as an irradiated peak and give a sense of the distance each is from the irradiated peak. As shown below, it involves a refocussing by a selective  $180^\circ$  pulse and a MLEV-17 spin-lock, during which the polarization migrates from the irradiated nucleus to the others within the spin system.



The gradient procedure below gives optimal results, but a non-gradient procedure is also given.

### Using Gradients

1. Acquire a reference spectrum and obtain the position of the peak to be irradiated, in Hz.
2. Create a new data set (edc).
3. rpar SELMLGP all to read in parameters. getprosol as usual.
4. Change the following parameters:

```
pulprog = selmlgp.2
01      = peak position (in Hz) noted in step 11
D9      = 40 ms (mixing time)
P12     = 30 ms (selective pulse length)
```

Type gpz and, in the resulting window, set the following gradient parameters:

```
gpz1    = 15
gpnam1  = SINE.100
gpz2    = 40
gpnam2  = SINE.100
```

5. Run and process the spectrum as usual. Adjust d9 as needed.

### Without Gradients

Similar to the setup above except:

- rpar SELMLZF1H (instead of SELMLGP)
- Do not change pulprog from the SELMLZF1H default
- Do not enter the gp... parameters

<sup>1</sup> Alternatively, leave 01 unchanged and set spoff to the difference between 01 and your measured value. (spoff2 for gradient sequence; spoff1 for non-grad)