

# Performing the 1D $^1\text{H}$ - $^{19}\text{F}$ NOE Experiment

This document describes how to set up a 1D, selective, steady-state NOE between  $^1\text{H}$  and  $^{19}\text{F}$  using the normal s2pul pulse sequence.

## A: Data Collection

- (1) In exp1, setup a normal  $^1\text{H}$  experiment, collect a  $^1\text{H}$  spectrum; probably save it
- (2) In exp2, setup a normal  $^{19}\text{F}$  experiment, collect a  $^{19}\text{F}$  spectrum; probably save it
- (3) Type **dn='F19'** on the command line
- (4) Put the cursor in the center of the  $^{19}\text{F}$  peak of interest and type **nl**
- (5) Type **sd** to give a dof value for  $^{19}\text{F}$ . **Write down the dof value.**
- (6) If you have more than one Fluorine peak that could potentially give the NOE to protons, repeat step 4 and 5 for all the peaks. **Write down all the dof values.**
- (7) Move the parameters in exp1 to exp3 by the commands: **mf(1,3) jexp3 wft**
- (8) In exp3, make sure  
**dn='F19'**  
**d1=2** (can be set longer e.g. 5s)  
**d2=5** (can be set longer e.g. 10s)  
**homo='y'**  
**gain='y'**  
**dm='nyn','nnn'**  
**dmm='ccc', dmf=200 dpwr=10**  
**dof**=the value found in exp2 (step 5 or 6 above)  
 Collect  $^1\text{H}$  spectrum with the desired nt (number of transients)
- (9) Type **da** to look at the arrayed values and **dg** to go back to the normal parameter display
- (10)  
 You can array **dpwr** value to find the best saturation power for obtaining NOE peaks (in this case, don't array dof, use just one dof value found above to avoid complications). Normally, dpwr=10 should be good enough and don't use a number larger than 10. Once you have a desired **dpwr** value, collect the selective NOE between  $^1\text{H}$  and  $^{19}\text{F}$  spectrum as in step (8).
- (11) Once you are done with the data collection, save the data which contains 2 spectra
- (12) If you have multiple  $^{19}\text{F}$  peaks and dof values, repeat step (7) to (11) in a different experiment window for instance, type **mf(1,4) jexp4 wft** (Don't use exp5).

## B: Data Processing

- (1) Process the arrayed data as usual (**proc** or **wft aph** or use manual phasing)
- (2) Type in the command line: **clradd ds(1) add ds(2) sub jexp5** (these macros delete experiment window 5 (exp5), and then create exp5, add spectrum #1 to exp5, then subtract the spectrum #2 from #1)
- (3) Go to exp5: **jexp5 wft** to see the resulting spectrum, only the  $^1\text{H}$  peaks which have NOE with the  $^{19}\text{F}$  signal show up, the other  $^1\text{H}$  peaks are subtracted to zero (note: there may be small residual peaks if the lower number of scans are used (more than 32 scans is recommended)).
- (4) You can save this resulting NOE spectrum as usual.
- (5) Note: above commands (**clradd ds(1) add ds(2) sub jexp5**) can be replaced by a macro by typing: **hfnoe** (vrx500 only)
- (6) Or you can process the NOE spectrum in Mnova using Arithmetic tool under Analysis menu