# NOESY1D-(DPFGSE NOE) Experiment for Measurement of Transient NOE's

NOESY1D is the Varian implementation of the 1D NOE experiment using pulsed field gradients and selective pulses developed by Shaka et al. At the heart of the experiment is a double-pulsed-field-gradient-spin-echo (DPFGSE) pulse sequence that achieves selective spin inversion effectively and cleanly.

It is important to recognize that NOESY1D is a <u>transient</u> NOE experiment, which differs from the traditional <u>steady-state</u> NOE experiment (cyclenoe on Varians). In a steady-state NOE experiment, like cyclenoe, a specific resonance (i.e. peak in the spectrum) is saturated by low power RF saturation for a designated time (sattime) and a percent increase (or decrease) in the spectrum relative to a reference spectrum is observed. During the saturation time, the NOE's will reach an equilibrium value. In contrast, NOESY1D employs a series of pulses and gradients to invert a particular spin, a mixing time to allow the NOE to develop, and a read pulse to obtain a NOE spectrum. Therefore, an equilibrium value for the NOE's is not obtained. NOE's obtained from transient NOE experiments are generally lower in intensity (38% max vs. 50% for small molecules) and their relative intensities are not directly comparable to steady-state NOE's.

An example of a steady-state cyclenoe and a NOESY1D experiment are shown below. With the same number of scans, cyclenoe gave 2 to 3 times the S/N. However,



because the cyclenoe experiment requires the saturation time to be greater than the longest 'H T<sub>1</sub>, the total time to acquire the spectrum was  $\sim$ 70 minutes for cyclenoe and  $\sim$ 11 minutes for NOESY1D. Furthermore, there are essentially no subtraction artifacts in

the NOESY1D spectrum while the cyclenoe spectrum has obvious artifacts (e.g. between 7 to 7.5 ppm and around 2 ppm). The combination of very low artifacts and faster per cycle repetition make the NOESY1D experiment a good choice for routine qualitative NOE determination.

Quantification of the % enhancement, which is a typical value reported for NOE's, is not readily obtained from NOESY1D. Since the target resonance is not saturated with NOESY1D but is inverted, there will be relaxation of that signal during the mixing time. Therefore, setting the inverted peak to 100% as is typical with traditional steady-state NOE's is not appropriate. Further experiments are necessary to obtain reliable quantitative numbers.

Comparison of Transient (NOESY1D) and Steady State NOE (cyclenoe) Experiments.

NOESY1D	cyclenoe
Optimum NOE when mixing time equals	Must have saturation time larger than
'H T <sub>1</sub> 's-quicker recycle time and more	longest 'H T <sub>1</sub> 's
scans per unit time	
% NOE is less than and not directly	Larger % NOE and better S/N for same
comparable to traditional steady-state	number of scans. Quantification is
techniques. Quantification is difficult	straightforward
Relatively insensitive to calibration	Careful calibration is recommended and a
errors and instrument instability	stable instrument is necessary
Essentially no subtraction or other	Subtraction artifacts are common and
artifacts, which allows for measurement	long range NOE's difficult to detect
of long range NOE's	
Setup is quick and straightforward	Setup is a little more complicated
Can generate NOE build-up curve for	Can not generate build-up curves
distance calculations	

\* \* \* \* \* \*

# **Explanation of Types of Commands Found in this Handout:**

1. The vnmr software and the UNIX operating system are both case sensitive. This means that the computer distinguishes whether the letters are entered in upper case (i.e. CAPITALS) or lower case. The user must be careful to type the correct case for each letter in a command.

### *EXAMPLE*: jexp1 is not the same as JEXP1

2. Some commands are line commands and are typed in by the user followed by a hitting the RETURN key.

# EXAMPLE: su

Hitting the RETURN key is assumed for all bold text.

3. Some commands are executed by clicking a mouse button with its pointer on a 'button' found on the screen. The execution of these commands is indicated by a two letter designation (LC {left click}, RC {right click}, or CC {center click}) followed by a word or words in bold that would appear in the 'button'.

# **EXAMPLE:** LC Main Menu

This means to click the left mouse button with its pointer on the 'button' that says "Main Menu".

4. Some commands are executed by the mouse itself. These commands are indicated by the two-letter designation (LC, RC, or CC) and a description of what the user should do in parentheses.

### *EXAMPLE*: LC (at 6 ppm)

This means to click the left mouse button with the mouse cursor at 6 ppm.

5. Parameters are entered by typing the parameter name followed by an equal sign, the value, and a return.

### EXAMPLE: nt=16 <rtn>

\* \* \* \* \* \* \*

# Running the NOESY1D Experiment

#### **Preliminaries:**

- It is recommended to degas your sample prior to running the experiment because dissolved oxygen will decrease the NOE. If necessary, you should also filter the sample.
- **Insert** and **Lock** as usual.
- Shim as usual.
- Since NOE's are temperature dependent, it is recommended to run the experiment at a slightly elevated temperature. Refer to VT operation handout for adjusting spectrometer temperature.

### **Creating the Selective Pulses:**

### jexp1

join experiment 1

Setup and run a standard 'H experiment.

(**OPTIONAL**): For optimal performance, you can determine your 90° proton pulse, and set pw90=*determined value*. See the handout for determination of the 90° proton pulse.

When the 'H experiment is completed:

move the 'H FID from exp1 to exp2
join experiment 2
Fourier transform FID
opens Experiment selector pane
sets up parameters for NOESY1D

Expand around first resonance to be inverted.

In the *Parameter Panel* (If not open, click <u>View</u>, **Parameter Panel**), click *Acquire*, *Defaults*. The display shouls look like the following:



# LC Box/Cursor

Selects Cursor mode

LC on spectrum to the LEFT of the region to be inverted, then RC (Right Click) to the RIGHT of the region to be inverted. Only invert one multiplet at a time. You should have two red cursors on the screen like the image above.

LC Select

selects current region between cursors for selective inversion

Repeat for all desired resonances. When completed,

If, at any point, you make an error, it is easiest to restart from **mf(1,2**).

set number of scans. MUST BE A
MULTIPLE OF 8
set mixing time. mix=500ms is the default
value and is optimal for short-range
interactions of small to medium-sized
molecules. If you are looking for long-range
interactions, mixing times comparable to the
'H T <sub>1</sub> 's can be used

Make sure the sample is not spinning as gradient selected experiments like NOESY1D do not work with spinning.

Click the green Acquire button in the top middle of the Parameter Panel. DO NOT use *ga* or *go*.

### Data Manipulation:

lb=1.0 wft

set line broadening to 1.5 Hz and Fourier transform your data

# vp=80 ds

move the spectrum up and display sepctrum

Phase the largest peak (this is the irradiated peak) so that it is negative (i.e. it points down). The small NOE peaks will be opposite phase to the irradiated peak.

### dssa

view all spectra in array

To view individual spectra, type ds(#), where # is the number of the desired spectrum. For example, I decided to do 5 different NOE experiments irradiating at 9, 6, 5, 3, and 1 ppm. I selected the peaks in the order given, which means that typing, ds(1), will give me the spectrum with irradiation at 9 ppm. Typing ds(4) would give me the spectrum with 3 ppm irradiation.

Integration and peak picking are done exactly like <sup>1</sup>H spectra. Remember that the integrated values are NOT directly comparable to traditional steady-state NOE's. It is best to use the values qualitatively.

# Printing your NOESY1D Spectra:

To print all spectra in the array:

dssa	look to see if the spectra overlap. If so, type <b>ds(1)</b> and reduce the scale of the spectrum. Type <b>dssa</b> . If necessary, readjust by typing <b>ds(1)</b> , adjusting scale, and typing <b>dssa</b> .
pl('all') pscale page	plots all spectra
To print individual spectra:	
ds(spectrum#)	for example, ds(1)
Phase, peak pick, and integrate as desired.	
pl pir pscale ppf page	print spectrum, scale, integrals, and peak

frequencies