## III. Performing <sup>1</sup>H T<sub>1</sub> Estimates

[updated: 13 July 2010]

- 1. Acquire a normal  ${}^{1}H$  1d spectrum, as described in Section I (sbs\_1H1d).
- 2. Calibrate and set **pw90**, as described in Section II (sbs\_1Hpw90).
- 3. The macro **T1est**, will automatically setup parameters as follows.

```
a) p1=pw90*2 d1=10 dps↓
```

manual checks:

```
set nt=1 ss=0 vp=80 d2=0.01 ↓
```

ga, J should show inverted magnetization everywhere.

**d2=10 ga**, should show positive magnetization everywhere, except perhaps for solvent (or other very slow relaxing protons) peaks.

Adjust d2 to give crossovers for important multiplets (see 4 below).

- b) Using an array for d2 is OK, but presents the problem of what to set d1 to
  - → d2=0.1,0.25,0.4,0.6,0.9,1.3,2.0,3.5 d1=10 go dsa is usually a good setup for most compounds, but beware that d1 may be too small, especially for samples sealed in non-O<sub>2</sub> containing atmospheres.

Acquire with: go, then dsa,

- 4. Note the crossovers for each important multiplet, and note these points as  $d_{null}$  times. Especially note the fastest and slowest relaxing protons of interest.
- 5.  $T_1 \approx 1.4 \times d_{null}$  for each proton. If  $T_1 > 1/3 \times d1 = 3$  sec, then d1 was too short during the experiment. Repeat with a longer d1 value.
- 6. See the  $T_I$  section in VUG for more information, especially how to obtain quantitative  $T_I$  values (rather than the estimates provided by  $d2_{null}$ ). A brief outline on how to analyze and plot quantitative data in VNMR is:
  - a) set **Th** (button on the **ds** menu) ;all peaks above line will be found by dpf
  - b) **dpf**, ;display peak frequencies
  - c) **fp**, ;writes list of peak amplitudes
  - d) t1, ;performs analysis and outputs T1 analysis
  - e) **expl**→ ;displays a plot of the T1 fits
- 7. To return to normal acquisitions, set:

T1off,⊣

or equivalently: p1=0 d2=0 vp=20 dps

 $\rightarrow$  useful related commands: da ff dc dg dssh dssl pl('all')