



KJM 9250

CarbonT₂ Spectra on the AVI600 and AVII Spectrometers

Version 5.0

Topspin 1.3 Windows XP AVI600

Topspin 2.1 Windows 7 AVII600



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April 2018

Carbon T₂ Spectra on the AVI-600 and AVII-600

1.0 Introduction

An **awcarbont2** parameter set and linked **VCLIST** file from which a **VDLIST** file *must* be calculated prior to processing a T₂ data set have been set up on the AVI-600 and AVII-600 spectrometers running under TS2.1 and TS3.2 respectively.

The parameter sets have **D1 = 10 sec** and 8 x **VCLIST** values. The T₂ data set is acquired using the smallest vclist value first. Prior to processing T₂ data sets variable constant (VC) values must be converted to time domain variable delay (VD) values using the **vctovd** command (see instructions 15 and 16.)

D1 and the longest **VD** time derived from the largest **VC value** should be 3-5 times the longest T₂ in the sample compound. T₂ is always shorter than T₁.

$$VD_{\max} = [P2 + (2 * D20)] * VC_{\max}$$

T₂ data sets can be processed on the spectrometer terminal or off line using any version of Topspin provided the linked **VCLIST** file is either resident in the experiment's top level folder (= default setting: see below) or recreated or copied into the off line terminal's *C:\Bruker\TopspinX.X\exp\stan\nmr\lists\vc* folder.

If required the **VCLIST** file that is present in the experiment's top level folder can be opened with **WordPad** and the **VC** values in it can be viewed and used to recreate the original named **VCLIST** file from which the **VDLIST** file can be created using the **vctovd** command.

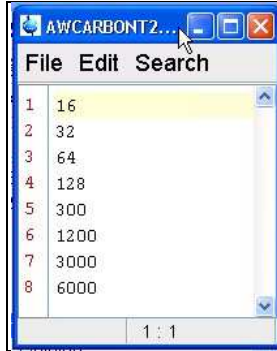
| | | | |
|------------|----------------------|-----------|-------|
| uxnmr.info | 2/01/2017 10:50 p... | INFO File | 4 KB |
| uxnmr.par | 2/01/2017 10:50 p... | PAR File | 24 KB |
| vclist | 8/02/2018 10:30 p... | File | 1 KB |

NB: The experimental copy of the variable constant file is named as **vclist** irrespective of the name of the source vclist file.

2.0 Experiment Set Up

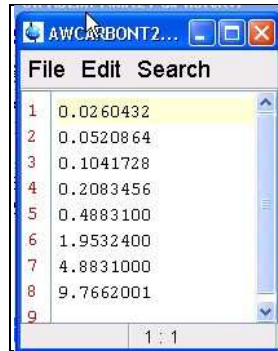
- 1) Create an experiment and read in the **awcarbont2** parameter set (+ **getprosol**).
Pulse programme = **awcpmgpg**
- 2) Review default settings. These settings can be adjusted if required.
TD(F2) = 32K, TD(F1) = 8
SW = 240 ppm, O1P = 110 ppm.
D1 = 10 sec.
D20 = 800 usec = 0.000800 sec
NS = multiple of 4 or 8, DS = 4 or 8.
- 3) Type **ased** (enter) and review other parameters used in the job including the linked **VCLIST** file = **AWCARBONT2-8**.

- 4) The **VCLIST** file should have the entries shown below. Do not alter the values in this file. An alternatively named **VCLIST** file should be created if different **VC** and **D1** values are required for a particular compound. The **VD** values calculated from the **AWCARBONT2-8** file's **VC** values using the **vctovd** command (see steps 15 and 16) when **P2** = (180 degree pulse) = **27.7 usec** on AVI-600 and **D20** = **800 usec** = 0.0008 sec are shown below.



| Line | Value |
|------|-------|
| 1 | 16 |
| 2 | 32 |
| 3 | 64 |
| 4 | 128 |
| 5 | 300 |
| 6 | 1200 |
| 7 | 3000 |
| 8 | 6000 |

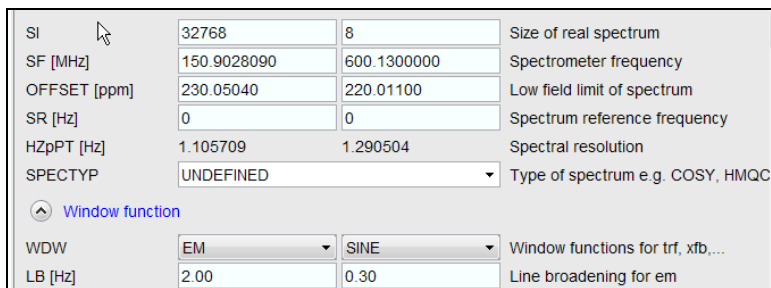
AWCARBONT2-8 VC file values



| Line | Value (sec) |
|------|-------------|
| 1 | 0.0260432 |
| 2 | 0.0520864 |
| 3 | 0.1041728 |
| 4 | 0.2083456 |
| 5 | 0.4883100 |
| 6 | 1.9532400 |
| 7 | 4.8831000 |
| 8 | 9.7662001 |

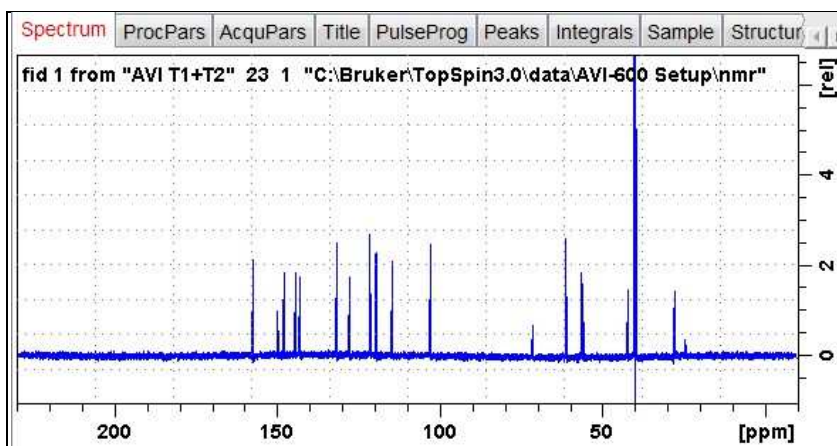
Calculated VD times (sec)

- 5) Set receiver gain using **RGA** (*important!*).
- 6) Type **edp** (enter) and check that **SI(F2) = 32K**, **SI(F1) = 8**, **WDW(F2) = EM**, **LB (F2) = 2 Hz** or other value of your choice.



| | | | |
|-----------------|-------------|-------------|-----------------------------------|
| SI | 32768 | 8 | Size of real spectrum |
| SF [MHz] | 150.9028090 | 600.1300000 | Spectrometer frequency |
| OFFSET [ppm] | 230.05040 | 220.01100 | Low field limit of spectrum |
| SR [Hz] | 0 | 0 | Spectrum reference frequency |
| HZpPT [Hz] | 1.105709 | 1.290504 | Spectral resolution |
| SPECTYP | UNDEFINED | | Type of spectrum e.g. COSY, HMQC |
| Window function | | | |
| WDW | EM | SINE | Window functions for trf, xfb,... |
| LB [Hz] | 2.00 | 0.30 | Line broadening for em |

- 7) Start the acquisition using the **ZG** command
- 8) When the experiment has run type **rser 1** (enter) to read in the first serial file which will appear in a TEMP screen display window.
- 9) Type **EFP** (return) to transform it and phase it as per a normal carbon spectrum.



- 10) Type **edp** (enter) and note the phase constants for this spectrum. As per AVI and AVII ^{13}C spectra a large negative PHC1 value will be require

| Phase correction | |
|------------------|----------|
| PHC0 [degrees] | -133.289 |
| PHC1 [degrees] | -603.435 |
| PH_mod | pk |

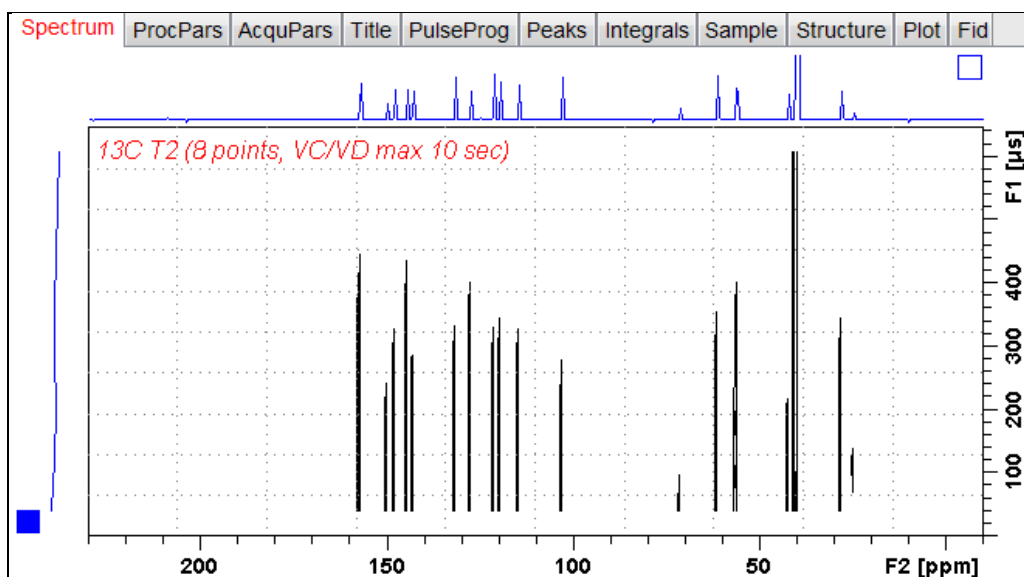
- 11) Close the **TEMP** window and reload the T₂ data set file.

- 12) Type **edp** (enter) and enter the phase constants noted in step 10 above into the **F2 PHC0** and **F2 PHC1** cells and check **PH_MOD = pk**. **F1** cell info is not used.

| Phase correction | | | |
|------------------|----------|----|----------------------|
| PHC0 [degrees] | -133.289 | 0 | 0th order correction |
| PHC1 [degrees] | -603.435 | 0 | 1st order correction |
| PH_mod | pk | mc | Phasing modes for t |

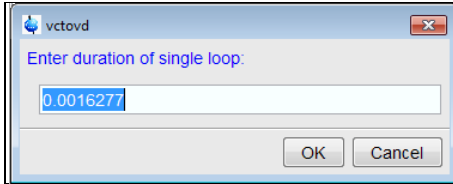
- 13) Type **xf2** (return) to transform the 2D data set followed by **abs2** (return) to baseline smooth it.

- 14) Provided phase constants have been correctly set up the transformed data set plot should resemble that shown below. Signals intensities go from their maximum positive value to zero intensity.

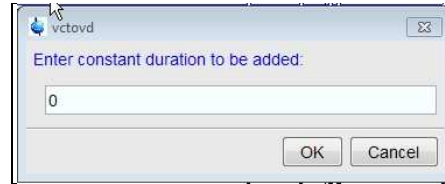


3.0 T₂ Data Set Processing

- 15) Prior to processing a T₂ data set it is essential that variable constant (VC) values are converted to time domain variable delay (VD) values.
- 16) Type **vctovd** (enter) and click OK on the display panels that appear.

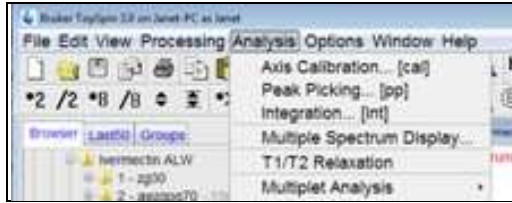


Single VC loop value = $p2 + 2 \times d20$

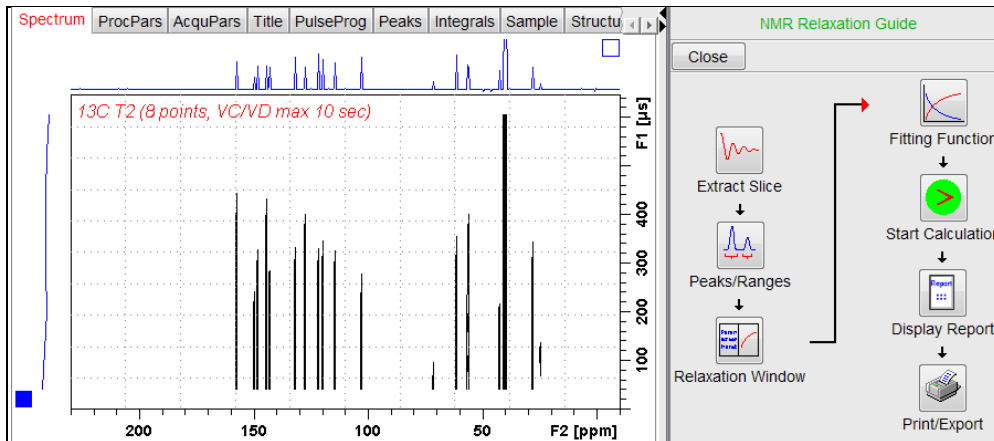


Not used. .

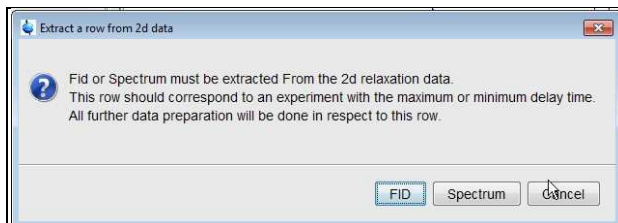
- 17) Open Topspin's **Analysis** menu and click its **T1/T2 Relaxation** tab. If other Bruker processing software has been installed on the spectrometer terminal, as may be the case on the AVII-600, select the **Analysis** menu's **Topspin T1/T2 module** tab and open its **T1/T2 Relaxation** sub-menu tab.



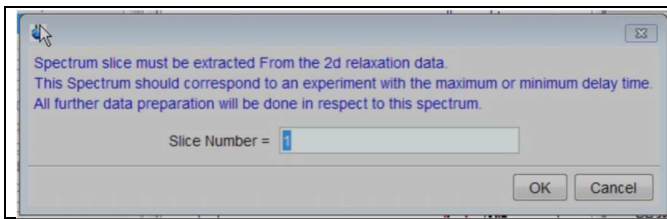
This will open up the screen display shown below.



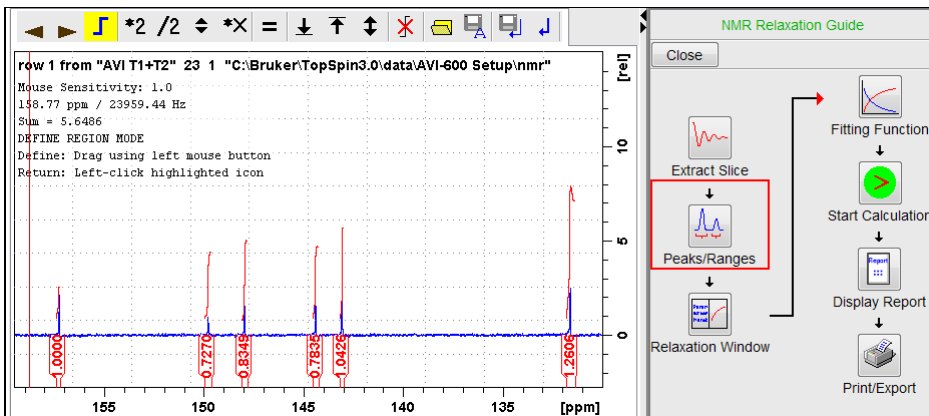
- 18) Click the **Extract Slice** button and then the click **Spectrum** button in the panel that appears.



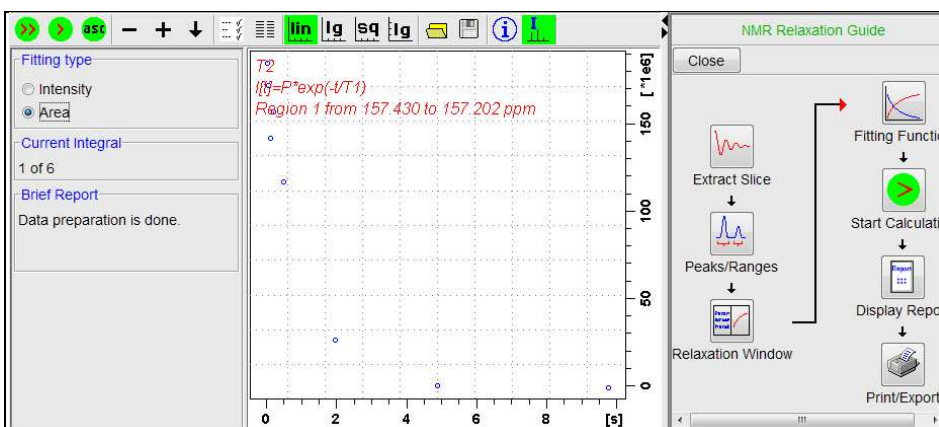
- 19) Enter **1** in the **Slice Number** cell and click the **OK** button. This will display the transformed spectrum ex the first (shortest) **VCLIST** → **VDLIST** value = the one that was phased via the **rser 1** routine in steps 8 and 9.



- 20) Expand the spectrum that appears in the usual way and integrate selected peaks in it. Integrals should start and terminate as close as possible to the edge of a peak.
- 21) Click the **"Save Region As"** icon button (= 3rd from the right in TS3.2's the upper menu bar, or towards the left hand side of the menu bar in TS2.1) = *the one with the floppy disk icon + A below it*) and then its **Export Regions to Relaxation Module and .ret. tab.**

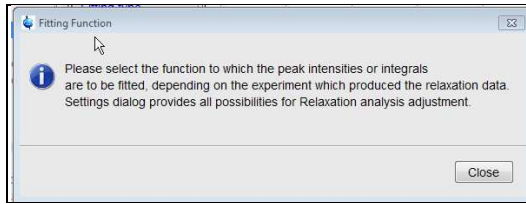


- 22) Click the **Relaxation Window** button and select **Intensity** or **Area** in the plot window that appears. **Area** is the best choice for carbon T₂ values.

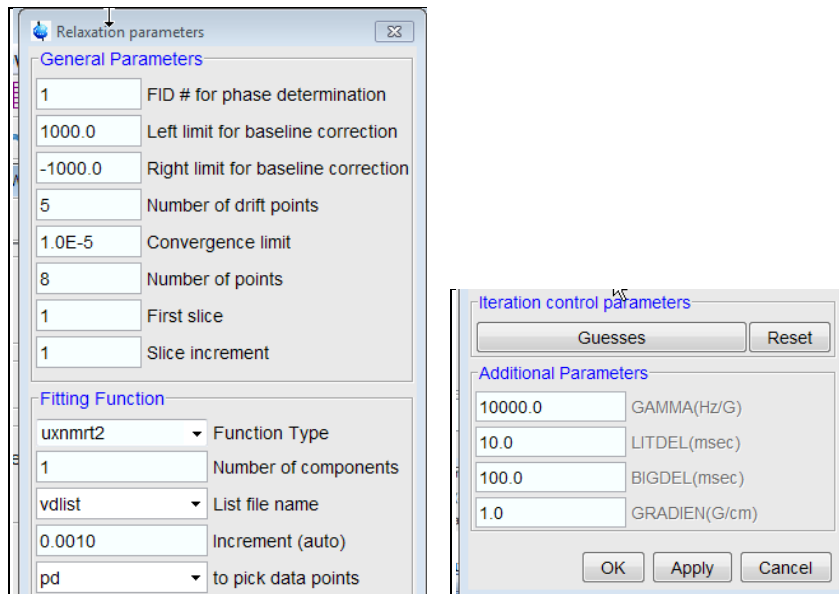


The plot's upper title line correctly has T₂, but the T₁ equation incorrectly appears below it. This is a long standing Topspin bug.

- 23) Click the **Fitting Function** button and note the comments about **intensity** or **area** options in the screen display that appears and close it.

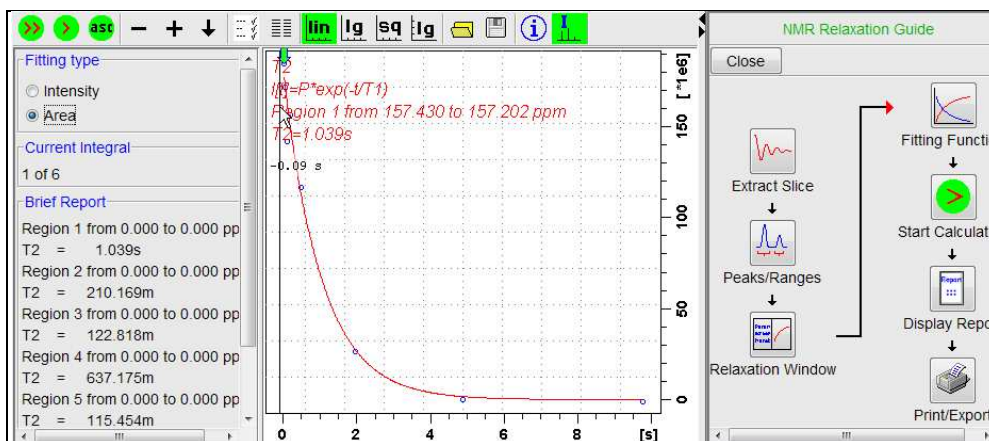


- 24) When the Fitting Function message screen is closed a panel with relaxation parameters (split into two screen captures below) will appear.



- 25) Check **Function Type = uxnmrt2** (*important!*) and **List file name = vdlist**. Other cells/values can be left as they are (= default settings). Click **Apply** and **OK** and close the window.

- 26) Click the **double red arrow** in the menu bar at the top left of the plot window and **NOT** the single red Start Calculation arrow button in the NMR Relaxation Guide menu below the Fitting Function button.



The appearance of the T_1 equation in this screen is a long standing Topspin error.

- 27) The + and - buttons in the upper menu bar can be used move through the series of T₂ plots.
- 28) Fitting type (**Intensity** or **Area**) can be changed in the plot display. If this is done clicking the **double red arrow** will recalculate the T₂ results and update their plots.
- 29) The NMR Relaxation Guide has buttons which can be used to display and/or print T₂ results. A sample report for one signal is shown below. Zero value points (= completely T₂ relaxed) or negative artifact peaks will be eliminated.

```

1 Dataset :
2 C:\Bruker\TopSpin3.0\data\AVI-600 Setup\rnmr\AVI T1+T2/23/pdata/1
3 AREA fit :
4 I[t]= P*exp(-t/T2)
5
6 8 points for Integral 1, Integral Region from 0.000 to 0.000 ppm
7 Results      Comp. 1
8
9 P      =    9.816e-001
10 T2     =     1.039s
11 SD     =    4.819e-002
12
13      tau      ppm      integral      intensity
14
15      26.043m  157.226  1.8464e+008  4.3523e+007
16      52.086m  157.225  1.7182e+008  4.3054e+007
17      104.173m 157.224  1.4193e+008  4.2111e+007
18      208.346m 157.226  1.5655e+008  3.6103e+007
19      488.310m 157.226  1.1662e+008  2.8708e+007
20      1.953s   157.225  2.6262e+007  8.0794e+006
21      4.883s   157.232 -2.8243e+005  eliminated
22      9.766s   157.232 -1.0611e+006  eliminated
23

```