#### 1. Introduction

The time constant  $T_2$  is the transverse or spin-spin relaxation time. Any process that causes loss of magnetization on the x-y plane contributes to  $T_2$ . The magnetization follows an exponential decay:

$$\mathbf{M} = \mathbf{M}_0 \mathbf{e}^{-t/T_2} \tag{1}$$

However, the inhomogeneity of the static field also causes a loss of the transverse magnetization, which is written as  $T_2^*$ . Since spin echoes remove the effects of inhomogeneity, the time constant that results from the decay envelope of a series of spin echoes approachs the real  $T_2$ . The classic experiment for this purpose is the Carr-Purcell-Meiboom-Gill (CPMG) experiment.

(Method 2)

## 2. Pulse sequence

In general, data acquisition time required for spectral resolution is longer then the delay between two  $\pi$  pulses required to cancel inhomogenity. Method 1 is only suitable for samples that result in one peak. Figure 2 shows a more general approach to implement CPMG. Again, a series of echoes is generated after the  $90^{\circ}$  pulse, but the signal is acquired only at the center of the last echo. The number of loops and thus number of echoes is given in a 2D table. Consequently, the specta with different numbers of echoes are stored as rows in the 2D matrix.

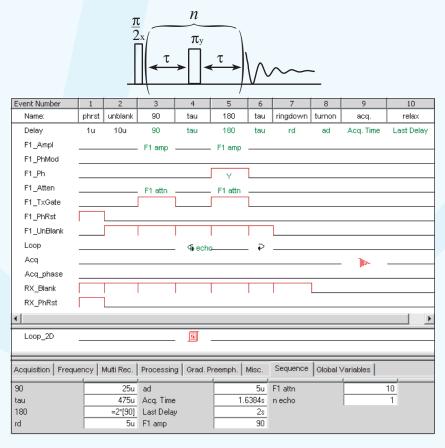


Fig. 3 The CPMG pulse sequence for Method 2.

# T<sub>2</sub> Measurements (II)

## 3. Experiment and results

Sample: dibromopropionic acid in  $D_6$ -benzene  $90^{\circ}$  pulse =  $25 \mu s$ 

 $\tau = 475 \,\mu s$ 

Max. number of echoes: 74

#### Data procesing:

- 1. Locate the cursor on the peak for  $T_2$  calculation.
- 2. Open the NTNMR data Analysis window, and select "Auto" mode to input "X Values".
- 3. Select "Real" for the real part of the FT data and "Intensity" as "Y Values", and click "Draw".

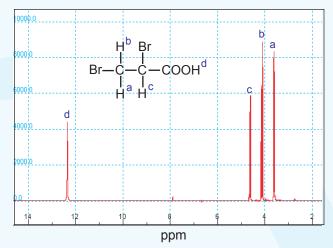


Fig. 2 <sup>1</sup>H NMR spectrum of dibromopropionic acid.

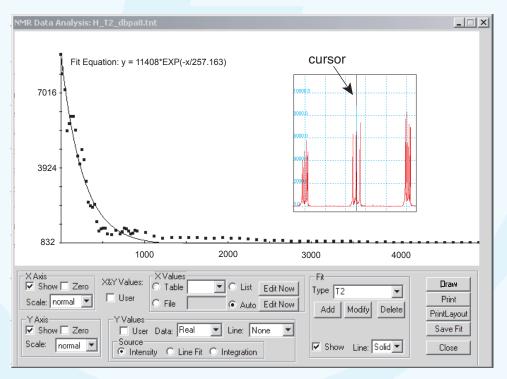


Fig. 3 NMR data Analysis window showing the  $T_2$  fitting for dibromopropionic acid and  $T_2 = 258$  ms. Insertion: Part of the spectrum of dibromopropionic acid. The cursor is located at a peak, and its intensity is used for  $T_2$  fitting.

## 4. Reference

1. Derome, A., "Modern NMR Techniques for Chemistry Research", Pergamon Press, New York, 1987.

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