## Introduction to 1D and 2D NMR Spectroscopy

(3) Diffusion experiments

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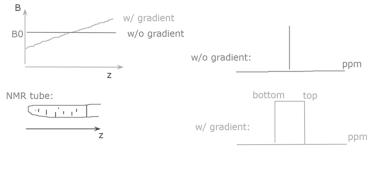
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### Objectives of this section

- Work with gradient pulses
- Work with a more complex experiment
- Learn ways to obtain high quality NMR data
  - Proper choice of acquisition parameters
  - Very careful phase and baseline corrections
- Learn a method to probe a physical attribute of your molecule/assembly

# A Simple MRI



 $\omega = \gamma B$ 

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Pulsed Field Gradients (PFG) in Modern NMR

- Gradient is nothing but the "evil" that you are trying to get rid of during shimming.
- Gradient pulses: magnetic field gradients applied for short durations, usually 0.5 to 2.5 ms; used along with RF pulses.
- Applications:
  - MRI
  - Topshim: automated shimming by applying pulsed gradients
  - Remove artifacts in many modern 2D experiments
  - Detect diffusion of molecules/assemblies

### **Diffusion NMR**

- NMR can detect the diffusion coefficients of molecules
  - The spatial location of the molecule is detected by applying a magnetic field gradient (<u>Use Larmor Equation to explain why</u>)
- Don't be overwhelmed by names these are all the same thing:
  - DOSY (Diffusion Ordered SpectroscopY)
  - PFG (Pulsed Field Gradient) NMR
  - PGSE (Pulsed Gradient Stimulated Echo) NMR
  - Diffusion NMR
- Good at detecting diffusion of small molecules and aggregates (R<sub>H</sub> <= 20 nm)</li>
  - Complementary with dynamic light scattering (DLS)

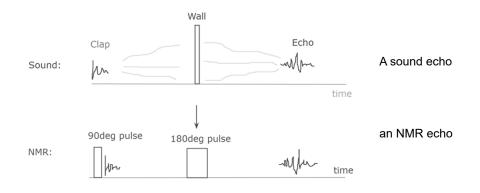
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### Principle of Diffusion NMR

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- Gradient pulses
  - Positive and negative gradients
  - Marks the spatial location of molecules
- Radio frequency (RF) pulses
  - The "echo" technique
  - "Wraps up" the spatial location information so that molecules can carry it to different locations by diffusion

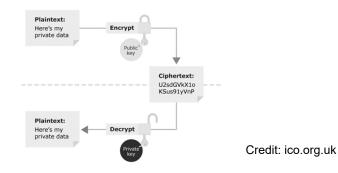




 A wall turns a sound wave by 180°, just like what a 180° pulse does to NMR signal



- Echo: a signal that is scrambled in the middle but reassembled at a later point
- We can use echoes to carry certain information

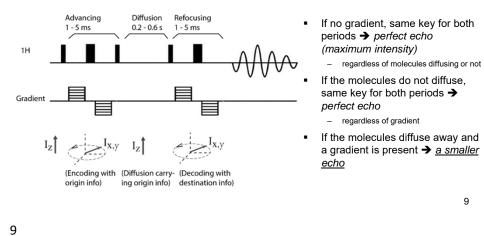


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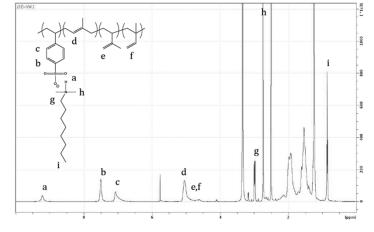
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#### Principle of Diffusion NMR

- Advancing period: a coherence is <u>scrambled/encrypted</u> with location info as key
   Location is defined by applied gradient
- The encrypted signal is then flipped back to z direction (by the 2<sup>nd</sup> 90deg pulse) to allow diffusion to occur
- Refocusing period: the coherence is <u>decrypted</u> with location info as key

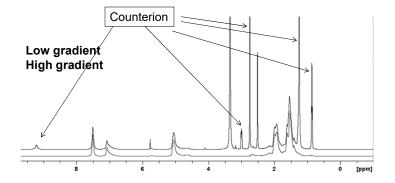


Is the Ionic Complexation Tight or Loose?



Solvent: DMSO

#### **Diffusion Behavior Can Tell!**



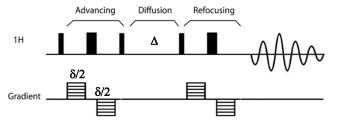
- Counterion diffuses much faster than the polymer
- The ionic association is quite loose even in DMSO
- You can easily tell whether a peak belongs to a large or a small molecule using the same technique

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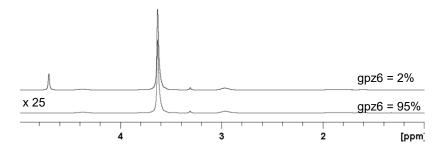
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### How to Run DOSY

- Parameters to adjust/optimize before you run
  - Calibration of 90° pulse length.
  - Gradient pulse duration ("small delta,  $\delta$ "): usually 500-2500  $\mu$ s
  - Diffusion time ("big delta, ∆"): usually 0.2-0.6 s
  - Longer gradient pulse and longer diffusion time → stronger attenuation
- Trial 1D runs with minimum (2-5%) and maximum (95%) gradient power
   Goal: adjust small delta and big delta such that A<sub>max</sub>/A<sub>min</sub> = 1 5%
- Parameter to increment (variable) during the run
  - Gradient pulse amplitude (in unit of percentage; 100% is the maximum), to increment between min and max



#### Example: A 1D Trial for PEO in D<sub>2</sub>O



- PEO signal decays to ca. 4% of original at high gradient
- Water signal decays much faster than PEO
- Proper choice of a decay range for your experiments:
  - A single component decay: 1 1.5 decades ( $A_{max}/A_{min} = 1 5\%$ )
  - A multi-component decay: 2 3 decades (A<sub>max</sub>/A<sub>min</sub> = 0.1 1%)
    - e.g. a polydisperse polymer

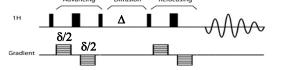
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### How to Choose Your Parameters

- Recycle delay (d1)
  - For quantitative 1D studies, should be 4 5 x T1
  - For diffusion studies, 3 x T1 is probably good enough
- Gradient Pulse duration (small delta)
  - Should be always <= 2500 μs</li>
  - If you set it too long (> 2500 µs) by mistake, you might burn the amplifier!
- Diffusion time (big delta)
  - T1 relaxation occurs during diffusion period, decaying from full amp to 0
  - If diffusion time is set too long, you will get less signal due to T<sub>1</sub> relaxation
    - Big delta < ½ x T<sub>1</sub> is recommended
- Usually:
  - First, use default small delta (p30) and adjust big delta (d20) between 0.05 s and 0.5 s to achieve desired decay range
  - If desired decay range cannot be achieved, increase p30 to a max of 2500 us



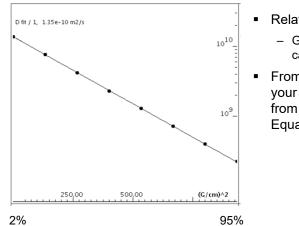
## **Data Processing**

• For a monodisperse object:

$$I = I_0 e^{-D\gamma^2 g^2 \delta^2 (\Delta - \delta/3)}$$

- I: peak intensity;
- D: diffusion coefficient;
- g: gradient amplitude
- Log(I) vs g<sup>2</sup> curve would be a straight line for a single component

Example: A Monodisperse Polystyrene



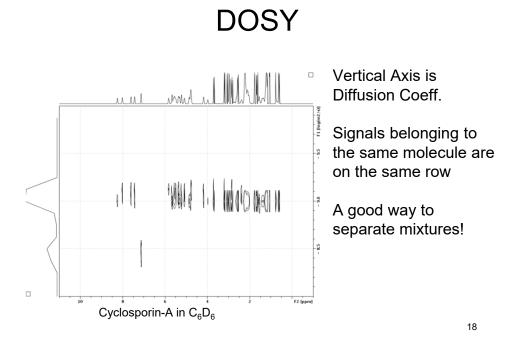
- Relative stdev for this fit < 1%</li>
  - Generally, relative stdev of < 2% can be easily achieved
- From D, hydrodynamic radius of your object can be calculated from the Stokes-Einstein Equation:

$$D = \frac{k_B T}{6\pi \eta r}$$

### Convection

- gives the same effect to signal intensity as diffusion
- Factors that contribute to convection:
  - Low viscosity (acetone, chloroform)
  - higher T
  - wider tube cross section
  - Long sample height (which results in high temperature gradient)
    2.5cm sample is recommended for CDCl3 or acetone-d6
- Sign of convection:
  - increasing D with longer big delta
  - Polymers and large particles seem to "diffuse" as fast as small molecules

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# **Transient Aggregation**

PCBM aggregates in 7.8 solution? 7.6 Diffusion NMR curve fit only finds one component 7.2 R<sub>H</sub> decreases with increasing T, why? Likely: PCBM spends time in both monomeric and aggregated states, and 6.4 NMR detects a weighted 250 260 270 280 290 300 average Temperature (K)

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## **Applications**

- Measure D (and thus R<sub>H</sub> (hydrodynamic radius)) of your target
  - Each peak on your <sup>1</sup>H spectrum can get its own D
- Determine molecular weight of polymers
  - Molecular weight distribution is possible
  - Any solvent is OK
- Separate mixture and remove impurity signals
- Probe interactions in solution: complexation; aggregation; dimerization; etc.
- Study single-molecule/assembly dynamic equilibrium