Week 5

Magnetic Resonance Spectroscopy

5.1 Lecture 8: Magnetic Resonance Spectroscopy

- 1/31: Refer to Chapter 14 of McQuarrie and Simon [1].
 - There is both NMR (nuclear magnetic resonance) and ESR (electron spin resonance) or EPR (electron paramagnetic resonance).
 - The last two are the same thing.
 - Two fields: The static magnetic field, and the probing electromagnetic field.
 - Derivation of quantized angular momentum.
 - In molecules, there is a multiplicity/degeneracy of states that grows as 2J+1. They have quantized angular momentum.
 - So is the orbital angular momentum of electrons!
 - Putting atoms into a magnetic field *creates* the anisotropy necessary for discussing the z-component (or any coordinate component) of angular momentum.
 - Spin angular momentum: Just means that the objects (e.g., electrons and nuclei) have a property that looks a lot like spin and/or angular momentum.
 - We say that each nucleon has a spin of 1/2. Protons and neutrons add separately.
 - Even number of protons and neutrons? Spin 0.
 - Mixed even/odd? We have actual nuclear spin.
 - ➤ We need a nucleus like this to detect!
 - Odd/odd? We have 0 spin again.
 - We focus on spin 1/2 particles. These have two degenerate energy states that split in a magnetic field.
 - Classical picture of spin angular momentum.
 - Picture a charged particle with angular momentum. The circulating charge produced a magnetic field which aligns along the direction of the angular momentum.
 - Indeed, a spinning charged particle behaves like a dipole.
 - $-\gamma = 2mc$ is the gyromagnetic ratio.
 - Quantum spin angular momentum.
 - Basically the same thing; we just rephrase everything from before in the language of operators.
 - **Zeeman effect**: Two energy levels split with increasing B.

- Larmor frequency: The frequency $\nu = \gamma B/2\pi$ in the radio frequency range that induces a shift.
- Typical operating conditions.
 - NMR vs. ESR: NMR has a stronger magnetic field, longer EM excitation radio waves, and significantly lower gyromagnetic ratios.
 - There is only a tiny difference between nuclear state occupation at room temperature; hence supercooling to get something detectable.
- What is the magnetic dipole doing in the magnetic field?
 - No constraints on the x and y components of I.
 - A dipole in a field experiences a torque.
 - Dipole precesses around B at the Larmor frequency ν .
 - FT-NMR spectrometers use pulsed rf fields to synchronize and detect the procession of spins.
- Lots of good extension material on NMR; also worth rewatching at some point!

5.2 Lecture 9: Magnetic Resonance Spectroscopy 2

- 2/2: Summary of last time.
 - The quantity that we're measuring is spin angular momentum \bar{I} , which is a vector quantity.

$$|\bar{I}| = \hbar \sqrt{I(I+1)}$$

where I = 1/2 is the nuclear spin quantum number.

- The other quantity of concern is the projection $I_z = m_I \hbar$ where $m_i = \pm 1/2$.
- In a magnetic field, we break degeneracy, getting $E(m_I) = -\gamma_N \hbar m_I B$ and $\Delta E = -\gamma_N \hbar B$.
- Electromagnetic resonance is achieved when the frequency ν of incident radiation satisfies $h\nu=\Delta E$.
- The interest in chemistry: Chemical shift.
 - There are small variations in the frequency for different types of protons depending on the surrounding electron density.
 - Measured frequency depends on effective magnetic field.
 - Shielding: The influence of electrons around the nucleus on the effective magnetic field.
 - The effective field is smaller than the applied field.
 - Shielding *decreases* the splitting (this is why nearby highly polar groups lead to large shifts, while alkanes have small shifts).
- Measuring the chemical shift.
 - We measure the difference in the Larmor frequency relative to a standard (TMS).
 - Shielding is a small effect (on the order of 10^{-6} , so we use ppm δ).
 - Example: 1 ppm at 500 MHz is $\nu = 500$ Hz, which is tiny (on the order of microjoules).
- Chemical shift charts (from OChem) are included in the slides.
- FT-NMR spectrometers.
 - How do we make these measurements?
 - NMR spectrometers are almost all working in FT mode these days.

- They use pulsed radiofrequency (r.f.) fields and detect the precession of spins.
- Precession of one spin in a magnetic field occurs at the Larmor frequency.
- Applying an excitation field creates a superposition of $m_s = \pm 1/2$ states. The net dipole is now perpendicular, and precesses that way (i.e., in the xy plane) in a mathematically describable fashion.
- Putting your superconducting coil along the x- or y-axis allows you to detect changes.
- Magnetization: The macroscopic alignment of magnetic dipoles $\bar{M} = \sum \bar{\mu}$.
 - At equilibrium, \bar{M} aligns along \bar{B} .
 - An rf field rotates magnetization to x: This is a $\pi/2$ -pulse or a 90°-pulse.
 - We detect precessing magnetization and return to equilibrium during the free-induction decay.
 An FT of the decay then generates our spectrum.
- Relaxation mechanism.
 - Spin state lifetime (T_1) .
 - "Spin-lattice" or "longitudinal" relaxation.
 - \blacksquare Recovery of the magnetization along z.
 - A molecular property.
 - Transfer of energy to the environment.
 - The return of magnetization to equilibrium has a characteristic time constant T_1 which appears in the time vs. relaxation plot $1 e^{-t/T_1}$.
 - Dephasing (T_2) .
 - "Transverse" relaxation.
 - \blacksquare Loss of magnetization in the xy-plane of many different sources.
 - We have the loss described by e^{-t/T_2} .
 - These two processes are not independent.
- There are numerous types of NMR experiments.
 - In our lab, we just scratch the surface.
 - We use a population inversion to measure T_1 .
 - \blacksquare π -pulse to invert magnetization.
 - Wait for relaxation.
 - Read out following another $\pi/2$ -pulse.
 - Two-dimensional spectroscopy.
 - Heteronuclear single quantum coherence spectroscopy (HSQC).
 - Excite ¹H; transfer its magnetization to ¹³C, which is nice because ¹³C is hard to excite on its own.
 - Transfer back to ¹H and detect.
 - Tells us which protons transfer magnetization.

5.3 Lab 3: NMR.

Lab Manual

- Modern NMR is performed in the time-domain, and frequency-domain spectra are obtained via Fourier Transformation of the data.
- Goal: Apply FT-NMR to investigate relaxation processes in liquids.

- Use an NMR technique called inversion recovery to determine the spin-lattice relaxation times T_1 for 13 C in n-hexanol, hexanoic acid, hexylamine, hexane thiol, or bromohexane.
- NMR basics (misc. notes).
 - Review from Chapter 14 of McQuarrie and Simon [1].
 - Definitions of the **Zeeman Hamiltonian** and **Zeeman levels** are new.
 - $-\Delta E = -\gamma \hbar B_0$ gives only a static picture of nuclear interaction with the magnetic field. In reality, a single nucleus remains in a certain state no longer than a time T_1 on average.
 - A nuclear spin exposed to a magnetic field B_0 tends to "align" with said field. It's not so much aligning, though, as it is acting like a gyroscope in a gravitational field; indeed, it will precess about the direction of B_0 with a characteristic frequency ω .
 - To rigorously treat the populations of the ground and excited spin states, we consider an ensemble
 of nuclei and the resultant thermodynamic argument.
 - The **total magnetization** takes the form of a vector aligned in the direction of B_0 (which is Z; there is no net magnetization in the XY-plane).
 - An alternating transmitter field B_1 "tips" the magnet from its equilibrium position by driving the Larmor procession. The angle α through which \mathbf{M} is tipped depends, via the following relation, on the strength of B_1 and the length of time t_p during which B_1 is applied.

$$\alpha = -\gamma B_1 t_p$$

- Immediately following this process, the transverse components M_X , M_Y decay to zero with a time constant T_2 , and the longitudinal component M_Z is restored to its equilibrium value with time constant T_1 .
- As with general time constants (which tend to be exponential factors), it will take much longer than T_1 for equilibrium to be "restored."
- **Zeeman Hamiltonian**: The Hamiltonian operator relevant to nuclear magnetic spins. *Denoted by* $\mathcal{H}_{\mathbf{Z}}$. Given by

$$\mathcal{H}_Z = -\mu \cdot B_0 = -B_0 I \gamma \hbar$$

- **Zeeman levels**: The 2I+1 energy levels $E(m)=-B_0\gamma\hbar m$, where $m=-I,-I+1,\ldots,I$.
- Spin-lattice relaxation time: The average time for which a single nucleus remains in a certain state. Denoted by T_1 .
- Spin-spin relaxation time: A type of interaction that occurs between different spins. Denoted by
- Larmor frequency: The characteristic frequency of a given nucleus at which it processes in a magnetic field. Denoted by ω .
- Total magnetization: The vector sum of all the magnetic moments. Denoted by M.
- FT-NMR (pulsed NMR).
 - Experiment: A strong transmitter field B_1 is applied for a short time t_n .
 - Specifically, since the transverse magnetization has its maximum when $\alpha = 90^{\circ}$, we want to choose t_p, B_1 such that $\gamma B_1 \cdot t_p = 90^{\circ}$. This is called a **90° pulse**. t_p is on the order of a few microseconds in general.
 - After the pulse, the relaxation contains spectroscopic information: The transverse magnetization will decay exponentially to zero due to spin relaxation in a manner that can be picked up by the receiver as a free induction decay.

- Differing transmitter and Larmor frequencies: M_Y and B_1 interfere, yielding a sine wave with exponentially decreasing amplitude:

$$M_Y(t) = M_{Y_0} \cos[(\omega_0 - \omega_1)t]e^{-t/T_2}$$

lacksquare M_{Y_0} is the transverse magnetization immediately after the pulse. It is given by

$$M_{Y_0} = M_{Z_0} \cos \alpha = M_{Z_0} \cos(\gamma B_1 t_p)$$

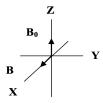
where M_{Z_0} is the equilibrium magnetization in the Z-direction.

- In general, a molecule will have multiple chemical environments, each with its own Larmor frequency. All of these will affect the FID, so we will need to use an FT to separate the different contributing frequencies.
- For more on pulsed NMR, see Bloch [2].
- 90° pulse: An application of a strong transmitter field B_1 over a time t_p sufficiently short such that $-\gamma B_1 t_p = \alpha = 90^\circ$.
- Free induction decay: A decay that occurs in the absence of an RF-field. Also known as FID.
- Chemical shifts and shielding.
 - Review from Chapter 14 of McQuarrie and Simon [1].
- ¹H vs. ¹³C.
- Signal processing considerations.
 - Real + imaginary parts of the FT data.
- Locking.
 - Making sure that the magnetic field is what the machine thinks it is.
 - This is like calibration, and we use a substance with a single sharp known NMR line (typically deuterium ²H) to do the locking.
- Spin decoupling.
 - Using a third magnetic field B_2 to resonate the nuclei to be decoupled.
 - We will use this technique to spin-decouple ¹H and ¹³C, greatly simplifying the latter spectrum.
- Spin-lattice relaxation.
 - $-T_1$ is the "lifetime" of the first-order rate process that returns the magnetization to the Boltzmann equilibrium along the +Z-axis.
 - The spin-lattice relaxation rate depends on the strength of intramolecular interactions and molecular motion.
 - Molecular mobility can be quantified by a **correlation time**.
 - Mechanisms involved in relaxation: Dipolar coupling, quadrupolar coupling, paramagnetic interaction, scalar coupling, chemical shift anisotropy, and spin rotation.
 - Protonated carbons: Dipole-dipole interactions with the attached protons are overwhelmingly dominant.
- Spin-lattice relaxation rate: The reciprocal of the spin-lattice relaxation time T_1 . Given by

- Correlation time: The time it takes for a molecule (or a molecular fragment) to reorient by a unit angle. Denoted by τ_C .
- Dipole-dipole coupling.
 - Depends on the strength of dipolar coupling, orientation of the interacting nuclei, distance between the interacting nuclei, and molecular motion.
 - Relaxation rate for a protonated carbon.

$$\frac{1}{T_1} = n \left(\frac{\mu_0}{4\pi}\right)^2 \frac{\hbar^2 \gamma_{\mathrm{C}}^2 \gamma_{\mathrm{H}}^2}{r_{\mathrm{CH}}^6} \tau_{\mathrm{C}}$$

- \blacksquare μ_0 is the permeability of a vacuum.
- \blacksquare γ_i is the gyromagnetic ratio of atom i ($i = {}^{13}C, {}^{1}H$).
- \blacksquare *n* is the number of bonded hydrogens.
- $r_{\rm CH}$ is the average C-H bond distance.
- The above equation only applies in the **extreme narrowing limit**.
- Extreme narrowing limit: The case where $1/\tau_{\rm C}$ is much greater than the resonance frequency.
 - Holds in liquids of low viscosity.
- Inversion recovery technique.



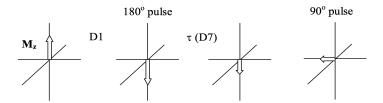


Figure 5.1: NMR inversion recovery.

- Use a multi-pulse NMR technique: Delay \to 180° pulse \to delay $\tau \to 90$ ° pulse \to acquisition (FID).
- This has the effect depicted in Figure 5.1.
- The decay of magnetization in the +Z-direction is

$$\frac{\mathrm{d}M_Z}{\mathrm{d}t} = -\frac{M_Z - M_{Z_0}}{T_1}$$

- Integrating the above yields

$$M_Z = M_0(1 - 2e^{-\tau/T_1})$$

which can be used to calculate T_1 .

- Implications of the above equation.
 - After a delay of T_1 , 63% of the magnetization is recovered along the +Z-axis.
 - To recover 99% of the magnetization, a delay of at least $5T_1$ is needed.

In-Lab Notes

- To start, create a dataset with proton NMR in CDCl₃. All of our ¹H NMR experiments will go into this dataset for the rest of the day.
- Create a dataset for carbon NMR as well.
- Any time you approach the instrument, do a mental metal check.
- Remove the dust cap, eject any previous samples, prepare yours, and sit it to float on top. Now click "Insert sample manually."
- Lock the machine onto the CDCl₃ deuterium peak. There are tons of solvents to choose from, among them fruit juice (for food chemists) and blood plasma (for biologists). Once it starts hovering in a certain region, you're good.
- We now do tuning. The tuner is made of adjustable capacitors, and it calibrates the receiver.
- We now do shimming. Think carpentry: Making everything level. Here, we introduce a homogeneous field to correct for any inhomogeneities.
- Sarah has a sample tuner and shimmer for us to hold.
 - They are very old devices from a previous, now disassembled NMR.
- Recall what's showing now from Tokmakoff's lectures.
- Control suite with some more adjustments.
- That's it for proton NMR. We now just collect our spectrum.
- Acquisition finished: See bottom left-hand corner.
- We now move on to data analysis. We can follow along with which scan we're on using the info in the bottom-left hand corner.
- Internal standard: TMS. Was developed here at UChi!
- Moving onto carbon NMR.
- We'll use 8 scans every time, though more would be better (96 is standard).
- Dummy scans are magnetic scans without data collection. They bring your sample back to a good place for collection after all of the calibration (24 is standard).
- \bullet After the scan, go to Process > Process Spectrum.
- We do see 6 peaks, as expected.
- As part of the short report, you'll need to assign each ¹³C peak to which carbon on the molecule it is, numerically label the peaks, and create a numerically labeled structure in chemdraw or something.
- Usually, it would take a long time to collect ¹³C spectra, but we've drastically decreased the number of scans, giving us quicker but worse data, and this NMR has a really strong probe that allows fewer scans to accomplish more.
- While we're waiting for the next spectrum to collect, we can set limits, noting which regions the instrument should measure.
- MNova for data analysis.
- We're going to plot I vs. τ and do a nonlinear data fitting to determine T_1 .

- After collecting the last few spectra, we'll split up: Some will be doing data analysis, finding our τ , I values. Others will be doing future experiments.
- There's also a 2D rendering of our data that we can do, using ¹H on one axis and ¹³C on the other.
- We'll start with $\tau = D7 = 0.001 \,\mathrm{s}$.
- For the lab report, we make the regression plot for each of the six carbons.
- The amount of time you need for a certain material is partially material dependent.
- Phase correction parameters are important.
 - PHC0 = -172.561.
 - PHC1 = -336.042.
- We're testing $\tau = 0.1 \,\mathrm{s}$ next.
- Strategy: Augment by 2 orders of magnitude twice and then fill in.
- Command line: "D7" and then enter your value.
- convbin2asc