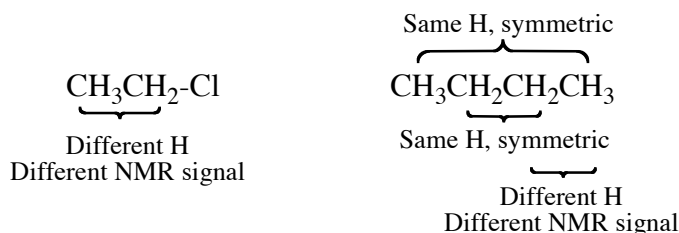


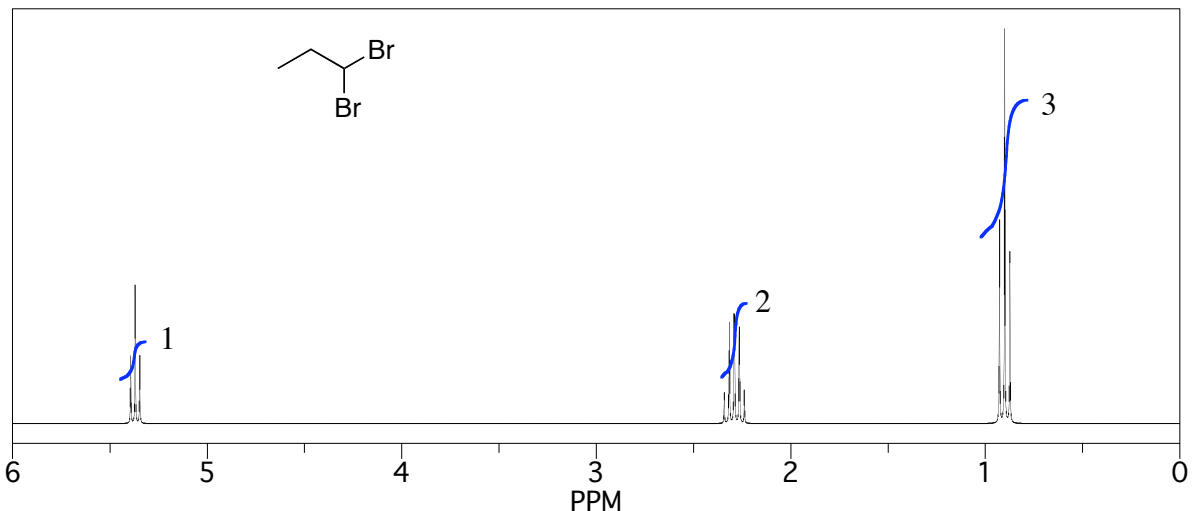
Nuclear Magnetic Resonance Spectroscopy (NMR)

- Spectrum represents the different interactions of **stereochemically different** protons (^1H) with the applied magnetic field.
- We will focus on ^1H NMR (proton, H^+)
- **4 general rules for ^1H NMR spectra**

1. Only stereochemically different ^1H s give different signals.



2. Area covered under the signal is proportional to the number of ^1H s causing the signal and is usually represented by integrals.



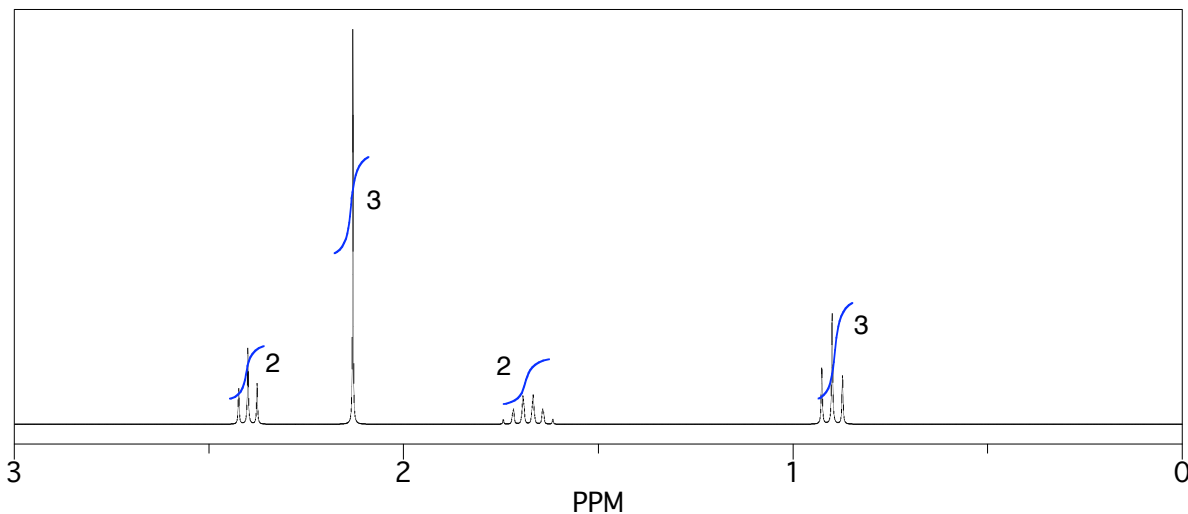
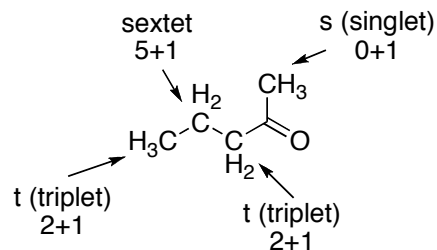
3. The **Chemical Shift** (where on spectrum each peak appears) depends on the “chemical environment” of each proton. (see above picture)

- ^1H s close to electronegative atoms (O, N, X (halogen)) or aromatics shift to the left (deshielded, downfield shifted)
- The larger the number of ^1H s on the same carbon the more to the right (shielded, upfield shifted) the NMR signal is.

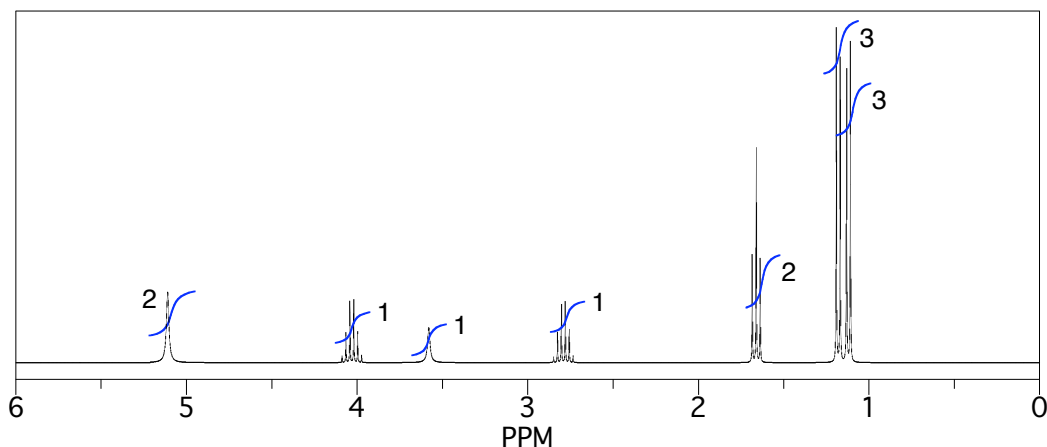
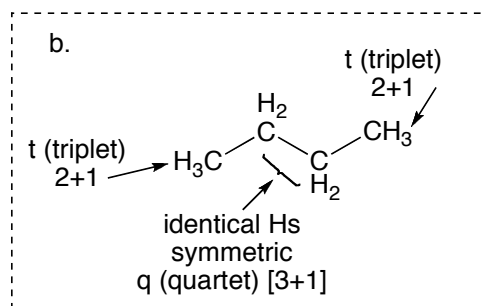
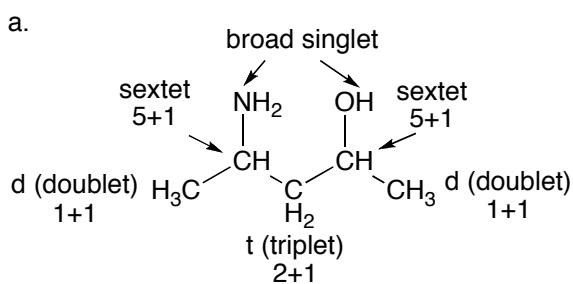
4. The multiplicity of the NMR peak depends on the number of ^1H s on neighboring carbons, NOT the same carbon. ^1H s attached to **adjacent carbons** split each other into:

$(n+1)$ peaks

n = number of ^1H s on adjacent carbons
(not the same C)



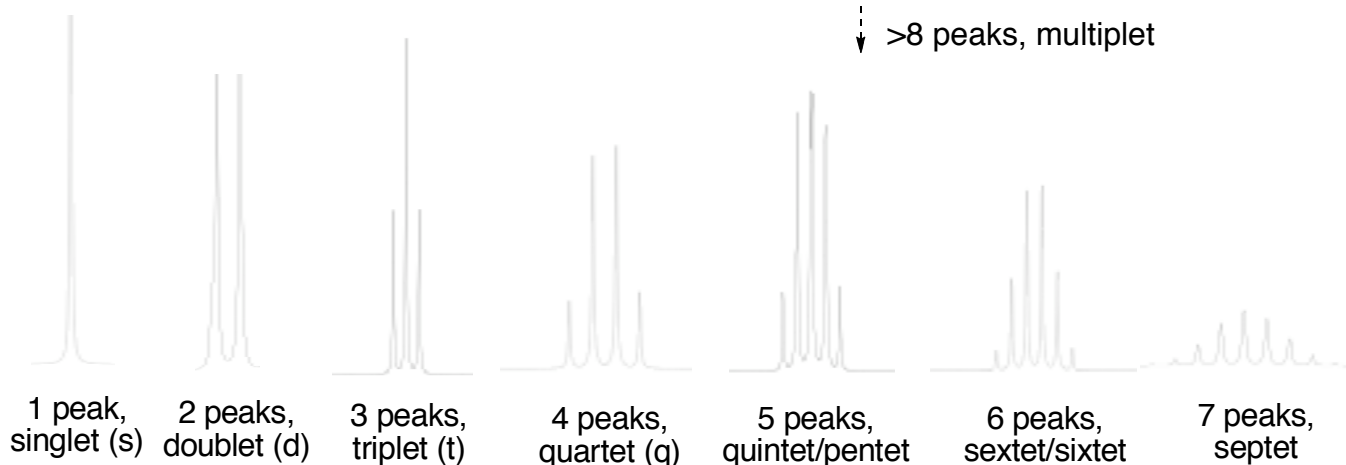
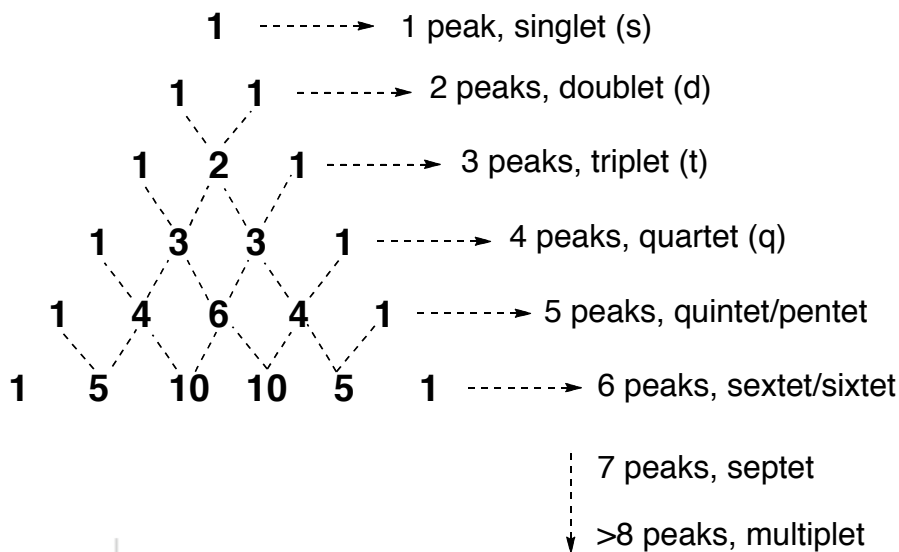
- a. Exchangeable, acidic ^1H ($-\text{OH}$, NH_2) DO NOT split ^1H s on adjacent carbons and show on the spectrum as broad singlets.



b. Only **non identical** ^1H s split each other.

The shape/relative intensity of the peaks follows the algorithm of **Pascal's Triangle**:

Pascal's Triangle



- Calculating the degree of unsaturation for a compound (number of RDBs, Rings and Double Bonds) when the molecular formula is known.

$$\#RDBs = \frac{2n + 2 - \#Hs - \#Halides + \# N \text{ atoms}}{2}$$

$n = \# \text{ Cs}$
 (Oxygens do not participate in this equation and can be ignored when RDBs are calculated)

i.e. $\text{C}_9\text{H}_9\text{OCl}$: $RDBs = (2 \times 9 + 2 - 9 - 1)/2 = 5$