

# 2023 NMR Users Training (II)

Basic NMR SOP for  
Small Molecules & Metabolomics Analysis

## Basic NMR Concept & Applications

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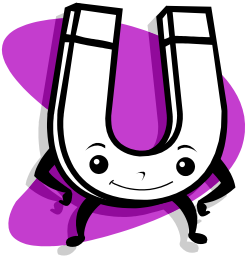
What is



Nuclear

Magnetic

Resonance?



## What you should know about **N**, **M**, and **R**?

( Just point out few "terms" here)

Properties of the **Nucleus** (原子核的特性)

*Nuclear spin & gyromagnetic ratio*

*Nuclear magnetic moments*

NMR Active Nucleus in a **Magnetic** Field (磁場中的原子核)

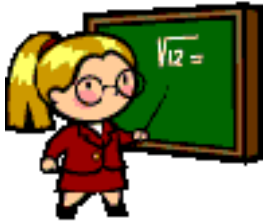
*Precession and the Larmor frequency*

*Nuclear Zeeman effect & Boltzmann distribution*

When the Spinning Nucleus receive the "Right" Energy

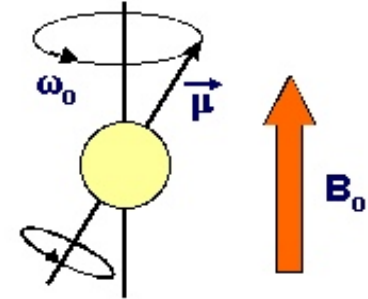
*Nuclear Magnetic **Resonance***

# (1) Properties of the Nucleus - Spin Parameters



$$\mu \text{ (Magnetic Moment)} = \gamma \mathbf{I} \left( \frac{h}{2\pi} \right)$$

$6.626 \times 10^{-34}$  joule-sec  
(Planck Constant)



$\mathbf{I}$ : spin number (property of nucleus)

$A$  (atomic mass) =  $Z$ (proton) +  $N$ (neutrons)

A	Z	N	Spin I
Even	Even	Even	$I=0$
Even	Odd	Odd	$I = \text{Integer}$ $1, 2, 3, \dots$
Odd	Even	Odd	$I = n/2,$ $n = 1, 3, 5, \dots$
Odd	Odd	Even	

$\gamma$ : gyromagnetic ratio (property of nucleus)

# Only those nucleus with spin I ≠ 0 are NMR Active!!



- 1/2 spin
- half-integer spin
- integer spin
- not observable
- radioactive

### JEOL ECZ NMR Frequencies

<sup>6</sup> Li	6.41	11.31	14.12
<sup>7</sup> Li	369.79	500.16	600.17
<sup>9</sup> Be	61.37	79.19	101.13
<sup>10</sup> B	426.42	533.49	640.15
<sup>11</sup> B	246.30	291.63	402.22
<sup>12</sup> C	58.83	73.50	89.32
<sup>13</sup> C	101.62	125.76	151.05
<sup>14</sup> N	50.31	62.58	76.32
<sup>15</sup> N	125.76	151.05	182.28
<sup>16</sup> O	54.25	67.51	81.25
<sup>17</sup> O	132.90	166.14	201.62
<sup>18</sup> O	101.62	125.76	151.05
<sup>19</sup> F	282.49	363.21	443.93
<sup>20</sup> F	112.27	142.41	172.55
<sup>21</sup> Ne	54.25	67.51	81.25
<sup>23</sup> Na	106.43	133.05	160.67
<sup>25</sup> Mg	125.76	151.05	182.28
<sup>27</sup> Al	101.62	125.76	151.05
<sup>29</sup> Si	50.31	62.58	76.32
<sup>31</sup> P	125.76	151.05	182.28
<sup>33</sup> S	50.31	62.58	76.32
<sup>35</sup> Cl	101.62	125.76	151.05
<sup>37</sup> Cl	125.76	151.05	182.28
<sup>39</sup> K	93.25	117.00	141.75
<sup>41</sup> K	125.76	151.05	182.28
<sup>43</sup> Ca	75.82	94.48	113.14
<sup>45</sup> Ca	112.27	142.41	172.55
<sup>47</sup> Ti	50.31	62.58	76.32
<sup>49</sup> Ti	125.76	151.05	182.28
<sup>51</sup> V	50.31	62.58	76.32
<sup>53</sup> Cr	50.31	62.58	76.32
<sup>55</sup> Cr	125.76	151.05	182.28
<sup>57</sup> Fe	50.31	62.58	76.32
<sup>59</sup> Co	50.31	62.58	76.32
<sup>61</sup> Ni	50.31	62.58	76.32
<sup>63</sup> Ni	125.76	151.05	182.28
<sup>65</sup> Ni	125.76	151.05	182.28
<sup>67</sup> Zn	50.31	62.58	76.32
<sup>69</sup> Ga	50.31	62.58	76.32
<sup>71</sup> Ga	125.76	151.05	182.28
<sup>73</sup> Ge	50.31	62.58	76.32
<sup>75</sup> Ge	125.76	151.05	182.28
<sup>77</sup> As	50.31	62.58	76.32
<sup>79</sup> Se	50.31	62.58	76.32
<sup>81</sup> Se	125.76	151.05	182.28
<sup>83</sup> Br	50.31	62.58	76.32
<sup>85</sup> Br	125.76	151.05	182.28
<sup>87</sup> Rb	50.31	62.58	76.32
<sup>89</sup> Y	50.31	62.58	76.32
<sup>91</sup> Zr	50.31	62.58	76.32
<sup>93</sup> Nb	50.31	62.58	76.32
<sup>95</sup> Mo	50.31	62.58	76.32
<sup>97</sup> Mo	125.76	151.05	182.28
<sup>99</sup> Tc	50.31	62.58	76.32
<sup>99</sup> Ru	50.31	62.58	76.32
<sup>101</sup> Ru	125.76	151.05	182.28
<sup>103</sup> Rh	50.31	62.58	76.32
<sup>105</sup> Pd	50.31	62.58	76.32
<sup>107</sup> Ag	50.31	62.58	76.32
<sup>109</sup> Ag	125.76	151.05	182.28
<sup>111</sup> Cd	50.31	62.58	76.32
<sup>113</sup> Cd	125.76	151.05	182.28
<sup>113</sup> In	50.31	62.58	76.32
<sup>115</sup> In	125.76	151.05	182.28
<sup>117</sup> Sn	50.31	62.58	76.32
<sup>119</sup> Sn	125.76	151.05	182.28
<sup>121</sup> Sb	50.31	62.58	76.32
<sup>123</sup> Sb	125.76	151.05	182.28
<sup>123</sup> Te	50.31	62.58	76.32
<sup>125</sup> Te	125.76	151.05	182.28
<sup>127</sup> I	50.31	62.58	76.32
<sup>129</sup> Xe	50.31	62.58	76.32
<sup>131</sup> Xe	125.76	151.05	182.28
<sup>133</sup> Cs	50.31	62.58	76.32
<sup>135</sup> Ba	50.31	62.58	76.32
<sup>137</sup> Ba	125.76	151.05	182.28
<sup>138</sup> La	50.31	62.58	76.32
<sup>139</sup> La	125.76	151.05	182.28
<sup>177</sup> Hf	50.31	62.58	76.32
<sup>179</sup> Hf	125.76	151.05	182.28
<sup>181</sup> Ta	50.31	62.58	76.32
<sup>183</sup> W	50.31	62.58	76.32
<sup>185</sup> Re	50.31	62.58	76.32
<sup>187</sup> Re	125.76	151.05	182.28
<sup>187</sup> Os	50.31	62.58	76.32
<sup>189</sup> Os	125.76	151.05	182.28
<sup>191</sup> Ir	50.31	62.58	76.32
<sup>193</sup> Ir	125.76	151.05	182.28
<sup>195</sup> Pt	50.31	62.58	76.32
<sup>197</sup> Pt	125.76	151.05	182.28
<sup>199</sup> Au	50.31	62.58	76.32
<sup>201</sup> Au	125.76	151.05	182.28
<sup>203</sup> Tl	50.31	62.58	76.32
<sup>205</sup> Tl	125.76	151.05	182.28
<sup>207</sup> Pb	50.31	62.58	76.32
<sup>209</sup> Pb	125.76	151.05	182.28
<sup>209</sup> Bi	50.31	62.58	76.32
<sup>209</sup> Po	125.76	151.05	182.28
<sup>210</sup> At	50.31	62.58	76.32
<sup>211</sup> At	125.76	151.05	182.28
<sup>212</sup> Rn	50.31	62.58	76.32
<sup>219</sup> Rn	125.76	151.05	182.28
<sup>235</sup> U	50.31	62.58	76.32
<sup>235</sup> Nd	50.31	62.58	76.32
<sup>235</sup> Pr	50.31	62.58	76.32
<sup>235</sup> Ce	50.31	62.58	76.32
<sup>235</sup> Lu	50.31	62.58	76.32
<sup>235</sup> Yb	50.31	62.58	76.32
<sup>235</sup> Tm	50.31	62.58	76.32
<sup>235</sup> Er	50.31	62.58	76.32
<sup>235</sup> Ho	50.31	62.58	76.32
<sup>235</sup> Dy	50.31	62.58	76.32
<sup>235</sup> Tb	50.31	62.58	76.32
<sup>235</sup> Gd	50.31	62.58	76.32
<sup>235</sup> Sm	50.31	62.58	76.32
<sup>235</sup> Pm	50.31	62.58	76.32
<sup>235</sup> Nd	50.31	62.58	76.32
<sup>235</sup> Pr	50.31	62.58	76.32
<sup>235</sup> Ce	50.31	62.58	76.32

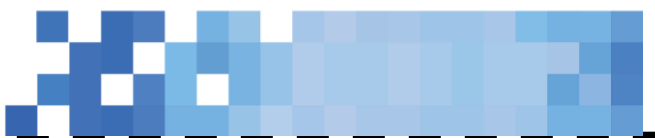
## Table of Isotopes and NMR Parameters

<b>Li</b>	<b>Be</b>	<b>H</b>	<b>H</b>	<b>B</b>	<b>C</b>	<b>N</b>	<b>O</b>	<b>F</b>	<b>Ne</b>																															
6 7 9 7.59 92.41 100 6.27 16.65 -5.98 3.79 1590 81.5 15 40	1 2 3 1/2 1 1/2 99.98 1.15E-02 42.58 6.54 5870 6.52E-03 10	10 11 3 3/2 19.9 80.1 4.58 13.66 23.2 777 150	13 14 15 1/2 1 -1/2 99.63 0.36 3.08 -4.32 1 1000	17 19 21 -5/2 -1/2 3/2 3.8E-02 100 0.27 -5.77 40.08 -3.36 4900 3.91E-02	27 29 31 5/2 -1/2 1/2 92 4.68 100 100 0.76 75.78 24.5 11.1 -8.47 17.25 1220 2.16 391 400 400 1100	33 35 37 3/2 3/2 3/2 50.69 49.31 11.49 10.70 11.54 -1.64 2.37 288 1.28	39 41 43 3/2 3/2 -7/2 93.25 6.73 0.13 1.99 1.09 -2.87 2.79 3.33E-02	45 47 49 7/2 -5/2 -7/2 10.36 -2.40 -2.41 0.918 1.2	50 51 6 7/2 0.25 99.75 4.25 11.21	53 55 -3/2 -5/2 9.50 10.68	57 59 1/2 7/2 2.11 10.08	61 63 65 -3/2 1/2 1/2 1.13 69.17 30.83 -3.81 11.32 12.10	67 69 71 3/2 3/2 3/2 4.10 60.1 39.89 10.25 13.02	73 75 -9/2 7/2 7.73 100 7.32 -1.49	77 79 81 7/2 3/2 3/2 7.63 50.69 49.31 8.16 10.70 11.54	83 85 87 5/2 3/2 3/2 79 81 83 50.69 49.31 11.49 2.37 288 1.28	89 91 93 5/2 3/2 -9/2 1.12 -1/2 -1/2 5.22 3.27 3.27 1.12 1.12 1.12	95 97 99 5/2 -5/2 -9/2 15.92 9.55 -- 9.62 -1.96 -2.19 0.848 1.59 0.186	101 103 105 1/2 3/2 3/2 12.76 17.06 100 -5.2 51.83 48.16 -1.96 -1.35 -1.96 1.49 0.205 0.29	107 109 111 113 -1/2 -1/2 -1/2 -1/2 107 109 111 113 51.83 48.16 12.8 12.22 -9.07 -9.49 9.37 9.39 20.8 26.6	115 117 119 -1/2 -1/2 -1/2 9/2 9/2 -1/2 -1/2 9.2 9.2 7.68 8.59 57.21 42.79	121 123 125 5/2 7/2 -1/2 -1/2 5/2 7/2 -1/2 -1/2 5.72 7.12 7.12 7.12 10.26 5.55 -11.23 -13.55	127 129 131 1/2 3/2 3/2 127 129 131 0.89 7.07 100 8.58 -11.86 3.52 560 33.6 3.5	133 135 137 3/2 3/2 3/2 133 135 137 1300	139 141 143 5 7/2 7/2 9.00E-02 99.91 18.6 13.62 99.98	147 149 151 153 7/2 -7/2 -7/2 14.99 13.82 47.81 52.19 -1.78 -1.46 10.59 4.67	155 157 -3/2 -3/2 14.8 15.65	159 161 163 3/2 5/2 5/2 18.91 24.9 10.24	165 167 1/2 1/2 16.87 13.18 29.52 70.47	169 171 173 -1/2 -1/2 -1/2 22.93 100 22.93 -1.23 -3.53	175 177 179 7/2 7/2 7/2 97.41 2.59	181 183 185 187 7/2 5/2 5/2 7.53 -2.07 4.86 3.45	191 193 195 197 -1/2 -1/2 -1/2 -1/2 0 0 0 0	199 201 203 205 1/2 3/2 3/2 3/2 16.87 13.18 29.52 70.47	207 209 211 213 1/2 1/2 1/2 1/2 22.1 100 22.1 100 8.88 6.96	209 210 211 212 1/2 5 -9/2 5/2 848	209 210 211 212 1/2 5 -9/2 5/2 848	209 210 211 212 1/2 5 -9/2 5/2 848	209 210 211 212 1/2 5 -9/2 5/2 848	209 210 211 212 1/2 5 -9/2 5/2 848

**Element**

- Isotope **13**
- Spin **1/2**
- Natural Abundance **1.07**
- Gamma (g/10<sup>6</sup> Hz/T) **10.71**
- Relative Receptivity to <sup>13</sup>C **1**
- Typical Chemical Shift Range (ppm) **200**

The rest of the actinide series is not commonly observed due to their radioactive nature.



# (2-1) NMR Active Nucleus in Magnetic Field - Larmor precession

有核磁共振特性的原子核, 在磁場中會像陀螺一樣自旋並產生磁矩(magnetic moment), 此自旋頻率稱之為 "Larmor frequency "

$$\mu = \gamma I \left( \frac{h}{2\pi} \right)$$

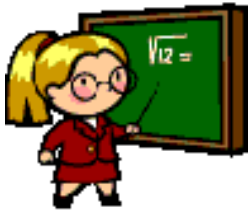
(magnet moment)       $6.626 \times 10^{-34}$  joule-sec (Planck Constant)

(gyromagnetic ratio)

$$\nu_{0,precession} = \left( \frac{\gamma B_0}{2\pi} \right)$$

or  $\omega_0 = \gamma B_0$

- ◆ Under the same magnetic field, precession frequency proportional to the gyromagnetic ratio  $\gamma$  of each atom (自旋頻率與原子核的特性有關)  
 $\gamma_{(1H)} = 26.7519 (10^7 \cdot \text{rad/T} \cdot \text{s})$        $\gamma_{(13C)} = 6.7624 (10^7 \cdot \text{rad/T} \cdot \text{s})$
- ◆ The higher the magnetic field, faster the spinning frequency (磁場愈高轉速愈快)



Differ "atom type" → different Larmor frequency

核種不同 (ie.磁旋比 gyromagnetic ratio  $\gamma$  不同), 自旋的頻率就不同!



14.1 Tesla

**1H**



I am the fast spinner in the world!!  
@14.1T ~ 600MHz!

**13C**



I can spin ~ 150MHz @14.1T

**15N**



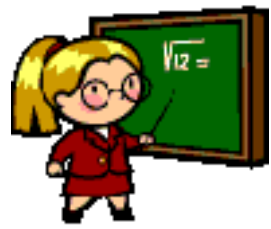
I prefer to spin slowly...  
@14.1T ~ 60MHz!

**1H**  $\gamma$ : 26.7519 ( $10^7 \cdot \text{rad/T} \cdot \text{s}$ )

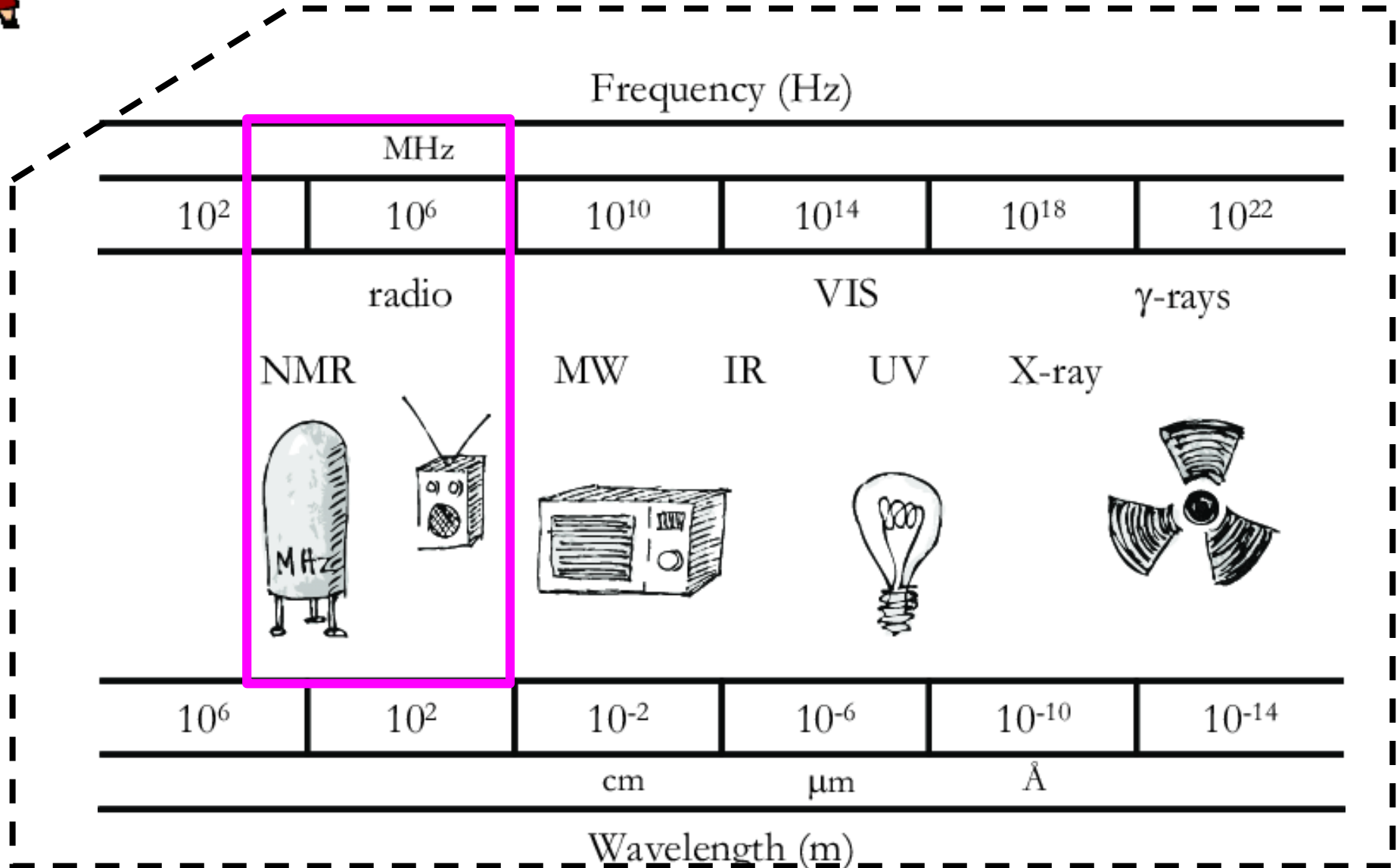
**13C**  $\gamma$ : 6.7262 ( $10^7 \cdot \text{rad/T} \cdot \text{s}$ )

**15N**  $\gamma$ : -2.7116 ( $10^7 \cdot \text{rad/T} \cdot \text{s}$ )

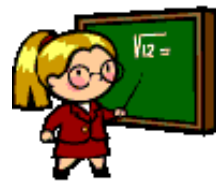
$$\nu_{0,\text{precession}} = (\gamma B_0 / 2\pi)$$



The precession frequency is @ radio frequency range  
自旋頻率接近無線電頻率 (radio frequency)







Same "atom type", but "under different chemical environment" could generate different signal. This makes NMR useful and interesting.

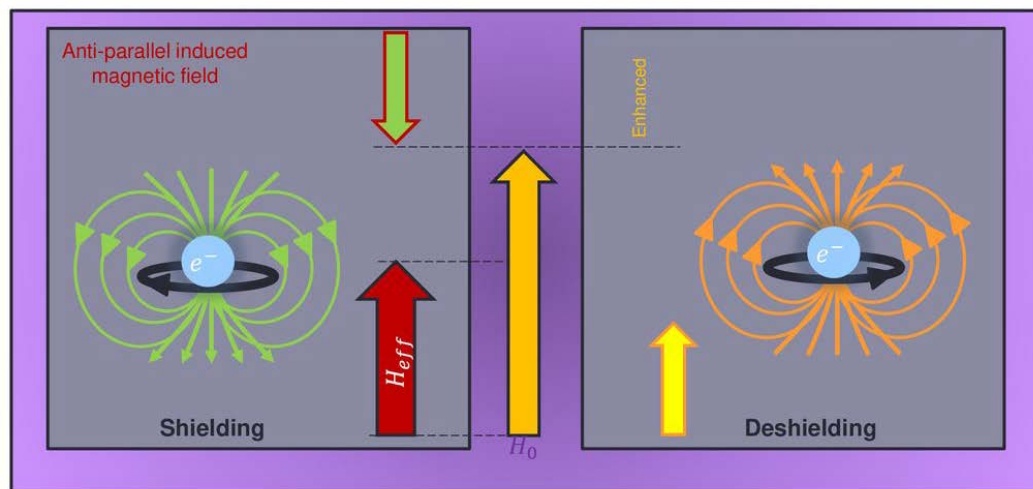
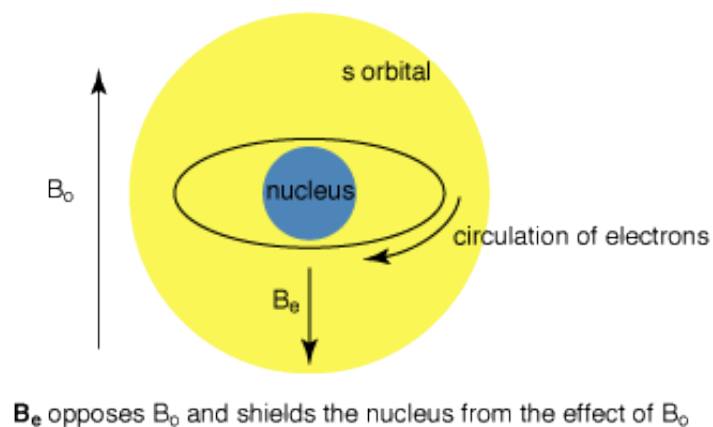
$$\nu_{0, \text{precession}} = (\gamma B_0 / 2\pi) \rightarrow \nu_{\text{precession}} = (\gamma B_{\text{eff}} / 2\pi)$$

where  $B_{\text{eff}} = (1 - \sigma) B_0$

$$\nu_{\text{precession}} = \nu_{0, \text{precession}} (1 - \sigma)$$

$\sigma > 0 \rightarrow$  nuclei is shielded by electron cloud

$\sigma < 0 \rightarrow$  electron around this nuclei is withdraw

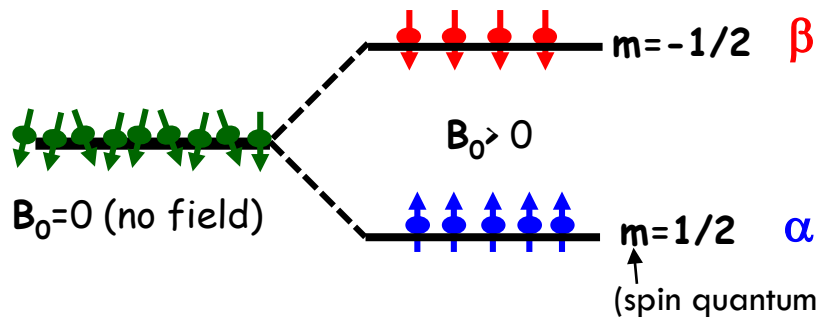


# (2-2) NMR Active Nucleus in a Magnetic Field - Zeeman Effect

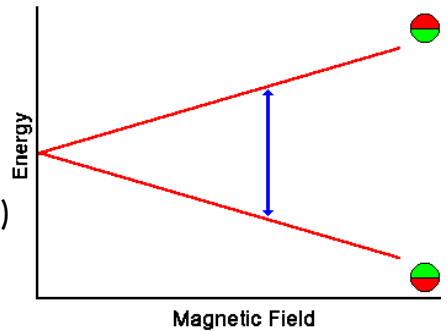
在磁場中, 核能階分裂(Nuclear Zeeman effect), 且核自旋以 Boltzmann distribution 分布



Example:  $I=1/2$  system, spins tend to generally aligned with or opposed to the external field, as "spin-up" or "parallel", and "spin-down" or "anti-parallel".



Nuclear Zeeman effect

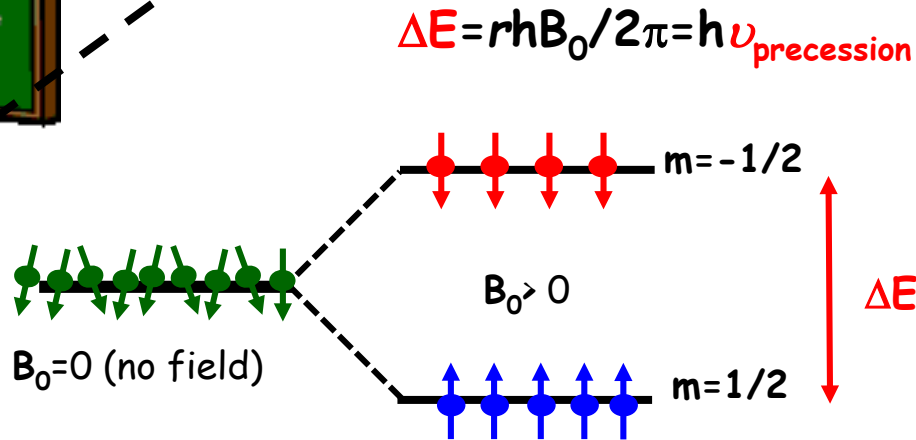
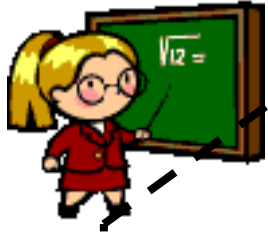


$$N_{\beta}/N_{\alpha} = \exp(-\Delta E/kT) = \exp[(\gamma h B_0)/(2\pi kT)]$$

Boltzmann distribution

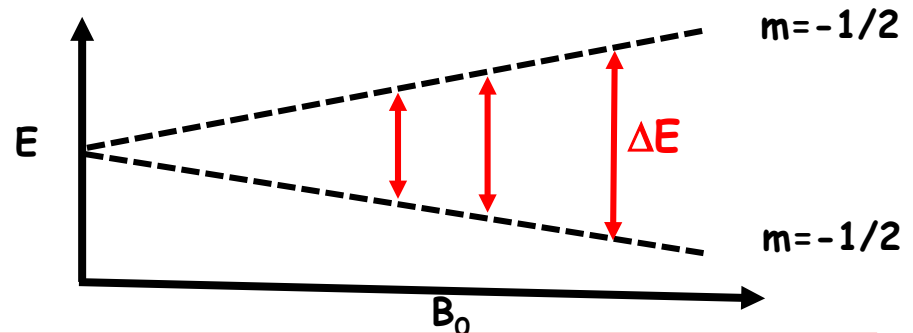
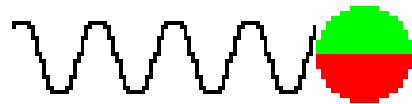
Different between  $N_{\beta}$  and  $N_{\alpha}$  just "ppm"  
, ie. NMR is an "insensitive" technique

### (3) When the Nucleus receive the "Right" Energy => Resonance



$E_{\text{induced}} = \Delta E$   
**On Resonance !!!**  
(ex:  $E_{\text{induced}} = h\nu_{\text{induced}}$ )

The spin at low energy level absorb the Induce Energy, then jump to the **higher energy level** ( $m=1/2 \rightarrow m= -1/2$ ), **but still spin at  $\nu_{\text{precession}}$**



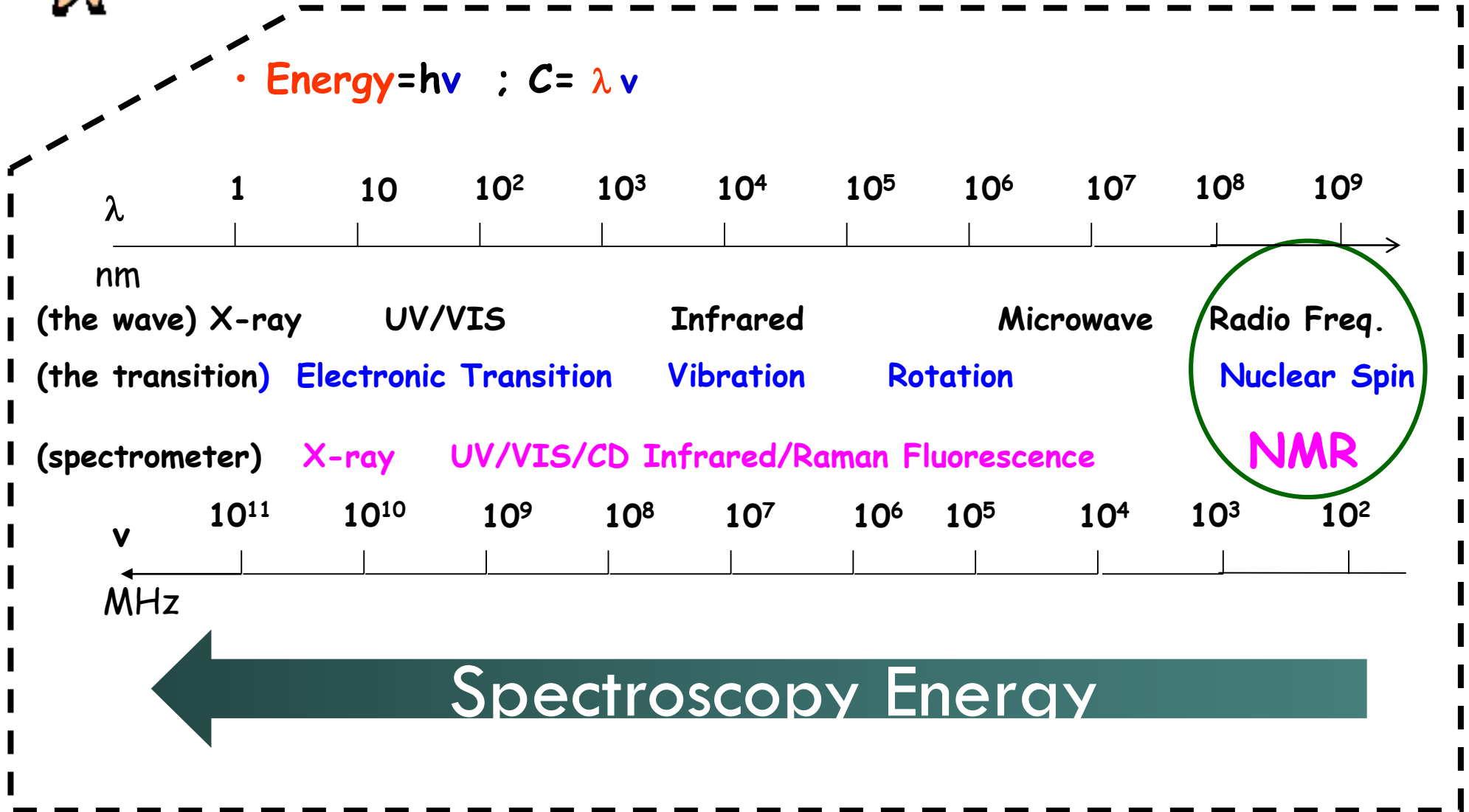
**Higher the field ( $B_0$ ), better the sensitivity and resolution**

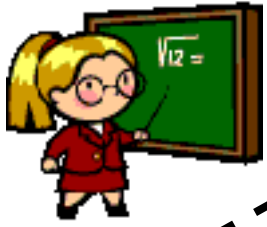


# Radiation energy for NMR spectroscopy

(頻率範圍約 80-1000MHz, 是能量弱的電磁波)

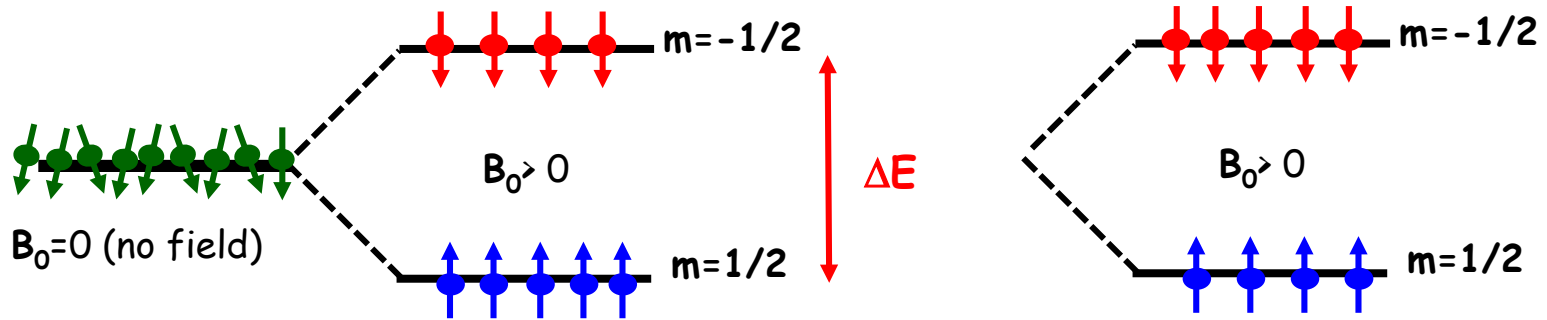
•  $\text{Energy} = h\nu$  ;  $c = \lambda \nu$





## During irradiation (current induced)

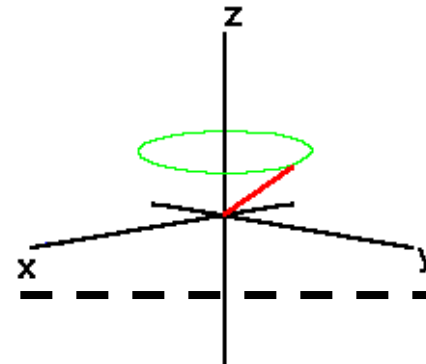
- (1) Some Nuclei in lower energy state will absorb induce energy , and jump to the **higher energy state** ( ie 能階躍遷 )



- (2) All of the individual nuclear magnetic moments become **phase coherent**, and the net M process around the z axis at a angle  $\alpha$

$$M_z = M \cos \alpha$$

$$M_{xy} = M \sin \alpha.$$

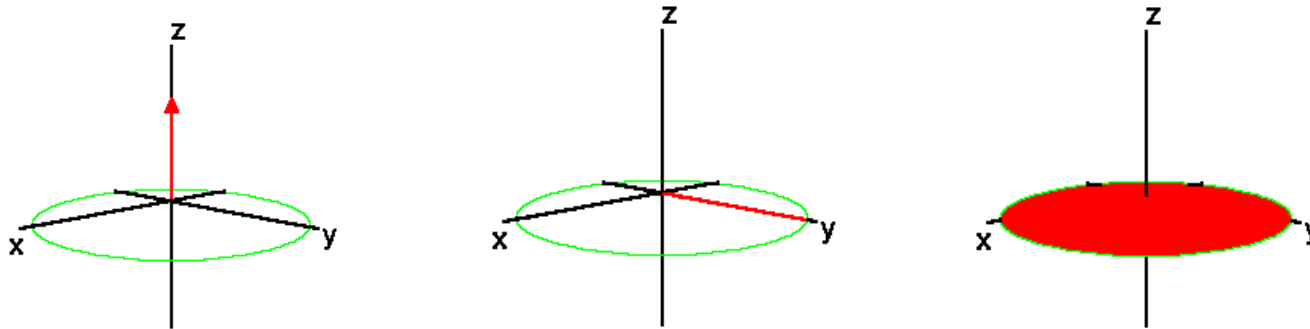




## After irradiation ceases

- (1) population of the states revert to a **Boltzmann distribution**
- (2) individual nuclear magnetic moments begin to lose their phase coherence and return to a **random** arrangement around the z axis.

- This process is called “relaxation process”
- There are two types of relaxation process :
  - T1 (spin-lattice relaxation) (ie. 回到最低能量)
  - T2 (spin-spin relaxation) (ie. 回到最大亂度)



**NMR signals will be collected during this relaxation process**

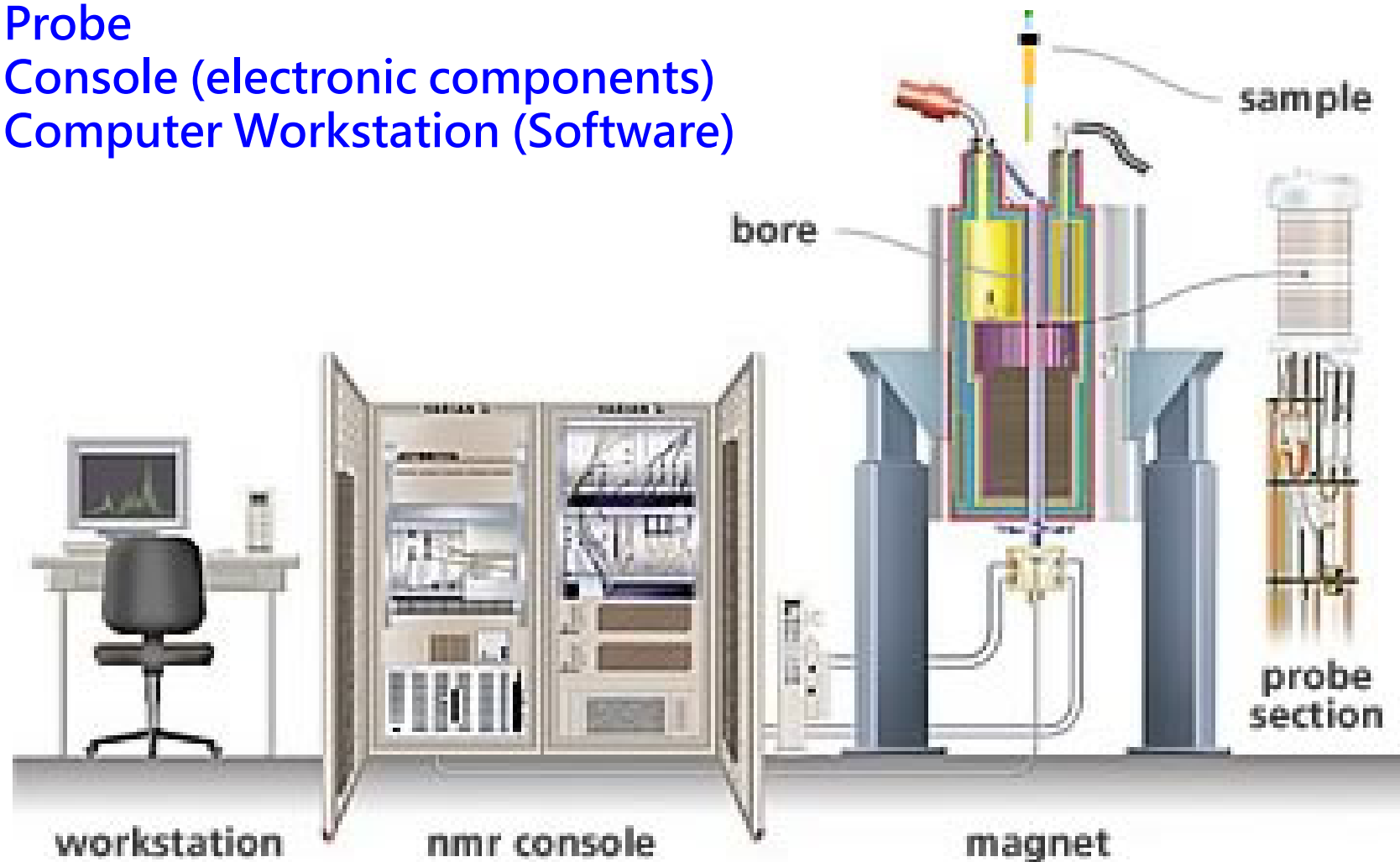
How to



Obtain NMR  
Signals  
(Spectrum)?

## NMR Hardware (核磁共振儀示意圖)

- (1) Magnet (superconducting magnet)
- (2) Probe
- (3) Console (electronic components)
- (4) Computer Workstation (Software)







# Steps for NMR Experiments

Prepare Samples

Collect NMR spectra

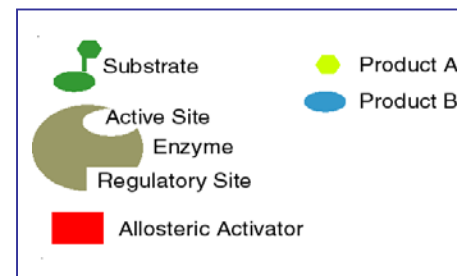
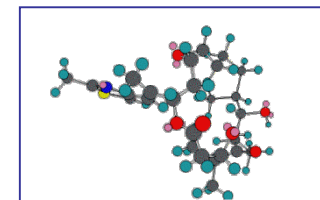
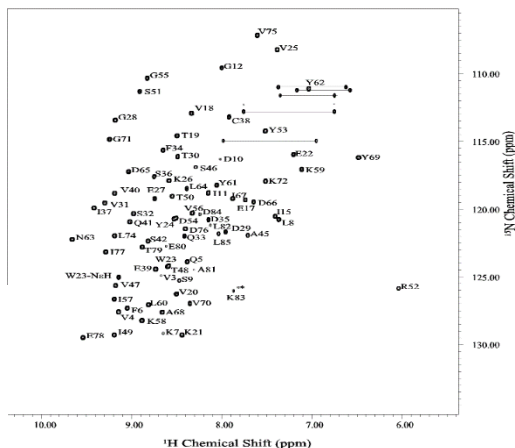
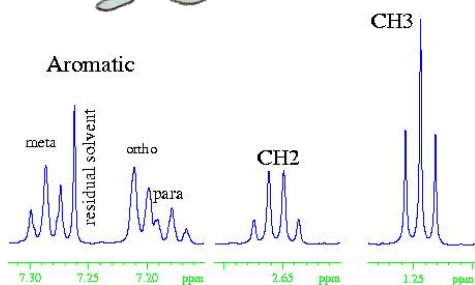
Data Analysis

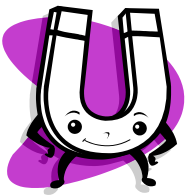
Expt. Methods

Processing / Analysis

Expt. Design

Assignment, Structure Calculation, Interpretation  
.....

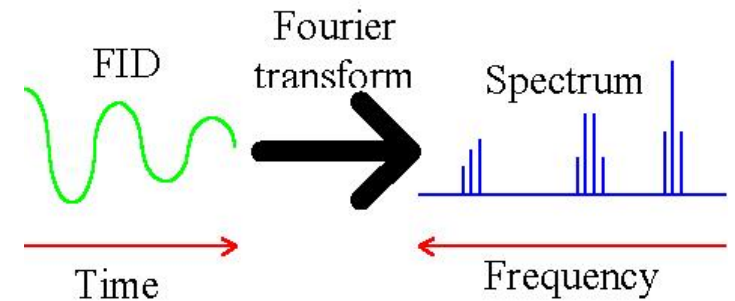
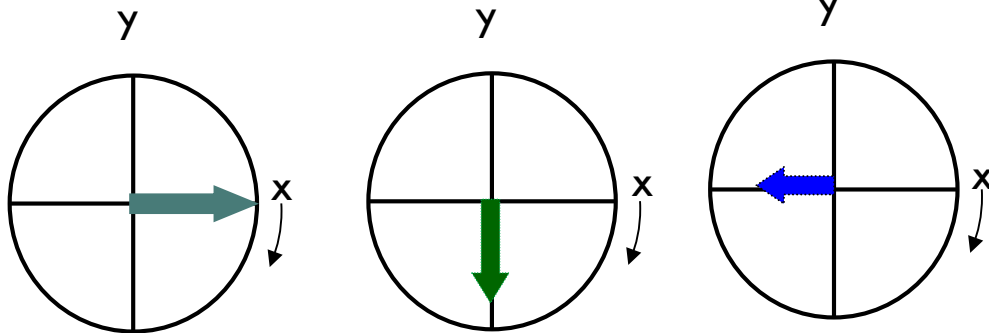
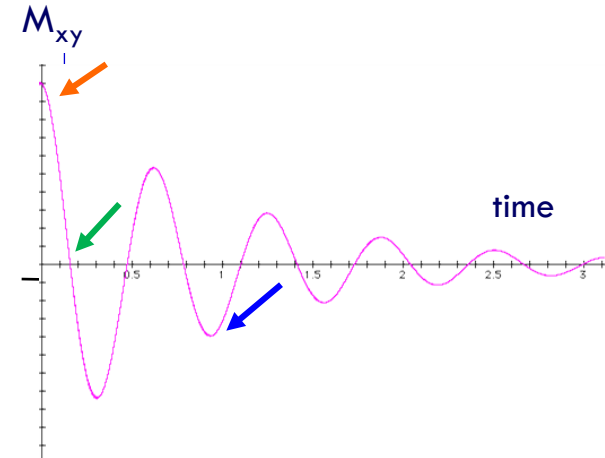
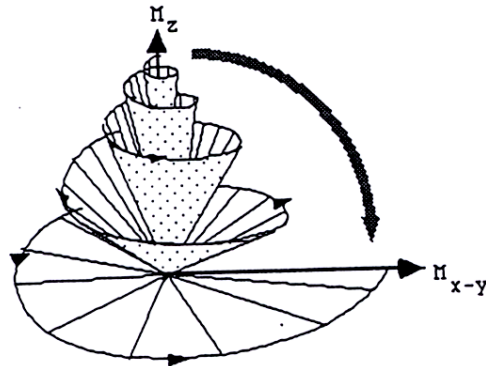




# (1) NMR Signal - FID

- FID is the **time-domain response** of the spin after a pulse (irradiation energy)
- Due to the **relaxation effect**, the response at receiver coil become a **exponentially decaying signal**, ie **Free Induction Decay (FID)**.

Relaxation process

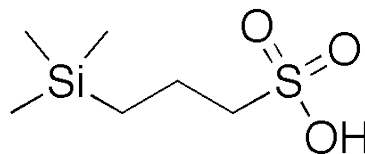
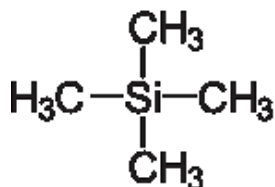


- The FID (free induction decay) **Fourier transform** to **frequency domain** to obtain  $\nu_{\text{precession}}$  for each different nuclei.



## (2) NMR Signal -Chemical Shift

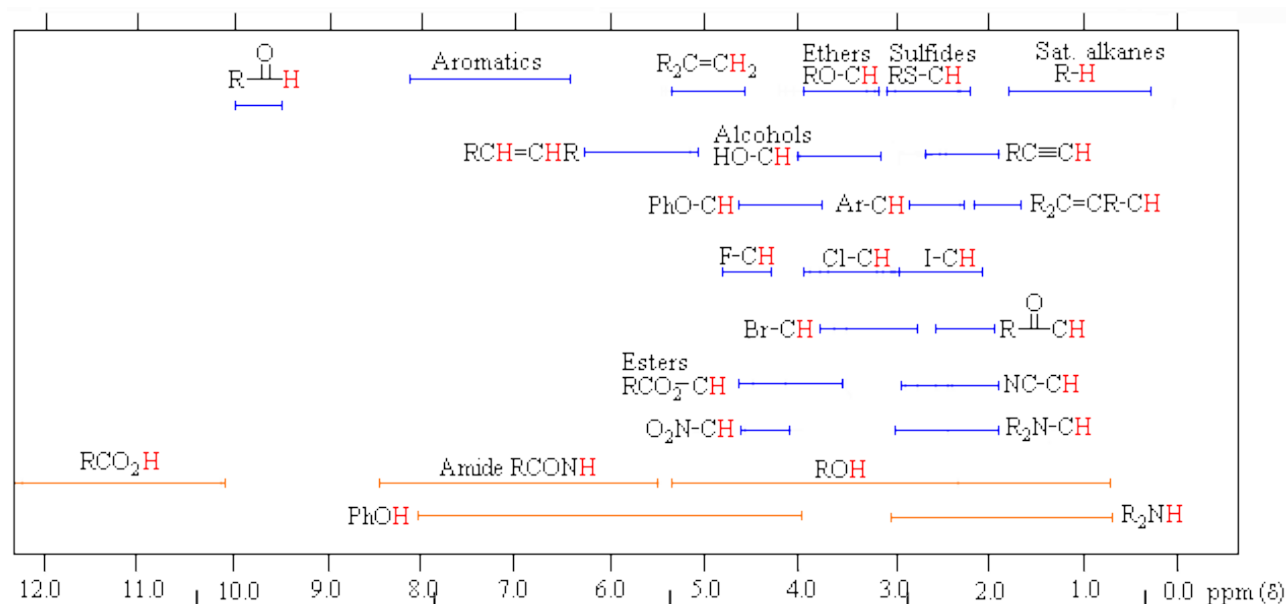
- The frequency of **absorption for a nucleus of interest** ( $\nu$ ) relative to the frequency of absorption of a **molecular standard** ( $\nu_{standard}$ ) is called the **chemical shift** of the nucleus (notice the frequency  $\nu$  is in unit of Hz).
- The standard molecule for both  $^1\text{H}$  and  $^{13}\text{C}$  NMR spectroscopy is tetramethylsilane (**TMS**). For biomolecular NMR, 2,2-dimethyl-2-silapentane-5-sulfonate (**DSS**) is used instead.



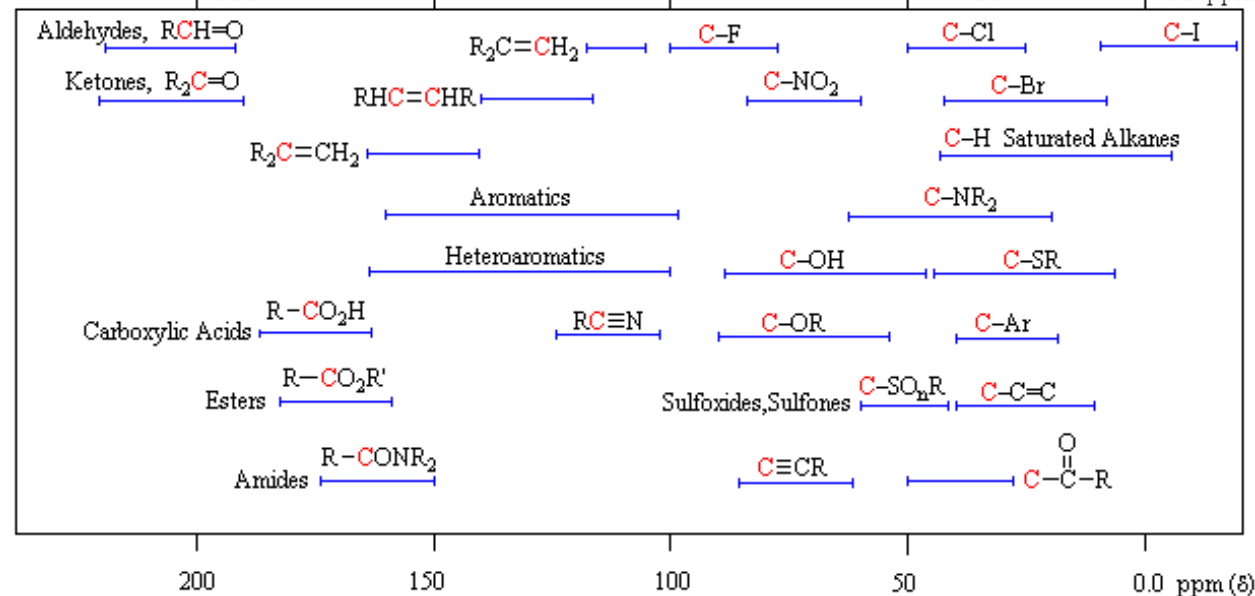
- Chemical shift is used to express as  $\delta$** , and  $\delta = (\nu - \nu_{\text{TMS}}) / \nu_0$  where  $\nu_0$  is the center frequency for magnet ( ex: 600MHz for  $^1\text{H}$ , 150MHz for  $^{13}\text{C}$  @14.1 T NMR)
- Chemical shift  $\delta$**  is in the ratio of **Hz/MHz**, the result is a ratio of one part per million, **ppm** (which is independent of external magnetic field )

- Notice that, the chemical shifts for a molecule **do not depend on the magnetic field**. For example, chemical shift for H<sub>2</sub>O in 100 MHz magnet (2.35 T) is the same as in a 600 MHz magnet (14.1 T).

<sup>1</sup>H

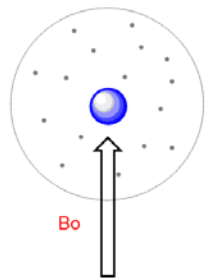


<sup>13</sup>C



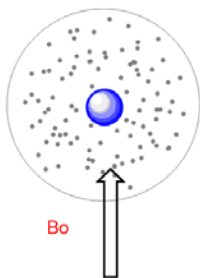
# Electron Cloud

low electron density



This nucleus experiences **stronger magnetic field** because it is not shielded as much.

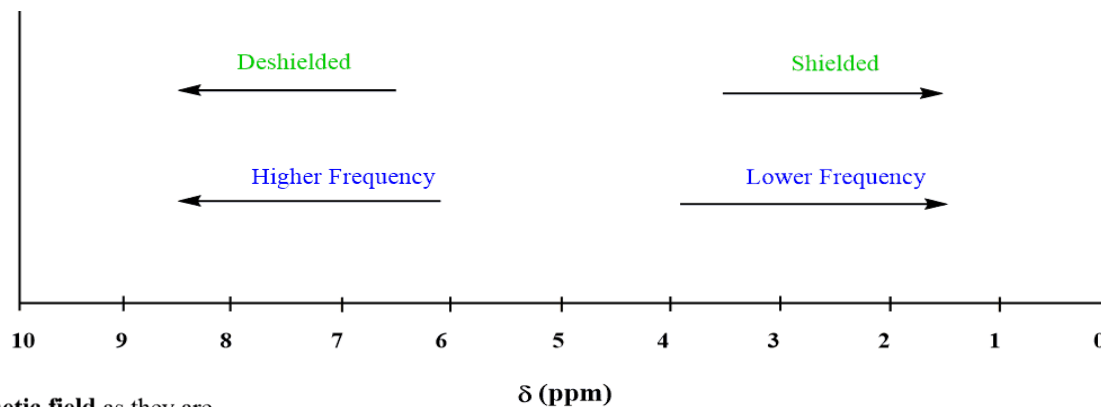
high electron density



This nucleus is shielded and experiences **weaker magnetic field** - will appear upfield (low energy).

Electron cloud shields the nucleus from the magnetic field

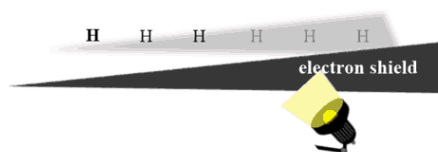
# Scale of NMR Spectrum



The more electron-withdrawing the group, the stronger the protons feel the magnetic field as they are less shielded by the electrons and more exposed to the field.

← Increasing electron-withdrawing power

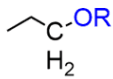
**Downfield**  
high frequency  
electron deshielded



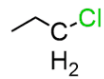
**Upfield**  
low frequency  
electron shielded



~ 4.5



~ 4



~ 3.5

ppm



~ 2.5



~ 1

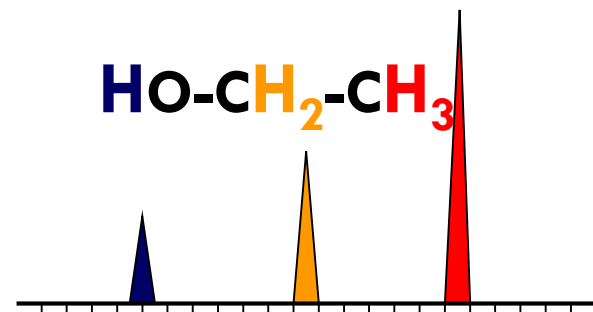
Fluorine, as the most electronegative element, deshields and shifts the signal of adjacent protons to ~ 4 ppm.

The oxygen and chlorine are not as electronegative and the ppm value is less downfield. I, Br, and some other groups also shift the signal to this region.

The carbonyl group is electron-withdrawing and the frequency of H's next to are shifted downfield by ~ 1.5 and found at 2.5 ppm.

This is the typical region of alkyl protons. Increasing the number of alkyl groups shifts the signal downfield.

# Ethanol

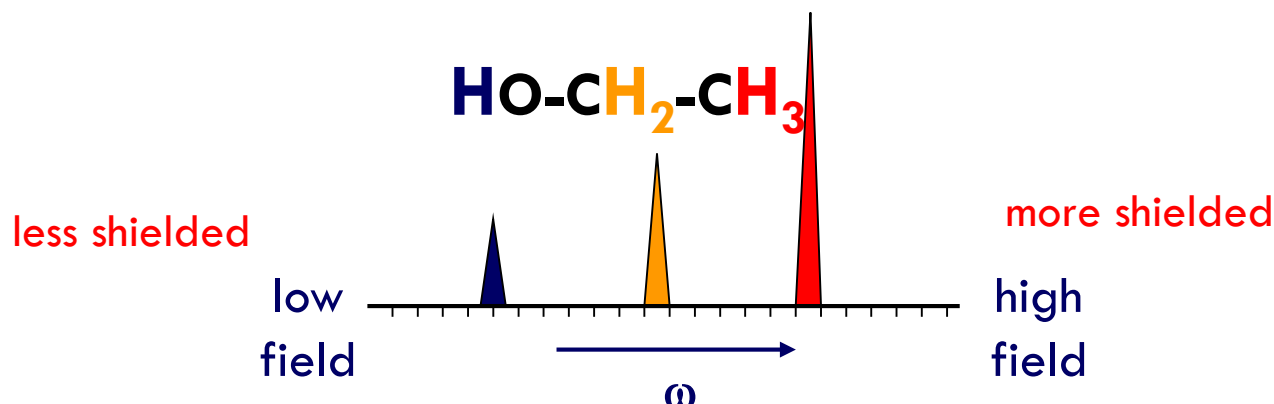


less shielded

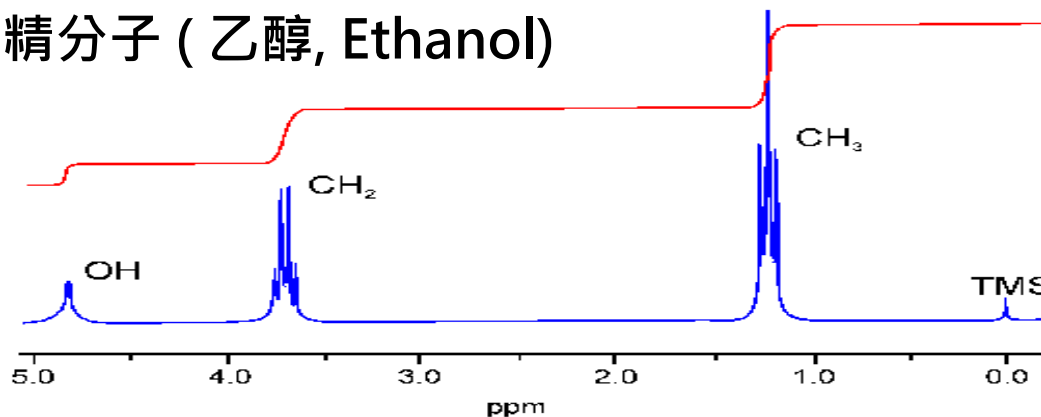
more shielded

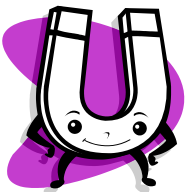
# What a simple 1D $^1\text{H}$ NMR spectrum tell us?

- The *number* of signals shows how many different **kinds of protons**
- The *location* of the signals shows how shielded/deshielded the proton
- The *intensity* of the signal shows **the number of protons** of that type.
- Signal *splitting* shows the number of protons on **adjacent atoms**.



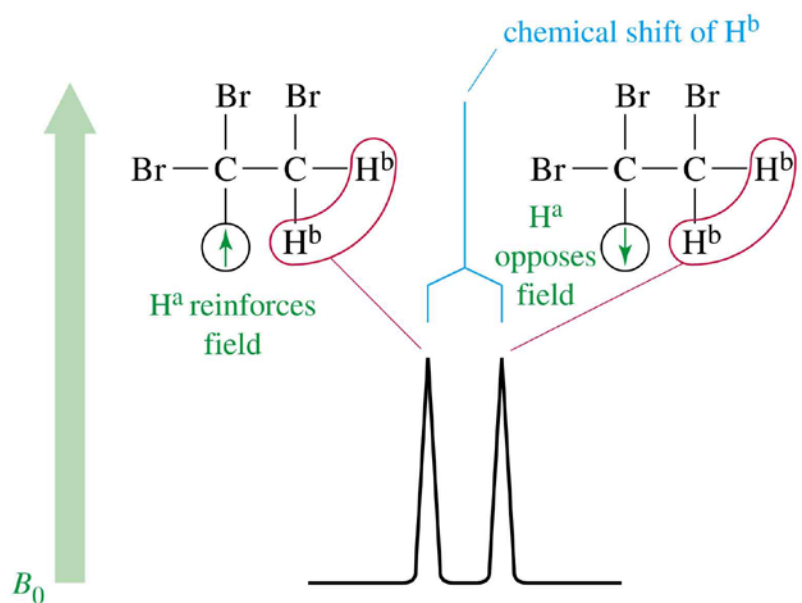
Example: 酒精分子 (乙醇, Ethanol)



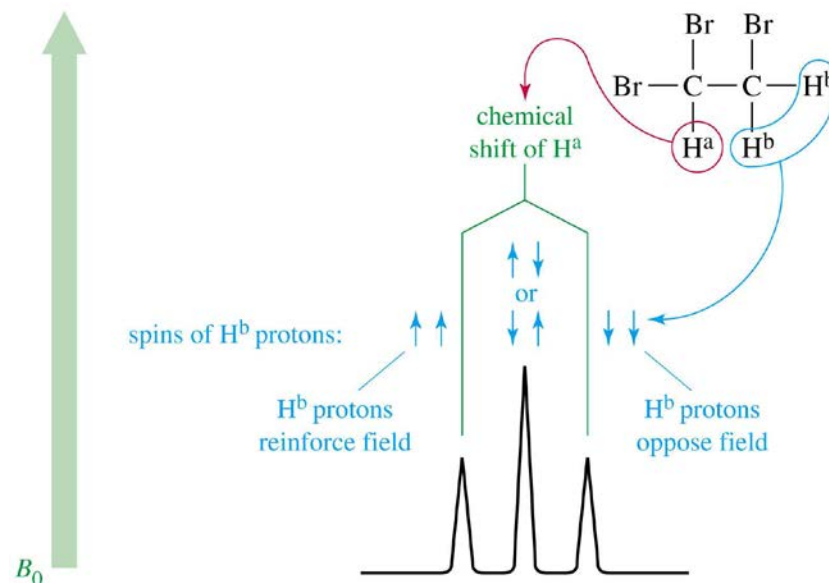


## (3) NMR Signal -Coupling Effect

- Nuclei which are close to one another could cause an influence on each other's effective magnetic field. If the distance between non-equivalent nuclei is less than or equal to **three bond lengths**, this effect is observable. This is called *spin-spin coupling* or *J coupling*.



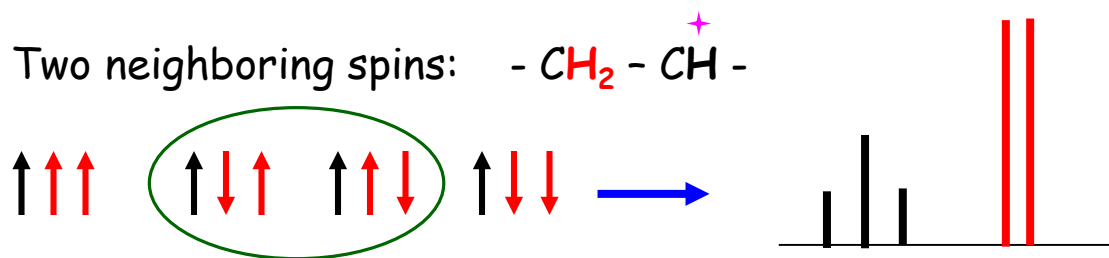
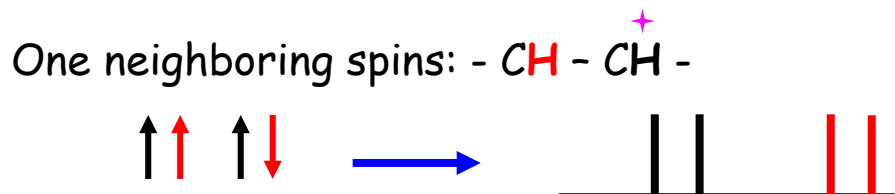
Chemical Shift of  $H^b$



Chemical Shift of  $H^a$

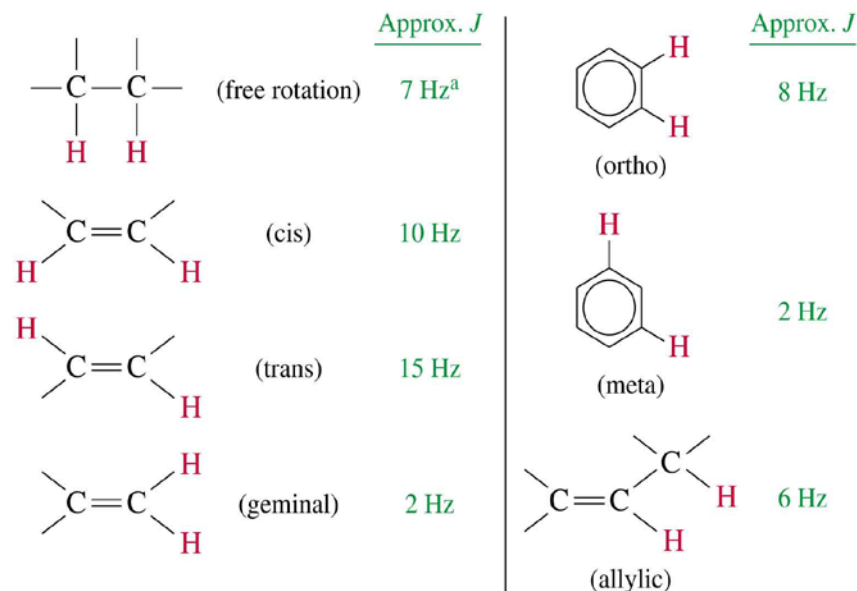
## Example of I=1/2 nuclei

•N neighboring spins: split into N + 1 lines



1  
 1 1  
 1 2 1  
 1 3 3 1  
 1 4 6 4 1  
 1 5 10 10 5 1

- Distance between the peaks of multiple is measured in Hz
- Coupling constant does not depend on strength of the external field



<sup>a</sup>The value of 7 Hz in an alkyl group is averaged for rapid rotation about the carbon-carbon bond. If rotation is hindered by a ring or bulky groups, other splitting constants may be observed.



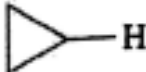

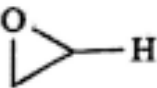
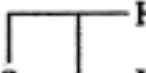

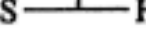

## Example of H-X Coupling Constant

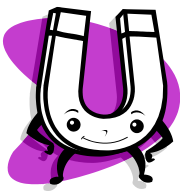
- Spin-spin coupling takes place **between all NMR active nuclei**, not just between protons. (EX: coupling to  $^{13}\text{C}$ ,  $^{31}\text{P}$ ,  $^{19}\text{F}$ ... but many other nuclei can couple. )
- Coupling constant does **not dependent on strength of the external field**

$$^1J_{XY}$$

$J_{\text{CD}}$	20
$J_{\text{CF}}$	-167
$J_{\text{CN}}$	6-8
$J_{\text{PH}}$	>600

$$^1\text{H}-^{13}\text{C}$$

Type	$J$ (Hz)	Type	$J$ (Hz)
$\text{CH}_3\text{-H}$	125	$\text{CH}_3\text{Li}$	98
$\text{Ph-CH}_2\text{-H}$	129	$\text{Cl}_2\text{CH-H}$	178
$\text{RC}\equiv\text{C-CH}_2\text{-H}$	132	$\text{O}_2\text{N-CH}_2\text{-H}$	147
$\text{R}_2\text{NCH}_2\text{-H}$	133	$\text{FCH}_3\text{-H}$	149
$\text{RSCH}_2\text{-H}$	138	$\text{ClCH}_2\text{-H}$	150
$\text{ROCH}_2\text{-H}$	140	$\text{ICH}_2\text{-H}$	151
$(\text{NC})_2\text{CH-H}$	145	$\text{BrCH}_2\text{-H}$	152
	161	$(\text{CH}_3\text{O})_2\text{CH-H}$	162
	134		180
	137		137
	150		150



## NMR could be helpful for diverse applications

◆ Due to **different chemical** or physical **environment** of the atom in the molecule, as well as the difference in atom number, NMR signals would be **different** !!

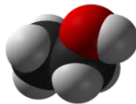
原子核週遭環境只要有一點不同, **核磁共振訊號就不同** !!

原子核個數不同, 訊號**強度也不同** !!

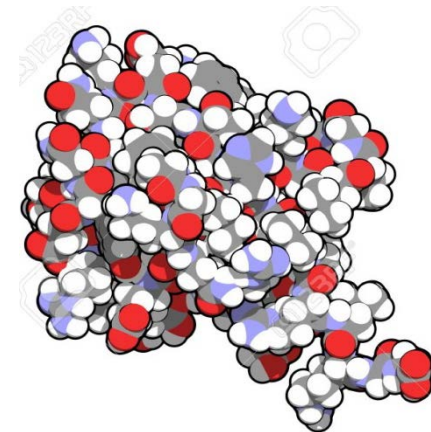
H<sub>2</sub>O



C<sub>2</sub>H<sub>5</sub>OH

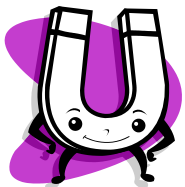


Ubiquitin

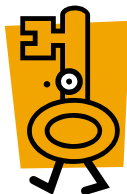


◆ Therefore, **NMR** could provide detail **atomic structural information** for molecules under specific condition, or provide useful information on **molecular interaction studies**.

不同的分子, 乃由原子以各種排列方式組合而成. 因此**核磁共振**方法是用來研究**分子結構/狀態**的重要工具之一, 對於“**分子間作用**”的相關課題也有極大的助益!!



## Applications for NMR in HFNMRC & GRC

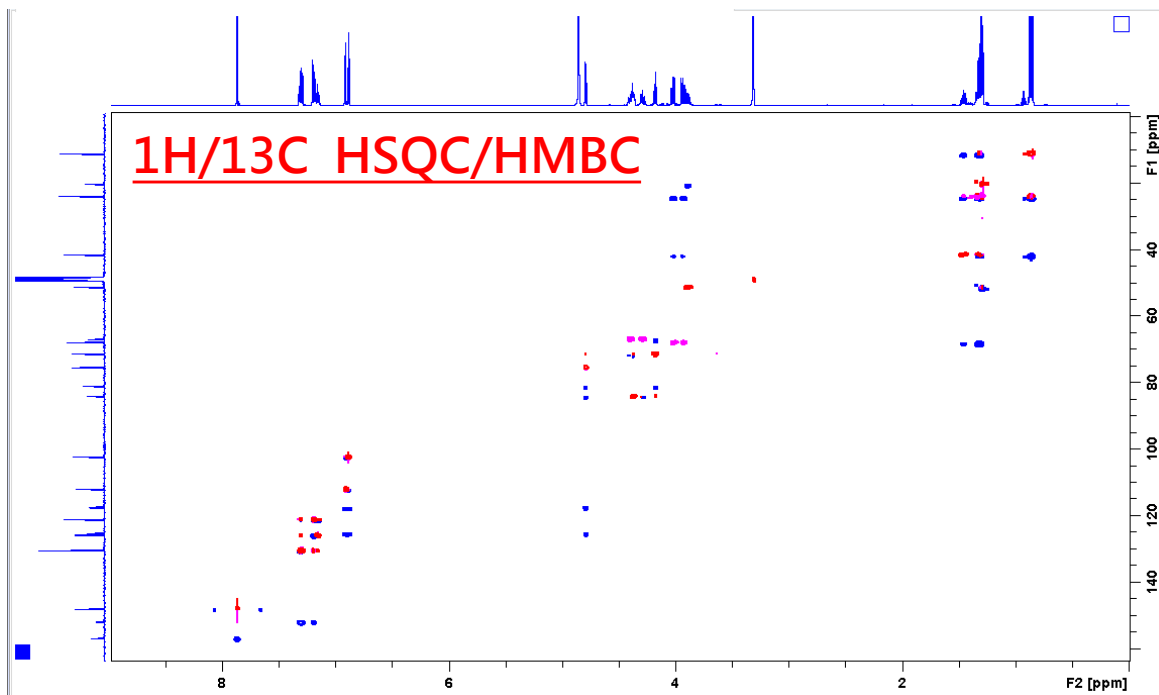
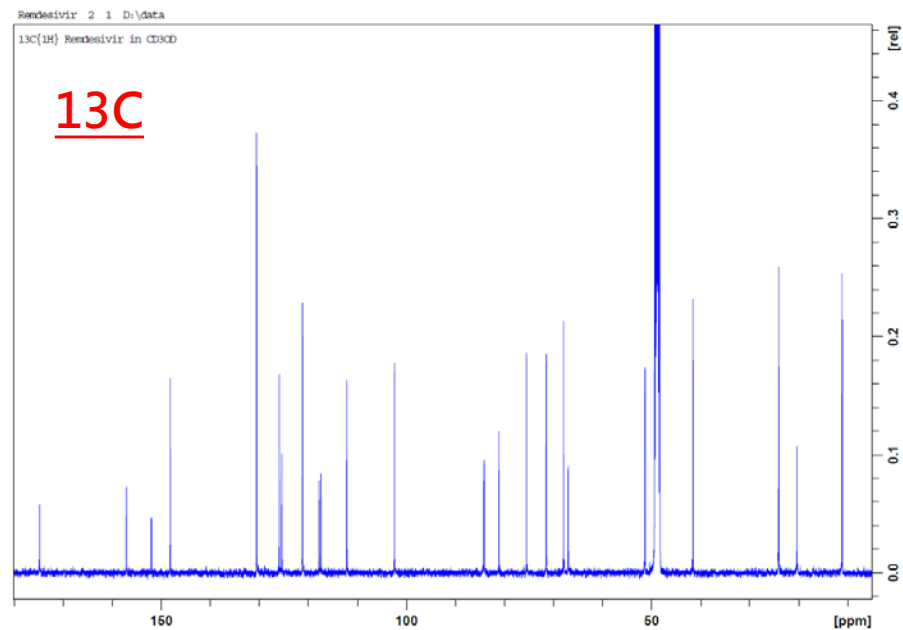
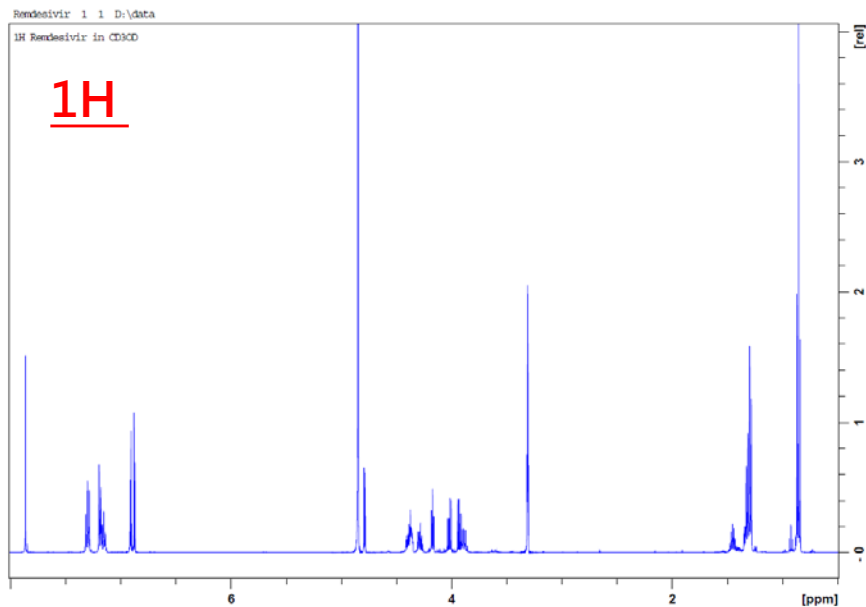


- ✓ Structure identification/ elucidation for **small molecules, nature products**.....
- ✓ **Bimolecular** structure /Dynamics (ex: nucleic acids, peptides, proteins)
- ✓ Molecule-Molecule interaction ( **protein-ligand, protein-protein, protein-DNA ....**)
- ✓ Fragment Based Drug Screening
- ✓ In cell NMR / Metabolomic NMR
- ✓ Others (qNMR....)

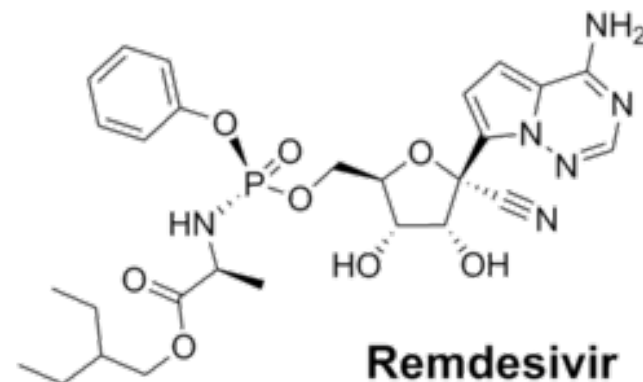
# Applications of NMR Spectroscopy



Example 1:  
Structure identification



Structure confirmed by NMR



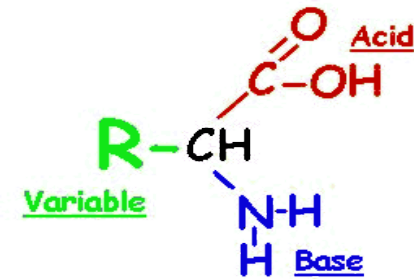
# Applications of NMR Spectroscopy



Example 2:  
Biomolecule Structure/Dynamics

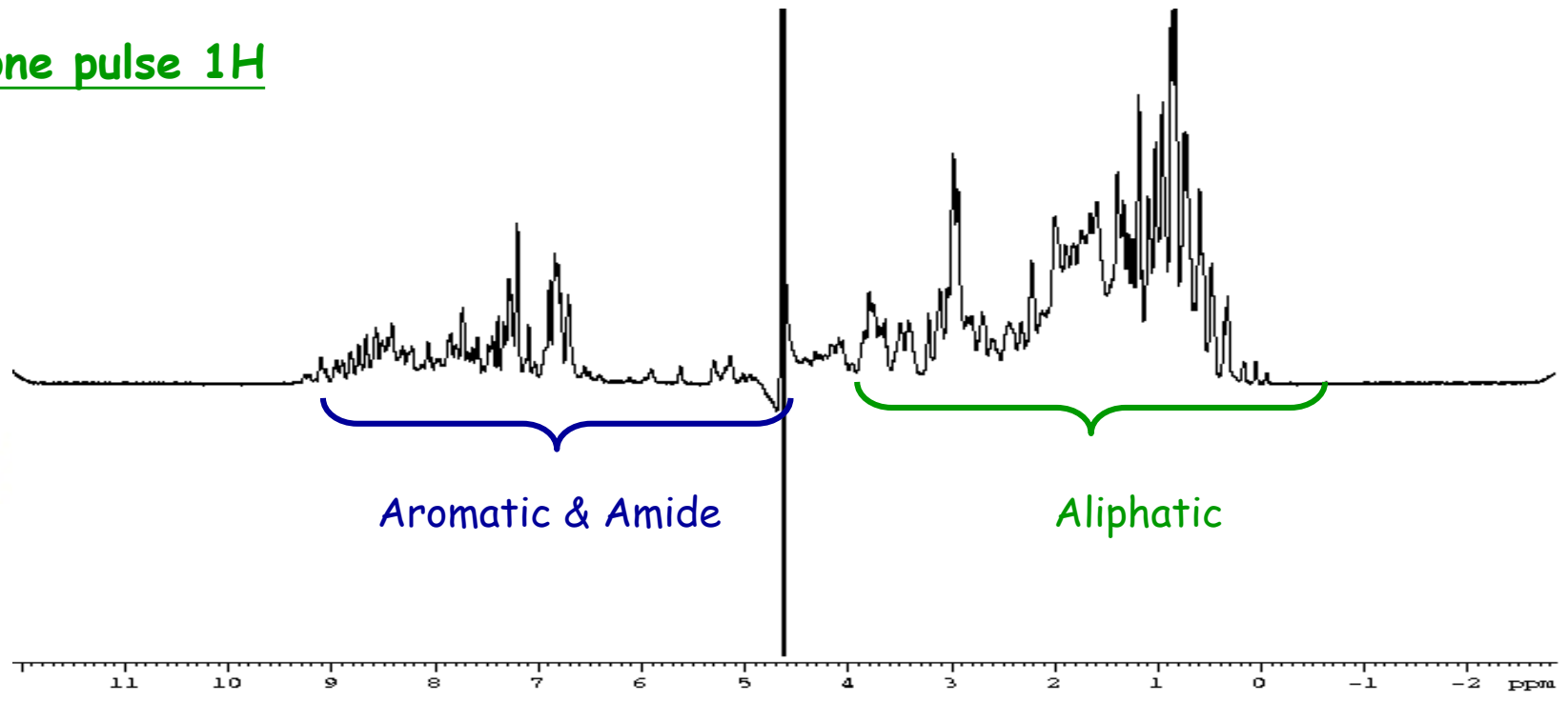


# Amino Acids to Proteins

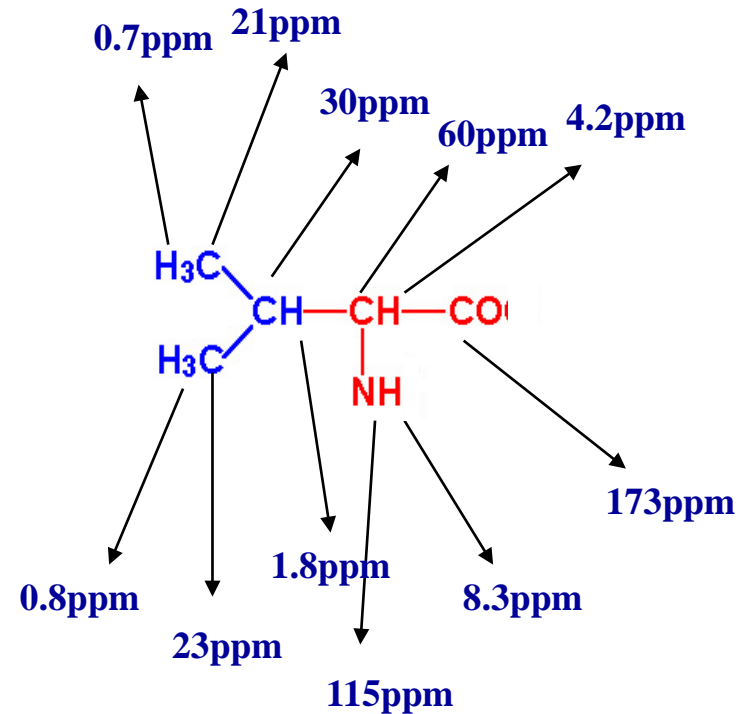
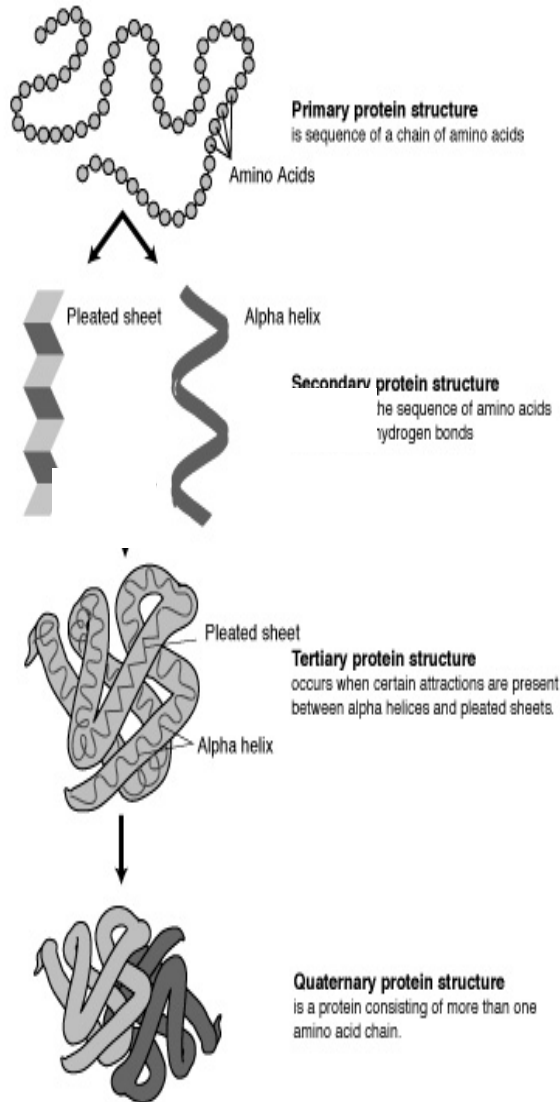


**A:** Ala, **C:** Cys, **D:** Asp, **E:** Glu, **F:** Phe, **G:** Gly, **H:** His,  
**I:** Ile, **K:** Lys, **L:** Leu, **M:** Met, **N:** Asn, **P:** Pro, **Q:** Gln,  
**R:** Arg, **S:** Ser, **T:** Thr, **V:** Val, **W:** Trp, **Y:** Tyr

## 1D one pulse 1H



# Sequence-specific and total resonance assignment



Try to assign Protein residues chemical shift as many as possible



Protein in solution 0.3~0.5ml, ~0.1~1mM

Non-labeled

$^{15}\text{N}$  labeled

$^{13}\text{C}/^{15}\text{N}$  labeled

$^2\text{H}/^{15}\text{N}$  labeled

$^2\text{H}/^{15}\text{N}/^{13}\text{C}$  labeled

Selected &  
Other specific labeled

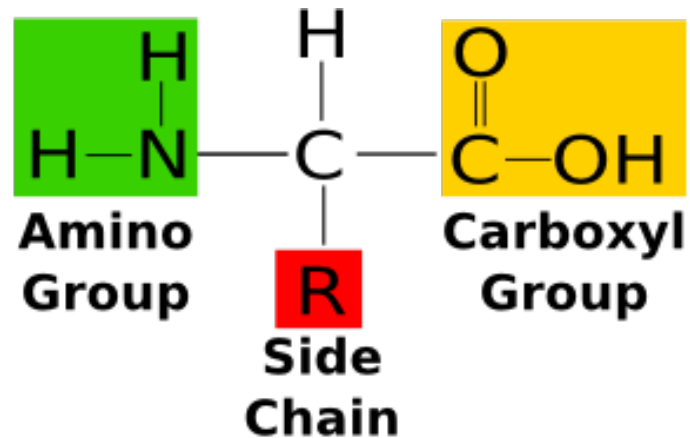
Protein Size Limitation :

(1) Slow tumbling problem

(2) Signal overlapping problem

$^{13}\text{C}$ : Nature abundance 1.1 %

$^{15}\text{N}$ : Nature abundance 0.4%



Understand the limitation, find the best condition!

# Workflow for Protein NMR Studies

Protein in solution 0.3~0.5ml, ~0.1~1mM

Labelled Proteins

NMR spectroscopy ( Select Useful/Specific NMR Experiments)

Sequence-specific/Backbone &  
Total resonance assignment

Secondary Structure  
Residue Specific Interaction  
Backbone Dynamics

Determining conformational constraints

Calculation of initial structure

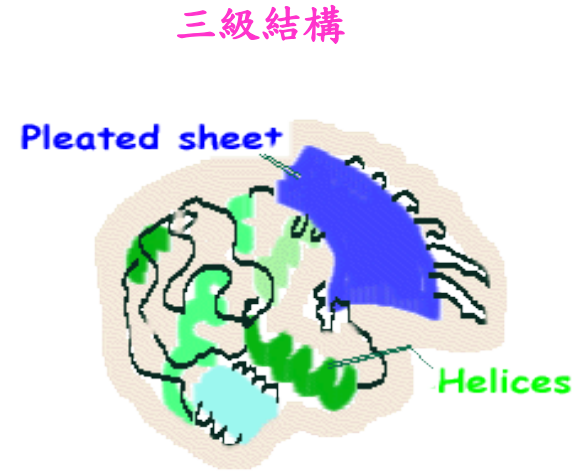
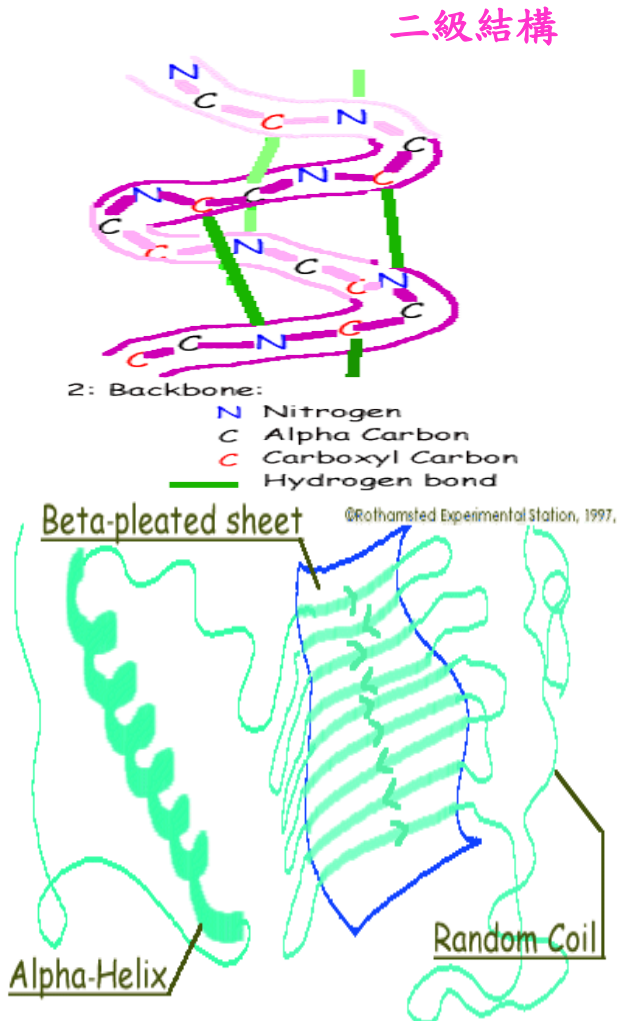
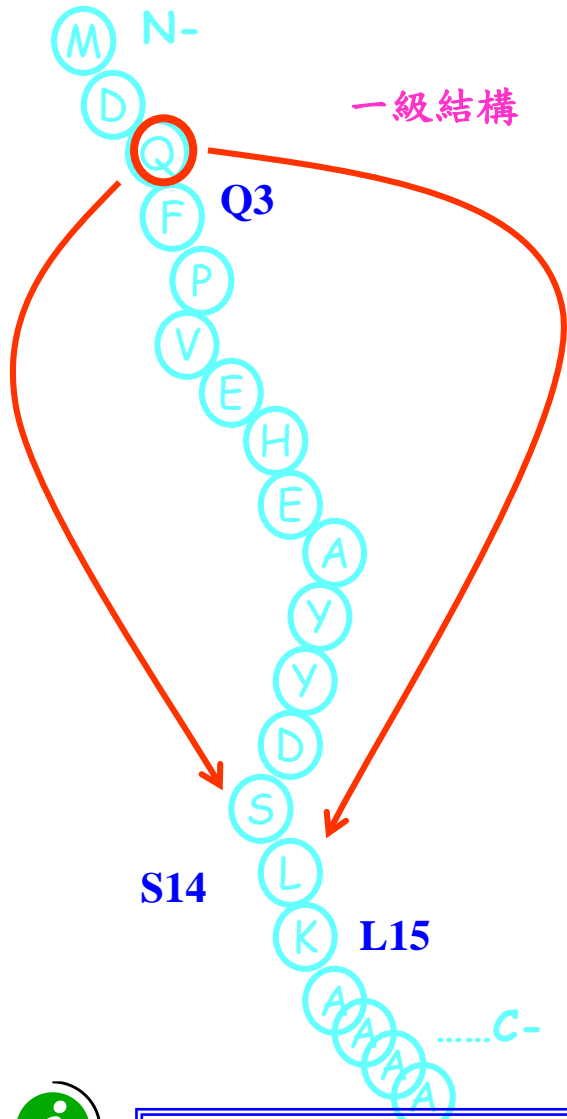
- NOE: distance restraints
- J-coupling: dihedral angle
- Others: H-bonds.....

Structure refinement

Structure calculation Soft wares

Final 3D Solution NMR Structure / Structure-Based Studies

# Determine conformational constraints



The tertiary structure is the way the secondary structures fold onto themselves to form a protein or a subunit of a more complex protein.

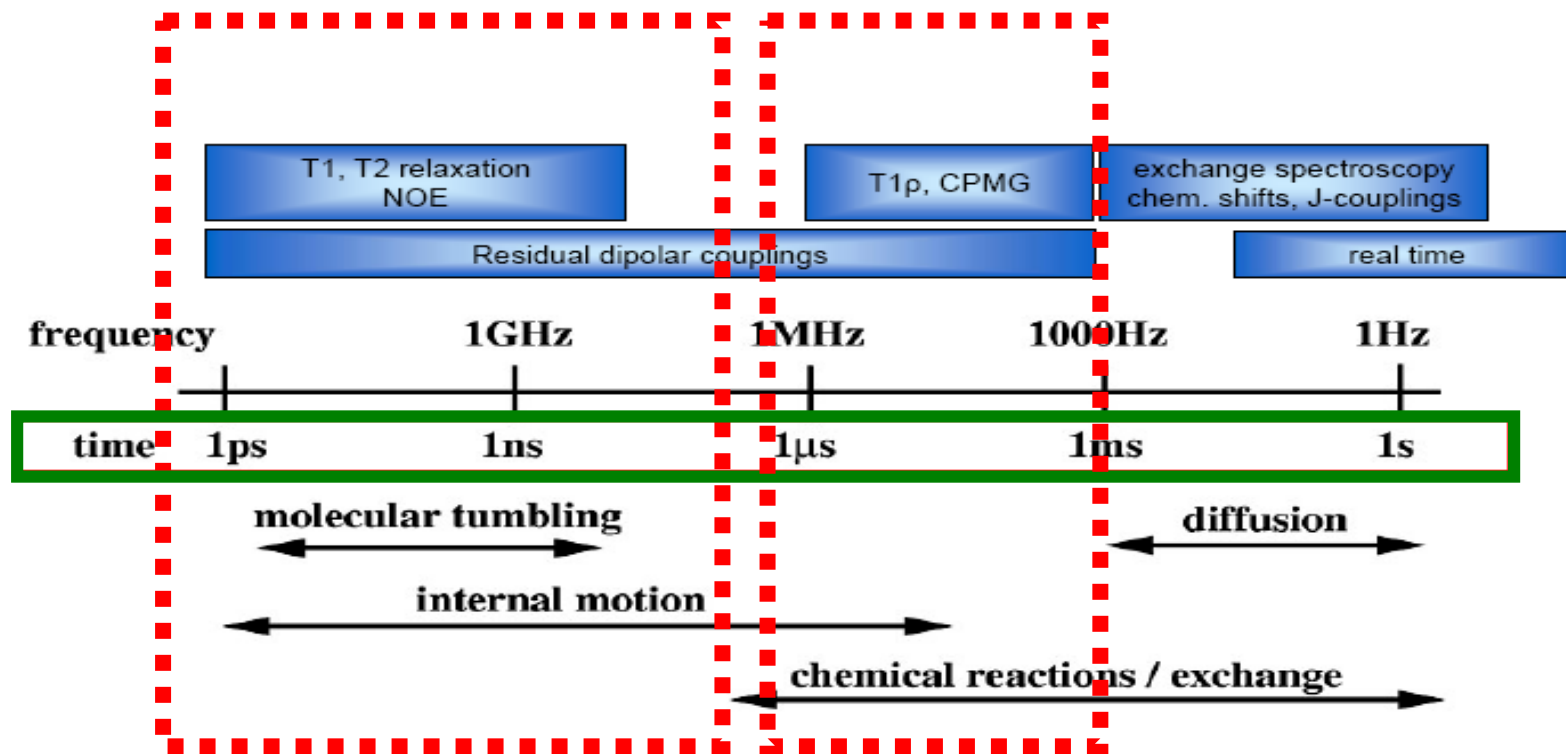
©Rothamsted Experimental Station, 1997, 1998



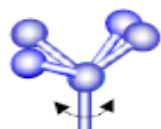
Obtain "distance or angle" information between atoms or bonds

# NMR Time Scales for Protein Dynamics Studies

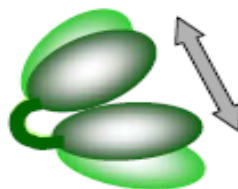
(From <http://www.embl.de/nmr/sattler/teaching>)



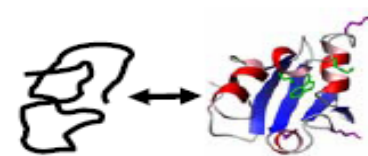
*bond vibrations*



*side chain rotations*



*domain movements*



*enzyme kinetics*

*folding, H/D exchange*

# Applications of NMR Spectroscopy



Example 3:

Molecule-Molecule interaction

/Fragment-based screening by NMR



## Our approach for NMR screening

"Free" Compound Groups

- 1D  $^1\text{H}$  spectrum
- $T_{1\rho}$  spectrum

- 0.2mM compounds
- PBD or d18-Tris Buffer @pH7.0

Compound Groups +  
Target Protein

- 1D  $^1\text{H}$  spectrum
- $T_{1\rho}$  spectrum
- WaterLOGSY spectrum
- STD spectrum

- 0.2mM compounds + ~0.01mM Protein
- PBD or d18-Tris Buffer @pH7.0

Find the "Hit Groups"

Analyze all spectrum  
Free vs. Complex

Narrow down to "individual Hits"



## Our approach for NMR screening

"Free" Protein NMR

- Backbone Assignments
- $^{15}\text{N}$  labelled protein

- 0.1mM  $^{15}\text{N}$ -labelled Protein
- PBD or d18-Tris Buffer @pH7.0

Target Protein +  
Hit Compounds

- 1D  $^1\text{H}$  spectrum
- 2D  $^{15}\text{N}$ - $^1\text{H}$  HSQC

- ~0.1mM Protein + compounds (titration)
- PBD or d18-Tris Buffer @pH7.0

Titration to find  
" Activity Site"

Analyze Chemical  
Shift perturbation  
Free vs. Complex

SAR (structure-activity relationship ) by NMR

Provide information for  
Compound Optimization

# Applications of NMR Spectroscopy



Example 4:  
Metabolomics



# Human related Metabolomics NMR Studies

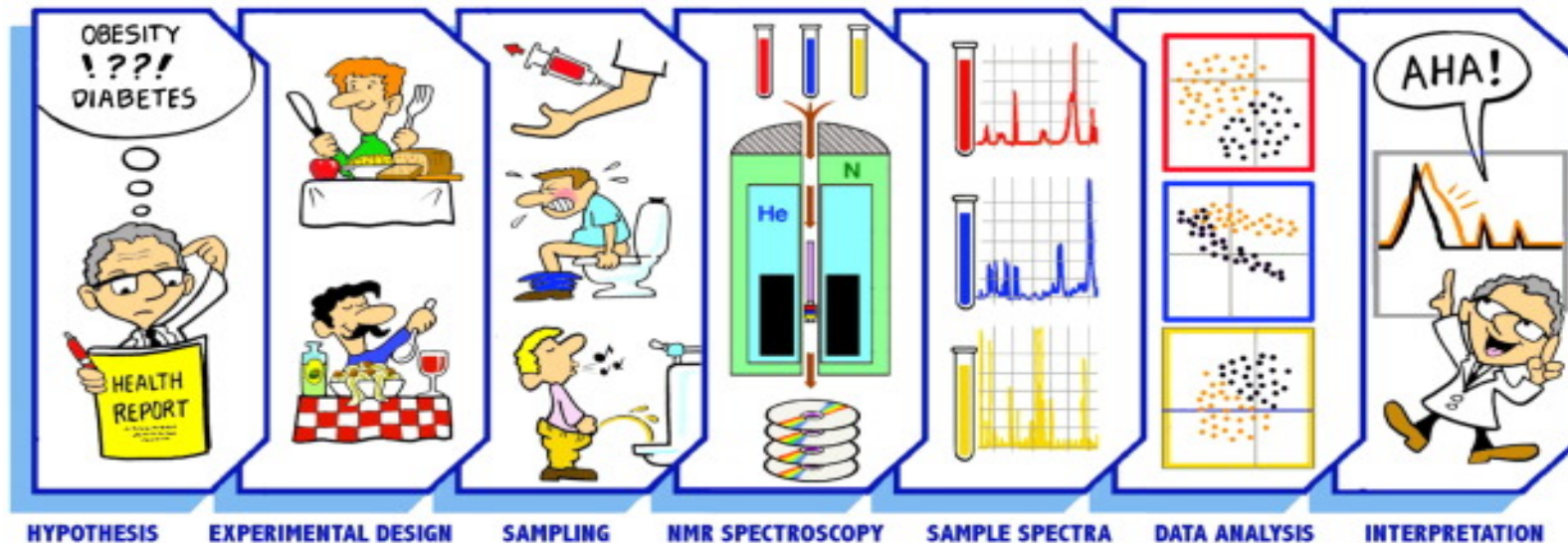


Fig from Food Research International 54 (2013) 1131–1145

# Plant Metabolomics Studies

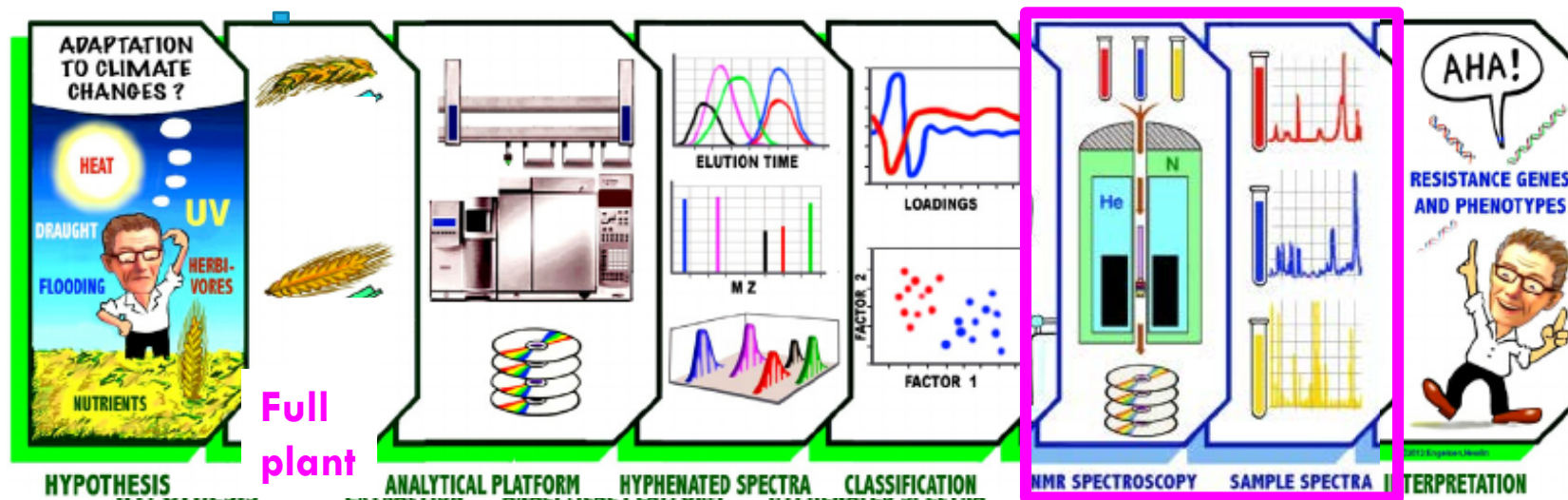


Fig from Journal of Cereal Science Volume 59, Issue 3, 393-418

# Different Techniques are Needed for Research

## NMR can provide information @ atomic resolution

