

Basic Operation for NMR Systems in Core Facility

Basic NMR Concepts & Experimental Set Up

by

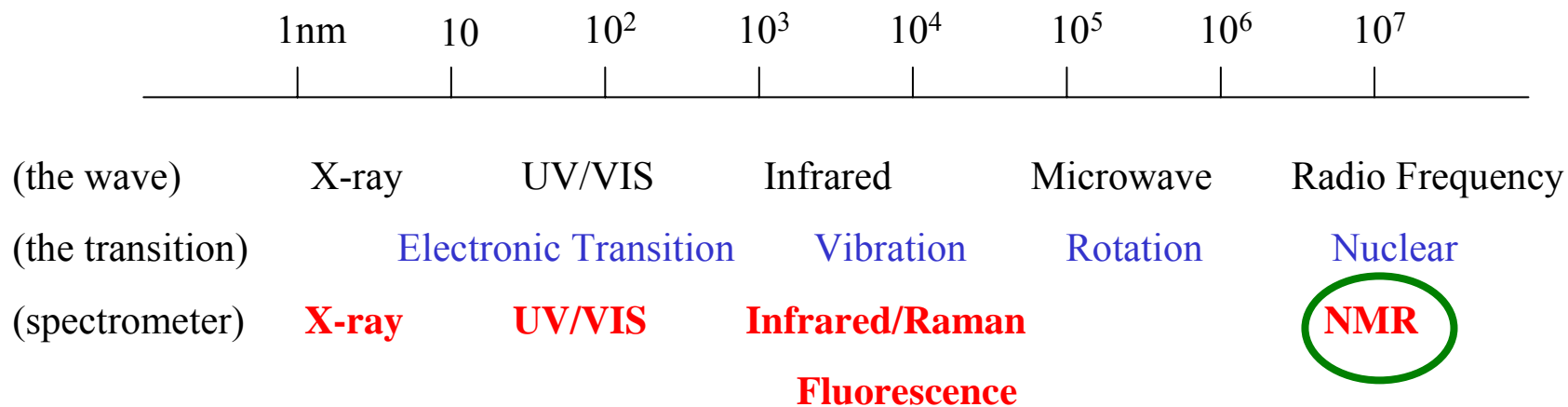
Chi-Fon Chang

09/29/2003

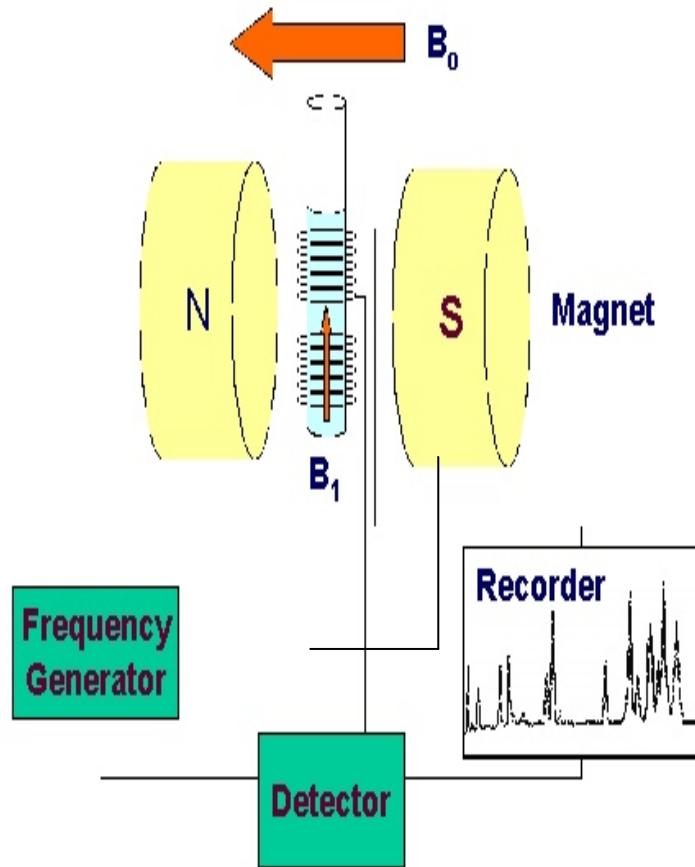
National Program for Genomic Medicine High Field NMR Core Facility,
The Genomic Research Center, Academia Sinica

NMR Spectroscopy

Where is it?



Nuclear Magnetic Resonance Spectrometer



B_0 : 光譜儀之磁場強度

B_1 : 外加小磁場 (來自樣品周圍之線圈)

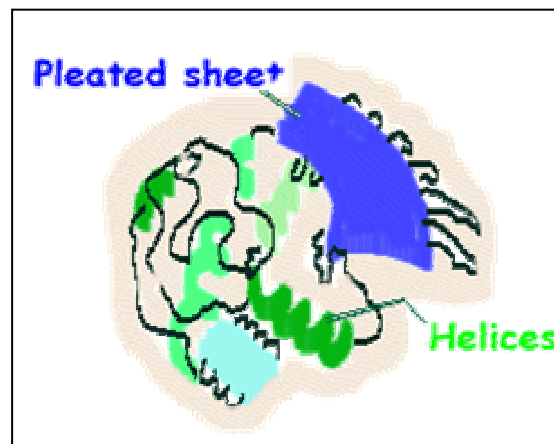
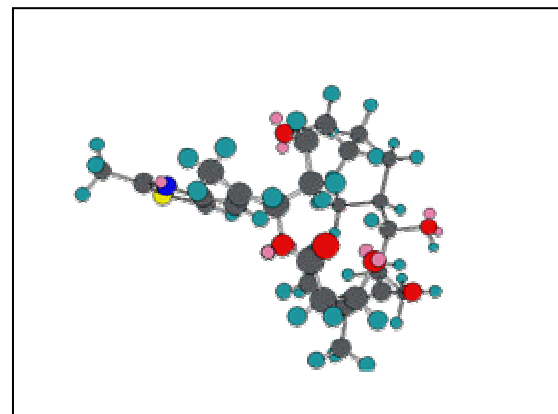
The problem the we want to solve

What we “really”
see



NMR

What we want to “see”



Steps for NMR Experiment

取得樣品

取得NMR圖譜

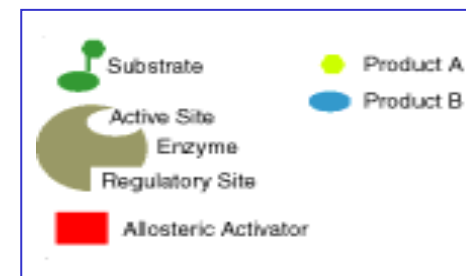
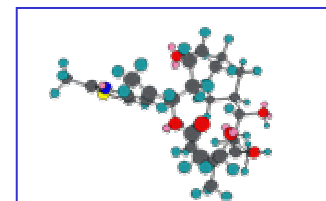
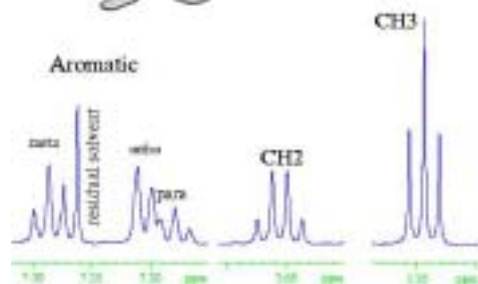
分析圖譜結果

適當的實驗方法

標定NMR譜線

鑑定化學(生化)分子

分子結構、動力學、
反應機制



Before using NMR
What's N, M, and R?

Properties of the Nucleus

Nuclear spin

Nuclear magnetic moments

The Nucleus in a Magnetic Field

Precession and the Larmor frequency

Nuclear Zeeman effect & Boltzmann distribution

When the Nucleus Meet the right Magnet

Nuclear Magnetic Resonance

◆ Properties of the Nucleus

Nuclear spin

- Nuclear spin is the total nuclear angular momentum quantum number. This is characterized by a quantum number I , which may be integral, half-integral or 0.
- Only nuclei with spin number $I \neq 0$ can absorb/emit electromagnetic radiation. The magnetic quantum number m_I has values of $-I, -I+1, \dots, +I$.
(e.g. for $I=3/2$, $m_I=-3/2, -1/2, 1/2, 3/2$)

1. A nucleus with an even mass A and even charge $Z \rightarrow$ nuclear spin I is zero

Example: ^{12}C , ^{16}O , $^{32}\text{S} \rightarrow$ No NMR signal

2. A nucleus with an even mass A and odd charge $Z \rightarrow$ integer value I

Example: ^2H , ^{10}B , $^{14}\text{N} \rightarrow$ NMR detectable

3. A nucleus with odd mass $A \rightarrow I=n/2$, where n is an odd integer

Example: ^1H , ^{13}C , ^{15}N , $^{31}\text{P} \rightarrow$ NMR detectable

Nuclear magnetic moments

Magnetic moment μ is another important parameter for a nuclei

$$\mu = \gamma I (h/2\pi)$$

I : spin number

h : Plank constant 6.626×10^{-34} joule-sec

γ : gyromagnetic ratio (property of a nuclei)

^1H : $I=1/2$, $\gamma = 267.512 \times 10^6 \text{ rad T}^{-1}\text{sec}^{-1}$

^{13}C : $I=1/2$, $\gamma = 67.264 \times 10^6$

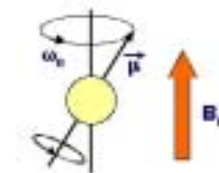
^{15}N : $I=1/2$, $\gamma = 27.107 \times 10^6$

Precession and the Larmor frequency

- The magnetic moment of a spinning nucleus precesses with a characteristic angular frequency called the Larmor frequency ω , which is a function of r and B_0

Larmor frequency $\omega = \gamma B_0$

Linear precession frequency $\nu = \omega/2\pi = \gamma B_0/2\pi$



Example: At what field strength do ^1H precess at a frequency of 600.13 MHz? What would be the precession frequency for ^{13}C at the same field?

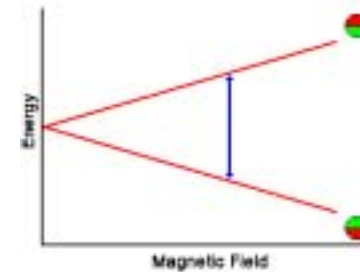
◆ The Nucleus in a Magnetic Field B_0

B_0 (the magnet of machine)

- (1) Provide energy for the nuclei to spin

$$E_i = -m_i B_0 \quad (\hbar/2\pi)$$

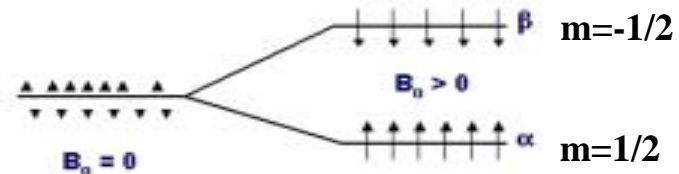
$$\text{Larmor precession } \nu = \omega/2\pi = \gamma B_0/2\pi$$



- (2) Induce energy level separation (Zeeman effect & Boltzmann distribution)

The stronger the magnetic field B_0 , the greater separation

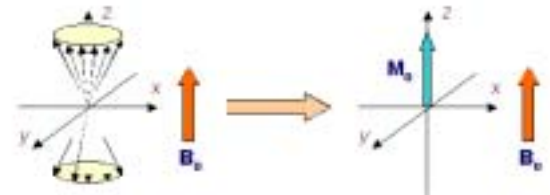
$$P_{m=-1/2} / P_{m=+1/2} = e^{-\Delta E/kT}$$



- (3) The nuclei in both spin states are randomly oriented around the z axis.

$$M_z = M_0, \quad M_{xy} = 0$$

(where M_0 is the net nuclear magnetization)

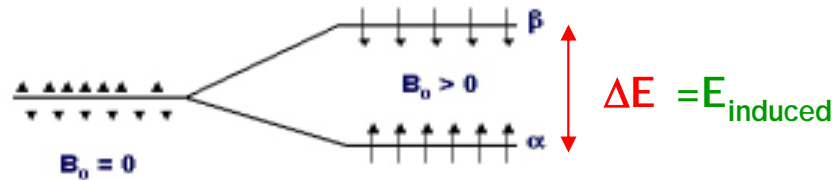


◆ When the Nucleus Meet the “right” Magnet: N. M. Resonance

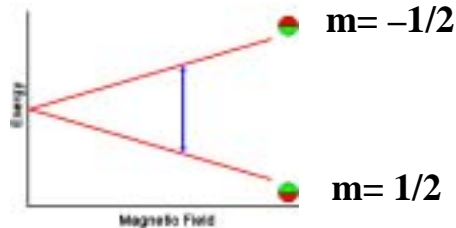
B_1 (the irradiation magnet, current induced)

(1) Induce energy for nuclei to absorb, but still spin at ω or $\nu_{\text{precession}}$

$$E_{\text{induced}} = \Delta E = \hbar \gamma B_0 / 2\pi = \hbar \nu_{\text{precession}}$$



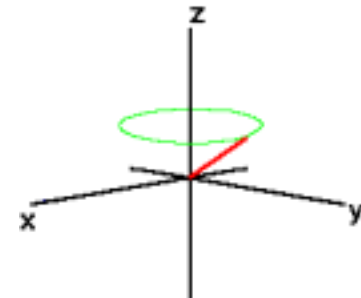
And now, the spin jump to the **higher energy** (from $m=1/2 \rightarrow m= -1/2$)



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

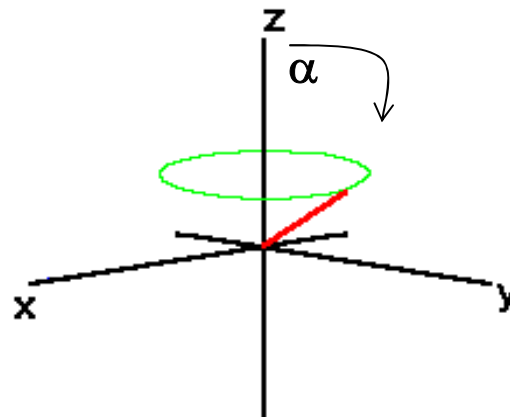
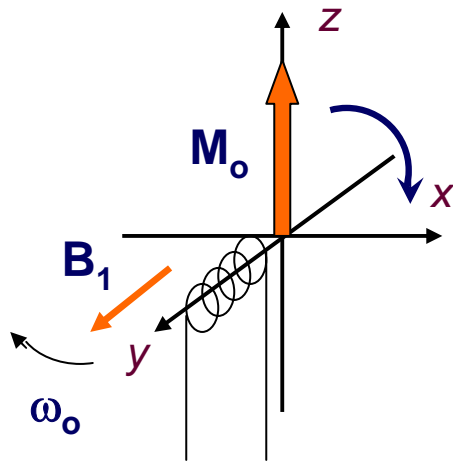
$$M_z = M \cos \alpha$$

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

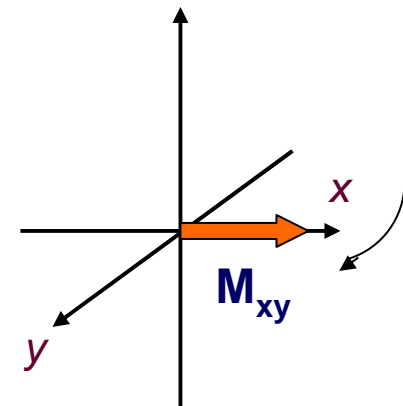


What happen during irradiation

When irradiation begins, all of the individual nuclear magnetic moments become **phase coherent**, and this phase coherence forces the net magnetization vector M_0 to process around the z axis. As such, M has a component in the x, y plan, $M_{xy}=M\sin\alpha$. α is the **tip angle** which is determined by the **power and duration of the electromagnetic irradiation**.



α deg pulse



90 deg pulse

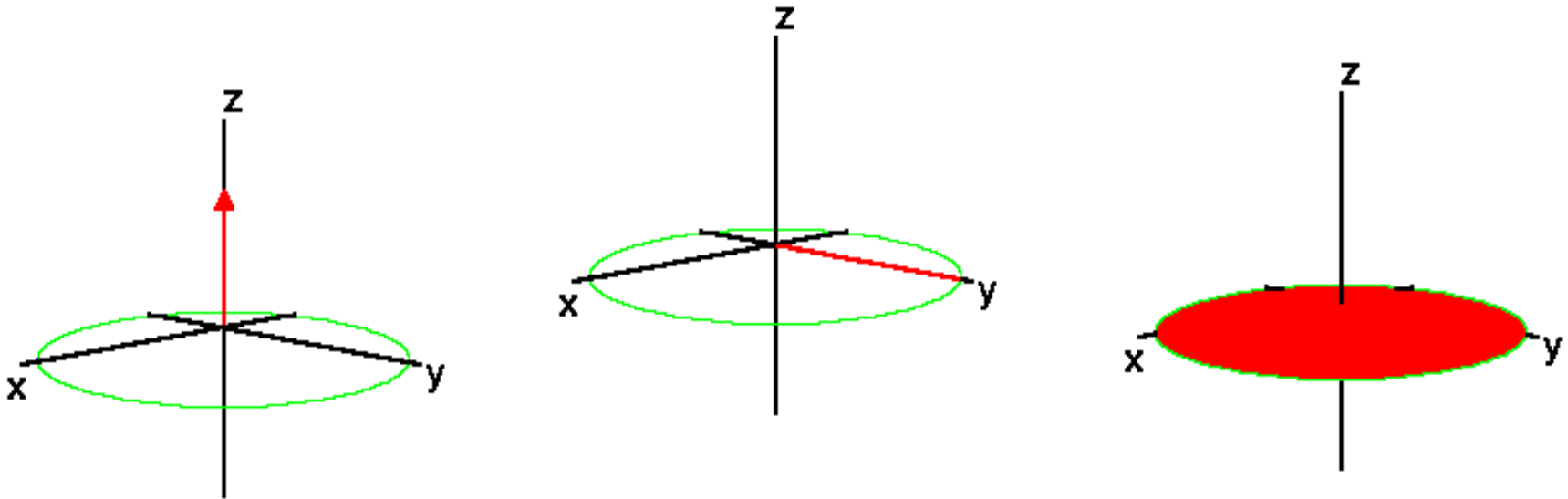
Hint: that's why we need to calibrate 90° pulse !!

What happen after irradiation ceases

- After irradiation ceases, not only do the population of the states revert to a **Boltzmann distribution**, but also the individual nuclear magnetic moments begin to lose their phase coherence and return to a **random** arrangement around the z axis.

(NMR 的光譜其實就是在紀錄這個過程!!)

- This process is called “relaxation process” (弛緩現象)
- There are two types of relaxation process :
 - T1 (spin-lattice relaxation)
 - T2 (spin-spin relaxation)



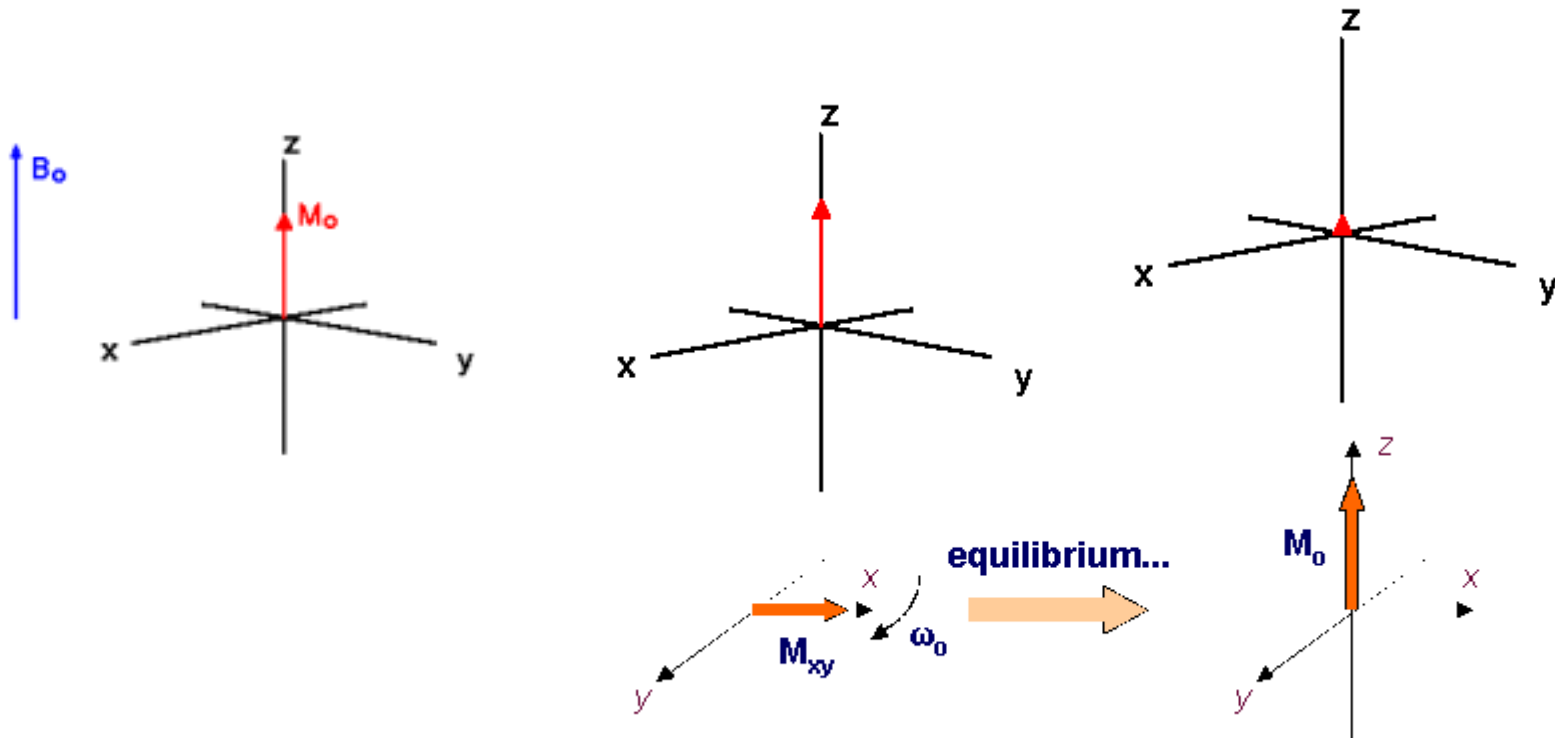
T1 (the spin lattice relaxation)

- How long after immersion in a external field does it take for a collection of nuclei to reach Boltzmann distribution is controlled by T1, the spin lattice relaxation time.

(考慮波茲曼分布的效應為主)

- Lost of energy in system to surrounding (lattice) as heat (能量釋放的過程)
- It's a time dependence exponential decay process of Mz components

$$dM_z/dt = -(M_z - M_{z,eq})/T1$$



T2 (the spin-spin relaxation)

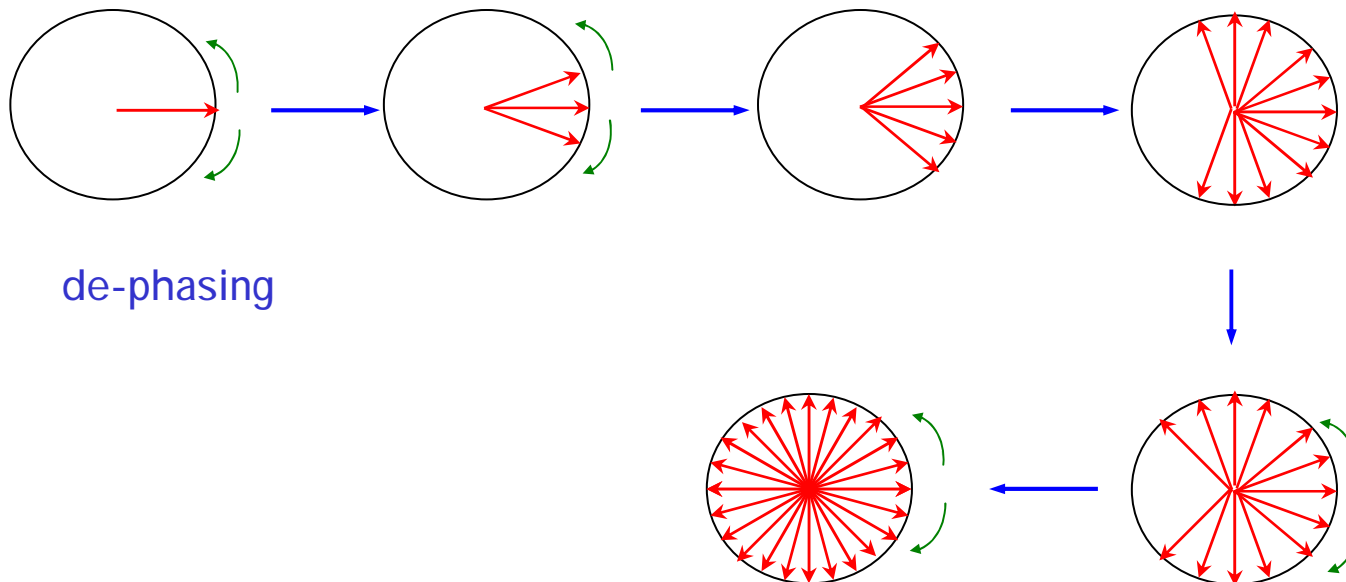
- This process for nuclei begin to lose their phase coherence and return to a random arrangement around the z axis is called spin-spin relaxation.

(考慮自旋方位由同一方向又回到 random 的過程)

- The decay of M_{xy} is at a rate controlled by the spin-spin relaxation time T2.

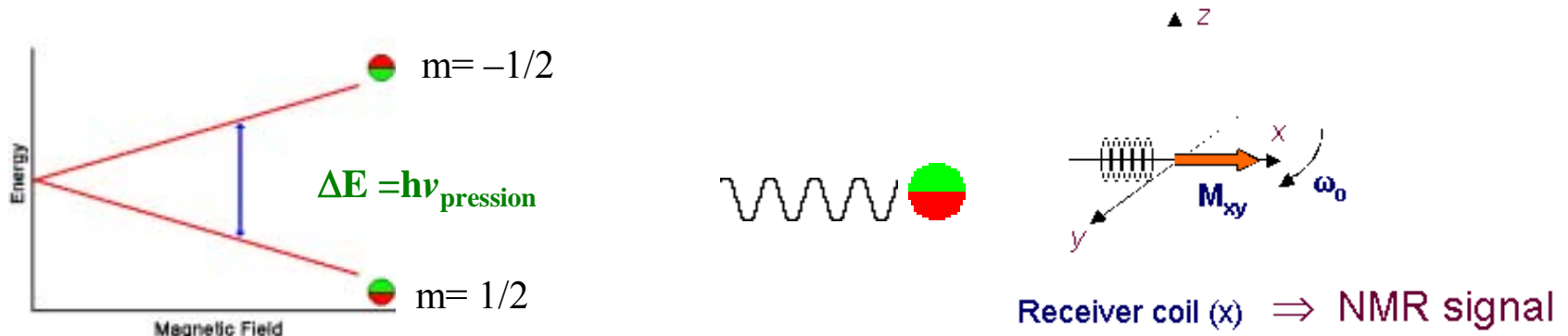
$$dM_x/dt = -M_x/T_2$$

$$dM_y/dt = -M_y/T_2$$

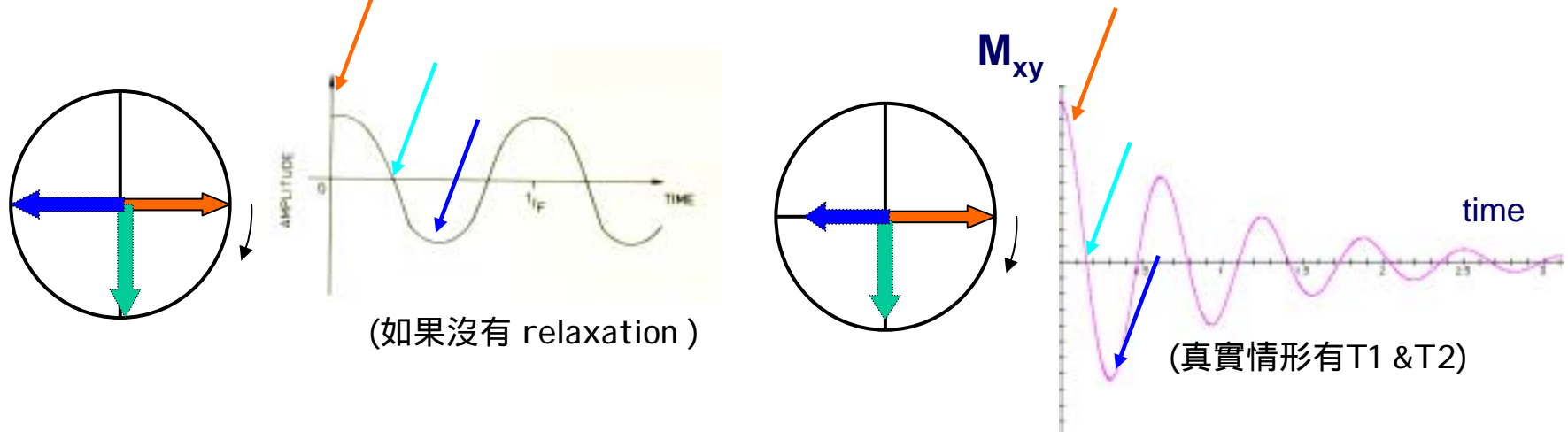


◆ Collecting NMR signals

• The detection of NMR signal is on the xy plane. The oscillation of M_{xy} generate a current in a coil, which is the NMR signal.

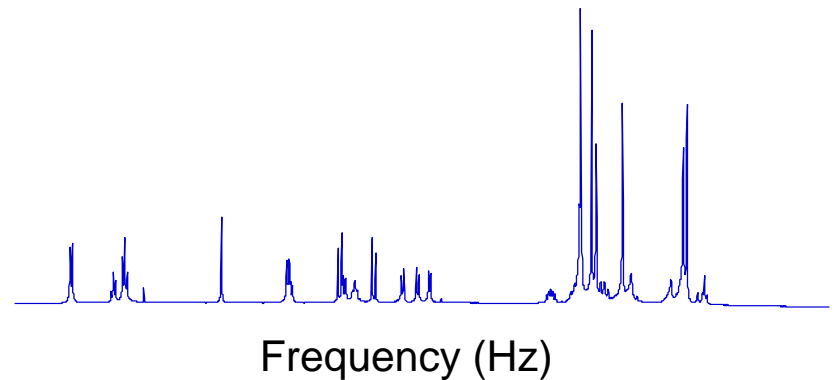
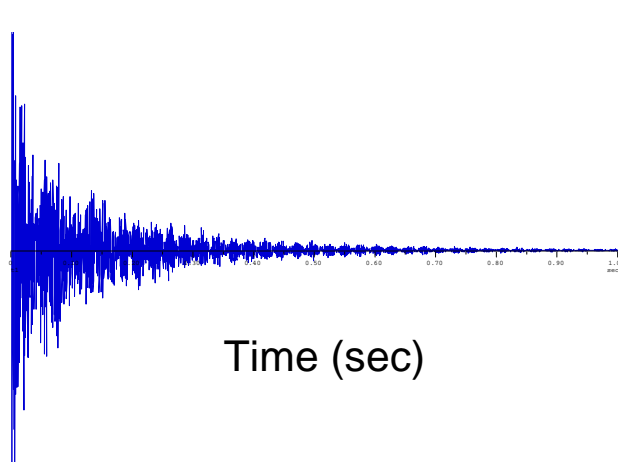
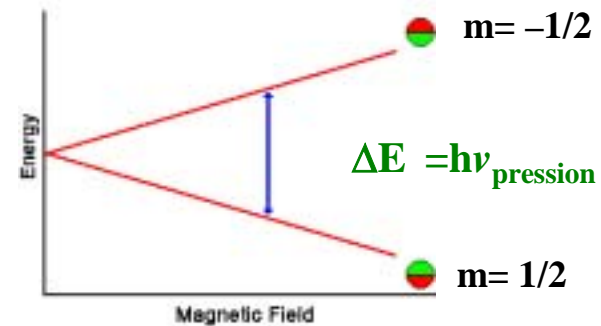
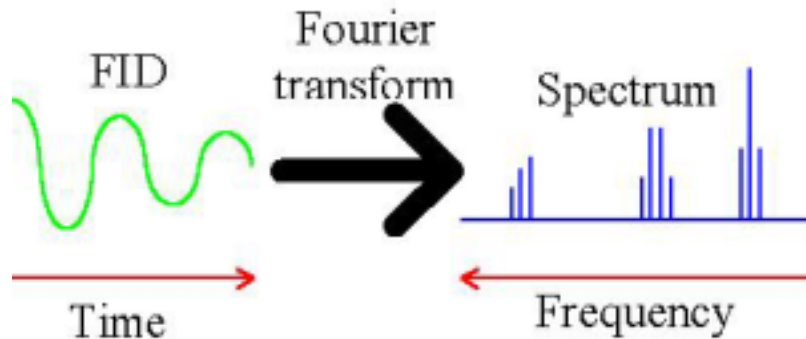


• Due to the "relaxation process", the time dependent spectrum of nuclei can be obtained. This time dependent spectrum is called "free induction decay" (FID)



NMR Data Processing

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



Example of NMR signals

- It's easy to understand that **different nucleus "type"** will **give different NMR signal**. ($\nu = \omega/2\pi = \gamma B_0/2\pi$, γ : gyromagnetic ratio is the property of a nuclei.)
- However, it is very important to know that for **same "nucleus type"**, but **"different nucleus"** could **generate different signal**. This is also what make NMR useful and interesting.
- **Electron surrounding each nucleus** in a molecule serves to **shield that nucleus** from the applied magnetic field. This shielding effect cause different ν in the spectrum

$B_{\text{eff}} = B_0 - B_i$ where B_i induced by cloud electron

$B_i = \sigma B_0$ where σ is the shielding constant

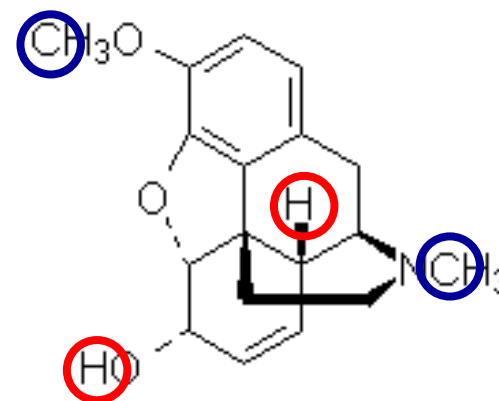
$B_{\text{eff}} = (1 - \sigma) B_0$

$\nu_{\text{precession}} = (r B_0 / 2\pi) (1 - \sigma)$

$\sigma = 0 \quad \rightarrow$ naked nuclei

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

$\sigma < 0 \quad \rightarrow$ electron around this nuclei is withdraw, i.e. deshielded



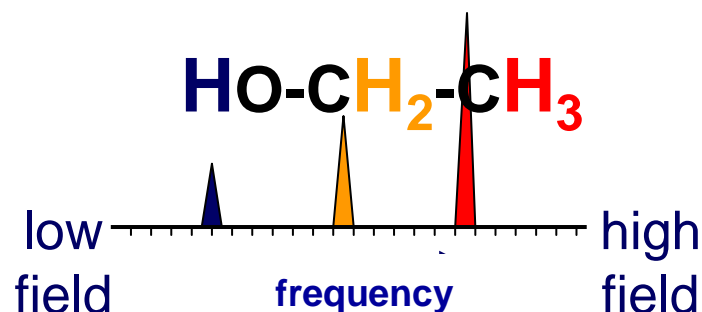
Basic of Assignment

$$\nu_{\text{precession}} = (\gamma B_0 / 2\pi) (1 - \sigma) = \nu_0 (1 - \sigma)$$

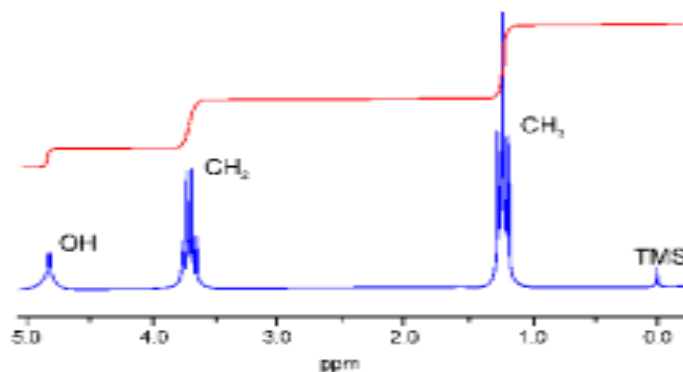
$\sigma = 0$ → naked nuclei

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

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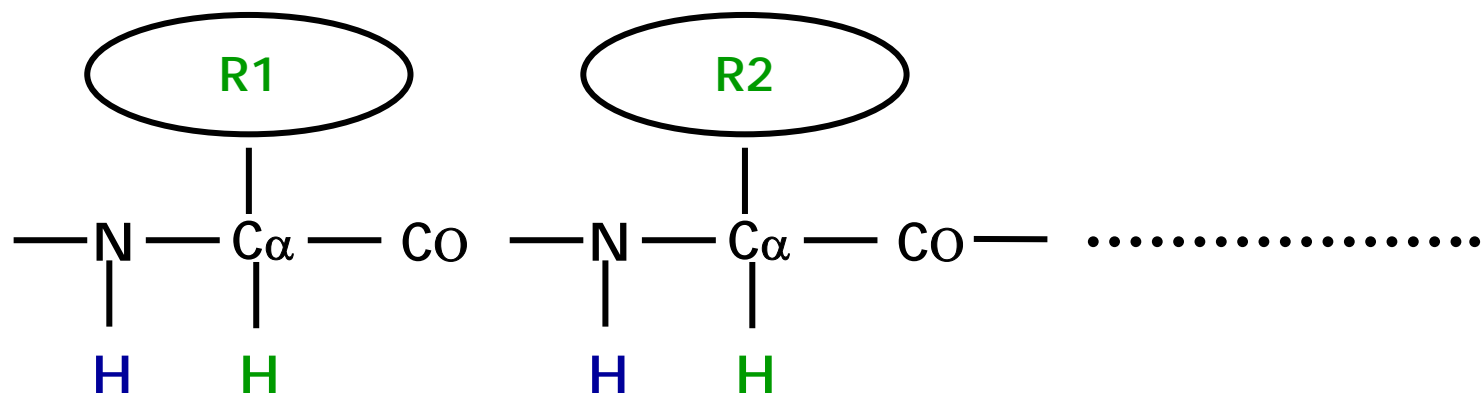
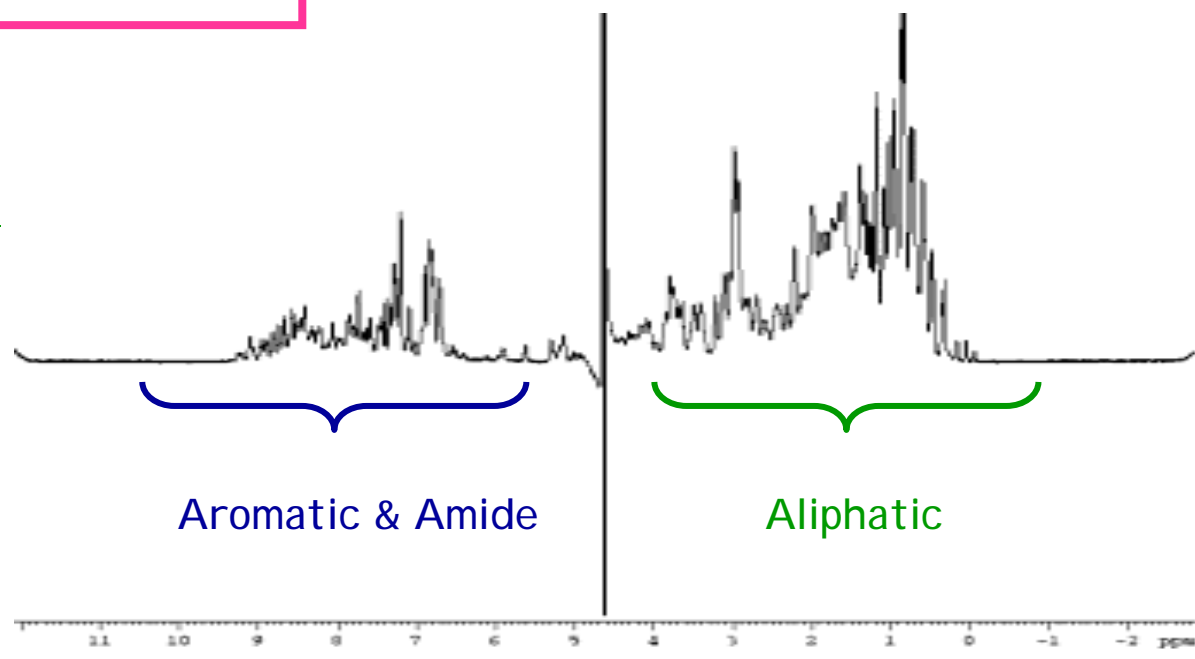
$$\text{ppm} \times 10^{-6} = \Delta\nu/\nu_0 = -\sigma$$



Introduction of NMR Experiments

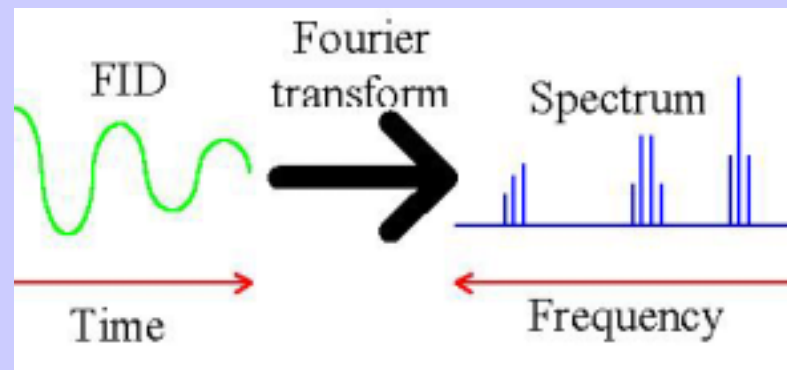
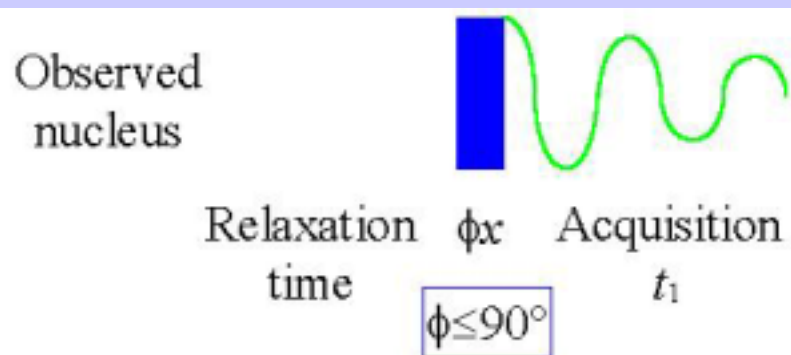
Homo Nuclear 1D NMR

1D one pulse 1H

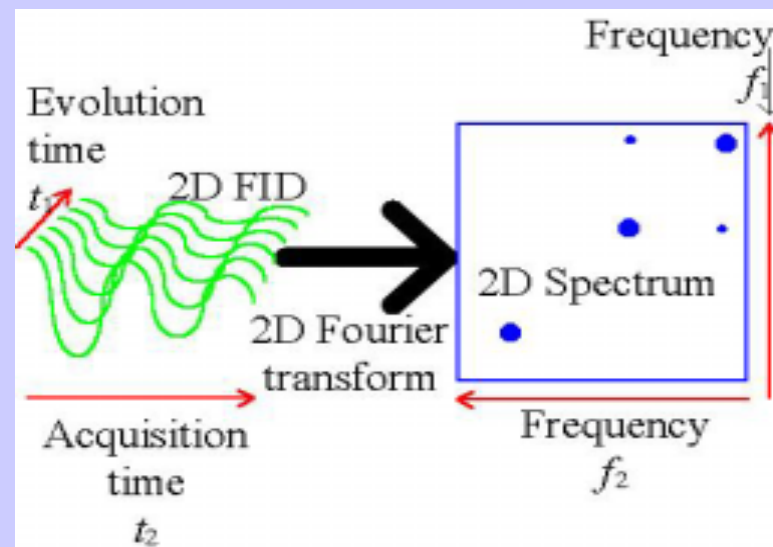
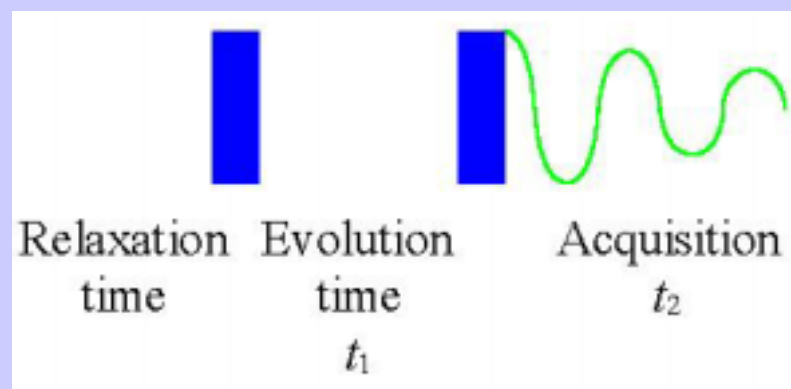


Homo/Hetro Nuclear 2D NMR

Basic 1D Experiment

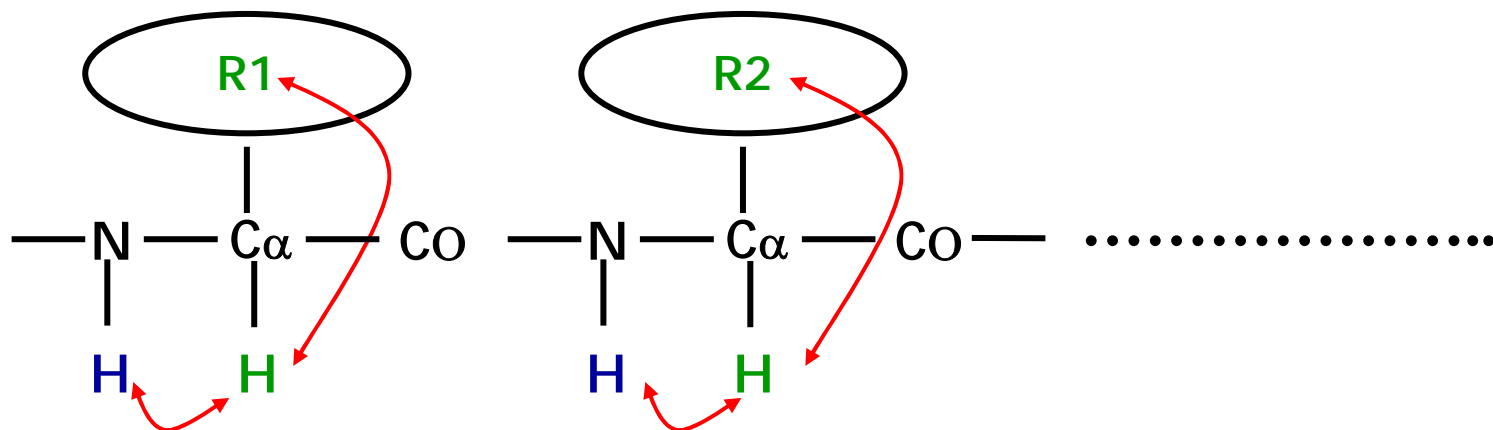
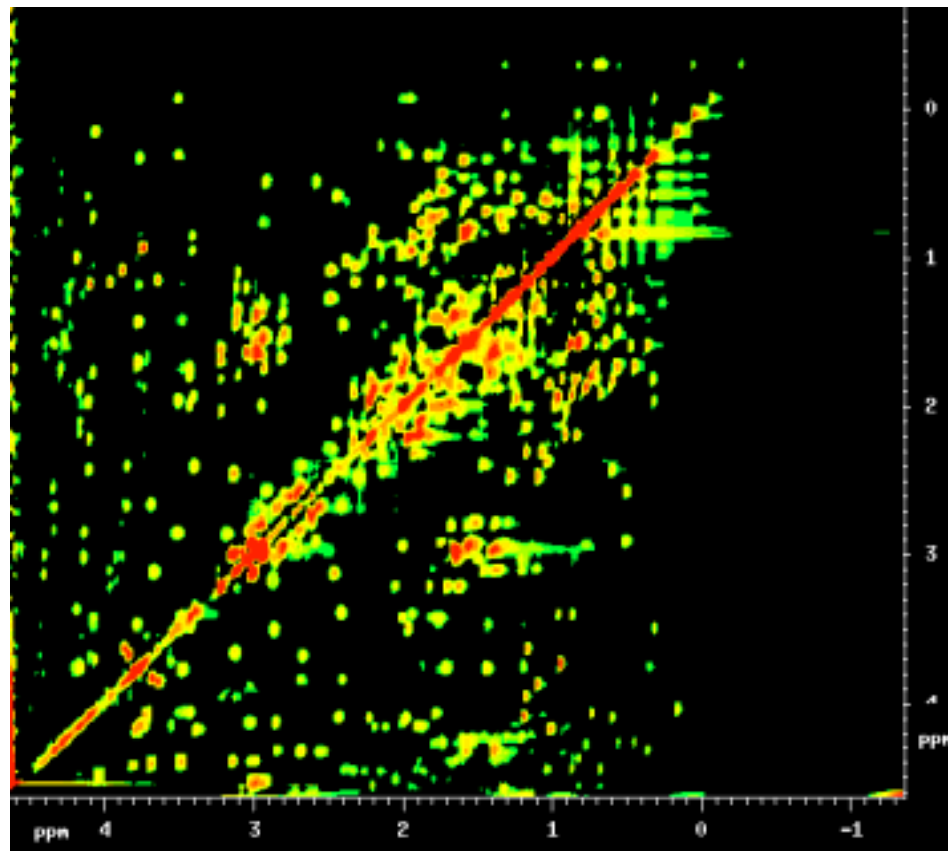
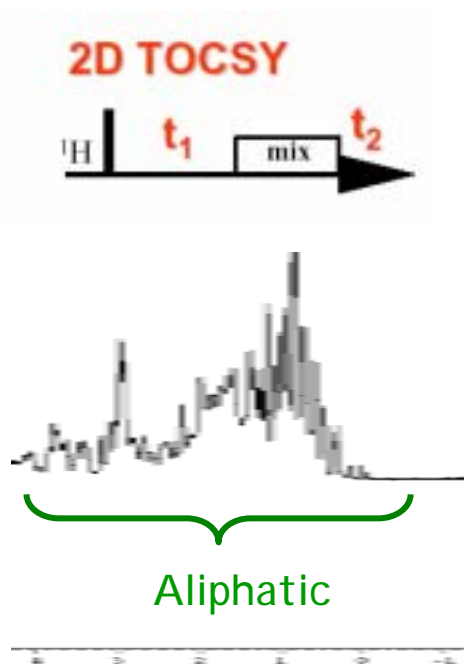


Basic 2D Experiment



2D Homo Nuclear

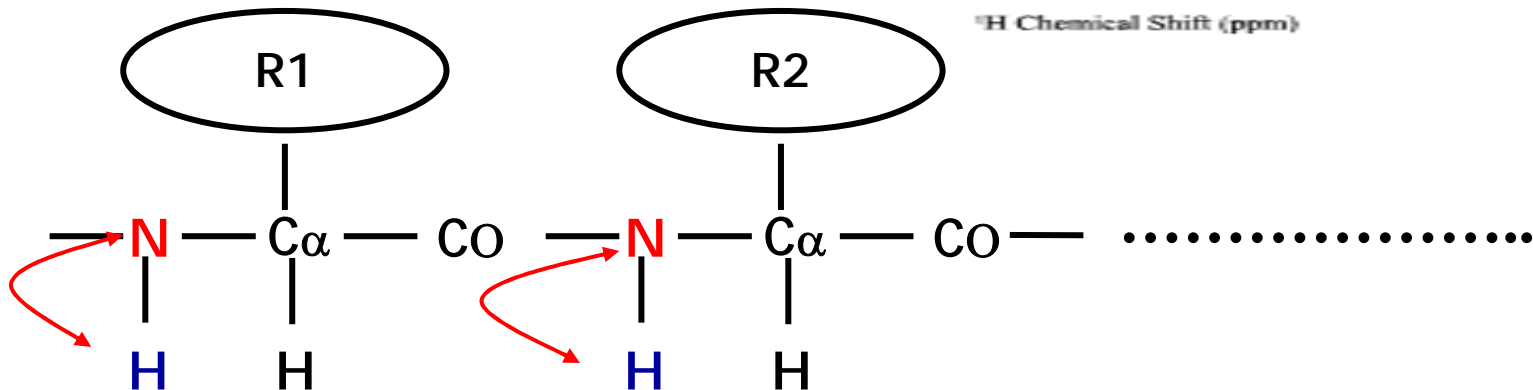
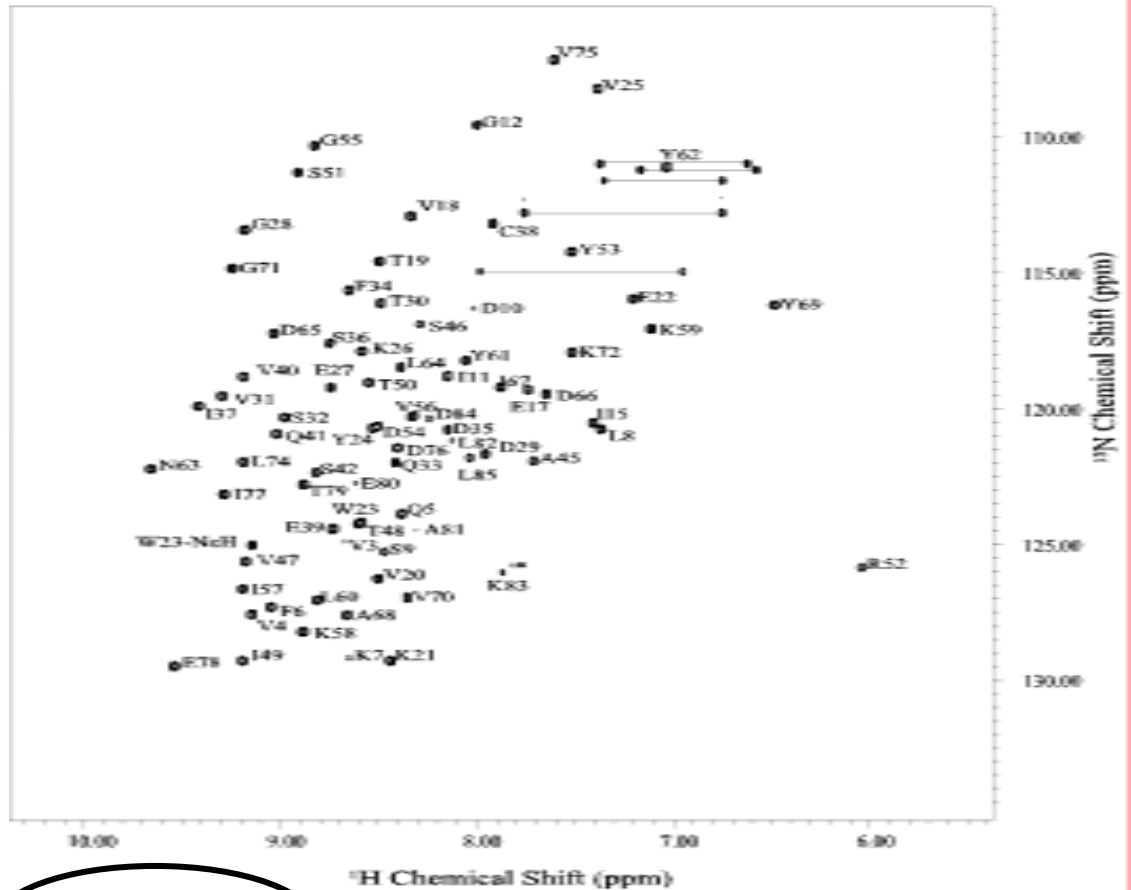
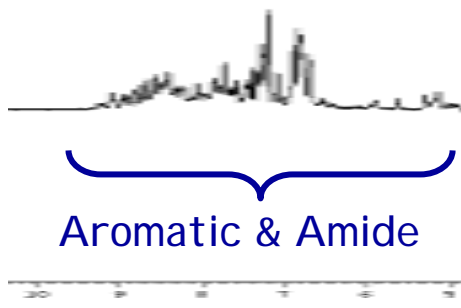
^1H - ^1H



2D Hetero Nuclear

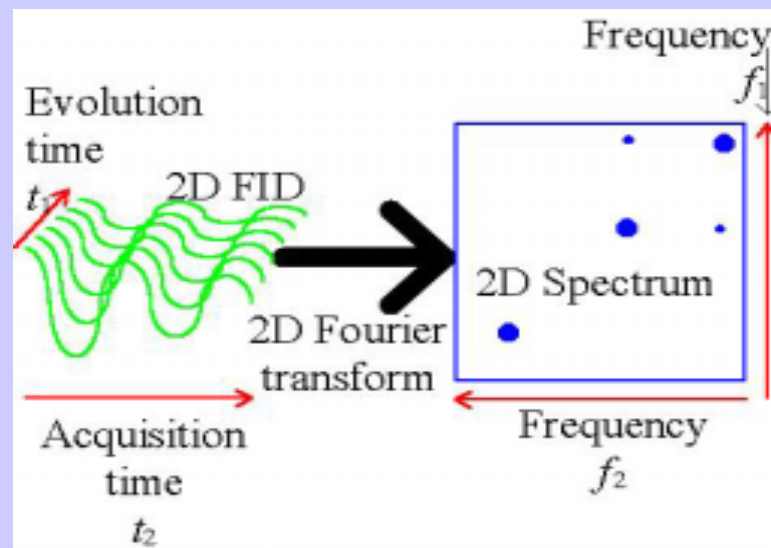
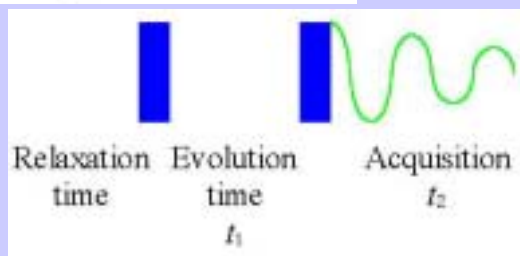
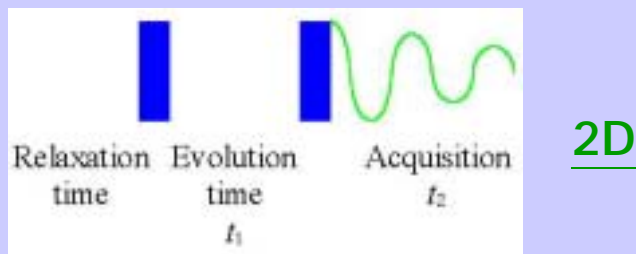
^1H - ^{15}N

2D HSQC

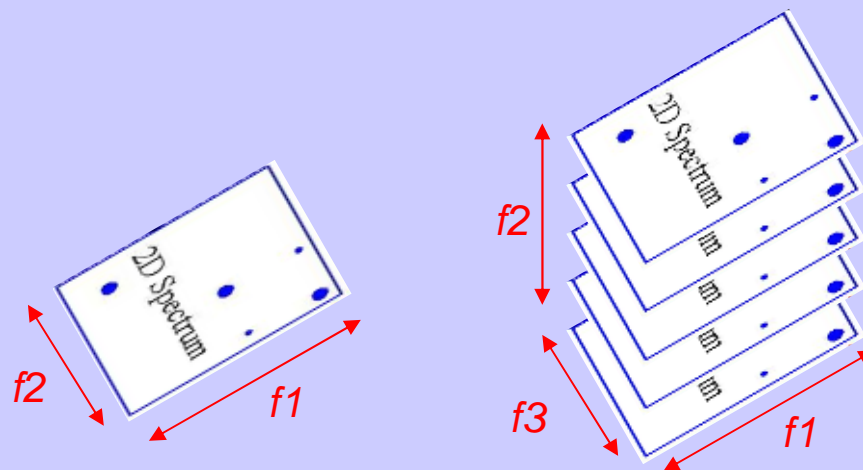
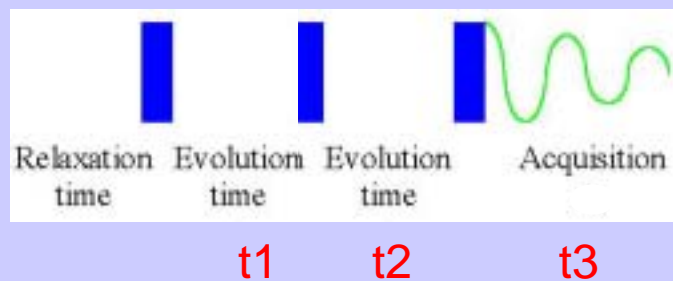


Multi-Dimensional NMR

Basic 3D Experiment



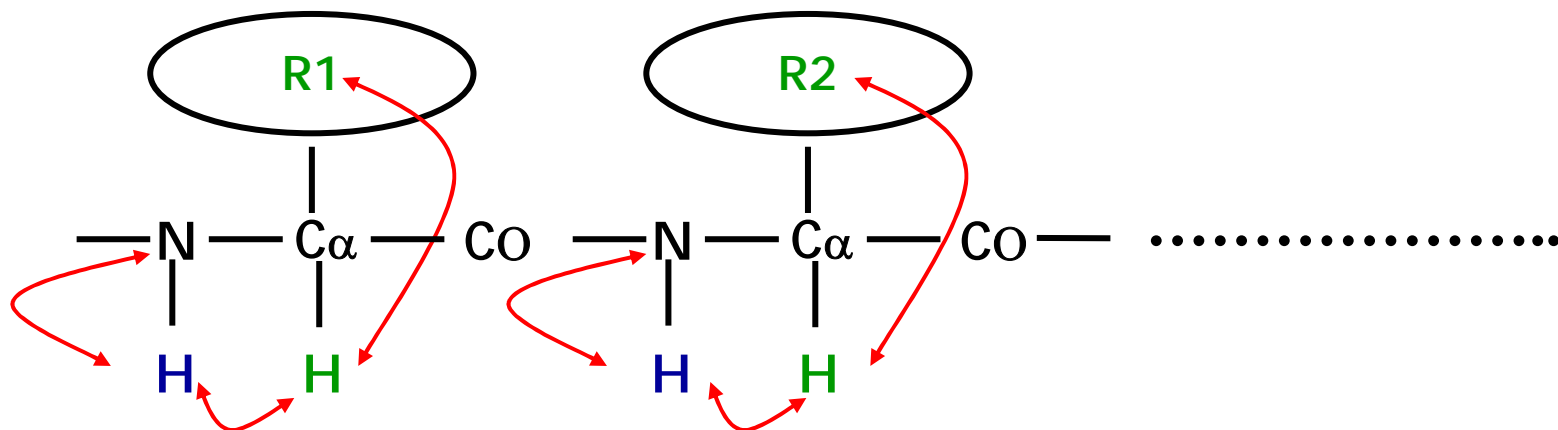
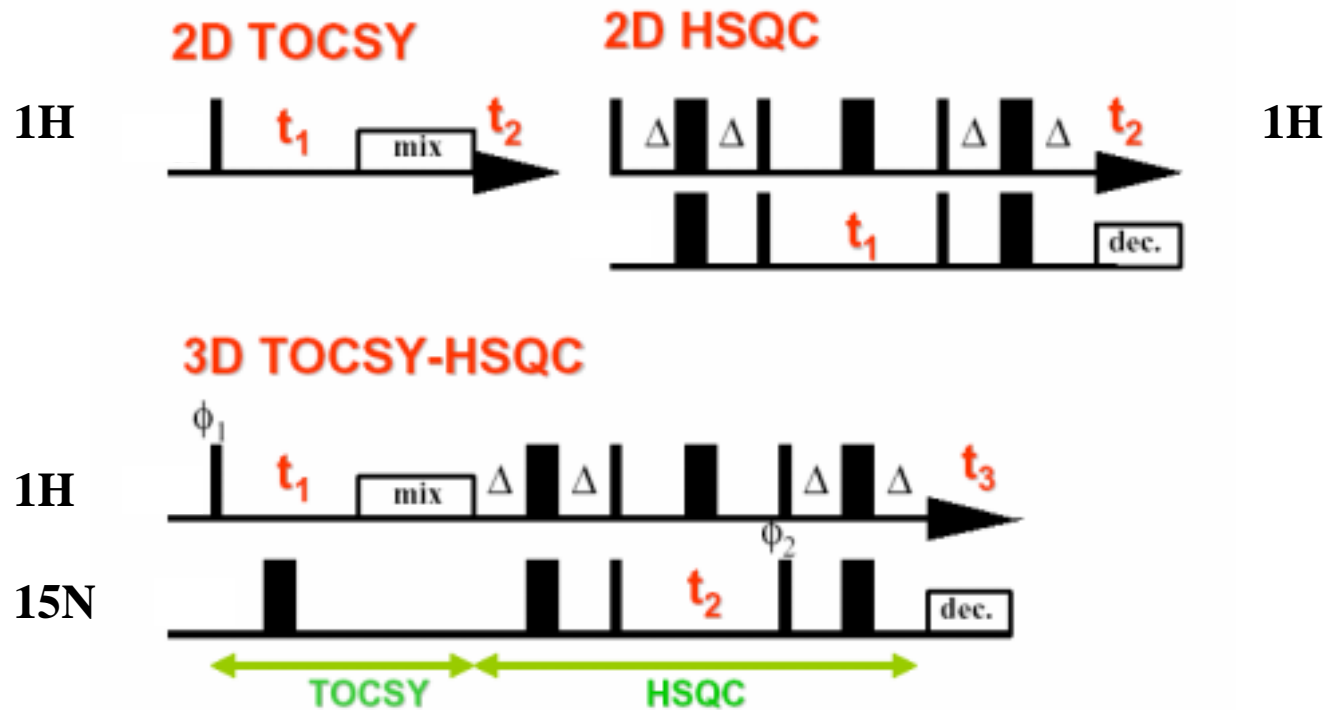
3D



3D Hetero Nuclear

^1H - ^{15}N - ^1H

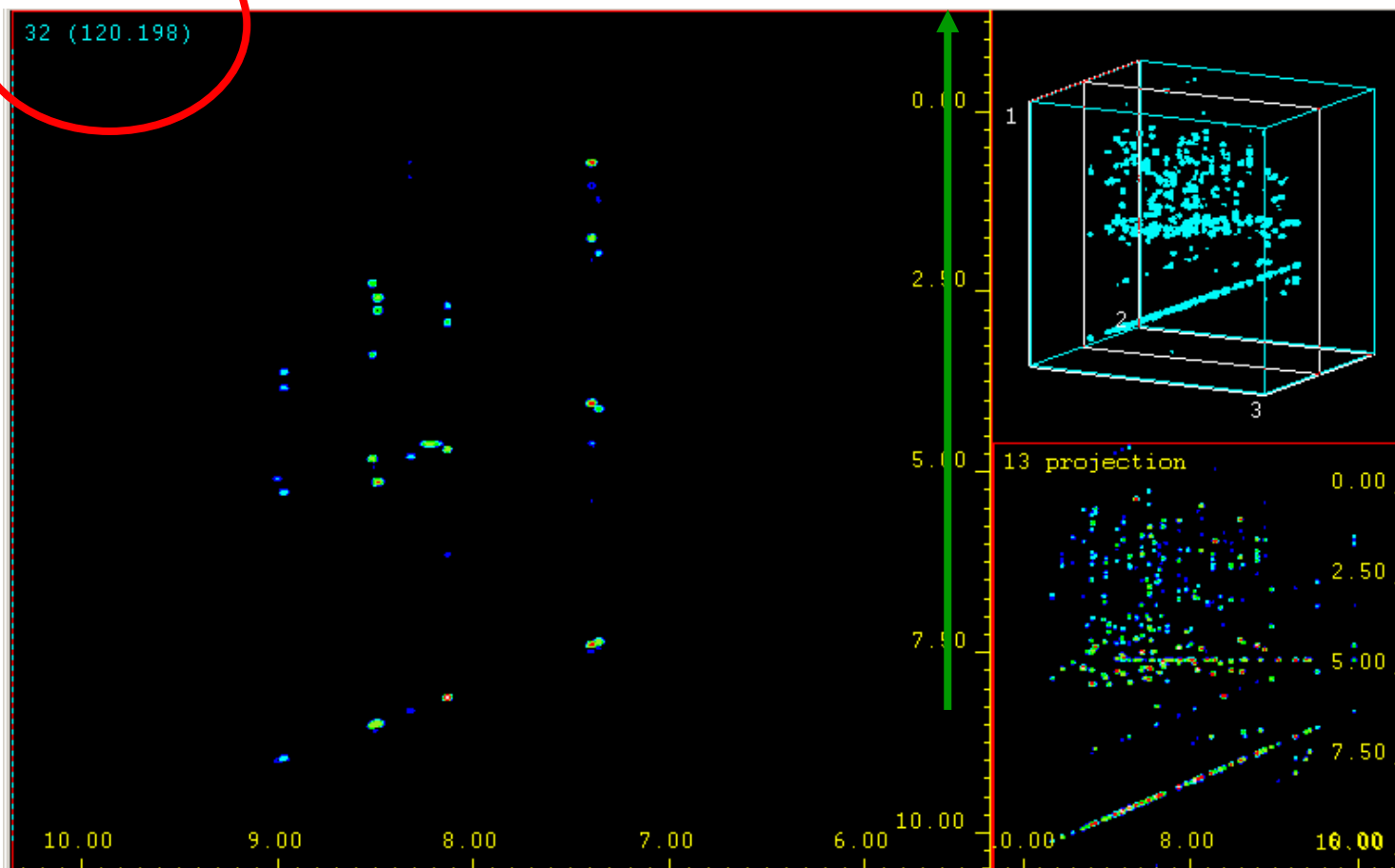
Fig. 1. The 3D pulse sequence.



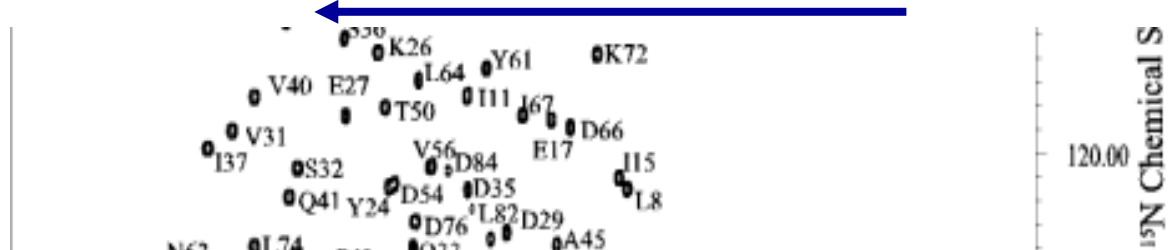
3D 15N-edit TOCSY-HSQC

15N

1H-all



1H-NH

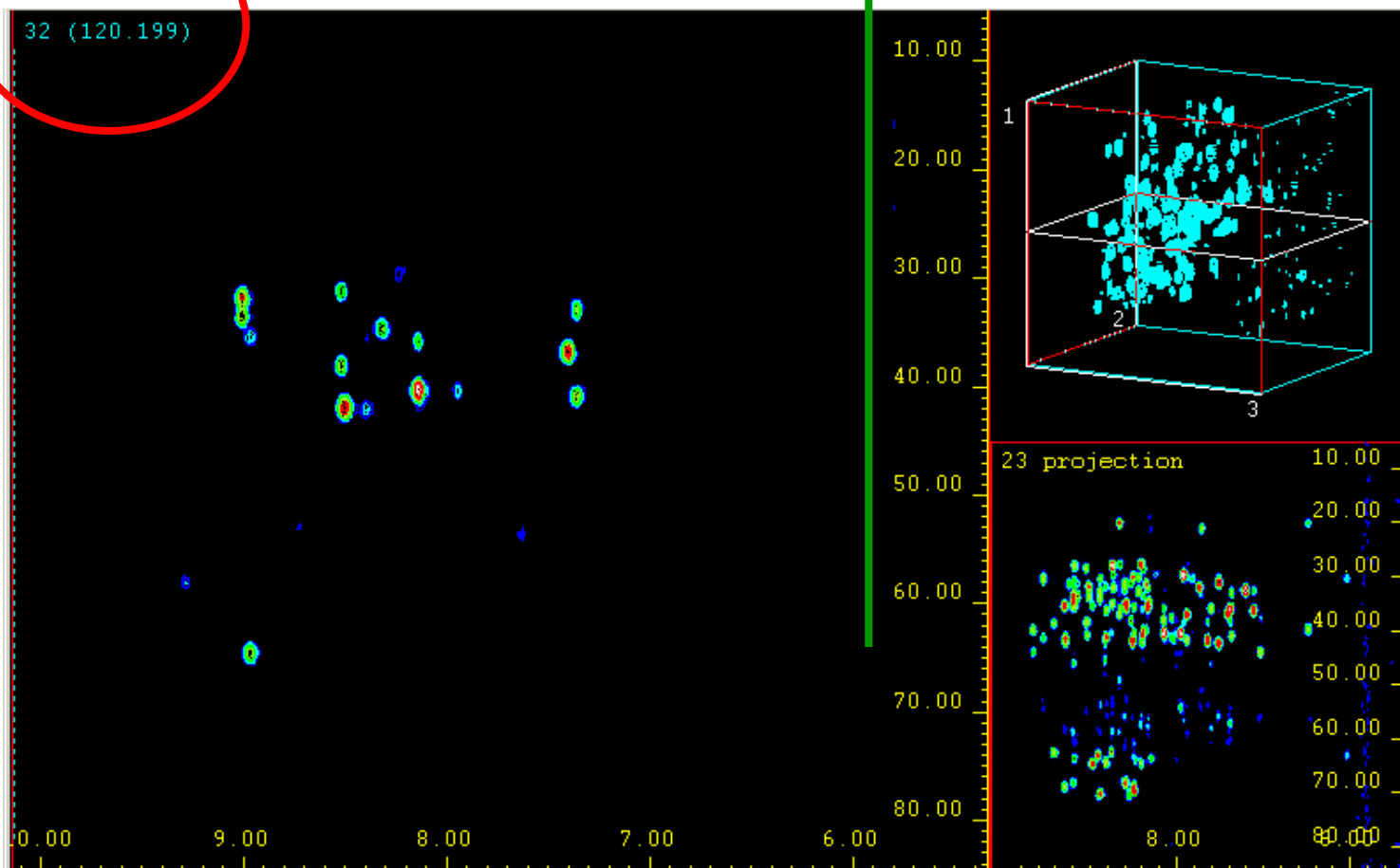


3D CBCA(CO)NH

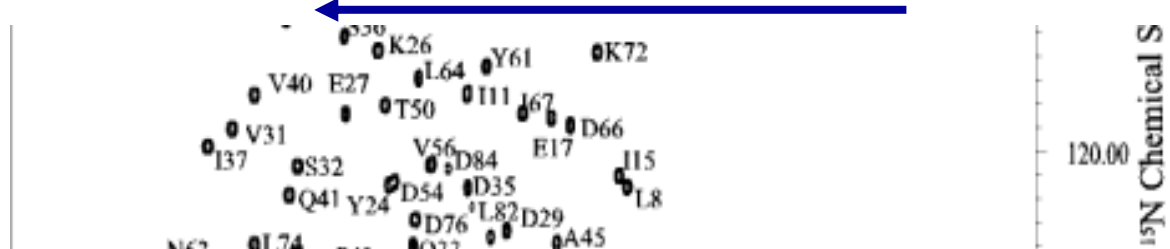
15N

32 (120.199)

13Ca & Cb



1H-NH



Set up 1D/homo Nuclear 2D (Bruker AV system)

Before go into NMR LAB

Overview of NMR Facility

AV500 in IBMS : 5mm TXI-Z ($^1\text{H}/^{15}\text{N}/^{13}\text{C}$, with Z gradient) only

5mm TXI-Z CryoProbe (not available yet)

AV600 in IBMS: 5mm QXI-Z ($^1\text{H}/^{15}\text{N}/^{13}\text{C} /^{31}\text{P}$)

5mm TXI-Z CryoProbe (not available yet)

AV600 in CHEM: 5mm BBO & TXI-Z ($^1\text{H}/^{15}\text{N}/^{13}\text{C}$, with Z gradient)

DRX600 in IBMS: 5mm TXI-XYZ ($^1\text{H}/^{15}\text{N}/^{13}\text{C}$, with XYZ gradient) and others

5mm : ^1H , $^1\text{H}/^{19}\text{F}$, BBO, TXI($^1\text{H}/^{15}\text{N}/^{13}\text{C}$) , TXI-Z ($^1\text{H}/^{13}\text{C}/^{31}\text{P}$)

8mm : TXI ($^1\text{H}/^{13}\text{C}/^{15}\text{N}$) 8mm with Z gradient

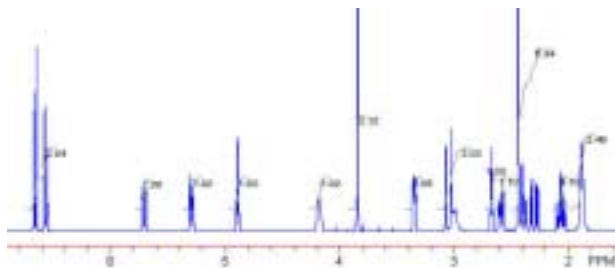
10mm: ^1H , $^1\text{H} /^{19}\text{F}$, BBO

AV800 in IBMS (not available yet) : 5mm TXI-Z & CryoProbe

Definition of some AQ Commands & parameters

edc,new	edit current data set or generate a new data set
eda.ased	edit AQ parameters (eda: shows all, ased: shows required only)
rga	auto optimize rg value
zg	zero memory, and start to collect FID (go)
go	start to collect FID and add signals to the previous memory
stop/halt	stop the active job (currently AQ job)
kill	kill active job (can choose several jobs)
o1.o2,o3	center frequency of the spectrum for nuclear at f1 channel (ex: 1H), f2 channel (ex:13C), and f3 channel (ex:15N)
sw	spectrum width (1 sw : F1 dimension, 2 sw: F2 dimension.....)
td	number of points for FID collection(1 td: F1 dimension, 2 td: F2 dimension....)
d1	relaxation time (usually $> 5 \cdot T_1$)
ns	number of scan
ds	dummy scan
rg	receiver gain (usually use the value calculated by rga)

1D

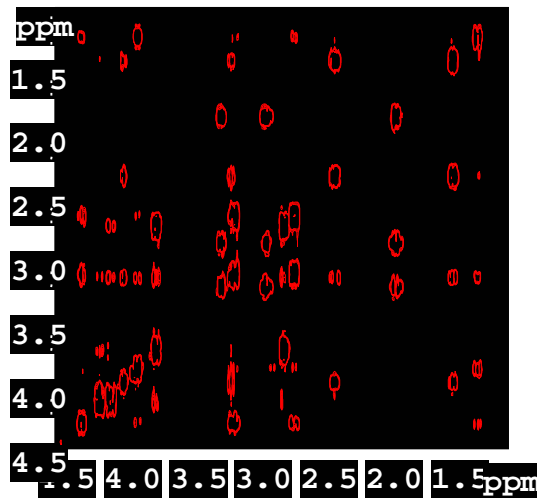


center (**o1**) at 4.5ppm

sw=6ppm

1.5ppm (=o1-sw/2) to 7.5ppm(=o1+sw/2)

2D/homo



F1 dimension !!!

F2 dimension !!!

F2: H (f1 channel)

O1(f1 channel):3ppm

2 sw(F2 dimension): 5ppm

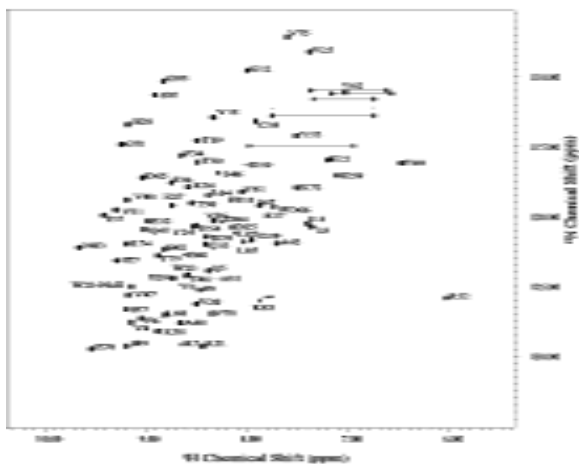
F1: H (f1 channel)

O1(f1 channel): 3ppm

1 sw(F1 diemnsion): 5ppm

Acquisition Parameters		F2	F1
PULPROG	zgpg30		
HLMod	zgpg		
TD	2048	128	
CHST	zgpg		
NUC1	edit	edit	
NUC2	edit	edit	
SAVE		Parameter	Plot

2D/hetero



F2: H (f1 channel)

O1(f1 channel):8ppm

2 sw(F2 dimension): 5ppm

F1: N (f3 channel)

O3(f3 channel): 120ppm

1 sw(F1 diemnsion): 40ppm

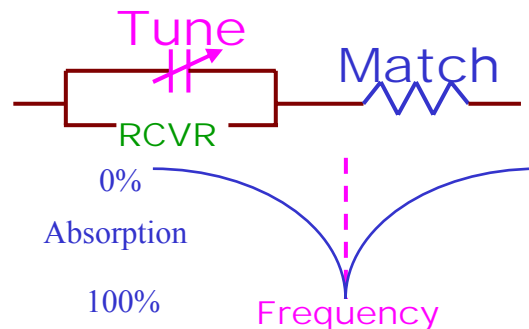
Loading Sample

1. The best condition for sample? → . Probes, Temperature, Sample position
Selecting the right probe : inner coil is observe coil and outer coil is decoupling
Example: better ^{13}C sensitivity should use BBO (^{13}C : inner, ^1H : outer), but if for better ^1H , "inversed probe" (^1H : inner, others: outer) is recommended

2. The best condition for NMR? → Wobble : Tune & Match

Tuning is the process of adjusting freq. until it coincides with the desired frequency

Matching is the process of adjusting the impedance of the resonant circuit



3. The best condition for field? → Lock and shim

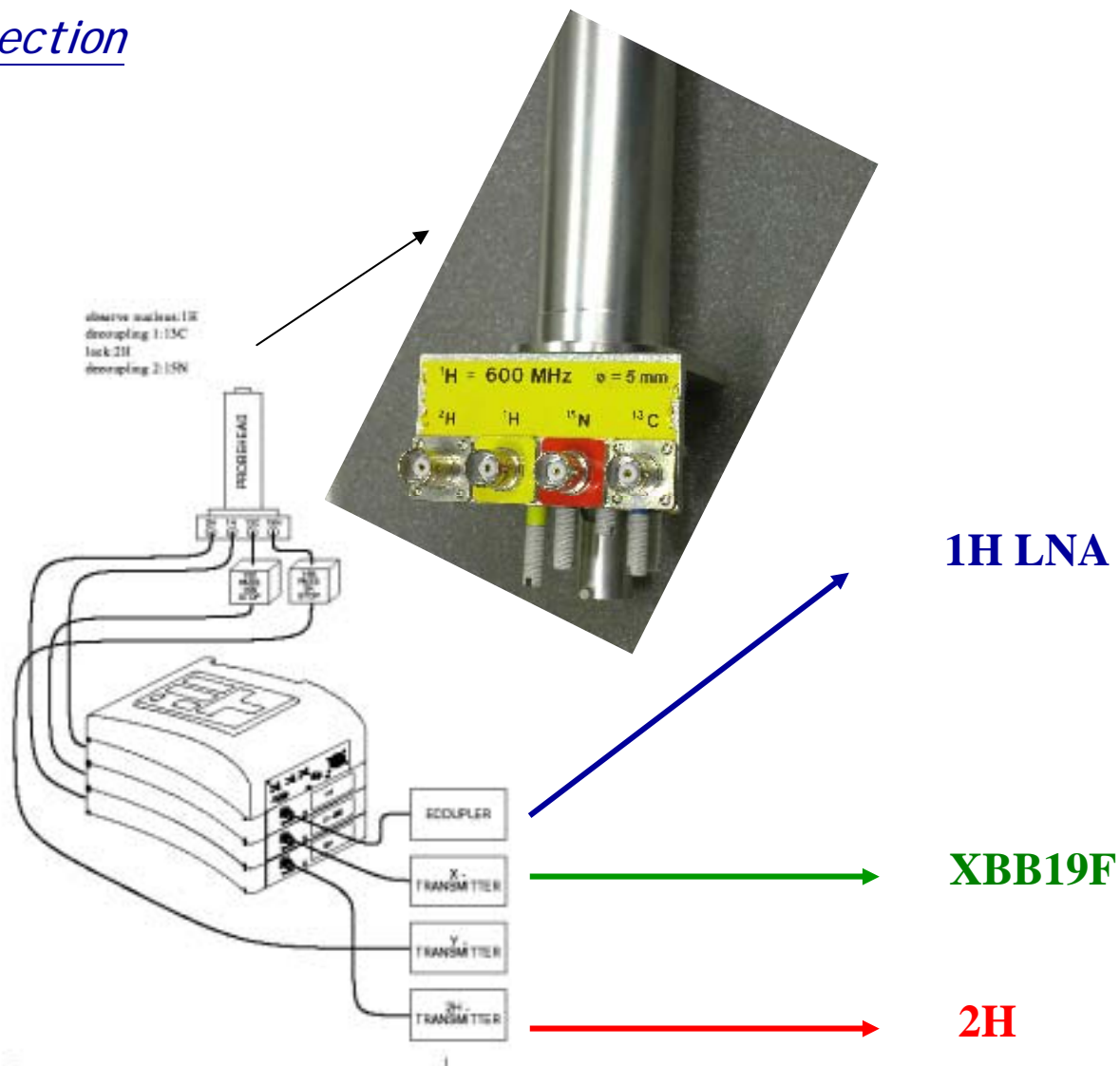
Deuterium lock means the long term stability of the magnetic field is achieved

The shims (coils) are small magnetic fields used to adjust the homogeneity of the field

4. Ready to go! (LAB work sheet)

In the NMR LAB

Hardware Connection

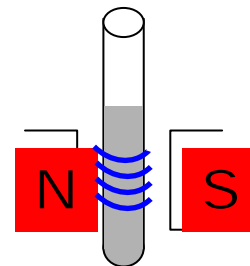


Loading Sample

1. Set up temperature : **edte** (edit temperature)



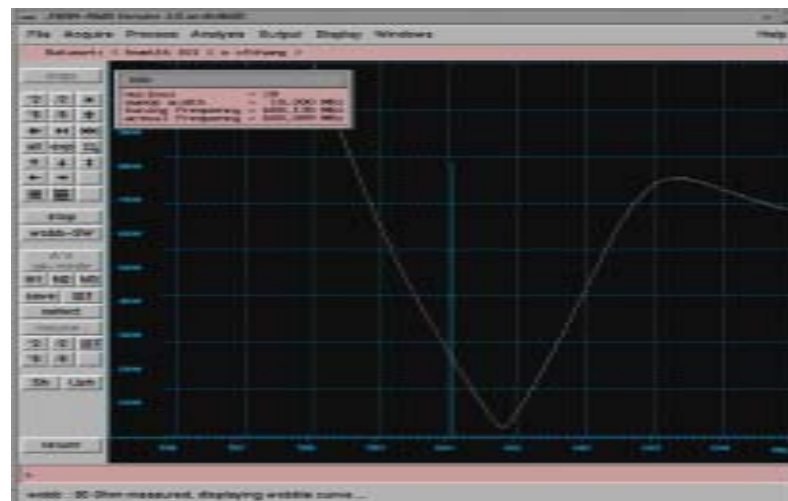
2. Adjust sample position



Hardware Adjustment

wobble

lock



gradshim : gradient shimming



Set up Experiment

Easy steps for beginner (1D):

1. Type "rpar" <enter> to load an appropriate parameter set
2. Type "ns" <enter> to input number of scan
3. Type "rga" <enter> to find appropriate receiver gain
4. Type "zg" <enter> to collect spectrum
5. Type "ft" <enter> to do Fourier Transfer
6. Click on phase to [phase spectrum](#)
7. Click on return, then save to [save the spectrum](#)
8. [Print out the spectrum](#)
9. Save your data on floppy

Define middle of the spectrum: O1

1. "rpar" <std_1D_1H_ZG>
2. "zg" → "ft" → <phase>
3. Click on <utility> to define O1 position at middle of the spectrum

Steps for optimize 90 deg pulse:

1. "rpar" <std_1D_1H_ZGPR> , and set O1 to H2O position (if not in H2O, just use std_1D_1H_ZG to determine 90 deg pulse)
2. Set pL1= 0db (or 10db or ? db), p1=5u
3. "rga" → "zg" → "ft" → <phase> → <save>
4. Keep pL1, increase p1= expected 180 deg
(or, keep p1=180 deg pulse length, decrease pL1=expected power level for p1 value)
5. "zg" → "fp" → check if the spectrum almost become null, if not, repeat step 4
6. Now, you should have a table with 1H pulse information of your sample

- How to optimize condition? → For users: Follow Experiment Guide

- Experiment Name: 2D 1H TOCSY MLEV17
- Experiment Type: H2O supression using 3-9-19 & gradient
- Standard Parameter Set: std_2D_TOCSY-MLEV
- Pulse Program: mlevgppl19
- AQ parameters to check

1H pulses

p1 (high power, ex: 0dB), **p1** (90o pulse at p1)
p110 (low power), **p6** (90o pulse at p110 for mixing pulse, ex: 25u)
p118 (low power for 3-9-19, ex: 10dB), **p0,p27** (90o pulse at p118)
p28 (trim pulse, ex:1m)

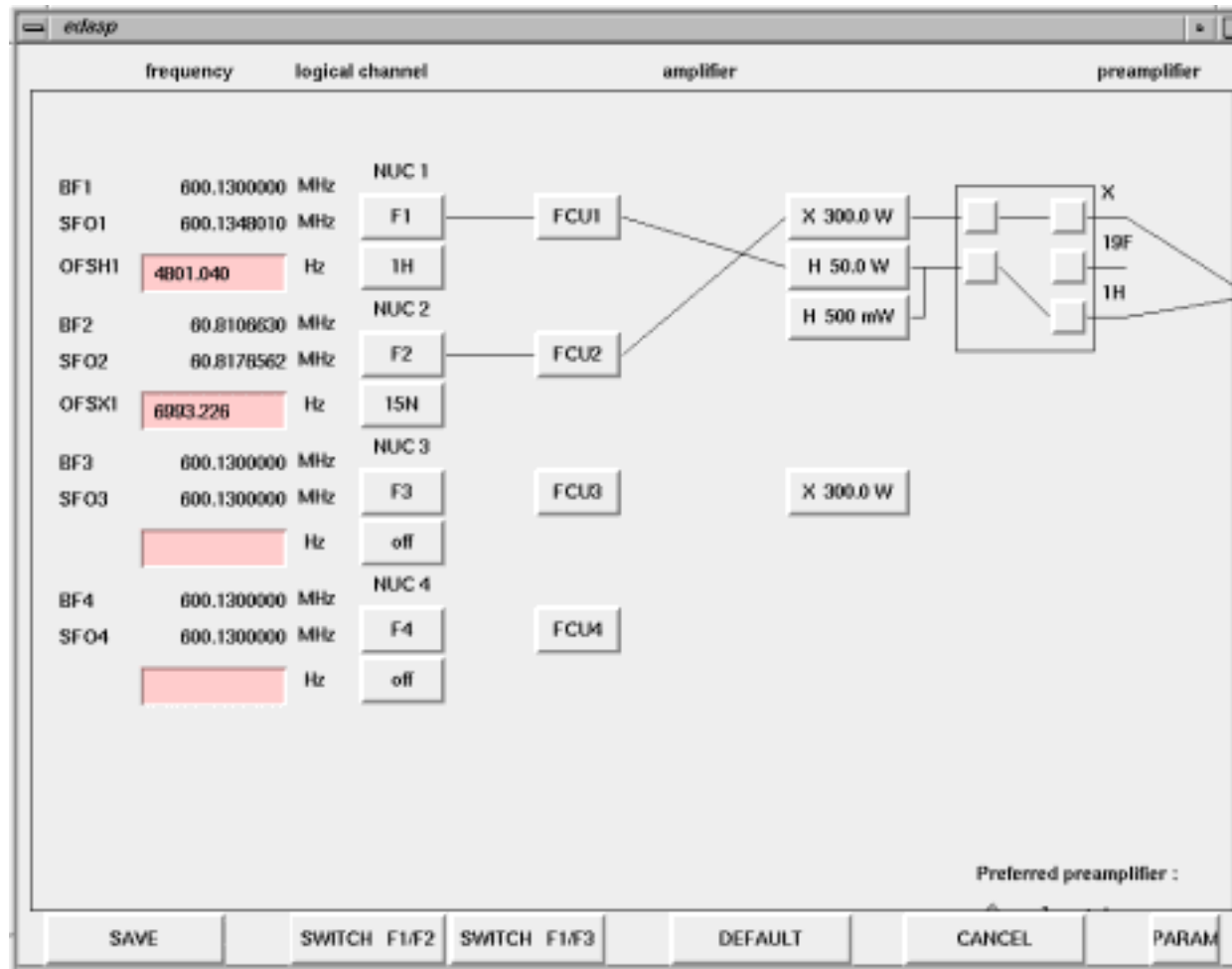
Others

d9 (mixing time,ex: 60-70ms)
d19 (=1/2d, d=distance of next null in Hz)
o1 (on H2O)
1 sw, 1td (for F1 dimension, H)
2 sw, 2 td (for F2 dimension, H)
d1, ns(=2*n), **ds**(>=16)
rg

Set up hetero nuclear 2D/3D (Bruker AV system)

Wobble all 3 channels: 15N → 13C → 1H

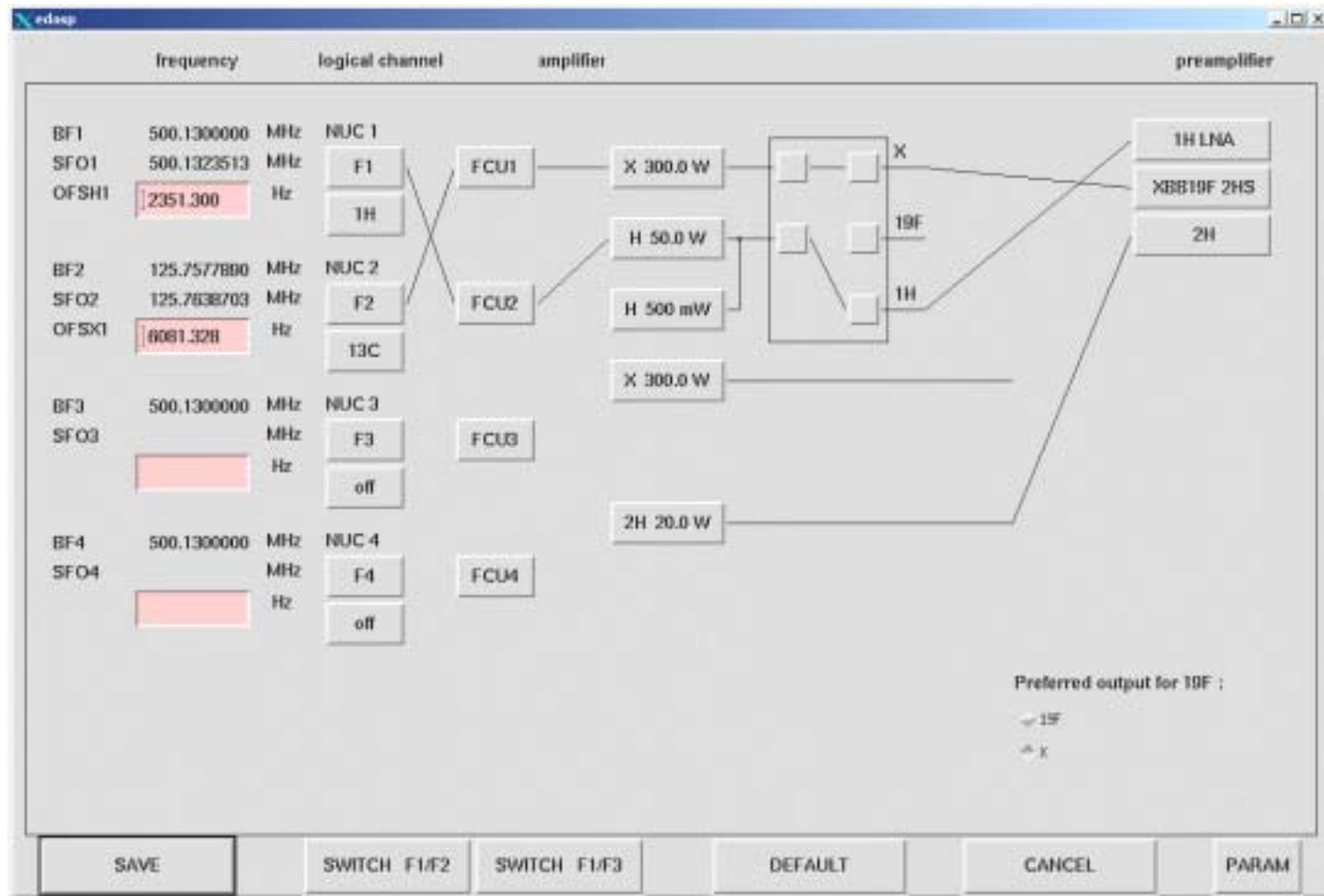
Step 1.1: (15N first) edasp to change setting and connection



Step 1.2: type wobble to wobble 15N, type stop after tune and match

Wobble all 3 channels: 15N → 13C → 1H

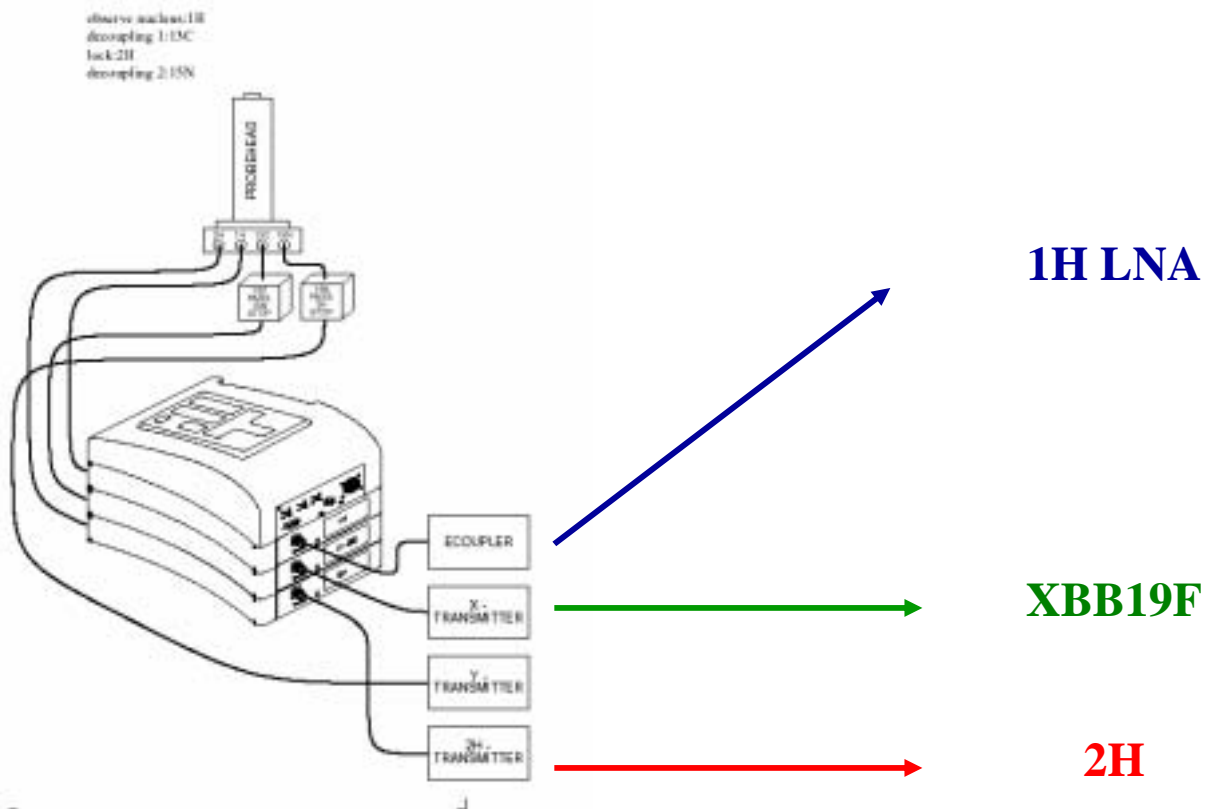
Step 2.1: (13C) edasp again to change setting and connection



Step 2.2: type wobble to wobble 13C,click any key on "HPPR" after 13C tuning and matching are done.

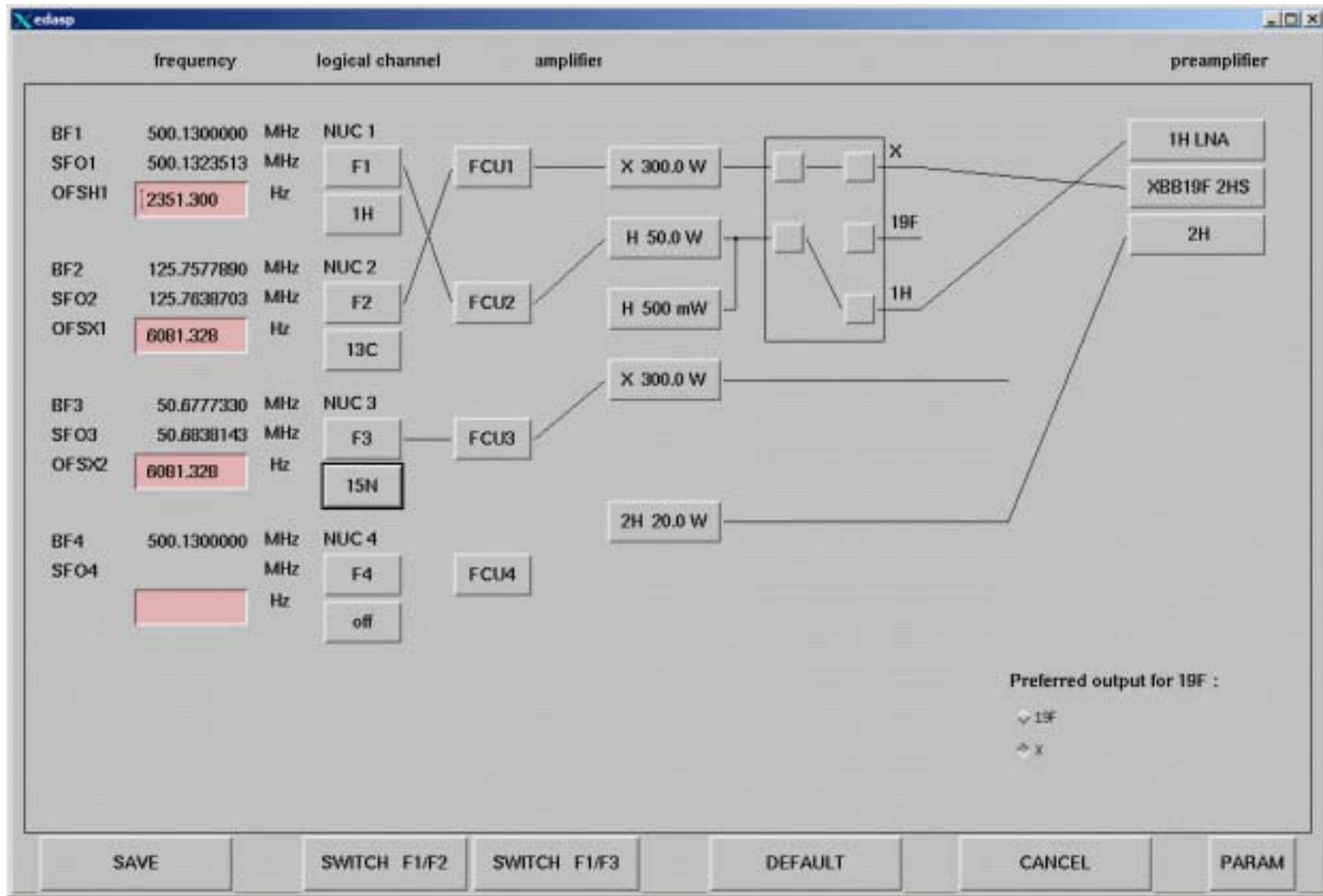
Wobble all 3 channels: 15N → 13C → 1H

Step 3.1: click "Chn ↑" on "HPPR" , then wait until wobble on 1H pop out (this might take 10-20 sec, please be patient!)



Step 3.2: type stop after tuning and matching are done for 1H.

Step 3.3: edasp to change setting and connection for the experiment



- How to optimize condition? → For users: Follow Experiment Guide

- Experiment Name: 2D 15N-1H HSQC
- Experiment Type: Using echo-antiecho, f1: H, f3:N
- Standard Parameter Set: std_2D_15N_HSQC_ET
- Pulse Program: invietf3gpsi
- AQ parameters to check

1H pulses

p1 (high power, ex: 0dB), **p1** (90o pulse at p1)
p28 (trim pulse, ex:1m)

Others

cnst4 (J H-N , ex: 90Hz)
d24 (1/4JH-N)
o1 (for 1H)
o3 (for 15N)
1 sw, **1td** (for F1 dimension, N)
2 sw, **2 td** (for F2 dimension, H)
d1
ns(=1*n)
ds(>=16)
rg

Users need to adjust parameters in “red” (meaning of the parameter in “green”)

- How to optimize condition? → For users: Follow Experiment Guide

•Experiment Name: **3D HNCO**

•Experiment Type: **Using Echo/antiecho , f1: H, f2:C, f3:N, F1(CO), F2(N), F3(H)**

•Standard Parameter Set: **std_3D_HNCO**

•Pulse Program: **hncogp3d.2**

•AQ parameters to check

1H pulses

pl1 (high power, ex: 0db), **p1**(90deg at pl1), **p2**(180deg at pl1)

pl19 (low power for dipsi2,pcpd1), **p26**(90deg at pl19), **pcpd1**(90deg ,ex: 40-50usec)

sp1 (shape pulse power for Sinc.1000) , **p11**(pulse length for sp1, ex: 2m)

Others

o1 (for 1H), **o2** (for 13CO), **o3** (for 15N)

1 sw, 1 td (for F1 dimension, ie: 13C)

2 sw, 2 td (for F2 dimension, ie:15N)

3 sw, 3 td (for F3 dimension, ie:1H)

d1, rg, ns(=8*n), **ds** (16)

Users need to adjust parameters in “red” (meaning of the parameter in “green”)

- How to optimize condition? → For operators : pulse program

(1) hard pulse calibration for hetero nuclei

(2) Shape pulse calibration for hetero nuclei

(3) other “uniform” (sample independent) parameters set up

ex: delays, decouple program, gradient program, frequency jump.....

```

;hncogp3d.2
;avance-version (01/05/09)
;HNCO
.....( skips)
;   F1(H) -> F3(N) -> F2(C=O,t1) -> F3(N,t2) -> F1(H,t3)
;
;on/off resonance Ca and C=O pulses using shaped pulse
;phase sensitive (t1)
;phase sensitive using Echo/Antiecho gradient selection (t2)
.....( skips)
;sp1: f1 channel - shaped pulse 90 degree (H2O on resonance)
;sp2: f2 channel - shaped pulse 90 degree (C=O on resonance)
;sp3: f2 channel - shaped pulse 180 degree (C=O on resonance)
;sp5: f2 channel - shaped pulse 180 degree (Ca off resonance)
;sp8: f2 channel - shaped pulse 90 degree (C=O on resonance)
;           for time reversed pulse
.....( skips)
;d21: 1/(2J(NH)) [5.5 msec]
;d23: 1/(4J(NCO)) [12 msec]
;d26: 1/(4J'(NH)) [2.3 msec]
.....( skips)
;cpds1: decoupling according to sequence defined by cpdprg1
;cpd3: decoupling according to sequence defined by cpdprg3
;pcpd1: f1 channel - 90 degree pulse for decoupling sequence
;pcpd3: f3 channel - 90 degree pulse for decoupling sequence
;use gradient ratio: gp 1 : gp 2 : gp 3 : gp 4 : gp 5
;                      60 : -40 : 10 : 80 : 8.1

```

Thank you!!

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