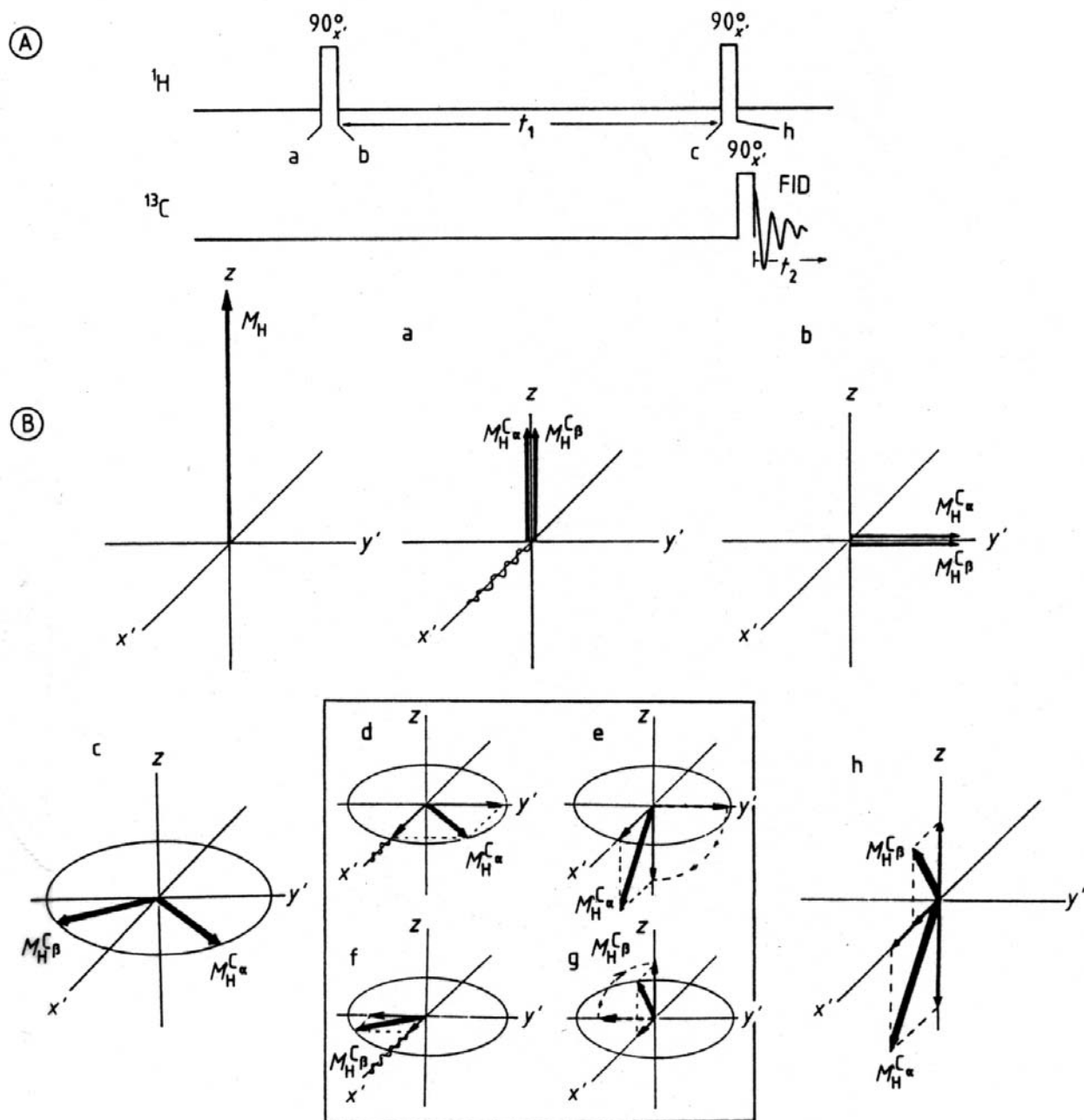


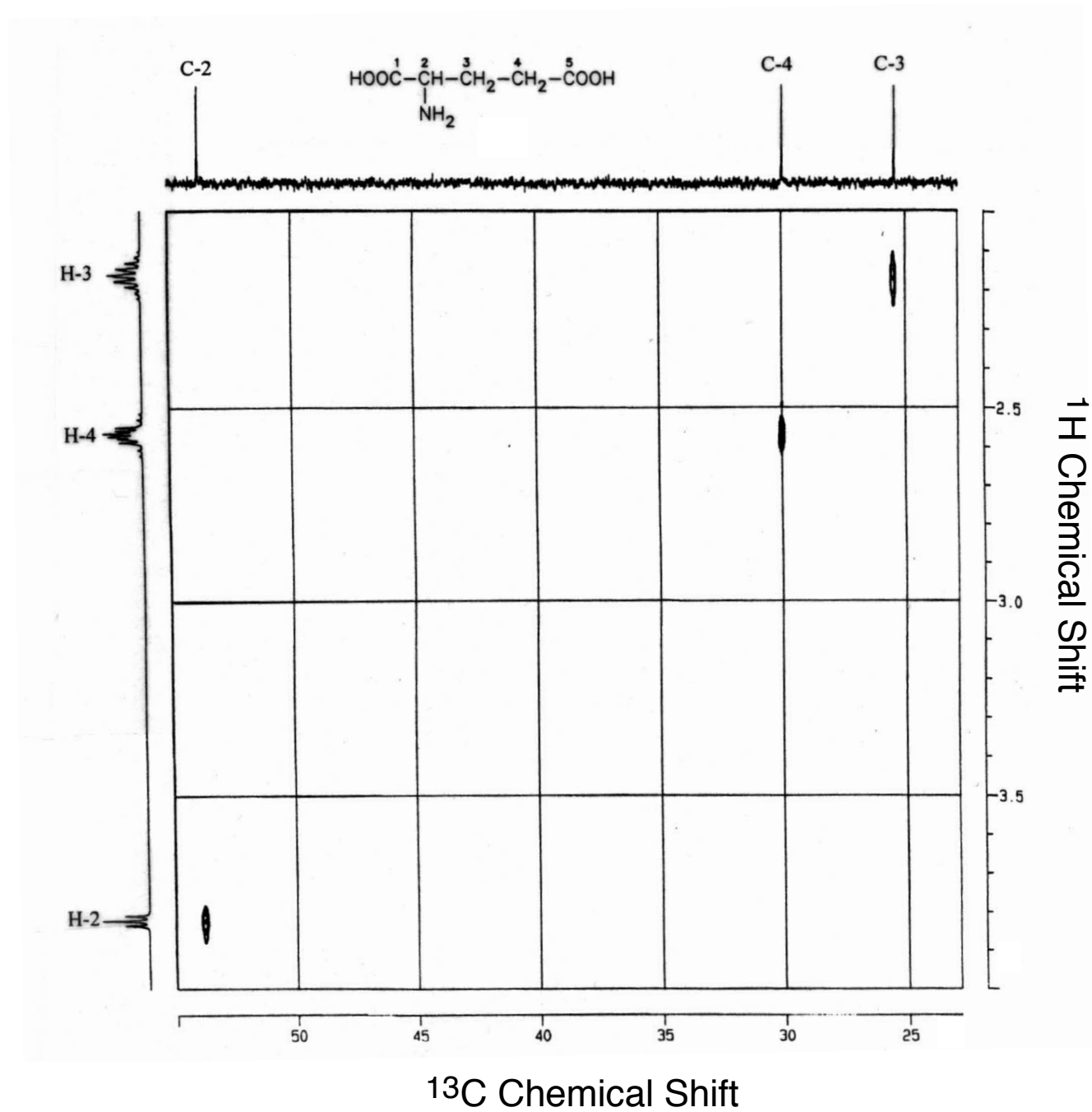
CHEM / BCMB 4190/6190/8189

Introductory NMR

Lecture 17

Last Time:
-HETCOR or C, H-COSY



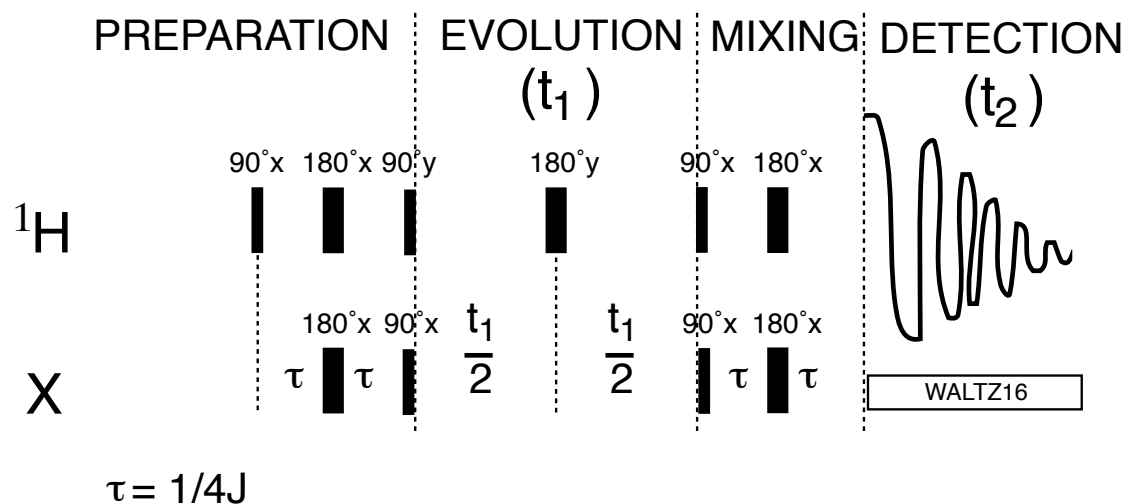


Two-dimensional Correlated NMR spectroscopy: The HSQC Experiment

1. HETCOR (or C,H-COSY) Versus HSQC:

- We have seen the C,H-COSY experiment where ^1H is detected in t_1 and ^{13}C is detected in t_2 . Although there is population transfer from ^1H to ^{13}C , the sensitivity of this experiment is poor because ^{13}C (not ^1H) is detected in t_2 .
- Higher sensitivity can be achieved by doing the "reverse" experiment i.e. by detecting ^1H in t_2 and ^{13}C in t_1 . The HSQC and HMQC are two experiments that achieve this heteronuclear H,C-correlation. Here, we will consider in more details the HSQC experiments.

2. The HSQC Pulse Sequence:

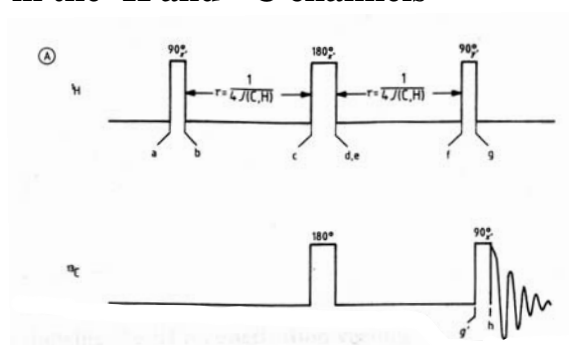


- In the ^1H - ^{13}C HSQC ($X = ^{13}\text{C}$):

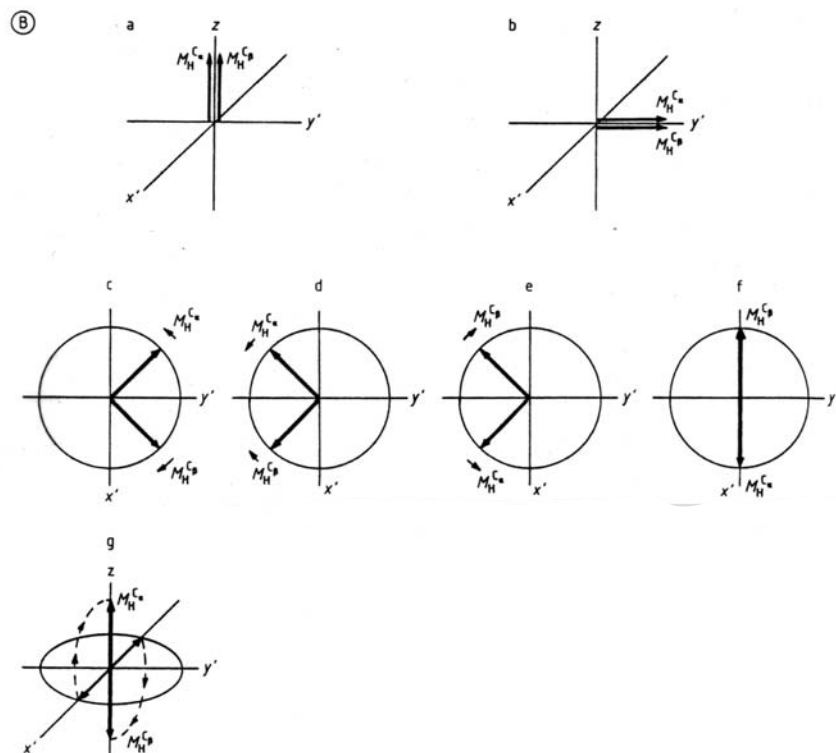
- 1) The PREPARATION period is an INEPT sequence (^1H to ^{13}C).
- 2) The t_1 EVOLUTION period allows for indirect ^{13}C chemical shift detection.
- 3) The MIXING period is a REVERSE INEPT sequence (^{13}C to ^1H).
- 4) The t_2 EVOLUTION period allows for direct ^1H chemical shift detection.

3. Review of the INEPT experiment:

A) Pulse sequence in the ^1H and ^{13}C channels



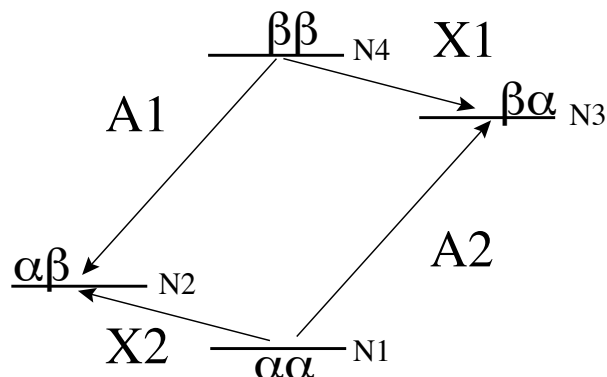
B) Vector diagrams showing the ^1H magnetization vectors ($^{13}\text{CHCl}_3$)



- At point g: ^1H 90° pulse rotates $M_H^{C\alpha}$ to $+z$ and $M_H^{C\beta}$ to $-z$

The populations of N2 and N4 are inverted:

before INEPT	after INEPT
$N4 = N$	$= N + \Delta H$
$N3 = N + \Delta C$	$= N + \Delta C$
$N2 = N + \Delta H$	$= N$
$N1 = N + \Delta C + \Delta H$	$= N + \Delta C + \Delta H$



X1 transition: $N3 - N4 = \Delta C - \Delta H = -3\Delta C$

X2 transition: $N1 - N2 = \Delta C + \Delta H = 5\Delta C$

- In the case of the HSQC, the contribution from the natural ^{13}C magnetization (ΔC) is unwanted and is removed using one of the selected methods that we have seen previously. The resulting populations and population differences are:

$$N4 = N + \Delta C/2 + \Delta H$$

$$N3 = N + \Delta C/2$$

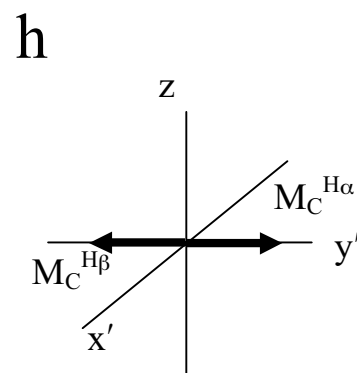
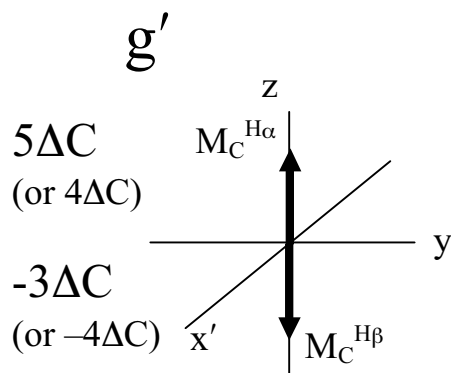
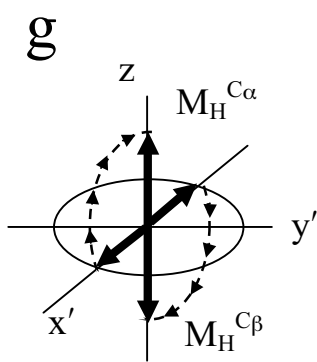
$$N2 = N + \Delta C/2$$

$$N1 = N + \Delta C/2 + \Delta H$$

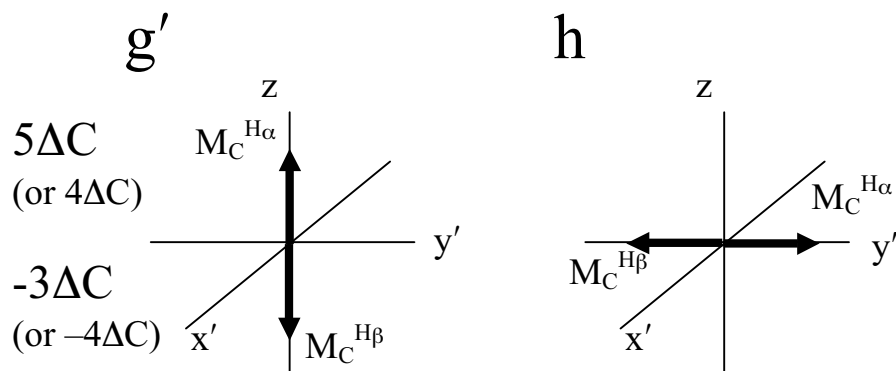
$$\text{X1 transition: } N3 - N4 = -\Delta H = -4\Delta C$$

$$\text{X2 transition: } N1 - N2 = \Delta H = 4\Delta C$$

- At point g': $M_C^{H\alpha}$ is in its original position, but $M_C^{H\beta}$ is inverted

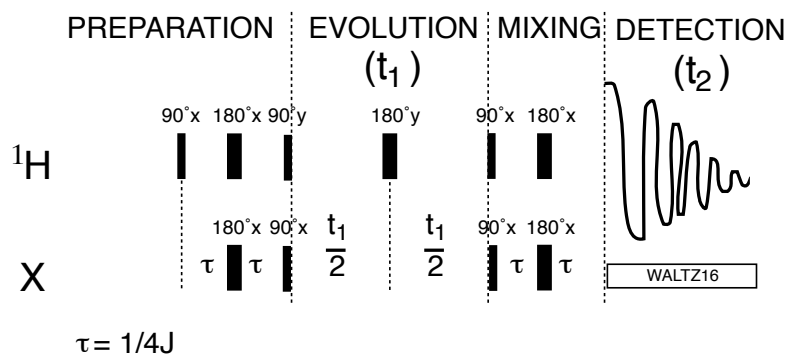


C) Vector diagrams showing the ^{13}C magnetization vectors

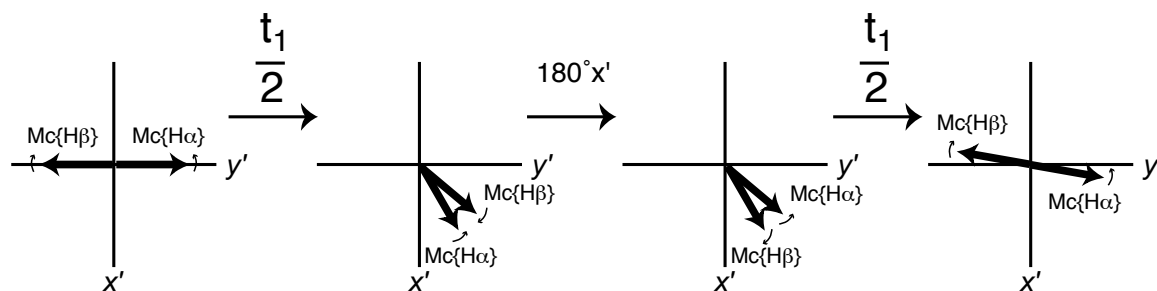


- At point h: The 90° pulse on ^{13}C creates transverse magnetization components that evolve during t_1 . Note that these ^{13}C magnetization components are 180° out of phase with each others at the beginning of t_1 .

4. The t_1 Evolution Period:

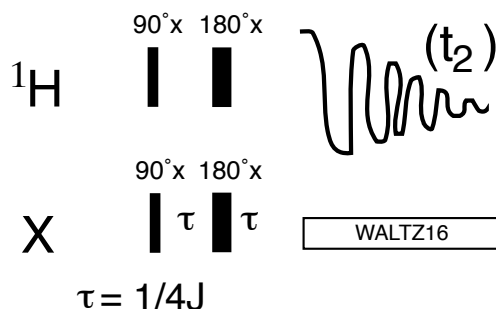


- The ^{13}C chemical shift evolves (to different points depending on the value of t_1).
- There is no net ^1H - ^{13}C coupling evolution. Note that the ^{13}C magnetization components are 180° out of phase with each others at the beginning and the end of t_1 .

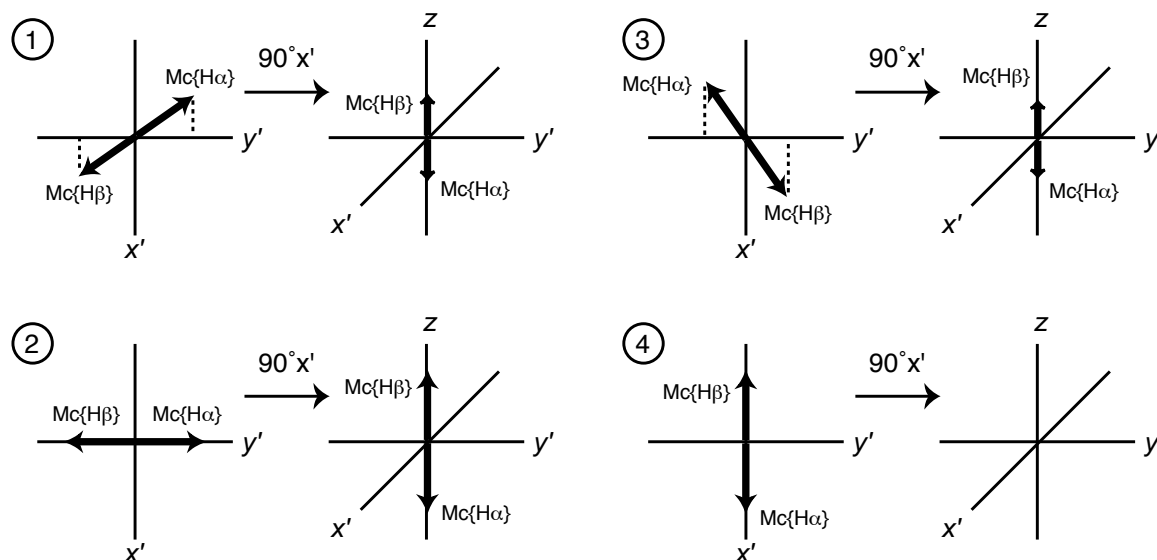


5. The Reverse-INEPT experiment:

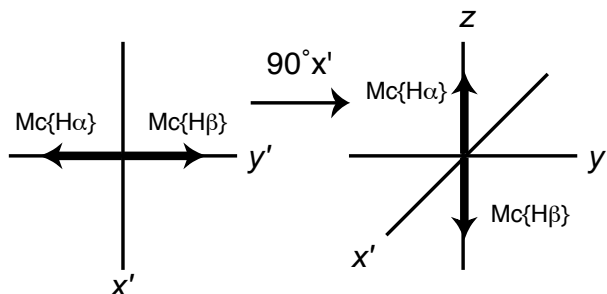
A) Pulse sequence in the ^1H and ^{13}C channels



B) Vector diagrams showing the effect of the first ^{13}C $90^\circ x'$ pulse on the ^{13}C magnetization vectors (Ex.: $^{13}\text{CHCl}_3$) for various t_1 values. Four cases are shown below:



We will only consider the following case, where after the first ^{13}C $90^\circ x'$ pulse, $\text{MC}^{\text{H}\beta}$ is in its original position, but $\text{MC}^{\text{H}\alpha}$ is inverted.



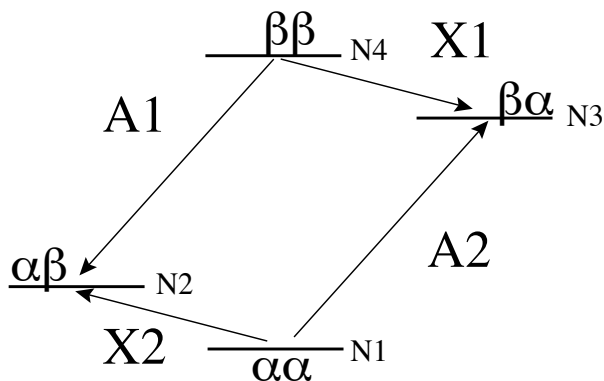
The populations are described as followed (same as after first ^{13}C 90° x):

$$N4 = N + \Delta C/2 + \Delta H$$

$$N3 = N + \Delta C/2$$

$$N2 = N + \Delta C/2$$

$$N1 = N + \Delta C/2 + \Delta H$$



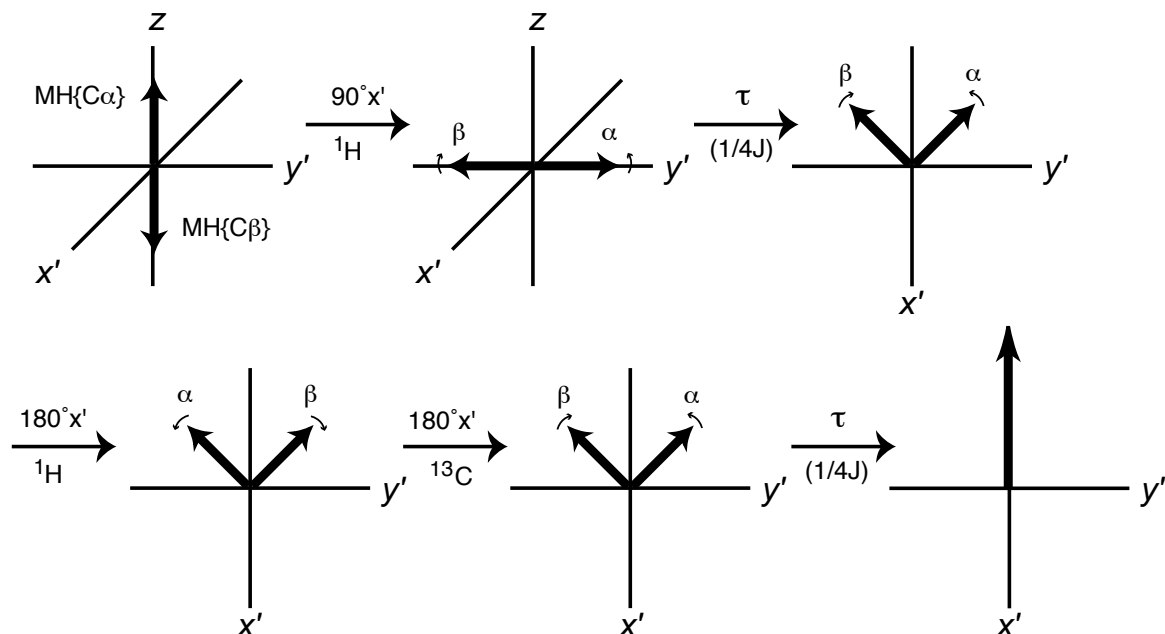
$$\text{A1 transition: } N2 - N4 = -\Delta H$$

$$\text{A2 transition: } N1 - N3 = \Delta H$$

$$\text{X1 transition: } N3 - N4 = -\Delta H = -4\Delta C$$

$$\text{X2 transition: } N1 - N2 = \Delta H = 4\Delta C$$

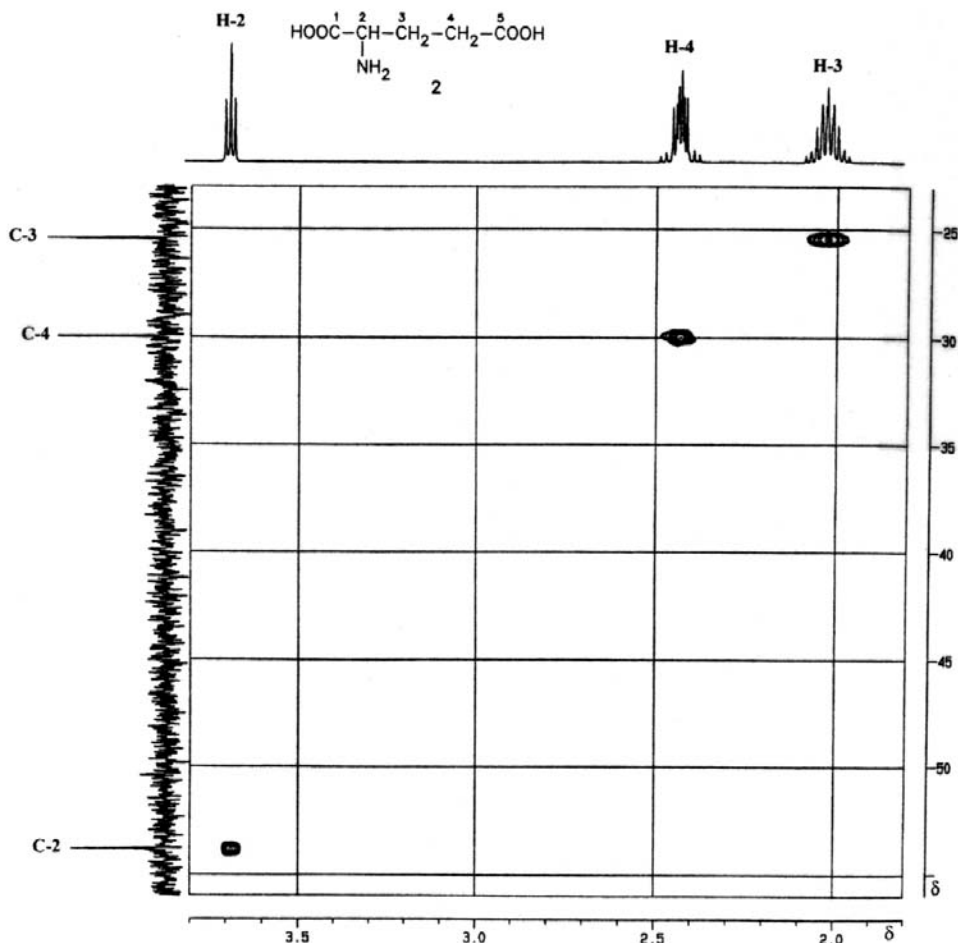
- The antiphase ^1H magnetization is refocused during the 2τ period. The two 180° pulse (on ^1H and ^{13}C) in the middle of the 2τ period allows J coupling evolution but refocuses the ^1H chemical shift evolution.
- For simplicity, one can ignore the effect of chemical shift, which is refocused during the reverse-INEPT period.



- At the end of the reverse-INEPT α and β are in phase, therefore we can turn on BB ^{13}C decoupling during acquisition.

6) HSQC Spectrum:

Example: ^1H - ^{13}C HSQC of Glutamic Acid

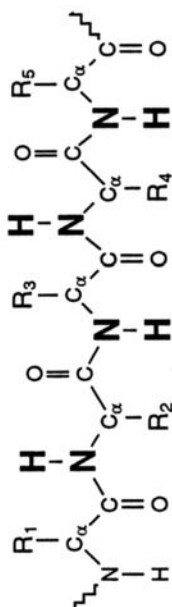


- Signals not obtained for ^1H that are not bound to ^{13}C
- Comparison with C,H COSY:
 - 1) Axes are interchanged (in HSQC, ^1H is detected, in HETCOR, ^{13}C is detected)
 - 2) HSQC is more sensitive and a good spectrum can be recorded quickly.

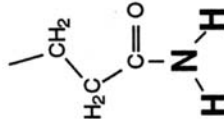
-Example: for 1 mM uniformly ^{13}C - (or ^{15}N -) isotopically labeled samples, typical recording times are:

- 15-30 minutes for HSQC
- 24 hours for C, H COSY

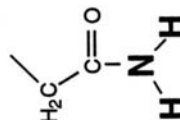
Proton (H) - Nitrogen (N) Correlations



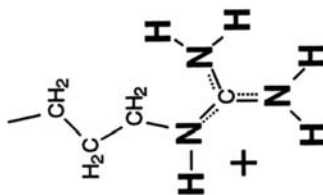
$\text{R}_1 = \text{Glutamine (Q)}$



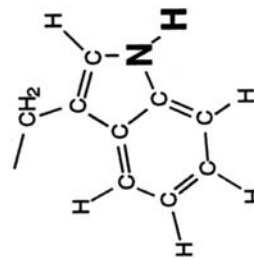
$\text{R}_2 = \text{Asparagine (N)}$



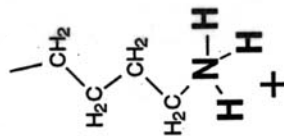
$\text{R}_3 = \text{Arginine (R)}$



$\text{R}_4 = \text{Tryptophan (W)}$



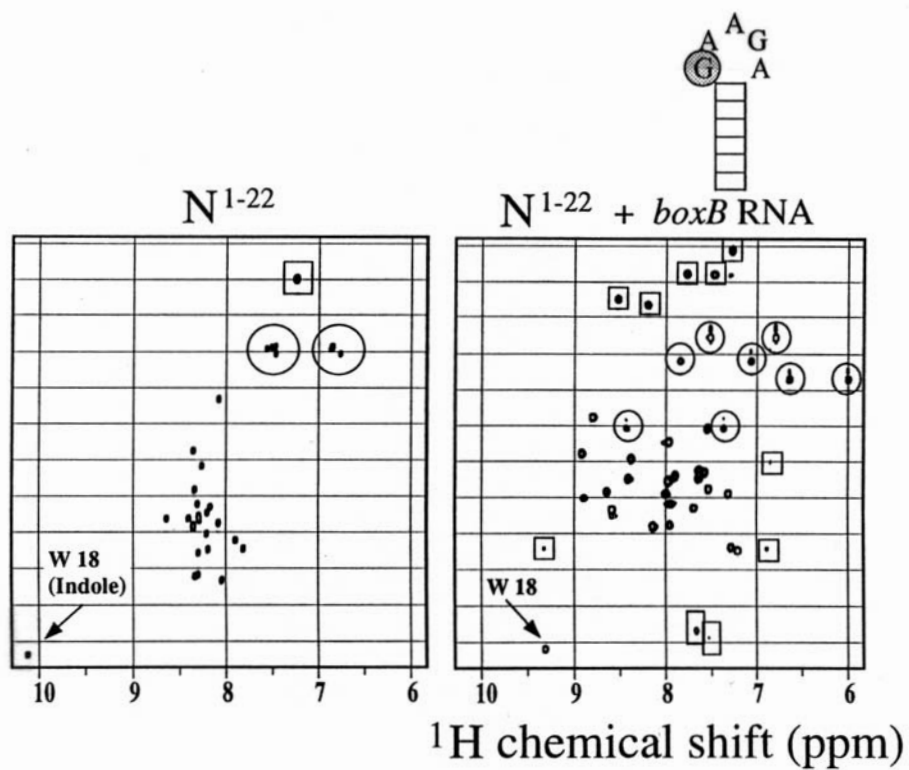
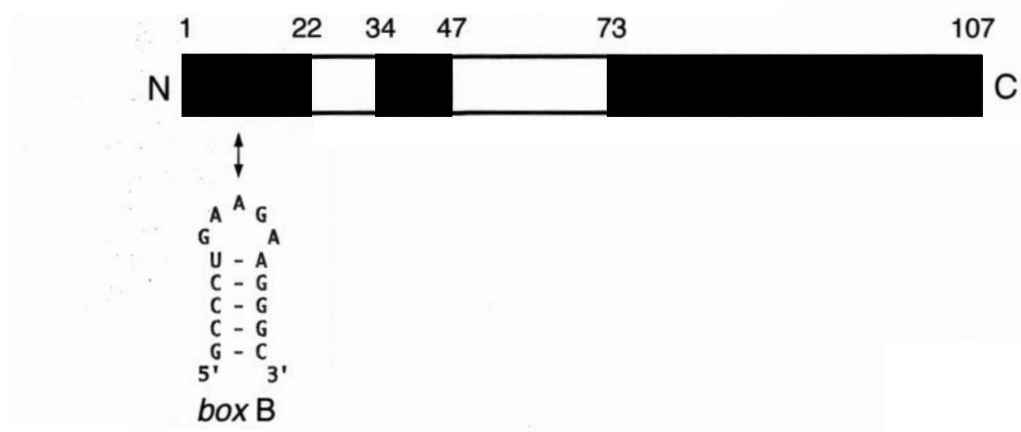
$\text{R}_5 = \text{Lysine (K)}$



7) Important “Biologically Biased” Applications: ^1H - ^{15}N HSQC for Analysis of Ligand Binding in Protein

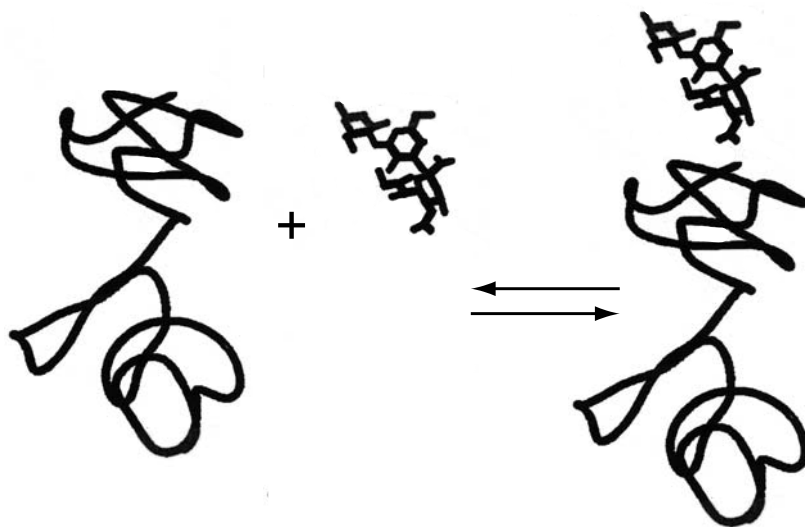
CASE 1: Protein Folding Upon Ligand Binding

Domains of the λ N Protein

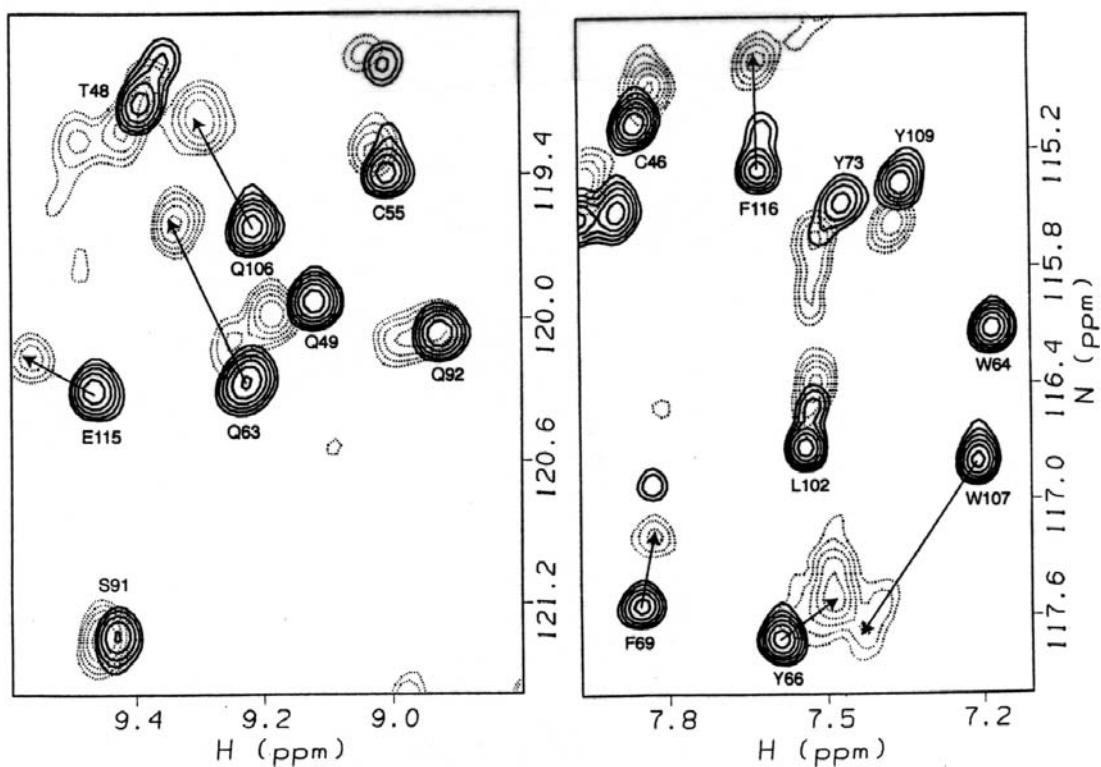


- ASN and GLN side chains
- ARG side chains (folded nitrogens)
- LYS side chains (folded nitrogens)

CASE 2: Mapping Ligand Binding Sites in Folded Proteins



- Chemical shifts of ligand binding site are perturbed:



The TOCSY Experiment

TOCSY: Total Correlation Spectroscopy

Also known as HOHAHA (HOmonuclear Hartmann-Hahn)

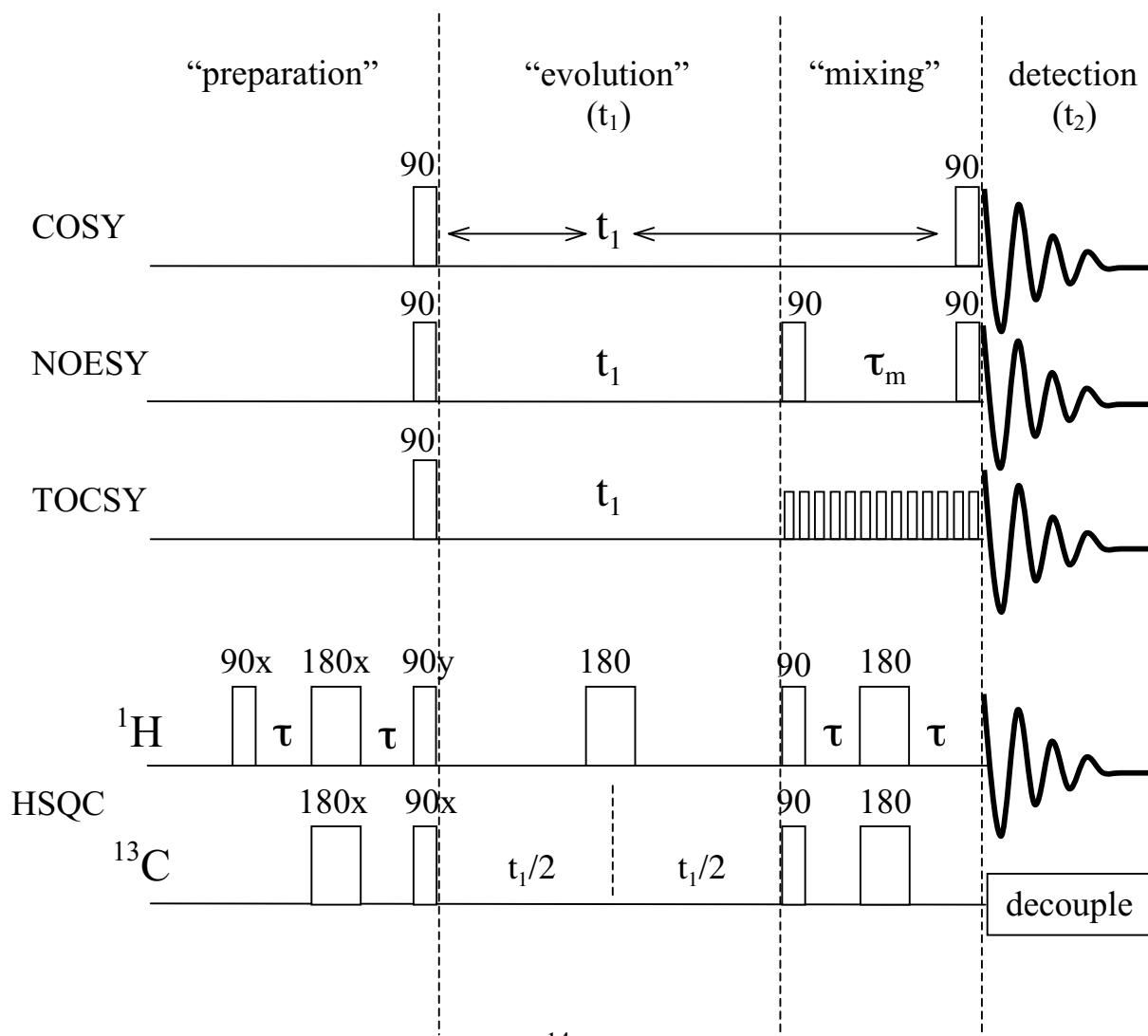
- Very useful experiment for determining the structures of oligosaccharides and peptides as well as many other organic compounds.

1. Pulse sequence:

The pulse sequence is similar to that of the COSY experiment except that the second ^1H 90° pulse is replaced by a spin-lock pulse train:

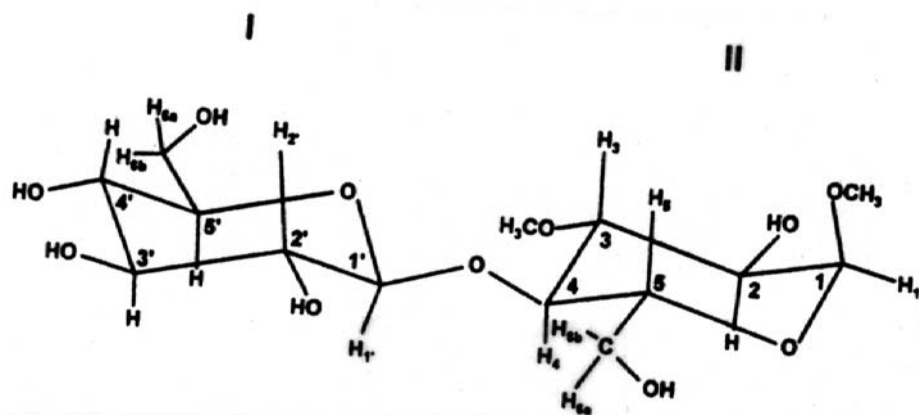
90° - t_1 - spin-lock(t_m) - FID

-The spin-lock allows polarization transfer from one selected proton to all protons within a coupled system (a.k.a. spin system).



2. Spin systems

Lets take α -methyl-3-O-methylcellobioside for example.



- In sugar I:
 - H1' is coupled to H2' (i.e. $J \neq 0$)
 - H2' is coupled to H3'
 - H3' is coupled to H4'
 - H4' is coupled to H5'
 - H5' is coupled to H6'a and H6'b ($J_{H5'-H6'a} > J_{H5'-H6'b}$)
 - H6'a is coupled to H6'b

H1', H2', H3', H4', H5', H6'a, and H6'b form a spin system
- In sugar II:
 - H1 is coupled to H2 (i.e. $J \neq 0$)
 - H2 is coupled to H3
 - H3 is coupled to H4
 - H4 is coupled to H5
 - H5 is coupled to H6a and H6b ($J_{H5-H6a} > J_{H5-H6b}$)
 - H6a is coupled to H6b

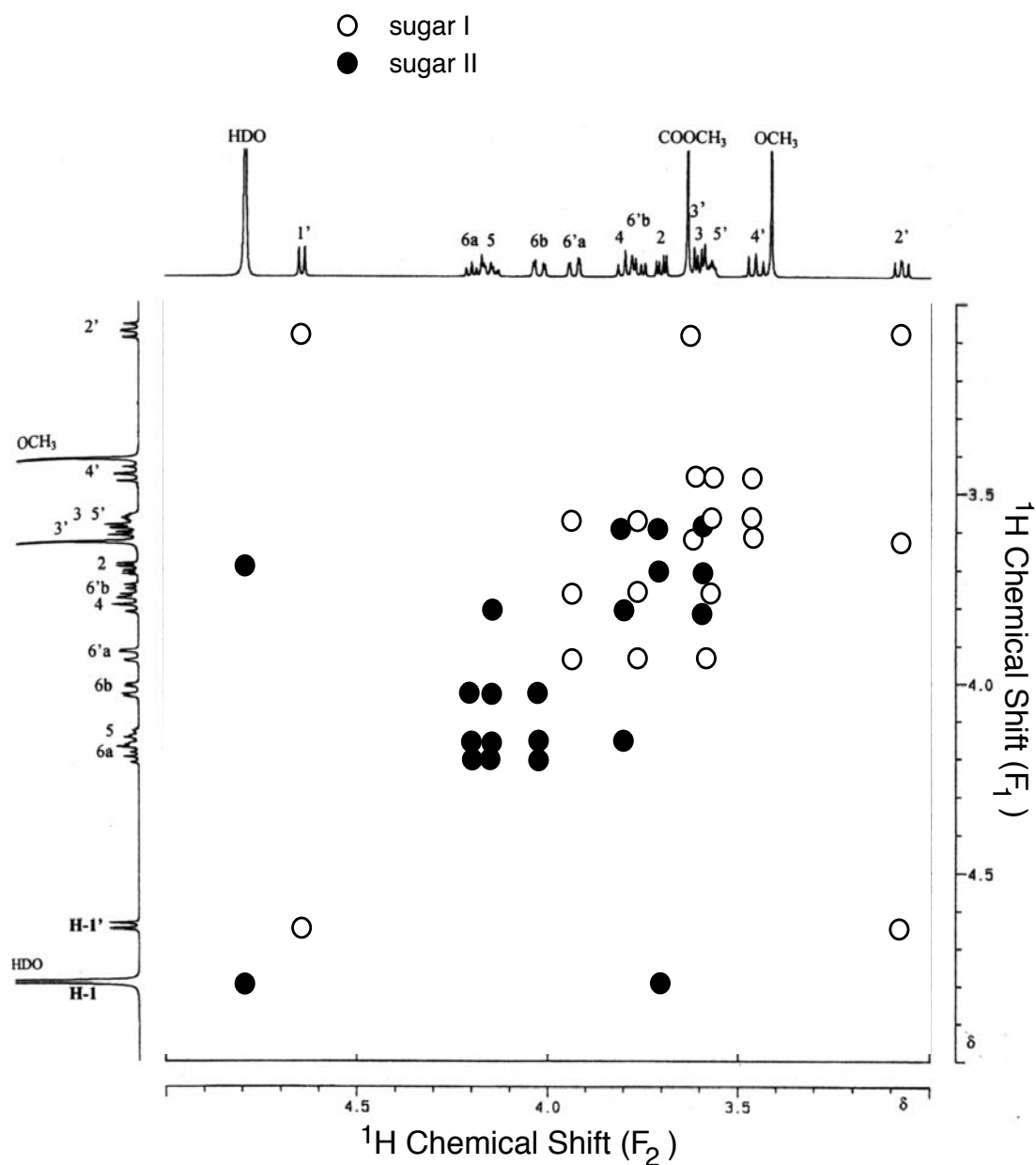
H1, H2, H3, H4, H5, H6a, and H6b form a spin system.
- Because there is no ^1H - ^1H J coupling between sugar I and sugar II, they both form independent spin systems.

3. Differences between COSY and TOCSY

- In a COSY spectrum, we observe one crosspeak for H1' (F2):

H1' (F2)-H2' (F1)

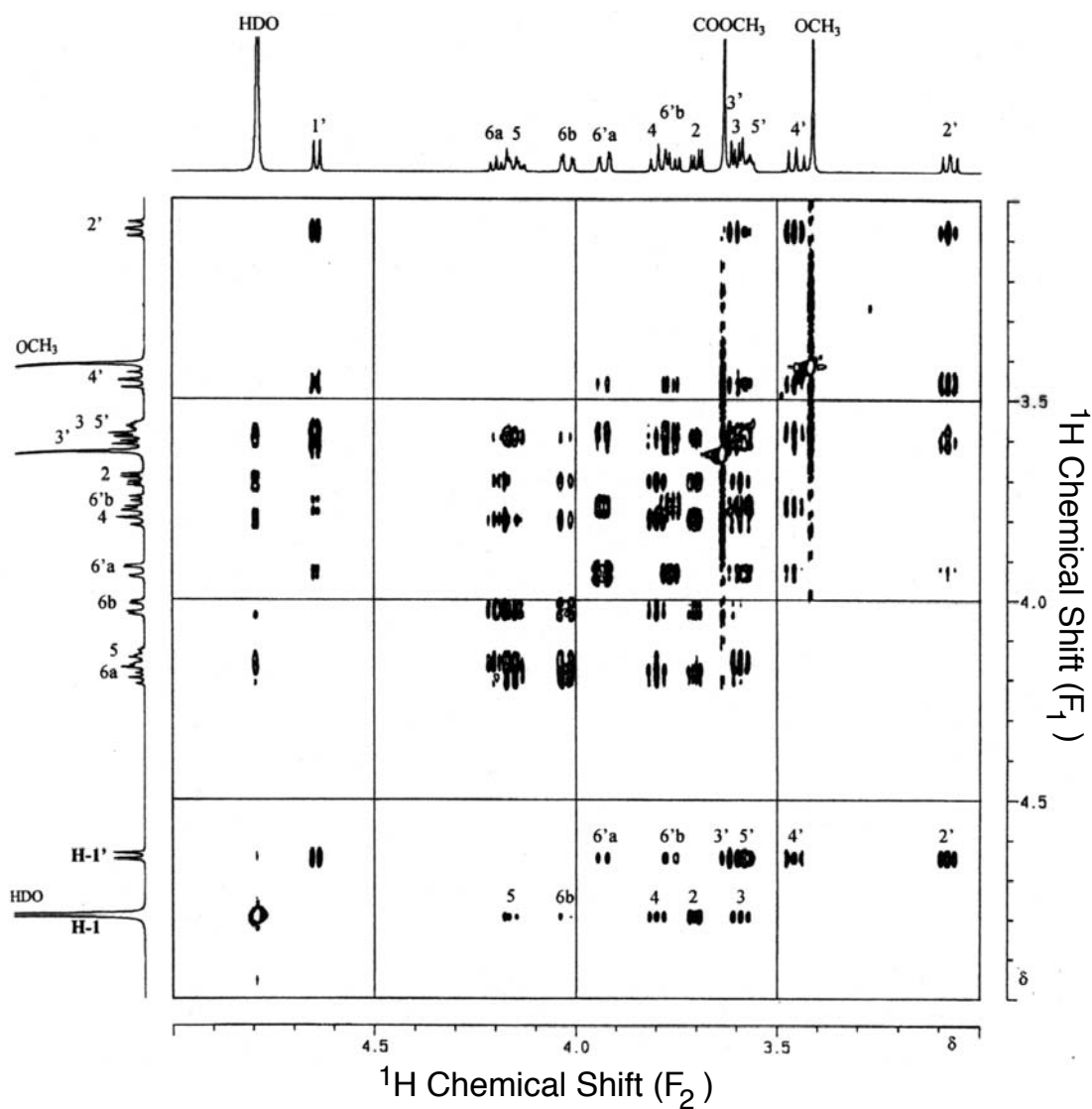
- We also observed the symmetrical peak: H2'(F2) -H1' (F1))
- Simulated COSY spectrum of α -methyl-3-O-methylcellobioside:



- In a TOCSY spectrum, we observe multiple crosspeaks for H1'

<u>F2</u>	<u>F1</u>
H1'	H2'
H1'	H3'
H1'	H4'
H1'	H5'
H1'	H6'a
H1'	H6'b

- We also observed the symmetrical peaks across the diagonal.
- TOCSY spectrum of α -methyl-3-O-methylcellobioside:



4. Importance of the spin-lock mixing period

- We also observe multiple crosspeaks for H1 in the TOCSY spectrum:

<u>F2</u>	<u>F1</u>
H1	H2
H1	H3
H1	H4
H1	H5
H1	H6b

But not H1 to H6a !!!! and H1 to H6b is very weak!

- The range to which crosspeaks can be detected depends on the length of the mixing time (τ_m) where the spin-lock is applied. τ_m usually varies from 10 ms - 200 ms.
- Very short τ_m gives few crosspeaks (more like the COSY) and very long τ_m gives a total correlation spectrum.