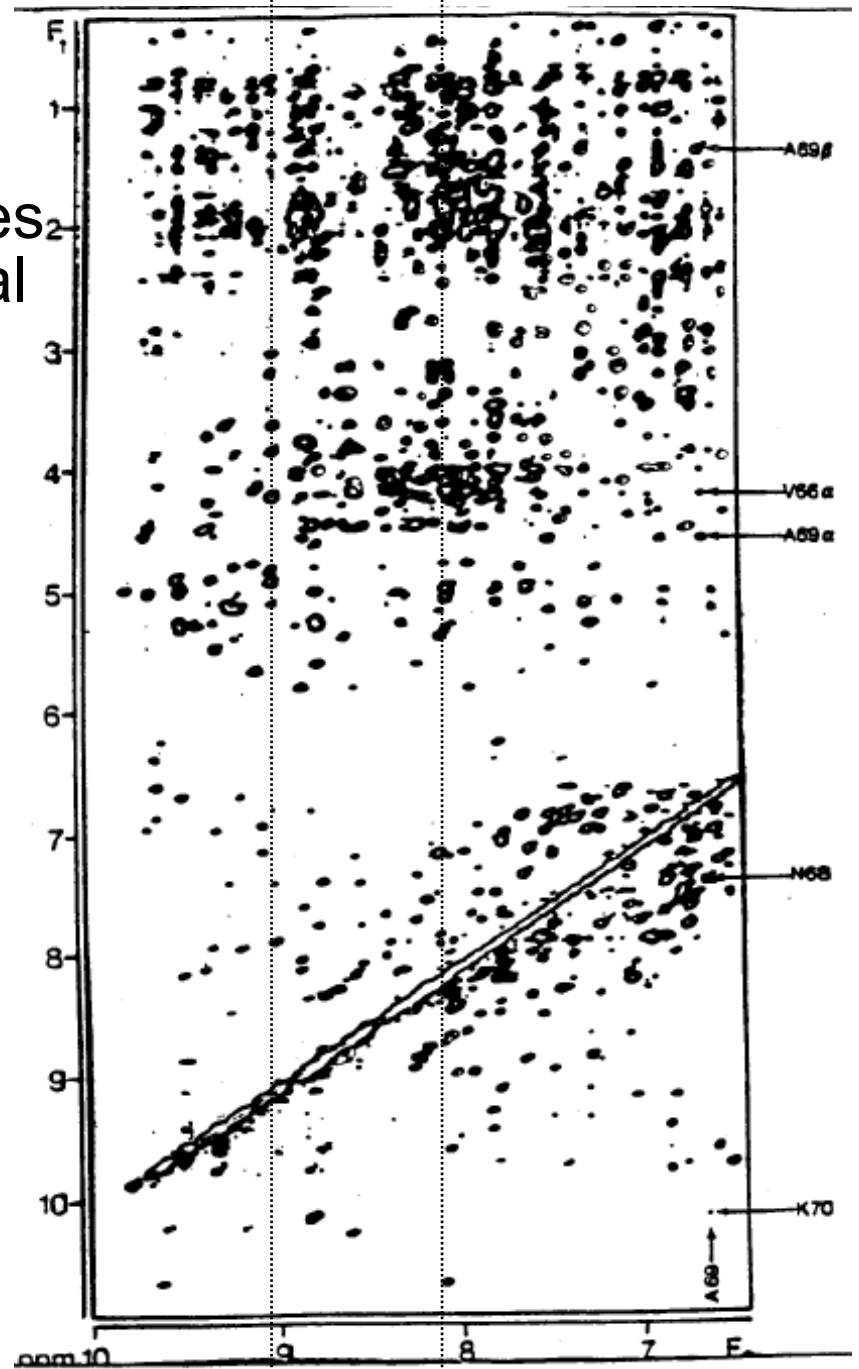


Extensions to 3D  
and  
Improving Efficiency with Pulsed Field Gradients

BCMB/CHEM 8190

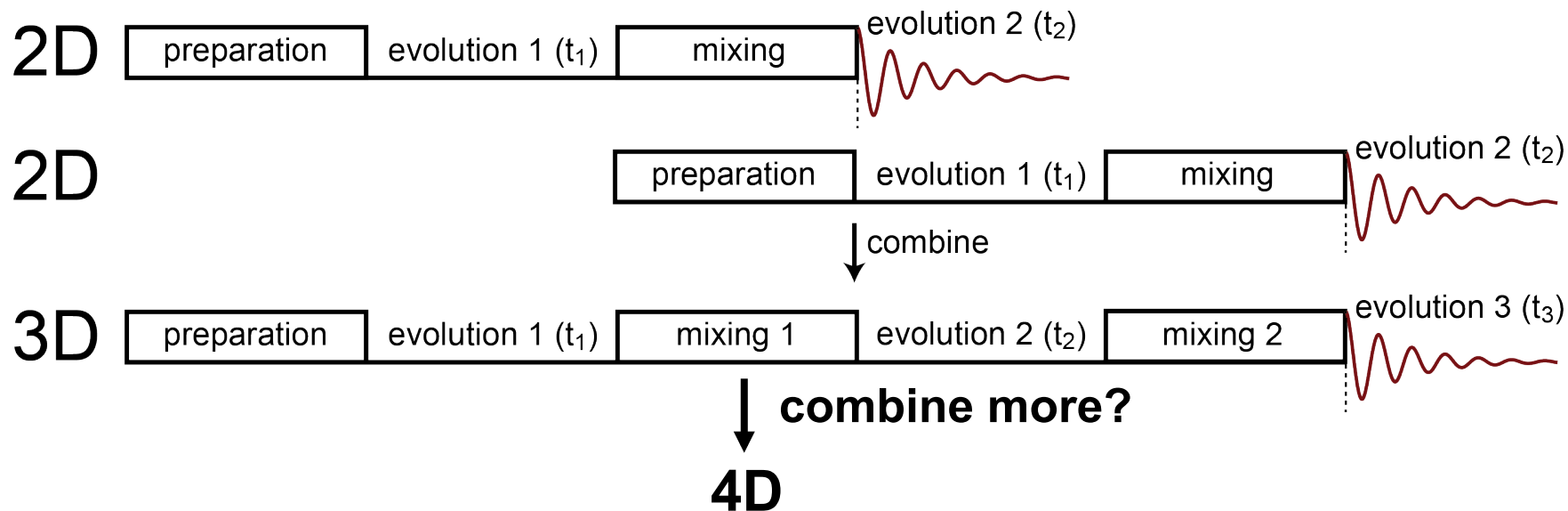
# Crowding in 2D NMR Spectra

- 2D NMR spectra of biomolecules are subject to problematic signal overlap
  - here (right) is the amide region of a 2D NOESY spectrum (Staph nuclease, 156 amino acids)
  - each peak represents a short  $^1\text{H}$ - $^1\text{H}$  distance between an amide  $^1\text{H}$  and some other  $^1\text{H}$
  - a single 'column' represents all  $^1\text{H}$  nuclei within  $\sim 5\text{\AA}$  of a particular amide  $^1\text{H}$  nucleus (dashed lines)
  - possible to unambiguously identify (assign) a small percentage of these in 2D spectra, but impossible to unambiguously assign a majority
  - this is the least crowded region (aliphatic region much worse!)
- Solution: 3D



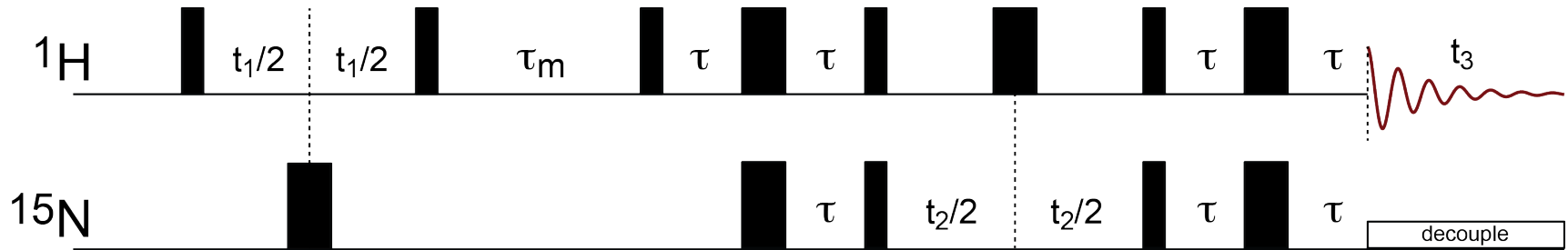
# One Strategy to Alleviate Signal Overlap: Heteronuclear Edited 3D Experiments

- 3D experiments can be built by combining 2D experiments
  - a 2D experiment can be used as the preparation element of another 2D experiment to create a 3D experiment



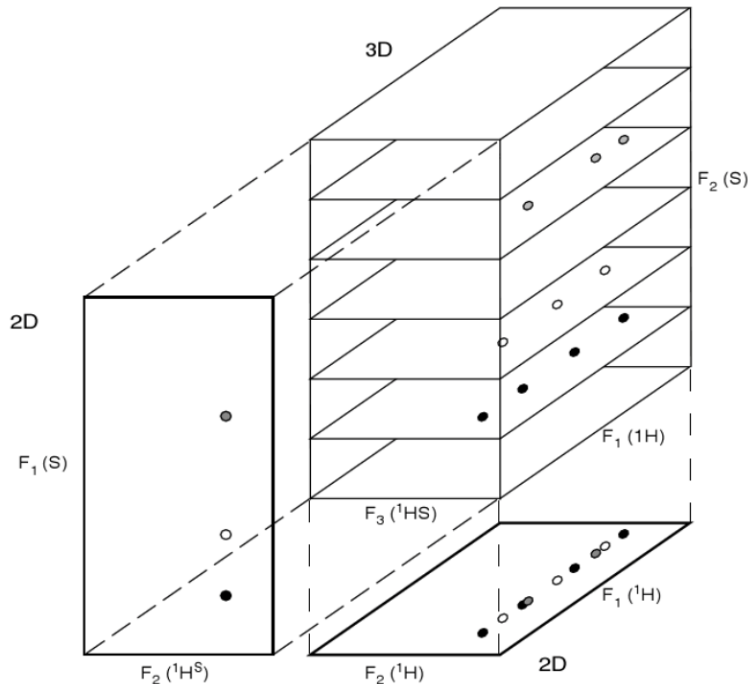
- Resolve signals in third dimension by correlations to  $^{15}\text{N}$  or  $^{13}\text{C}$ 
  - example: for NOESY signals involving amide  $^1\text{H}$  nuclei add another dimension correlating the amide  $^{15}\text{N}$
  - signals are separated in the 3<sup>rd</sup> dimension based on this  $^{15}\text{N}$  chemical shift
  - "NOESY-HSQC"
- Resolution and S/N great, but acquisition is long (hours to days)

# 3D NOESY-HSQC

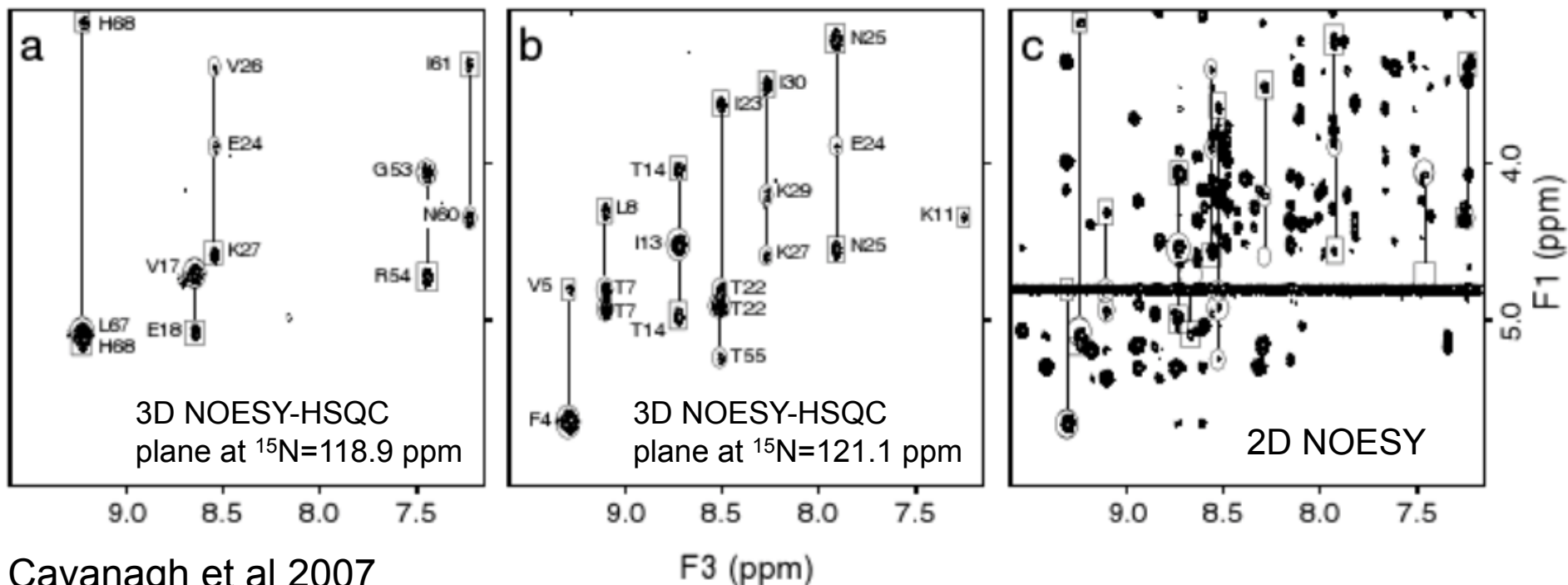


- NOESY sequence serves as preparation period for HSQC
  - the final pulse in the NOESY experiment generates the transverse magnetization for the initial INEPT of the HSQC experiment
  - only  $^1\text{H}$  NOE crosspeaks where one of the  $^1\text{H}$  nuclei is directly bonded to a  $^{15}\text{N}$  nucleus are observed
- Prerequisites
  - need uniform  $^{15}\text{N}$  (or  $^{13}\text{C}$ ) labeling (more on this in later lectures)
- Importance
  - NOE-based distance restraints are the most important restraints for traditional studies for determining high resolution protein structures by NMR
  - It is critical to be able to unambiguously assign the NOE crosspeaks, so 3D (and higher dimension) experiments are essential
- Time consuming
  - if each 2D NOESY experiment takes 1 hour, and you need 64 complex points in the  $^{15}\text{N}$  dimension, that's 128 2D experiments (128 hours)!

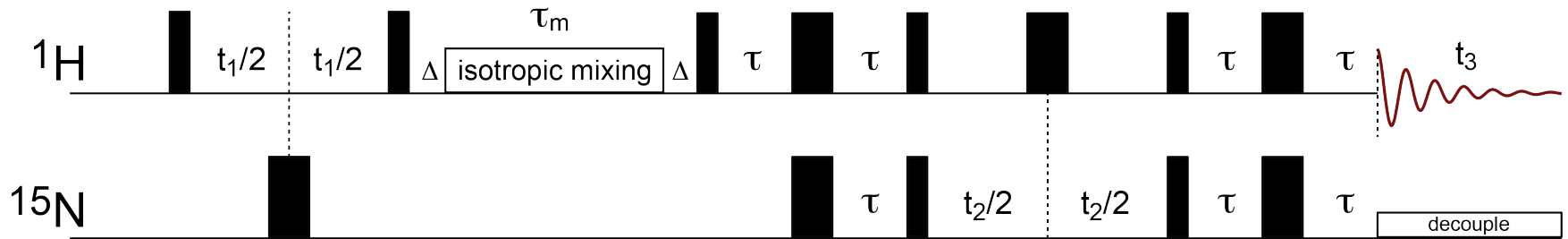
# 3D NOESY-HSQC



- 2D planes of 3D NOESY-HSQC much easier to analyze
  - data shown are for the very small (76 amino acids) protein ubiquitin
  - significant reduction in peak overlap in the planes compared to the 2D spectrum

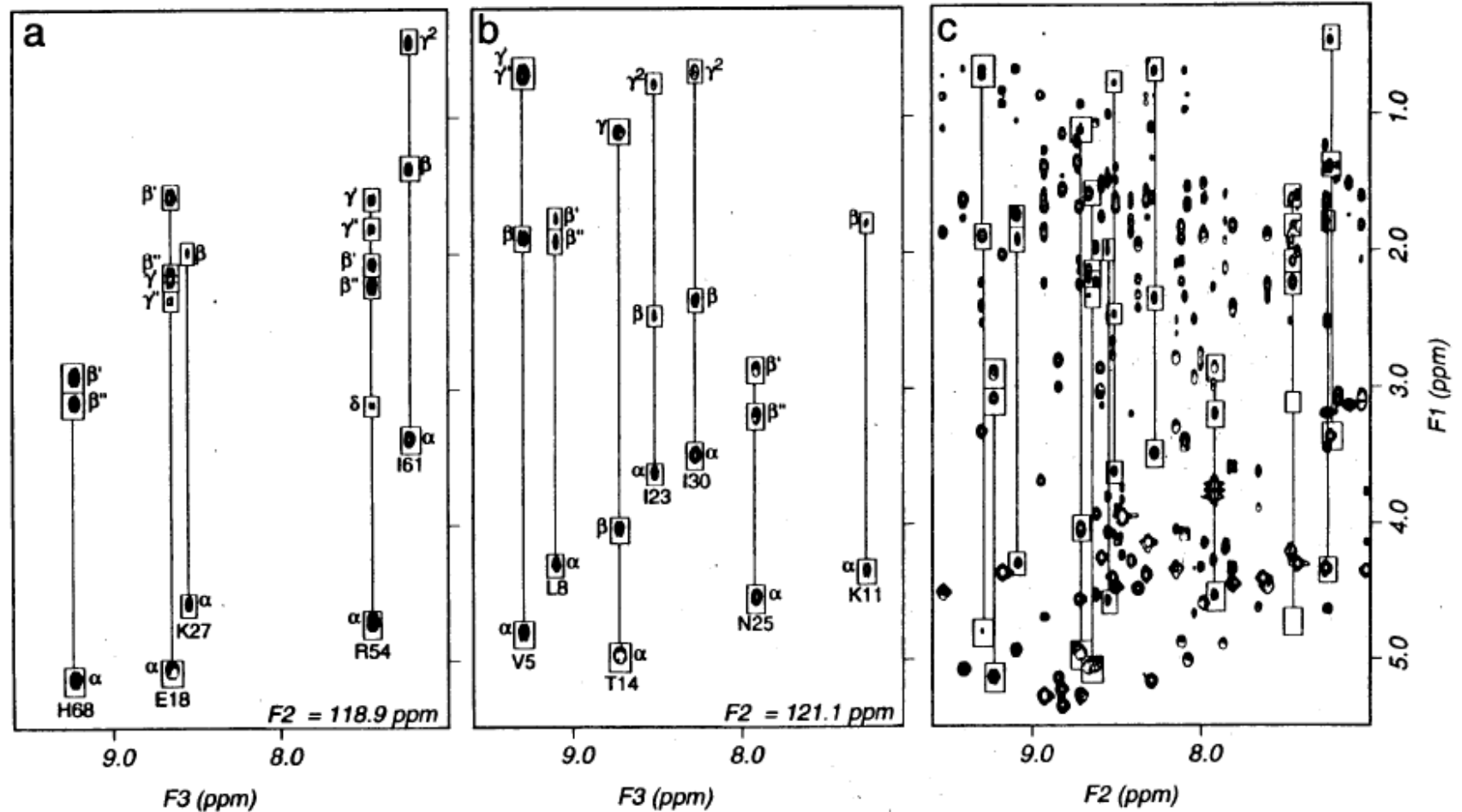


# 3D TOCSY-HSQC



- TOCSY sequence serves as preparation period for HSQC
  - the final pulse in the TOCSY experiment generates the transverse magnetization for the initial INEPT of the HSQC experiment
  - only TOCSY spin systems where one of the  $^1\text{H}$  nuclei is directly bonded to a  $^{15}\text{N}$  nucleus are observed
- Prerequisites
  - need uniform  $^{15}\text{N}$  (or  $^{13}\text{C}$ ) labeling
- Importance
  - identifying spin systems and correlating all chemical shifts in each
  - helpful in identifying amino acid type (side chain chemical shifts are often diagnostic of the amino acid type)
- Time consuming
  - like NOESY-HSQC

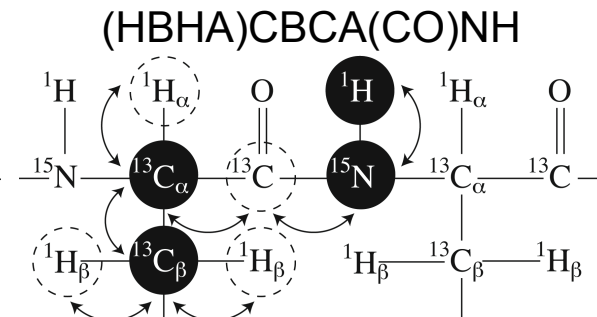
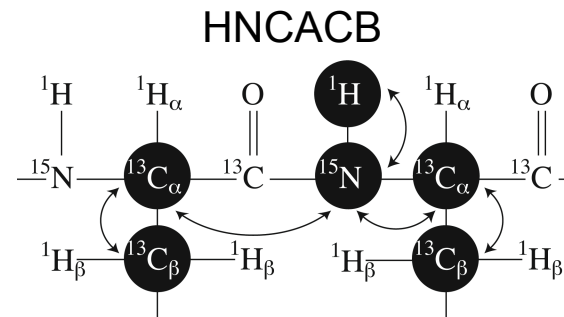
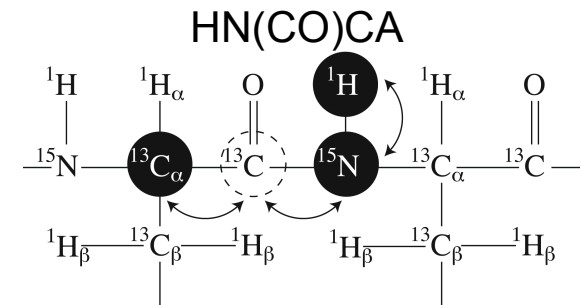
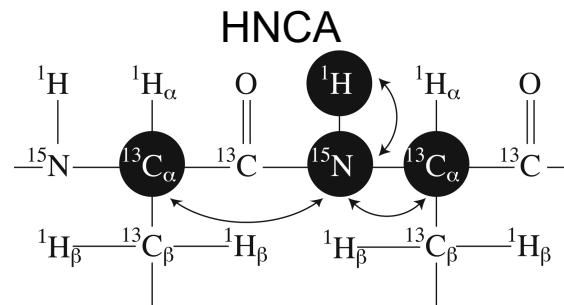
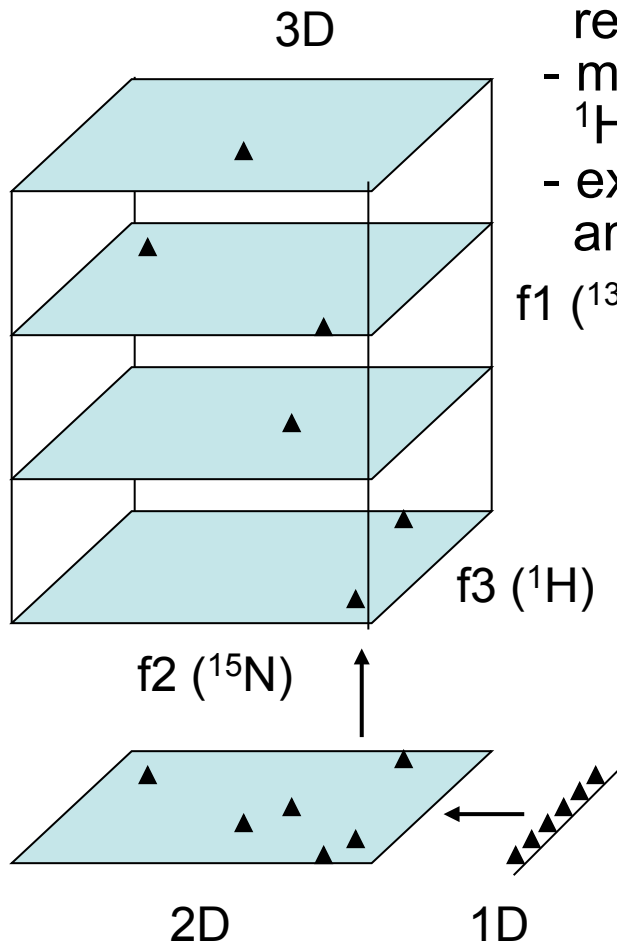
# 3D TOCSY-HSQC for $^{15}\text{N}$ -Labeled Ubiquitin



# Cavanagh et al 1996

# 3D Triple Resonance Experiments

- Triple resonance experiments correlate multiple nuclear types (typically  $^1\text{H}$ ,  $^{13}\text{C}$  and  $^{15}\text{N}$ , each in its own dimension)
  - these aren't necessarily simple combinations of 2D experiments, but they employ similar pulse sequence elements (INEPT, COSY, etc.)
  - are dozens of these experiments, critical for assigning resonances of biomolecules
  - magnetization transferred via scalar couplings, with  $^1\text{H}$  detection (inverse detect experiments)
  - examples: (names have meaning, HNCA correlates amide  $^1\text{H}$ , amide  $^{15}\text{N}$  and alpha carbon  $^{13}\text{C}$ )



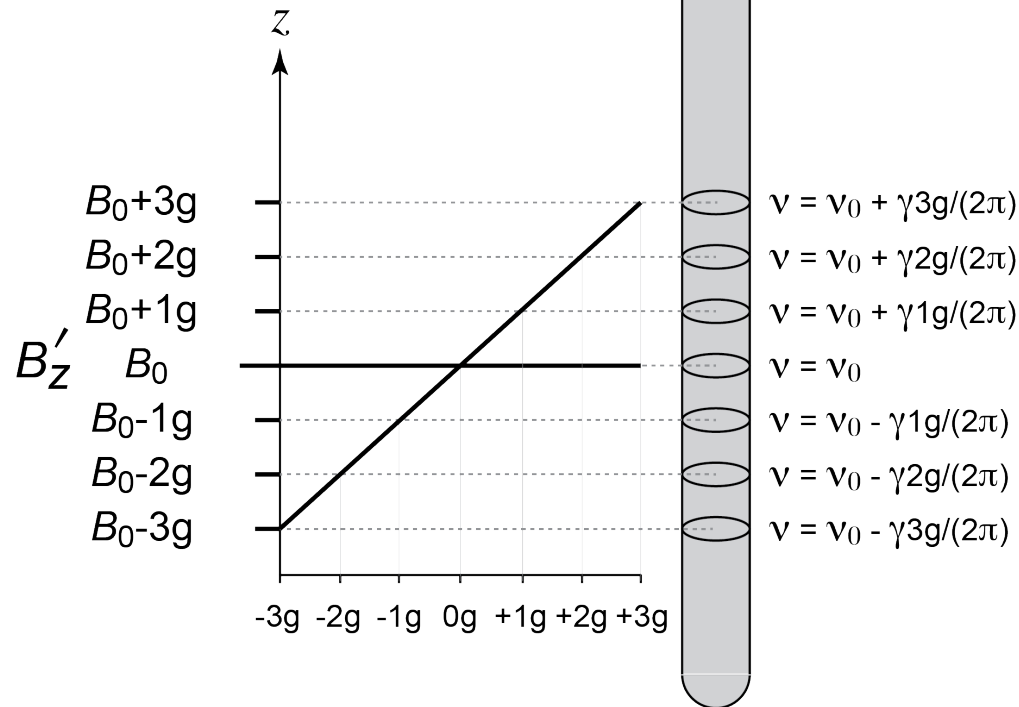
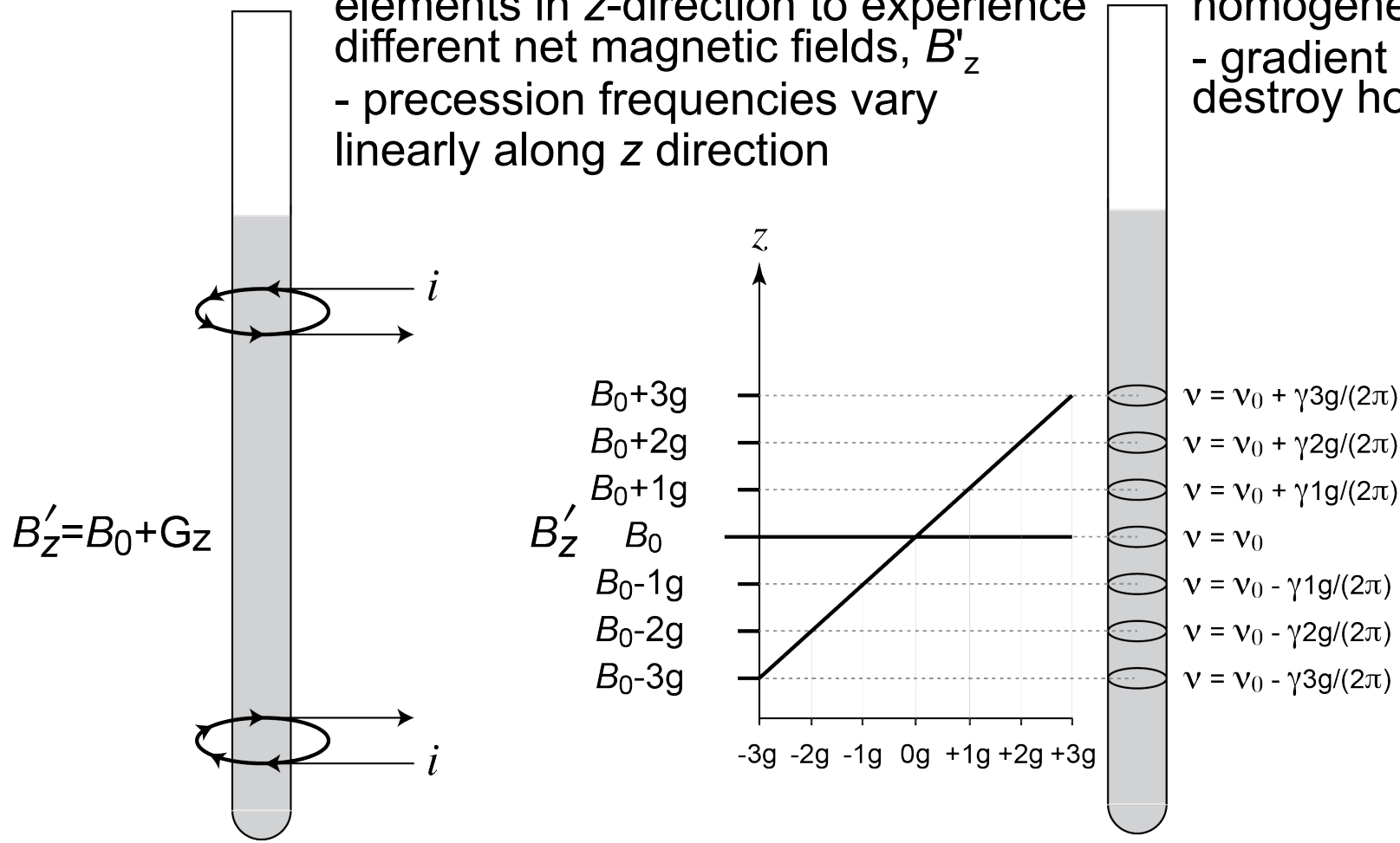


# Pulsed Field Gradients: Improved Efficiency in Multidimensional Spectra

- Pulsed field gradients are used in many important ways in modern biomolecular NMR experiments
  - they can be used in a very simple way to remove spurious signals and unwanted magnetization
  - they can be used to reduce phase cycling (thus significantly decreasing overall time necessary to acquire an experiment)
  - they can be used to suppress water signals (and other signals as well)
  - they can be used for coherence selection
  - they can be used to measure diffusion
- Coherence selection using pulsed field gradients. J. R. Tolman & J. H. Prestegard, *Concepts in Magnetic Resonance*, **7**, 247-262 (1995).
- Water suppression (WATERGATE), M. Piotto, V. Saudek & V. Sklenar, *J. Biomol. NMR*, **2**, 661-665 (1992).
- Diffusion measurements. Altieri, Hinton & Byrd, *J. Am. Chem. Soc.*, **117**, 7566-7567 (1995).

# Pulsed Field Gradients – How they Work

- Similar to a shim coil
  - simplest 'Z' shim coil, two loops of wire with current running in opposite directions, sets up a linear magnetic field gradient along the z-axis
  - Z gradient coil is similar, but used to apply transient current (pulses)
  - Z-gradient pulse causes volume elements in z-direction to experience different net magnetic fields,  $B'_z$
  - precession frequencies vary linearly along z direction
  - shim coils: improve homogeneity
  - gradient pulses, destroy homogeneity



# Shim Coils and Gradient Coils

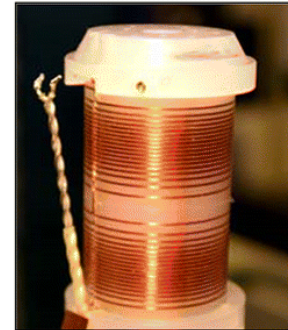


Dewar



room temperature  
shim coils

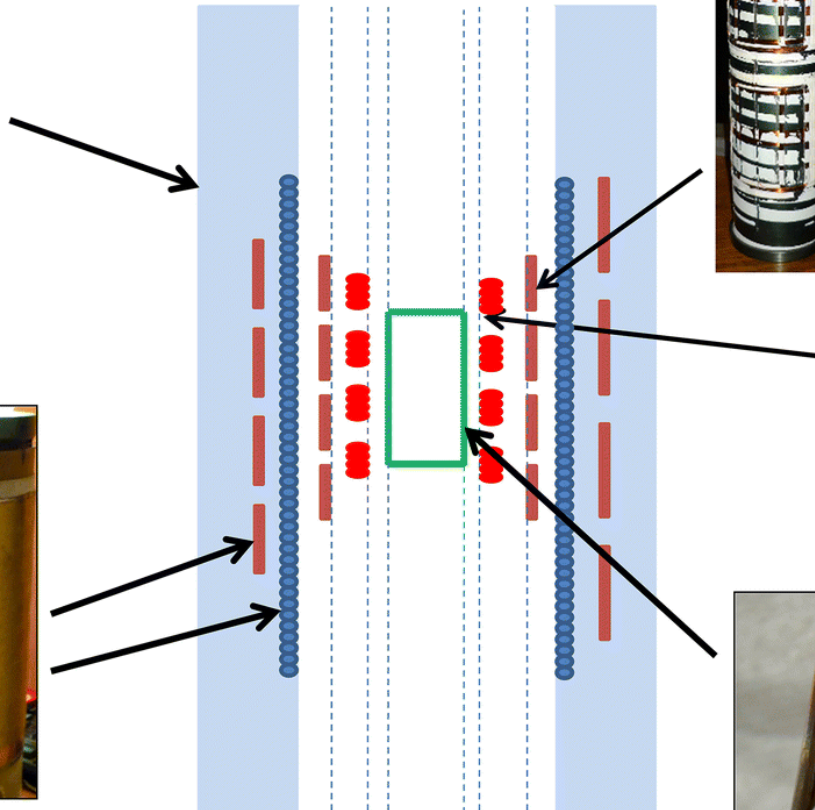
gradients



superconducting  
magnet + shims

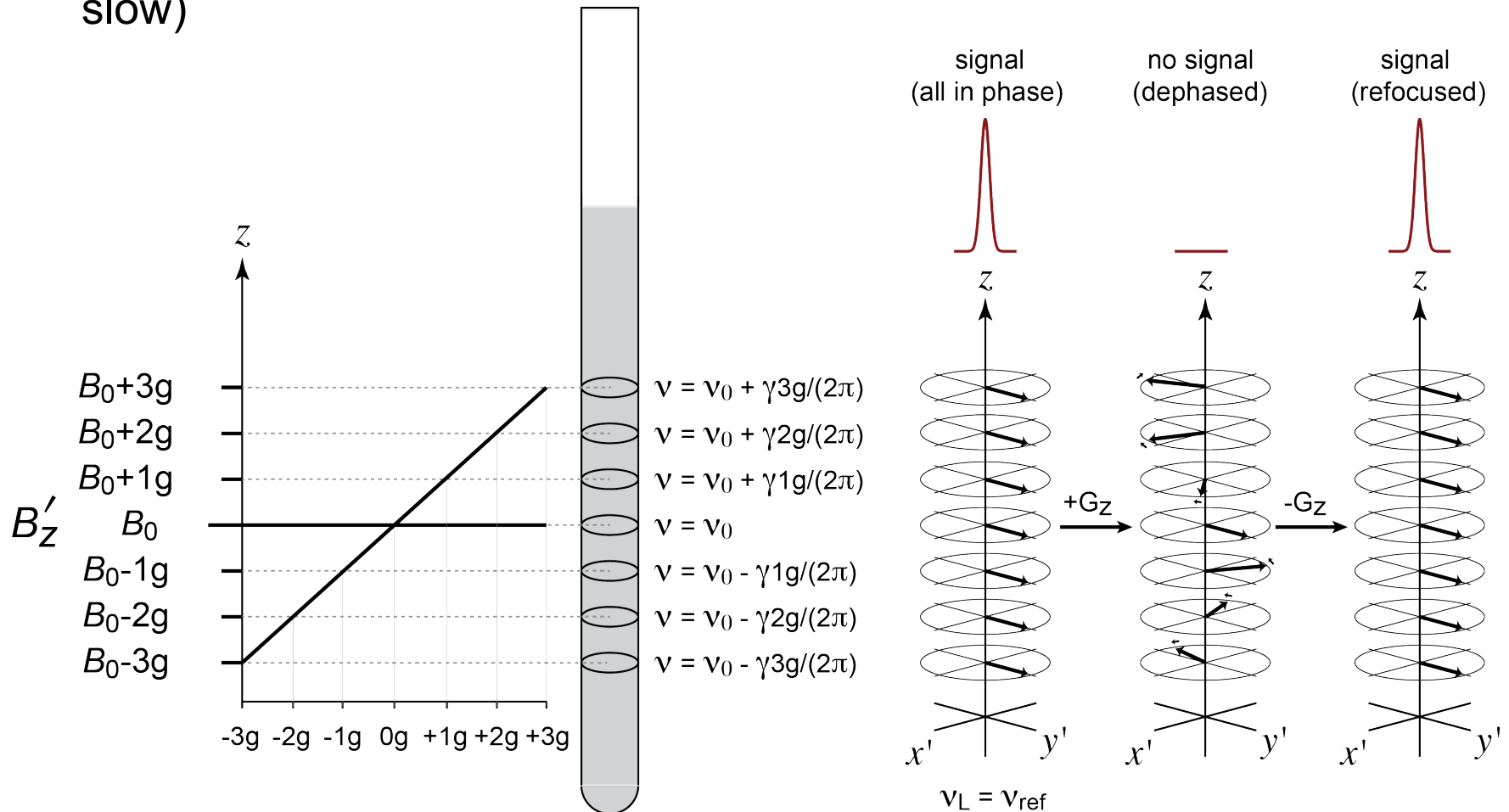


RF coil



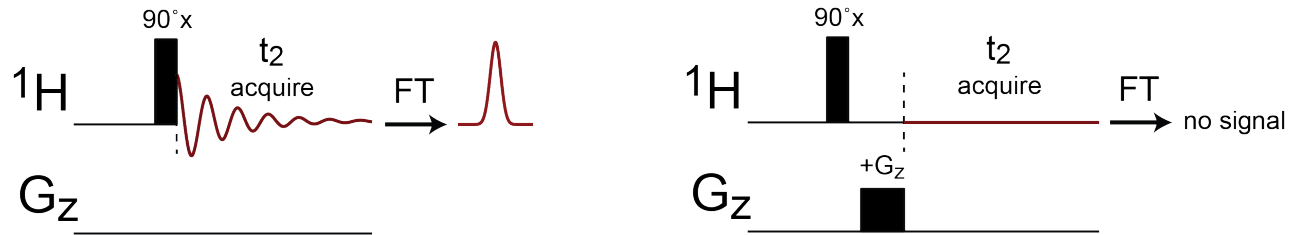
# Pulsed Field Gradients – How they Work

- Gradient pulses defocus and refocus magnetization
  - applying a Z gradient pulse on coherent transverse magnetization causes defocusing (dephasing) of magnetization in the transverse plane
  - however, the magnetization can be refocused by applying a second, identical gradient pulse with opposite polarity (works because diffusion is slow)

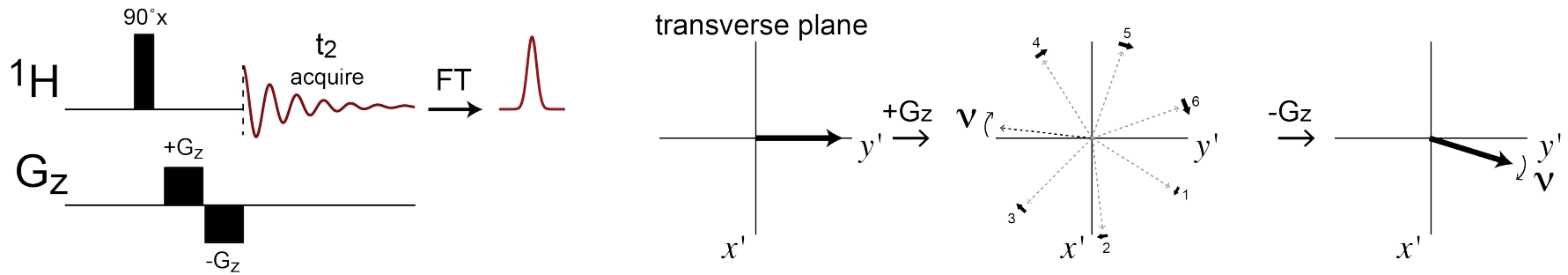


# Defocusing/Refocusing, Encoding/Decoding

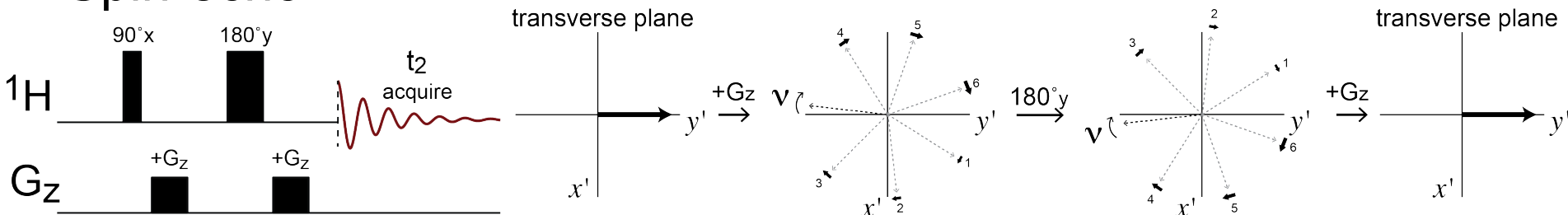
- Field gradient pulses encode a spatially dependent phase, that can be decoded by another gradient pulse
  - as on the previous page, a simple example is here, whereby application of a gradient pulse encodes a spatially dependent phase, which is decoded by the second gradient pulse of opposite polarity



- Simple defocus/refocus (spatial encoding/decoding)



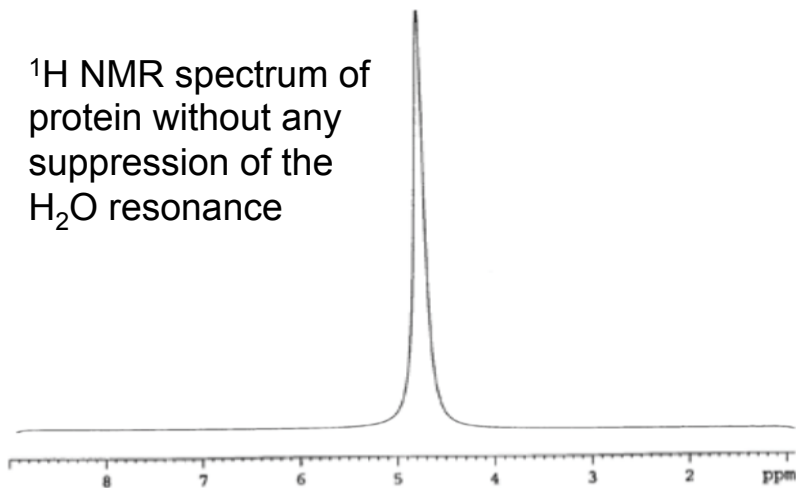
- Spin-echo



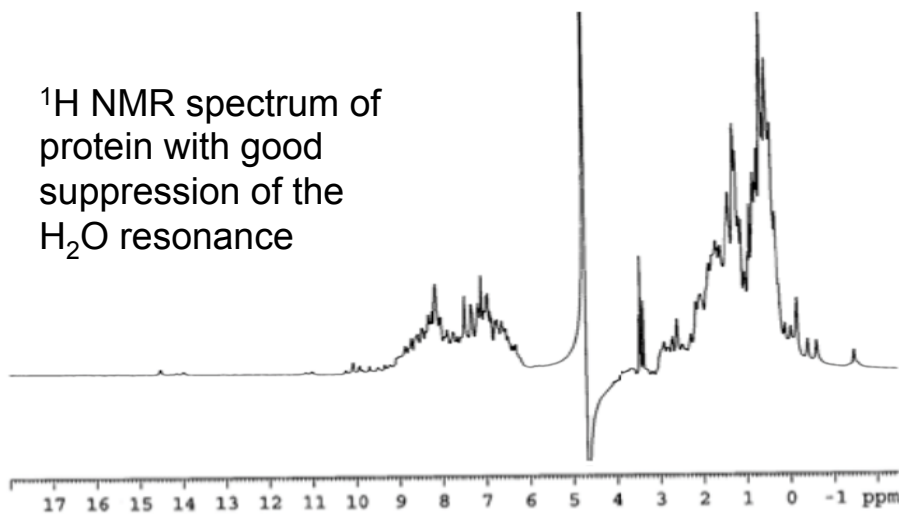
# Field Gradients Application: Water Suppression

- Much of biomolecular NMR is done in  $\text{H}_2\text{O}$ 
  - this is the natural milieu for proteins, nucleic acids, etc.
  - replacing with  $\text{D}_2\text{O}$  can cause some subtle effects on stability, dynamics, etc.
  - amide  $^1\text{H}$  hydrogens exchange with solvent: if D replaces H, can't detect amide  $^1\text{H}$
- Many experiments detect amide  $^1\text{H}$  magnetization
  - very convenient: signals are further downfield than most other  $^1\text{H}$  signals in proteins, well resolved, 1 per amino acid residue
  - so, NOESY-HSQC, TOCSY-HSQC, and many, many triple resonance experiments detect amide  $^1\text{H}$  magnetization
- Problem: protein in  $\text{H}_2\text{O}$  only observable with good suppression of large  $\text{H}_2\text{O}$  signal
  - $\text{H}_2\text{O}$  is 55 M  $\text{H}_2\text{O}$  (110 M protons), protein 1 mM (110,000 fold difference)
  - with good  $\text{H}_2\text{O}$  suppression, protein signals easily observed

$^1\text{H}$  NMR spectrum of protein without any suppression of the  $\text{H}_2\text{O}$  resonance

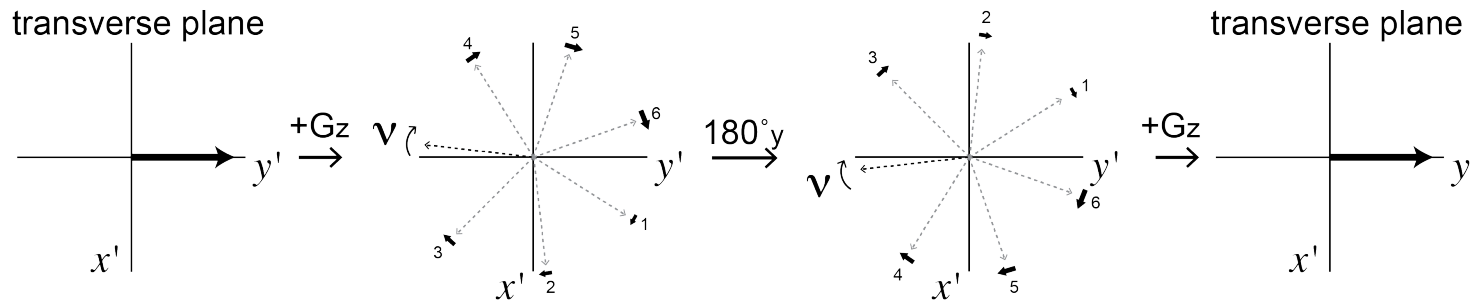
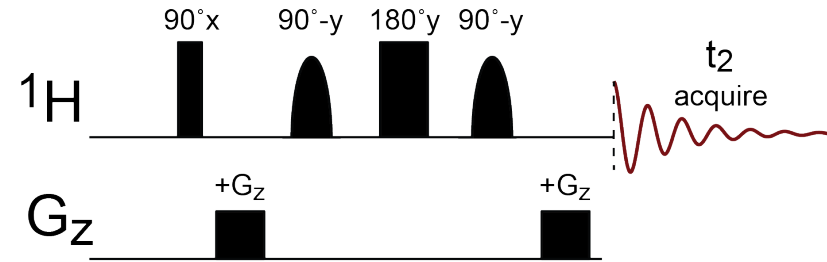


$^1\text{H}$  NMR spectrum of protein with good suppression of the  $\text{H}_2\text{O}$  resonance

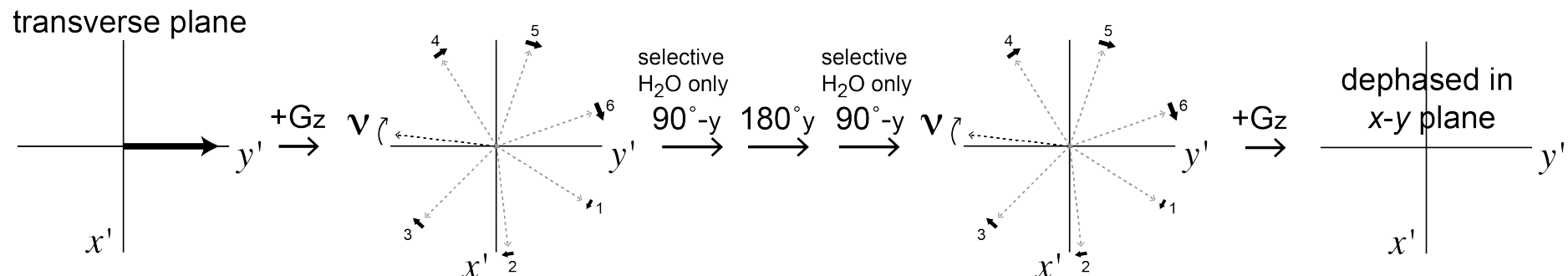


# Field Gradients Application: Water Suppression

- This application uses field gradient pulses, and frequency selective pulses (selective for  $\text{H}_2\text{O}$ ) to reduce in great measure the signal from  $\text{H}_2\text{O}$
- First, consider all signals other than  $\text{H}_2\text{O}$ 
  - these will not be affected by the pulses selective for  $\text{H}_2\text{O}$ , so the result is just the spin-echo we saw previously



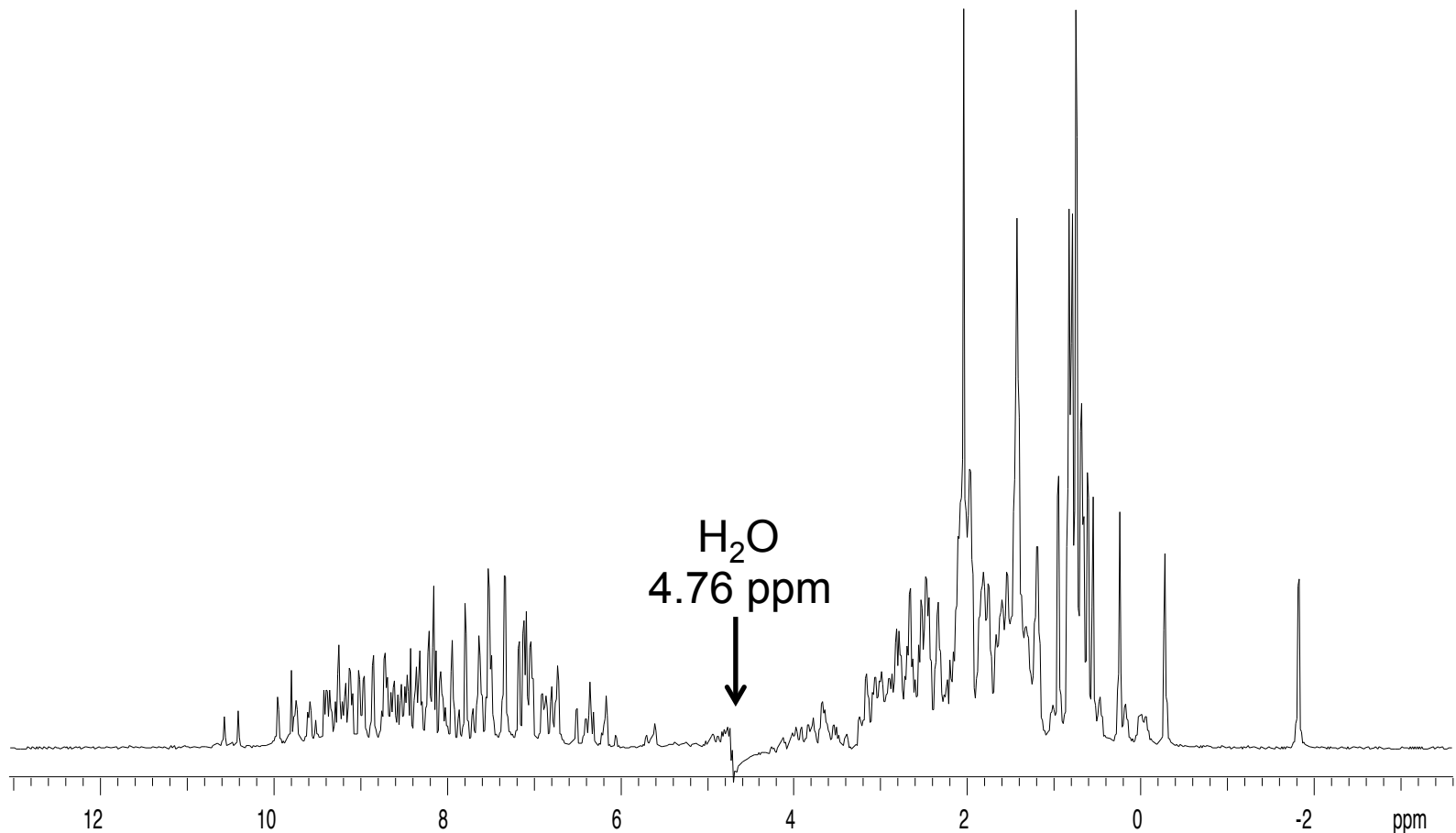
- For  $\text{H}_2\text{O}$ , the  $90^\circ -y$ ,  $180^\circ y$ ,  $90^\circ -y$  pulses move all  $\text{H}_2\text{O}$  magnetization back to where it started (dephased)
  - the second gradient pulse then just continues dephasing the  $\text{H}_2\text{O}$  (so, not observed)





# 1D $^1\text{H}$ Water-Suppressed Spectrum *Pf*-Rubredoxin in $^1\text{H}_2\text{O}$

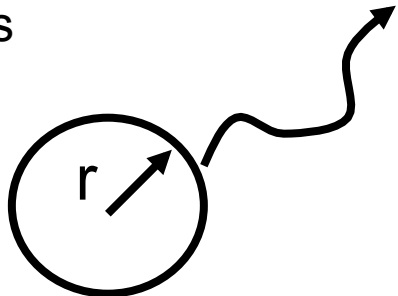
- Works remarkably well (combined with very good shimming, good phase cycling, very accurate pulse angle calibrations, etcetera)





# Translational Diffusion Constants for (Macro)molecules

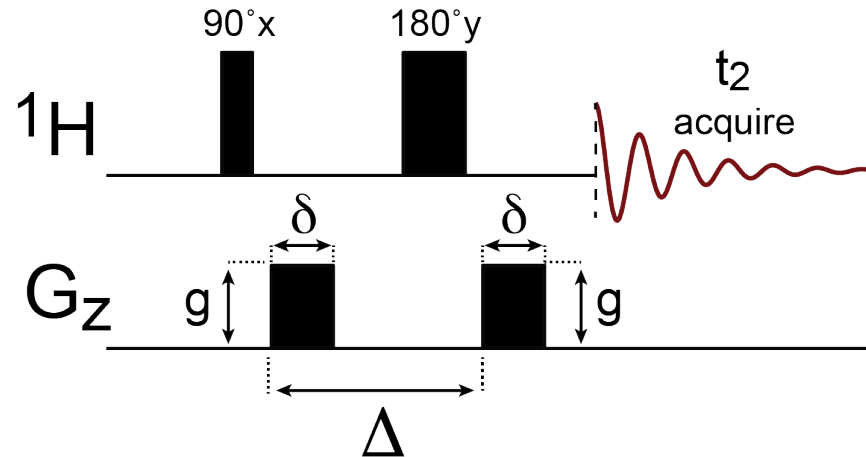
- NMR methods employing pulsed field gradients (PFG) are commonly used to study molecular diffusion
  - the PFG experiments we've seen thus far depend on the fact the diffusion is very slow on the NMR timescale
  - however, it is not insignificant, and PFG methods can be employed to measure diffusion constants and relative translational diffusion rates
- Diffusion studies can be used for many purposes:
  - measure the effective size of a protein
  - study protein complexes or aggregates
  - analyze/detect protein aggregation
  - identify and study protein-protein interactions
  - screen for ligand binding and otherwise study interactions of proteins with ligands
- The average displacement (squared) is proportional to the diffusion constant
  - the diffusion constant,  $D$ , is dependent on viscosity ( $\eta$ ) and molecular size ( $r$ , radius)



$$(Z_1 - Z_0)^2 / N = Dt \quad D = kT / (6\pi\eta r) \text{ (Stokes formula)}$$
  - $D \propto \sqrt[3]{MW}$  ( $MW \propto r^3$ )
- PFG methods for measuring diffusion rely on the fact that magnetization dephased with a gradient pulse will not rephase fully if the molecule moves (remember, phases are spatially encoded)

# Measuring Diffusion Using Field Gradient Pulses

- Stejskal and Tanner pulse sequence for diffusion measurement



- This is just the gradient 'spin-echo' experiment we saw previously
  - if the molecule moves significantly (translationally, z-direction) during the time  $D$  (time between beginning of one gradient pulse and beginning of second), then the magnetization is not refocused fully and the signal decreases
  - thus, diffusion is detected by signal intensity decrease

- Intensity of signal decreases as molecule diffuses according to:

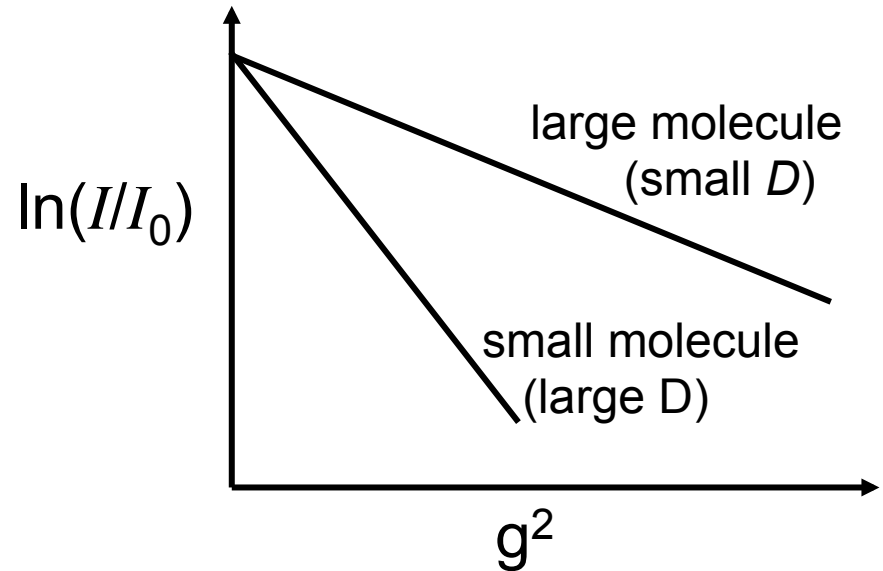
$$I = I_0 e^{[(-2\tau/T_2) - \gamma^2 g^2 \delta^2 D(\Delta - \delta/3)]} \quad \text{linear form, } \ln[I/I_0] = (-2\tau/T_2) - \gamma^2 g^2 \delta^2 D(\Delta - \delta/3)$$

- can measure change in intensity as a function of either  $\delta$ ,  $\Delta$ , or  $g$
- changing  $\delta$  or  $\Delta$  introduces additional complications due to signal loss from relaxation, so the nearly universal practice is to change  $g$

# Measuring Diffusion Using Field Gradient Pulses

$$I = I_0 e^{[(-2\tau/T_2) - \gamma^2 g^2 \delta^2 D(\Delta - \delta/3)]} \quad \text{linear form, } \ln[I/I_0] = (-2\tau/T_2) - \gamma^2 g^2 \delta^2 D(\Delta - \delta/3)$$

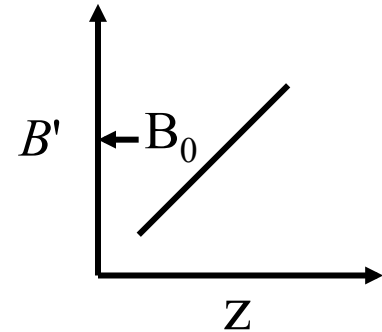
- In linear form, the slope of  $\ln(I/I_0)$  versus  $g^2$  is proportional to  $D$ 
  - units:  $g$  (gauss/cm),  $\gamma$  ( $\text{s}^{-1}\text{gauss}^{-1}$ ),  $D$  ( $\text{cm}^2\text{s}^{-1}$ )
- Small molecules diffuse fast, so plot drops off fast with  $g^2$ , not as fast for large molecules
- Diffusing measurements are limited by  $T_2$ 
  - the simple Stejskal and Tanner pulse sequence is prone to  $T_2$  related problems when working with large molecules
  - newer, improved methods perform better overall, in particularly for larger molecules (for instance, Alteiri et al. (1995) *J. Am. Chem. Soc.* 117, 7566-7567)



# Coherence Selection Using Pulse Field Gradients

- Consider the simple Hamiltonian (neglecting chemical shift) including  $B_z$  (field due to gradient pulse)

$$B'_z = B_0 + B_z \quad \hat{H}(r) = -\sum_k \gamma_k [B_0 + B_z(r)] \hat{I}_{kz}$$



- In order to understand how to select coherence using field gradient pulses, it is convenient to demonstrate their effects on our product operators

- first, recall that we can write operators in terms of raising and lowering operators

$$\hat{I}_x = (\hat{I}_+ + \hat{I}_-)/2 \quad \hat{I}_{kx} = (\hat{I}_{k+} + \hat{I}_{k-})/2$$

- consider the + and - operators as counter-rotating magnetization components

$$I_{kz} \xrightarrow{-\gamma_k B_z(z) I_{kz} \tau} I_{kz}$$

$z$ -magnetization doesn't precess, isn't in transverse plane, so isn't defocused (affected) by gradient pulse

$$I_k^+ \xrightarrow{-\gamma_k B_z(z) I_{kz} \tau} \exp(i\gamma_k B_z(z) \tau) I_k^+$$

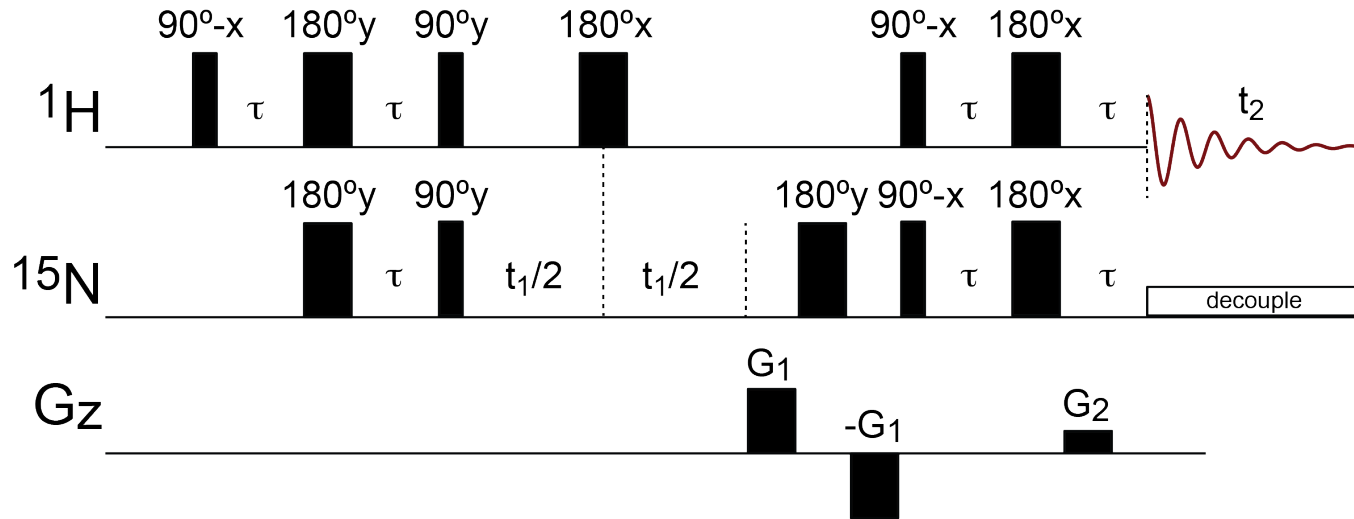
precession due to the field gradient in one direction (i.e. gradient increases precession frequency)

$$I_k^- \xrightarrow{-\gamma_k B_z(z) I_{kz} \tau} \exp(-i\gamma_k B_z(z) \tau) I_k^-$$

precession due to the field gradient in the other direction (i.e. gradient decreases precession frequency)

- For linear ( $z$ ) gradients, we know  $B_z(z)$  (the gradient strength,  $G_z$ )
- Observables ( $M_x, M_y$ ) are integrals over  $z$ 
  - these are zero: proportional to sum of  $I_k^+$  and  $I_k^-$  (why signals go away, no net difference in + and - components, so no net transverse magnetization in any direction)

# Gradient Selected HSQC



- At the end of the  $t_1$  evolution period, the  $^{15}\text{N}$  magnetization is dephased (spatially encoded) twice by  $G_1$  ( $G_1$ ,  $180^\circ$ ,  $-G_1$ ) (i.e. phase encoded by  $\gamma_N 2G_1$ )
- In order to decode (in order to get observable magnetization in  $t_2$ ), after transferring to  $^1\text{H}$ ,  $\gamma_N 2G_1$  must equal  $\gamma_H G_2$  (for maximal signal)

$$I^+(t_2) \propto \int_z \{S^+(t_1) \exp[i\gamma_N 2G_1 z] \exp[-i\gamma_H G_2 z]\}$$

$$\propto \int_z \{S^+(t_1) \exp[i(\gamma_N 2G_1 z - \gamma_H G_2 z)]\}$$

- the only magnetization observed is that defocused by  $G_1/-G_1$  and refocused by  $G_2$
- all spurious magnetization components are defocused by the gradients and never refocused (so, spectra are very clean, no spurious signals, without phase cycling)
- quadrature achieved by cycling gradients, or final  $90^\circ$   $^{15}\text{N}$  pulse ( $+x/-x$ ) and  $G_2$  ( $+/-$ )