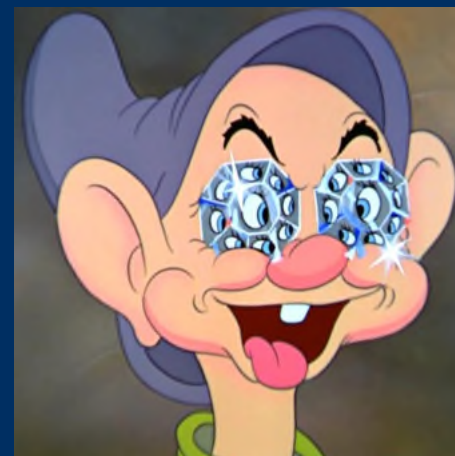


DEPARTMENT OF CHEMISTRY

Polishing GEMSTONES

Optimisation of Ultra-Selective NMR
Experiments

22/06/2023

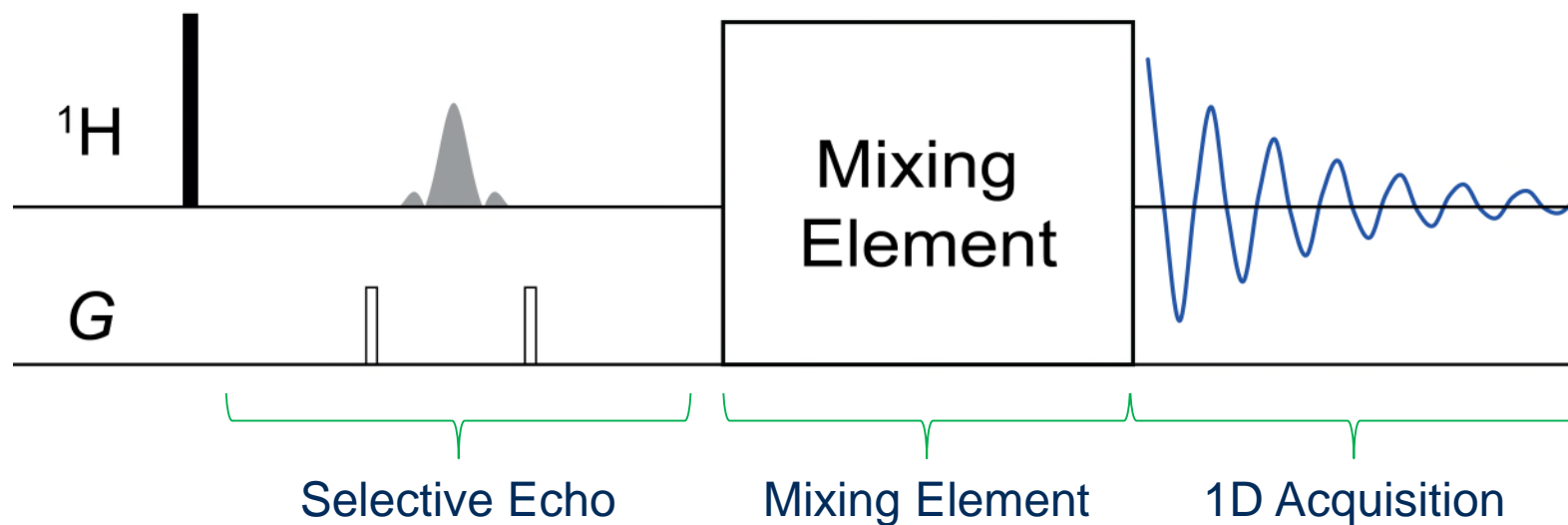


Overview



- Conventional selective experiments & CSSF
- Gradient enhanced selective excitation (i.e GEMSTONE)
- Road to success – implementation (Bruker systems)

Selective NMR Experiments



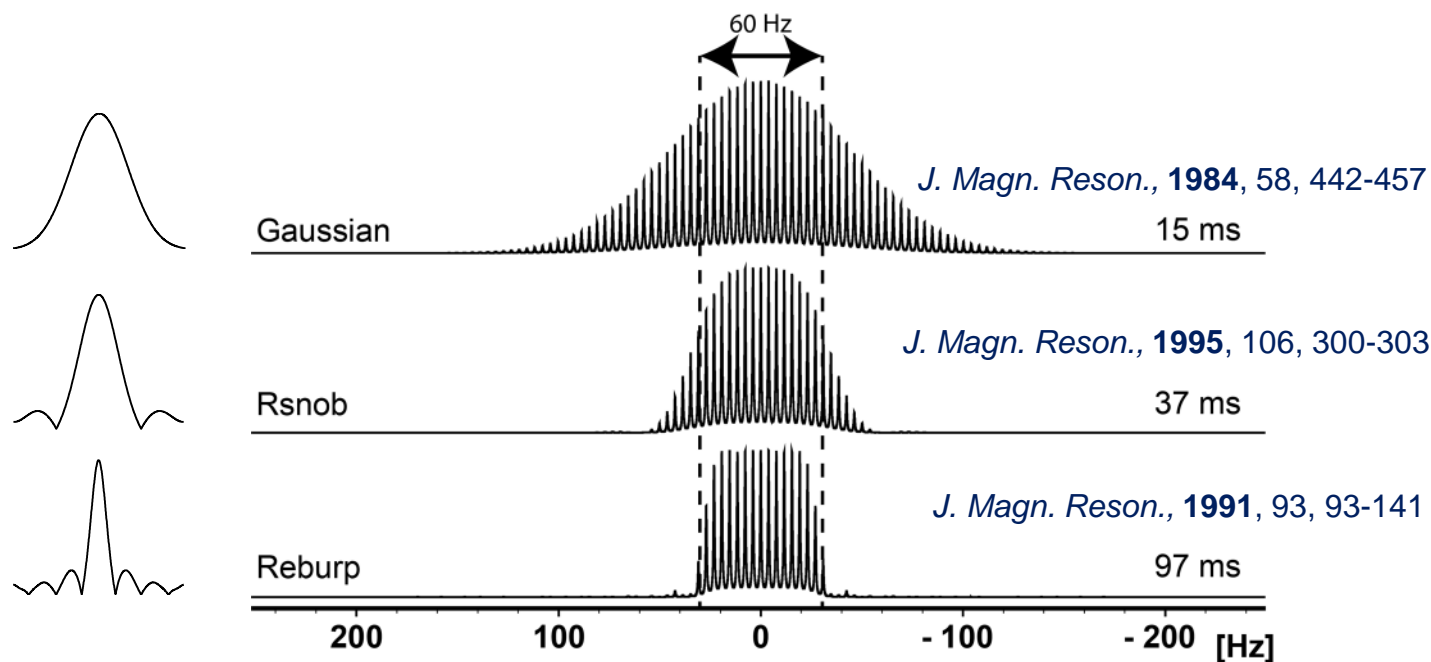
✓ Shorter experiments

✓ Targeted analyses

✓ Simplified spectra

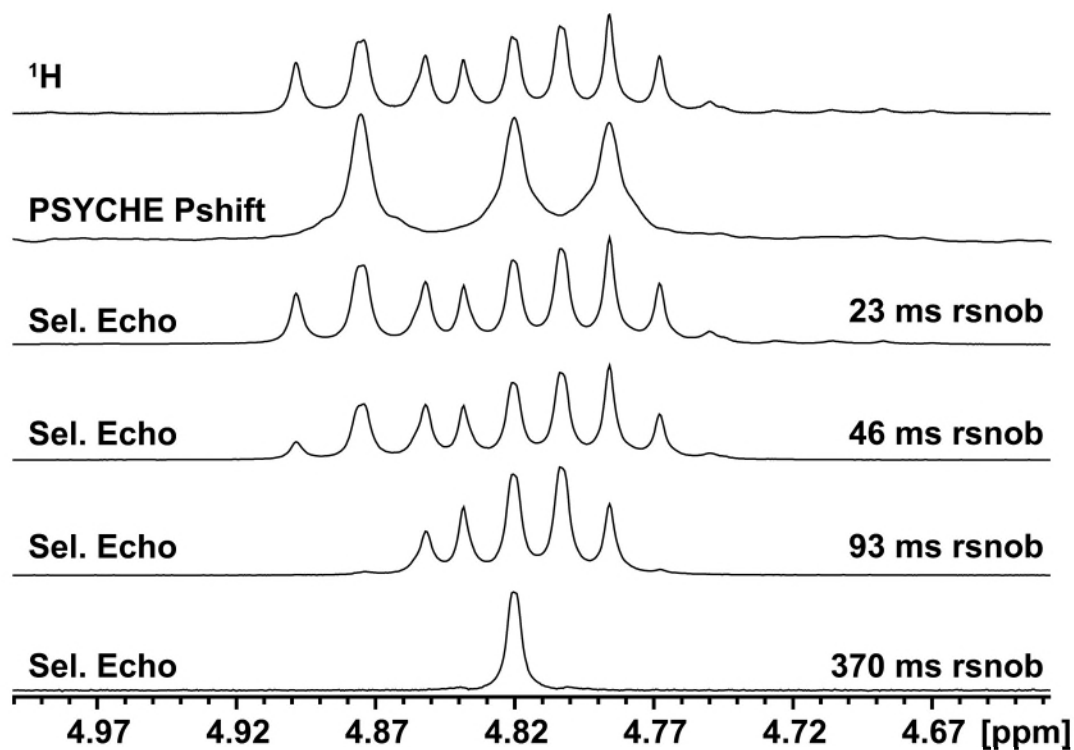
Selective Pulses – Refocusing Profiles

Shapes of pulses determine the refocusing profiles

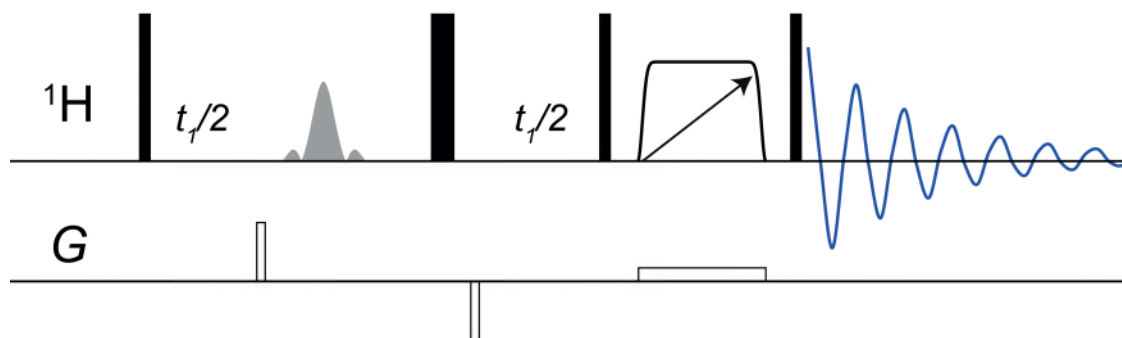


Cannot discriminate between overlapping multiplets !!!

Selective Pulses – Multiplet Overlap



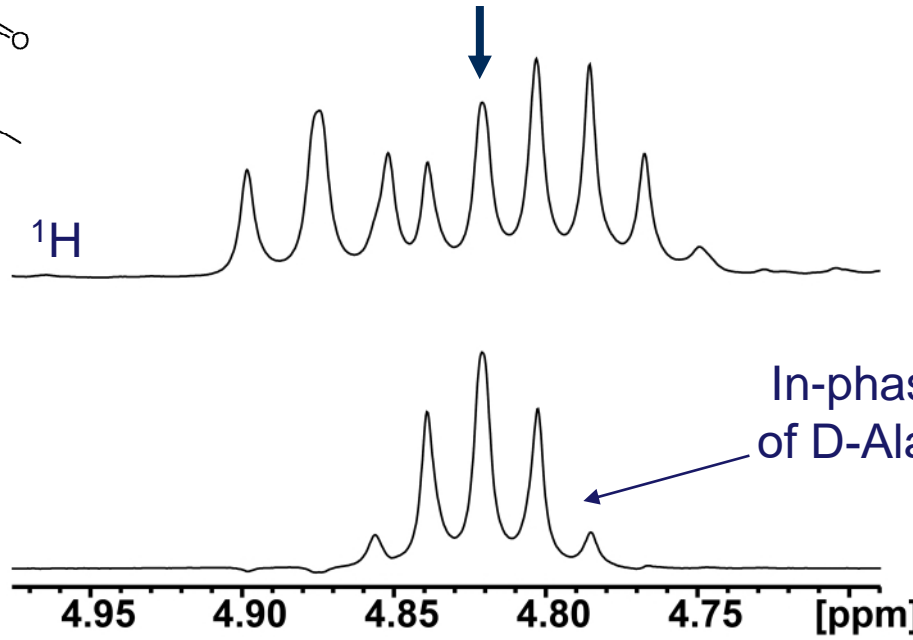
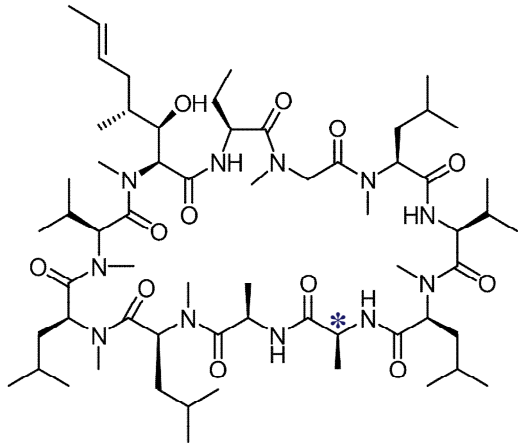
Chemical Shift Selective Filter



Pseudo 2D experiment

- Increment an evolution delay
- Off resonance spins develop a phase as a function of t_1
 - Average to zero by summing FIDs in 2D array

CSSF – Cyclosporin Example



Gradient *E*nhanced *M*ultiplet *S*elective *T*argeted *O*bservation *NMR* *E*xperiments



Uses similar ideas to CSSF

Achieves ultra-selective observation in a single scan

GEMSTONE Family



Communications



How to cite: *Angew. Chem. Int. Ed.* **2021**, *60*, 666–669
International Edition: doi.org/10.1002/anie.202011642
German Edition: doi.org/10.1002/ange.202011642

NMR Spectroscopy **Hot Paper**

Single-Scan Selective Excitation of Individual NMR Signals in Overlapping Multiplets

Peter Kiraly,* Nicolas Kern, Mateusz P. Plesniak, Mathias Nilsson, David J. Procter, Gareth A. Morris, and Ralph W. Adams*

ChemComm



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Single-scan ultra-selective 1D total correlation spectroscopy†

Cite this: *Chem. Commun.*, **2021**, *57*, 2568
Received 10th December 2020.
Accepted 26th January 2021

Peter Kiraly, ‡ Mathias Nilsson, Gareth A. Morris and Ralph W. Adams *

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Check for updates

Ultra-selective, ultra-clean 1D rotating-frame Overhauser effect spectroscopy†

Emma L. Gates, ^a Marshall J. Smith, ^b Jonathan P. Bradley, ^b Myron Johnson, ^b Göran Widmalm, ^c Mathias Nilsson, ^a Gareth A. Morris, ^a Ralph W. Adams *^a and Laura Castañar *^{ad}

Cite this: *Chem. Commun.*, **2023**, *59*, 5854
Received 6th February 2023.
Accepted 29th March 2023

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Check for updates

Ultra-selective 1D clean in-phase correlation spectroscopy†

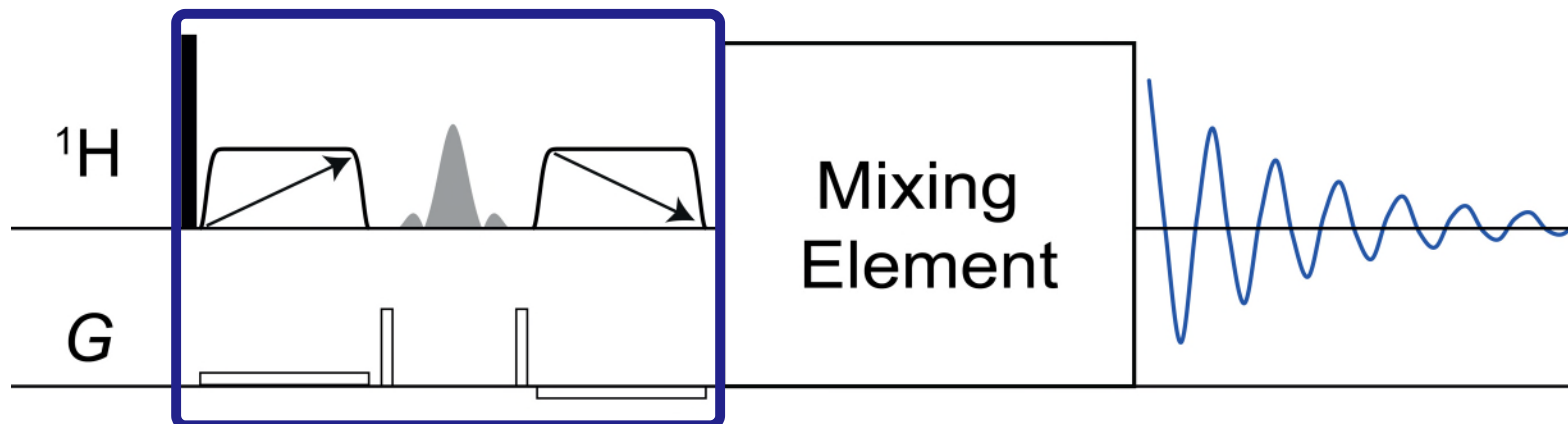
Cite this: *Chem. Commun.*, **2023**, *59*, 6734
Received 17th March 2023.
Accepted 19th April 2023

Daniel A. Taylor, ^a Peter Kiraly, ^b Paul Bowyer, ^c Mathias Nilsson, ^a Laura Castañar, ^{ad} Gareth A. Morris ^a and Ralph W. Adams *^a

More to come...

GEMSTONE Experiments

Initial selective element replaced with GEMSTONE element



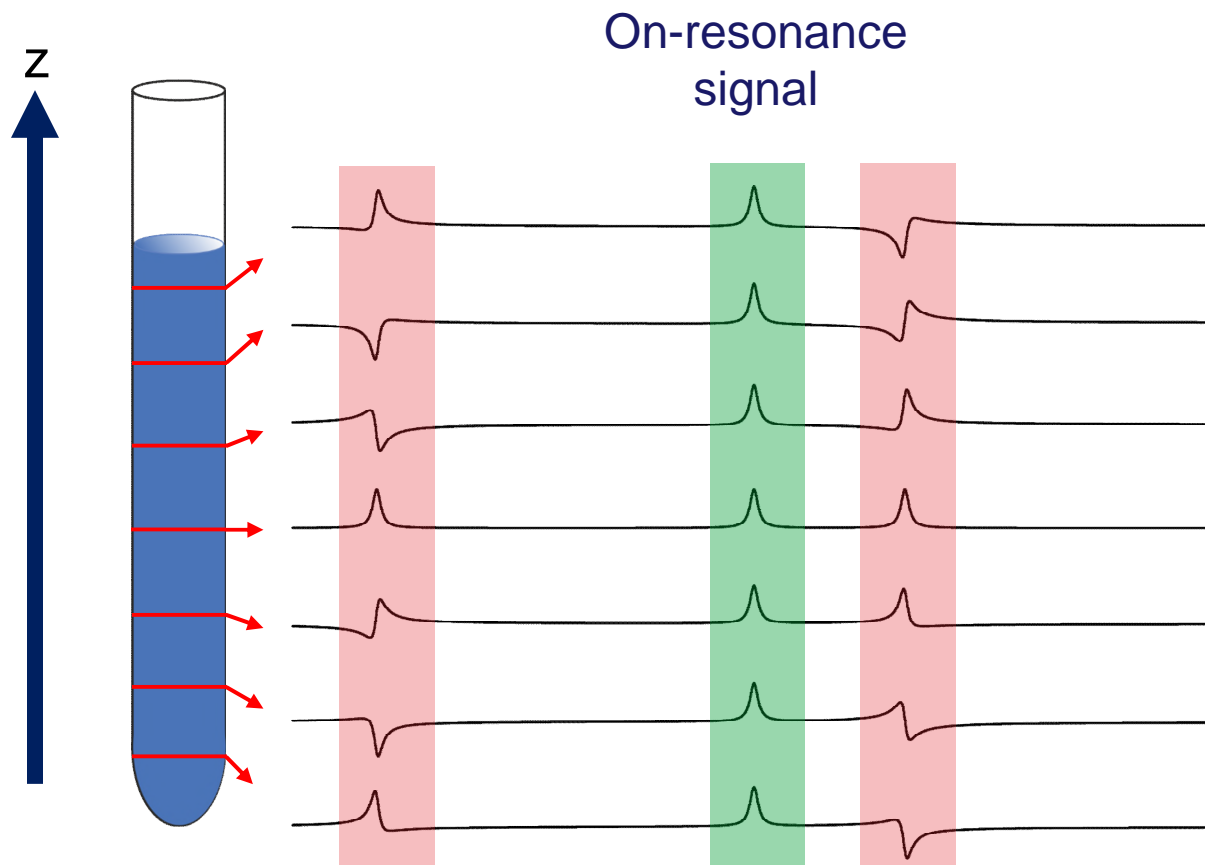
Want to observe individual multiplets

Want to observe in phase multiplets with absorption mode line shapes

Want J_{HH} evolution to be refocused

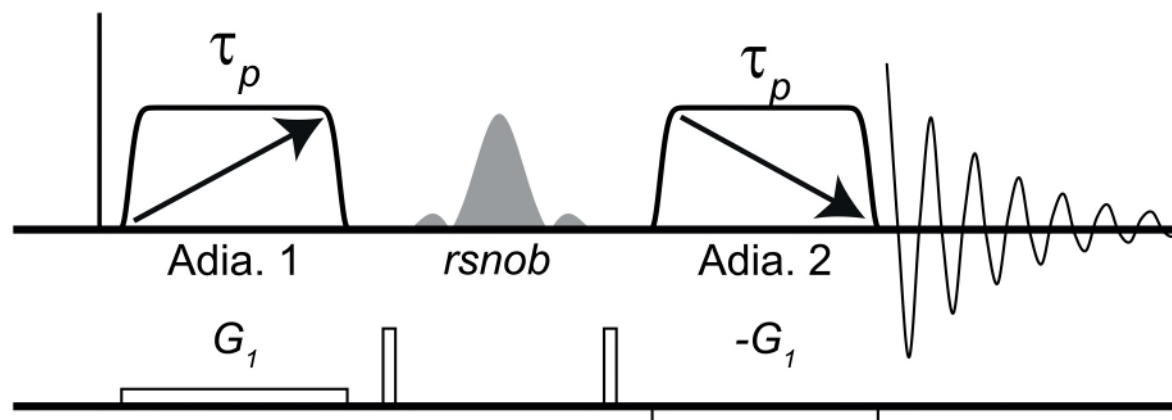


Aim Of The Game



- **On-resonance** signal retains the same phase throughout the NMR tube
 - **Off-resonance** signals acquire a spatially-dependent phase
- ⇒ Off-resonance signals average to zero over the length of the NMR tube

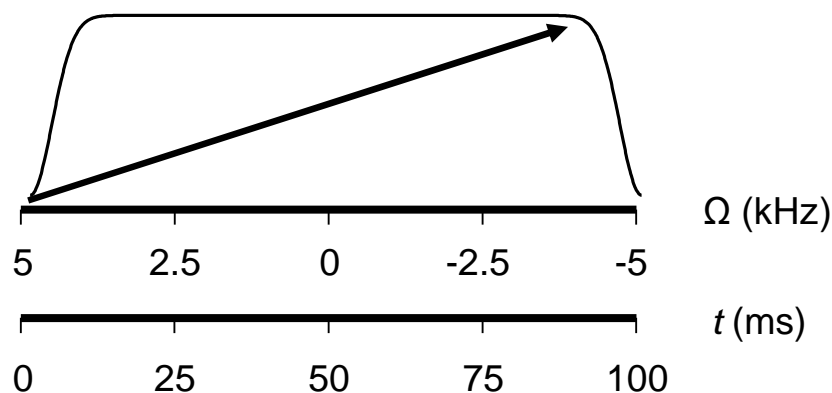
GEMSTONE Pulse Sequence



- Semi-selective $rsnob$
 - Refocuses J_{HH}
 - Suppression of signals far from resonance
 - Inversion of active spins between adiabatic pulses
- Adiabatic pulses and pulsed field gradient
 - Spatial encoding of signals – samples a continuous array of t_1 evolution times

Frequency Swept Pulses

Maps frequency onto time



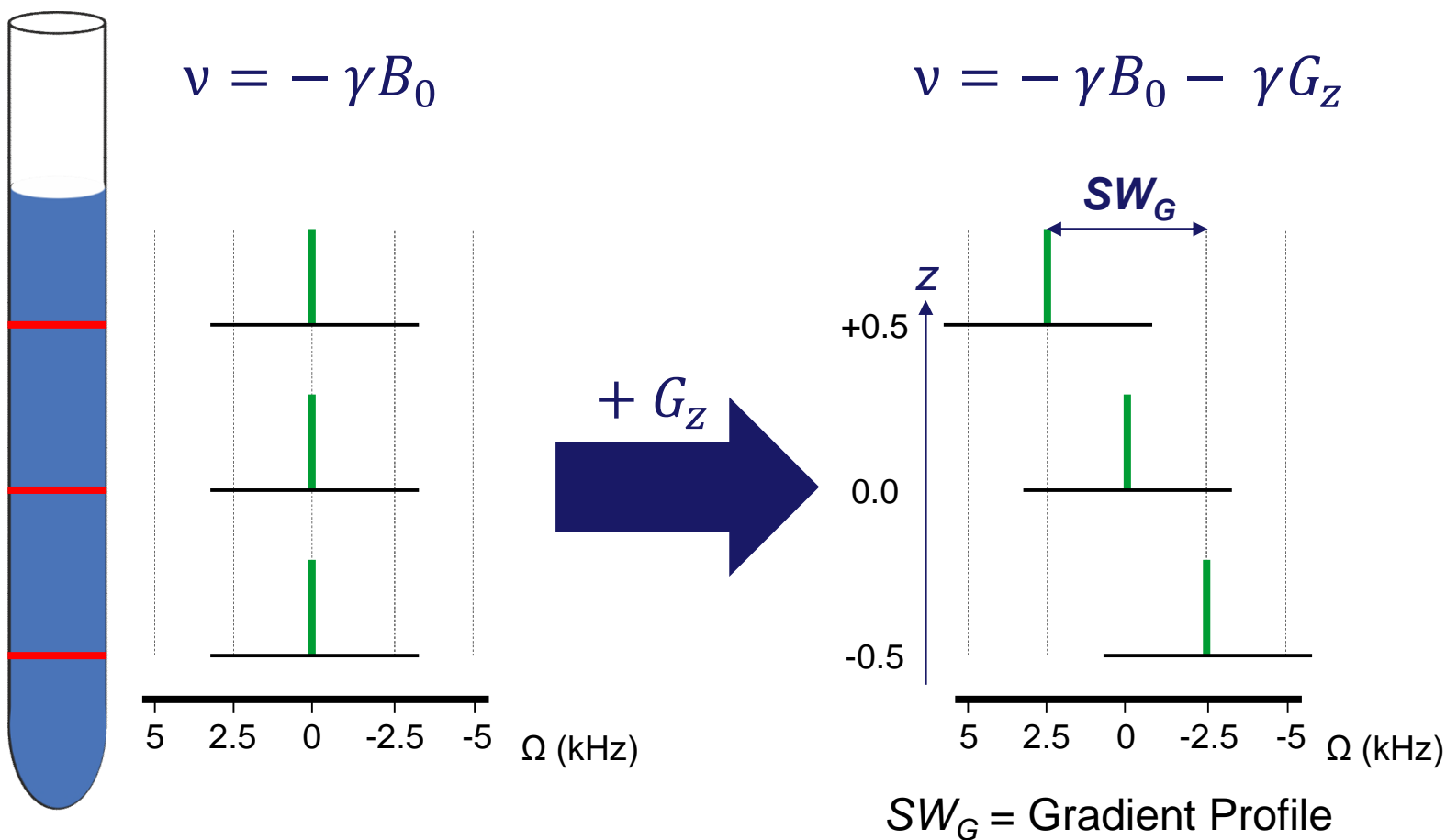
Pulse linearly sweeps from $+BW/2$ to $-BW/2$ for the duration of the pulse

Inversion of a spin depends on its frequency from resonance

Assuming the instantaneous flip approximation

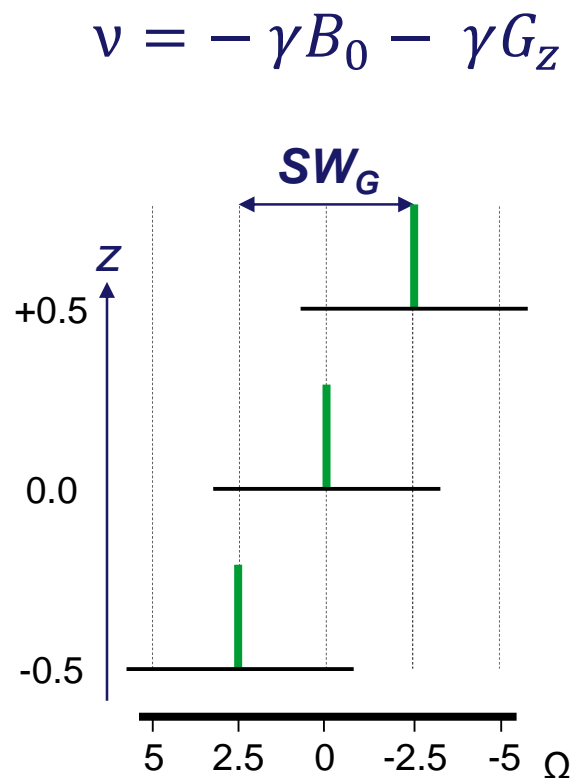
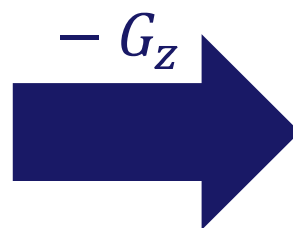
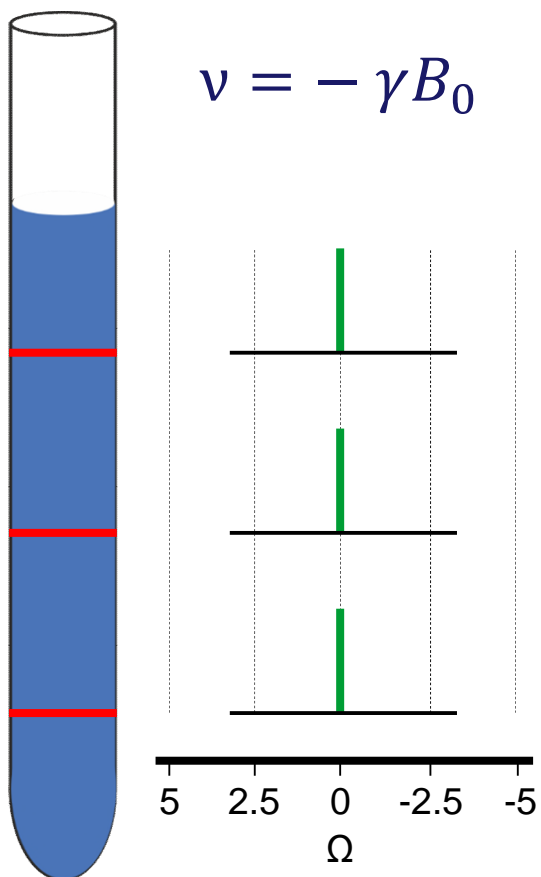
Gradient Profile

Frequency of spins becomes spatially dependent during a gradient



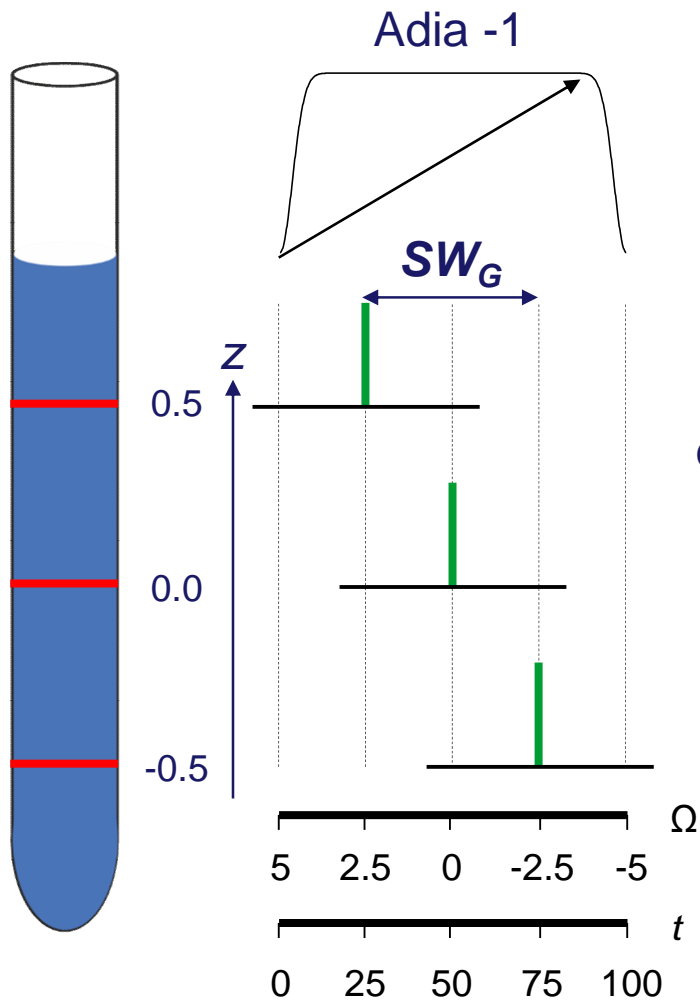
Gradient Profile

Frequency of spins becomes spatially dependent during a gradient



$SW_G = \text{Gradient Profile}$

Spatio-Temporal Averaging – on resonance spins

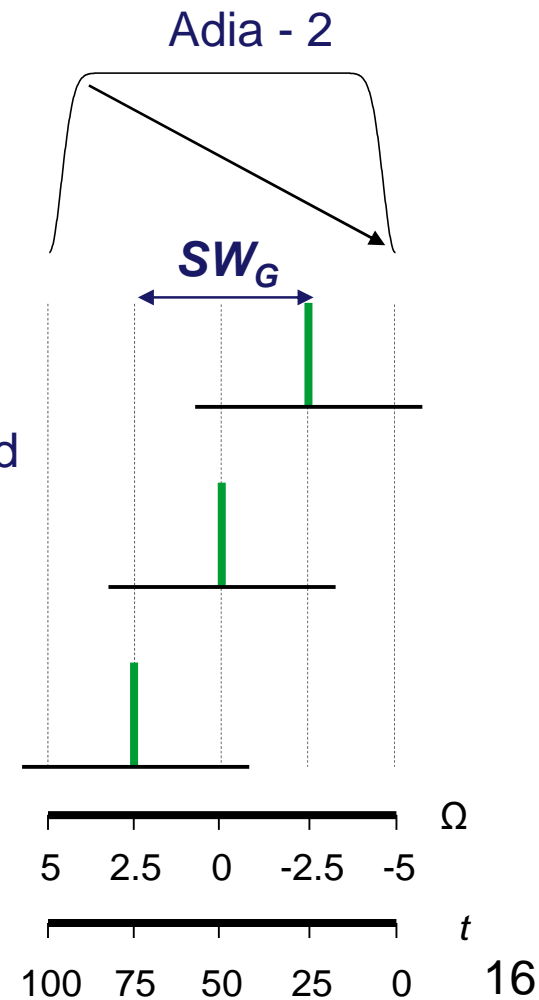


Continuous series of t_1 evolution times are sampled

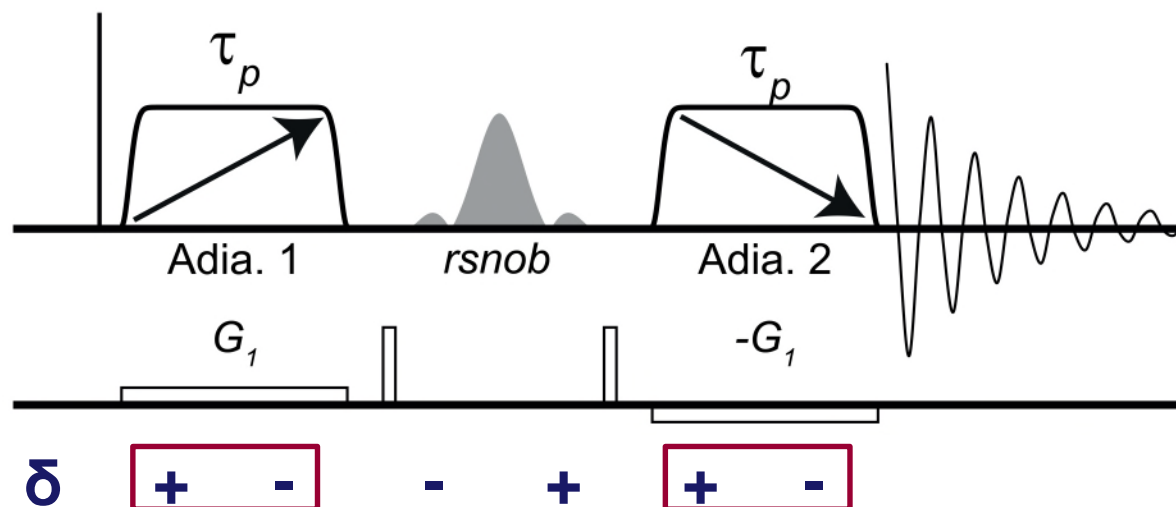
Profiles are centred within pulse duration

Bandwidth: 10 kHz

Duration: 100 ms

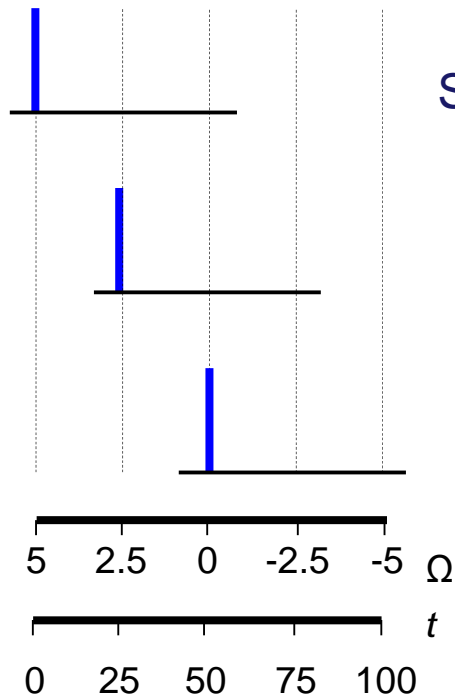


Desired Spin Evolution – Chemical Shifts



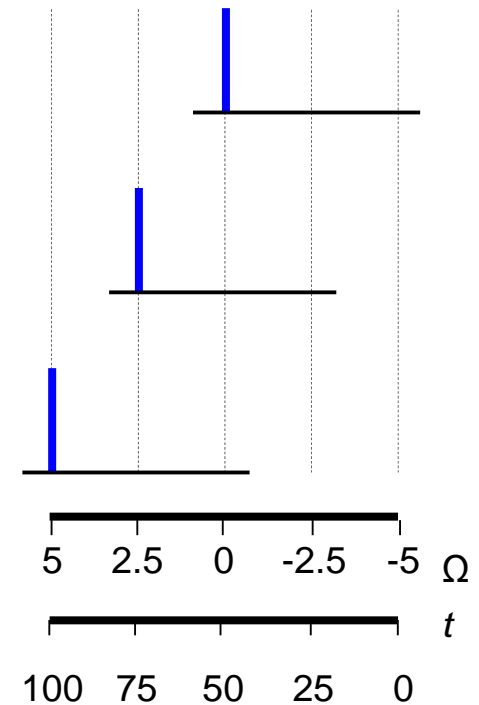
Same sense of chemical shift evolution entering both adiabatic pulses

Spatio temporal averaging – off resonance spins

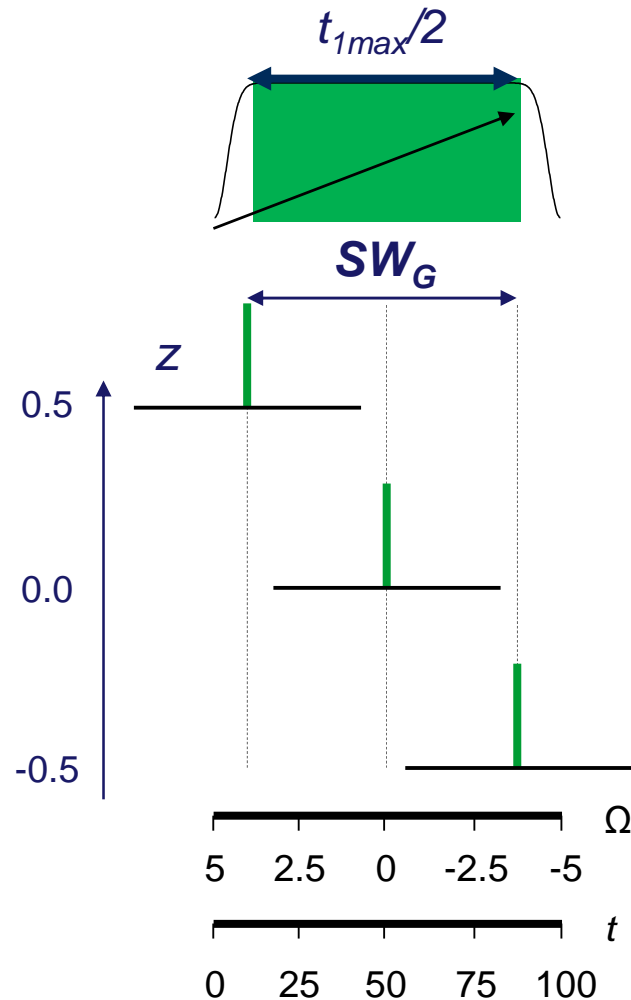


Spin profiles not centred
on pulse duration

Spin evolution is
spatially dependent !!!



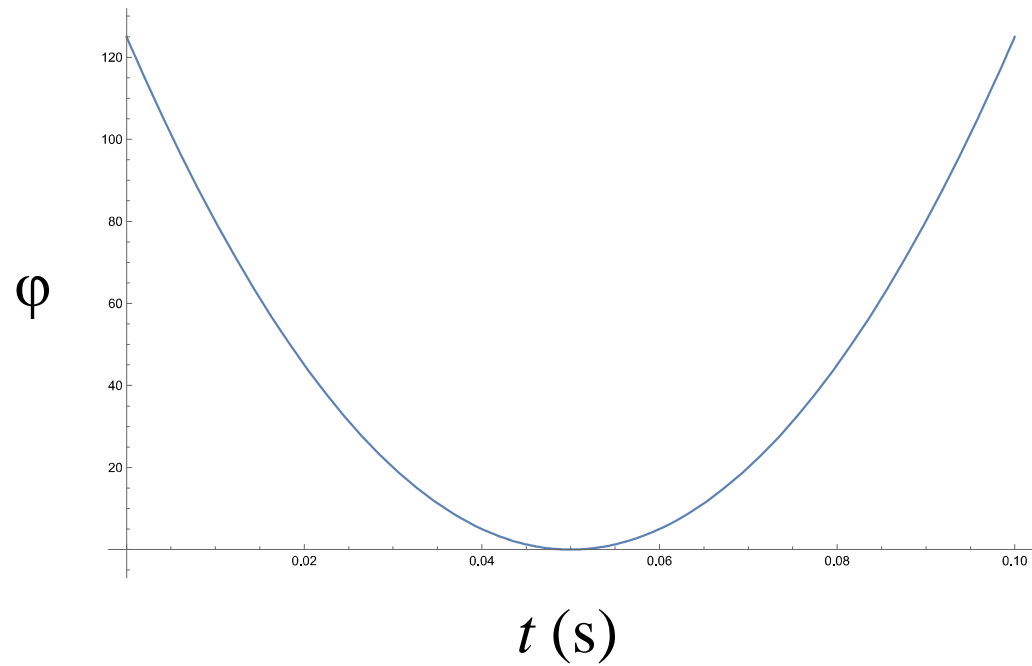
Selectivity determined by t_{1max}



Longer $t_{1max} \rightarrow$ greater selectivity

Matching SW_G to full bandwidth wastes magnetisation – smooth regions of the pulse imperfectly flip spins

Pulse Phase Modulation

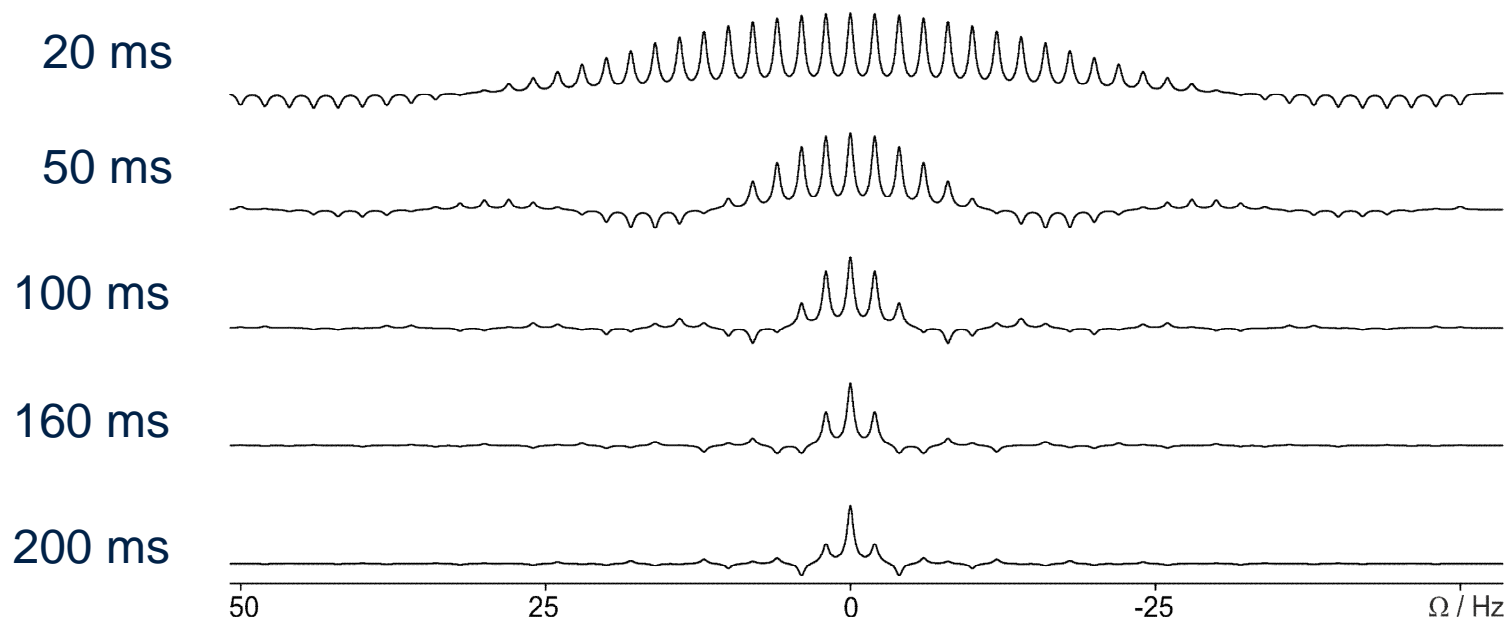


Imparts a time dependent phase shift on spins

Excitation Profile

Excitation profile takes the form of a sinc function

Recommended
pulse durations:
70-150 ms

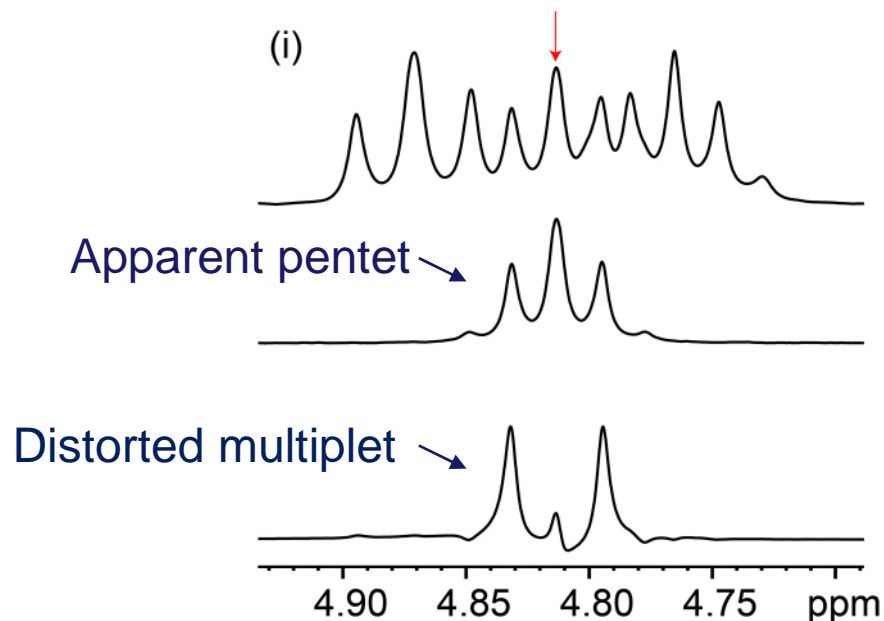


Width of central lobe dependent on t_{1max} (set by adiabatic pulse duration)

Observation of an Individual Multiplet – Cyclosporin Example

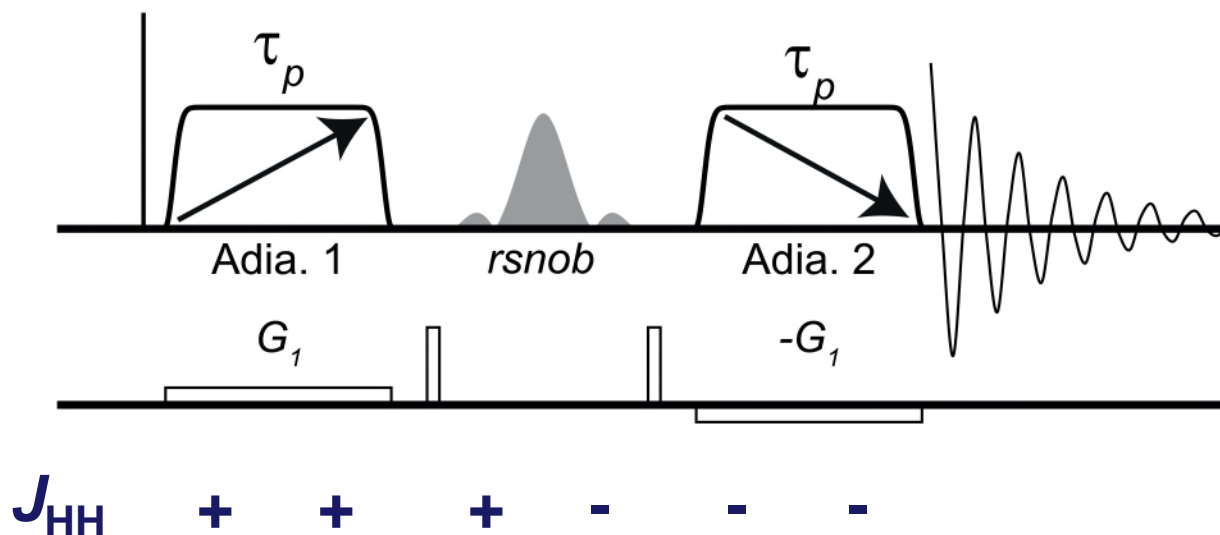
Targeting H_{α} of D-Alanine

V(H_{α}) D-A(H_{α}) L-A(H_{α})



Region from Cyclosporin A spectrum

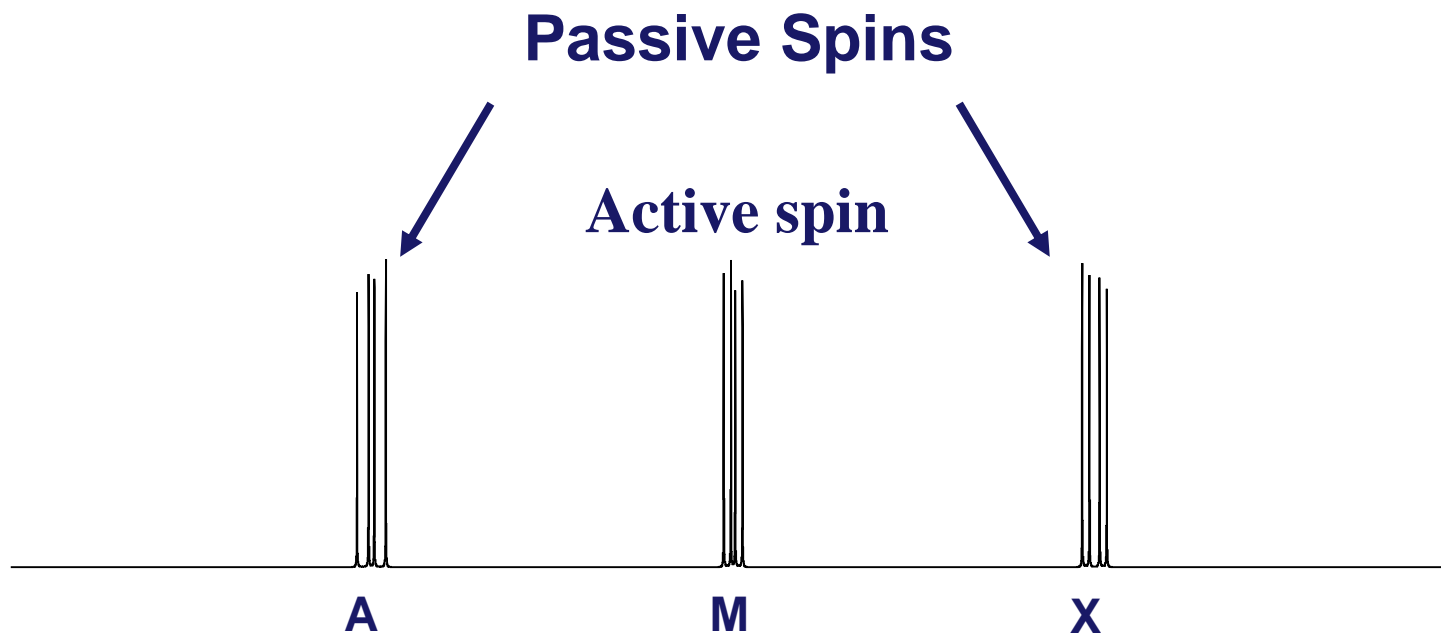
Scalar coupling evolution



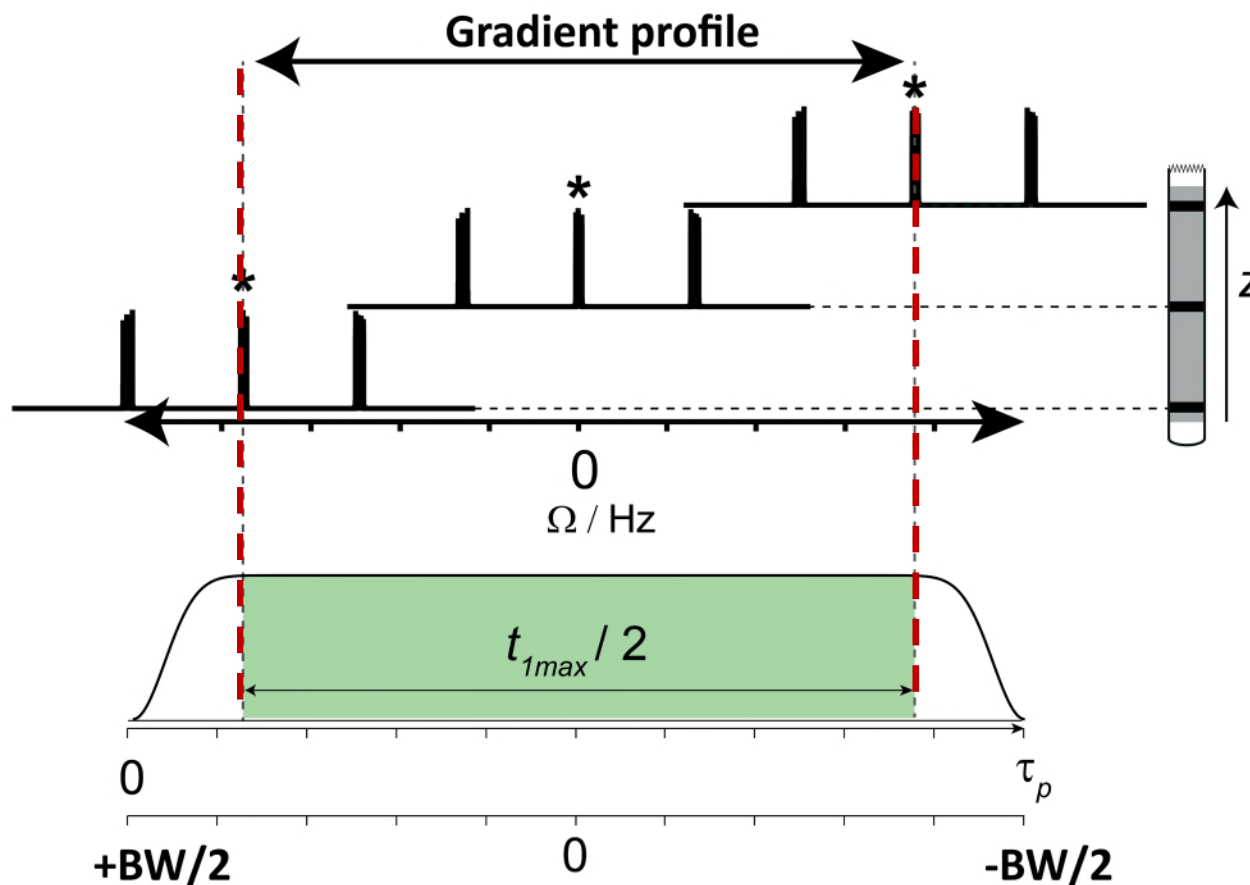
J_{HH} becomes refocused at the beginning of acquisition in perfect scenario

Practically, a compromise needs to be made

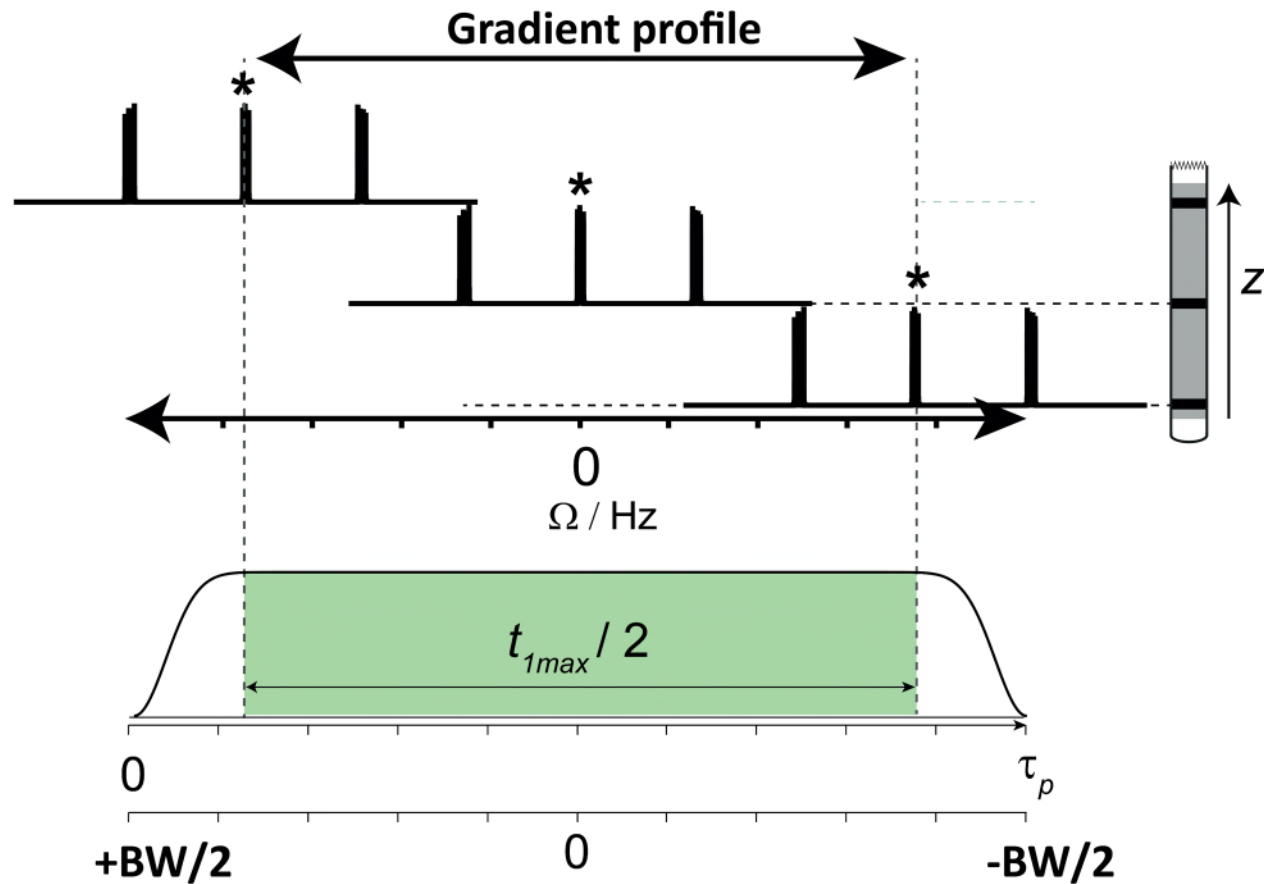
AMX Spin System



Effect of a Gradient – AMX Spin System



Inversion Error – Second Adiabatic pulse

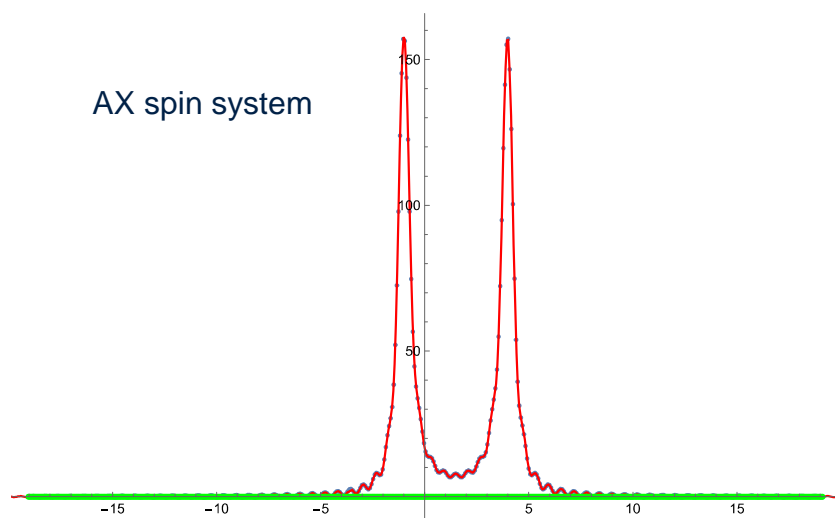


Error will appear in the top and bottom of the sample

Line shape analysis of doublet

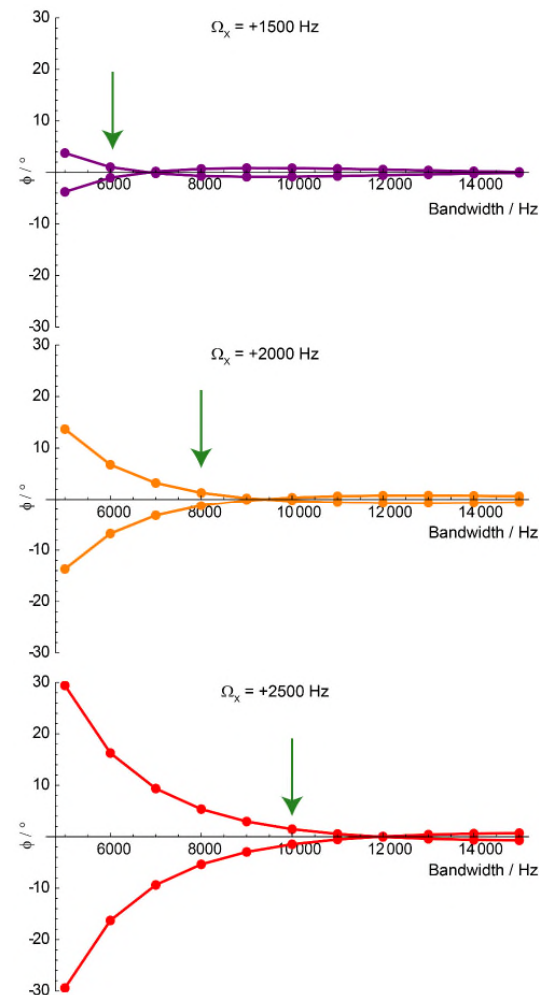
Measure phase of observed multiplet lines as a function of pulse bandwidth

AX spin system



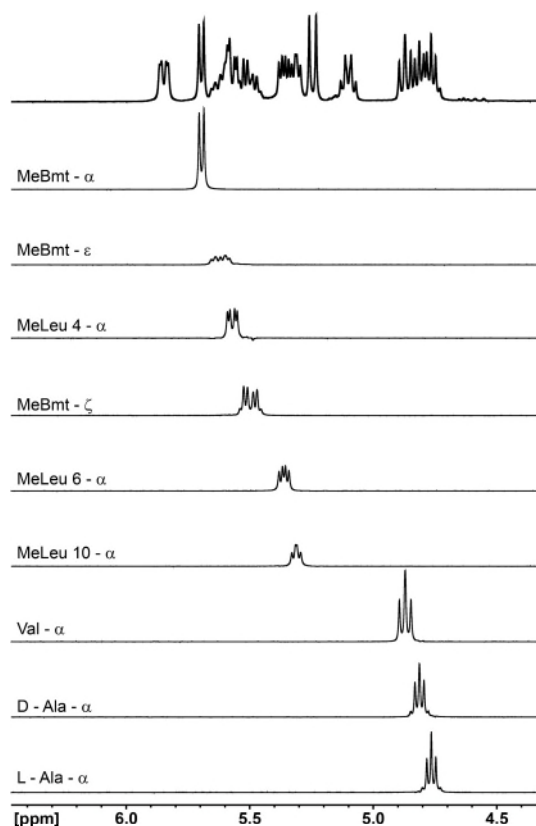
Appropriate SW_G was used for each bandwidth Hz

Phases of doublet components extracted



Cyclosporin A H α Region

Rule of thumb: use pulse with bandwidth 3 times the frequency range in a given spectrum

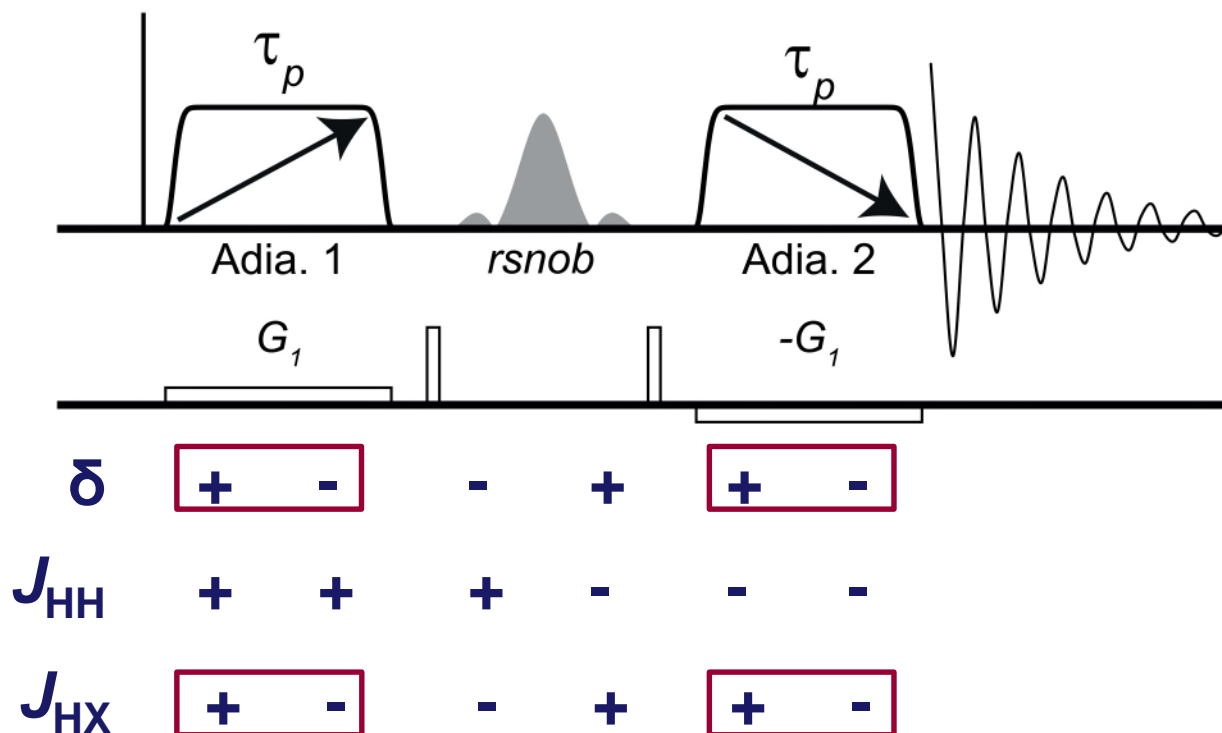


Multiplets are all in phase

Residue	Passive spin Ω_s / Hz
Me-Bmt 1 Hα	H β -600
Me-Bmt 1 Hϵ	H δ -1550
	H ζ -50
Me-Leu 4 Hα	2x H β -1600
Me-Bmt 1 Hζ	H ϵ +50
	H η -1500
Me-Leu 6 Hα	H β -1250
Me-Leu 10 Hα	H β -1250
Val 5 Hα	NH +1050
	H β -900
D-Ala 7 Hα	NH +1100
	H β -1500
L-Ala 8 Hα	NH +1300
	H β -1250

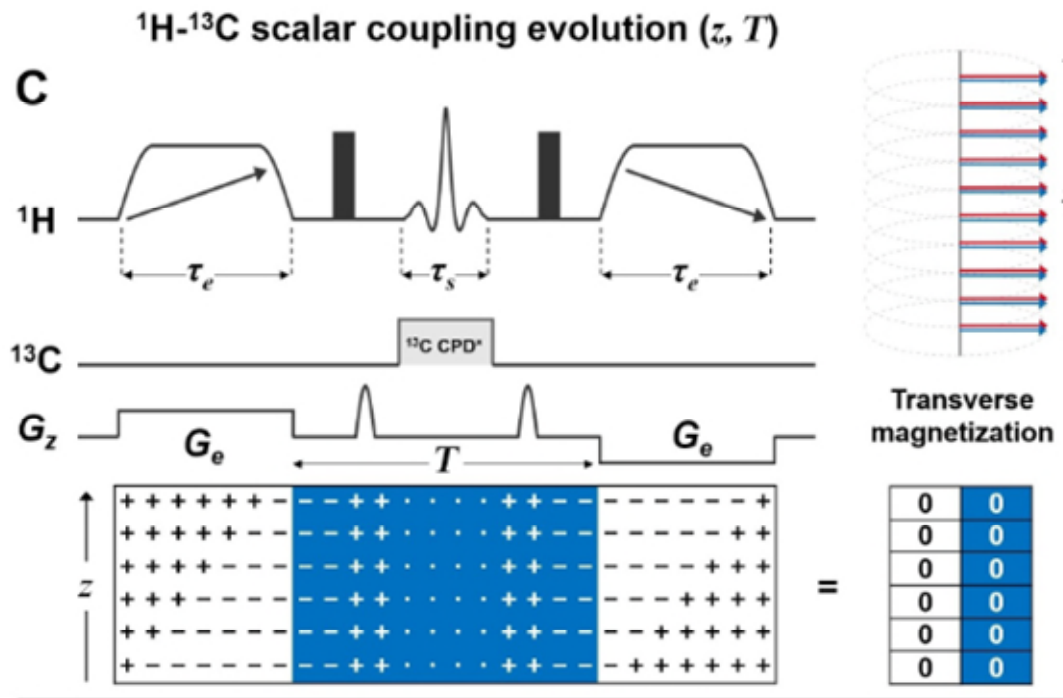
Spectra acquired @ 400 MHz

Note on Heteronuclear Couplings



Heteronuclear couplings behave like chemical shifts

Heteronuclear coupling solution



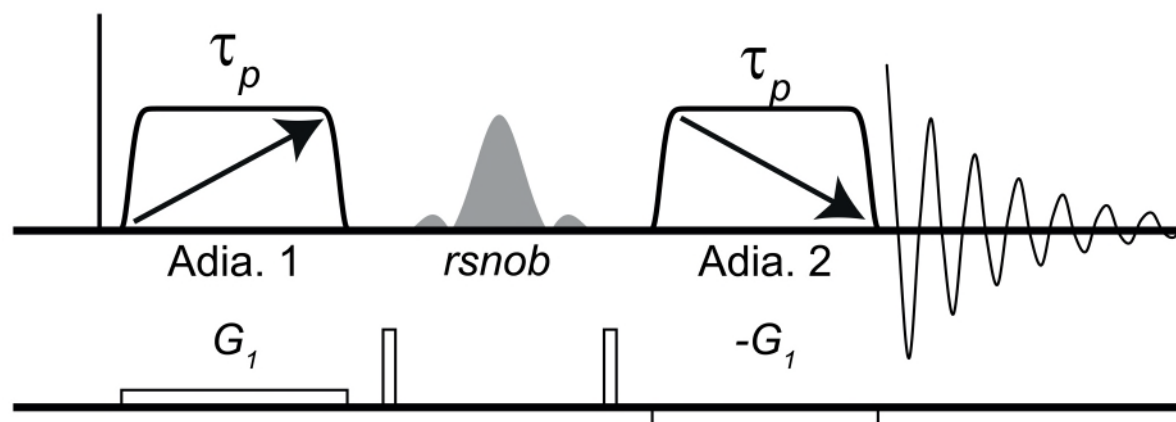
^1H - ^{13}C scalar coupling evolution (z, T)

^{13}C CPD*: Adiabatic decoupling sequence using an odd number of inversion pulses

Spatial position (z)
dependent evolution

Time delay (T)
dependent evolution

A General Parameter Set – ≤ 500 MHz




Rsnob:

- ✓ Must discriminate between active (targeted) and coupled spins
- ✓ Must be wide enough to cover full multiplet width
- ✓ Typically has a 50-100 Hz bandwidth

Frequency swept pulses:

- ✓ High enough bandwidth to minimise imperfect inversion of passive spins
- ✓ Long enough duration so only targeted chemical shift appears in central lobe of sinc function
- ✓ Typically 100 ms duration and 10 kHz bandwidth

Pulse sequences



Pulse Sequences | Manchester

https://www.nmr.chemistry.manchester.ac.uk/?q=node/255

- Relaxation sequences**
 - PROJECT
 - REST
 - PUREST
 - SCALPEL
- Ultra-selective sequences**
 - GEMSTONE
 - GEMSTONE-NOESY
 - GEMSTONE-TOCSY
 - GEMSTONE-ROESY
 - GEMSTONE CLIP-COSY
- Ultra-broadband sequences**
 - CHORUS
- 19F related sequences**
 - 1D heteronuclear decoupled
 - FESTA
 - 19F Oneshot45
- 31P related sequences**
 - 1D heteronuclear decoupled
- 13C satellite suppression sequences**
 - DISPEL
 - ODYSSEUS
- Perfect echo sequences**
 - PPROJECT
 - PE-WATERGATE

spectroscopy

- Ultra-selective, ultra-clean 1D rotating-frame Overhauser effect spectroscopy
- Quinone voltammetry for redox-flow battery applications

More...

GEMSTONE Experiments

<https://www.nmr.chemistry.manchester.ac.uk/?q=node/255>

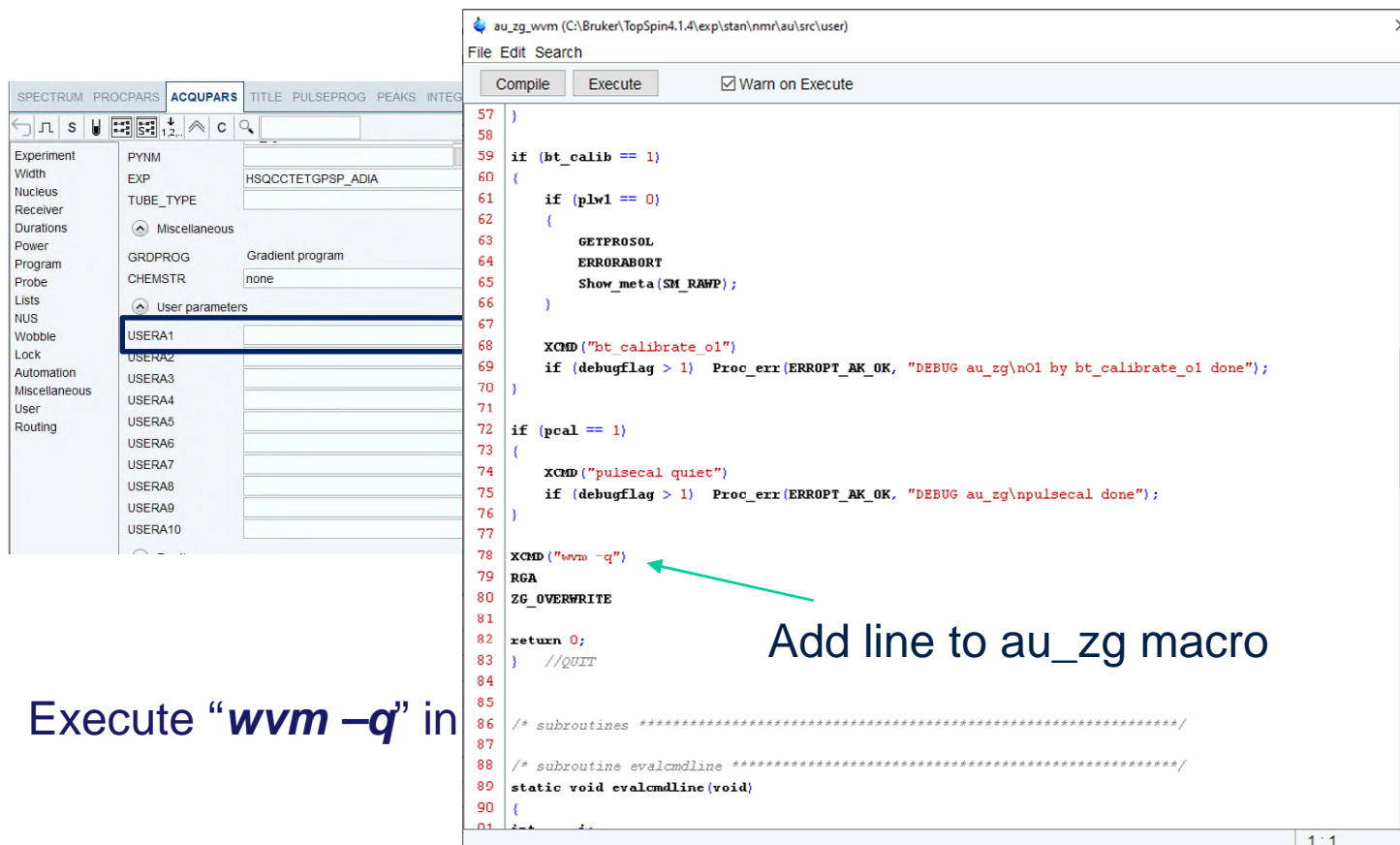
Setting GEMSTONE Pulse Parameters

Easy adjustment of GEMSTONE waveforms using wavemaker

General

PULPROG	GEMSTONE	...	E	Pulse program for acquisition
TD	65536			Time domain size
SWH [Hz, ppm]	7812.50	19.5248		Sweep width in acquisition direction
AQ [sec]	4.1943040			Acquisition time
RG	101			Receiver gain
DW [µsec]	64.000			Dwell time
DE [µsec]	6.50			Pre-scan-delay
CNST50	100.0000000			Band-width of the band-selective RSNOB pulse [Hz]
CNST51	10000.0000000			Sweep-width of the adiabatic pulse [Hz]
CNST52	100.0000000			Duration of the adiabatic pulse [t1max/2: 30-100 ms]
D1 [sec]	2.000000000			Relaxation delay; 1-5 * T1
d11 [sec]	0.0299999993			d11=30m
D16 [sec]	0.001000000			Delay for homospoil/gradient recovery
DS	2			Number of dummy scans
NS	2			Scans to execute
TD0	1			Number of averages in 1D

Setting GEMSTONE Pulse Parameters



The image shows two windows from the Bruker software. The left window is the 'ACQUPARS' parameter editor, with the 'User parameters' section expanded to show fields for USERA1 through USERA10. The right window is a code editor for the 'au_zg_wvm' macro, showing a series of conditional statements and commands. A green arrow points to line 78, which contains the command `XCMD("wvm -q")`. Below the code editor, the text 'Add line to au_zg macro' is written in blue. At the bottom left, the text 'Execute "wvm -q" in' is written in blue.

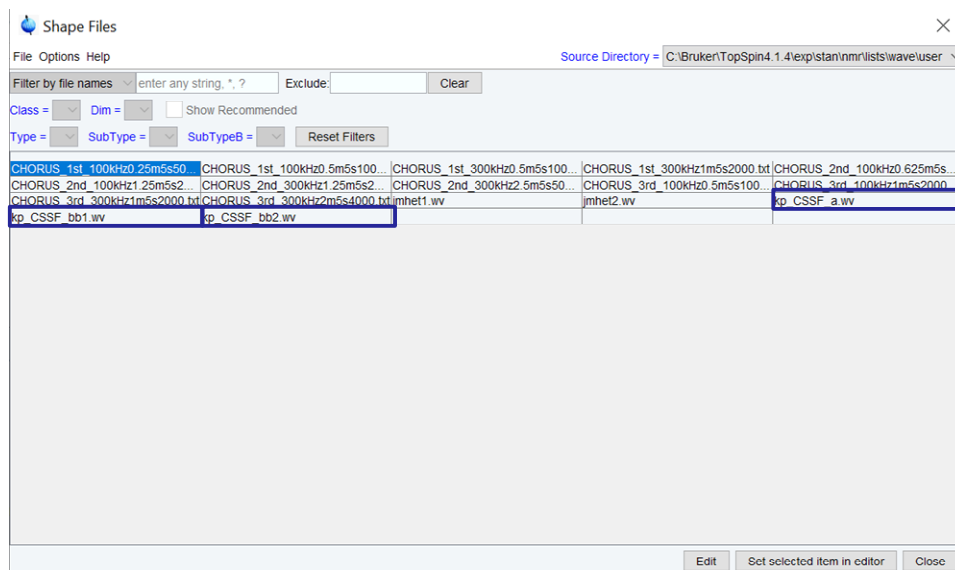
```
57 }
58
59 if (bt_calib == 1)
60 {
61     if (plvl == 0)
62     {
63         GETPROSOL
64         ERRORABORT
65         Show_meta(SM_RAMP);
66     }
67
68     XCMD("bt_calibrate_01")
69     if (debugflag > 1) Proc_err(ERRROPT_AK_OK, "DEBUG au_zg\n01 by bt_calibrate_01 done");
70 }
71
72 if (pcal == 1)
73 {
74     XCMD("pulsecal quiet")
75     if (debugflag > 1) Proc_err(ERRROPT_AK_OK, "DEBUG au_zg\npulsecal done");
76 }
77
78 XCMD("wvm -q")
79 RGA
80 ZG_OVERWRITE
81
82 return 0;
83 } //QUIT
84
85
86 /* subroutines *****/
87
88 /* subroutine evalcmdline *****/
89 static void evalcmdline(void)
90 {
91
```

Execute "wvm -q" in

Add line to au_zg macro

Prevents pop-up wavemaker output window

Waveform File Location



“C:\Bruker\TopSpin4.1.4\exp\stan\nmr\lists\wave\user\”

SPNAM 2	kp_CSSF_a.wv	...	E	File name for SP2
SPOAL2	0.500			Phase alignment of freq. offset in SP2
SPOFFS2 [Hz]	0			Offset frequency for SP2
SPW2 [W]	0.00038076			F1 channel - shaped pulse
SPNAM 41	kp_CSSF_bb1.wv	...	E	File name for SP41
SPOAL41	0.500			Phase alignment of freq. offset in SP41
SPOFFS41 [Hz]	0			Offset frequency for SP41
SPW41 [W]	0.0041617			wvm:kp_CSSF_bb1:f1 wurst-80(cnst51 Hz, cnst52 ms; L2H, Q=11) ss=5.0us;
SPNAM 42	kp_CSSF_bb2.wv	...	E	File name for SP42
SPOAL42	0.500			Phase alignment of freq. offset in SP42
SPOFFS42 [Hz]	0			Offset frequency for SP42
SPW42 [W]	0.0041617			wvm:kp_CSSF_bb2:f1 wurst-80(cnst51 Hz, cnst52 ms; H2L, Q=11) ss=5.0us;

GEMSTONE Gradient Strength

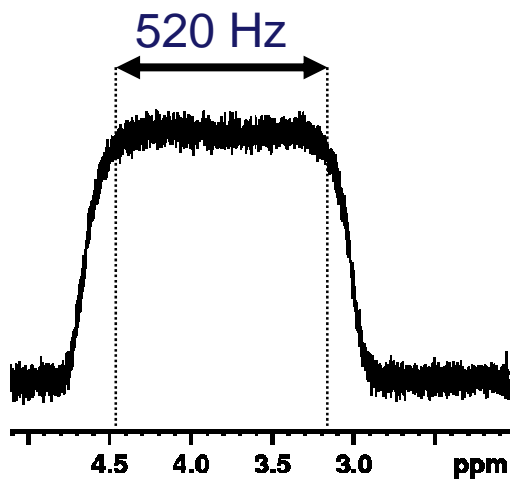
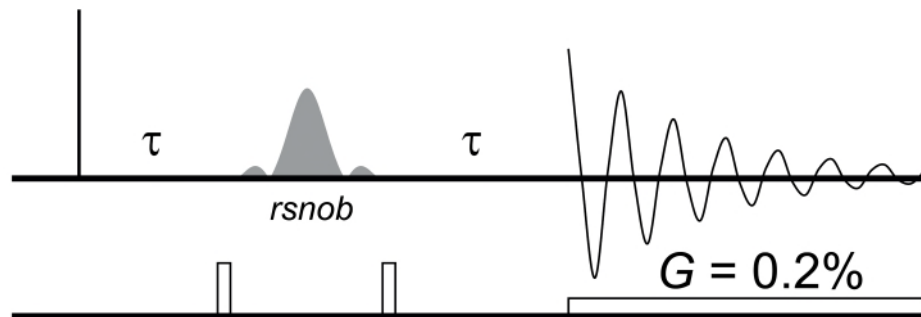
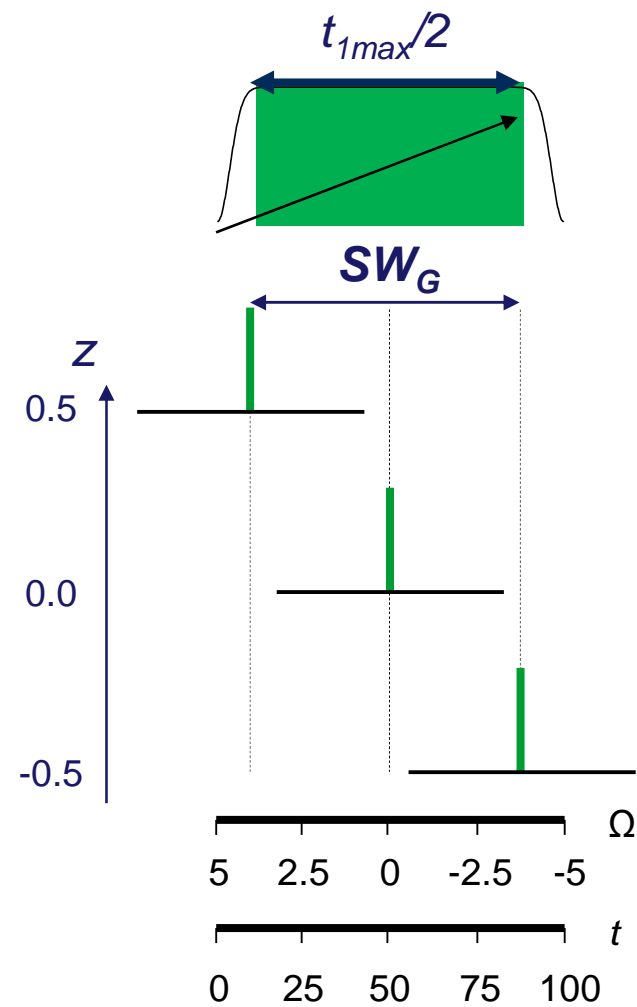
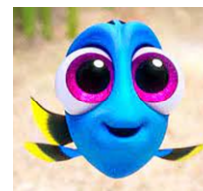


Image of sample



Things to look out for

- Targeted multiplet must be put exactly on resonance
- Strong coupling
- Low viscosity solvents – avoid acetone etc.
- Aliphatic aldehydes
- Other spatio-temporal averaging elements
- Just keep shimmming



Acknowledgments



Peter Kiraly



Ralph Adams



Gareth Morris



Mathias Nilsson



Laura Castañar



Emma Gates



Daniel Taylor



The University of Manchester

Over to you



DISCUSS !!!