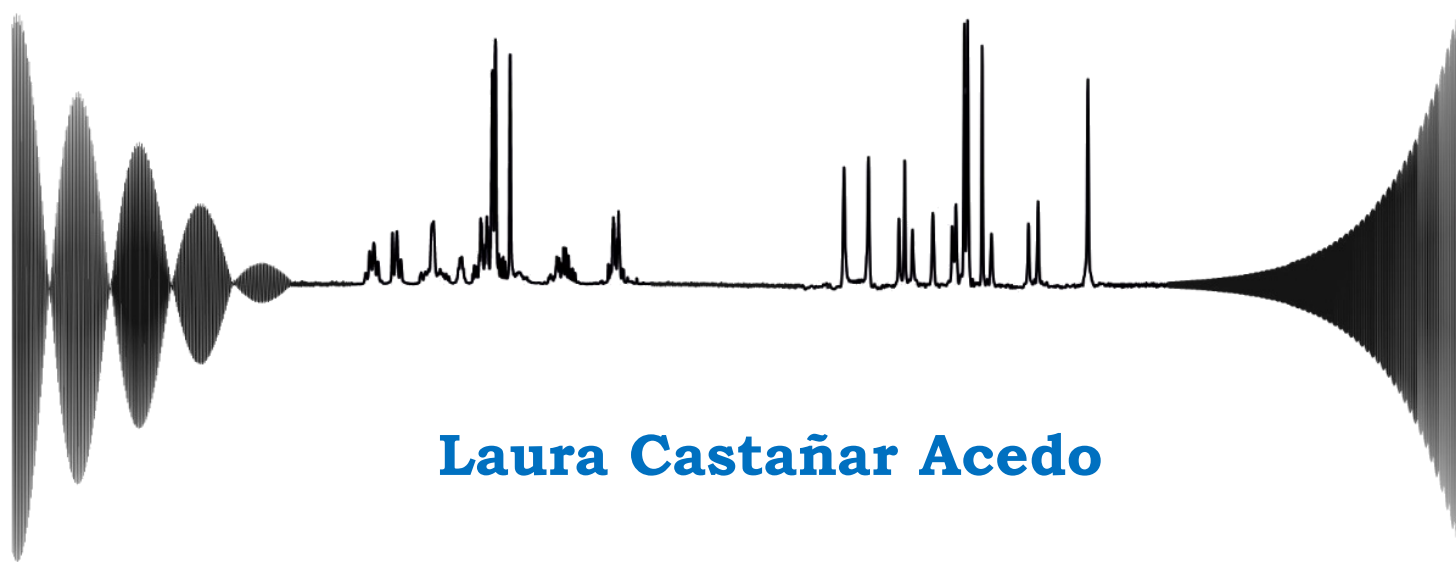


Pure Shift NMR Spectroscopy



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NMR Methodology group
The University of Manchester

Introduction

Setting the scene

Key concepts

J-refocussing

Chunking acquisition

ASR elements

Interferogram

Real-time

Acquisition methods

Band-selective

Zangger-Sterk

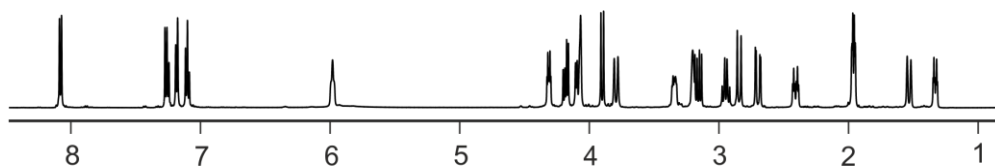
BIRD

PSYCHE

Applications

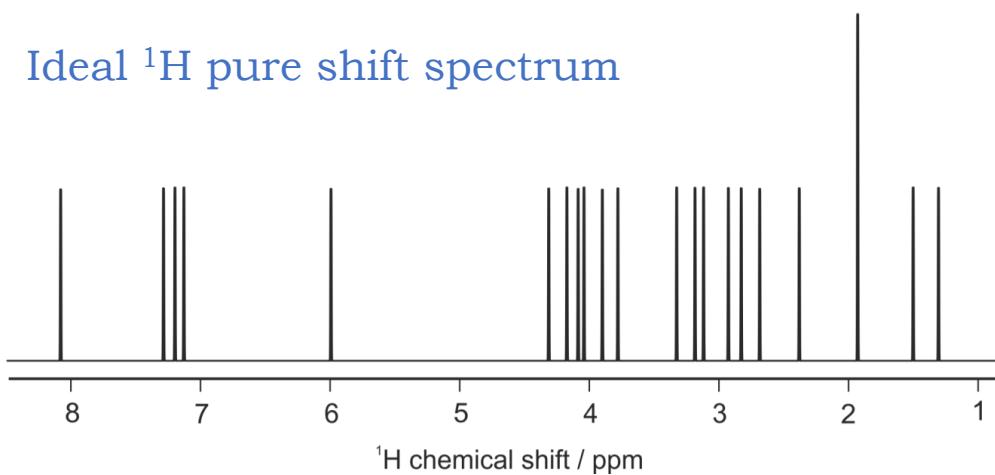
What is pure shift NMR?

Conventional ^1H spectrum



- ✓ Chemical shifts (δ_{H})
- ✓ Homonuclear couplings (J_{HH})
- ✓ Heteronuclear couplings (J_{HX})

Ideal ^1H pure shift spectrum

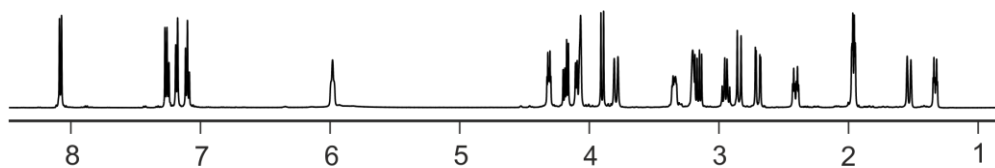


- ✓ Chemical shifts (δ_{H})
- ✗ Homonuclear couplings (J_{HH})
- ✗ Heteronuclear couplings (J_{HX})

A pure shift spectrum is one in which peak positions are determined solely by chemical shifts

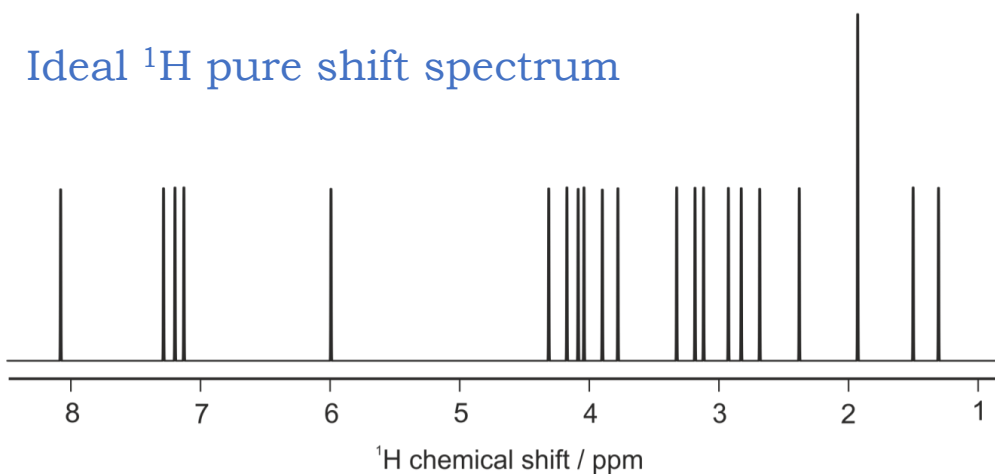
Why is pure shift NMR useful?

Conventional ^1H spectrum



- ☹️ Signal overlap
- ☹️ Poor spectral resolution
- ☹️ Challenging spectral analysis

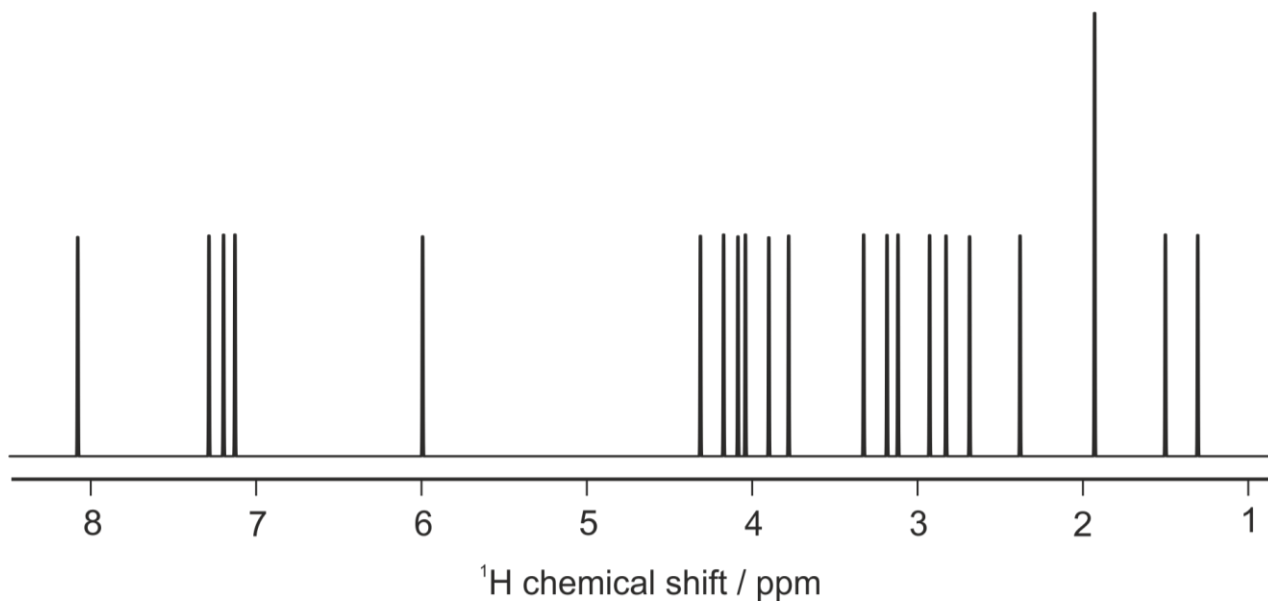
Ideal ^1H pure shift spectrum



- 😊 Reduced spectral complexity
- 😊 Enhanced signal resolution
- 😊 Easier spectral analysis

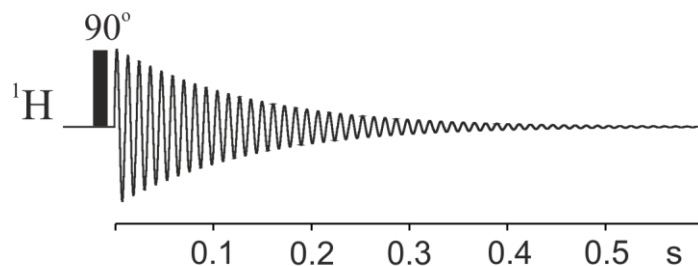
Spectral simplification

How could we get a “perfect” pure shift spectrum?



How do we get a conventional NMR spectrum?

^1H NMR
experiment

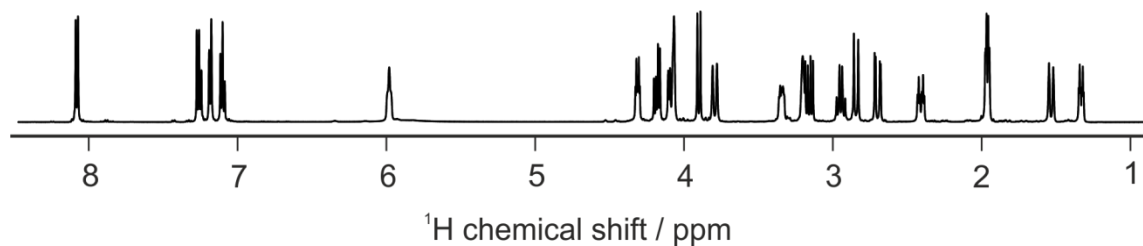


Evolution during FID

δ_{H}	J_{HX}	J_{HH}
✓	✓	✓

^1H NMR
spectrum

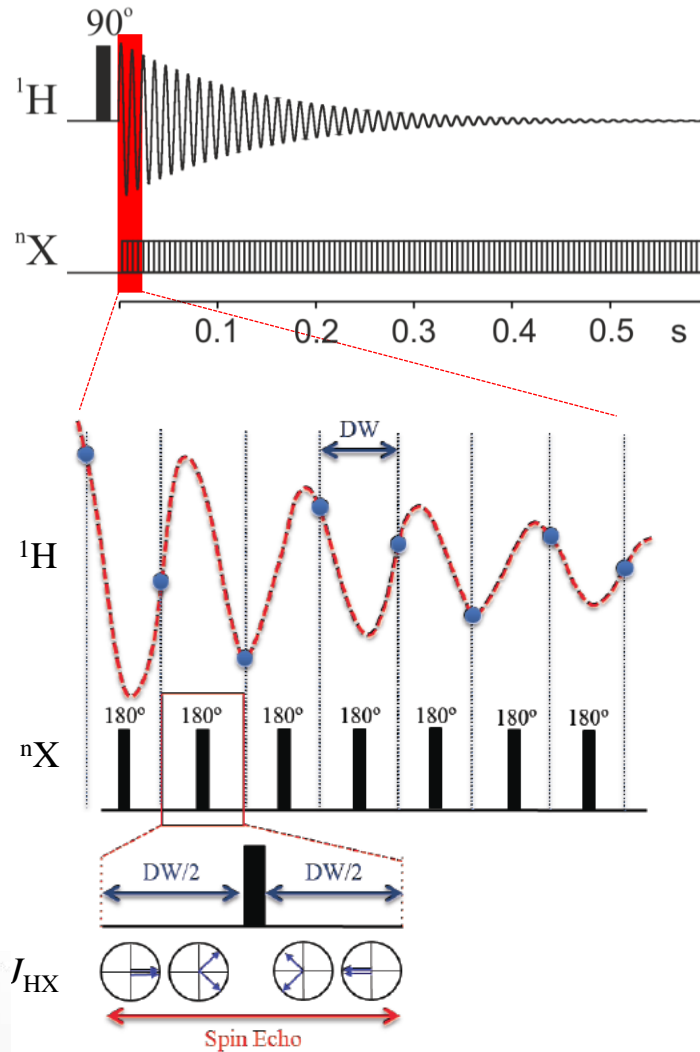
↓ FT



- ✓ Chemical shifts (δ_{H})
- ✓ Homonuclear couplings (J_{HH})
- ✓ Heteronuclear couplings (J_{HX})

How do we get heteronuclear decoupled NMR spectrum?

$^1\text{H}\{^n\text{X}\}$ NMR experiment

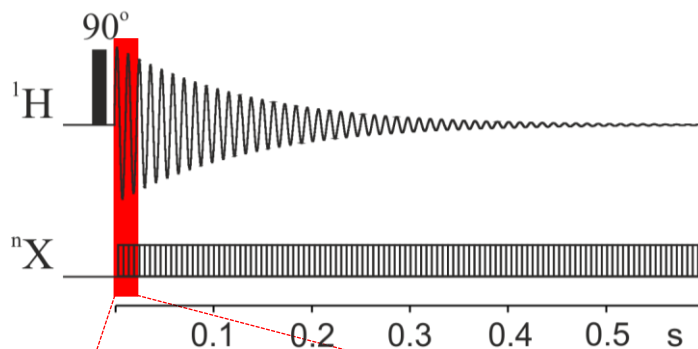


L6
6.9.4



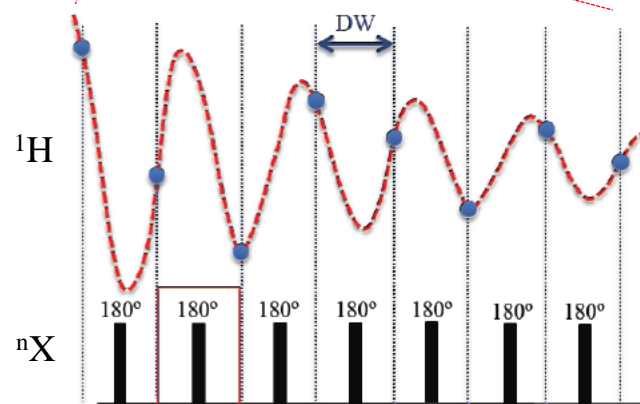
How do we get heteronuclear decoupled NMR spectrum?

$^1\text{H}\{^n\text{X}\}$ NMR experiment



Evolution during FID

δ_{H}	J_{HX}	J_{HH}
✓	✗	✓



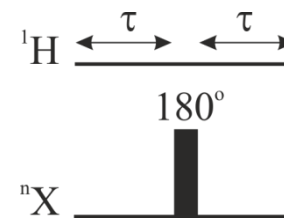
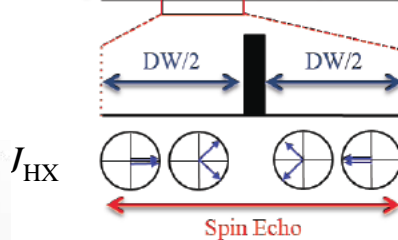
Sampling interval (DW) $\sim 50 \mu\text{s}$

Hard 180° ^nX pulse $\sim 20 \mu\text{s}$

L2
2.4.3

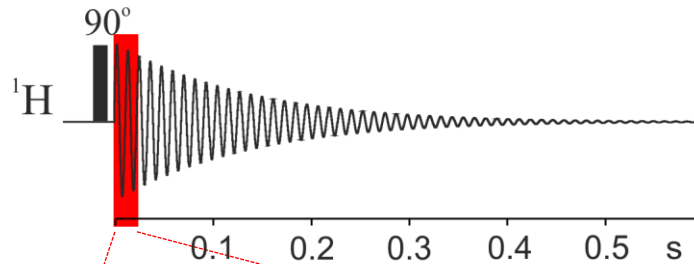
Heteronuclear spin echo

- refocuses J_{HX}
- δ_{H} and J_{HH} evolve



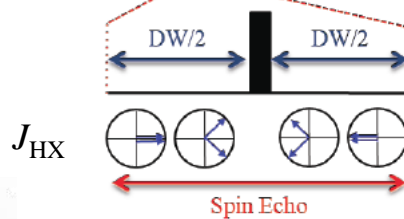
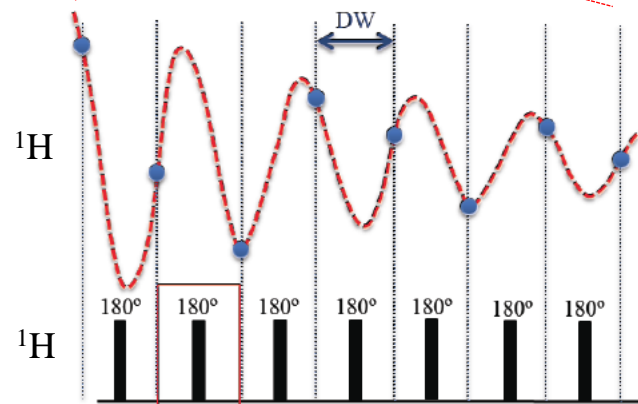
How could we get a “perfect” homodecoupled NMR spectrum?

Ideal
 $^1\text{H}\{^1\text{H}\}$ NMR
 experiment

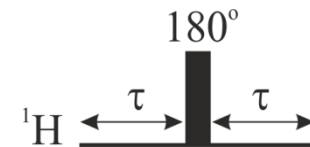


Evolution during FID

δ_{H}	J_{HH}
✓	✗



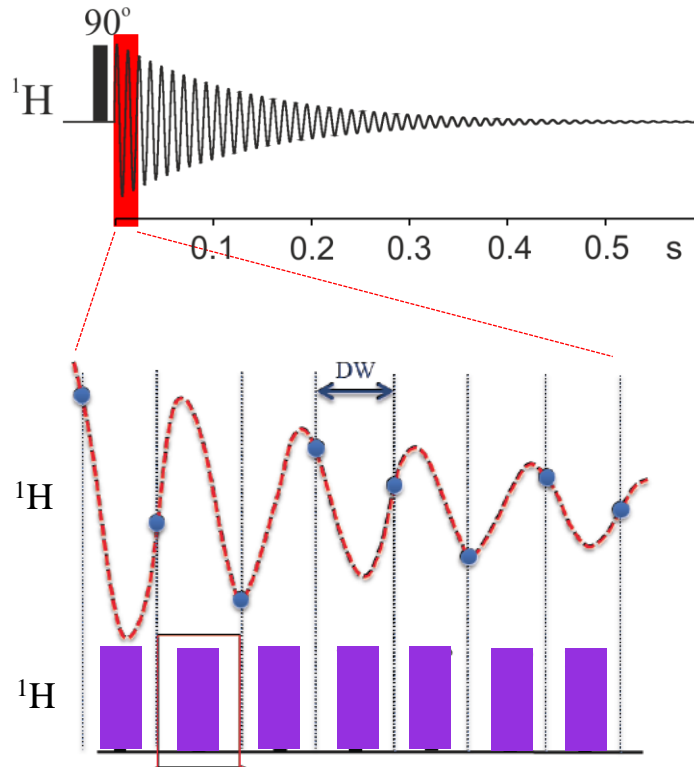
Homonuclear spin echo
 - refocuses δ_{H}
 - J_{HH} evolve



L2
2.4.2

How could we get a “perfect” homodecoupled NMR spectrum?

Ideal
 $^1\text{H}\{^1\text{H}\}$ NMR
 experiment



Evolution during FID

δ_{H}

J_{HH}

✓

✓

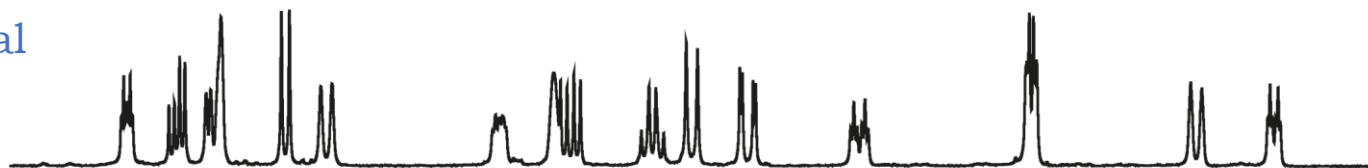
Not
 (currently)
 available

We would need a decoupling element:

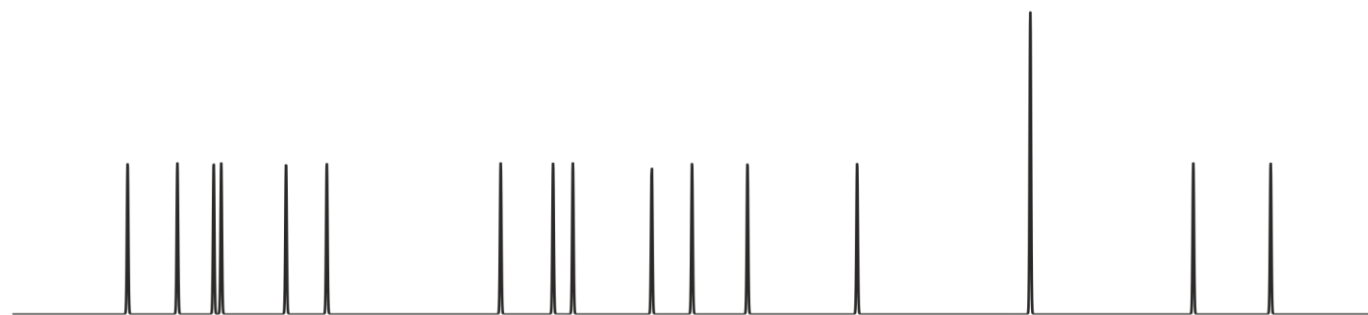
- refocuses J_{HH}
- leaves δ_{H} to evolve
- last few μs

How far are we from a “perfect” pure shift spectrum?

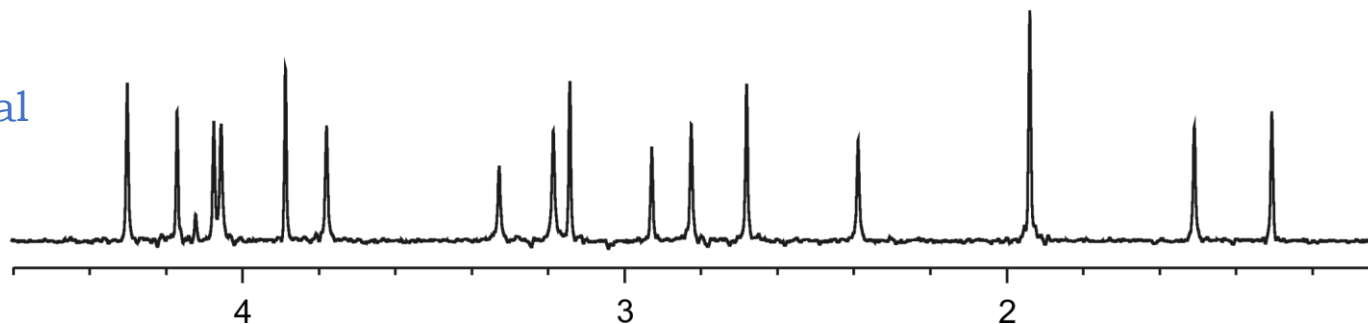
Conventional
 ^1H NMR



Ideal
pure shift



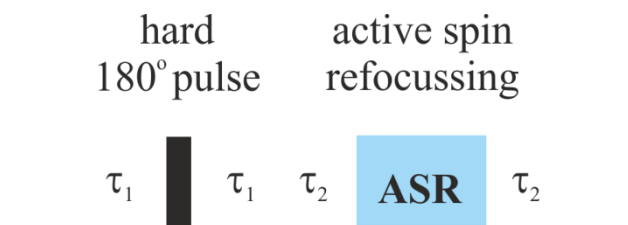
Experimental
pure shift



How do we get an experimental pure shift NMR spectrum?

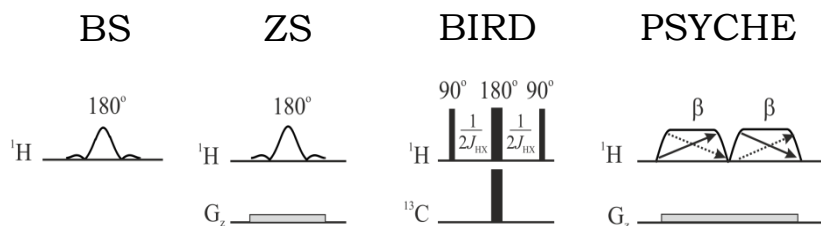
Homodecoupling element

J-refocussing element



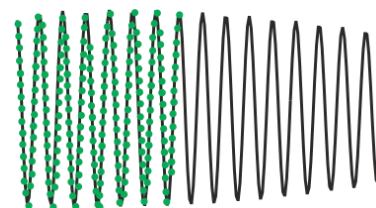
J_{HH} is refocused while δ_{H} evolves
(for *active* spins only)

Active spin refocussing



Homodecoupled FID

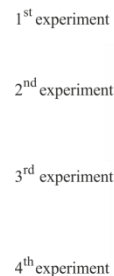
Chunking data acquisition



Chunk duration (τ_c) $\ll 1/J_{\text{HH}}$

Acquisition methods

Interferogram



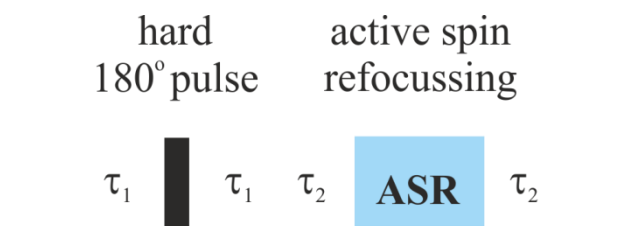
Real-time



How do we get an experimental pure shift NMR spectrum?

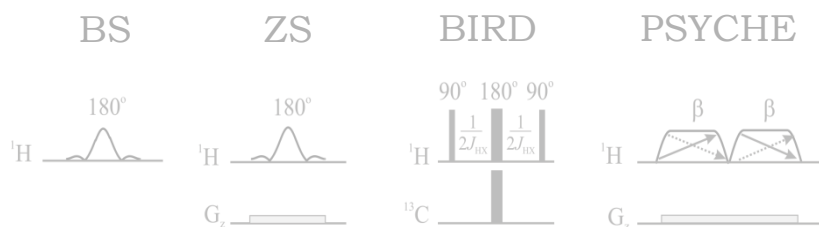
Homodecoupling element

J -refocussing element



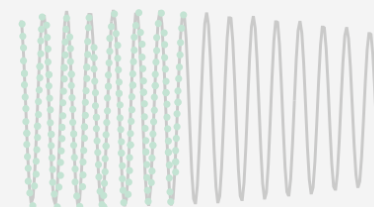
J_{HH} is refocused while δ_{H} evolves
(for *active* spins only)

Active spin refocussing



Homodecoupled FID

Chunking data acquisition



Chunk duration (τ_c) $\ll 1/J_{\text{HH}}$

Acquisition methods

Interferogram

1st experiment

2nd experiment

3rd experiment

4th experiment



Real-time



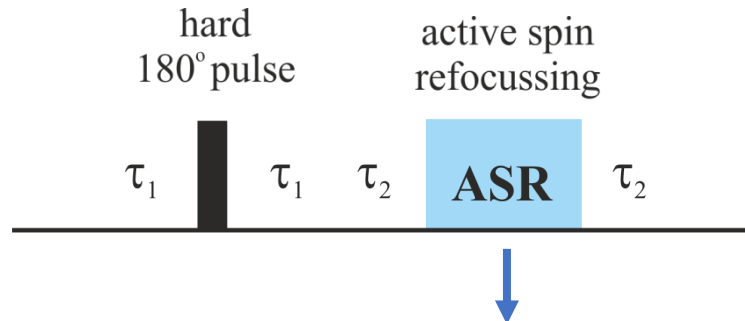
How do we remove the effect of J_{HH} couplings only?



We need a **decoupling element** that:

- refocuses J_{HH}
- leaves δ_H to evolve

J-refocussing element

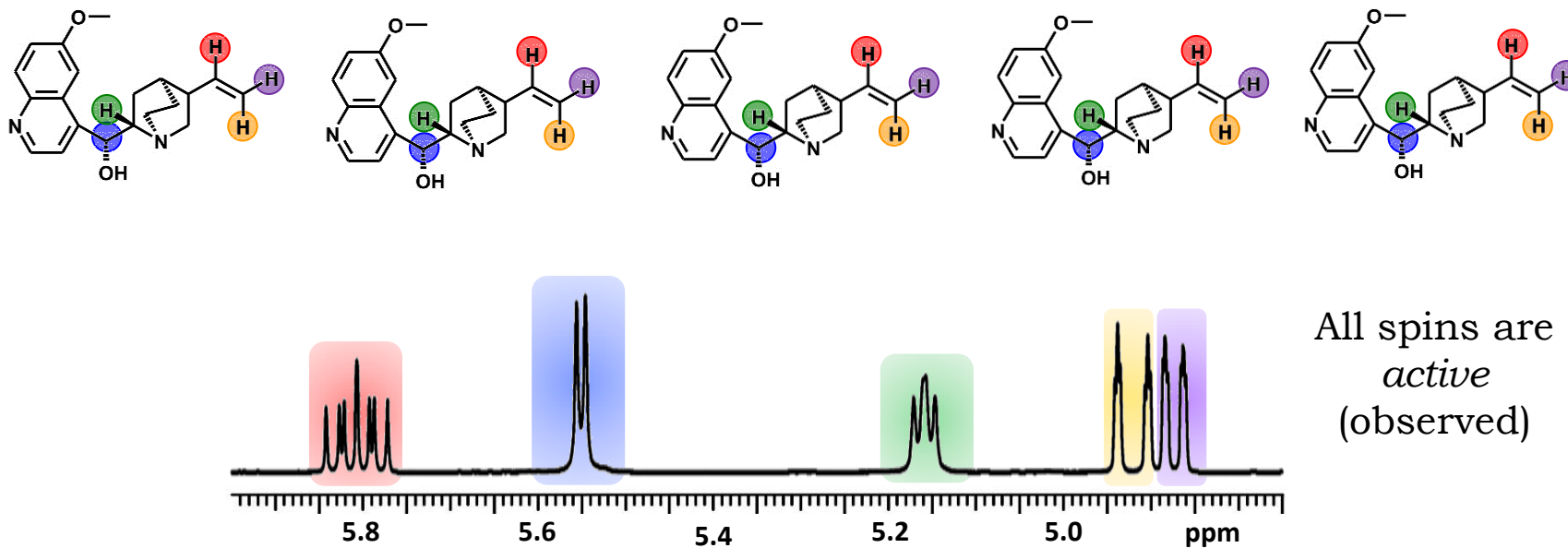


Divides the available spins into:

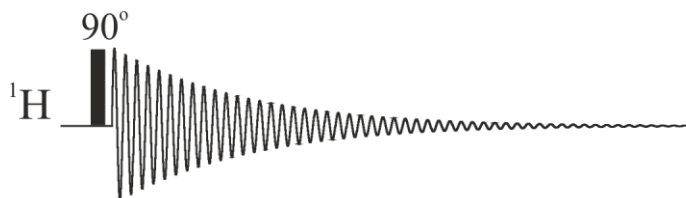
- *active* (observed)
- *passive* spins (non observed)

The concept of *active* and *passive* spins

Conventional ^1H NMR

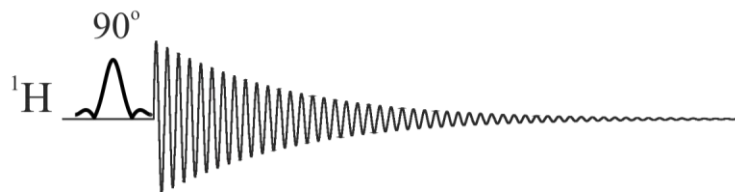
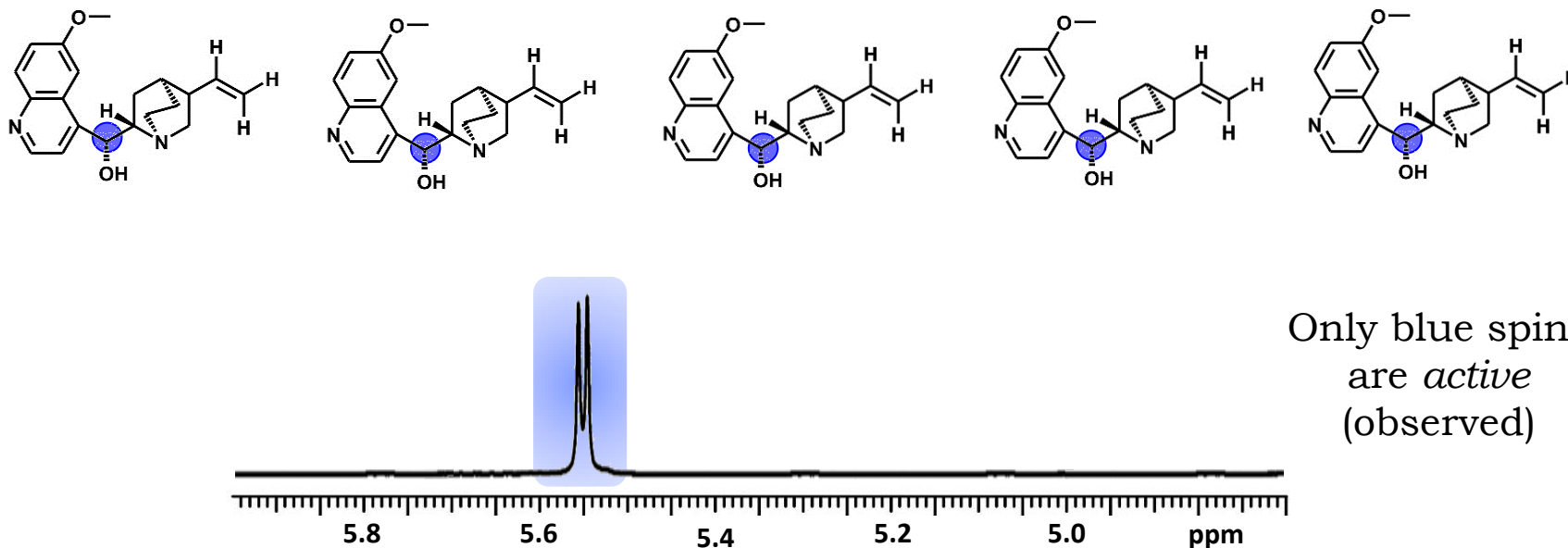


All spins are *active*
(observed)



The concept of *active* and *passive* spins

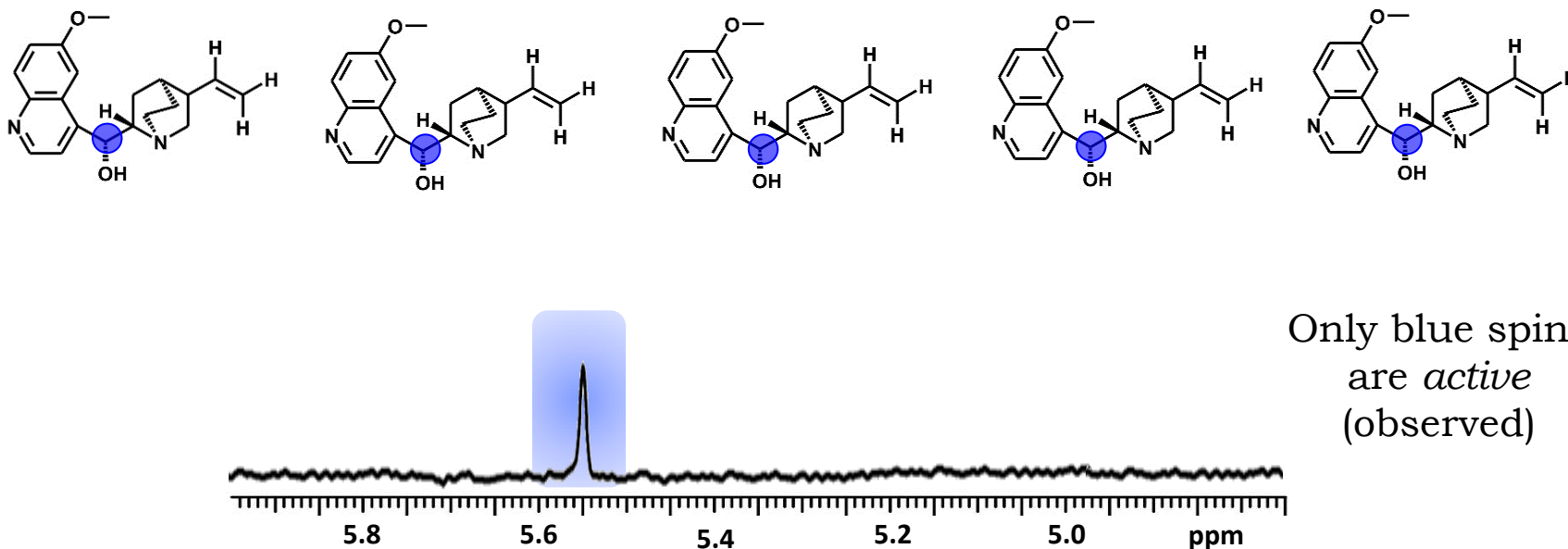
Band-selective ^1H NMR



- Divides the available spins into
- *active* (observed)
 - *passive* spins (non observed)

The concept of *active* and *passive* spins

Band-selective pure shift ^1H NMR

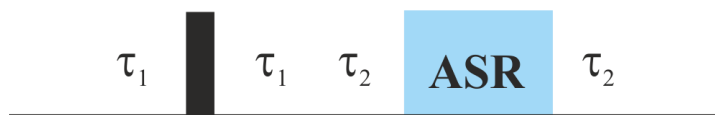


hard
 180° pulse

active spin
refocussing

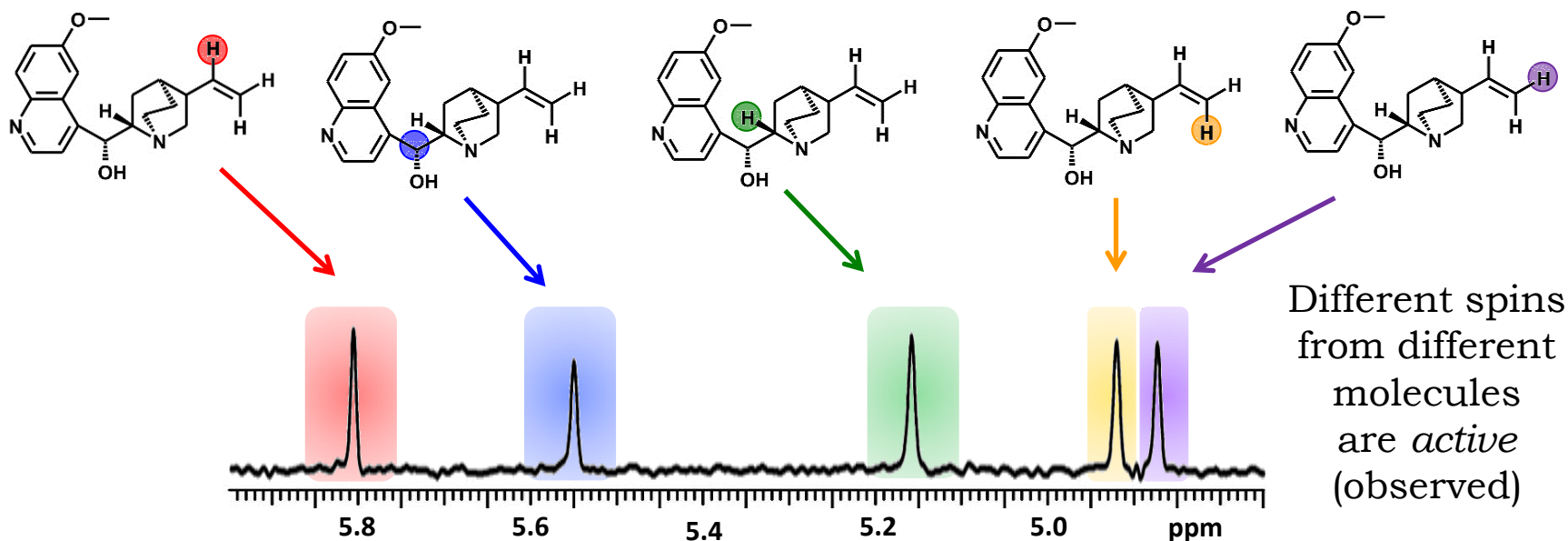
Divides the available spins into

- *active* (observed)
- *passive* spins (non observed)



The concept of *active* and *passive* spins

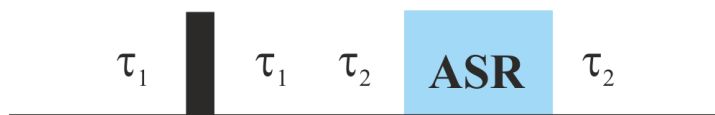
Pure shift ^1H NMR



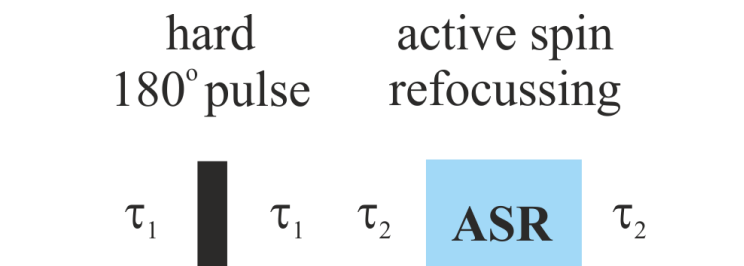
hard
 180° pulse

active spin
refocussing

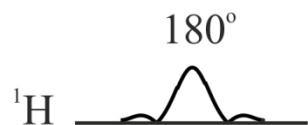
Divides the available spins into
- *active* (observed)
- *passive* spins (non observed)



Active spin refocussing (ASR) elements

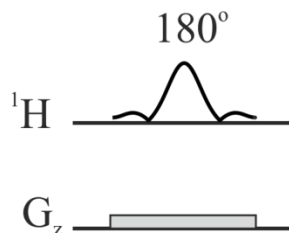


Band-selective



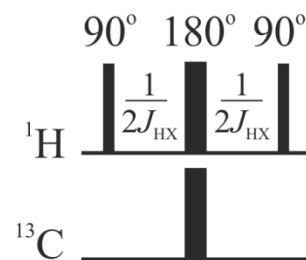
Shift selective
180° rotation

Zangger-Sterk



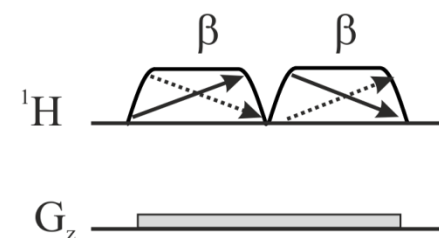
Slice and shift
selective
180° rotation

BIRD



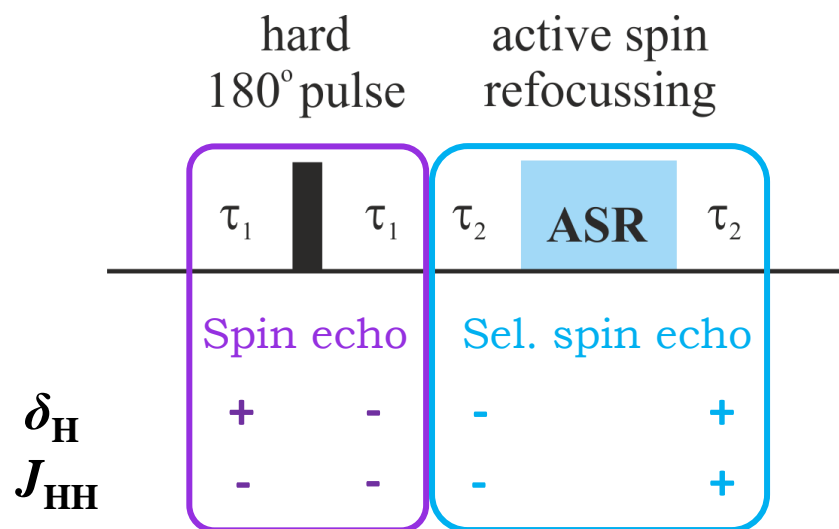
180° rotation
of protons
coupled to ^{13}C

PSYCHE

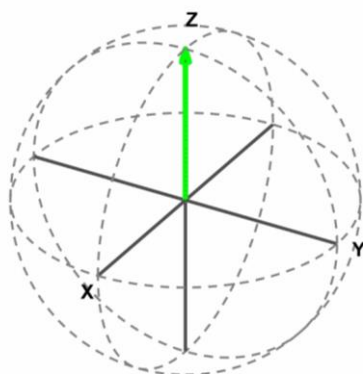


Flip a fraction
 $\sin^2\beta$ of spins

General mechanism for J-refocussing

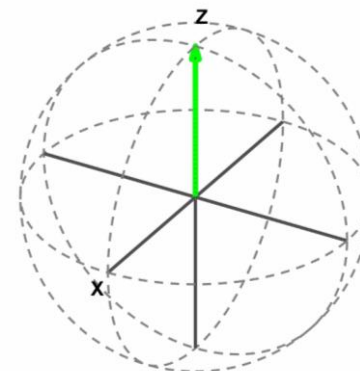


Spin echo



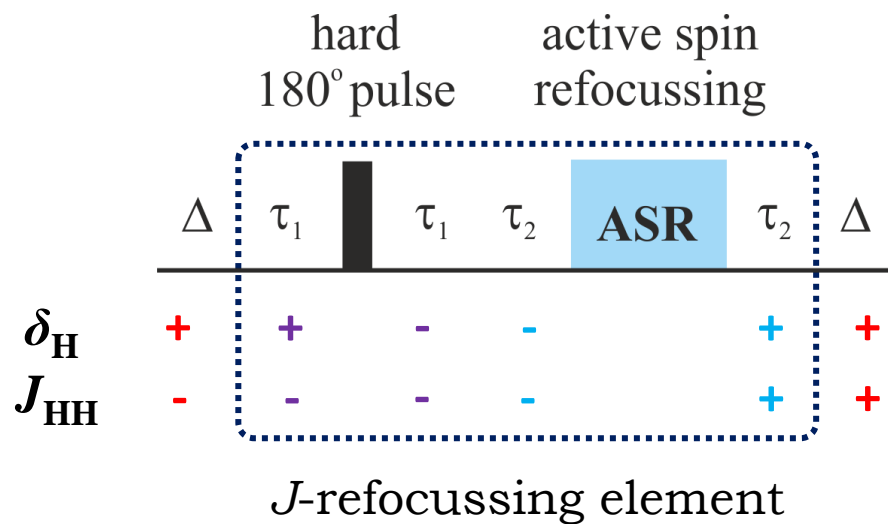
refocuses δ_H
but
not J_{HH}

Sel. spin echo



refocuses both
 δ_H and J_{HH}
(for active
spin only)

General mechanism for J-refocussing

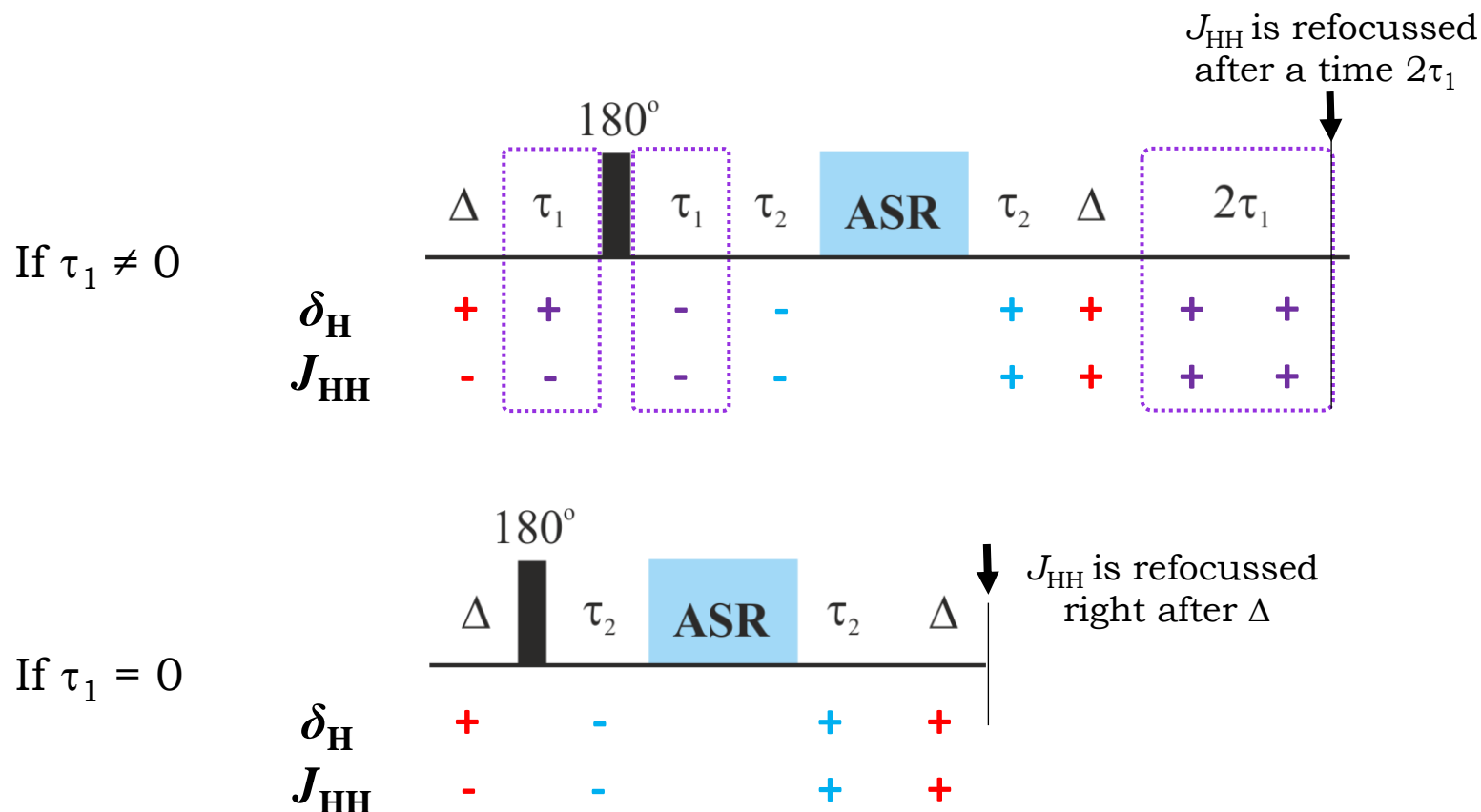


δ_H evolves

J_{HH} is refocused
(for active spin only)



Controlling J-refocussing time position



The timing of the J -refocussing element is carefully designed to have full control of when J_{HH} is refocussed (by changing τ_1)

How do we get an experimental pure shift NMR spectrum?

Homodecoupling element

J-refocussing element



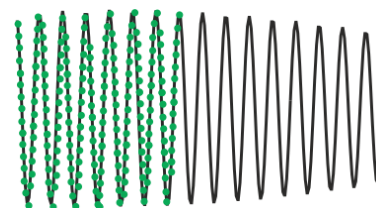
J_{HH} is refocused while δ_H evolves (for *active* spins only)

Active spin refocussing



Homodecoupled FID

Chunking data acquisition



Chunk duration (τ_c) $\ll 1/J_{HH}$

Acquisition methods

Interferogram

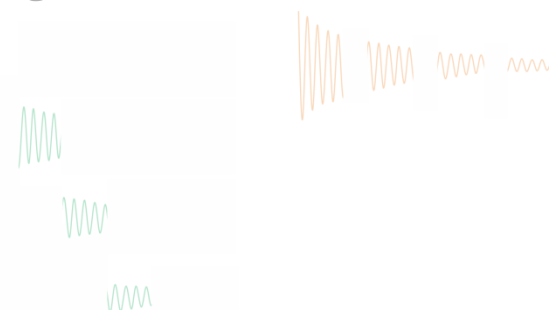
Real-time

1st experiment

2nd experiment

3rd experiment

4th experiment

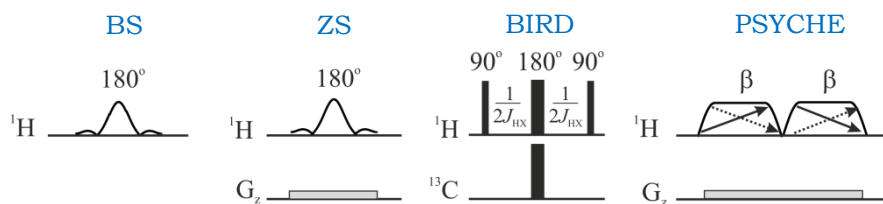
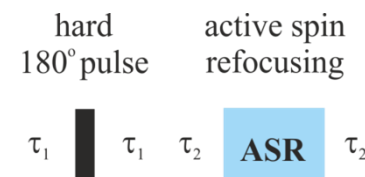


How do we remove the effect of J_{HH} coupling during FID?



We need to apply a homodecoupling element **after each acquisition point**

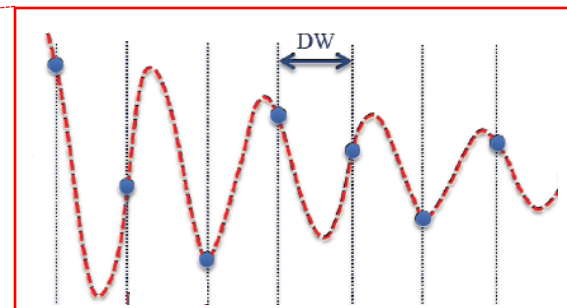
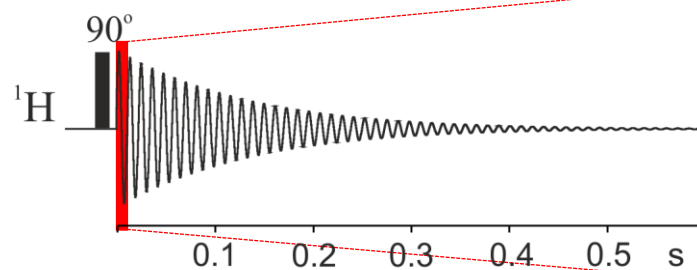
J -refocussing element



ASR duration (ms) 1-100 1-100 5-10 20-40

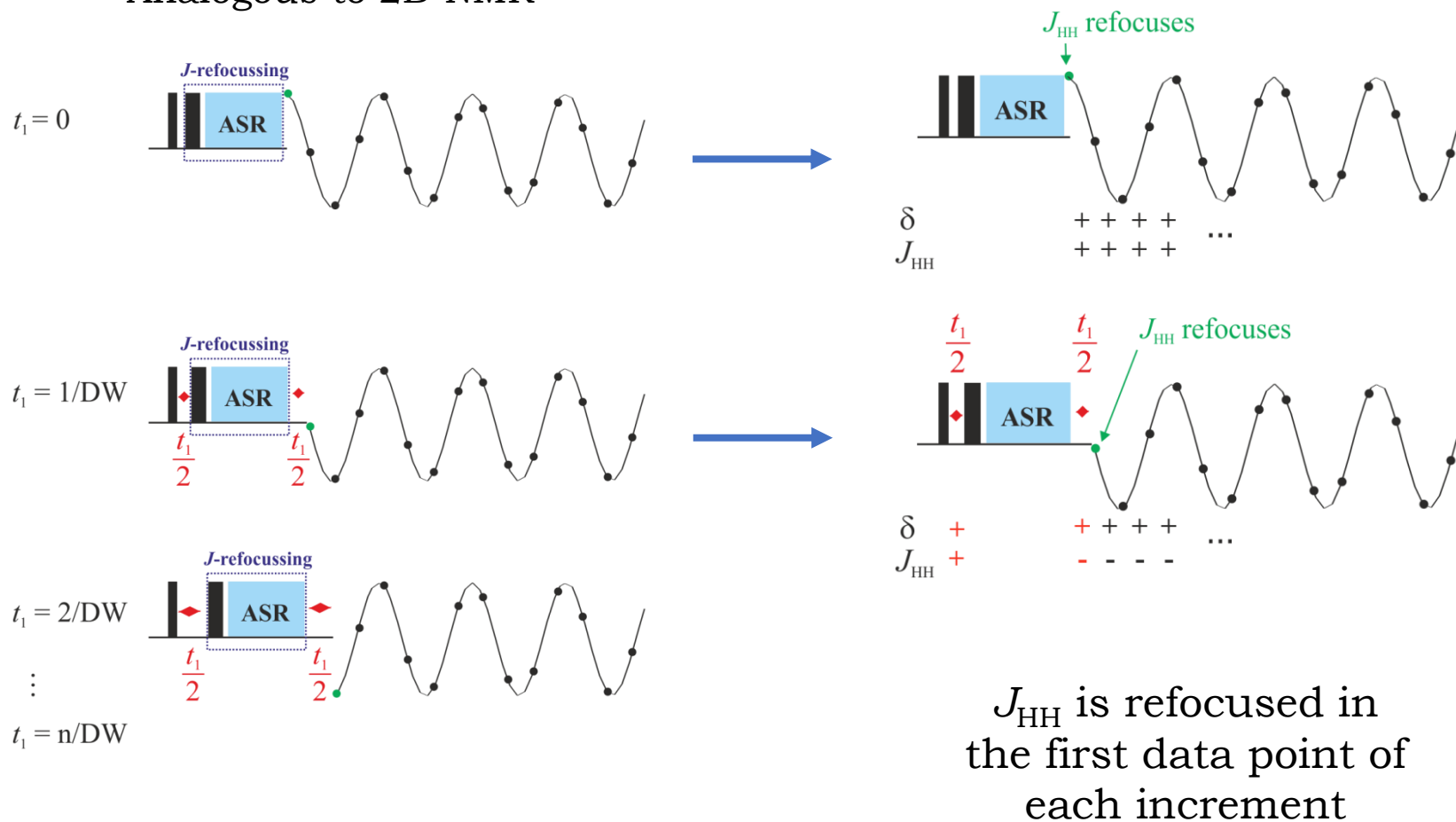
Sampling interval (dwell time) $\sim 50 \mu\text{s}$

**Not
(currently)
possible**



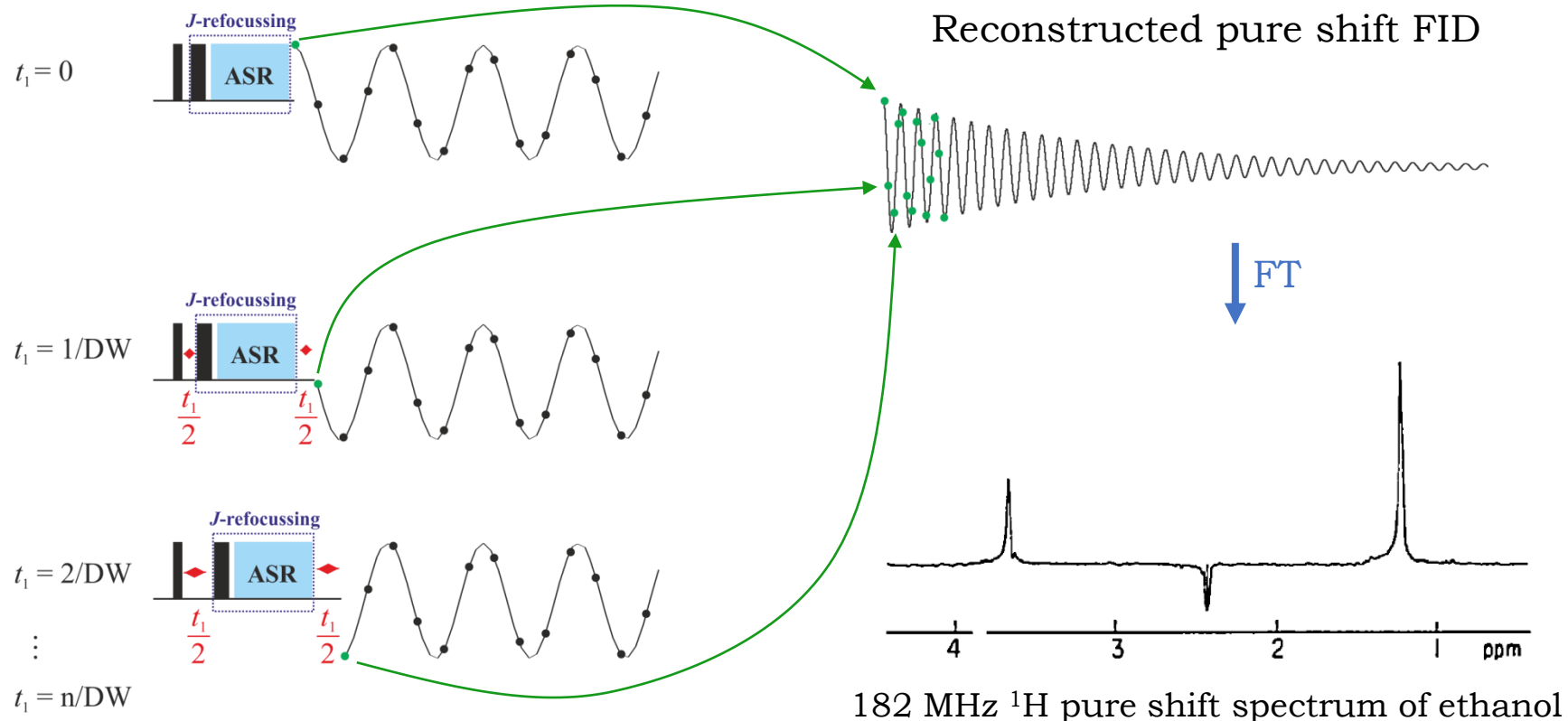
Point-by-point data acquisition (first approach)

Analogous to 2D NMR



Point-by-point data acquisition (first approach)

Analogous to 2D NMR



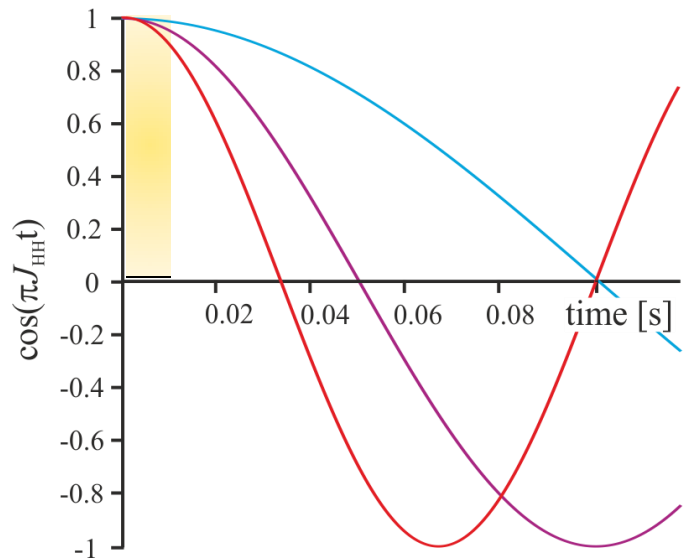
**Extremely
long
(impractical)**

Experiment time $\sim 24\text{h}$
(8192 t_1 increments)

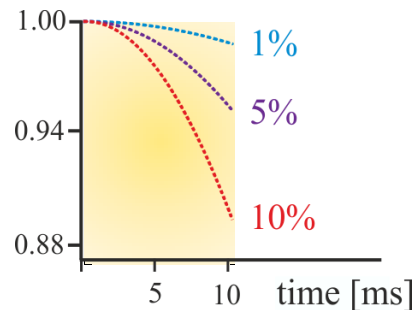
FID data points = 8192
(2.5 Hz resolution)

Exploiting the relative slowness of J_{HH} evolution

J_{HH} evolution for a AX spin system

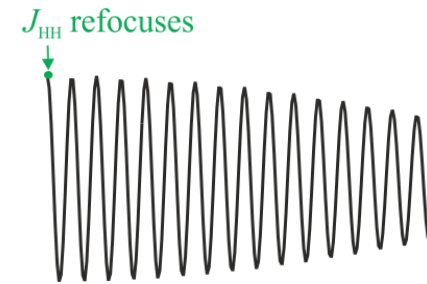


$J_{HH} = 5$ Hz
 $J_{HH} = 10$ Hz
 $J_{HH} = 15$ Hz

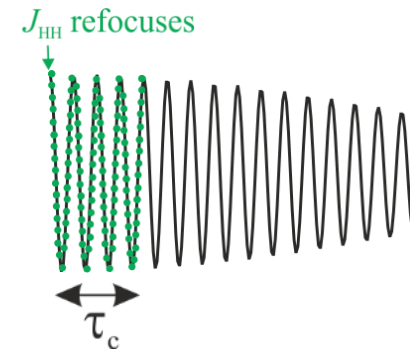


**J_{HH} evolution
 during the chunk
 is negligible**

Point-by-point data acquisition

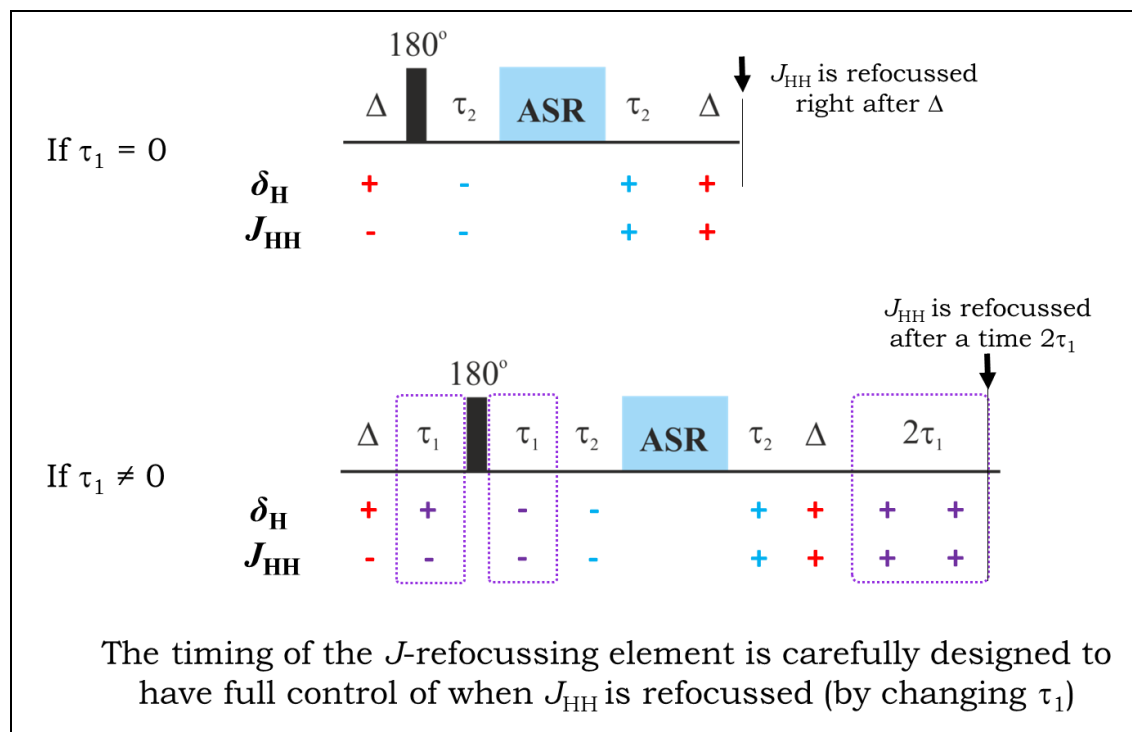
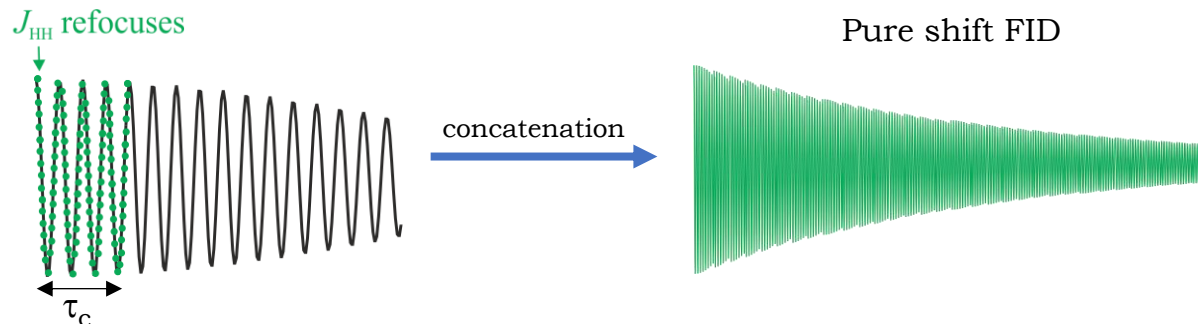
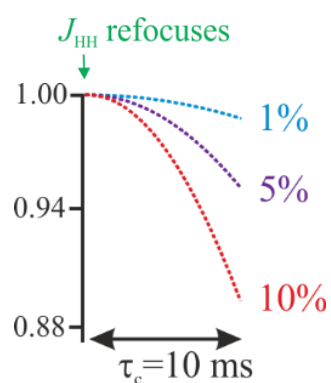


Chunking data acquisition

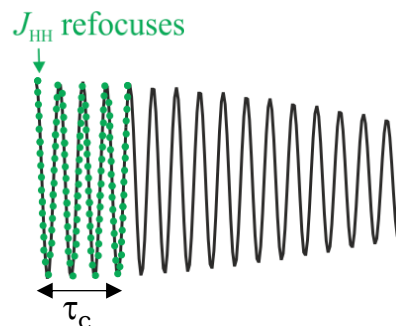
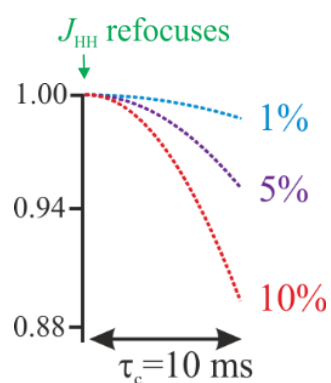


← chunk duration (τ_c) $\ll 1/J_{HH}$
 (~200 data points per chunk)

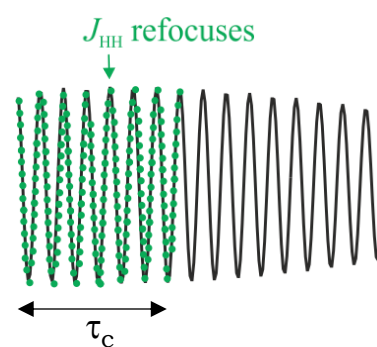
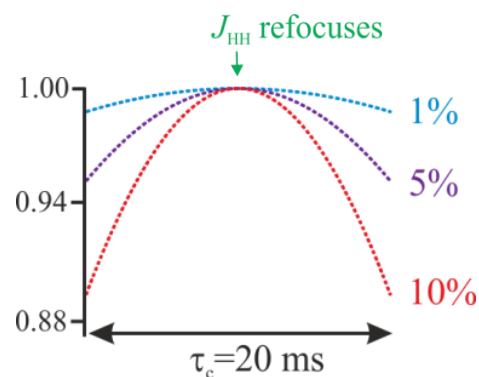
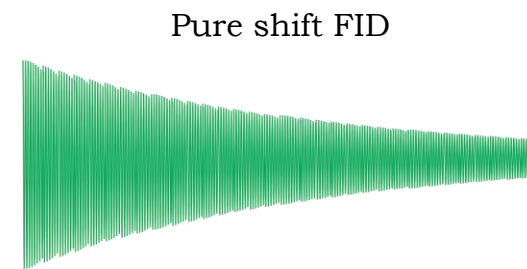
Chunking data acquisition (speeding things up)



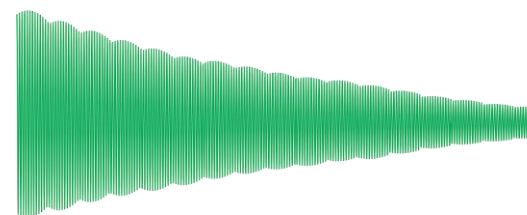
Chunking data acquisition (speeding things up)



concatenation



concatenation



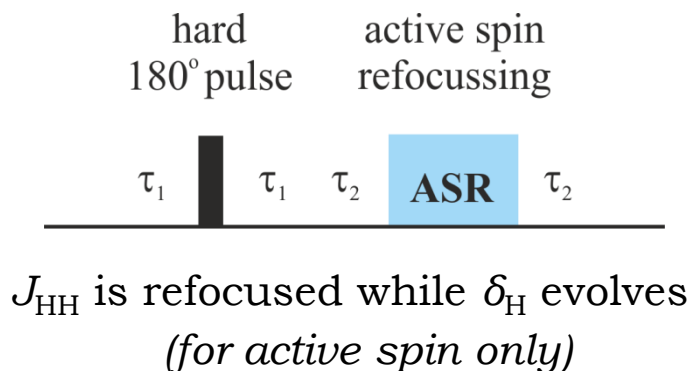
Refocussing J_{HH}
in the middle
of the chunk

Allows to acquire
chunks twice
longer

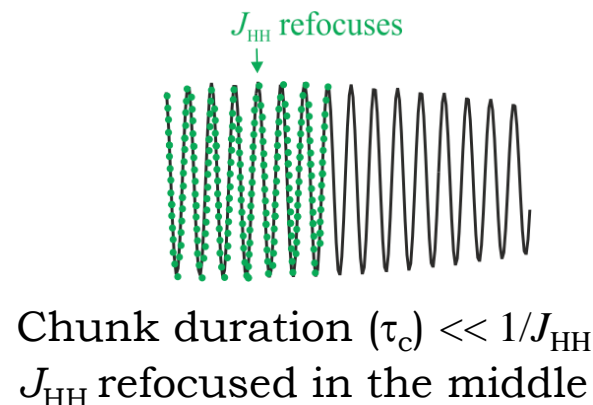
Reducing experiment time
(half number of chunks are
needed for achieving the same
FID resolution)

Pure shift key ideas

J -refocussing element

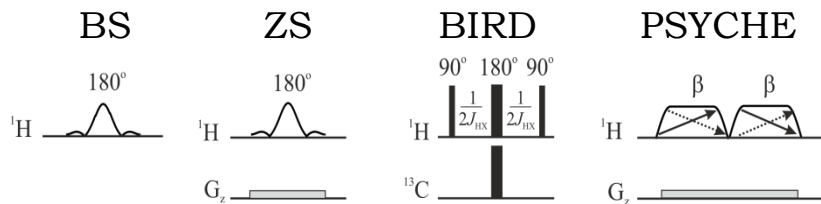


Chunking data acquisition



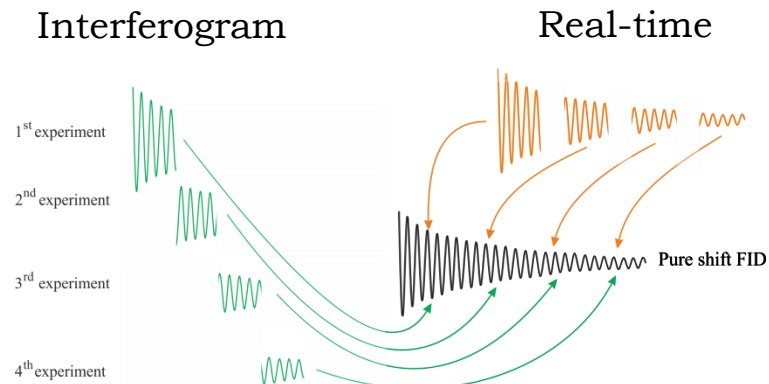
Pure shift methods

Active spin refocussing



Divides the spins into *active* (observed) and *passive* spins (non observed)

Acquisition methods

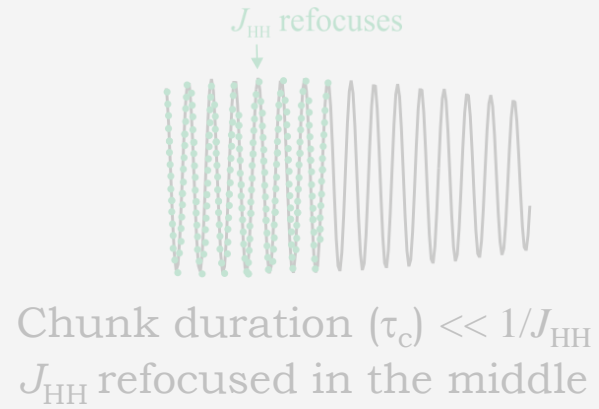


Pure shift key ideas

J-refocussing element



Chunking data acquisition



Pure shift methods

Active spin refocussing

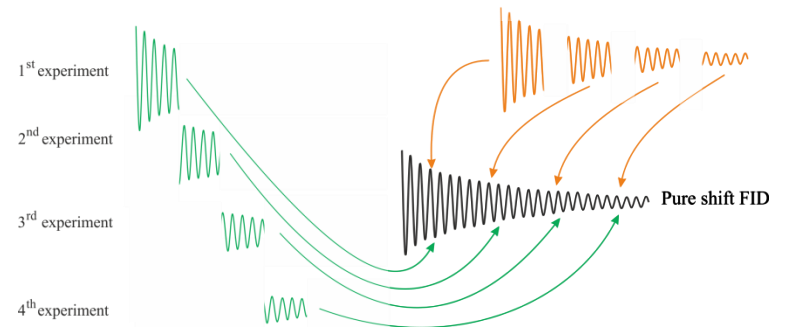


Divides the spins into *active* (observed) and *passive* spins (non observed)

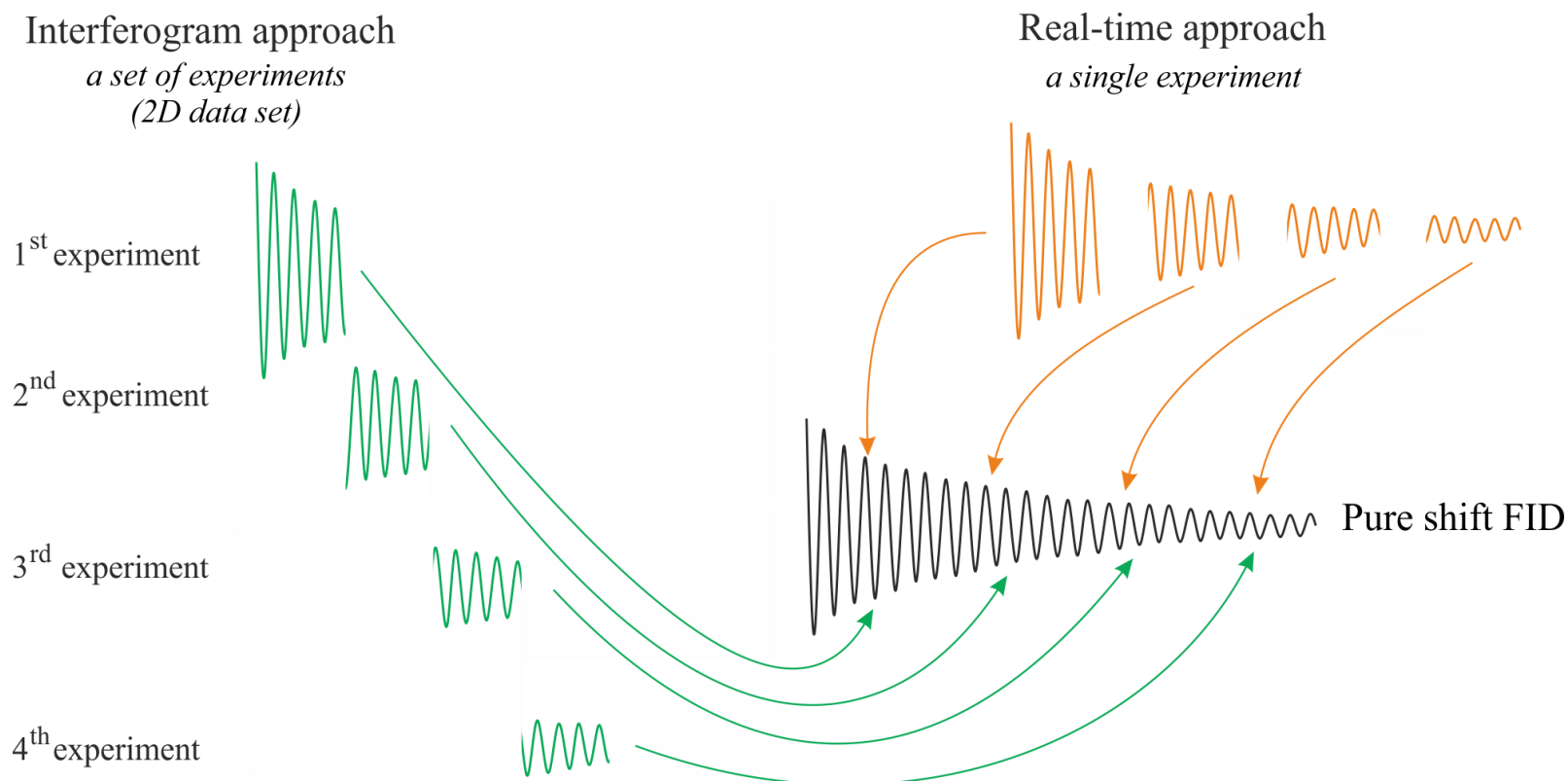
Acquisition methods

Interferogram

Real-time



Pure shift acquisition methods

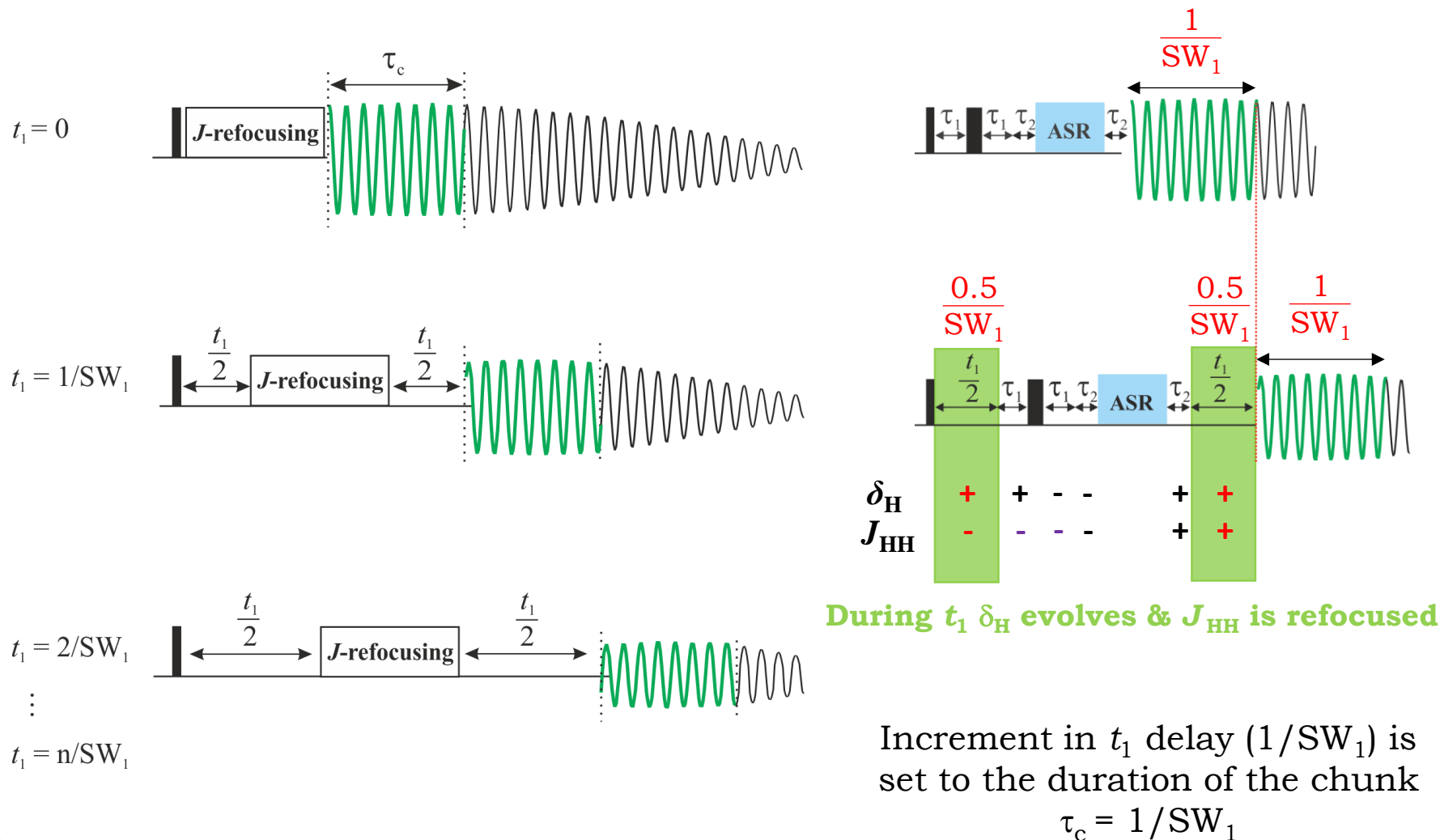


Chunk duration (τ_c) $\ll 1/J_{\text{HH}}$ ($\tau_c \sim 10\text{-}20$ ms)

J_{HH} -evolution refocused in the middle of the chunk

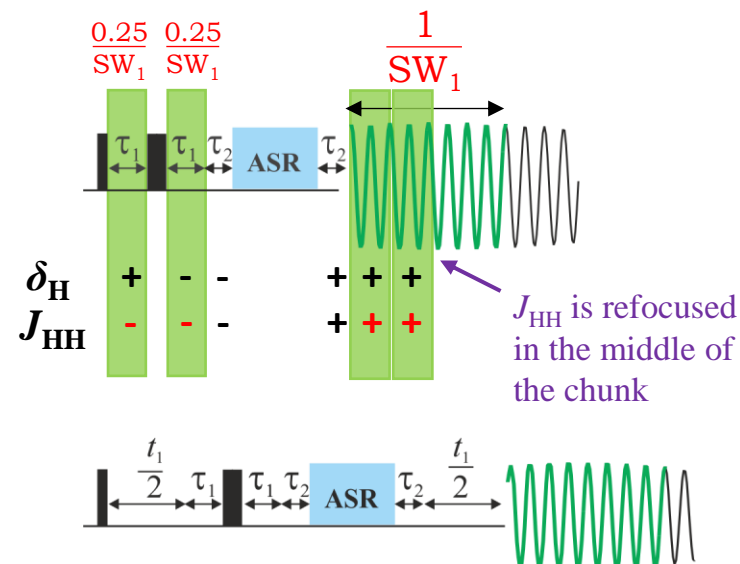
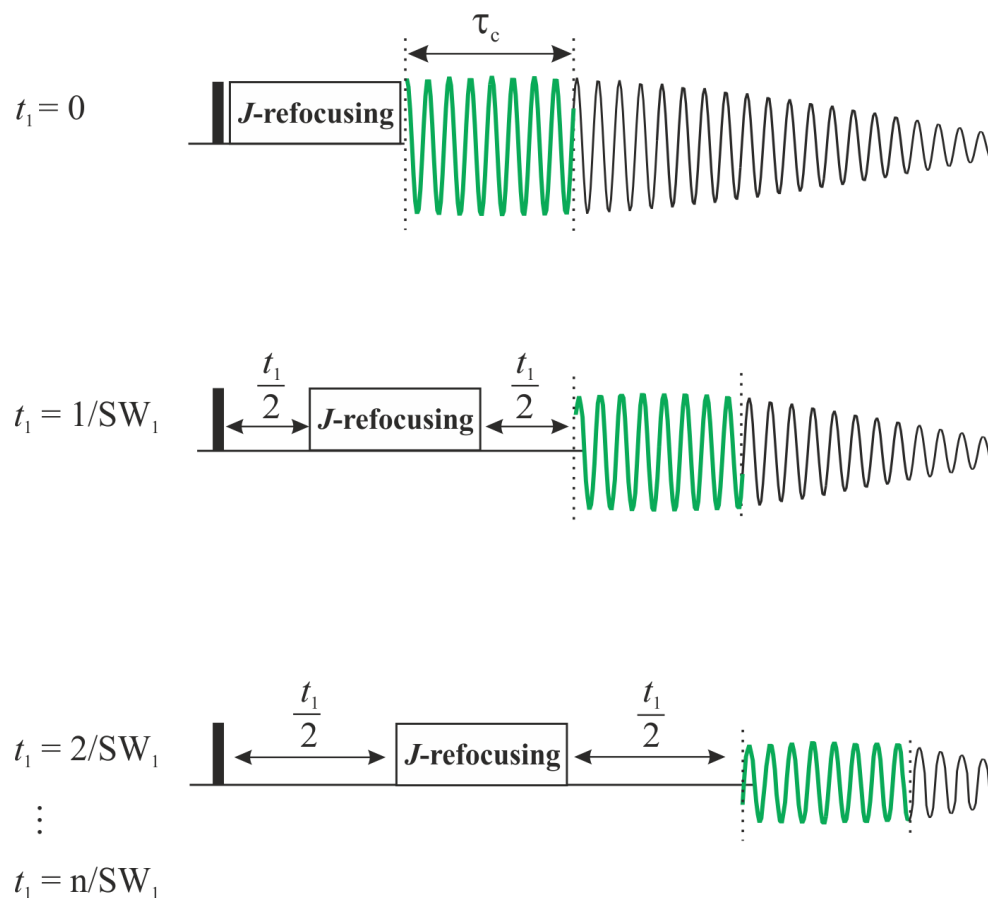
Interferogram pure shift experiments – 2D acquisition

Analogous to 2D NMR



Interferogram pure shift experiments – 2D acquisition

Analogous to 2D NMR

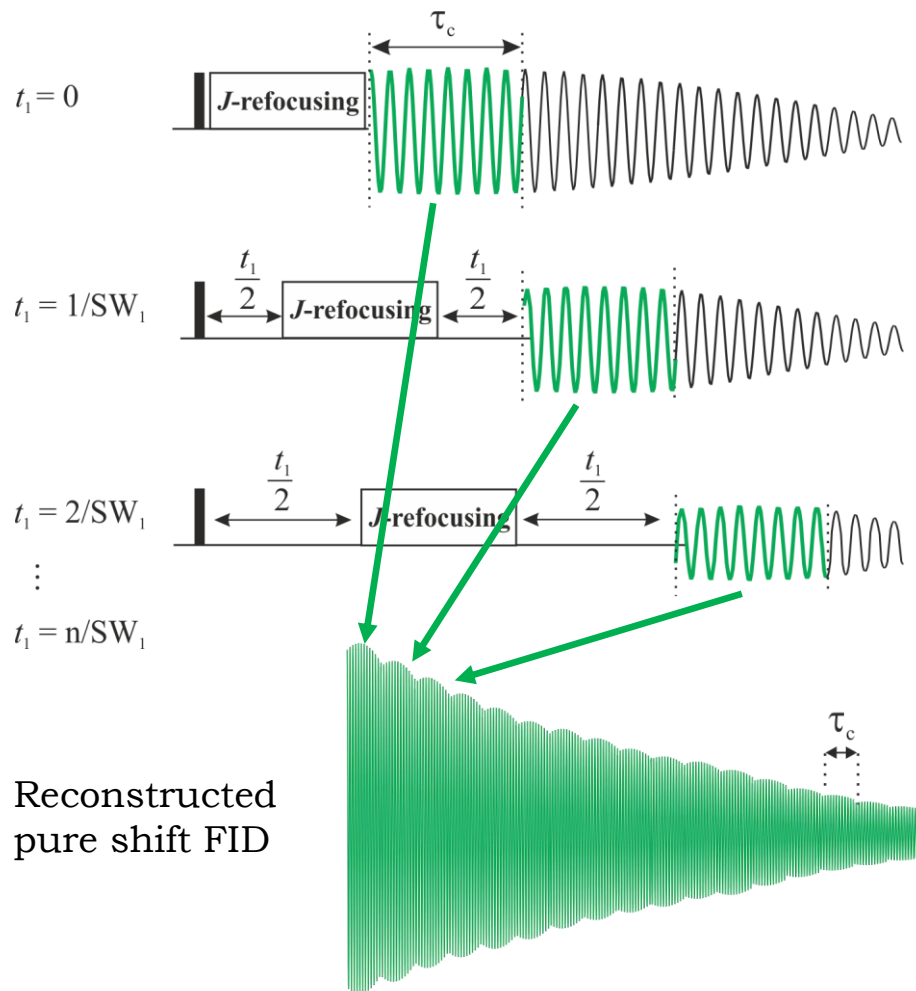


τ_1 is normally set to ensure that J_{HH} is refocused at the midpoint of each chunk

Increment in t_1 delay ($1/SW_1$) is set to the duration of the chunk

$$\tau_c = 1/SW_1$$

Interferogram pure shift experiments – 2D acquisition



Special data processing is needed

Extract first chunk of
each increment



Assemble them sequentially
(reconstructed pure shift FID)

Pure shift
signal

chunking artefacts

Acquiring pure shift data in chunks of duration τ_c gives rise to J -sidebands (“chunking artefacts”) with a spacing $1/\tau_c$ in the spectrum

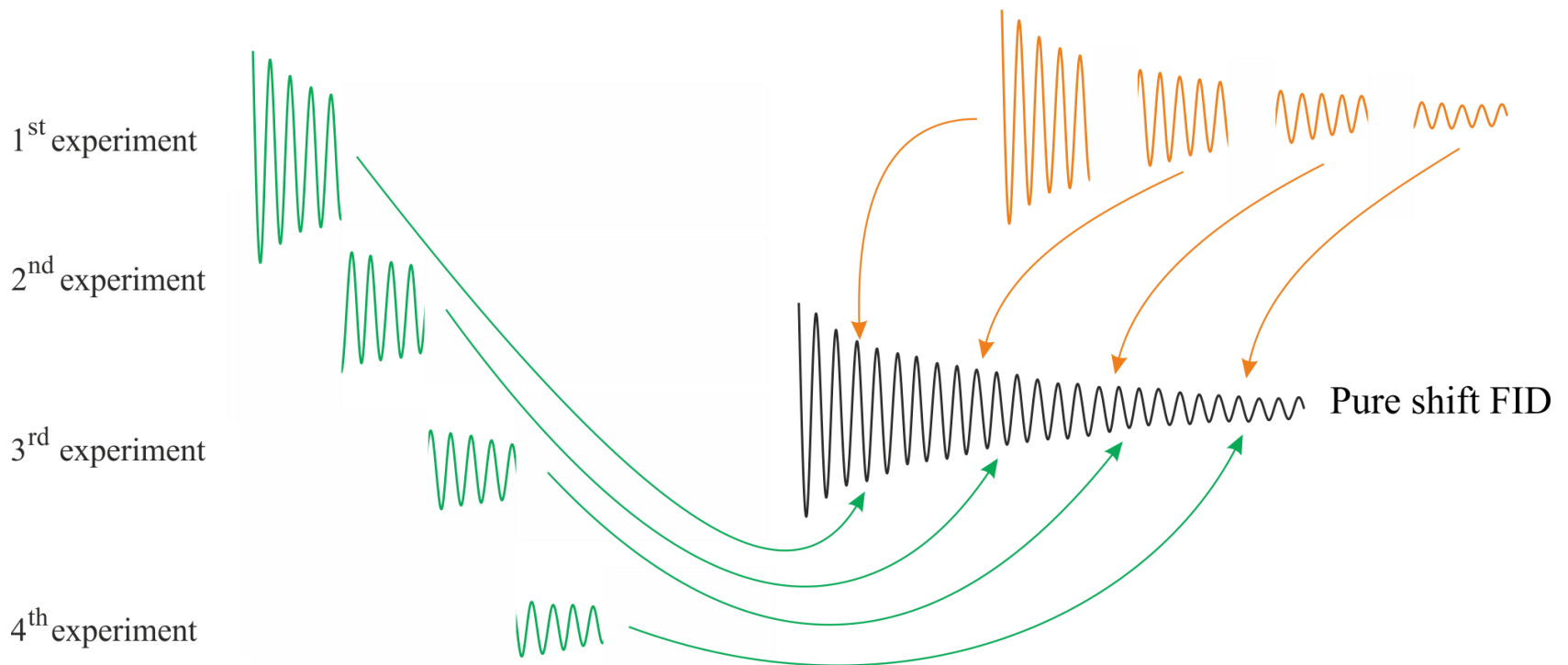
Speeding things up

Interferogram approach

a set of experiments

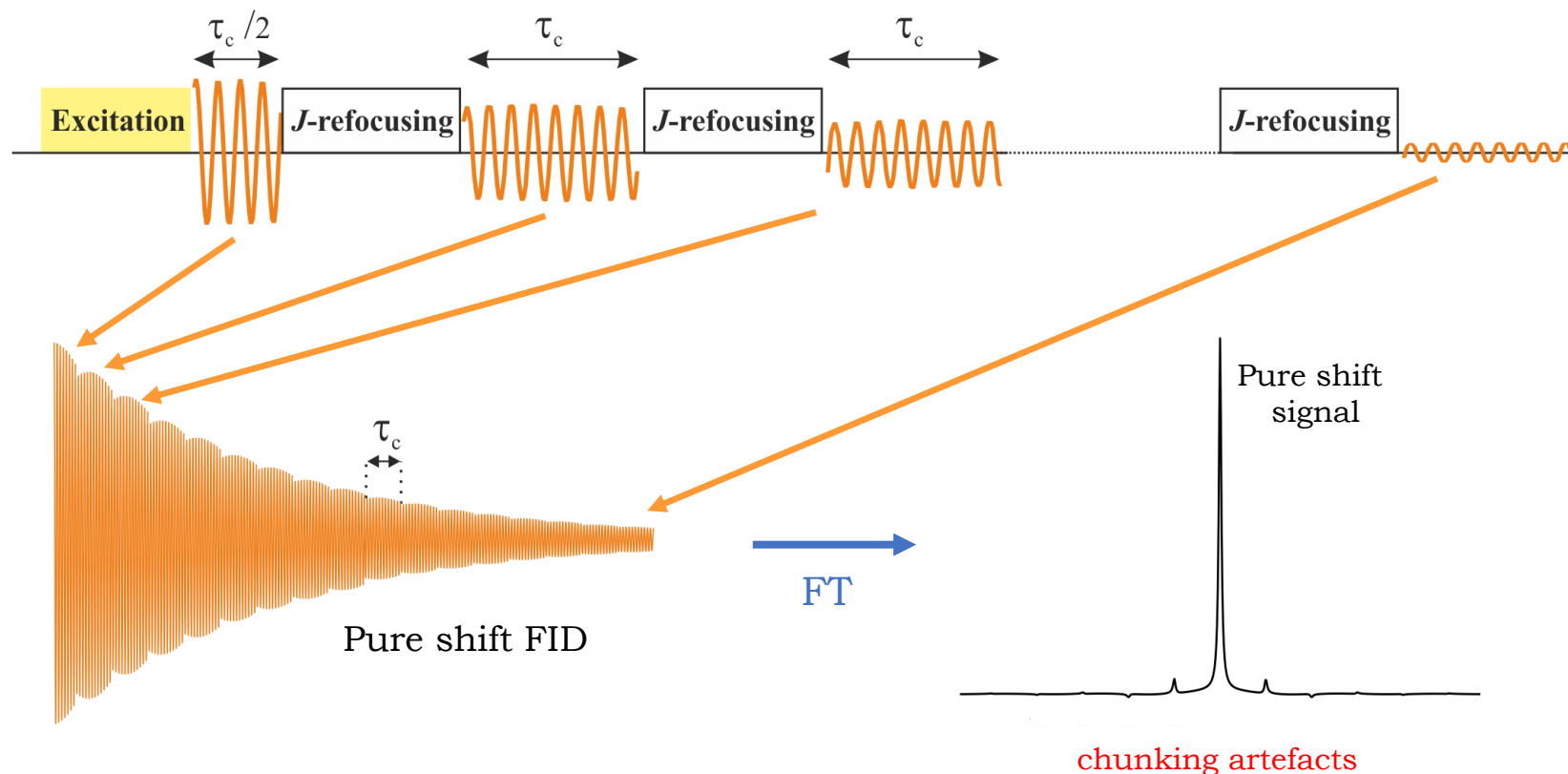
Real-time approach

a single experiment



Real-time pure shift experiments – 1D acquisition

Single shot acquisition

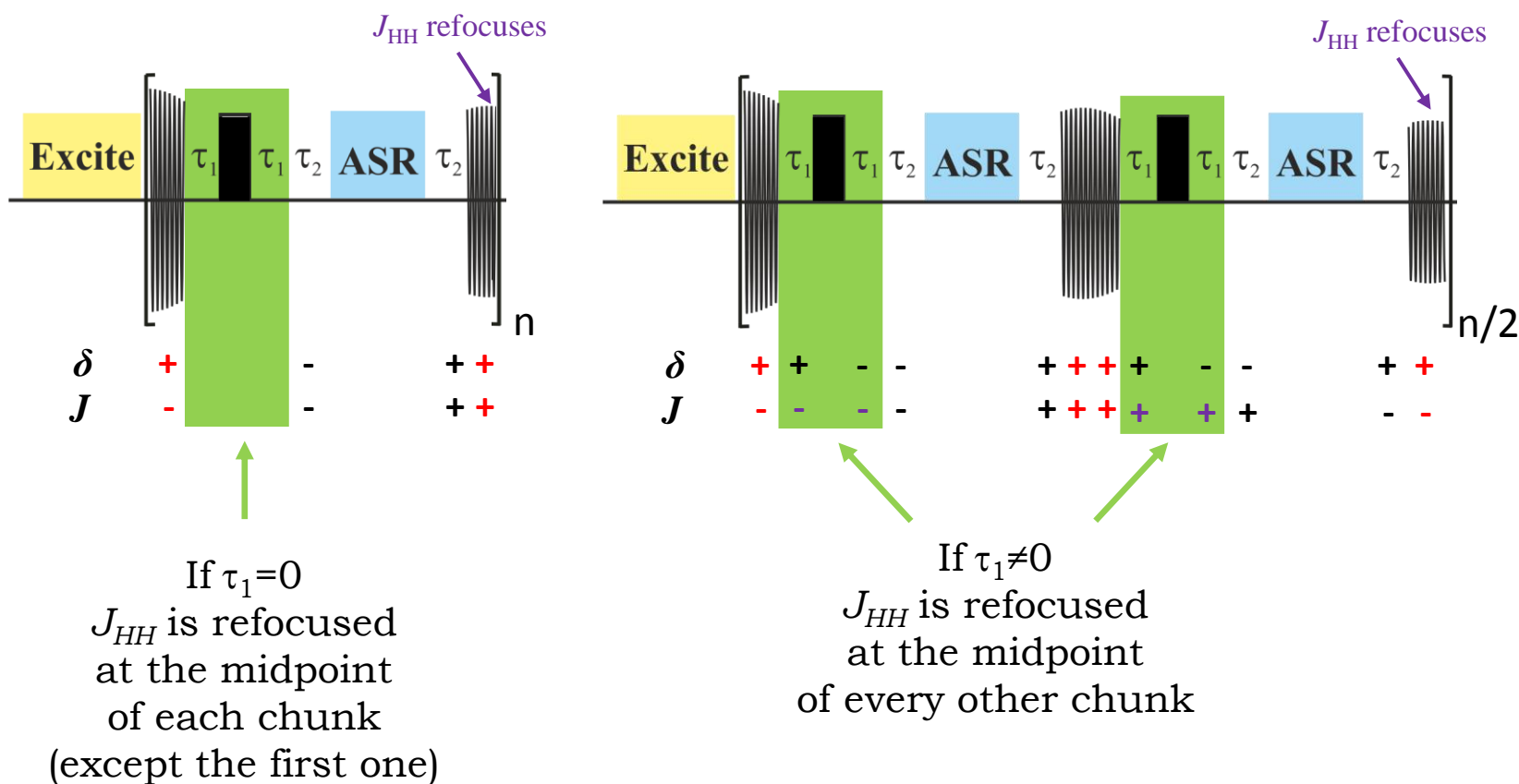


Pure shift FID acquired directly
(special data processing is **NOT** needed)

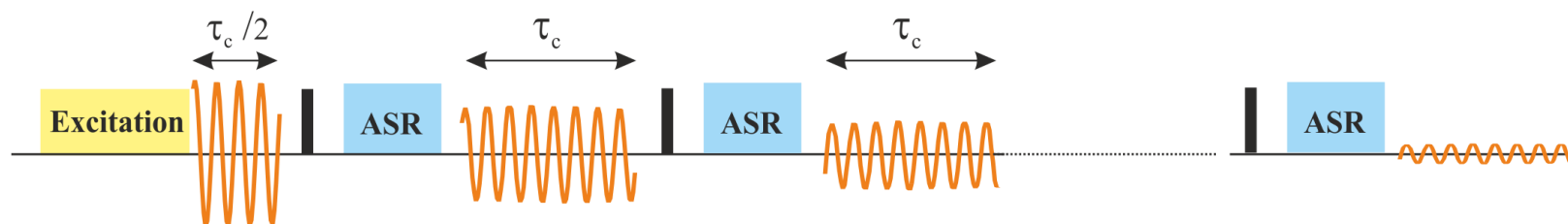
Acquiring pure shift data in chunks of
duration τ_c gives rise to J -sidebands
with a spacing $1/\tau_c$ in the spectrum

Real-time pure shift experiments – 1D acquisition

J -refocussing in real time



Resolution in real-time pure shift spectra



Digital resolution

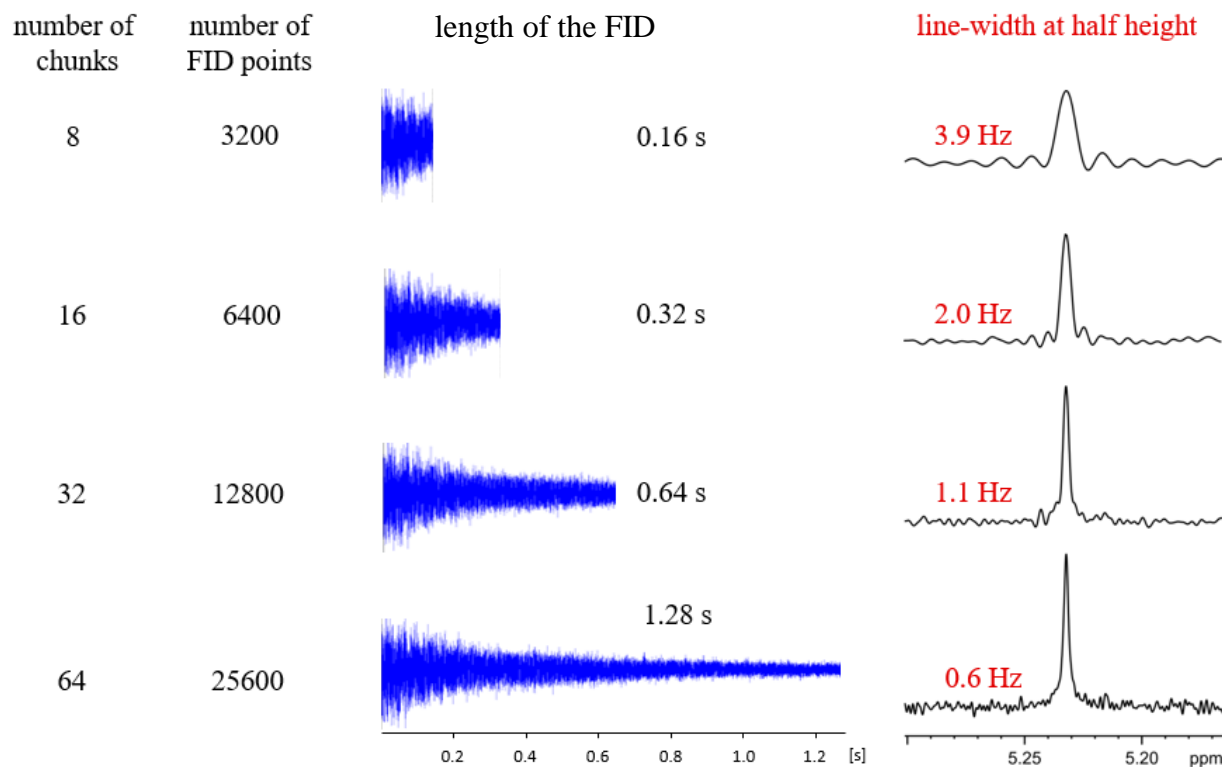
Increasing the number of chunks



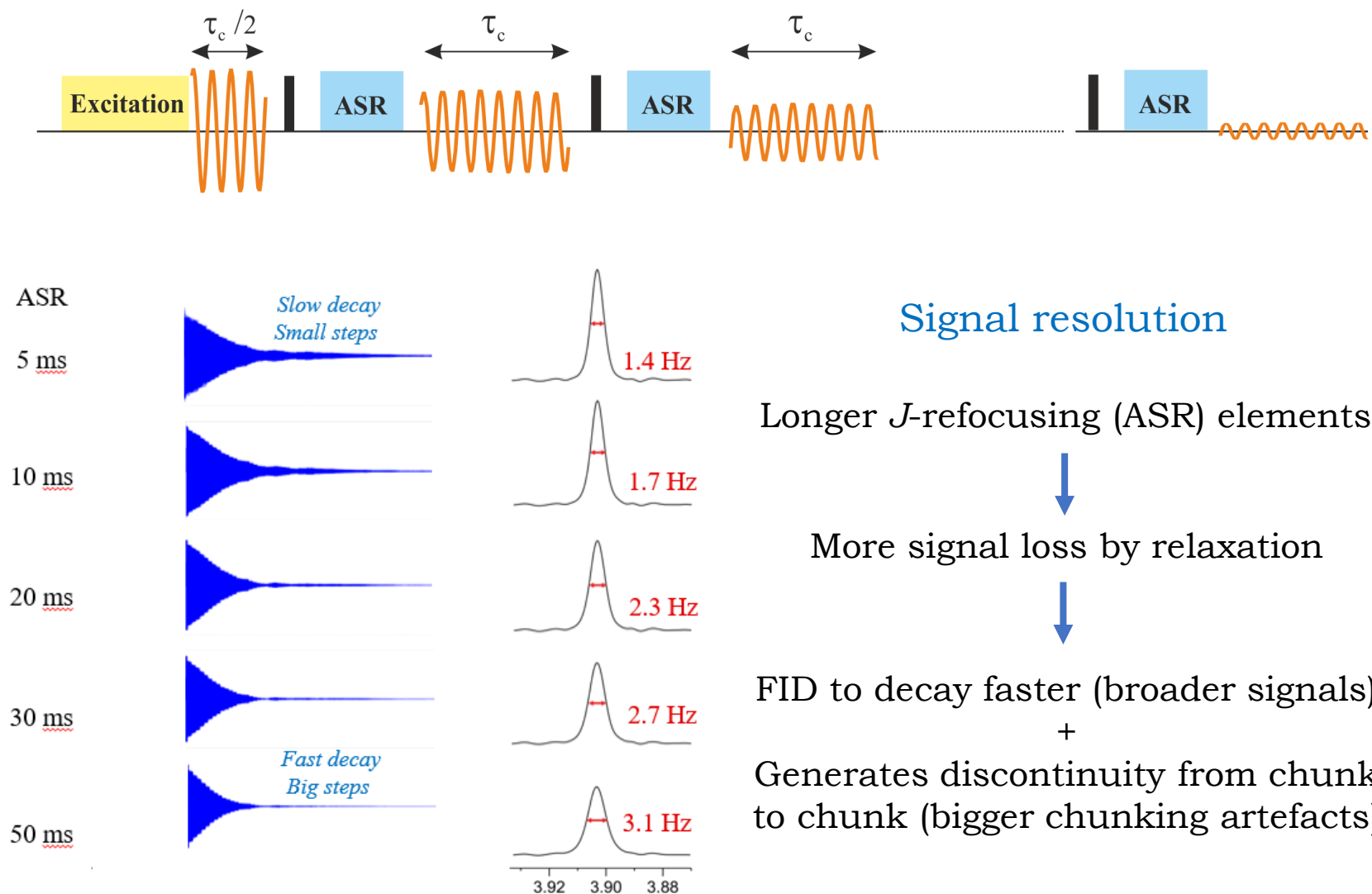
Improves digital resolution



Problems with irreproducibility from chunk to chunk occurs



Resolution in real-time pure shift spectra



Pros and cons

Interferogram approach

- ☹️ A set of experiments
- ☹️ Low SNR /experiment time
- ☹️ Special data processing
- ☹️ Chunking artefacts
- 😊 Smooth FID decay
- 😊 High spectral quality

Real-time approach

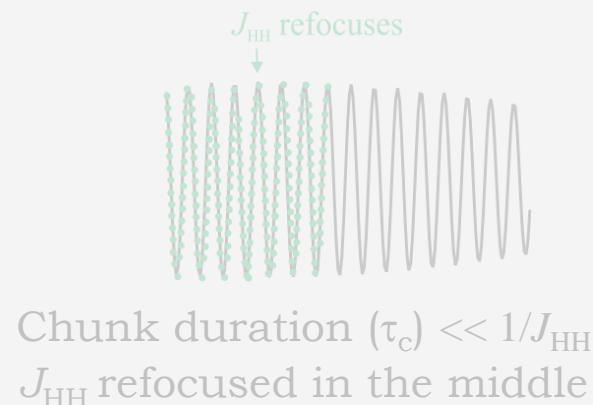
- 😊 A single experiment
- 😊 High SNR /experiment time
- 😊 Direct Fourier transformed
- ☹️ Chunking artefacts
- ☹️ FID discontinuities
- ☹️ Variable spectral quality

Pure shift key ideas

J -refocussing element

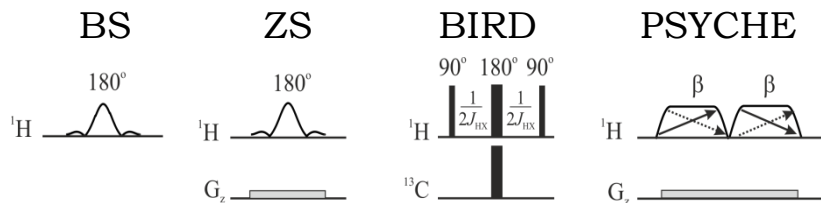


Chunking data acquisition



Pure shift methods

Active spin refocussing

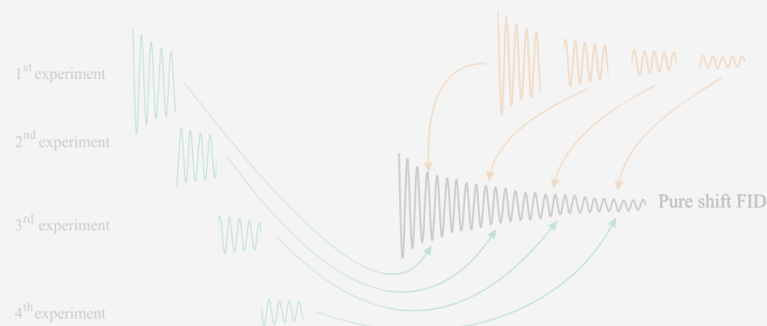


Divides the spins into *active* (observed) and *passive* spins (non observed)

Acquisition methods

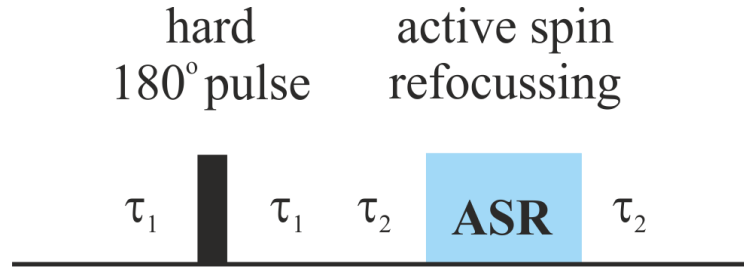
Interferogram

Real-time

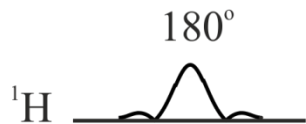


Active spin refocussing (ASR) elements

Divides the spins into *active* (observed) and *passive* spins (non observed)

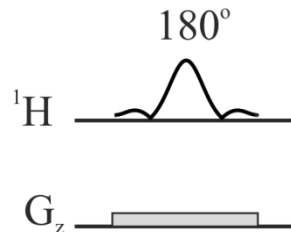


Band-selective



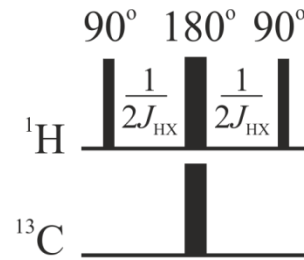
Shift selective
 180° rotation

Zangger-Sterk



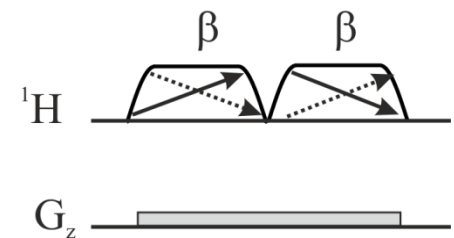
Slice and shift
selective
 180° rotation

BIRD



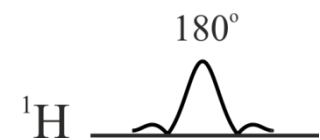
180° rotation
of protons
coupled to ^{13}C

PSYCHE

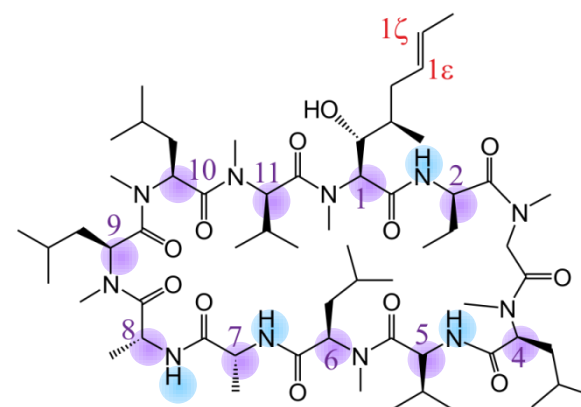
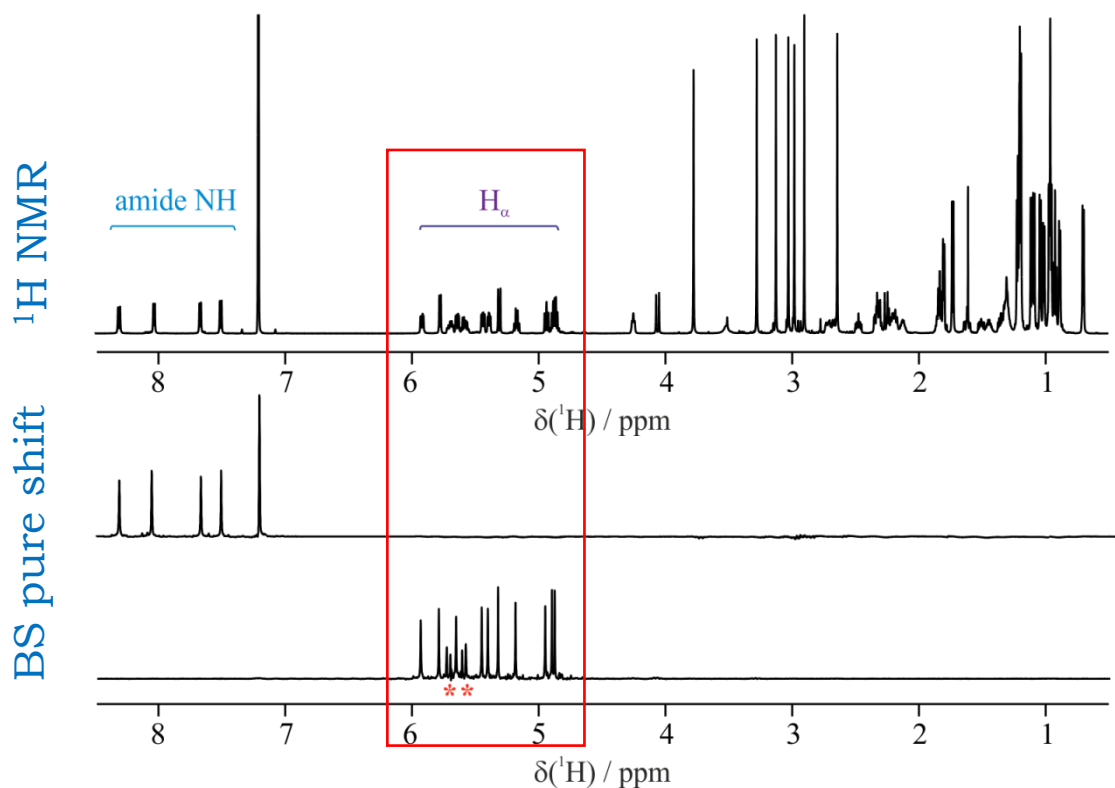


Flip a fraction
 $\sin^2\beta$ of spins

Band-selective – BS, HOBS, BASH and BASHD

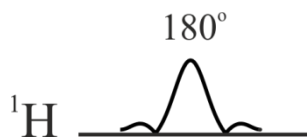


- The selective 180° pulse refocuses only protons within its bandwidth (*active* spins)
- One (frequency selection) or several resonances (band-selection or multiple-frequency selection) can be homodecoupled

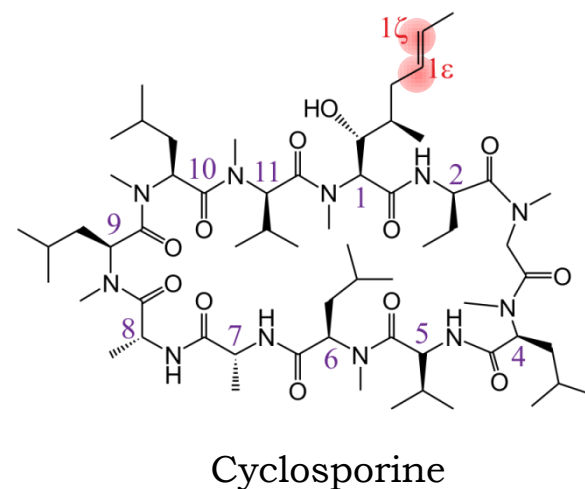
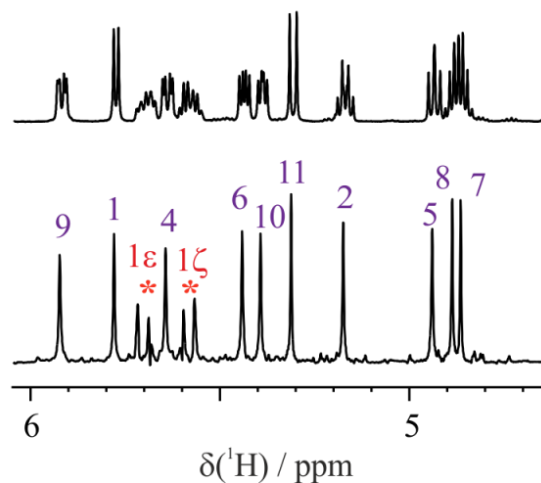


Cyclosporine

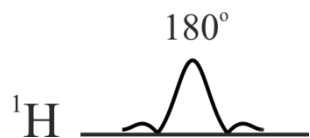
Band-selective – BS, HOBS, BASH and BASHD



- The selective 180° pulse refocuses only protons within its bandwidth (*active* spins)
- One (frequency selection) or several resonances (band-selection or multiple-frequency selection) can be homodecoupled
- Need to avoid exciting coupled protons
- Suitable for the analysis of molecules with well-separated regions (peptides and proteins) and isomeric mixtures

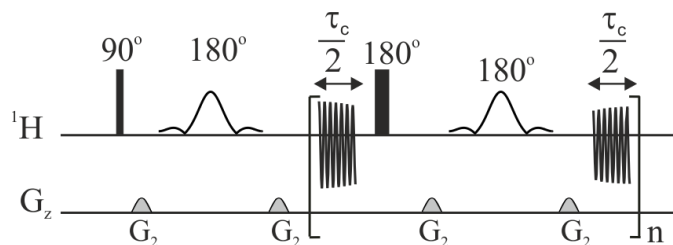


Band-selective pure shift methods

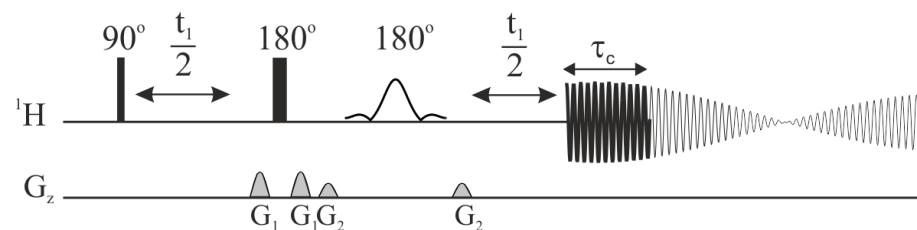


- Compatible with both real-time and interferogram acquisition

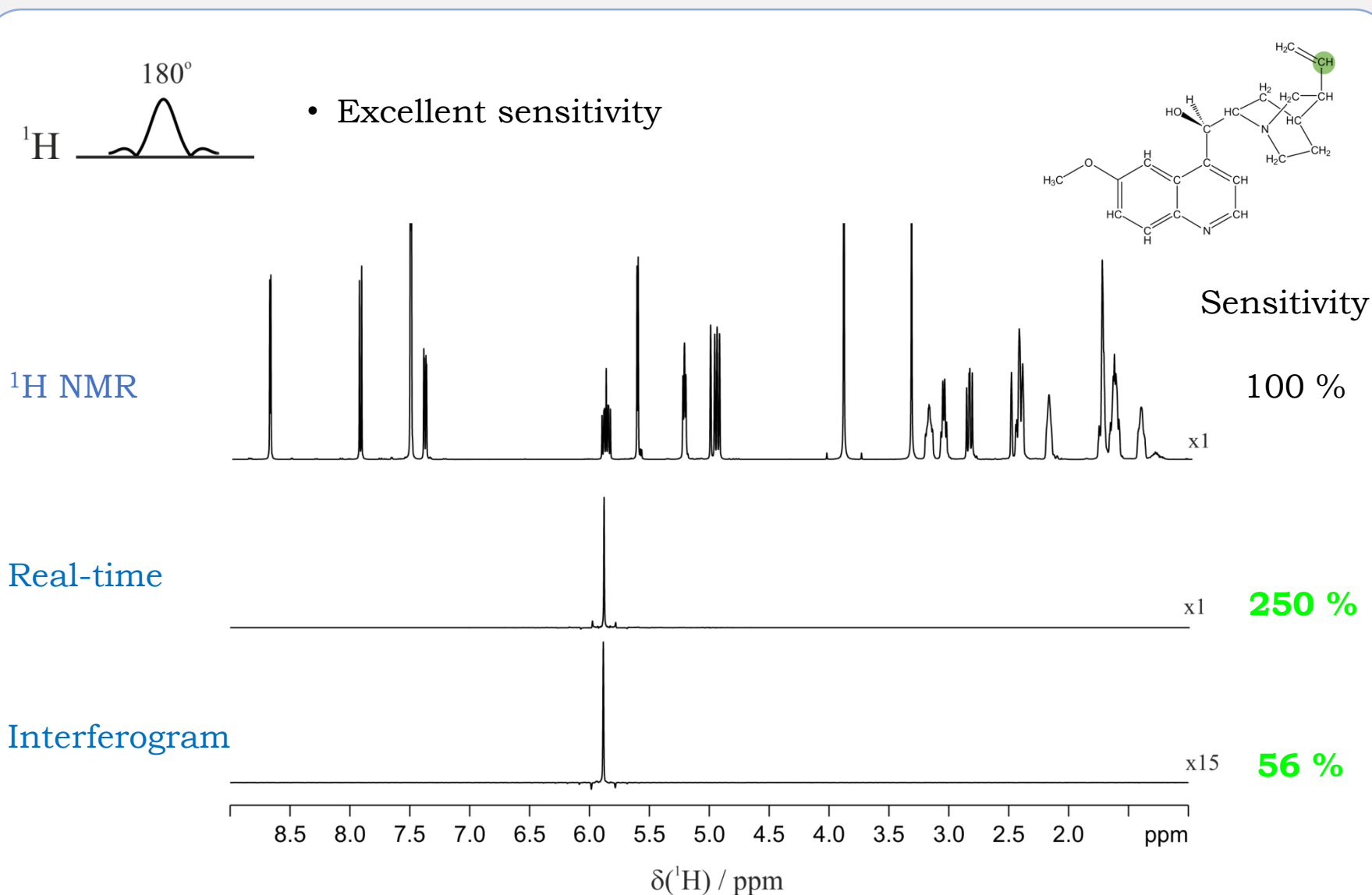
Real-time BS experiment



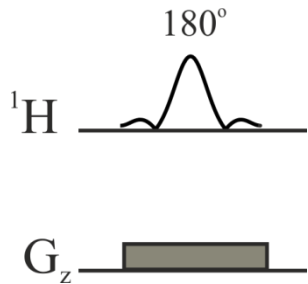
Interferogram BS experiment



Sensitivity in band-selective pure shift spectra

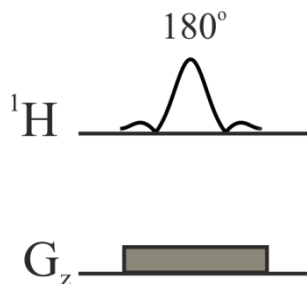


Zangger-Sterk (ZS)



- Simultaneous application of a selective 180° pulse and a weak pulse field gradient
- Slice and shift selection
- Each *active* spin is refocused in a narrow region (slice) of the sample
- Coupled spins cannot be refocused within the same slice

Zanger-Sterk (ZS)

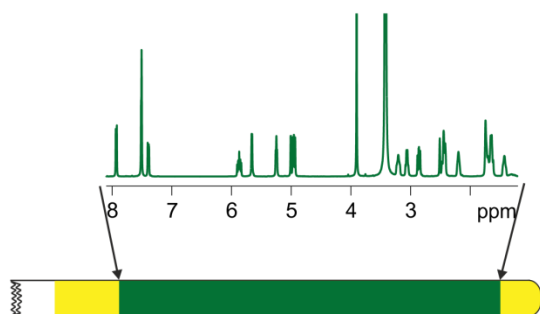
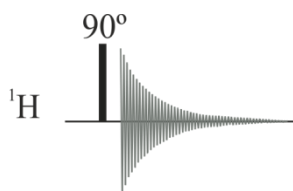


- Simultaneous application of a selective 180° pulse and a weak pulse field gradient
- **Slice and shift selection**
- Each *active* spin is refocused in a narrow region (slice) of the sample
- Coupled spins cannot be refocused within the same slice



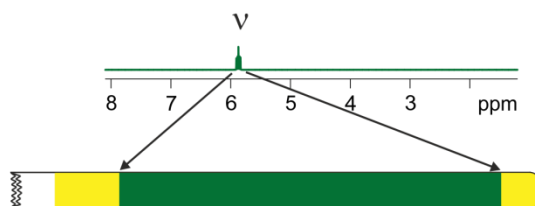
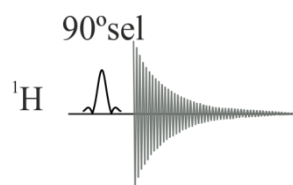
Broadband excitation – shift selection – band selection

Broadband



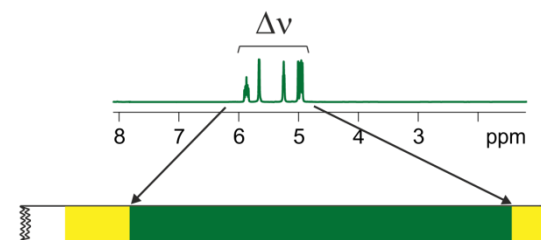
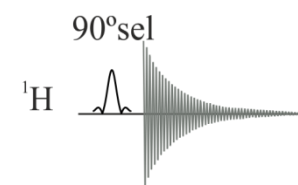
All frequencies are excited in the whole active volume

Frequency selection



A narrow range of frequencies is excited in the whole active volume

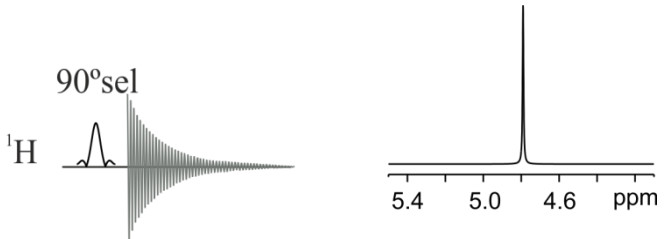
Band selection



A wide range of frequencies is excited in the whole active volume

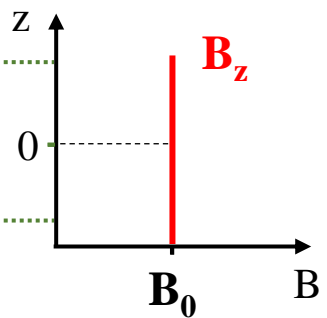
Shift selection – slice and shift selection

Shift selection



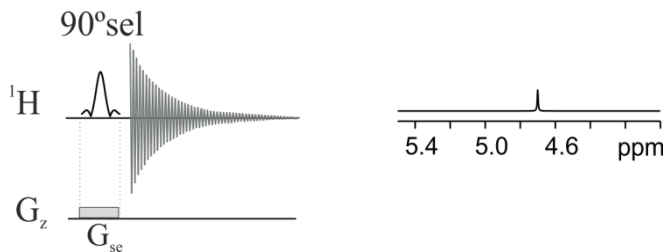
$$\nu_i \propto B_z$$

$$B_z = B_0$$



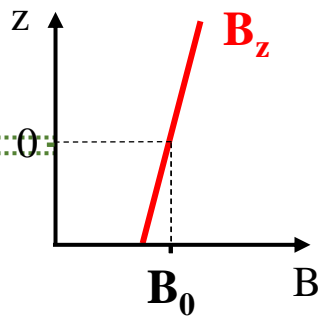
In absence of PFG B_z is constant along z-axis

Slice and shift selection



$$\nu_i \propto B_z$$

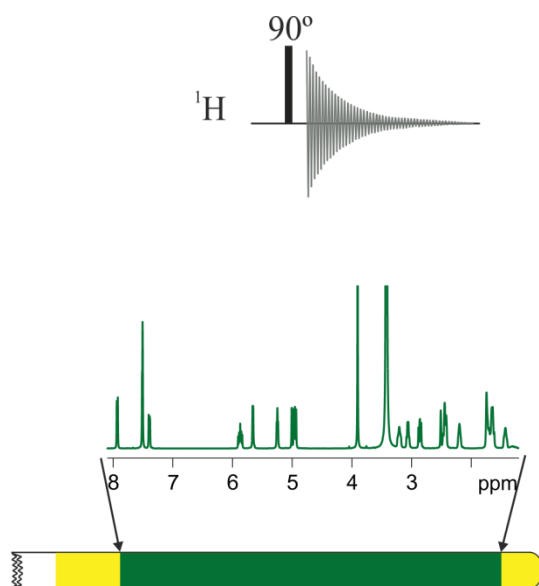
$$B_z = B_0 + zG$$



In presence of PFG B_z varies linearly along z-axis

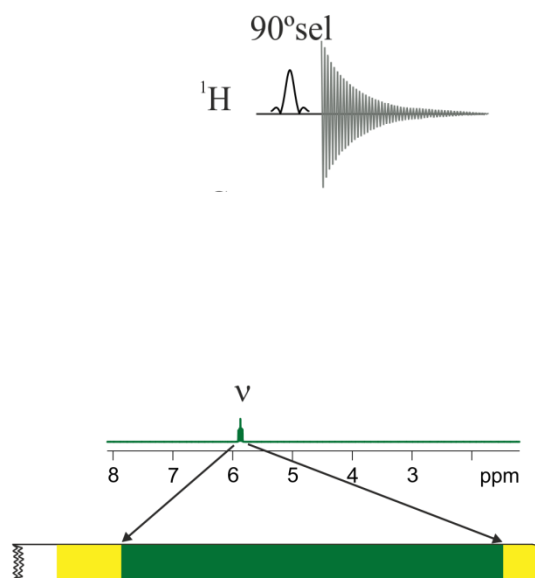
Broadband excitation – shift selection – band selection

Broadband



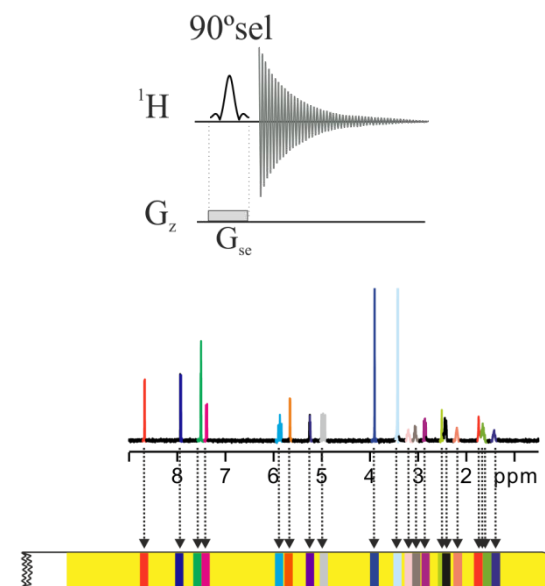
All frequencies are excited in the whole active volume

Shift selection



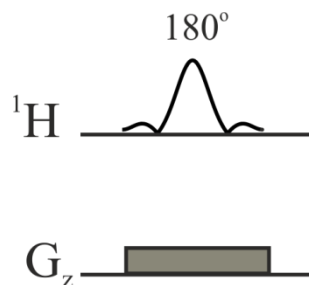
A narrow range of frequencies is excited in the whole active volume

Slice and shift selection



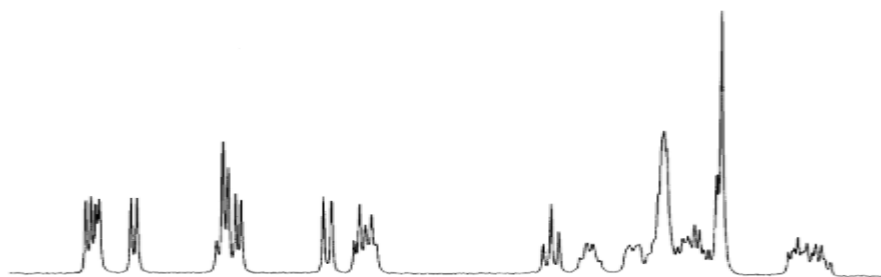
Different frequencies are excited in different parts of the active volume

Zangger-Sterk (ZS)

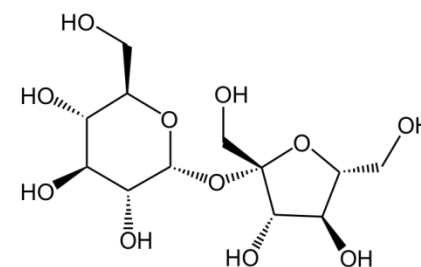
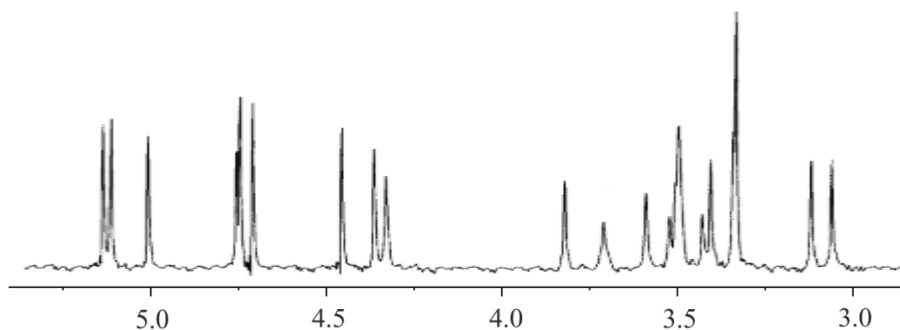


- Simultaneous application of a selective 180° pulse and a weak pulse field gradient
- Slice and shift selection
- Each active spin is excited in a narrow region (slice) of the sample
- Coupled spins cannot be refocused within the same slice
- Broadband pure shift spectrum

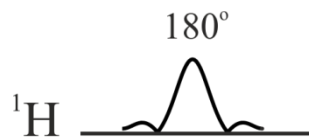
^1H NMR



ZS pure shift

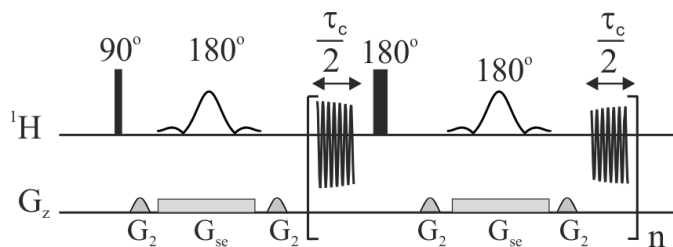


Zangger-Sterk pure shift methods

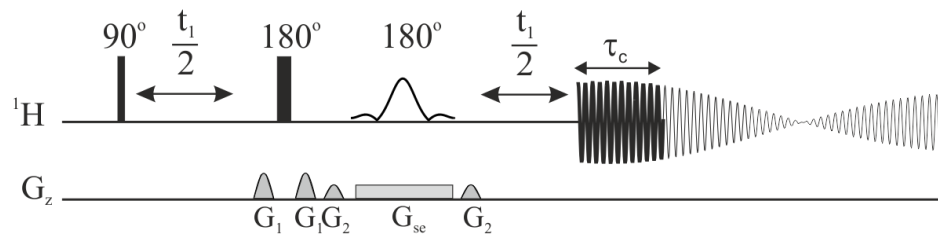


- Compatible with both real-time and interferogram acquisition

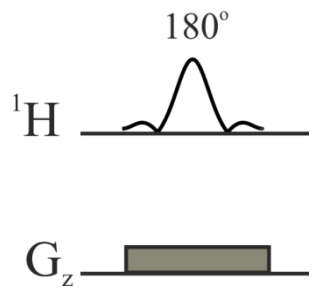
Real-time ZS experiment



Interferogram ZS experiment



Sensitivity in Zangger-Sterk pure shift spectra



- *Active* spins only come from a slice
- Low sensitivity
- Proportional to the slice thickness

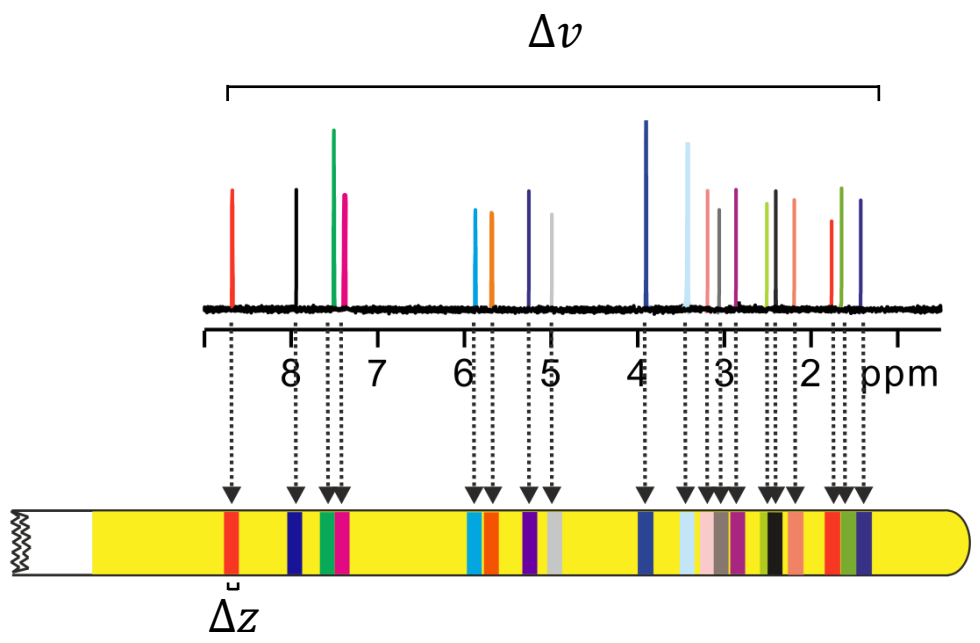
Slice thickness:

$$\Delta z = \frac{\Delta\omega}{\gamma G}$$

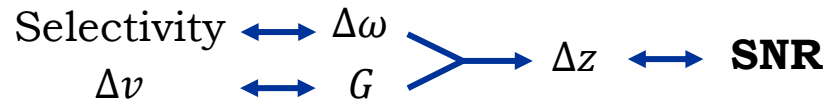
Gradient strength:

$$G = \frac{\Delta\nu}{L\gamma}$$

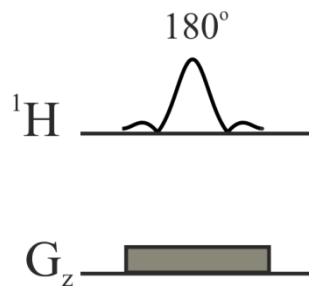
Bandwidth of the selective pulse ($\Delta\omega$) dictated by the separation of the closest coupled spins



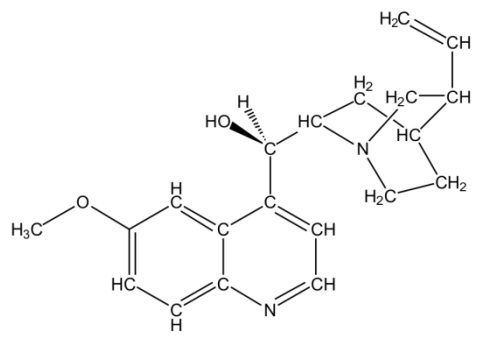
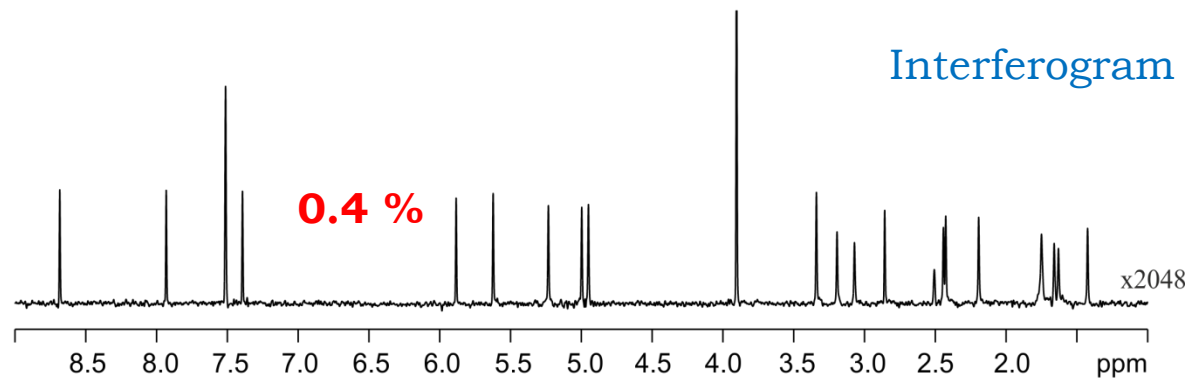
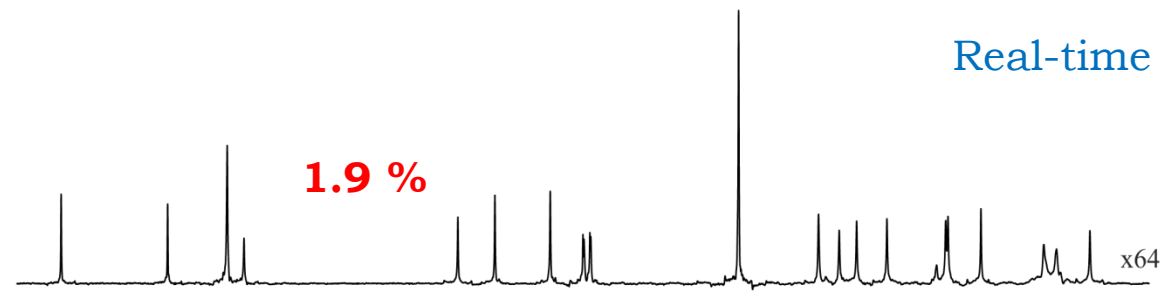
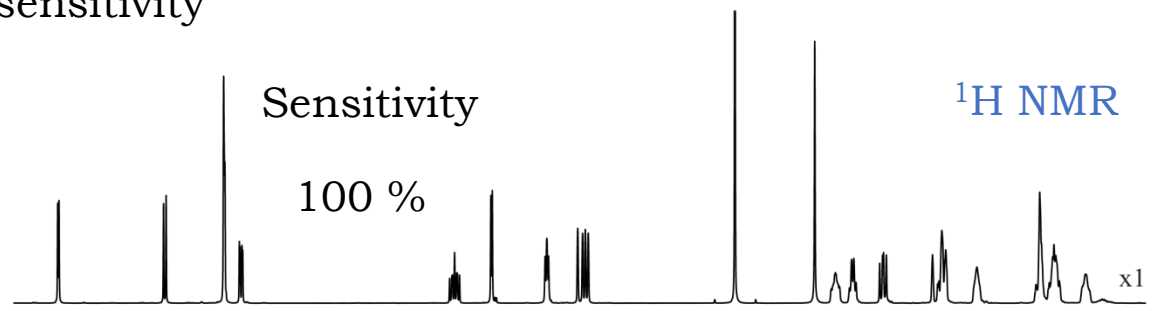
Selectivity and spectral range required will determine the sensitivity



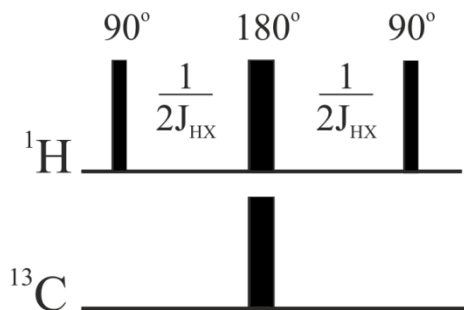
Sensitivity in Zangger-Sterk pure shift spectra



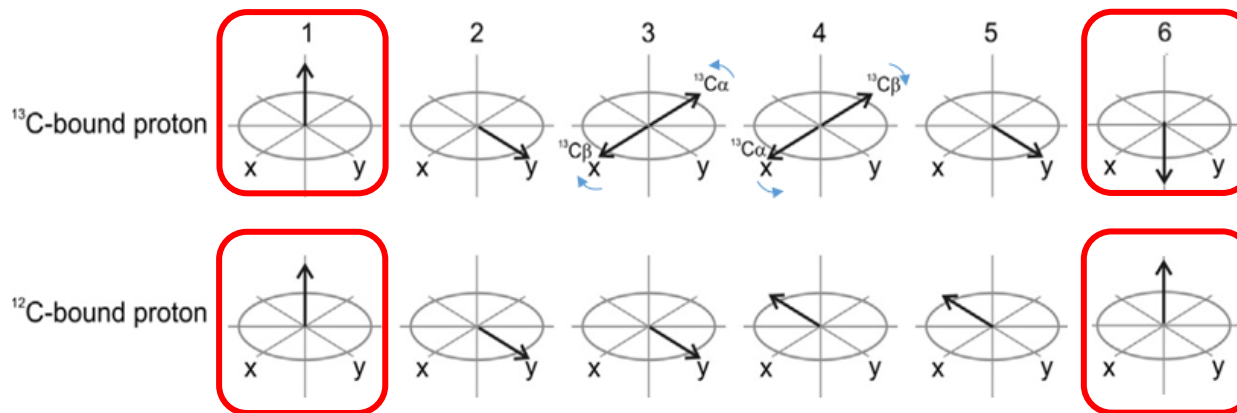
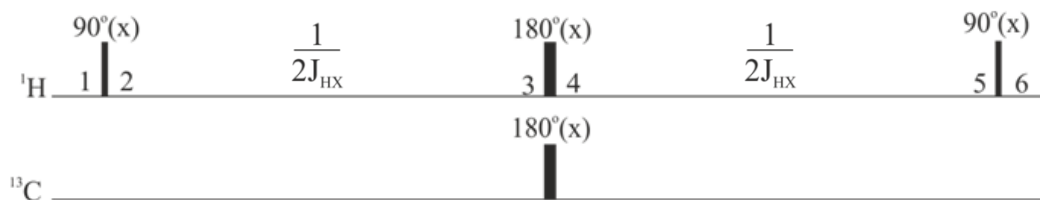
• Low sensitivity



Bilinear rotation decoupling (BIRD)



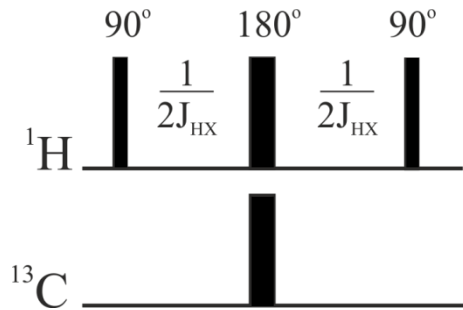
- Protons are *active* if attached to ^{13}C (or ^{15}N) and *passive* if not
- BIRD^x inverts only protons directly coupled to ^{13}C (or ^{15}N)



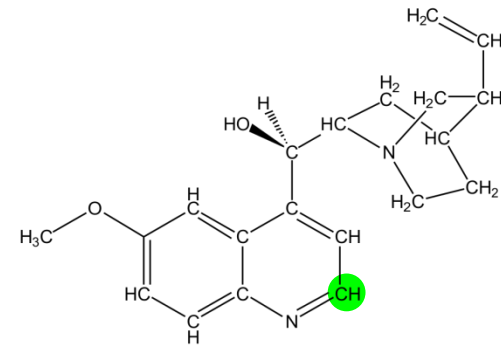
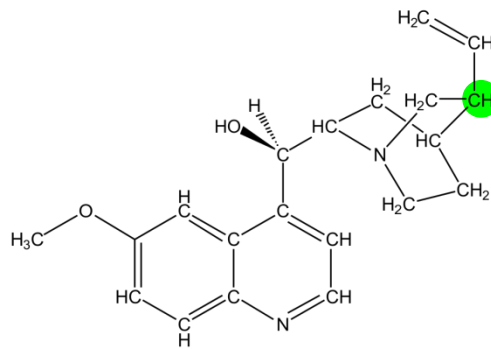
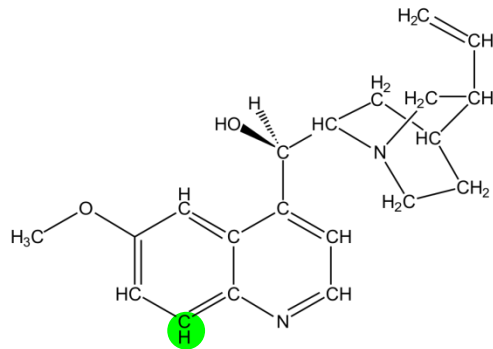
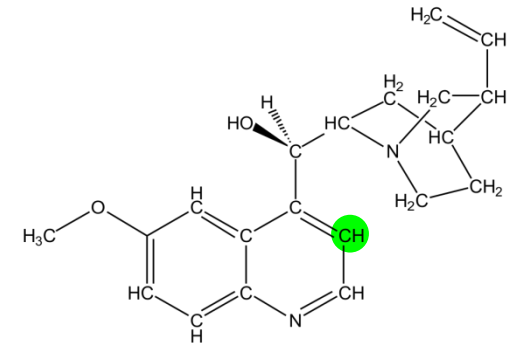
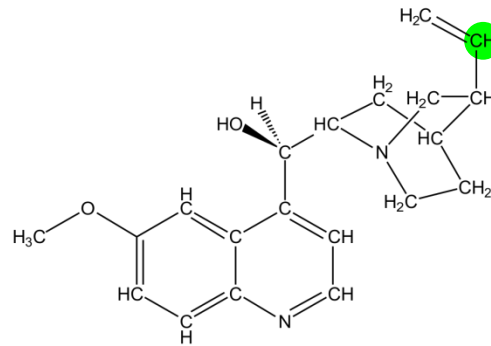
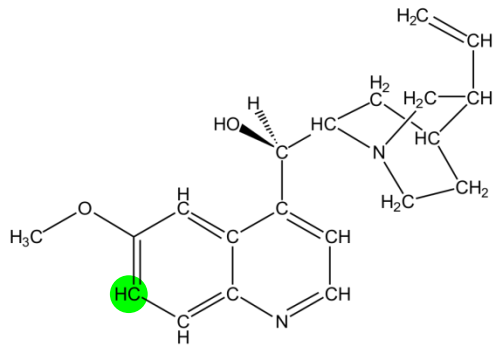
Active spins
inverted

Passive spins
“unaffected”

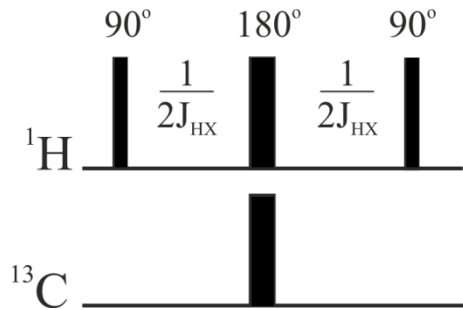
Bilinear rotation decoupling (BIRD)



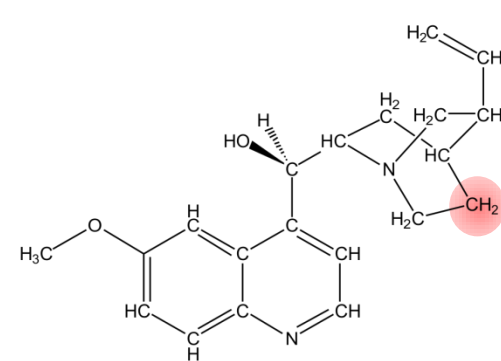
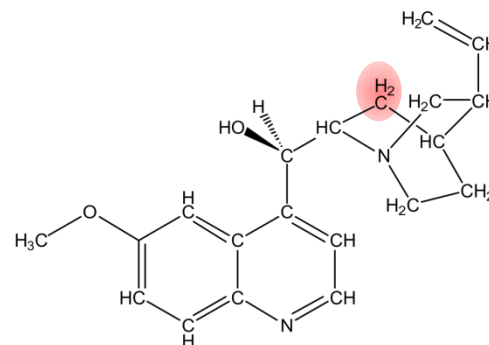
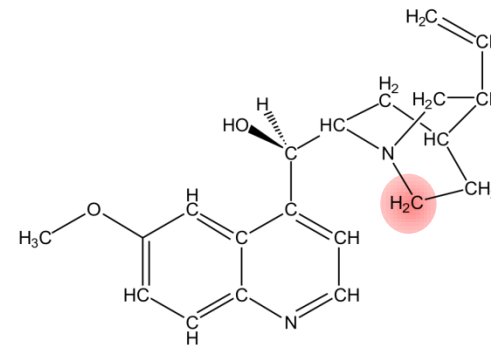
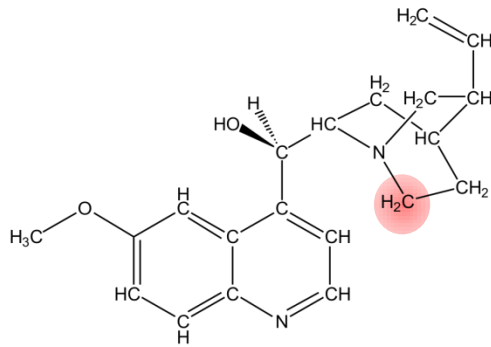
- Protons are *active* if attached to ^{13}C (or ^{15}N) and *passive* if not
- BIRD^x inverts only protons directly coupled to ^{13}C (or ^{15}N)
- Isotopic dilution ensures that their coupled partners are not inverted



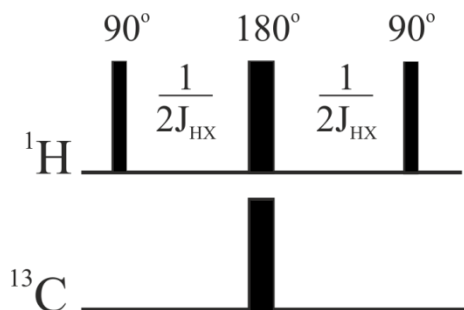
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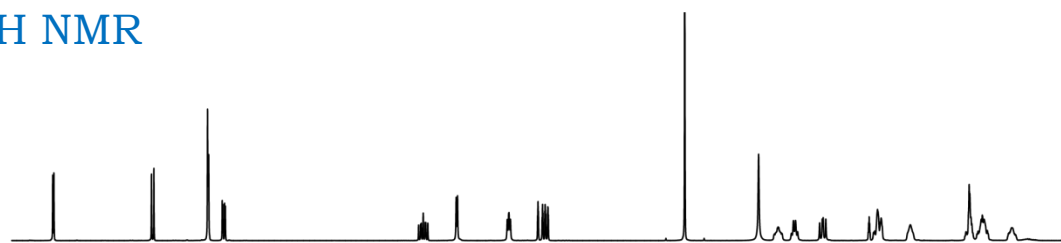


Bilinear rotation decoupling (BIRD)

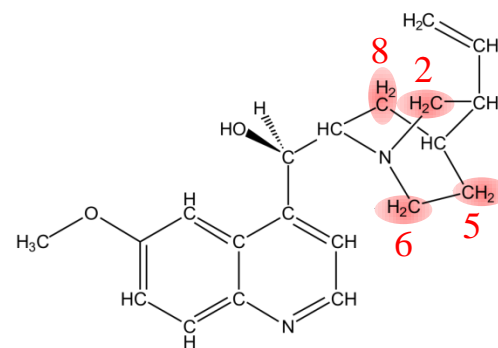
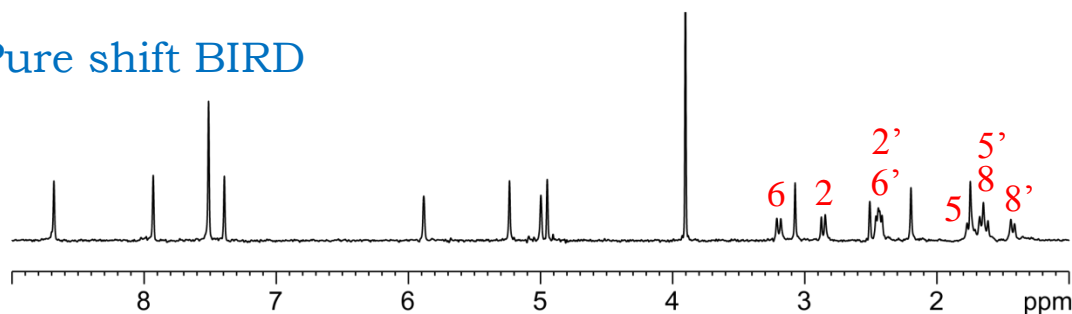


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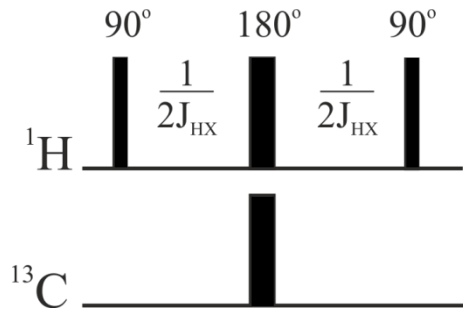
^1H NMR



Pure shift BIRD

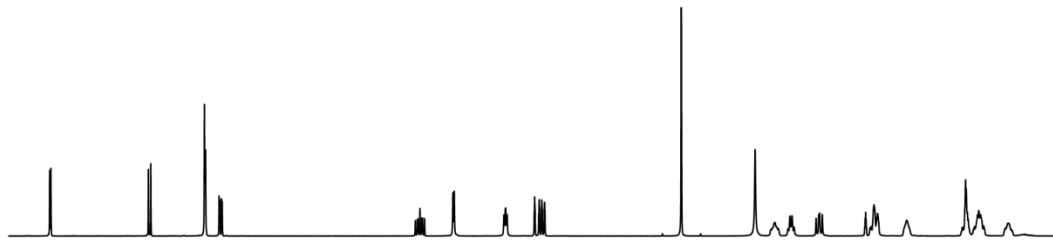


Sensitivity in BIRD pure shift spectra



- Compatible with both real-time and interferogram acquisition
- Low sensitivity
- Limited by the natural abundance at 1.1 % ^{13}C (0.37 % in ^{15}N)

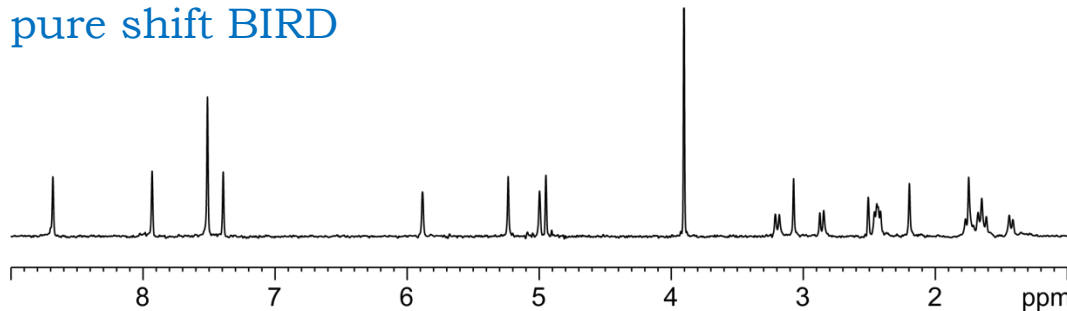
^1H NMR



Sensitivity

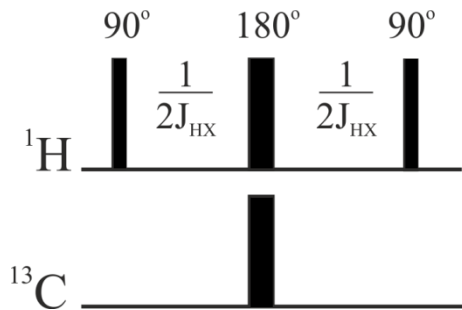
100 %

Real-time pure shift BIRD



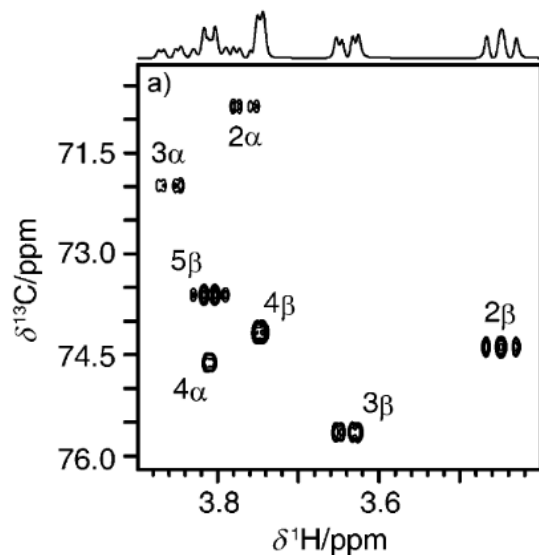
0.8 %

Sensitivity in BIRD pure shift spectra



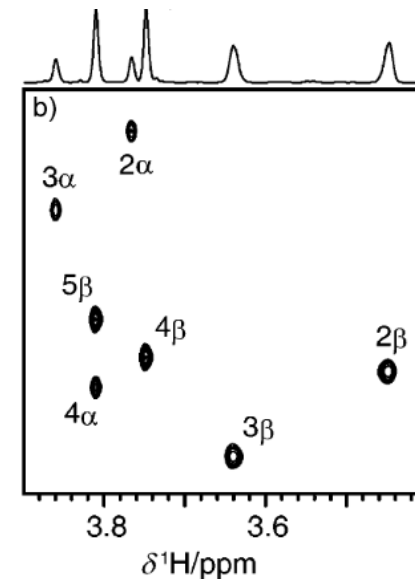
- Compatible with both real-time and interferogram acquisition
- Low sensitivity
- Limited by the natural abundance at 1.1 % ^{13}C (0.37 % in ^{15}N)
- No sensitivity penalty in HSQC experiments

Conventional ^1H - ^{13}C HSQC

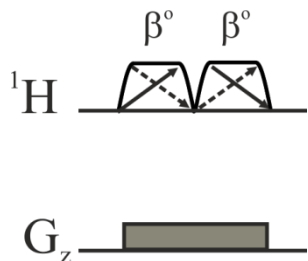


Real-time pure shift ^1H - ^{13}C HSQC

Sensitivity
100 % **180 %**

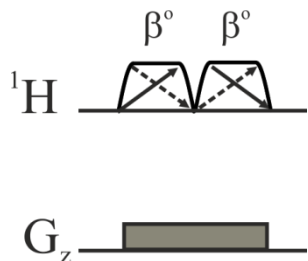


Pure shift yield by chirp excitation (PSYCHE)



- Simultaneous application of a low-flip angle (β) saltire chirp pulse and a weak pulse field gradient
- The refocussing of the J_{HH} couplings is achieved by the simplification of coupling patterns using low flip angle pulses (similar to anti-z-COSY experiment)
- Only a fraction $\sin 2\beta$ of spins (*active spins*) are “refocused”

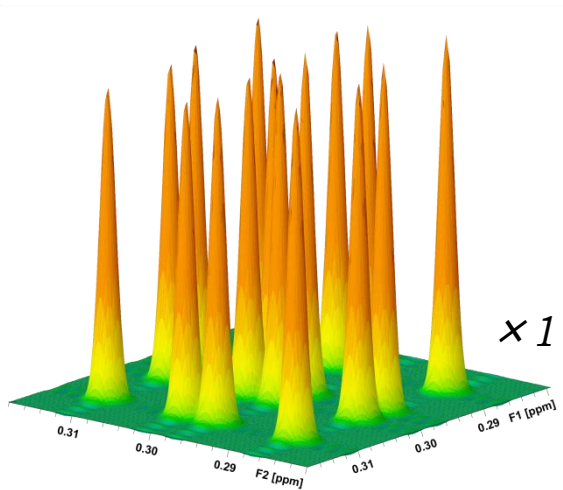
Pure shift yield by chirp excitation (PSYCHE)



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- The refocussing of the J_{HH} couplings is achieved by the simplification of coupling patterns using low flip angle pulses (similar to anti-z-COSY experiment)
- Only a fraction $\sin 2\beta$ of spins (*active spins*) are “refocused”

The effect of flip angle

$$\beta = 90^\circ$$



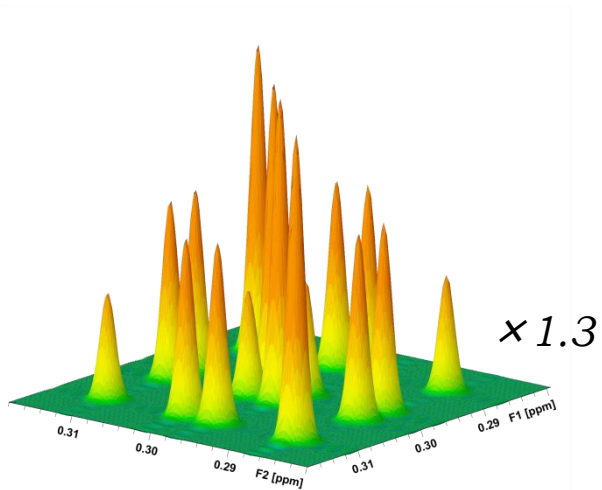
Diagonal: 100 %

Off-diagonal: 100 %

Decoupling factor = 1

The effect of flip angle

$$\beta = 60^\circ$$



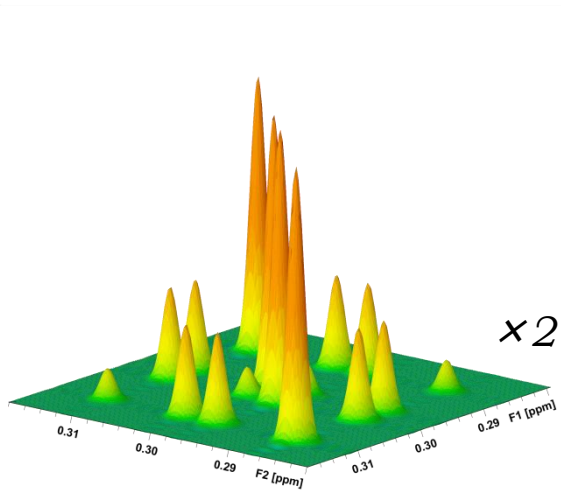
Diagonal: 60 %

Off-diagonal: 36 %

Decoupling factor = 1.7

The effect of flip angle

$$\beta = 45^\circ$$



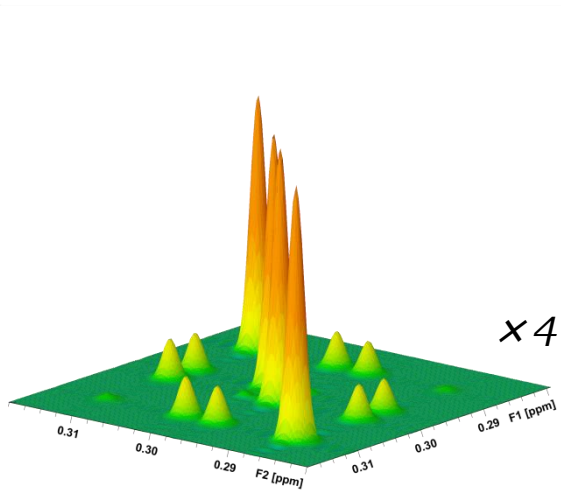
Diagonal : 33.3 %

Off-diagonal : 11.1 %

Decoupling factor = 3

The effect of flip angle

$$\beta = 30^\circ$$



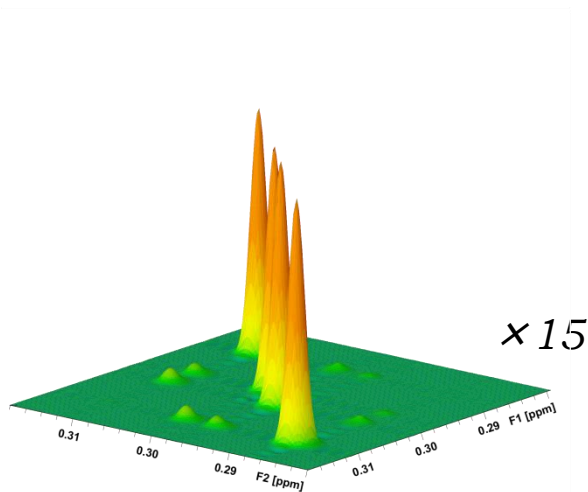
Diagonal : 14.3 %

Off-diagonal : 2 %

Decoupling factor = 7

The effect of flip angle

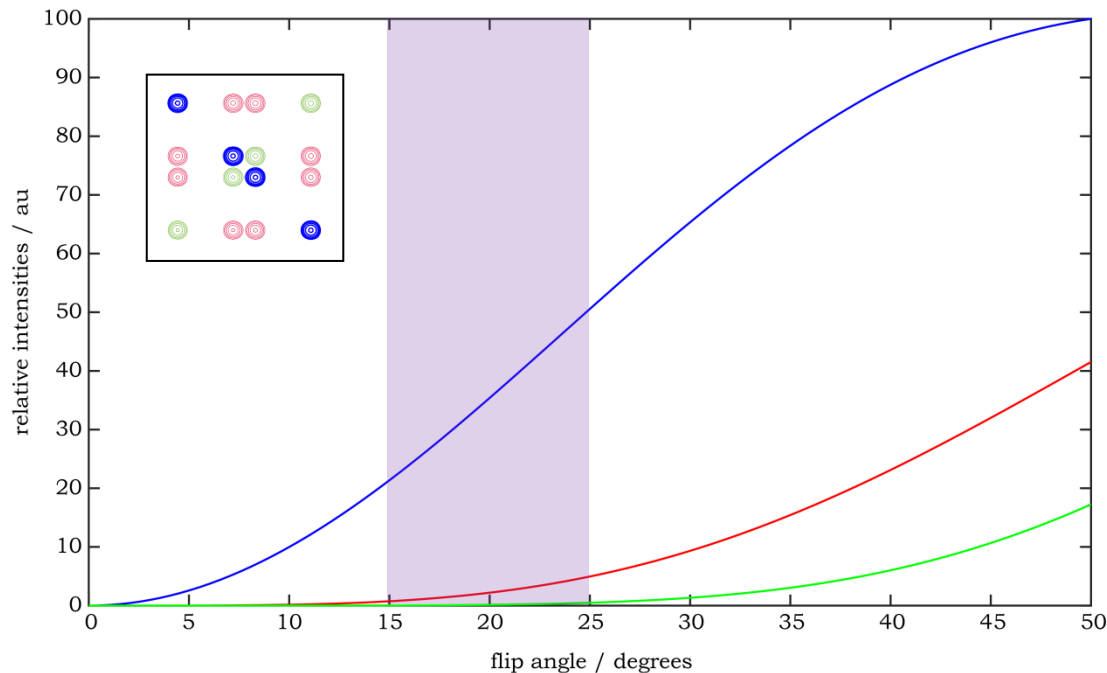
$$\beta = 15^\circ$$



Diagonal : 3.5 %

Off-diagonal : 0.12 %

Decoupling factor = 28.6



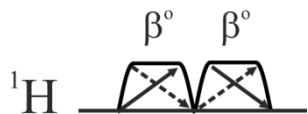
Signal of interest

- pure shift signal

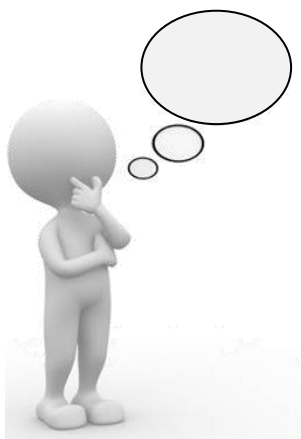
Unwanted signals

- Recoupling artefacts

Pure shift yield by chirp excitation (PSYCHE)



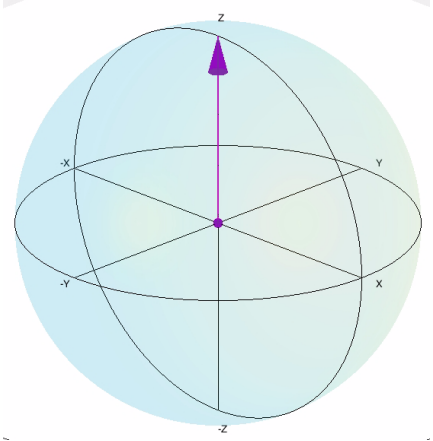
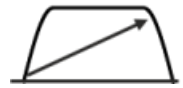
- Simultaneous application of a low-flip angle (β) **chirp pulse** and a weak pulse field gradient
- The refocussing of the J_{HH} couplings is achieved by the simplification of coupling patterns using low flip angle pulses (similar to anti-z-COSY experiment)
- Only a fraction $\sin 2\beta$ of spins (*active spins*) are “refocused”



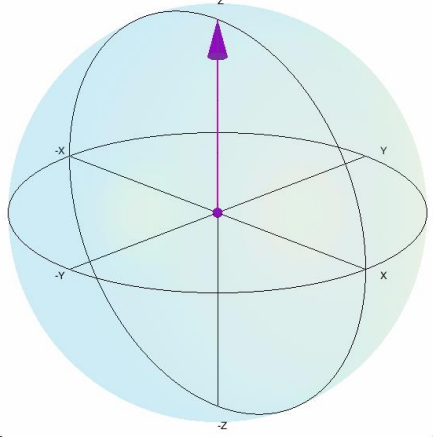
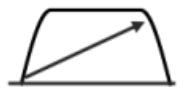
Swept-frequency (chirp) pulses

Generally they sweep the frequency across the spectral window in a linear fashion

180° chirp



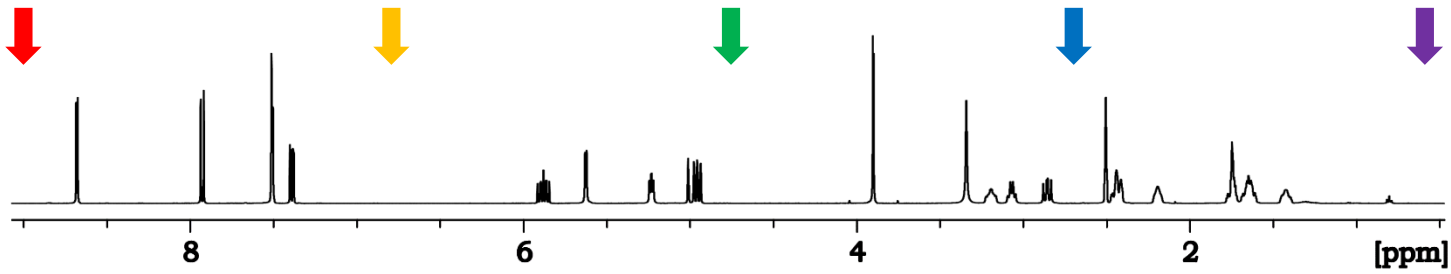
Low-flip angle chirp



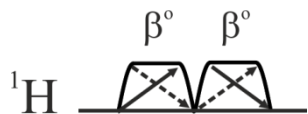
Low frequency

On-resonance

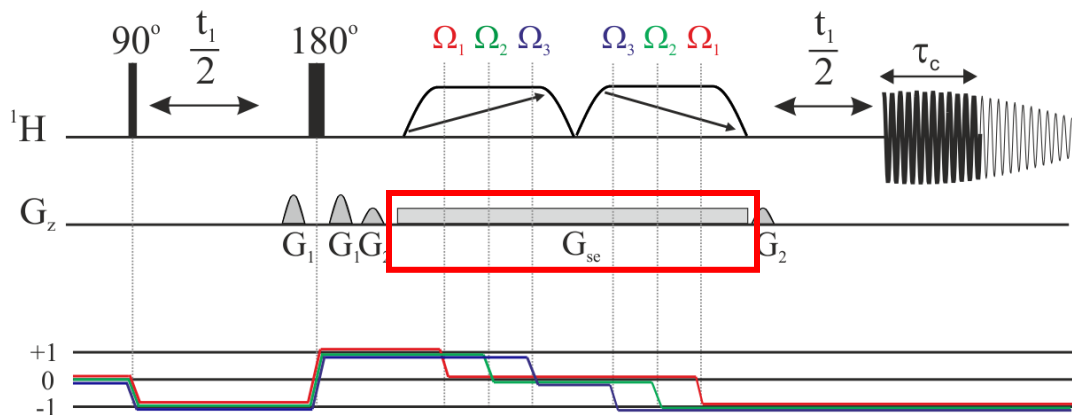
High frequency



Pure shift yield by chirp excitation (PSYCHE)



- Simultaneous application of a low-flip angle (β) saltire chirp pulse and a weak pulse field gradient
- The refocussing of the J_{HH} couplings is achieved by the simplification of coupling patterns using low flip angle pulses (similar to anti-z-COSY experiment)
- Only a fraction $\sin 2\beta$ of spins (*active spins*) are “refocused”



Signal of interest

- pure shift signal

Unwanted signals

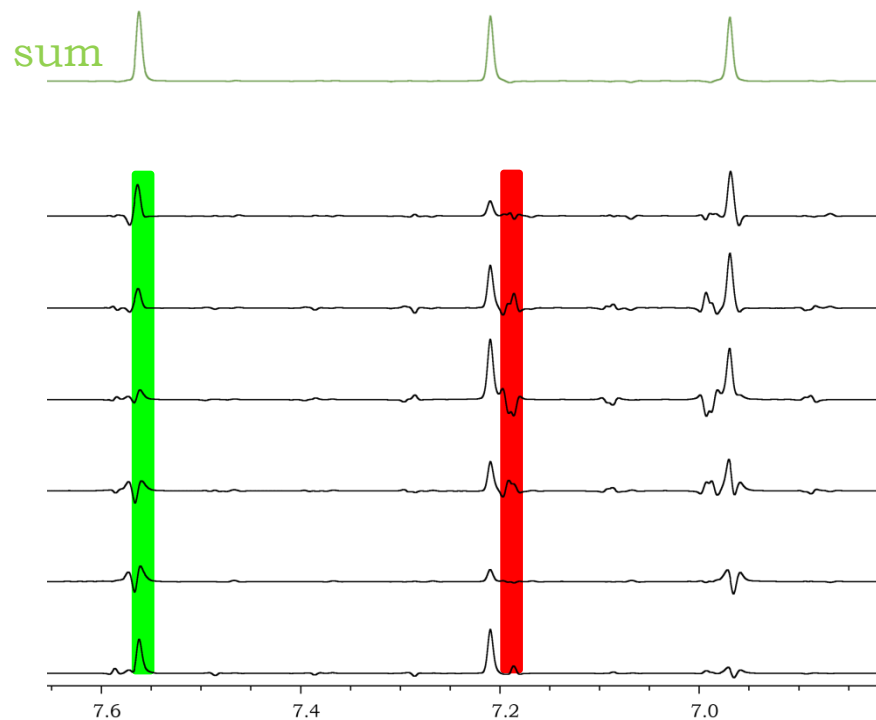
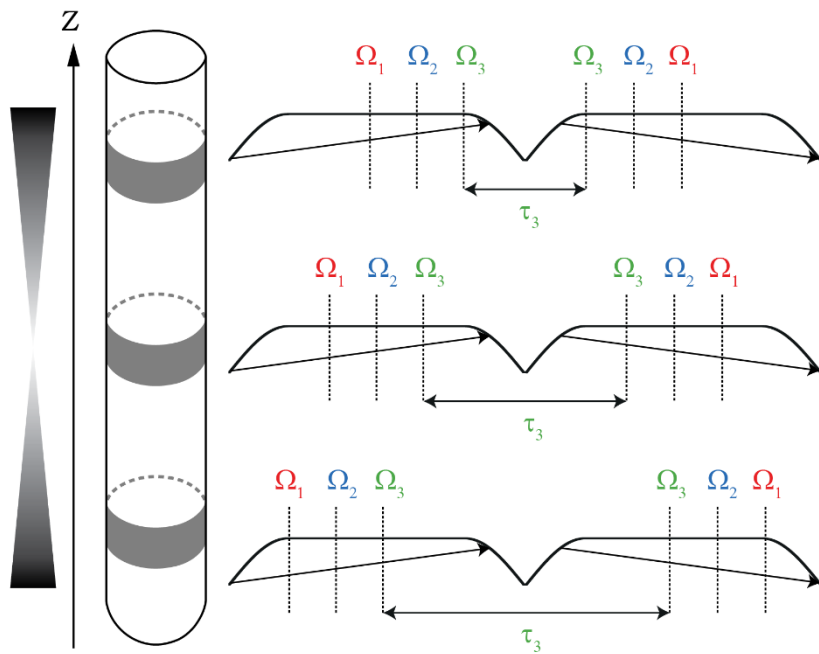
- Recoupling artefacts
- Zero Quantum signals
- CO signals
- Strong coupling signals

Spatial-temporal averaging

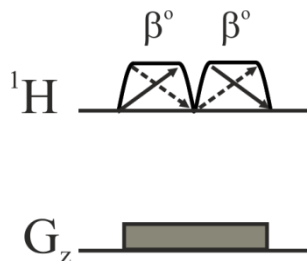
Spatio-temporal averaging

- Wanted signals (pure shift singlets) have time-invariant phase
- Unwanted signals have time-dependent phases

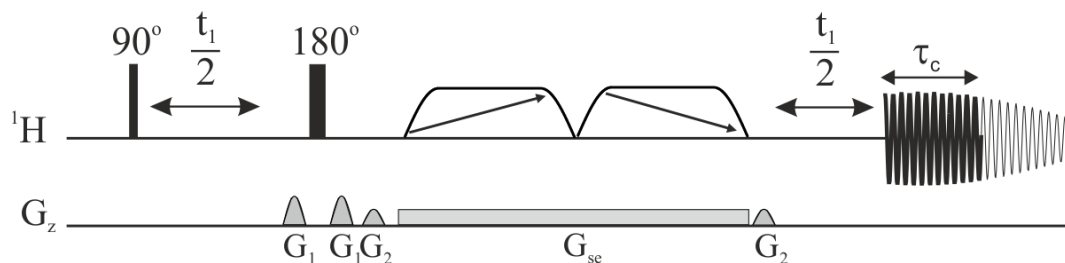
$$B(z) = B_0 + G_z(z - z_0)$$



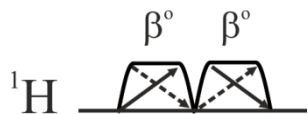
Pure shift yield by chirp excitation (PSYCHE)



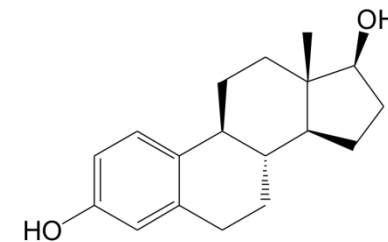
- Simultaneous application of a low-flip angle (β) saltire chirp pulse and a weak pulse field gradient
- The refocussing of the J_{HH} couplings is achieved by the simplification of coupling patterns using low flip angle pulses (similar to anti-z-COSY experiment)
- Only a fraction $\sin 2\beta$ of spins (*active spins*) are “refocused”
- Only compatible with interferogram acquisition (statistical excitation)



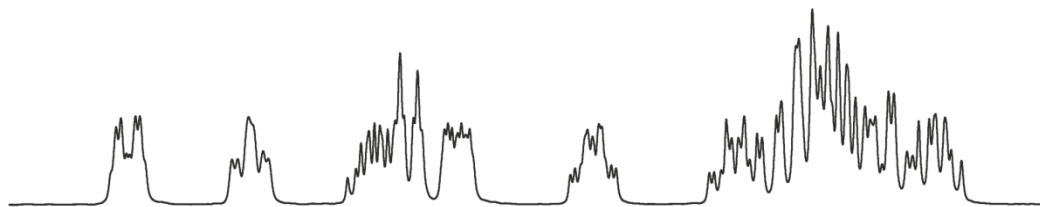
Sensitivity in PSYCHE pure shift spectra



- Sensitivity depends on the fraction of spins (*active* spins) that have been “refocused”
- It depends on the low-flip angle: $\sin 2\beta$



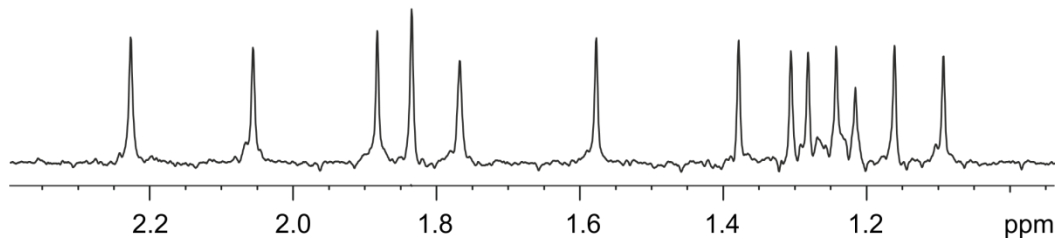
^1H NMR



Sensitivity

100 %

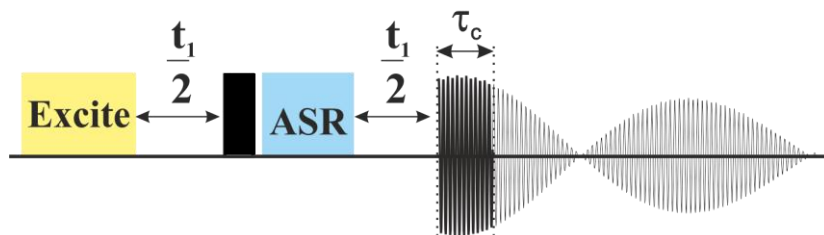
Interferogram pure shift PSYCHE



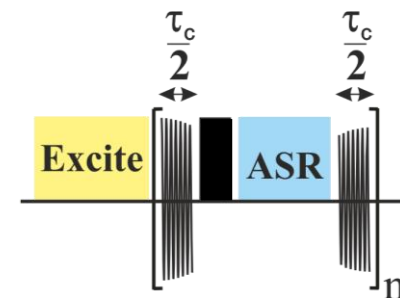
3.8 %

How can we combine all these elements?

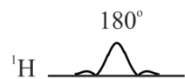
Interferogram



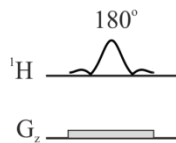
Real-time



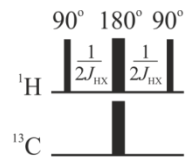
BS



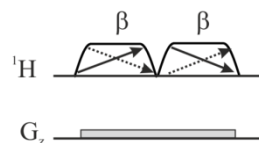
ZS



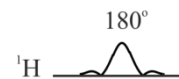
BIRD



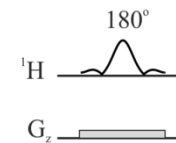
PSYCHE



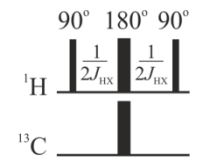
BS



ZS

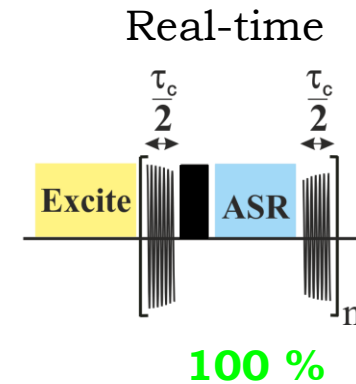
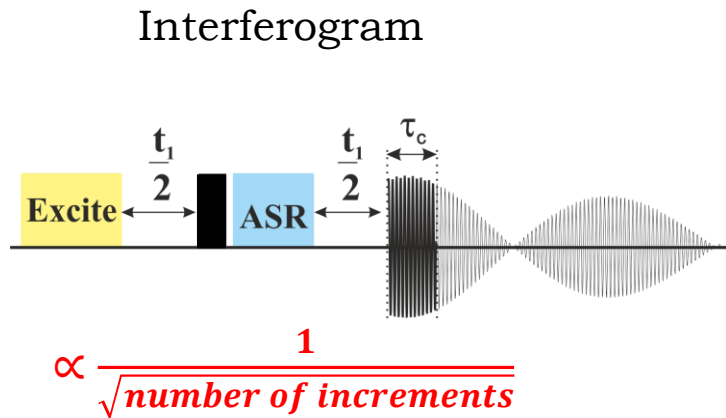


BIRD



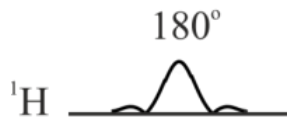
General overview: sensitivity of each element

Acquisition mode



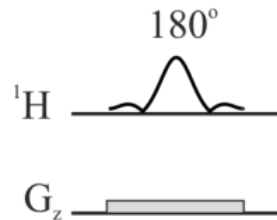
Active spin refocusing methods

BS



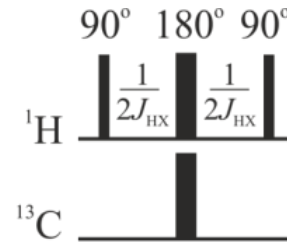
$\geq 100\%$

ZS



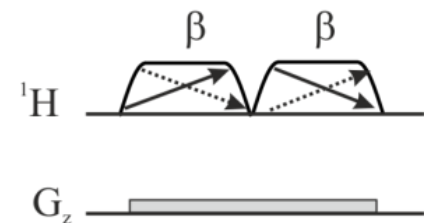
0.5 - 10 %

BIRD



1.1 %

PSYCHE



3 - 20 %

Introduction

Setting the scene

Key concepts

J-refocussing

Chunking acquisition

ASR elements

Interferogram

Real-time

Acquisition methods

Band-selective

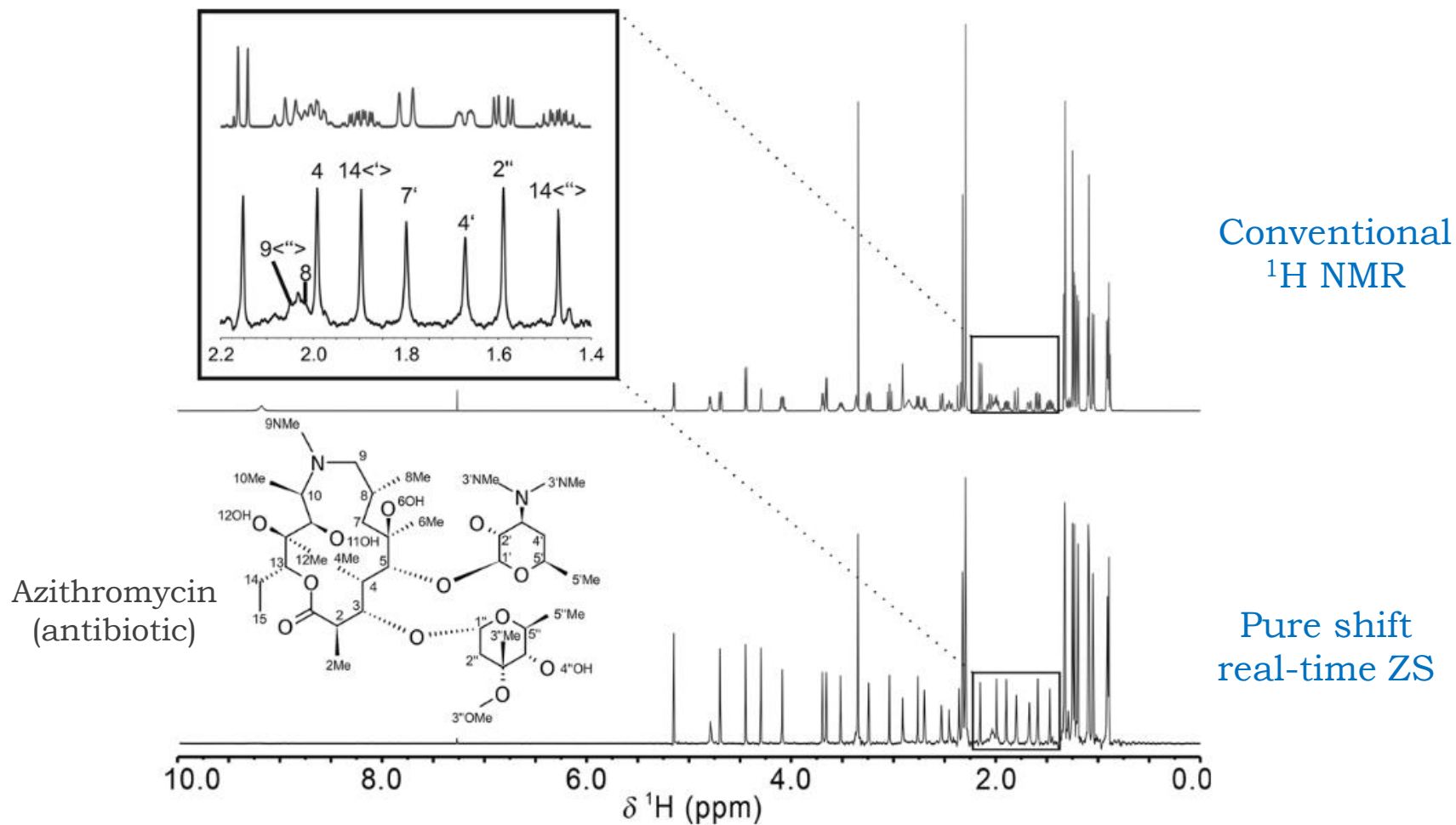
Zangger-Sterk

BIRD

PSYCHE

Applications

1D pure shift NMR experiments for structure analysis of small and medium sized molecules



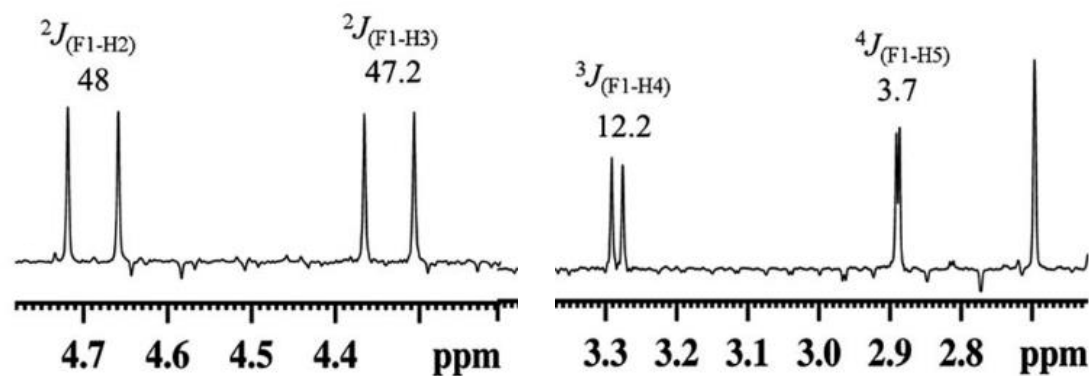
Conventional
 ^1H NMR

Pure shift
real-time ZS

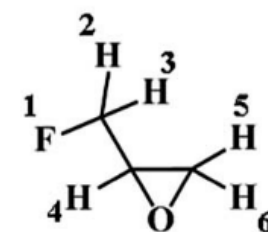
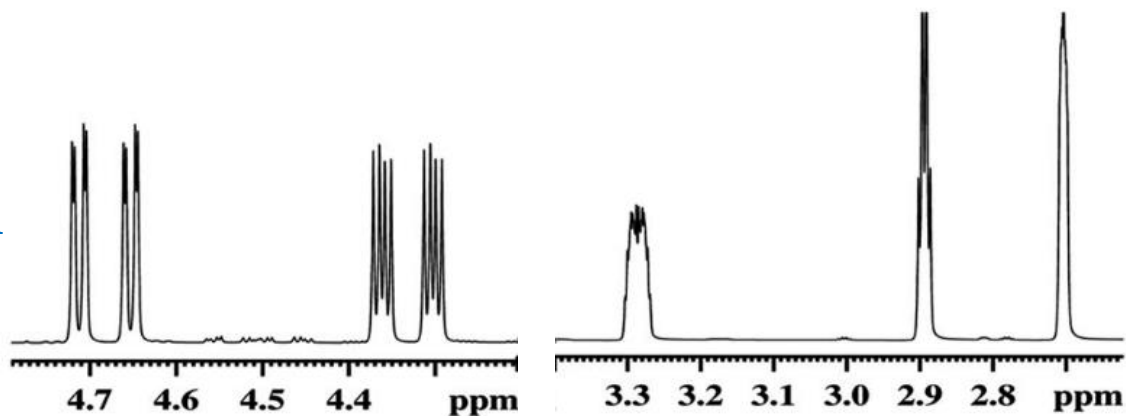
Azithromycin
(antibiotic)

1D Pure shift NMR experiments for fast and accurate extraction of heteronuclear coupling constants

Pure shift
interferogram
ZS

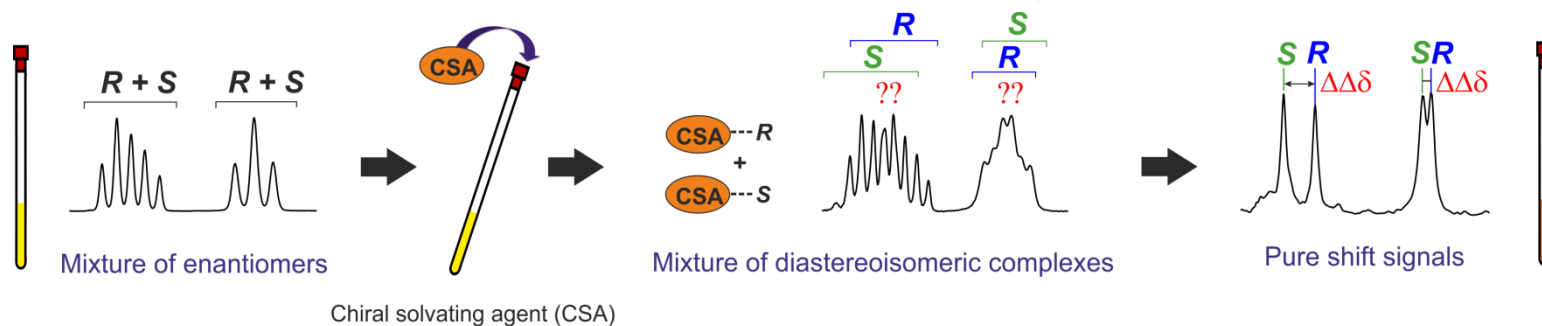


Conventional
 ^1H NMR



Epifluorohydrin

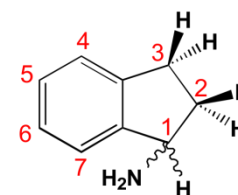
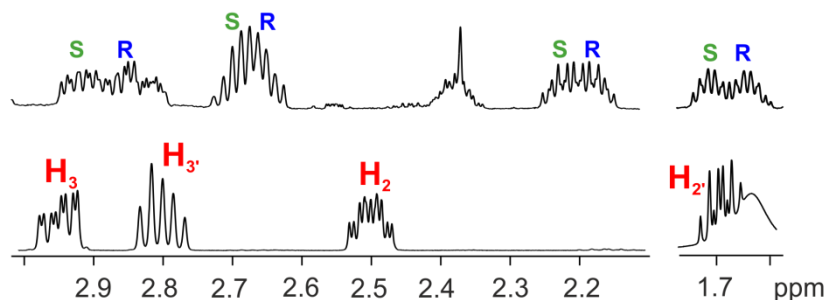
1D pure shift NMR experiments for enantiomer and diastereomer studies



Pure shift
real-time BS



Conventional
 ^1H NMR

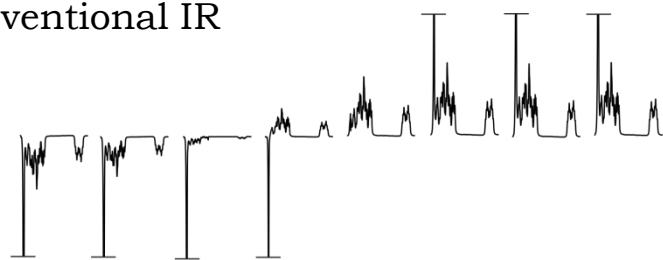


1-aminoindan
(racemic mixture)

Pure shift NMR experiments to measure relaxation times in overlapped regions

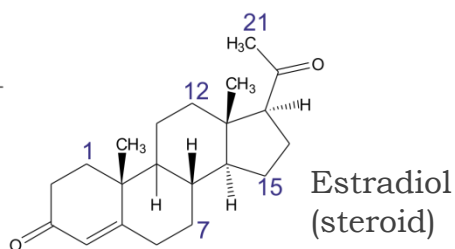
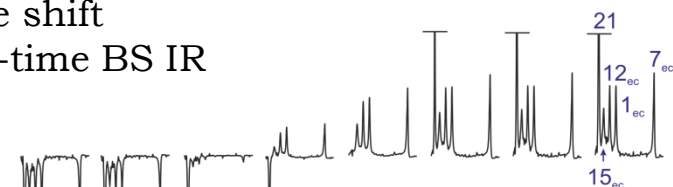
Inversion-Recovery: T_1 measurement

Conventional IR



τ 0.05s 0.1s 0.25s 0.5s 1s 2s 4s 8s

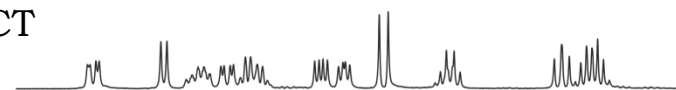
Pure shift
real-time BS IR



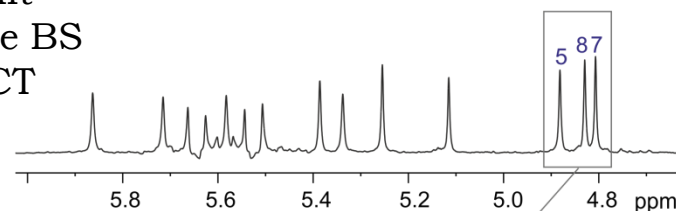
Estradiol
(steroid)

PROJECT: Determining T_2 measurement

Conventional
PROJECT



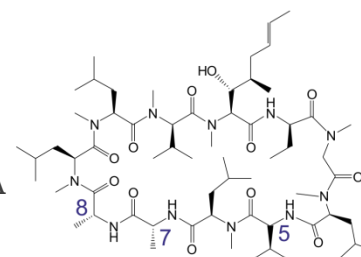
Pure shift
real-time BS
PROJECT



12ms 66ms 126ms 156ms 246ms 306ms 456ms 606ms 906ms 1.2s 1.5ms

Total echo time (τ_e)

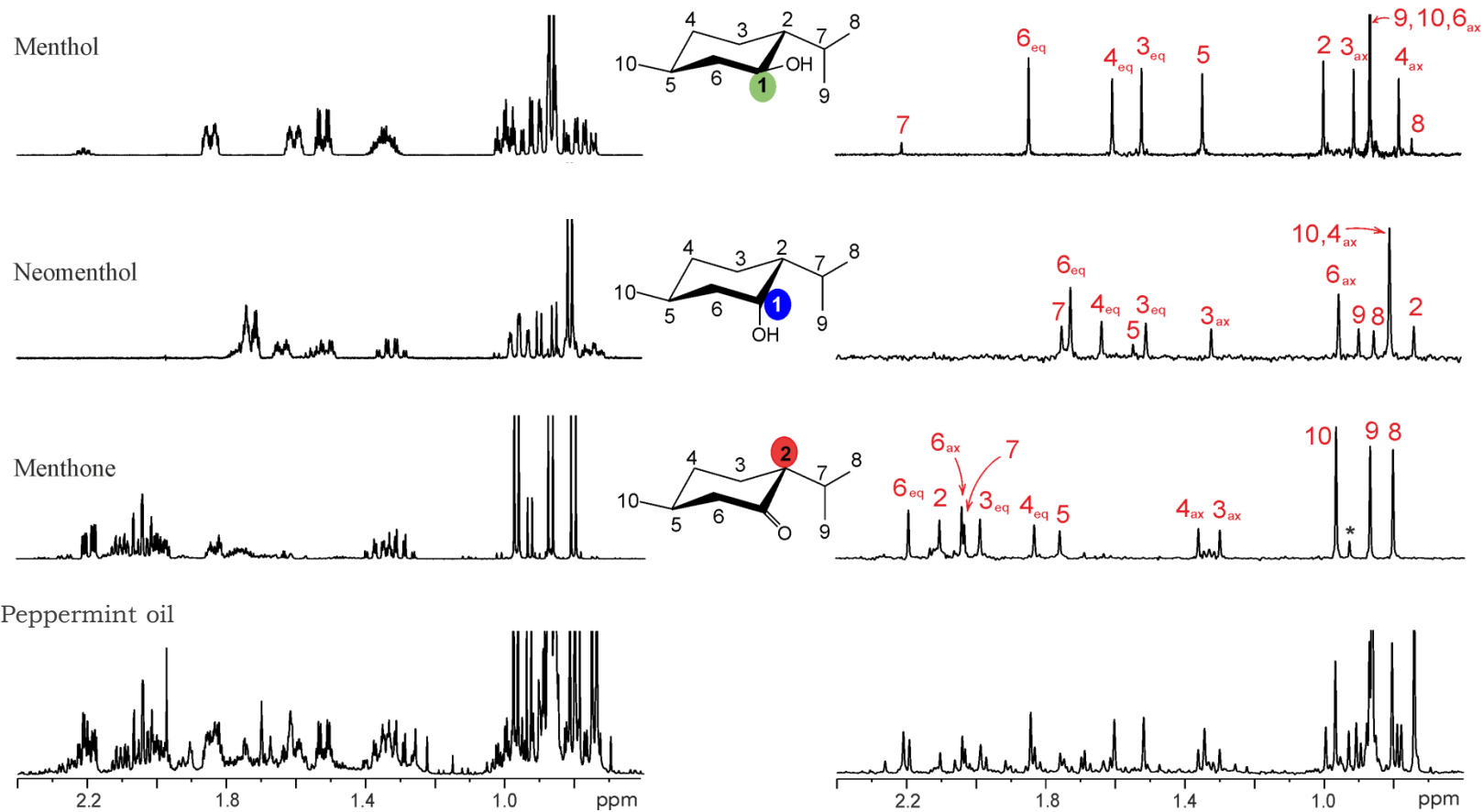
Cyclosporine A
(peptide)



1D pure shift NMR experiments for the study of complex mixtures

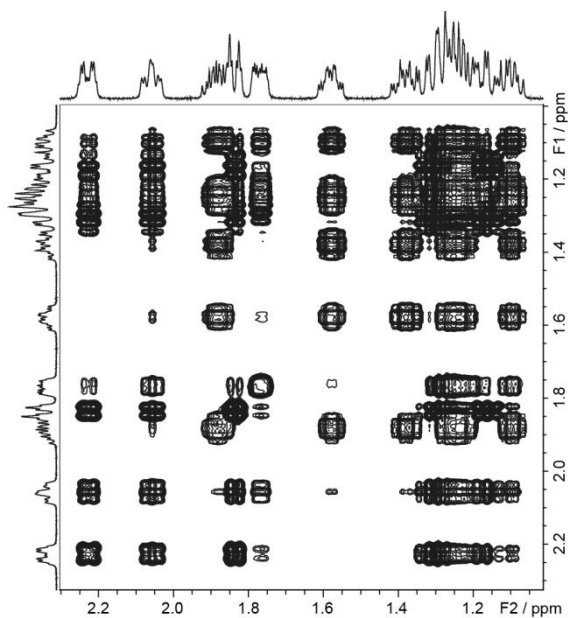
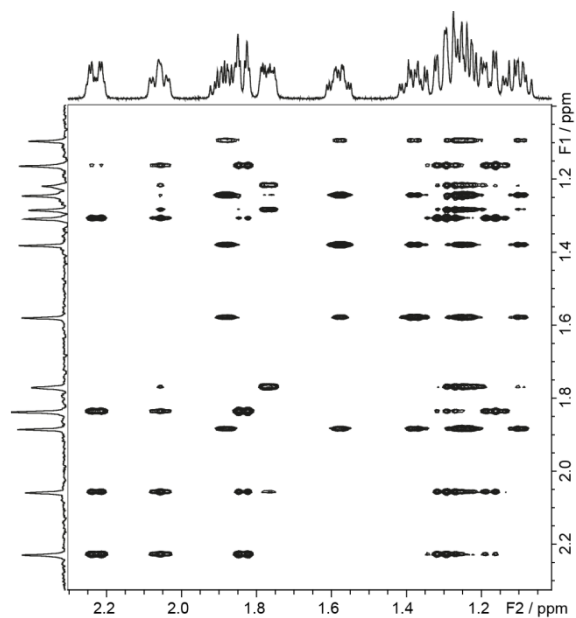
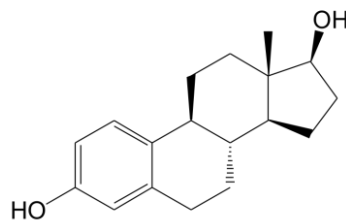
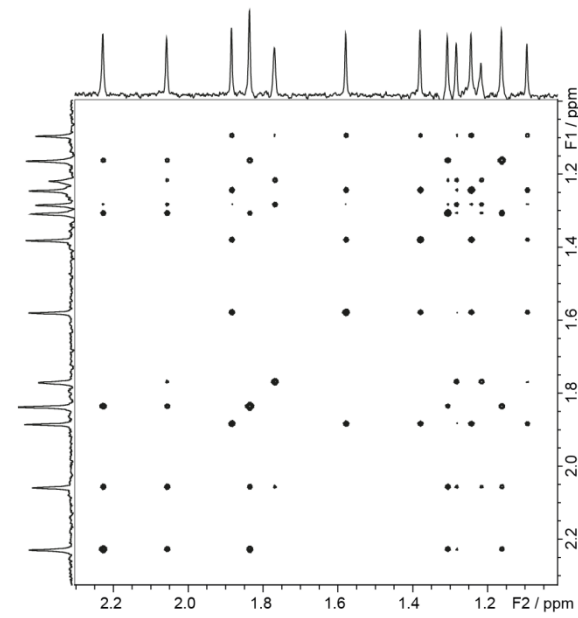
Conventional selective 1D TOCSY

Pure shift interferogram selective 1D PSYCHE-TOCSY



Homonuclear 2D pure shift NMR experiments for structure analysis of small and medium sized molecules

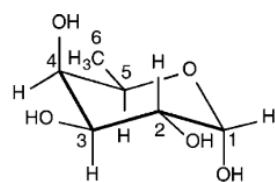
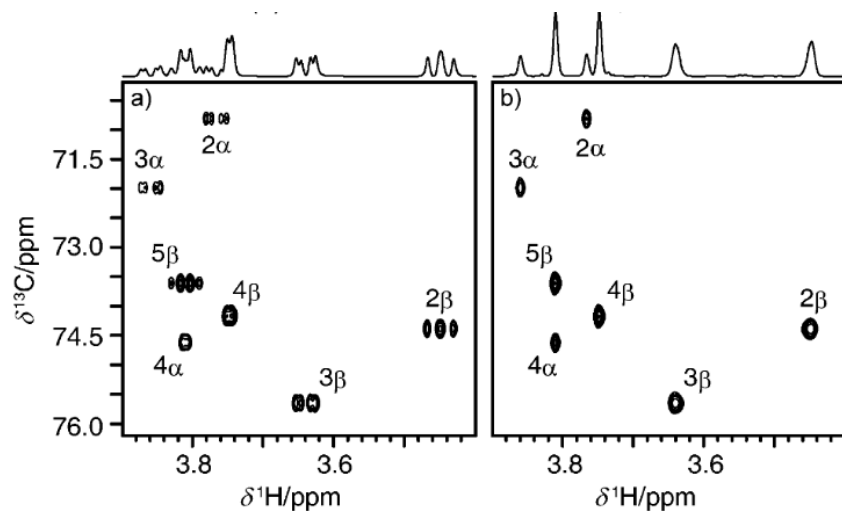
Conventional TOCSY

Pure shift F_1 -PSYCHE TOCSYPure shift F_1 -PSYCHE TOCSY and indirect covariance processingEstradiol
(steroid)

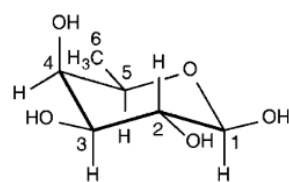
Heteronuclear 2D pure shift NMR experiments for structure analysis of small and medium sized molecules

Conventional
 ^1H - ^{13}C HSQC

Pure shift real-time
 ^1H - ^{13}C HSQC



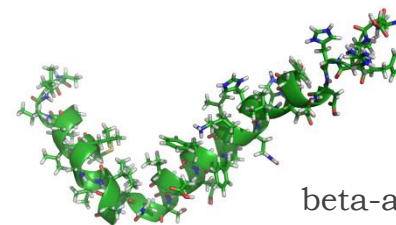
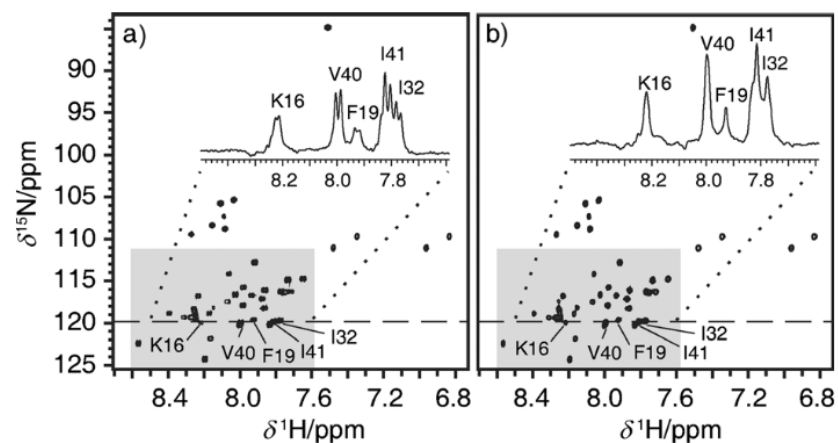
α -D(+)-fucose



β -D(+)-fucose

Conventional
 ^1H - ^{15}N HSQC

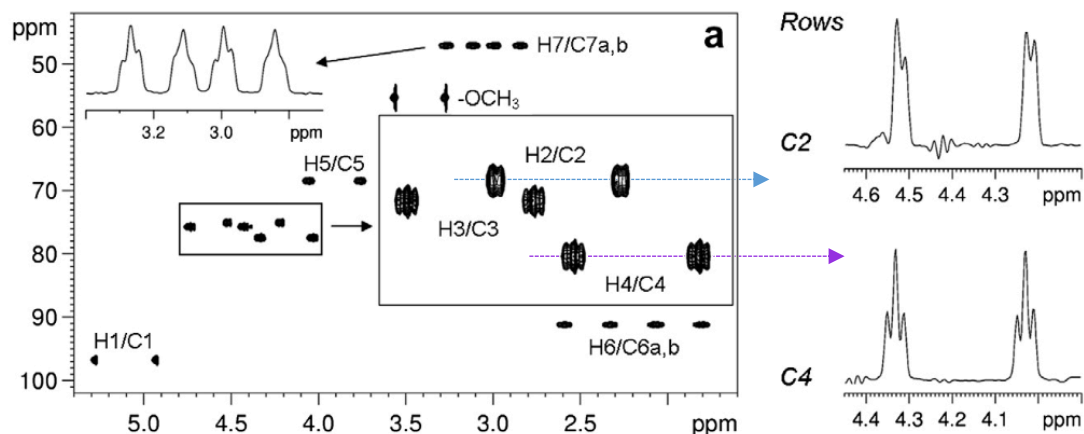
Pure shift real-time
 ^1H - ^{15}N HSQC



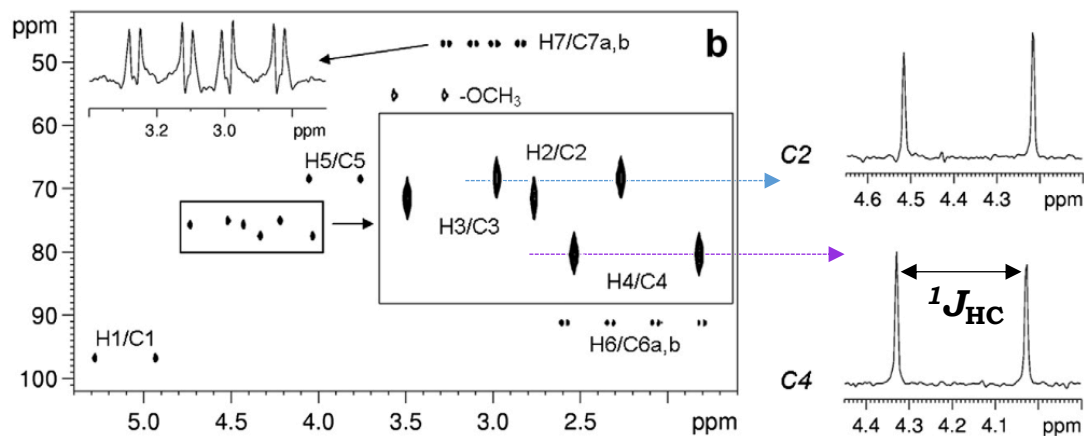
^{15}N -labeled
beta-amyloid peptide 1-42

2D Pure shift NMR experiments for accurate determination of one-bond heteronuclear coupling constants

Conventional
 1H - ^{13}C CLIP-HSQC

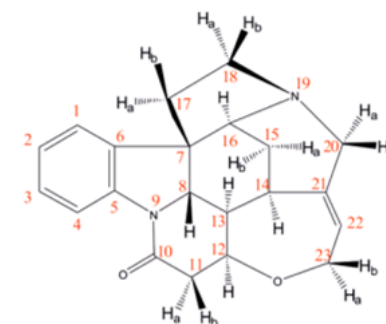
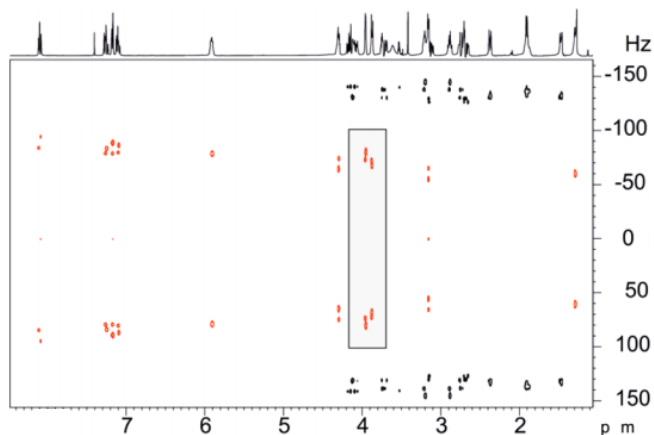


Pure shift
interferogram
 1H - ^{13}C CLIP-HSQC

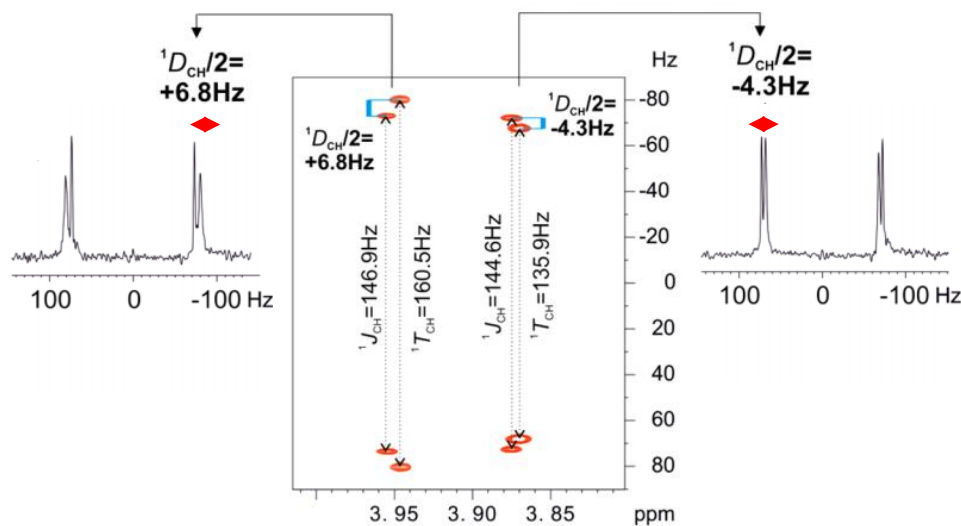


One-shot determination of residual dipolar coupling (RDCs) using pure shift NMR experiments

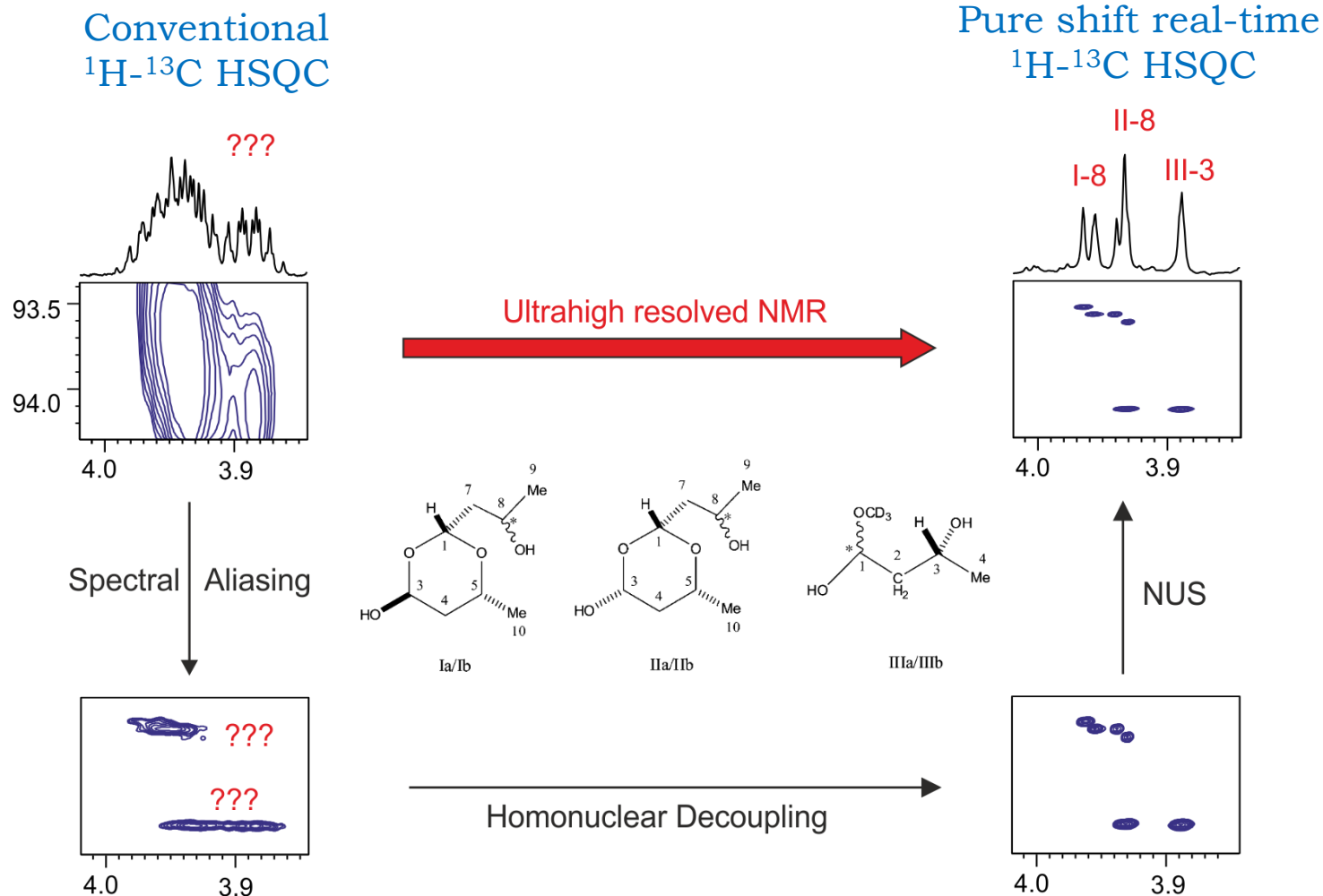
Pure shift
real-time
 J -resolved
 1H - ^{13}C HSQC



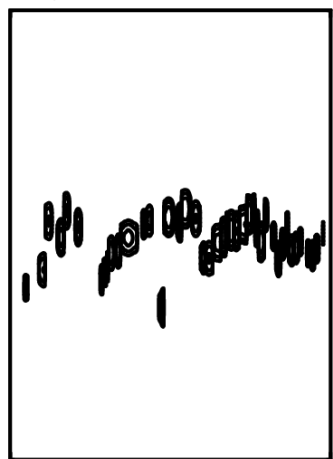
Strychnine



2D pure shift NMR experiments for the study of complex mixtures



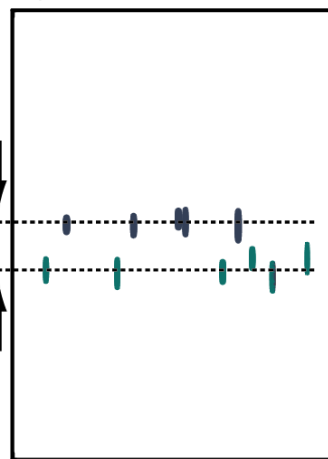
Ultrahigh-resolution diffusion-ordered spectroscopy

Conventional
DOSY

Chemical shift

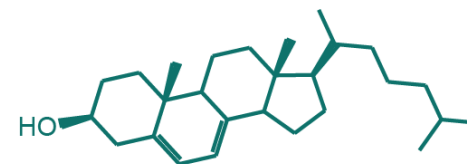
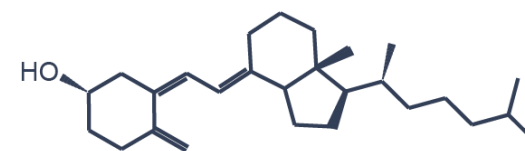
Diffusion

PSYCHE

Pure shift
interferogram
PSYCHE-iDOSY

Chemical shift

Diffusion

Vitamin D₃
(secosteroids)Provitamin D₃
(zoosterol)

Review articles

Pure shift NMR spectroscopy

R. W. Adams, *EMagRes*, **2014**, 3, 295-310

Broadband ^1H homodecoupled NMR experiments: recent developments, methods and applications

L. Castañar and T. Parella, *Magn. Reson. Chem.*, **2015**, 53, 399-426

Pure shift NMR

K. Zangger, *Prog. Nucl. Magn. Reson. Spectrosc.*, **2015**, 86-87, 1-20

Pure shift ^1H NMR: what is next?

L. Castañar, *Magn. Reson. Chem.*, **2017**, 55, 47-53

PSYCHE pure shift NMR spectroscopy

M. Foroozandeh, G. A. Morris, M. Nilsson, *Chem. Eur. J.* **2018**, 24, 13988-14000

Book chapter

High-resolution NMR techniques in organic chemistry

T. D. W. Claridge, Elsevier Science, 3rd Edition, **2016**, chapter 8.

Example data: download from our website

<http://nmr.chemistry.manchester.ac.uk/pureshift>



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Workshop on pure shift NMR

Copies of slides for the talks given at the *Workshop on pure shift NMR*, Manchester, 12th Sept 2017 can be accessed via this [link](#).

A [data archive](#) containing pure shift pulse sequences, processing software and sample experimental data is available for download via this [link](#).

Workshop on pure shift NMR - slides

Gareth Morris - **Welcome, introduction and history** - [pdf](#) - [pptx](#)

Peter Kiraly - **Interferogram and real-time acquisition methods** - [pdf](#) - [pptx](#)

Laura Castañar - **Zangger-Sterk and band-selective methods** - [pdf](#) - [pptx](#)

Mohammadali Foroozandeh - **PSYCHE** - [pdf](#) - [pptx](#) - [zip including avi videos](#)

Ralph Adams - **Other pure shift and related methods** - [pdf](#) - [pptx](#)

Mathias Nilsson - **Practical implementation** - [pdf](#) - [pptx](#)

Adolfo Botana - **JEOL pure shift implementation** - [pdf](#) - [pptx](#)

Vadim Zorin - **Mestrelab pure shift implementation** - [pdf](#) - [pptx](#)

Ēriks Kupče - **Bruker shaped pulse implementation** - [pdf](#) - [pptx](#)

Workshop on pure shift NMR - downloads

Data Archives, including instructions, sequences, parameter files and example data.

Bruker

Software only (< 1 Mb): [Pure_shift_archive_Bruker_software_only.zip](#).

Full (262 Mb): [Pure_shift_archive_Bruker.zip](#).

Varian

Software only(< 1 Mb): [Pure_shift_archive_Varian_software_only.zip](#).

Full (26 Mb): [Pure_shift_archive_Varian.zip](#).

Manual: [UoM_PureShiftNMR_Varian_Manual_rev1.pdf](#).

External Contributions

DIAG package(< 1 Mb): [DIAG_package_Geneva.zip](#).

The Bruker and Varian/Agilent pure shift data and software archives can also be downloaded from [DOI:10.17632/w9nz44cyft.1](https://doi.org/10.17632/w9nz44cyft.1) and [DOI:10.17632/rgj4jwcsnz.1](https://doi.org/10.17632/rgj4jwcsnz.1) respectively

Example data: download from our website

<http://nmr.chemistry.manchester.ac.uk/pureshift>

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Manchester NMR Methodology Group

Home

The NMR methodology group is jointly supervised by [Gareth Morris](#) and [Mathias Nilsson](#), and currently has 13 [members](#). Our [research](#) concerns the development of novel techniques in high resolution NMR spectroscopy, and their application to problems in chemistry, biochemistry, and medicine. In many cases this work leads to new pulse sequences and software tools, some of which are freely available [here](#).

Download from our website:

<http://nmr.chemistry.manchester.ac.uk>

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Pulse Sequences

We are currently preparing many of our pulse sequences, parameter sets, example datasets and processing macros for the website. Some are available [here](#) but if you would like to use any of the other the sequences, as described in the [publications](#) section, please email us. The majority of sequences are available for Varian systems and we are gradually writing the Bruker variants.

The pulse sequences and any macros required for data conversion can be accessed from [this](#) part of the website.

Workshops and presentations

The slides from some of the workshops and presentations given by group members are available from [this](#) part of the website. There is a pure shift NMR package available for download as part of our [2017 workshop on pure shift NMR](#).



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Manchester NMR Methodology Group

Workshops and Conference

Workshops

[UNICAMP 2018 - Introduction to diffusion NMR](#)

[SMASH 2017 - Pure Shift NMR Workshop](#)

[Manchester 2017 - Pure Shift NMR Workshop](#)

[SMASH 2014 - Pure Shift NMR Workshop](#)

Conference tutorials

[EUROMAR 2018 - Pure Shift NMR](#)

Acknowledgments

Jaca 2011



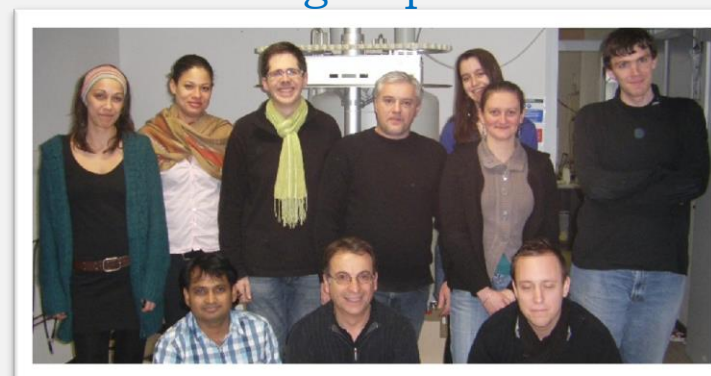
Jaca 2019



SeRMN group – UAB



CEISAM group – Nantes



mancNMR group – UoM



Thank you very much for your attention

A horizontal NMR spectrum is displayed across the middle of the slide. It features a series of peaks of varying heights and widths, characteristic of a complex organic molecule. The spectrum is centered on a horizontal baseline.

laura.castanaraedo@manchester.ac.uk



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