

Diffusion Ordered SpectroscopY

Advanced Operation Training Course



1. DOSY of Theory

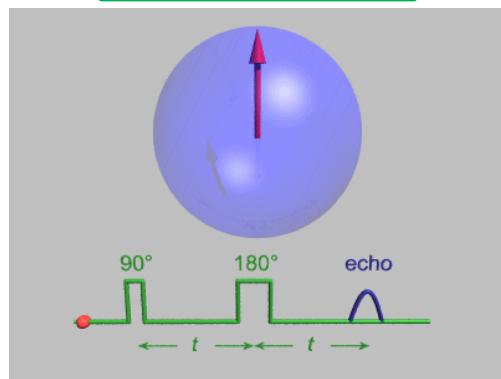


UNIST

ULSAN NATIONAL INSTITUTE OF
SCIENCE AND TECHNOLOGY

Spin echo (Hahn-echo) vs Gradient echo

SE (Spin Echo)

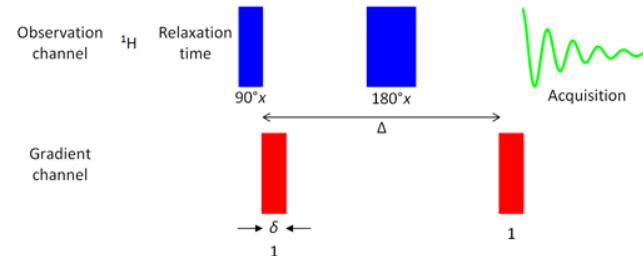


- Dephasing/Refocusing : Related T2
- Experiment time: Long

T2 time을 따라 dephasing이 일어나므로 실험시간이 길어짐

Origin pulse sequence of DOSY

PGSE (Pulsed Gradient Spin Echo)



- Dephasing/Refocusing : Use Gradient
- Experiment time: Middle
- Limited Diffusion Delay (**T2***)
 $T2^* = T2 \text{ in rotating frame}$

Gradient를 이용하여 강제로 Dephasing / Refocusing을 진행하므로 실험시간이 짧음, 대신 $T2^*$ 시간이내로 Diffusion delay가 제한됨

STE (STimulated Echo)

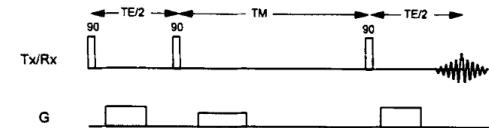


Figure 3 Pulse sequence used to generate a stimulated echo.

- Dephasing/Refocusing : Use Gradient
- Experiment time: Short
- Limited Diffusion Delay (**T1**)

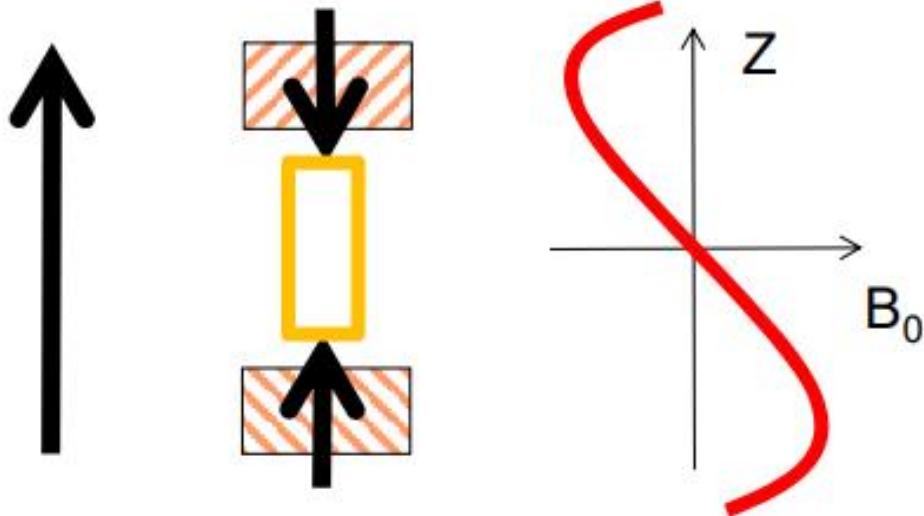
3개의 90도 펄스와 Gradient를 이용하여 T2보다 상대적으로 긴 T1을 Diffusion delay로 사용

What is Gradient?

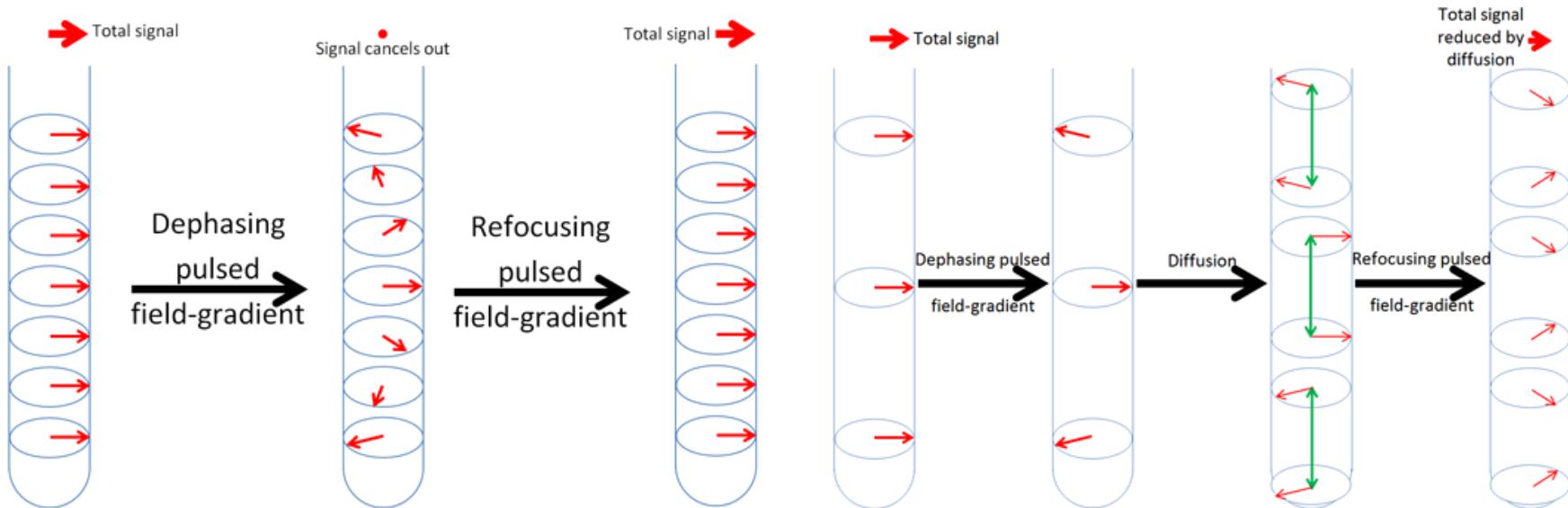
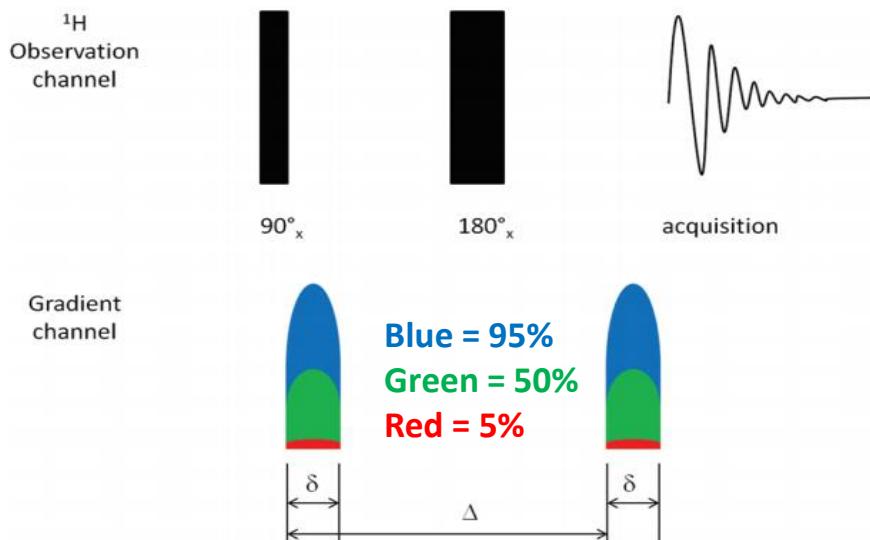
Use Gradient coil -> Make field gradient

- > Force Dephasing / Refocusing for Nuclear spin
- > Save time, Correct good data

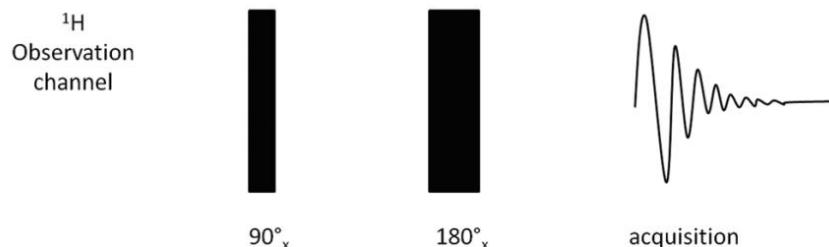
Gradient coil을 이용하여 Magnetic field gradient 형성할 수 있음, 이를 이용 강제적으로 Nuclear spin의 Dephasing / Refocusing을 할 수 있음 -> 실험시간 단축, 정확한 데이터 획득 가능



Pulsed field gradient spin-echo, PGSE

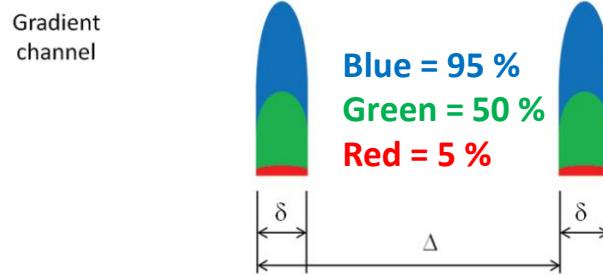


Gradient power vs I value

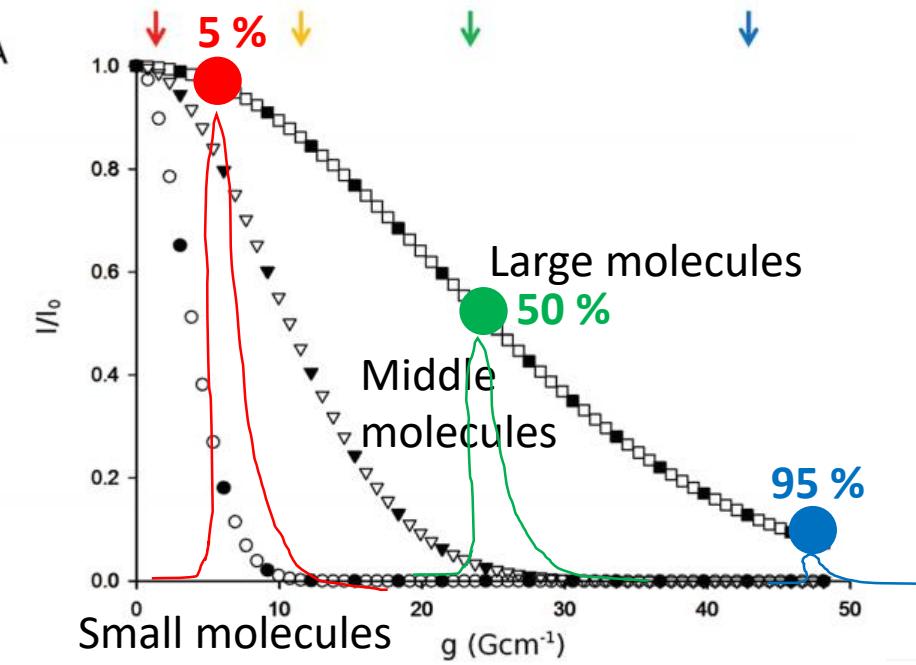
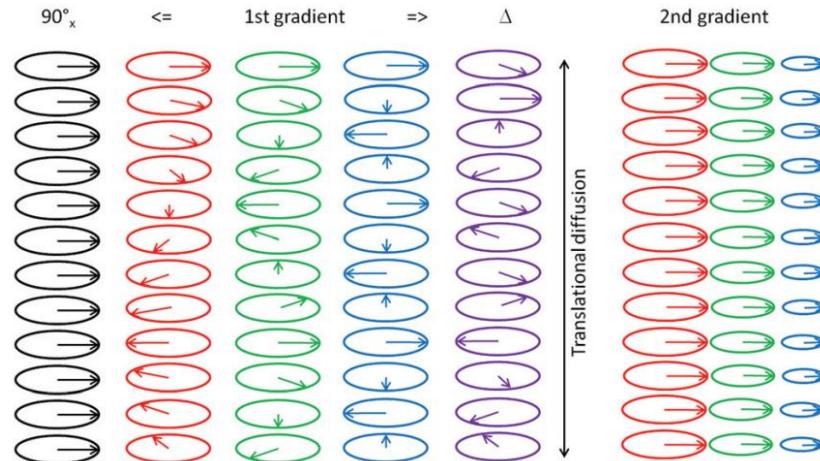


Δ diffusion time
 δ gradient pulse length

Gradient induce dephasing (like grab)
 Powerful grab makes small signal.



Small molecules = Diffuse quickly = Rapidly recovered magnetization
 Large molecules = Diffuse slowly = Slowly recovered magnetization



Equation

$$I/I_0 = e^{(-Dg^2\delta^2\sigma^2g^2\Delta')}$$

I -> Measure DOSY 2D
I₀ -> Measure DOSY 2D
D20 -> Calibrate DOSY 1D
P30 -> Calibrate DOSY 1D

D diffusion coefficient ($\text{m}^2 \text{ s}^{-1}$) Calculation value

g gyro-magnetic ratios of the studied nuclei

σ gradient shape factor

δ gradient pulse lengths Little Delta, **P30**

Δ diffusion time

Δ' diffusion delay Big Delta, **D20**

Peak Intensity
(or integration)
of 95% Gradient power **I**

Peak Intensity
(or integration)
of 5% Gradient power **I₀**

2. DOSY Pulse sequence



UNIST

ULSAN NATIONAL INSTITUTE OF
SCIENCE AND TECHNOLOGY

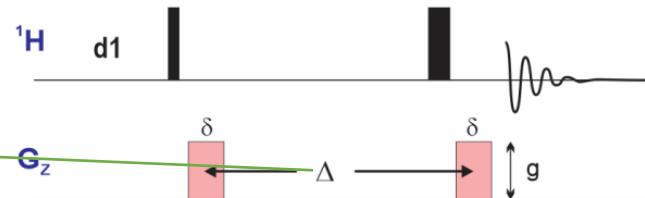
Stimulated Echo (STE)

PGSE (Pulsed field Gradient Spin-Echo)

Disadvantage

1. Poor sensitivity
 - = Related transverse magnetization($T2^*$)
 $(D20 < T2^*)$
2. Chemical shift and J coupling dependence
(180° pulse isn't perfect for sample by sample)

PGSE – Pulsed (field) Gradient Spin-Echo

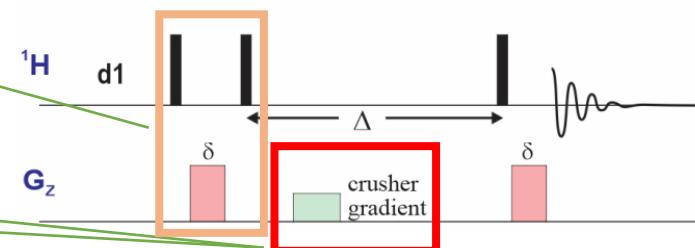


STE (STimulated Echo)

Advantage

1. Remove transverse magnetization($T2$) and
Only Store longitudinally magnetization($T1$)
 $(D20 < T1)$ [$T2 \leq T1$]
2. Eliminate a lot of artifact
3. Reduce chemical shift and J coupling dependence

STE – STimulated Echo



(Homo) Spoil gradient
(Crusher gradient)

Disadvantage

1. Eddy current

Eddy current

DOSY
Experiment

USE

Field Gradient

Gradient Coil



Kieran McKenzie



Metal
of the magnet

Eddy current
Interaction of Metal & Field Gradient

Interaction
Side effects

Interaction
Side effects

Metal
of the probe

Longitudinal Eddy current Delay (LED)

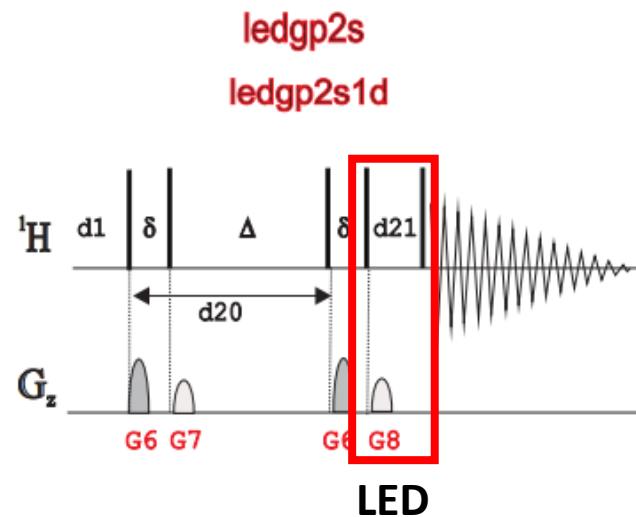
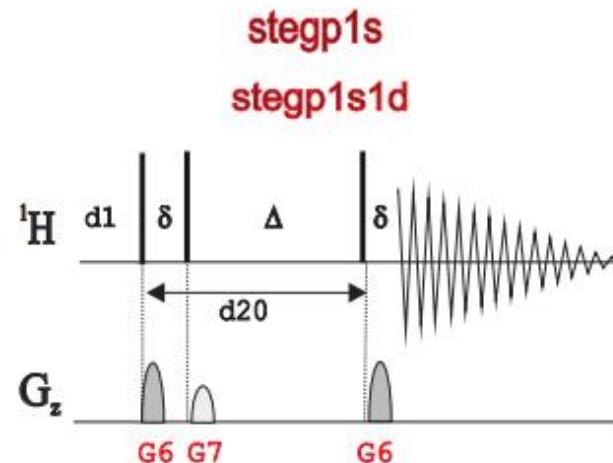
LED (Longitudinal Eddy current Delay)

D21 = 5 ms (typically)

Eddy current = some ms to tens of ms

Advantage

1. Reduce **eddy-currents**
2. Reduce Disturbance of field-frequency **Lock**



Bipolar Pulse Pair (BPP)

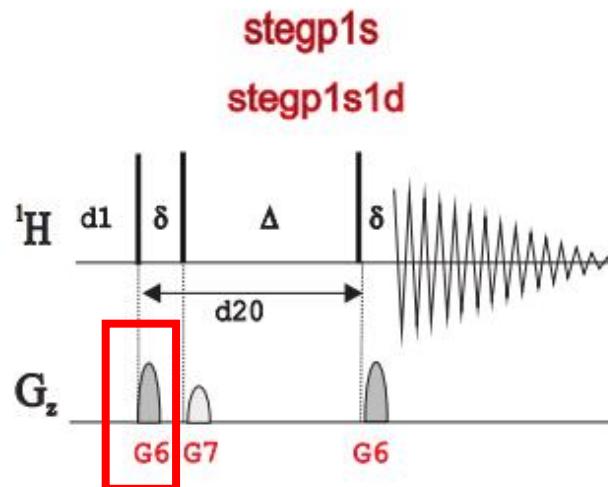
BP (BiPolar)

Advantage

1. Reduce **eddy-currents** (95%)
2. Reduce Gradient imperfections
3. Reduce Disturbance of field-frequency Lock

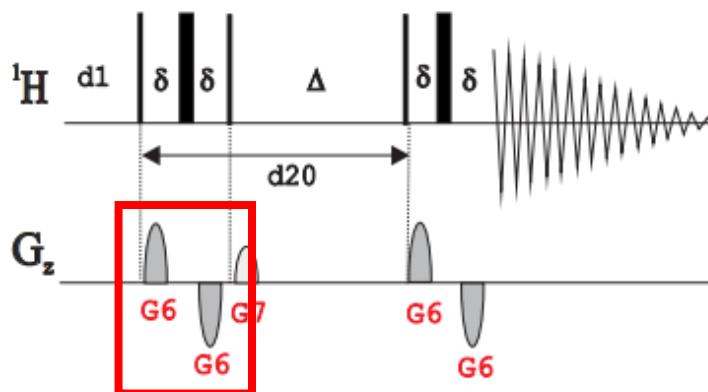
Disadvantage

1. Phase Cycling
(Minimum NS=8 / Preferably NS=16)



MP(Mono Polar)

stebpgp1s
stebpgp1s1d



BP(BiPolar)

Diff (kappa in VNMR)

1. Unbalancing of the bipolar pair reduces reliance on EXORCYCLE phase cycling
2. Diff = 0.2 (typically)

EXORCYCLE: Remove the effects of an imperfect refocusing pulse

(Phase cycling: at least NS = 8)

$$gpz1 = -gpz3 = 1, 20, 37, 49, 58, 66, 73, 80$$

$$diff = \boxed{0.1} \text{ to } 0.2$$

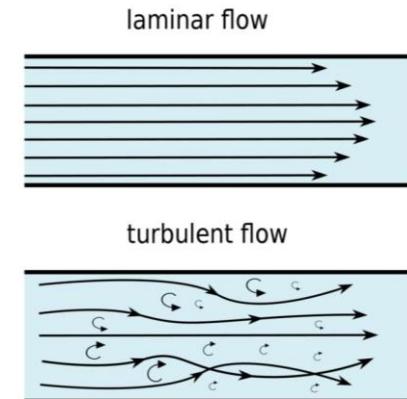
$$\begin{aligned} gpz1 &= 1.1, 22, 40.7, 53.9, 63.8, 72.6, 80.3, 88 \\ gpz3 &= -0.9, -18, -33.3, -44.1, -52.2, -59.4, -65.1, -72 \end{aligned}$$

Convection compensation (CC)

DSTE (Double STimulated Echo)

-> CC(Convection compensation)

- Canceling all constant-velocity effect (laminar flow only)
- Convection cause
 - 1) Temperature gradients
 - 2) RF pulse heating effect
 - 3) Cryo-probe occur convection on room temp



Disadvantage

1. Reduce 50% of the magnetization is lost compared to the non-cc version

Minimize convection

1. Use 3 mm sample tube
2. Disconnect all air flow into the probe sample area
3. Turn off temperature regulation
4. Use a shorter sample with less volume.

Stimulated Echo pulse, STE / LED

ste_gp1s = STE

ste_{bpgp}1s = STE + BP

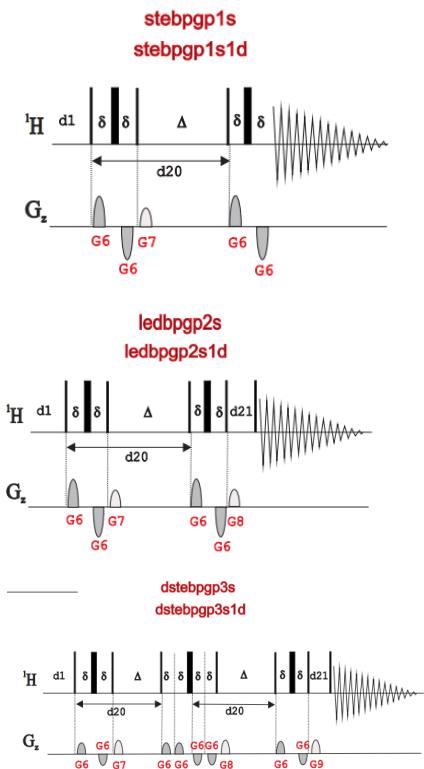
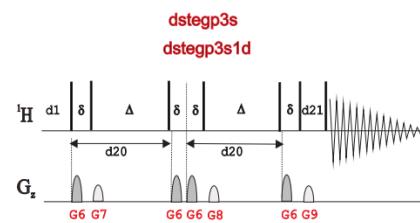
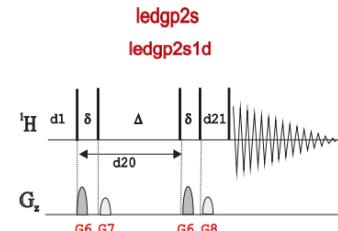
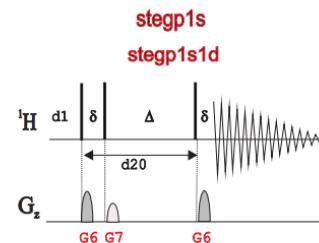
ste_{bpgp}1s19 = STE + 3-9-19 watergate

led_{bpgp}2s = LED

led_{bpgppr}2d = LED + preset

dste_{bgp}3s = STE + CC

dste_{bpgp}3s = STE + BP + CC



Pulsed field Gradient Spin-Echo (PGSE)

STimulated-Echo pulse (STE)

Longitudinal Eddy current Delay (LED)

Double STimulated-Echo (DSTE)

BiPolar gradient (BP)

Origin of DOSY sequence

Minimization of the short T2 effects

Minimization of eddy currents

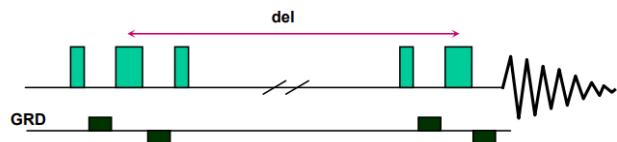
Convection Compensation (CC)

Reduce eddy-currents (95%)

& Inhomogeneous background gradient

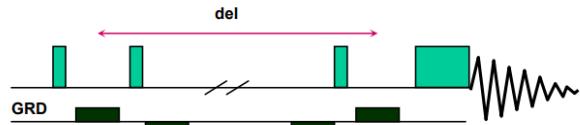
DOSY sequence in Varian

4.2.1 Dbppste (DOSY Bipolar Pulse Pair STimulated Echo) Experiment



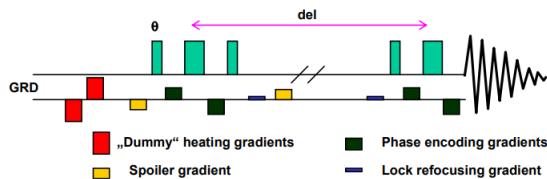
Reference: Wu, D.; Chen, A.; Johnson, C.S., Jr., J. Magn. Reson. 1995, 115, Series(A), 260-264.

4.2.2 DgcsteSL (DOSY Gradient Compensated Stimulated Echo with Spin Lock) Experiment

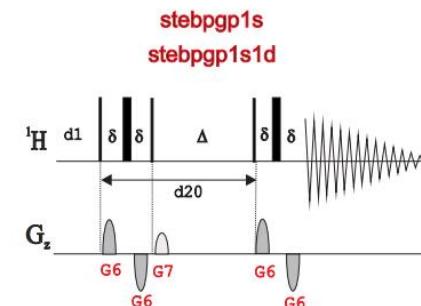


Reference: Pelta, M.D.; Barjat, H.; Morris, G.A.; Davis, A.L., Hammond, S.J. Magn. Reson. Chem. 1998, 36, 706.

4.2.3 The “Doneshot” Experiment



Reference: M. D. Pelta, G. A. Morris, M. J. Tschedroff and S. J. Hammond: MRC 40, 147-152 (2002)
For eliminating radiation damping: M. A. Conell, A. L. Davis, A. M. Kenwright and G. A. Morris: Anal. Bioanal. Chem. 378, 1568-1573, (2004).



SL(Spin Lock):
Keep magnetization in the **transverse plane**

Oneshot DOSY: Possible NS=1 (Non phase cycling)

CC(Convection compensation) in Varian

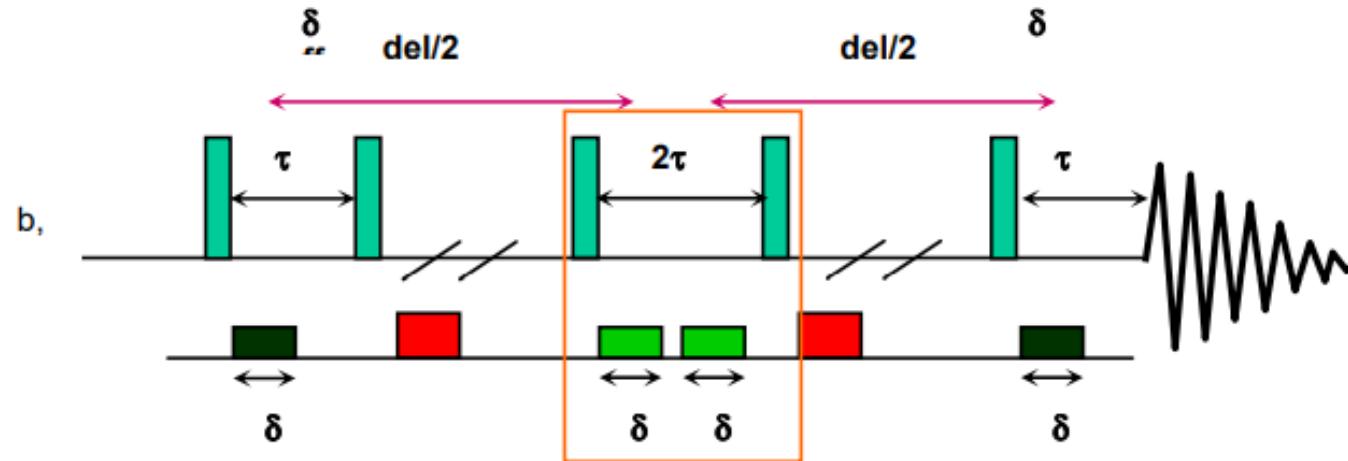


Figure 10 Modification of a gradient stimulated echo experiment with convection compensation.

DgsteSL_cc (Gradient STimulated Echo with Spin-Lock and Convection Compensation)

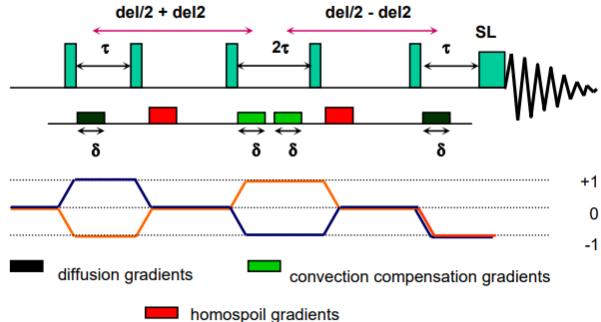
DgcsteSL_cc (Gradient Compensated STimulated Echo with Spin-Lock and Convection Compensation)

Dbppste_cc (Bipolar Pulse Pair STimulated Echo with Convection Compensation)

Dpfgdste (Pulse Field Gradient Double STimulated Echo)

CC(Convection compensation) in Varian

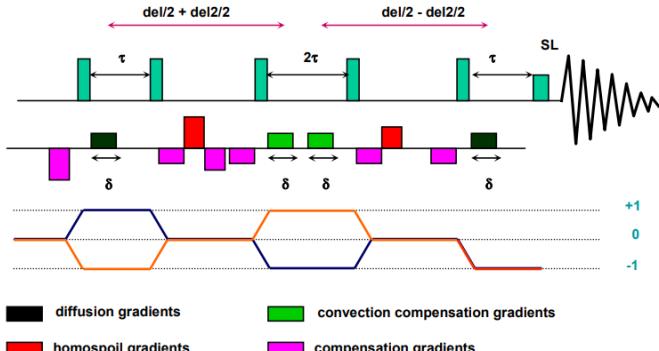
4.4.1.2. DgsteSL_cc (Gradient STimulated Echo with Spin-Lock and Convection Compensation)



Reference: A. Jerchow and N. Müller, J. Magn. Reson. 125, 372-375 (1997).

SL(Spin Lock): O
Gradient compensation: O
Convection compensation: O
Z-filter(LED): X

4.4.1.3 DgcsteSL_cc (Gradient Compensated STimulated Echo with Spin-Lock and Convection Compensation)



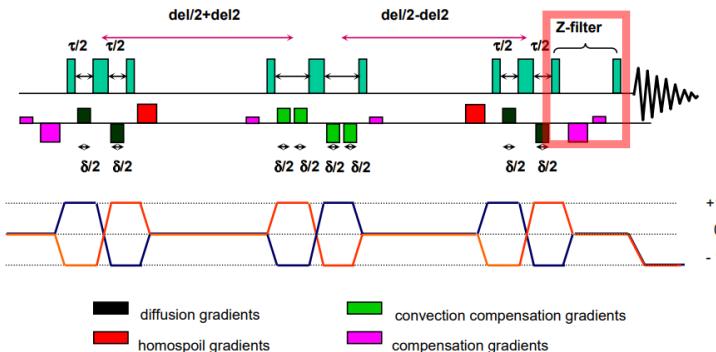
Reference: A. Jerchow and N. Müller, J. Magn. Reson. 125, 372-375 (1997).

SL(Spin Lock): O
Gradient compensation: O
Convection compensation: O
Z-filter(LED): X

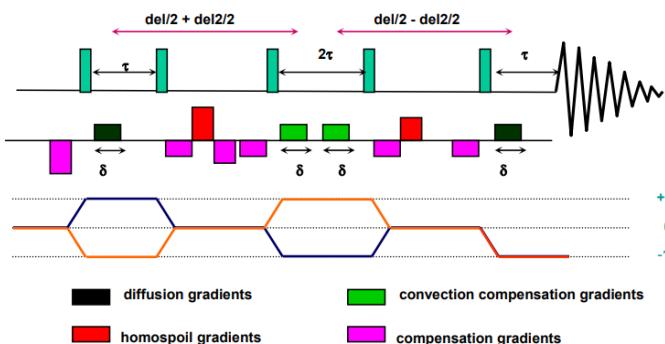
CC(Convection compensation) in Varian

4.4.1 Pulse Sequences with Convection Compensation

4.4.1.1. Dbppste_cc (Bipolar Pulse Pair STimulated Echo with Convection Compensation)



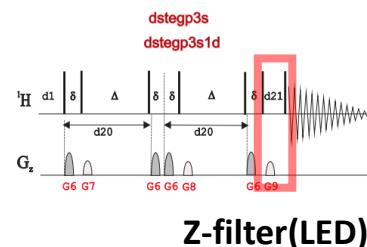
4.4.1.4. Dpfgdste (Pulsed Field Gradient Double STimulated Echo)



Reference: Nilsson M, Gil AM, Delgadillo I, Morris GA. Anal Chem 2004;76:5418-5422



SL(Spin Lock): X
 Gradient compensation: O
 Convection compensation: O
 Z-filter(LED): O



SL(Spin Lock): X
 Gradient compensation: O
 Convection compensation: O
 Z-filter(LED): X

One-shot DOSY

MAGNETIC RESONANCE IN CHEMISTRY
Magn. Reson. Chem. 2002; **40**: S147–S152
Published online in Wiley InterScience (www.interscience.wiley.com). DOI: 10.1002/mrc.1107

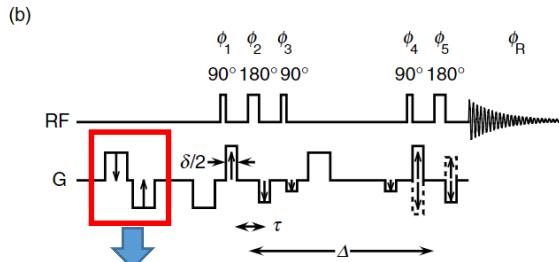
MRC

A one-shot sequence for high-resolution diffusion-ordered spectroscopy

BPPSTE(stebpgp1s, ledbpgp2s, dstebpgp3s)

-> Required EXORCYCLE phase cycling (at least NS=8)

EXORCYCLE: Remove the effects of an imperfect refocusing pulse



Remove the need for EXORCYCLE

-> Cyclops cycle

Retain the refocusing of chemical shifts and background field inhomogeneity

keep constant the **net energy** supplied to the gradient coil by the sequence for a **single transient (NS=1)**

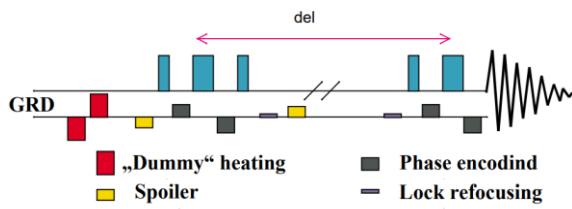


Figure 8. Oneshot DOSY Experiment

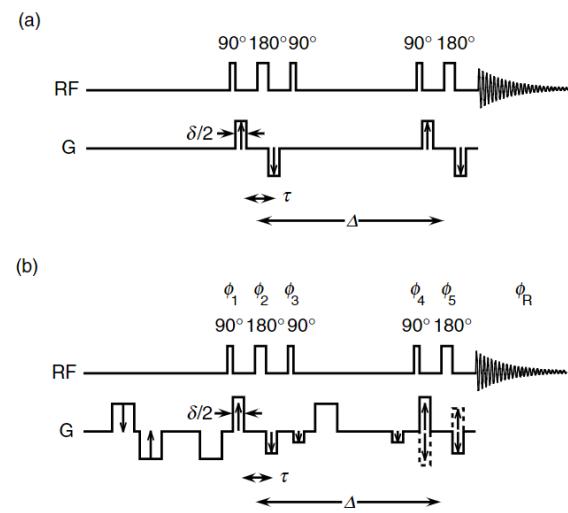


Figure 1. Radiofrequency (RF) and gradient (G) pulse sequences for DOSY. (a) Bipolar pulse pair stimulated echo (BPPSTE) sequence; (b) the proposed one-shot sequence using (solid line) the stimulated antiecho and (dotted line) the stimulated echo. The diffusion delay Δ is the time between the

Table 1. Phase cycling for the pulse sequences of Fig. 1(b)

ϕ_1	$0_{414} + 02$
ϕ_2	$0_{128}2_{128}$
ϕ_3	$0_{32}2_{32}$
ϕ_4	$0_{22}2 + 0_{81}8_{23}8$
ϕ_5	$0_{64}1_{64} + 0_{16}2_{16}$
ϕ_R	$\phi_1 - 2\phi_2 + \phi_3 - \phi_4 + 2\phi_5$ (solid line)
ϕ_R	$-\phi_1 + 2\phi_2 - \phi_3 - \phi_4 + 2\phi_5$ (dotted line)

Phases are notated as multiples of 90° ($0 = 0^\circ$, $1 = 90^\circ$, $2 = 180^\circ$, $3 = 270^\circ$), with subscripts denoting repetition; thus the cycle $0_{414} + 02$ corresponds to the sequence of phases $0^\circ, 180^\circ, 0^\circ, 180^\circ, 90^\circ, 270^\circ, 90^\circ, 270^\circ$ on successive transients.

3. Standard Parameter Calibration



UNIST

ULSAN NATIONAL INSTITUTE OF
SCIENCE AND TECHNOLOGY

Parameters for Diffusion NMR

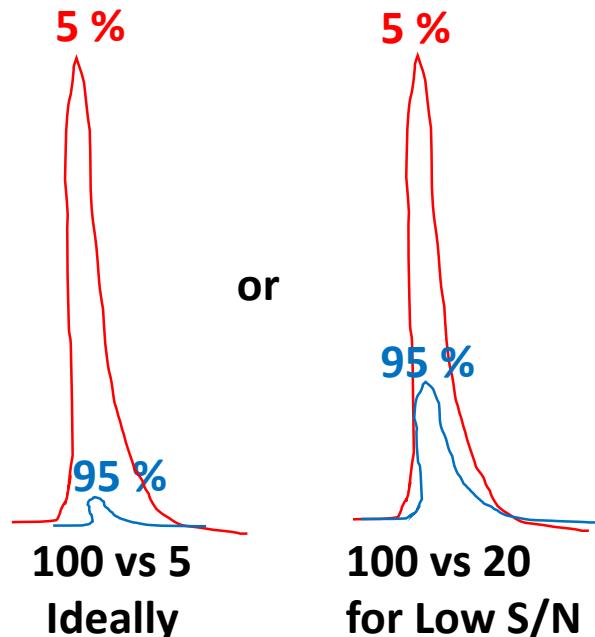
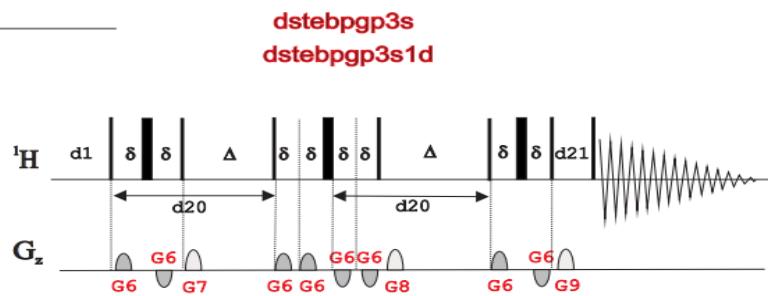
P1

Pulse width

Maximum Peak intensity

Phasing / Dephasing

DOSY sequence use many $\pi/2$ (90°), π (180°)



T1

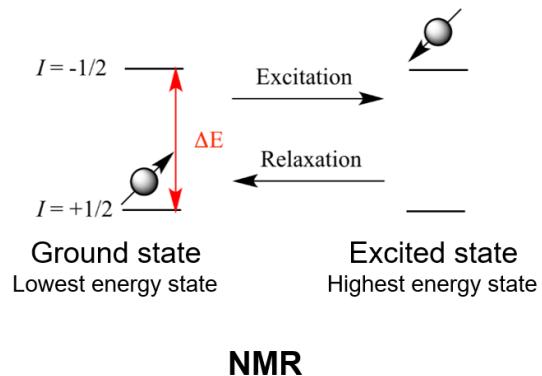
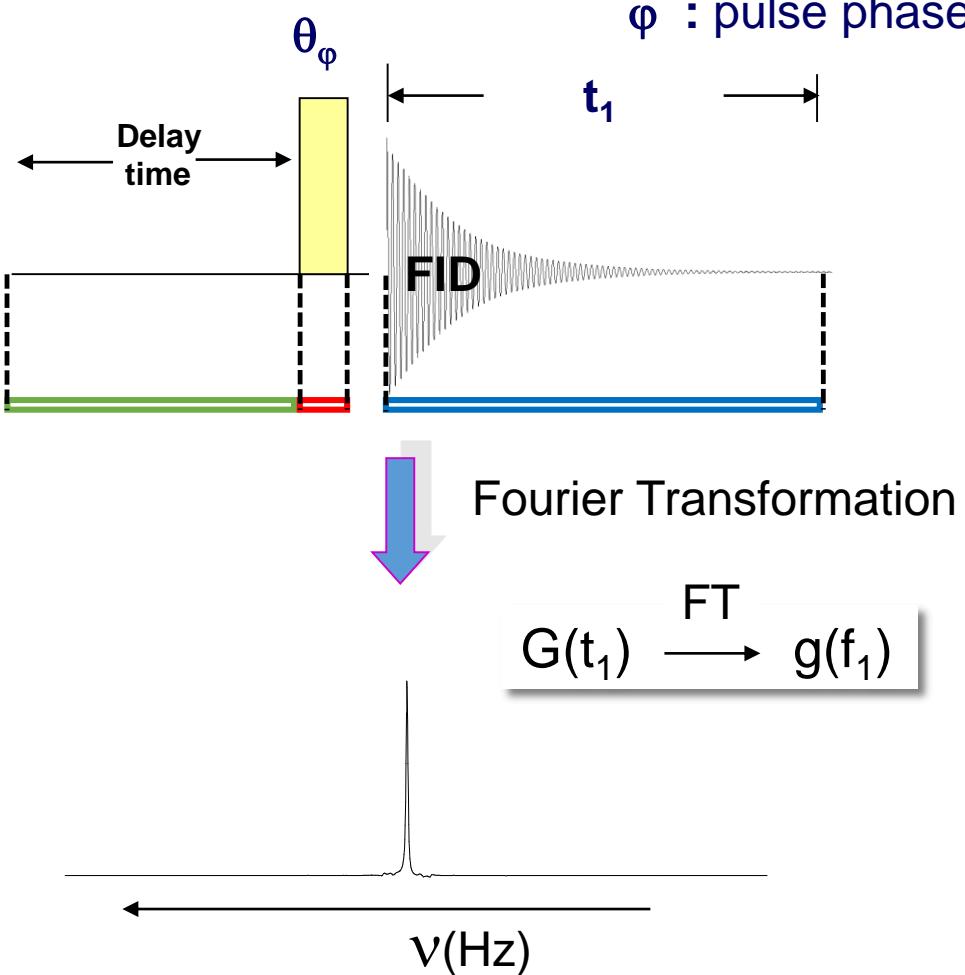
$$AQ + D1 = T1 * 5$$

$$T1 > D20$$

Big Delta, **D20**

Little Delta, **P30**

Basic 1D NMR

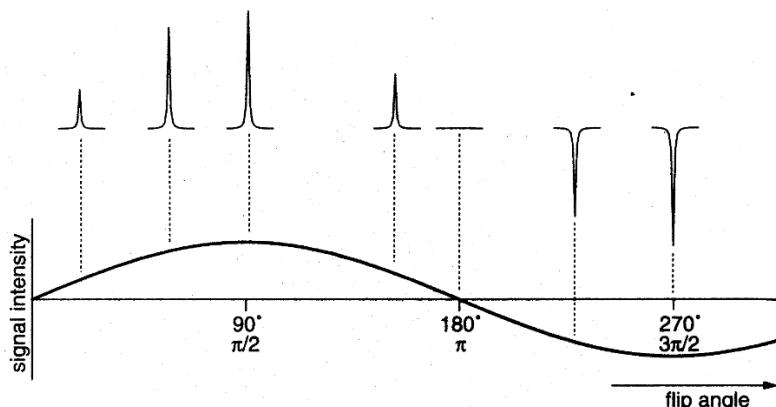


Parameter

- D1 (Relaxation delay time)
- P1 (Pulse width)
PLW1 (Power level)
- AQ (Acquisition time)

1. Find P1

- 1) Type edc – create experiment room
- 2) Change PULPROG zg30 to **zg**
- 3) Set D1=10, P1=3, DS=0, NS=1
- 4) Type rga; zg
- 5) ProcPars -> Change pHmod no to pk
- 6) Type efp
- 7) Correct phase and save
- 8) Set P1=20 (About 180 °) or P1=40 (About 360 °)
- 9) Type efp -> Check peak phase (negative or positive)
- 10) Change P1 (negative -> decrease P1, positive -> increase P1)
and find symmetry peak intensity (Exact 180 °)
- 11) Set P1/2 = 90 degree pulse (=Highest intensity)



1-1. Find P1 (with pulsecal)

- 1) Type edc – create experiment room
- 2) Change PULPROG zg30 to **zg**
- 3) Set D1=10, P1=3, DS=0, NS=1
- 4) Type **pulsecal**

2. Find **T1**

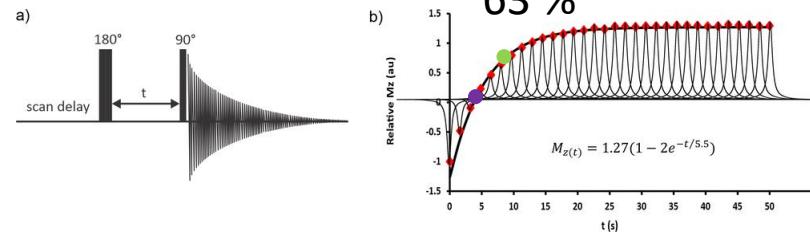
- 1) Type wrpa (ex. exp# 900)
- 2) Change PULPROG **t1ir**
- 3) Click AcquPars and change 1D to 2D
- 4) Set F1 domain TD=10 and set vdlist (ex, T1=20s)
- 5) Type ased
- 6) Set DS=0, NS= 1 or 2
- 7) Type rga; zg
- 8) Type xf2; abs2
- 9) Type rser 1 and correct phase and then **save 2d**
- 10) Type xf2; abs2
- 11) Click Analyse -> Dynamics -> T1/T2
- 12) Click Spectrum and put 1 at Slice number
- 13) Click Peak/Range and click manual integration
- 14) Export Regions o relaxation module and .ret
- 15) Click Relaxation and click fitting

2-1. Find **D7null(T1)**

- 1) Type wrpa (ex. Exp# 901)
- 2) Change PULPROG **t1ir1d**
- 3) Type ased
- 4) Set D7null=0.002(or 2m)
- 5) Type zg and efp
- 6) Check negative peak.
- 7) Find D7null value
(zero intensity)

$$T_1 * \ln(2) = D7null$$

$$T_1 = D7null * 1.443$$



4. DOSY Parameter Calibration



UNIST

ULSAN NATIONAL INSTITUTE OF
SCIENCE AND TECHNOLOGY

DOSY parameters

D20	Diffusion delay	100 ms < T1 (shortest)
D21	Eddy-current delay	5 ms (typically)
P30	Gradient duration	500 – 3000 us (Usually 2500 us), P30/(D1+AQ) < 0.05 s
D16	Gradient recovery delay	200 – 1000 us (typically)
P19	Gradient length	600 us (0.6ms)
gpz6	Gradient ratio	5 – 95 %
gpz7	Spoil gradient	-17.13 %
GPNAM 6		SMSQ10.100 (smoothed rectangular shape)
GPNAM 7		SMSQ10.100 (smoothed rectangular shape)
NS	number of scans	8*n (Related Phase cycling for BP)
DS	dummy of scans	4*m
D1	Delay time	T1 * 3 – 5
TD1	Steps	16 (usually 16 ~ 32, Use \geq 7)

P30 (Gradient duration)

P30 Gradient duration

Protect the probe -> Max gradient time is 10 ms

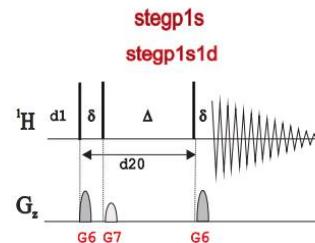
stepp1s

- 1 spoil gradient (0.6 ms)
- 2 diffusion gradients (4.95 ms * 2)
- GPZ6 : GPZ7(Spoil)

100 : -17.13

$$10 \text{ ms} - 0.6 \text{ ms} * 1 * 17.13 \% = 9.90 \text{ ms}$$

$$9.9 \text{ ms} / 2 = \mathbf{4.95 \text{ ms}}$$



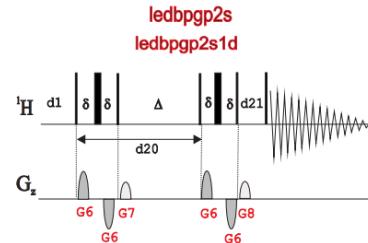
ledbpgp2s

- 2 spoil gradients (0.6 ms * 2)
- 4 diffusion gradients (2.41 ms * 4)
- GPZ6 : GPZ7 (Spoil) : GPZ8 (Spoil)

100 : -17.13 : -13.17

$$10 \text{ ms} - 0.6 \text{ ms} * 2 * (17.13 \% + 13.17\%) = 9.64 \text{ ms}$$

$$9.64 \text{ ms} / 4 = \mathbf{2.41 \text{ ms}}$$



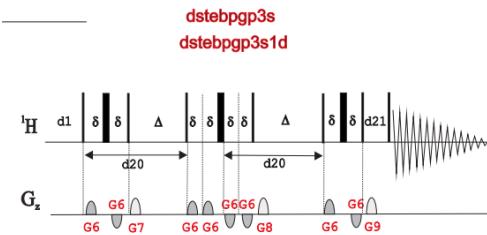
dstebpgp3s

- 3 spoil gradients (0.6 ms * 3)
- 8 diffusion gradients (1.15 ms * 8)
- GPZ6 : GPZ7 (Spoil) : GPZ8 (Spoil) : GPZ9 (Spoil)

100 : -13.17 : -17.13 : -15.37

$$10 \text{ ms} - 0.6 \text{ ms} * 3 * (13.17 \% + 17.13 \% + 15.37 \%) = 9.18 \text{ ms}$$

$$9.18 \text{ ms} / 8 = \mathbf{1.15 \text{ ms}}$$



1. Find P1, AQ, D1(T1)
2. **Find D20 and P30 (Measure 1D DOSY)**
3. Measure 2D DOSY
4. Processing

- 1) Type ***wrpa*** exp # 10
- 2) Change pulprog to stepgp1s1d or dstepgp1s1d (1D DOSY)
- 3) Type ***gppp*** (automatically insert GPNAM6, GPANM7, etc..)
- 4) Set parameter D20, P30
- 5) Set parameter gpz6=95 (95%)
- 6) Type ***wrpa*** exp # 11
- 7) Set parameter gpz6=5 (5%)
- 8) Type ***re 10; zg***
- 9) Type ***re 11; zg***
- 10) Type ***ef*** or click processing spectrum – change ph mode to pk – type efp
- 11) Type ***re 10; efp; .md*** / Double click exp #11
- 12) Click right mouse button → choice match intensities
→ check intensity ratio 5% (0.05 or 20 times),
Poor S/N ratio sample could 20% (0.2 or 5 times)
- 13) Change D20 or P30 until intensity ratio 5%

1. Find P1, AQ, D1(T1)
2. Find D20 and P30 (Measure 1D DOSY)
3. **Measure 2D DOSY**
4. Processing

- 1) Type ***wrpa exp # 100***
- 2) Change pulprog to stepgp1s
- 3) Click AcquPars and change 1D to 2D
- 4) FnMODE = QF
- 5) Type ***gppp*** (Automatically write gradient parameters)
- 6) Set parameter D20, P30
- 7) Type ***xau dosy 5 95 16 l y y***

xau: macro

xau dosy: execute dosy macro

5: First gradient (gpz6)

95: Last gradient (gpz6)

16: TD (for F1), 16 steps

l: Linear lamp (Could choice q:square lamp, e: exponential lamp)

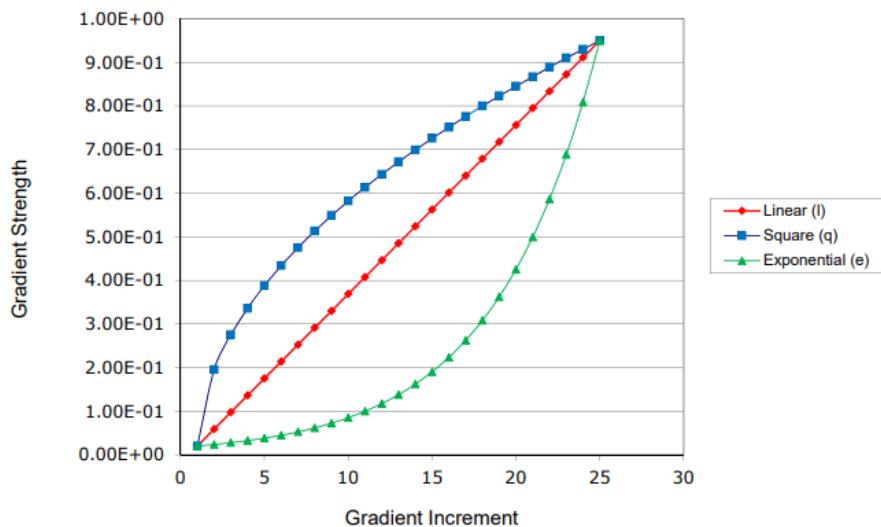
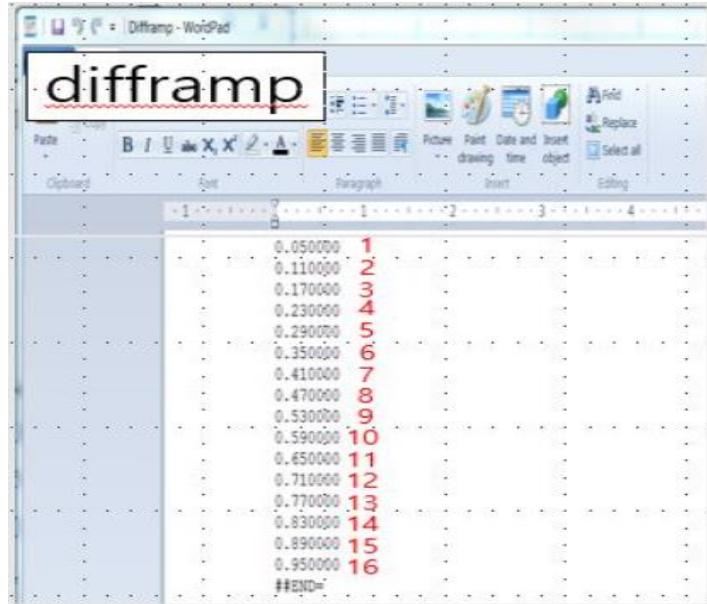
y: start acquisition

y: rga

Class of gradient lamp

Class of gradient lamp

- l: Linear lamp
 - q: Square lamp
 - e: Exponential lamp
-
- Diffraamp -> Normal gradient
 - DiffraampR -> Reverse gradient
- (C/bruker/topspin/exp/stan/nmr/list/gp/user)



Special technique

Special technique

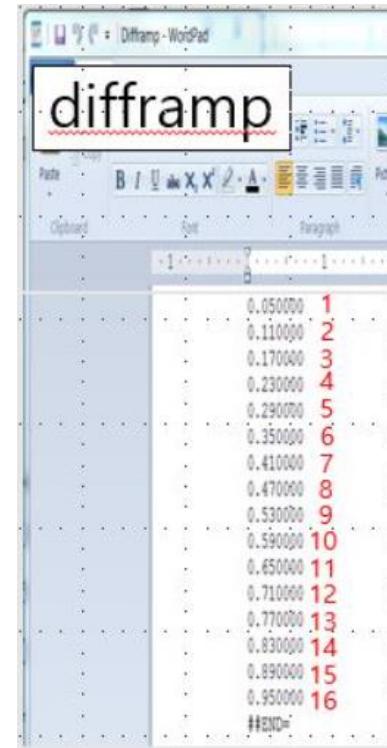
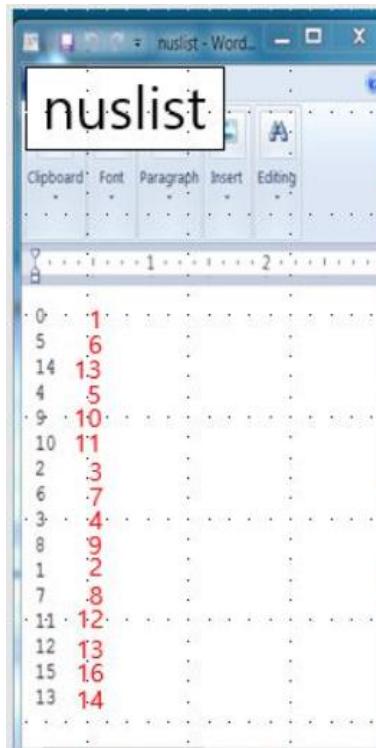
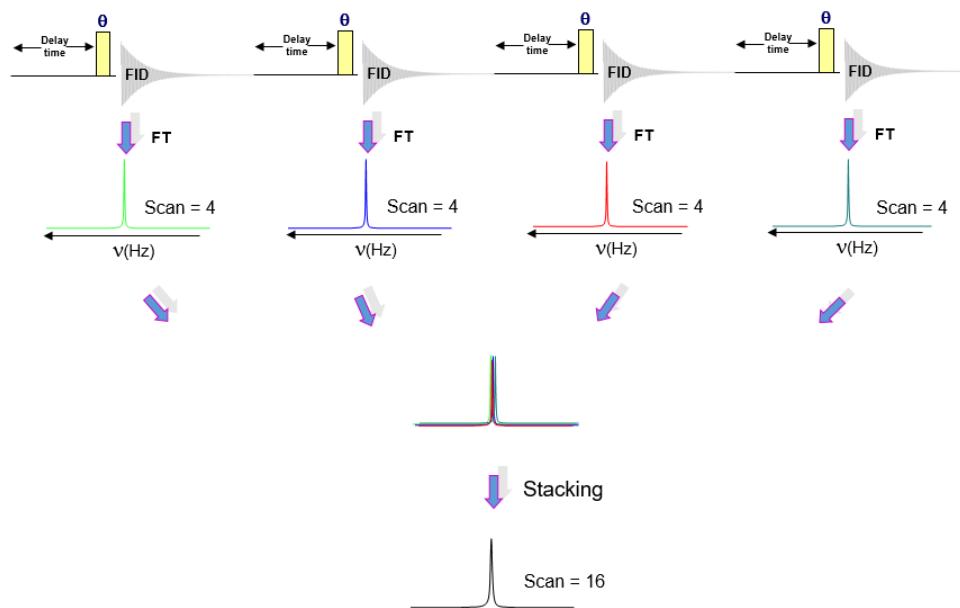
1. Random diffusion (Use NUS)

- Set FnType to “Non-uniform sampling”
- Set NUSamount to 100%
- Set NusSPTYPE to 3

2. Tdav (= interleaved in Varian)

If we set Tdav = 4, NS=16, it works like

-> Improve diffusion data accuracy



1. Find P1, AQ, D1(T1)
2. Find D20 and P30 (Measure 1D DOSY)
3. Measure 2D DOSY

4. Processing

- 1) Type *xf2; abs2; setdiffparm*
- 2) Analyse -> T1/T2 -> FID -> Spectrum -> 1
- 3) T1/T2/ -> Peaks/FID -> Integration -> Save to relaxation module
- 4) Relaxation -> >>

5. Temperature Calibration



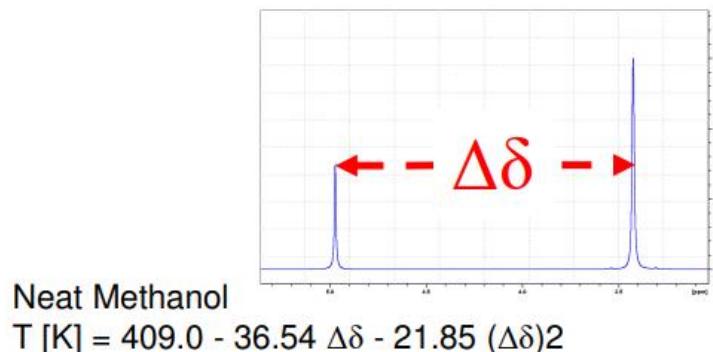
UNIST

ULSAN NATIONAL INSTITUTE OF
SCIENCE AND TECHNOLOGY

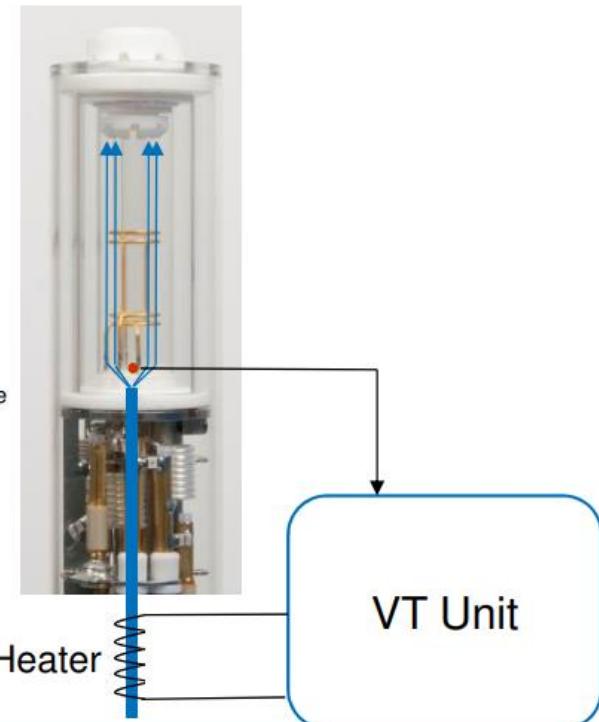


Temperature Calibration

- The VT Sensor can be calibrated for accurate sample temperature values
- The -OH of MeOH is temperature sensitive

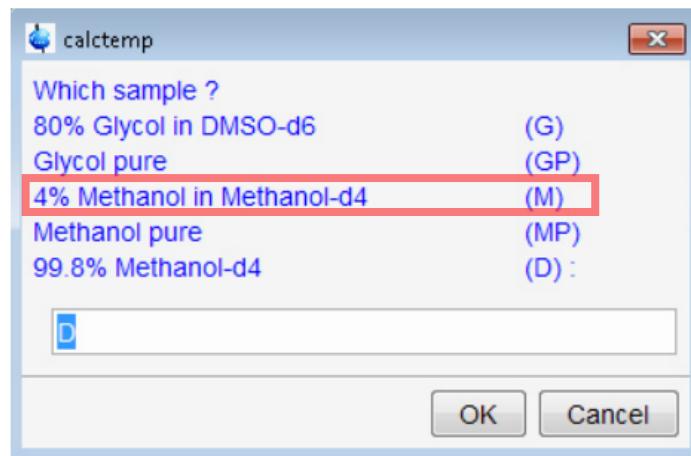


- The AU program **calctemp** measures the splitting and determines actual sample temp



J. Magn. Reson. 1982, 46, 319-321

Temperature Calibration - calctemp



- High Temp = Ethylene Glycol
 - ✓ 300-380 K
- Room/Low Temp = Methanol
 - ✓ 180 – 330K
- Cryoprobes especially 99.8% Methanol-d4

Temperature Calibration

Ethylene Glycol is useful for temperatures from ~300K to 380K

Source: Bruker Instruments, Inc. VT-Calibration Manual

80% Ethylene Glycol (DMSO-d6): $T = (4.218 - \Delta)/0.009132$

100% Ethylene Glycol: $T = (4.637 - \Delta)/0.009967$

(Δ is the shift difference (ppm) between CH_2 & OH peaks)

Methanol is useful for temperatures from ~180K to 300K

Source: Bruker Instruments, Inc. VT-Calibration Manual

100% Methanol: $T = -23.832\Delta^2 - 29.46\Delta + 403.0$

(Δ is the shift difference (ppm) between CH_3 & OH peaks)

Source: Bruker Instruments, Inc. VT-Calibration Manual

4% Methanol in Methanol-d4:

180-300K, $T = (3.86 - \Delta)/0.00782$ (approximate)

for more accurate values, use the following, depending on T:

180-230K, $T = (3.72 - \Delta)/0.007143$

230-270K, $T = (3.92 - \Delta)/0.008$

270-300K, $T = (4.109 - \Delta)/0.008708$

(Δ is the shift difference (ppm) between the CH_3 and OH peaks)

100% d4-Methanol, Deuterium Observe

(Source: Experimental Data: Univ. Nebraska, 2001)

$$T = -23.1902\Delta^2 - 31.1062\Delta + 399.081$$

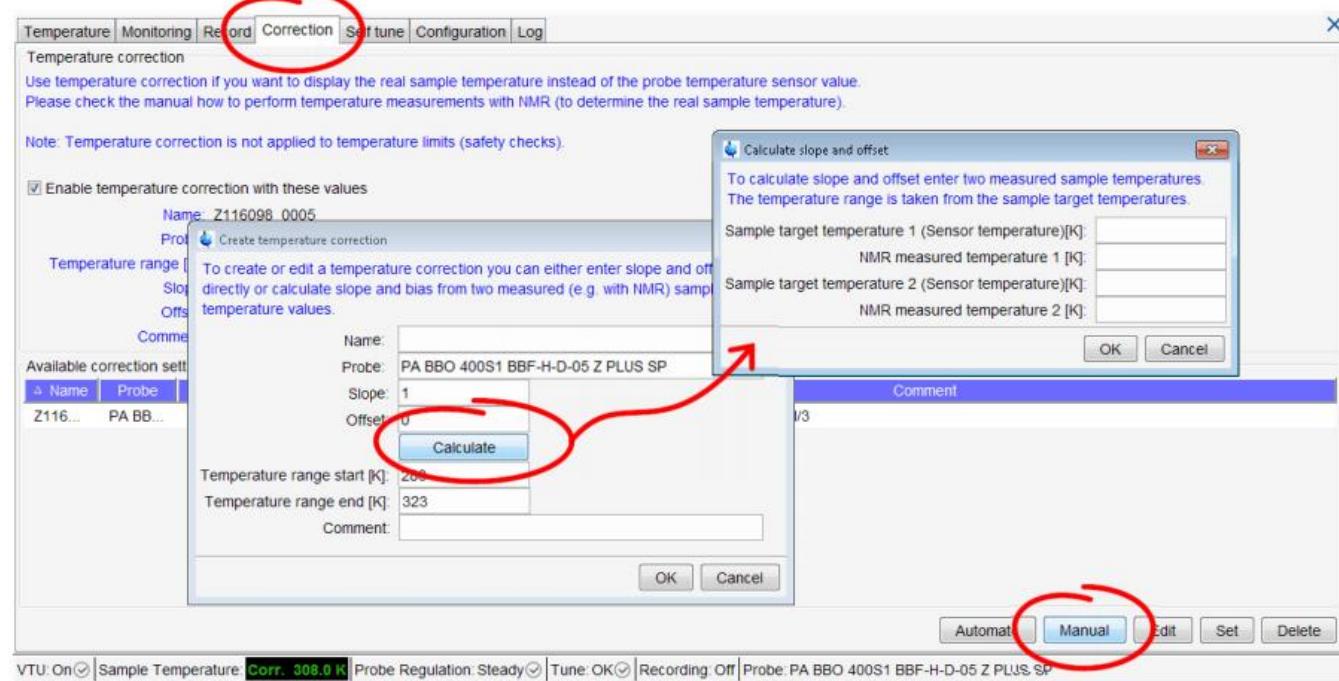
where Δ = ppm(OD) vs. ppm (CD_3)

Temperature Calibration / Correction



A simple 2 point correction can be created and enabled

1. Set and measure the temperature at 2 targets
2. Suggest 30 – 40 degrees apart.
3. This is the correction for this range
4. Add another correction for other ranges



Temperature Calibration / Correction



Enable the correction appropriate for your temperature range

Screenshot of the Bruker NMR software interface showing the Temperature Correction dialog box.

The dialog box has tabs: Temperature, Monitoring, Record, Correction, Self tune, Configuration, and Log. The Correction tab is selected.

Temperature correction

Use temperature correction if you want to display the real sample temperature instead of the probe temperature sensor value.
Please check the manual how to perform temperature measurements with NMR (to determine the real sample temperature).

Note: Temperature correction is not applied to temperature limits (safety checks).

Enable temperature correction with these values

Name: Z116098_0005
Probe: PA BBO 400S1 BBF-H-D-05 Z PLUS SP

Temperature range [K]: 298 - 308
Slope: 0.87832
Offset: 36.7218
Comment: created by NPT_1H_tempcalib_998meod/3

Available correction settings

Name	Probe	Temperature Range [K]	Slope	Offset	Comment
Z116098_	PA BBO	298 - 308	0.87832	36.7218	created by NPT_1H_tempcalib_998meod/3

Buttons at the bottom: Automatic, Manual, Edit, Set (highlighted with a red circle), Delete.

Bottom status bar: VTU: Off (radio button), Sample Temperature: **Corr. 301.1 K**, Probe Regulation: Transient (radio button), Tune: OK (radio button), Recording: Off, Probe: PA BBO 400S1 BBF-H-D-05 Z PLUS SP

6. Gradient Calibration



The logo of Ulsan National Institute of Science and Technology (UNIST) features the letters "UNIST" in a bold, blue, sans-serif font. A bright blue light source is positioned at the top right of the letter "I", casting a glow across the letters.

ULSAN NATIONAL INSTITUTE OF
SCIENCE AND TECHNOLOGY

Calibration Procedure for a Single Axis Gradient System

$$GCC_{new} = GCC_{old} \sqrt{\frac{D}{D_{Literature}}}$$

Type gradpar

Initial value = 5.35 G/cm

Find proper gradpar value

(Doped water = 1.913 on RT)

2.4.6 Preparation for NMR Adjustments

It is highly recommended to use the doped water sample (Z10906) for all setup experiments. If this sample is not available, a similar sample consisting of 1% H₂O in D₂O, plus 0.1% CuSO₄ should be prepared. The exact concentration is not important for the pre-emphasis adjustment, but it is useful for the later gradient calibration. Low concentrations of H₂O are important in preventing radiation damping. The CuSO₄ is mainly used to save time, but it also helps prevent radiation damping. The sample filling height should be about 40 mm.

1.1. Product identifier

NMR-SAMPLE 0,1 mg GdCl₃/ml D₂O + 1% H₂O + 0,1% CH₃OH 13C (DOPED WATER)

Further trade names

NMR-SAMPLE 0,1 mg GdCl₃/ml D₂O + 1% H₂O + 0,1% CH₃OH 13C (DOPED WATER):
Z100933, Z10727, Z10046, Z10083, Z10906, Z10085

Z10082, Z10084, Z10916

NMR-SAMPLE 0,1 mg GdCl₃/ml D₂O + 1% H₂O (DOPED WATER):

Z10047, Z10088, Z10090

Z10087, Z10089

Nucleus	Sample	Temperature [C]	D [$10^{-9}m^2/s$]
1H	H ₂ O	20	2.031
1H	H ₂ O	25	2.299
1H	D ₂ O	25	1.872
1H	“Doped Water“	25	1.91
1H	DMSO	25	0.730
7Li	0.25 m LiCl in H ₂ O (ca. 10 g LiCl / l H ₂ O)	25	0.960
^{23}Na	2 m NaCl in H ₂ O (ca. 117 g NaCl / l H ₂ O)	25	1.135

Table 2.3: Some useful diffusion coefficients

$$D(x_D) = (2.3 - 0.4652 \cdot x_D + 0.0672 \cdot x_D^2) \cdot 10^{-9} \frac{m^2}{s}$$

x_D = mole fraction of deuterons

Figure 2.25: Equation 1

[* Calibration in Accurate Spin-Echo Self-Diffusion Measurements Using 1H and Less-Common Nuclei, Holz et al., JMR 92, 115-125 (1991)].

Non-linearity of the gradient

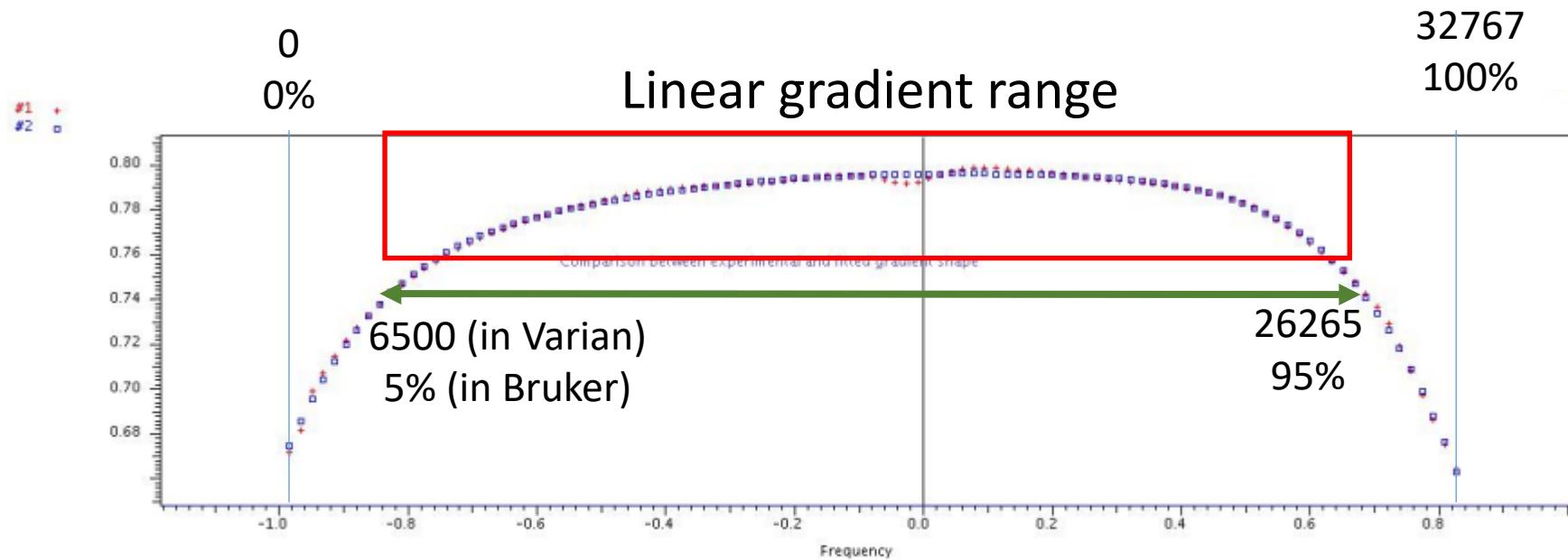


Figure 7 Typical comparison between the experimental and fitted gradient shape produced by the probe.

Limitation of measurement

Probe		Gradient	T1, T2 limit	Estimation limit
BBO	Bruker	0.05 T/m		$< 3 \times 10^{-11} \text{ m}^2/\text{s}$
AutoX-DB	Varian	0.06 T/m		$< 2.8 \times 10^{-11} \text{ m}^2/\text{s}$
DiffBB	Bruker	17 T/m		$< 1 \times 10^{-14} \text{ m}^2/\text{s}$
Diff30	Bruker	18 T/m	$T1 > 1 \text{ s}, T2 > 100 \text{ ms}$	$< 9.4 \times 10^{-15} \text{ m}^2/\text{s}$
Diff50	Bruker	30 T/m	$T1 > 0.56 \text{ s}, T2 > 56 \text{ ms}$	$< 5 \times 10^{-15} \text{ m}^2/\text{s}$
DOTY Diff	DOTY	33 T/m	$T1 > 0.504 \text{ s}, T2 > 50.4 \text{ ms}$	$< 4.5 \times 10^{-15} \text{ m}^2/\text{s}$

PROBES

DiffBB

The DiffBB is a broadband gradient probe with the Automatic Tuning Module (ATM) for NMR diffusion investigations of a wide range of diffusivities. 10-8 to 10-13 m²/s (10-15 m²/s under favorable sample conditions).



PFG/Diffusion Liquids Z Gradient Probes-Pulsed Field

Gradient Coils Over 3,300 G/cm (33 T/m)



We have a new version of our PFG probe that can extend sample temperatures up to +300°C. For recent experimental results: "Temperature Dependence of the ¹³C NMR of Nylon 6,6—Liquids- Z Gradient Probe up to 300°C"

For Information on Diffusion Techniques: "A Practical Guide to Setting Up Diffusion Measurements Utilizing Pulsed Field Gradients"

Introduction Specifications Oscillating Gradient Spin Echo (OGSE)

Highest Efficiency, Ultra-Shielded, Fastest Switching

- Measure the lowest diffusion coefficients — to $10^{-15} \text{ m}^2/\text{s}$



Before Your DOSY,

1. Check Temperature Calibration !!

<DOSY Standard sample>

- Doped (0.1mg/ml GdCl₃) 1% H₂O in D₂O sample
- Diffusion coefficient (different temp.)
25.0 °C = 1.913 * 10⁻⁹ m²/s
25.1 °C = 1.918 * 10⁻⁹ m²/s
25.2 °C = 1.923 * 10⁻⁹ m²/s

2. Check Gradient Calibration

3. Check Solvent

- CDCl₃ = Low viscosity = Strong Convection