

# Water suppression NMR

**Advanced Operation  
Training Course**



# Solvent suppression

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The logo of Ulsan National Institute of Science and Technology (UNIST) is displayed in a stylized, glowing blue font. The letters are bold and have a slight perspective, giving them a three-dimensional appearance.

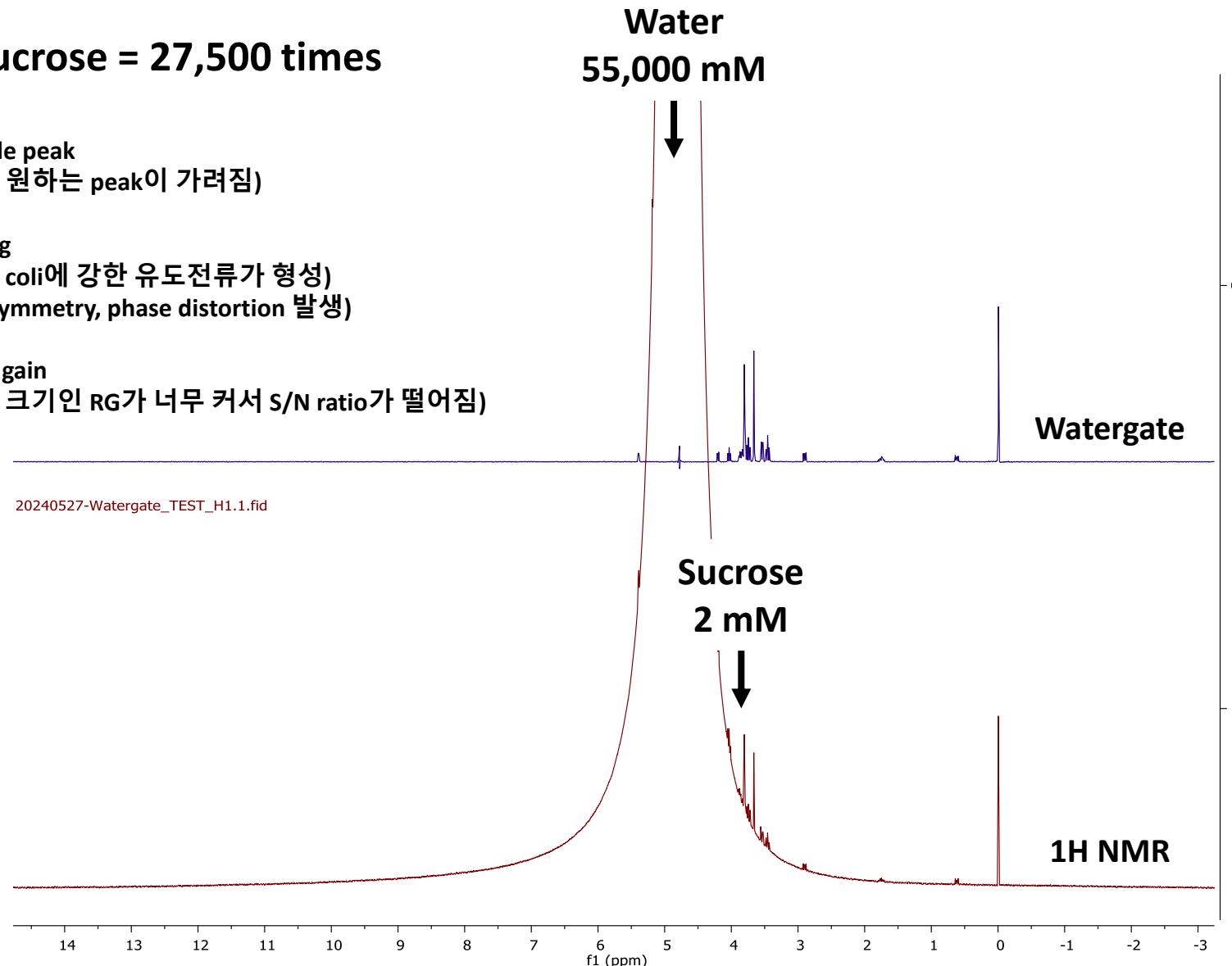
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ULSAN NATIONAL INSTITUTE OF  
SCIENCE AND TECHNOLOGY

# Why need to suppress solvent?

Water / sucrose = 27,500 times

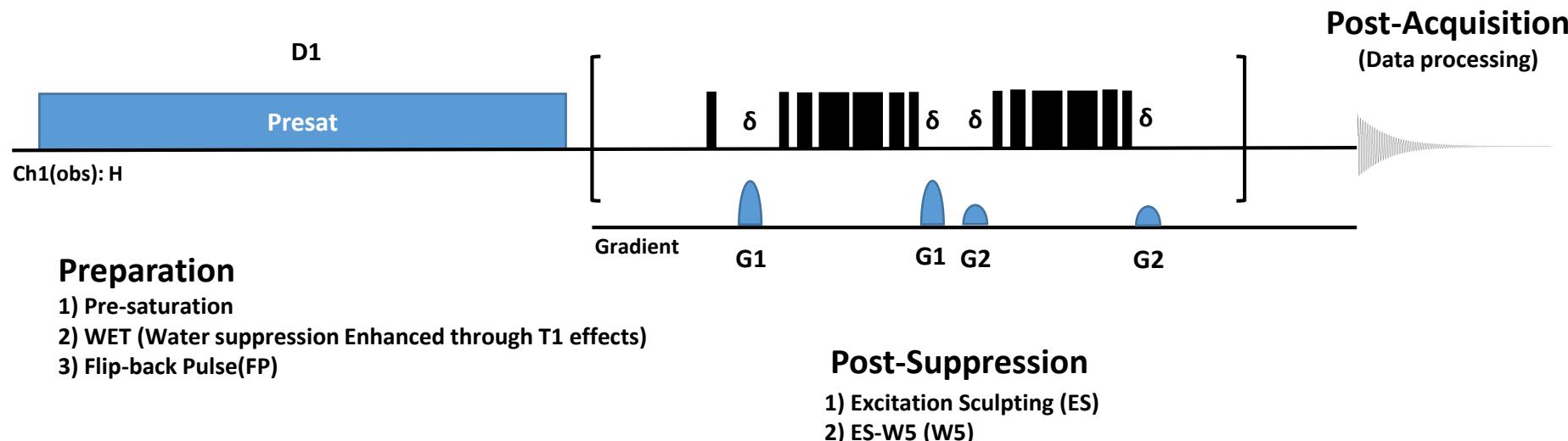
- 1) Overlapped sample peak  
(peak간 겹침으로 원하는 peak이 가려짐)
- 2) Radiation damping  
(고농도로 인하여 coli에 강한 유도전류가 형성)  
(-> broadening, asymmetry, phase distortion 발생)
- 3) Too high Receiver gain  
(Detector window 크기인 RG가 너무 커서 S/N ratio가 떨어짐)



# Type of solvent suppression

## Excitation

- 1) Jump-Return
- 2) Watergate (WG) (+ Field gradient)
- 3) Pulsed Gradient Spin-Echo (PGSE) through T2\*
- 4) Pulsed Gradient Stimulated-Echo (PGSTE) through T1



# Building blocks

## Preparation blocks

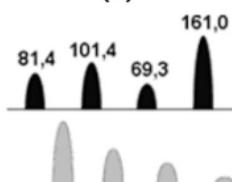
(a) Saturation



(b) DANTE



(c) WET



(d) Flip-back

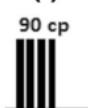


## Excitation blocks

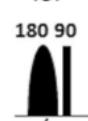
(e) Basic 90° excitation



(f) CP block



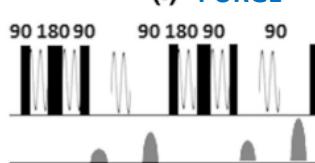
(g) Sat-180 block



(h) NOESY-1D



(i) PURGE

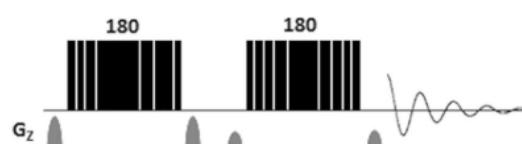


## Readout blocks

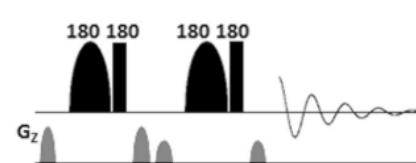
(j)



(k) ES-W5



(l) ES-Sel



**Fig. 1** Basic building blocks constituting water suppression sequences. **a–c** Preparation blocks, **e–i** excitation blocks and **(j–k)** readout blocks. **a** Sat block, **b** DANTE block, **c** WET block, **d** Flip-back block, **e** basic 90° excitation, **f** CP block, **g** Sat-180 block, **h** NOESY-1D block, **i** PURGE block, **k** ES-W5 block and **l** ES-Sel block

## Presat (Pre-saturation)

zgpr: conventional presaturation

- pr (presaturation):

- 1) P1(RF pulse) 이전인 D1 동안 H<sub>2</sub>O만 선택적으로 CW irradiation을 통하여 Excitation (Saturation)
- 2) Saturation된 H<sub>2</sub>O은 Net magnetization이 감소하여 signal이 작아짐

- Ernst angle: 주어진 D1 동안 최단시간에 최대 S/N ratio를 얻을 수 있는 Flip angle

- p19 (Presat power를 가하는 시간)

- plw9 (Presat power level)

- plw9의 값을 조절하여 Saturation peak이 최대한 작아지는 방향으로 조절

- Baseline 기준으로 대칭에 가깝도록!

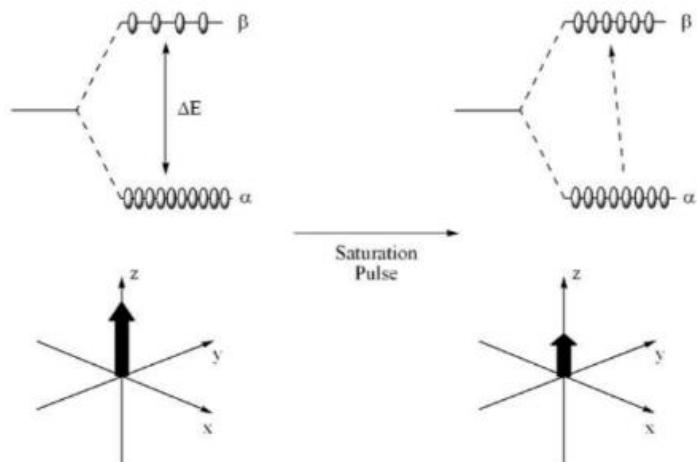
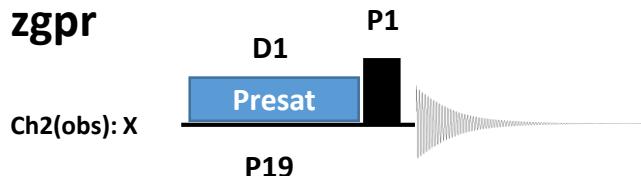
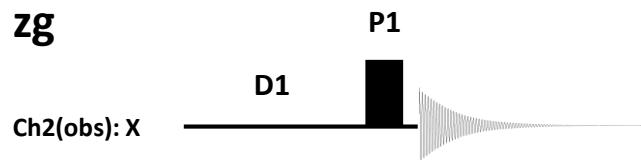
- Saturation peak이 작아질 수록 RG(Receiver Gain)을 열어 S/N 를 증가시킬 수 있음

- 최소값

plw9 ≤ 0.0003 W (0.3 mW)

(pldb9 ≥ 35 -dBW)

plw9과 pldb9은 서로 연동 되어있는 값



## Presat (Presaturation) Origin technique

**zgpr:** conventional presat.

**zg0pr:** find ernst angle

**zgcppr:** presat. with cp(composite pulse)

■ PR (PResaturation):

- 1) P1(RF pulse) 이전인 D1 동안 H<sub>2</sub>O만 선택적으로 CW irradiation을 가함
- 2) Saturation된 H<sub>2</sub>O은 Net magnetization이 감소하여 signal이 작아짐

■ CP (Composite Pulse): 연속적인 RF 입력보다 pulse 형식으로 여러 번 조개서 RF pulse를 통하여 B<sub>1</sub> field의 불균일성을 보정하고 더 완벽한 H<sub>2</sub>O peak 포화(제거)를 할 수 있음

■ Ernst angle: 주어진 D1 동안 최단시간에 최대 S/N ratio를 얻을 수 있는 Flip angle

■ P19 (Presat power를 가하는 시간)

■ plw9 (Presat power level)

■ plw9의 값을 조절하여 Saturation peak이 최대한 작아지는 방향으로 조절

■ Baseline 기준으로 대칭에 가깝도록!

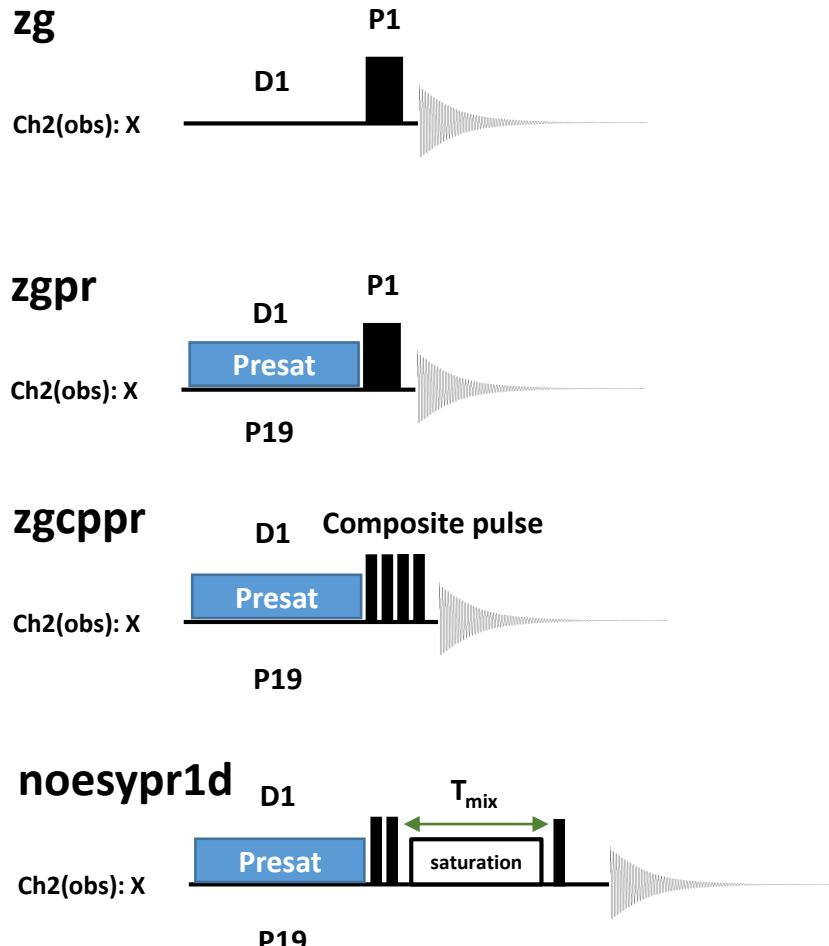
■ Saturation peak이 작아질 수록 RG(Receiver Gain)을 열어 S/N 를 증가시킬 수 있음

■ Minimum Value (최소값)

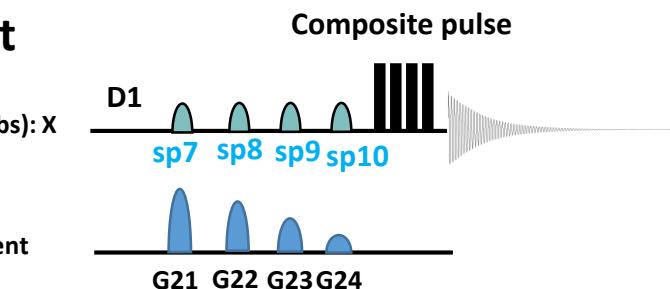
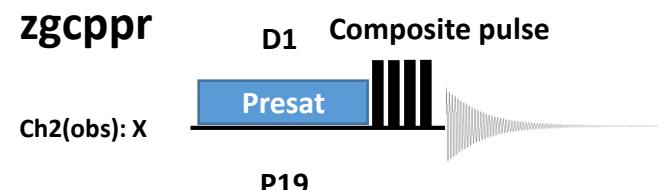
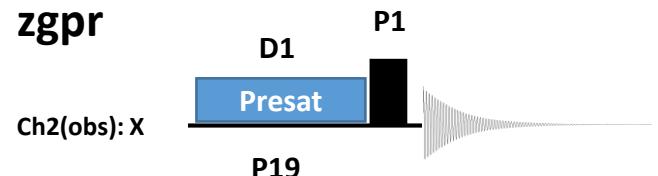
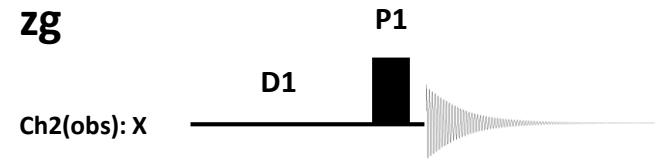
plw9 ≤ 0.0003 W (0.3 mW)

(pldb9 ≥ 35 -dBW)

plw9과 pldb9은 다른 단위로 연동 되어있는 값



# WET (Water suppression Enhanced through T1 effects)

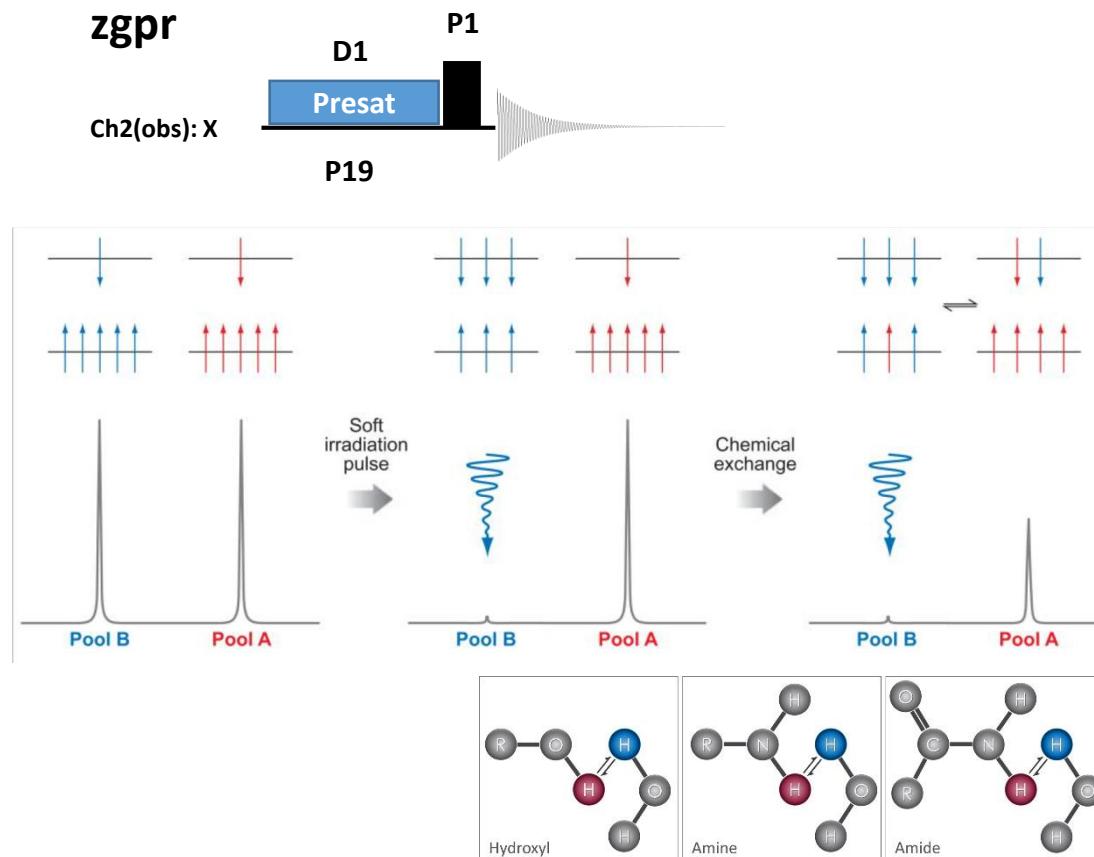


wet: Pre-saturation with gradient + CP (Composite Pulse)  
Minimizing reduced exchangeable proton effect  
Need to control sp7 / sp10

# Exchangeable proton

## Signal attenuation effect

- Saturation된 H<sub>2</sub>O의 H(Proton)이 다른 OH/NH group과 exchange 되어 해당 peak의 신호 감소 효과를 가져옴
- Protein sample의 경우 pH를 7.5이하로 낮춰 Exchange rate을 낮출 필요가 있음

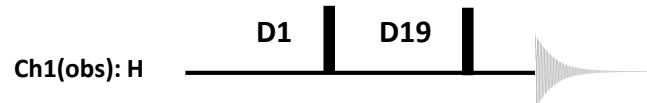


## Jump-Return

**P11 (1-1 scheme)**

**P1331 (1-3-3-1 scheme)**

### P11



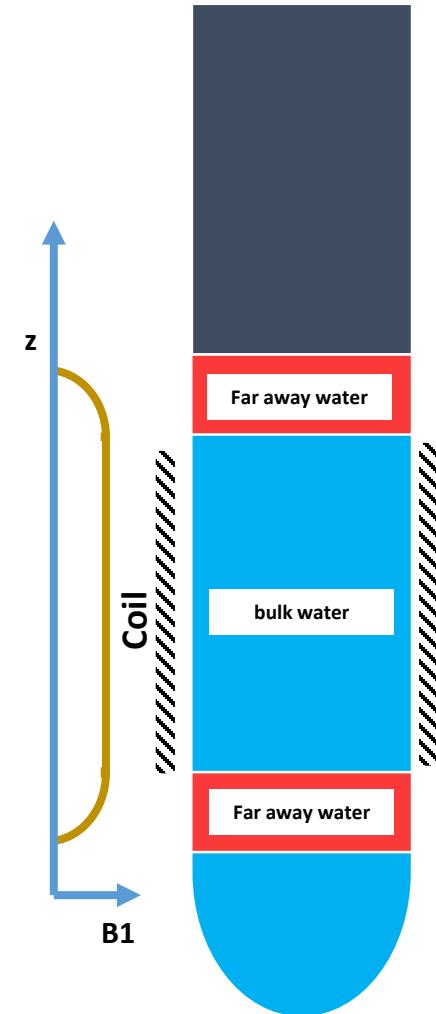
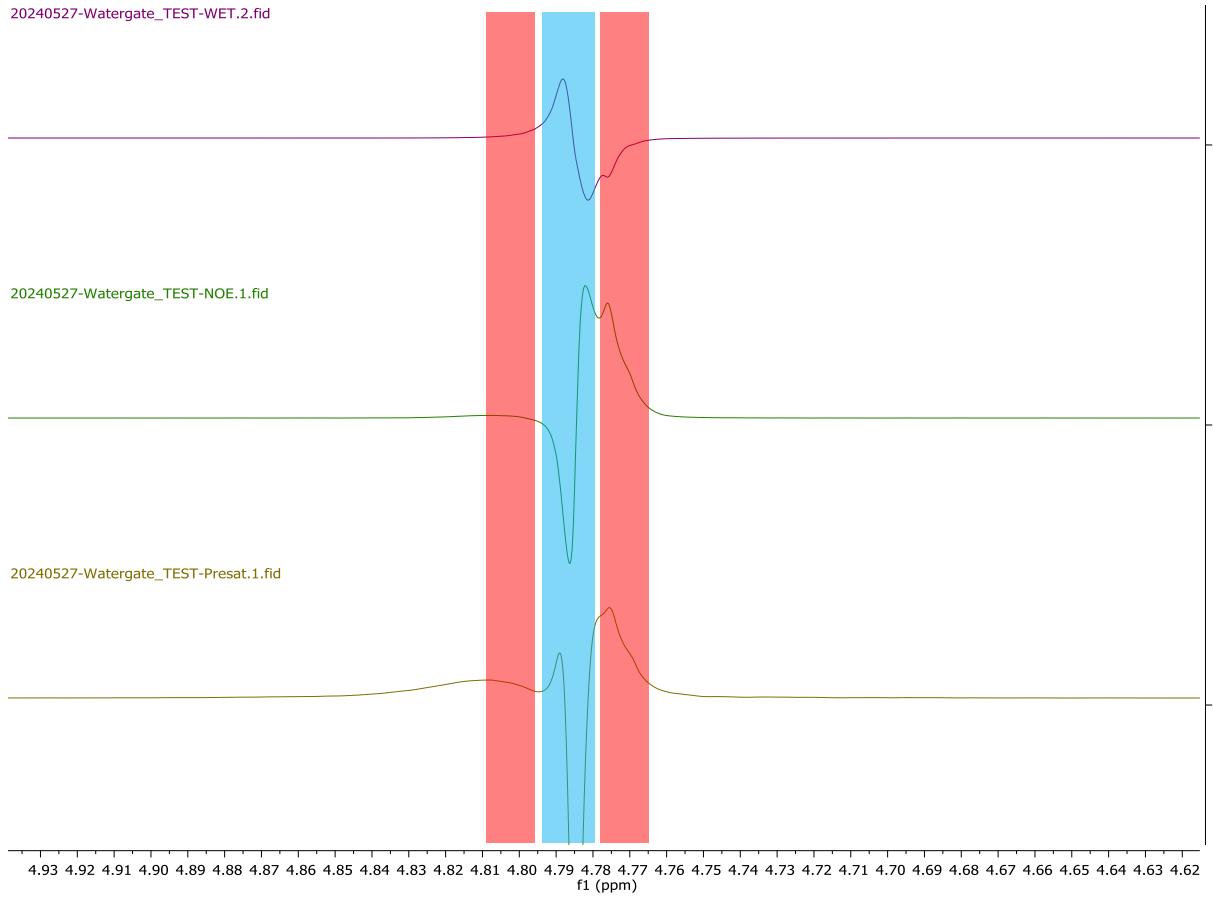
### P1331



# Far away water

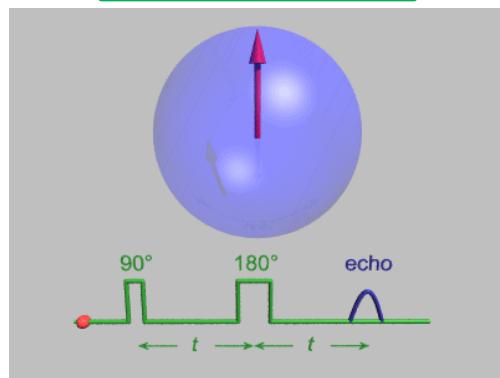
## ■ Far way water:

Coil center에서 멀어질수록 B1 field(직접 가해주는 RF pulse의 영향이 감소하므로 Residual peak이 나타남)



# Spin echo (Hahn-echo) vs Gradient echo

## SE (Spin Echo)



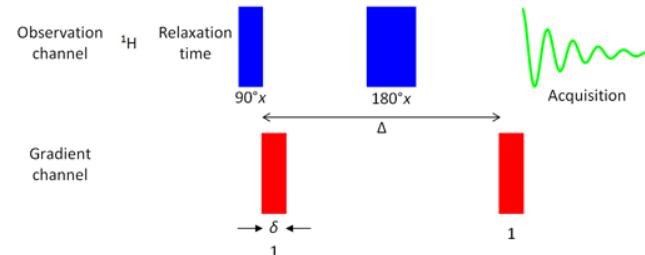
- Dephasing/Refocusing : Related T2
- Experiment time: Long

T2 time을 따라 dephasing이 일어나므로 실험시간이 길어짐

## Origin pulse sequence of DOSY

Jump-return

## 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)

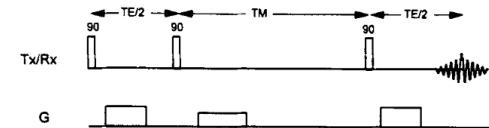


Figure 3 Pulse sequence used to generate a stimulated echo.

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

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

Watergate

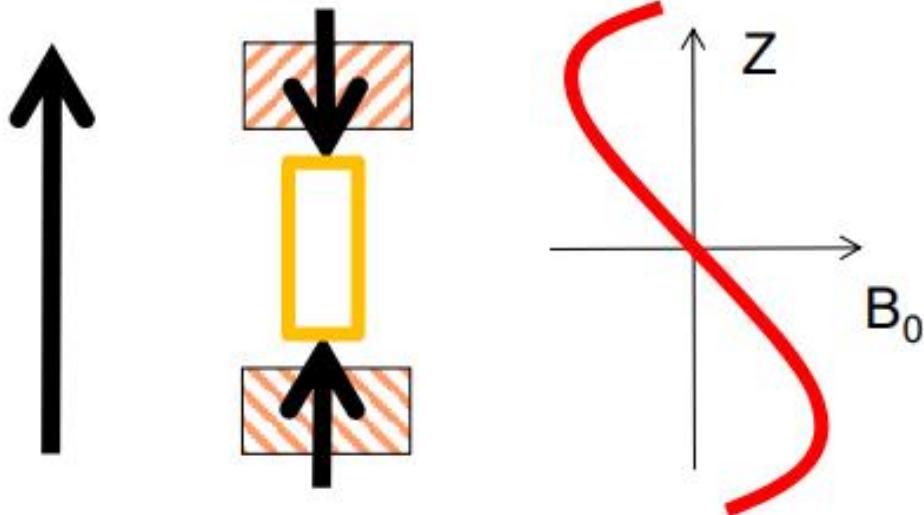
PGSTE

## 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



# Post suppression

## WATERGATE (WATER suppression by GrAdient Tailored Excitation)

p3919gp: 3-9-19 Watergate + gp (gradient pulse)

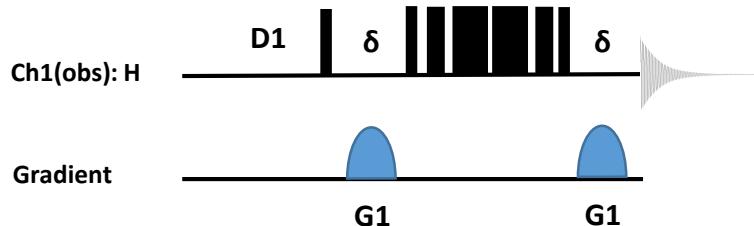
p3919fpgp: 3-9-19 Watergate + fp (flip back pulse) + gp (gradient pulse)

zggpwg: 90 degree water selective pulse

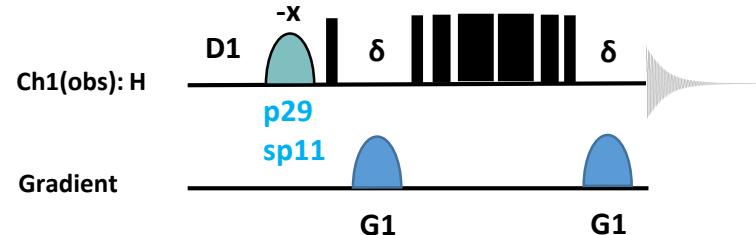
zggpw5: W5 (Similar 3-9-19, but narrow notch)

- Flip back Pulse: 실험 전 단계에  $-x$  방향으로 water의  $M_z$  값만 선택적으로 이동시킴  
**Saturation Transfer**에 의한 Exchangeable proton의 감소를 억제

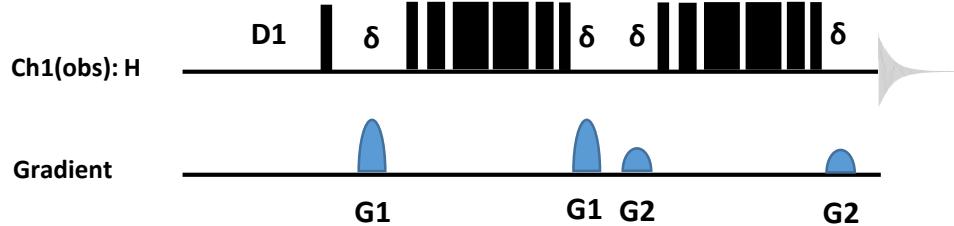
**p3919gp**



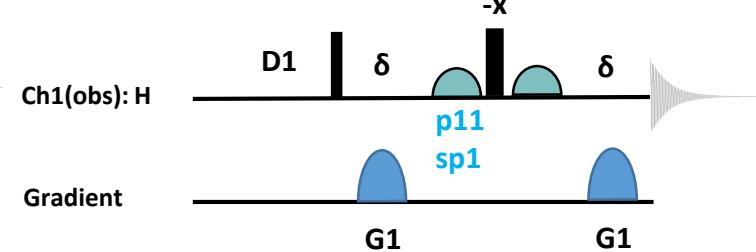
**p3919fpgp**



**zggpw5**



**zggpwg**



# ES (Excitation Sculpting)

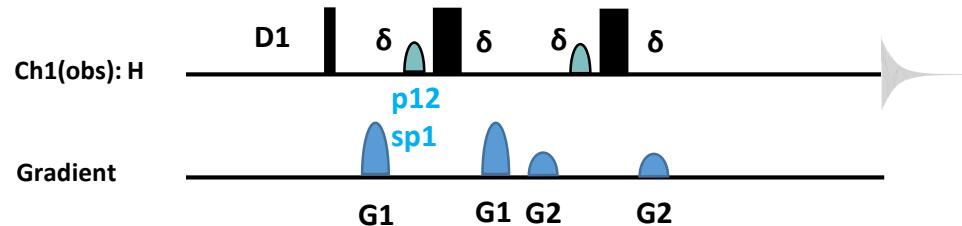
## 1D ES (Excitation Sculpting)

**zgesgp:** zg + es (excitation sculpting) + gp (gradient pulse)

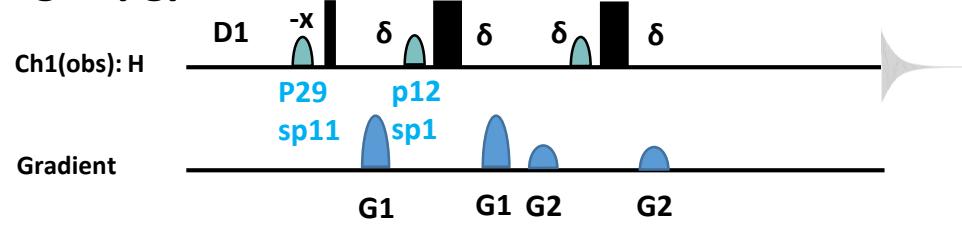
**zgesfpgp:** zg + es (excitation sculpting) + fp (flip back pulse) + gp (gradient pulse)

**zggpw5:** W5 (Similar 3-9-19, but narrow notch)

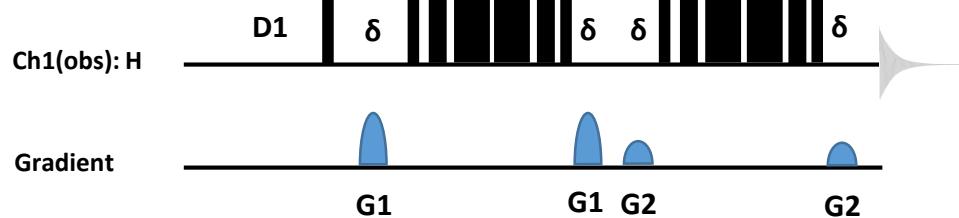
**zgesgp**



**zgesfpgp**



**zggpw5**



## Nanoscale Organisation and Dynamics

» Group Member Profiles

» Research Candidates

» Biomedical Magnetic  
Resonance Facility

» Research Projects

## NMR Pulse Sequences

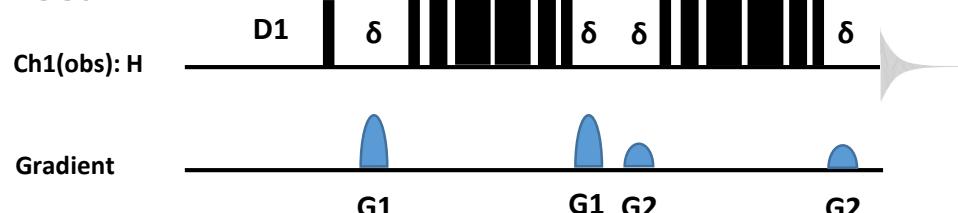
The following sequences are freely available for use. They have been developed by the Nanoscale Research Group for Bruker systems, although they can be ported to other spectrometer brands. It would be appreciated if you would cite the appropriate reference / references, when using any of the following sequences.

Expand all

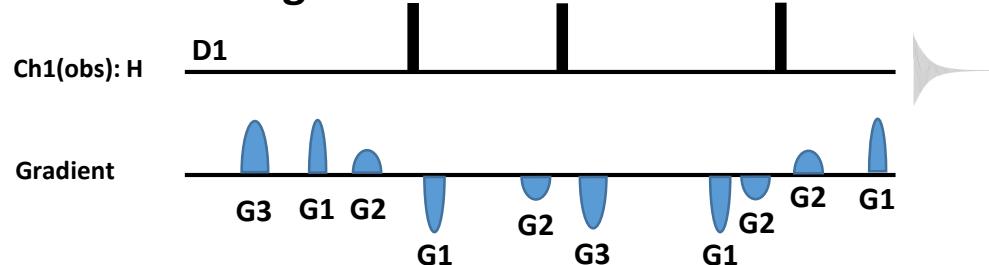
### PGSTE-WATERGATE: An STE-based PGSE NMR sequence with excellent solvent suppression

This is a stimulated-echo based pulsed gradient spin-echo NMR diffusion sequence incorporating selective inversion based solvent signal suppression. The sequence provides superb solvent suppression without any phase distortion. The sequence is simple to set up and particularly suited to measuring diffusion coefficients in aqueous solution such as is commonly required in biomolecular NMR experiments (e.g., probing drug-protein interactions).

## zggpw5



## PGSTE-Watergate



Manchester NMR Methodology Group

- Home Research People Publications Downloads Vacan

Pulse Sequences

- 1D PSYCHE
- 1D TSE-PSYCHE
- 1D Zanger-Stern
- 1D Selective TOCSY-PSYCHE
- 1D SAPPHIRE
- 1D semi-real-time pure shift
- 2DJ-ZOS-PSYCHE
- 2D F1-PSYCHE-TOCSY
- 2D rtPS-gHSQC
- GEMSTONE
- GEMSTONE-CLIP-COSY
- GEMSTONE-NOESY
- GEMSTONE-ROESY

**PE WATERGATE**

**Summary:**  
Suppression of  $J$  modulation can be achieved using the Perfect Echo to give pure absorption mode Lorentzian lineshapes. Combining the perfect echo with WATERGATE allows long, selective solvent suppression pulses to be used, giving a narrow suppression band while retaining clean NMR spectra.

## zggpw5



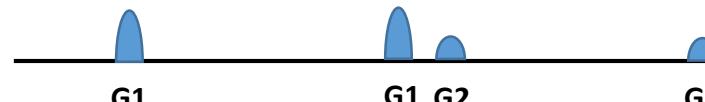
## Gradient



## pew5



## Gradient



# 1D Solvent suppression

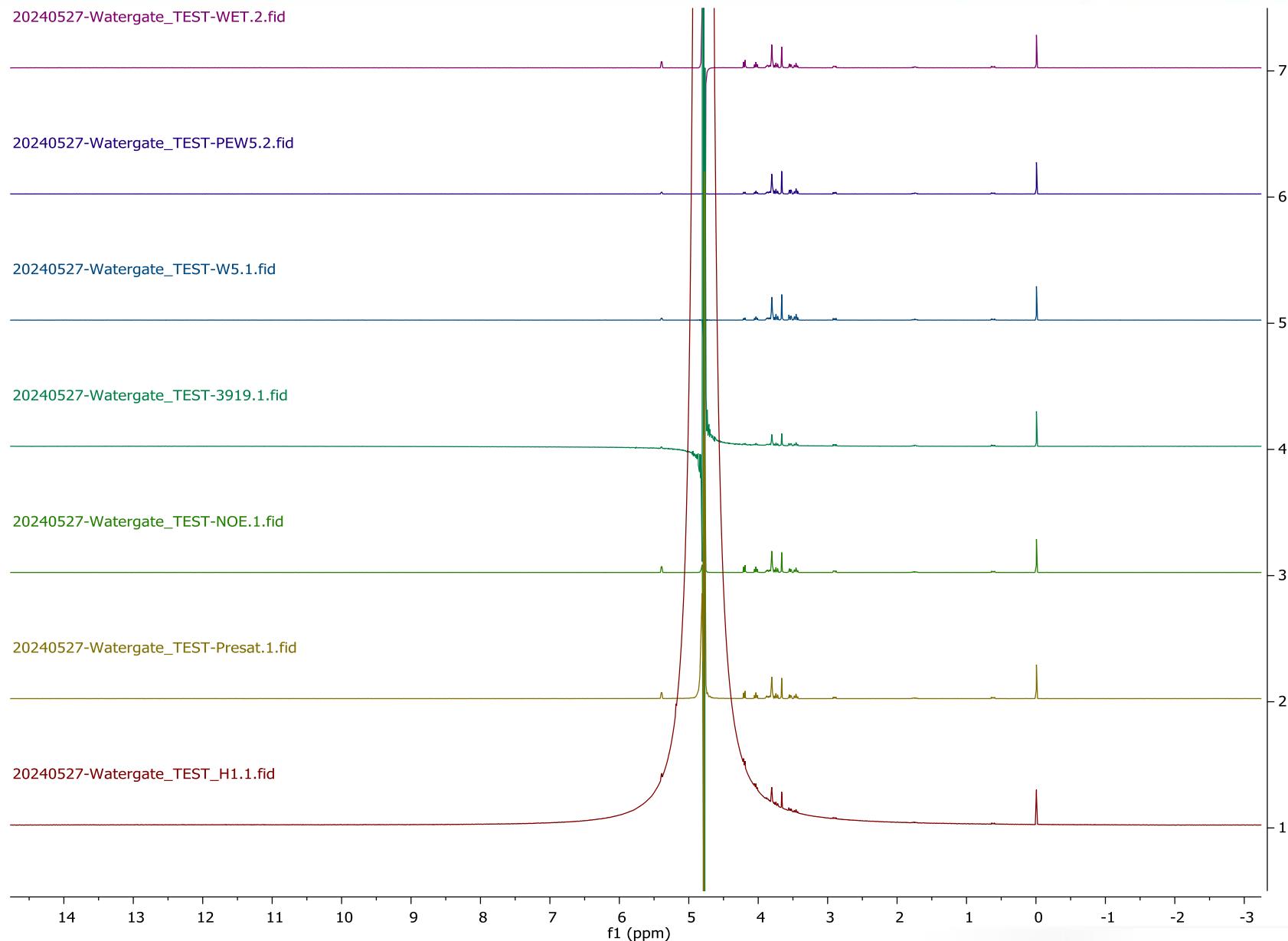
Pulse sequence	Advantage	Exchangeable Proton	Notch	Key parameter	Remark
Presat	<ul style="list-style-type: none"> <li>▪ Cleanest</li> <li>▪ Minimized saturation peak</li> <li>▪ Possible multi suppression</li> </ul>	Highly Reduced	Narrow	O1 $\pi_{lw9} \leq 0.3\text{mW}$	<ul style="list-style-type: none"> <li>▪ Solvent suppression</li> <li>▪ Follow T2</li> </ul>
noesy1d (noesygppr1d)	<ul style="list-style-type: none"> <li>▪ Clean baseline</li> </ul>	Reduced	Narrow	O1 $D8 \leq 1\text{ms}$	<ul style="list-style-type: none"> <li>▪ Common purpose</li> <li>▪ for metabolite</li> </ul>
Watergate (3-9-19)	<ul style="list-style-type: none"> <li>▪ Water suppression</li> </ul>	No effect (+ Flip back)	Slightly Narrow		<ul style="list-style-type: none"> <li>▪ for NH4+, Acid, Amine</li> <li>▪ Check D19</li> <li>(Delay for binomial water suppression)</li> </ul>
Watergate (W5)	<ul style="list-style-type: none"> <li>▪ Water suppression</li> </ul>		Narrow		
Purge	<ul style="list-style-type: none"> <li>▪ Quantitative</li> </ul>				<ul style="list-style-type: none"> <li>▪ for Quantitative NMR</li> </ul>
WET	<ul style="list-style-type: none"> <li>▪ Multi suppression</li> </ul>	Possibly reduced			<ul style="list-style-type: none"> <li>▪ Follow T1 effects</li> </ul>
ES (Excitation Sculpting)	<ul style="list-style-type: none"> <li>▪ Use Gradient/spin echo</li> <li>▪ Excellent suppression</li> </ul>		Broad		
PGSTE (Pulsed Gradient Stimulated Echo)	<ul style="list-style-type: none"> <li>▪ Use Gradient/spin echo</li> <li>▪ Excellent suppression</li> </ul>	No effect	Excellent Narrow		<ul style="list-style-type: none"> <li>▪ Advanced sequence</li> </ul>
PEW5	<ul style="list-style-type: none"> <li>▪ Clean spectra</li> </ul>	No effect	Excellent Narrow		<ul style="list-style-type: none"> <li>▪ for NH4+, Acid, Amine</li> <li>▪ Advanced sequence</li> </ul>

[https://ethz.ch/content/dam/ethz/special-interest/biol/mol-biol/bnsp-am/NMRSeminar2018HS/Water\\_suppression\\_pwu\\_20181017.pdf](https://ethz.ch/content/dam/ethz/special-interest/biol/mol-biol/bnsp-am/NMRSeminar2018HS/Water_suppression_pwu_20181017.pdf)

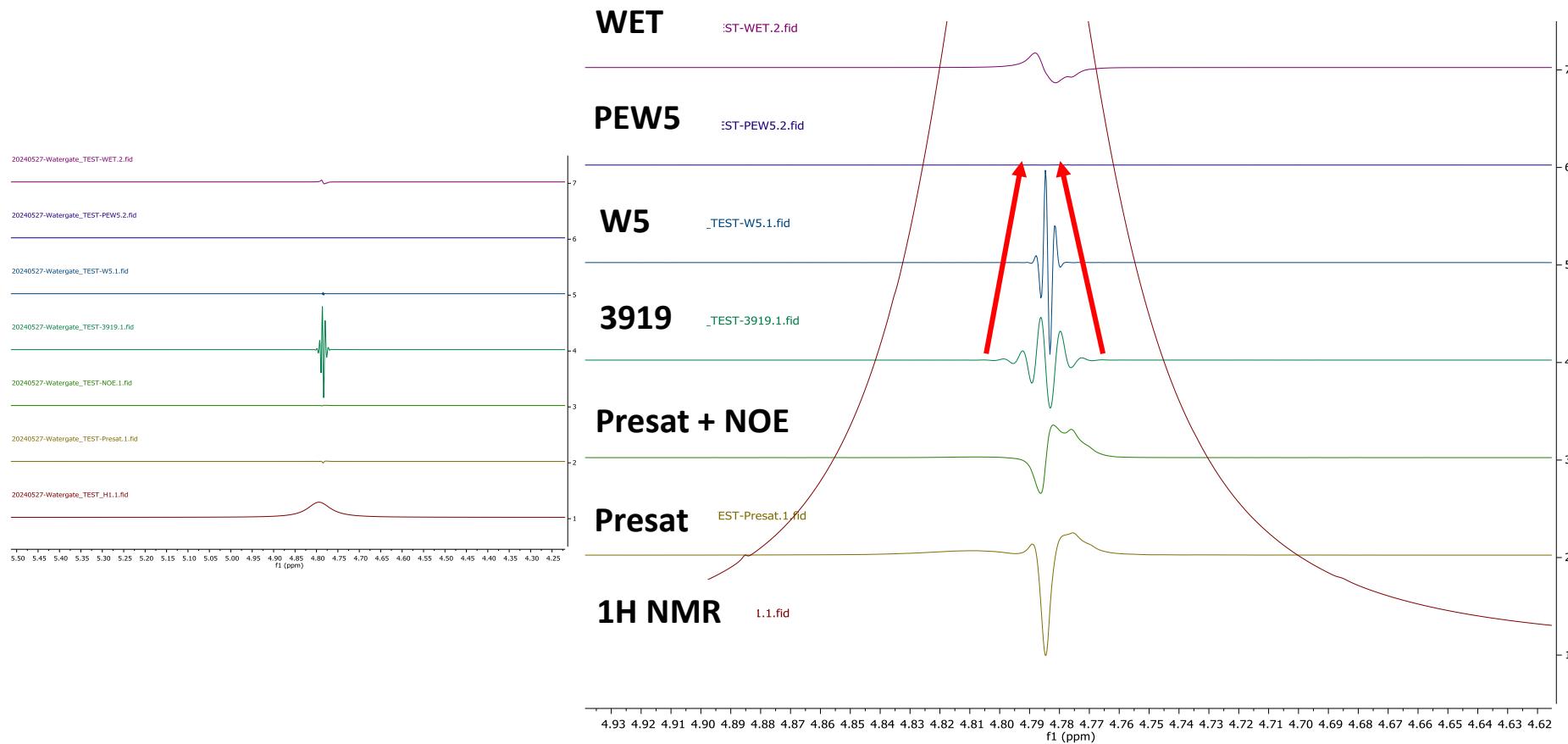
<https://www.nmr.chemistry.manchester.ac.uk/?q=node/285>

[https://www2.chem.wisc.edu/~cic/nmr/Guides/Ba3vug/AV3\\_SolventSuppression.pdf](https://www2.chem.wisc.edu/~cic/nmr/Guides/Ba3vug/AV3_SolventSuppression.pdf)

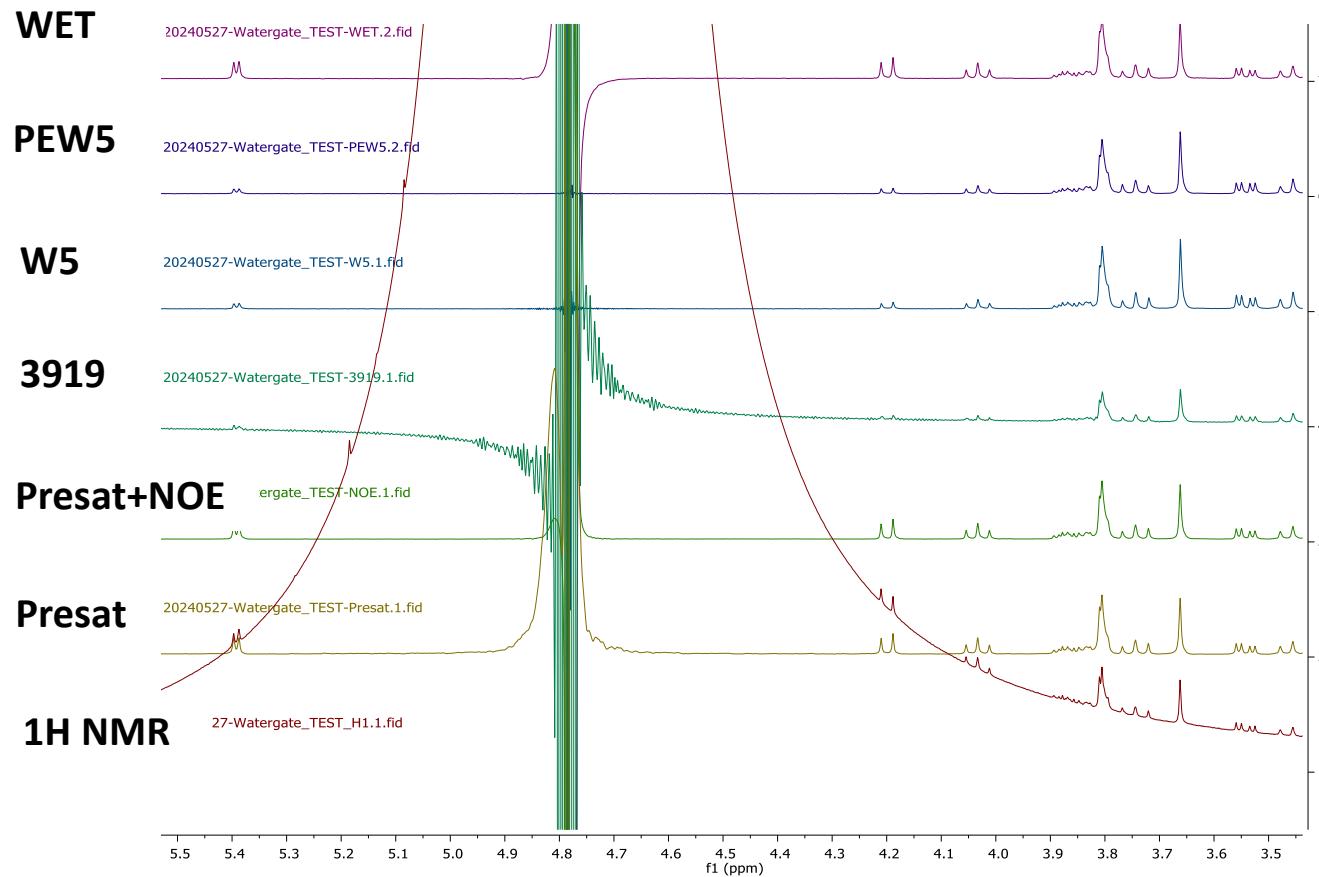
# 1D Solvent suppression



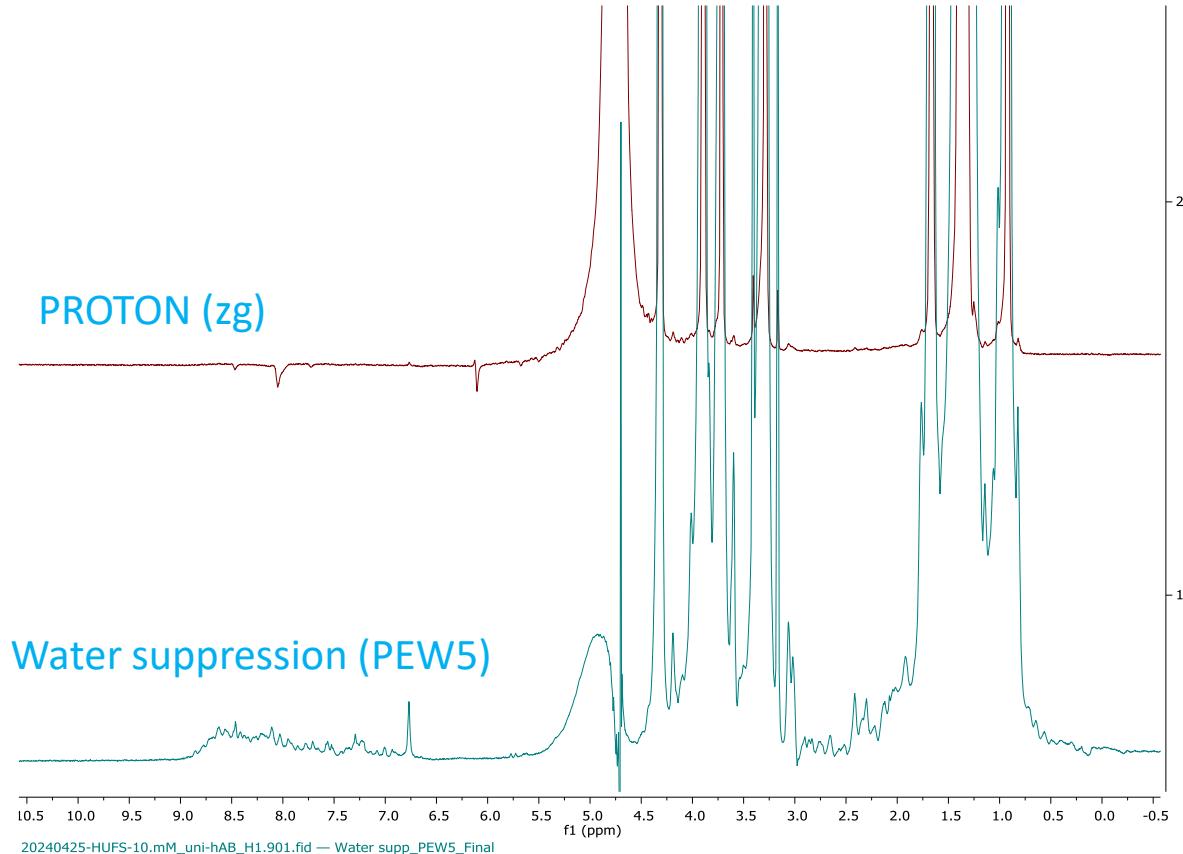
# Spectrum for notch



# Spectrum for signal intensity



# Protein for water suppression



# Pulse sequence in UNIST

N Presat+NOE_UNIST	Solvent suppression (Com)			=				UCRF
<b>N PROTON_UNIST      1H experiment</b>								
N T1_short	5min							
N T1_Long	1hr							
N SST1IR	Single scan_T1_experiment							
N C13CPD_UNIST	90degree+_NOE(Signal Enhanced)_for_Intensity							
N C13IG_UNIST	90_degree_-NOE_for_Integration							
N C13DEPT135	13C DEPT135, CH3/CH positive, CH2 negative, 256 scans, 160 ppm							
N PROTON_PRESAT	Solvent_Suppression							
N Presat+NOE_UNIST	Solvent suppression (Common purpose)							
N Watergate_3919_UNIST	Water suppression w 3-9-19							
N WG-3919FP_UNIST	with FP (for exchange)							
N Watergate_W5_UNIST	Water suppression for narrow notch							
N WET_UNIST	Water suppression for quantitative NMR							
N PEW5_UNIST	Perfect echo + W5 for eliminated suppression peak							
C COSYGPSW	Gradient selected COSY							
C COSYGPDPHWSW	Gradient selected double quantum filtered phase sensitive COSY							
C FASTLANE_HSQC	Combined analysis of 1D1H and HSQC data							
C FASTLANE_HSQC_HMBC	Combined analysis of 1D1H, HSQC and HMBC data							
C FASTLANE_HSQC_13C_HMBC	Combined analysis of 1D1H, HSQC, 13C and HMBC data							
C HMBCGP	1H-13C HMBC with gradient selection							

Name in Bruker NMR	Pulse sequence	Purpose	Remarks
Proton_PRESAT	Presaturation	Solvent suppression (with water suppression)	Simple, any solvent, multi suppression
Presat+NOE_UNIST	noesy1d (noesygppr1d)	Solvent suppression (with water suppression)	More effective sequence
Watergate_3919_UNIST	Watergate (3-9-19)	Water suppression	
WG-3919FP_UNIST	Watergate (W5)	Water suppression	Flip back (No effect exchangeable H)
WET_UNIST	WET	Solvent suppression (with water suppression)	Good baseline, any solvent, multi suppression
PEW5_UNIST	Perfect echo W5	Water suppression	Smallest notch!

# Important parameter (O1P)

The screenshot shows the IconNMR software interface. At the top, there is a toolbar with icons for File, Run, Holder, View, Parameters (highlighted with a yellow box and a red number '1'), Options, Tools, and Help. Below the toolbar is a menu bar with the same items. A dropdown menu for 'Parameters' is open, showing options: Edit all Acquisition Parameters, Edit Processing Parameters, Pertinent Acquisition Parameter Editor, and User Specific Commands. The main window is titled 'NMR - Auto'. It contains an 'Experiment Table' with columns for Holder, T., Status, Name, No., and Solvent. There are eight rows of data, each with a status icon (e.g., 'Finished', 'Available') and a name like '240805JK\_1mM\_LiF\_H2O'. Row 7 has a different name ('dsfsdfsdfs'). The 'Solvent' column shows values like 'None' or 'no'.

A screenshot of a software dialog box titled 'Automation'. It contains four entries:

AUNM	au_watersc supp_opt=1	...	E	Acquisition AU program
PYNM		...	E	Acquisition Jython program
EXP	PEWS_UNIST			Experiment performed
TUBE_TYPE				Type of used sample tube

**au\_watersc : acquisition program in auto**  
**bt\_calib=1 : Automatically find O1**  
**pcal=1 : Automatically find P1**  
**supp\_opt=1 : Automatically find O1 and P1 standard set is supp\_opt=1**

**O1P(O1 ppm) related to suppression effectivity!**  
**[Solvent peak center = O1P]**

**Already known solvent chemical shift..**

- 1) Just set au\_watersc
- 2) Directly change O1P (ex. H<sub>2</sub>O+D<sub>2</sub>O = 4.7 ppm)

