

# MRI PRACTICALS ON CIBM PRECLINICAL IMAGING SYSTEMS

## BASICS OF MRSI

Cristina Cudalbu

*CIBM PCI EPFL*

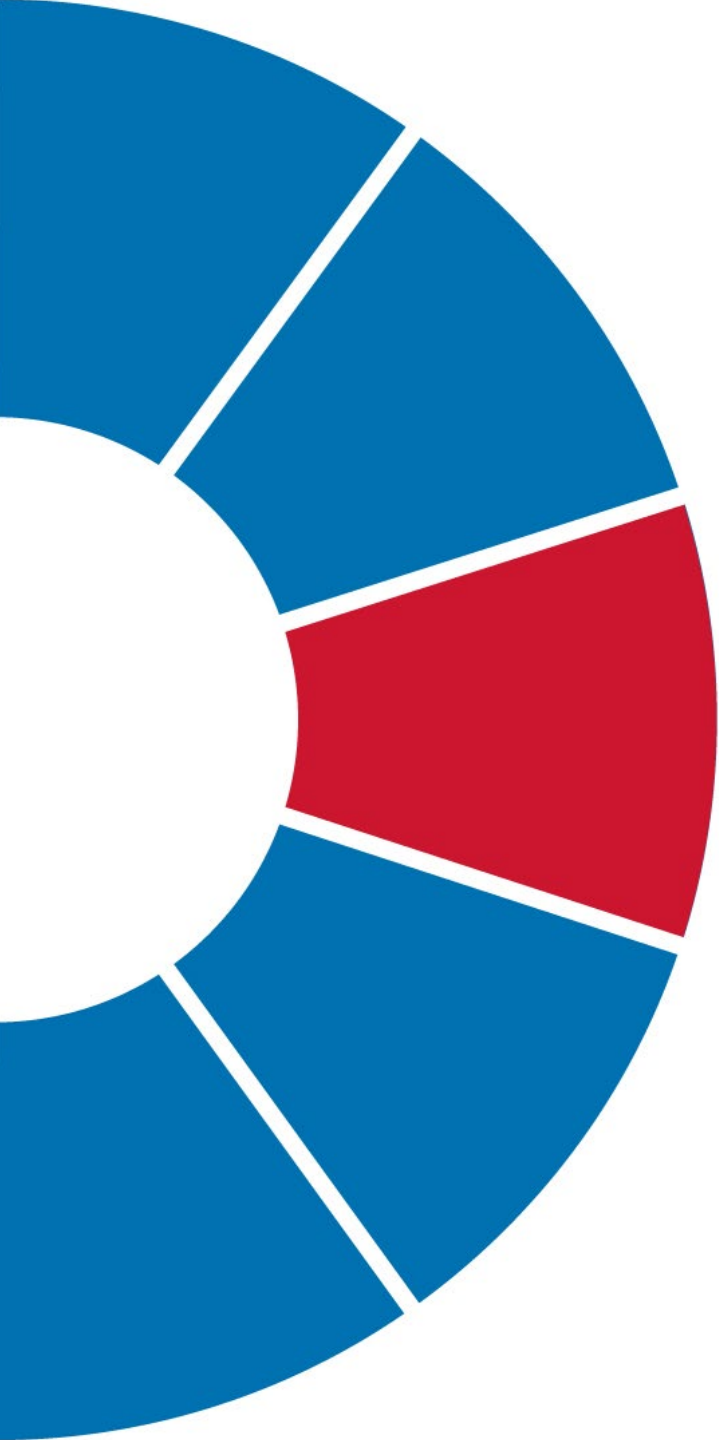
*Master Course 2025-2026*

*[cristina.cudalbu@epfl.ch](mailto:cristina.cudalbu@epfl.ch)*



# OUTLINE

- From MRS to MRSI – the magic formula
- Spatial encoding for MRSI
- MRSI advantages and challenges
- From square maps to full brain coverage

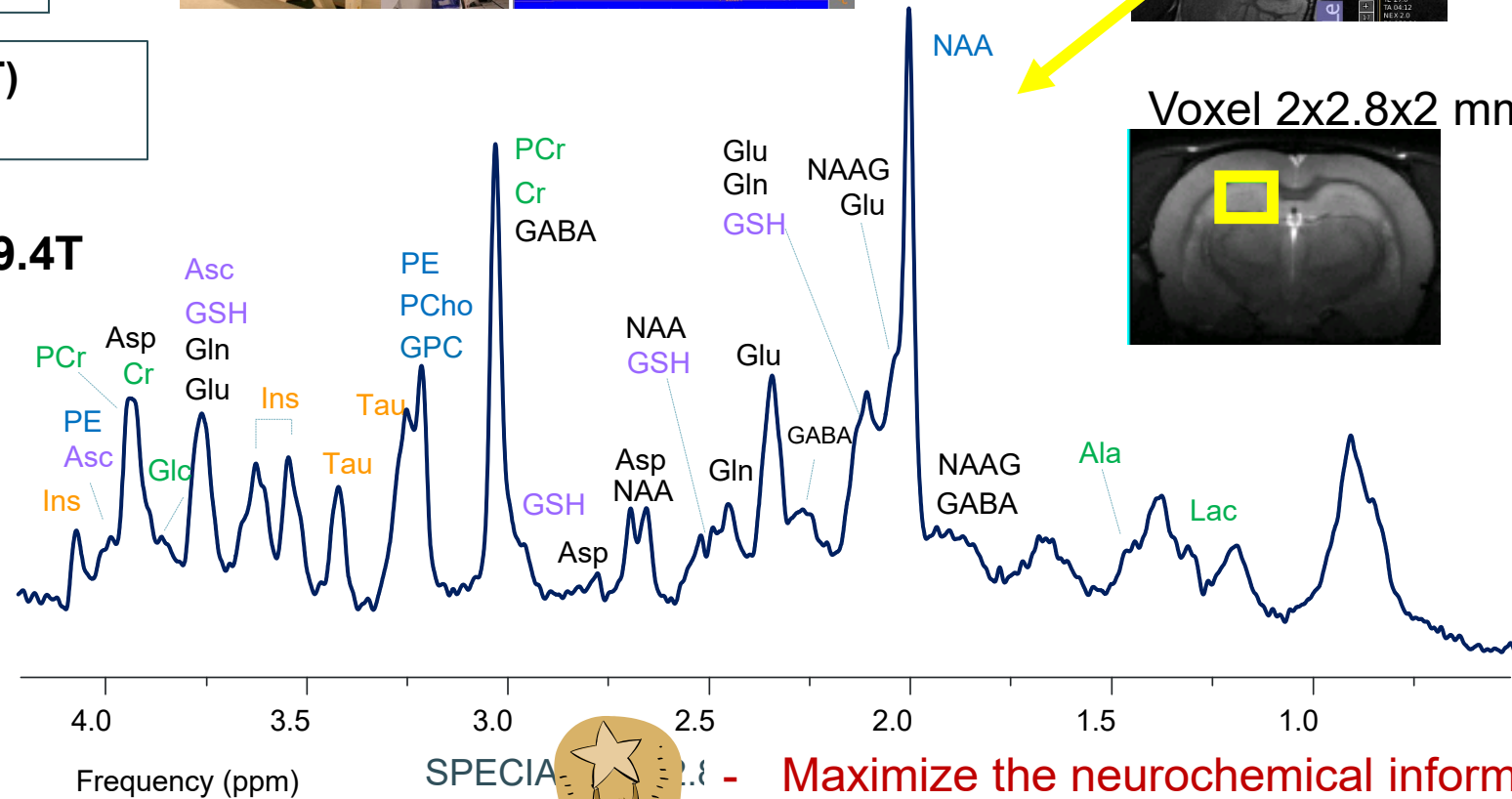
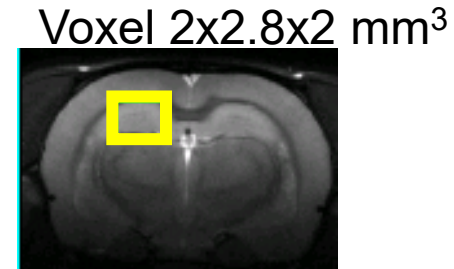
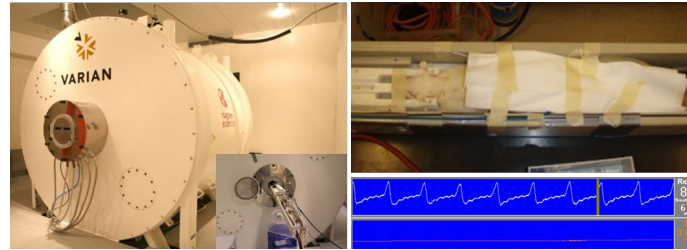


## FROM MRS TO MRSI – THE MAGIC FORMULA

# WHAT DO WE MEASURE: $^1\text{H}$ MRS ?

- *in vivo*
- non invasively
- localized in hippocampus

- at high magnetic field (9.4T)
- ultra short TE (2.8ms)



## Neurochemical Profile at 9.4T

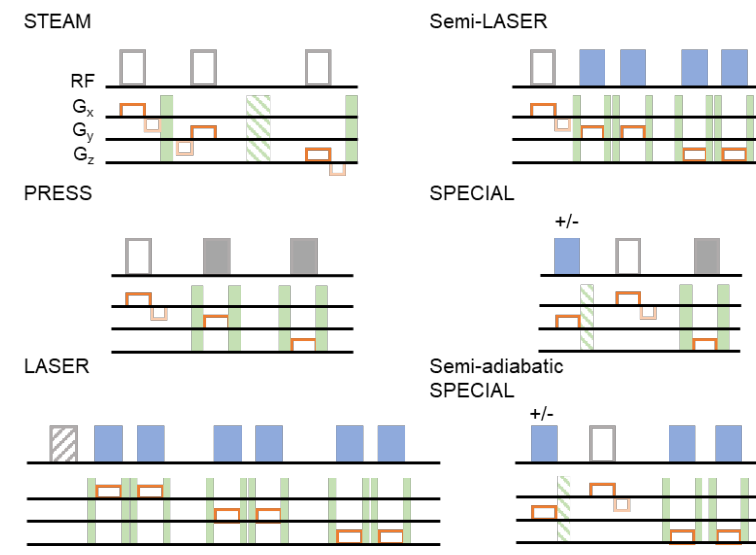
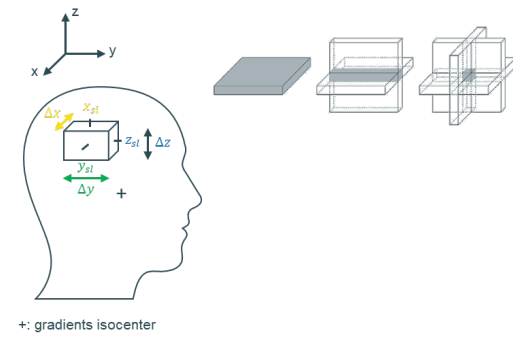
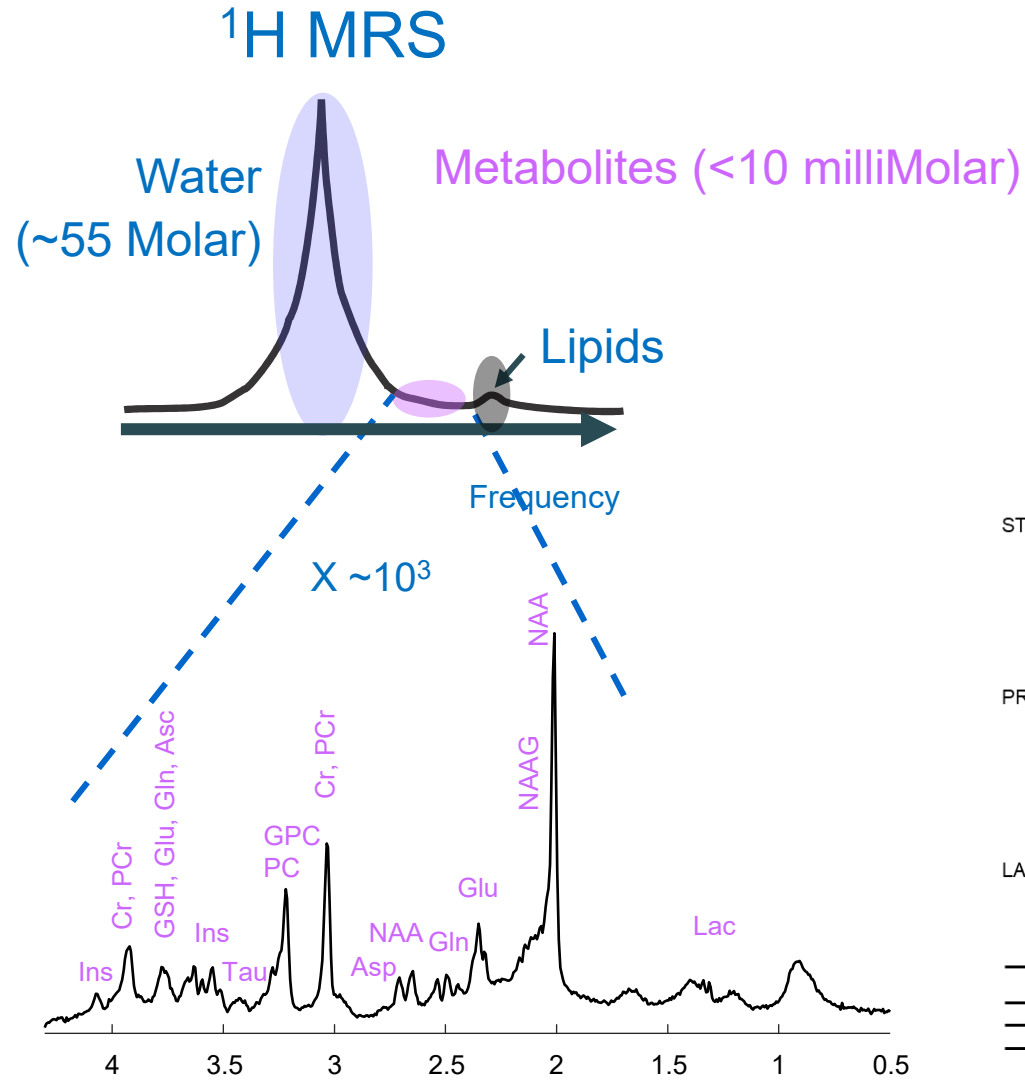
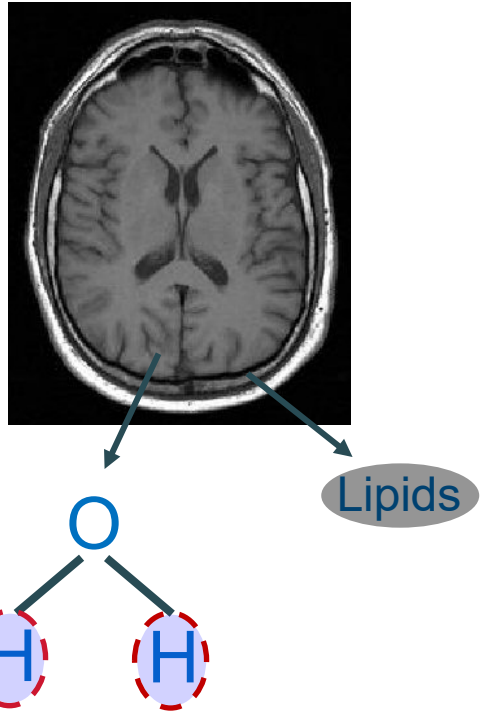
- >18 Markers of :
- Myelination/Cell proliferation
  - Energy metabolism
  - Osmoregulation
  - Neurotransmitter metabolism
  - Antioxidants



- Maximize the neurochemical information
- Increase precision, accuracy – quantification
- Increase the reliability of obtained concentrations

# BASICS OF MRS

## MRI



## BRAIN METABOLISM – *in-vivo* and non-invasively

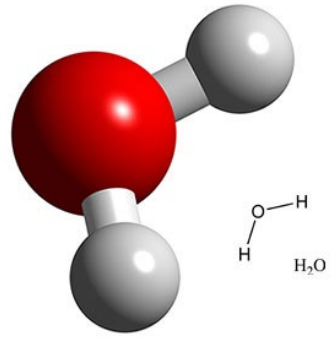
J Mosso

CIBM.CH

Ivan Kirov, ENIGMA-U  
[https://www.youtube.com/playlist?list=PL1T aL2CueZ-DOznKgKWdg\\_mHRK9UUiQoJ](https://www.youtube.com/playlist?list=PL1T aL2CueZ-DOznKgKWdg_mHRK9UUiQoJ)

# MAGNETIC RESONANCE IMAGING -> SPECTROSCOPY

MRI:  
Imaging  $^1\text{H}$  of water



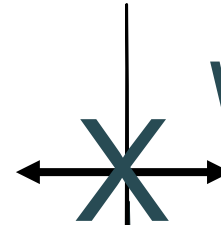
WATER IN THE HUMAN BODY

Brain	75% Water
Blood	83% Water
Heart	79% Water
Bones	22% Water
Muscles	75% Water
Liver	85% Water
Kidneys	83% Water

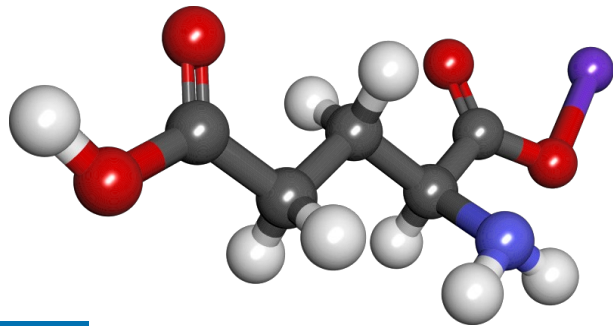
usgs.gov

~ 55 M

Water suppression

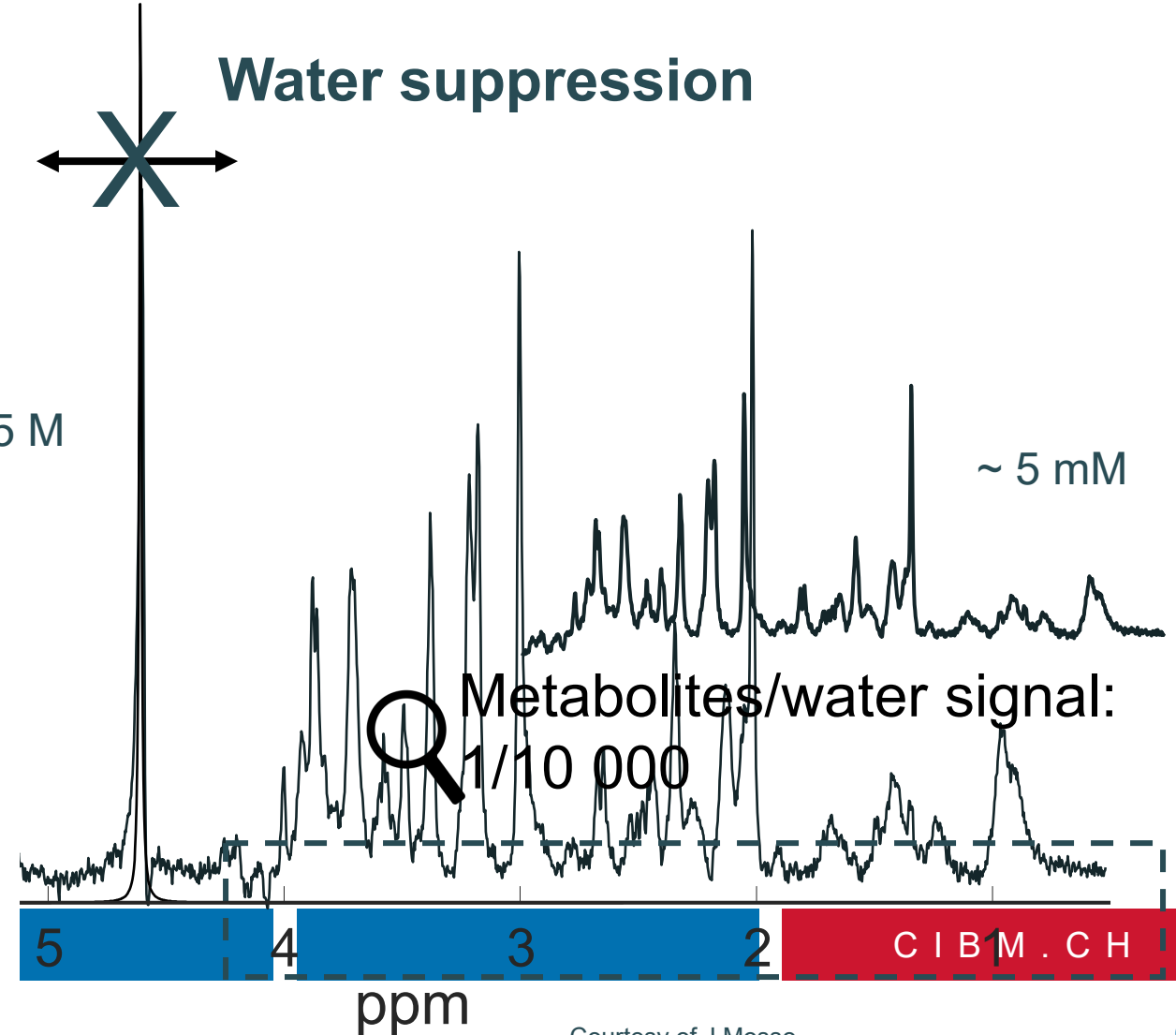
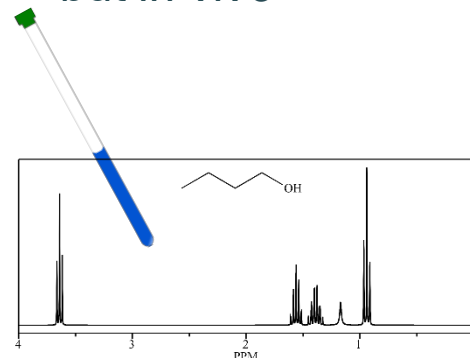


MRS:  
Measuring  $^1\text{H}$  of biomolecules

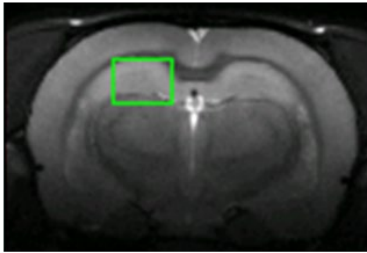


glutamate

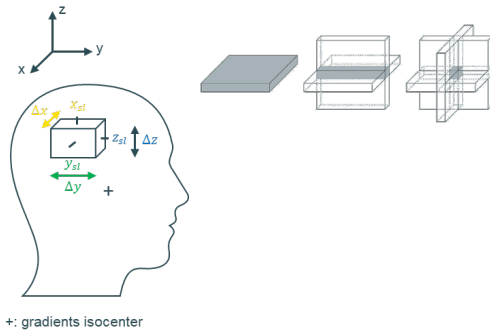
As chemical NMR...  
but *in vivo*



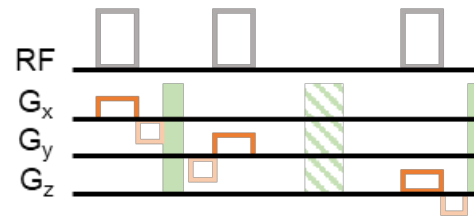
# UNTIL NOW



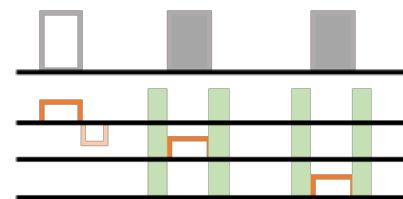
T<sub>2</sub> weighted image of the rat brain: VOI in the hippocampus



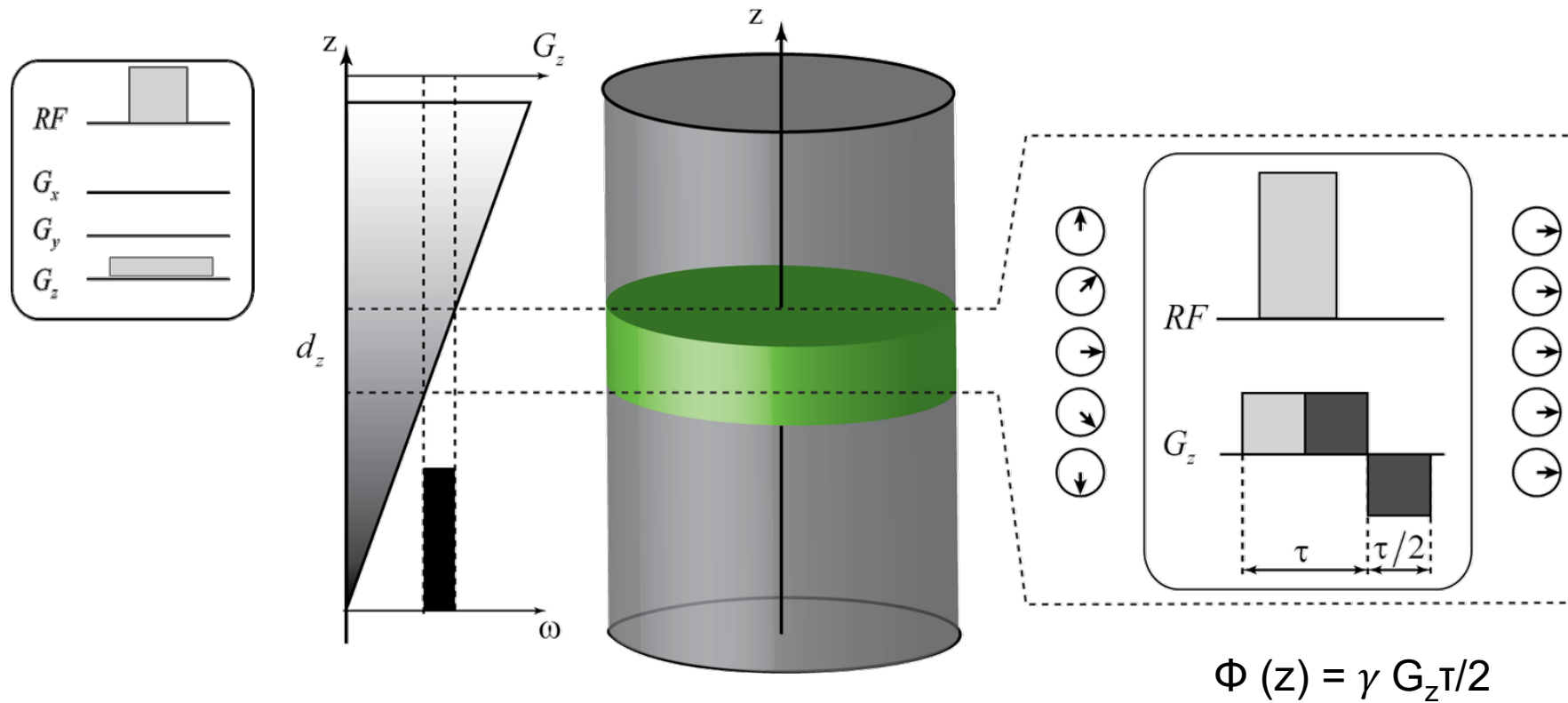
## STEAM



## PRESS



# SPATIAL ENCODING – SLICE SELECTION



$$S \propto \iint_{\text{slice}} \rho(x, y) dx dy$$

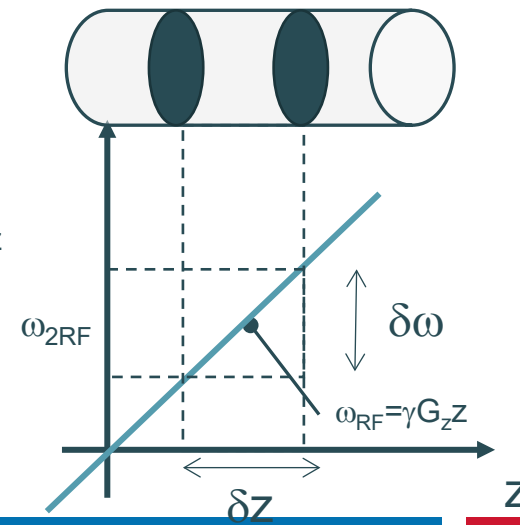
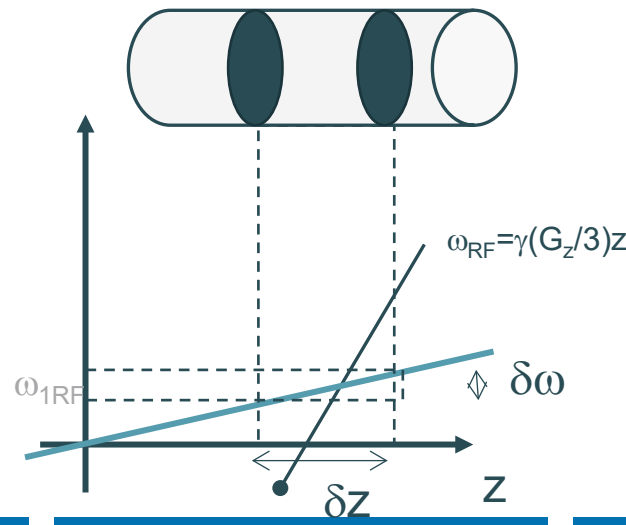
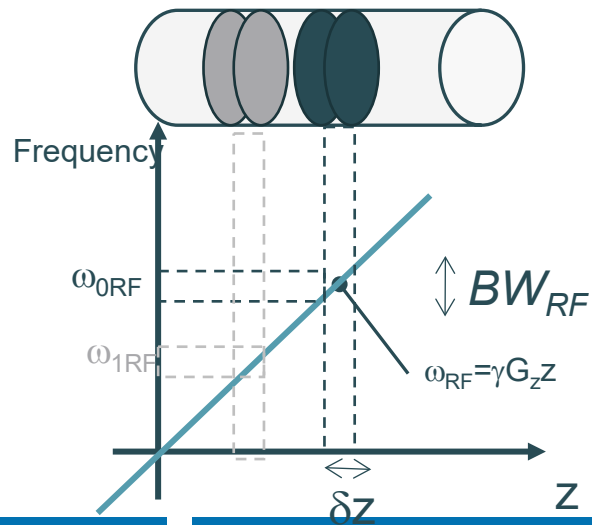
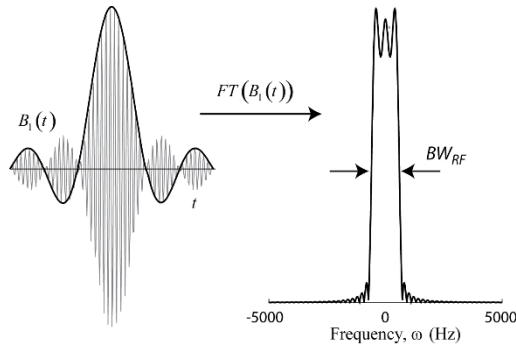
$$\delta z = \frac{BW_{RF}}{\gamma G_z}$$

# SLICE SELECTION

- Gradient is applied during an RF-pulse, which is characterized by its excitation bandwidth ( $BW_{RF}$ )

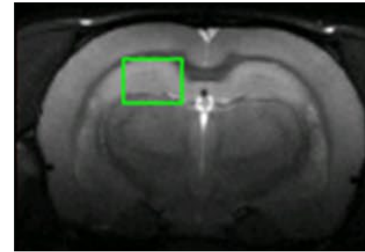
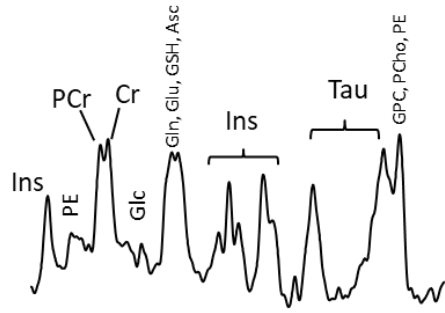
$$\delta Z = \frac{BW_{RF}}{\gamma G_z}$$

- The slice thickness can be changed either by adjusting the pulse length or the gradient amplitude



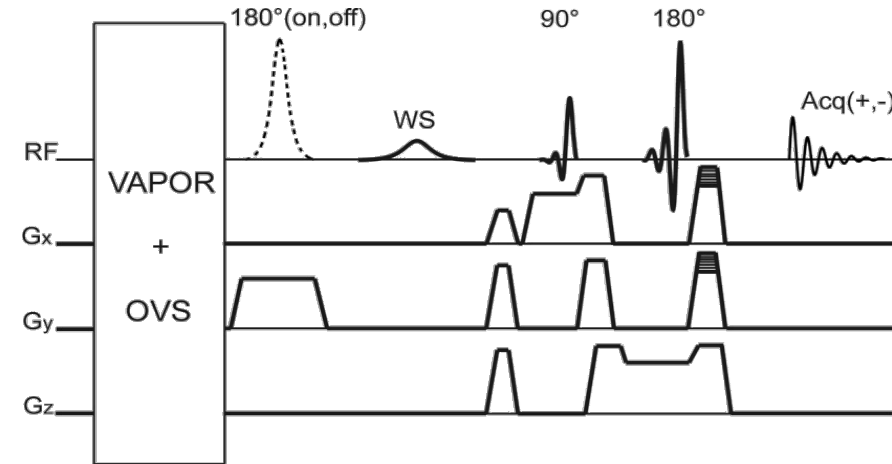
# UNTIL NOW

High spectral resolution

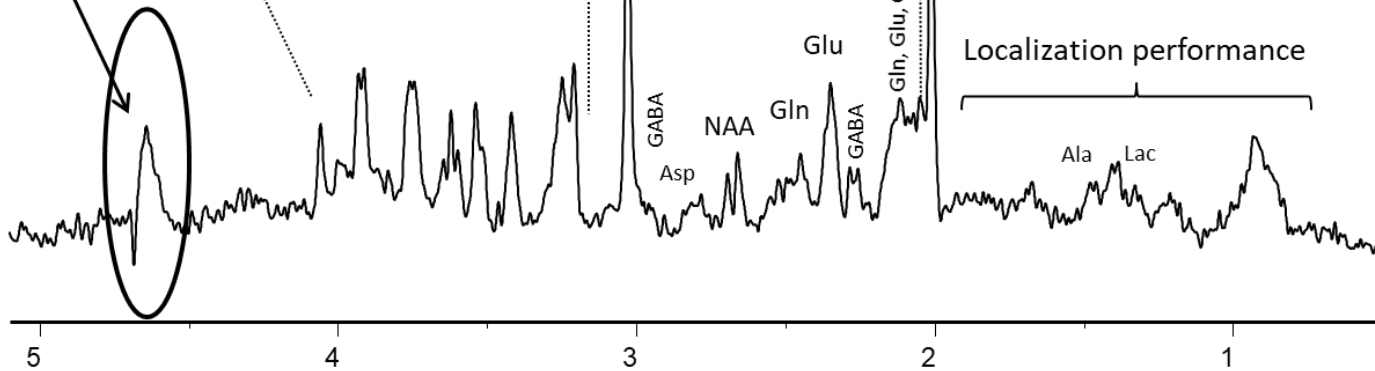


T<sub>2</sub> weighted image of the rat brain: VOI in the hippocampus

SPECIAL sequence



Efficient water suppression

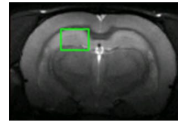
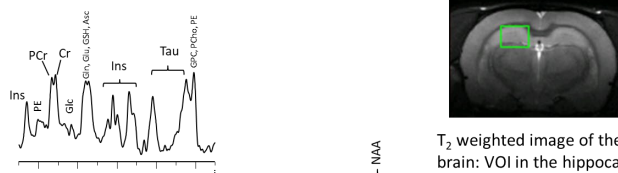


Localization performance

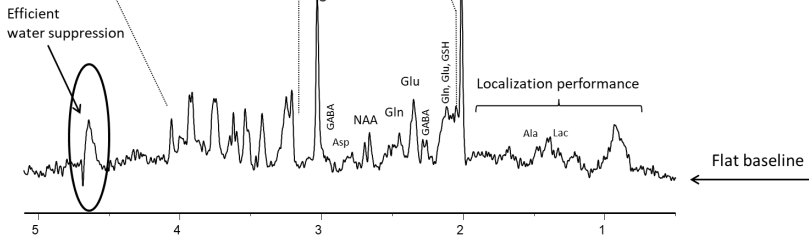
Flat baseline

# UNTIL NOW

High spectral resolution

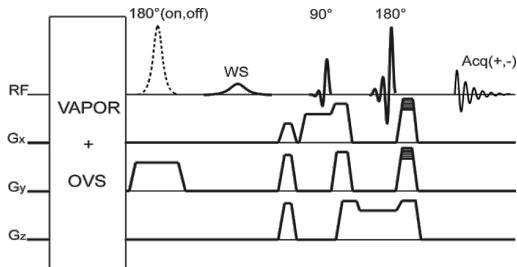


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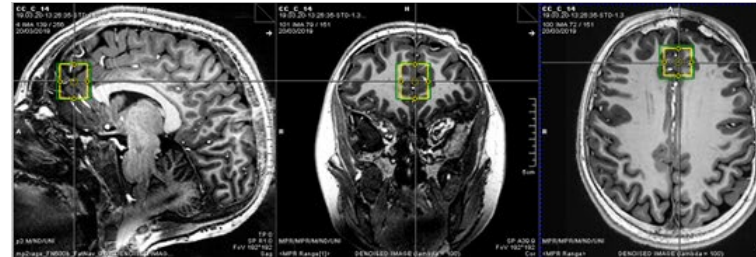


Localization performance

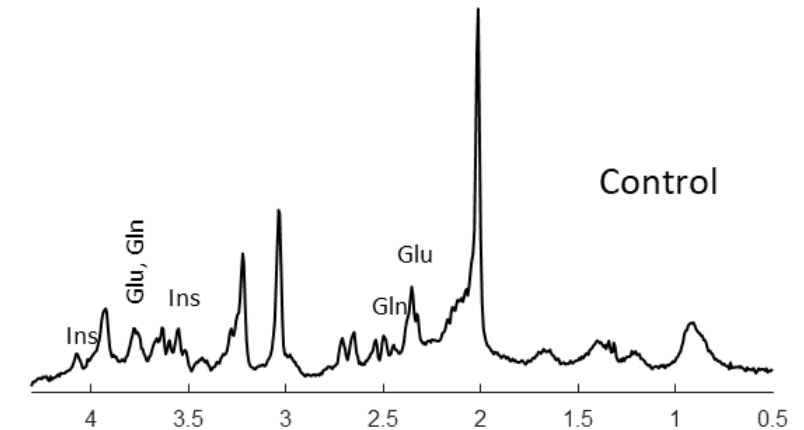
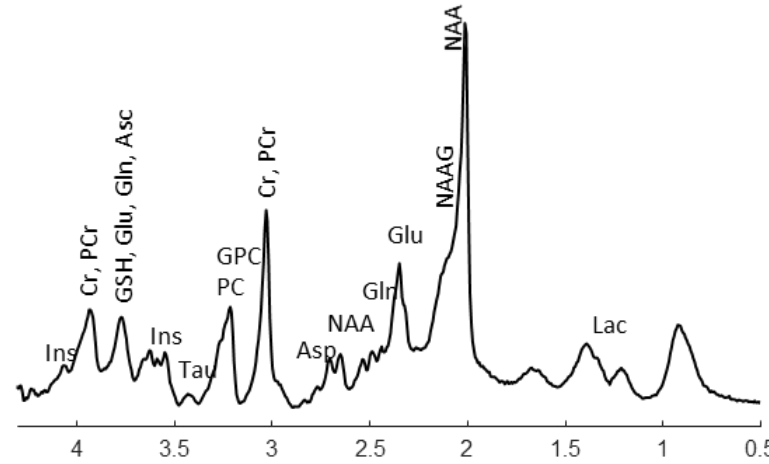
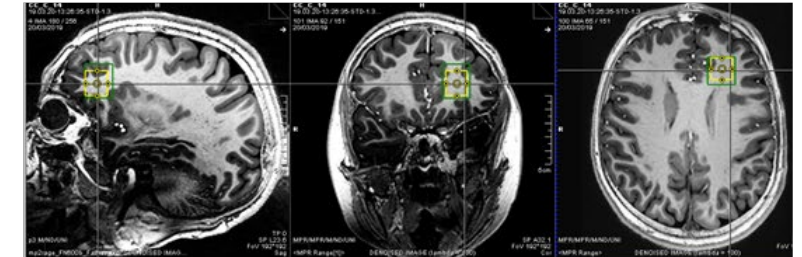
Flat baseline



## Gray Matter

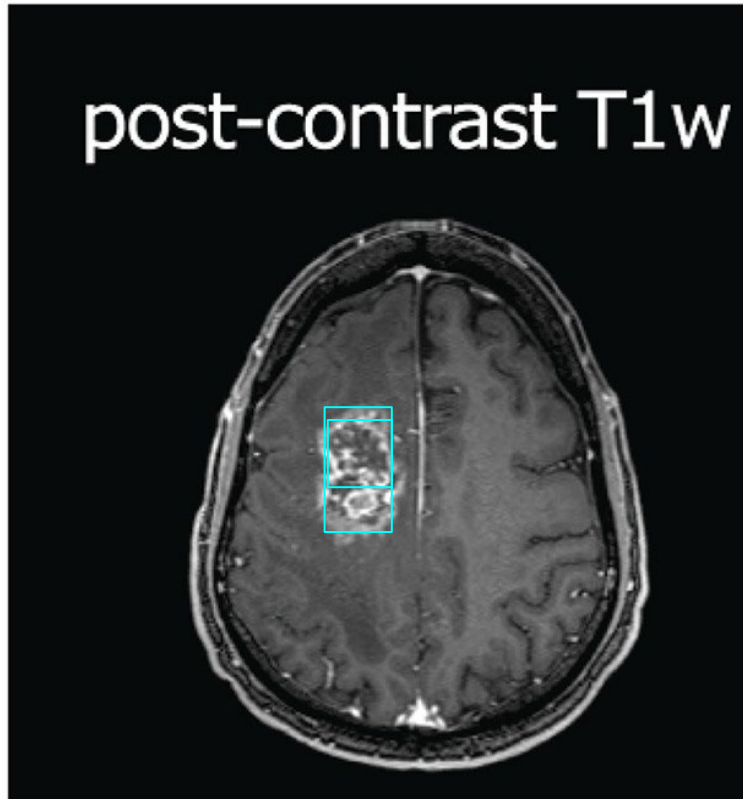


## White Matter



Control

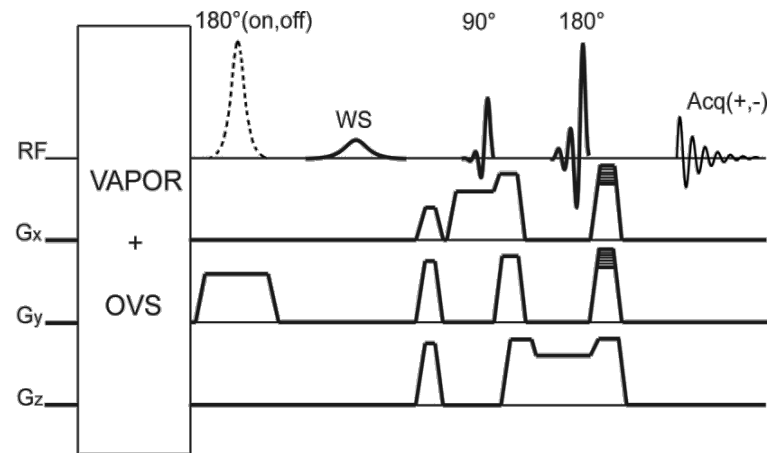
# BUT WHAT IF .....



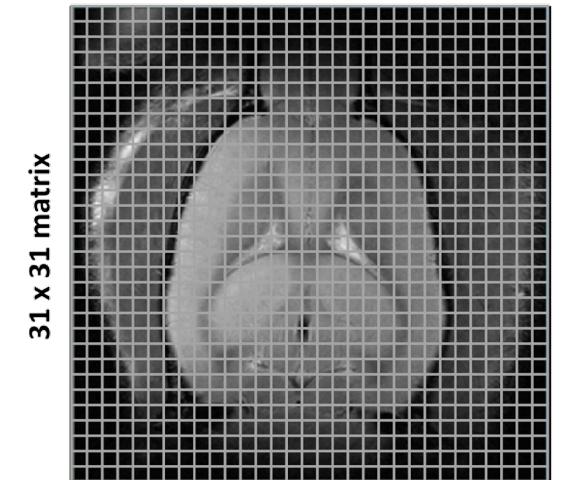
What to do –any idea ?

Multiple voxels? With multiple spectra inside ?

Extract each metabolite and have a color map



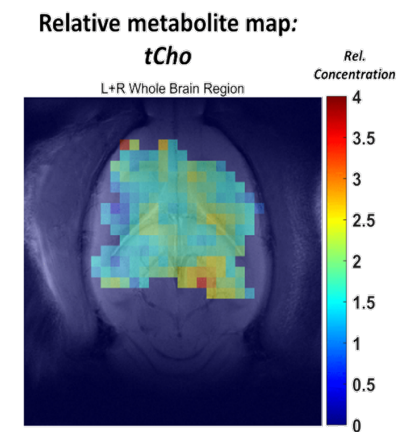
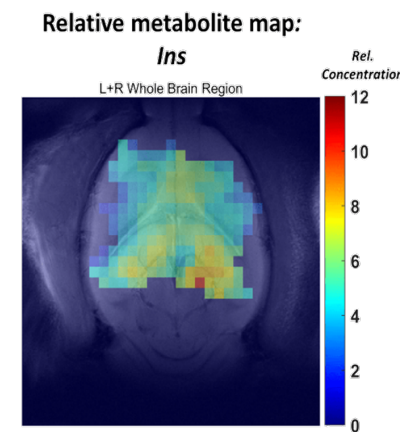
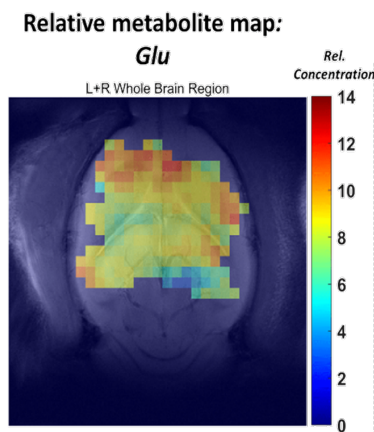
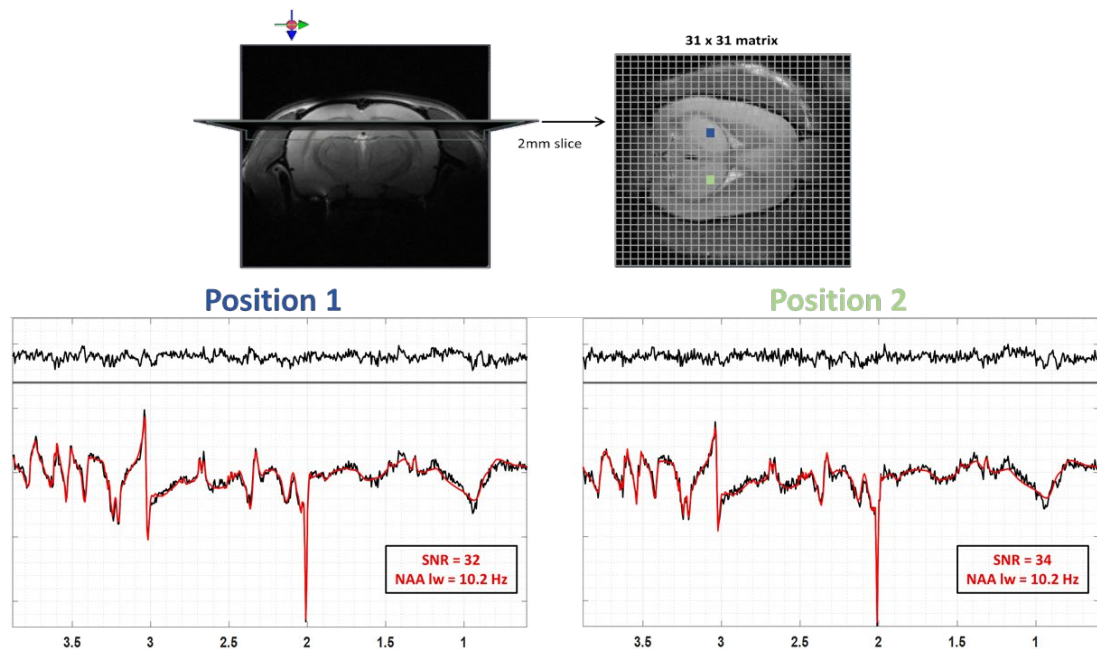
Mlynárik V, et al, MRM 2008

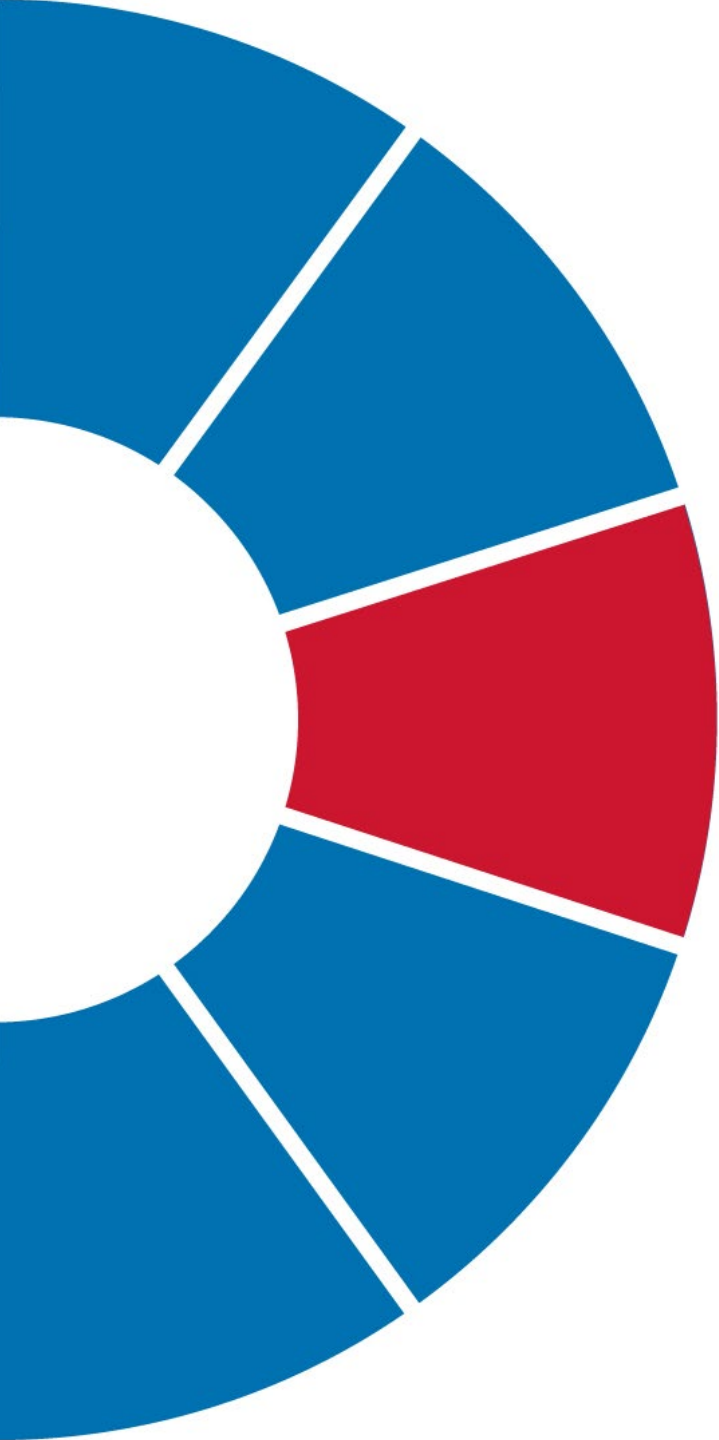


[Advanced magnetic resonance spectroscopic neuroimaging: Experts' consensus recommendations - Maudsley - 2021 - NMR in Biomedicine - Wiley Online Library](#)

# BUT WHAT IF .....

## Extract each metabolite and have a color map



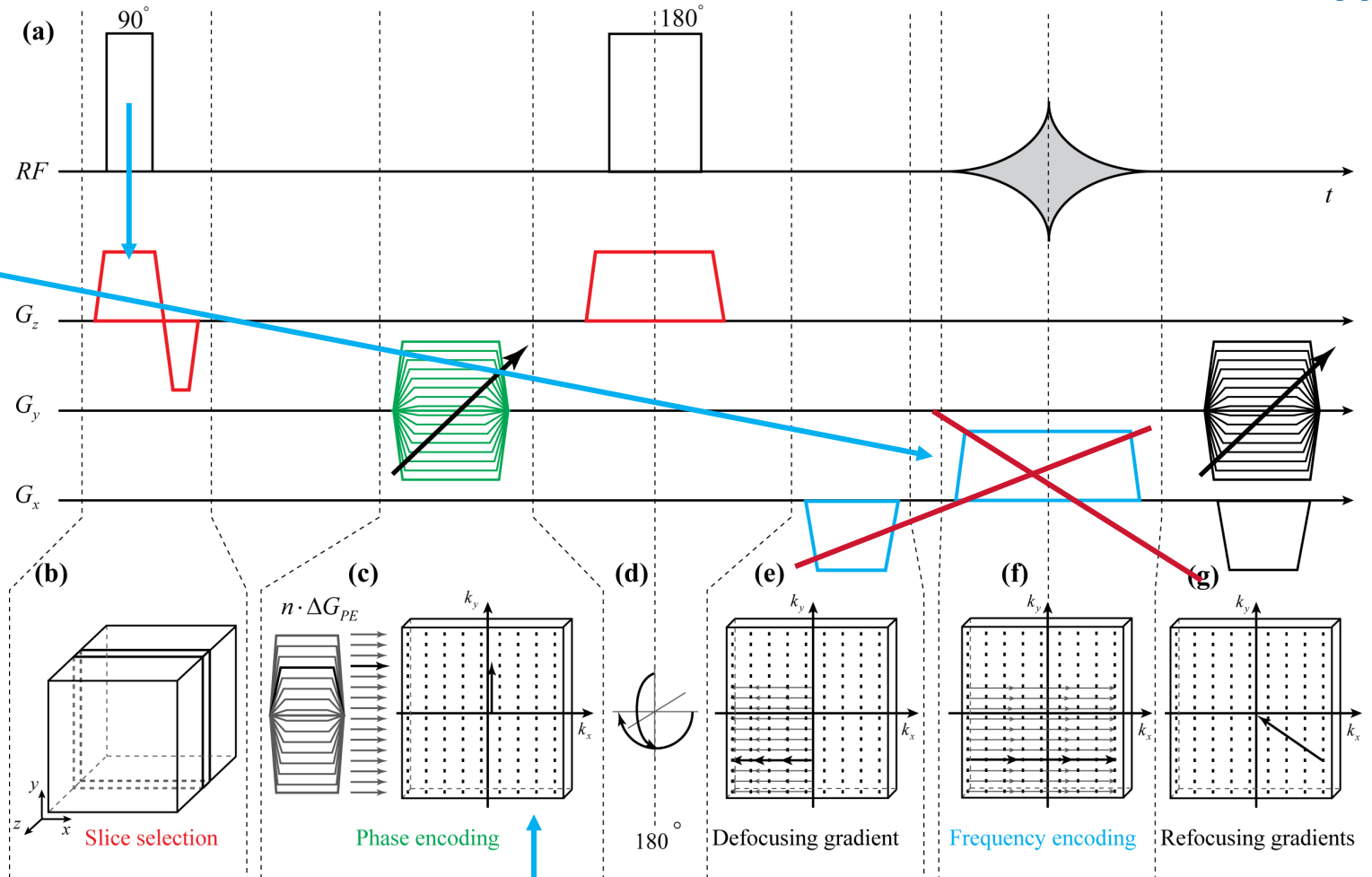
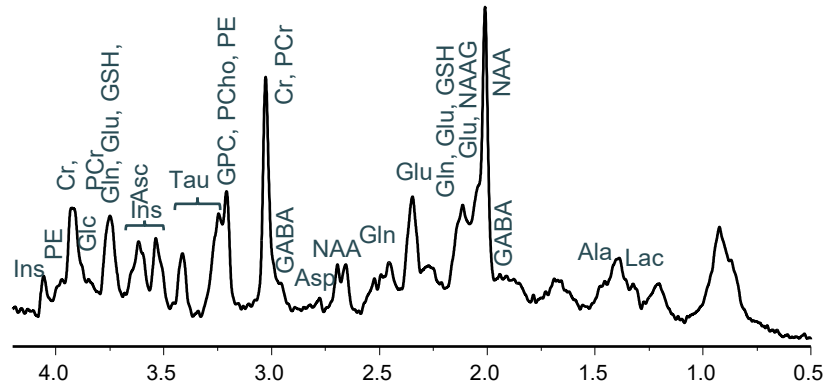


## SPATIAL ENCODING FOR MRSI

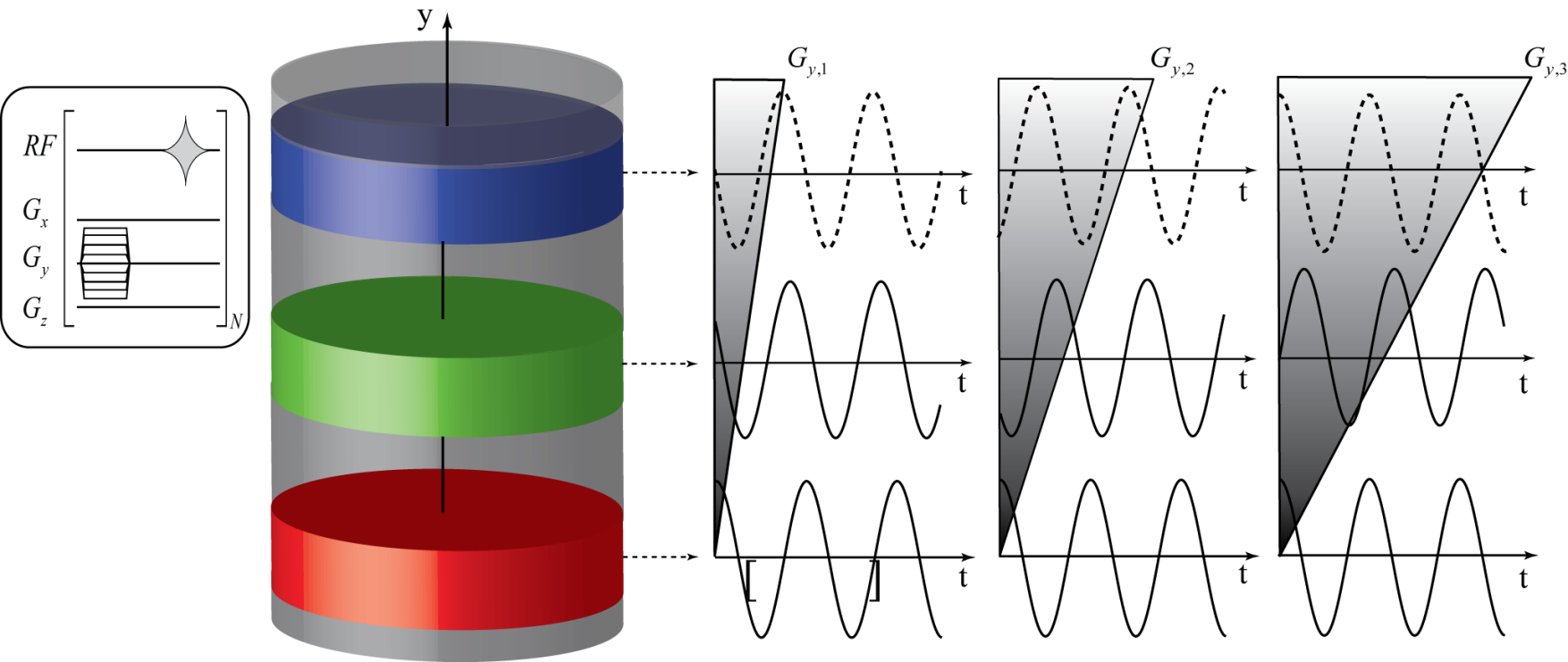
# BASICS OF MRI

## Conventional 2D acquisition

### MR spectrum – frequency encoded



# SPATIAL ENCODING – PHASE ENCODING

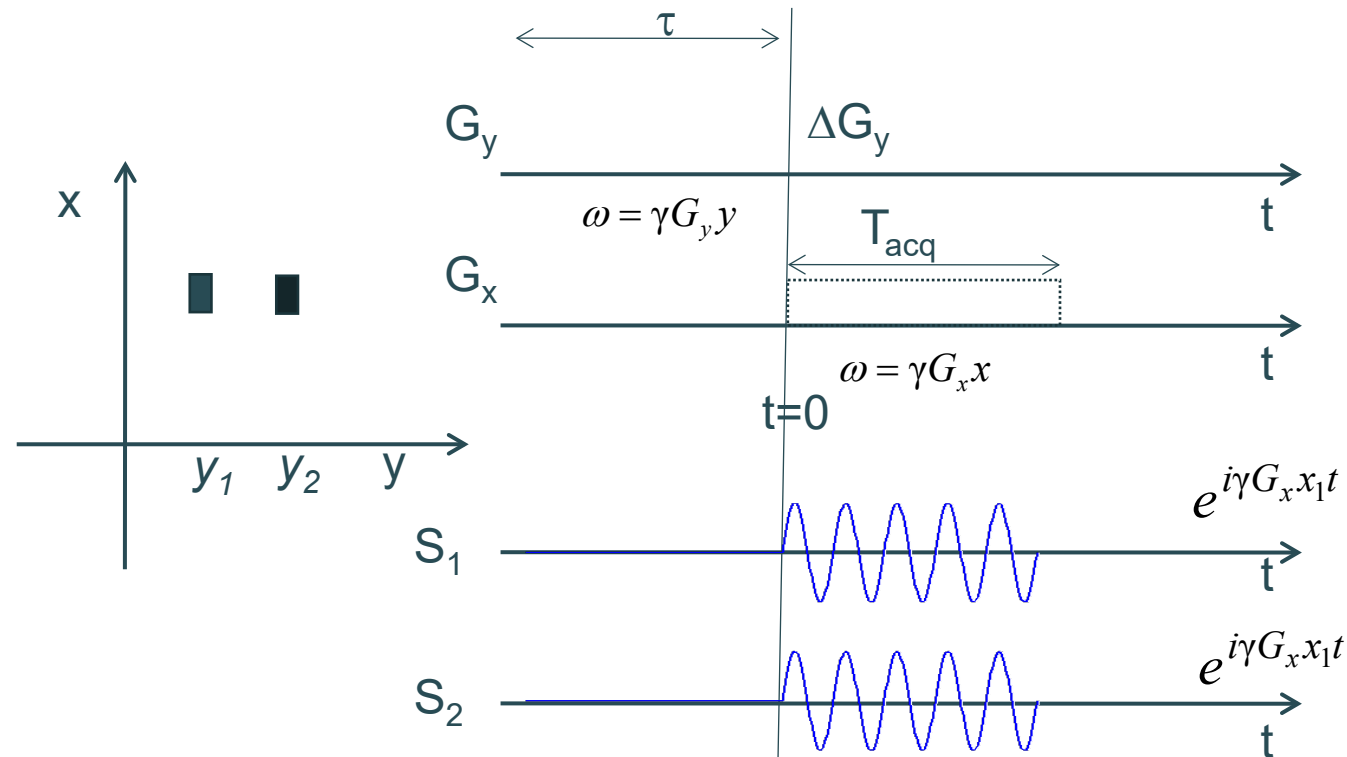


$$\Phi(y) = \omega_y \cdot \tau_{pe} = \gamma \cdot y \cdot G_y \cdot \tau_{pe}$$

$$S(G_y, \tau_{pe}) \propto \iint_{slice} \rho(x, y) e^{-j\gamma G_y y \tau_{pe}} dx dy$$

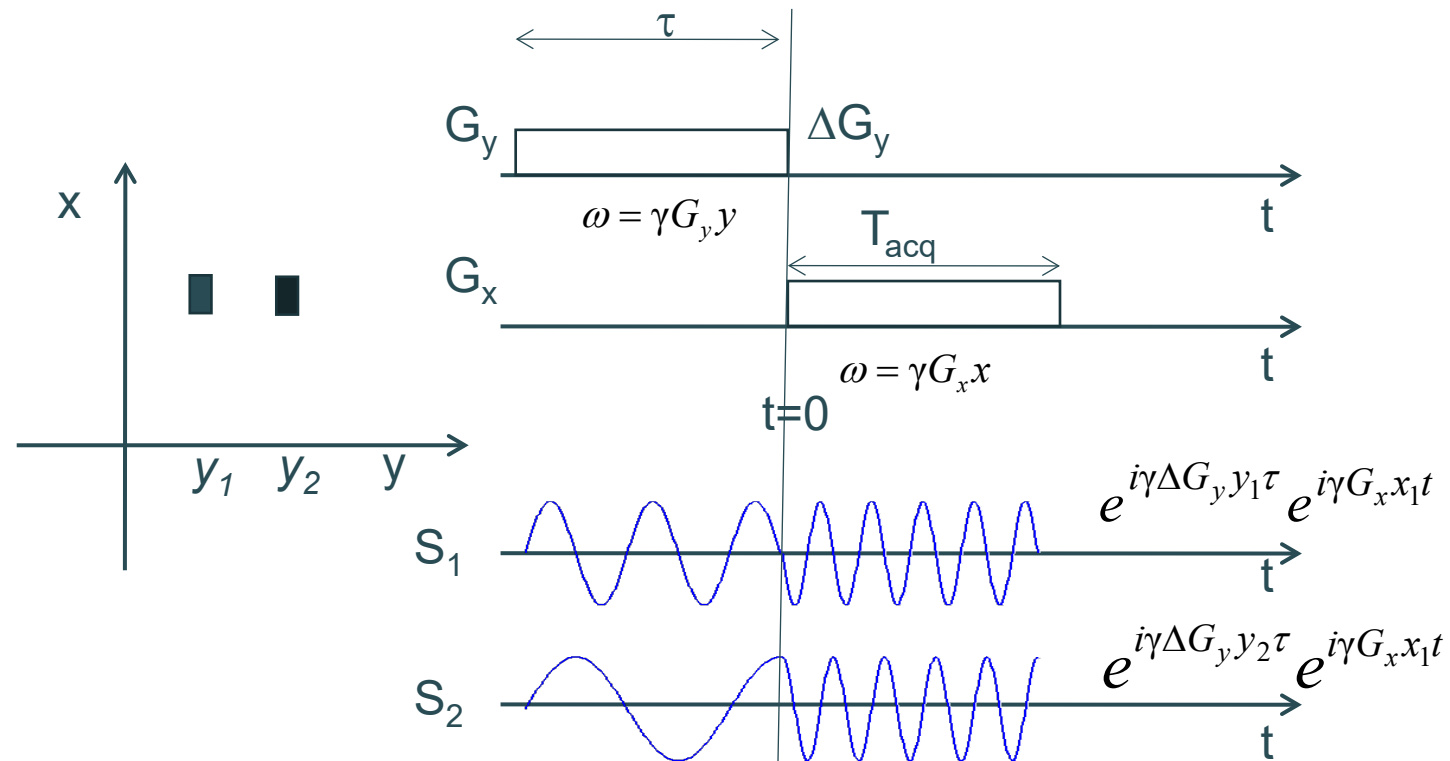
# PHASE ENCODING

- In a single acquisition, frequency encoding indicates the position in one dimension. Encoding 2<sup>nd</sup> and 3<sup>rd</sup> dimension is usually accomplished via phase encoding (position is encoded in the phase of the NMR signal)

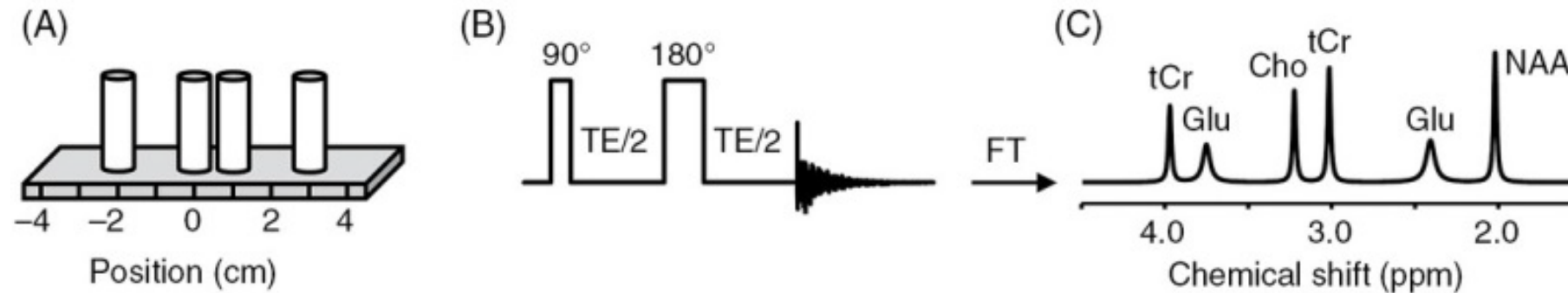


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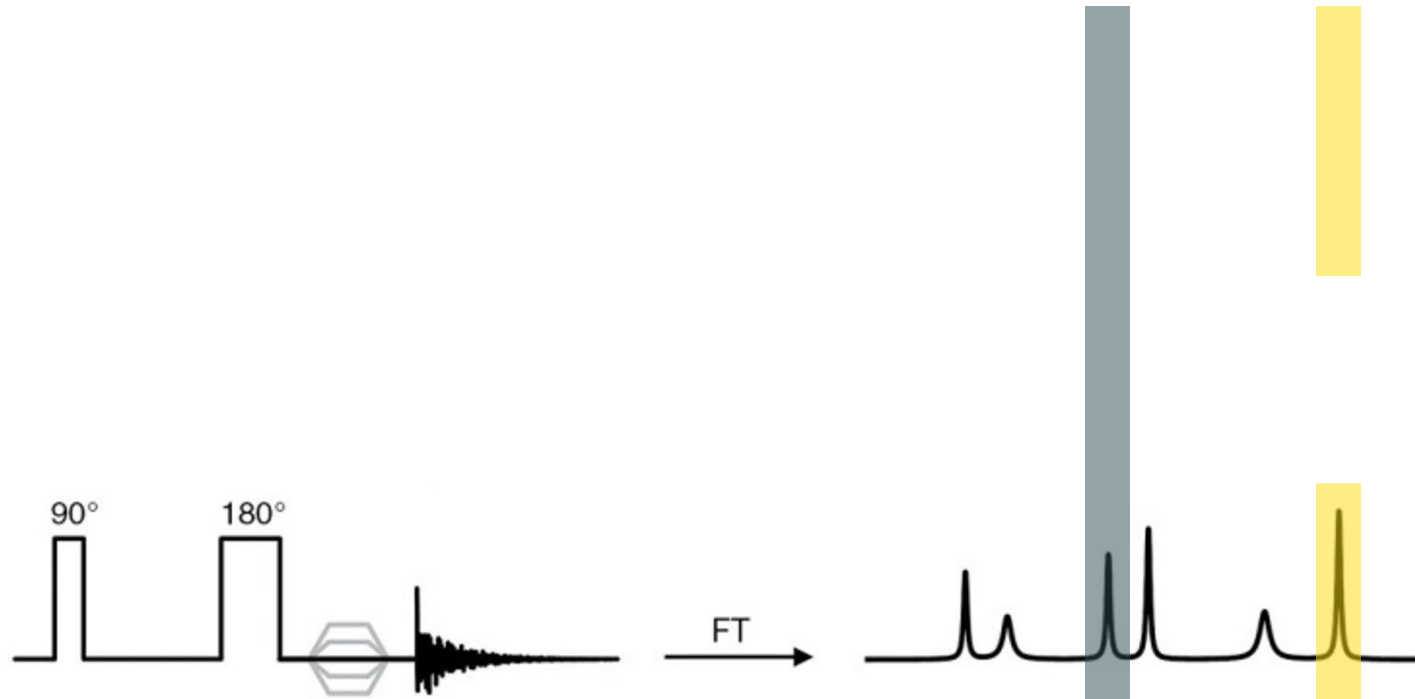


# PHASE ENCODING FOR MRSI



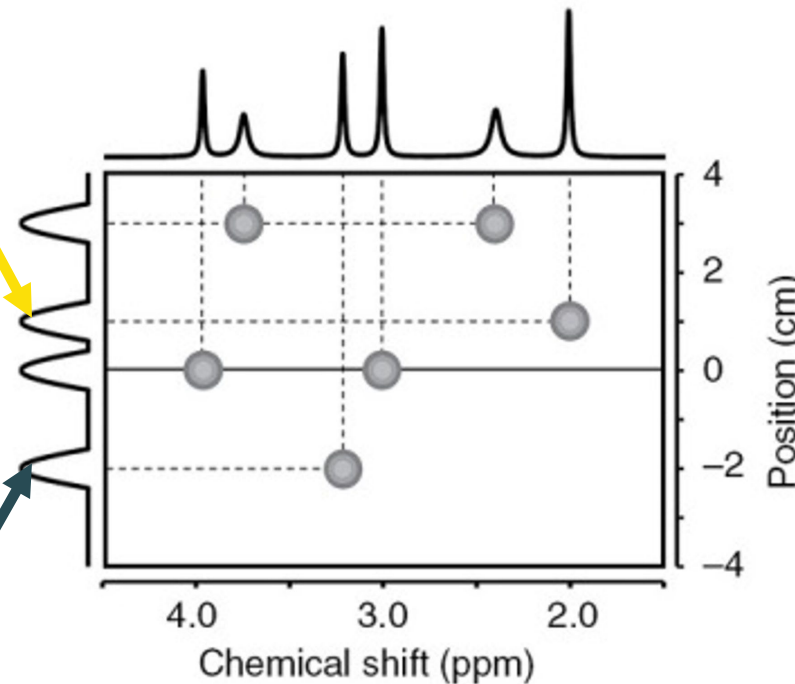
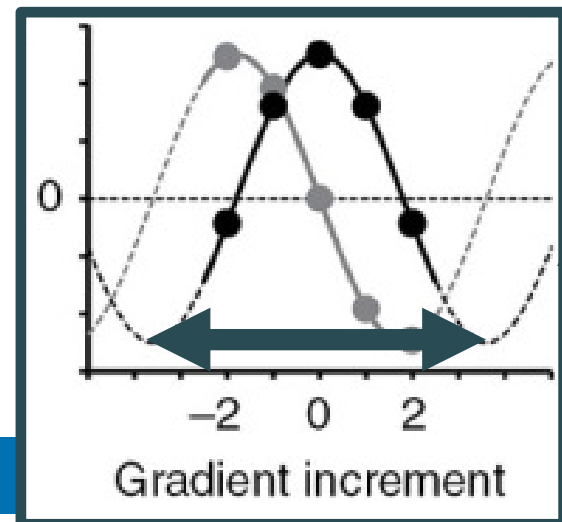
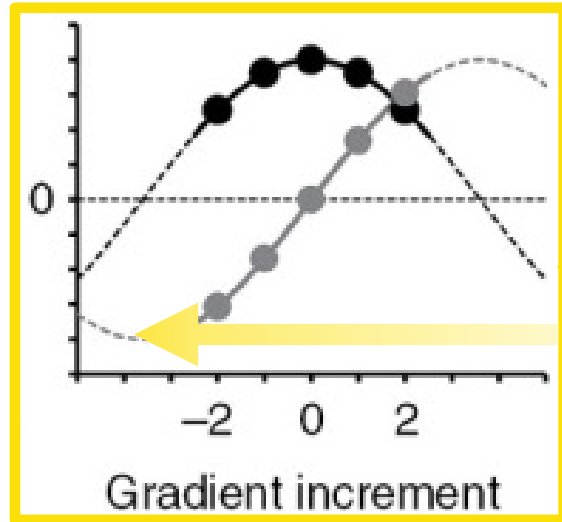
- Frequency encodes metabolites – we cannot encode spatial position with frequency, but ...
- ... we can convert any MRS sequence into an MRSI sequence by adding phase encoding gradients!

# PHASE ENCODING FOR MRSI- QUALITATIVE EXPLANATION



$$\phi(y) = \gamma y G_y t$$

# 2D FOURIER TRANSFORM



$$F(G, \omega) = \int_{-\infty}^{+\infty} f(r, \omega) e^{i\gamma r G t} dr$$

$k = \gamma G t$   $\downarrow$   $k$ -space formalism

$$F(k, \omega) = \int_{-\infty}^{+\infty} f(r, \omega) e^{ikr} dr$$

$\downarrow$  Fourier transformation

$$f(r, \omega) = \int_{-\infty}^{+\infty} F(k, \omega) e^{-ikr} dr$$

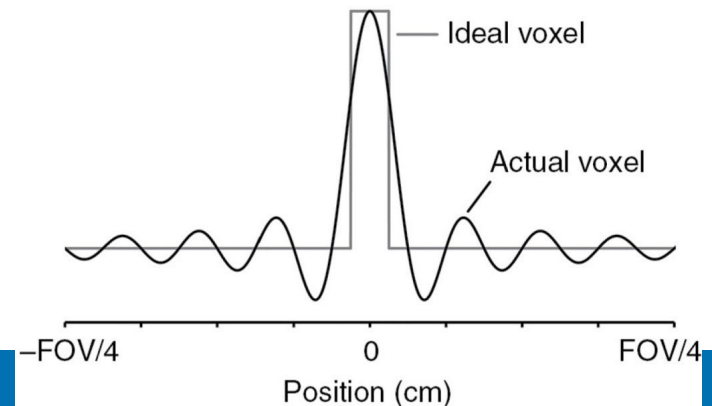
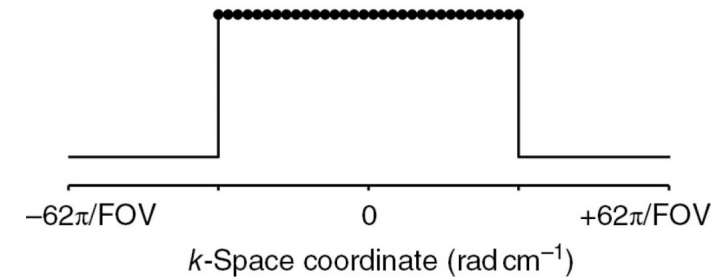
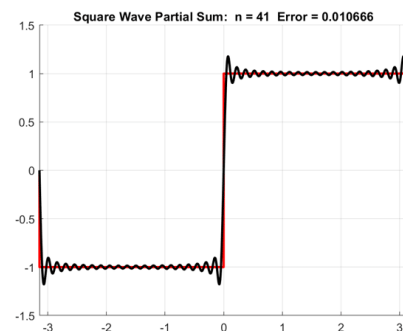
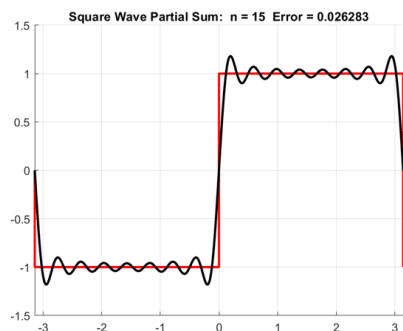
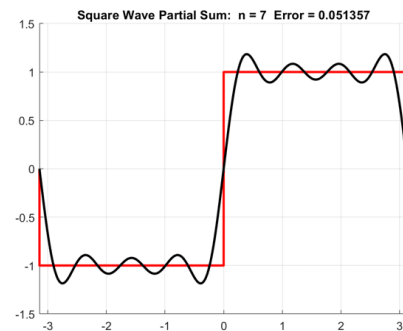
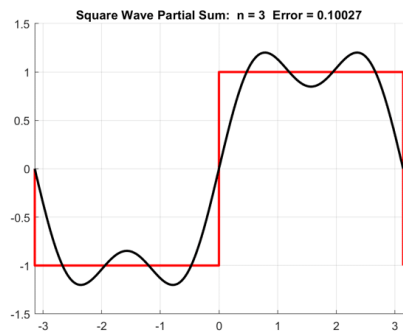
# SAMPLING – $k_{\text{SPACE}}$ – FT

- The signal is not continuously sampled in the k-space
- Digitalization rate in the spatial frequency– phase encoding steps -- if too many – too much time ☹️
- Nyquist sampling criterion – avoid aliasing maximum phase shift between two gradient increments over the whole FOV needs to be  $2\pi$
- $2\pi = \gamma \text{FOV} \Delta G t \quad \rightarrow \quad \text{FOV} = \frac{2\pi}{\gamma \Delta G t} = \frac{2\pi}{\Delta k}$
- $\Delta V = \frac{\text{FOV}}{N}$  - nominal voxel size

# SPATIAL RESOLUTION & PSF

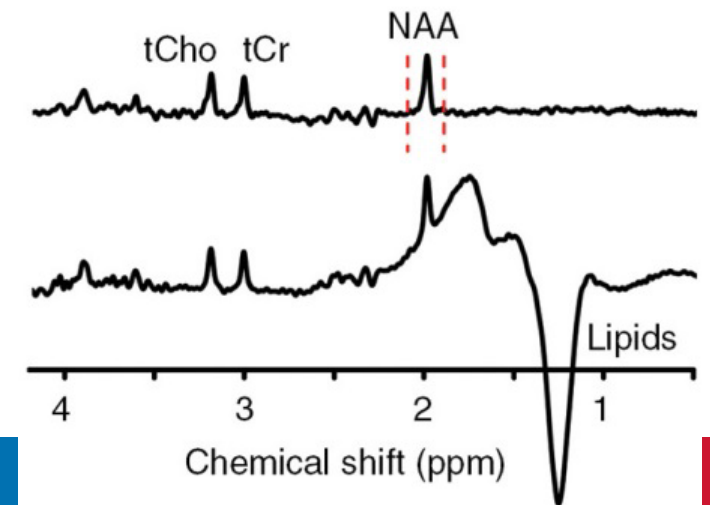
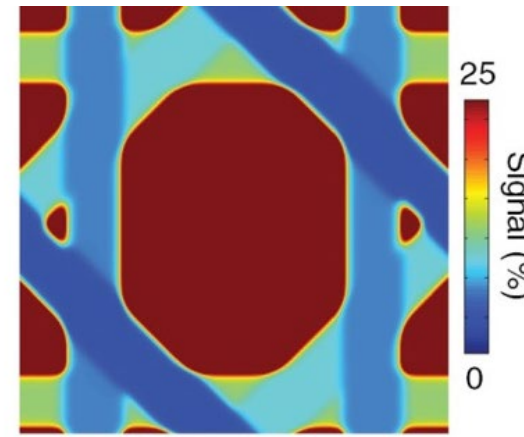
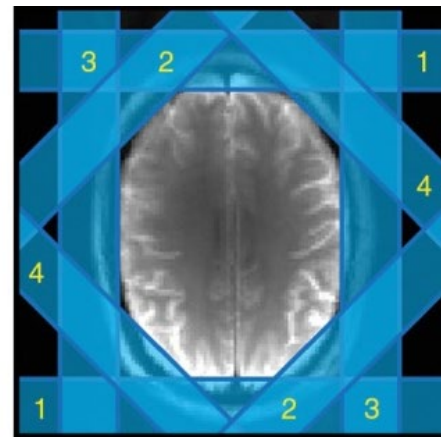
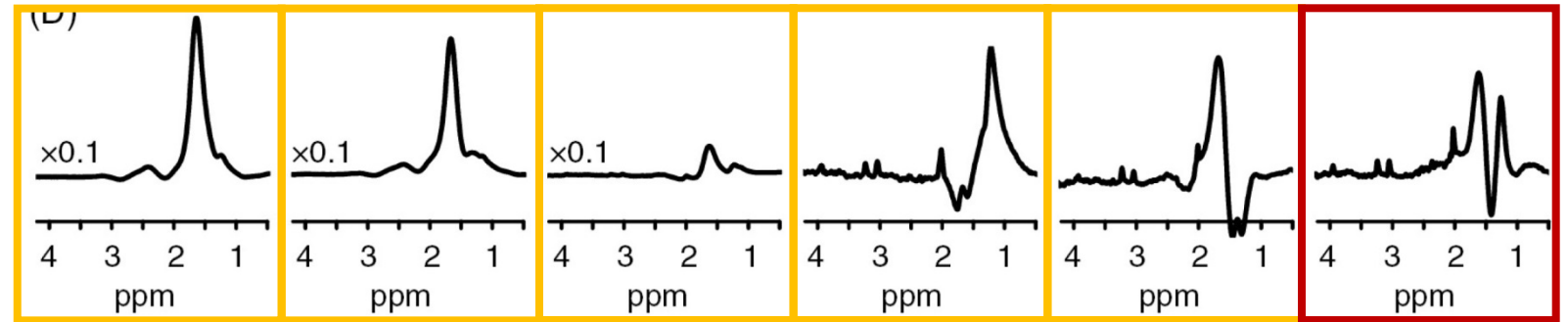
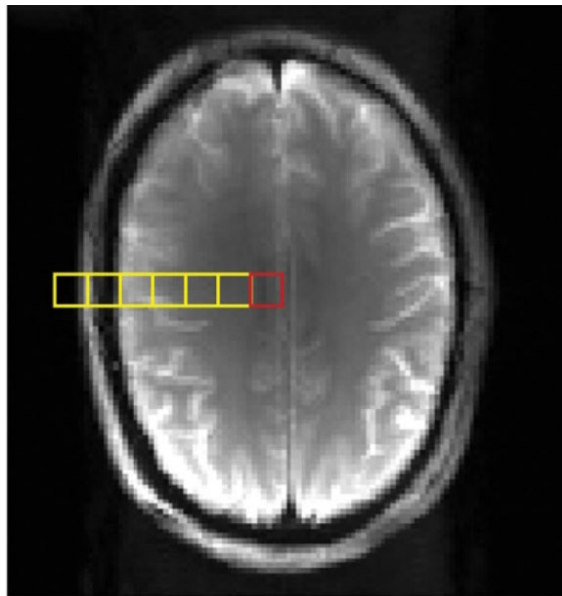
- We cannot efficiently collect a large  $k$  space grid in a reasonable time
- Small number of  $k$  space frequency components cause ripples in spatial localization (described by the Point Spread Function)
- signal is contaminated with the signals from other nominal voxels

Image credit: gaussianwaves.com



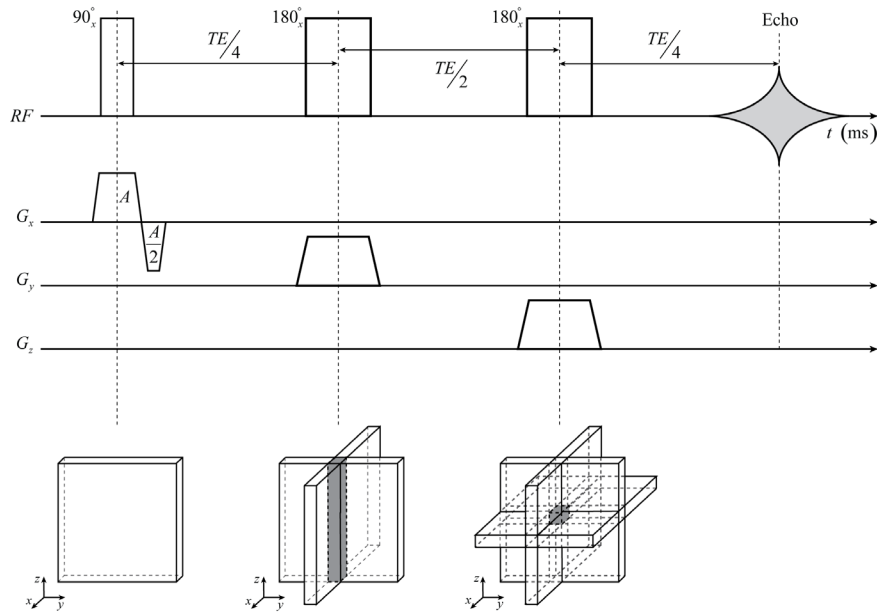
# SPATIAL RESOLUTION

Extracranial lipids 'leak' even into center-brain voxels due to PSF!

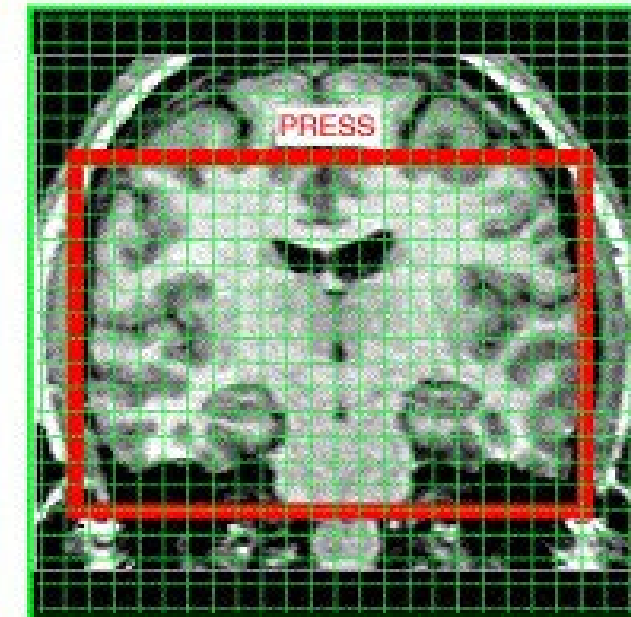
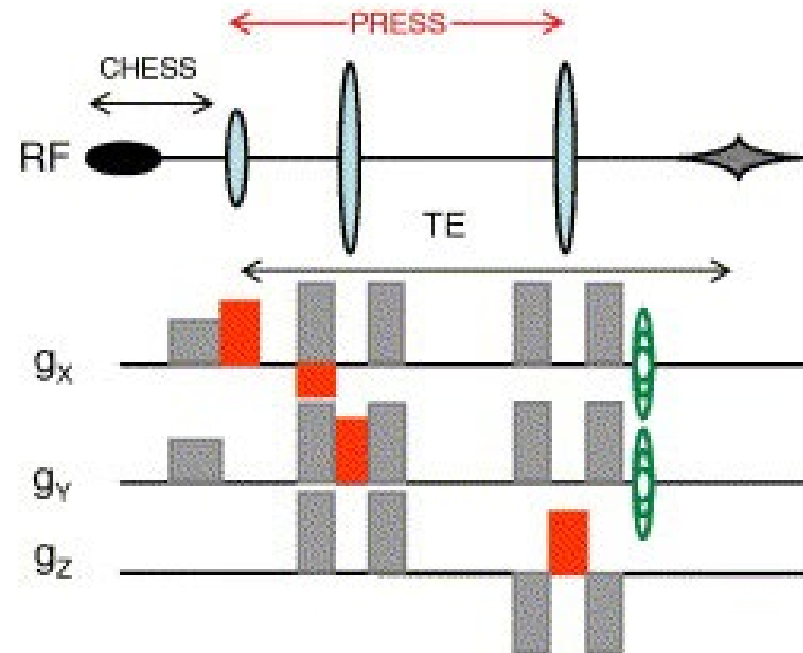


# WHAT SEQUENCES TO USE

Any type of sequence to which you add phase encoding gradients



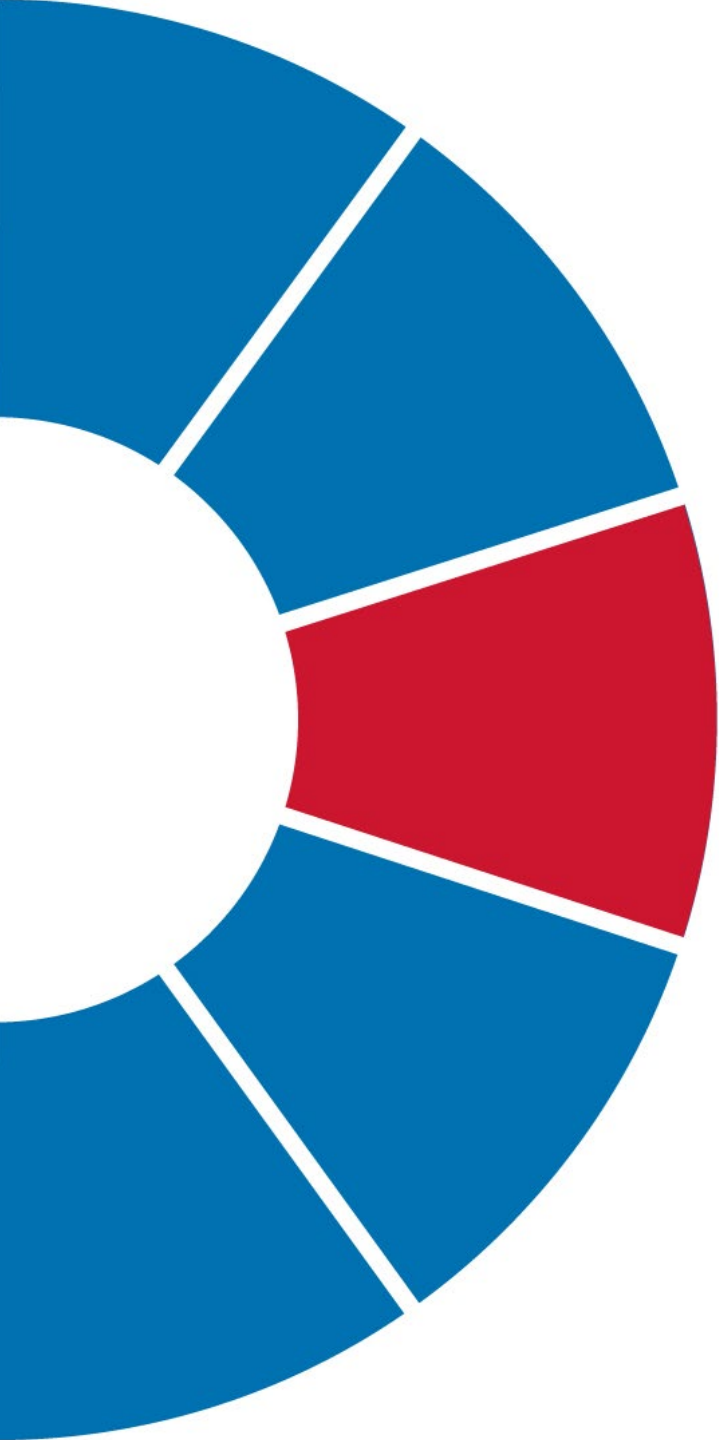
- 3RF pulses – traditional SVS localization techniques
- 2 RF pulses – what will you be selecting here?
- 1RF pulse – what will you be selecting here?
- 2 pe gradients but you can also have 1pe gradient ☺



# TEMPORAL RESOLUTION

- $T_{measurement} = NA \times N_x \times N_y \times TR$ 
  - 1.5T, 16x16x16, NA=1, TR=2000ms --- 136min
  - 32x32 – 34min
- SNR is defined as signal per one SD of the noise
  - Effective voxel volume and no of averages
  - No of phase encoding x NA --- 32x32 with 1NA = 1024 averages as the entire VOI is excited each time
- SNR unit of time =  $\frac{SNR}{\sqrt{T_{tot}}}$

Terminology and concepts for the characterization of in vivo MR spectroscopy methods and MR spectra: Background and experts' consensus recommendations - Kreis - 2021 - NMR in Biomedicine - Wiley Online Library



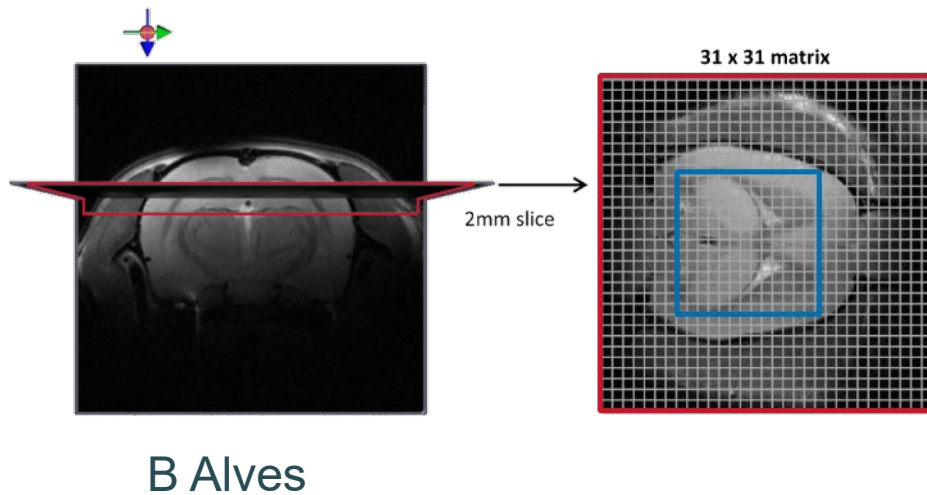
## MRSI ADVANTAGES AND CHALLENGES

# ADVANTAGES

- Multiple locations in the same time / simultaneously
- Direct comparison of metabolic information from multiple regions of interest in a single scan
- $^1\text{H}$  but also  $^{31}\text{P}$ ,  $^2\text{H}$ , ....fMRSI, DW-MRSI, ....
- phase encoding gradient – no CSDA
- Can be full brain coverage and very fast – next talk

# CHALLENGES

- Bo shimming – shim larger regions – significant Bo inhomogeneities



ParaVision 360 V1.1

File View Programs Window Help

Palette Workspace Explorer MRI Acq/Reco Data x Dataset Browser x Examination - 19042021\_rat\_testCC / 19042021\_rat\_testCC x Viewing - 19042021\_rat\_testCC / 19042021\_rat\_testCC x

job0 Navigator

STUDY 1  
19 Apr 2021  
09:25:23 +0000  
MR E:6 P:1  
IMA 7/20  
Bruker:RARE

EPFL Lausanne  
BAP141/26

cm 35605.1  
100%  
1  
0 47.43  
0%

Rt

SP Cd2.07  
SL 0.80 / 0.80  
FoV 25.00x25.00  
MTX 256x256 (a)

STUDY 1  
19 Apr 2021  
09:25:23 +0000  
MR E:6 P:1  
IMA 10/20  
Bruker:RARE

EPFL Lausanne  
BAP141/26

cm 34754.7  
100%  
1  
0 59.46  
0%

Rt

SP Ro0.33  
SL 0.80 / 0.80  
FoV 25.00x25.00  
MTX 256x256 (a)

STUDY 1  
19 Apr 2021  
09:25:23 +0000  
MR E:6 P:1  
IMA 15/20  
Bruker:RARE

EPFL Lausanne  
BAP141/26

cm  
1  
0

Rt

SP Ro4.33  
SL 0.80 / 0.80  
FoV 25.00x25.00  
MTX 256x256 (a)

Slice 7/20  
TR 3000.0  
TE 27.0  
TA 04:12  
NEX 2.0  
FA 180.0

Slice 10/20  
TR 3000.0  
TE 27.0  
TA 04:12  
NEX 2.0  
FA 180.0

Slice 15/20  
TR 3000.0  
TE 27.0  
TA 04:12  
NEX 2.0  
FA 180.0

SP Cd  
SP Ro0.33  
SP Ro4.33

Frequency Offset: 0.01 Hz  
Line Width: 17.91 Hz

Ref	Instruction Name	Status	Duration
1	1_Localizer (E1)	SUCCEEDED	00:00:00
2	1_Localizer (E2)	SUCCEEDED	00:00:00
3	2_Localizer (E4)	SUCCEEDED	00:00:00
4	3_Localizer_multi_slice_10_s	SUCCEEDED	00:00:00
5	4_T2_TurboRARE_6_54K_180	SUCCEEDED	00:00:00
6	6_STEAM_highres_JM_11092	SUCCEEDED	00:00:00
7	6_STEAM_highres_JM_11092	SUCCEEDED	00:00:00
8	5_T2_TurboRARE_6_54K_180	SUCCEEDED	00:00:00
9	6_STEAM_highres_JM_11092	SUCCEEDED	00:00:00
10	6_STEAM_highres_JM_11092	SUCCEEDED	00:00:00
11	6_STEAM_highres_JM_11092	SUCCEEDED	00:00:00
12	6_STEAM_highres_JM_11092	SUCCEEDED	00:00:00

Drop Here to Append Σ 00:00:24

Open Apply Cancel

Main Fat Sup Fov Sat Sel IR SatTrans Flow Sat Black Blood Auto Shim Shim

Requested Shim  Current...  Study\_Shim  Map\_Sh  Automatic Shim Volume

Iterative Correction

Status  Shape of shim volume  C...  Cylinder  Ellipsoid

Success

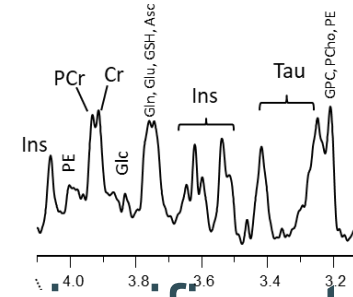
<https://www.epfl.ch/labs/mrs4brain/links/standardized-preclinical-mrs-a-multi-center-study/>

Sequence Setup System Reconstruction Single Parameter Instruction  
Routine Contrast Resolution Geometry

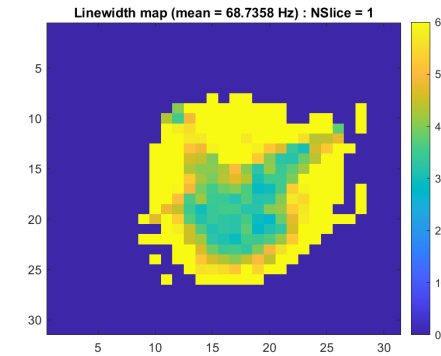
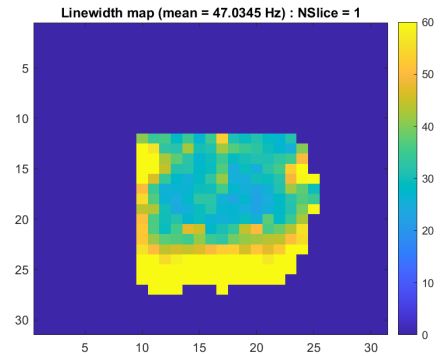
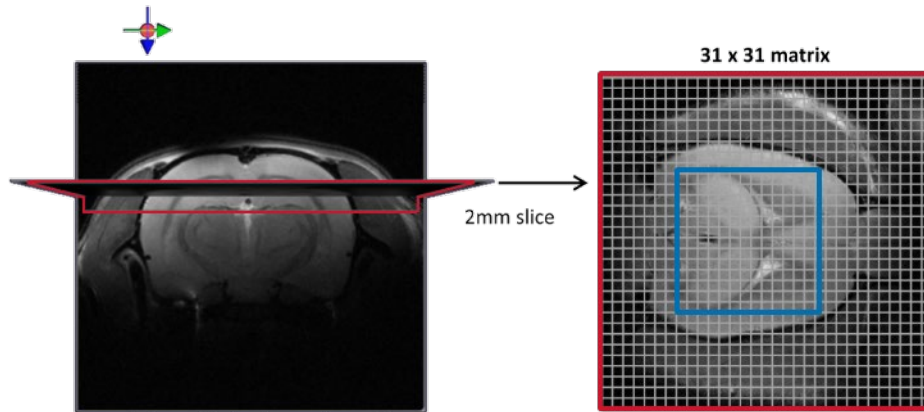
Continue Scan Setup

# CHALLENGES

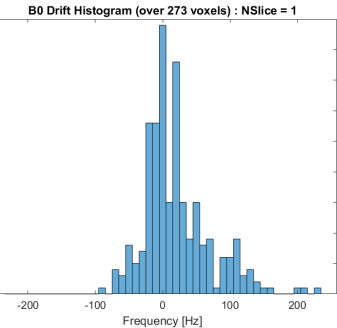
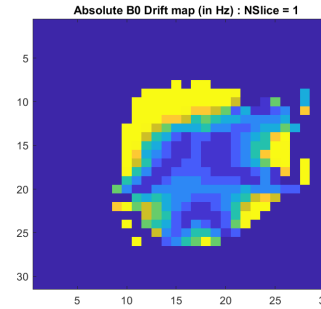
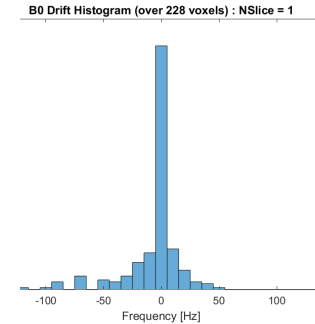
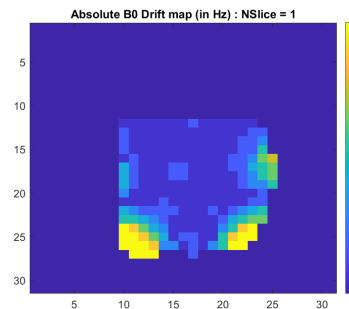
High spectral resolution



- B<sub>0</sub> shimming – shim larger regions – significant B<sub>0</sub> inhomogeneities

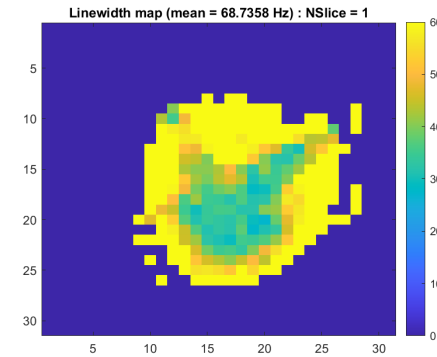
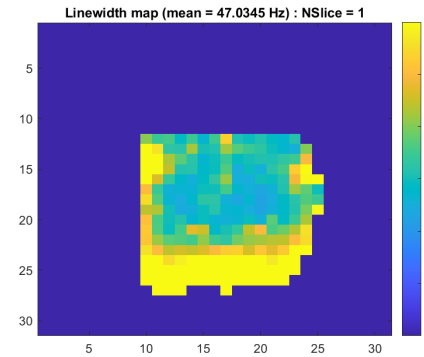
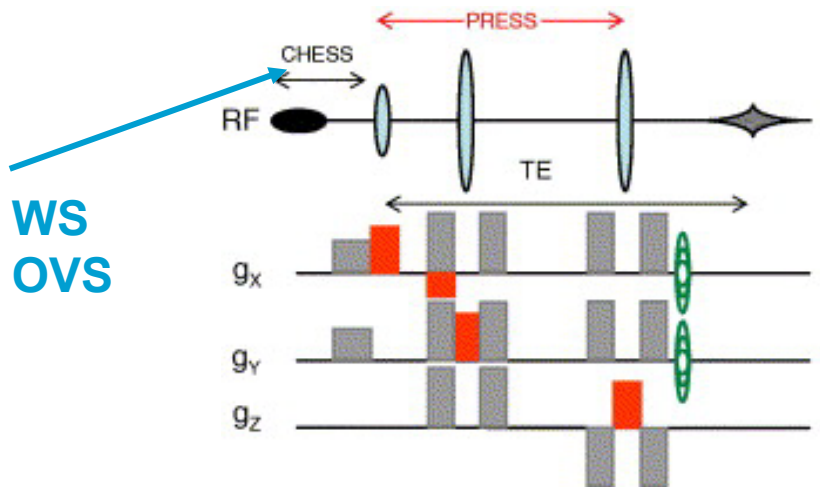


B<sub>0</sub> Altes

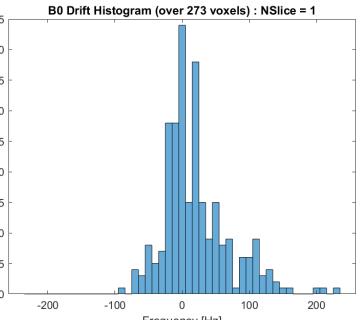
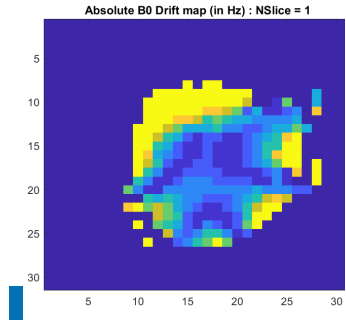
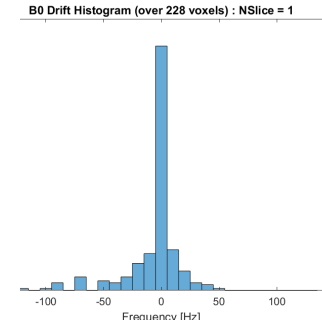
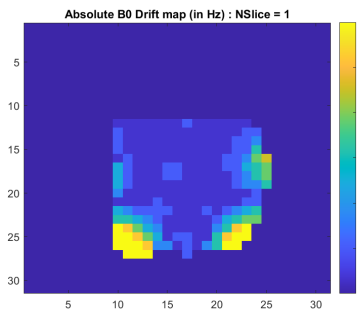


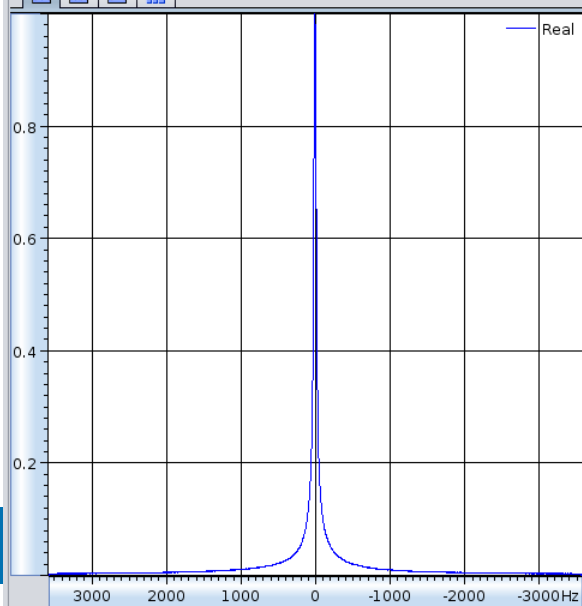
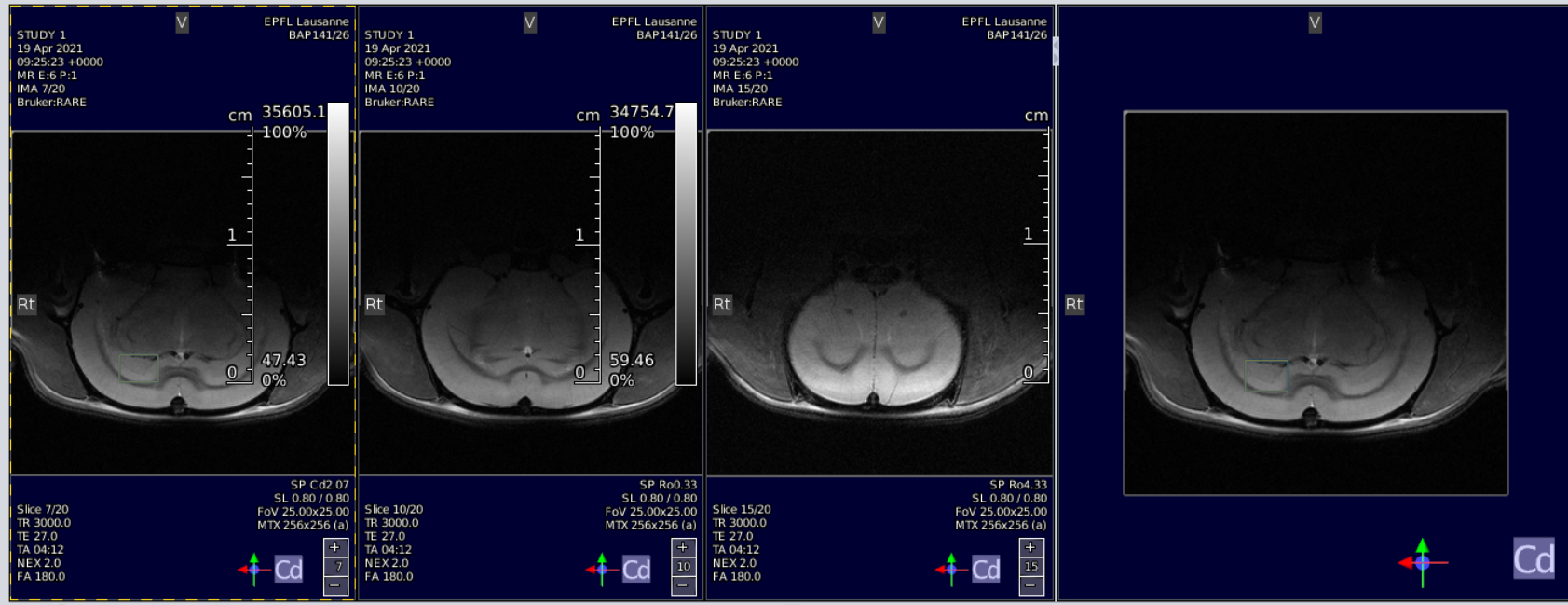
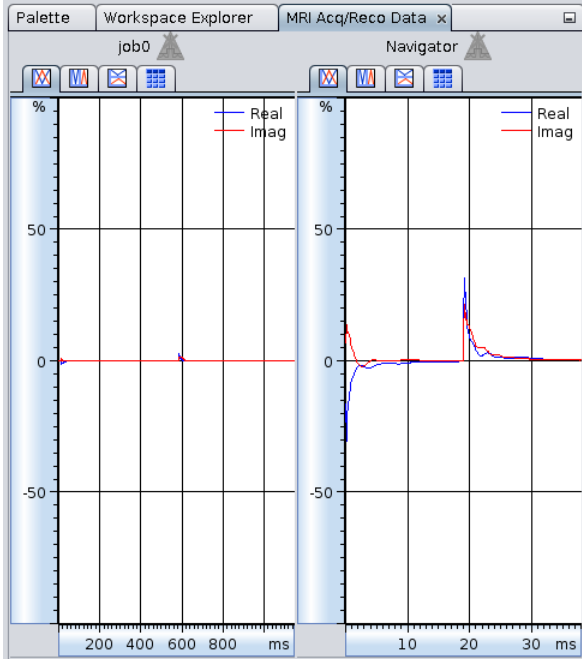
# CHALLENGES

- Water suppression – is done on the signal coming from the entire organ where the resonant frequency (due to  $B_0$  inhom,  $B_0$  drifts, ..) can not be the best one



B Alves





Ref	Instruction Name	Status	Duration
3	2_Localizer (47)	SUCCEEDED	00:00:00
4	3_Localizer_multi_slice_10_s	SUCCEEDED	00:00:00
5	4_T2_TurboRARE_6_54K_180	SUCCEEDED	00:00:00
6	6_STEAM_highres_JM_11092	SUCCEEDED	00:00:00
7	6_STEAM_highres_JM_11092	SUCCEEDED	00:00:00
8	5_T2_TurboRARE_6_54K_180	SUCCEEDED	00:00:00
9	6_STEAM_highres_JM_110	SUCCEEDED	00:00:00
10	6_STEAM_highres_JM_11092	SUCCEEDED	00:00:00
11	6_STEAM_highres_JM_11092	SUCCEEDED	00:00:00
12	6_STEAM_highres_JM_11092	SUCCEEDED	00:00:00
13	7_STEAM_highres_JM_11092	SUCCEEDED	00:00:00
14	7_STEAM_highres_JM_11092	SUCCEEDED	00:00:00
15	8_STEAM_highres_JM_11092	SUCCEEDED	00:00:00

Main Water Sup OVS Auto Shim Shim

Requested Shim  Current\_Shim  Study\_Shim  Map\_Shim

Status Mapshim ready.

Determines when RX array phases are adjusted.  
Parameter PVM\_ArrayPhaseAdjMode

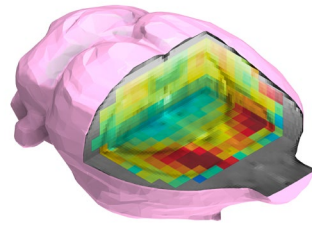
Sequence Setup System Reconstruction Single Parameter Instruction

Routine Spectroscopy Preparation Optimization

Imaging

H

# 3D METABOLIC MAP



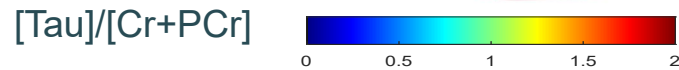
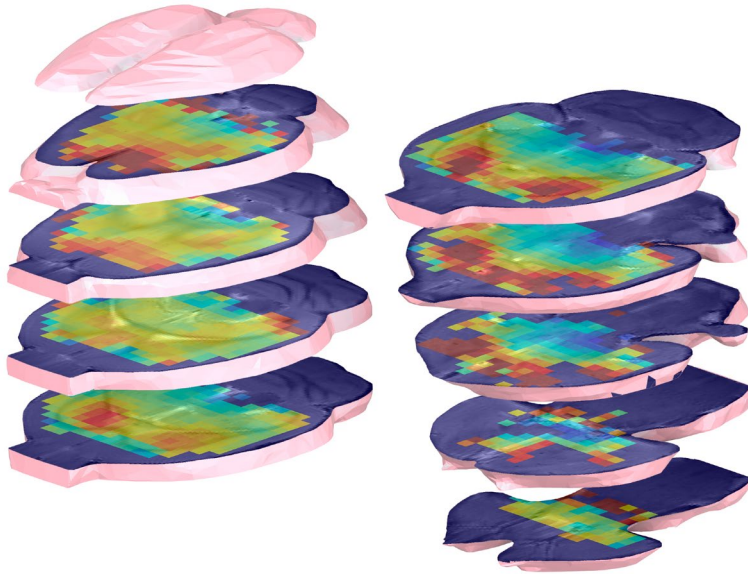
Tan Toi Phan



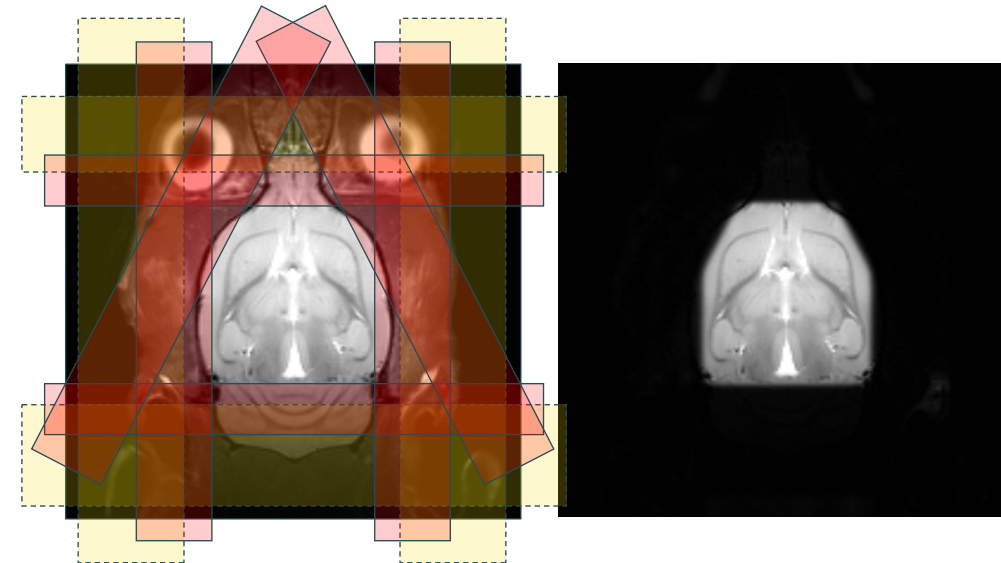
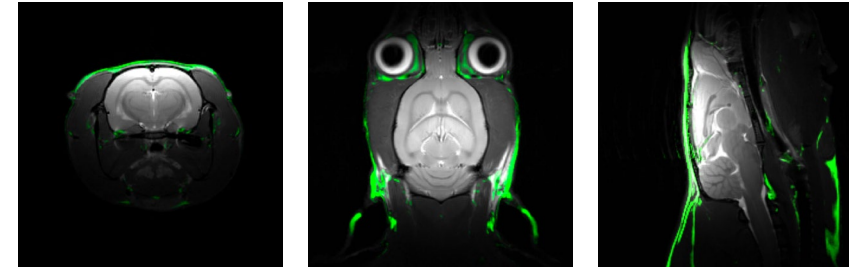
B Alves



T Le



- Time – CS (AF=4, 119min→28min), 1.19  $\mu$ L to 0.59  $\mu$ L
- PSF & Resolution
- Lipid contamination -- Saturation bands
- Post processing
- ...

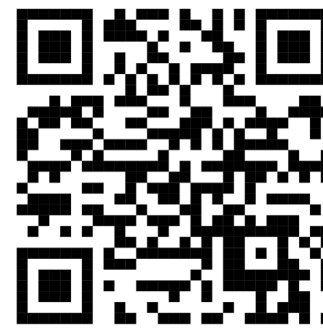


# CHALLENGES

- low concentration of metabolites, long measurement times,
- low signal-to-noise ratio (SNR)
- hardware limitations ( $B_0$  and gradient strength, RF coils,  $B_0$  inhomogeneities)
- requirement for advanced pulse sequences that need to be developed in-house

# CHALLENGES

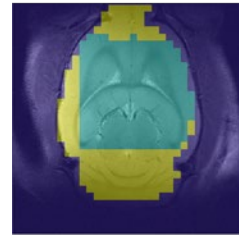
- CSDA is present as we localize the signal at some extent using slice selection gradients
- PSF - contaminations
  - Hamming filter
  - Increase the resolution
  - Weighted acquisitions, not necessary post-acquisition as they decrease the spatial resolution (increases the FWHM of PSF) and sensitivity per unit time (high Kspace coordinates are suppressed)
- Duration !!!!!
  - Low concentration of metabolites
  - Phase encoding steps



1. Read the Bruker MRSI data format

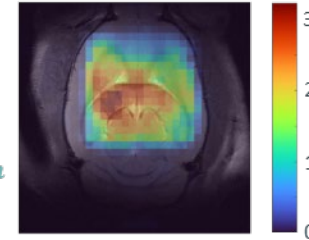
2. Brain mask – Water power mask

3. HSVD water removal



$$Power_{x,y} = \sum_{i=1}^{1024} |spectral\ point|$$

$$0 < Power_{x,y} - \overline{Power} \leftarrow Selection$$



4. Lipid suppression

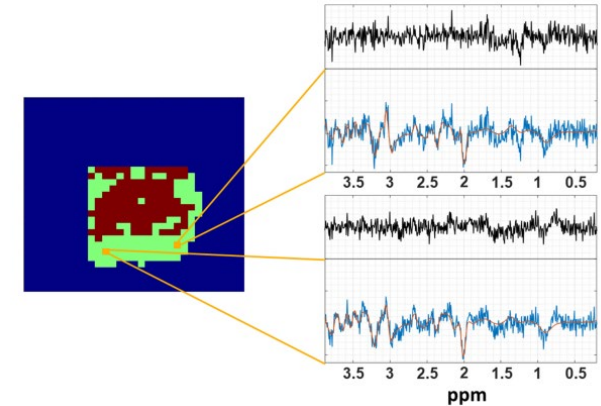
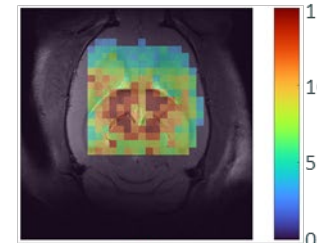
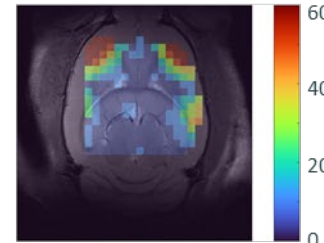
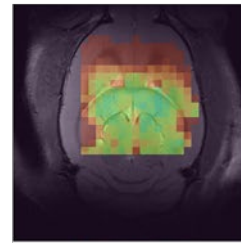
5. Quick data assessment maps

→ linewidth and  $\Delta B_0$  map (water signal)

→ SNR map (NAA peak height /  $\sigma$  noise)

→  $\Delta B_0$  and linewidth map (water signal)

→ SNR map (NAA peak height /  $\sigma$  noise)



6. LCmodel (Version 6.3-1N)

7. Quality selection criteria

→ SNR (75% of  $\overline{SNR}$ ), FWHM (125% of  $\overline{FWHM}$ ) and CRLBs (<40%)

8. Metabolic maps and atlas based automatic segmentation (SIGMA atlas)

**Display settings**

LCModel QC (Only for SNR)

Quality controls

SNR    Mean SNR: 5.922

Minimum SNR: 3

FWHM    Mean FWHM: 0.05442

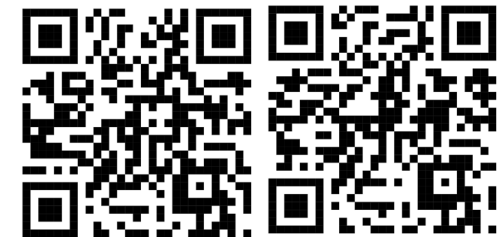
Maximum threshold: 1.25

Max CRLB limit [%]: 30

Interpolation: Off  On

Manual max concentration:

Max concentration: 14



B Alves & G Briand

CIBM.CH

cristina.cudalbu@epfl.ch

MRS4Brain Toolbox

Menu MRSI SVS DWS

**CIBM** **MRS4BRAIN Toolbox**

**Data management**

Folder: Bruker data folder

Folder: Result folder

Study name: 31102023\_FullySampled\_HomeFilter

**Processing steps**

3D MRSI Number slices: 1

Lipid suppression: [button] Fillgaps: [button]

Denoising: None

Reconstruction: Cartesian

**Data specific**

9.4T (for MRI) Reco. Nb

Experiment number folder: [button] PDF document:  Slice range: 2

Metabolites:  Ref: 16 Image: 17 Slice range: 8 1

MRI Slice	Number
MRI central slice 1	6

**Options**

[button] [button] **MRSI Analysis**

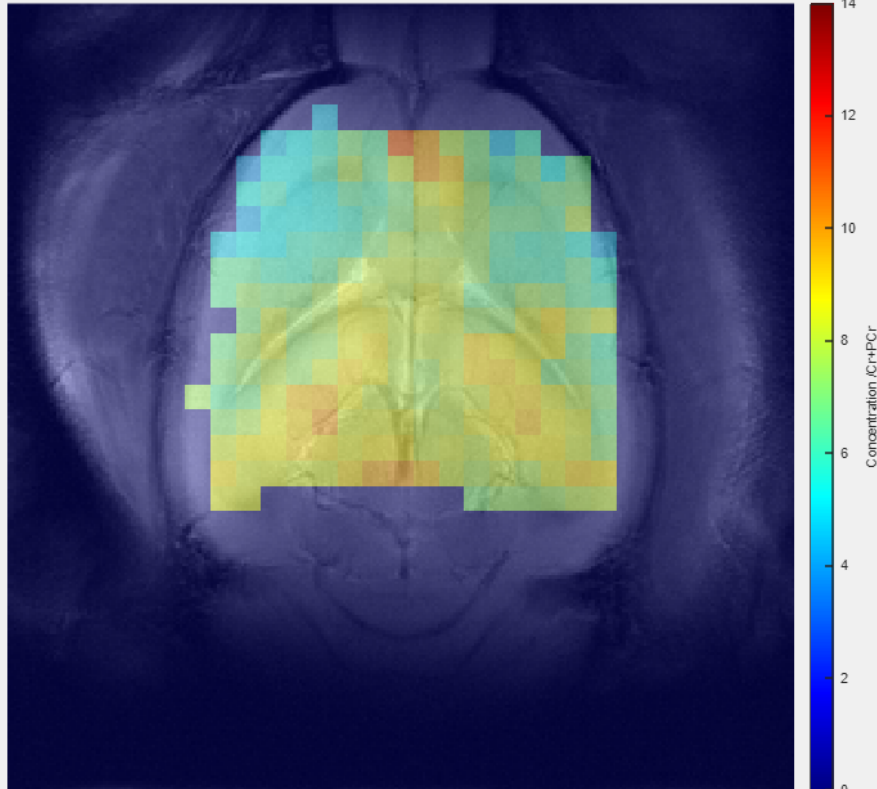
**Show Results**

Slice: 1  Custom regions  Job0 Recon.

Region: Whole Brain Side: L+R Concentrations: [button] [button] [button]

Metabolite: Ins Relative: [button] [button] [button]

Overlap: 0 0.1 0.2 0.3 0.4 0.5 0.6 0.7 0.8 0.9 1 [0.8] [button] [button] [button]



Concentration iCr+PCr: 0 to 14



### MRSI parameters

NRATIO

LCModel path:  Basis set:

PPM start:  PPM end:  Configs:  Original:

DKNTMN:   NSIMUL  VITRO

NRATIO:  WCONC:  DEGZER:  SDDEGZ:  DEGPPM:  SDDEGP:

Relative metabolite:  Relative concentration:

Combination: NCOMB:

Omission: NOMIT:

Use: NUSE:

Index	Combination	Index	Omitted	Index	Used
1	NAA+NAAG	1	-CrCH2	1	NAA
2	Glu+Gln	2	Gua	2	Gln
3	GPC+PCho	3	Ser	3	PCr
4	Cr+PCr	4	Lip13a	4	Cr
		5	Lip13b	5	Ins
		6	Lip09	6	Tau
		7	MM09	7	PCho

Registration:

### Data management

Broker data folder:

Result folder:

Study name:

### Processing steps

3D MRSI Number slices:

Denoising:  Reconstruction:

### Data specific

9.4T (for MRI) Reco. Nb:

Experiment number folder:

Metabolites:  Ref:  Image:  Slice range:

MRI Slice	Number
MRI central slice 1	6

Options:

Concentration (Cr+PCr)

### Show Results

Slice:   Custom regions  Job0 Recon.

Region:  Side:  Concentrations:

Metabolite:  Overlap:

### Volumetry table

Brain region	Left	Right	L+R
Olfactory bulb	52.82	54.01	106.83
Prelimbic cortex	20.92	20.4	41.32
Frontal Association Cortex	2.32	2.81	5.13
Cingulate cortex	25.78	22.74	48.52
Retrosplenial Cortex	29.5	27.54	57.04
Primary Motor Cortex	0	0	0
Secondary Motor Cortex	13.76	12.25	26.01
Primary Somatosensory Cortex	97.28	68.74	166.01
Secondary Somatosensory Cortex	7.67	7.99	15.67
Orbital Cortex	9.49	9.72	19.21
Insular Cortex	56.97	54.3	111.28
Amygdalopiriform Cortex	8.11	8.85	16.96
Entorhinal Cortex	49.49	59.08	108.57
Ectorhinal Cortex	12.35	12.71	25.06
Perirhinal Cortex	12.2	11.65	23.85
Primary Auditory Cortex	23.29	21.69	44.99
Secondary Auditory Cortex	11.77	10.46	22.23

### Concentration table

Metabolite	Mean	Std	N voxels
Mac	0.00	0.00	143
Cr	4.43	1.51	140
PCr	4.05	1.24	130
Ins	9.51	1.60	144
NAA	10.88	1.63	144
Tau	7.52	1.56	144
PCho	1.38	0.58	124
GPC	0.94	0.43	72
Glu	8.80	1.76	144
Gln	2.80	0.92	142
Ala	1.29	0.44	17
Asc	3.65	1.38	98
Asp	1.57	0.47	105
GABA	1.19	0.39	135
Glc	1.05	0.27	135
GSH	1.20	0.43	136
Lac	1.78	1.01	21

### Statistics MRSI

Data specific:  Slice:

On/Off:  MRSI data  Hippocampus\_L+R  Striatum\_L+R

Metabolite list:  Mac  Cr  PCr  Ins  NAA  Tau

Concentrations:

Two-sided p-value:  ANOVA stats:

Ins, Factor B : Region

Region	Mean	F-value	P-value
Factor A	7.06   6.64   7.29	4.99	0.0076
Factor B	7.92   6.08	92.48	0
Interaction	7.10	0.06	0.9380

### Display settings

LCModel QC (Only for SNR)

Quality controls:

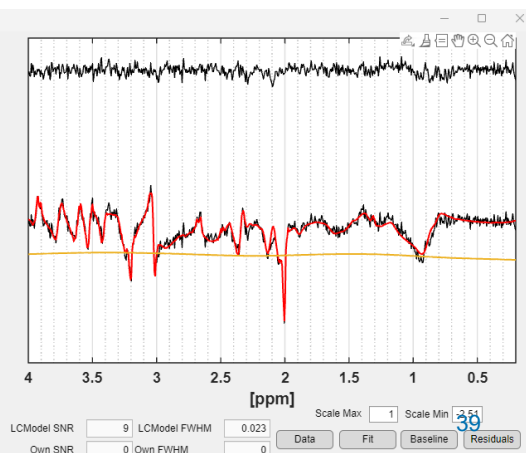
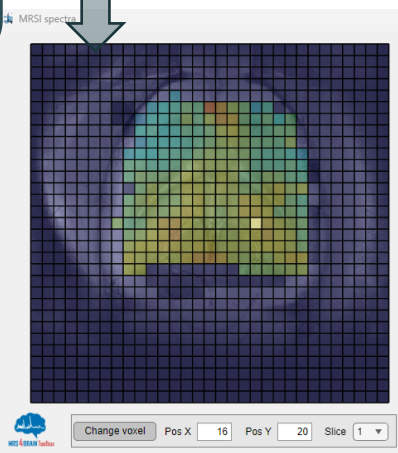
SNR Mean SNR:  Minimum SNR:

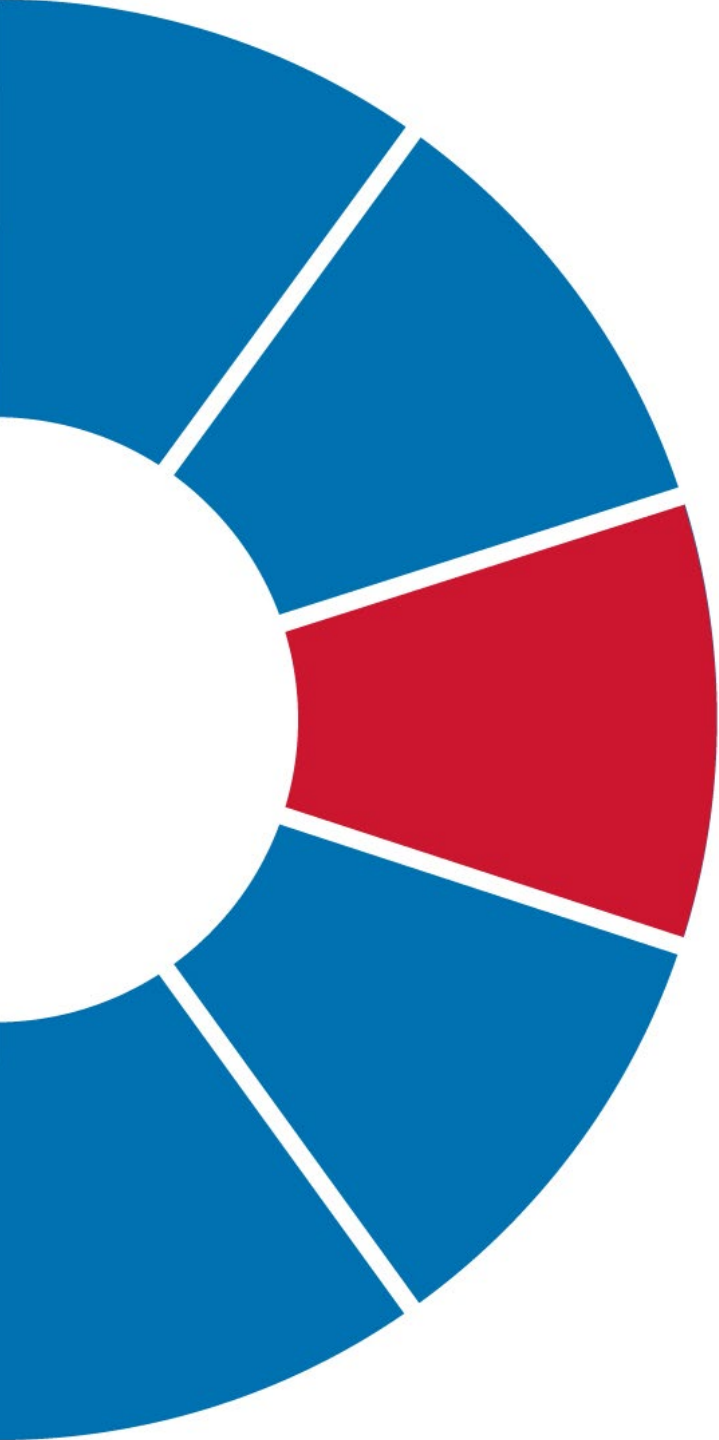
FWHM Mean FWHM:  Maximum threshold:

Max CRLB limit [%]:

Interpolation:  Off  On

Manual max concentration:



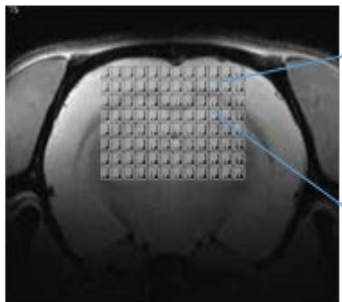


# FROM SQUARE MAPS TO FULL BRAIN COVERAGE

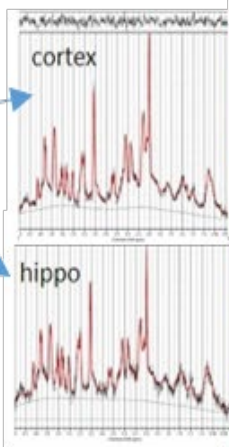
# CHALLENGES

- Overall we still do not get full brain coverage and we are limited to squares inside the brain as we are using SVS localization techniques

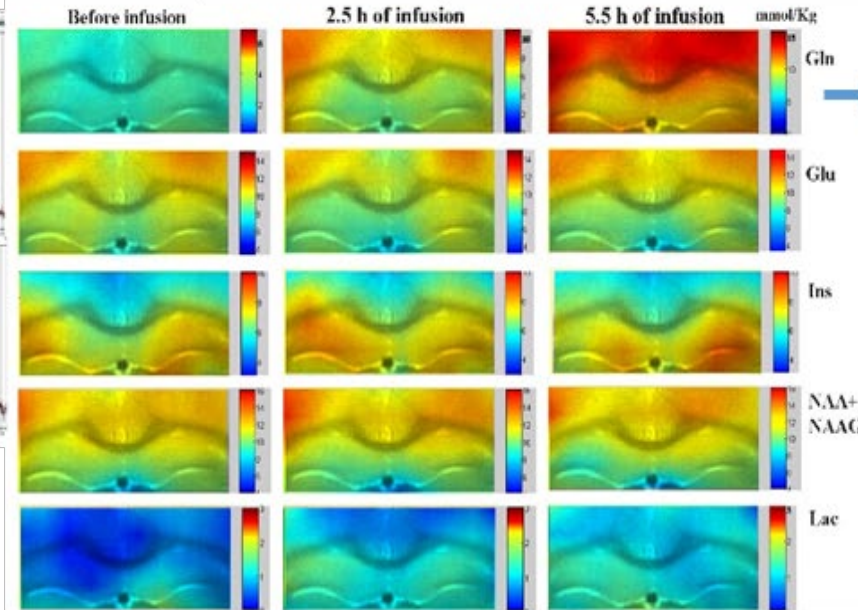
9.4 T Anatomical image +  
Spectroscopic Imaging grid



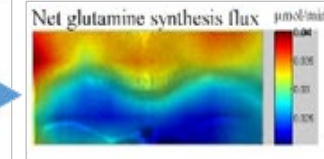
Acquired spectra



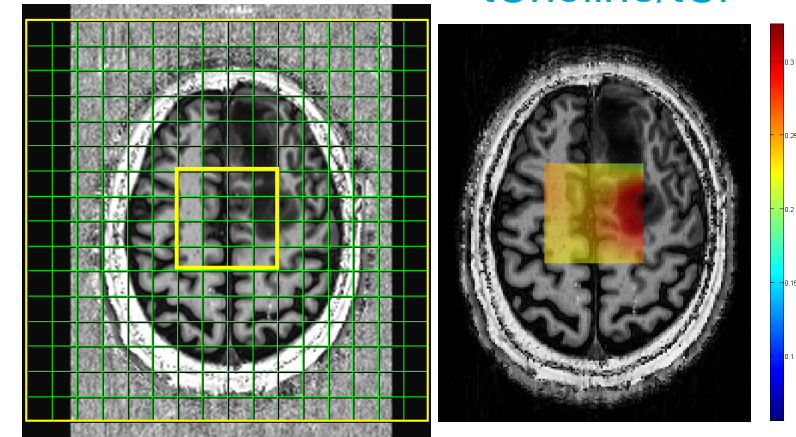
Computed metabolic maps



Computed metabolic fluxes



tCholine/tCr



FOV=24x24mm<sup>2</sup>, Matrix:32x32

Resolution: 0.75x0.75x2mm<sup>3</sup> – 1.1μl

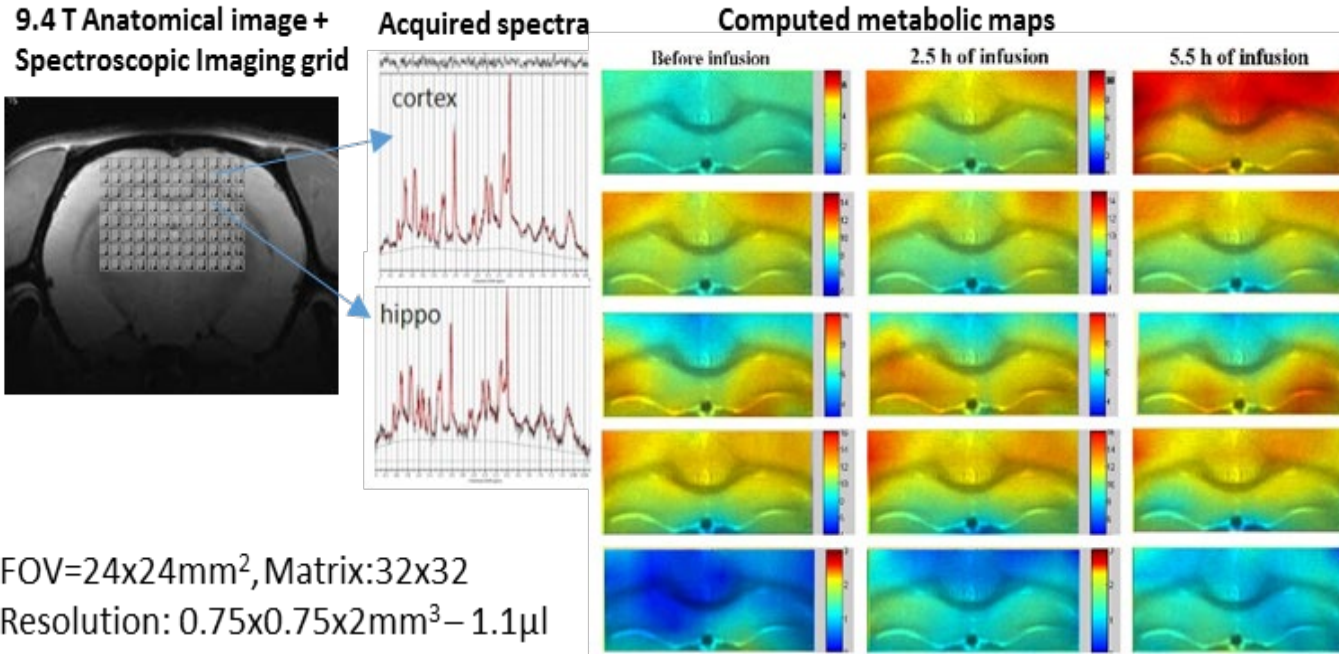
1 axial slice

Acquisition time (metabolites+H<sub>2</sub>O) = ~4h

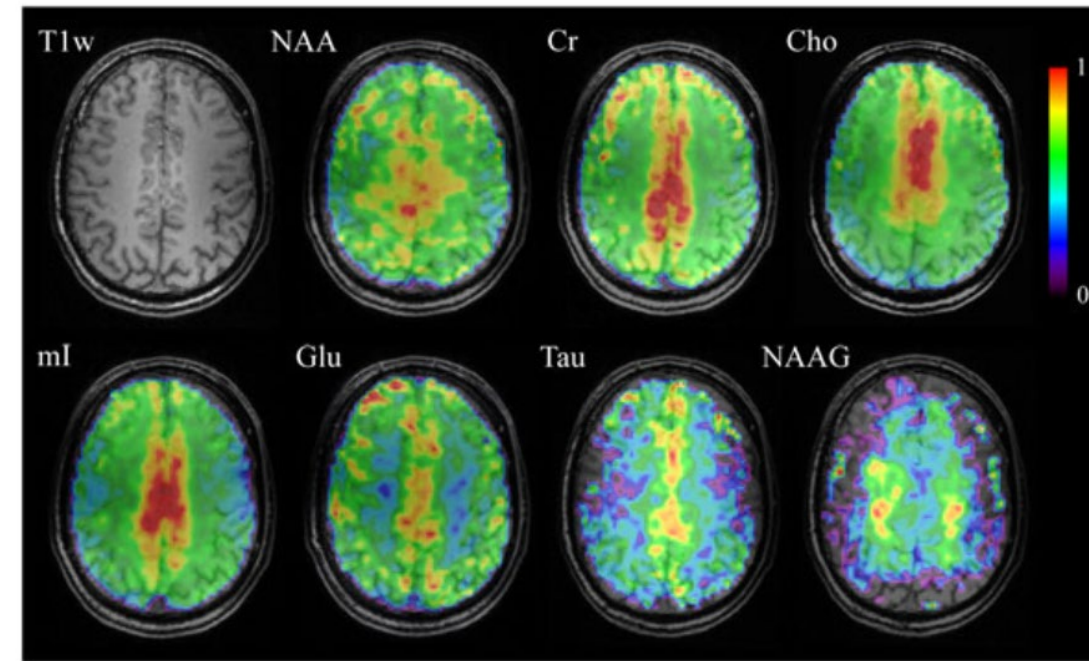
Lijing Xin

C I B M . C H

# FROM SQUARE MAPS TO FULL BRAIN COVERAGE



FOV=24x24mm<sup>2</sup>, Matrix:32x32  
 Resolution: 0.75x0.75x2mm<sup>3</sup>– 1.1μl  
 1 axial slice  
 Acquisition time (metabolites+H<sub>2</sub>O) = ~4h



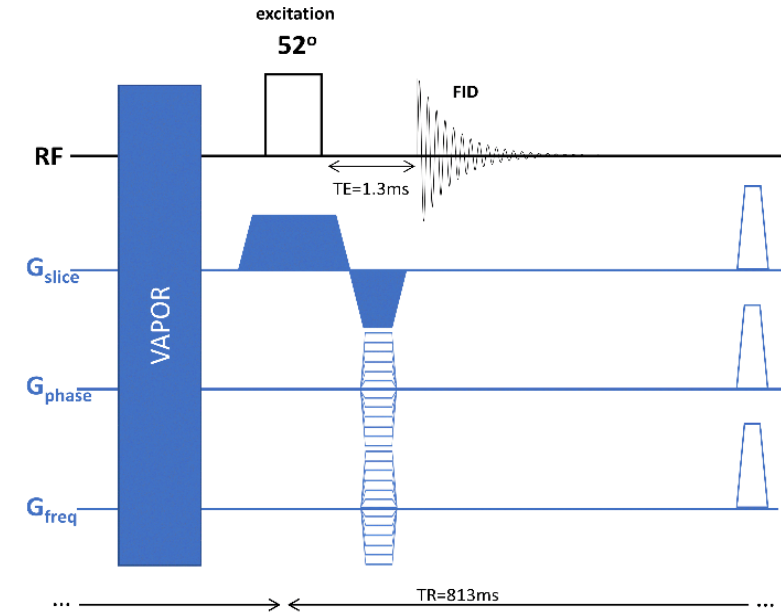
W Bogner et al, NMR Biomed , 2011  
 3.4x3.4x12 mm<sup>3</sup>, within 30 min

<https://www.epfl.ch/labs/mrs4brain/ressources/live-demos/>

**FID-MRSI: negligible J-coupling and T<sub>2</sub> related signal loss, better suitability for fast MRSI at UHF in preclinical settings**

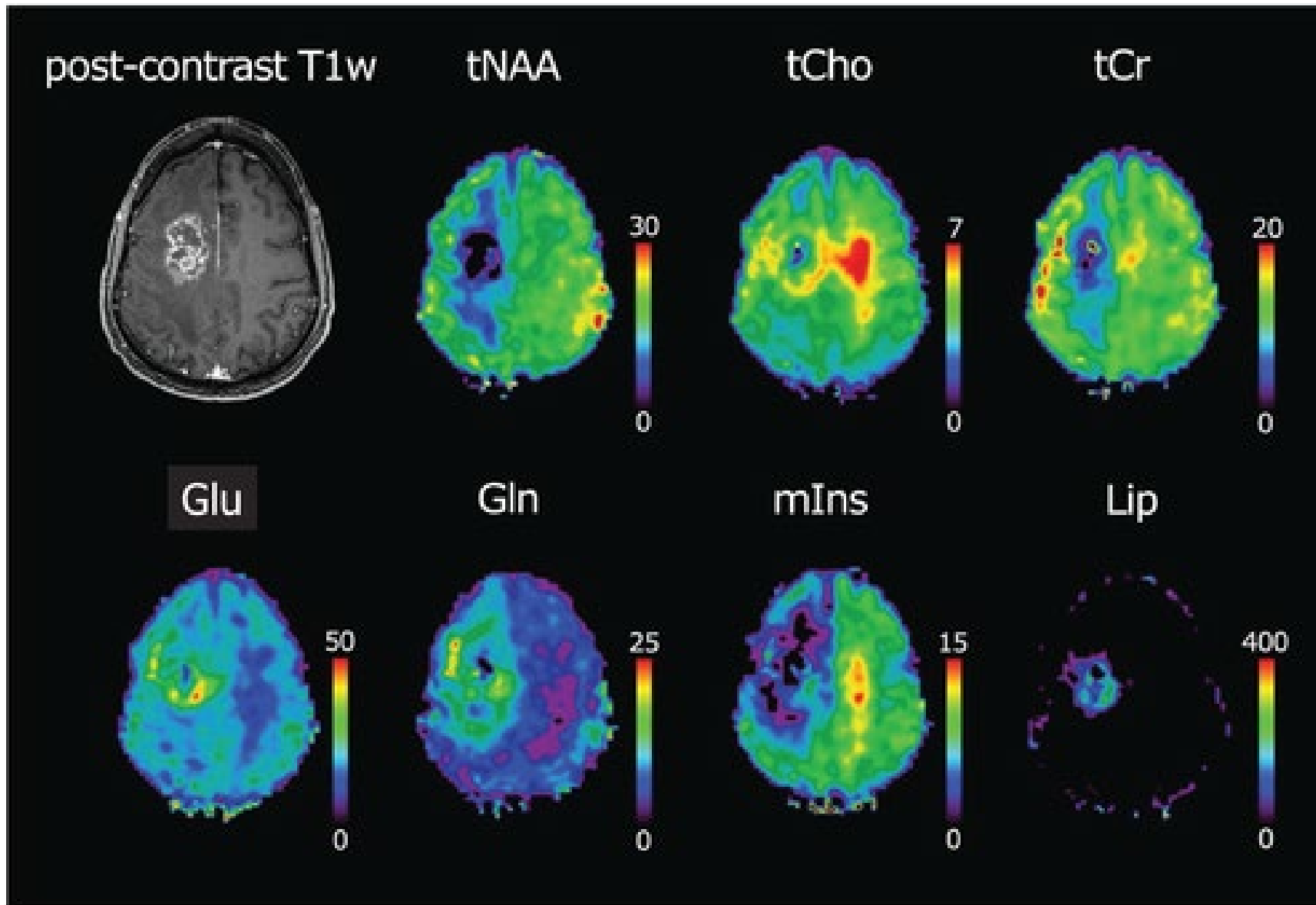
# FID-MRSI

- Around 2009??
- Short TR (200-800ms) ----Low Excitation Flip Angle
- Ultra short TE ( $\sim 1-1.3$  ms) – no J evolution, not too much signal loss due to T2 relaxation, but 1<sup>st</sup> order phase in the spectra
- Short acquisitions – 13min (31x31 with TR=811ms) but can go even lower ---- increase number of phase encoding steps – better PSF
- in plane spatial resolutions of 1.7-10 mm for humans



<https://www.epfl.ch/labs/mrs4brain/ressources/live-demos/>

**FID-MRSI: negligible J-coupling and  $T_2$  related signal loss, better suitability for fast MRSI at UHF in preclinical settings**



The single-slice FID-MRSI was acquired in 6 min with 6-fold accelerated phase-encoding, voxel size of 3.4 x 3.4 x 8 mm<sup>3</sup> and TR/AD of 600/1.3 ms.

[Key clinical benefits of neuroimaging at 7 T - ScienceDirect](#)

# TEMPORAL RESOLUTION

## MRSI

- $T_{measurement} = NA \times N_x \times N_y \times TR$ 
  - 1.5T, 16x16x16, NA=1, TR=2s → 136min

## SVS

- $T_{measurement} = NA \times TR$ 
  - ~ 5min – human (cm)
  - ~10-15min – preclinical (mm, SE sequence)

Received: 26 November 2019 | Revised: 24 March 2020 | Accepted: 30 March 2020  
DOI: 10.1002/nbm.4314

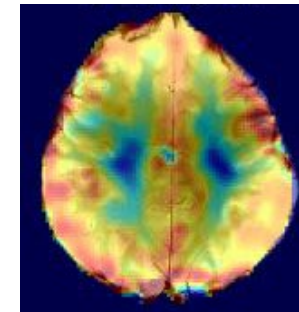
**SPECIAL ISSUE REVIEW ARTICLE**

**NMR**  
IN BIOMEDICINE WILEY

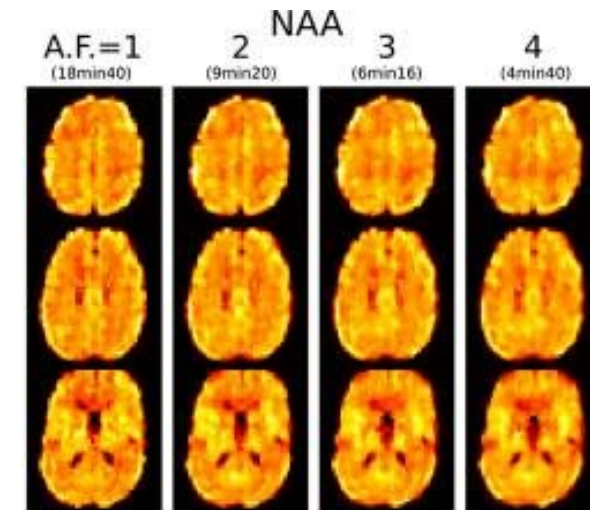
## Accelerated MR spectroscopic imaging—a review of current and emerging techniques

Wolfgang Bogner<sup>1</sup> | Ricardo Otazo<sup>2</sup> | Anke Henning<sup>3,4</sup>

- short TR
- acquisition of multiple k-space points per TR (e.g. spatial-spectral encoding)
- k-space undersampling (e.g. PI, CS)
- data reconstruction using spectral or spatial prior knowledge



Nassirpour S, et al, NeuroImage 2017



3.4x3.4x3.4mm<sup>3</sup>

A Klauser, et al, 2024.

# METABOLIC MAPPING via MRSI



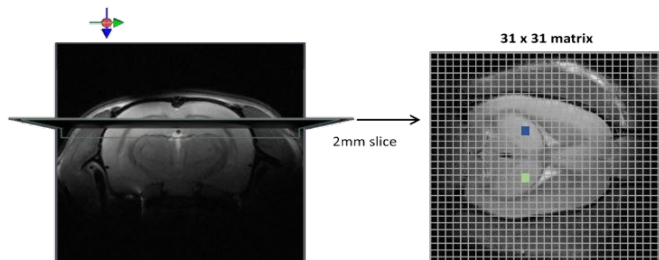
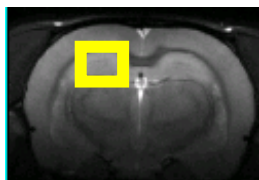
Tan Toi Phan

B Alves

A Siviglia

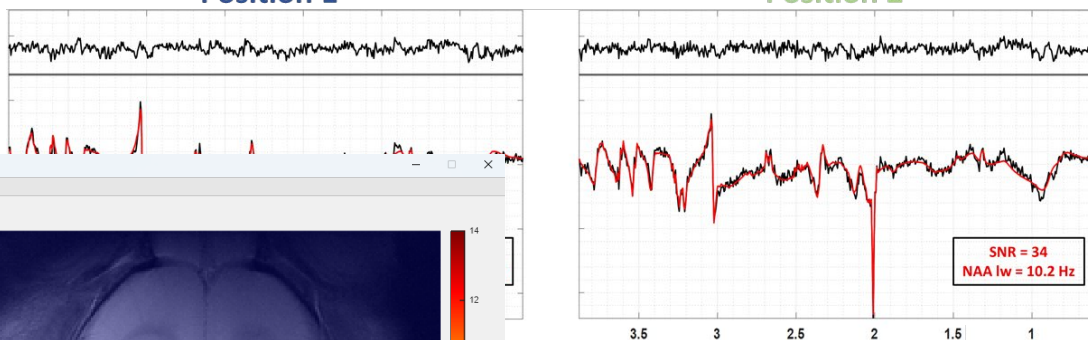


Voxel  $2 \times 2.8 \times 2 \text{ mm}^3$



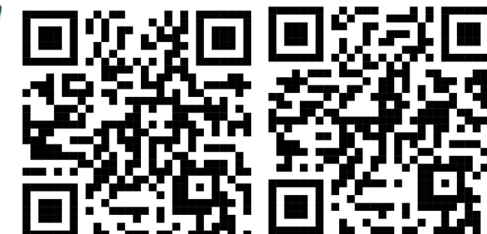
Position 1

Position 2

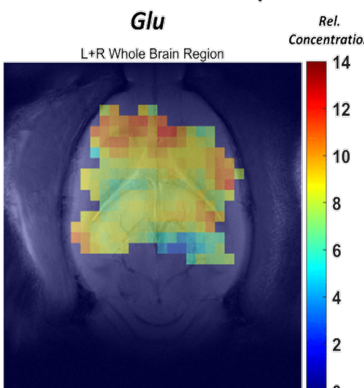


FID- MRSI  
 $0.7 \times 0.7 \times 2 \text{ mm}^3$   
 13 min @14.1T

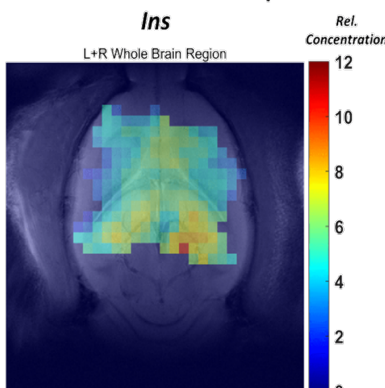
<https://www.epfl.ch/labs/mrs4brain/ressources/mrs4brain-toolbox/>



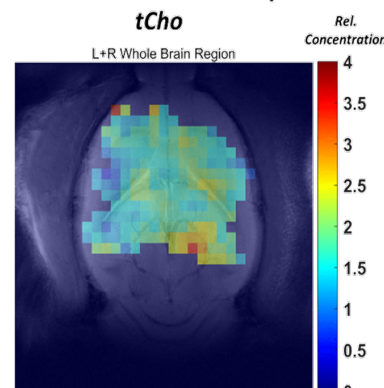
Relative metabolite map:



Relative metabolite map:



Relative metabolite map:

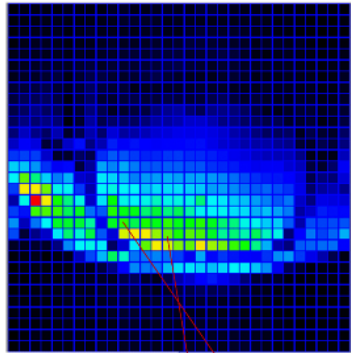


C I B M . C H

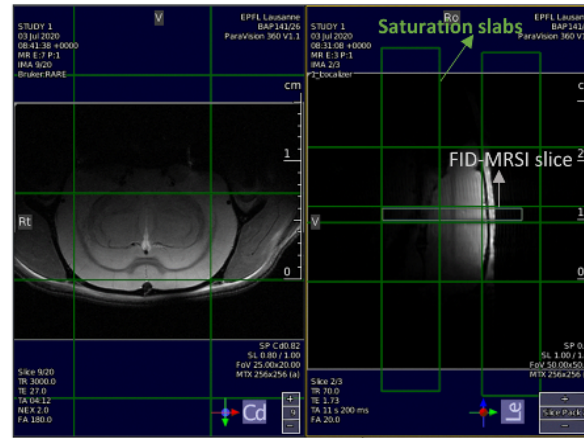
cristina.cudalbu@epfl.ch

B Alves, et al, NMR Biomed, 2024; D Simicic et al, NMR Biomed 2025

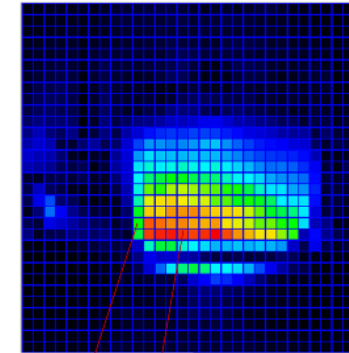
Water map  
NO FOV saturation



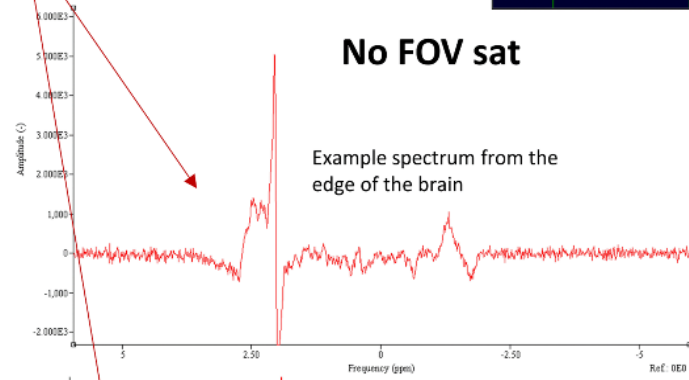
Addition of six 10mm thick saturation slabs



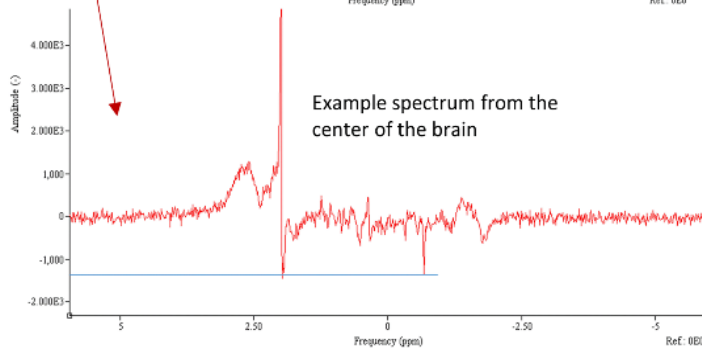
Water map  
With FOV saturation



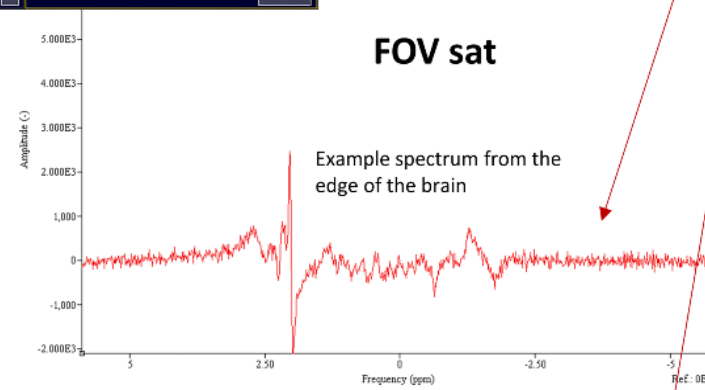
No FOV sat



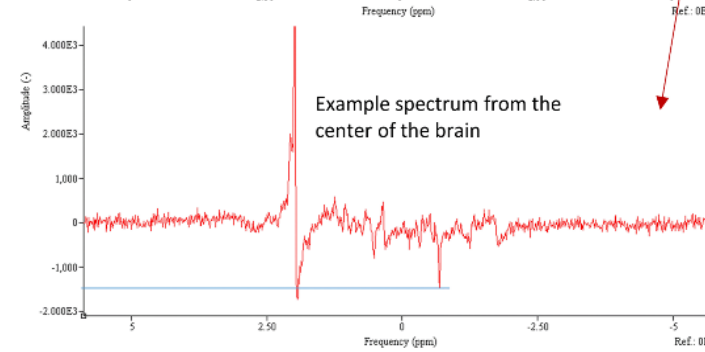
Example spectrum from the center of the brain



FOV sat

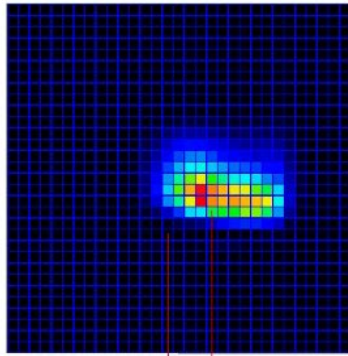


Example spectrum from the center of the brain

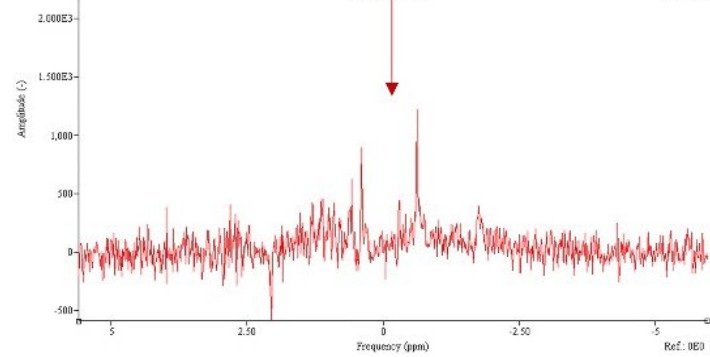
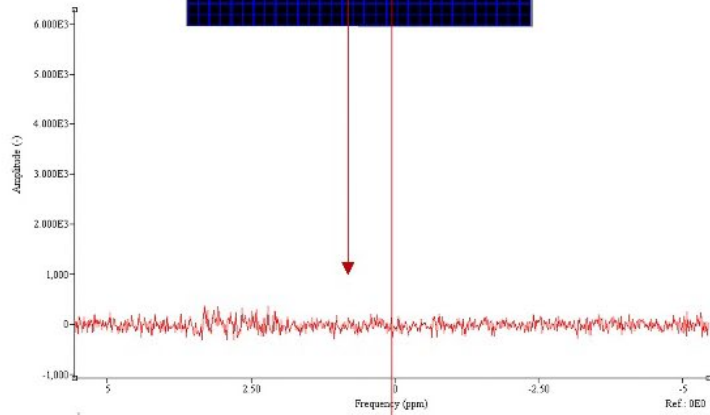
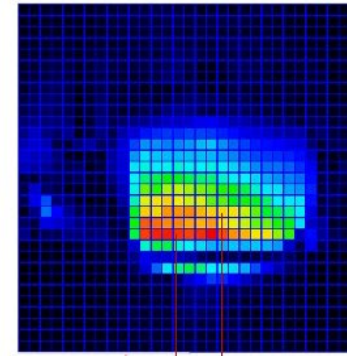


D Simicic  
<https://infoscience.epfl.ch/record/296066>

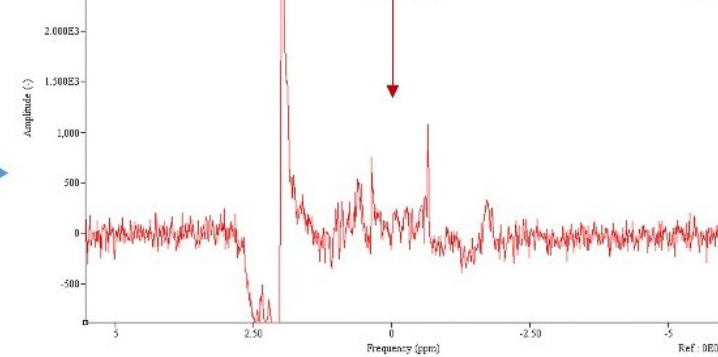
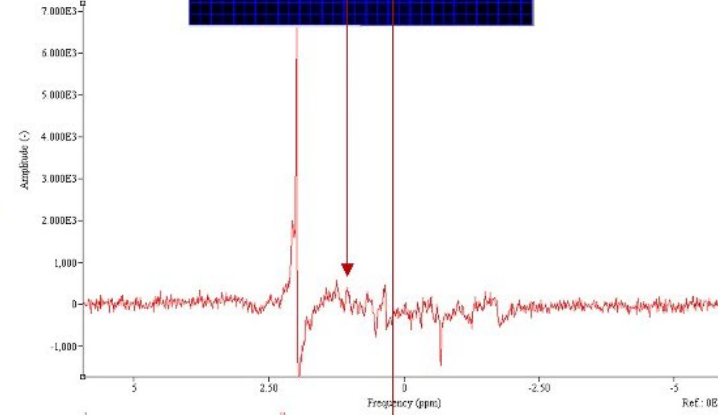
### Water Map PRESS-MRSI



### Water Map FID-MRSI



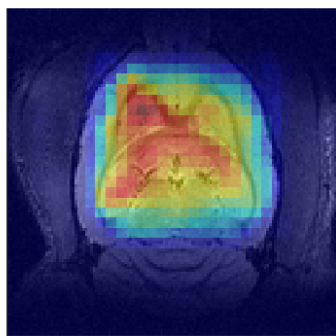
The same positions in the matrix



D Simicic  
<https://infoscience.epfl.ch/record/296066>

# FID-MRSI

Water Power Map

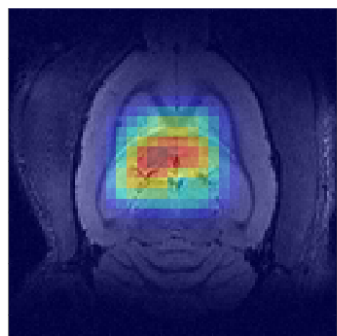


$\times 10^{16}$

15  
10  
5

# PRESS-MRSI

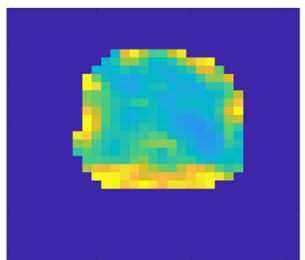
Water Power Map



$\times 10^{17}$

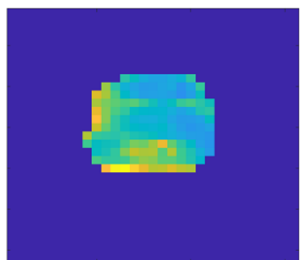
7  
6  
5  
4  
3  
2  
1

Linewidth [Hz]



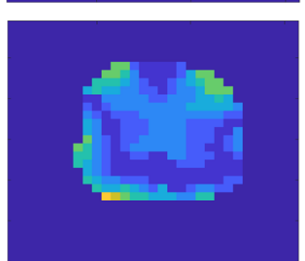
60  
50  
40  
30  
20  
10  
0

Linewidth [Hz]



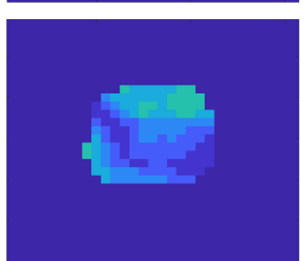
60  
50  
40  
30  
20  
10  
0

$|\Delta B_0|$  shift [Hz]



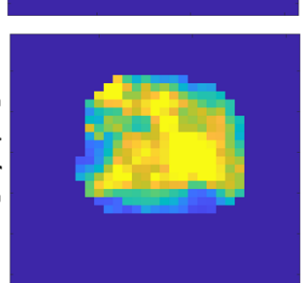
60  
50  
40  
30  
20  
10  
0

$|\Delta B_0|$  shift [Hz]



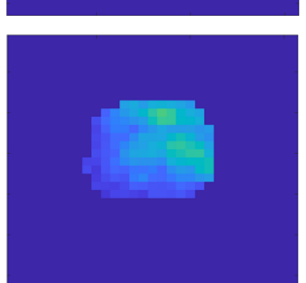
60  
50  
40  
30  
20  
10  
0

SNR per unit time  
[ $1/\sqrt{s}$ ]

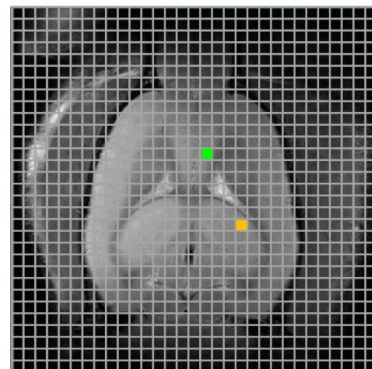


0.5  
0.4  
0.3  
0.2  
0.1  
0

SNR per unit time  
[ $1/\sqrt{s}$ ]



0.5  
0.4  
0.3  
0.2  
0.1  
0



FID-MRSI

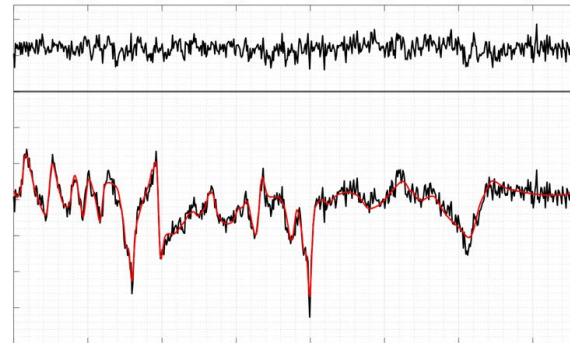
PRESS-MRSI

Voxel  
1

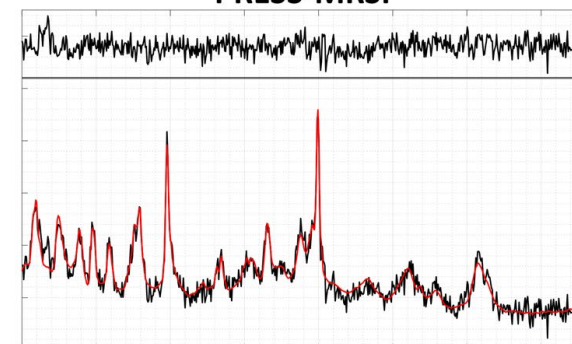


4 3.5 3 2.5 2 1.5 1 0.5

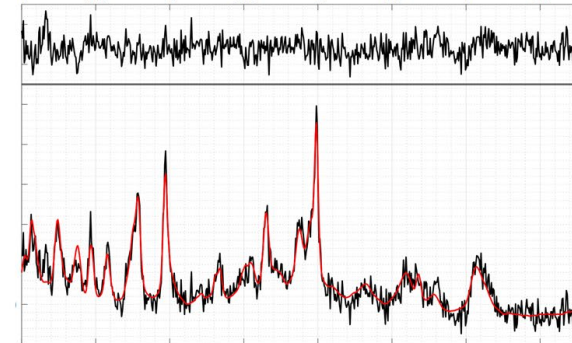
Voxel  
2



4 3.5 3 2.5 2 1.5 1 0.5  
ppm

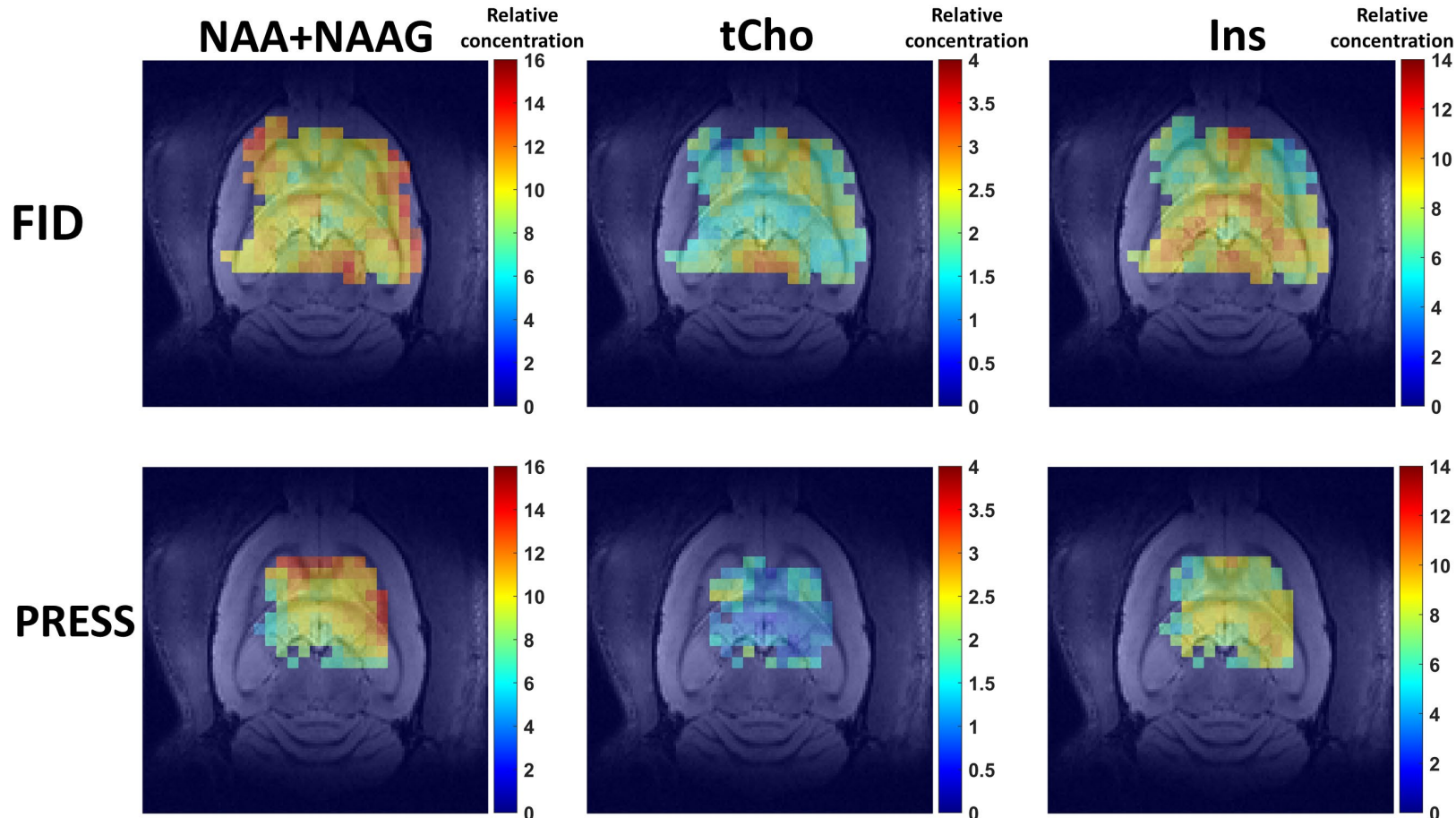


4 3.5 3 2.5 2 1.5 1 0.5



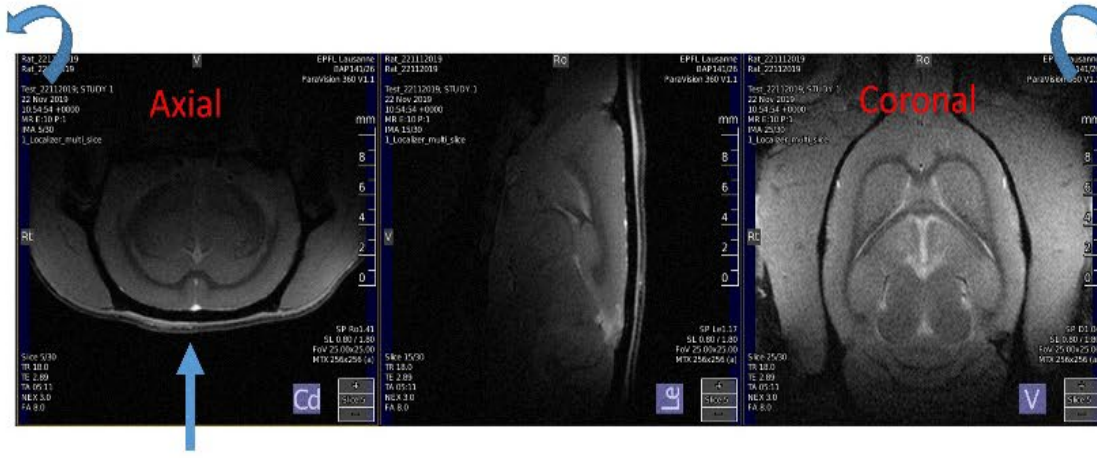
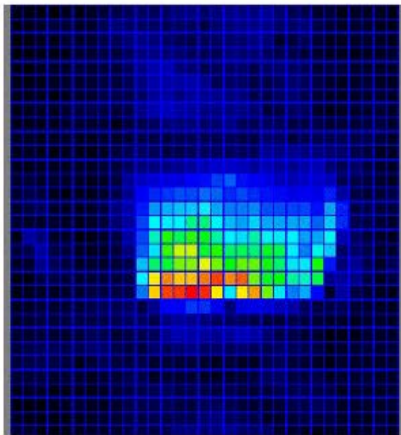
4 3.5 3 2.5 2 1.5 1 0.5  
ppm

# RESULTS : METABOLITE MAPS



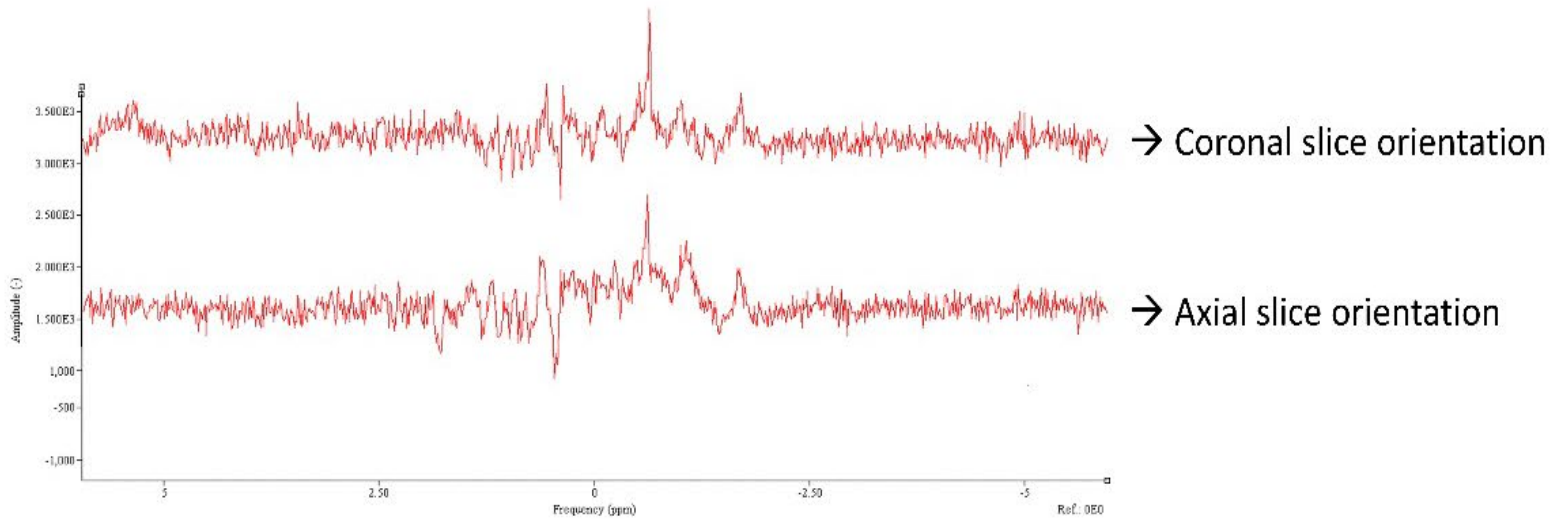
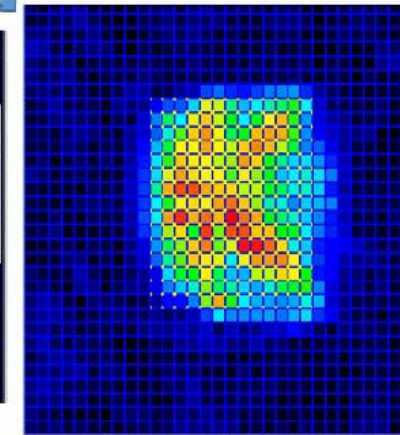
- FID-MRSI provided an increased brain coverage even after the application of quality control criteria (min. SNR, max. FWHM, max. CRLBs limits)
- Minor differences in metabolite concentrations were noted between the two sequences (underestimation for PRESS-MRSI) which might be related to the lack of metabolite  $T_1$  and  $T_2$  corrections, etc, something requiring further investigations.

Axial slice – Water map



Surface coil

Coronal slice – Water map



# SUMMARY & ACKNOWLEDGMENTS

MRS & MRSI is incredibly rich & versatile 😊

THANK YOU – D Simicic and B Alves; B Lanz and G Briand

Thank you for listening! Questions?

For any question you might have later on, please write me an email:

[cristina.cudalbu@epfl.ch](mailto:cristina.cudalbu@epfl.ch)



THANK YOU FOR YOUR ATTENTION



Avec le soutien de:

