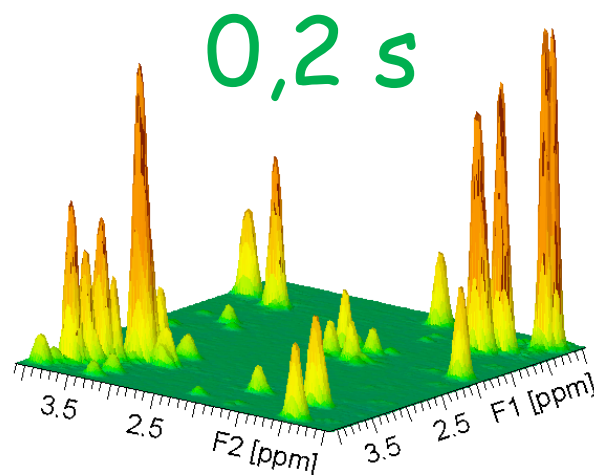


# Analytical Ultrafast 2D NMR



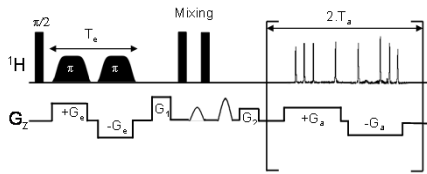
**Patrick GIRAUDEAU**

CEISAM, Université de Nantes, CNRS, Nantes, France

# Analytical ultrafast 2D NMR

## Main subjects in our research group

### NMR method development



- High precision quantitative pulse sequences
- Quantitative nD methods
- Ultrafast (UF) NMR**
- Localized spectroscopy

### Analytical aspects

- Robustness
- Linearity
- Accuracy



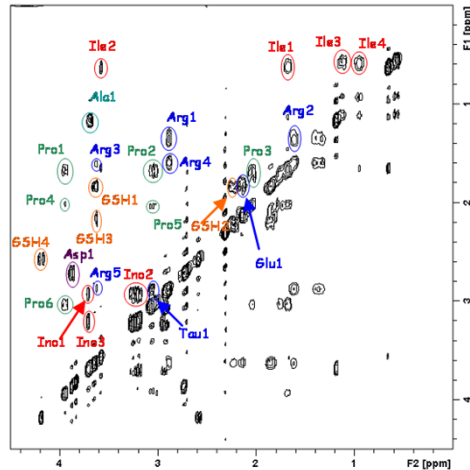
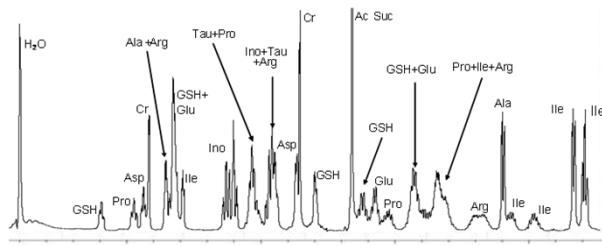
## Quantitative NMR methods

### Quantitative applications

- Quantitative analysis of biological samples (understanding metabolism)
- Isotopic analysis @ natural abundance (authentication of food, drugs...)
- Isotopic analysis in enriched media (fluxomics)

# Analytical ultrafast 2D NMR

## Quantitative 2D NMR

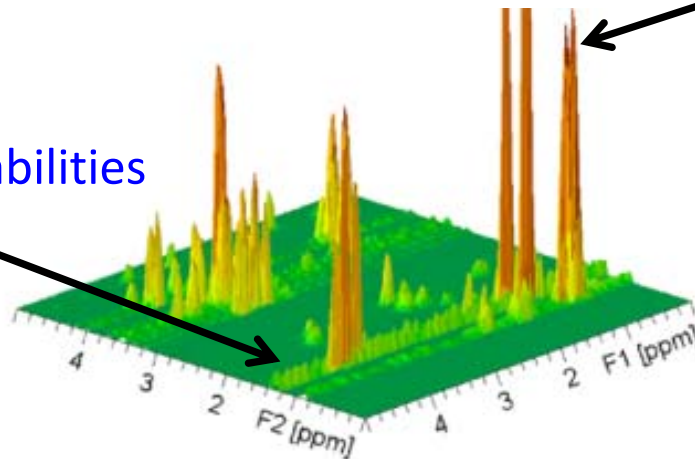


Unambiguous quantification with higher precision thanks to reduced overlap

## Challenges

### Long experiment duration

- timetable constraints
- instable samples
- sensitivity to hardware instabilities



### 2D response

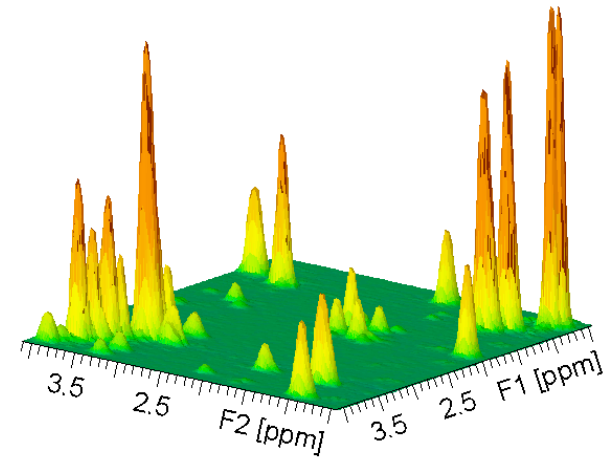
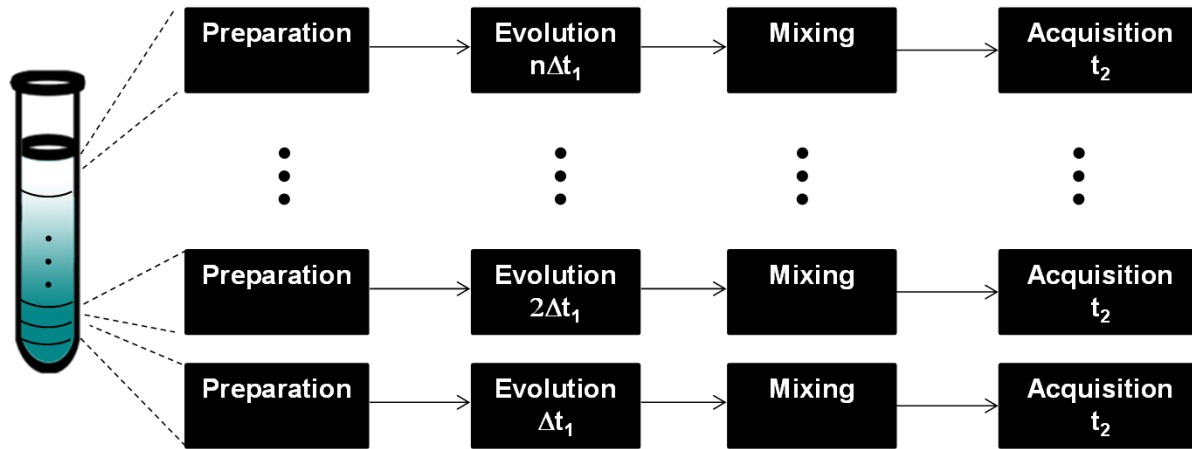
is peak dependant ( $T_2$ ,  $J$ ...)



Calibration is needed !



## Ultrafast 2D NMR for quantitative analysis?



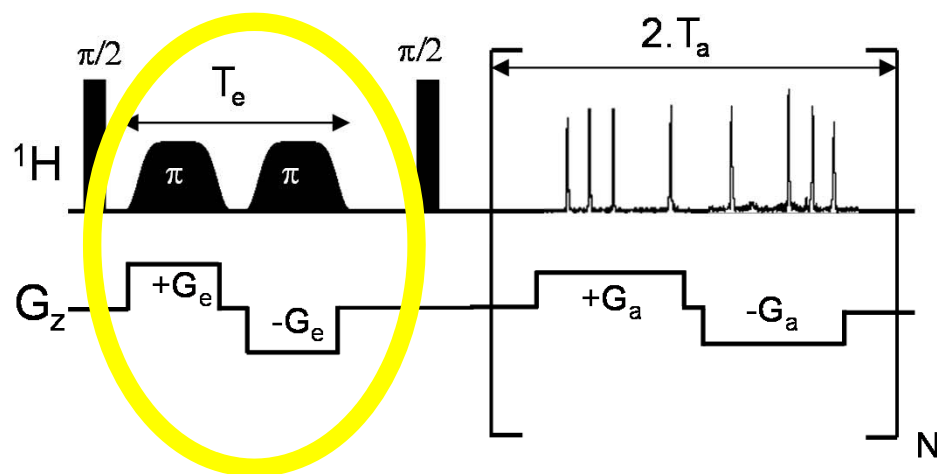
1. Increasing the analytical performance
2. Applications in quantitative metabolomics
3. Applications in fluxomics
4. Other recent analytical applications

## Limitations of UF 2D NMR

Compromise SW/resolution

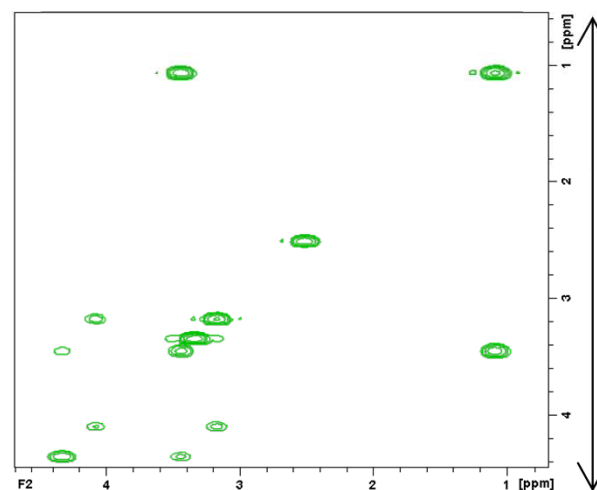
$$\gamma \cdot G_a \cdot L = \frac{2 \cdot SW_1 \cdot SW_2}{\Delta v_2}$$

Compromise  
Sensitivity/Resolution



Ultrafast dimension:  $SW_2$

$\Delta v_2$ : Peak width -  $F_2$

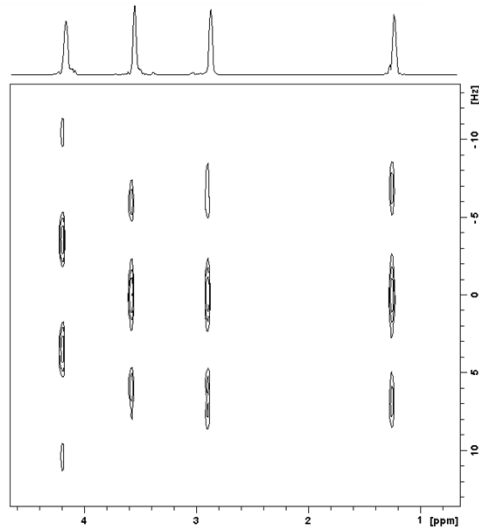


Conventional  
dimension:  $SW_1$

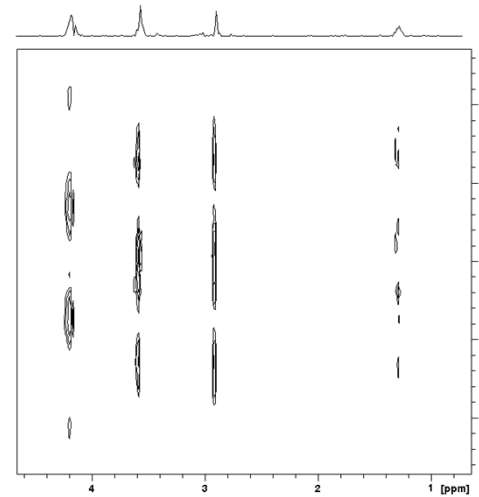
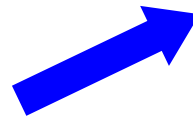
$\Delta v_1$ : Peak width -  $F_1$

# Increasing the analytical performance

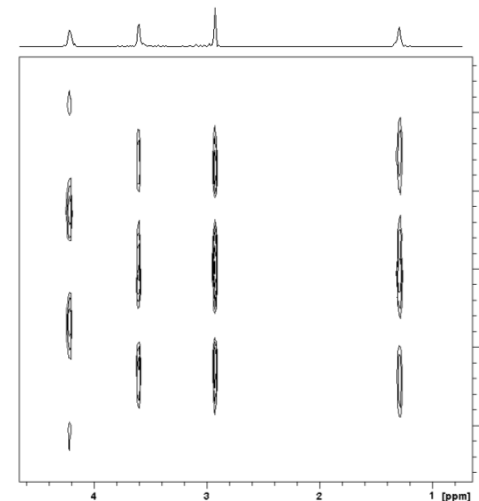
## Reducing diffusion effects - multi-echo encoding



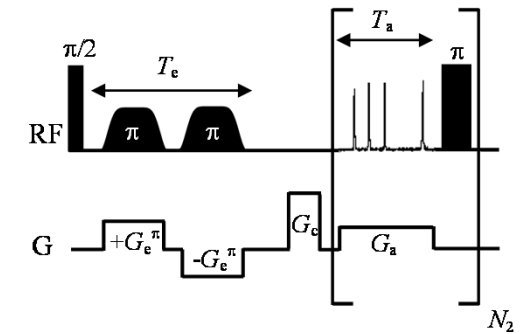
UF Jres,  $T_e = 60$  ms  
 $\Delta\nu = 17$  1 Hz



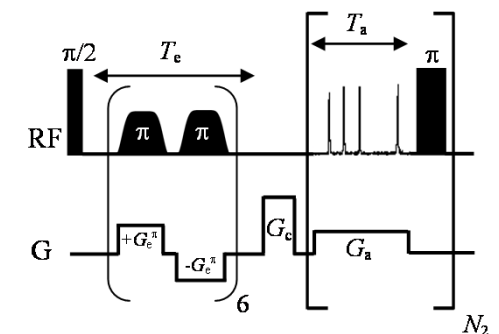
UF Jres,  $T_e = 120$  ms  
 $\Delta\nu = 13$  1 Hz



UF Jres multi-écho,  $T_e = 120$  ms  
 $\Delta\nu = 13$  1 Hz



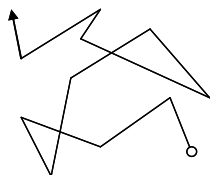
Better resolution  
 Sensitivity losses  
 (diffusion)



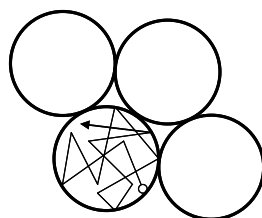
Better resolution  
 Better sensitivity

# Increasing the analytical performance

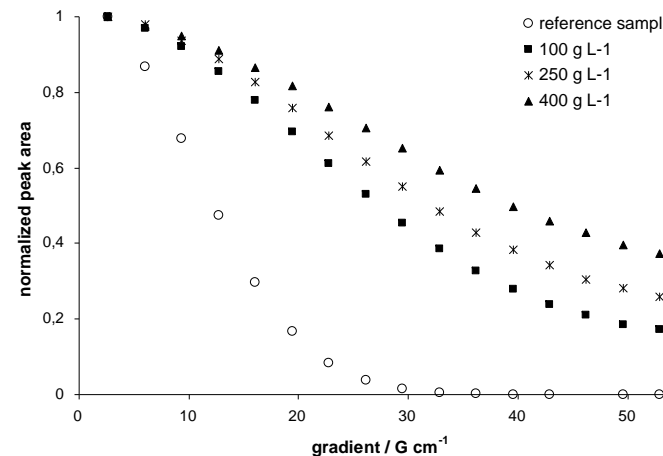
## Reducing diffusion effects - sample preparation



Free diffusion

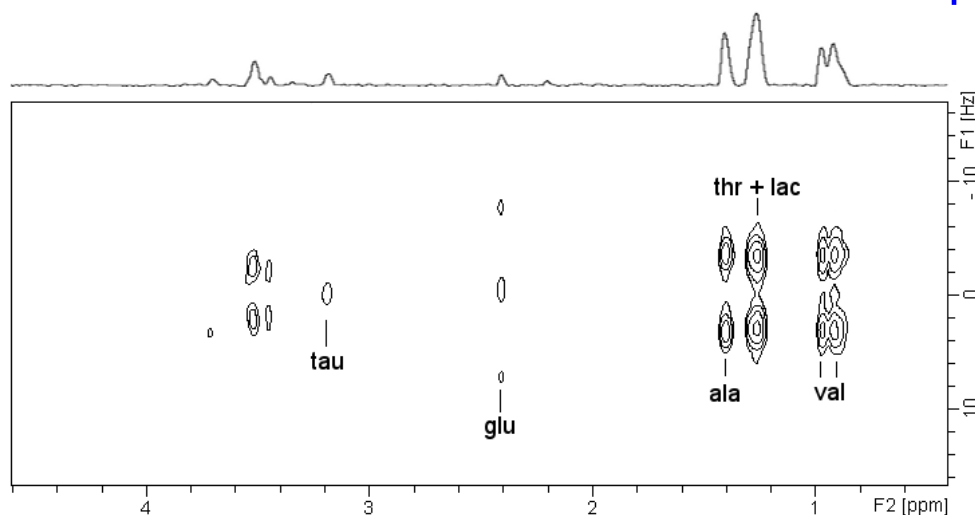


Restricted diffusion  
In liposomes

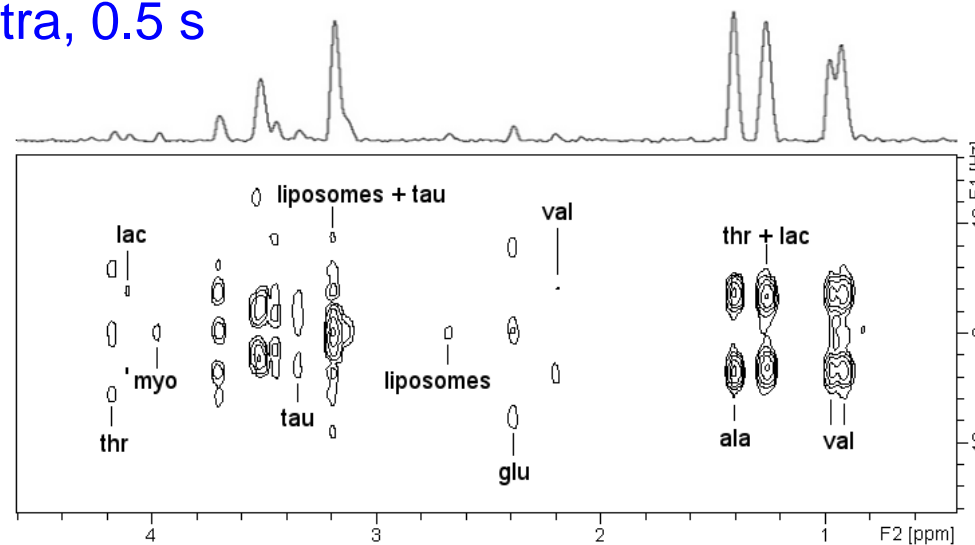


Diffusion curves - DSTE

UF Jres spectra, 0.5 s



Metabolic sample in D<sub>2</sub>O

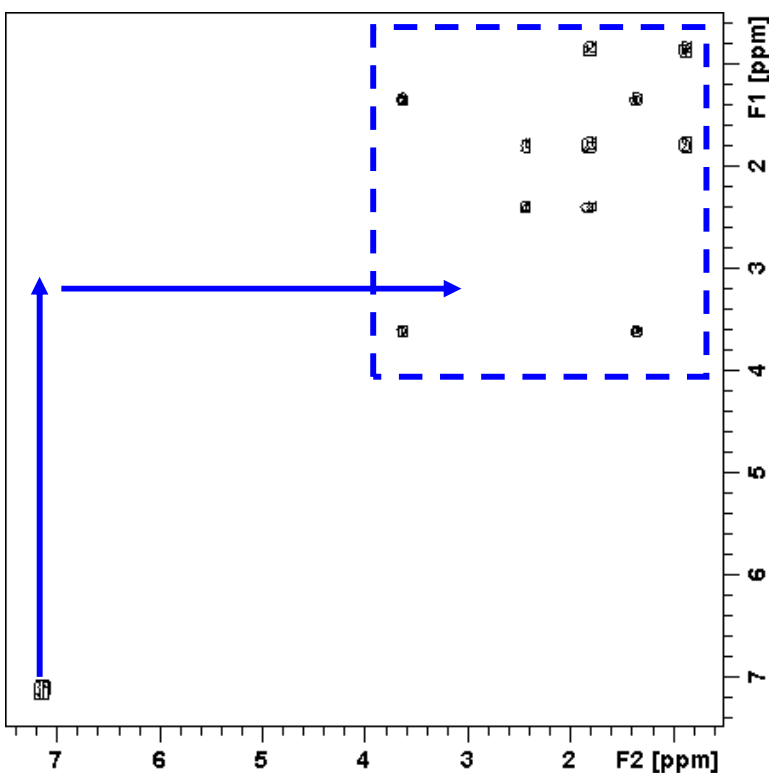
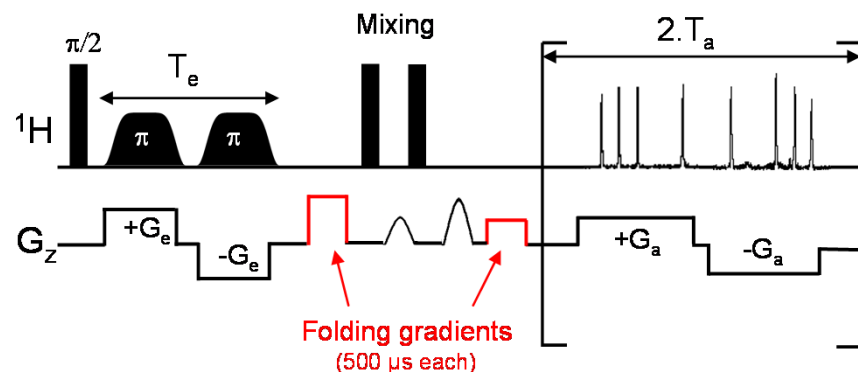
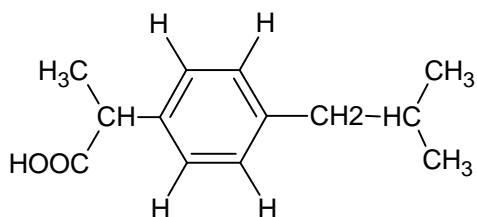


Metabolic sample in D<sub>2</sub>O + liposomes

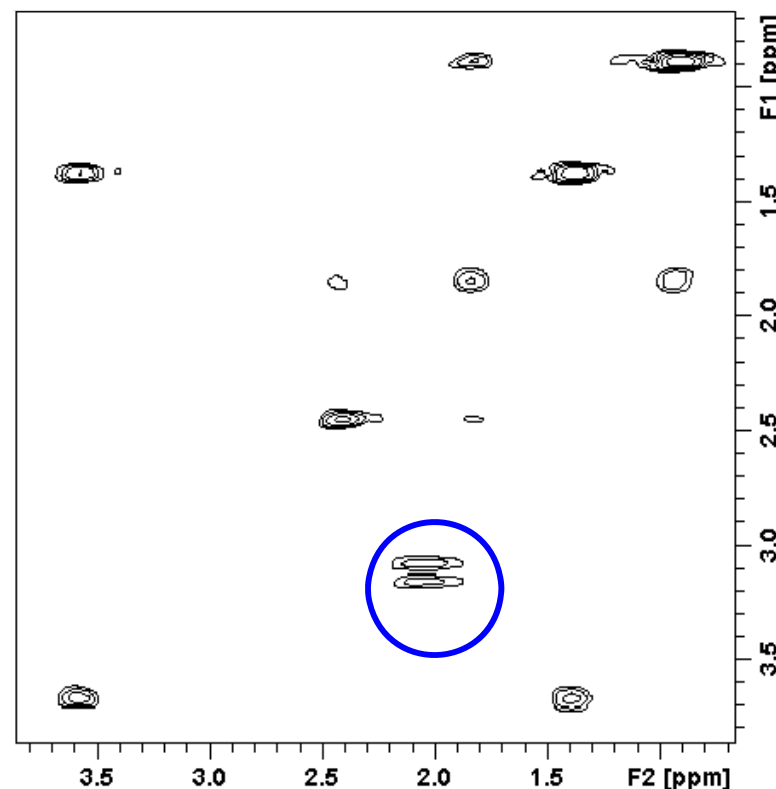


# Increasing the analytical performance

## Increasing the spectral range - folding



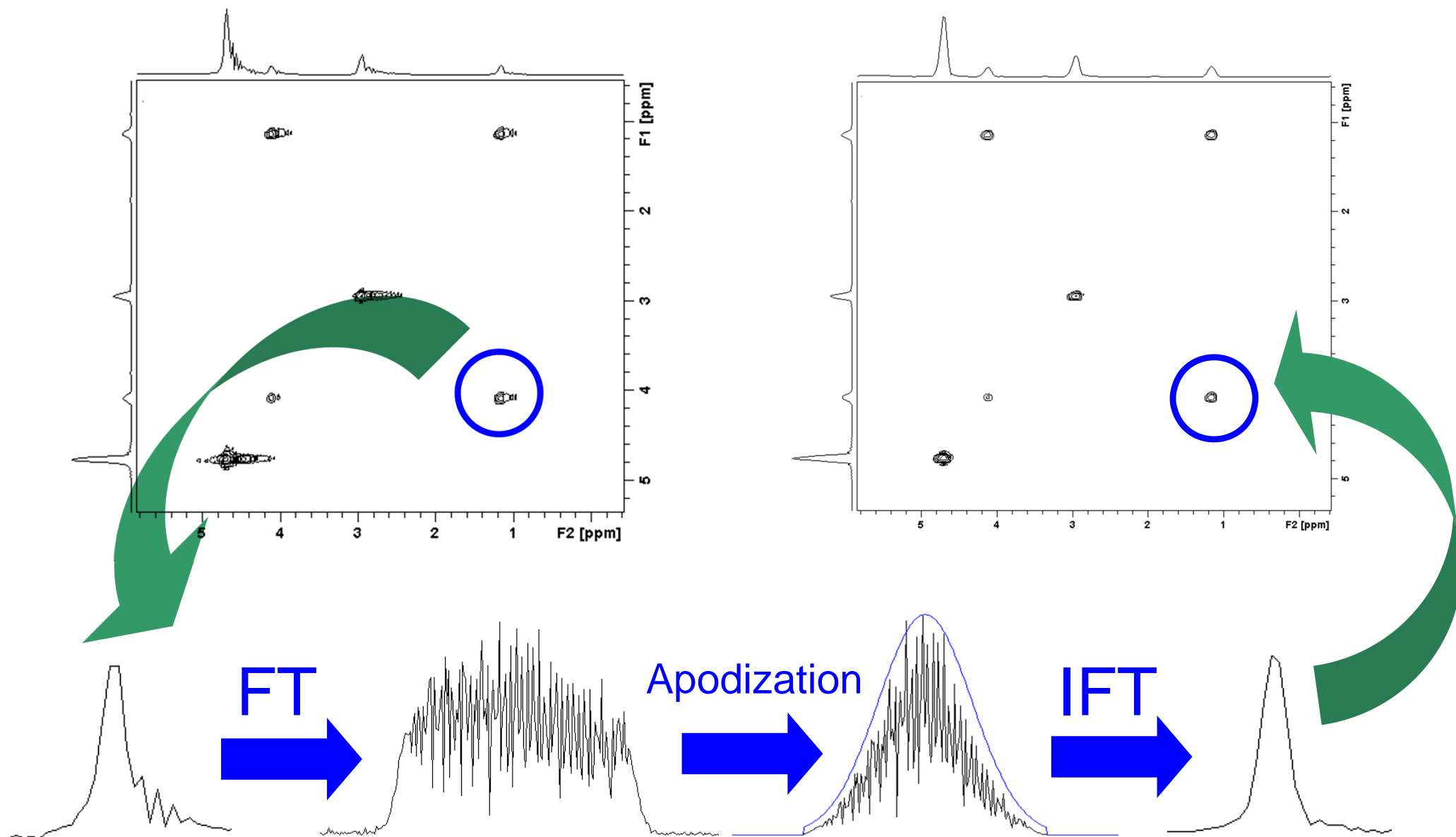
Conventional DQF-COSY – 13 min



Ultrafast folded DQF-COSY – 0.2 s

# Increasing the analytical performance

## Improving the lineshape by processing





## Making UF 2D NMR more accessible

Dedicated webpage including:

- pulse sequences
- implementation protocol
- processing routine
- web interface for parameter setting

UNIVERSITÉ DE NANTES

Centre Et Interdisciplinaire  
Synthèse, Analyse,  
Catalyse

UNIVERSITÉ DE NANTES

### Implementing ultrafast 2D NMR experiments on a Bruker Avance Spectrometer

Patrick Giraudeau, Benoît Charrier,  
Meerakhan Pathan, Serge Akoka

EBSI group – CEISAM laboratory  
[http://www.sciences.univ-nantes.fr/CEISAM/en\\_ebsi.php](http://www.sciences.univ-nantes.fr/CEISAM/en_ebsi.php)  
[patrick.giraudeau@univ-nantes.fr](mailto:patrick.giraudeau@univ-nantes.fr)

last updated on 17/01/2013

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www.sciences.univ-nantes.fr/CEISAM/ebsi/ultrafast/uf\_acq.html

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## Acquisition Parameters in UltraFast NMR

Implementation of Ultrafast 2D NMR experiments:

The protocol, pulse sequences and processing program to implement ultrafast 2D NMR experiments on your spectrometer are available for download here:  
<http://madoc.univ-nantes.fr/course/view.php?id=24710>

Please first read carefully the protocol and download the files. Once the ultrafast experiments are implemented on your spectrometer, use the following pages to set up your acquisition parameters:

Type of experiment:

Pulse sequence : [UFCOSY](#)  
[UFHSQC](#)

[Troubleshooting](#)  
You can [here](#) download the pages to run on a local computer.

[QUANTUM project](#): Quantitative Ultrafast Analysis by 2D NMR To Unravel Metabolic complexities 2011-2014 (ANR grant 2010-JCJC-0804-01)

*Important remark: Ultrafast 2D NMR has been patented by the Weizmann Institute of Science, Israel. Its use for commercial purposes requires a licence from the Weizmann Institute.*

Rechercher : NMR

Suivant Précédent Tout surligner Respecter la casse

# Increasing the analytical performance

2006      2007      2008      2009      2010      2011      2012      2013



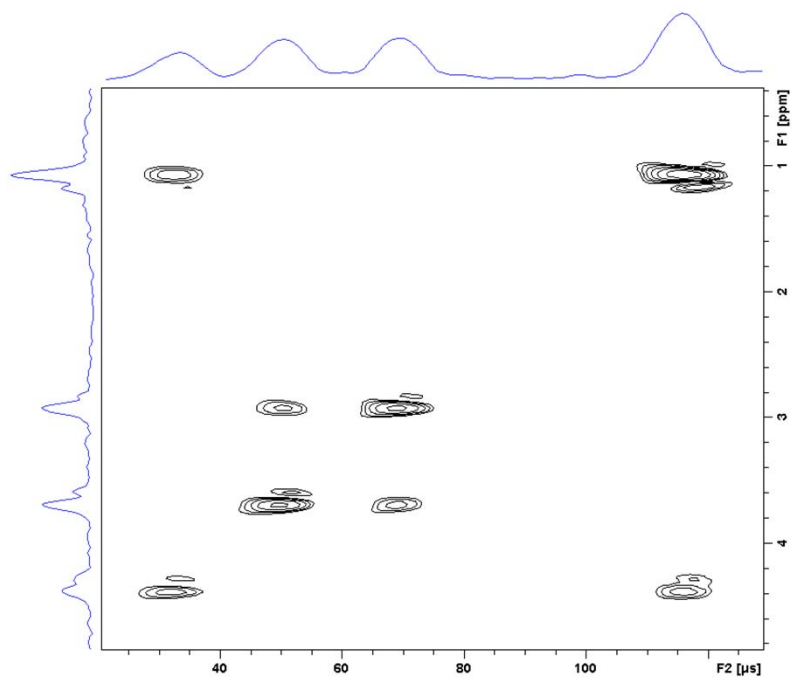
1<sup>st</sup> ultrafast  
experiment  
at CEISAM

Sensitivity  
and resolution  
improvements

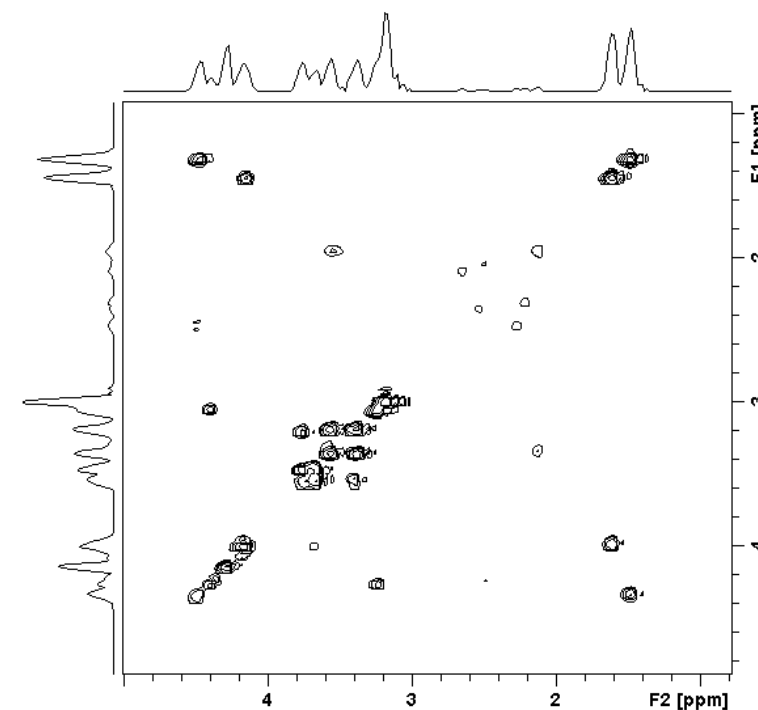
Spectral  
width  
improvements

Lineshape  
improvement

Automated  
acquisition and  
processing



UF COSY 0.1 s - model sample



UF COSY 0.1 s - metabolites

## Context: NMR metabolomics

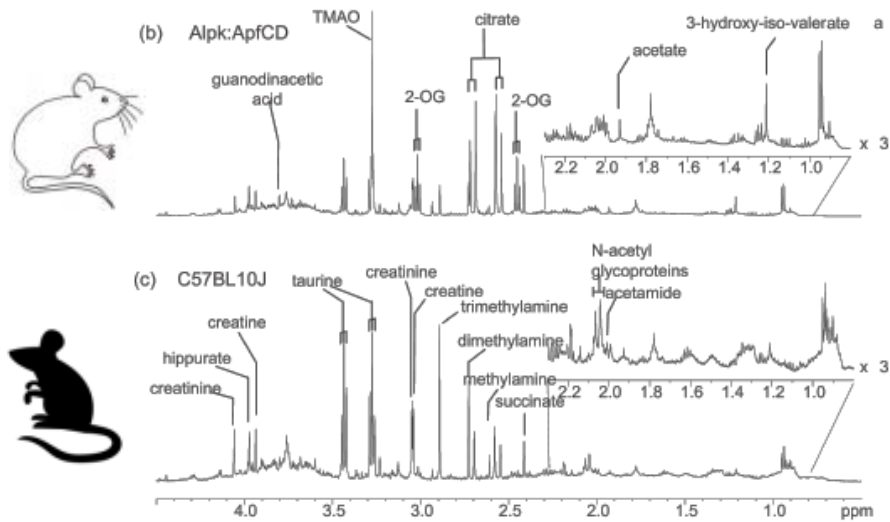
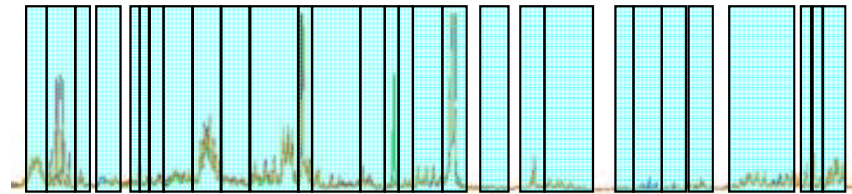
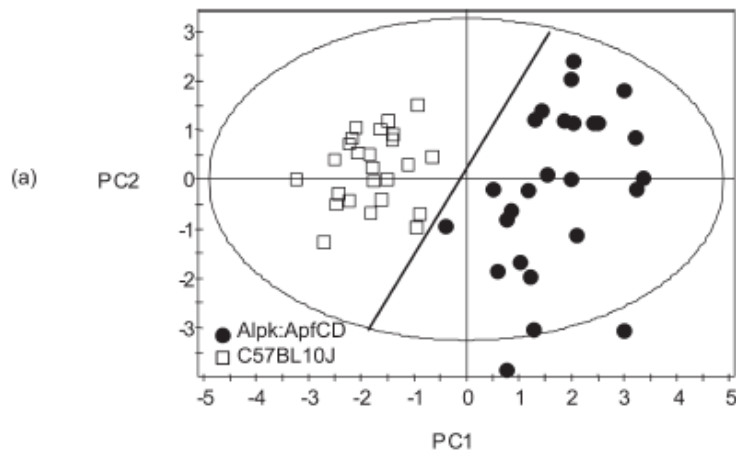


Figure 4. (a) PC scores plot derived from the 500 MHz  $^1\text{H}$  NMR spectra of urine samples from (b) Alpk:ApfCD (white) mice and (c) C57BL10J (black) mice. 2-OG, 2-oxoglutarate; TMAO, trimethylamine-*N*-oxide

Spectral decomposition (bucketing)



Statistical treatment



Individual quantification is difficult due to overlap

# Applications to quantitative metabolomics

## Ultrafast quantitative 2D NMR ?

*Anal. Chem.* 2009, 81, 479–484

### Evaluation of Ultrafast 2D NMR for Quantitative Analysis

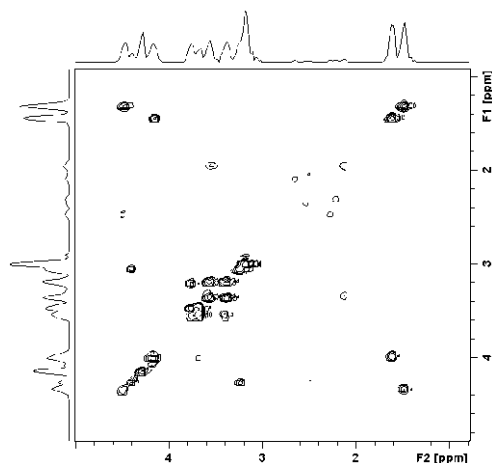
Patrick Giraudeau,\* Gérald S. Remaud, and Serge Akoka

Université de Nantes, CNRS, CEISAM UMR 6230, B. P. 92208, 2 Rue de la Houssinière,  
F-44322 Nantes Cedex 03, France



High  
repeatability  
and linearity

## What about metabolic mixtures ?



UF COSY 0.1 s  
metabolites

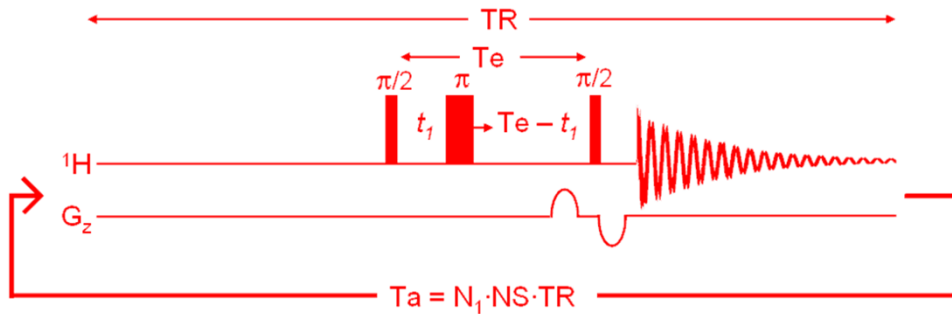
LOQ (1 scan) = 10 mM (500 MHz cryo)



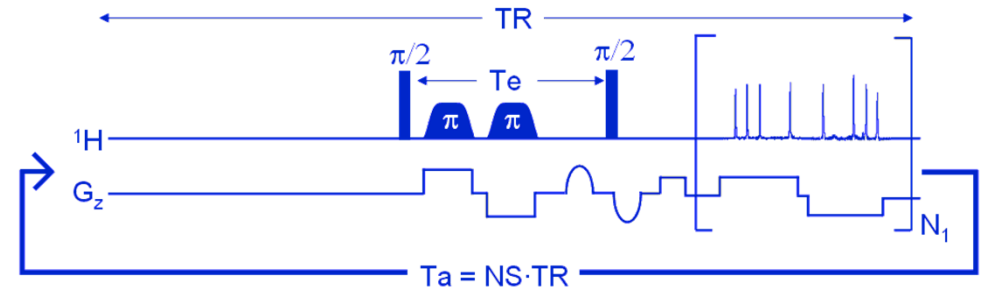
Signal averaging necessary !!!



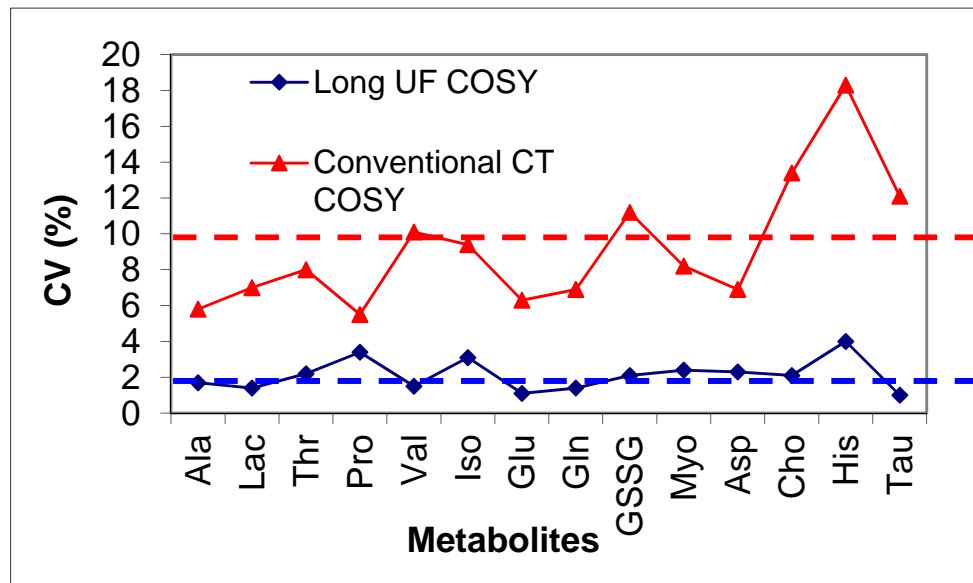
## « Long-ultrafast » vs conventional 2D NMR



Conventional CT COSY



« Long ultrafast COSY »



Repeatability of  
**Conventional CT COSY**  
 VS  
 « Long ultrafast COSY »

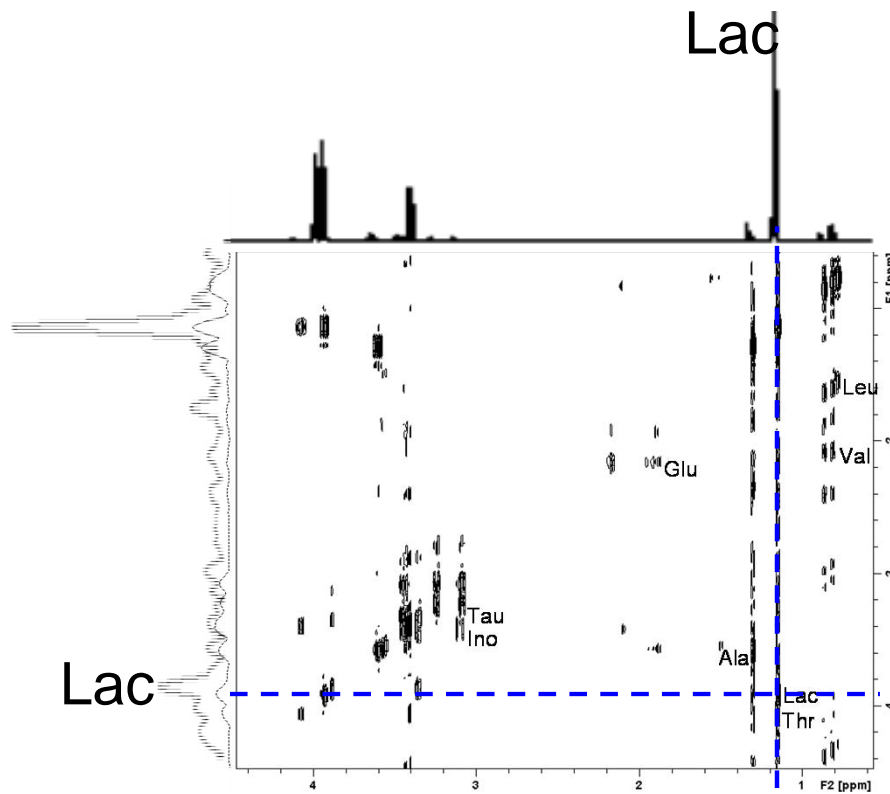


**Better  
 precision**

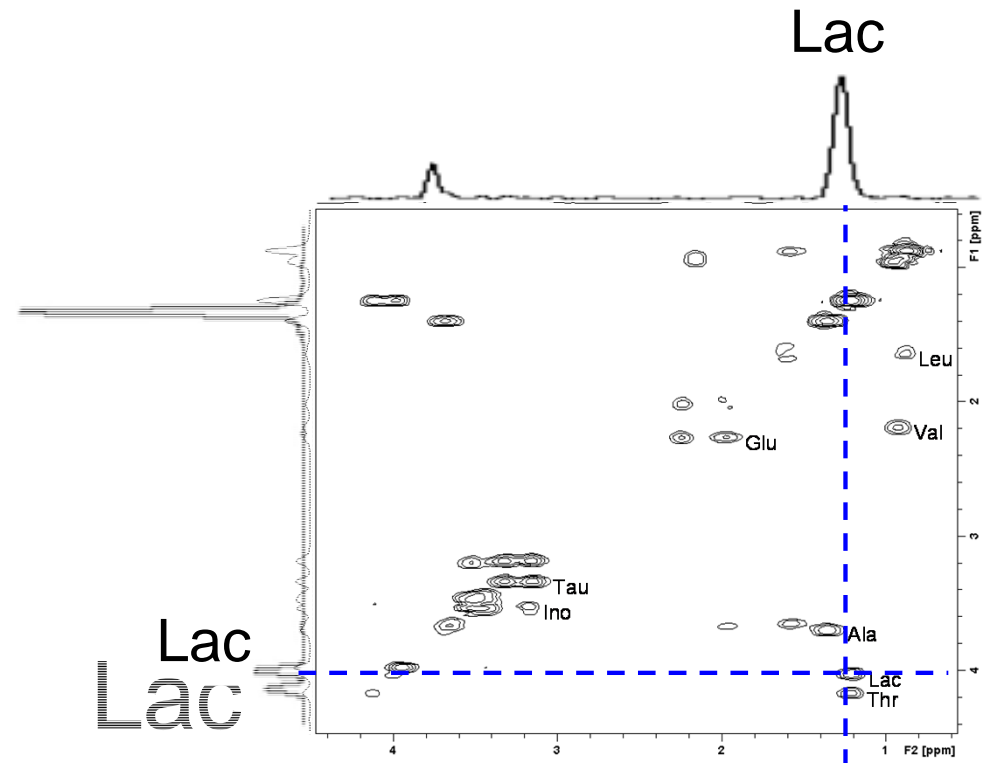
# Applications to quantitative metabolomics

## Why is long UF more repeatable ?

UF 2D NMR is more immune to hardware instabilities  
This is particularly true in quantitative conditions



Conv. COSY, 34 min  
NS = 1,  $N_1 = 64$



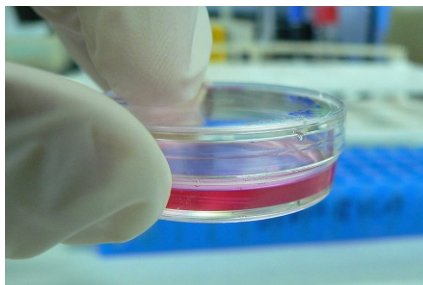
« Long ultrafast » COSY  
34 min  
NS = 64





## Application to biological samples

1. Growth of various breast cancer cell lines

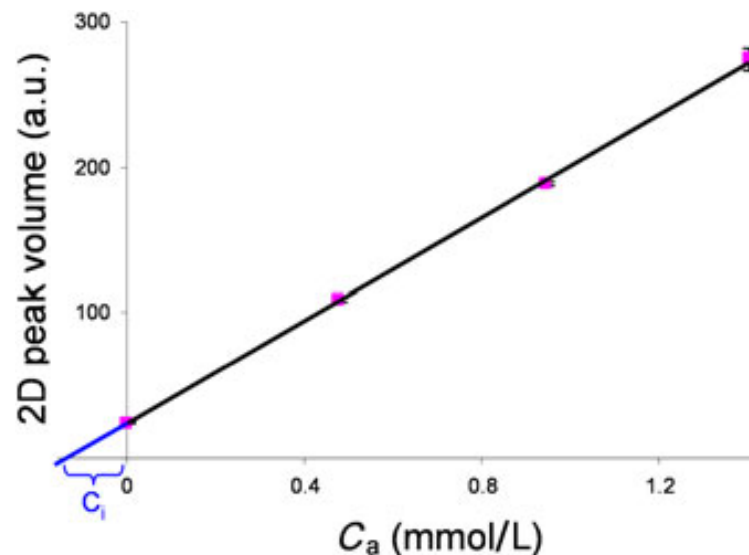


2. Optimized metabolite extraction



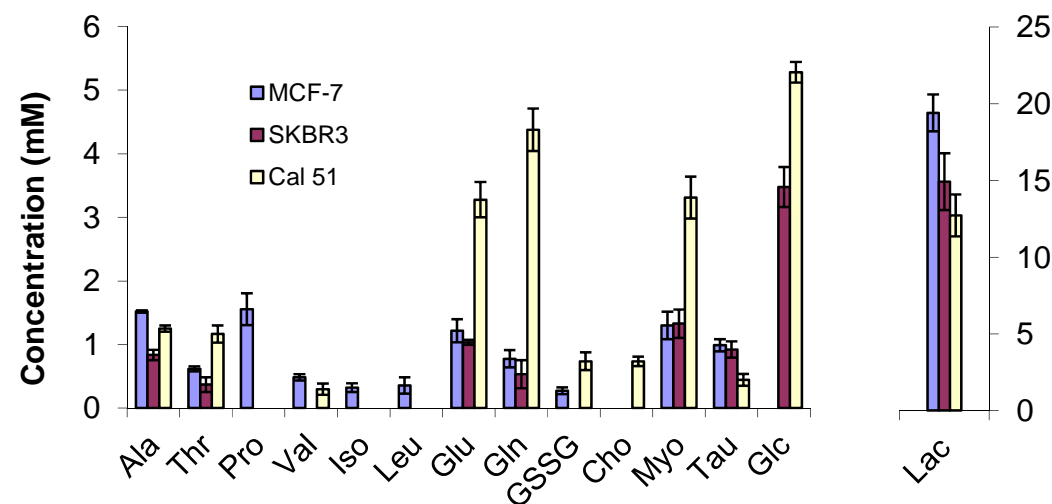
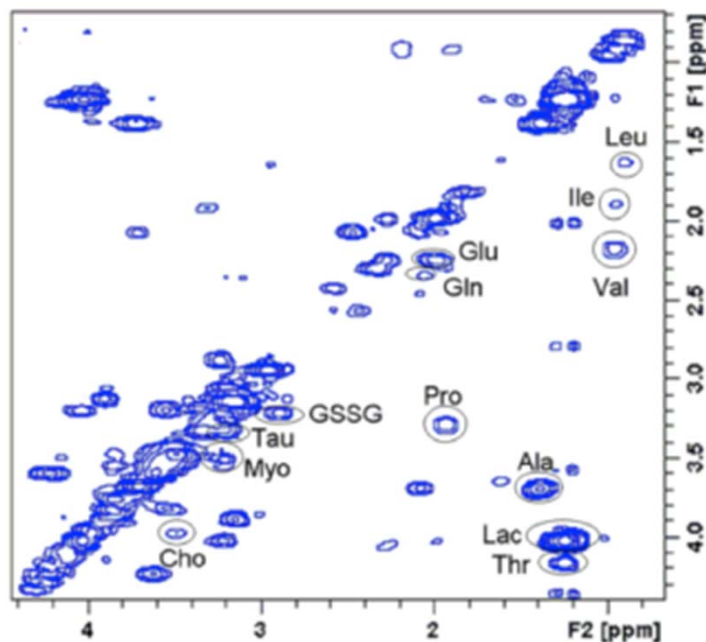
E. Martineau, I. Tea, P. Giraudeau, S. Akoka, *Anal. Bioanal. Chem.* 2011

3. NMR quantification with standard addition procedure





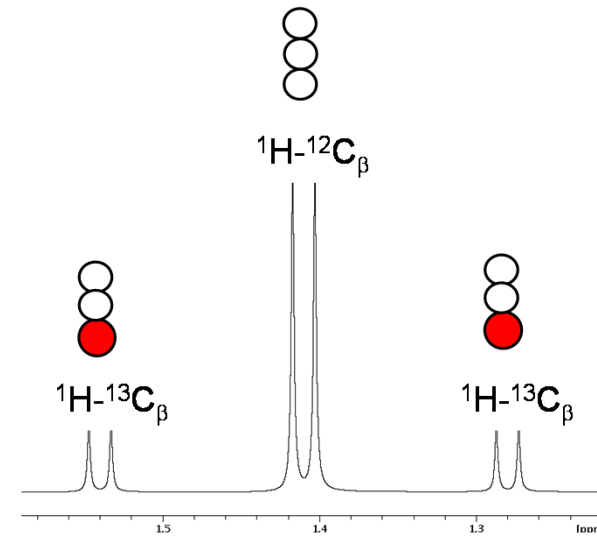
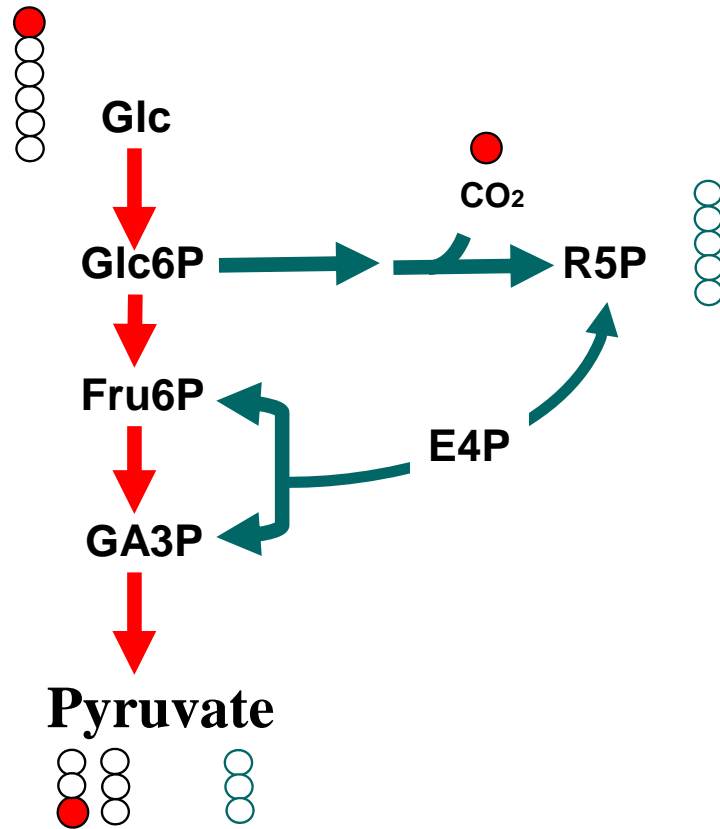
## Application to biological samples



- Comparison of 3 breast cancer cell lines
- 20 min / spectrum
- Standard addition procedure

# Applications in fluxomics

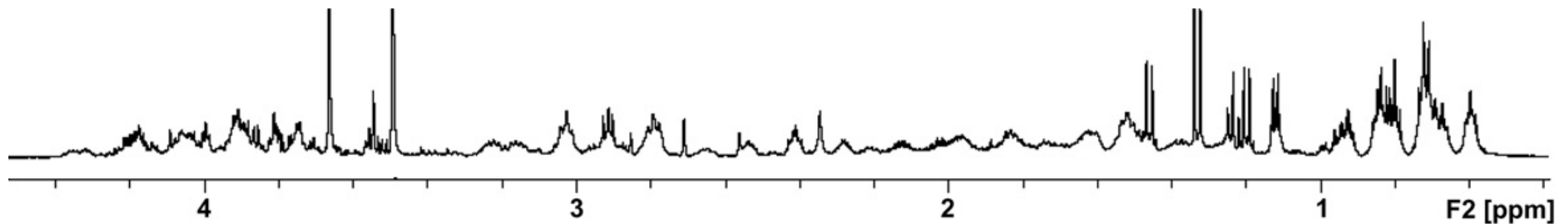
Context : measurement of  $^{13}\text{C}$  isotopic enrichments in fluxomics



Simple cases: 1D  $^1\text{H}$  NMR

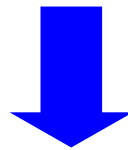
$$\text{Site-specific } ^{13}\text{C} \text{ isotopic enrichment (IE)} = S^{\text{satellites}} / S^{\text{total}}$$

## Limitations of 1D NMR - biological mixtures



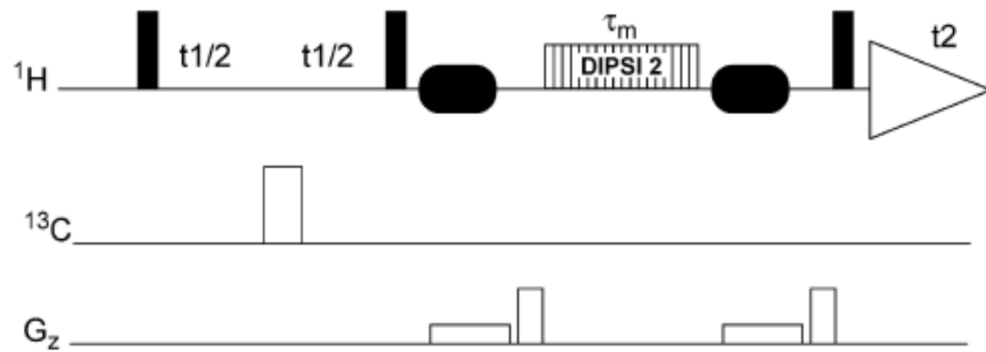
biomass hydrolysate from E. coli cells  
grown on 20% [U-<sup>13</sup>C]-glucose +80% [1-<sup>13</sup>C]-glucose

Measurement of specific isotopic enrichments  
**IMPOSSIBLE**



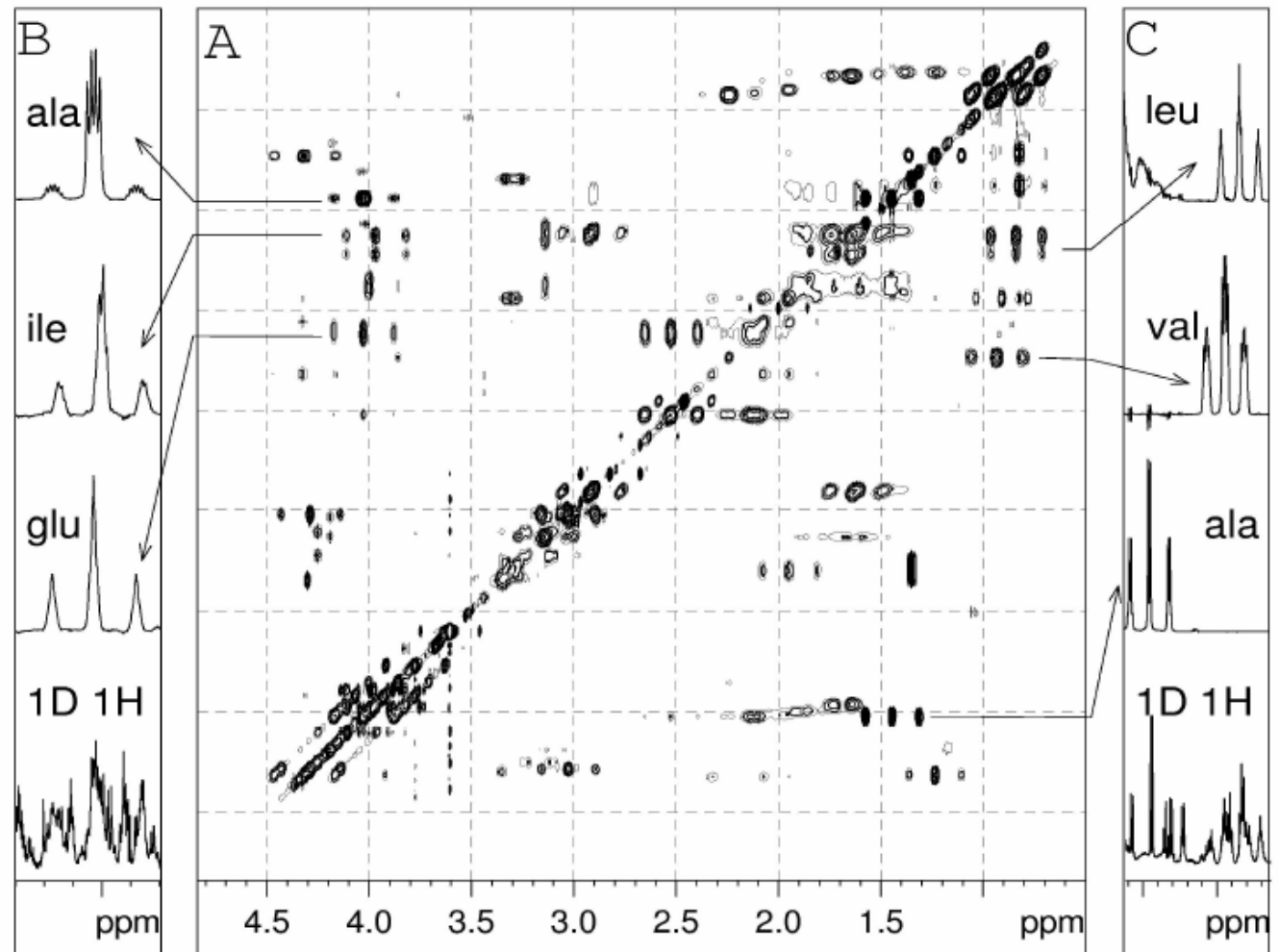
**2D NMR**

# Applications in fluxomics



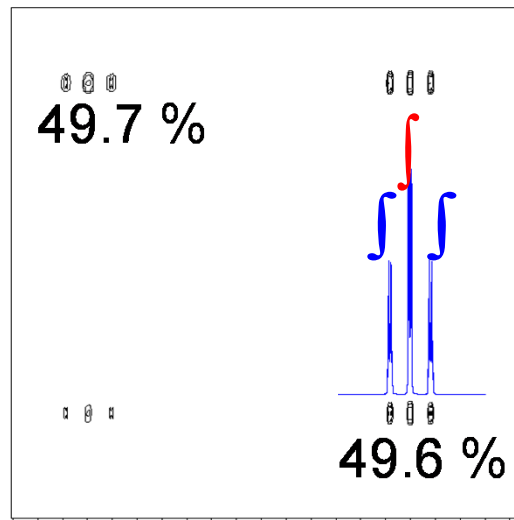
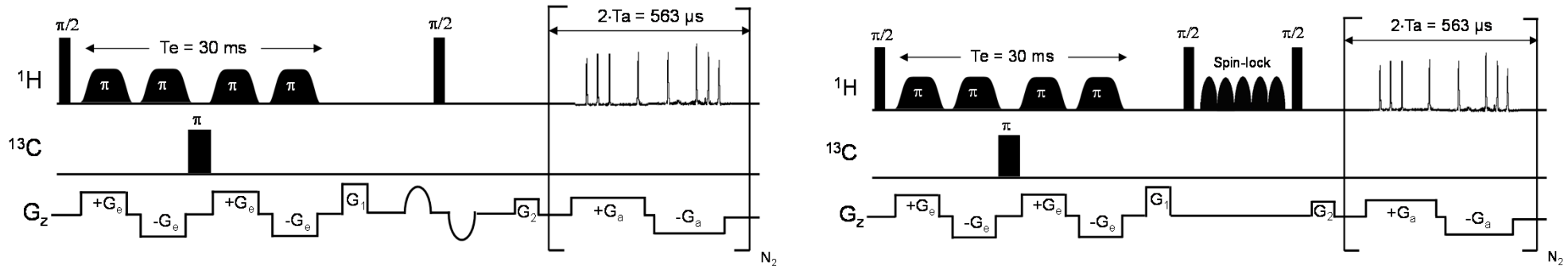
Conv. 2D ZQF TOCSY  
<sup>13</sup>C decoupling in F<sub>1</sub>

Acquisition time:  
5-10h  
in quantitative  
conditions  
( $TR \geq 5 \cdot T_1^{\max}$ )

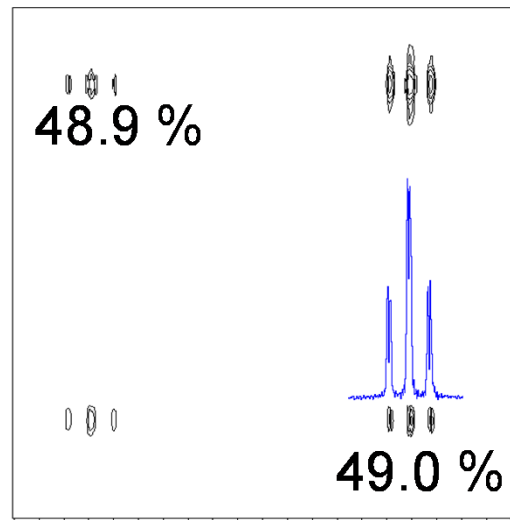


# Applications in fluxomics

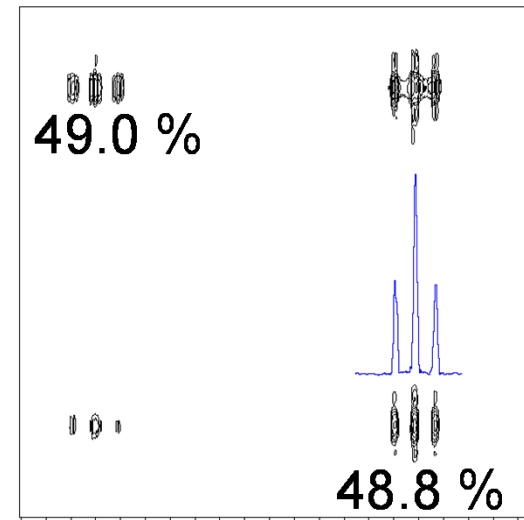
## Ultrafast COSY and zTOCSY



Conv. zTOCSY, 2.5 h



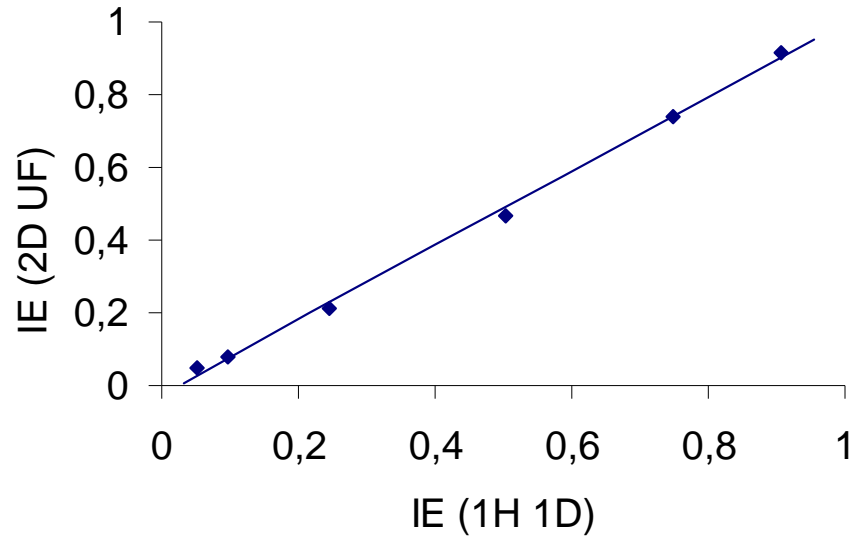
UF zTOCSY, 3 s



UF COSY, 3 s

Mixture of variously labeled alanines, 400 MHz

## Analytical performance - $^{13}\text{C}$ glucose samples

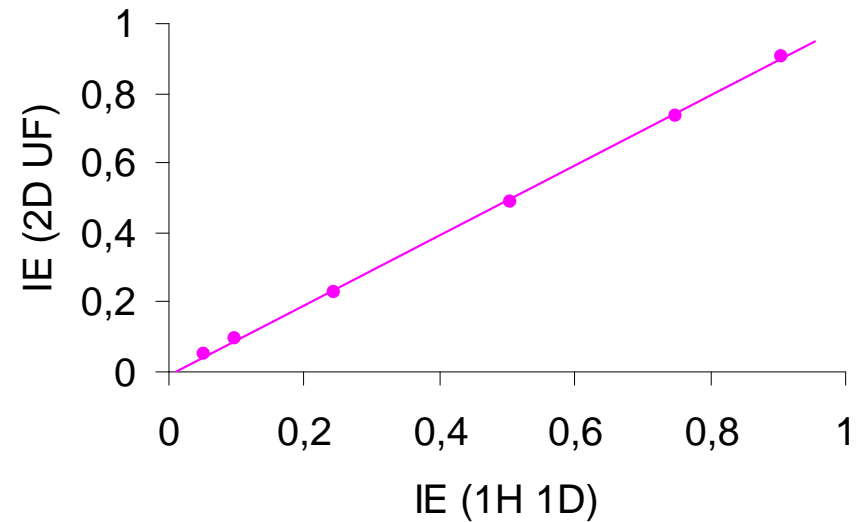


UF TOCSY

$$y = 1.018x + 0.023$$

$$R^2 = 0,998$$

Precision 3.7 %

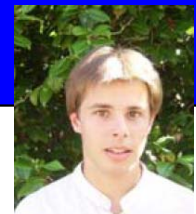


UF COSY

$$y = 1.000x + 0.008$$

$$R^2 = 0,999$$

Precision 3.0 %

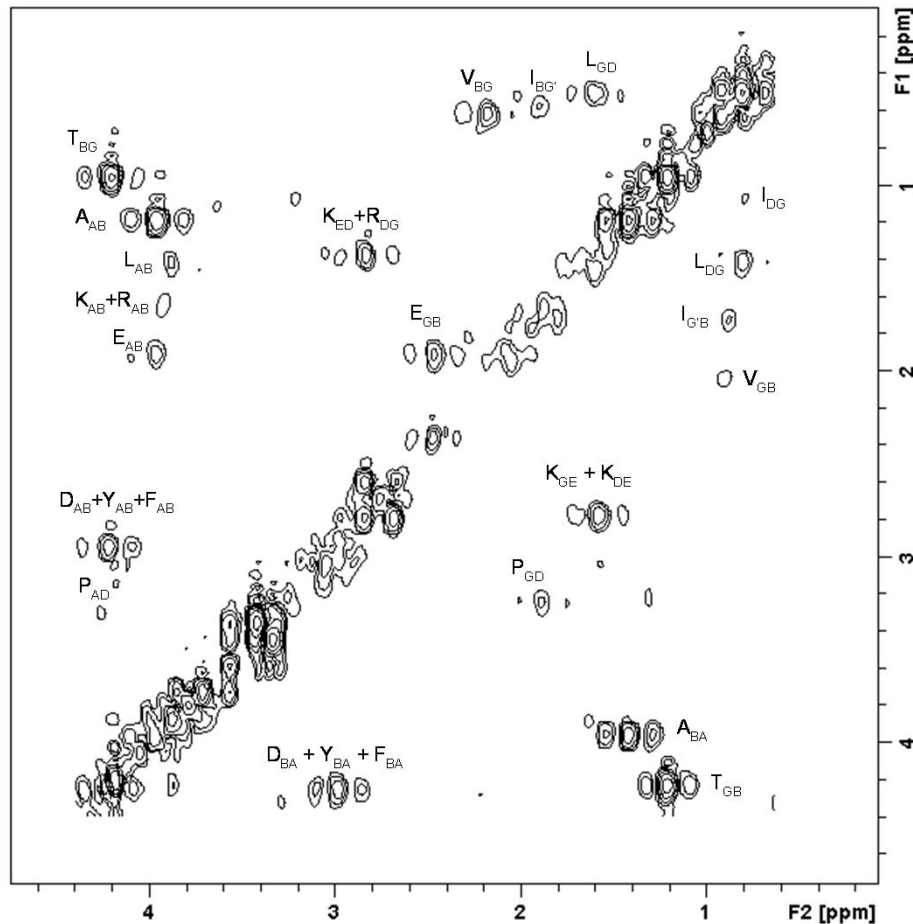


## Application to a biological sample

### COSY UF

### $T_a = 40$ s

Biomass hydrolyzate (*E. coli*)  
Grown on  
50 % [U-13C]-glucose  
and 50 % *n.a.* glucose



Isotopic enrichments measured from 2D spectra

	TOCSY Conv.	TOCSY UF	COSY UF
$A_{AB}$	44.5	45.7	41.2
$A_{BA}$	46.8	51.3	50.7
$E_{AB}$	47.9	44.7	45.2
$L_{AB}$	48.3	43.8	43.4
$L_{DG}$	47.9	42.7	45.1
$P_{AD}$	47.0	43.1	42.0
$T_{BG}$	26.9	23.0	25.8
$T_{GB}$	24.5	23.3	24.7
$V_{GG'}$	47.9	42.3	47.6

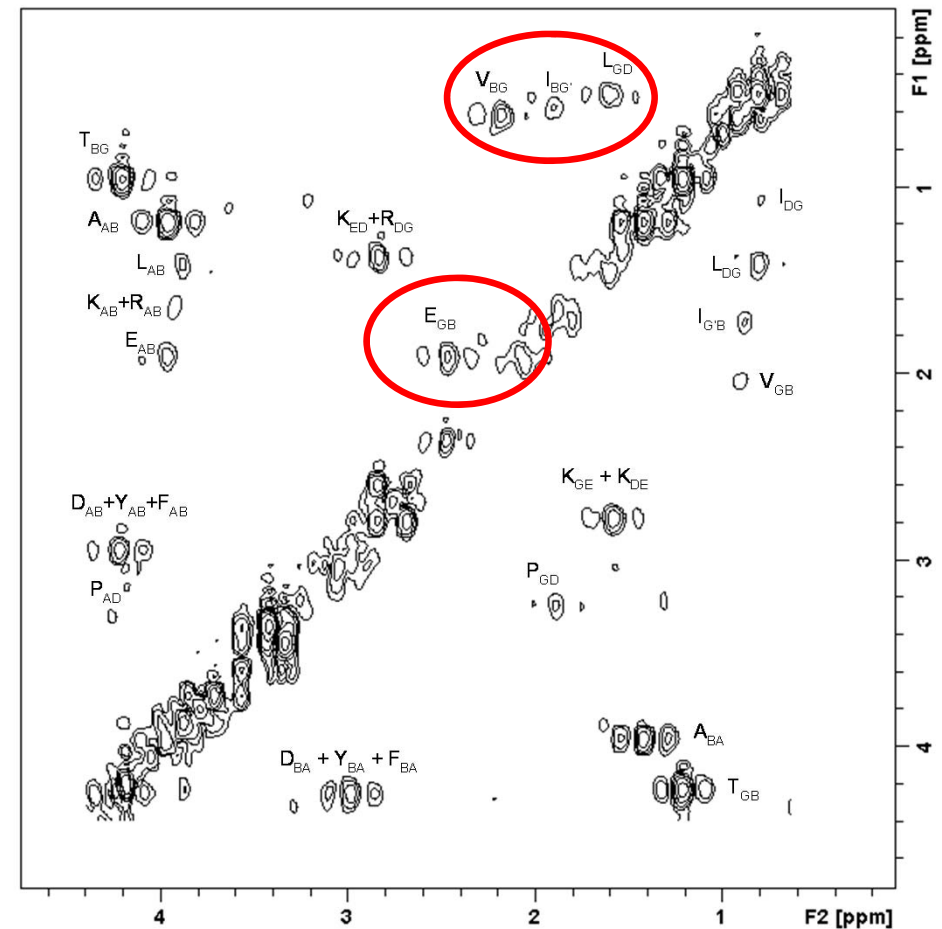
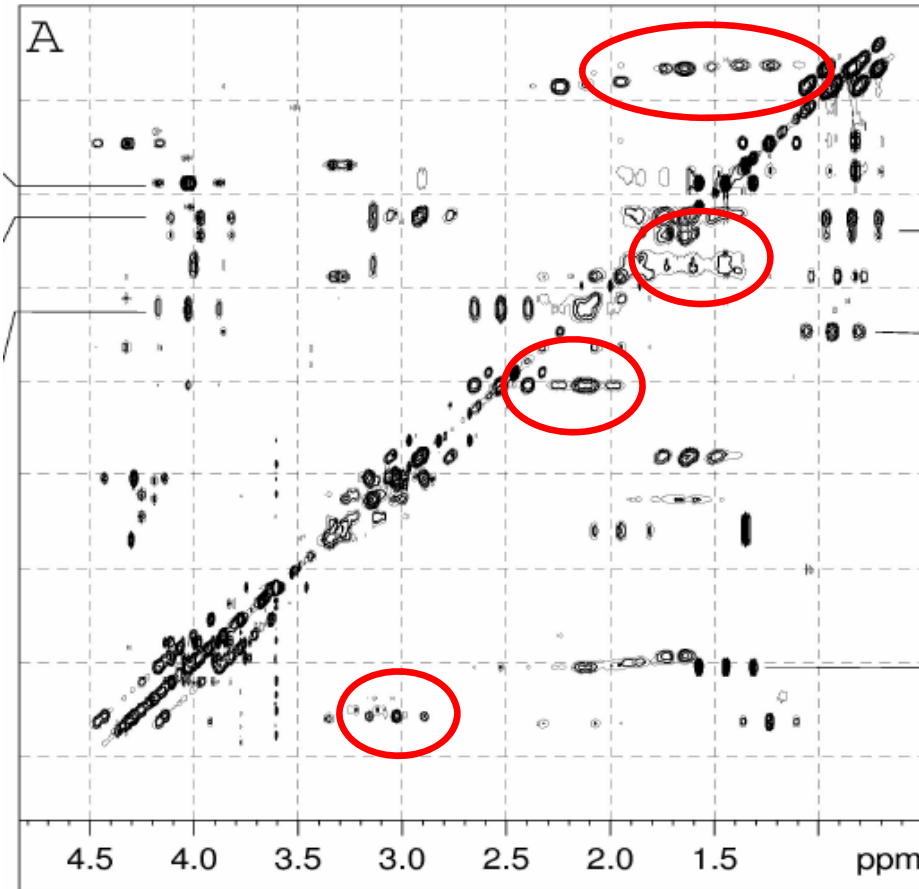


# Applications in fluxomics

## Limitations of 2D NMR

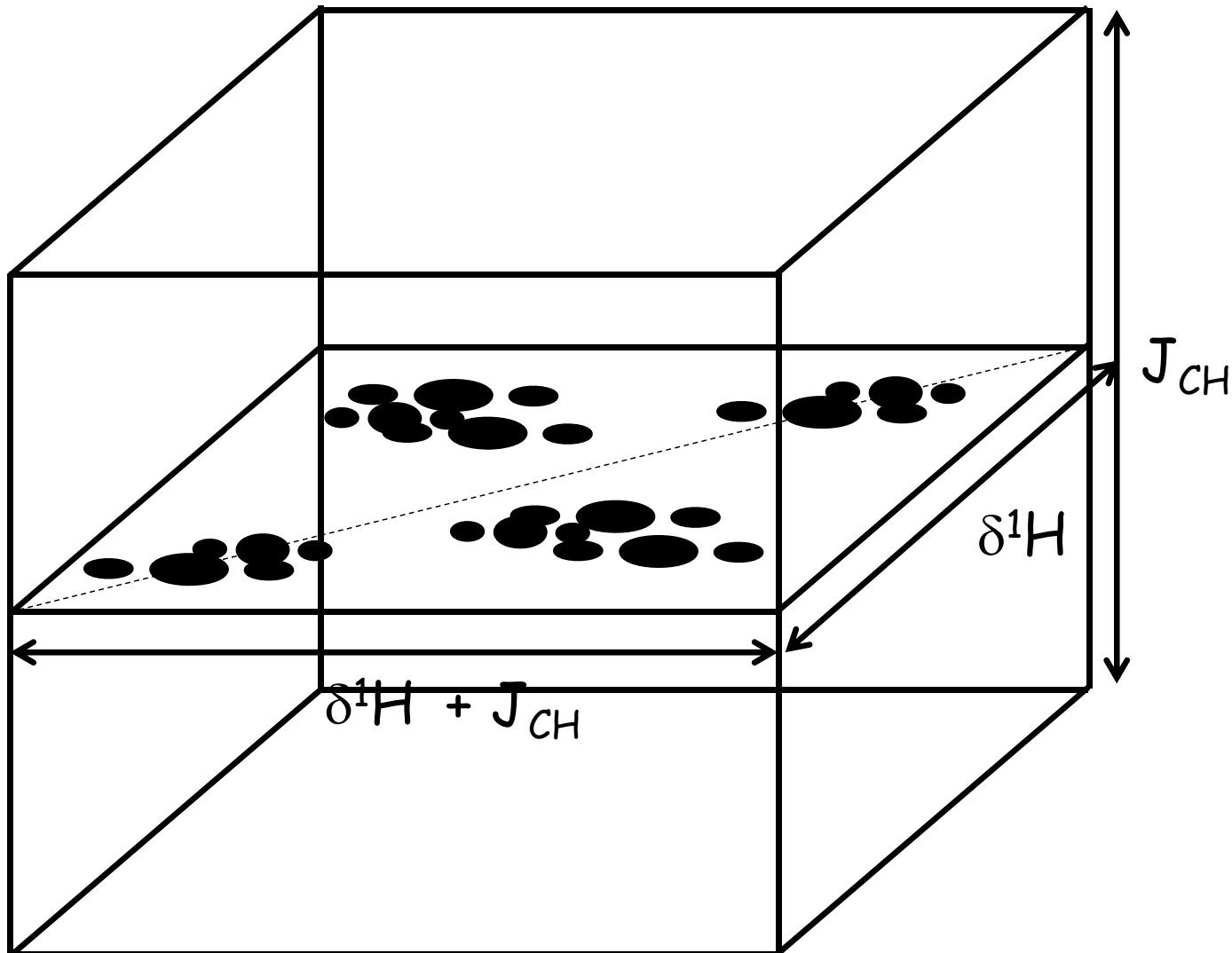
### Conventional 2D

### 2D UF



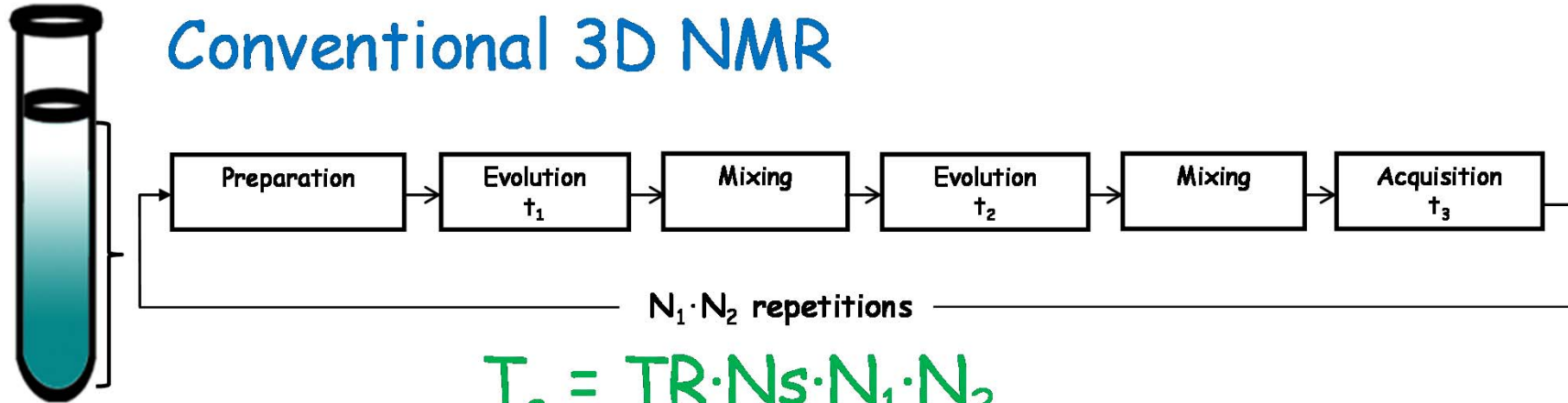
Peak overlaps still prevent accurate quantification

# 3D NMR



## 3D NMR

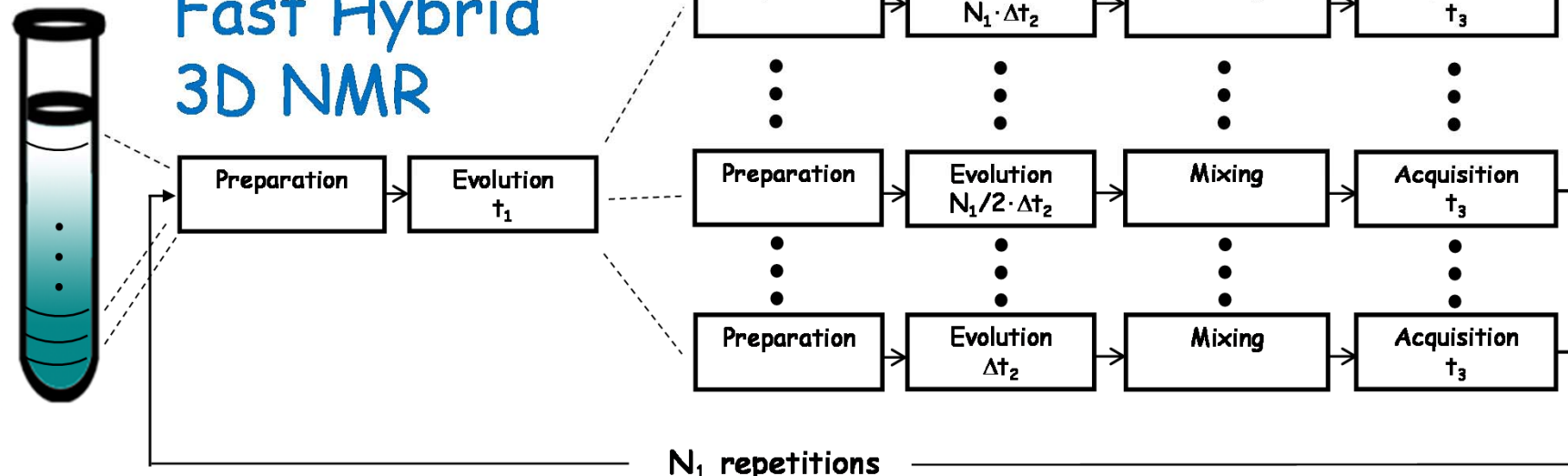
### Conventional 3D NMR



$$T_a = TR \cdot N_s \cdot N_1 \cdot N_2$$

Spatial encoding (1 scan)

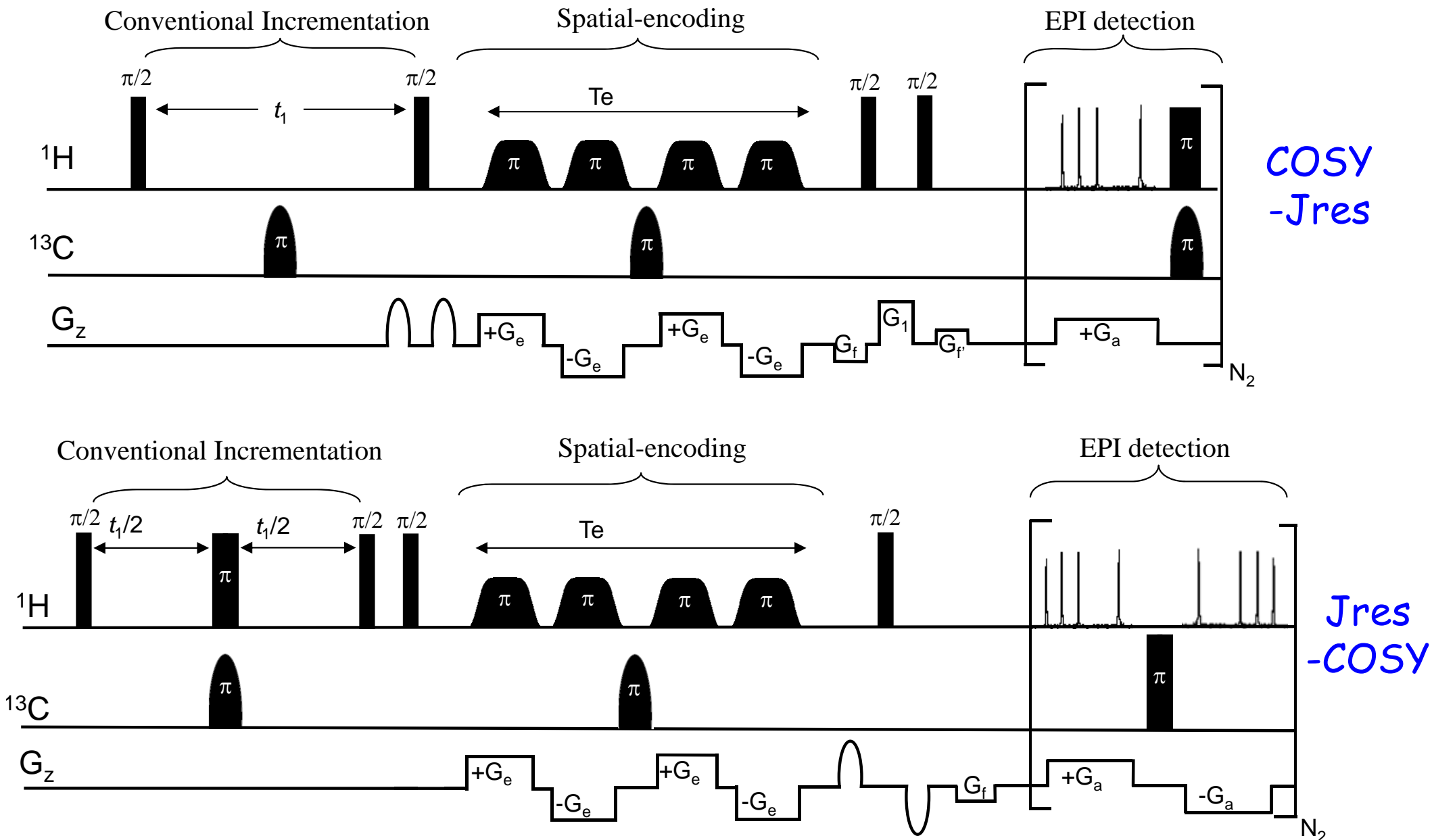
### Fast Hybrid 3D NMR



$$T_a = TR \cdot N_s \cdot N_1$$

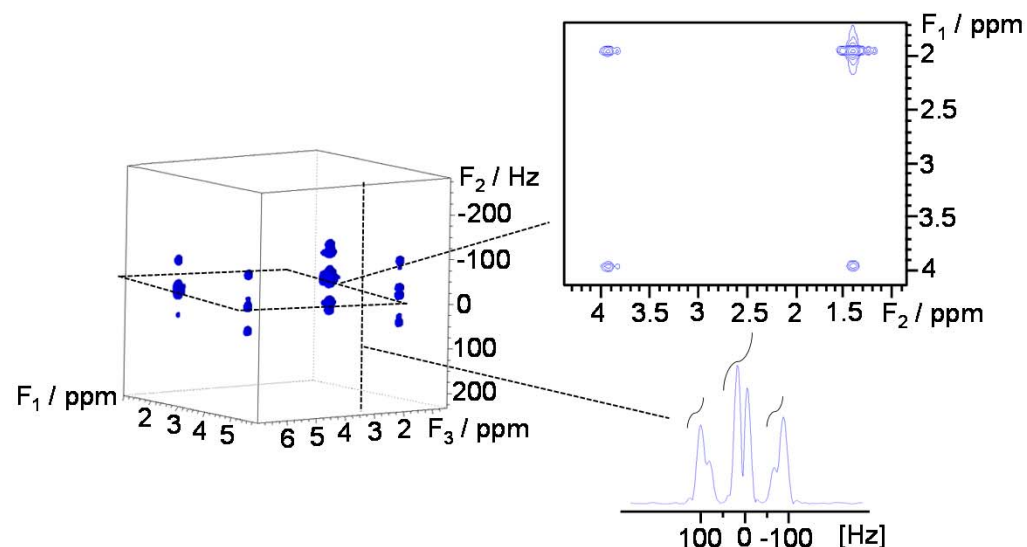
# Applications in fluxomics

## Fast-Hybrid 3D NMR

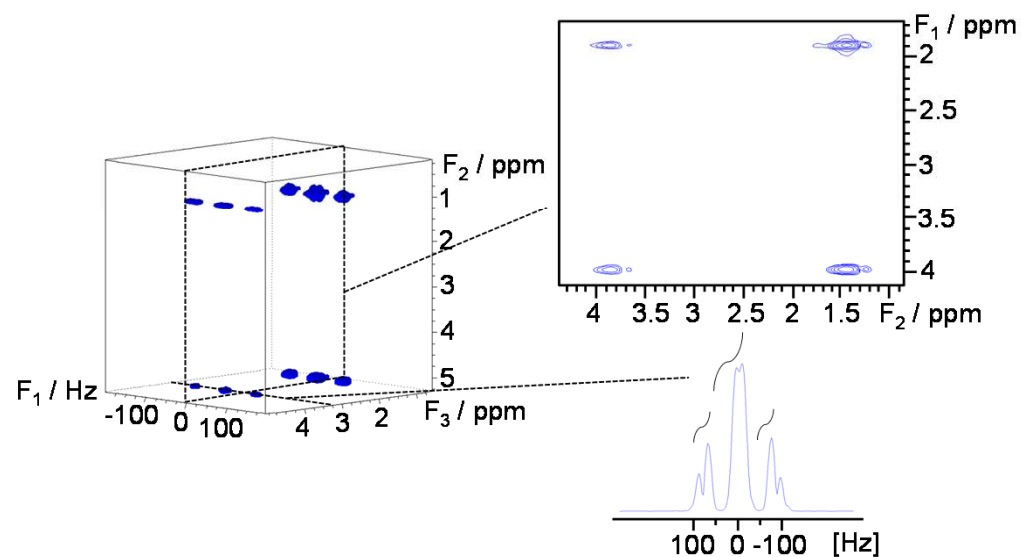


# Applications in fluxomics

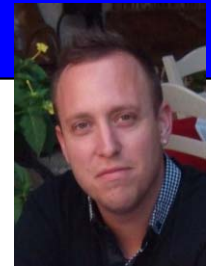
## Fast-Hybrid 3D NMR (12 min, labeled alanine sample)



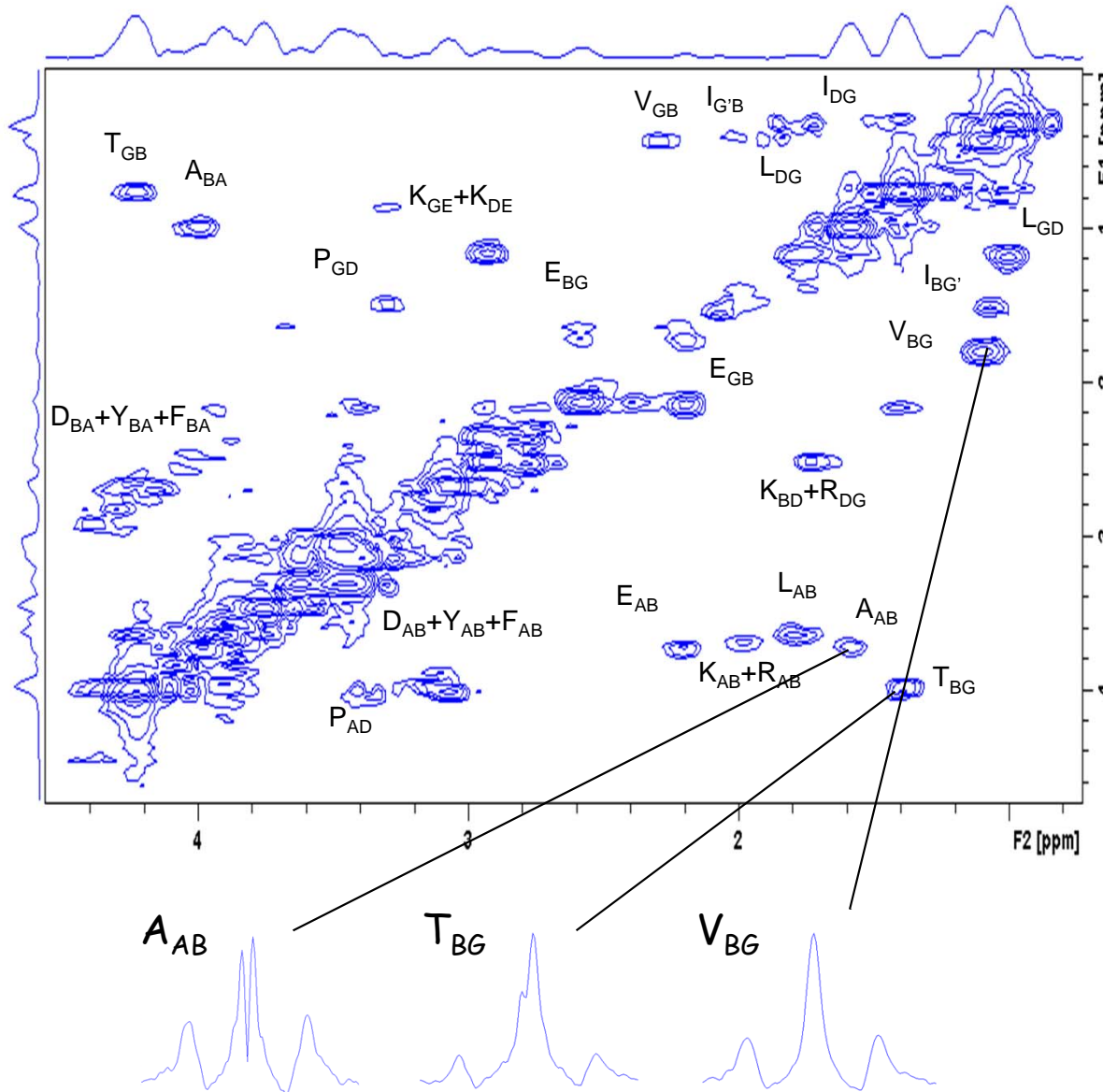
**Jres-COSY**



# Applications in fluxomics



## Fast-Hybrid 3D NMR: application to a biological sample (12 min)

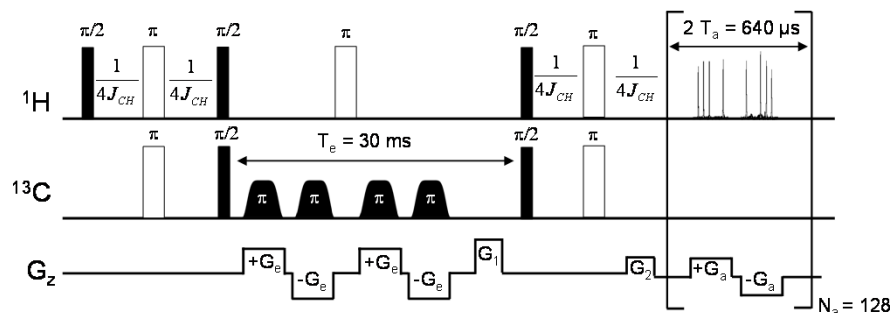


### Site-specific IEs

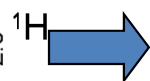
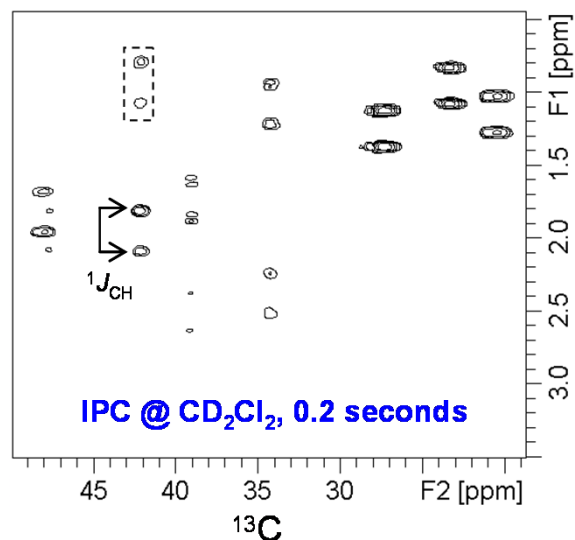
peak	2D UFCOSY	3D COSY-Jres
$A_{AB}$	41,2	44,7
$A_{BA}$	50,7	47,1
$E_{AB}$	45,2	47,9
$E_{BG}$	-	48,6
$E_{GB}$	37,6	49,9
$I_{G'B}$	44,8	46,9
$L_{AB}$	43,4	48,5
$L_{DG}$	45,1	47,9
$P_{AD}$	42	47,2
$T_{BG}$	25,8	27,1
$T_{GB}$	24,7	24,9
$V_{BG}$	47,6	48,0
$V_{GB}$	-	46,8
$P_{GD}$	-	41,4



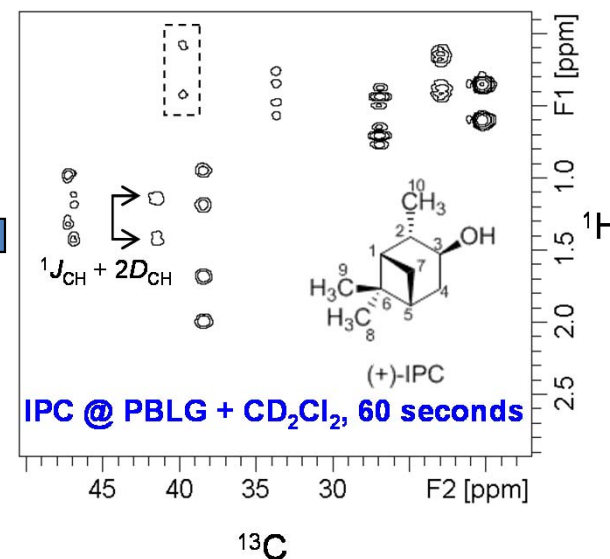
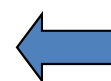
## Measurement of Residual Dipolar Couplings (RDCs) in oriented media



F<sub>2</sub>-coupled UF HSQC



Residual Dipolar Couplings (RDCs)

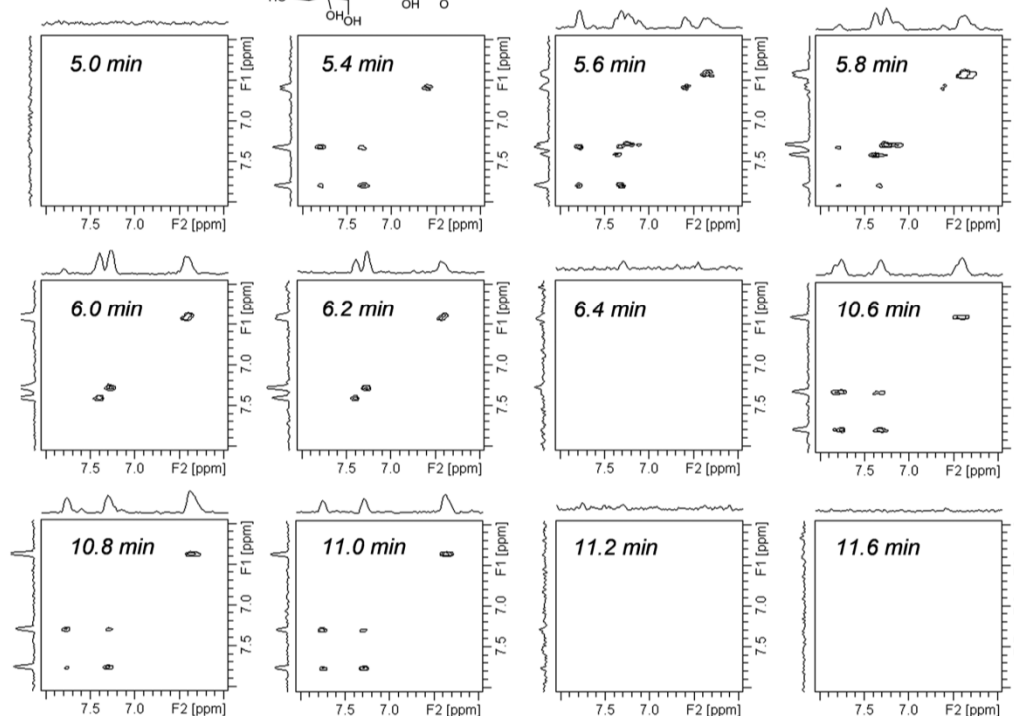
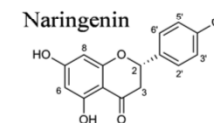
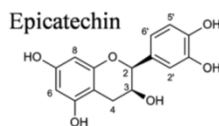
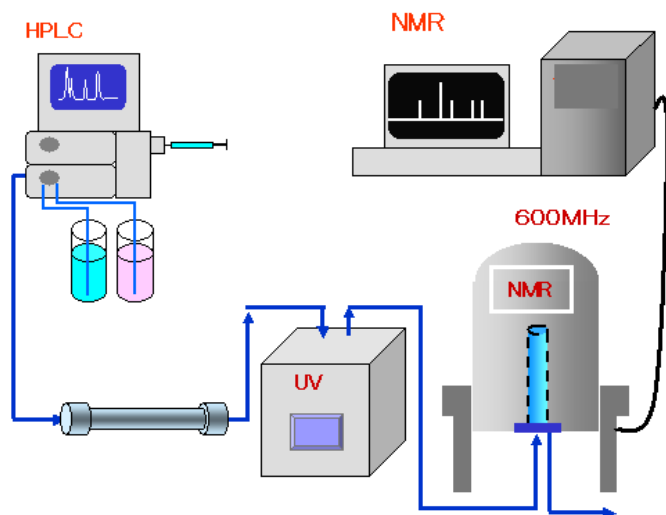


Faster structural analysis



## Coupling with on-line HPLC

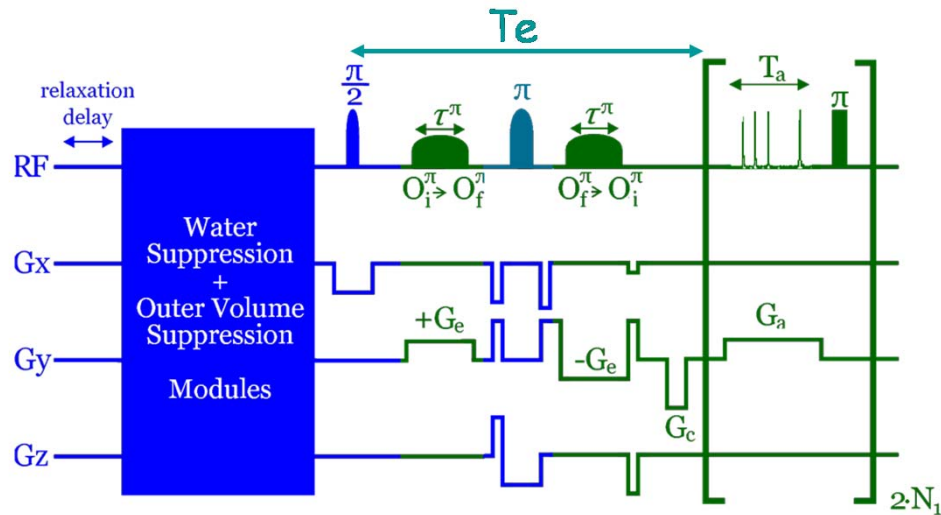
### Real-time separation of natural products





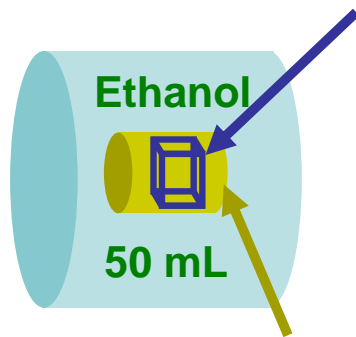


## Towards ultrafast *in vivo* spectroscopy



PRESS + UF Jres

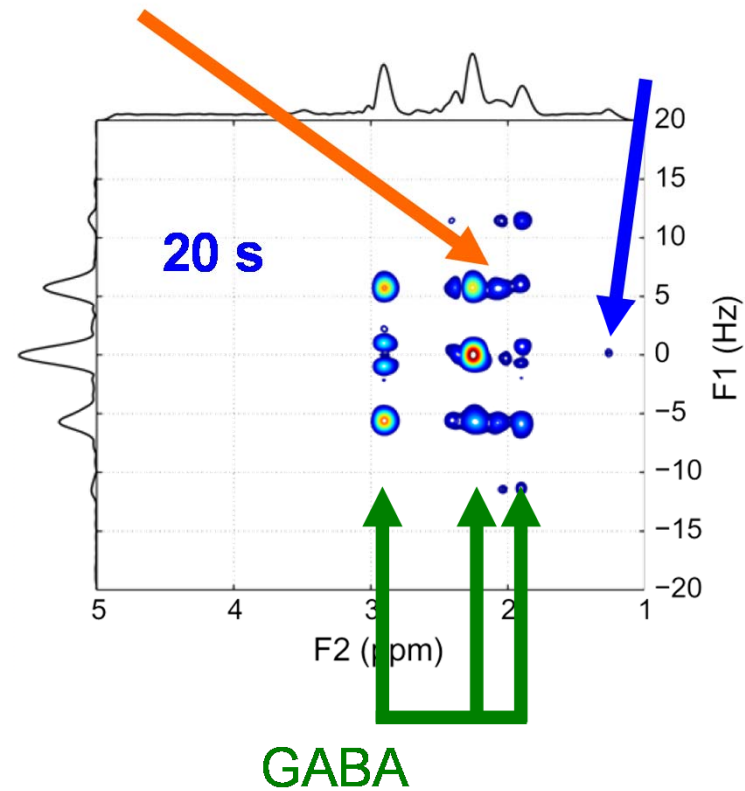
Voxel 5mm.5mm.5mm



GABA 10% @ H<sub>2</sub>O  
1.5 mL

Strong coupling artefacts

Residual signal from outer volume



# Conclusion & Perspectives

- Improved analytical performance of ultrafast experiments
- Immunity to spectrometer temporal instabilities
- High potential for quantitative analysis
- Application to samples of increasing complexity
- Next steps : *in vivo* and hyperpolarization

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