

Comparison of New 2D High Resolution Experiments for Quadrupolar Nuclei Based on STMAS Concept

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INTRODUCTION

STMAS is a technique introduced in 2000 [1] using satellite transitions (usually inner ST₁), instead of Multiple Quanta coherences (3Q/5Q). Gan found in 2003 [2] a way to eliminate the CT-CT unwanted signal by using a double quantum filter (STMAS-DQF). He also expressed the concept of DQ-STMAS, where DQ coherences (instead of ST₁) are correlated with CT.

We present experimental proofs of the DQ-STMAS concept and compare its performance (sensitivity, resolution etc...) to STMAS-DQF for several reference compounds such as RbNO₃, AlPO₄ berlinite and VPI-5.

We also introduce an original t₁-split STMAS sequence specific to 3/2 spins. Instead of splitting the t₁ evolution time between MQ and CT, we split t₁ between ST₁ and DQ coherences. t₁-split STMAS performs as well as DQ-STMAS and STMAS-DQF, concerning homonuclear broadening [3] or efficiency.

We finally show how these experiments can be combined with the SPAM technique and the sensitivity gain that can be expected.

[1] Z.H. Gan, *J. Am. Chem. Soc.* 122 (2000) 3242-3243.

[2] H.T. Kwak, Z.H. Gan, *J. Magn. Res.* 164 (2003) 369-372.

[3] J.P. Amoureux, J. Trébosc, *J. Magn. Res.* 179 (2006) 311-316.

RESOLUTION COMPARISON

In addition to distributions (inhomogeneous broadening), in STMAS isotropic projections are affected by several parameters

- magic angle setting and spinning-speed stability,
- relaxation homogeneous effects in t₁ (T_p) and t₂ (T_{CT}),
- R ratio and p quantum level used during t₁

The magic angle has the same effect on all STMAS versions.

We have observed that T_{DQ} and T_{ST1} are often very similar.

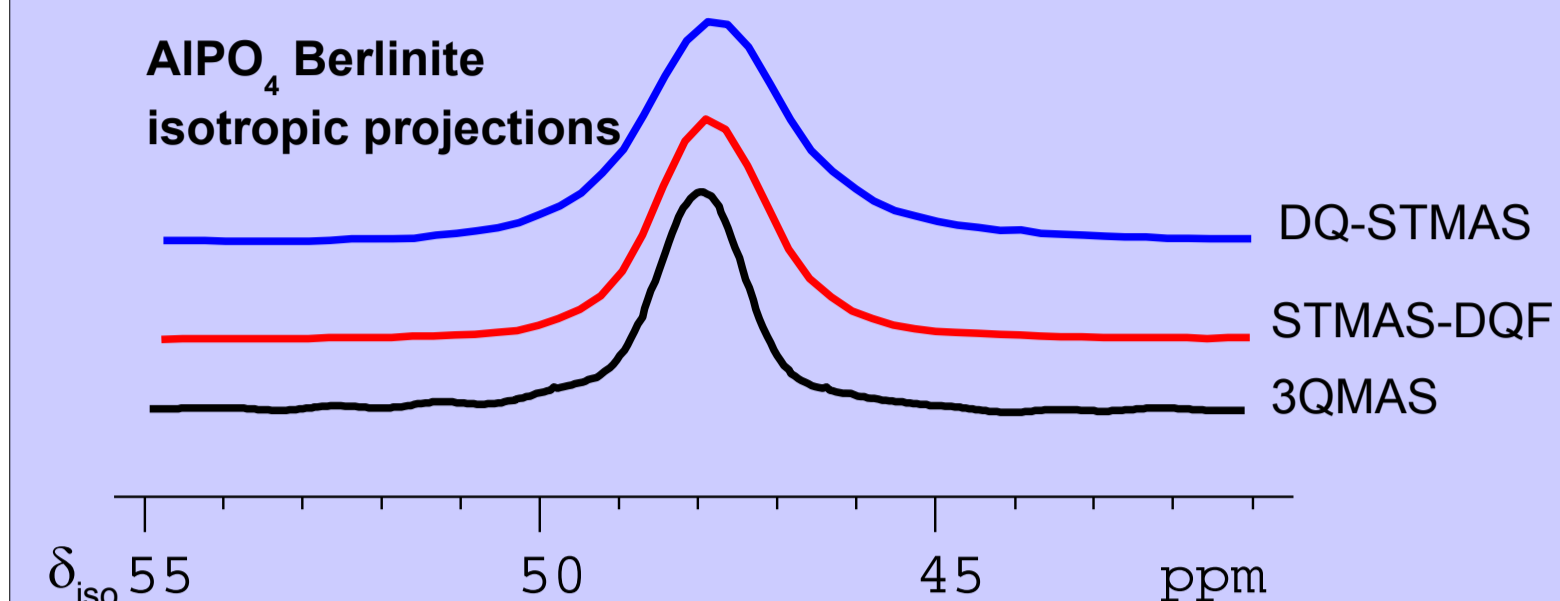
Homogeneous broadening (B) in isotropic dimension due to relaxation can be calculated with :

$$B = \frac{(1+R) \cdot 10^6}{\pi \nu_0 |R-p| T_{tot}} \quad \frac{1}{T_{tot}} = \frac{R}{T_{CT}} + \frac{1}{T_p}$$

S	1D-MAS	3QMAS	5QMAS	STMAS-DQF	DQ-STMAS	t ₁ -split-STMAS
3/2	1	8/17		17/17	10/17	9/17
5/2	1	31/17	37/85	31/17	55/17	
7/2	1	73/17	50/85	73/17	118/17	
9/2	1	127/17	131/85	127/17	199/17	

Broadening (B), given in 10⁶/πν₀T unit, calculated with T_p = T_{CT}

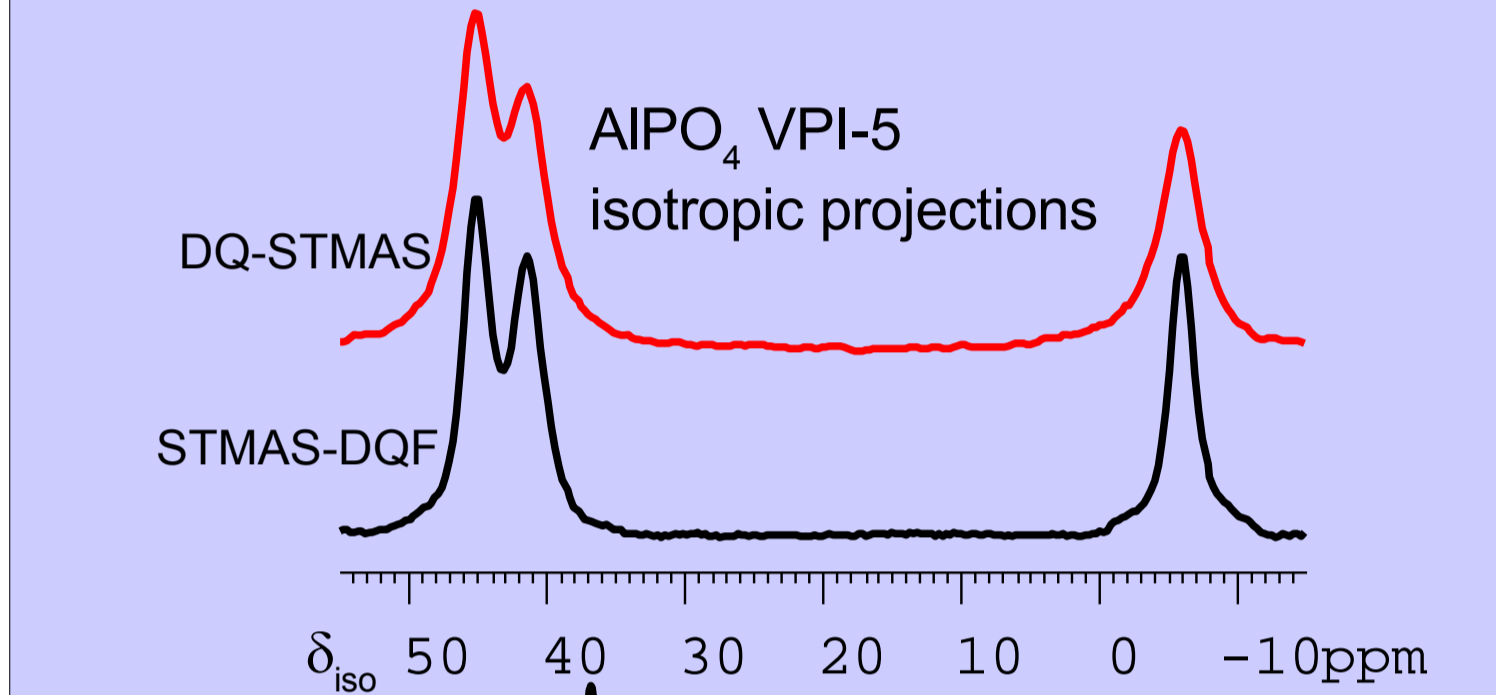
SPIN 5/2 (²⁷Al)



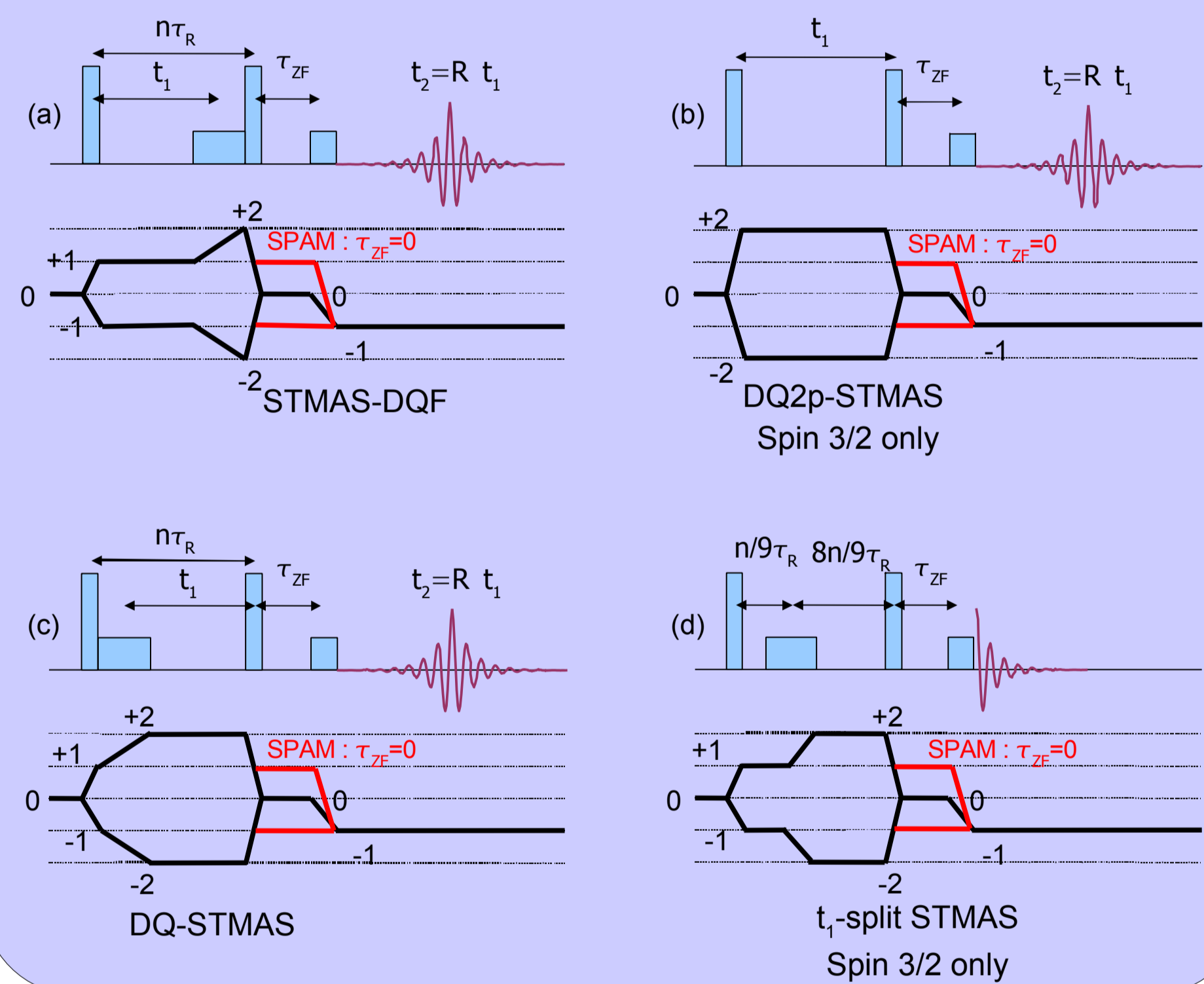
3QMAS is better resolved than STMAS-DQF due to its robustness towards magic angle and spinning speed stability, but also because ST₁ and DQ coherences have smaller T₂ than CT or 3Q coherences.

On berlinite or VPI-5, despite similar T₂ for ST₁ and DQ coherences, DQ-STMAS has unfavorable B factor (see table).

T₂ Measurements on Berlinite (see poster P 189)

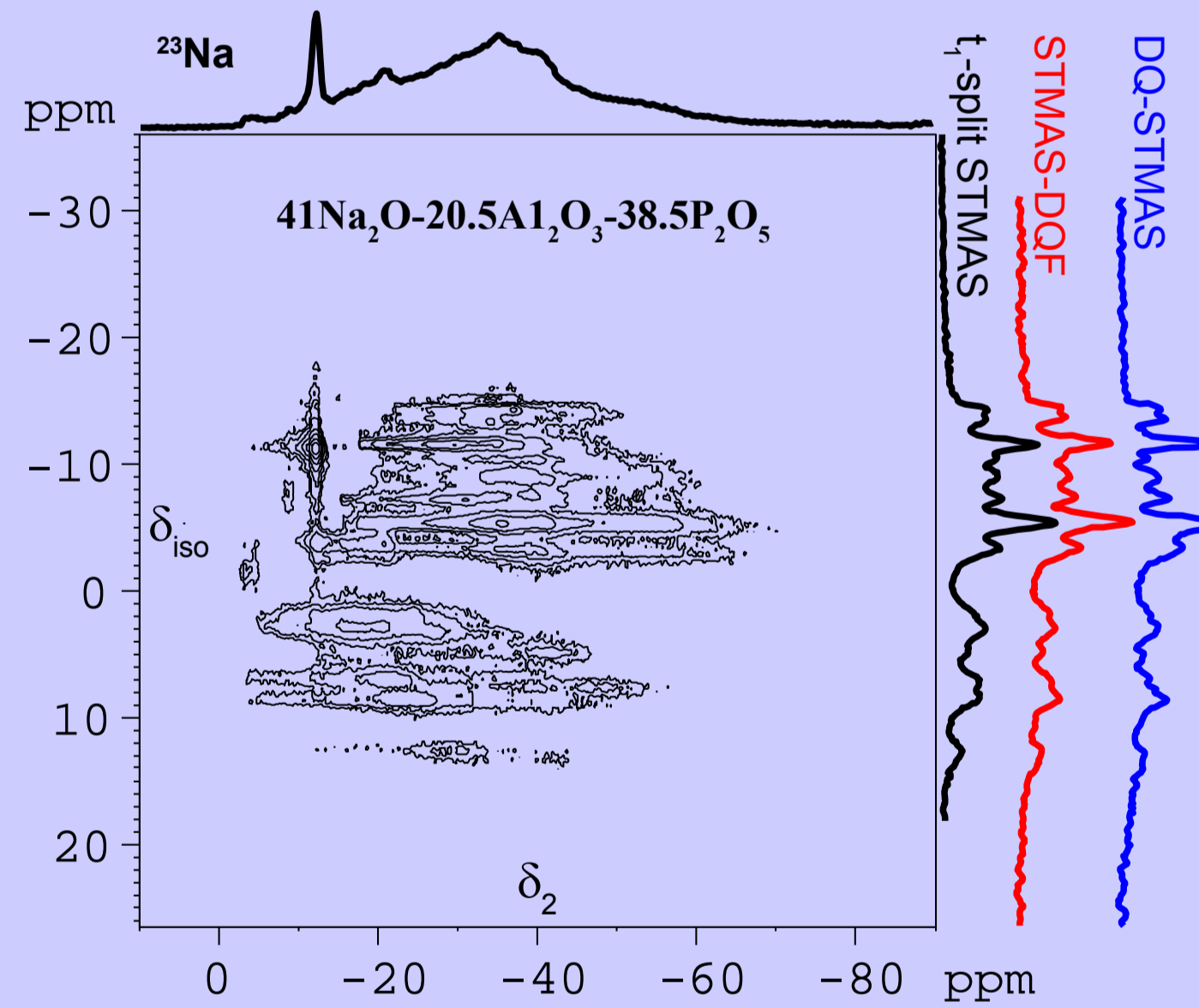
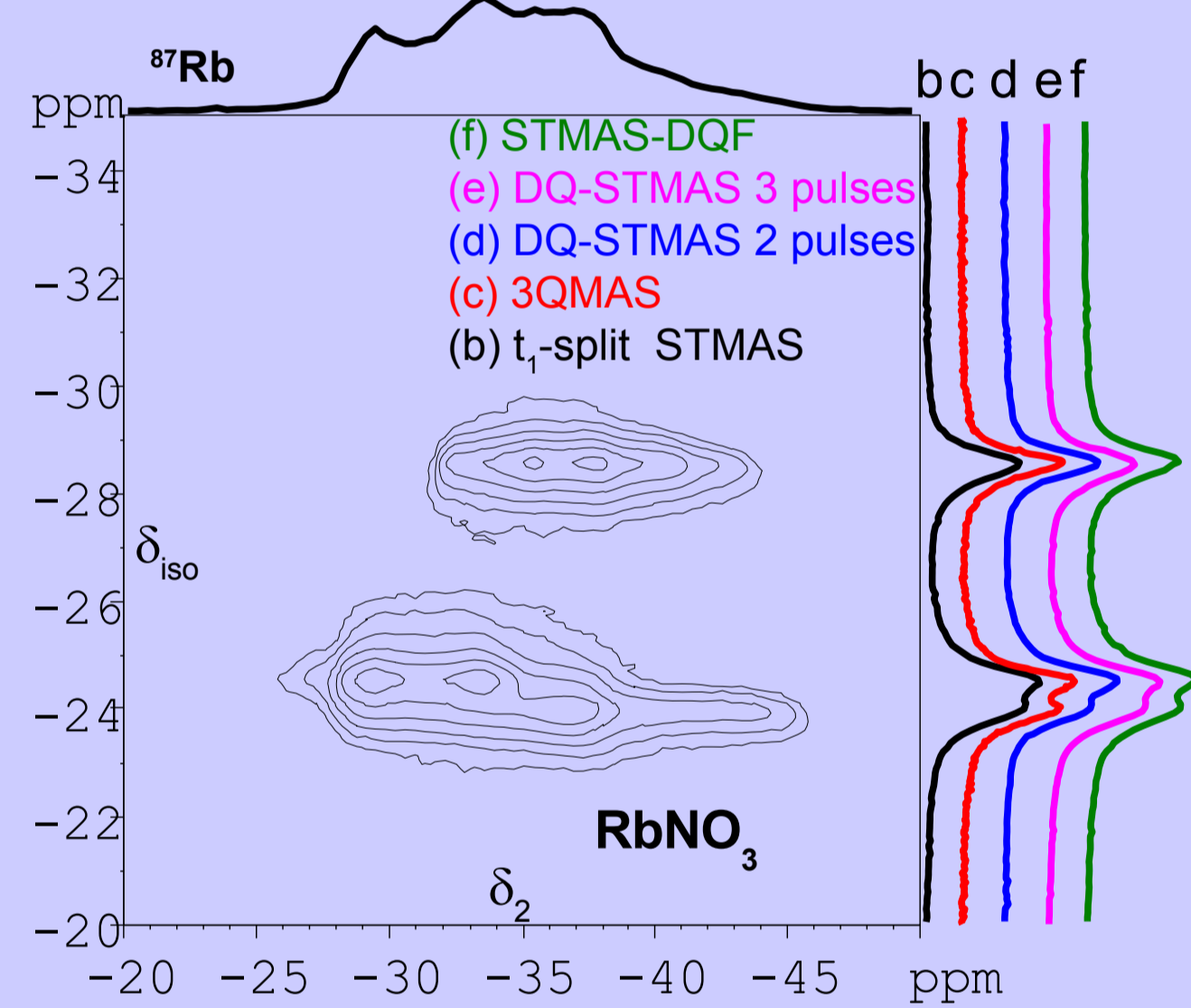


PULSE SEQUENCES



SPIN 3/2

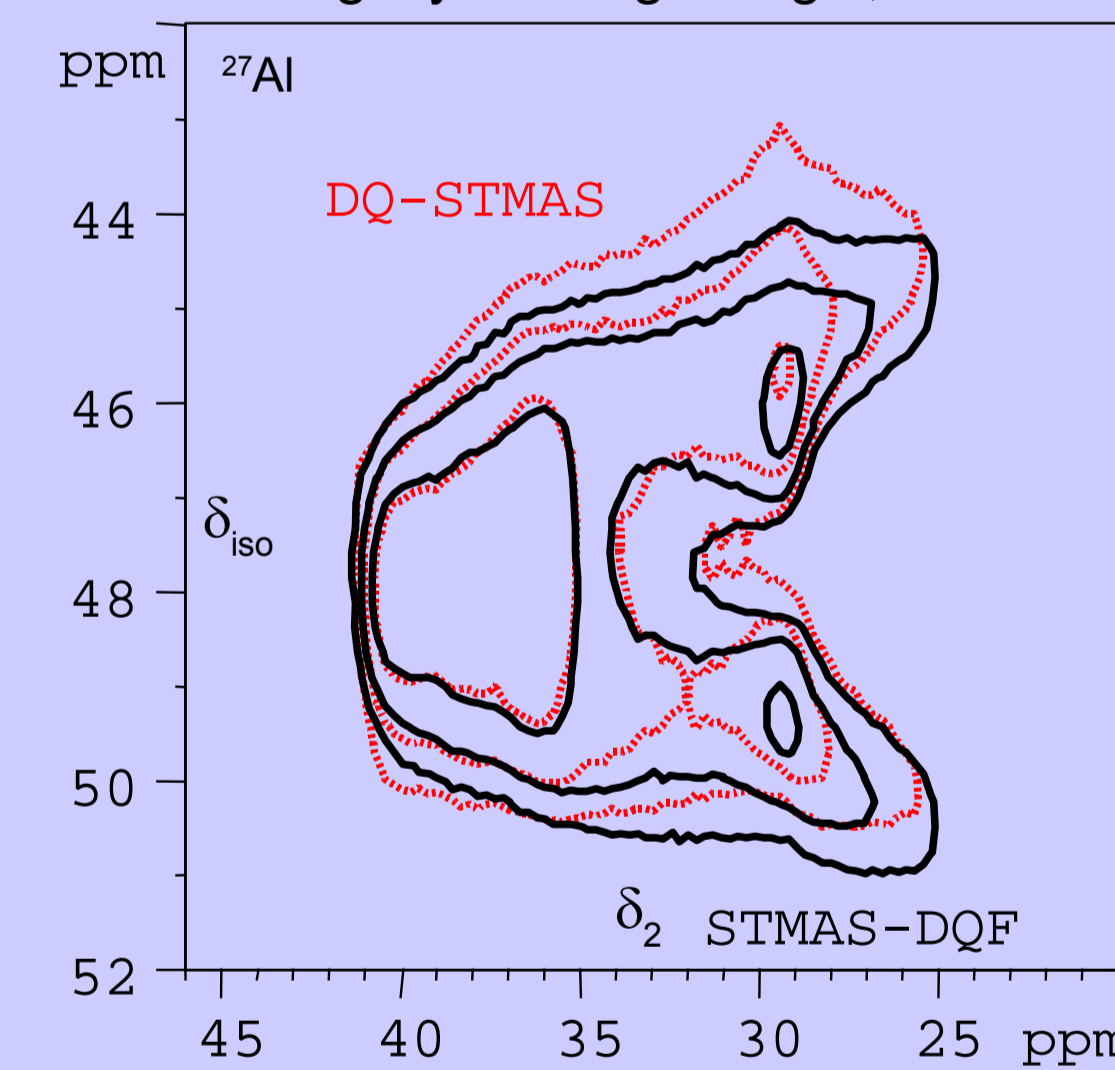
In RbNO₃ all relation times are very large. Thus all experiments have the same resolution



This recrystallised glass also has large T_{2s} : no difference in resolution is noticed here

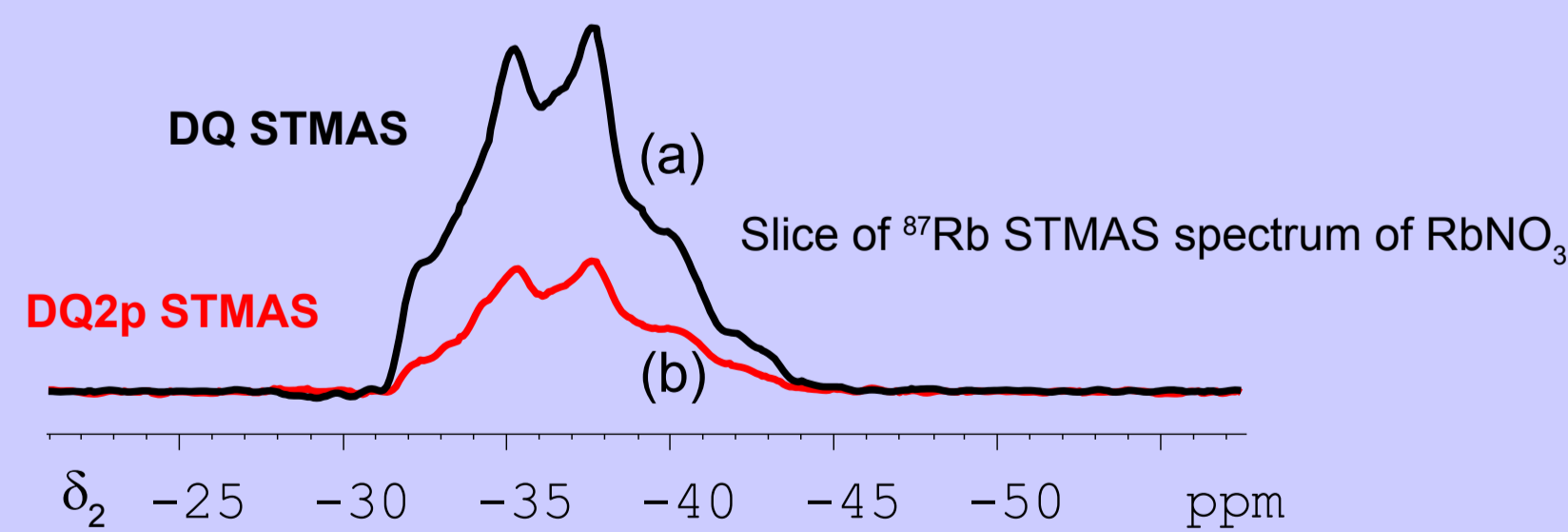
MAGIC ANGLE SETTING

DQ-STMAS or STMAS-DQF experience the same splitting due to 1st order quadrupolar interaction: slightly off magic-angle,

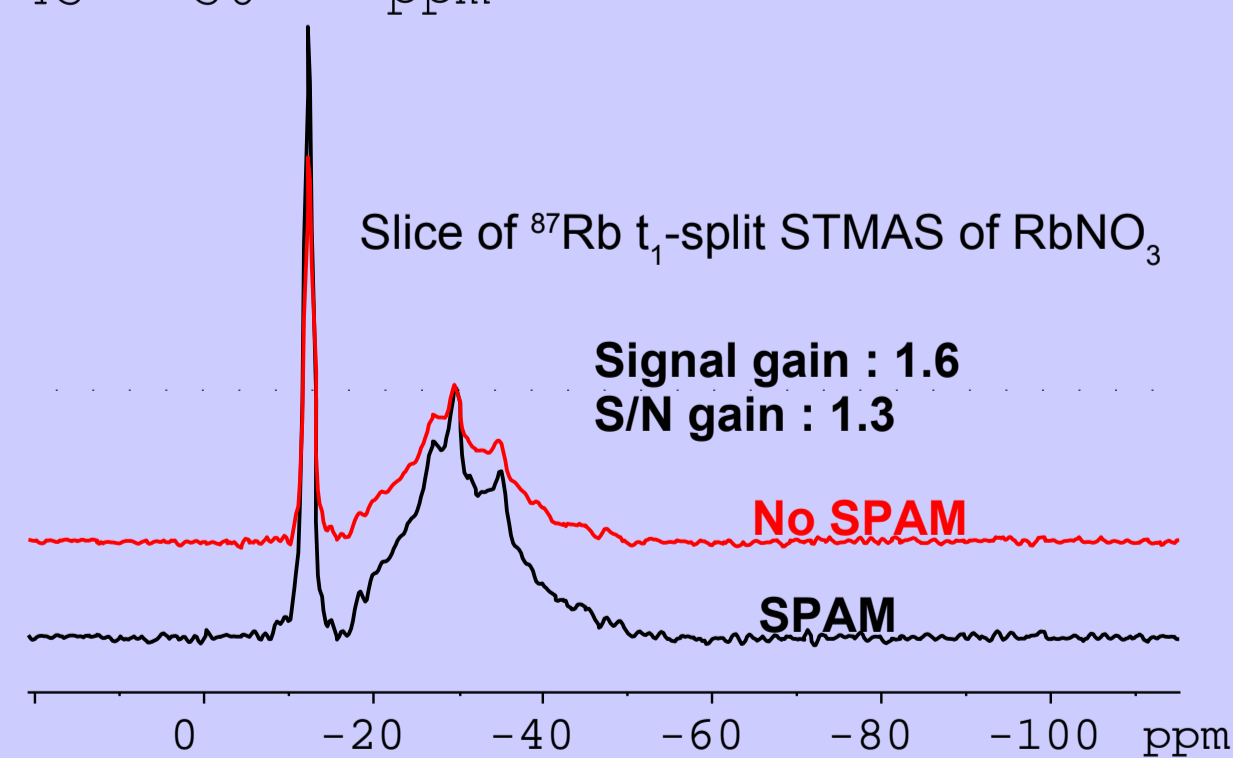


EFFICIENCY

Since they flow the same pathway, all STMAS pulse sequences, but (b), have the same efficiency. Direct excitation of DQ coherences proves to be twice less efficient than the indirect way (note: this is not true for the reconversion pulse).



The SPAM technique can be applied to all STMAS sequences and produce a S/N gain of at least 1.3. This gain depends on the number of anti-echo rows that are acquired,



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CONCLUSION

DQ-STMAS and STMAS-DQF have the same efficiency and constraints related to observation of satellite transition and 1st-order quadrupole interaction.

S > 3/2: due to homogeneous broadening, STMAS-DQF achieves better resolution than DQ-STMAS.

S = 3/2: DQ-STMAS has a better resolution than STMAS-DQF. In that case, t₁-split STMAS can also be a very good option (for Q-CPMG) as it balances completely echo and anti-echo in time, thus leading to very clean spectra.

SPAM concept can be applied to all these pulse sequences with a S/N gain 1.3 to 2.5, depending on the number of anti-echo registered,

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