

Emerging Trends in Solution NMR

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Limitations of solution NMR

Molecular size (< 100 kDa?)

Larger membrane proteins

Need deuteration for sharper lines

Express in *E.coli* with ^{15}N , ^{13}C , ^2H

$^2\text{H} \rightarrow ^1\text{H}$ amide back exchange

Can ^{15}N detection help?

^{15}N detection is intrinsically much less sensitive than other nuclei
Are we stupid?

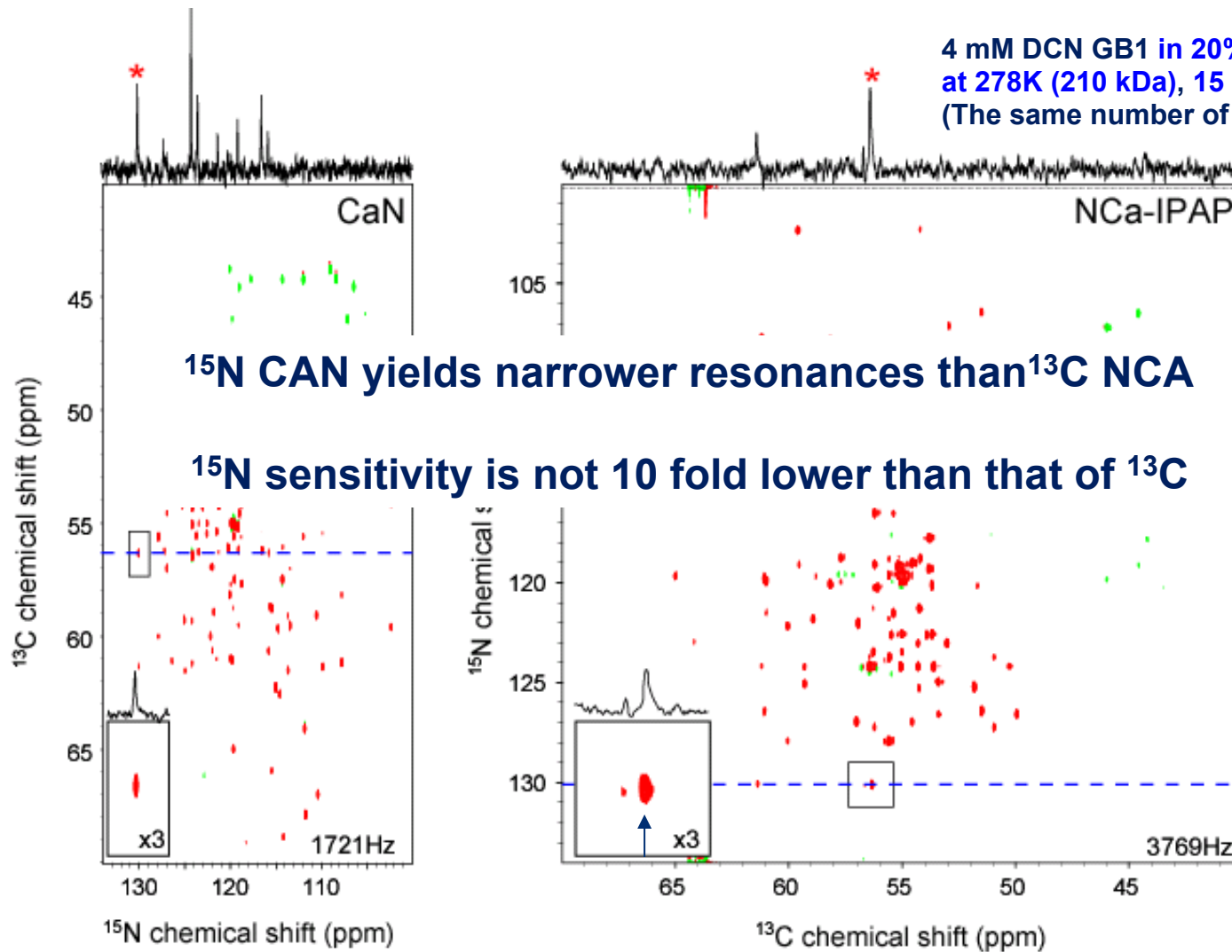
Intrinsic sensitivity (S_0) $\propto \gamma_e \cdot \gamma_d^{3/2} \cdot B_0^{3/2}$

Intrinsic sensitivity of ^{15}N against ^1H : 0.0032

Intrinsic sensitivity of ^{15}N against ^{13}C : 0.10

^{15}N CaN vs ^{13}C NCa

4 mM DCN GB1 in 20% glycerol/ D_2O
at 278K (210 kDa), 15 hrs @500 MHz
(The same number of indirect points)



Why ^{15}N instead of ^1H detection

^{15}N has a much smaller gyromagnetic ratio and is of course much less sensitive ($I = \text{intensity} \sim A = \text{area}$)

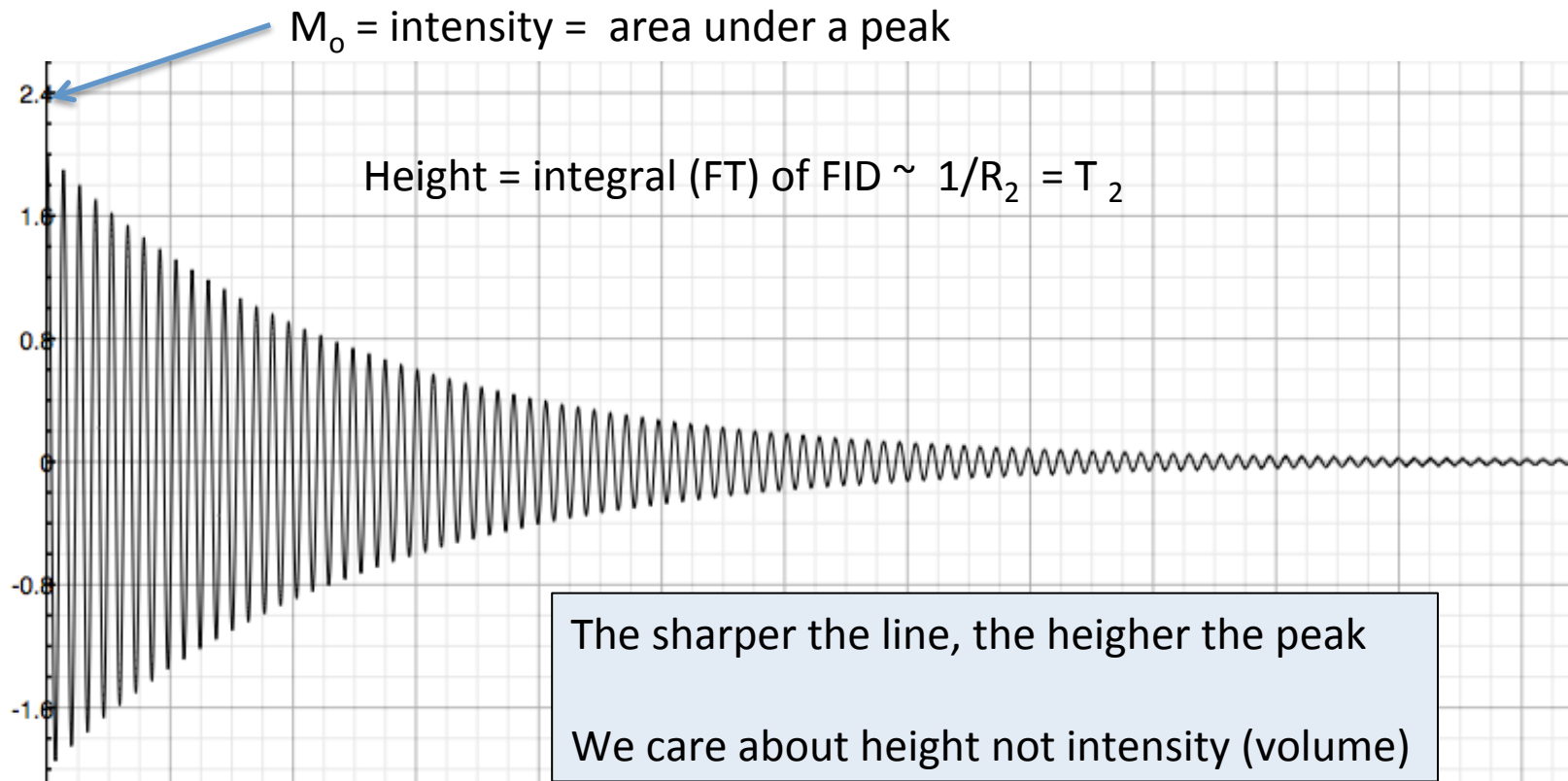
$$I \propto A \propto \gamma_e \cdot \gamma_d^{3/2} \cdot B_0^{3/2}$$

$$H \propto \gamma_e \cdot \gamma_d^{3/2} \cdot B_0^{3/2} / R_2$$

Peak height is proportional to integral (FT) over FID and is proportional to inverse of relaxation rate R_2

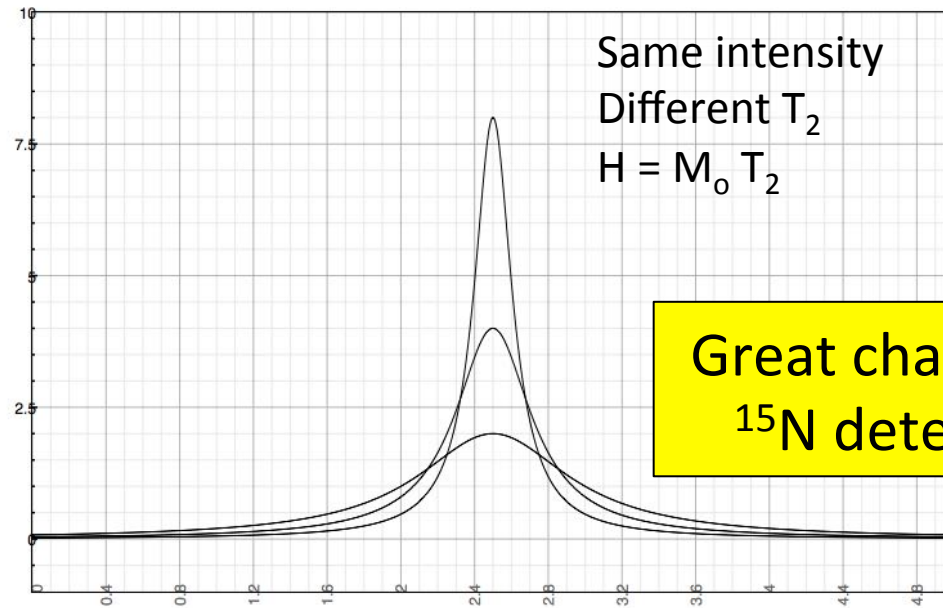
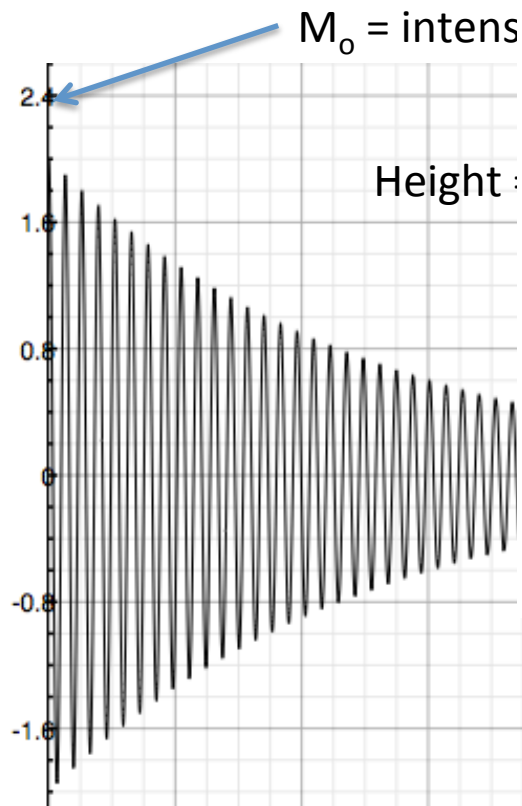
Intensity and Peak height

$$S_c^c(\omega) = \int_0^{\infty} M_0^A \cos\omega_A t \cos\omega t e^{-t/T_2^A} dt = M_0^A \frac{T_2^A}{1 + 4(T_2^A)^2 (\omega - \omega_A)^2}$$



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Great chance for ^{15}N detection

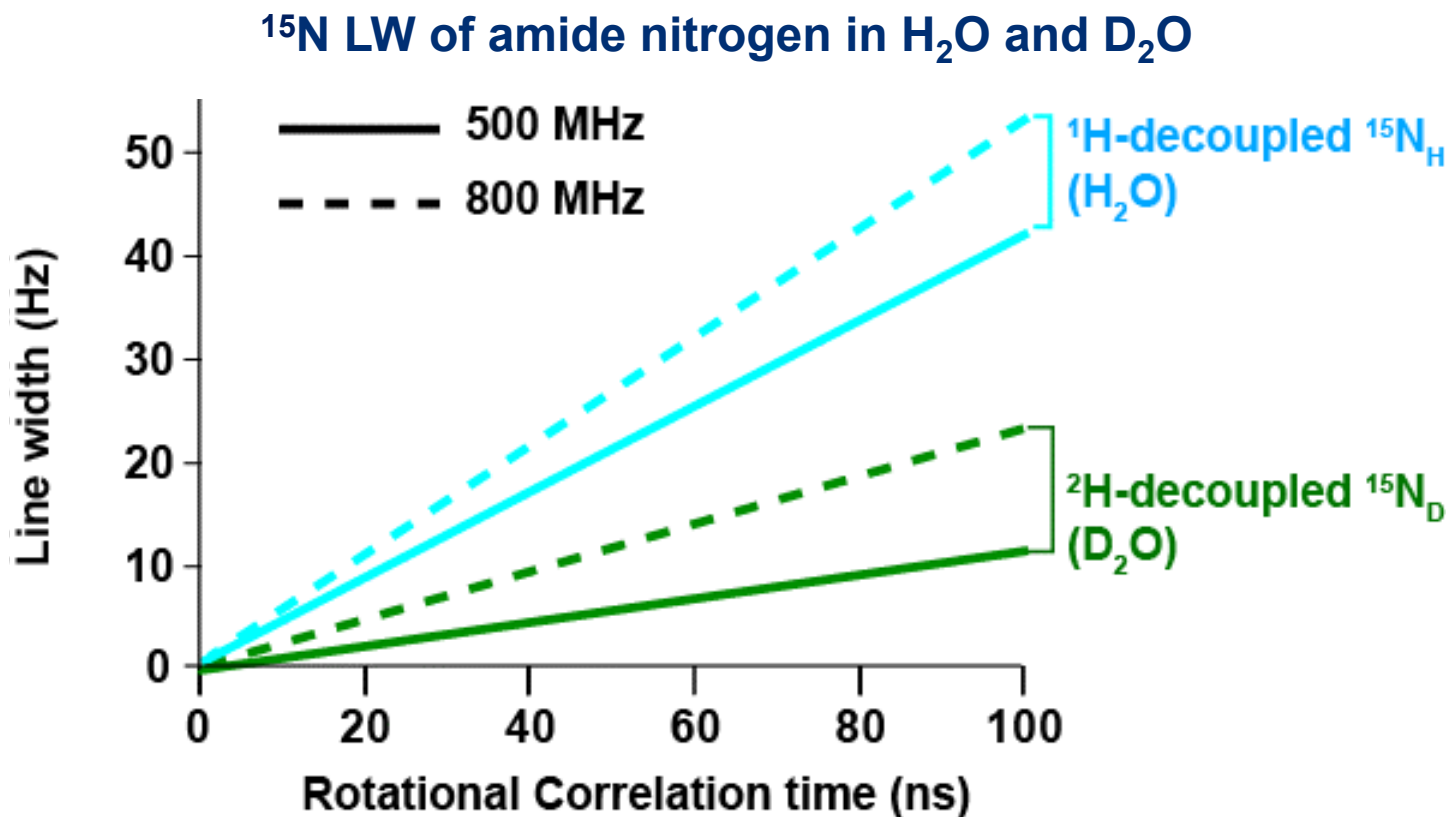
The sharper the line, the heigher the peak

We care about height not intensity (volume)

Very promising at Avance 500 ☺
cryoprobe for C,N-detection cold ^{15}N preamp

But no gain at Avance 800 ☹

Conventional ^{15}N detection uses deuterium-bound amides ($^{15}\text{N}_\text{D}$)



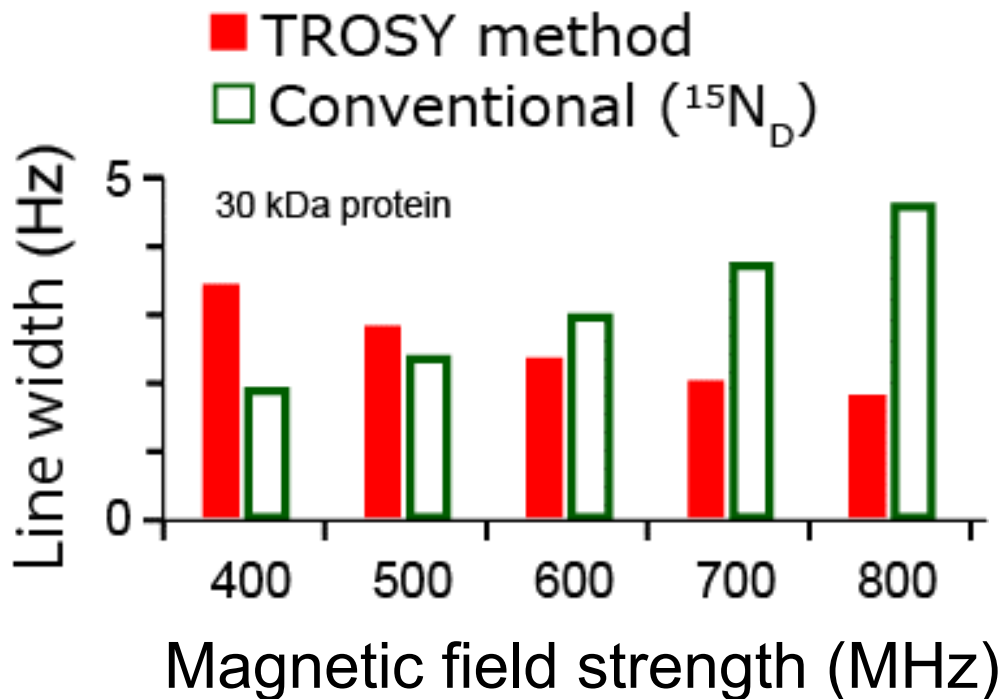
- ^{2}H -decoupled $^{15}\text{N}_\text{D}$ in D_2O is narrower than ^{1}H -decoupled $^{15}\text{N}_\text{H}$ in H_2O
- An expensive higher field magnet would simply broaden our resonance.

Reconsideration of detection strategy is needed to benefit from high-field.

$^{15}\text{N}_\text{D}$ line width increases with field strength

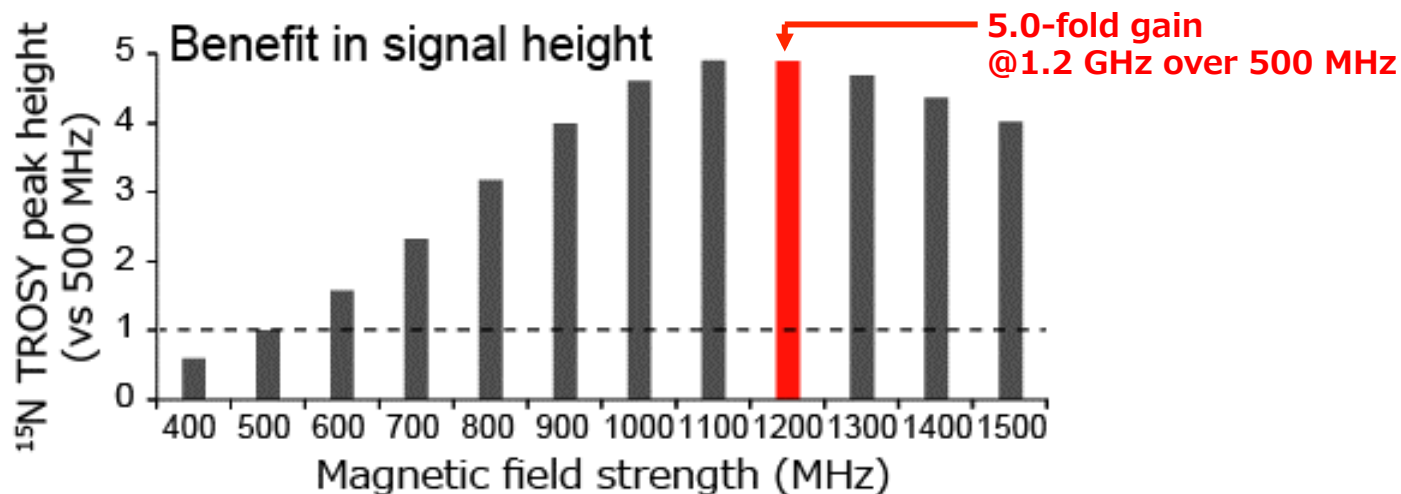
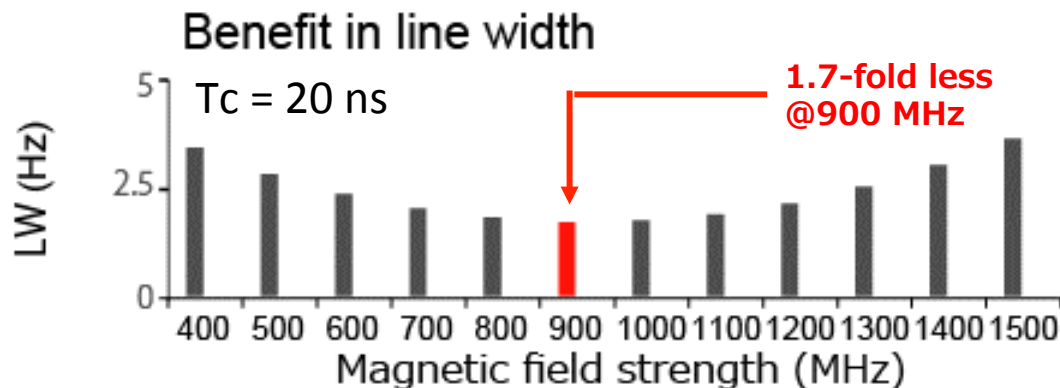
$^{15}\text{N}_\text{H}$ TROSY line width should decrease with field strength

LW of conv. $^{15}\text{N}_\text{D}$ and TROSY $^{15}\text{N}_\text{H}$ resonances



TROSY will enhance resolution and sensitivity of ^{15}N detection at higher magnetic fields

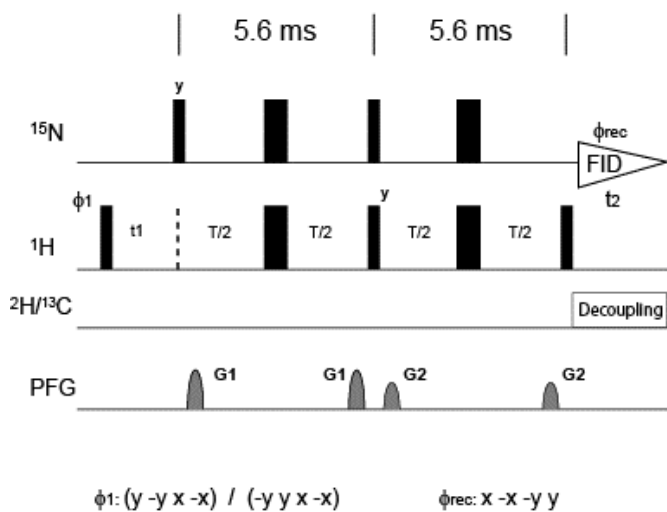
^{15}N TROSY provides maximal benefits @900 MHz for resolution and @1.2 GHz for sensitivity



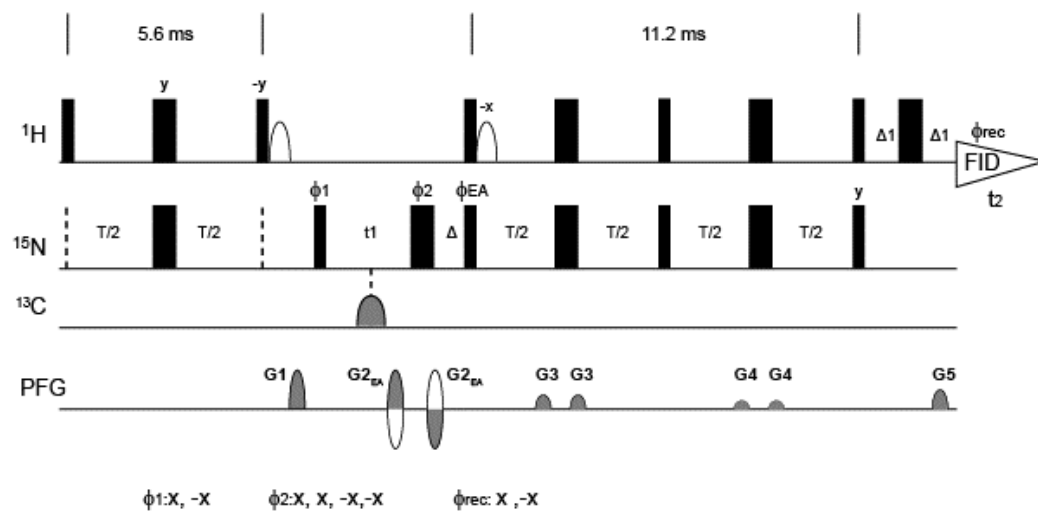
Combination of ^{15}N TROSY & higher magnetic fields will provide additional gains.

^{15}N -detected TROSY is shorter than the ^1H detected version

^{15}N -detected TROSY-HSQC

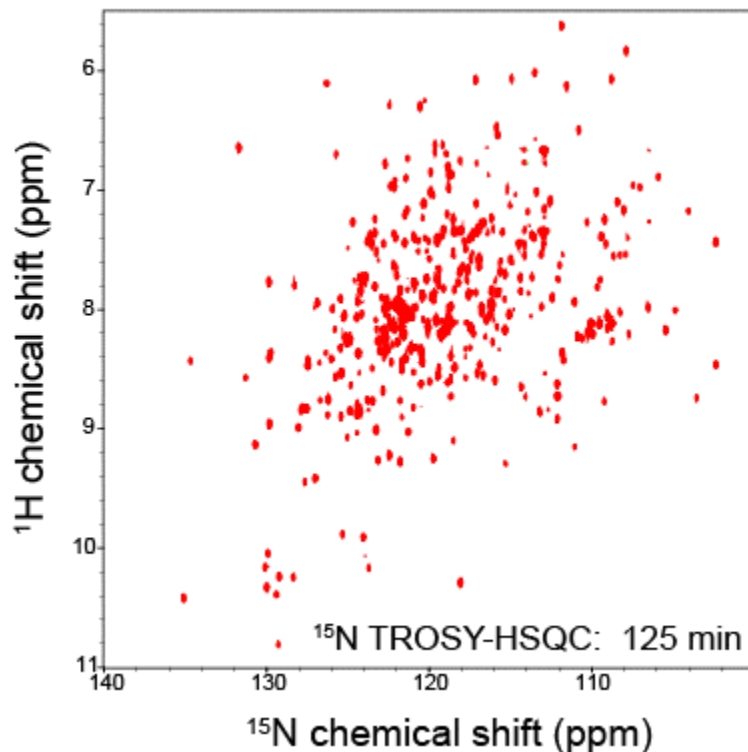


^1H -detected TROSY-HSQC

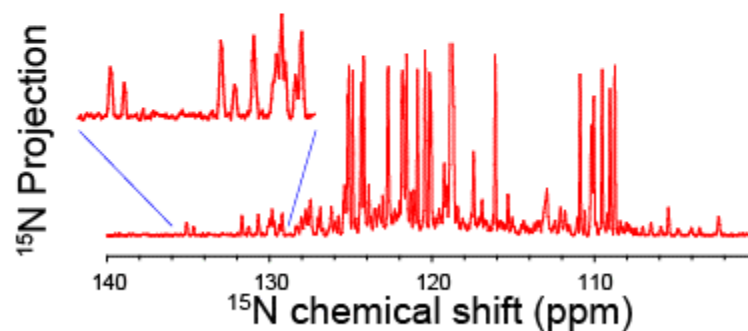


^{15}N -detected 2D TROSY experiment of 70 kDa protein

$^2\text{H}^{15}\text{N}$ MBP 1mM, 278K: $T_c = 40$ ns ($\sim 70\text{K}$)

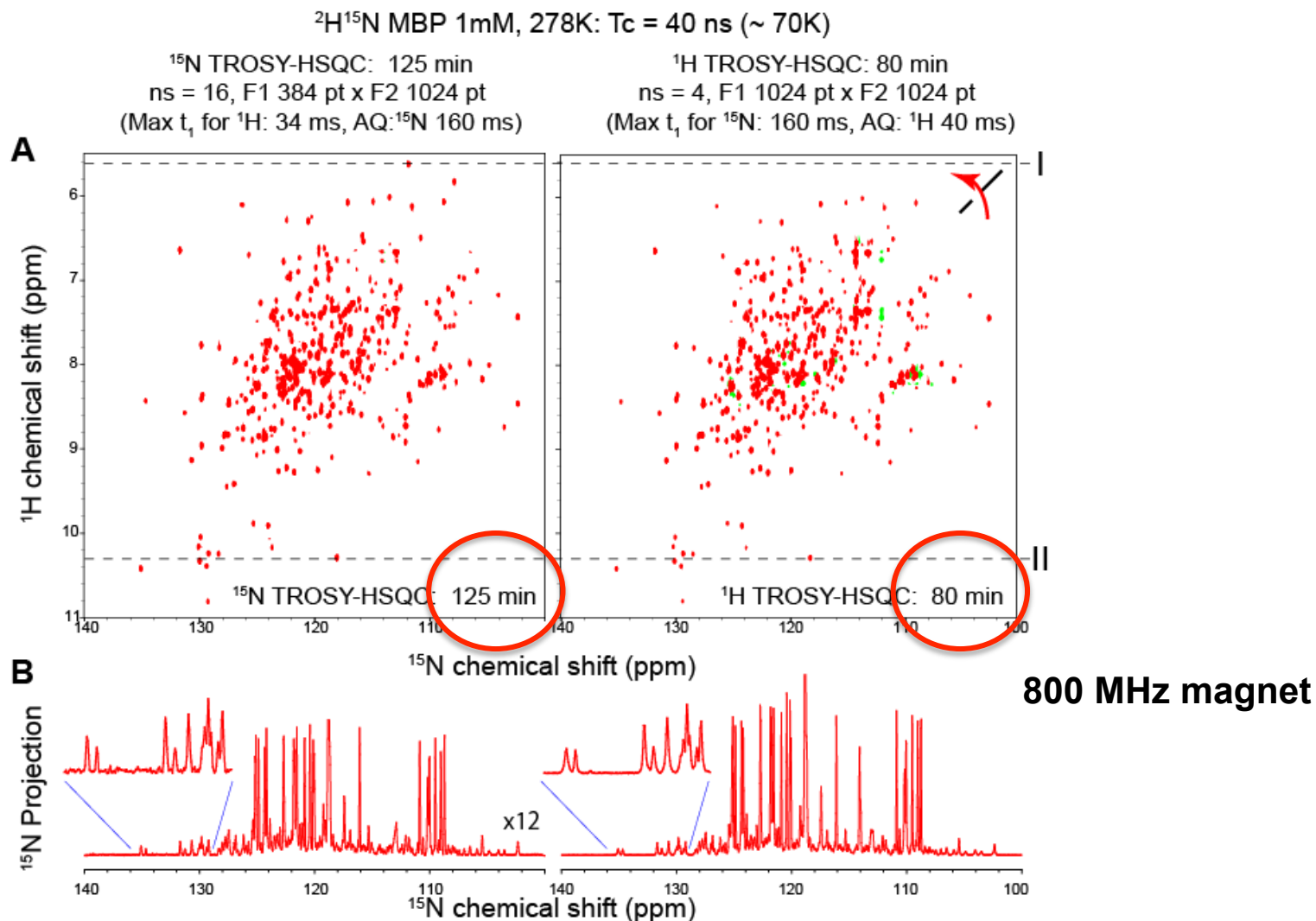


^{15}N TROSY-HSQC: 125 min
ns = 16, F1 384 pt x F2 1024 pt
(Max t_1 for ^1H : 34 ms, AQ: ^{15}N 160 ms)
800 MHz magnet



^{15}N -detected TROSY-HSQC yield nice spectra with reasonable amount of time.

^{15}N TROSY-HSQC vs ^1H TROSY-HSQC



$^{15}\text{N}_\text{H}$ TROSY-HSQC showed similar quality as $^1\text{H}_\text{N}$ TROSY-HSQC.

Do we need to deuterate?

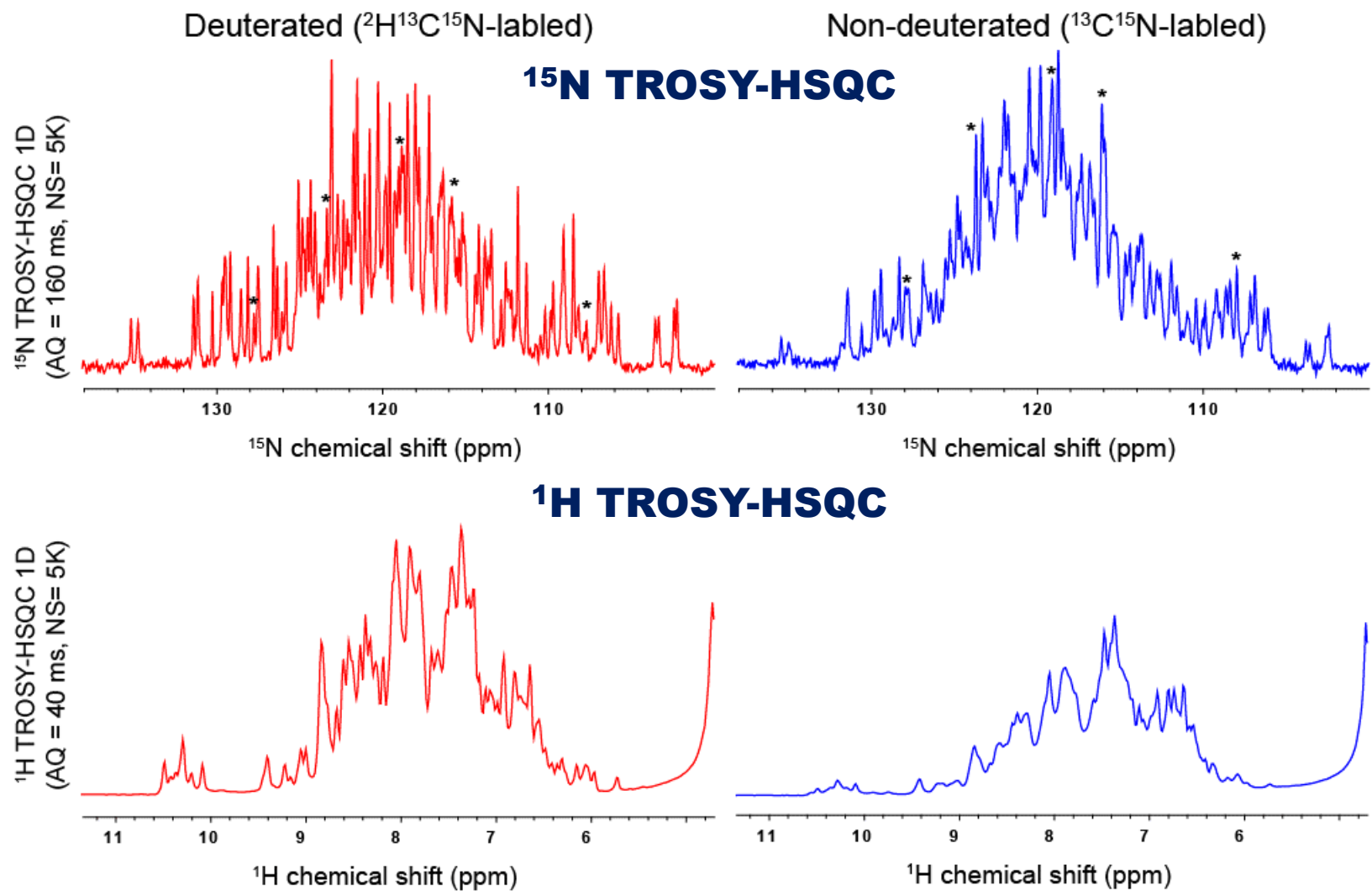
^{15}N has very weak dipolar interactions with nearby ^1H and shouldn't be much broadened

Would alleviate the need for amide back-exchange in deuterated proteins

Reduce cost of making deuterated samples!

Would not be limited to *E.coli* expression!

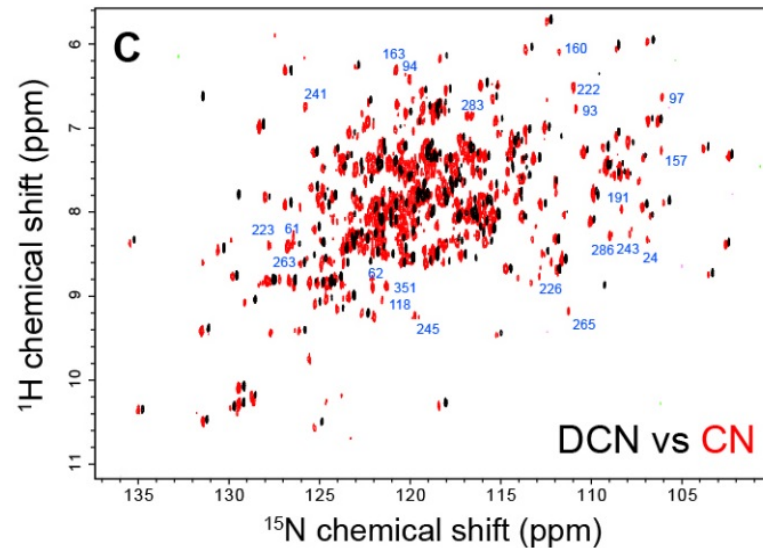
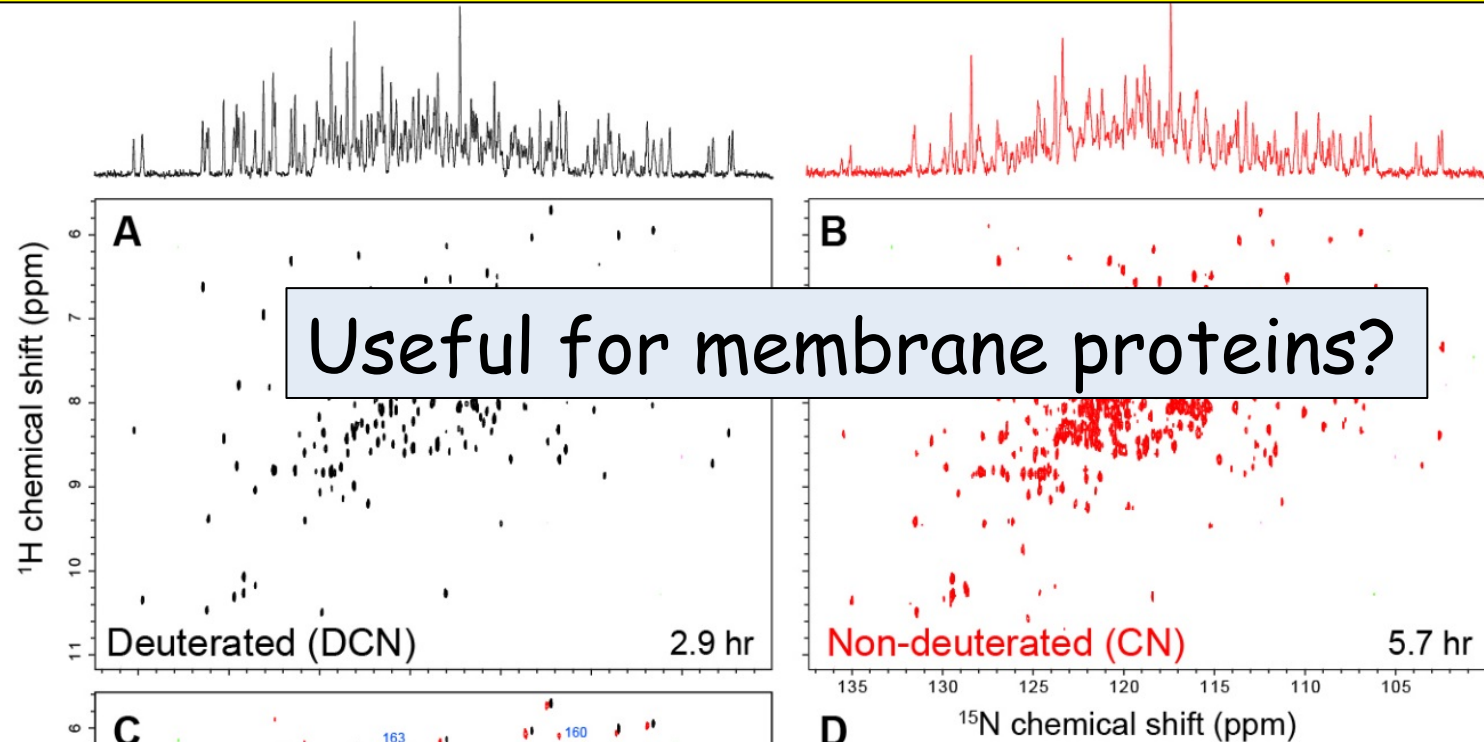
^{15}N and ^1H TROSY without deuteration



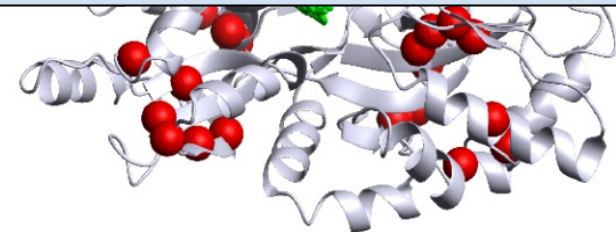
67K protein @800 MHz

^{15}N TROSY would work as well for protonated proteins.

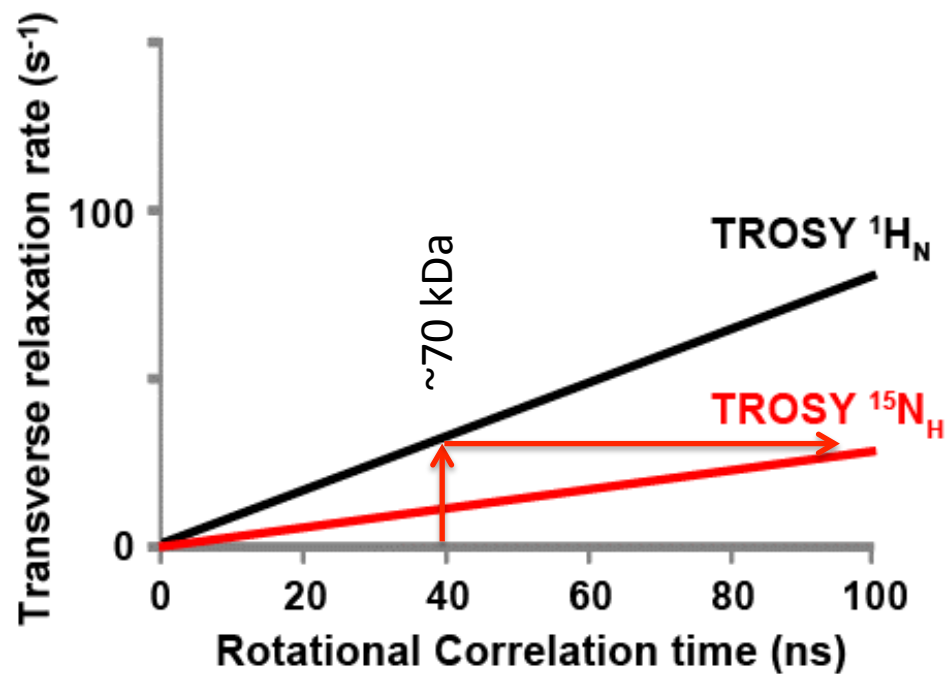
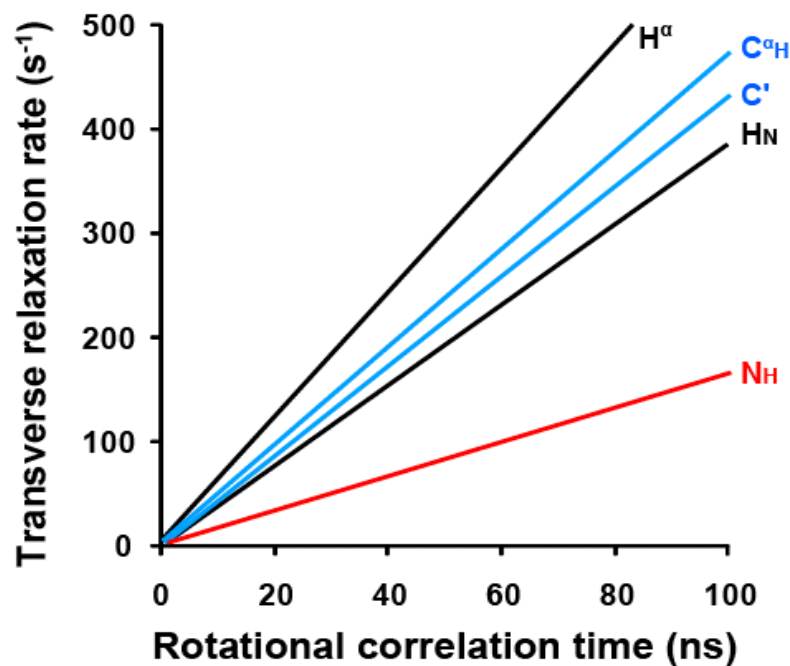
Comparison of $^{15}\text{N}_\text{H}$ detected TROSY using deuterated vs non-deuterated MBP



MBP can be refolded – no problem with incomplete amide back exchange



$^{15}\text{N}_\text{H}$ -detected TROSY is least deteriorated when going to higher molecular weight



Summary

TROSY selection provides resolution and sensitivity of ^{15}N detection in high-magnetic field

$^{15}\text{N}_\text{H}$ TROSY-HSQC spectra can be recorded at only moderately lower sensitivity than $^1\text{H}_\text{N}$ TROSY HSQC

$^{15}\text{N}_\text{H}$ TROSY is least sensitive to molecular weight and promises to extend the size range of NMR to high-molecular weight systems

^{15}N TROSY would work as well for protonated proteins and will be beneficial for proteins that can be obtained only by eukaryotic expression

^{15}N TROSY with non-deuterated protein eliminates the incomplete-amide back-exchange problem

These results illustrate the potential of ^{15}N TROSY as a means to exploit the high resolution offered by high field magnets near and above 1 GHz

^{15}N detected TROSY will benefit assignments of crowded spectra of unstructures regulatory domains with prolines in phosphorylation sites and **much more complex systems could be assigned at higher field**

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