



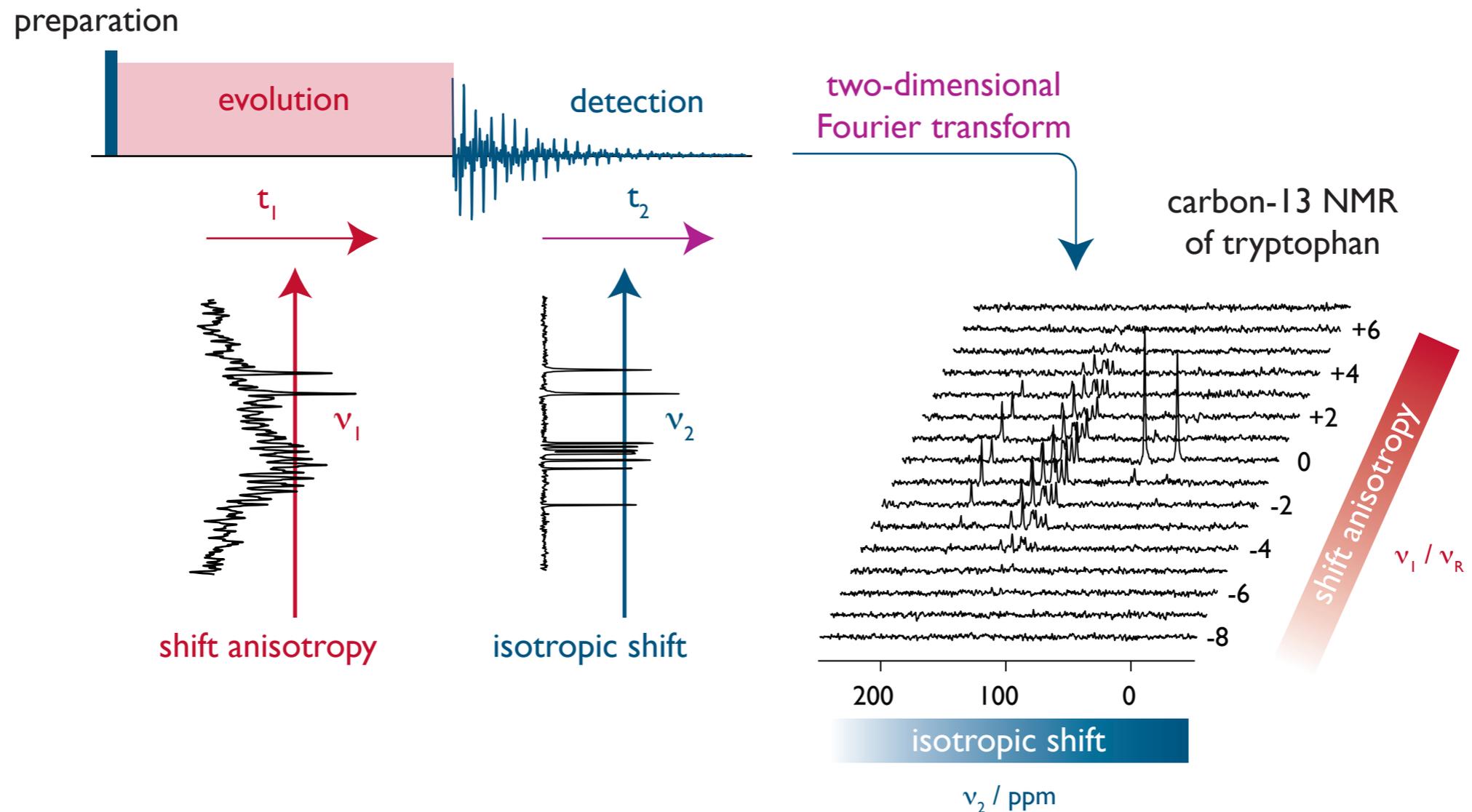
# Two-dimensional separation experiments: measuring anisotropic interactions

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# Two-dimensional separation experiments

A two-dimensional separation experiment consists of three time periods: **Preparation, evolution and detection.**

- ★ The evolution and detection Hamiltonians are different.
- ★ The two-dimensional spectrum **separates** the spectra which result from the two Hamiltonians.



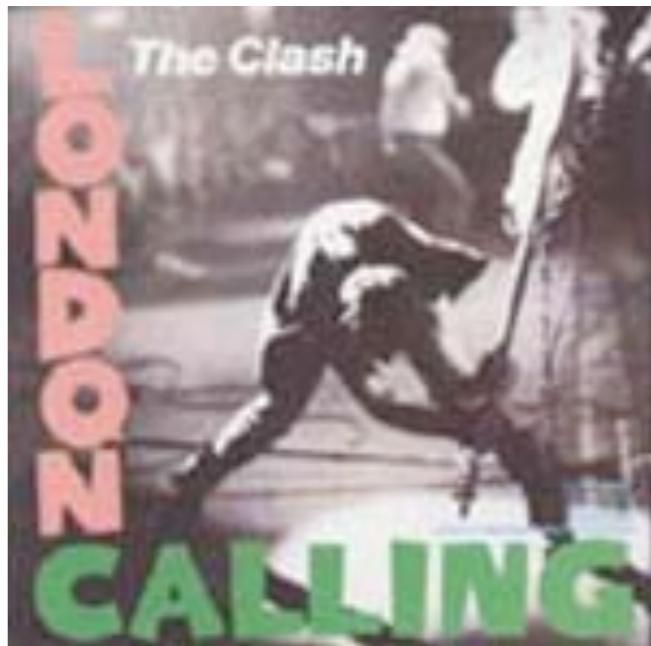
# Philosophy

## “Solution-like” approach:

Rapid MAS, Efficient Decoupling

$$H = H_{\text{iso}}$$

No orientational information; recoupling

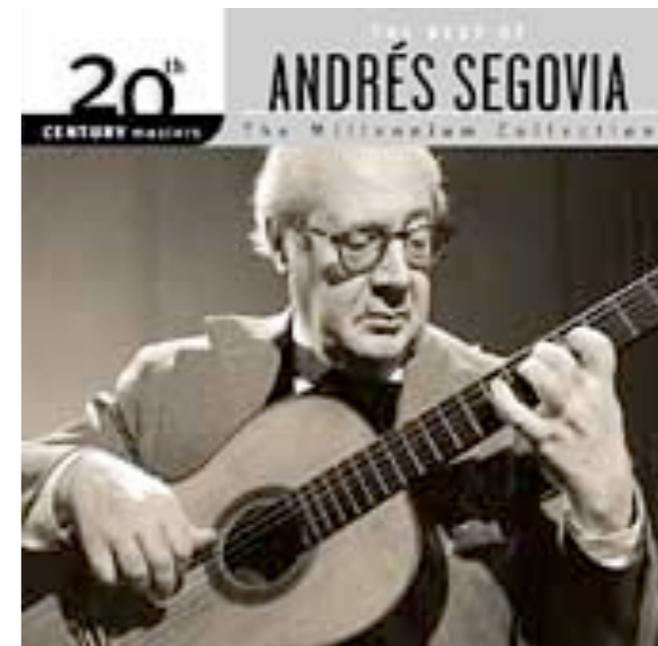


## “Classical solids” approach:

Slow (or no) MAS

$$H = ?$$

Retain orientational information



# Types of separation

## Shift anisotropy and isotropic shift

- ★ Many experiments have been designed which separate the **isotropic** and **anisotropic** parts of the chemical shift Hamiltonian.
- ★ These are used to obtain the shift anisotropies of complicated molecules where overlap of powder patterns or sideband manifolds from different chemical sites prevents their measurement using a one-dimensional spectrum.

## Heteronuclear dipolar coupling and chemical shift

- ★ The archetypal two-dimensional separation method is the “**separated local field**” experiment
- ★ This is used to establish the **orientation** of the principal axis system of the chemical shift tensor relative to the molecule.

# Experimental approaches

## Wideline separation

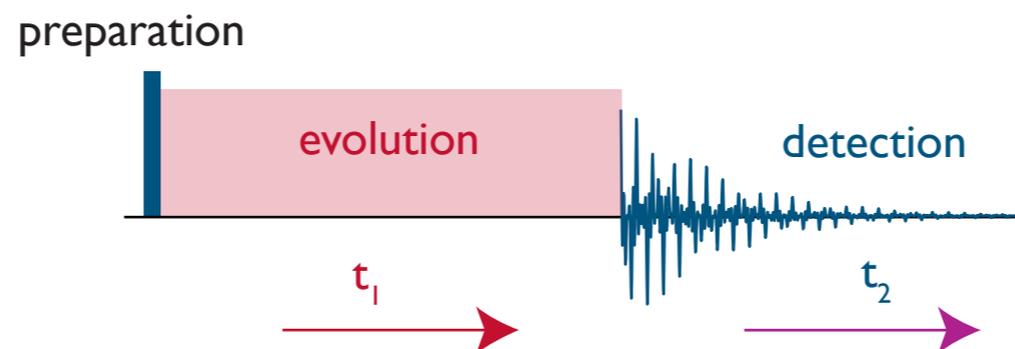
- ★ A non-spinning experiment gives a powder lineshape in two dimensions

## Recoupled Hamiltonian

- ★ MAS gives resolution in  $\nu_2$  while a recoupling sequence reintroduces the desired Hamiltonian to give a “powder pattern” in  $\nu_1$ .

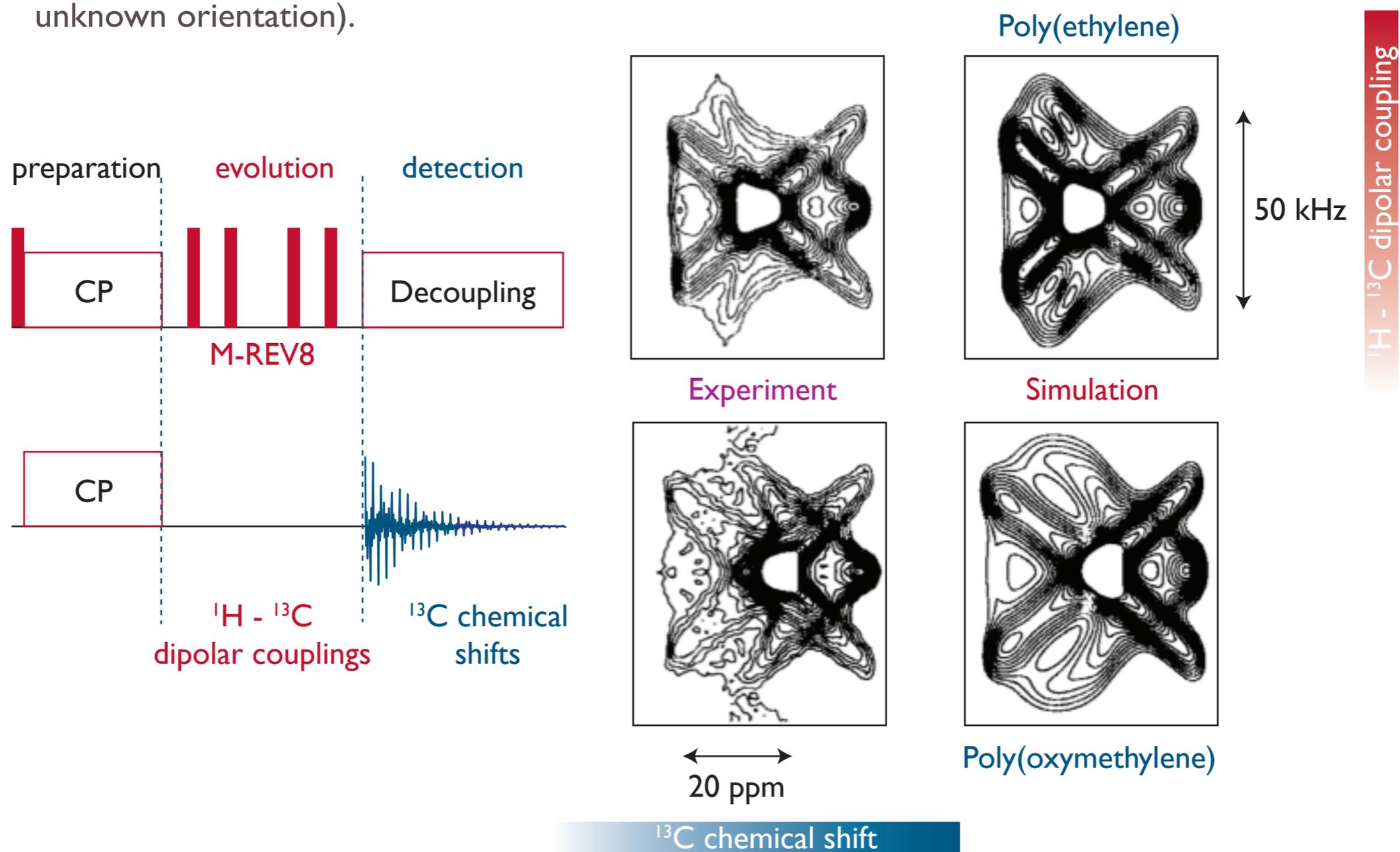
## Separation of MAS sideband manifolds

- ★ Sideband manifolds in  $\nu_1$  are separated by the isotropic shift in  $\nu_2$ .
- ★ With very slow spinning (“magic angle turning”) sequences of  $\pi$  pulses are used to give the isotropic shift in  $\nu_1$  and sideband manifolds in  $\nu_2$ .



# “Separated local field” experiment

- ★ This separates the heteronuclear dipolar coupling in  $\nu_1$  and the chemical shift in  $\nu_2$ .
- ★ The resulting two-dimensional spectrum depends on the relative orientation of the dipolar coupling tensor (with the z axis along the internuclear vector) and the chemical shift tensor (usually with an unknown orientation).

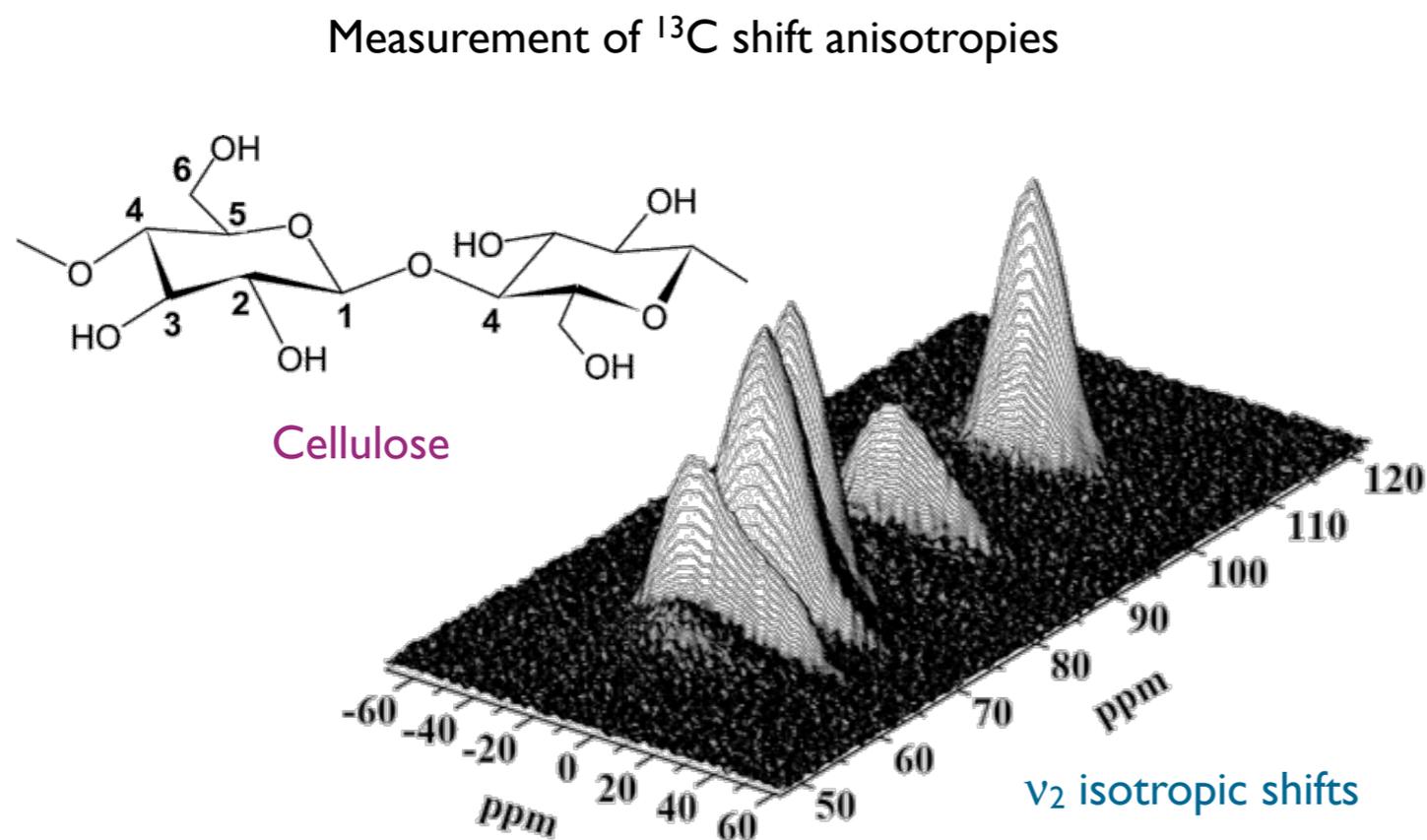


# Separation of “recoupled” powder patterns

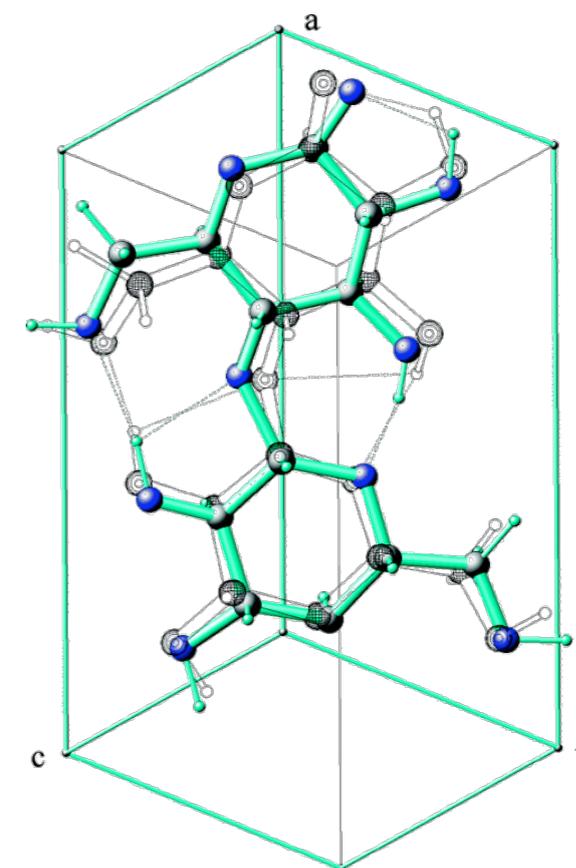
These experiments separate a recoupled powder pattern in  $\nu_1$  and the isotropic shift in  $\nu_2$ .

## Example: cellulose

The carbon-13 chemical shift tensors measured in this way have been used as restraints in a molecular dynamics refinement of the neutron structure.



$\nu_1$  shift anisotropy  
via powder pattern

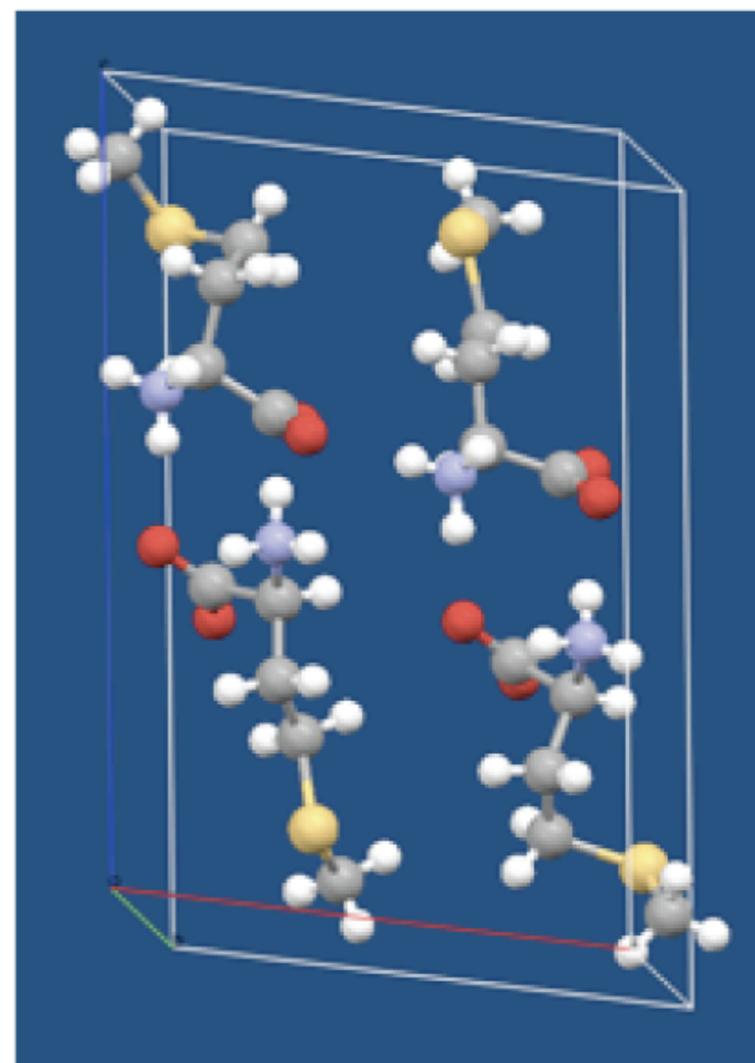
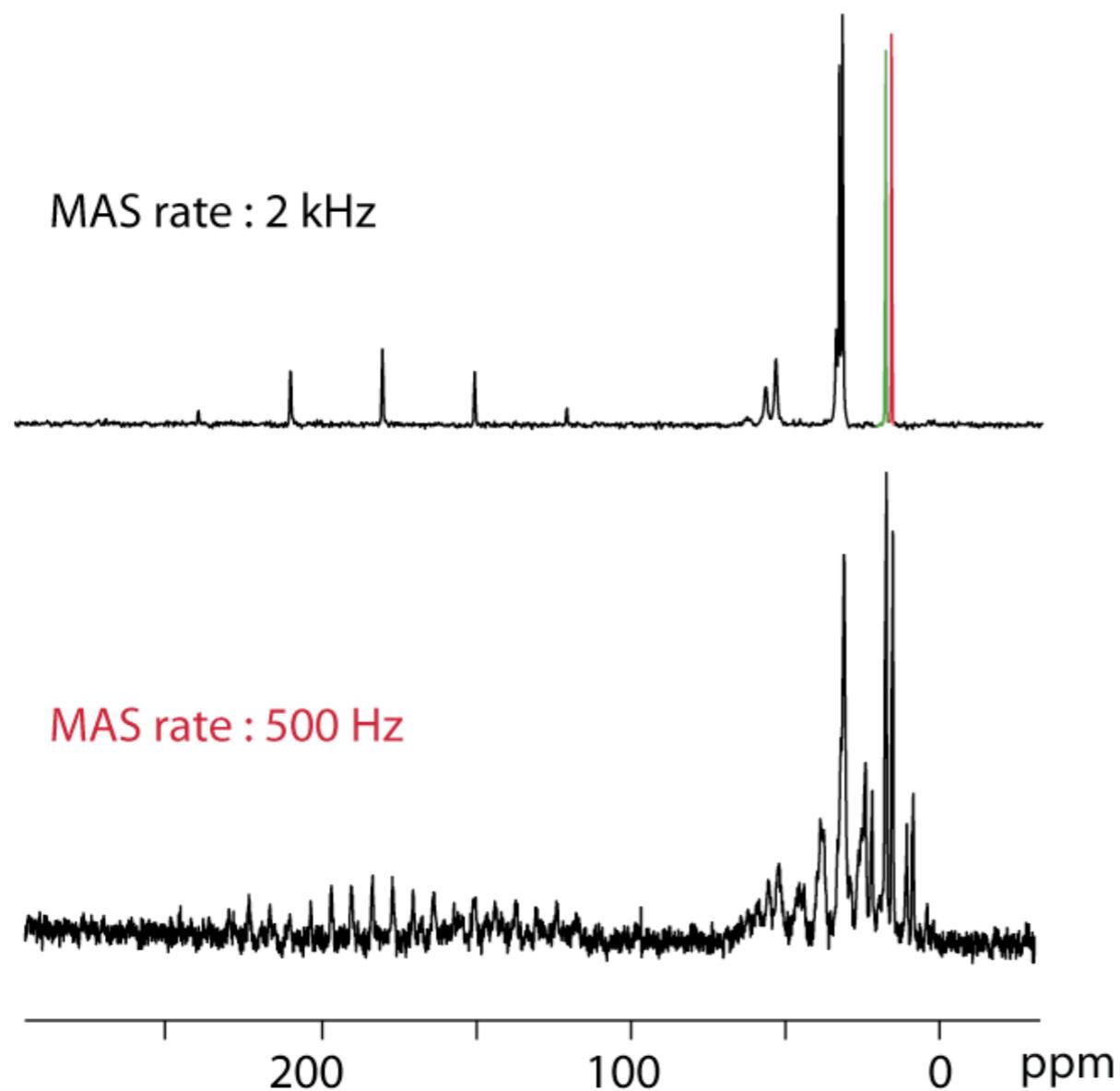


Chemical shift optimized structure  
compared with neutron diffraction data

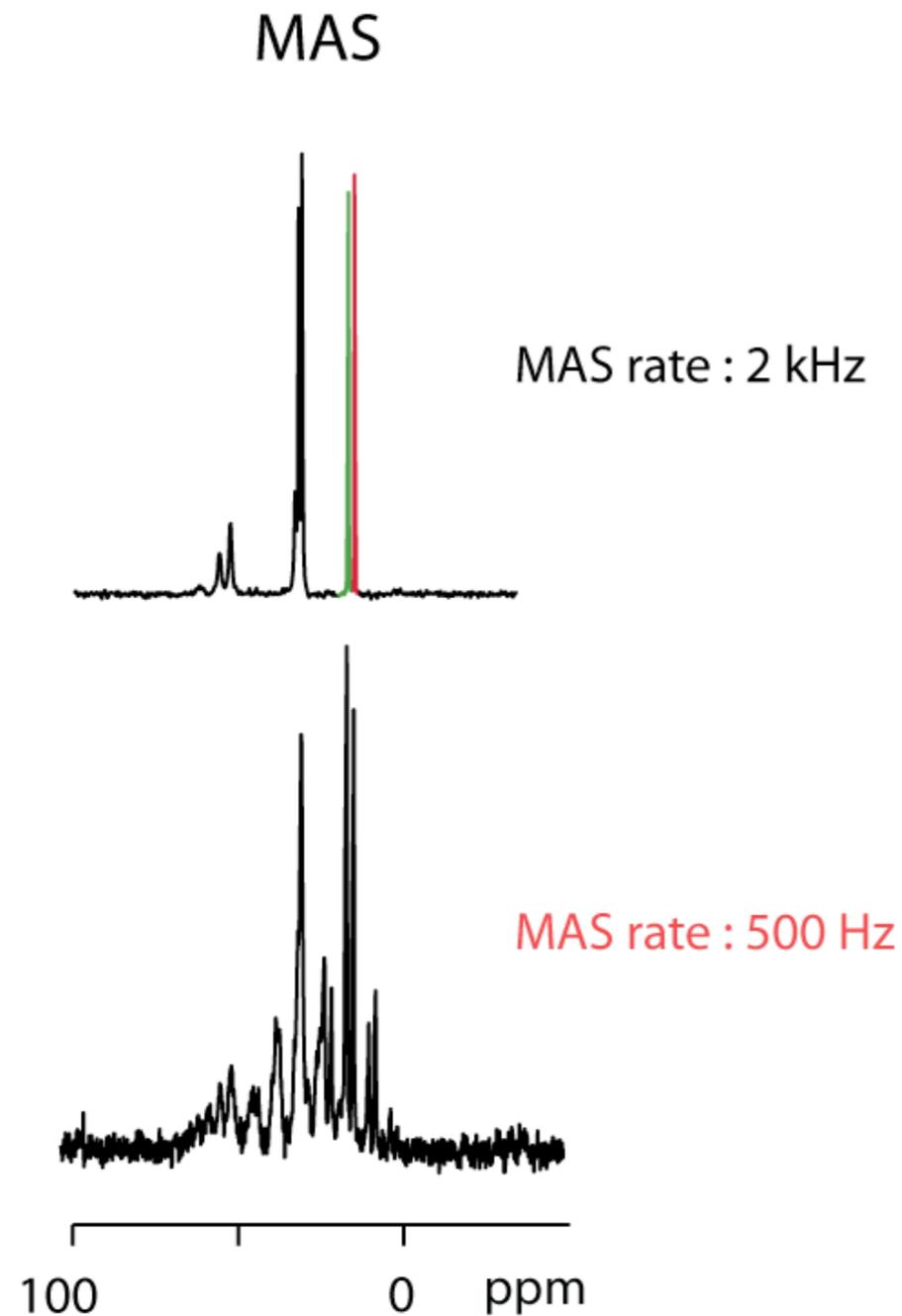
# Shift anisotropies by MAS sidebands

- ★ MAS allows the resolution of different chemical environments via their isotropic shifts.
- ★ With slow MAS the **envelope** of the sideband intensities reflects the shift anisotropy.
- ★ A few sidebands give a more **reliable** result for the anisotropy than a wideline spectrum.

## methionine carbon-13 NMR



# Problems with MAS sideband measurements



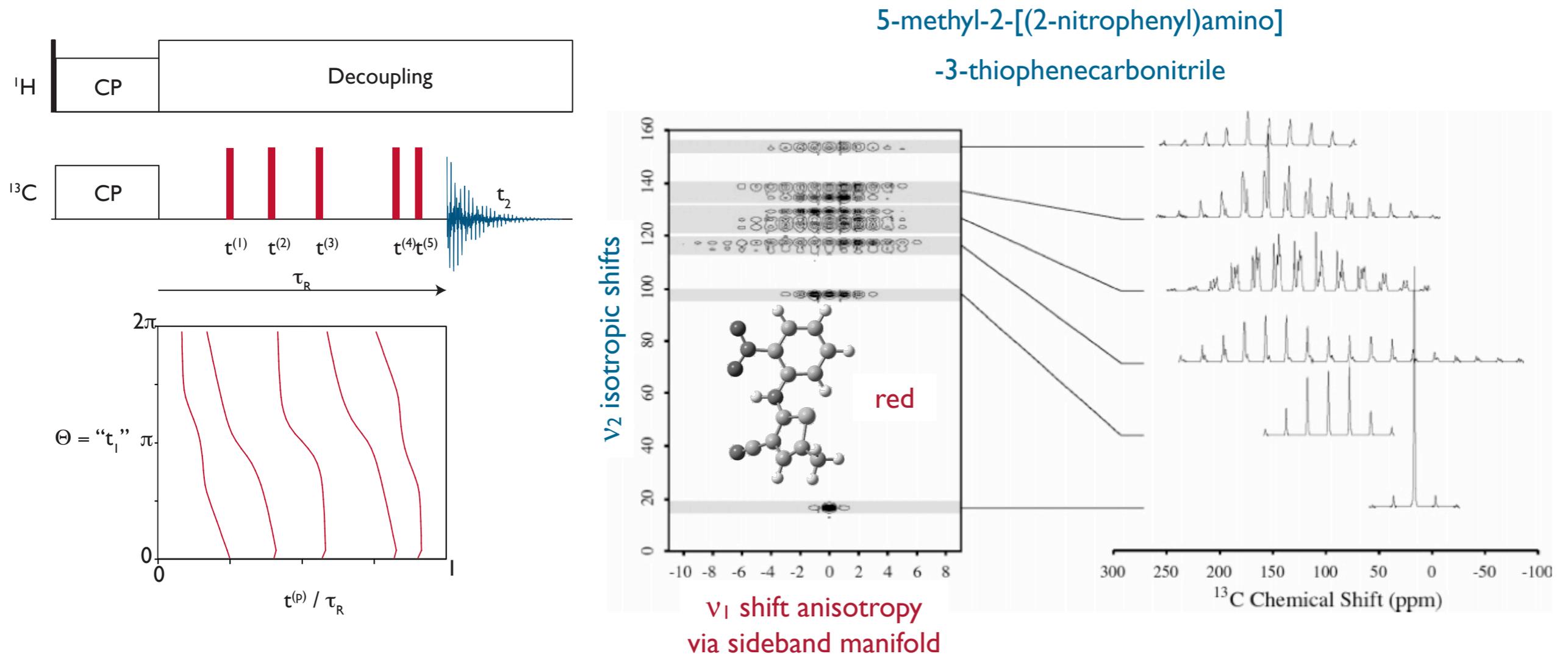
There are many problems when measuring sideband intensities:

- ★ sideband manifolds from different sites **overlap**
- ★ for small shift tensors even at the slowest stable MAS rates there are not enough sidebands
- ★ modern decoupling methods require fast MAS
- ★ there are sideband pattern distortions due to homonuclear dipolar couplings

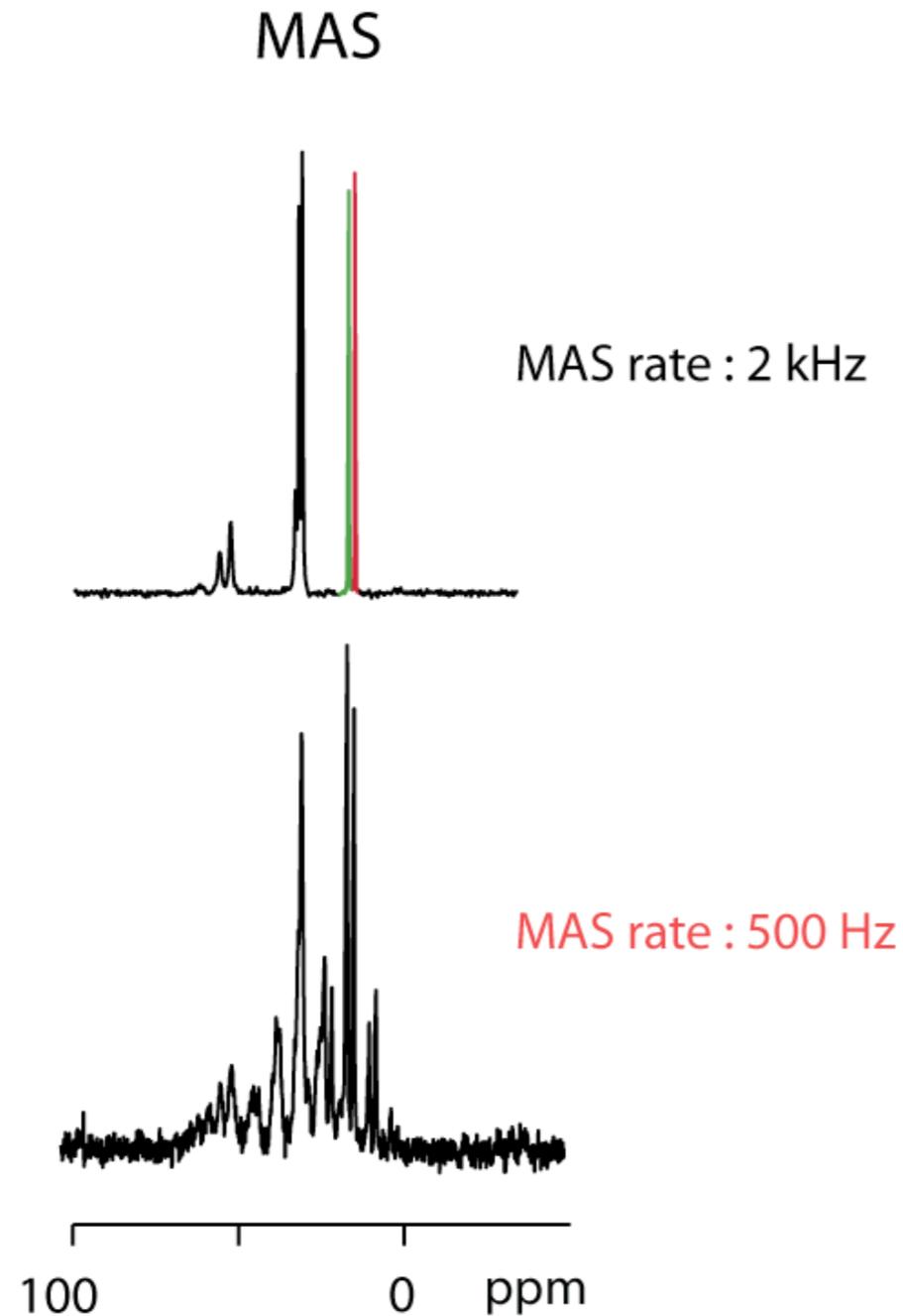
# 2D-PASS

This experiment separates the anisotropic part of the chemical shift measured via the sideband manifold in  $\nu_1$  and the isotropic shift in  $\nu_2$ . Advantages of 2D-PASS include:

- ★ a **minimal** number of  $t_1$  increments is required
- ★ the  $\nu_1$  sideband pattern is **identical** to the MAS pattern
- ★ the use of short sequences and few  $\pi$  pulses



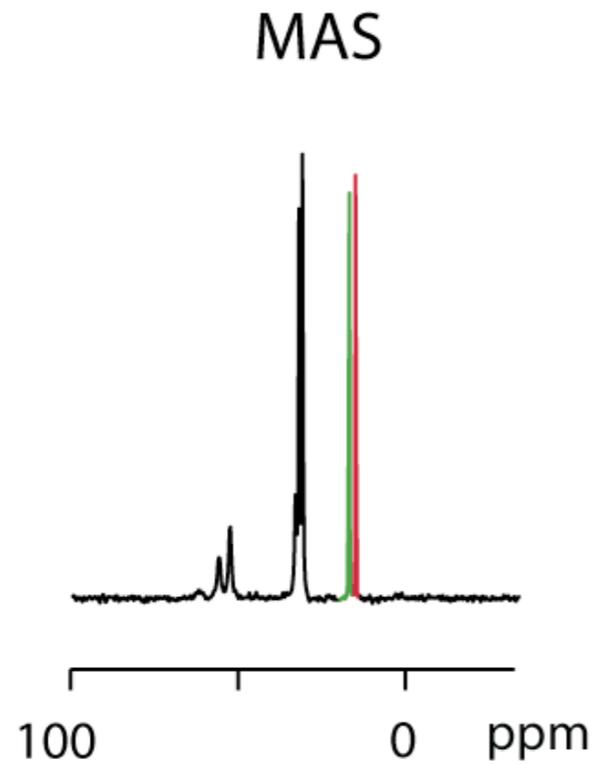
# Problems with MAS sideband measurements



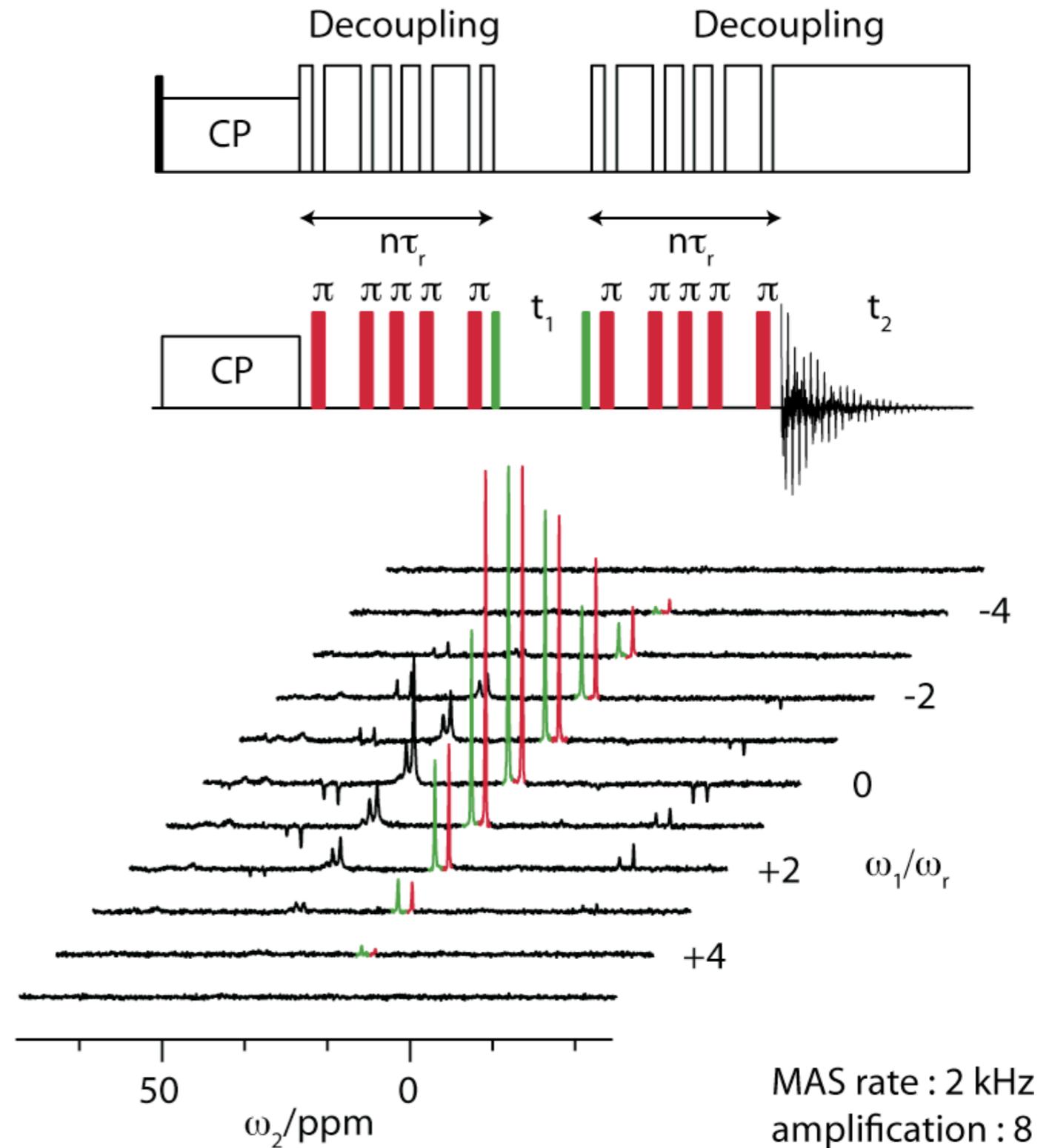
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# CSA amplification: method



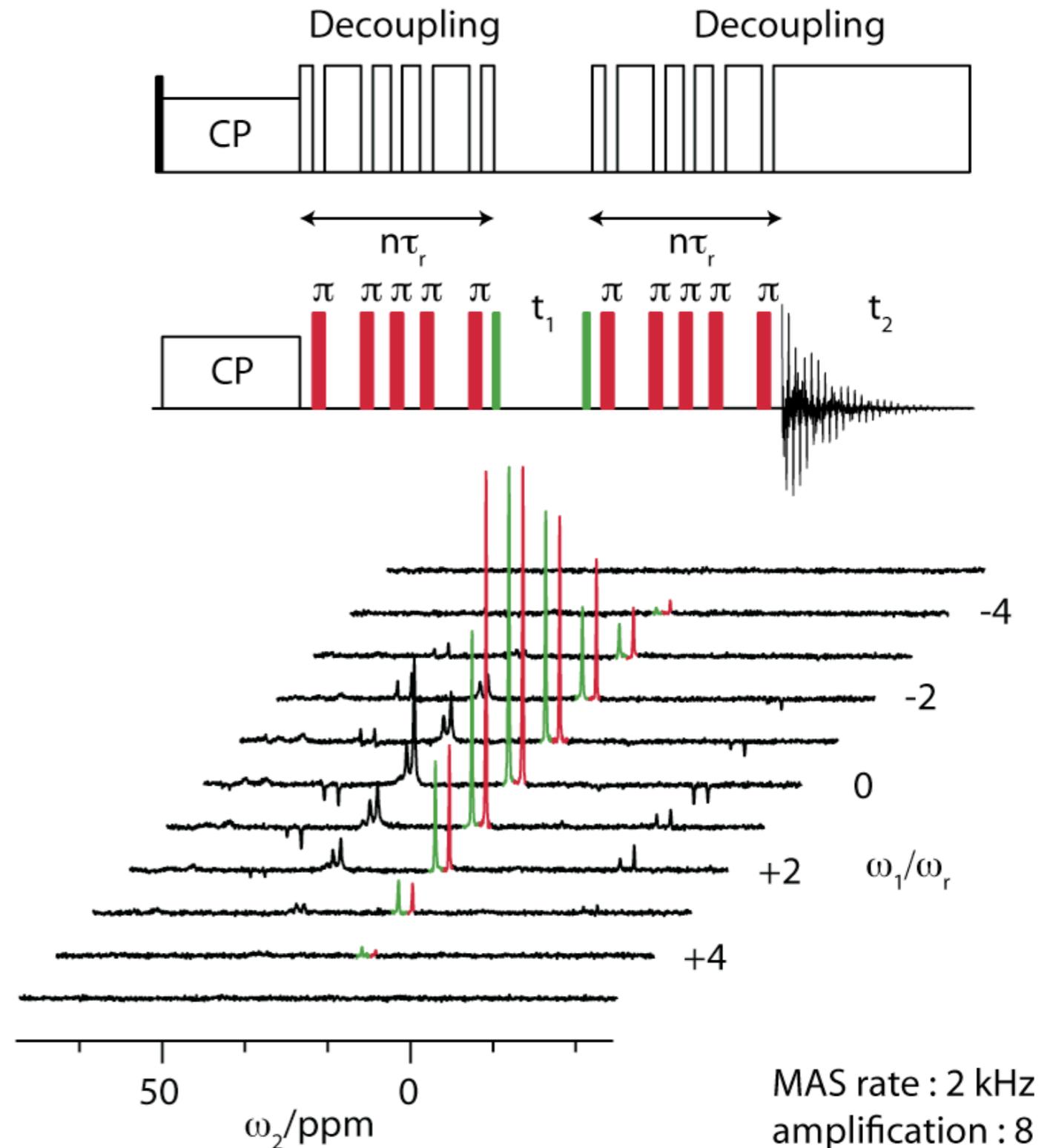
The CSA amplification method separates the fast MAS spectrum in  $\nu_2$  from a sideband pattern in  $\nu_1$  identical to that observed at some fraction of the actual rate.



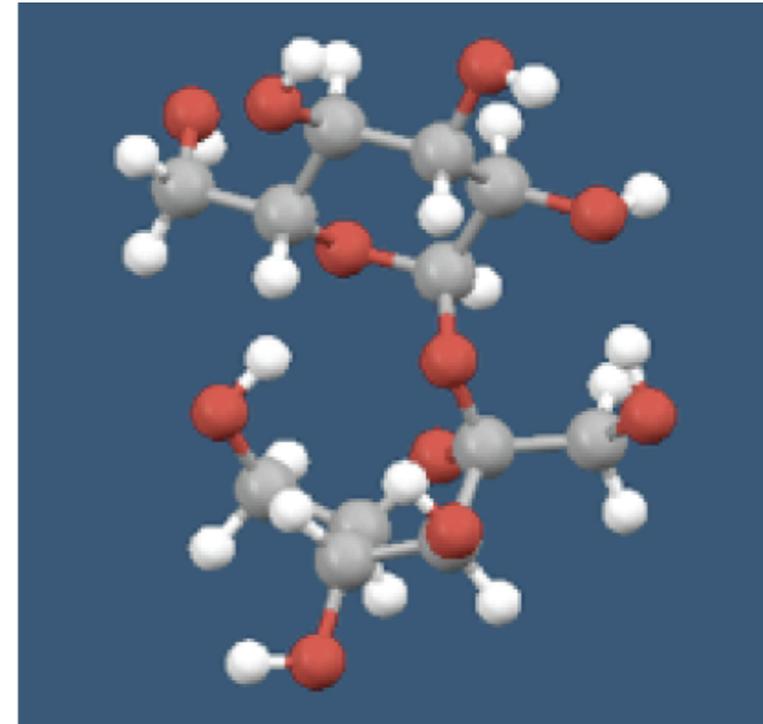
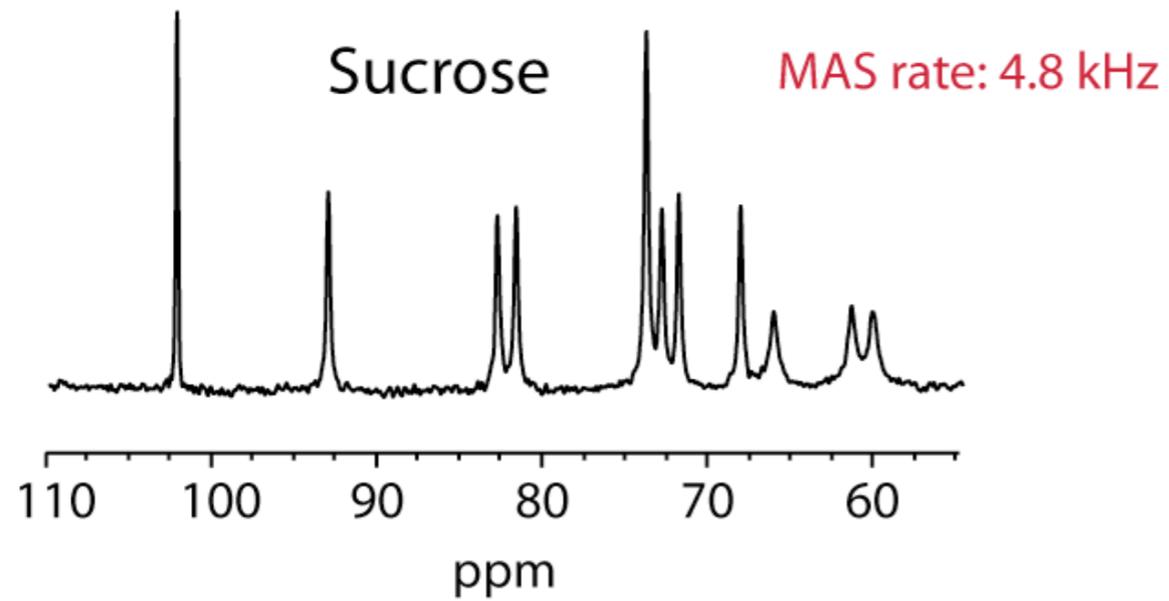
# CSA amplification: advantages

Advantages of the CSA amplification method include:

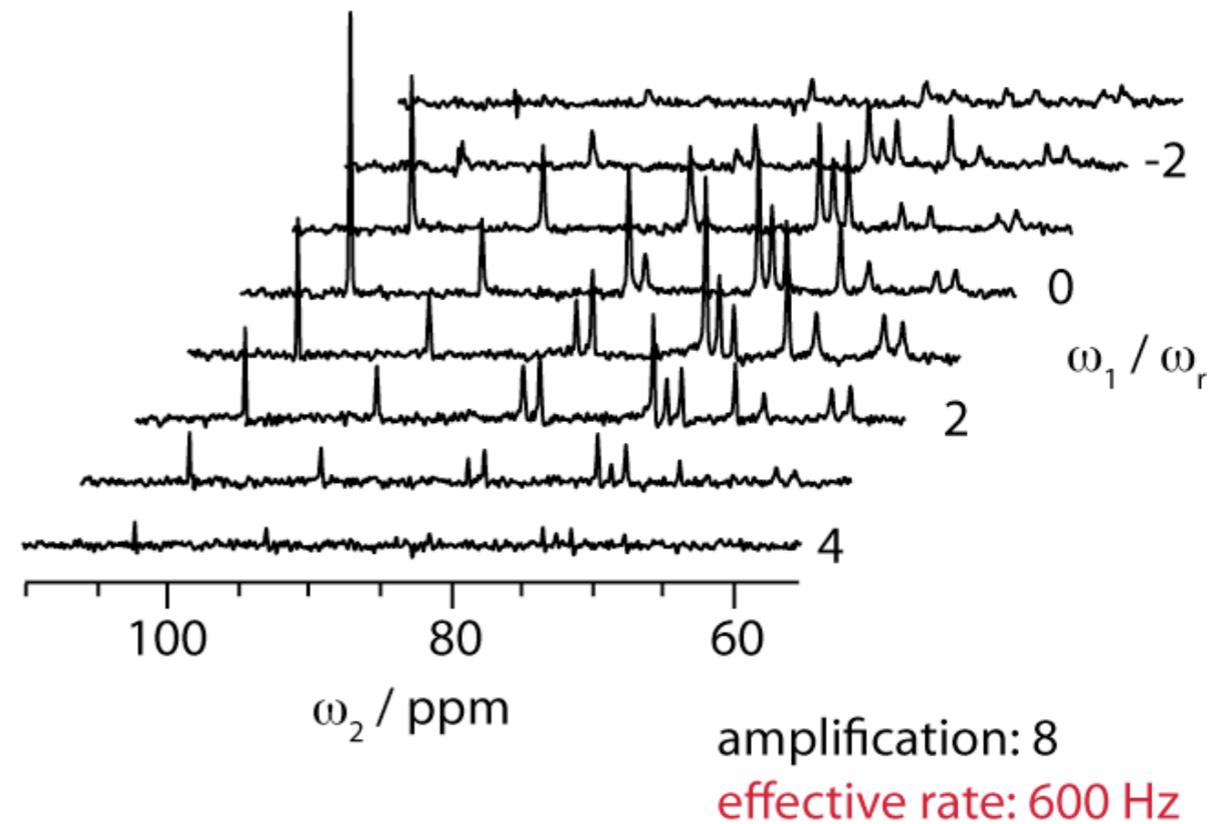
- ★ a minimal number of  $t_1$  increments is required
- ★ the  $\nu_1$  sideband pattern is identical to the **slow** MAS pattern
- ★ short sequences and few  $\pi$  pulses
- ★ no need for slow unstable spinning even for **small shift tensors**
- ★ amplification independent of sequence length



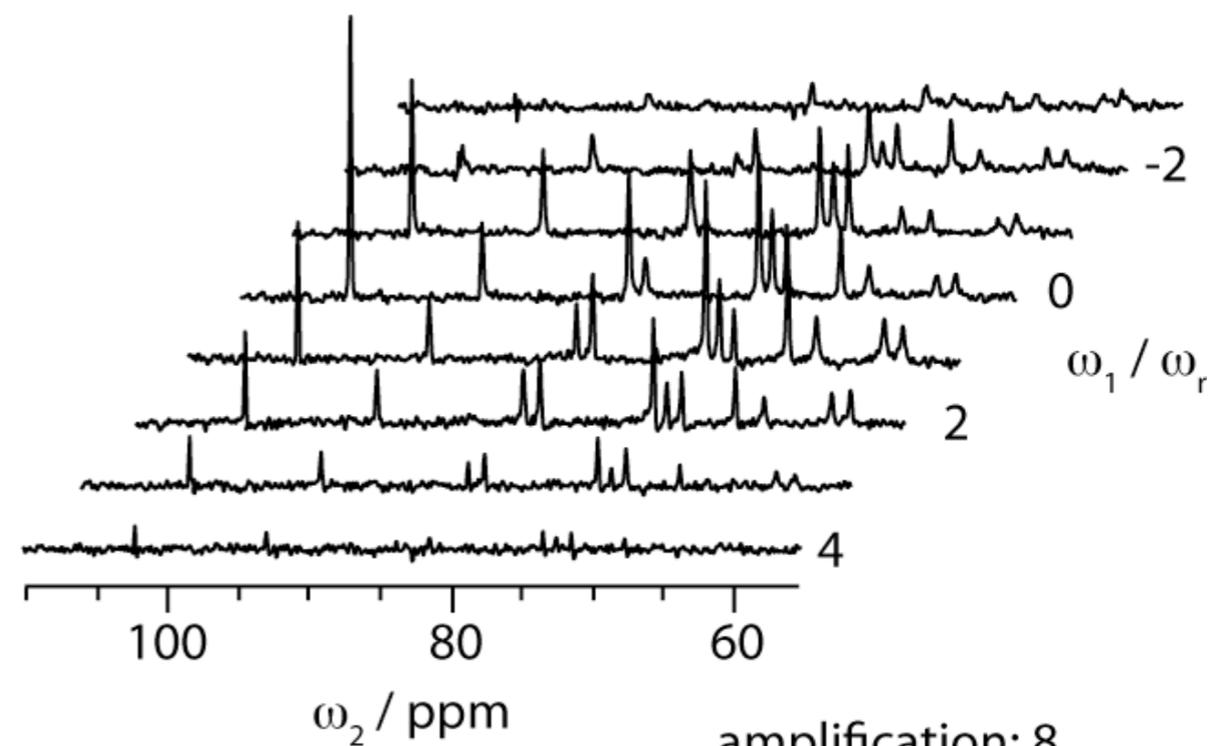
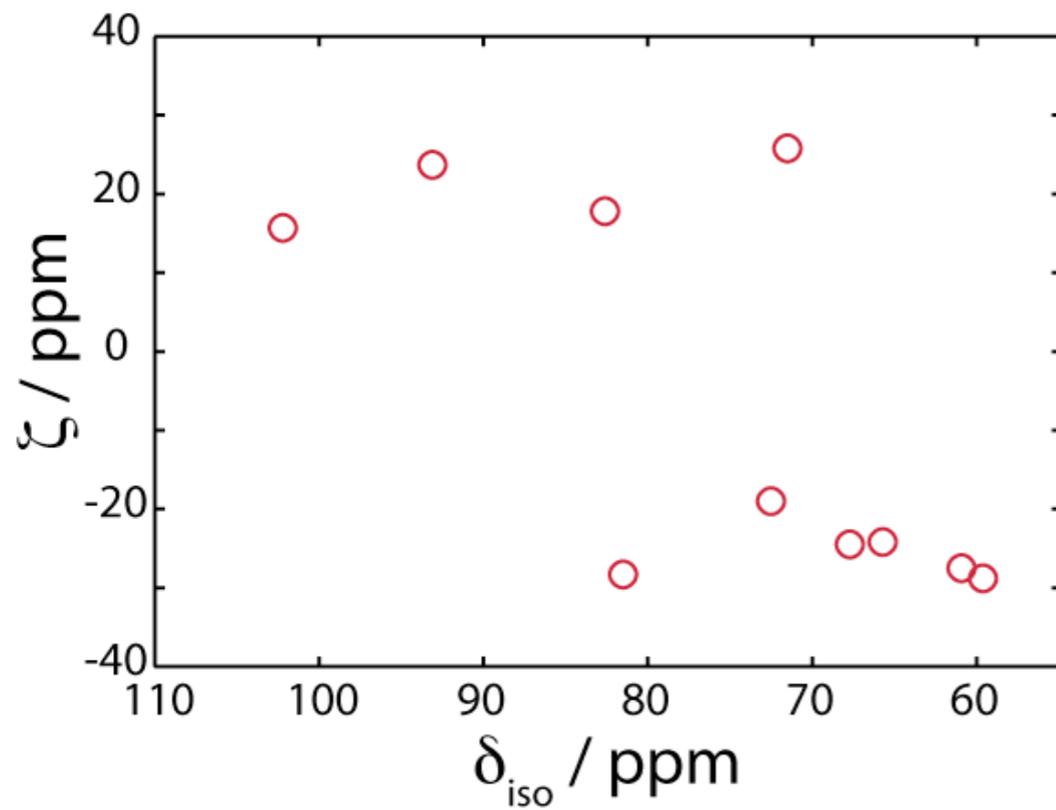
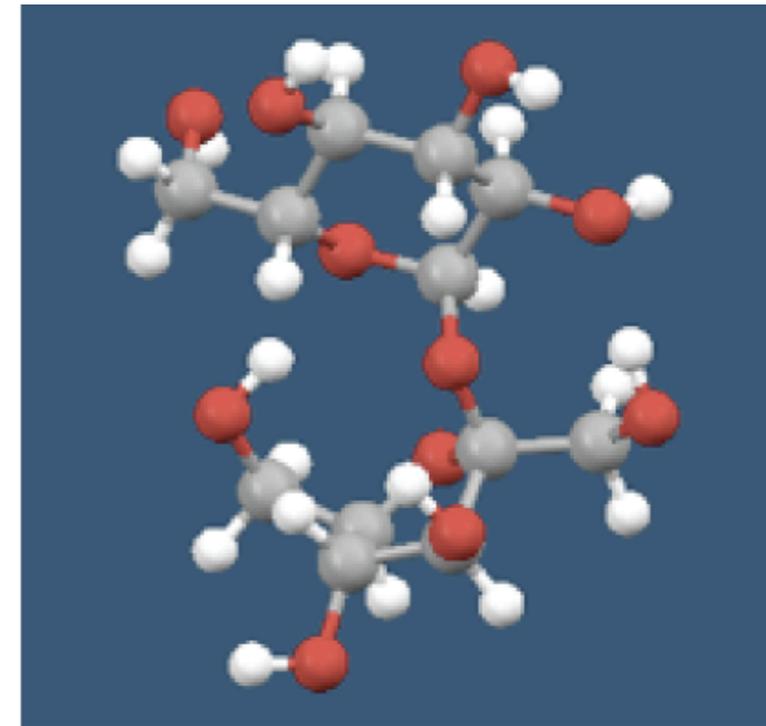
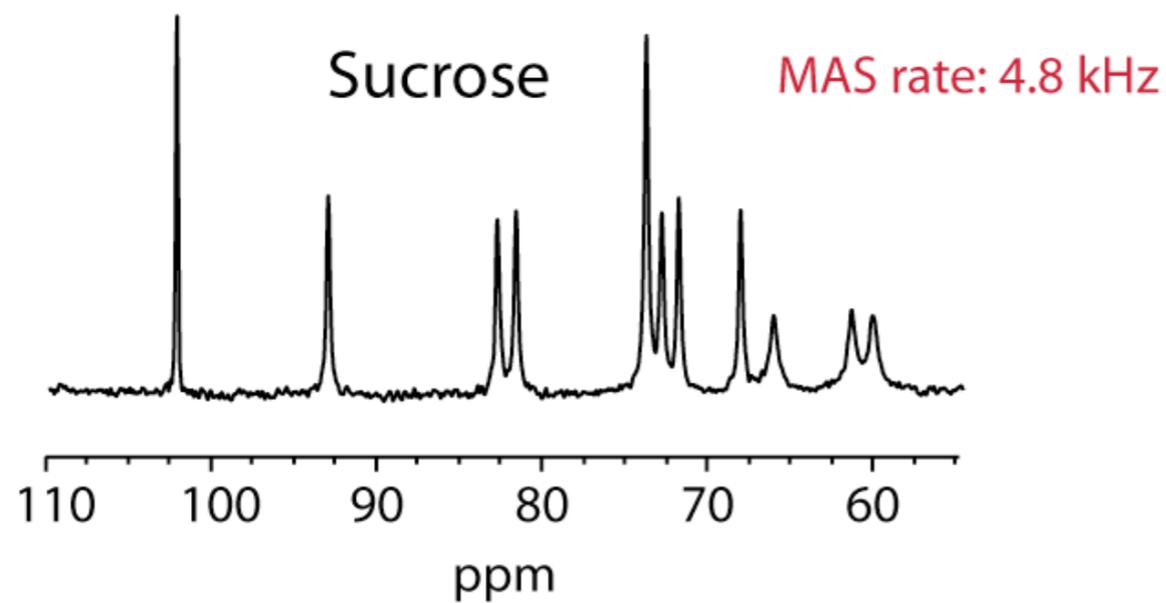
# Sucrose



Optimum number of sidebands for  
measurement of shift anisotropy

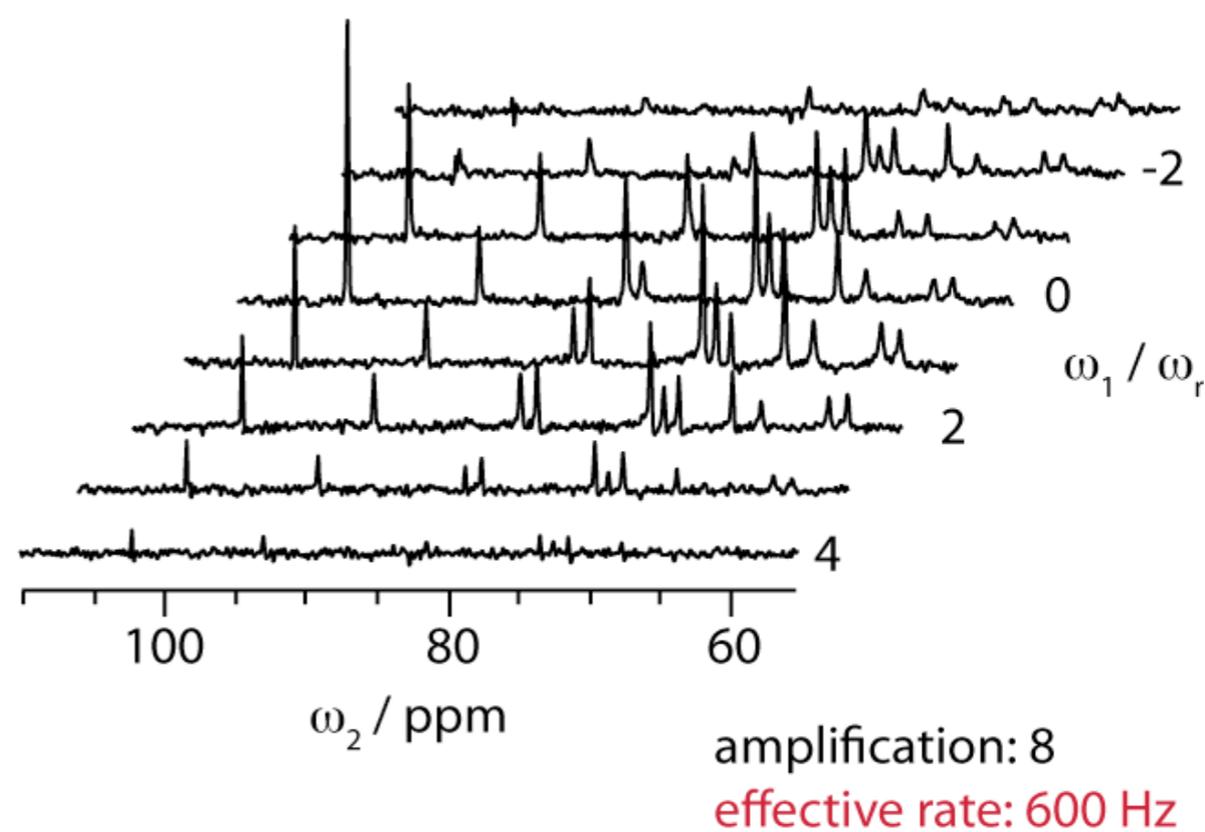
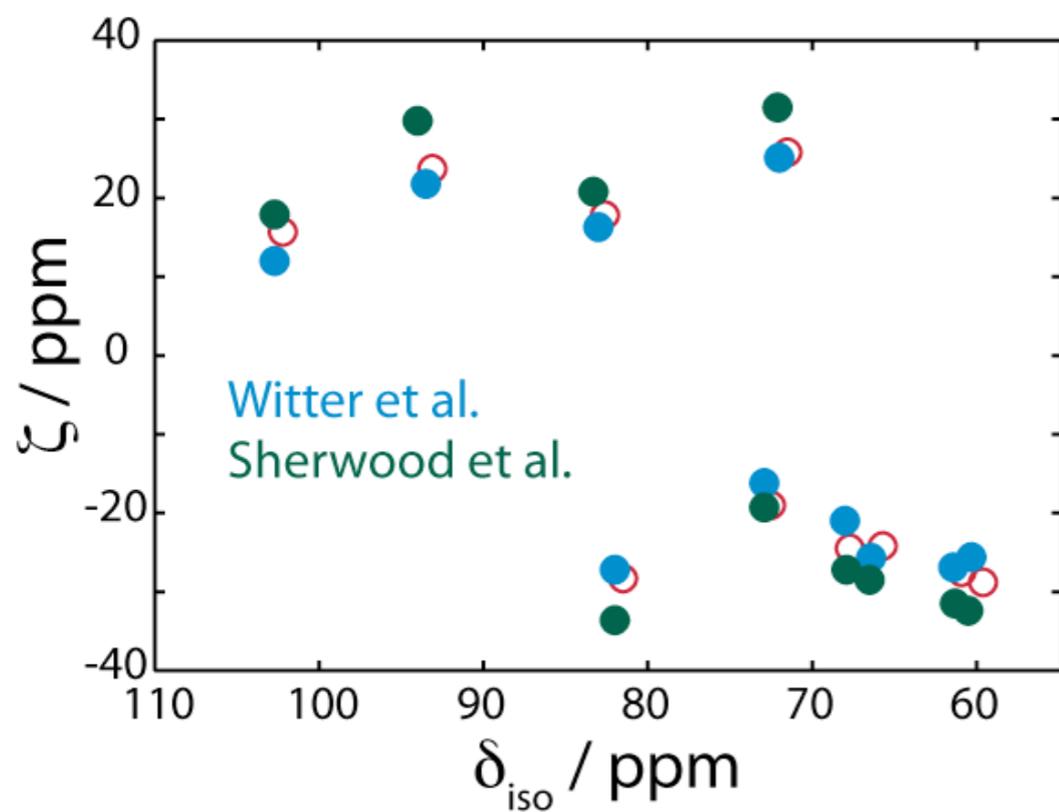
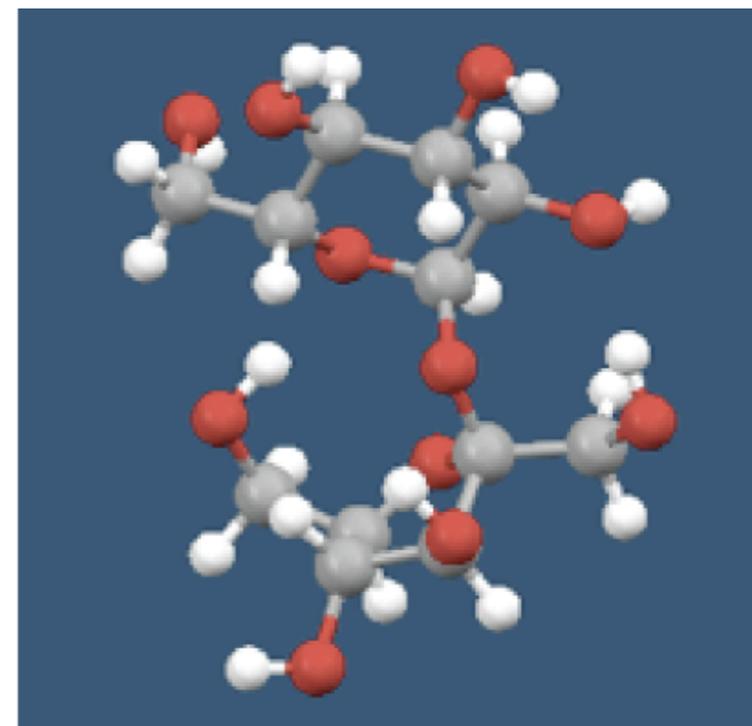
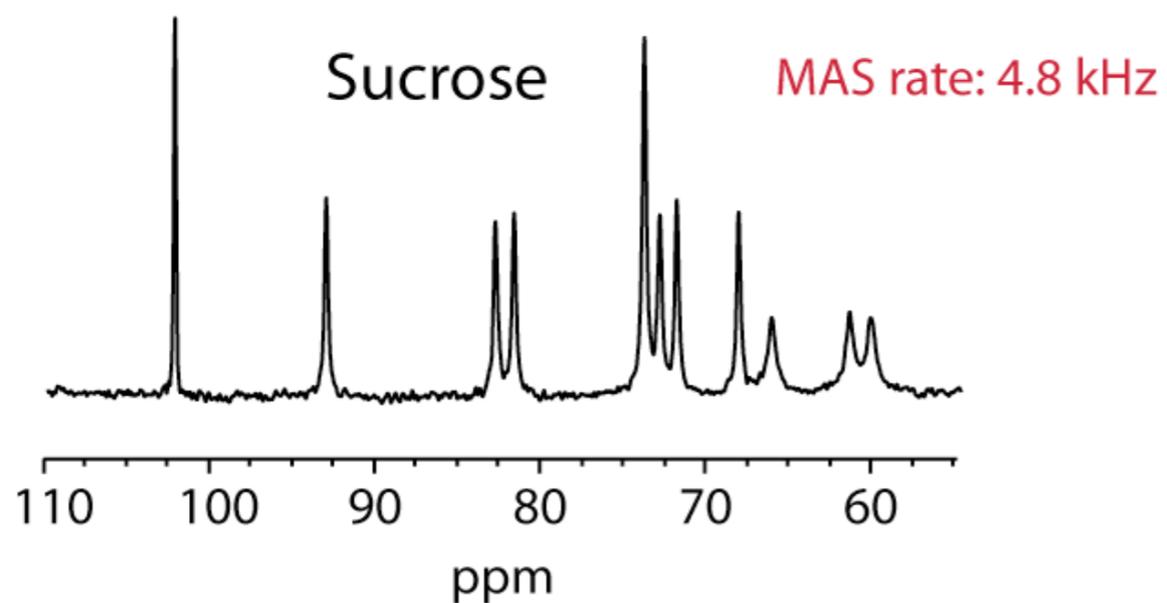


# Shift anisotropies by CSA amplification

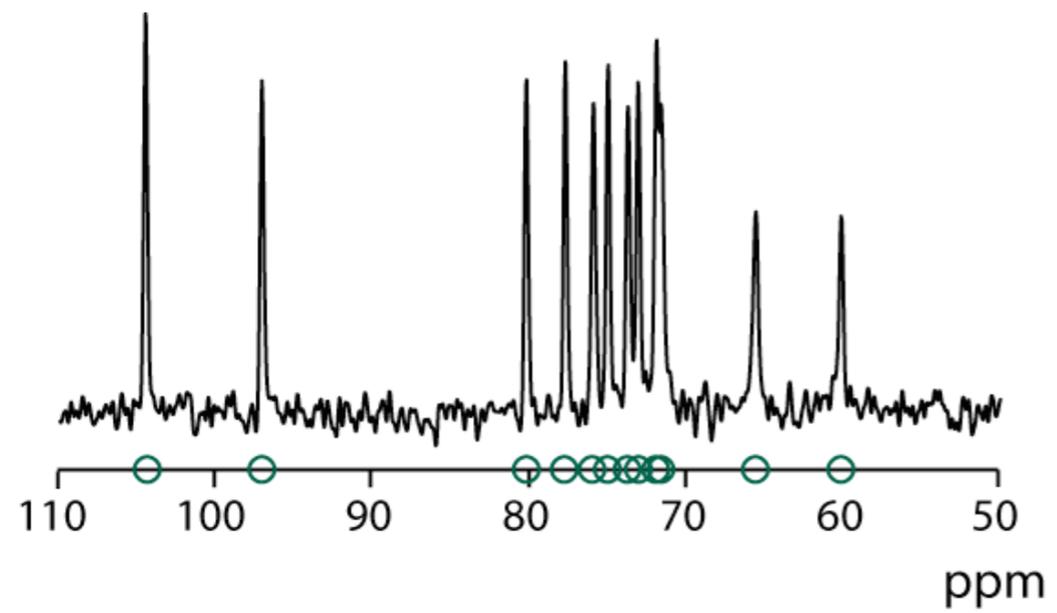


amplification: 8  
effective rate: 600 Hz

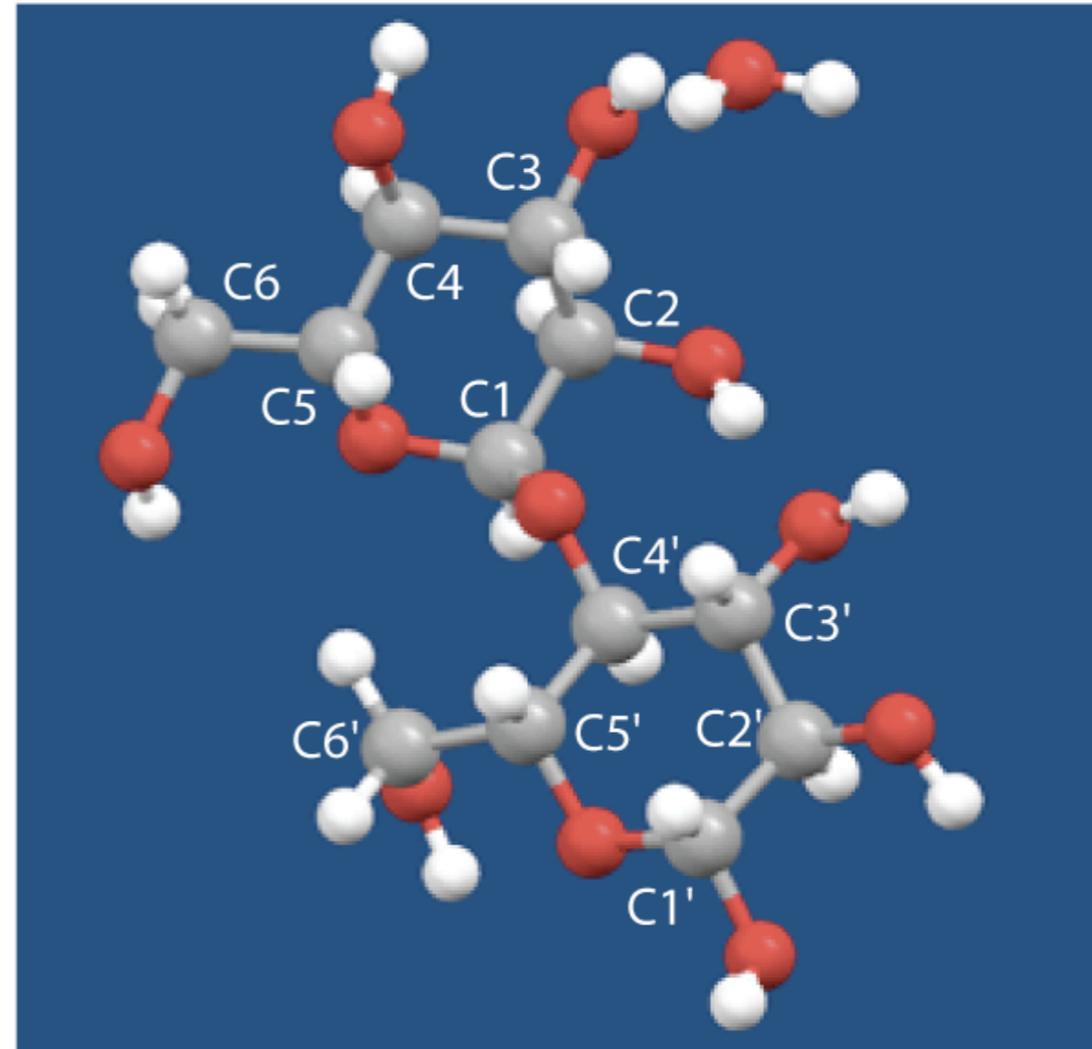
# Validation of the CSA amplification method



# Maltose

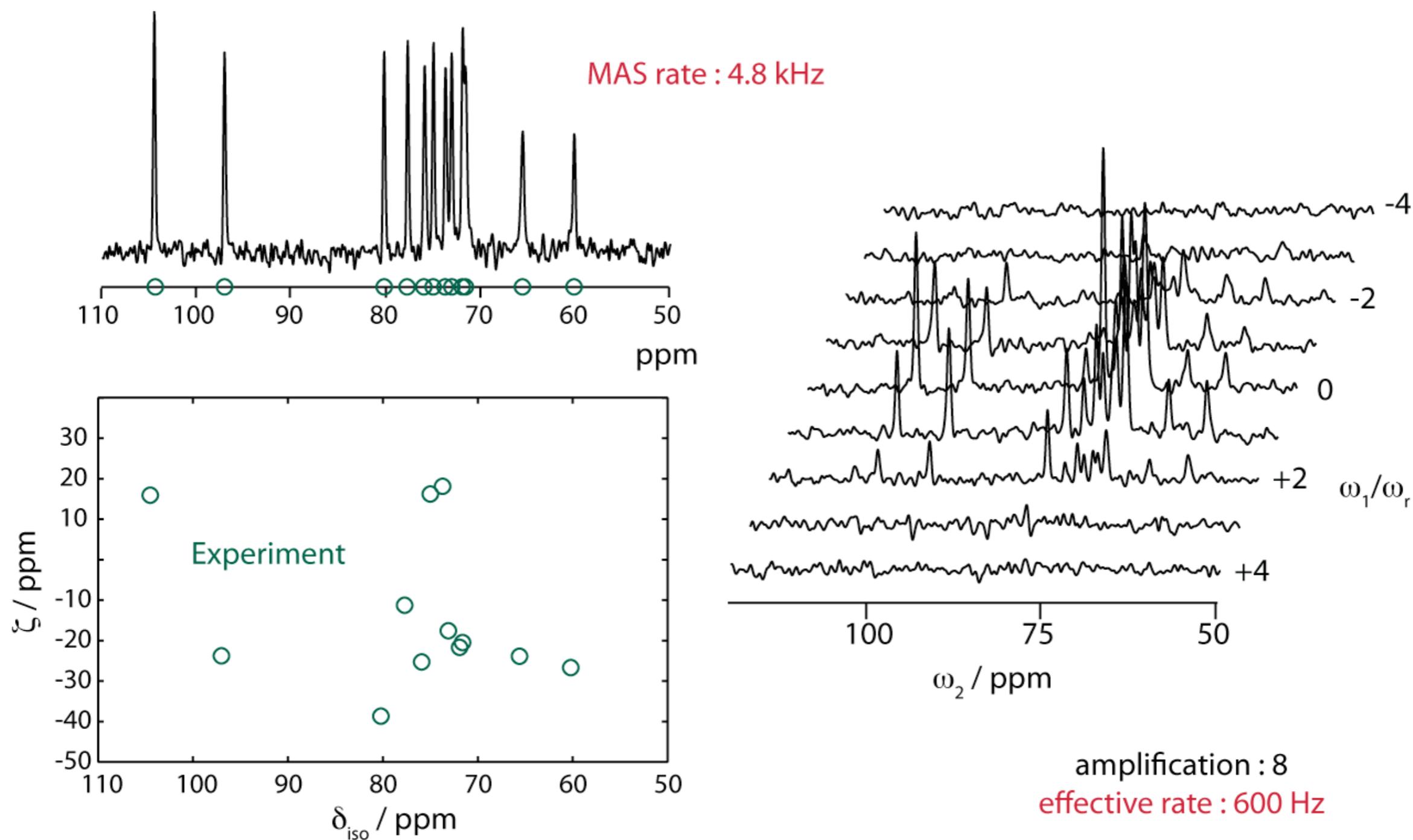


MAS rate : 4.8 kHz



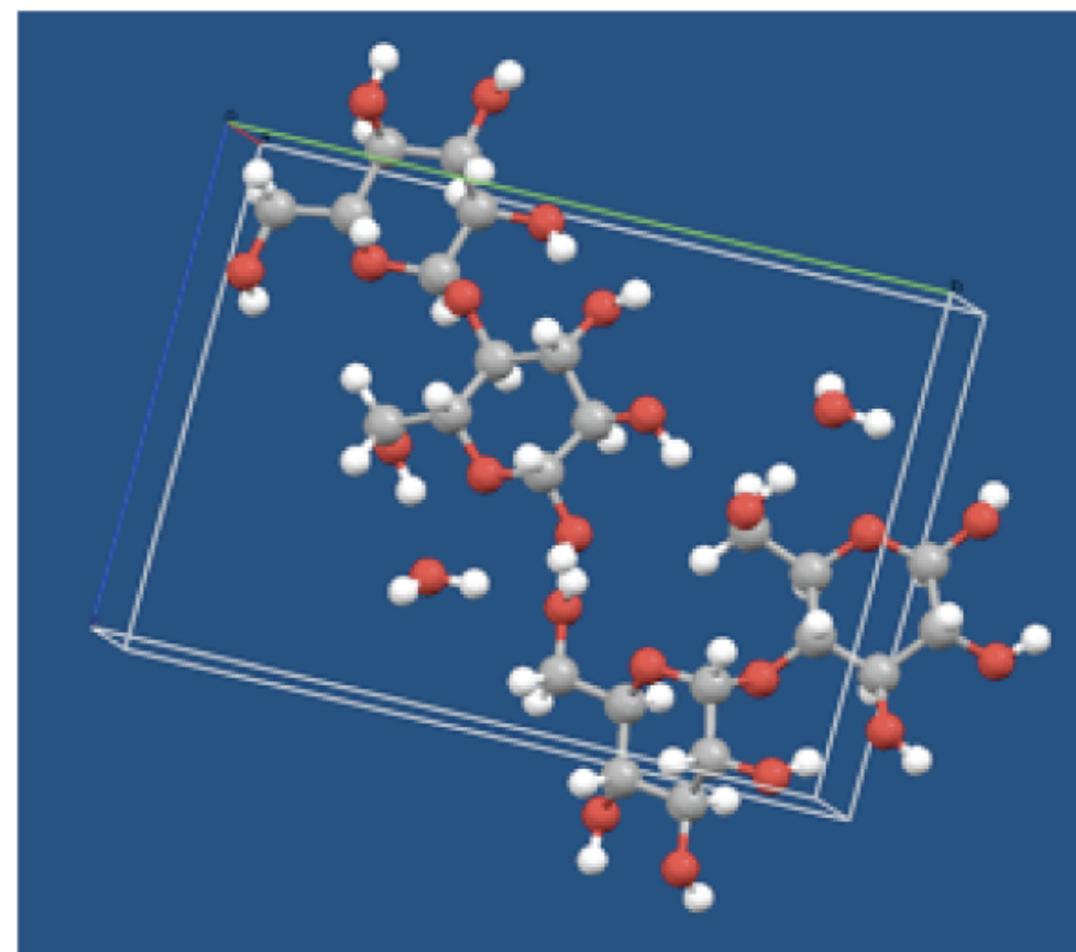
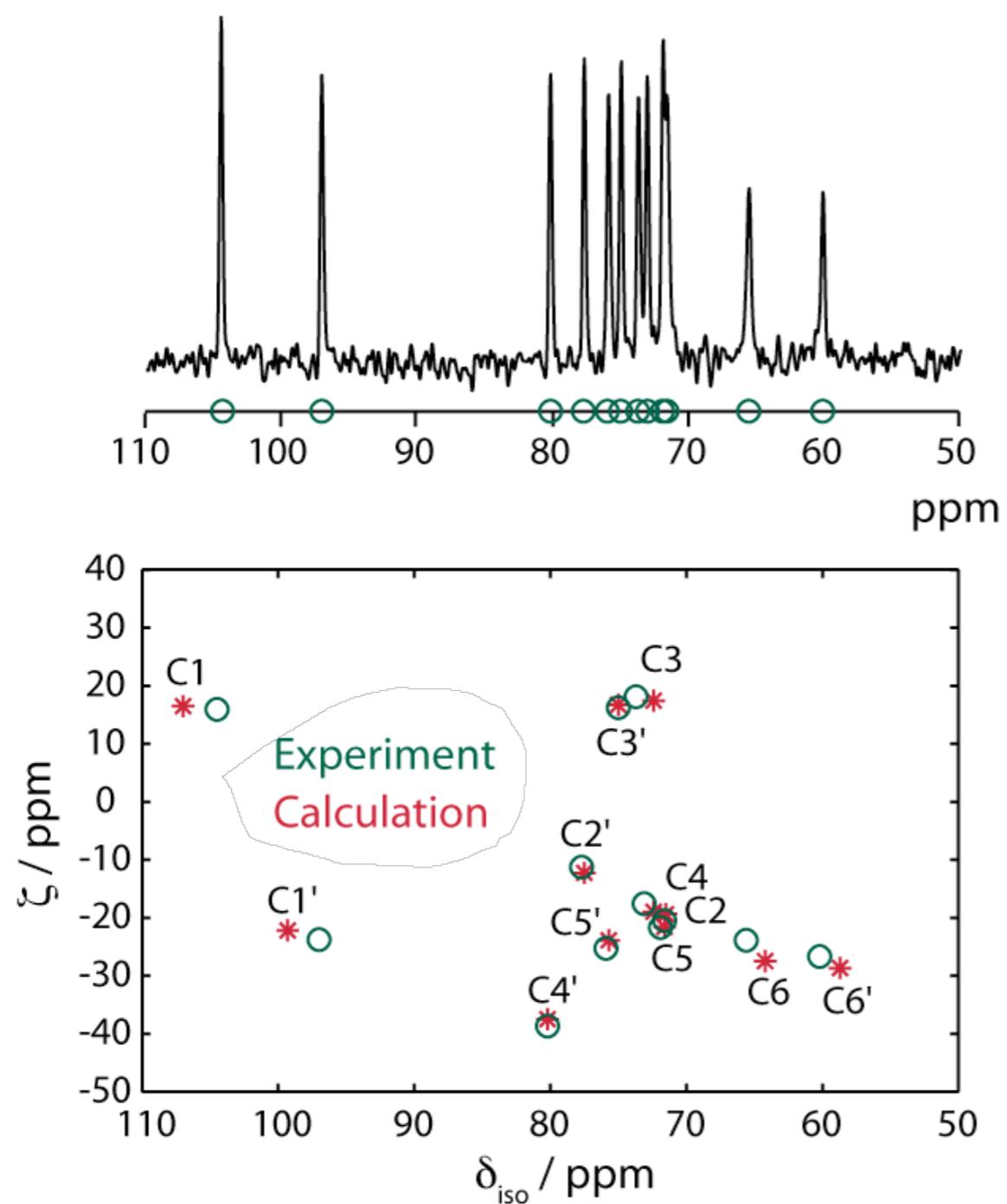
Poor shift dispersion in saccharides prevents assignment of carbon-13 spectrum

# Shift anisotropies by CSA amplification



Correlation of two shift parameters gives better resolution

# Maltose: calculations and assignment



CASTEP/GIPAW

Combination with calculations allows assignment

# Summary

Two-dimensional **separation** experiments show spectra in the two dimensions which result from two different Hamiltonians.

**Isotropic/anisotropic** separation experiments are used to obtain the shift anisotropies of complicated molecules where **overlap** of powder patterns or sideband manifolds from different chemical sites prevents their measurement using a one-dimensional spectrum.

**Separated local field** experiments are used to establish the **orientation** of the principal axis system of the chemical shift tensor relative to the molecule.