

Supporting Information

Determination of Relative Tensor Orientations by γ -encoded Chemical Shift Anisotropy/Heteronuclear Dipolar Coupling 3D NMR Spectroscopy in Biological Solids

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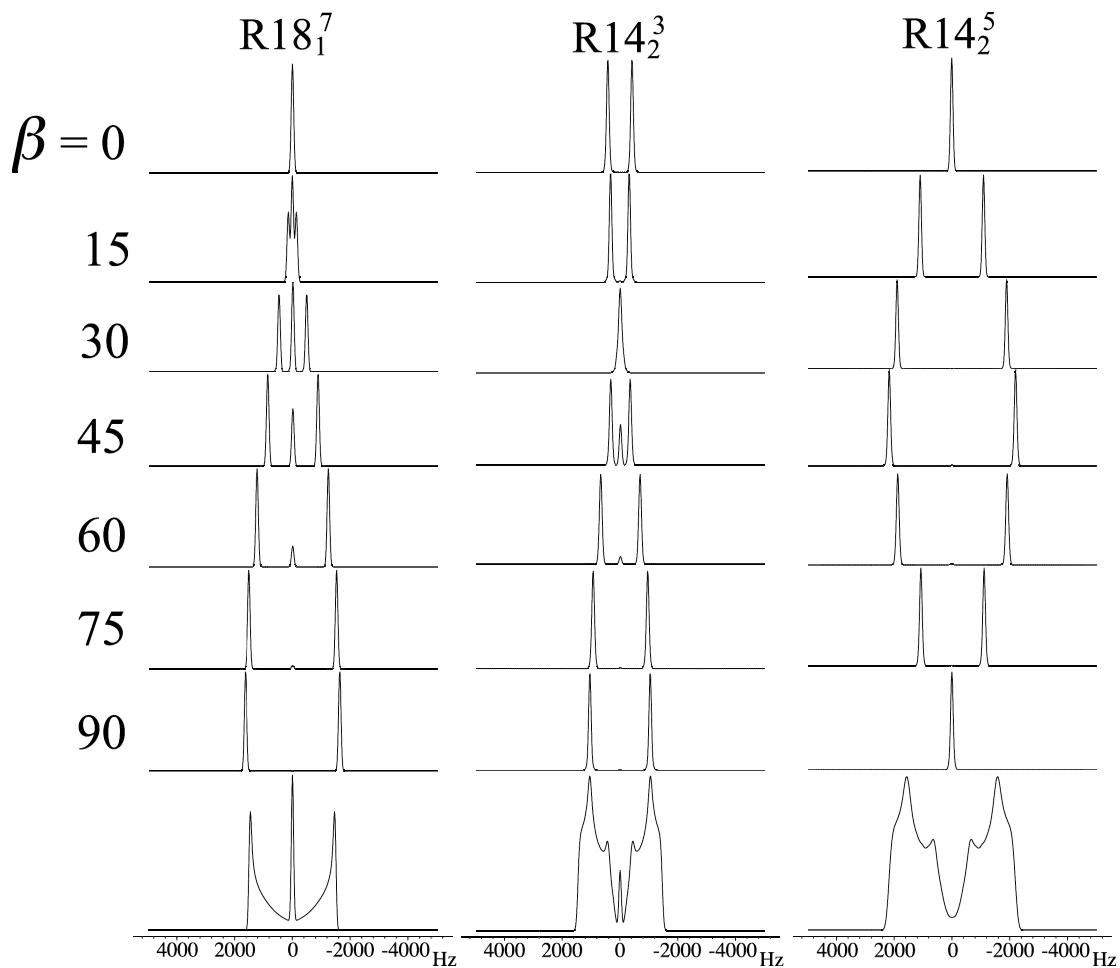


Figure S1. The recoupled ¹H-¹⁵N dipolar and ¹⁵N CSA lineshapes for different crystallite orientations in the laboratory frame produced by γ -encoded R18₁⁷ and R14₂^{3/5}. Two-spin system (¹H, ¹⁵N) is used in all simulations. The recoupled powder patterns are illustrated in the bottom figure; the powder averaged were generated using 986 ZCW crystallite orientations (α , β). As shown in the Figure, there is only one splitting corresponding to each crystallite and it is independent of the number of gamma angles. This demonstrates that R18₁⁷ and R14₂^{3/5} recoupling techniques are γ -encoded.

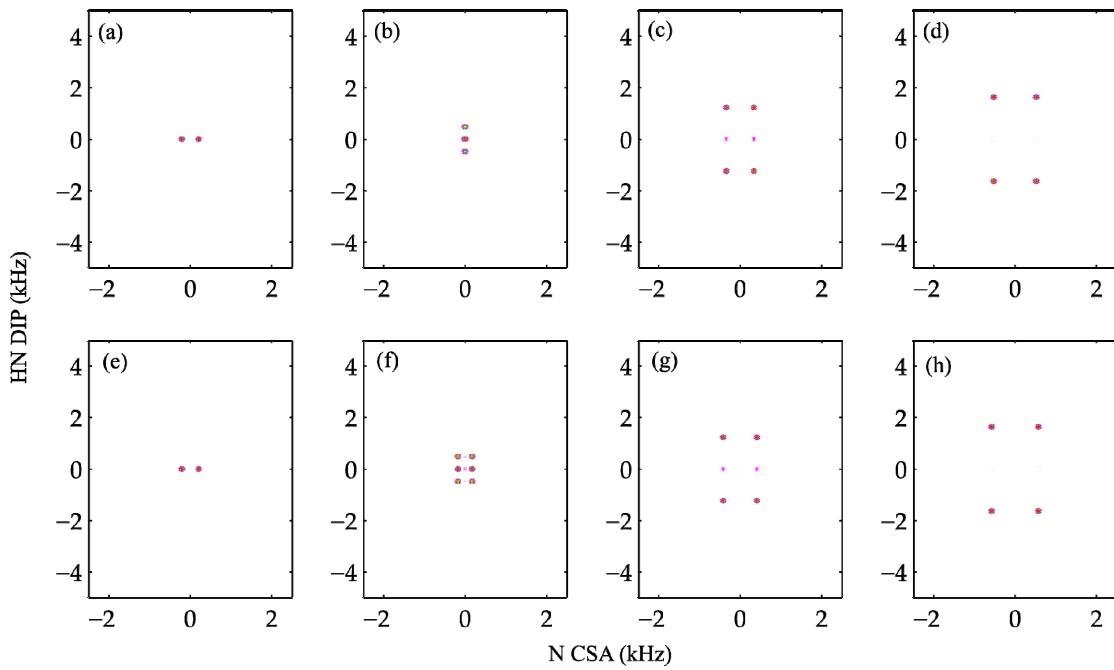


Figure S2. The simulated ^1H - ^{15}N dipolar and ^{15}N CSA correlation spectra by the combined R18₁⁷/R14₂³ symmetry scheme for different crystallite orientations. The same DIP/CSA relative orientation was used for all simulations. The crystallite orientations expressed in terms of the Euler angles are (a) $\alpha = 0^\circ$, $\beta = 0^\circ$; (b) $\alpha = 0^\circ$, $\beta = 30^\circ$; (c) $\alpha = 0^\circ$, $\beta = 60^\circ$; (d) $\alpha = 0^\circ$, $\beta = 90^\circ$; (e) $\alpha = 30^\circ$, $\beta = 0^\circ$; (f) $\alpha = 30^\circ$, $\beta = 30^\circ$; (g) $\alpha = 30^\circ$, $\beta = 60^\circ$; (h) $\alpha = 30^\circ$, $\beta = 90^\circ$. It can be seen that the combined R-type symmetry scheme shows high sensitivity to the molecular orientation. Only one splitting can be found along DIP and CSA dimension for each crystallite orientation.

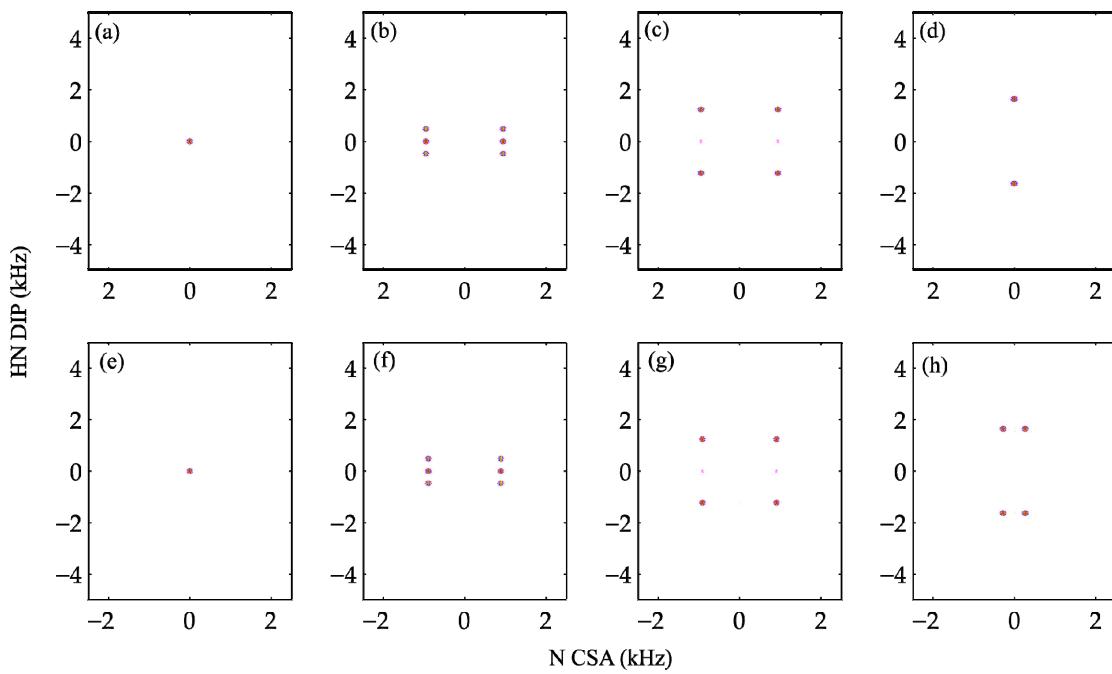


Figure S3. The simulated ^1H - ^{15}N dipolar and ^{15}N CSA correlation spectra by the combined $\text{R}18_1^7/\text{R}14_2^5$ symmetry scheme for different crystallite orientations. The same DIP/CSA relative orientation was used for all simulations. The crystallite orientations expressed in terms of the Euler angles are (a) $\alpha = 0^\circ$, $\beta = 0^\circ$; (b) $\alpha = 0^\circ$, $\beta = 30^\circ$; (c) $\alpha = 0^\circ$, $\beta = 60^\circ$; (d) $\alpha = 0^\circ$, $\beta = 90^\circ$; (e) $\alpha = 30^\circ$, $\beta = 0^\circ$; (f) $\alpha = 30^\circ$, $\beta = 30^\circ$; (g) $\alpha = 30^\circ$, $\beta = 60^\circ$; (h) $\alpha = 30^\circ$, $\beta = 90^\circ$. As with the $\text{R}18_1^7/\text{R}14_2^3$ symmetry scheme in Figure S2, the combined $\text{R}18_1^7/\text{R}14_2^5$ symmetry scheme also shows high sensitivity to the molecular orientation.

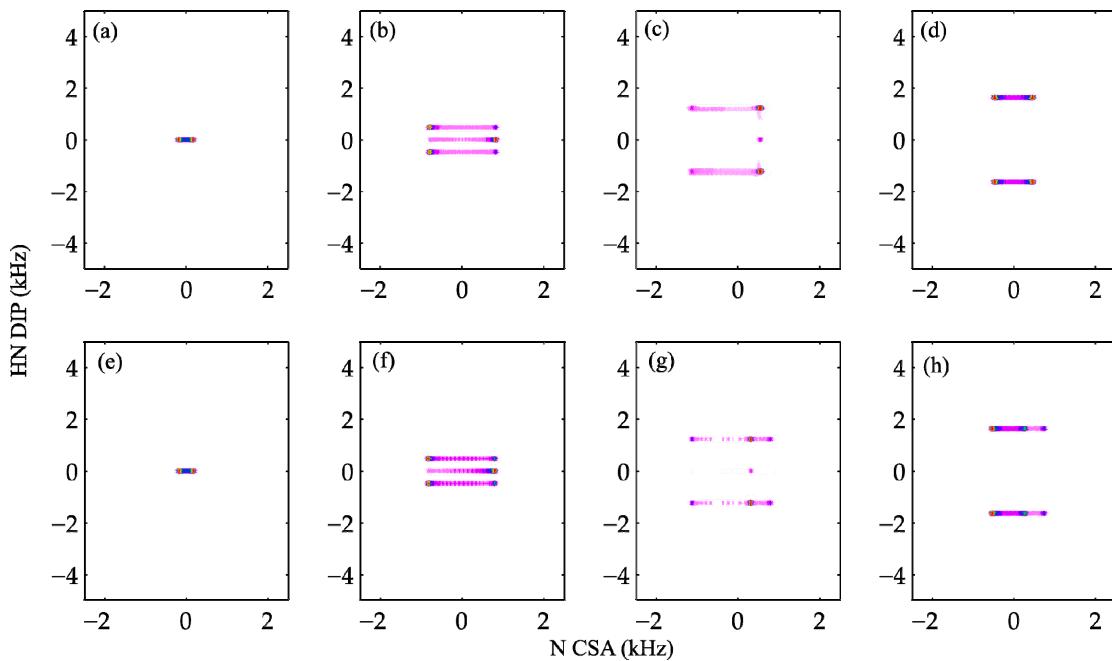


Figure S4. The simulated ^1H - ^{15}N dipolar and ^{15}N CSA correlation spectra by the combined γ -encoded R18₁⁷ / non γ -encoded ROCSA¹ scheme for different crystallite orientations. The same DIP/CSA relative orientation was used for all the simulations. The crystallite orientations expressed in terms of the Euler angles are (a) $\alpha = 0^\circ$, $\beta = 0^\circ$; (b) $\alpha = 0^\circ$, $\beta = 30^\circ$; (c) $\alpha = 0^\circ$, $\beta = 60^\circ$; (d) $\alpha = 0^\circ$, $\beta = 90^\circ$; (e) $\alpha = 30^\circ$, $\beta = 0^\circ$; (f) $\alpha = 30^\circ$, $\beta = 30^\circ$; (g) $\alpha = 30^\circ$, $\beta = 60^\circ$; (h) $\alpha = 30^\circ$, $\beta = 90^\circ$. Compared with the R-symmetry based schemes, the non γ -encoded ROCSA sequence results in broad powder pattern along the CSA dimension for each orientation (α , β). The lineshape of the DIP/CSA powder pattern also depends on the relative orientation, but the sensitivity is much lower than in the R-symmetry schemes illustrated in figures S2 and S3. It is therefore necessary to employ γ -encoded recoupling scheme for determination of the CSA tensor orientation in the molecular frame.

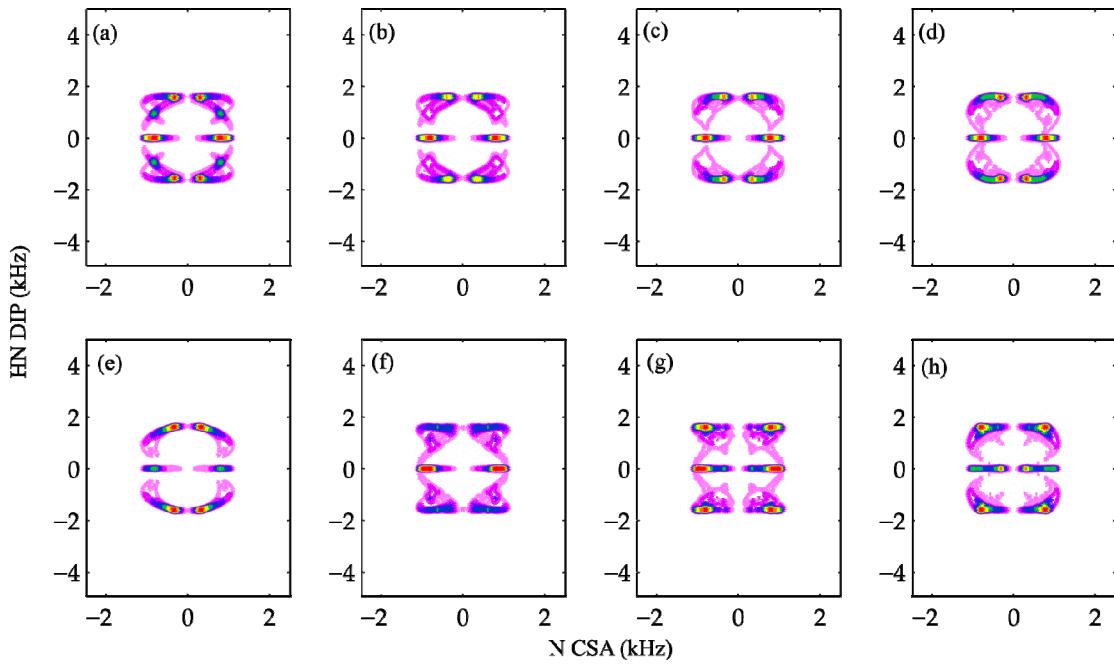


Figure S5. The simulated ^1H - ^{15}N dipolar and ^{15}N CSA correlation spectra by the combined γ -encoded R18₁⁷/R14₂⁵ scheme for different relative tensor orientations. In a)-d), the Euler angle β_{NH} (the angle between the N-H vector and the CSA δ_{zz} principal value) was 20°, and the Euler angle α_{NH} (the angle between Cα-N-H peptide plane and the CSA δ_{xx} principal value) was 0°, 30°, 60°, and 90°, respectively. In e)-h), the Euler angle α_{NH} was 20°, and β_{NH} was 0°, 30°, 60°, and 90°, respectively. This demonstrates that both Euler angles α_{NH} and β_{NH} affect the 2D correlation spectrum patterns, but the effect of α_{NH} on the lineshape is much less pronounced.

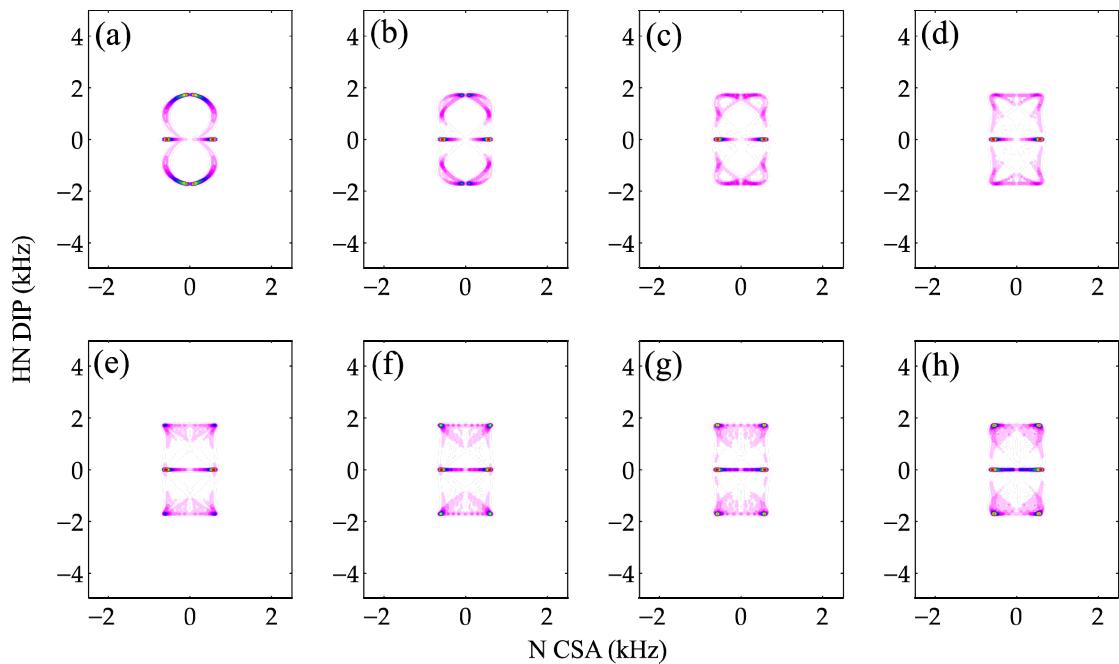


Figure S6. Simulated R18₁/R14₂ CSA/DIP spectra for different relative tensor orientations. In these simulations, the δ_{xx} principal value CSA is considered to lie in the C α -N-H peptide plane ($\alpha_{\text{NH}} = 0^\circ$), and the angle β_{NH} between the H-N vector and the CSA δ_{zz} principal value was changed from 0 to 70 degrees with 10-degree increments from (a) to (h), respectively. 64 t_1 increments and 128 t_2 increments were used for each simulation. A powder average was generated with 986 ZCW angles and 3 γ angles.

References:

1. Chan, J. C. C.; Tycko, R., *J. Chem. Phys.*, **2003**, *118*, 8378-8389.