

Unveiling Salt-Bridge Interactions Between GAGs and Collagen Protein in Cartilage by DNP-Enhanced ssNMR Spectroscopy

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The extracellular cartilage matrix (ECM) is a complex biological structure, consisting of highly mobile glycosaminoglycans (GAGs) alongside rigid collagen proteins. The interaction between GAGs and collagen is crucial for cartilage stability, yet direct evidence of these interactions in their native state has not been observed.

To address this, we applied novel magic-angle spinning/dynamic nuclear polarization (MAS-DNP) solid-state NMR (ssNMR) methodologies to study the ECM in its natural ¹³C-abundance state. For the first time, these approaches enabled the direct observation of dipolar-coupled ¹⁵N-¹³C and SQ-SQ ¹³C-¹³C interactions in 2D correlation spectra—previously unattainable in natural-abundance ¹³C/¹⁵N samples.

¹⁵N-¹³C 2D ssNMR experiments (Fig. 1, left) provide the first atomic-level confirmation of direct bonding interactions between hydroxyproline (Hyp) in collagen and GAGs, which contributes to the structural stability of the cartilage extracellular matrix. ¹³C-¹³C 2D correlation experiments (Fig. 1, right) identified ¹³C-¹³C correlations between GAGs and collagen at distances of up to 4–5 Å, revealing direct molecular interactions, including charge-pair salt-bridge interactions between the sulfate groups of GAGs and the guanidinium groups of arginine in collagen.

This approach has provided a powerful new method for characterization of biological systems without isotopic enrichment.

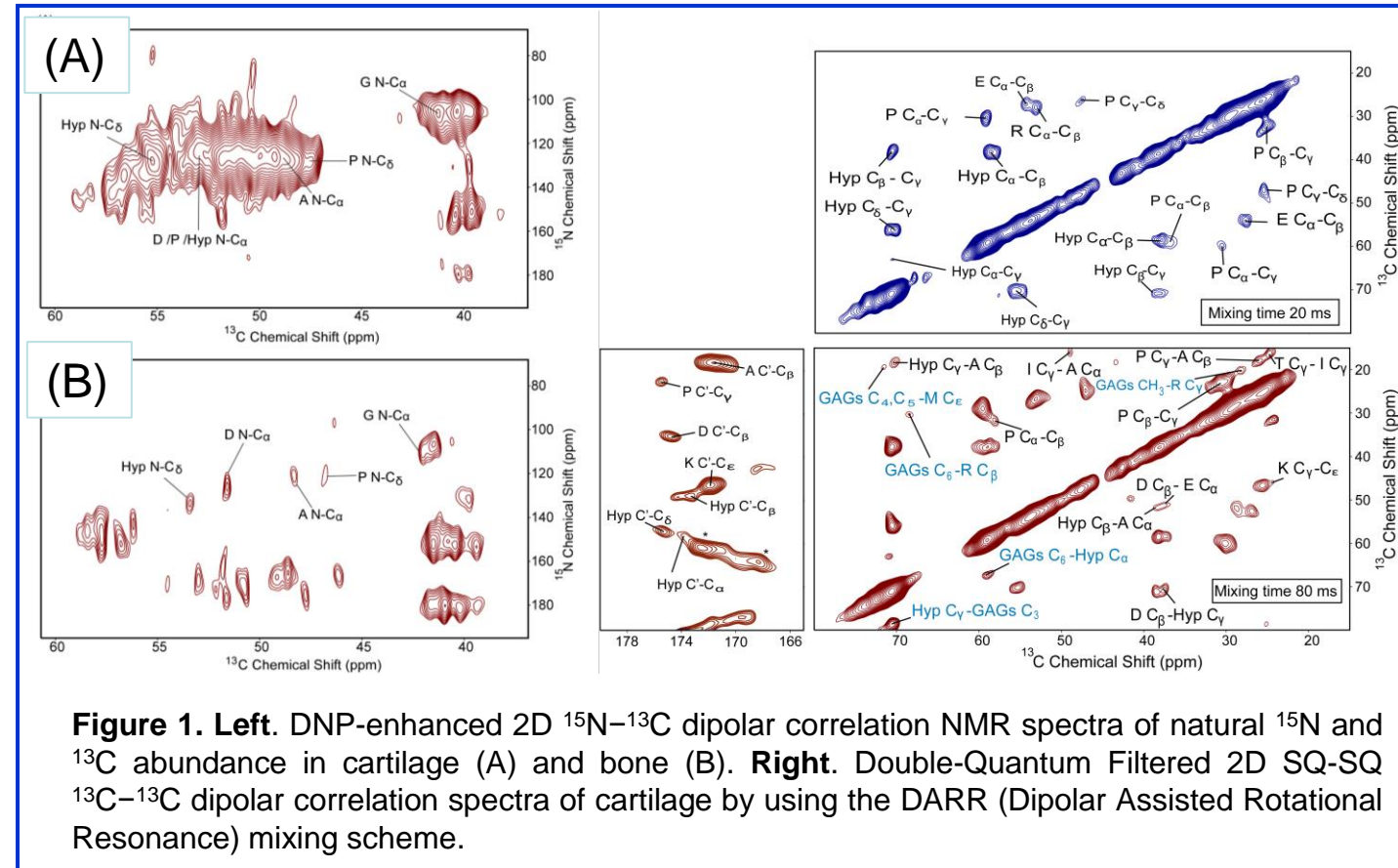


Figure 1. Left. DNP-enhanced 2D ¹⁵N-¹³C dipolar correlation NMR spectra of natural ¹⁵N and ¹³C abundance in cartilage (A) and bone (B). Right. Double-Quantum Filtered 2D SQ-SQ ¹³C-¹³C dipolar correlation spectra of cartilage by using the DARR (Dipolar Assisted Rotational Resonance) mixing scheme.

Facilities and instrumentation used: Solid-State NMR Facility - Wide-bore 600 MHz Bruker Avance III spectrometer configured with a 395 GHz gyrotron microwave source, microwave quasi-optical table, and a Bruker 3.2 mm MAS DNP ¹H-X-Y triple-resonance probe.

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