

Assessment of Degradation of Proteoglycans and Matrix Proteins in Intervertebral Disc Degeneration by Diffusion Weighted MRS

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INTRODUCTION: Intervertebral disc degeneration (IVDD) is considered to be the root cause of back pain. It is a common disease of high morbidity in aging population (1-3). Currently, correlation between MRI and CT morphologic findings and patient symptoms in disc degenerative disease was poor (4). Previous high-resolution HR-MAS studies have revealed the relationship between spectroscopy derived metabolite concentrations in 3.5-4.2ppm, namely the carbohydrate (Carb) region and levels of disc degeneration (2,4). However, in clinical MRS study, the quantification of Carb region resonances is technically challenging due to the relatively poor shimming in vivo and broad spectral overlap by water peak (3). In this preliminary study, we hypothesize that diffusion of water is much faster than that of macromolecules which generate the Carb region resonances, thus we propose to suppress the water peak contamination by diffusion weighting. Furthermore, we hypothesize that diffusion characterization of macromolecules resonances including Carb region resonances can provide a sensitive marker for IVDD.

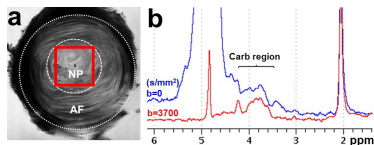


Fig.1 (a) Axial T2-weighted image of a normal bovine disc (NP, nucleus pulposus; AF, annulus fibrosus; VOI selection shown as the red rectangle) (b) Diffusion weighting (DW) suppresses the large overlapping water contamination and allows the reliable measurement of Carb region (3.5-4.2ppm) resonances in a normal disc.

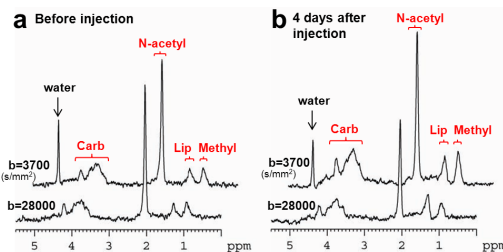


Fig.2 Representative DW spectra acquired from a bovine disc before (a) and 4 days after papain injection (b). Four major resonances were observed and assigned as Carb region (3.5-4.2ppm): C-H resonances of GAG, ethanolamine and glycine; N-acetyl region (1.9-2.1ppm): N-acetyl group of GAG, proline and glutamate; Lip region (1.2-1.4ppm): aliphatic chain of lipid and lactate; Methyl region (0.8-1.0ppm): isoleucine, leucine and valine (2,4,5).

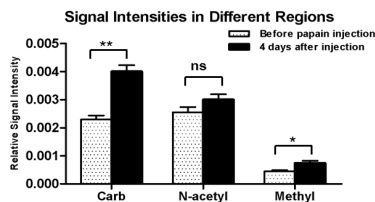


Fig.3 Signal intensities in different regions (N=6, normalized by water signal) before and after papain injection.

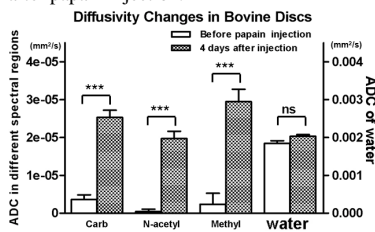


Fig.4 Diffusivity changes in bovine discs before and after papain injection (N=6).

regions were below 10^{-5} mm²/s while ADC in N-acetyl region was even smaller. However, 4 days after papain injection, ADC values in all these three regions increased by more than ten folds ($p < 0.0001$) (Fig. 4), indicating that these macromolecules now became significantly more mobile. In the same discs, water ADCs were also measured and seen to increase slightly from $1.90 \pm 0.11 \times 10^{-3}$ mm²/s to $2.03 \pm 0.10 \times 10^{-3}$ mm²/s after papain injection. Note that this water diffusion increase was comparable to the finding in a previous human study (6).

DISCUSSIONS AND CONCLUSION: Previous biochemistry study indicated that degradation proteoglycans (PGs) of plays a major role in IVDD progression, which is the main cause of disc dehydration and reduction of osmotic pressure in disc matrix (1,7). Papain digestion is a widely used model of disc degeneration in IVDD research because papain is a relatively non-specific protease that can break down both PGs and the disc matrix proteins (3,7). In this preliminary study, the resonances in Carb and Methyl region increased substantially after papain injection, which was in agreement with the earlier HR-MAS findings showing that the number and intensity of free amino acids resonances increased in the degenerated human discs (2,4). A slight signal intensity increase was also observed in N-acetyl region, but not as notable as Carb and Methyl regions. This observation may arise from the fact that the resonance intensity of amino acids (proline, glutamate) in 1.9-2.1 ppm was low, thus difficult to yield statistically significant difference (Fig. 2). More importantly, apparent diffusion coefficients (ADCs) of Carb, N-acetyl and Methyl groups were observed to increase drastically after papain injection, indicating the great potential of these macromolecular diffusion measurements as robust and sensitive markers in detecting disc degeneration. We reason that the byproducts of disc degeneration, including both amino acids and degraded glycosaminoglycans (GAGs), had much smaller molecular weights than PGs and matrix proteins. At the same time, GAGs was known to be highly hydrophilic. Once no longer anchored within the disc matrix, GAGs dissolved into the water, thus exhibiting greatly increased mobility. These factors together led to the drastic ADC increases observed in this study. In conclusion, diffusion weighting provides an effective method to separate and quantify carbohydrate resonances by suppressing the large overlapping water peak. More importantly, DW-MRS provides a new and clinically translatable approach to probe the biophysical processes in IVDD. In particular, MR characterization of carbohydrate, N-acetyl and Methyl macromolecular groups holds great potential to sensitively and quantitatively detect and assess the degradation of PGs and matrix proteins during disc degeneration. Future studies will include the longitudinal examination of papain injection and other disc degeneration models, optimization of DW-MRS sensitivity, and clinical implementation and assessment.

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