

# Developing Protein-based MRI Biomarkers for Alzheimer's Disease

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## INTRODUCTION:

The hallmarks of Alzheimer's disease (AD) are the gradual accumulation and aggregation of toxic soluble and insoluble beta-amyloid (A $\beta$ ) and tau species in the brain. Amide proton transfer (APT) imaging is a relatively new protein-based molecular MRI technique that is based on endogenous mobile proteins and peptides in tissue. It is known that both extracellular A $\beta$  and intracellular tau first exist as soluble monomers and oligomers,<sup>1,2</sup> which are APT-detectable. This study was performed to examine whether APT signal can be a surrogate biomarker for early AD diagnosis.

## METHODS:

MRI experiments were carried on a Phillips 3T MRI scanner. Coronal 3D APT images (RF saturation power = 1.5  $\mu$ T; saturation time = 1.5 sec; 24 offsets; 15 slices) were acquired. WASSR images were acquired for correcting B0 inhomogeneity. The conventional APT-weighted image is defined as:  $APT_w = MTR_{asym}(3.5ppm) = [S_{sat}(-3.5ppm) - S_{sat}(+3.5ppm)] / S_0 = Z_{exp}(-3.5ppm) - Z_{exp}(+3.5ppm)$ . Based on the extrapolated semi-solid magnetization transfer reference (EMR) approach, the Z-spectrum data of seven large, positive offsets (80~8 ppm) were fitted with a super-Lorentzian line shape, leading to a reference curve ( $Z_{EMR}$ ). Then,  $APT^\#$  and  $NOE^\#$  were calculated:  $APT^\# = Z_{EMR}(3.5ppm) - Z_{exp}(3.5ppm)$  and  $NOE^\# = Z_{EMR}(-3.5ppm) - Z_{exp}(-3.5ppm)$ . T1 and T2 maps were also acquired. ROIs (hippocampus) were manually drawn on both sides. The mean values of T1, T2,  $APT_w$ ,  $APT^\#$ ,  $NOE^\#$  and MTR at 20ppm in the ROIs were counted. Group-based analysis of these values were performed.

## RESULTS:

31 participants were included, 23 cognitively normal controls (CN), 5 mild cognitive impairment (MCI) and 3 AD. MCI had higher  $APT^\#$  signal intensity values than CN in multiple brain regions while AD had even higher values (in hippocampus,  $p = 0.11$  and  $< 0.01$ , respectively, based on the limited sample size).  $APT_w$  values were smaller than  $APT^\#$ , but there was a similar trend. This may be attributed to the abnormal accumulation of various mobile proteins, including soluble A $\beta$  and tau oligomers, with the progress of AD. Besides,  $APT^\#$  signal values showed a correlation coefficient of -0.524 with the mini mental state examination (MMSE) score and 0.496 with the Clinical dementia rating (CDR). Patients with more severe cognitive impairment tended to have higher  $APT^\#$  values. Inclusion of more MCI and AD cases and correlation analysis with amyloid and tau biomarkers in CSF are needed to further validate the results.

Based on the definition,  $APT_w = APT^\# - NOE^\#$ . This suggests that the EMR-APT approach can achieve purer and higher APT signals than simply using the conventional  $APT_w$  metric.<sup>3,4</sup> In addition, our recent study demonstrated that the deep-learning-based EMR method achieved high reproducibility and reliability in the quantification.<sup>5</sup>

## DISCUSSION AND CONCLUSION:

The early results showed that APT imaging may detect an increased signal that may reflect soluble A $\beta$  and tau proteins in MCI and AD and have the potential to track decreases in these abnormal protein levels during anti-amyloid immunotherapy against AD.

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