

## Comparison between 7T T2\* and 3T MTR in the *in vivo* human cortex.

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**Purpose.** Characterization of cortical myelo-architecture with MRI is an active field of research<sup>1</sup>, which can give insights on the structural and functional organization of the brain. It is however challenging to image the cortex due to its convoluted and thin geometry (2-4mm). Recently, ultra-high field MRI (7T) combined with T2\* was shown to reveal features of myelin density<sup>2</sup>. However, several confounds hamper the specificity of T2\* measures such as iron content and blood vessels<sup>3</sup>. An independent measure with different contrast mechanisms would increase the specificity to myelin. Magnetization Transfer Ratio (MTR) imaging at 3T was shown to be sensitive to myelin content<sup>4</sup> and thus would be an excellent complementary measure. The goal of this study was to evaluate the relationship between T2\* at 7T and MTR at 3T, and show their respective sensitivity and specificity to myelin content.

**Methods.** *Data acquisition.* Healthy subjects (N=6, age = 36 +/- 5 years) were recruited and scanned with a 7T whole-body scanner (Siemens Healthcare, Erlangen, Germany) to measure T2\* and with a 3T scanner (Siemens TIM Trio) to measure MTR. Both scanners were equipped with a 32-channel coil. Parameters at 7T were: TR = 2020ms, TE = 6.34+3.2n [n=1...12], resolution = 0.33x0.33x1mm3. Parameters at 3T were: 3D FLASH, TR/TE = 30/2.49ms, matrix = 192x192, resolution=1.2x1.2x1.2 mm3, with and without Gaussian MT pulse (7:45min each). *Data processing.* T2\* and MTR data were registered to individual cortical surfaces, sampled at the mid-cortical distance and registered to a common template surface<sup>5</sup>. Data were first averaged in the common space and SD maps were computed to assess inter-subject variability. A linear regression between the mean T2\* and MTR maps was performed for each hemisphere, as well as within Brodmann regions with different myelin content.

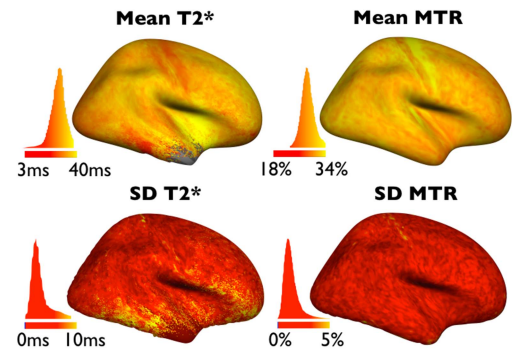
**Results.** Average and SD maps of T2\* and MTR are shown in **Fig. 1**. SD map of T2\* show high variability in the lower brain, likely due to poor shimming in this region. Conversely, SD of MTR shows fairly good reproducibility (mean SD = 1.59%). **Fig. 2** shows the relationship between T2\* and MTR. Strong correlations in the right (r=-0.77) and left (r=-0.75) hemispheres were detected. To verify if partial volume effect affected our measures, cortical thickness was correlated with these measures and showed low effect (r=0.14 and r=-0.09 for T2\* and MTR, respectively). **Fig. 3** shows the mean values of T2\* and MTR for the Brodmann regions (B1, B2, B3, B4, B43 and B44), sorted by T2\* values. Once again, T2\* and MTR are highly anti-correlated.

**Discussions:** Our results show within the same subjects an increase of T2\* and a decrease of MTR, in regions that are known to be heavily myelinated (e.g., B4, B1). These trends were expected given the sensitivity of T2\* and MTR to myelin content. However, this is the first time these two metrics are combined within the same subjects, providing a framework to isolate confounding parameters affecting T2\* (iron, issue orientation, poor shimming) and MTR (B1 inhomogeneities, T1). Combining other metrics (quantitative T1, diffusion, T2w/T1w) within the same methodological framework could potentially bring more insight into cyto- and myeloarchitecture than if these metrics were studied separately.

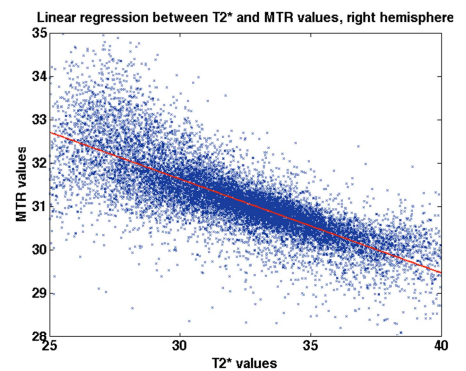
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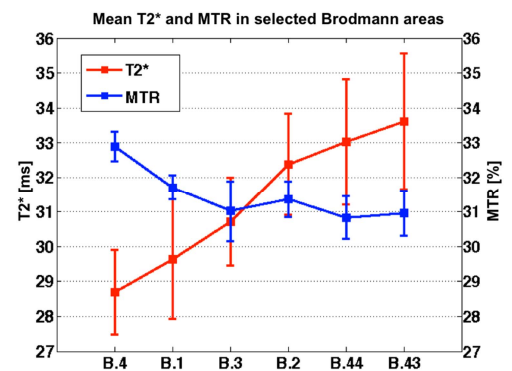
**Acknowledgments:** This work was supported by the National Multiple Sclerosis Society [FG 1892A1/1, 4281-RG-A-1], by the National Center for Research Resources [P41-RR14075], by the Fonds de Recherche du Québec - Santé (FRQS), by the Quebec BioImaging Network (QBIN) and by the Natural Sciences and Engineering Research Council of Canada (NSERC) and by the Polytechnique MEDITIS program.



**Fig. 1.** Mean and SD maps between the six controls for the T2\* metric (left) and the MTR values (right). Mean maps (top) shows an increase in MTR and a decrease in T2\* in the central sulcus.



**Fig. 2.** Linear regression between T2\* and MTR vertices values. The Pearson's correlation coefficient for this regression is r=-0.77.



**Fig. 3.** Mean values of T2\* and MTR across Brodmann regions with different myelin content. Data are averaged between the six controls. Error bars represent SD across subjects.