# Evidence of distributed subpial T2* signal changes at 7T in multiple sclerosis: an histogram based approach. 

C. Mainero ${ }^{1}$, C. Lima ${ }^{1}$, J. Cohen-Adad ${ }^{1}$, D. Greve $^{1}$, A. Radding ${ }^{1}$, T. Benner ${ }^{1}$, R. P. Kinkel ${ }^{2}$, B. Fischl ${ }^{1}$, and B. R. Rosen ${ }^{1}$

${ }^{1}$ A. A. Martinos Center for Biomedical Imaging, Massachusetts General Hospital, Charlestown, MA, United States, ${ }^{2}$ Neurology, Beth Israel Deaconess Medical Center, Boston, MA, United States

Introduction. Subpial lesions are the most frequent type of cortical lesion in multiple sclerosis (MS), and are thought to be closely associated with poor clinical outcome. Neuropathological studies report that subpial lesions may come in two major types: they may appear as circumscribed, focal lesions, or extend across multiple adjacent gyri leading to a phenomenon termed "general subpial demyelination" [1]. The in vivo evaluation of diffuse subpial disease is challenging - signal changes may be subtle, and extend across large regions where signal inhomogeneities due to $B_{1}$ and $R F$ receive coil non-uniformities become more pronounced. Here, we investigate whether a histogram-based analysis of $\mathrm{T}_{2}{ }^{*}$ signal intensity in the cortex, at 7 T MRI, can show evidence of distributed subpial cortical changes in patients with MS, as described histopathologically. We hypothesized that this phenomenon would be associated with significantly increased $T_{2}{ }^{*}$ signal intensity in patients compared to age-matched controls.

Methods. Fourteen MS patients (nine with relapsing-remitting MS, RRMS; five with secondary progressive MS, SPMS; mean $\pm$ SD age=38.9 $\pm 12.9$ years; median Expanded Disability Status Scale=3.0, range=1.0-6.5; mean $\pm$ SD disease duration=10.2 $\pm 7.7$ years) and eight age-matched controls were scanned twice on a human 7 T Siemens scanner using an in-house developed 8- or 32-channel phased array coil, and on a 3 Siemens Tim Trio scanner using the Siemens 32-channel coil. The 7 T protocol included acquisition of 2 D FLASH- $\mathrm{T}_{2}{ }^{*}$ spoiled gradient-echo weighted images (TR/TE=1000/22 ms, 20, $0.33 \times 0.33 \times 1 \mathrm{~mm}^{3}$ slices). For each modality two to three slabs were acquired, allowing coverage of the supratentorial brain. A 3D MPRAGE (TR/TE/TI=2600/3.26/1100ms, $0.60 \times 0.60 \times 1.5 \mathrm{~mm}^{3}$ slices) with the same orientation as the FLASH-T ${ }_{2}$ * scans was also acquired. During the $3 T$ session we acquired a high- structural 3 D scan with a magnetization-prepared rapid acquisition with multiple gradient echoes (MEMPR) sequence resolution $\left(0.9 \times 0.9 \times 0.9 \mathrm{~mm}^{3}, \mathrm{Tl}=1200 \mathrm{~ms}\right.$, $\mathrm{TR}=2530 \mathrm{~ms}$, flip angle $=7^{\circ}, \mathrm{TE}=1.7+n .1 .88 \mathrm{~ms}$ where $n=0, . ., 3$, FoV=230 mm , bandwidth $=651$ $\mathrm{Hz} / \mathrm{px}$ ). Prior performing the histogram quantitative analysis of $\mathrm{T}_{2}{ }^{*}$ signal in the cortex, all 7 T images were corrected for coil sensitivity profiles using a non-uniformity correction algorithm [2].

Pial and white matter (WM) surfaces generated by FreeSurfer on the 3T MEMPR (http://surfer.nmr.mgh.harvard.edu) were overlaid on the 7T FLASH-T ${ }_{2}{ }^{*}$ scans. Registration between FLASH-T ${ }_{2}{ }^{*}$ scans and the FreeSurfer anatomical reconstructed whole-brain 3T MEMPR was performed in several stages: 1) the whole-brain MPRAGE collected during the 7T session was registered to the FreeSurfer anatomical using the FSL FLIRT registration tool (www.fmrib.ox.ac.uk/fsl); 2) an initial registration between the partial FoV of the FLASH-T2* slabs and the 7T MPRAGE was computed from the geometry information found in the DICOM header; 3) a registration between the FreeSurfer anatomical and FLASH-T ${ }_{2}{ }^{*}$ slabs was performed by concatenating these volumes to the 7T MPRAGE. Because this is only accurate if the subject does not move, the final registration was computed based on the actual intensity values. For this purpose a new registration procedure, Boundary-Based-Registration (BBR), was used [3]. FLASH-T ${ }_{2}{ }^{*}$ partial volumes were then combined into a $0.33 \times 0.33 \times 0.33 \mathrm{~mm}^{3}$ single volume in the anatomical space. In places where the partial volumes overlap, they were averaged together. $T_{2}{ }^{*}$ intensities were normalized to the mean CSF intensity ( $T_{2}{ }^{*} / C S F$ ) and then sampled 1 mm inside the pial surface.

Histograms of $\mathrm{T}_{2}{ }^{*} / \mathrm{CSF}$ intensity in the subpial cortical volume across the whole right or left hemisphere, or in selected cortical regions were normalized by the total number of voxels included to correct for between-participant variability in brain volumes. For each histogram the following metrics were derived: a) Relative Peak Height (RPH), which measures the percent of voxels at the most common $T_{2}{ }^{*} / C S F$ value; b) Peak Position (PP), which measures the most common $T_{2}{ }^{*} / C S F$ value; c) $T_{2}{ }^{*} /$ CSF $_{25}, \mathrm{~T}_{2}{ }^{*} / \mathrm{CSF}_{50}, \mathrm{~T}_{2}{ }^{*} / \mathrm{CSF}_{75}$, which indicate the $\mathrm{T}_{2}{ }^{*} / \mathrm{CSF}$ at which the respective integrals of the histograms are $25 \%, 50 \%$ and $75 \%$ of the total area under the curve; d) the Average of $T_{2}{ }^{*} / C S F$ (Av $T_{2}{ }^{*} / C S F$ ) of the region analyzed. Histogramderived metrics were compared between all patients, and in SPMS only vs controls using Student's t-test for unpaired data. For this preliminary study we focused our analysis in the whole right or left hemisphere and in different frontal regions including the frontal pole, superior frontal gyrus, rostral middle frontal gyrus, and cingulate as pathology data demonstrated that these areas are greatly affected by diffuse subpial demyelination.

Results. With the exception of RPH, all $\mathrm{T}_{2}{ }^{*} / \mathrm{CSF}$ histogram-derived metrics for the whole cortex in the right hemisphere were greater in the group of all patients vs controls ( $\mathrm{PP}=38.7 \pm 5.7$ vs $36.5 \pm 4.9 ; \mathrm{T}_{2}{ }^{*} / \mathrm{CSF}_{25}=33.8 \pm 5.5$ vs $31.2 \pm 2 ; \mathrm{T}_{2}{ }^{*} / \mathrm{CSF}_{50}=39.4 \pm 5.2$ vs $36.9 \pm 5.2 ; \mathrm{T}_{2}{ }^{*} / \mathrm{CSF}_{75}=44.7 \pm 4.5 \mathrm{vs} 42.2 ; \mathrm{Av}$ $\mathrm{T}_{2}{ }^{*} / C S F=0.64 \pm 0.1$ vs $0.57 \pm 0.03$ ) but only $A v T_{2}{ }^{*} / C S F$ difference reached statistically significance ( $\mathrm{p}<0.04$ ). Similarly, Av $\mathrm{T}_{2}{ }^{*} / C S F$ in the left hemisphere was significantly higher in patients vs controls ( $0.63 \pm 0.1$ vs $0.57 \pm 0.03, \mathrm{p}<0.04$ ). When we assessed $\mathrm{T}_{2}{ }^{*} / \mathrm{CSF}$ histogram-derived metrics in frontal cortical regions, we found that the greatest increase in $T_{2}{ }^{*} / C S F$ was in the right rostral middle frontal gyrus (Fig. 1, Table 1).

Fig. 1. Right Rostral Middle Frontal Gyrus


Table 1. $\mathrm{T}_{2}{ }^{*} /$ CSF histogram-derived metrics in the right rostral middle frontal gyrus

|  | Controls | 14 MS patients | p-value |
| :--- | :--- | :--- | :--- |
| $\mathrm{RPH}(\mathrm{SD})$ | $6.8(1.4)$ | $6.0(1)$ | 0.13 |
| $\mathrm{PP}(\mathrm{SD})$ | $40.5(7.2)$ | $41.6(12.7)$ | 0.8 |
| $\mathrm{~T}_{2}{ }^{*} / \mathrm{CSF}_{25}(\mathrm{SD})$ | $36.7(3.8)$ | $40.9(6.2)$ | 0.07 |
| $\mathrm{~T}_{2}{ }^{*} / \mathrm{CSF}_{50}$ | $41.4(5.2)$ | $45.9(6.1)$ | 0.08 |
| $\mathrm{~T}_{2}{ }^{*} / \mathrm{CSF}_{25}$ | $45.7(5.4)$ | $51.1(6.0)$ | 0.04 |
| $\mathrm{Av} \mathrm{T}_{2}{ }^{*} / \mathrm{CSF}$ | $0.6(0.05)$ | $0.7(0.1)$ | 0.04 |

Subpial $T_{2}{ }^{*} / C S F$ changes were more evident in the subgroup of patients with SPMS, and involved mostly all the frontal regions analyzed in both hemispheres, though changes were greater in the right one. An example is shown in Fig. 2, Table 2.

Fig, 2. Right Cingulate


Table 2. $T_{2}{ }^{*} / C S F$ histogram-derived metrics in the right cingulate

|  | Controls | SPMS | p-value |
| :--- | :--- | :--- | :--- |
| $\mathrm{RPH}(\mathrm{SD})$ | $4.9(0.8)$ | $3.6(0.7)$ | 0.004 |
| $\mathrm{PP}(\mathrm{SD})$ | $62.1(7.2)$ | $70.20(3.7)$ | 0.01 |
| $\mathrm{~T}_{2}{ }^{*} / \mathrm{CSF}_{25}(\mathrm{SD})$ | $53.0(6.3)$ | $55.4(.7)$ | 0.5 |
| $\mathrm{~T}_{2}{ }^{*} \mathrm{CSF}_{50}$ | $60.7(6.6)$ | $65.6(3.4)$ | 0.09 |
| $\mathrm{~T}_{2}{ }^{*} / \mathrm{CSF}_{25}$ | $66.9(6.5)$ | $74.2(2.3)$ | 0.008 |
| ${\mathrm{~A} v \mathrm{~T}_{2}{ }^{*} / \mathrm{CSF}}^{2}$ | $0.6(0.04)$ | $0.61(0.05)$ | 0.4 |

Conclusions. The histogram-based analysis showed significant, diffuse $T_{2}{ }^{*} / C S F$ signal increases in MS patients vs matched controls, particularly evident in frontal areas and in SPMS. The observed changes may underlie diffuse subpial demyelination reported by several neuropathology examinations. This hypothesis can be validated by correlating histological evidence of diffuse cortical demyelination to the presence of diffuse cortical MR changes in ex vivo MS samples
References. 1. Stadelmann C. Curr Opin Neurol 21:229-34 (2008). 2. Sled, JG. IEEE Trans Med Imaging 17:87-97 (1998). 3. Greve D, Fischl. Neuroimage 48:3-72 (2009).
Acknowledgments. This study was partially supported by NIH (5P41 RR14075 08) and NMSS (4281-RG-A-1).

