Advanced MRI unravels the nature of tissue alterations in early multiple sclerosis

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Introduction

Multiple sclerosis (MS) is an inflammatory and neurodegenerative disease of the central nervous system (CNS) characterized by the presence of focal lesions in white matter (WM) and gray matter (GM) and also by diffuse inflammation and degeneration in normal-appearing (NA) tissue.1,2 Conventional MRI plays a major role in identifying focal inflammation and diagnosing MS, but has important limits in assessing underlying pathology. As a consequence, this method provides only modest correlations with patient functional performance, particularly during early phases of the disease.3

In this context, quantitative and semiquantitative (q/sq) MRI techniques1,4,5 may provide new biomarkers of disease severity and help to improve the clinical–radiological mismatch in MS treatment. To this end, pathological processes such as demyelination, edema formation, tissue...
loss, and iron accumulation lead to variable changes in quantitative measures of proton relaxation times (T1, T2, and T2*) as well as in semiquantitative parameters such as the magnetization transfer ratio (MTR).\textsuperscript{6-10} Thanks to recent MRI developments,\textsuperscript{11,12} it is now possible to combine multiple q/sq MRI sequences in a clinically applicable protocol and gather more specific information about the nature of tissue pathology in MS.

In this work, we investigated whether the combination of advanced T1, T2, and T2* relaxometry and magnetization transfer imaging may be employed (1) to assess the nature of brain tissue changes occurring early in MS and (2) to improve the correlation between imaging and clinical performance.

**Methods**

**Study population**

We enrolled 36 patients with relapsing-remitting MS (RRMS), 24 women/12 men, age = 34.8 ± 9.2 years (mean ± standard deviation [SD]) and 18 age-matched healthy controls (HC), 9 women/9 men, age = 33 ± 9.7 years. All patients were <6 years from initial symptoms (33.3 ± 21 months, range 2–70 months) and disease diagnosis (27.1 ± 18 months, range 0–59 months). Thirty patients (83%) were under immunomodulatory treatment (high dosage interferon beta or fingolimod) for at least 3 months. No patient had received corticosteroid therapy within the 3 months preceding the enrollment. The study was approved by the ethics committee of the Lausanne University Hospital (CHUV). Written, informed consent was obtained from each subject.

**Clinical assessment**

Each subject underwent a neurological examination including the following cognitive and behavioral tests: (1) Brief Repeatable Battery of Neuropsychological Tests (BRB-N),\textsuperscript{13} which examine verbal and spatial memory, sustained attention, information processing speed, and verbal fluency on semantic cues; (2) the Hospital Anxiety and Depression scale (HAD)\textsuperscript{14}; and (3) the Fatigue Scale for Motor and Cognitive functions (FSMC)\textsuperscript{15} which quantifies depressive mood symptoms and fatigue. The Expanded Disability Status Scale (EDSS\textsuperscript{16}) and the Multiple Sclerosis Functional Composite (MSFC\textsuperscript{17}) scores were assessed by a certified neurologist (C. Granziera, CG) to quantify motor performance.

**MRI techniques**

All MR images were acquired on a 3T Siemens Trio (Siemens, Erlangen, Germany) equipped with a 32-channel head coil. The acquisition protocol consisted of: (1) high-resolution 3D magnetization-prepared acquisition with gradient echo (MPRAGE) (TR/TE = 2300/2.98 ms, voxel size = 1.0 × 1.0 × 1.2 mm\(^3\), FoV = 256 × 240 × 192 mm\(^3\), acquisition time = 5:12 min) for automatic brain tissue, and atlas-based segmentation as reported previously\textsuperscript{18-20}; signal-to-noise ratio (SNR) measurements on a MPRAGE image were performed based on\textsuperscript{21,22} and reported in detail in the supplementary data (2) high-resolution 3D fluid attenuated inversion recovery (FLAIR) (TR/TE/TI = 5000/394/1800 ms, voxel size = 1.0 × 1.0 × 1.2 mm\(^3\), FoV = 256 × 240 × 212 mm\(^3\), acquisition time = 6:27 min); (3) high-resolution 3D double inversion recovery (DIR) (TR/TE/TI = 10,000/218/3650 ms, voxel size = 1.1 × 1.0 × 1.2 mm\(^3\), FoV = 256 × 240 × 192 mm\(^3\), inversion times 450/3652 ms, acquisition time = 12:52 min); (4) Magnetization-Prepared 2 Rapid Acquisition Gradient Echoes MP2RAGE\textsuperscript{12} (TR/TE = 5000/2.89 ms, voxel size = 1.0 × 1.0 × 1.2 mm\(^3\), FoV = 256 × 240 × 212 mm\(^3\), acquisition time = 8:22 min) for lesion count,\textsuperscript{23} and whole-brain T1 relaxometry; (5) T2 relaxometry (TR/TE = 5850/9 ms, 21 echos, 30 slices: voxel size = 1.0 × 1.0 × 4.0 mm\(^3\), FoV = 210 × 175 × 120 mm\(^3\), acquisition time = 3 min) using a new nonlinear inverse reconstruction algorithm that directly estimates a T2 and spin-density map from a train of undersampled spin echoes\textsuperscript{18}; and finally, (6) T2* relaxometry (TR/TE = 47/1.23 ms, 32 gradient echoes, voxel size = 1.6 × 1.6 × 1.6 mm\(^3\), FoV = 217 × 217 × 179 mm\(^3\), acquisition time = 11:16 min) with and without magnetization transfer (MT) pulse (MT pulse flip angle: 220°C; duration: 4000 ms; pulse offset: 2000 Hz; spoiler moment: 25,000 us × mT/m\textsuperscript{24}). In order to correct for susceptibility induced macroscopic field inhomogeneities, which were already diminished by isotropic high-spatial resolution, we used a 3D Sinc Correction\textsuperscript{25} that was extended to include a nonlinear correction term based on the underlying B0 map\textsuperscript{26}. The B0 map was calculated as the weighted mean phase difference\textsuperscript{27} of the temporally unwrapped phase followed by a median and Gaussian filters to remove phase inconsistencies.\textsuperscript{28} R2* maps were computed from T2 and T2* maps according to

\[
R^{*2} = \frac{1}{T^{*2}} - \frac{1}{T^{2}}
\]

MTR maps were derived from the T2* data by

\[
MTR = \frac{M_0 - M_T}{M_0}
\]

with \(M_0\) and \(M_T\) the images acquired without and with MT pulse, respectively. MT images were registered to images without MT pulse. Before any processing, image quality was assessed for each modality by visual inspection.
Figure 1 provides an example of all images and maps in one HC and one MS subject. Raw data from a HC are available in Data S1. Total scan time was ~47 min.

**MRI contrasts**

T1 relaxation time (rt) in brain tissue is mainly influenced by free water protons and the degree of structural
organization (i.e. amount of macromolecules such as myelin, lipids, proteins). In this context, an increase in T1\textsubscript{rt} may indicate a loss of structure and/or an increase in water content. Conversely, greater density of macromolecules and reduced water content as well as iron accumulation tend to reduce T1.29

T2\textsubscript{rt} measures the loss of spin coherence, and therefore, mainly reflect the dynamic state of water protons and their interaction with macromolecules. An increase in T2\textsubscript{rt} characterizes a loss of macromolecules and/or increased water content. On the contrary, a decrease in T2\textsubscript{rt} reflects an increase in protons bound to macromolecules. As for T1, iron accumulation also causes a shorter T2\textsuperscript{0} (Fig. 1).

The effective T2\textsuperscript{0} transverse rt describes the loss of transverse magnetization due to T2 relaxation and magnetic field inhomogeneities (R2’ component\textsuperscript{31}). Possible sources are tissue-dependent differences in magnetic susceptibility or the presence of paramagnetic or ferromagnetic ions like iron. For these reasons, an increase in T2\textsuperscript{0} most often indicates a loss of macromolecules, while a decrease suggests an increase in macromolecular compounds or iron that translate into an increase in R2’.

MT images are based on the interaction between free protons and immobilized protons bound to macromolecules, so that a lower MTR indicates a reduced spin exchange between macromolecules and surrounding bulk water suggesting neuroaxonal damage or myelin breakdown\textsuperscript{32} and/or water increase.

**Image analysis and tissue segmentation**

We used the Elastix c++ library\textsuperscript{33} to perform (1) rigid registration with BSpline interpolation of the T2 maps to the T1 maps (from the MP2RAGE); and (2) rigid registration of T2\textsuperscript{0} maps, MPRAGE, FLAIR, and DIR images to one of the inverted contrasts of the MP2RAGE sequence. By doing this, we obtained all images in the MP2RAGE space.

Regions of interest (ROIs) were derived from the MPRAGE image using in-house software based on variation-maximization tissue classification.\textsuperscript{34} The following ROIs were automatically segmented: whole-brain WM and cortical GM, thalamus, and basal ganglia (caudate, putamen, and globus pallidus), cerebellar WM and GM. In addition, we computed lobar WM and GM (temporal, occipital, frontal, parietal areas).

An experienced neurologist (CG) and a radiologist (D. Rotzinger, DR) manually counted MS lesions by consensus in 3D FLAIR, 3D DIR, and MP2RAGE images for all MS subjects and HC, as performed previously.\textsuperscript{20} A trained technician generated manual contours for each lesion in the three different contrasts (rechecked by DR). In order to maximize the sensitivity of lesion count and volume, lesion masks from each contrast were merged into a single mask (lesion union mask), as reported by Kober et al.\textsuperscript{20}

The lesion union mask and the ROIs masks were then registered to the T1, T2, T2\textsuperscript{0}, and MTR maps to obtain parametric values in lesions and NA tissue in each ROI.

The volume of each ROI was also automatically obtained using the in-house software based on a previous report\textsuperscript{34} and normalized by total intracranial volume.

**Statistical analyses**

**Between-groups comparisons of subjects’ demographics and clinical scores**

Differences in age, gender, education, and clinical performance were assessed using a nonparametric ANOVA (Kruskal–Wallis test) among HC and MS patients.

**Between-groups comparisons of multicontrast MRI data**

To assess NA tissue differences in mean T1, T2, T2\textsuperscript{0}, and MTR of patients and controls, we performed a permutation-based Hotelling test with 10,000 permutations, age and gender as covariates, and family-wise correction for multiple comparisons.

The following null hypotheses were tested: (1) there are no differences in WM and GM of temporal, parietal, occipital, and frontal lobes; (2) there are no differences in cerebellum WM and GM; and (3) there are no differences in thalamus and basal ganglia.

Lobar assessment was chosen, instead of whole brain, to take into account the local variation in quantitative relaxation measures, as reported previously.\textsuperscript{35,36}

In order to determine the strength of the significance, we also calculated the Cohen’s d effect size as follow:

\[
d = \frac{\bar{x}_1 - \bar{x}_2}{s}
\]

with \(\bar{x}_1\) and \(\bar{x}_2\) the mean of the group 1 (HC) and group 2 (RRMS), and \(s\) defined as follows:

\[
s = \sqrt{\frac{(n_1 - 1)s_1^2 + (n_2 - 1)s_2^2}{n_1 + n_2 - 2}}
\]

Parameters \(s_1\) and \(s_2\) refer to the standard deviation of group 1 (HC) and group 2 (RRMS), while \(n_1\) and \(n_2\) are the number of samples of group 1 and 2.

In order to compare lesional tissue, MRI properties in patients with the corresponding healthy tissue in HC, we calculated a z-score for each lesion and then averaged the
lesion z-scores across all lesions in each subject as follows (i.e. for T1 data):

\[ z_{T1} = \frac{1}{N} \sum_{l \in \text{Lesions}} \sum_{v \in f} \frac{I_{T1}(v) - \mu_{T1}(L, T)}{\sigma_{T1}(L, T)} \]

where \( z_{T1} \) corresponds to the average of T1 lesion z-scores in one patient, \( N \) to a normalization term, \( I_{T1} \) the T1 map, \( \mu_{T1}(L, T) \) and \( \sigma_{T1}(L, T) \) to the mean and the standard deviation of T1 in the lobe \( L \) and tissue \( T \) (i.e. WM or GM) in the HC group, corresponding to the lesion location. Averages of each patient’s T1, T2, T2*, and MTR lesion z-scores were also performed in the whole MS group.

This approach was chosen instead of the permutation-based test applied for NA tissue to account for spatial variation in relaxometry values.\(^{35,36}\) A permutation test was not feasible for each lobe as not all patients exhibited lesions in all lobes.

**Between-groups comparison of volumes**

To assess volumetric differences in ROIs’ between patients and controls, we performed a permutation-based Hotelling test with 10,000 permutations, age and gender as covariates, and family-wise correction for multiple comparisons.

**Linear regression of MRI parameters with clinical scores**

All regression analyses were performed using R software (http://www.R-project.org).

A multivariate linear regression of clinical scores was performed using a general linear model (GLM) applied (1) T2*, T2, T1, and MTR in the ROIs that differed between patients and HC, (2) T1, T2, T2*, and MTR lesion z-scores and (3) cortical/subcortical lesion count and volume. Age, gender, educational years, anxiety, and depression scores (HAD) were considered as covariates, since they have been reported to be linked to functional performance.\(^{37,38}\) Cognitive scores were adapted using Box–Cox transformation to satisfy model assumption for normality.\(^{39}\) EDSS scores were not considered, as they were positive only in patients.

We performed seven regressions, where we used a backward stepwise approach to select the best prediction model for each dependent variable (clinical scores). Bonferroni correction was applied for multiple comparisons (seven tests).

Cook’s distance (Cd) was computed to assess the influence of each observation on the regression process, using \( 4/n \) (n: number of observations) as the threshold of significance. Robust regression was used to reduce influence of the outliers identified by Cook’s distance analysis. “Leave-one-out” (LOO) cross-validation was applied to assess the prediction quality and robustness of each model. A P < 0.05 was considered statistically significant.

**Results**

**Between-groups comparisons of subject demographics and clinical scores**

No significant differences were observed between HC and MS patients in terms of age (\( P = 0.3 \)) or gender (\( P = 0.8 \)); however, HC had slightly higher education levels (17 ± 4 years, mean ± standard deviation) than MS patients (15 ± 3 years, \( P = 0.04 \)).

Mean EDSS in patients was 1.6 ± 0.3 (interval: 1–2). The FSMC motor score was significantly higher in MS patients (23.1 ± 10.5) than in HC (14.8 ± 5.8, \( P < 0.02 \)). The FSMC cognitive scores, cognitive performance, MSFC scores, as well as anxiety and depression scores (HAD) were not significantly different between groups (\( P > 0.1 \)).

**Between-groups comparison of multicontrast MRI data**

In temporal NAWM, mean T2* and T2 were significantly higher in RRMS patients compared to HC (T2*: rt: 55.1 ± 1.55 msec in patients and 53.4 ± 1.35 msec in HC, \( d = 1.17, P = 0.004 \), Fig. 2; T2: rt: 82.0 ± 2.38 msec in patients and 79.8 ± 2.0 msec in HC, \( d = 1, P = 0.03 \), Fig. 2).

In order to assess whether the observed T2* increase in temporal NAWM depended on local field inhomogeneities, we also compared temporal NAWM R2’ between groups and found no significant differences.

Additionally, parietal NAWM and cerebellar NAWM exhibited a trend toward higher T2 values in patients compared to HC (parietal NAWM T2: 83.5 ± 2.44 msec in patients compared to 81.8 ± 2.62 msec in HC; \( d = 0.7, P = 0.05 \); and cerebellar WM T2: 85.90 ± 1.69 msec in patients compared to 85.48 ± 1.47 msec in HC; \( d = 1.62, P = 0.07 \)).

Further, no differences were seen for T1 and MTR in NAWM and cortical NAGM, nor for T2 and T2* in cortical NAGM, frontal or occipital NAWM. Finally, no significant differences between groups were found for T1, MTR, T2, or T2* in the thalamus or basal ganglia.

Results of microstructural analysis of lesions are reported in Figure 3.

In the MS cohort, MS lesions showed a strong increase in T1 mean z-score (4.42) and an important decrease in MTR mean z-score (−4.09). T2 and T2* mean z-scores slightly increased (2.33 and 2.25, respectively).
Figure 2. (A) (Top): $T_2^*$ and $T_2$ mean histograms in NAWM (temporal lobe) for HC (blue) and MS patients (red); (B) (Below): Boxplot of $T_2^*$ and $T_2$ in NAWM (temporal lobe) for HC (left) and MS patients (right).
Between-groups comparison of volumes

No significant differences were observed in volumes between MS patients and HC, however, there was a trend toward smaller normalized thalamic volumes in patients (absolute volume $15.31 \pm 1.36 \text{ mm}^3$, normalized volume $0.01 \pm 0.0006$) compared to HC (absolute volume $16.52 \pm 2.04 \text{ mm}^3$, normalized volume $0.01 \pm 0.0003$ $P = 0.07$).

Linear regression of MRI parameters with clinical scores

GLM using backward, stepwise regression revealed a highly significant association, confirmed by cross-validation results, between multicontrast MRI features and four clinical scores (Table 1):

1. Cortical lesions count and volume, T1, T2, and T2* mean z-score of lesions, T1, T2*, and MTR mean of temporal NAWM together with gender predicted the Symbol Digit Modalities Test (attention function) score (adjusted $R^2$: 0.6, $P = 0.0005$).

2. T2, T2*, and MTR mean in temporal NAWM in conjunction with T1 and T2 mean z-score in lesions as well as subcortical lesion volume and educational years, gender and HAD scores predicted the MSFC (general disability) score (adjusted $R^2$: 0.4, $P = 0.03$).

3. T1 and T2 mean in temporal NAWM combined with cortical lesion volume, subcortical lesion count, and

![Figure 3. T1, MTR, T2, and T2* mean z-scores per patient (columns) in MS lesion.](image)

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<th>Table 1. Multiple regression analysis between lesional and temporal NAWM MRI characteristics, covariates, and clinical scores.</th>
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Top part: each line corresponds to the $P$-values, corrected $P$-values, and adjusted-$R^2$ of each model ($n = 7$) subjected to regression and cross-validation analysis. Bottom part: each line corresponds to the $P$-values of each predictor for every regression model performed. The color scheme signifies the difference in significance: dark orange = highest significance ($P < 0.001$), light orange = middle range significance ($P < 0.01$), and yellow = low significance ($P < 0.05$).
HADD score predicted the FSMC cognitive score (adjusted R2: 0.4, P = 0.01).

T1 and T2 mean in temporal NAWM combined with cortical lesion volume, subcortical lesion count, and volume with HADD score predicted the FSMC motor score (adjusted R2: 0.4, P = 0.01).

**Discussion**

The present results demonstrate that combining multiple advanced MRI techniques, it is possible to unravel the nature of subtle tissue alterations in early MS. Moreover, MRI markers of inflammation and neurodegeneration may substantially improve clinical–radiological correlations compared to conventional measures.

The RRMS patients enrolled in our study exhibited significant increases in T2 and T2*rt in temporal NAWM, and to a lesser extent in parietal and cerebellar NAWM. These changes hint to an accumulation of extracellular water (microedema) and/or a reduction in macromolecular content (myelin) in affected brain tissue (Fig. 2). In the absence of significant changes in MTR and T1rt, which would support the structural explanation, the increase in both T2 and T2*rt most likely indicates the presence of subtle edema. Iron loss might also be responsible for a prolongation of T2 and T2*rt, but appears to be a less probable cause as no differences were observed in R2*, which reflects local field inhomogeneities.

By combining multiple q/sq MRI measures, our study confirms work reporting T2 increase in NAWM in early MS and extends these findings by providing new insights into the pathology underlying those changes. However, our data contradict studies showing a measured unimodal MTR decrease in NAWM of early MS patients attributed to myelin loss. Nevertheless, these studies focused mainly on untreated patients and applied MT imaging at lower spatial resolution and lower field strength (1.5 T) than ours. Furthermore, unimodal MTR studies in MS should be considered with caution, as MT imaging alone cannot discriminate between myelin alterations and variation in water content in tissues.

Axonal degeneration in NAWM of early MS patients was also suggested by unimodal diffusion tensor imaging (DTI) studies, showing reduced fractional anisotropy (FA). Nevertheless, this interpretation may be misleading since a decrease in anisotropy can derive from the loss of the branched-shape of microglia cells that is typical of their activated-inflammatory form. Thus, another possible explanation is that reduced FA might point to inflammatory rather than degenerative phenomena.

Several studies tried to address the limitation of unimodality studies by combining T2 relaxometry, MTR measurements, and DTI; however, these studies focused on selected brain structures (i.e. corpus callosum and corticospinal tract) in patients with advanced stages of MS. Our approach overcomes the above-mentioned limits by performing a whole-brain analysis of multiple q/sq assessments in early MS.

Our data also showed that both cortical and subcortical lesions were characterized by a strong increase in T1rt and decrease in MTR with relatively modest positive variations of T2 and T2*rt (Fig. 3). These findings are consistent with previous MRI literature and histopathological studies showing significant neurodegeneration in MS plaques.

No significant microstructural alterations were found in NA tissue belonging to the basal ganglia or thalamus in our MS cohort. Still, volumetric analyses revealed a trend toward lower regional volumes in patients (P = 0.07), which is consistent with thalamic atrophy reported in larger and more heterogeneous patient groups.

Last, we showed that MRI findings of microstructural alterations in NA tissue and lesions substantially improved the clinical–radiological correlation obtained with conventional measures, even in the presence of minor functional deficits. In fact, a variable combination of relaxometry and MTR values significantly ameliorated the prediction of cognitive performance (attention), cognitive fatigue, and general disability obtained with traditional measures of disease burden and patient covariates (Table 1).

Conventional MRI measures of MS disease impact provide only modest correlations with clinical performances, a phenomenon that is known as the clinico-radiological paradox. Multivariate analyses and multicontrast, tract-specific measures were proposed to alleviate this paradox but suffered from the limitations of conventional protocols and partial brain analyses. Recently, ultra-high field MRI at 7 T has been used to identify subtypes of cortical lesions, whose numbers showed good correlations with disability and cognitive performance in MS. Extending such work, the multicontrast approach presented here emerges as a whole-brain MRI method at a clinically-compatible magnetic field, which produces strong clinical–radiological correlations for both cognition and disability.

Future developments should aim at reducing the number of sequences required for optimal lesion detection (i.e. MP2RAGE and 3DFLAIR) as well as at applying accelerated T1-T2* relaxometry sequences to achieve a well-suited protocol for the clinical workflow.

In summary, combining multiple recent MRI techniques at 3T we found (1) increased T2 and T2* in temporal NAWM, suggesting subtle microedema; (2) a strong increase in T1rt and decrease in MTR in lesions, indicating prevalent tissue degeneration; and (3) improved correlations between MRI data and measures of cognition and disability in early and minimally impaired MS patients. Additional studies extending the current methods to...
patients at later disease stages and containing larger cohorts will be necessary in the future.

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Conflict of Interest

Dr A. Roche and G. Krüger work for Siemens AG Schweiz.

References


Supporting Information

Additional Supporting Information may be found in the online version of this article:

Data S1. Supplementary data.