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White Matter Abnormalities in Multiple Sclerosis Evaluated by Quantitative Synthetic MRI, Diffusion Tensor Imaging, and Neurite Orientation Dispersion and Density Imaging

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ABSTRACT

BACKGROUND AND PURPOSE: A number of MR-derived quantitative metrics have been suggested to assess the pathophysiology of MS, but the reports about combined analyses of these metrics are scarce. Our aim was to assess the spatial distribution of parameters for white matter myelin and axon integrity in patients with relapsing-remitting MS by multiparametric MR imaging.

MATERIALS AND METHODS: Twenty-four patients with relapsing-remitting MS and 24 age- and sex-matched controls were prospectively scanned by quantitative synthetic and 2-shell diffusion MR imaging. Synthetic MR imaging data were used to retrieve relaxometry parameters (R1 and R2 relaxation rates and proton density) and myelin volume fraction. Diffusion tensor metrics (fractional anisotropy and mean, axial, and radial diffusivity) and neurite orientation and dispersion index metrics (intracellular volume fraction, isotropic volume fraction, and orientation dispersion index) were retrieved from diffusion MR imaging data. These data were analyzed using Tract-Based Spatial Statistics.

RESULTS: Patients with MS showed significantly lower fractional anisotropy and myelin volume fraction and higher isotropic volume fraction in widespread white matter areas. Areas with different isotropic volume fractions were included within areas with lower fractional anisotropy. Myelin volume fraction showed no significant difference in some areas with significantly decreased fractional anisotropy in MS, including in the genu of the corpus callosum and bilateral anterior corona radiata, whereas myelin volume fraction was significantly decreased in some areas where fractional anisotropy showed no significant difference, including the bilateral posterior limb of the internal capsule, external capsule, sagittal striatum, fornix, and uncinate fasciculus.

CONCLUSIONS: We found differences in spatial distribution of abnormality in fractional anisotropy, isotropic volume fraction, and myelin volume fraction distribution in MS, which might be useful for characterizing white matter in patients with MS.

ABBREVIATIONS: AVF = axon volume fraction; EDSS = Expanded Disability Status Scale; FA = fractional anisotropy; ICVF = intracellular volume fraction; ISO = isotropic volume fraction; MNI = Montreal Neurological Institute; MVF = myelin volume fraction; NAWM = normal-appearing white matter; NODDI = neurite orientation dispersion and density imaging; ODI = orientation dispersion index; QRAPMASTER = quantification of relaxation times and proton density by multiecho acquisition of a saturation-recovery using turbo spin-echo readout

MS is a demyelinating disorder that mainly affects young individuals and involves inflammatory demyelination accompanied by axonal degeneration. Various quantitative

metrics, including myelin water imaging,¹ quantitative synthetic MR imaging,^{2,3} diffusion tensor imaging,⁴ and neurite orientation dispersion and density imaging (NODDI),⁵ have revealed abnormalities in both WM lesions and the WM that appears normal on conventional T2WI.


Quantitative synthetic MR imaging simultaneously measures longitudinal T1 and transverse T2 relaxation times (and their inverse, R1 and R2 relaxation rates) and proton density in


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multiple slices,⁶ with high repeatability and reproducibility across scanners.⁷ It allows creation of any contrast-weighted image, including T1-weighted, T2-weighted, and FLAIR images,⁸ without requiring additional scanning time. Furthermore, a 4-compartment model was developed to estimate myelin volume fraction (MVF) in each voxel in the brain, based on the measured R1, R2, and proton density.⁹ The estimated myelin correlates with histology in healthy populations¹⁰ and in patients with MS¹¹ and with other MR imaging-based myelin measurements.¹² Even though synthetic myelin maps have shown tissue abnormalities in plaques and periplaque WM in patients with MS,² the voxel-level difference in the estimated myelin between patients with MS and healthy controls has not been investigated to date.

Diffusion-weighted imaging also provides information on microstructure, such as fiber density and orientation, by altering the diffusion-sensitization strength and direction of the encoding gradients.¹³ Although used as a standard diffusion MR imaging technique, diffusion tensor imaging lacks specificity for individual tissue microstructural features.¹⁴ Advanced diffusion models have been developed to quantify specific neurite morphology,¹³ such as NODDI, which assumes 3 compartments in the brain: an intracellular compartment with restricted anisotropic non-Gaussian diffusion (intracellular volume fraction [ICVF]), an extracellular compartment with hindered anisotropic Gaussian diffusion, and a water compartment with free isotropic Gaussian diffusion (isotropic volume fraction [ISO]).¹⁵ ICVF is attributed to the axon and dendrite density and can be used for calculating axon volume fraction (AVF) when combined with myelin imaging.¹⁶ Notably, the myelin signal is negligible in diffusion MR imaging, and an additional technique is required to retrieve information about axons and myelin.¹⁶ NODDI provides directional neurite information as an orientation dispersion index (ODI).

Despite the recent development of multiple advanced MR imaging techniques, reports of combined analyses are scarce. Granberg et al⁵ found that NODDI was more sensitive than myelin-sensitive imaging to changes in normal-appearing WM (NAWM) in patients with MS. However, they used the T1-weighted/T2-weighted ratio as a measure of myelin, which has been reported to be less than optimal for evaluating myelin in WM.^{12,17} Moreover, no previous study has performed voxelwise whole-brain analysis of tissue-damage distribution in patients with MS using NODDI or myelin-imaging techniques. Even though our previous work³ combined quantitative synthetic MR imaging and NODDI for evaluating WM damage in patients with MS, we did not compare quantitative values in patients with MS with those in healthy controls.

Therefore, this study aimed to evaluate the distribution of WM damage in patients with MS compared with healthy controls by combining quantitative synthetic MR imaging and NODDI. We used Tract-Based Spatial Statistics (TBSS; <http://fsl.fmrib.ox.ac.uk/fsl/fslwiki/TBSS>),¹⁸ which allows voxelwise analysis of the whole brain within core WM tracts, to compare patients with MS and healthy controls, and we used ROI analysis to compare NAWM and lesions in patients with MS with WM in healthy controls.

MATERIALS AND METHODS

Study Participants

We prospectively recruited 24 patients with relapsing-remitting MS from August to November 2016 who were diagnosed according to the 2010 revised McDonald diagnostic criteria.¹⁹ Disability was assessed using the Expanded Disability Status Scale (EDSS) score.²⁰ These patients were on stable disease-modifying treatment or no treatment for at least 3 months and had been free of clinical relapse within the 3 months and corticosteroid use within 1 year before MR imaging. As a control group, we also recruited 24 age- and sex-matched healthy subjects without neurologic and psychological symptoms or a history of neuropsychological disorders. Acquired images were confirmed not to include abnormalities such as moderate-to-severe WM ischemic lesions (Fazekas grade II or higher²¹), brain infarction, or tumor. This study was approved by the institutional review board of Juntendo University Hospital, Tokyo, Japan, and written informed consent was obtained from all participants. The patients in this study partially overlapped with a previously published study population.²²

Image Acquisition and Processing

MR imaging was performed on a 3T scanner (Discovery MR750w; GE Healthcare, Milwaukee, Wisconsin) with a 19-channel head coil. All participants were scanned with a 2D axial quantification of relaxation times and proton density by multiecho acquisition of a saturation-recovery using turbo spin-echo readout (QRAPMASTER) pulse sequence and 2-shell diffusion MR imaging.

QRAPMASTER is a multislice, multiecho, multisaturation-delay saturation-recovery turbo spin-echo acquisition method in which images are collected with combinations of 2 TEs and 4 saturation-delay times. We used TEs of 16.9 and 84.5 ms and delay times of 146, 546, 1879, and 3879 ms. The other parameters were as follows: TR = 4.0 seconds, FOV = 240 × 240 mm, matrix = 320 × 320, echo-train length = 10, bandwidth = 31.25 kHz, section thickness/gap = 4.0 mm/1.0 mm, slices = 30, and acquisition time = 7 minutes 12 seconds. The 8 complex images acquired per section were postprocessed with SyMRI software (Version 8.0; SyntheticMR, Linköping, Sweden) to derive longitudinal R1 relaxation and transverse R2 relaxation rates and proton density and to estimate MVF per voxel. The myelin estimation model assumes 4 brain compartments: myelin, excess parenchymal water, cellular water, and free-water volume fractions.⁹ The R1, R2, and proton density of each volume fraction in a voxel presumably contribute to the effective R1, R2, and proton density of the voxel as a whole. Synthetic FLAIR images and T1WI were also created with the same software using the R1, R2, and proton density maps, with the following postprocessing parameters: TR = 15,000 ms, TE = 100 ms, TI = 3000 ms for FLAIR, and TR = 500 ms, TE = 10 ms for T1WI.

Diffusion MR imaging was performed with single-shot echo-planar imaging along 30 and 60 motion-probing gradient directions for b-values of 1000 and 2000 s/mm², respectively. Additionally, a volume of non-diffusion-weighted images was also acquired. The other sequence parameters were the following: TR = 5000 ms, TE = 88.2 ms, FOV = 256 × 256 mm, matrix size =

256 × 256, echo-train length = 128, bandwidth = 1953.12 kHz, section thickness/gap = 4.0 mm/1.0 mm, slices = 30, and acquisition time = 7 minutes 40 seconds. All datasets were visually inspected for artifacts. We corrected in-plane and through-plane distortions of diffusion-weighted images caused by eddy currents and motion using affine brain registration to non-diffusion-weighted images.²³ Processed images were further denoised using multishell position-orientation adaptive smoothing based on the propagation-separation approach.^{24,25} On denoising, diffusion data with b-values of 1000 and 2000 s/mm² were handled simultaneously to improve denoising stability.²⁵ Maps of fractional anisotropy (FA), mean diffusivity, axial diffusivity, and radial diffusivity were computed using diffusion data with b-values of 0 and 1000 s/mm² by fitting a tensor model. A NODDI model¹⁵ was applied to the whole 2-shell diffusion data to produce ICVF, ISO, and ODI maps, while processing was accelerated using an Accelerated Microstructure Imaging via Convex Optimization algorithm.²⁶

The AVF and g-ratio maps were acquired using the following equations:

$$AVF = (1 - MVF)(1 - ISO)ICVF,$$

$$g - ratio = \sqrt{\frac{AVF}{MVF + AVF}},$$

assuming that the signal of myelin is almost negligible in diffusion-weighted imaging but that the volume of myelin is not,¹⁶ — that is, the volume fractions calculated by NODDI correspond to nonmyelinated tissues. Postprocessing was performed with an in-house program in Matlab (MathWorks, Natick, Massachusetts). Affine transformation was performed to register the acquired images using Statistical Parametric Mapping software (SPM12; <http://www.fil.ion.ucl.ac.uk/spm/software/spm12>).

TBSS Analysis

We performed whole-brain voxelwise analysis of the quantitative maps using TBSS implemented in the FMRIB Software Library (FSL; <http://www.fmrib.ox.ac.uk/fsl>).^{18,27} First, FMRIB58_FA (https://fsl.fmrib.ox.ac.uk/fsl/fslwiki/FMRIB58_FA) in the Montreal Neurological Institute (MNI) common space was used as the target image for nonlinear registration of all subjects' FA maps, using the FMRIB Nonlinear Registration Tool (FNIRT; <http://fsl.fmrib.ox.ac.uk/fsl/fslwiki/FNIRT/>). The transformed FA images were averaged and skeletonized, representing the centers of all WM tracts common to all subjects. The mean FA skeleton was thresholded at 0.2 to include major WM tracts and to exclude peripheral tracts and gray matter.¹⁸ Each participant's aligned FA map was then projected onto this skeleton by assigning to each voxel the maximum FA in a line perpendicular to the local skeleton. The R1, R2, proton density, MVF, mean diffusivity, axial diffusivity, radial diffusivity, ICVF, ISO, ODI, AVF, and g-ratio maps were projected onto the mean FA skeleton after applying the warping registration field of each subject to the standard space.

Comparisons between patients with MS and healthy controls were performed by voxelwise statistics of the skeletonized

quantitative maps using nonparametric statistical thresholding (FSL Randomise permutation algorithm; <https://fsl.fmrib.ox.ac.uk/fsl/fslwiki/Randomise>). The thresholded mean FA skeleton was used as a mask. Five thousand permutations and statistical inference using threshold-free cluster enhancement were performed,²⁸ with *P* values < .05 after family-wise error correction for multiple comparisons considered significant. Age and sex were used as covariates. The anatomic locations of regions with significant group differences on the WM skeleton were identified from the Johns Hopkins University WM labels atlas.²⁹ For parameters showing significant differences, correlations with disease duration and EDSS were also examined using Randomise (*P* < .05, corrected for age and sex).

Lesion Maps

For all patients, hyperintense lesions were automatically segmented on synthetic FLAIR images using the lesion-prediction algorithm³⁰ implemented in the Lesion Segmentation Toolbox, Version 2.0.15 (<http://www.applied-statistics.de/1st.html>)³¹ running in SPM 12. All lesion maps were visually inspected and manually corrected by an experienced neuroradiologist (A.H.). The whole-brain WM lesion volume in each patient was calculated by multiplying the lesion area by the section thickness. Synthetic T1WI in each patient was spatially normalized to MNI space, and warping fields were saved and subsequently applied to lesion maps. Because synthetic images derived from QRAPMASTER are inherently aligned,³² no prior registration between synthetic T1WI and lesion maps created on synthetic FLAIR was required. Normalized lesion maps in all patients were aggregated to create group lesion maps.

ROI Analysis

To investigate tissue damage in NAWM and lesions separately, we performed ROI analysis using the WM skeleton and lesion maps. We registered the skeleton in the MNI space to each subject's space by applying the warping field created for TBSS analysis after inversion of the field. Next, we enlarged each patient's lesion map by 4 voxels. We segmented the synthetic T1WI for each subject to extract WM segmentation maps using FMRIB's Automated Segmentation Tool (FAST; <http://fsl.fmrib.ox.ac.uk/fsl/fslwiki/fast>). WM segmentation maps were then thresholded at 0.9 to minimize the partial volume effects and were used as WM masks. We created NAWM masks by subtracting the enlarged lesion maps from the thresholded WM masks for each patient. Overlapping areas between the warped skeleton and WM masks in healthy controls, NAWM masks in patients, or lesion maps in patients were used as ROIs to extract metrics from each quantitative map for the WM of healthy controls, the NAWM of patients with MS, or the lesion areas of patients with MS, respectively.

Statistical Analysis

Statistical analyses of demographic data were performed using a 2-sample *t* test for age and a χ^2 test for sex. For ROI data, non-normality of synthetic MR imaging metrics has been reported previously^{2,3}; therefore, we used the nonparametric Steel test for multiple comparisons to compare R1, R2, proton density, MVF,

FA, mean diffusivity, axial diffusivity, radial diffusivity, ICVF, ISO, ODI, AVF, and g-ratio values between the WM of healthy controls and the NAWM or lesion areas of patients with MS. A 2-sided $P < .05$ was considered statistically significant. These statistical analyses were performed with the software package R, Version 3.2.1 (<http://www.r-project.org/>).

RESULTS

The demographic and clinical data of patients with MS and of healthy controls are provided in the Table. Age and sex did not differ significantly between groups.

Demographic and clinical details of all participants

	MS	Healthy Control	P Values
No. of subjects	24	24	
Mean age (yr)	39.83 \pm 8.25	39.50 \pm 11.13	.91 ^a
Sex (male/female)	5:19	5:19	1 ^b
Disease duration (mean) (yr)	11.82 \pm 5.99	NA	
EDSS score (range)	1 (0–7)	NA	
White matter lesion volume (mean) (mL)	10.05 \pm 10.00	NA	

Note:—NA indicates not applicable.

^a Two-sample t test.

^b χ^2 test.

On TBSS analysis, patients with MS showed significantly lower FA and MVF and higher ISO values in the corpus callosum, cingulate gyri, and corona radiata (Fig 1 and On-line Table 1). The MVF differed between patients with MS and healthy controls in the most widespread areas, and only MVF showed significant differences (lower values in patients with MS) in the inferior longitudinal fasciculus, inferior fronto-occipital fasciculus, uncinate fasciculus, fornix, and external capsule. Conversely, only FA showed significantly different values (lower in patients with MS) in the genu of the corpus callosum and anterior corona radiata. All areas with significant ISO differences were included within areas that showed a significant FA difference. No significant correlation was found between EDSS or disease duration and FA, ISO, or MVF. R1, R2, proton density, mean diffusivity, axial diffusivity, radial diffusivity, ICVF, ODI, AVF, and g-ratio maps did not differ statistically significantly in the WM between patients and controls.

Figure 2 and On-line Table 2 show ROI analysis results. All quantitative metrics except ISO differed significantly between the healthy control WM and the WM lesions in patients with MS. R1, FA, mean diffusivity, radial diffusivity, ISO, ODI, and MVF differed significantly between healthy control WM and NAWM of patients with MS.

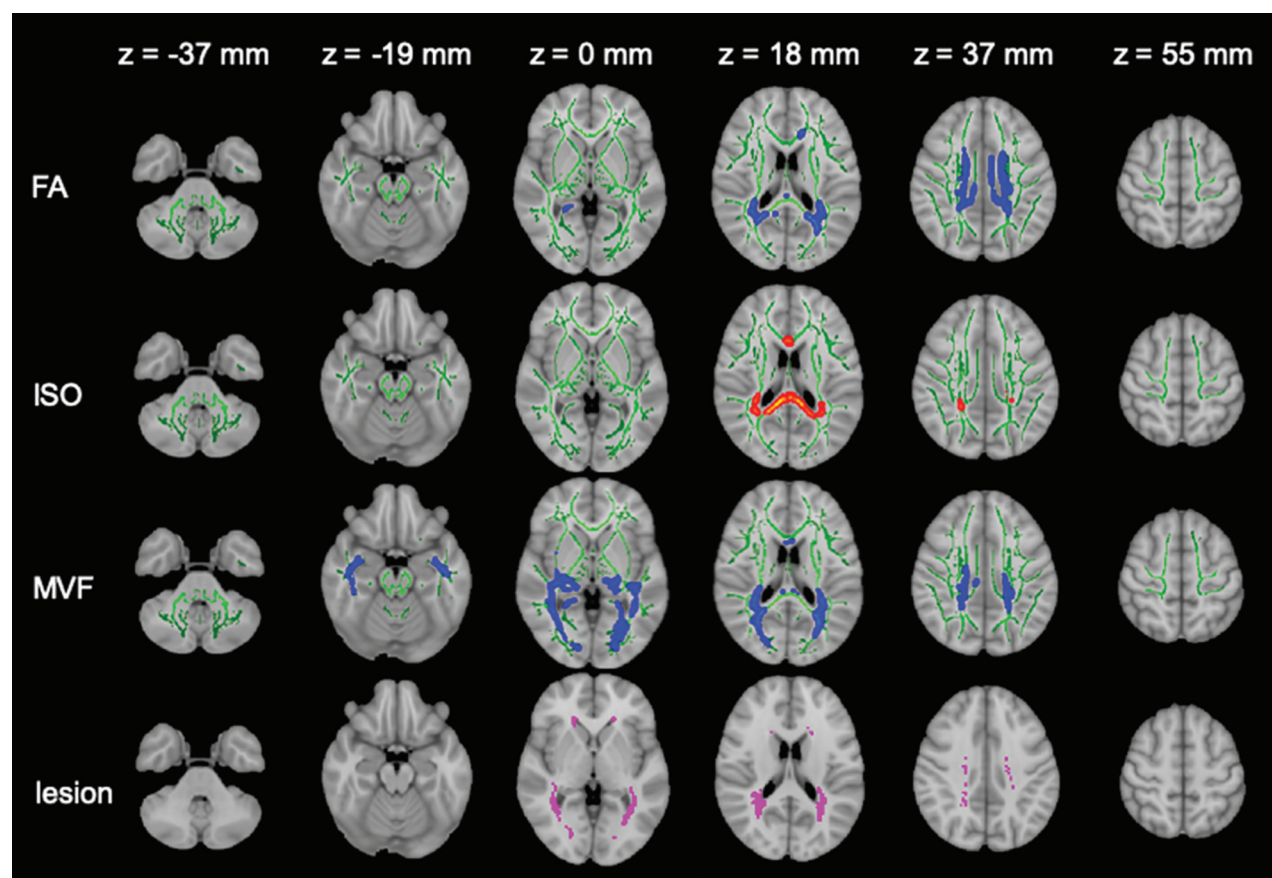


FIG 1. TBSS results are shown for FA, ISO, and MVF maps, which show significant differences between patients with MS and healthy controls. Green represents the mean FA skeleton of all participants thresholded at 0.2. Blue–light blue represents lower values in patients with MS compared with healthy controls in FA (first row) and MVF (third row) maps; red–yellow represents higher values in ISO maps (second row) (family-wise error-corrected $P < .05$). The significant regions are thickened for better visibility. The mean lesion probability distribution in patients with MS, thresholded at 20%, is shown in the fourth row in purple.

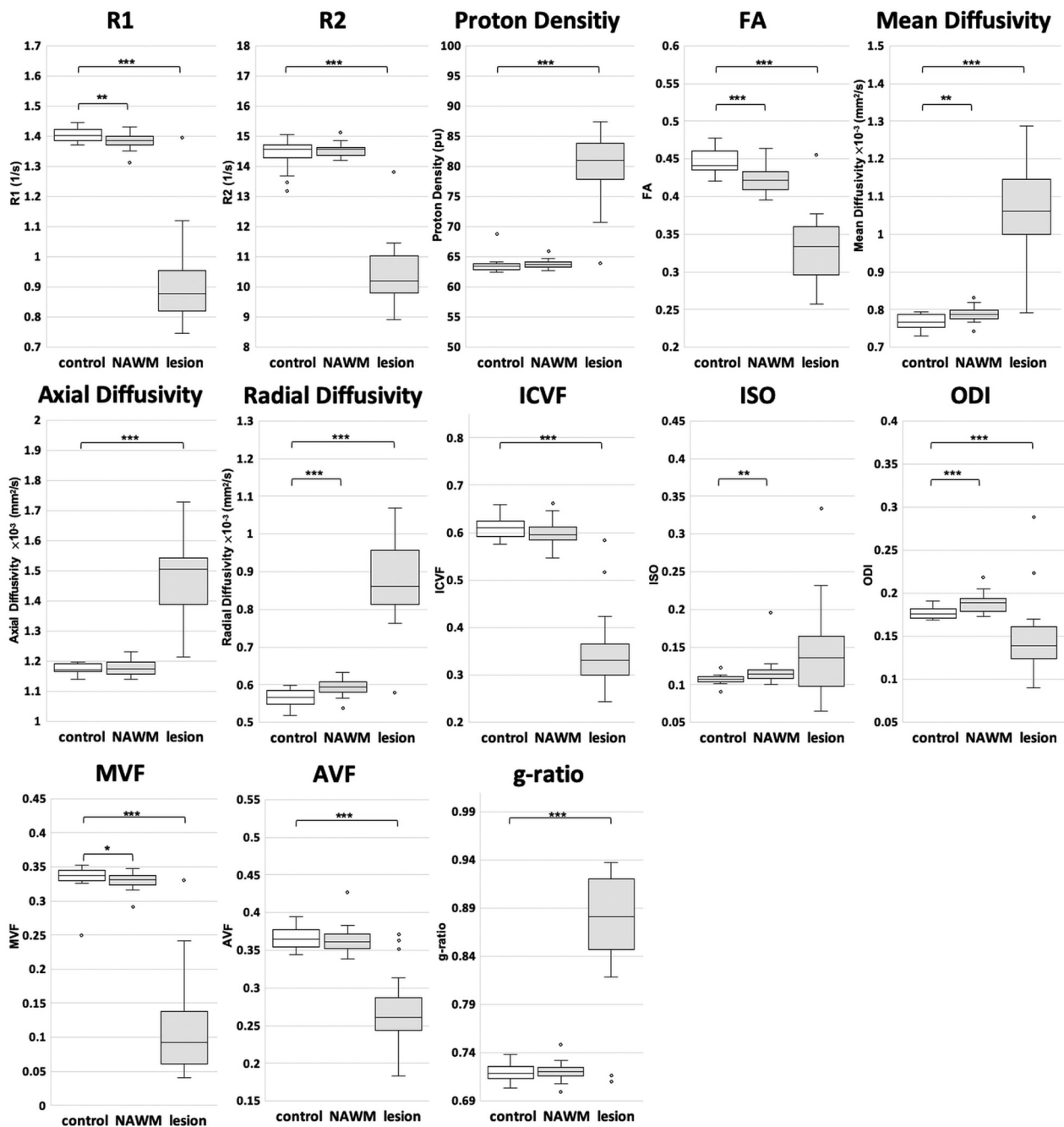


FIG 2. Quantitative metrics compared among the WM of healthy controls, NAWM of patients with MS, and WM lesions of patients with MS. The median and 25th and 75th percentiles are marked in *boxplots*, with outliers plotted by *open circles*. Asterisks indicate significant differences among the groups. The *asterisk* indicates $P < .05$; *double asterisks*, $P < .01$; *triple asterisks*, $P < .001$.

DISCUSSION

We here compared the WM of patients with MS and healthy controls using a combination of synthetic MR imaging and diffusion MR imaging by implementing TBSS and ROI analysis. To our knowledge, no previous study has performed voxelwise analysis of whole-brain WM using synthetic MVF or NODDI. We showed a lower FA in widespread WM areas in patients with MS, consistent with previous findings.³³⁻³⁵ The distribution of abnormality in MVF value in patients differed from

those of FA and ISO values. Even though the mechanisms underlying these differences are unclear, we can assume that various disease processes such as demyelination, axonal degeneration, gliosis, and edema³⁶ play a role. Because FA is nonspecific for demyelination and other pathologic processes³⁷ and ISO reportedly shows inflammatory free-water or edema,⁵ this combination of synthetic MR imaging and diffusion MR imaging may allow more detailed evaluation than is achieved using FA only.

TBSS analysis did not show statistically significant differences in the AVF, ICVF, and g-ratio between patients with MS and healthy controls, while MVF differed significantly. Furthermore, in a comparison between the NAWM of patients and the WM of healthy controls, ROI analysis showed significant differences in MVF, but not in the AVF, ICVF, and g-ratio. These results agreed with those of our previous study³ reporting that MVF was more sensitive than AVF to WM damage in plaque and periplaque WM than in NAWM. Even though TBSS analysis showed no significant difference in axial and radial diffusivity between patients with MS and healthy controls in our study, ROI analysis revealed higher radial diffusivity in the NAWM of patients with MS than in healthy control WM, whereas axial diffusivity did not differ significantly, even in ROI analysis. Our observation was consistent with previously reported findings that radial diffusivity showed more widespread abnormalities in the WM of patients with MS than axial diffusivity,³⁴ where radial diffusivity and axial diffusivity are considered surrogate markers for myelin and axon integrity, respectively. The widespread decrease in MVF, without a significant AVF decrease in this study, supported the notion that demyelination is a key pathologic element in MS and that MVF can be a biomarker for monitoring progression and treatment response in MS.

In ROI analysis, the ODI was higher in the NAWM of patients with MS than in the healthy control WM, but it was lower in WM lesions in patients with MS than in healthy control WM. This was congruent with the findings of Schneider et al³⁸ involving 5 patients with relapsing-remitting MS, which indicated a loss of fiber coherence (ie, increased dispersion) in the NAWM. They found a lower ODI in WM lesions, as in our study, even though another study involving early-stage MS (disease duration, <5 years) showed higher ODI in WM lesions.⁵ These results might suggest that NAWM and early plaques show a loss of fiber coherence with relatively maintained neuronal fiber density, leading to an increase in ODI, whereas loss of neuronal fibers, with fewer signals from fibers, may cause a decrease in ODI in chronic plaques.

ISO values were higher in the WM of patients with MS on TBSS analysis and in the NAWM of patients with MS on ROI analysis, than in healthy control WM, possibly indicating edema with free diffusion in patients' NAWM. However, patients' WM lesion values were not significantly higher than those of healthy control WM, in contrast to the findings of Schneider et al.³⁸ This difference could be attributed to the high variability of ISO in lesions in our study (Fig 2), which is partly explained by the partial volume effects of lesion ROIs. Chung et al³⁹ previously revealed that NODDI metrics showed higher variability at tissue boundaries. Hence, future investigations with higher resolution are warranted.

In TBSS, R1, R2, and proton density, as opposed to the MVF derived from these metrics, did not differ significantly between the patients with MS and healthy controls, agreeing with a previous observation showing the higher sensitivity of MVF than R1, R2, and proton density for revealing WM damage in plaque and periplaque WM.² On ROI analysis, only R1 differed significantly between the NAWM of patients with MS and the WM of healthy

controls. Previous reports have inconsistently reported differences between healthy WM and NAWM revealed by these metrics,^{40–43} and 1 previous study showed significant differences in T2 and proton density, but not in T1, between healthy WM and NAWM.⁴³ These metrics are thus nonspecifically affected by pathologic processes; hence, MVF and AVF may be more suitable for capturing changes in tissue microstructure in patients with MS.

In our study, no correlation was found between quantitative metrics and EDSS or disease duration. This could be due to the small sample size and the low EDSS scores (median, 1). Larger studies including a wider range of clinical disabilities are needed.

This study had some limitations. First, the study design was cross-sectional. Future longitudinal studies are required to understand intraindividual variation and the trajectory of metric changes during disease progression. Second, we used a 4-mm section thickness with a 1-mm gap. A future combination of 3D synthetic MR imaging⁴⁴ and high-resolution diffusion-weighted imaging in a larger study population would facilitate a more robust analysis.

CONCLUSIONS

We compared the metrics acquired by synthetic MR imaging and NODDI in the WM between patients with MS and healthy controls. Compared with healthy controls, patients with MS showed significantly lower FA, lower MVF, and higher ISO in large WM areas. Furthermore, a decrease in FA and MVF showed different spatial distributions; combined, these quantitative values can facilitate better WM characterization in patients with MS.

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