

## On-line Appendix

### MR Imaging Acquisition

Conventional MR imaging and diffusion tensor MR imaging scans of the brain were acquired from all subjects using identical 1.5T magnetic field strengths (Avanto; Siemens, Erlangen, Germany). The following MR imaging sequences of the brain were acquired from all subjects: a) axial dual-echo (DE) turbo spin-echo (TSE) (TR = 2650 ms, TE = 28–113 ms, echo-train length = 5, number of sections = 50, section thickness = 2.5 mm with no gap, matrix size =  $512 \times 512$ , FOV =  $250 \times 250$ -mm<sup>2</sup>), b) axial pulsed-gradient spin-echo (PGSE) echo-planar diffusion MR imaging sequence (TR = 6400 ms, TE = 93 ms, number of sections = 40, section thickness = 2.5 mm with no gap, matrix size =  $128 \times 128$ , FOV =  $240 \times 240$  mm<sup>2</sup>), with diffusion-encoding gradients applied in 12 non-collinear directions (b factor = 1000 sec/mm<sup>2</sup>, number of averages = 8), and c) sagittal 3D T1-weighted magnetization-prepared rapid acquisition gradient echo (MPRAGE) (TR = 2000 ms, TE = 3.93 ms, TI = 1100 ms, number of sections = 208, section thickness = 0.9 mm, matrix size =  $256 \times 224$ , FOV =  $236 \times 270$  mm<sup>2</sup>). All the scans were positioned following published guidelines.<sup>1</sup> Diffusion-weighted images were positioned with the same orientation as the DE scan, with the central section positioned to match the central section of the DE set.

In center B, using the same scanner, the following pulse sequences of the cervical cord were also acquired using a tailored cervical spine phased array coil for signal reception: a) sagittal T2-weighted TSE (TR = 4130 ms, TE = 104 ms, echo-train length = 25, number of sections = 12, section thickness = 3.0 mm, intersection gap = 0.3 mm, matrix =  $448 \times 336$ , FOV =  $280 \times 280$  mm<sup>2</sup>); b) sagittal 3D T1-weighted MPRAGE (TR = 1160 ms, TE = 4.24 ms, TI = 600 ms, number of sections = 88, section thickness = 0.9 mm, matrix size =  $256 \times 128$ , FOV =  $115 \times 230$  mm<sup>2</sup>); and c) axial 2D gradient-echo (TR = 640 ms, TE = 12 ms, number of averages = 2, number of sections = 20, section thickness = 5.0 mm, intersection gap = 0 mm, matrix size =  $256 \times 256$ , FOV =  $250 \times 250$  mm<sup>2</sup>) with and without an off-resonance radio-frequency saturation pulse (magnetic field strength = 1.5 kHz, flip angle = 20°).

### MR Imaging Postprocessing

All images were assessed by consensus of 2 experienced observers blinded to patient identity. Brain T2 lesion loads (LL) were measured using a local thresholding segmentation technique (Jim; Xinapse System, Leicester, United Kingdom). On brain MPRAGE scans, normalized brain volume (NBV) was measured using the cross-sectional version of the fully automated Structural Imaging Evaluation of Normalized Atrophy (SIENAX) software.<sup>2</sup>

Using an in-house software, from diffusion-weighted images, the diffusion tensor was estimated for each voxel using linear regression,<sup>3</sup> and mean diffusivity (MD) and fractional anisotropy (FA) maps derived.<sup>4</sup> MD histograms of the normal-appearing white matter (NAWM) and gray matter (GM) and FA histograms of the NAWM were produced as previously described.<sup>5</sup> FA histograms were derived only for the NAWM, because no preferential direction of water molecular motion is

expected to occur in the GM, due to the absence of a microstructural anisotropic organization of this tissue compartment. For each histogram, the average MD and FA were measured.

A FA atlas was created based on data from 24 healthy subjects (15 women and 9 men, mean age = 31.8 years, range = 21–40), with no previous history of neurologic dysfunction (reference group).<sup>6–8</sup> Then, using diffusion tensor MR imaging tractography, probability maps of the corpus callosum, corticospinal tract, thalamocortical connection (TCC), inferior fronto-occipital fasciculus (IFOF), uncinate fasciculus, cingulum, arcuate fasciculus, inferior longitudinal fasciculus (ILF), optic radiation (OR), superior cerebellar peduncle (SCP), and middle cerebellar peduncle (MCP) were produced, as described elsewhere.<sup>6–8</sup> MD and FA maps from controls, as well as MD, FA and lesion maps from patients were non-linearly transformed<sup>9</sup> to the FA atlas, using the FA maps to calculate the transformation. T2-visible lesions were removed from MD and FA maps, and WM fiber bundle probability maps were applied to patients' data to obtain average MD and FA of the tracts of interest. For white matter (WM) fiber bundles with a bilateral location in the brain, the averages of the MD and FA values measured in the right and left hemisphere entered the analysis. Volumes of T2-visible lesion in the different WM fiber bundles were derived by applying fiber bundle probability maps, obtained using diffusion tensor MR imaging tractography, and calculating the volume of lesions located inside each of them.<sup>6</sup> Figure 1 in the text shows an example of WM fiber bundle reconstruction in an individual subject prior to constructing the probability maps.

Cervical cord hyperintense lesions were identified on the sagittal T2-weighted scans. The original cervical cord MPRAGE data were reformatted and a set of 5 contiguous, 3-mm-thick axial sections were reconstructed using the center of the C2-C3 disk as the caudal landmark. Then a semiautomated technique was used to segment the cord tissue and to measure the cross-sectional cord area at the level of each section.<sup>10</sup> Values from the 5 sections were averaged to obtain a single value for each subject.

After coregistration of the 2 cord gradient-echo scans (ie, with and without the magnetization transfer > saturation pulse), MT ratio (MTR) images were derived pixel by pixel, using an automated technique based on pixel similarity measures. Extra cord tissue was then removed from the MTR maps using a local thresholding segmentation technique (Jim; Xinapse System) and the corresponding average MTR values obtained.<sup>11</sup>

### Statistical Analysis

Between-group differences were assessed using the nonparametric Kruskal-Wallis test. Correlations between clinical and MR-derived quantities were estimated using the Spearman rank correlation coefficient.

Random forest analysis was performed to classify clinically impaired vs. unimpaired patients using a set of MR imaging covariates (including measures of global brain and cord damage as well as selective damage to brain WM fiber bundles). Because the number of observations for each value of the functional system and EDSS scales was small, potentially leading to unreliable results, we dichotomized the FS into impaired (FS

≥ 1) and nonimpaired (FS = 0) and the EDSS scale according to a cutoff of 4.0 (which identifies fully ambulatory patients). According to the random forest technique, 100 000 trees were built to classify compromised patients.<sup>12</sup> The training set used to grow each tree was a .632+ bootstrap resample of the observations.<sup>13</sup> Trees were allowed to grow to their full size without pruning; nodes with at least 1 event and minimum total size of 5 were used as stopping rules. The best split at each node was selected from a random subset of covariates. The left-out observations (ie, “out of bag” observations) were then predicted to obtain the classification error rate of the considered tree. Predictive ability of the classifier was assessed aggregating the single tree error rates. Furthermore, the random forest framework allowed us to estimate the importance of a variable by looking at how much the classification error increases when “out of bag” data for that variable are permuted while all others are left unchanged. We followed Strobl et al<sup>14</sup> to avoid possible bias in variable selection: individual classification trees were built using subsampling without replacement and adopting a conditional permutation scheme.<sup>15</sup>

The goodness of the fit of the classifier is reported as an “error rate,” which is the rate of misclassified patients. When the random forest shows a large error rate, this may be due to at least 2 reasons: 1) the covariates do not explain well the patient’s status, 2) the sample design is very unbalanced (eg, 95% compromised and 5% non-compromised patients). An output of the random forest corresponds to variable importance reported as a ranking: each covariate gets a score according to its ability to classify correctly the patient according to the decrease of classification accuracy. When the random forest achieved an error rate close to 100% in 1 of the outcome classes, the derived variable importance was not considered reliable.

Because cervical cord MR imaging was acquired in 1 center only, subgroup analysis was also required. Therefore, all the random forest analyses were re-run, including data from cen-

ter B only, in order to confirm and/or understand better the importance of cord MR-derived variables.

A *P* value < .05 was considered as significant. All analyses were performed using SAS Release 9.1. For the random forest analysis, we used the package ‘randomForest’ version 4.5 implemented in R software (A language and environment for statistical computing; version 2.12.0).

## References

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**On-line Table 1: Main demographic and clinical findings from the cohorts of patients with MS studied**

	CIS (22)	RRMS (51)	SPMS (44)	BMS (20)	PPMS (35)
Men	27.3%	25.5%	47.7%	20.0%	51.0%
Women	72.7%	74.5%	52.3%	80.0%	49.0%
Mean age (range) (yr)	29.5 (19–43)	41.4 (22–68)	49.5 (28–67)	45.0 (31–59)	46.7 (26–70)
Mean disease duration (range) (yr)	0.06 (0.01–0.25)	10.3 (1–30)	17.8 (4–38)	23.3 (15–40)	9.7 (1–29)
Median EDSS score (range)	2.0 (0.0–4.0)	2.5 (0.0–4.0)	6.5 (3.5–8.5)	2.0 (0.0–3.0)	5.5 (2.0–7.5)

**Note:**—CIS indicates clinically isolated syndromes.

**On-line Table 2: Number of patients with impairment in the different functional systems of the EDSS**

Functional systems		CIS	RRMS	SPMS	BMS	PPMS
Pyramidal	0	5	5	0	3	0
	1	9	11	1	12	2
	2	2	7	2	5	4
	3	6	18	11	0	13
	4	0	10	23	0	13
	5	0	0	7	0	3
	6	0	0	0	0	0
Cerebellar	0	14	18	1	8	10
	1	3	6	3	6	2
	2	4	11	14	6	10
	3	1	12	24	0	11
	4	0	4	2	0	2
Brain stem	0	15	18	5	15	9
	1	4	7	8	3	7
	2	3	18	17	2	14
	3	0	7	13	0	5
	4	0	1	1	0	0
Sensory	0	7	12	6	11	7
	1	4	12	7	6	6
	2	10	17	14	3	14
	3	1	9	15	0	8
	4	0	1	2	0	0
	5	0	0	0	0	0
Bowel and bladder	0	19	27	9	17	10
	1	3	11	14	2	12
	2	0	9	11	1	7
	3	0	2	7	0	4
	4	0	1	2	0	1
	5	0	1	1	0	1
Visual	0	17	46	29	10	26
	1	3	2	7	9	5
	2	2	3	8	1	4
	3	0	0	0	0	0
	4	0	0	0	0	0
	5	0	0	0	0	0
Cerebral (mental)	0	22	40	25	18	27
	1	0	5	5	2	2
	2	0	6	11	0	6
	3	0	0	3	0	0
	4	0	0	0	0	0
	5	0	0	0	0	0
Ambulation Index	0	15	24	10	14	0
	1	2	12	10	4	8
	2	4	14	5	2	6
	3	1	1	3	0	10
	4	0	0	1	0	3
	5	0	0	2	0	1
	6	0	0	2	0	1
	7	0	0	4	0	1
	8	0	0	4	0	2
9	0	0	3	0	3	

**Note:**—CIS indicates clinically isolated syndromes.

**On-line Table 3: Structural MR imaging findings from healthy controls and patients with MS studied<sup>2</sup>**

	Healthy Controls	All Patients	CIS Patients	RRMS Patients	SPMS Patients	BMS Patients	PPMS Patients	<i>P</i> <sup>b</sup>	
T2 LL (ml) (SD)	–	25.3 (27.0)	4.1 (6.1)	19.2 (17.9)	43.8 (29.5)	30.9 (35.4)	21.5 (23.9)	<0.0001	
NBV (ml) (SD)	1562.9 (22.9)	1499.7 (113.9)	1621.3 (96.6)	1519.4 (107.2)	1425.4 (89.7)	1480.2 (84.4)	1498.8 (104.3)	<0.0001	
AL FA (SD)	–	0.31 (0.04)	0.33 (0.05)	0.31 (0.04)	0.28 (0.03)	0.34 (0.02)	0.32 (0.04)	<0.0001	
AL MD (SD)	–	–	1.00 (0.1)	1.07 (0.1)	1.18 (0.1)	1.03 (0.1)	1.07 (0.1)	<0.0001	
NAWM Average FA (SD)	0.40 (0.03)	0.37 (0.04)	0.41 (0.02)	0.37 (0.05)	0.34 (0.03)	0.38 (0.3)	0.37 (0.04)	<0.0001	
NAWM Average MD (SD)	0.73 (0.02)	0.78 (0.05)	0.75 (0.02)	0.76 (0.05)	0.81 (0.06)	0.78 (0.05)	0.77 (0.03)	<0.0001	
GM Average MD (SD)	0.89 (0.04)	0.96 (0.09)	0.89 (0.04)	0.95 (0.08)	1.02 (0.13)	0.96 (0.06)	0.96 (0.07)	<0.0001	
Number of cervical cord lesions (range)	0	1.1 (0–6)	0.5 (0–3)	1.5 (0–6)	2 (0–5)	1 (0–4)	1 (0–5)	n.s.	
Cervical cord area (ml) (SD)	68.1 (8.5)	62.9 (9.9)	69.2 (8.8)	68.6 (7.6)	56.7 (8.0)	64.2 (7.8)	56.7 (10.0)	<0.0001	
Cervical cord MTR (%) (SD)	48.7 (1.5)	46.1 (2.2)	47.8 (1.2)	47.2 (1.4)	44.6 (2.2)	45.7 (2.0)	45.1 (2.3)	<0.0001	
CC	T2 LL (ml) (SD)	–	0.82 (0.8)	0.14 (0.2)	0.66 (0.8)	1.50 (0.9)	0.74 (0.6)	0.68 (0.7)	<0.0001
	AL FA (SD)	–	0.45 (0.08)	0.55 (0.10)	0.45 (0.07)	0.40 (0.06)	0.47 (0.06)	0.47 (0.08)	<0.0001
	AL MD (SD)	–	0.99 (0.14)	0.86 (0.12)	0.98 (0.14)	1.10 (0.14)	0.96 (0.10)	0.98 (0.13)	<0.0001
	NAWM FA (SD)	0.63 (0.05)	0.58 (0.07)	0.63 (0.03)	0.59 (0.07)	0.52 (0.09)	0.60 (0.05)	0.60 (0.05)	<0.0001
Arcuate fasciculus	NAWM MD (SD)	0.74 (0.03)	0.85 (0.10)	0.77 (0.03)	0.83 (0.17)	0.95 (0.20)	0.83 (0.08)	0.82 (0.08)	<0.0001
	T2 LL (ml) (SD)	–	0.2 (0.20)	0.02 (0.04)	0.2 (0.20)	0.4 (0.30)	0.3 (0.03)	0.1 (0.20)	<0.0001
	AL FA (SD)	–	0.38 (0.09)	0.29 (0.13)	0.39 (0.09)	0.39 (0.07)	0.40 (0.07)	0.38 (0.11)	n.s.
	AL MD (SD)	–	0.84 (0.2)	0.58 (0.2)	0.82 (0.2)	0.95 (0.2)	0.89 (0.2)	0.80 (0.2)	<0.0001
Cingulum	NAWM FA (SD)	0.50 (0.04)	0.47 (0.04)	0.50 (0.03)	0.48 (0.04)	0.45 (0.04)	0.48 (0.04)	0.48 (0.03)	<0.0001
	NAWM MD (SD)	0.71 (0.03)	0.77 (0.06)	0.74 (0.02)	0.75 (0.06)	0.80 (0.06)	0.78 (0.07)	0.76 (0.04)	<0.0001
	T2 LL (ml) (SD)	–	0.01 (0.10)	0.01 (0.07)	0.01 (0.06)	0.02 (0.30)	0.01 (0.07)	0.01 (0.06)	<0.0001
	AL FA (SD)	–	0.25 (0.07)	0.23 (0.05)	0.25 (0.07)	0.25 (0.07)	0.27 (0.10)	0.26 (0.07)	n.s.
IFOF	AL MD (SD)	–	0.53 (0.20)	0.42 (0.02)	0.51 (0.20)	0.65 (0.30)	0.46 (0.20)	0.47 (0.09)	0.01
	NAWM FA (SD)	0.56 (0.05)	0.52 (0.06)	0.56 (0.03)	0.53 (0.06)	0.48 (0.07)	0.54 (0.05)	0.55 (0.05)	<0.0001
	NAWM MD (SD)	0.73 (0.03)	0.78 (0.07)	0.76 (0.03)	0.76 (0.05)	0.81 (0.10)	0.78 (0.05)	0.77 (0.04)	<0.0001
	T2 LL (ml) (SD)	–	0.09 (0.10)	0.01 (0.03)	0.08 (0.10)	0.2 (0.10)	0.09 (0.10)	0.07 (0.10)	<0.0001
ILF	AL FA (SD)	–	0.38 (0.09)	0.38 (0.10)	0.38 (0.10)	0.38 (0.07)	0.36 (0.10)	0.37 (0.10)	n.s.
	AL MD (SD)	–	0.94 (0.3)	0.73 (0.3)	0.93 (0.3)	1.07 (0.2)	0.86 (0.3)	0.90 (0.3)	<0.0001
	NAWM FA (SD)	0.55 (0.05)	0.50 (0.05)	0.56 (0.04)	0.51 (0.06)	0.46 (0.05)	0.51 (0.04)	0.52 (0.04)	<0.0001
	NAWM MD (SD)	0.77 (0.03)	0.85 (0.07)	0.79 (0.04)	0.83 (0.07)	0.90 (0.07)	0.85 (0.07)	0.84 (0.05)	<0.0001
MCP	T2 LL (ml) (SD)	–	0.8 (0.90)	0.01 (0.02)	0.06 (0.08)	0.14 (0.10)	0.08 (0.10)	0.06 (0.08)	<0.0001
	AL FA (SD)	–	0.34 (0.09)	0.33 (0.08)	0.30 (0.10)	0.35 (0.06)	0.39 (0.08)	0.34 (0.10)	n.s.
	AL MD (SD)	–	0.92 (0.3)	0.76 (0.2)	0.81 (0.3)	1.07 (0.2)	0.97 (0.2)	0.87 (0.3)	0.0002
	NAWM FA (SD)	0.47 (0.04)	0.43 (0.05)	0.48 (0.04)	0.44 (0.04)	0.40 (0.06)	0.44 (0.04)	0.44 (0.04)	<0.0001
SCP	NAWM MD (SD)	0.76 (0.04)	0.86 (0.08)	0.81 (0.05)	0.83 (0.08)	0.90 (0.08)	0.88 (0.09)	0.85 (0.07)	<0.0001
	T2 LL (ml) (SD)	–	0.03 (0.05)	0.01 (0.03)	0.03 (0.05)	0.05 (0.07)	0.02 (0.06)	0.03 (0.04)	<0.0001
	AL FA (SD)	–	0.34 (0.01)	0.32 (0.10)	0.34 (0.10)	0.39 (0.10)	0.32 (0.20)	0.32 (0.20)	n.s.
	AL MD (SD)	–	0.59 (0.2)	0.54 (0.2)	0.55 (0.2)	0.68 (0.2)	0.51 (0.2)	0.57 (0.2)	n.s.
Uncinate fasciculus	NAWM FA (SD)	0.55 (0.04)	0.53 (0.04)	0.56 (0.02)	0.54 (0.03)	0.50 (0.04)	0.54 (0.03)	0.54 (0.03)	<0.0001
	NAWM MD (SD)	0.71 (0.02)	0.74 (0.05)	0.72 (0.02)	0.74 (0.05)	0.77 (0.05)	0.75 (0.05)	0.74 (0.04)	<0.0001
	T2 LL (ml) (SD)	–	0.01 (0.01)	0.01 (0.01)	0.01 (0.01)	0.01 (0.01)	0.01 (0.01)	0.01 (0.01)	<0.0001
	AL FA (SD)	–	0.30 (0.01)	0.35 (0.1)	0.26 (0.1)	0.31 (0.1)	0.26 (0.1)	0.31 (0.1)	n.s.
CST	AL MD (SD)	–	0.6 (0.2)	0.62 (0.2)	0.58 (0.2)	0.76 (0.3)	0.57 (0.2)	0.62 (0.3)	n.s.
	NAWM FA (SD)	0.55 (0.04)	0.53 (0.04)	0.55 (0.03)	0.53 (0.04)	0.51 (0.05)	0.56 (0.05)	0.54 (0.03)	0.0004
	NAWM MD (SD)	0.87 (0.04)	0.93 (0.07)	0.90 (0.06)	0.91 (0.07)	0.97 (0.08)	0.93 (0.07)	0.93 (0.05)	<0.0001
	T2 LL (ml) (SD)	–	0.01 (0.001)	0.01 (0.007)	0.01 (0.02)	0.02 (0.02)	0.01 (0.008)	0.01 (0.006)	<0.0001
CST	AL FA (SD)	–	0.22 (0.08)	0.16 (0.05)	0.20 (0.08)	0.23 (0.07)	0.21 (0.10)	0.25 (0.08)	n.s.
	AL MD (SD)	–	0.74 (0.3)	0.61 (0.2)	0.70 (0.4)	0.82 (0.4)	0.66 (0.2)	0.71 (0.2)	n.s.
	NAWM FA (SD)	0.42 (0.04)	0.40 (0.04)	0.43 (0.03)	0.40 (0.04)	0.37 (0.04)	0.41 (0.03)	0.41 (0.03)	<0.0001
	NAWM MD (SD)	0.78 (0.03)	0.84 (0.07)	0.82 (0.04)	0.83 (0.07)	0.88 (0.08)	0.85 (0.09)	0.83 (0.05)	<0.0001
TCC	T2 LL (ml) (SD)	–	0.06 (0.05)	0.01 (0.01)	0.06 (0.06)	0.1 (0.08)	0.07 (0.07)	0.04 (0.04)	<0.0001
	AL FA (SD)	–	0.44 (0.08)	0.44 (0.05)	0.44 (0.09)	0.41 (0.05)	0.48 (0.04)	0.45 (0.10)	0.05
	AL MD (SD)	–	0.87 (0.1)	0.76 (0.2)	0.88 (0.1)	0.91 (0.1)	0.85 (0.09)	0.83 (0.08)	0.001
	NAWM FA (SD)	0.63 (0.03)	0.62 (0.03)	0.62 (0.02)	0.61 (0.02)	0.61 (0.04)	0.62 (0.02)	0.61 (0.02)	n.s.
TCC	NAWM MD (SD)	0.70 (0.03)	0.74 (0.05)	0.72 (0.02)	0.73 (0.05)	0.76 (0.05)	0.75 (0.05)	0.73 (0.03)	<0.0001
	T2 LL (ml) (SD)	–	0.02 (0.05)	0.001 (0.001)	0.03 (0.04)	0.05 (0.04)	0.03 (0.03)	0.02 (0.02)	<0.0001
	AL FA (SD)	–	0.46 (0.09)	0.47 (0.1)	0.43 (0.07)	0.43 (0.06)	0.48 (0.09)	0.48 (0.1)	0.03
	AL MD (SD)	–	0.91 (0.1)	0.81 (0.06)	0.91 (0.1)	0.95 (0.1)	0.88 (0.1)	0.88 (0.1)	0.001
OR	NAWM FA (SD)	0.60 (0.04)	0.58 (0.03)	0.59 (0.02)	0.58 (0.02)	0.58 (0.04)	0.58 (0.03)	0.58 (0.03)	0.02
	NAWM MD (SD)	0.71 (0.03)	0.74 (0.05)	0.74 (0.02)	0.74 (0.06)	0.78 (0.05)	0.77 (0.05)	0.75 (0.03)	<0.0001
	T2 LL (ml) (SD)	–	0.05 (0.06)	0.01 (0.01)	0.05 (0.06)	0.10 (0.06)	0.06 (0.06)	0.05 (0.06)	<0.0001
	AL FA (SD)	–	0.40 (0.08)	0.42 (0.10)	0.39 (0.09)	0.38 (0.06)	0.43 (0.06)	0.40 (0.08)	n.s.
OR	AL MD (SD)	–	1.06 (0.2)	0.86 (0.3)	1.02 (0.3)	1.21 (0.2)	1.02 (0.2)	1.02 (0.3)	<0.0001
	NAWM FA (SD)	0.58 (0.05)	0.50 (0.07)	0.57 (0.04)	0.50 (0.07)	0.45 (0.07)	0.50 (0.06)	0.51 (0.06)	<0.0001
	NAWM MD (SD)	0.82 (0.04)	0.96 (0.10)	0.86 (0.05)	0.95 (0.10)	1.05 (0.20)	0.97 (0.10)	0.94 (0.07)	<0.0001

**Note:**—MS indicates multiple sclerosis; CIS, clinically isolated syndromes; B, benign form; PP, primary-progressive; RR, relapsing-remitting; SP, secondary-progressive; LL, lesion load; SD, standard deviation; NBV, normalized brain volume; MD, mean diffusivity; FA, fractional anisotropy; NAWM, normal-appearing white matter; GM, gray matter; MTR, magnetization transfer ratio; AL, average lesion; CST, corticospinal tract; TCC, thalamocortical connection; CC, corpus callosum; IFOF, inferior fronto-occipital fasciculus; ILF, inferior longitudinal fasciculus; MCP, middle cerebellar peduncle; SCP, superior cerebellar peduncle; OR, optic radiation; n.s., not significant.

<sup>a</sup> Average MD is expressed in units of  $\text{mm}^2\text{s}^{-1} \times 10^{-3}$ , FA is a dimensionless index.

<sup>b</sup> Non-parametric Kruskal-Wallis test.

**On-line Table 4: Results of the random forest analysis<sup>a,b</sup>**

Clinical Variable	MS			CIS			RRMS			SPMS			BMS			PPMS		
EDSS	18.0	26.2	10.9	13.6	5	100	14.2	100	0	39.2	25.0	63.2				6.8	100	0
	NAWM FA <sup>c</sup>			Cingulum AL FA			Cingulum AL FA			SCP NAWM MD <sup>c</sup>			-			CC NAWM MD		
	CC NAWM FA <sup>c</sup>			SCP NAWM FA			Cingulum AL MD			CC T2 LL <sup>c</sup>						OR NAWM MD		
	CC T2 LL <sup>c</sup>			NAWM MD			MCP AL MD			Cingulum NAWM FA <sup>c</sup>						Uncinate NAWM MD		
	MCP NAWM FA <sup>c</sup>			NBV			CC NAWM MD			SCP NAWM FA <sup>c</sup>						CC NAWM FA		
Ambulation Index	30.7	41.4	22.8	35.0	23.1	57.1	42.8	21.4	85.7	-	-	-	50.0	28.6	100	-	-	-
	NAWM FA <sup>c</sup>			CST AL FA <sup>c</sup>			Cord MTR <sup>c</sup> MCP <sup>c</sup>			-			NAWM FA			-		
	Cord area MCP <sup>c</sup>			SCP NAWM FA <sup>c</sup>			NAWM MD <sup>c</sup>			-			NAWM MD			-		
	NAWM FA <sup>c</sup>			CST T2 LL <sup>c</sup>			NBV <sup>c</sup>			-			CST NAWM MD			-		
	SCP NAWM MD <sup>c</sup>			CST AL MD <sup>c</sup>			NAWM FA <sup>c</sup>			-			MCP NAWM FA			-		
Brain stem FS	21.5	43.5	9.1	36.4	26.7	57.1	13.7	27.8	6.0	13.6	100	2.5	30.0	20.0	60.0	22.8	66.7	7.7
	NAWM FA <sup>c</sup>			CST T2 LL <sup>c</sup>			CST NAWM <sup>c</sup> FA <sup>c</sup>			SCP NAWM MD			GM MD <sup>c</sup>			CST NAWM <sup>c</sup> FA <sup>c</sup>		
	MCP NAWM FA <sup>c</sup>			MCP T2 LL <sup>c</sup>			NBV <sup>c</sup>			SCP NAWM FA			NAWM FA <sup>c</sup>			NBV <sup>c</sup>		
	NBV <sup>c</sup>			T2 LL <sup>c</sup>			SCP NAWM MD <sup>c</sup>			OR NAWM FA			NAWM MD <sup>c</sup>			SCP NAWM MD <sup>c</sup>		
	MCP T2 LL <sup>c</sup>			NAWM MD <sup>c</sup>			CST T2 LL <sup>c</sup>			OR NAWM MD			SCP NAWM FA <sup>c</sup>			CST T2 LL <sup>c</sup>		
Cerebellar FS	19.3	43.5	5.8	27.3	14.3	50.0	11.7	16.7	9.1	-	-	-	25.0	37.5	16.7	22.8	60.0	8.0
	T2 LL <sup>c</sup>			MCP NAWM <sup>c</sup> MD <sup>c</sup>			NBV <sup>c</sup>			-			NAWM MD <sup>c</sup>			T2 LL <sup>c</sup>		
	CST NAWM <sup>c</sup> MD <sup>c</sup>			CST NAWM FA <sup>c</sup>			CST NAWM <sup>c</sup> MD			-			SCP NAWM FA <sup>c</sup>			SCP NAWM MD <sup>c</sup>		
	SCP NAWM FA <sup>c</sup>			SCP NAWM MD <sup>c</sup>			T2 LL <sup>c</sup>			-			MCP NAWM FA <sup>c</sup>			OR NAWM MD <sup>c</sup>		
	NAWM FA <sup>c</sup>			SCP NAWM FA <sup>c</sup>			SCP NAWM MD <sup>c</sup>			-			SCP NAWM MD <sup>c</sup>			MCP T2 LL <sup>c</sup>		
Pyramidal FS	7.5	100	0	22.7	100	0	9.8	100	0	-	-	-	20.0	100.0	5.8	-	-	-
	NBV			CST AL MD			NBV			-			CST T2 LL			-		
	CST T2 LL			CC NAWM MD			MCP NAWM FA			-			CST NAWM FA			-		
	MCP NAWM FA			CC NAWM FA			OR NAWM MD			-			CC NAWM FA			-		
	CC NAWM FA			CST T2 LL			OR NAWM FA			-			NAWM FA			-		
Visual FS	27.3	5.5	91.0	22.7	0	100	9.8	0	100	50.0	31.0	86.7	30.0	30.0	30.0	28.6	7.7	88.9
	CST NAWM <sup>c</sup> MD <sup>c</sup>			NAWM MD			OR T2 LL			SCP NAWM <sup>c</sup> MD <sup>c</sup>			NAWM MD <sup>c</sup>			SCP NAWM <sup>c</sup> MD <sup>c</sup>		
	CC NAWM MD <sup>c</sup>			CC NAWM MD			MCP NAWM FA			OR NAWM FA <sup>c</sup>			CC NAWM MD <sup>c</sup>			OR NAWM FA <sup>c</sup>		
	OR NAWM FA <sup>c</sup>			GM MD			SCP NAWM FA			OR AL MD <sup>c</sup>			CST NAWM MD <sup>c</sup>			NAWM MD <sup>c</sup>		
	OR NAWM MD <sup>c</sup>			NBV			NAWM MD			NAWM MD <sup>c</sup>			CST NAWM FA <sup>c</sup>			OR NAWM <sup>c</sup> MD <sup>c</sup>		
Sensory FS	25.5	88.4	4.6	9.0	0	13.0	17.6	33.3	12.8	18.2	100	5.2	70.0	63.6	66.6	25.7	100	7.1
	TCC AL FA <sup>c</sup>			TCC AL FA <sup>c</sup>			Cord MTR <sup>c</sup>			NAWM FA			MCP NAWM FA <sup>c</sup>			NAWM FA		
	NAWM FA <sup>c</sup>			TCC AL MD <sup>c</sup>			Cord area <sup>c</sup>			SCP NAWM MD			Cord MTR <sup>c</sup>			NAWM MD		
	TCC T2 LL <sup>c</sup>			NBV <sup>c</sup>			TCC NAWM <sup>c</sup> FA <sup>c</sup>			TCC AL MD			NBV <sup>c</sup>			MCP NAWM MD		
	NBV <sup>c</sup>			TCC T2 LL <sup>c</sup>			NBV <sup>c</sup>			NBV			MCP NAWM MD <sup>c</sup>			Arcuate NAWM MD		
Bowel and bladder FS	22.1	21.9	22.2	13.6	0	100	15.6	14.8	16.6	20.4	100	0	20.0	5.8	100	25.7	80.0	8.0
	Cord area <sup>c</sup>			CST AL FA			Cord area <sup>c</sup>			CST AL FA			CST NAWM FA			Uncinate <sup>c</sup> NAWM FA <sup>c</sup>		
	Cord MTR <sup>c</sup>			CST AL MD			Cord MTR <sup>c</sup>			Cord area			Uncinate NAWM FA			CST NAWM MD <sup>c</sup>		
	CST AL FA <sup>c</sup>			CST T2 LL			IFOF T2 LL <sup>c</sup>			NAWM MD			Cingulum NAWM FA			Cingulate <sup>c</sup> NAWM MD <sup>c</sup>		
	CC NAWM FA <sup>c</sup>			MCP NAWM MD			NBV <sup>c</sup>			T2 LL			IFOF NAWM FA			Uncinate T2 LL <sup>c</sup>		
Cerebral FS	21.5	6.8	70.0	-	-	-	27.4	7.5	100	29.5	28.0	31.6	15.0	5.5	100	25.7	3.7	100
	NAWM FA <sup>c</sup>			-			Cord MTR			CST T2 LL <sup>c</sup>			Cingulum T2 LL			Cingulum T2 LL		
	Cingulum T2 LL <sup>c</sup>			-			Cord area			Cord area <sup>c</sup>			CC NAWM MD			Arcuate NAWM MD		
	CC T2 LL <sup>c</sup>			-			T2 LL			T2 LL			ILF NAWM FA			NAWM FA		
	ILF T2 LL <sup>c</sup>			-			ILF NAWM FA			IFOF T2 LL <sup>c</sup>			IFOF NAWM FA			CST NAWM MD		

**Note:**—MTR indicates magnetization transfer ratio; IFOF, inferior fronto-occipital fasciculus; ILF, inferior longitudinal fasciculus; CIS, clinically isolated syndromes; FS, functional system; LL, lesion load; OR, optic radiation; AL, average lesion; TCC, thalamocortical connection; GE, global error; UE, unimpaired error; IE, impaired error.

<sup>a</sup> The divided cells represent the variable importance of GE (%), UE (%), and IE (%), respectively.

Empty cells are due to the absence of a group of patients (eg, 100% impaired and 0% unimpaired).

<sup>b</sup> Variables whose importance was considered reliable (the others were not considered reliable due to an error rate of 100% in 1 of the outcome classes).