Supplementary Information:

MRI-derived g-ratio and lesion severity in newly diagnosed multiple sclerosis

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Supplementary Methods

MR Imaging Data Acquisition

MRI Acquisition Parameters are outlined in supplementary table 1.

MR Preprocessing

Hyperintense white matter lesions (WML) on T2 FLAIR (Fluid Attenuated Inversion Recovery) automatically **FSL** were segmented using а combination of (5.0.1,http://www.fmrib.ox.ac.uk/fsl) and an in-house thresholding method (Figure 1C). Intracranial volumes (ICV) were segmented with FSL and cerebral normal-appearing white matter (NAWM) segmentation masks obtained with FreeSurfer (6.0,were https://surfer.nmr.mgh.harvard.edu/) from the 3D T1-weighted MPRAGE image. Along with the 3D T1-weighted MPRAGE, masks were registered to DW space with FSL FLIRT epi reg, binarised and NAWM masks were eroded by one voxel to reduce partial volume effects. Segmentations were manually checked and edited where necessary (Figure 2A). The cerebellum was not included due to technical inaccuracies.

MTsat (δ_{app}) and T1app parametric maps were calculated according to Helms et al¹, with inhouse code in MATLAB (R2018b, see https://doi.org/10.7488/ds/2965), and registered to the first volume (b0) of the diffusion data with the FSL FLIRT epi_reg script. The magnetisation transfer ratio (MTR) was calculated as:

$$MTR = 100 \cdot \frac{(S_{PD} - S_{MT})}{S_{PD}}$$

The diffusion data were processed with FSL (6.0.1) to extract the brain from the skull, remove bulk motion and eddy-current induced distortions, and register all volumes to the first diffusion volume. The NODDI toolbox (v1.0, mig.cs.ucl.ac.uk, MATLAB R2016b) was used to determine signal fractions from the diffusion MR-visible compartments.

Myelin Volume Fraction: Calculation of k from healthy control data

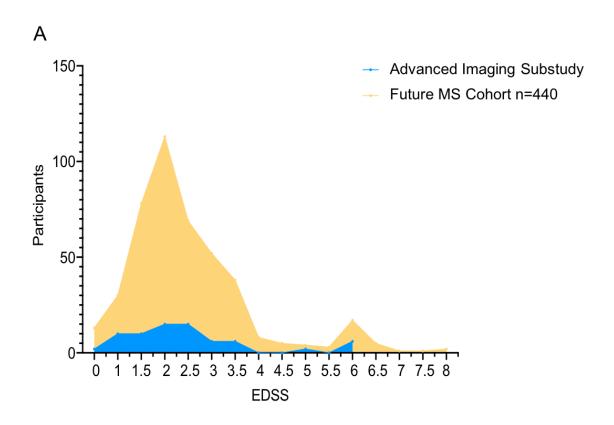
The constant, *k*, was derived by assuming a g-ratio of 0.7 in the splenium of the corpus callosum from two healthy controls. Healthy controls were scanned twice, one week apart. Data processing followed the same steps as patient data (without WML masks). The splenium mask was derived with FreeSurfer (https://surfer.nmr.mgh.harvard.edu/). The calibration factor was calculated as:

$$k = \frac{1}{\delta_{app}} * (1 - (\frac{1}{(1 + ((\frac{1}{0.7^2}) - 1)(1 - v_{iso})v_{ic})}))$$

The resulting mean k value across both sessions and subjects was 13.921 (Subject 1: 12.1717 and 14.24; Subject 2: 15.237 and 13.488, at time-point 1 and 2, respectively).

G-ratio Flat Lesion Maps

In order to assess the impact of lesion location on g-ratio, we created a 2D g-ratio distribution "flat" lesion maps for visual assessment. G-ratio parametric maps were masked with white matter lesion segmentations (FSLmaths), and imported into MATLAB (R2018b) with SPM12 functions. G-ratio values were averaged along the z-axis, creating a 2D array, which roughly approximates an individual's cerebral lesion distribution in the superior-to-inferior axis. Although, anatomically, this distribution will vary to some degree depending on the subject's position in the MR scanner, it allows quick visual assessment of any anterior-posterior bias in the distribution of g-ratio values. A colour-map was overlaid onto the 2D maps and the distribution of lesions and z-direction-averaged g-ratios across the four groups (high vs low lesion load/high vs. normal g-ratio) were compared visually.

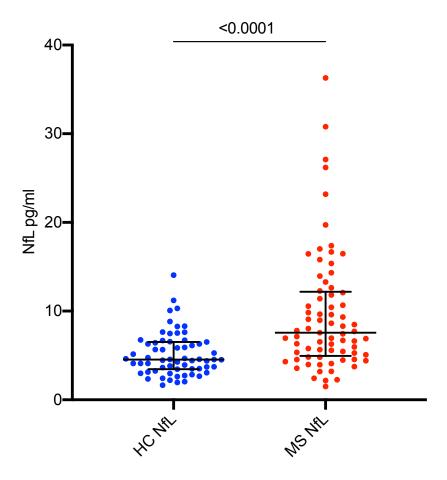


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	This cohort	Future MS
Participants	73	440
Sex ratio, F:M	3.6	2.82
Age at diagnosis	33 (28-45)	37 (30-45)
Time from symptom onset to study inclusion	34 (12-69)	22 (10-62)
	N=67/73	N=422/440
EDSS at baseline	2 (1.5-3)	2 (1.5-3)

Supplementary Figure 1. Baseline disability and demographics of Advanced Imaging Substudy of the Future MS cohort

EDSS distribution and demographic details of the Future-MS extended imaging substudy (n=73) compared to the Future MS (FMS) study (n=440, which include the 73 imaging substudy participants)



Supplementary Figure 2. Plasma neurofilament levels are elevated in newly diagnosed MS patients compared to age and sex-matched healthy controls (HC) (p<0.001, Mann-Whitney U-test)

High Lesion Load, Low g-ratio

Supplementary Figure 3. Flat map images of individual study participants with high lesion load (Corrected WML volume> 0.5%) and low g-ratio

In order to demonstrate 2D lesion distribution of patients with high lesion load and normal gratio in the axial plane, g-ratios within WML were averaged in the z-direction; red indicates an abnormally elevated g-ratio and blue is indicative of a normal g-ratio. There is no apparent difference in the distribution of lesion location in the axial plane between low and high g-ratio individuals (Supplementary Figure 4) with a high lesion load.

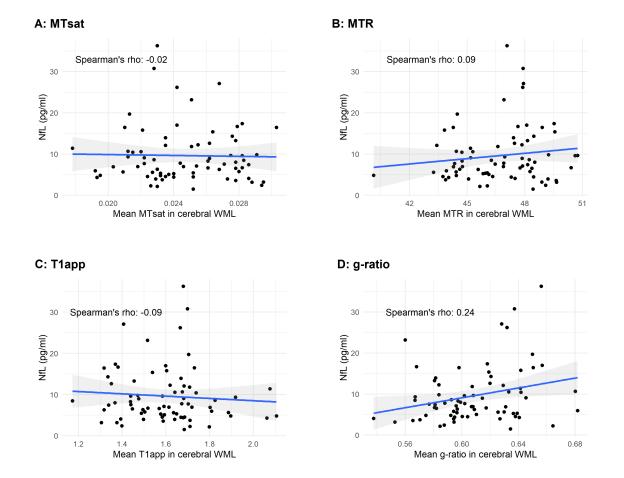
There was no significant difference in lesion volume between individuals with high and normal WML g-ratios for either the high lesion load group (0.998 vs 1.158, W=185, p=0.72, Wilcoxon Rank Sum Test) or the low lesion load group (0.299 vs 0.180, W=196, p=0.07).

Within the high lesion load group, the anatomical location of lesions did not differ, when assessed visually on 2D "flattened" lesion distribution maps

High Lesion Load, High g-ratio

Supplementary Figure 4. Flat map images of individual study participants with high lesion load (Corrected WML volume> 0.5%) and high g-ratio

As in Supplementary Figure 3 distribution of WML g-ratio in patients with high lesion load and abnormal g-ratio in the axial plane is illustrated by averaging WML g-ratio in the z-direction. Again, red indicates an abnormally elevated g-ratio and blue is indicative of a normal g-ratio. There is no apparent difference in the distribution of lesion location in the axial plane between low (Supplementary Figure 3) and high g-ratio individuals with a high lesion load.



Supplementary Figure 5. Correlation of MTsat, MTR, T1app and g-ratio with serum NFL

In nonparametric tests of correlation (Spearman's rho) g-ratio is significantly associated with NFL (r_s =0.24, p<0.05). This was not observed in MTSat, MTR and T1app analyses

Supplementary Table MRI Acquisition Parameters

3.0-T MR system (Prisma; Siemens Healthcare, Erlangen, Germany) with a 32-channel head coil

Seq	uence	FOV	Acquisition Matrix	Slices	di	Voxel imensions	TE (ms)		TR (ms)	α (deg)		TI (ms)		-plane x slice cceleration factor	Acquisition time (mm:ss)	
sagittal 3D T1-	weighted MPRAGE	256 x 256	256 x 256	:	176	1mm isotropio	2.	26	250	00	7		1100	2 x 1	5:59	
	axial 2D T2 FLAIR	250 x 250	250 x 250		60	1 x 1 x 3mm	1	20	950	00	150		2400	2 x 1	4:47	
	proton density (no saturation pulse)	224 (sagittal-							30	1	5				6:14	
sagittal 3D FLASH spoiled gradient echo	proton density (with saturation pulse)	inferior) x 241 (anterior- posterior)	2 12	1.4 mm isotropic		1.54/ 4.55/ 8.49	55/				-	2 x 1	6:14			
	T1-weighted (no saturation pulse)										15	5	18			
	magnetisation transfer gaussian off-resonance saturation pulse (α: 500°; duration: 9.984 ms; offset: 1.2kHz)															
	diffusion-weighted cho-planar imaging	256 x 256	128 x 128		74	2.0 mm isotropic	7	'4	430	00	-		-	2 x 2	11:12	
	acquired with 151 diffusion directions comprising interleaved b-values of 0 (14), 200 (3), 500 (6), 1000 (64), 2000 (64) s/mm², in addition to 3 x b0 volumes with reversed phase encoding				on to 3 x b0 volumes											

FOV: field of view; SI: sagittal-inferior; AP: anterior-posterior; TE: echo time; TR: repetition time; MPRAGE: Magnetisation Prepared Rapid Gradient Echo; FLASH: fast low-angle shot spoiled gradient echo

Supplementary Table 1. MRI Acquisition Parameters

	LOW LESIO	ON VOLUME	HIGH LESIO	HIGH LESION VOLUME		
	LOW g-ratio	HIGH g-ratio	LOW g-ratio	HIGH g-ratio		
Baseline EDSS	2	2.5	2	2.5		
1-year EDSS	2	3	3	3		

Supplementary Table 2. Median baseline and 1-year EDSS scores for study individuals, according to lesion volume and g-ratio

There was no correlation between g-ratio and baseline EDSS (Spearman's rho=0.08, p=0.51), and no correlation between NfL and EDSS (Spearman's rho=0.06, p=0.63)

Fisher's Exact Test: Threshold: 0.4%					
	High Lesion L	oad (n=43)	Low Lesion Lo	oad (n=30)	
	Normal g-ratio	High g-ratio	Normal g-ratio	High g-ratio	
Normal NfL	15	15	17	9	
High NfL	2	11	3	1	
p-value	0.045		1		

Fisher's Exact Test: Threshold: 0.5%					
	High Lesion L	oad (n=38)	Low Lesion Lo	oad (n=35)	
	Normal g-ratio	High g-ratio	Normal g-ratio	High g-ratio	
Normal NfL	13	12	19	12	
High NfL	2	11	3	1	
p-value	0.039		1		

Fisher's Exact Test: Threshold: 0.6%					
	High Lesion L	oad (n=33)	Low Lesion Lo	oad (n=40)	
	Normal g-ratio	High g-ratio	Normal g-ratio	High g-ratio	
Normal NfL	12	10	20	14	
High NfL	1	10	4	2	
p-value	0.022		1		

Supplementary Table 3. Sensitivity analysis for lesion volume threshold

Appendix - Members of the Future-MS Consortium

The FutureMS consortium includes clinicians, data processors, or other FutureMS team members that have contributed to acquisition of data, access to patients, or whom have been relied upon for their input.

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Gwen Kennedy	NHS Tayside
Tracey Hopkins	NHS Tayside
Aberdeen	
Margaret Ann MacLeod	NHS Grampian
James MacDonald	NHS Grampian
Beverly Maclennan	NHS Grampian
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