

Supplemental Materials

Increasing serum neurofilament and glial fibrillary acidic protein after discontinuing multiple sclerosis treatment predicts future disease activity

Gauruv Bose, MD ^{1,2,3} ORCID: 0000-0002-5204-6348

Brian C. Healy, PHD ^{1,2}

Shrishti Saxena, MSC ¹

Fermisk Saleh¹

Bonnie I. Glanz, PHD ^{1,2}

Rohit Bakshi, MD, MA ^{1,2}

Howard L. Weiner, MD ^{1,2}

Tanuja Chitnis, MD ^{1,2}

1. Brigham and Women's Hospital, Department of Neurology, Boston, MA, USA

2. Harvard Medical School, Boston, MA, USA

3. The University of Ottawa and Ottawa Hospital Research Institute, Ottawa, ON, Canada

eTable 1. Comparison of eligible patients with and without serum samples available

	Serum available (included) (N=78)	Serum not available (excluded) (N=59)	p-value
Treatment-stop, calendar year, median (min, max)	2013 (2005, 2019)	1995 (1980, 2018)	<0.001
Female, No. (%)	72 (92%)	46 (78%)	0.016
Age at first symptoms, median (IQR)	32.1 (25.2, 41.8)	27.7 (22.5, 34.2)	0.003
Age at treatment-stop, median (IQR)	48.7 (38.8, 55.7)	40.6 (32.6, 47.6)	<0.001
Disease duration at treatment-stop, years, median (IQR)	12.9 (7.5, 19.2)	11.0 (6.0, 16.0)	0.16
Years from closest attack to treatment-stop, median	-6.5 (-12.6, -4.2)	-10.0 (-16.0, -5.5)	0.036
EDSS at treatment-stop, median (IQR)	1.5 (1.0, 2.5)	4.0 (3.0, 6.5)	<0.001

eTable 2. Multivariate time-to-event outcomes including MRI Lesion Burden (N=76)

	6-month CDW	New attack	New MRI activity
Pre-treatment-stop sNfL	1.37 [0.59, 3.20]; p=0.458	1.54 [0.55, 4.13]; p=0.418	2.02 [0.84, 4.90]; p=0.119
Post-treatment-stop sNfL	6.36 [2.27, 17.8]; p<0.001	1.48 [0.54, 4.05]; p=0.440	3.26 [1.43, 7.47]; p=0.005
Percent-change in sNfL	1.88 [1.22, 2.89]; p=0.004	1.13 [0.69, 1.85]; p=0.633	1.36 [0.96, 1.92]; p=0.082
Pre-treatment-stop sGFAP	1.17 [0.04, 3.43]; p=0.771	2.98 [0.87, 10.2]; p=0.082	0.99 [0.37, 2.64]; p=0.979
Post-treatment-stop sGFAP	1.85 [0.53, 6.51]; p=0.337	2.73 [0.72, 10.4]; p=0.140	2.49 [0.77, 8.07]; p=0.128
Percent-change in sGFAP	4.21 [0.78, 22.7]; p=0.094	0.88 [0.17, 4.52]; p=0.873	5.89 [1.39, 24.9]; p=0.016

Data shown as hazard ratio [95% confidence interval], and p-value.

Pre- and post-treatment-stop biomarker levels were log-transformed.

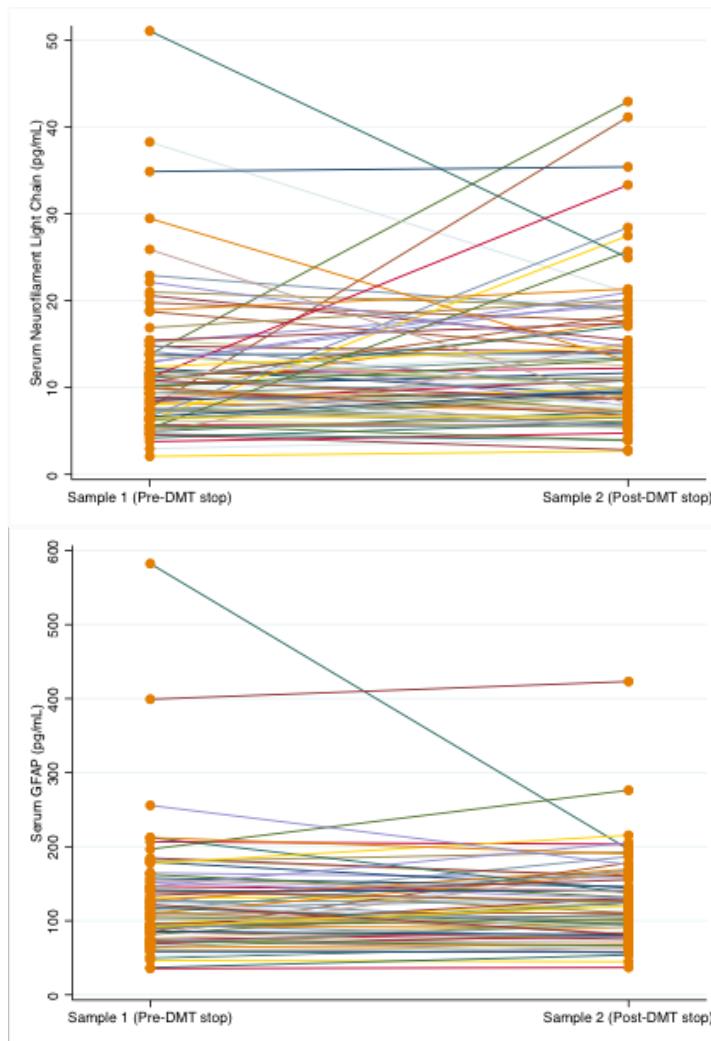
Percent-change was calculated from non-transformed levels where reported hazard ratios represent a doubling, or 100% increase.

*Adjusted for covariates: age, disease duration, EDSS, and duration from last attack (at treatment-stop date), T2LV at time of discontinuation, as well as batch effect

CDW = confirmed disability worsening; EDSS = expanded disability status scale; sNfL = serum neurofilament light chain; sGFAP = serum glial fibrillary acidic protein

eTable 3. Area Under Receiver Operating Characteristic Curve (AUC) analyses

Liu cutpoint:		Sensitivity	Specificity	AUC
Confirmed Disability Worsening				
Pre NFL (pg/mL)	9.336	0.59	0.45	0.52
Post NFL (pg/mL)	11.04	0.56	0.59	0.57
Change in NFL (%)	1.22	0.63	0.51	0.57
Pre GFAP (pg/mL)	116.34	0.41	0.51	0.46
Post GFAP (pg/mL)	99.77	0.56	0.39	0.47
Change in GFAP (%)	-1.81	0.63	0.51	0.57
New Attack				
Pre NFL (pg/mL)	10.050	0.53	0.51	0.52
Post NFL (pg/mL)	7.27	0.74	0.27	0.50
Change in NFL (%)	19.05	0.42	0.63	0.52
Pre GFAP (pg/mL)	95.19	0.53	0.39	0.46
Post GFAP (pg/mL)	88.28	0.74	0.29	0.51
Change in GFAP (%)	3.20	0.63	0.58	0.60
MRI Activity				
Pre NFL (pg/mL)	9.336	0.42	0.37	0.39
Post NFL (pg/mL)	14.95	0.35	0.73	0.54
Change in NFL (%)	19.94	0.42	0.67	0.55
Pre GFAP (pg/mL)	93.86	0.54	0.37	0.45
Post GFAP (pg/mL)	97.34	0.62	0.37	0.49
Change in GFAP (%)	2.47	0.69	0.62	0.65
Empirical optimal cutpoint determined by Liu test				



eFigure 1. Profile plots.

Each patient's pre-treatment-stop (sample 1) and post-treatment-stop (sample 2) levels of sNfL and sGFAP are shown. Out of 78 patients, 43 (55%) had an increase in sNfL, 41 (53%) had an increase in sGFAP, and 30 (38%) had an increase in both biomarkers (chi-squared P=0.001). There were 7 patients with a 100% increase in sNfL, and 5 patients with a 50% increase in sGFAP, of whom 3 patients had both.