Serum lipid profile changes predict neurodegeneration in interferon-β1a-treated multiple sclerosis patients

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Abstract The purpose of this work was to determine whether changes in cholesterol profiles after interferon-β (IFN-β)1a treatment initiation following the first demyelinating event suggestive of multiple sclerosis are associated with clinical and MRI outcomes over 4 years. A group of 131 patients (age: 27.9 ± 7.8 years, 65% female) with serial 3-monthly clinical and 12-monthly MRI follow-ups over 4 years were investigated. Serum cholesterol profiles, including total cholesterol (TC), HDL cholesterol (HDL-C), and LDL cholesterol (LDL-C) were obtained at baseline, 1 month, 3 months, and every 6 months thereafter. IFN-β1a initiation caused rapid decreases in serum HDL-C, LDL-C, and TC within 1 month of IFN-β1a initiation (all P < 0.001) that returned slowly toward baseline. In predictive mixed model analyses, greater percent decreases in HDL-C after 3 months of IFN-β1a treatment initiation were associated with less brain atrophy over the 4 year time course, as assessed by percent brain volume change (P = 0.001), percent gray matter volume change (P < 0.001), and percent lateral ventricle volume change (P = 0.005).

Decreases in cholesterol biomarkers following IFN-β1a treatment are associated with brain atrophy outcomes over 4 years. Pharmacological interventions targeting lipid homeostasis may be clinically beneficial for disrupting neurodegenerative processes.—Uher, T., K. Fellows, D. Horakova, R. Zivadinov, M. Vaneckova, L. Sobisek, M. Tyblova, Z. Seidl, J. Krasensky, N. Bergland, B. Weinstock-Guttman, E. Havrdova, and M. Ramanathan. Serum lipid profile changes predict neurodegeneration in interferon-β1a-treated multiple sclerosis patients. J. Lipid Res. 2017. 58: 403–411.

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Interferon-β (IFN-β) treatment is one of the most widely used disease-modifying treatments for multiple sclerosis (MS). The efficacy of recombinant human IFN-β for relapsing MS has been established by multiple double-blind placebo-controlled multi-center trials (1–3). IFN-β can delay the conversion to clinically definite MS (CDMS) in patients with a first demyelinating event (4–6), which supports early intervention with IFN-β.

MRI atrophy measures of brain volume and gray matter, white matter, and lateral ventricle volumes provide quantitative measures of global and tissue-specific brain volumes (7–9). Brain atrophy is useful for evaluating disease progression and therapeutic efficacy in MS (7) because it is a predictor of physical disability, cognitive dysfunction, and quality of life (10). However, there is a lack of effective serum biomarkers capable of predicting atrophy in MS patients.

There is considerable inter-individual variability of IFN-β1a effectiveness among MS patients. Phase III studies of IFN-β indicate a 30–40% general clinical benefit, although the response varies significantly among MS patients. On MRI, approximately 40% of patients show complete suppression of new contrast-enhancing lesions (CELs), whereas 20% of patients have less than 70% suppression.

Abbreviations: CDMS, clinically definite multiple sclerosis; CEL, contrast-enhancing lesion; EDSS, Expanded Disability Status Scale; HDL-C, HDL cholesterol; HDL-C%, percent changes in HDL cholesterol; IFN-β, interferon-β; LDL-C, LDL cholesterol; LDL-C%, percent changes in LDL cholesterol; MS, multiple sclerosis; NAB, neutralizing antibody; PBVC, percent brain volume change; PGmVC, percent gray matter volume change; PLVVC, percent lateral ventricle volume change; TC, total cholesterol; ΔTC%, percent changes in total cholesterol.

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patients. The associations, if any, of cholesterol changes with MRI confirmed (24–26). MS disease progression have since been independently with a 7.4%, 5.9%, and 16% increase in the number of new (LDL-C), total cholesterol (TC), and apoB was associated that each 10 mg/dl of greater baseline LDL cholesterol 1 IFN treatment caused coordinated changes in the expres-

sion of genes involved in sterol synthesis (21). We found 1a-treated patients and up to 38% of IFN-β1b-treated patients (12, 13), provide a clinically useful biologically intuitive mechanistic explanation for partial responsive-

ness to IFN-β therapy. The presence of persistently high levels of NAB abrogates IFN-induced signaling responses and is associated with decreases in clinical and MRI effectiveness of IFN-β therapy. The majority of MS patients who are partially responsive to IFN-β tend to be NAB-negative (2, 14, 15).

IFN-β has antiviral, anti-proliferative, and immunomodulatory effects (16, 17). Diverse biomarkers, ranging from single nucleotide polymorphisms, mRNA (e.g., MxA, Stat1, TRAIL, and others), proteins (e.g., oligoadenylate synthetase activity, Mx protein, and β2 microglobulin), and metabolites (e.g., neopterin) to immune cell populations, have been investigated as potential IFN-β biomarkers (18–20). However, they were not useful predictors of MRI and clinical outcomes.

We investigated cholesterol profiles as a potential biomarker for IFN-β therapy in MS based on a report that type 1 IFN treatment caused coordinated changes in the expression of genes involved in sterol synthesis (21). We found that each 10 mg/dl of greater baseline LDL cholesterol (LDL-C), total cholesterol (TC), and apoB was associated with a 7.4%, 5.9%, and 16% increase in the number of new T2 lesions over 2 years of IFN-β treatment, respectively (22, 23). Our findings on cholesterol and apolipoproteins in MS disease progression have since been independently confirmed (24–26).

The goals of this study were to investigate the effects of IFN-β therapy on cholesterol profile changes for a 72 month period following IFN-β1a initiation and to examine the associations, if any, of cholesterol changes with MRI measures over a 48 month period in IFN-β-treated MS patients.

METHODS

Study population

The SET study (27) was a prospective longitudinal observational clinical study that involved eight centers in the Czech Republic (clin.gov # NCT01592474). The Ethics Committees of all participating centers approved the study protocol. All patients gave their written informed consent.

Inclusion and exclusion criteria. This multi-center study enrolled patients within 4 months after their first clinical event suggestive of MS. The inclusion criteria were: age 18–55 years, Expanded Disability Status Scale (EDSS) score of 3.5 or less, presence of two or more T2-hypointense lesions on diagnostic MRI, and two or more oligoclonal bands in CSF obtained at the screening visit prior to steroid treatment. Exclusion criteria were occurrence of a second relapse before the baseline visit and pregnancy.

Study design and assessments

Assessments and treatment algorithms. The study included clinical visits every 3 months during 48 months of follow-up in routine clinical practice. The clinical outcomes of the study were new relapses associated with conversion to CDMS and sustained disability progression, defined as an increase in EDSS by 1.0 point (if baseline EDSS > 0) or 1.5 points (if baseline EDSS = 0), confirmed after 12 months.

MRI was obtained using standardized protocols on a single 1.5 Tesla MRI scanner at baseline, at 6 months, and yearly thereafter until the 48 month time point. Information on the MRI acquisition and analysis protocols can be found in the supplemental data.

All patients started with 30 mcg of intramuscular IFN-β1a once-weekly (AVONEX®, Biogen-Idec) at baseline. The mean time between clinical onset and baseline was 81.8 ± 22.8 (SD) days (median 78.0 days; range 33–122 days). All patients were treated with 3–5 g of methylprednisolone for the first clinical symptoms before study entry. None of the patients were on statins.

Treatment change criteria. The SET study protocol defines once-weekly IFN-β1a treatment failure as ≥2 relapses during 12 months or a 6 month sustained disability progression.

Treating physicians had full discretion to individualize treatment changes for once-weekly IFN-β1a treatment failures. Protocol-recommended options for treatment failures included the following: subcutaneous IFN-β1a (14 μg three times weekly), intravenous natalizumab (300 mg per month), or the addition of azathioprine (50 mg twice-daily to once-weekly IFN-β1a). Patients who failed the initial treatment change were treated with subcutaneous glatiramer acetate (20 mg daily) or mitoxantrone (10 mg).

Serum cholesterol profiles

For biochemical assessment, 5 ml of blood was obtained at screening, baseline, 1 month, every 3 months for the first year, and every 6 months thereafter. Serum for lipid analyses was obtained in the nonfasted state at the central site. Examination at screening was performed before any steroid treatment, and baseline examinations were at least 30 days after. Diagnostic reagent kits (Cholesterol Liquicolor and HDL Liquicolor; Human Gesellschaft für Biochemica und Diagnostica mbH, Germany) were used to measure serum TC, HDL cholesterol (HDL-C), and triglycerides. The intra-assay coefficient of variation for TC and HDL-C serum levels was 0.85%, and the inter-assay coefficient of variation was 1.46–1.86%. LDL-C was obtained from the Friedewald equation (28).

Statistical analysis

SPSS 22.0 (IBM, Armonk, NY) statistical software was used for statistical analyses. Percent changes in TC (ΔTC%), HDL-C (ΔHDL-C%), and LDL-C (ΔLDL-C%) were computed at each time point using baseline TC, HDL-C, and LDL-C values as the reference. A zero value for percent change represents no change compared with baseline, a positive value represents an increase in levels, and a negative value represents a decrease in levels compared with baseline.

The statistical significance of ΔTC%, ΔHDL-C%, and ΔLDL-C% values at 1, 3, and 6 months was assessed using a one-sample t-test.
The time courses of clinical and MRI outcome measure-dependent variables [percent brain volume change (PBVC), percent gray matter volume change (PGMVC), percent lateral ventricle volume change (PLVVC), new/enlarging T2 and CEL, EDSS, and cumulative relapses] were analyzed with longitudinal adjusted linear mixed-effect model analysis with a random intercept for time and each patient. In the longitudinal analyses, the fixed effects included age, gender, BMI, and baseline values of the clinical or MRI outcome measure and the time course of the individual cholesterol variable (either ∆HDL-C%, ∆LDL-C%, or ∆TC%) of interest.

The same linear mixed-effect model method was used for the predictive analysis. The fixed effects included age, gender, BMI, baseline values of the clinical or MRI outcome measure, and individual cholesterol variable of interest (∆HDL-C%, ∆LDL-C%, or ∆TC%) at a single early time point (either 1, 3, or 6 months).

The Mann-Whitney test was used to assess differences in cumulative relapses, EDSS change, PBVC, and ∆HDL-C% at 3 months between patients who remained on once-weekly intramuscular IFN-β1a or switched to no treatment versus patients who switched to subcutaneous IFN-β, glatiramer acetate, natalizumab, or other treatments.

A Benjamini-Hochberg correction with \( P < 0.05 \) was used to control the false discovery rate (\( q \) value) due to multiple testing for predictive associations.

## RESULTS

### Baseline and follow-up clinical characteristics

Of the 220 patients enrolled in the SET study, 131 (age: 27.9 ± 7.8 years, 63% female) had lipid, clinical, and MRI follow-up data at 4 years. A subset of patients also had lipid measures available over 6 years that were included in the analysis of the time course of lipid profile changes. Aside from a modest difference in baseline EDSS (\( P = 0.018 \), Mann-Whitney test) between the included patients (median EDSS ± interquartile range = 1.5 ± 0.5) and the excluded patients (median EDSS ± interquartile range = 1.5 ± 1.0), the subset of patients was representative of the larger SET study cohort on demographic, clinical, and MRI characteristics. Supplemental Table S1 summarizes the number of lipid profiles available at each time-point.

Table 1 summarizes patient demographic, clinical, laboratory, and MRI characteristics at baseline, 24 months, and 48 months. At 4 years, 13% of patients reached 12 month sustained disability progression.

### Time course of cholesterol profile changes

TC, HDL-C, and LDL-C levels declined from baseline at 1 month following IFN-β1a treatment and gradually returned to baseline over 72 months (Fig. 1). The corresponding percent changes, ∆TC%, ∆HDL-C%, and ∆LDL-C%, followed the same pattern.

At 1 month following IFN-β1a initiation, the average ∆TC% was an 8.7% decrease, the average ∆HDL-C% was a 6.1% decrease, and the average ∆LDL-C% was a 13.4% decrease. ∆TC%, ∆HDL-C%, and ∆LDL-C% at the 1, 3, and 6 month time points following IFN-β1a initiation were negative relative to baseline (all \( P < 0.001 \)). The decreases in ∆TC% were sustained through 4 years (all \( P < 0.007 \)) with a return to baseline occurring at month 54. ∆HDL-C% decreases remained significant compared with baseline through the first year (all \( P < 0.002 \)), but returned to baseline levels at 18 months. The decreases in ∆LDL-C% were sustained over the observation period (all \( P < 0.05 \)), with a return to baseline at the end of the 6 year period.

There was substantial inter-patient variability in the lipid changes over time. Supplemental Fig. S1 shows histograms of ∆TC%, ∆HDL-C%, and ∆LDL-C% at 1, 3, and 6 months of IFN-β1a treatment. The peaks of percent change were negative, indicating that the majority of patients exhibit...
decreases in $\Delta$TC%, $\Delta$HDL-C%, and $\Delta$LDL-C% at 1, 3, and 6 months.

The time courses for the mean values of $\Delta$TC%, $\Delta$HDL-C%, and $\Delta$LDL-C% by the quartiles of $\Delta$HDL-C% at 3 months are summarized in supplemental Fig. S2. The lowest quartile of $\Delta$HDL-C% at 3 months was associated with sustained decreases in mean $\Delta$HDL-C% and $\Delta$TC% over the 4 year period. The patterns of $\Delta$LDL-C% were not sufficiently segregated by quartiles of $\Delta$HDL-C%.

### Longitudinal associations of cholesterol profile with MRI and clinical measures

We investigated the associations of cholesterol profiles at baseline and longitudinal $\Delta$TC%, $\Delta$HDL-C%, $\Delta$LDL-C% changes occurring after IFN-β1a treatment with the longitudinal clinical and MRI measures. The results are summarized in Table 2.

**Baseline cholesterol profiles.** Baseline LDL-C ($P = 0.025$) and TC ($P = 0.001$) levels were associated with a greater occurrence of new T2 lesions and a greater number of new/enlarging T2 lesions ($P = 0.003$ for TC and $P = 0.015$ for LDL-C). Increases in baseline TC were associated with a greater cumulative number of relapses ($P = 0.039$). We did not obtain evidence for associations for changes in brain volumes with baseline TC, LDL-C, and HDL-C (data not shown).

**Longitudinal percent decreases.** Greater longitudinal decreases in $\Delta$TC% and $\Delta$HDL-C% were associated with a smaller number of new lesions ($P = 0.027$ for $\Delta$TC%, $P = 0.009$ for $\Delta$HDL-C%) and new/enlarging T2 lesions ($P = 0.038$ for $\Delta$TC%, $P = 0.009$ for $\Delta$HDL-C%).

Greater longitudinal decreases in $\Delta$TC% ($P < 0.001$) and $\Delta$HDL-C% ($P = 0.001$) were associated with smaller decreases in PBVC. Greater longitudinal decreases in $\Delta$HDL-C% ($P = 0.002$) were associated with smaller decreases in PGMVC. In contrast, greater longitudinal decreases in $\Delta$LDL-C% were associated with greater decreases in PGMVC ($P < 0.001$). Longitudinal decreases in $\Delta$TC% were not associated with PGMVC. Thus the changes in $\Delta$LDL-C% and $\Delta$HDL-C% following IFN-β1a treatment have opposing effects on PGMVC that offset each other. Likewise, greater decreases in $\Delta$HDL-C% were associated with smaller increases in PLVVC ($P = 0.001$).

These results suggest that greater decreases in $\Delta$HDL-C% are associated with less brain atrophy.
Predictive associations of cholesterol profile changes with clinical and MRI measures

We assessed whether the associations of ΔHDL-C% changes at a single early time point (at 3 months) following IFN-β1a treatment initiation could be deployed predictively as a possible biomarker for longitudinal clinical and MRI measures.

For completeness, similar analyses were done with ΔHDL-C% changes at 1 and 6 months and for ΔLDL-C% and ΔTC% changes at 1, 3, and 6 months. The ΔHDL-C% at 1 and 3 months exhibited associations with a greater number of clinical measures than ΔLDL-C% and ΔTC% changes at 1, 3, and 6 months. In the interest of brevity, we report only our ΔHDL-C% results at 1 and 3 months in detail.

Lesion measures. Table 3 summarizes the associations of MRI measures (number of new lesions, number of new/enlarging T2 lesions, and CEL number) with ΔHDL-C% after 1 and 3 months.

Following correction for multiple testing, there were no significant associations of the ΔHDL-C% with these lesional measures.

Atrophy measures. In adjusted mixed model analyses (Table 3), a significant association was observed between ΔHDL-C% after 1 month with PBVC (P < 0.001) and PG-MVC (P = 0.004). Table 3 also shows the slopes that provide a measure of the strength of the associations with ΔHDL-C% at 1 month with all other variables held constant. A 1% decrease in HDL-C at 1 month was associated with a 2.55% improvement in brain volume and a 1.67% improvement in gray matter volume.

ΔHDL-C% at 3 months was associated with PBVC (P < 0.001), PG-MVC (P < 0.001), and PLVVC (P = 0.005). The slopes in Table 3 indicate that a 1% decrease in HDL-C at 3 months is associated with a 1.9% improvement in brain volume, a 1.7% improvement in gray matter volume, and a 5.5% improvement in lateral ventricle volume.

These results are graphically summarized by quartiles of ΔHDL-C% at 3 months in Fig. 2A–C. The lowest quartile represents the patients with the greatest decrease in ΔHDL-C% at 3 months, whereas the highest quartile represents the patients with the smallest decreases in ΔHDL-C% at 3 months. The lowest quartile subgroup shows better atrophy outcomes compared with the highest quartile subgroup.

For completeness, we also investigated the associations of ΔLDL-C% and ΔTC% at 1, 3, and 6 months with the same atrophy measures. PBVC, PG-MVC, and PLVVC were not associated with ΔLDL-C% at 1, 3, or 6 months. PBVC was significantly associated with ΔTC% at 1 month

TABLE 2. Associations of longitudinal percent changes in cholesterol variables with MRI and clinical outcomes

| Clinical/MRI Variable | ΔTC% | ΔLDL-C% | ΔHDL-C% |
|-----------------------|------|---------|---------|
| PBVC                  | −1.90 (0.001) | 0.0113 (0.95) | −0.835 (0.001) |
| PGMVC                 | 0.037 (0.98)  | 1.2 (0.001)    | −1.34 (0.002)  |
| PLVVC                 | 1.30 (0.51)   | −4.48 (0.001)  | 6.10 (0.001)   |
| New T2 lesions        | 2.90 (0.027)  | 0.838 (0.35)   | 3.29 (0.009)   |
| New/enlarging T2 lesions | 3.69 (0.038) | 1.18 (0.33)    | 4.47 (0.009)   |
| CEL                   | 5.16 (0.001)  | 3.51 (0.001)   | 2.39 (0.023)   |
| EDSS                  | 0.145 (0.36)  | 0.105 (0.35)   | 0.0054 (0.97)  |
| Cumulative relapses   | 0.289 (0.059) | 0.0456 (0.674) | 0.252 (0.089)  |

Slope (P values) from mixed effect models are summarized.

*In MS decreases in normalized brain volume, decreases in gray matter volume, increases in lateral ventricle volume, increases in the number of T2 lesions, increases in contrast enhancing lesions (CEL), increases in EDSS, and increases in cumulative relapses are adverse MRI/clinical outcomes.

TABLE 3. Predictive associations between HDL-C percent changes at 1 month and 3 months and clinical and brain imaging outcome measures

| Clinical/MRI Variable | ΔHDL-C% 1 Month | ΔHDL-C% 3 Months |
|-----------------------|-----------------|-----------------|
|                       | Slope | P     | q     | Slope | P     | q     |
| PBVC                  | −2.55 | 1.2 × 10^-8 | 1.2 × 10^-7 | −1.90 | 4.9 × 10^-8 | 4.9 × 10^-7 |
| PGMVC                 | −1.67 | 0.004 | 0.02 | −1.70 | 1.4 × 10^-4 | 4.7 × 10^-4 |
| PLVVC                 | 3.39  | 0.16  | 0.23 | 5.5  | 0.005 | 0.013 |
| New T2 lesions        | −3.13 | 0.16  | 0.27 | −0.644 | 0.71 | 0.71 |
| New/enlarging T2 lesions | −6.39 | 0.035 | 0.07 | −1.45 | 0.54 | 0.68 |
| CEL                   | 1.12  | 0.47  | 0.52 | 0.687 | 0.56 | 0.62 |
| EDSS                  | −0.01 | 0.96  | 0.96 | 0.394 | 0.054 | 0.077 |
| Cumulative relapses   | 0.187 | 0.42  | 0.52 | 0.391 | 0.024 | 0.04 |

Slope, P value, and q-value (false discovery rate) are provided.

*In MS decreases in normalized brain volume, decreases in gray matter volume, increases in lateral ventricle volume, increases in the number of T2 lesions, increases in CEL, increases in EDSS, and increases in cumulative relapses are adverse MRI/clinical outcomes.

*ΔHDL-C% at 1 month and ΔHDL-C% at 3 months are positive when HDL levels are increased relative to baseline, zero when HDL is unchanged relative to baseline, and negative when HDL is decreased relative to baseline.
(\(P = 0.001\)), \(\Delta TC\% \) at 3 months (\(P < 0.001\)), and \(\Delta TC\% \) at 6 months (\(P = 0.002\)). We attribute the associations of \(\Delta TC\% \) to the contributions of \(\Delta HDL-C\% \).

**Effects of T2 lesions on predictive associations.** In additional mixed effect analyses, we included baseline T2 lesion volume as an additional predictor to determine whether the associations of \(\Delta HDL-C\% \) at 3 months with atrophy measures remained significant. The associations of \(\Delta HDL-C\% \) at 3 months with PBVC, PGMVC, and PLVVC remained significant (all \(P \leq 0.003\), data not shown) in these analyses.

We also performed similar analyses that included the number of new T2 lesions as an additional predictor. Again, the associations of \(\Delta HDL-C\% \) at 3 months with PBVC, PGMVC, and PLVVC remained significant (all \(P \leq 0.001\)).

The statistical significance of these associations indicates that \(\Delta HDL-C\% \) at 3 months explains variance in brain atrophy that is not explained by T2 lesions. Thus, the predictive associations of \(\Delta HDL-C\% \) at 3 months with brain atrophy may be clinically useful, because they complement MRI measures of lesion burden and activity.

**Disability and relapses.** Greater decreases in \(\Delta HDL-C\% \) at 3 months were associated with a lower cumulative number of relapses (\(P = 0.024\)). \(\Delta HDL-C\% \) at 1 month was not associated with cumulative number of relapses (Table 3). Figure 2F compares the mean cumulative number of relapses for the quartiles of \(\Delta HDL-C\% \) at 3 months. The patients in the highest quartile of \(\Delta HDL-C\% \) at 3 months had a larger number of cumulative relapses compared with the lowest quartile. The \(\Delta HDL-C\% \) after 1 and 3 months was not associated with EDSS.

These findings are consistent with the hypothesis that the magnitude of decreases in \(\Delta HDL-C\% \) 3 months after IFN-\(\beta1a\) initiation may be a biomarker for individuals at risk for increased whole brain atrophy and relapses.

**Cholesterol profiles in patients receiving treatment changes**

Group 1 comprised the patients who remained on once-weekly IFN-\(\beta1a\) and those who switched to no treatment. Group 2 consisted of patients whose treatment was changed to other treatments (subcutaneous IFN-\(\beta\), glatiramer acetate, natalizumab, and other therapies). The number of patients requiring treatment change at 2, 3, and 4 years is summarized in Fig. 3D.

Figure 3A–C shows the cumulative number of relapses, the EDSS change from baseline, and the PBVC for the treatment change groups at 2 and 4 years. At 2 years,
Lipid metabolic changes predict neurodegeneration

Initial decrease of serum cholesterol biomarkers followed by a gradual return to baseline. HDL-C returned to baseline within 18 months, TC returned to baseline within 54 months, but LDL-C did not return to baseline until the end of the 6 year observation period. Greater longitudinal decreases in \( \Delta \text{HDL-C}\% \) were associated with smaller decreases in PBVC and PGMVC, and greater increases in PLVVC. Changes in \( \Delta \text{LDL-C}\% \) and \( \Delta \text{HDL-C}\% \) following IFN-\( \beta \)-1a treatment have opposing effects on PGMVC that offset each other. \( \Delta \text{HDL-C}\% \) at 3 months was associated with the time course of atrophy outcomes over 4 years.

The strengths of the study include the prospective longitudinal study design and the availability of cholesterol biomarkers and clinical and MRI data over a 4 year period. This enabled us to investigate the role of cholesterol biomarkers at the early stages of MS in a relatively young patient cohort that was not on statin treatments. The limitations include the lack of a suitable control population that was not on IFN-\( \beta \)-1a treatment. We also did not have data on dietary and lifestyle changes in the patient population.

The mechanisms responsible for the associations between brain volume changes and early decreases in HDL-C levels following IFN initiation are unknown. A plausible explanation is that decreases in HDL-C are particularly effective as a clinically integrative measure of the same initial decrease of serum cholesterol biomarkers followed by a gradual return to baseline. HDL-C returned to baseline within 18 months, TC returned to baseline within 54 months, but LDL-C did not return to baseline until the end of the 6 year observation period. Greater longitudinal decreases in \( \Delta \text{HDL-C}\% \) were associated with smaller decreases in PBVC and PGMVC, and greater increases in PLVVC. Changes in \( \Delta \text{LDL-C}\% \) and \( \Delta \text{HDL-C}\% \) following IFN-\( \beta \)-1a treatment have opposing effects on PGMVC that offset each other. \( \Delta \text{HDL-C}\% \) at 3 months was associated with the time course of atrophy outcomes over 4 years.

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The mechanisms responsible for the associations between brain volume changes and early decreases in HDL-C levels following IFN initiation are unknown. A plausible explanation is that decreases in HDL-C are particularly effective as a clinically integrative measure of the same
pathways that mediate IFN-β treatment effects on brain volume changes. Previously investigated biomarkers, e.g., MxA, reflect single pathways, whereas serum cholesterol levels are largely dependent on multiple biochemical homeostatic pathways. Cholesterol is also required for important brain functions, including myelin formation, immune functioning, neuronal signaling, and vascular function; therefore, changes may reflect not only treatment effects, but also pathophysiological effects. Interestingly, the MS-STAT study of the cholesterol-lowering drug, simvastatin, reported reduction of brain atrophy in secondary progressive MS (32). This may suggest that reductions in cholesterol may affect accumulation of atrophy in MS.

Vascular comorbidities were reported to be associated with more rapid disability progression in an MS patient-reported registry (33). However, the mean age of this study was 52.7 ± 10.4 years compared with our cohort, which had an average age of 27.9 ± 7.8 years. The venous insufficiency hypothesis for MS has been discredited (34). Our results confirm and extend the findings reported by Morra et al. (35), who reported that the percent decrease in TC at 2 months after IFN initiation was 8.2% and at 12 months following IFN initiation was 7%. We observed an 8.7% decrease in TC at 1 month, a 7.8% decrease at 6 months, and a 6% decrease at 12 months after IFN initiation. Morra et al. (35) did not fully assess whether the decreases in TC following IFN-β treatment were sustained over the long-term and did not include longitudinal outcome measures.

The IFN-β1a effects on cholesterol homeostasis that we report are likely shared by other type 1 IFNs. Once-daily intramuscular recombinant and leukocyte-derived IFN-α induced an approximately 25% decrease in cholesterol and a 16% decrease in HDL-C within a few days of initiating treatment in metastatic breast carcinoma and nodular lymphoma patients (36). Similar results were reported in patients treated with IFN-α2a for hepatitis (37). We anticipate that PEGylated-IFN (PEGinterferon-β1a, PLEGRIDY), which has been approved for MS, will also exhibit effects on cholesterol profiles.

The reason for the return of cholesterol levels to baseline remains unclear. The initial decrease may be attributed to the multiple mechanisms by which IFN-β acts on the cholesterol pathway. These include downregulation of the sterol pathway through decreases in SREBP2 (21), the primary transcription factor in sterol synthesis, as well as induction of 25-hydroxylase (38), which promotes the metabolism of cholesterol to 25-hydroxycholesterol. The return of cholesterol levels to baseline occurs more slowly compared with the IFN-β1a-induced decreases and may reflect compensatory mechanisms. It is also possible that patients are more susceptible to dyslipidemia as they age or that responses to IFN-β1a may be attenuated over time.

IFN-β remains an important first-line MS therapy alongside glatiramer acetate. The identification of HDLC decrease as a potential biomarker of long-term atrophy following IFN-β treatment could find clinical utility if validated in larger well-controlled studies, because the underlying methodology is familiar and inexpensive. Our results could thus be a first step toward individualizing MS therapy by identifying IFN-β nonresponders who could be switched to other therapies.

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