## **P037**



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**Prognostic Value of Serum** Neurofilament Light Chain for Disease Activity in **Patients With Relapsing Nultiple Sclerosis: Results** From Subgroup Analysis Based on Body Mass Index and Age From the Phase 3 **ASCLEPIOS I/II Trials** 

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# **KEY FINDINGS & CONCLUSIONS**

- The use of a single sNfL threshold, as defined in the ASCLEPIOS I/II study protocols, was prognostic of future lesion formation in the overall study population, including subgroups defined by BMI and age
- Overall, these data support further work on the optimization of sNfL for prognostication of future MS disease activity that would be applicable for patients with RMS

Please refer to these related posters for more details on the prognostic value of sNfL:

- Prognostic Value of On-Treatment Serum Neurofilament Light Chain for New or Enlarging T2 Lesions in People With Relapsing Multiple Sclerosis: Pooled Analysis of the ASCLEPIOS I/II Trials (**P032**)
- Baseline Serum Neurofilament Light Chain Levels Predict Future Disease Activity Irrespective of Race/Ethnicity: Results From the Phase 3 ASCLEPIOS I/II Trials (P036)

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## Table at ba

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# **Prognostic value of sNfL for neT2 lesions by BMI and age subgroups**

# INTRODUCTION

- A challenge encountered in clinical practice with relapsing multiple sclerosis (RMS) is the difficulty in prognosticating the risk of future disease activity because of the variable disease course across patients<sup>1</sup>
- Inflammatory disease activity mostly occurs in the younger RMS population and declines with age<sup>2</sup>
- A biomarker that can prognosticate disease activity may help optimize individualized patient care and limit irreversible neurological damage, even in the absence of overt clinical symptoms or radiological signs<sup>3</sup>
- In the phase 3 ASCLEPIOS I/II trials (ofatumumab versus teriflunomide in people with RMS [pwRMS]), a pre-planned analysis of baseline serum neurofilament light chain (sNfL) levels based on being above or below the baseline median found that sNfL levels were prognostic for on-study lesion formation and brain volume loss in the overall population and in recently diagnosed treatment-naive participants<sup>4</sup>
- The prognostic value of sNfL was also observed when participants were categorized by baseline sNfL concentration quartiles - Irrespective of treatment, participants in the lowest quartile (Q1) had the lowest risk for on-study new or enlarging [ne] T2 lesions, with the risk for neT2 lesions increasing sequentially up to the highest sNfL quartile (Q4)<sup>5</sup>
- Some factors such as age and/or body mass index (BMI) may influence sNfL levels; however, in younger pwRMS without comorbidities, the impact of these factors on sNfL levels is expected to be limited compared to the effect of MS neuroinflammation itself<sup>6-10</sup>

## OBJECTIVE

• To evaluate the prognostic value of low versus high sNfL (based on a pre-planned analysis of baseline sNfL being above or below the baseline median) for future disease activity according to BMI and age in pwRMS

## RESULTS

## **Participant characteristics**

• Of the 1882 participants randomized in ASCLEPIOS I/II trials, 1746 (92.8%) had baseline sNfL data, including 1678 (96.1%) participants with neT2 and sNfL data available for BMI and age subgroups

• Baseline characteristics were similar between sNfL groups, except the number of Gd+ lesions and T2 lesion volume, which were considerably higher in participants with high versus low baseline sNfL (**Table 1**)

e 1.	<b>Demographic and</b>	disease	characteristics	by sNfL	category
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racteristic	Low sNfL category (<9.3 pg/mL) N=876 <sup>a</sup>	High sNfL category (≥9.3 pg/mL) N=870 <sup>a</sup>
, years	38.6±8.5	37.8±9.7
, kg/m²	27.0±6.4	24.7±5.5
nale, n (%)	588 (67.1)	602 (69.2)
duration since first symptom,	8.3±7.2	7.9±6.9
viously treated with DMT, n (%)	537 (61.3)	507 (58.3)
nber of relapses in the year ore the study	1.2±0.7	1.3±0.7
e since onset of most recent pse, months	7.8±13.5	7.0±9.3
SS score	2.8±1.3	2.9±1.4
malized brain volume, cm <sup>3</sup>	1447.6±74.8	1437.2±81.0
nber of Gd+ T1 lesions	0.4±1.2	2.6±5.4
icipants free of Gd+ T1 lesions, )	679 (77.5)	383 (44.0)
esion volume, cm <sup>3</sup>	9.4±10.6	16.7±15.0
dian sNfL, pg/mL	6.76	14.23

<sup>a</sup>Only participants with non-missing baseline sNfL values are included.

Data are expressed as mean±standard deviation unless specified otherwise.

BMI, body mass index; DMT, disease-modifying therapy; EDSS, Expanded Disability Status Scale; Gd+, gadolinium-enhancing; MS, multiple sclerosis; sNfL, serum neurofilament light chain.

 Across BMI and age subgroups, the mean annualized rate of neT2 lesions was significantly higher in participants with high sNfL (≥9.3 pg/mL) versus low sNfL (<9.3 pg/mL) levels (**Figure 1**)

### References

- 1. Oh J, et al. *Curr Opin Neurol.* 2018;31:752–759.
- **2.** Dahlke F, et al. *Mult Scler.* 2021;27:2062–2076. **3.** Thebault S, et al. *Mult Scler.* 2022;28:1491–1497.
- **4.** Ziemssen T, et al. *Front Immunol.* 2022;13:852563.
- **5.** Hauser SL, et al. *N Engl J Med.* 2020; 383:546–557.
- 6. Consortium of Multiple Sclerosis Centers. Int J MS Care. 2021;23:1–36
- 7. Bittner S, et al. *Brain.* 2021;144:2954–2963.
- 8. Bar-Or A, et al. *EBioMedicine*. 2023;93:104462. **9.** Kapoor R, et al. *Neurol.* 2020;95:436–444.
- **10.** Barro C, et al. *Ann Clin Transl Neurol.* 2020;7:2508–2523. **11.** Bar-Or A, et al. *Neurol Ther.* 2023 Feb;12(1):303–317.

**Abbreviations** 

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Figure 2A (age, left side): Black line is from linear regression of log (baseline sNfL) on age; blue line is when applying a non-linear regression of log (baseline sNfL) on age; blue line is from linear regression of log (baseline sNfL) on age; blue line is when applying a non-linear regression of log (baseline sNfL) on age; blue line is when applying a non-linear regression of log (baseline sNfL) on age; blue line is from linear regression of log (baseline sNfL) on age; blue line is when applying a non-linear regression of log (baseline sNfL) on age; blue line is from linear regression of log (baseline sNfL) on age; blue line is when applying a non-linear regression of log (baseline sNfL) on age; blue line is from linear regression of log (baseline sNfL) on age; blue line is when applying a non-linear regression of log (baseline sNfL) on age; blue line is when applying a non-linear regression of log (baseline sNfL) on age; blue line is when applying a non-linear regression of log (baseline sNfL) on age; blue line is when applying a non-linear regression of log (baseline sNfL) on age; blue line is when applying a non-linear regression of log (baseline sNfL) on age; blue line is when applying a non-linear regression of log (baseline sNfL) on age; blue line is when applying a non-linear regression of log (baseline sNfL) on age; blue line is when applying a non-linear regression of log (baseline sNfL) on age; blue line is when applying a non-linear regression of log (baseline sNfL) on age; blue line is when applying a non-linear regression of log (baseline sNfL) on age; blue line is when applying a non-linear regression of log (baseline sNfL) on age; blue line is when applying a non-linear regression of log (baseline sNfL) on age; blue line is when applying a non-linear regression on age; blue line is when applying a non-linear regression of log (baseline sNfL) on age; blue line sNfL) on age; blue line is when applying a non-linear regression of log (baseline sNfL) on age; blue line sNfL) on age; blue line sNfL on on BMI; blue line is when applying a non-linear relationship; dashed red line indicate the 5<sup>th</sup> percentile of BMI (18.78) and the corresponding mean sNfL (12.10 pg/mL). The vertical and horizontal green lines indicate the 95<sup>th</sup> percentile of BMI (37.74) and the corresponding mean sNfL (7.86 pg/mL). BMI, body mass index; MS, multiple sclerosis; RMS, relapsing multiple sclerosis; sNfL, serum neurofilament light chain.

## LIMITATIONS

- Based on the pre-planned nature of the analysis, participants were stratified by baseline median sNfL value into "high" or "low" with the intention to divide a phase 3 trial RMS population into groups of equal size with higher versus lower than median sNfL
- The use of a single sNfL threshold may be mostly applicable to relatively young RMS population included in these trials; for understanding sNfL changes outside of this age range, the use of sNfL as a continuous variable may be preferable
- While results support the applicability of sNfL for most patients across the BMIs (<18.5 and >40.0), prognostication using a single sNfL value should take into account the effect of BMI on sNfL
- Results reported here are based on the protocol-defined single sNfL as a prognostic factor could be optimized with a specific target and population in mind • The data presented in this study are based on a population that was selected according to the ASCLEPIOS inclusion/exclusion criteria to represent a population suitable for phase 3 trials/regulatory purposes, and it may not reflect the broader population of individuals with RMS seen in everyday clinical practice Recent findings suggest that sNfL has comparable prognostic value to Gd-enhancing lesions on MRI scans for future lesion formation<sup>11</sup>. The relative prognostic value of sNfL and baseline MRI characteristics should be further confirmed

## METHODS Study design

### Assessments

- (<24.5 vs  $\geq$ 24.5 kg/m<sup>2</sup>) and age (<38.0 vs  $\geq$ 38.0 years) subgroups

### **Statistical analysis**





BMI, body mass index; DMT, disease-modifying therapy; EDSS, Expanded Disability Status Scale; Gd+, gadolinium-enhancing; MRI, magnetic resonance imaging; NS, multiple sclerosis; entry, ended Disability Status Scale; Gd+, gadolinium-enhancing; MRI, magnetic resonance imaging; MS, multiple sclerosis; entry, ended Disability Status Scale; Gd+, gadolinium-enhancing; MRI, magnetic resonance imaging; MS, multiple sclerosis; entry, ended Disability Status Scale; Gd+, gadolinium-enhancing; MRI, magnetic resonance imaging; MS, multiple sclerosis; entry, ended Disability Status Scale; Gd+, gadolinium-enhancing; MRI, magnetic resonance imaging; MS, multiple sclerosis; entry, ended Disability Status Scale; Gd+, gadolinium-enhancing; MRI, magnetic resonance imaging; MS, multiple sclerosis; entry, entry

• ASCLEPIOS I/II were two phase 3, double-blind, active-controlled trials in which participants with RMS were randomized to receive either of atumumab or teriflunomide for up to 30 months

 Participants aged 18–55 years with a diagnosis of RMS, Expanded Disability Status Scale (EDSS) score 0–5.5, ≥1 relapse in the year before screening or  $\geq 2$  relapses in the last 2 years before screening, or  $\geq 1$  gadolinium-enhancing (Gd+) lesion on magnetic resonance imaging (MRI) in the year before randomization were included

• The baseline sNfL cut-off was predefined in the clinical study protocol (i.e., before measuring sNfL or any clinical or radiological outcomes), as the median sNfL value for the overall population across ASCLEPIOS I/II (9.3 pg/mL)

• Participants were stratified as having high (≥9.3 pg/mL) versus low (<9.3 pg/mL) based on the median baseline sNfL concentration

• Quantification of sNfL levels was performed centrally (Navigate BioPharma Services, Carlsbad, CA, USA), as a single batch at the end of the trials, using a validated Quanterix Simoa<sup>®</sup> NF-light advantage kit

• MRI scans were performed at baseline, Months 12 and 24, and end of treatment/end of study

• The prognostic value of high versus low baseline sNfL for the annualized rate of neT2 lesions was assessed in baseline BMI

• Negative binomial regression model adjusting for sNfL group and sNfL group by BMI/age subgroup interaction was used to estimate the lesion rate ratio for high versus low sNfL levels in each BMI/age subgroup

• In Figure 2B, based on data from participants aged 18–55 years in ASCLEPIOS I and II, there is an association between BMI and sNfL levels, mainly driven by extreme values of BMI

 Underweight individuals (BMI<18.5) were associated with</li> higher mean sNfL levels, while extremely obese individuals (BMI>40) were associated with lower mean sNfL levels

- However, the magnitude of individual variability in sNfL driven by MS disease activity by far exceeded the magnitude of change explained by BMI (mean NfL ranges from 7.9 to 12.1 pg/mL, i.e., 4.2 pg/mL difference within the inner 90% of the BMI range)

• Thus, for most RMS patients in these studies,  $sNFL \ge 9.3 pg/mL$ was prognostic of future MS disease activity irrespective of BMI (Figure 1B)