



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

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- 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 vs 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 showed 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 T2 (neT2) 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

**BMI**, body mass index; **pwRMS**, people with relapsing multiple sclerosis; **Q**, quartile; **RMS**, relapsing multiple sclerosis; **sNfL**, serum neurofilament light chain.

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6. Consortium of Multiple Sclerosis Centers. *Int J MS Care*. 2021;23:1–36.
7. Bittner S, et al. *Brain*. 2021;144:2954–2963.
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- ASCLEPIOS I/II were two phase 3, double-blind, active-controlled trials in which participants with RMS were randomized to receive either ofatumumab 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,  $\geq 1$  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 into **high ( $\geq 9.3$  pg/mL)** and **low ( $< 9.3$  pg/mL)** sNfL groups based on the median baseline sNfL concentration

RMS, relapsing multiple sclerosis; sNfL, serum neurofilament light chain.  
Hauser SL, et al. *N Engl J Med*. 2020;383:546–557.

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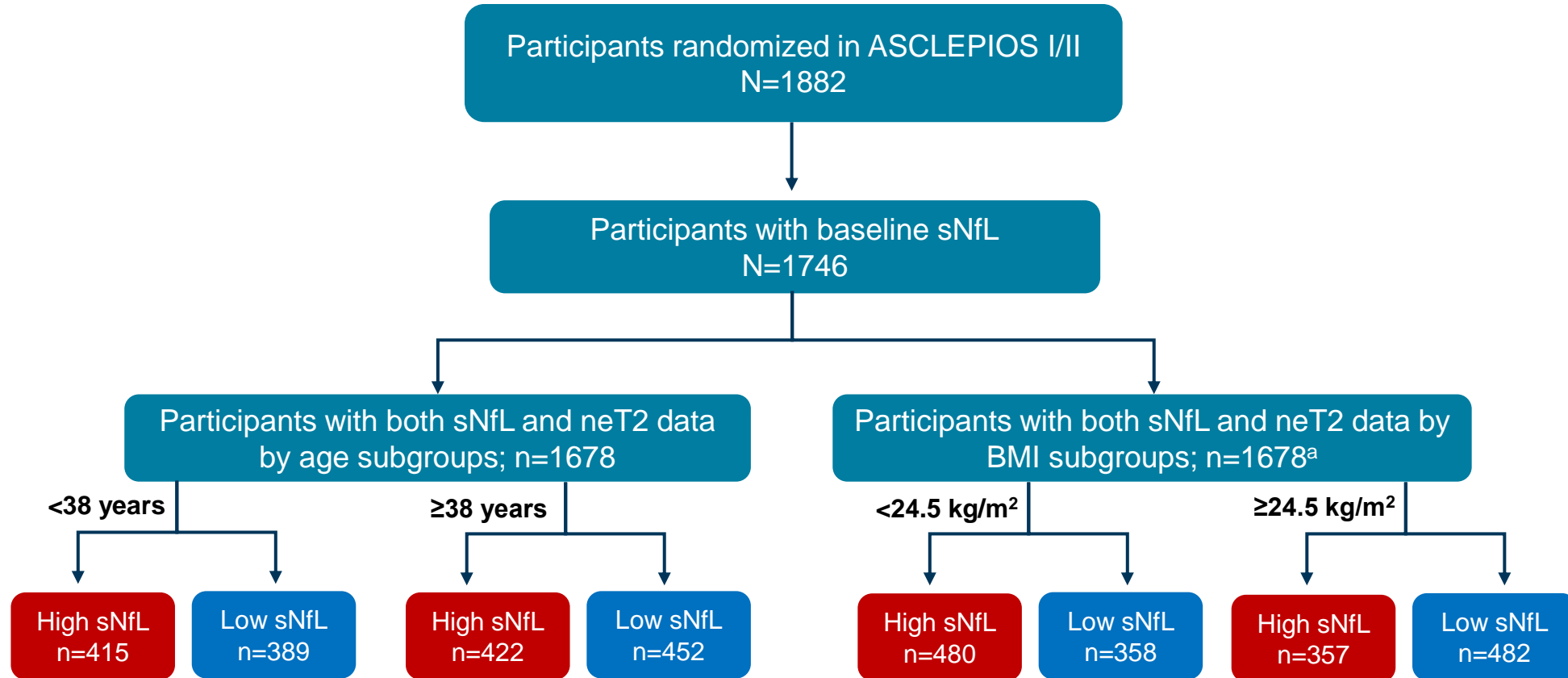


- Quantification of sNfL levels was performed centrally (Navigate BioPharma Services, Carlsbad, CA, USA), as a single batch at the end of the trials, using a validated Quanterix Simoa<sup>®</sup> NF-light advantage kit (Billerica, MA, USA)
- MRI scans were performed at baseline, Months 12 and 24, and 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 (<24.5 vs ≥24.5 kg/m<sup>2</sup>) and age (<38.0 vs ≥38.0 years) subgroups
- 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

**BMI**, body mass index; **MRI**, magnetic resonance imaging; **neT2**, new or enlarging T2; **sNfL**, serum neurofilament light chain.

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# Participants with sNfL and neT2 data available



<sup>a</sup>BMI data was not available for one patient.

**BMI**, body mass index; **neT2**, new or enlarging T2; **sNfL**, serum neurofilament light chain.

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# Demographic and disease characteristics by sNfL category at baseline



Characteristic	Low sNfL category (<9.3 pg/mL); N=876 <sup>a</sup>	High sNfL category (≥9.3 pg/mL); N=870 <sup>a</sup>
Age, years	38.6±8.5	37.8±9.7
BMI, kg/m <sup>2</sup>	27.0±6.4	24.7±5.5
Female, n (%)	588 (67.1)	602 (69.2)
MS duration since first symptom, years	8.3±7.2	7.9±6.9
Previously treated with DMT, n (%)	537 (61.3)	507 (58.3)
Number of relapses in the year before the study	1.2±0.7	1.3±0.7
Time since onset of most recent relapse, months	7.8±13.5	7.0±9.3
EDSS score	2.8±1.3	2.9±1.4
Normalized brain volume, cm <sup>3</sup>	1447.6±74.8	1437.2±81.0
Number of Gd+ T1 lesions	0.4±1.2	2.6±5.4
Patients free of Gd+ T1 lesions, n (%)	679 (77.5)	383 (44.0)
T2 lesion volume, cm <sup>3</sup>	9.4±10.6	16.7±15.0
Median sNfL, pg/mL	6.76	14.23

- Baseline demographic and disease characteristics were similar between sNfL groups, except the **mean number of Gd+ lesions** and **T2 lesion volume**, which were considerably higher in participants with high versus low baseline sNfL

<sup>a</sup>Only participants with non-missing baseline sNfL values are included. Data are expressed as mean±SD unless specified otherwise.

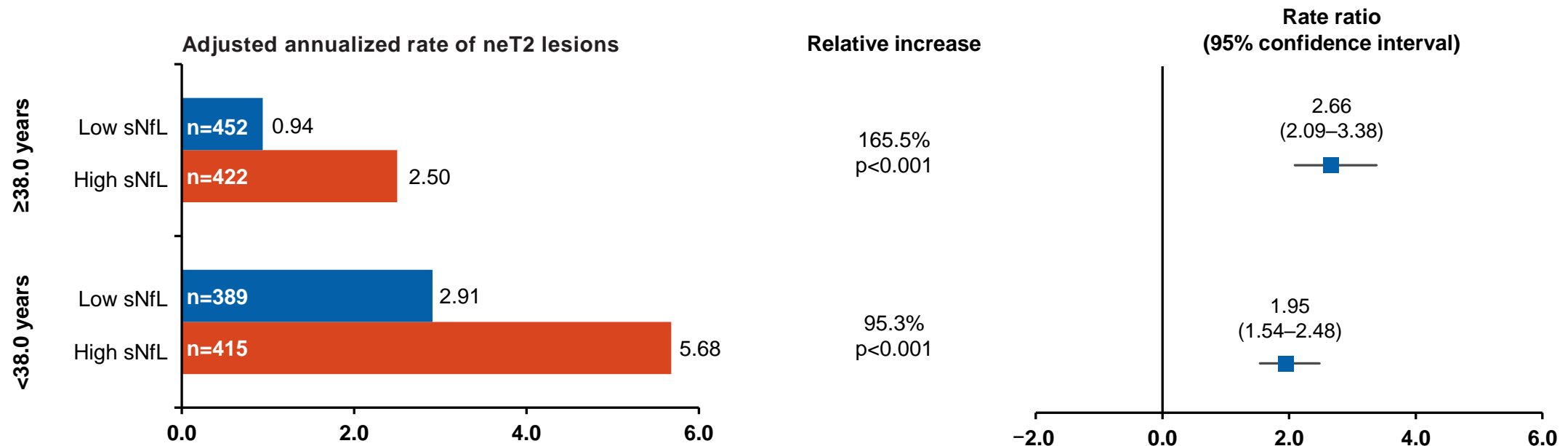
**BMI**, body mass index; **DMT**, disease-modifying therapy; **EDSS**, Expanded Disability Status Scale; **Gd+**, gadolinium-enhancing; **MS**, multiple sclerosis; **neT2**, new or enlarging T2; **SD**, standard deviation; **sNfL**, serum neurofilament light chain.

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# Prognostic value of sNfL for neT2 lesions by age subgroups



Mean annualized rate of neT2 lesions by age<sup>a</sup>



- Across age subgroups, the mean annualized rate of neT2 lesions was **significantly higher** in participants with high ( $\geq 9.3$  pg/mL) versus low sNfL ( $< 9.3$  pg/mL) levels

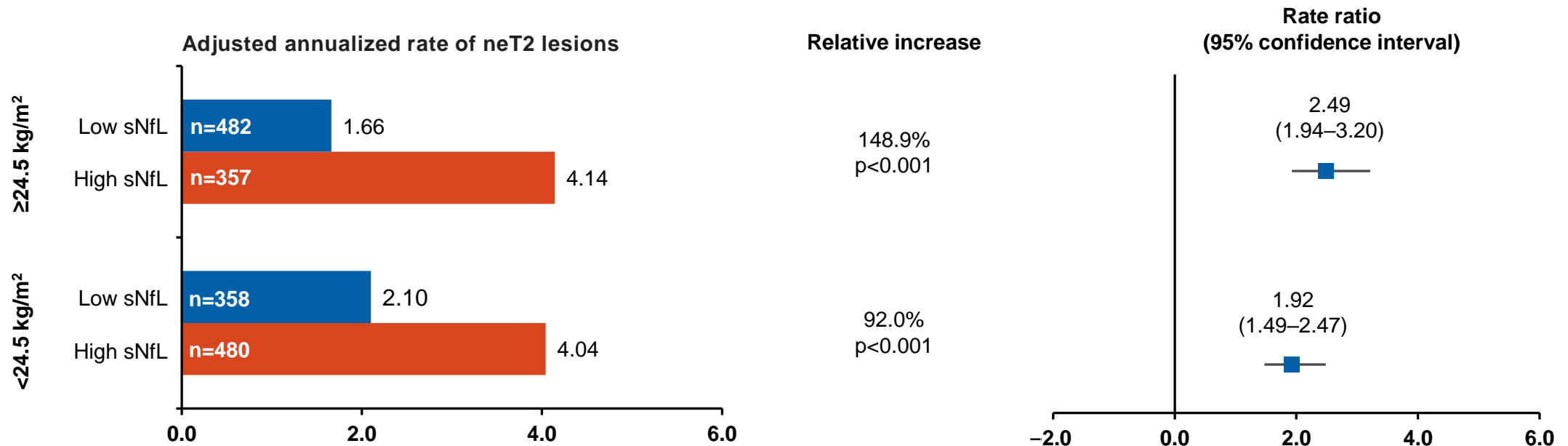
<sup>a</sup>Analyses were based on the population that had both baseline sNfL and neT2 data available.  
neT2, new or enlarging T2; sNfL, serum neurofilament light chain.



# Prognostic value of sNfL for neT2 lesions by BMI subgroups



Mean annualized rate of neT2 lesions by BMI<sup>a</sup>

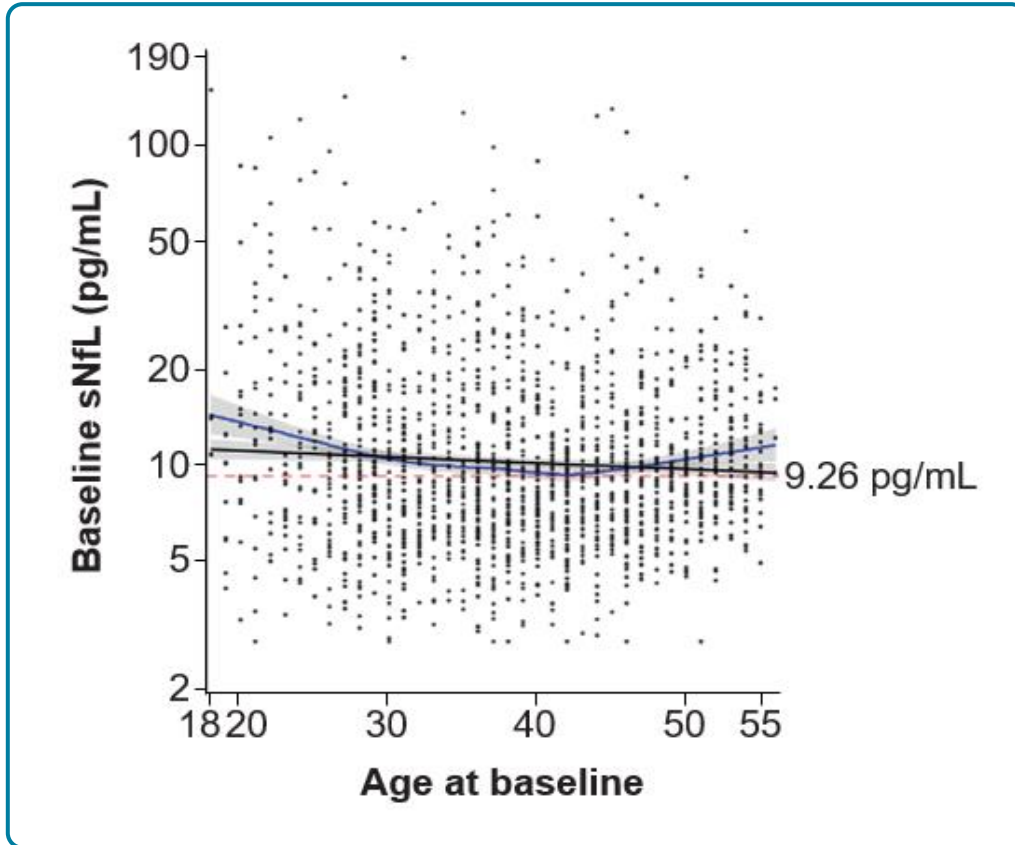


- Across BMI subgroups, the mean annualized rate of neT2 lesions was **significantly higher** in participants with high ( $\geq 9.3$  pg/mL) versus low sNfL ( $< 9.3$  pg/mL) levels

<sup>a</sup>Analyses were based on the population that had both baseline sNfL and neT2 data available.

**BMI**, body mass index; **neT2**, new or enlarging T2; **sNfL**, serum neurofilament light chain.

# Additional analyses of data using a single sNfL threshold by age



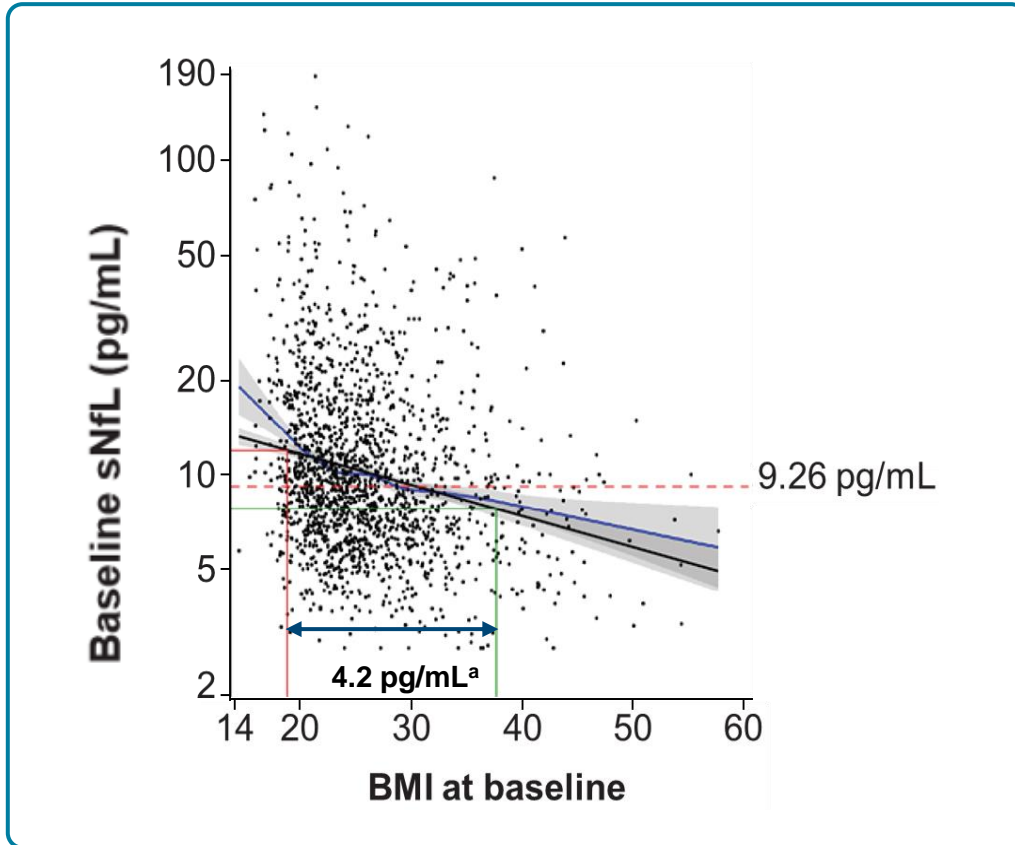
- Based on data from participants aged 18–55 years in ASCLEPIOS I and II, the variability in sNfL levels associated with age is much lower than the observed overall variability in sNfL (likely due to MS disease activity) in patients with RMS
- Compared to inflammatory-associated sNfL elevations, the age dependency of sNfL may not be a relevant confounder in clinical practice in younger RMS patients without comorbidities<sup>1</sup>

Black line is from linear regression of log (baseline sNfL) on age; blue line is when applying a non-linear relationship; dashed red line indicates median baseline sNfL.

**MS**, multiple sclerosis; **RMS**, relapsing multiple sclerosis; **sNfL**, serum neurofilament light chain.

1. Bittner S, et al. *Brain*. 2021;144:2954–2963.

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- 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 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 far exceeded the magnitude of change explained by BMI (mean sNfL 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  $\geq 9.3$  pg/mL was prognostic of future MS disease activity irrespective of BMI

<sup>a</sup>Difference in mean sNfL within the inner 90% of the BMI range.

Black line is from linear regression of log (baseline sNfL) on BMI; blue line is when applying a non-linear relationship; dashed red line indicates median baseline sNfL. The vertical and horizontal red lines 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.

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- 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 populations** (18–55 years) such as the 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 BMI range**, in those with extreme BMI (<18.5 and >40.0), prognostication using a single sNfL value should take into account the effect of BMI on sNfL
- The results reported here are based on the **protocol-defined single sNfL threshold**; future work should evaluate how 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 gadolinium-enhancing lesions on MRI scans for future lesion formation.<sup>1</sup> The relative prognostic value of sNfL and baseline MRI characteristics should be further confirmed

**BMI**, body mass index; **MRI**, magnetic resonance imaging; **RMS**, relapsing multiple sclerosis; **sNfL**, serum neurofilament light chain.

1. Bar-Or A, et al. *Neurol Ther*. 2023;12:303–317.

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- 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 pwRMS

**BMI**, body mass index; **MS**, multiple sclerosis; **pwRMS**, people with relapsing multiple sclerosis; **sNfL**, serum neurofilament light chain.

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