# CSF biomarker correlations with primary outcome in NurOwn Phase 3 clinical trial

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# **Background**

MSC-NTF cells (NurOwn®) are autologous bonemarrow derived mesenchymal stem cells (MSC) induced to secrete high levels of neurotrophic factors (NTFs). MSC-NTF cells were administered in three bimonthly intrathecal injections to ALS participants in a US Phase 3 multicenter double-blind placebocontrolled trial to evaluate safety and efficacy (NCT03280056).

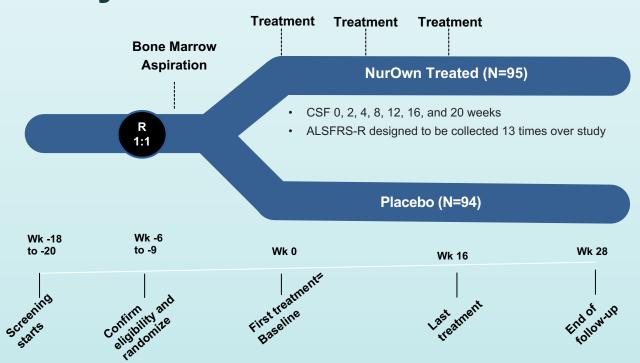
# **Objective**

To relate CSF biomarkers with primary ALSFRS-R outcome in the Phase 3 placebo-controlled trial.

# **Design/Methods**

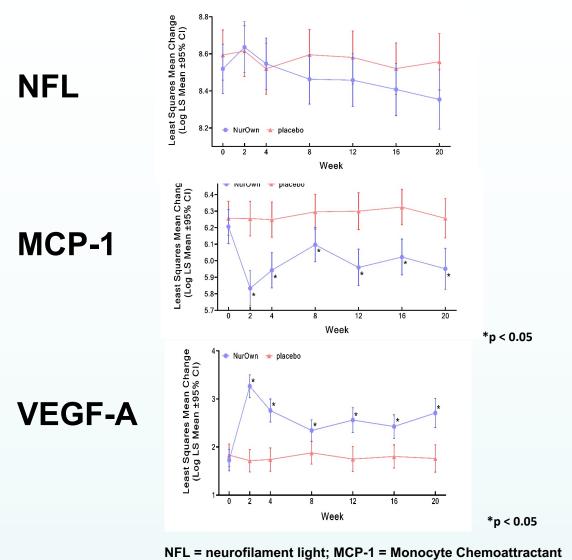
The primary endpoint was a responder analysis of the change in rate of ALSFRS-R decline, defined by an improvement of ≥1.25 points/month compared to the pre-treatment ALSFRS-R slope. CSF was collected at 7 time points, at baseline (before treatment) and through week 20. CSF biomarker selection (n=37) and biomarker statistical analyses were prespecified in a statistical analysis plan that was completed prior to unblinding. An unbiased stepwise logistic regression analysis was performed using log values (biomarker values and changes, and ENCALS model\*) to identify the minimum set of CSF biomarkers in addition to ENCALS (independent variables) that predicted the primary clinical outcome (dependent variable).

# **Study Schematic**



#### Results

CSF Neurodegenerative (NFL), Neuroinflammatory (MCP-1), and Neuroprotective (VEGF) biomarker changes over time (LSMean, 95% CI).



NFL = neurofilament light; MCP-1 = Monocyte Chemoattractant Protein-1; VEGF = Vascular Endothelial Growth Factor; MSR1 = Macrophage Scavenger Receptor Type 1

#### **Stepwise linear regression model:**

Predictive Biomarkers Show 82.5% Accuracy in ROC Analysis (NurOwn and Placebo Treatment Arms)

Stepwise logistic model  ENCALS	,			<i>'</i>
Model* Neurodegeneration   MCP-1 ✓, Δ   NFL ✓   Fetuin A ✓, Δ   VEGF-A ✓   S100B ✓   Neurodegeneration Neuroprotection ✓ : biomarker values Δ : change in biomarker	logistic			
NFL  Fetuin A  VEGF-A  S100B  ✓  ✓ : biomarker values  △ : change in biomarker		✓	✓	<u> </u>
Fetuin A  VEGF-A  S100B  ✓ , Δ  ✓ : biomarker values  Δ : change in biomarker	MCP-1		√, Δ	Neuroprotection
Fetuin A  VEGF-A  S100B  ✓, Δ  ∴ change in biomarker	NFL	✓	✓	
S100B ✓	Fetuin A		√, Δ	
	VEGF-A		✓	
MSR1 ✓	S100B	✓		
	MSR1		✓	

 Reflective of overall baseline health of participants.
 ENCALS Model uses terms: Age of onset, FVC, Duration from onset of symptoms to first treatment, Bulbar onset, ALSFRS-R slope, 'Definite' ALS

#### **Discussion**

- ➤ ALS is a complex disease in which neurodegeneration, neuroinflammation and failure of intrinsic neuroprotective mechanisms may play an important role¹
- ➤ NurOwn treatment resulted in consistent changes in CSF neurodegenerative (NFL), neuroinflammatory (MCP-1), and neuroprotection (VEGF-A) biomarkers, while placebo levels remained stable.
- ➤ A detailed analysis of CSF biomarkers in this phase 3 trial suggests that the primary clinical responder outcome may be predicted by a combination of neurodegenerative, neuroinflammatory and neuroprotection biomarkers.
- ➤ Statistical Modeling highlights biomarkers that are predictive of NurOwn treatment response with good accuracy (82.5%)

#### **Conclusions**

➤ Predictive CSF biomarkers identified in the stepwise logistic regression analysis will be an important step in furthering our understanding of the mechanism of action of NurOwn in ALS as well as in future studies that use CSF biomarkers to advance ALS science.

1. Brown, Al-Chalabi NEJM 2017