



NurOwn Targets Multiple Disease Pathways in ALS Phase 3 Trial: A Biomarker Analysis

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Phase 3 Trial of NurOwn® in ALS Patients : Protocol

Double-blind, Placebo-controlled, Randomized Trial



Inclusion criteria:

- ALSFRS-R ≥ 25 at Screening
- Upright SVC ≥ 65% of predicted for gender, height, and age at Screening
- Onset of ALS disease symptoms within 24 months of Screening
- Decline in ALSFRS-R total score of 3 or more points in the 12 weeks before randomization.
- ALS diagnosed by revised El Escorial criteria as possible, laboratory-supported probable, probable, or definite

The primary and secondary endpoints failed to reach statistical significance in the overall study population Pre-specified and post-hoc analyses suggest a potential treatment effect with MSC-NTF across the primary and secondary efficacy

Three Pathways Interrogated by Biomarkers

• Neurodegeneration (5)

NfL, UCH-L1, pNfH, DR6, Fetuin-A

• Neuroprotection (7)

Follistatin, VEGF, Clusterin, BDNF, OPG, GDF-15, LIF

• Neuroinflammation (9)

CHI3L1, Chit-1, IP-10, MCP-1, Trem-2, SDF1a, S100B, MSR1, NMNAT1

Three Approaches to Mapping Biomarkers onto Disease Pathways

- Analysis of Individual Biomarkers
- Principle component analysis of all biomarkers simultaneously within each pathway
- Regression modeling using all biomarkers to predict outcome

NurOwn significantly lowers pro-inflammatory biomarkers over time compared to placebo

MCP-1: Statistically significant reduction with NurOwn at each visit, and significant overall treatment effect

SDF-1a: Statistically significant reduction with NurOwn, significant overall treatment effect



NurOwn lowers some biomarkers of neurodegeneration over time compared to placebo



NfL: NurOwn values lowered 82% compared to Placebo at week 20

DR6: Statistically significant reduction with NurOwn, significant overall treatment effect

NurOwn significantly elevates neuroprotective biomarkers over time compared to placebo

VEGF: Statistically significant increase with NurOwn at each visit, and significant overall treatment effect (2-fold increase)

BDNF: Statistically significant increase with NurOwn, with overall significant treatment effect



Three Approaches to Mapping Biomarkers onto Disease Pathways

- Analysis of Individual Biomarkers
- Principle component analysis of all biomarkers simultaneously within each pathway
- Regression modeling using all biomarkers to predict outcome

Statistical Analysis of Biomarker data through Principal Components

NurOwn therapy reduces markers of Neuroinflammation and Neurodegeneration, and increases Neuroprotection

Trend with individual biomarkers confirmed across biomarkers with PCA

- Neuroinflammation decreased with NurOwn treatment over 20 weeks
 - Placebo Neuroinflammation levels remain high
- Neurodegeneration decreased with NurOwn
 - Placebo Neurodegeneration levels decrease across the study, remain elevated relative to NurOwn.
- Neuroprotection increased with NurOwn treatment and is maintained over 20 weeks
 - Placebo Neuroprotective levels remain low

Biomarkers driving PCA results:

- Neuroinflammation (5/9): CHI3L1, Chit-1, IP-10, MCP-1, Trem2
- Neurodegeneration (3/4): NfL, UCH-L1, pNfH
- Neuroprotection (4/7): Follistatin, VEGF, Clusterin, BDNF



Three Approaches to Mapping Biomarkers onto Disease Pathways

- Analysis of Individual Biomarkers
- Principle component analysis of all biomarkers simultaneously within each pathway
- Regression modeling using all biomarkers to predict outcome (Stepwise forward regression model machine learning algorithm)

Final Model identifies Biomarkers predictive of Primary Endpoint with high accuracy

Leveraging model with biomarkers, treatment effect on Primary Endpoint is statistically significant (p=0.003)

| Stepwise logistic model | Post baseline |
|----------------------------|---------------|
| ENCALS Model [*] | \checkmark |
| MCP-1 | √, Δ |
| NfL | \checkmark |
| Fetuin A | √, Δ |
| VEGF | \checkmark |
| MSR1 | \checkmark |

Neuroinflammation
Neurodegeneration
Neuroprotection



- Baseline health was important to predicting clinical response
 - ENCALS model term
- Biomarker data was important to the prediction of clinical response
 - Markers of neuroinflammation, neurodegeneration and neuroprotection were all in the final model
 - Highlights the importance of all 3 pathways
- Information about the magnitude of change for MCP-1 and Fetuin-A contributed to the accuracy of the model prediction
 - ➢ larger change in MCP-1 more likely to respond
 - Iarger change in Fetuin-A more likely to respond

* ENCALS model Reflective of overall baseline health of participants.

Model uses terms: Age of onset, FVC, Duration from onset of symptoms to first treatment, Bulbar onset, ALSFRS-R slope, 'Definite' ALS

NurOwn response: Stepwise linear regression model to select predictive biomarkers 82.5% accuracy of model predicting clinical response



Receiver Operating Curve

- ➤ Measures accuracy of the model
- ➢Area Under the Curve (AUC)
- ≻ Values range from 0-100%
 - ≻50% is equivalent to a random guess
 - >80% represents high classification accuracy

Conclusions

- NurOwn significantly and consistently elevates markers of neuroprotection, and significantly lowers markers of neuroinflammation and neurodegeneration over time compared to placebo
 - > The same conclusion is drawn from individual markers and summaries of markers using PCA
- Prespecified statistical modeling leveraging machine learning highlights biomarkers that are predictive of treatment response with NurOwn with high accuracy (>80%)
 - Markers of neuroinflammation, neurodegeneration and neuroprotection were <u>all selected</u> in the final model
- These results enhance our understanding of NurOwn's mechanism of action in ALS
- NurOwn cell therapy targets multiple disease pathways and provides additional evidence linking the mechanism of action to NurOwn's impact on ALS progression (primary endpoint).

Manufacturing Sites



City of Hope Center for Biomedicine & Genetics



Dana-Farber Harvard Cancer Center Cell Manipulation Core















Funding







Thank You !!

Extra slides

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Biomarkers with significant explanatory value for Clinical Response, NurOwn and Placebo

| | NurOwn Phase 3 | | Description | Published relationships |
|---------------------------|-------------------|------------------|---|---|
| | Baseline value | Post baseline | Mechanism of Action | CSF levels correlate with ALS survival |
| ENCALS Model [*] | \checkmark | \checkmark | | \checkmark |
| MCP-1 | | √, Δ | B-chemokine Neuroinflammation | \checkmark |
| NfL | \checkmark | \checkmark | Specific marker of neuronal injury, possible biomarker of neurological disease activity | \checkmark |
| Fetuin-A | | √, Δ | Inhibits proinflammatory mediator pathways | |
| S100b | \checkmark | | Astrocyte inflammatory mediator, RAGE activation | \checkmark |
| VEGF | | \checkmark | Neuroprotection Growth factor | |
| MSR1 | | \checkmark | Anti-inflammatory, Macrophage scavenger receptor DAMP clearance | |

Neuroinflammation, Neurodegeneration,

Neuroprotection

: biomarker values

* Reflective of overall baseline health of participants.

Model uses terms: Age of onset, FVC, Duration from onset of symptoms to first treatment, Bulbar onset, ALSFRS-R slope, 'Definite' ALS | Δ

: change in biomarker

NurOwn MOA: Lessons learned from Phase 2, Replicated in Phase 3

Neuroinflammation, Neurodegeneration, and Neuroprotection Biomarkers

