

# QurAlis

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## **Biomarkers for Patient Stratification and Target Engagement in Patients with TDP-43 Pathology**

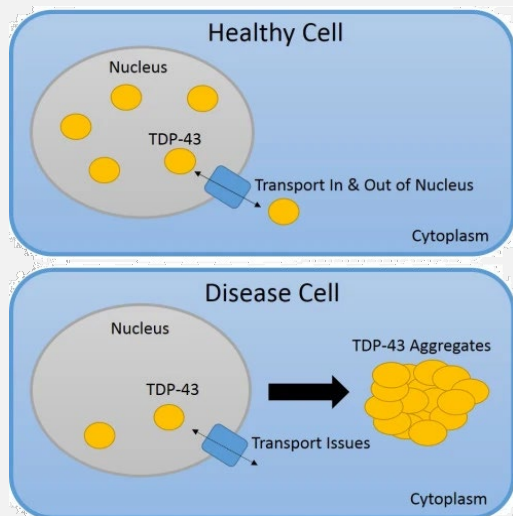
**Sandy Hinckley, Ph.D.**  
**Head of Biology**



# STMN2: A Genetic Target For The Sporadic ALS Population and TDP43opathies

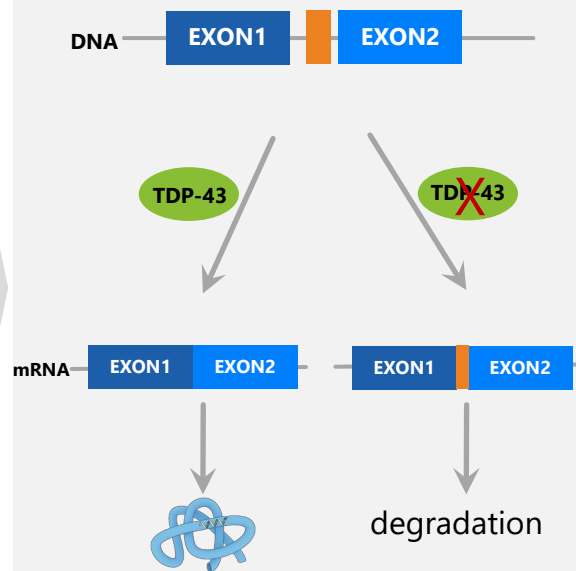
## QurAlis Therapeutic Strategy

In ALS motor neurons  
TDP-43 leaves the nucleus



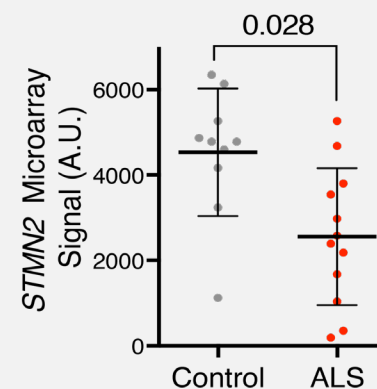
QurAlis niche

Loss of TDP-43 controlled  
cryptic exon splicing



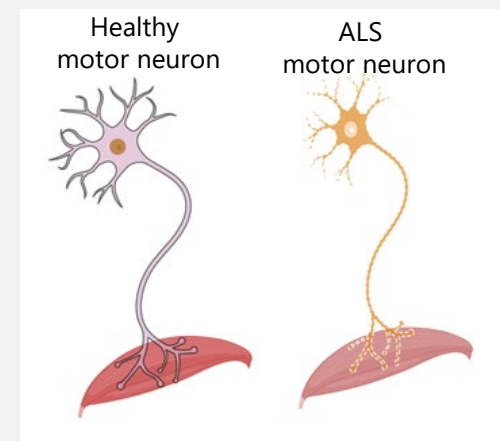
TDP-43 regulation of *STMN2*

Loss of full  
length *STMN2*



Cryptic splicing-ASO approach  
to restore *STMN2*

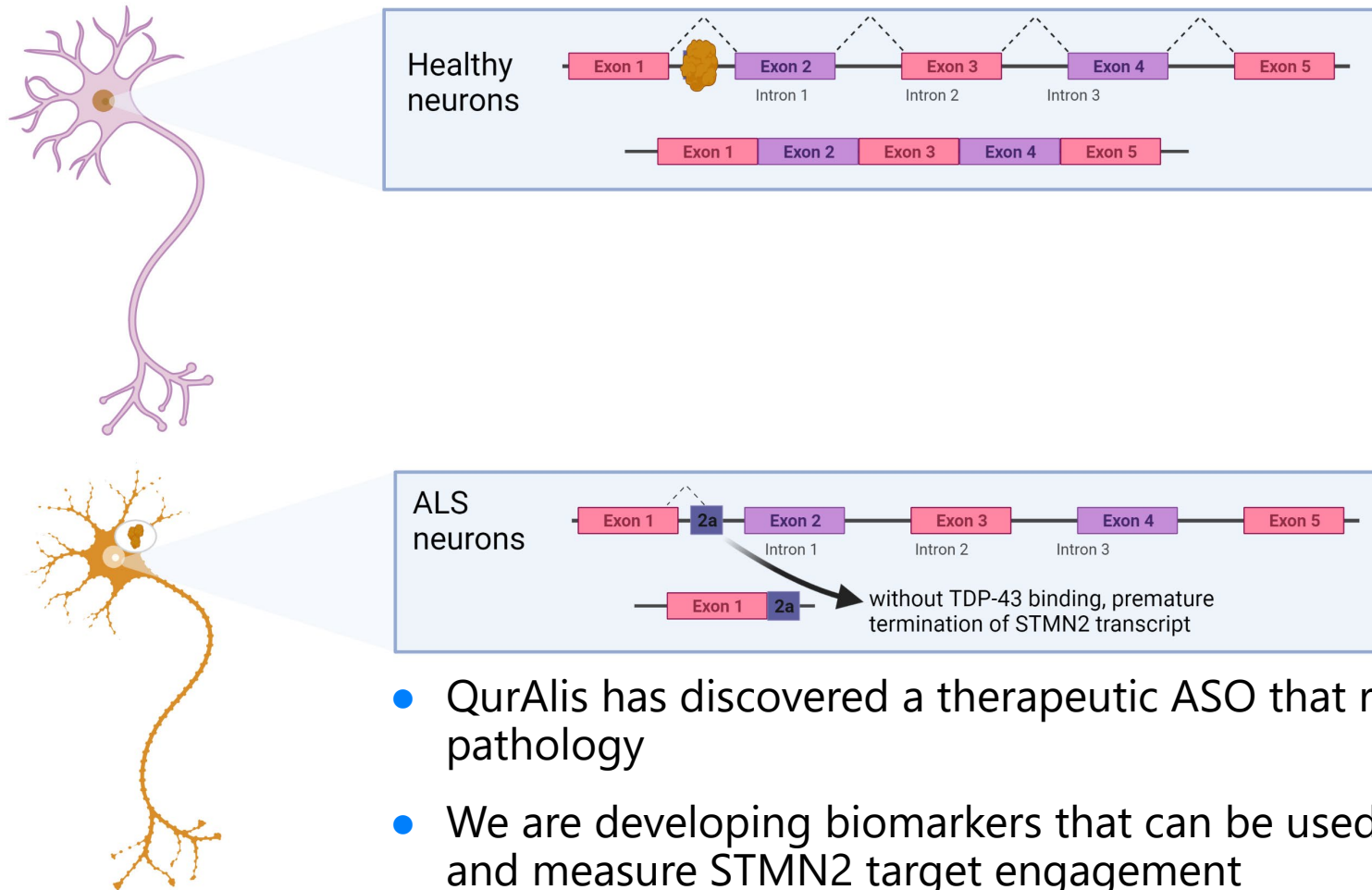
Axonal degeneration  
and impaired repair



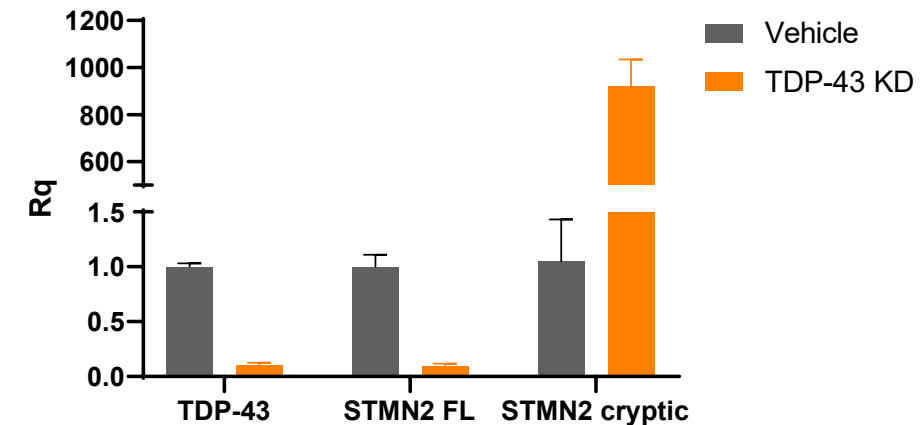
Rescue of axonal stability and  
repair

# STMN2 is a downstream target of TDP-43 function

- TDP-43 binding to exon 2a in the STMN2 pre-mRNA controls splicing and expression of STMN2 full length transcripts

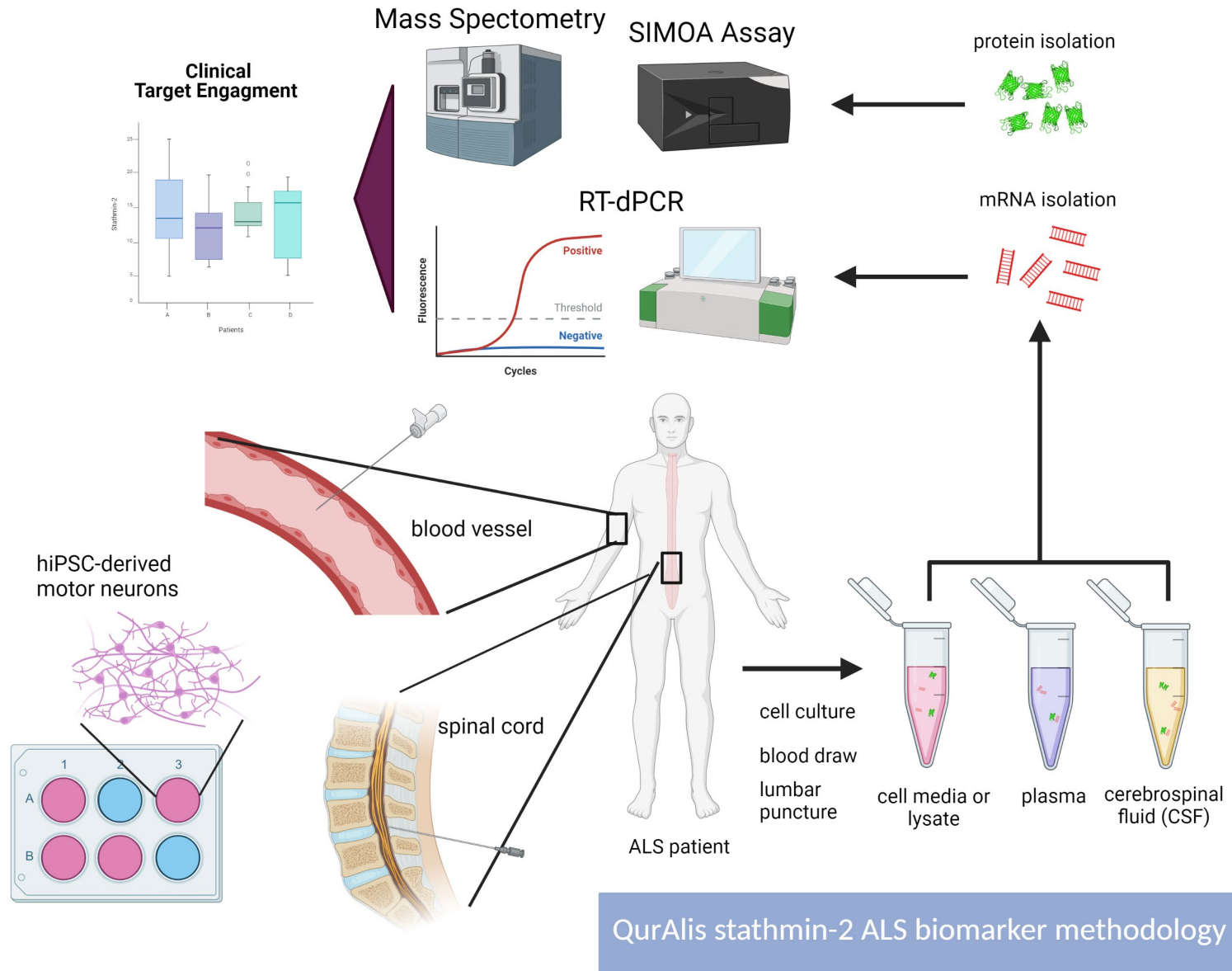


TDP-43 loss of function causes STMN2 mis-splicing

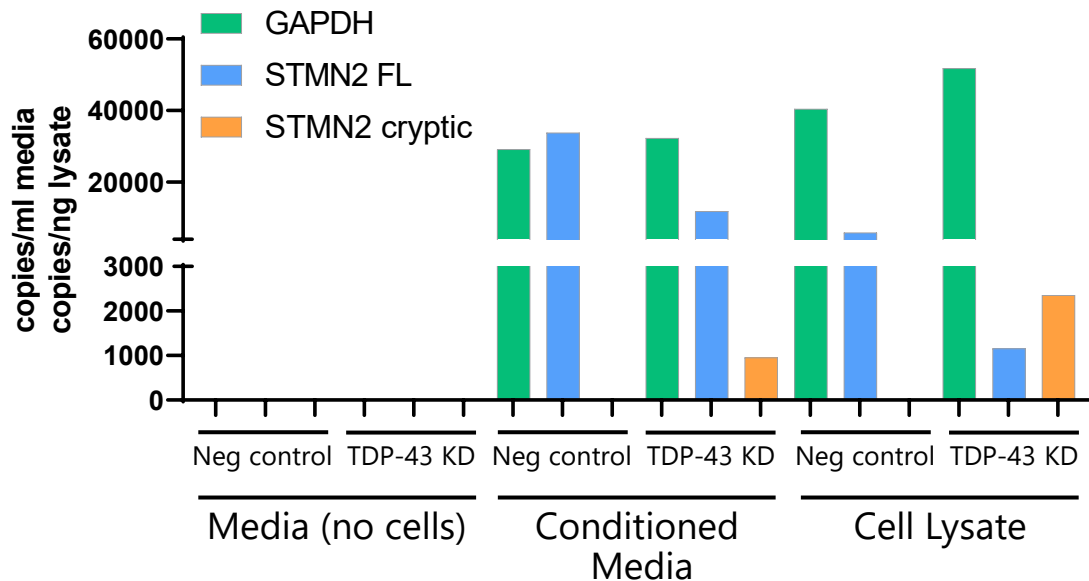
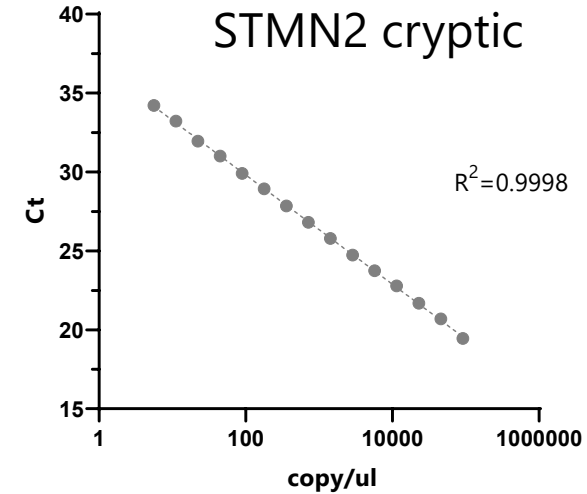
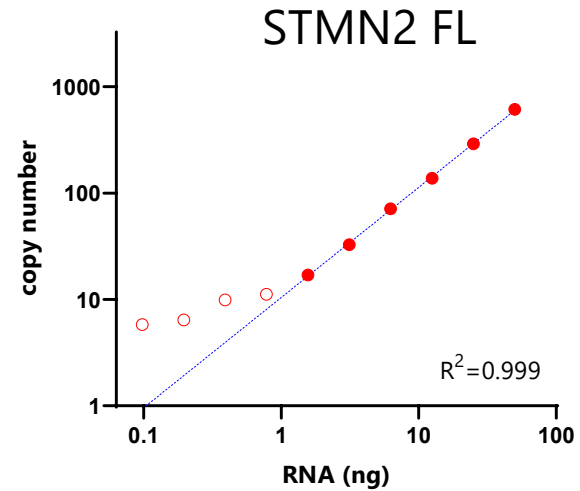
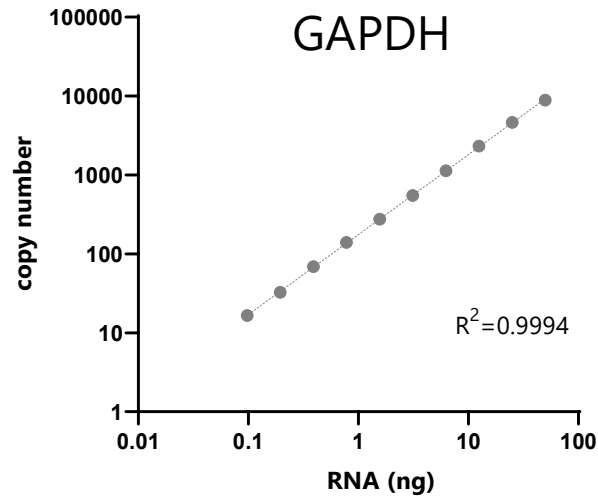


- QurAlis has discovered a therapeutic ASO that restores *STMN2* mis-splicing due to TDP-43 pathology
- We are developing biomarkers that can be used to identify patients with TDP-43 pathology and measure *STMN2* target engagement

# STMN2 biomarker development

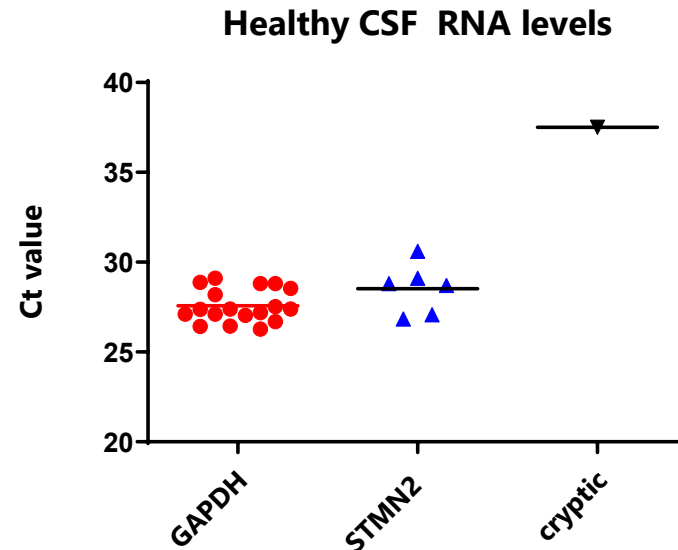
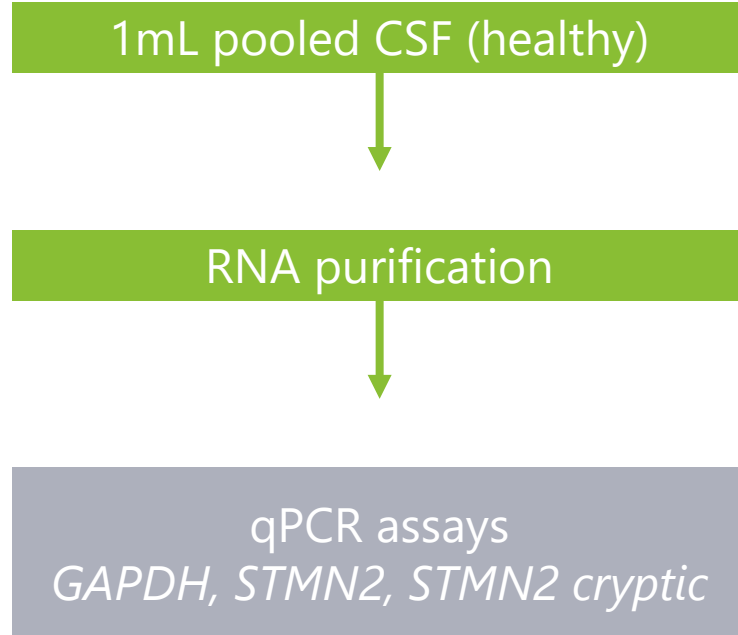


# Detection of STMN2 transcripts in hiPSC-derived motor neuron cultures



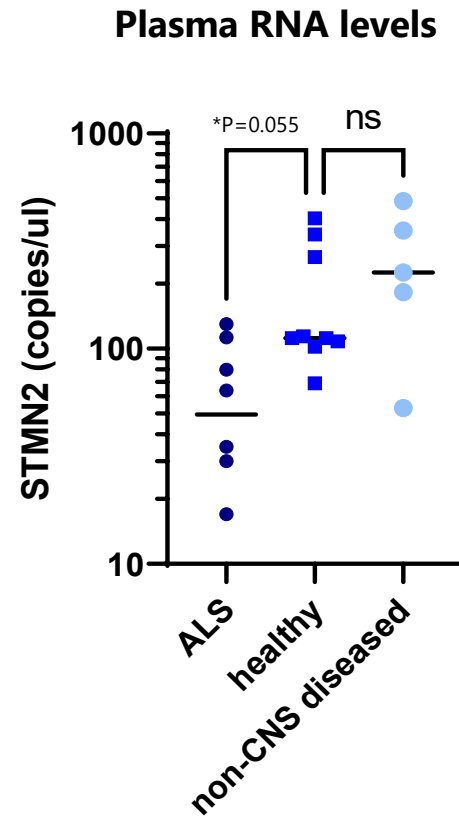
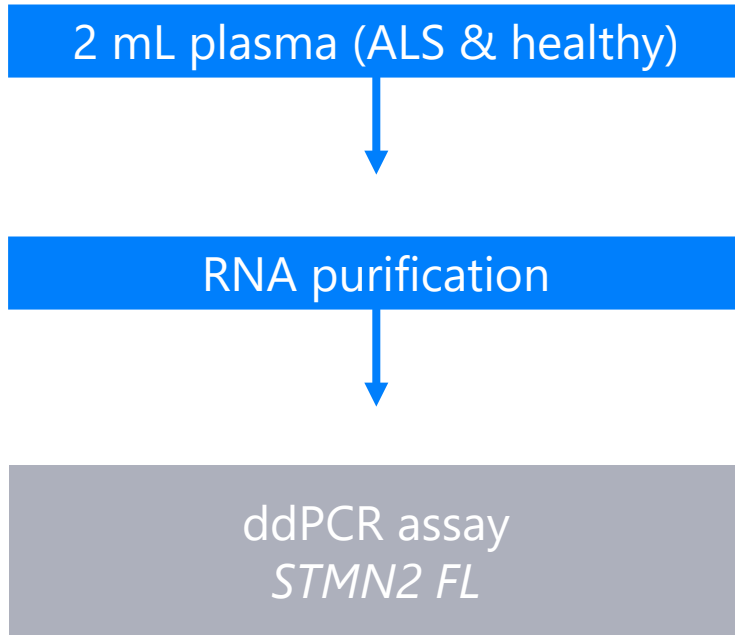
- ddPCR for GAPDH and STMN2 FL shows assay linearity down to 10 copies
- STMN2 cryptic assay was evaluated using G-block DNA and qPCR testing shows assay linearity
- Quantitation of STMN2 FL and cryptic exon STMN2 mRNA isolated either from conditioned media or cell pellets of hiPSC-derived motor neurons
- Both STMN2 FL and STMN2 cryptic exon mRNA is detectable in conditioned media
- Fluid STMN2 FL and cryptic levels represent the cellular biology and show STMN2 FL decrease and STMN2 cryptic increase upon mis-splicing induction

# STMN2 RNA quantification in human cerebrospinal fluid (CSF)



- Healthy CSF was assayed for the presence of *STMN2* RNA
- *STMN2* RNA was detected at high quantity in CSF about the same as the reference *GAPDH*
- Cryptic exon containing transcripts were mainly not detected, as expected in healthy samples

# STMN2 RNA quantification in human plasma



- ALS, healthy, and non-CNS disease plasma was assayed for the presence of *STMN2* RNA
- *STMN2* RNA was detected in 21/22 samples. 1/8 ALS sample had no measurable *STMN2*
- Trend toward decreased *STMN2* RNA level in ALS plasma

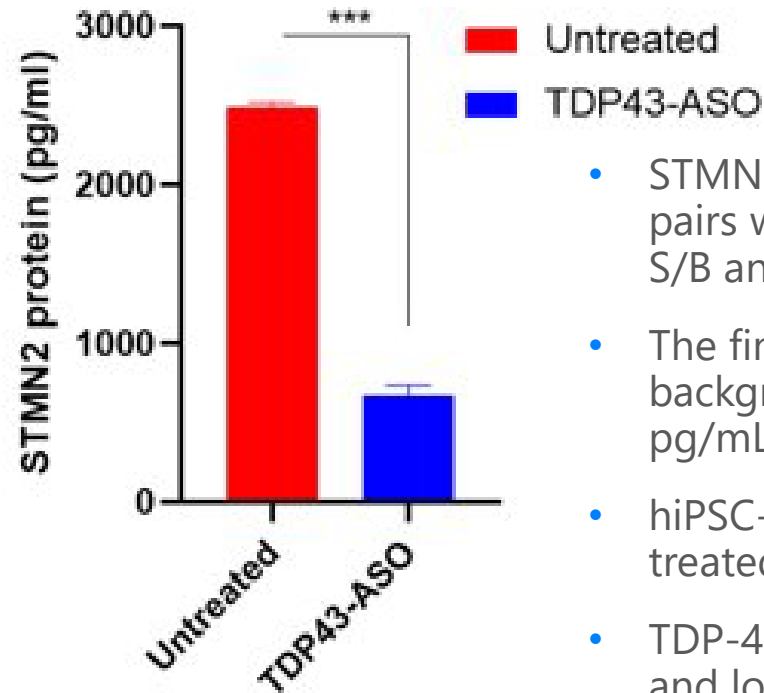
N=8 ALS  
N=9 healthy  
N=5 non-CNS disease controls  
Kruskal-Wallis test  
ANOVA  
Dunn's MC test

# SIMOA assay for detection of STMN2 protein

Calibration Curve			
[stathmin-2] pg/mL	Average AEB	%CV	S/B
0.0	0.005	22%	
0.4	0.012	2%	2
1.5	0.034	8%	6
5.9	0.120	10%	22
23.4	0.450	13%	82
93.8	2.282	13%	417
375.0	8.023	0%	1466
1500.0	22.500	2%	4111

LOD: 0.087 pg/ml  
LLOQ: 0.33 pg/ml

## Motor Neuron Lysates

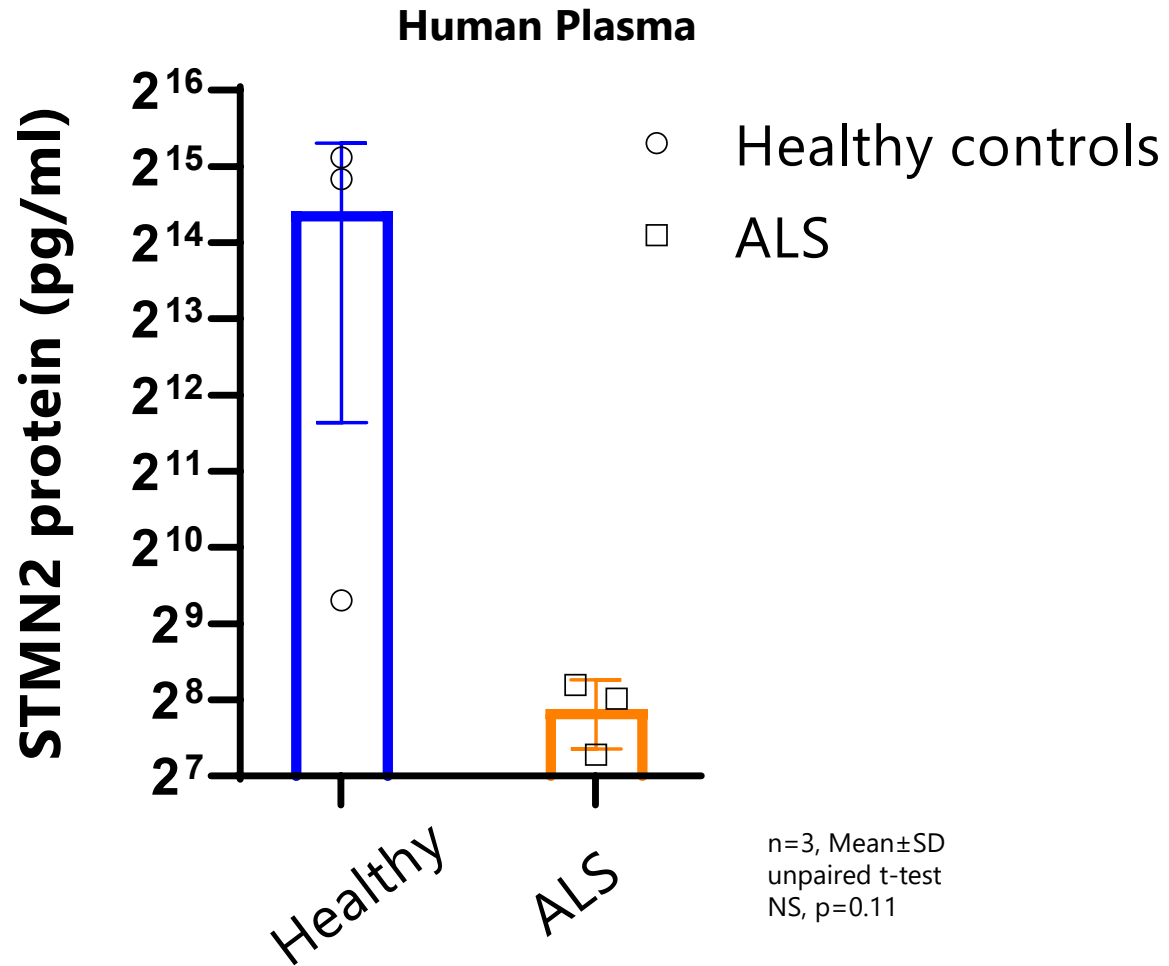


n=2, Mean±SD  
unpaired t-test  
\*\*\*p<0.005

- STMN2 capture and detection antibody pairs were tested and the 2 with lowest S/B and dilution linearity were selected
- The final assay format shows low background levels (AEB<0.01) and sub-pg/mL LLOQ
- hiPSC-derived motor neurons were treated with/out TDP-43 gapmer ASO
- TDP-43 KD results in STMN2 mis-splicing and loss of STMN2 protein levels
- STMN2 SIMOA assay detects changes in human STMN2 levels due to TDP-43 loss of function



# STMN2 protein quantification in human plasma



- Healthy and ALS plasma was tested in the STMN2 SIMOA assay (n=3 individuals each)
- Healthy control and ALS samples within the dynamic range of the assay
- ALS STMN2 levels are 50% reduced compared to the lowest healthy sample and 98% reduced compared by mean values

# Conclusions

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- We have developed both RNA and protein quantitation methods to detect STMN2 in human biofluids
- STMN2 biomarker assay allows the characterization of TDP-43 pathology and downstream target biology in living ALS patients
- Next steps include:
  - Assay qualification and validation for use as a clinical exploratory Biomarker
  - Characterization of larger patient cohorts for protein assay from annotated clinical libraries
  - Expansion of human data set into other TDP43opathies (FTD, AD, PD)

\*\*check out QurAlis's other presentation by Taylor Gray on STMN2 biology

# Acknowledgements

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Contributions by:

Taylor Gray, Scientist

Hafiz Mohmmadabdul, Senior Scientist

Daniel Elbaum, Chief Scientific Officer

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