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MeiraGTx | Gene therapy for Amyotrophic Lateral Sclerosis (ALS) and Frontotemporal Dementia (FTD) Preclinical efficacy of AAV-hUPF1 optimized for clinical translation with improved vector genome and novel CNS capsid.

ESGCT October 2023 | T.D. Barbara Nguyen-Vu, MBA, PhD



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Amyotrophic Lateral Sclerosis (ALS)

Program Overview

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- ALS is a severe neurodegenerative disease affecting motor neurons with mean survival of 2-5 years.
- Only 5-10% of ALS cases are inherited, familial ALS (fALS), whereas the rest are sporadic (sALS).
- MeiraGTx is developing a first-in-class gene therapy to potentially address **both** forms of ALS :

Therapeutic Target: AAV-hUPF1

Targets underlying cellular defects associated with different forms of ALS

Validated in multiple in-vivo and in-vitro models of ALS, in the context of AAV



TPD43 Proteinopathy is a Cellular Hallmark of ALS

Only 5-10% of ALS is genetically driven



- Only 10% of ALS cases are inherited, familial ALS (fALS); remaining 90% of cases are sporadic (sALS)
- More than 20 genes were found to be associated with fALS

TDP43 cellular mislocalization and aggregation is present in 97% of ALS cases



Feneberg et al. 2018

- TDP43 mis-localization and aggregation is a cellular hallmark of ALS (*except for SOD1driven ALS*)
- TDP43 cytoplasmic granules are also observed in ~50% of frontotemporal dementia (FTD) patients

hUPF1 Identified as a Potential Therapeutic Target for ALS in a Yeast Genome-Wide Screen

ECM32, yeast homologue of UPF1, protects against FUS (and TDP43) cytotoxicity



Human UPF1 and UPF2 rescue FUS (and TDP43) cytotoxicity in yeast



- Overexpression of TDP43 or FUS recapitulates cellular hallmarks of ALS (i.e., mislocalization, aggregation and cytotoxicity)
- Only 5 out of 5,500 genes rescued all were <u>RNA binding proteins</u>
- ECM32, a homologue of UPF1, was most potent
- <u>human UPF1</u> (or UPF2) also rescued the toxic phenotype of FUS and TDP43 in yeast

UPF1 is a Key Player in RNA Metabolism and Homeostasis Pathways Dysregulated in ALS

UPF1 plays a central role in RNA metabolism & quality control; mechanisms dysregulated in ALS:

- UPF1 is a core component of the Nonsense-Mediated Decay (NMD) pathway
- UPFI has multiple roles beyond NMD, including targeting misfolded proteins to the 'aggresome' for clearance
- Dysregulation of RNA metabolism is known to contribute to ALS pathogenesis:
 - Mutations such as C9orf72, TDP43 and FUS, have been found to disrupt RNA processing
 - Aberrant RNA species & impaired clearance may disrupt cellular homeostasis and contribute to pathological protein aggregates, such as TDP43 - a cellular hallmark of ALS.

AAV-UPF1 potentially targets cellular defects underlying ALS, providing a way to address both familial and sporadic forms of the disease (>95% of patients)



UPF1 Decreases Risk of Neuronal Death by TDP-43 and FUS

In primary rodent neurons, **hUPF1 expression rescues** FUS & TDP43 cytotoxicity in a dose-dependent manner



Protective Role of UPF1 is Specific to TDP43 proteinopathy

hUPF1 expression rescues **specific** to FTD and ALS associated with TDP43 proteinopathy (>95% of ALS cases)



 UPF1 unable to rescue <u>SOD1 ALS</u> mutant cytotoxicity, a form of fALS which lacks cytoplasmic inclusions of TDP43 mHTT (Huntington's)



 UPF1 unable to rescue cytotoxicity induced by <u>mutant Huntington</u> (mHTT)



 No effect in WT primary neurons of UPF1 overexpression

Barmada, S. J. et al. (2015). Amelioration of toxicity in neuronal models of amyotrophic lateral sclerosis by hUPF1. Proc Natl Acad Sci, 112(26), 7821-7826.

UPF1 Validation in Multiple in-vivo Models of ALS

Collaborations with leading ALS researchers to <u>validate UPF1 in multiple in-vivo models</u> of ALS provide **strong evidence** for the **therapeutic potential of UPF1**



In-vivo validation: Functional Evidence for Therapeutic Potential

In-vivo model 1: TDP43 neonate rat model

AAV-UPF1 in **neonate rescues escape reflex** impairments induced by TDP43



Jackson et al. Gene Ther. 2015

In-vivo model 3: conditional FUS mouse model

AAV-UPF1 in FUS mouse model **protects against motor neuron loss** & improves hindlimb grip strength

Unpublished using model in Korobeynikov et al.Nat.Med. 2022



In-vivo model 2: TDP43 adult rat model

AAV-UPF1 in **adult rat rescues escape reflex** impairments induced by TDP43



In-vivo model 4: mouse C9orf72 model

AAV-UPF1 in (G4C2)66 mouse model **protects against neuron loss** & improves locomotor activity

Unpublished using model in <u>Chew et al. Science 2015</u>



UPF1 Construct Optimization for the Clinic

Optimized to decrease size, increase production yield, and increase potency



Optimized hUPF1 Constructs Validated in iNeuron Models of ALS

All optimized AAV-UPF1 **constructs rescued**, in parallel testing, of optimized hUPF1 constructs in human iNeuron TDP-43 Disease Model



Optimized hUPF1 Decrease Risk of Death in iNeuron Models of ALS

hUPF1 expression **promotes survival** of human iPSC-derived neurons (iNeurons) with isogenic mutations of **TDP43 or C9orf72**



Optimized hUPF1 Promotes Neuronal Survival In-Vivo in FUS mouse

Optimized AAV-UPF1 constructs **rescued loss of motor neurons in-vivo** in FUS mouse model



- 36% AAV transduction efficiency of motor neuron via neonate ICV was sufficient for in-vivo rescue
- AAV-hUFP1 gene therapy protective against motor neuron loss in FUS mouse model

Novel AAV Capsids for Broad CNS Transduction



• AAV2-retro variants transduced both upper and lower motor neurons by ICM injection in mice



Novel AAV Capsid Transduces Spinal Motor Neurons in NHP

AAV2-retro/rh10 mosaic produced broad CNS **transduction** of neurons **in both brain and spinal cord** of NHPs via intra-cisterna magna (ICM) administration





Transduction in NHP Neurons in Brain



Transduction in spinal motor neurons at levels
sufficient to drive in-vivo rescue in mice

Confirmed transduction in cortical
neurons via ICM route of administration

Clinical AAV-hUPF1 Validated in ALS Patient-Derived Neurons

ALS Patient-derived iPSC Model: Myotubes & Neurons

Myotubes: ALS3 C9 patient

Neurons: ALS3 C9 patient



Efficacy of AAV-hUPF1 in C9 ALS Patient-Derived Neurons

Clinical AAV-hUPF1 candidates demonstrate **reproducible target engagement in C9** ALS patient-derived neurons



• Exp. 1: 2 weeks after AAV transduction



• Exp. 2 : 1.5 weeks after AAV transduction

AAV-hUPF1 Rescued in Multiple ALS Patient-Derived Lines

AAV-hUPF1 efficacy in ALS patient-derived lines of different genetic context: C9 & TDP43



TDP43 (A382T) Patient Line







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AAV-UPF1 Program **HIGHLIGHTS** AAV-UPF1 is a novel gene therapy for ALS with the potential to treat both familial and sporadic ALD (>95% of patients) and FTD:

- UPF1 is a novel therapeutic target for ALS discovered in an unbiased genetic screen in yeast and validated in multiple in-vitro and in-vivo models of ALS
- AAV-UPF1 targets an underlying cellular defect of ALS- RNA metabolism and homeostasis. UPF1 is known to play a central role in RNA regulation, including in Nonsense-Mediated Decay (NMD).
- In-vivo studies in multiple models of ALS have demonstrated the ability of AAV-UPF1 to reduce neuronal death and ameliorate ALS symptoms related to limb strength and mobility
- A proprietary capsid, AAV2-retro, demonstrates favorable transduction of upper and lower motor neurons

Status:

- Vector optimization reduced size for enhanced packaging efficiency, and improved potency over original academic construct
- Initiation of IND-enabling studies planned for 2024

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