# Effects of LM11A-31 on longitudinal cerebrospinal fluid biomarkers of pathophysiology in Alzheimer's disease

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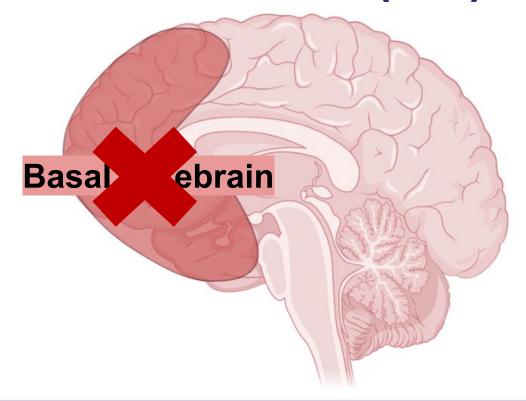




# Introduction

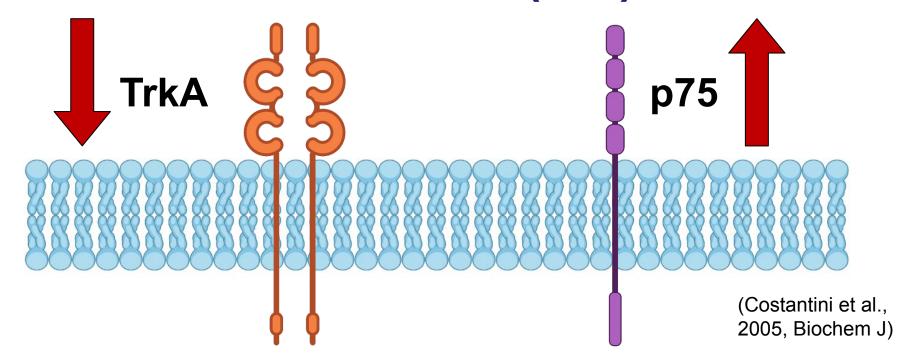






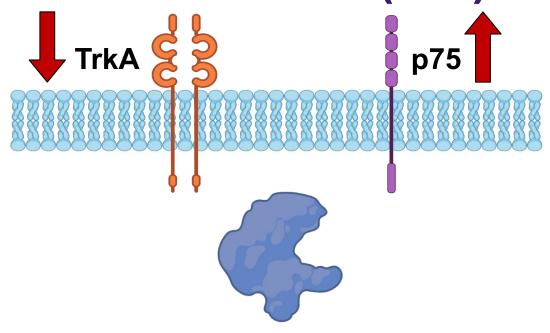










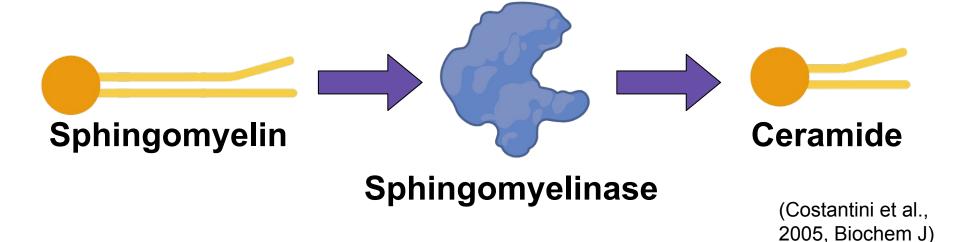


**Sphingomyelinase** 

(Costantini et al., 2005, Biochem J)











**β** secretase





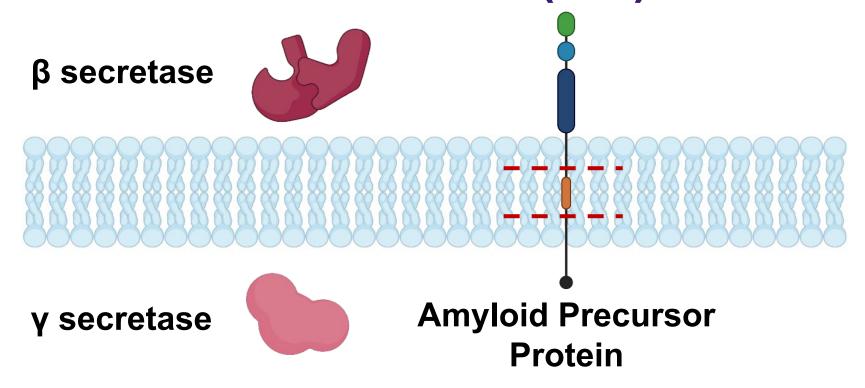
γ secretase



(Takasugi et al., 2015, Biochem Biophys Res Commun)

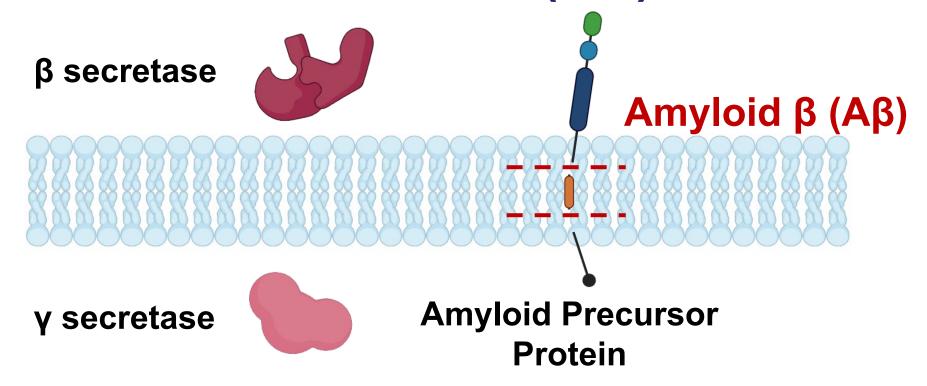






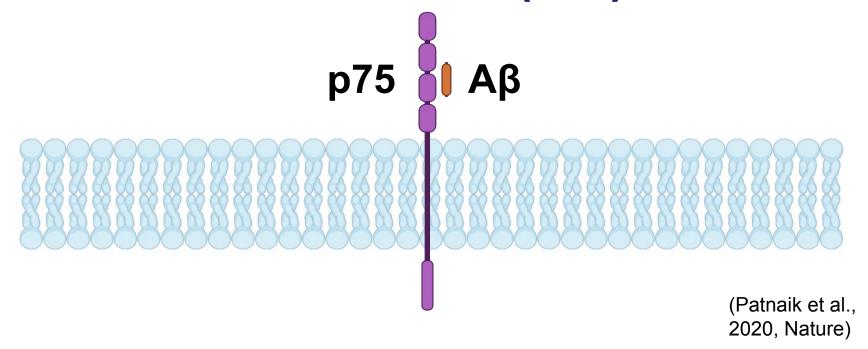






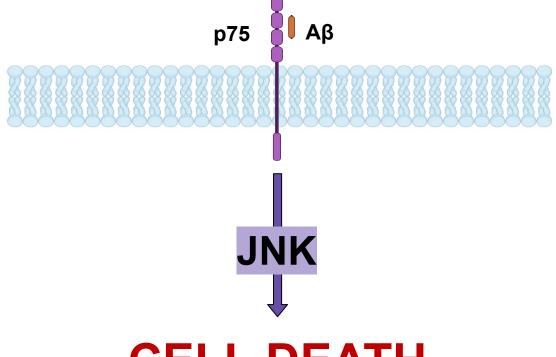










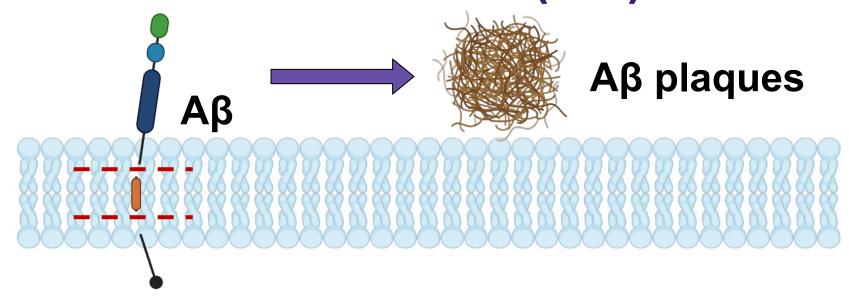


(Patnaik et al., 2020, Nature)





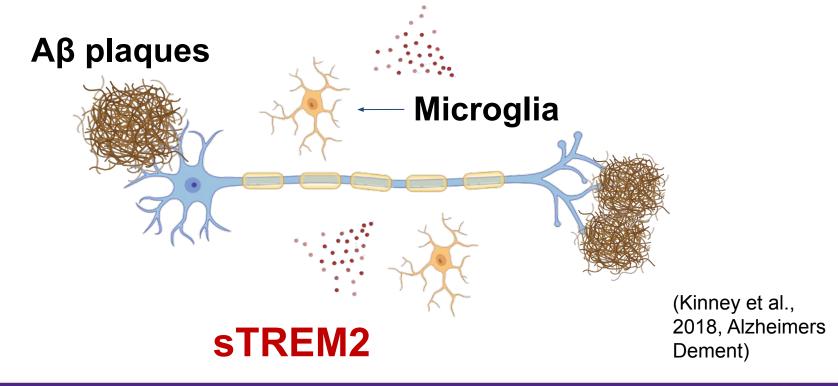




Amyloid Precursor Protein

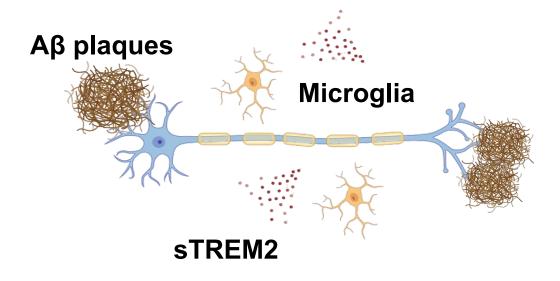










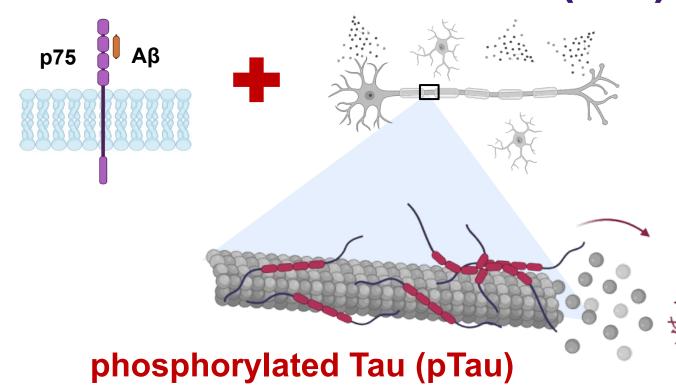


**CELL DEATH** 

(Kinney et al., 2018, Alzheimers Dement)





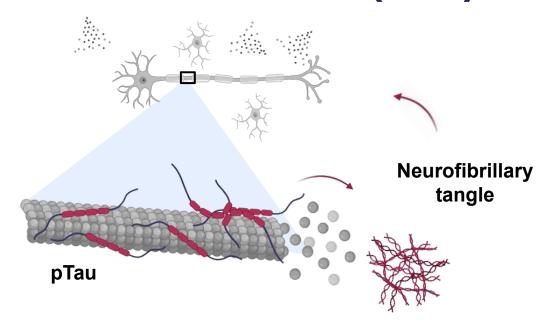


(Shen et al., 2019, Neurobio dis)

Neurofibrillary tangle



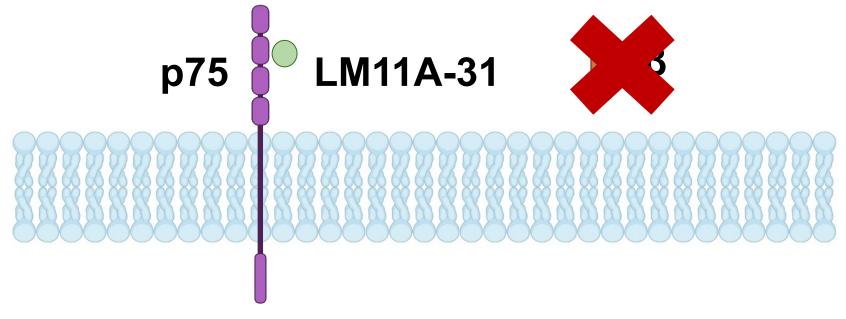




**CELL DEATH** 







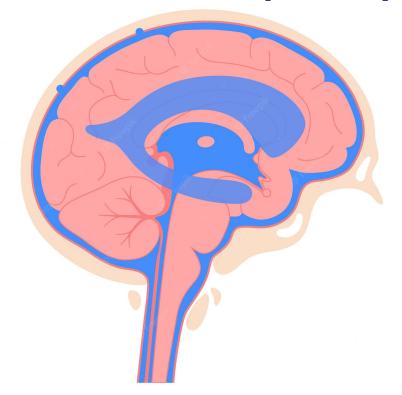
**CELL SURVIVAL** 

(Simmons et al., 2014, PLoS One)





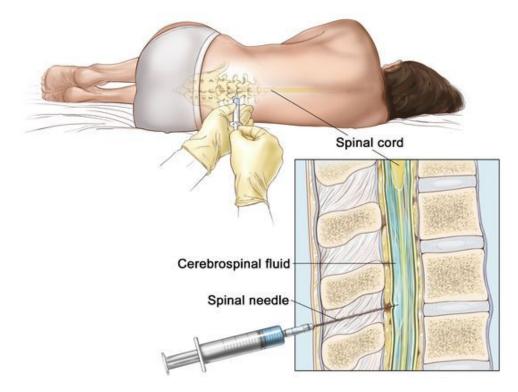
# Cerebrospinal fluid (CSF)







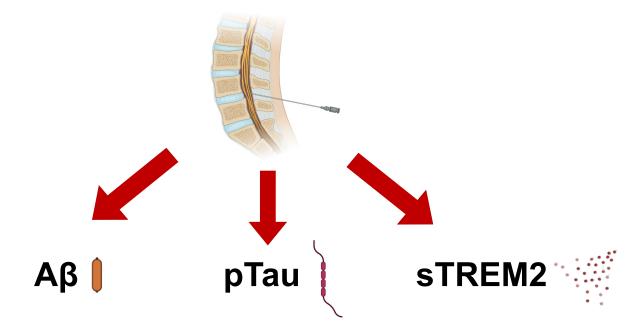
# Cerebrospinal fluid (CSF)







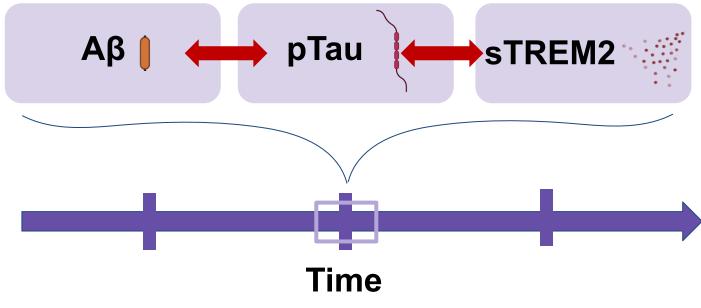
# Cerebrospinal fluid (CSF)



#### TREATMENT EFFECTIVENESS



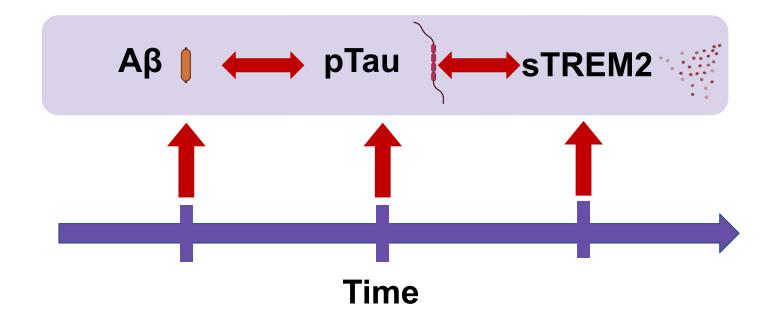




**UNIVARIATE & CROSS-SECTIONAL** 

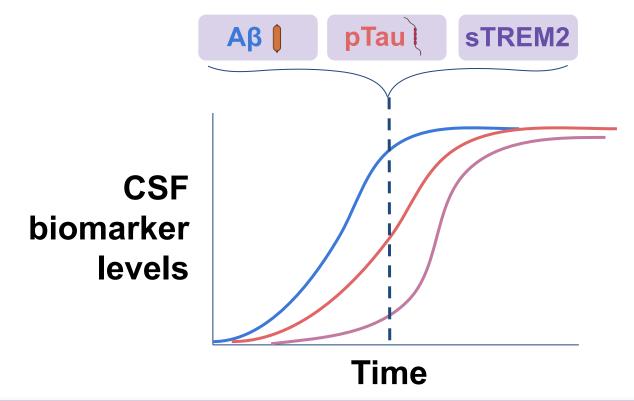








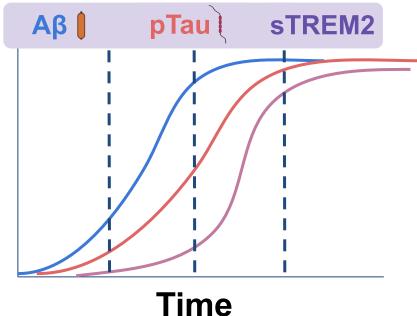












**MULTIVARIATE & LONGITUDINAL** 





# **Hypothesis**

Longitudinal multivariate analysis of CSF biomarkers will detect distinct phenotypes of biomarker interactions and trajectories associated with the treatment <u>responder</u> group in comparison to the treatment <u>non-responder</u> group.

LM11A-31



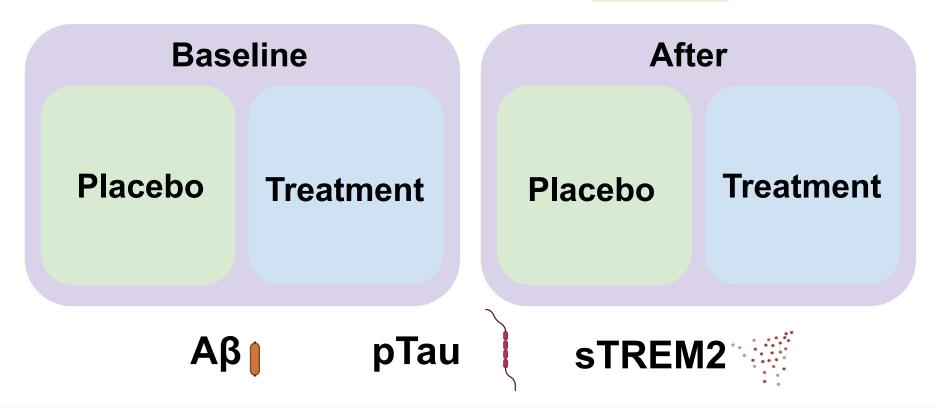


# Methods





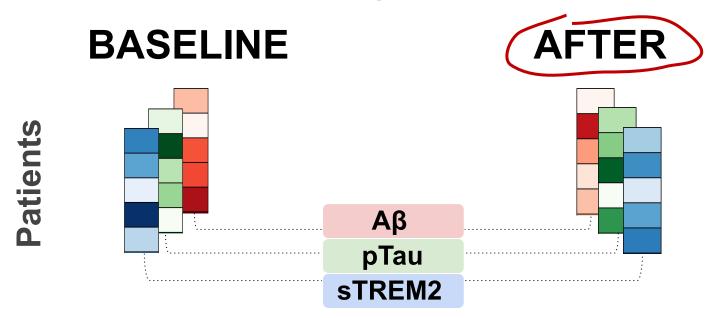
#### Clinical Trial Dataset n = 150







**Step 1: Original Data** 







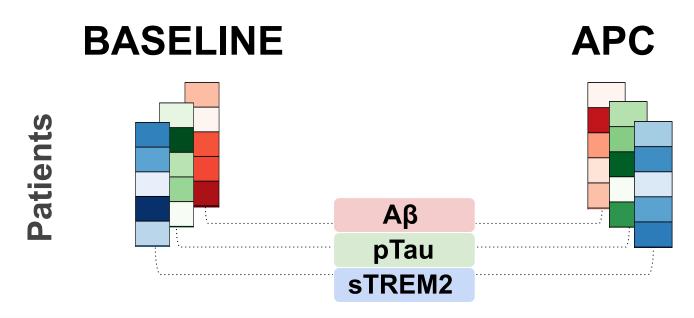
**Step 1: Original Data** 

$$APC = \left[ \left[ \frac{\text{Time 2}}{\text{Time 1}} \right]^{\frac{365}{\text{Intervals}}} - 1 \right] \times 100$$





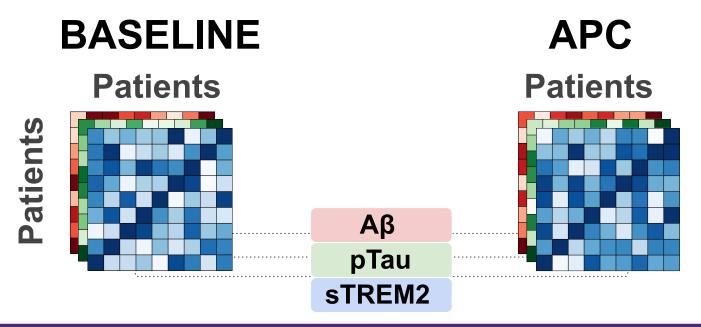
**Step 1: Original Data** 







**Step 2: Patient Similarity Matrices** 



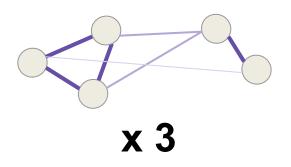


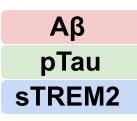


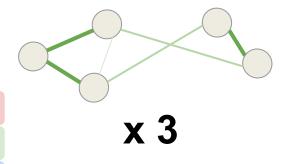
**Step 3: Patient Similarity Networks** 

**BASELINE** 

**APC** 



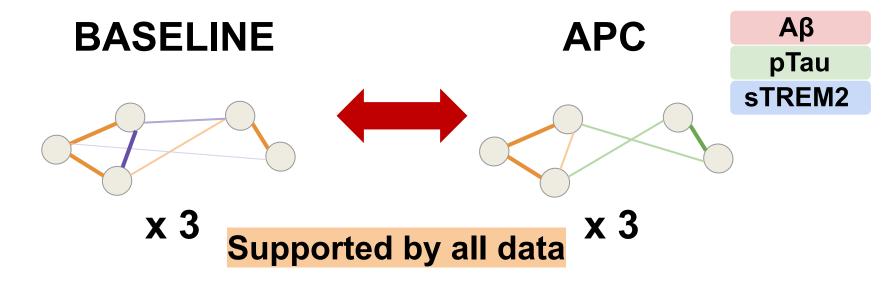








**Step 4: Patient Network Fusion** 







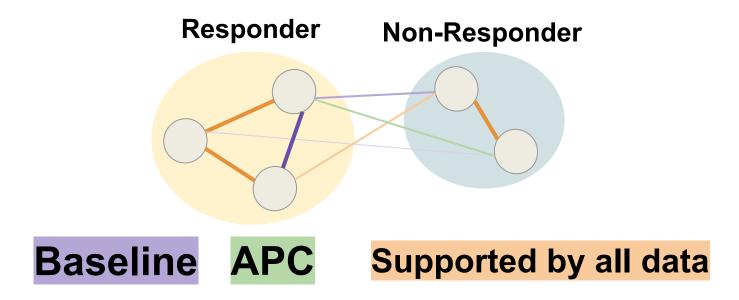
**Step 5: Fused Patient Similarity Network** 

Responder Non-Responder **Baseline** Supported by all data





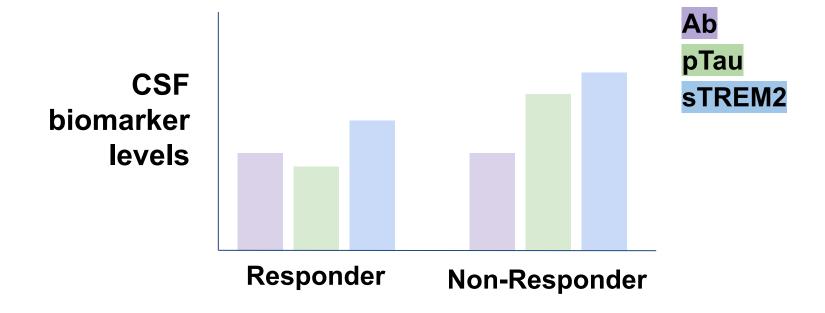
**Step 6: Spectral Clustering** 







# **Post-Hoc Analysis**







# Anticipated Results





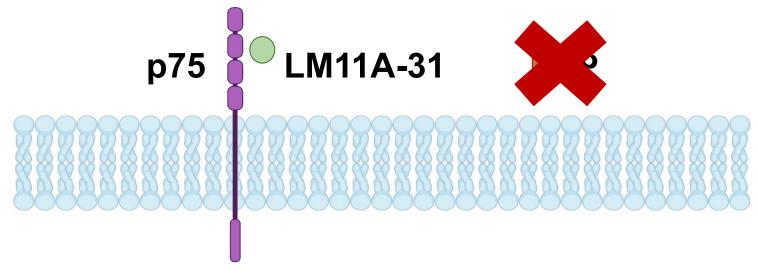
#### Clustering

(Blennow, 2005, Front. Neurosci) Responder Non-Responder **Before After** Supported by all data





#### Αβ

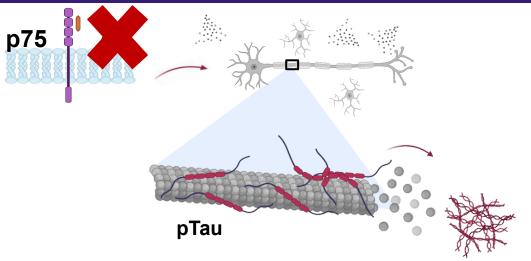


No significant difference between treatment response groups





#### pTau



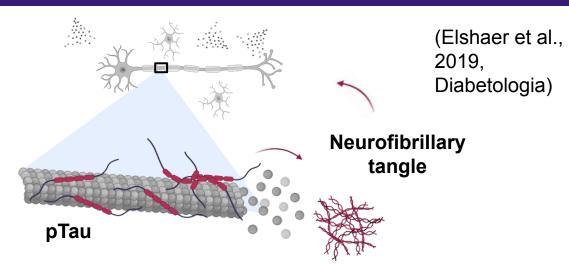
(Manucat-Tan et al., 2019, Aging)

# Significantly decreased in response group





#### sTREM2



# Significantly decreased in response group



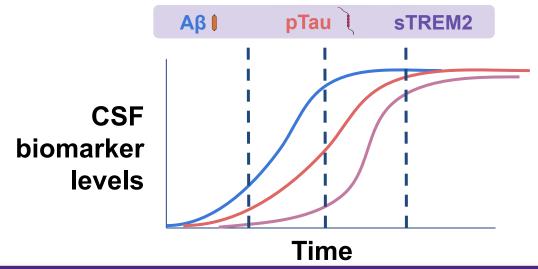






1

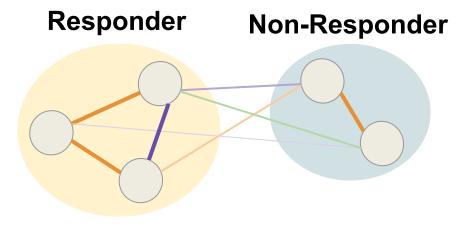
#### Longitudinal & Multivariate







2 Novel Phenotypes







3

#### **Advance Precision Medicine**

