

# Gene Therapy in Alzheimer's Disease: Understanding the Role of Apolipoprotein E in Late Onset Alzheimer's Disease

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## Background

### What is Alzheimer's Disease (AD)?

- A progressive neurodegenerative disease that causes memory loss and cognitive decline
- Late onset Alzheimer's disease (LOAD) is the most common form (affects people ages 65 and older)
- Pathological hallmarks of AD are amyloid plaques and neurofibrillary tangles, associated with amyloid-β protein clustering and tau protein misfolding respectively

### Why Study It?

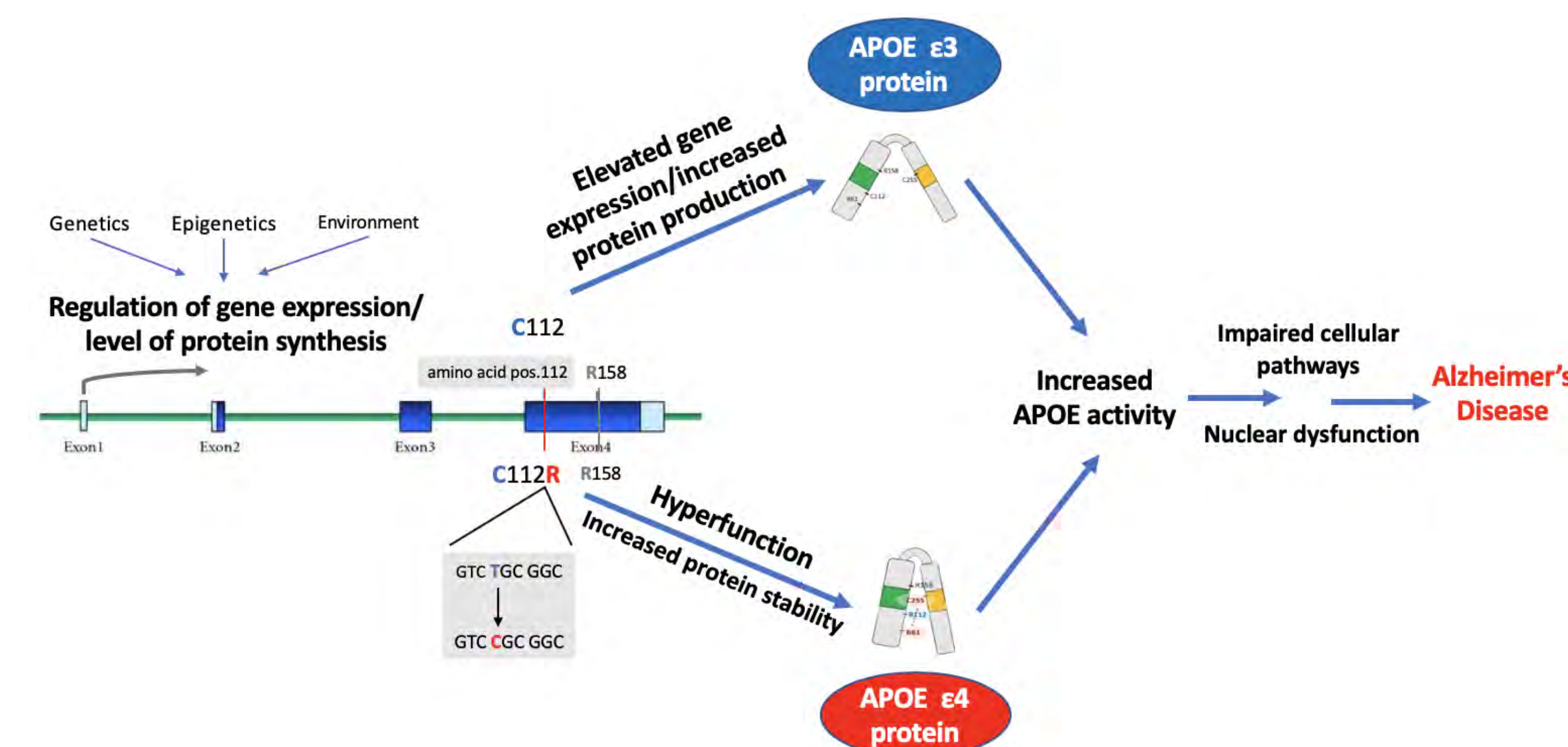
- AD affects nearly 6 million people in the U.S. and is projected to affect 14 million by 2050
- Causative factors and the mechanism behind its onset and progression remain unknown; no cure currently exists

### The APOE gene can modify your risk of developing Alzheimer's

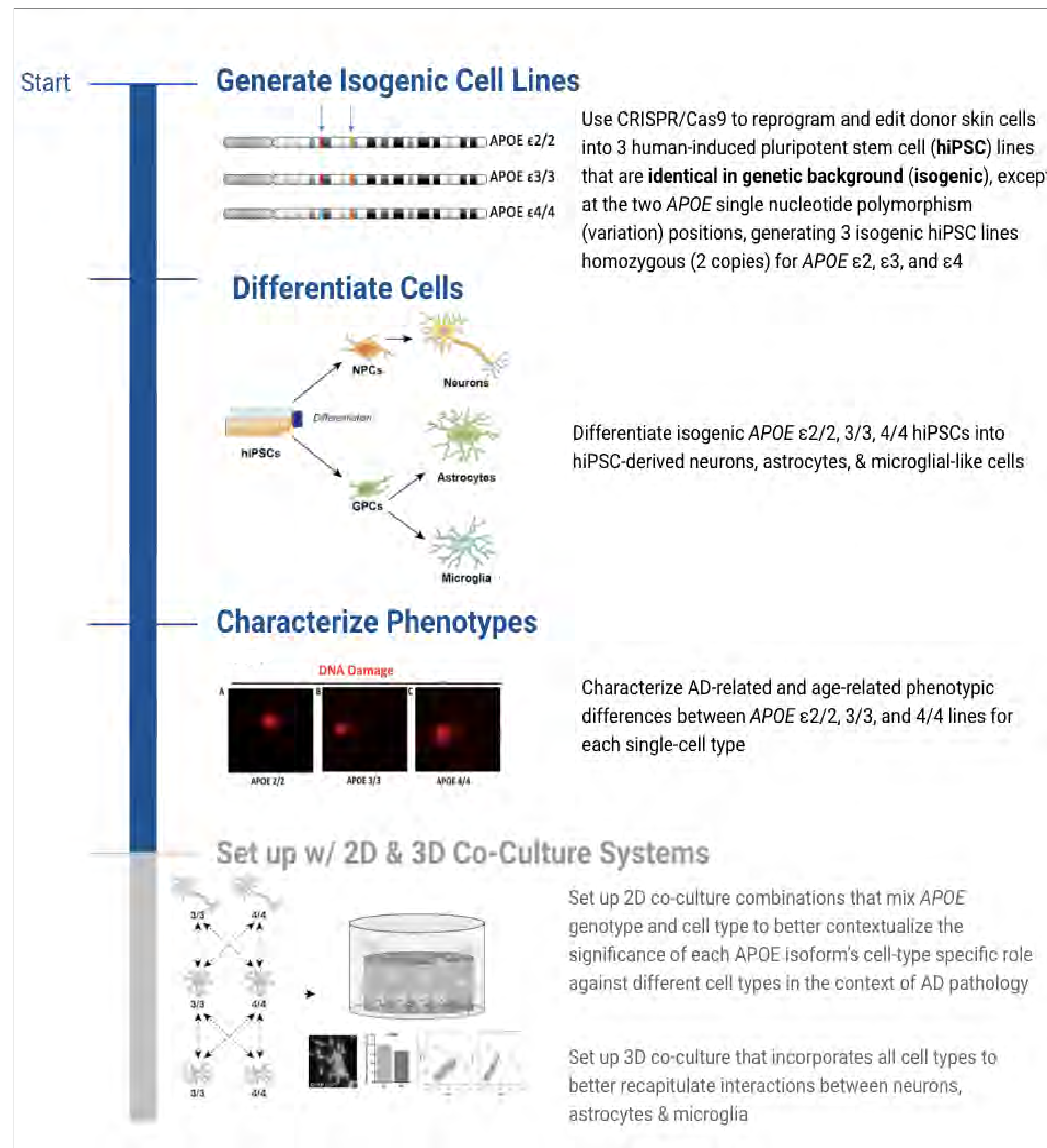
- APOE ε4 has been identified as the strongest and most reproducible genetic risk factor for LOAD
- APOE has 3 protein variants (isoforms) that vary by a single nucleotide in two possible positions in their DNA sequence, results in the coding of different amino acids in the protein

Isoform	Frequency	LOAD Risk
ε4 (R112, R158)	Less common (~14)	Higher (risk) 1 copy = 3-4 x 2 copies = 8-12 x
ε3 (C112, R158)	Most common (~79)	Normal (neutral)
ε2 (C112, C158)	Rare (~4)	Lower (resilience)

## Our Hypothesis



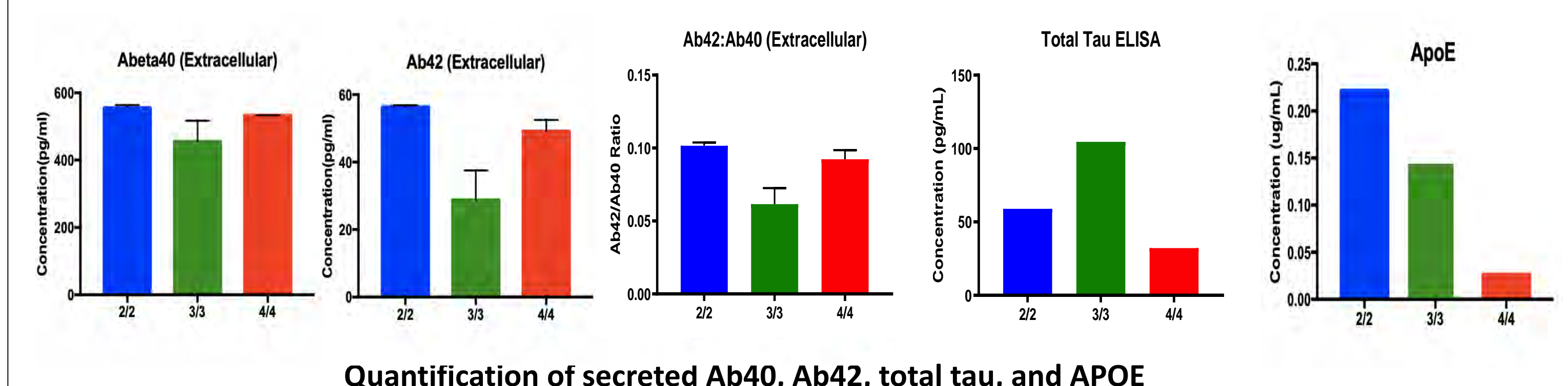
## Methods



## Results

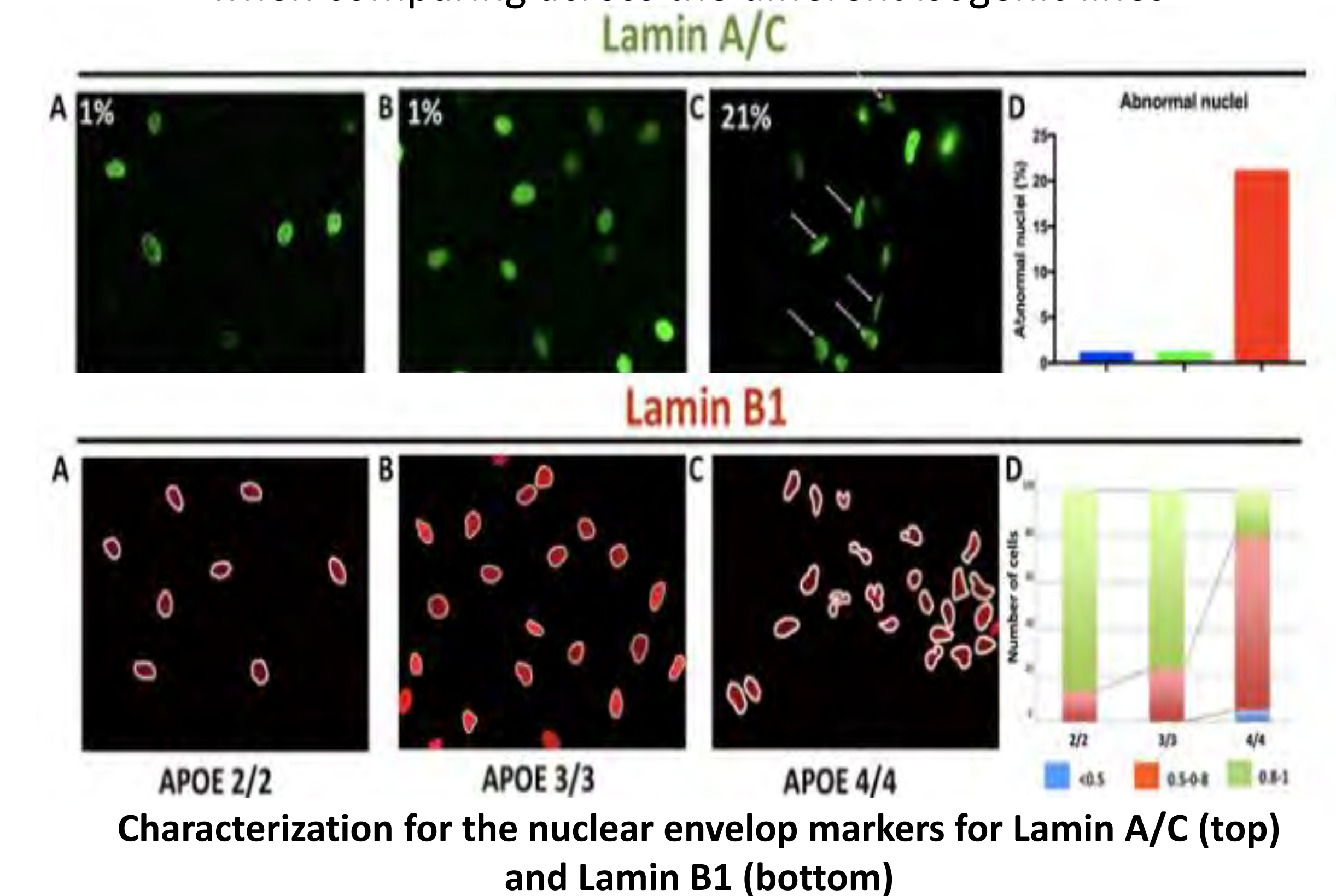
### AD-Related Phenotypes

APOE ε4 hiPSC-derived neurons show increased levels of Ab40, Ab42, and lower levels of total tau and APOE

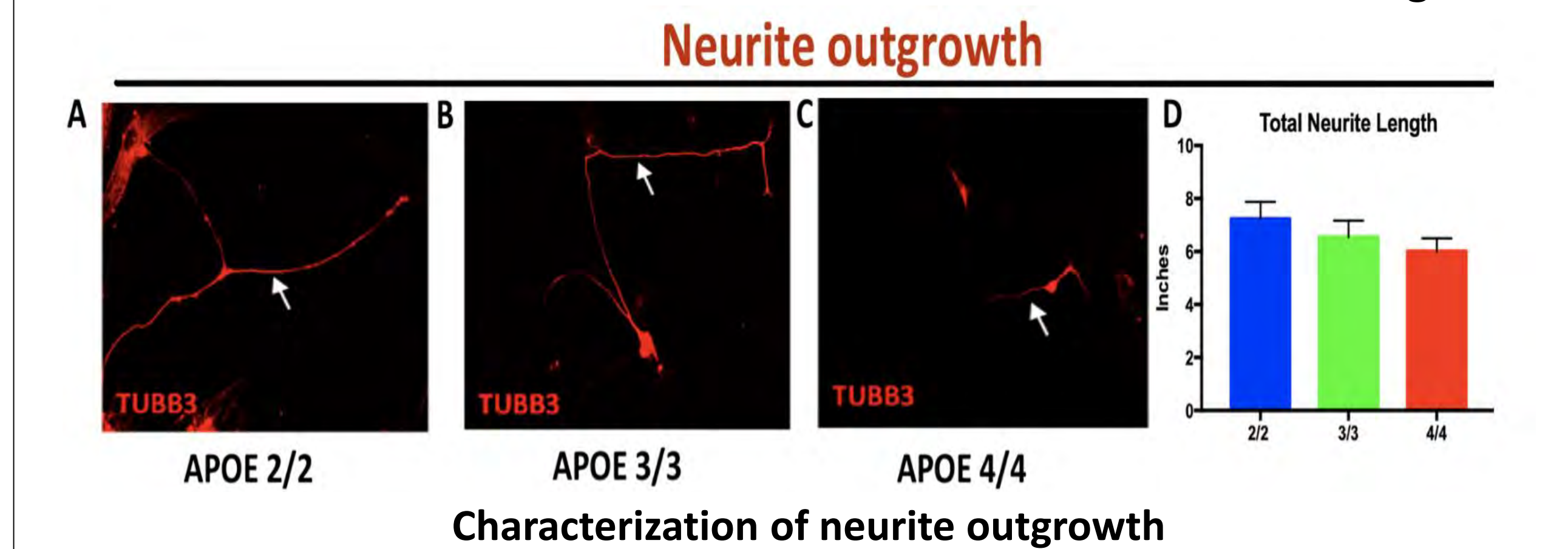


### Age-Related Phenotypes

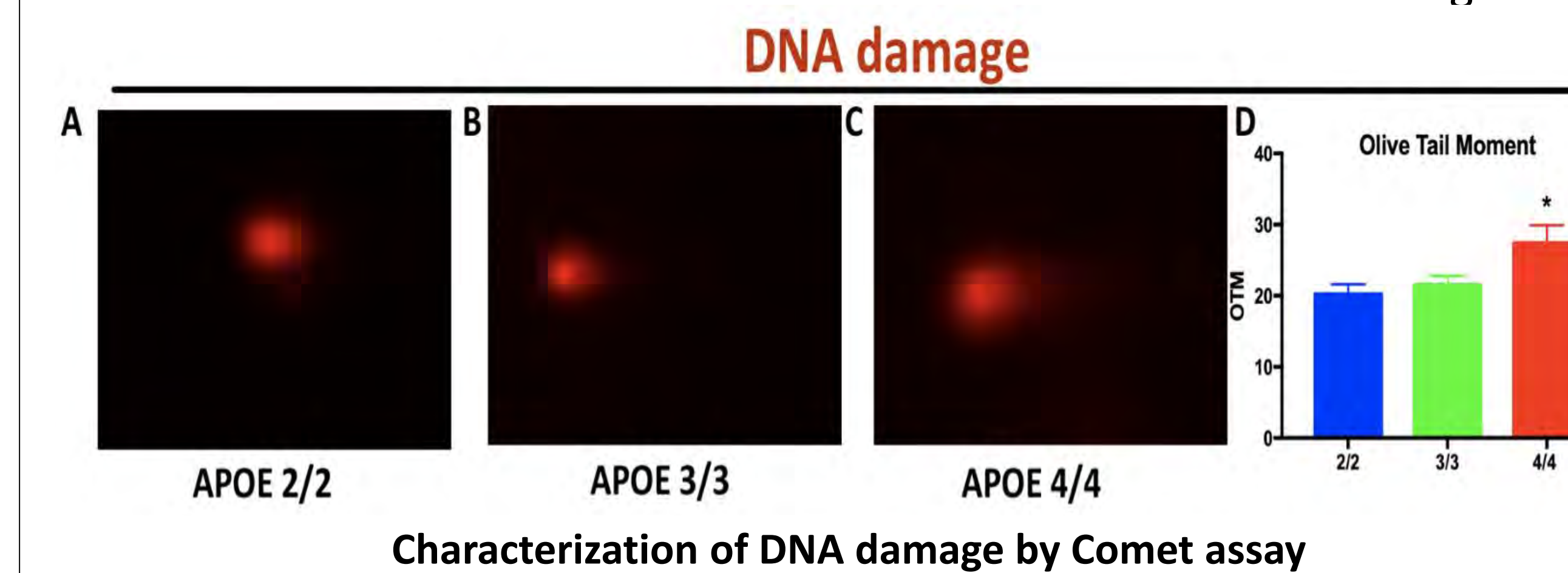
The hiPSC-derived neurons showed differences in nuclear envelope markers when comparing across the different isogenic lines



APOE ε4 hiPSC-derived neurons show decreased neurite outgrowth



APOE ε4 hiPSC-derived neurons show increased DNA damage



## Conclusion

Our data suggests a role of APOE in the maintenance of the nuclear envelope integrity that may lead to alterations in the cell homeostasis and cause neuronal loss.

We conclude our model provides a strong & malleable system for evaluating novel therapeutic gene therapy approaches in the APOE gene.