

Conserved astroglial toxicity promotes synaptic degeneration in frontotemporal dementia with *GRN* mutations

Elise Marsan, Dmitry Velmeshev, Arren Ramsey, Ravi K. Patel, Jiasheng Zhang, Mark Koontz, Madeline G. Andrews, Martina de Majo, Jessica Blumenfeld, Alissa N. Li, Salvatore Spina, Lea T. Grinberg, William W. Seeley, Bruce L. Miller, Erik M. Ullian, Matthew F. Krummel, Arnold R. Kriegstein, Eric J. Huang

Mutations in the human *Progranulin (GRN)* gene are a leading cause of frontotemporal lobar degeneration (FTLD). While previous studies implicate aberrant microglial activation as a disease-driving factor in neurodegeneration in the thalamocortical circuit in *Grn*^{-/-} mice, the exact mechanism for neurodegeneration in FTLD-*GRN* remains unclear. By performing comparative single-cell transcriptomics in the thalamus and frontal cortex of *Grn*^{-/-} mice and patients with FTLD-*GRN*, we have uncovered a highly conserved astroglial pathology characterized by upregulation of gap junction protein GJA1, water channel AQP4, and lipid-binding protein APOE, and downregulation of glutamate transporter SLC1A2, that promotes profound synaptic degeneration across the two species. This astroglial toxicity can be recapitulated in mouse astrocyte-neuron cocultures and by transplanting human induced pluripotent stem cell-derived astrocytes to cortical organoids, where Progranulin-deficient astrocytes promote synaptic degeneration, neuronal stress, and TDP-43 proteinopathy. Together, these results reveal previously unappreciated astroglial pathology as a key mechanism in neurodegeneration in FTLD-*GRN*.

Conflicts of interest

A.R.K. is a founding member of Neurona Therapeutics. All other authors declare no conflicts of interest.