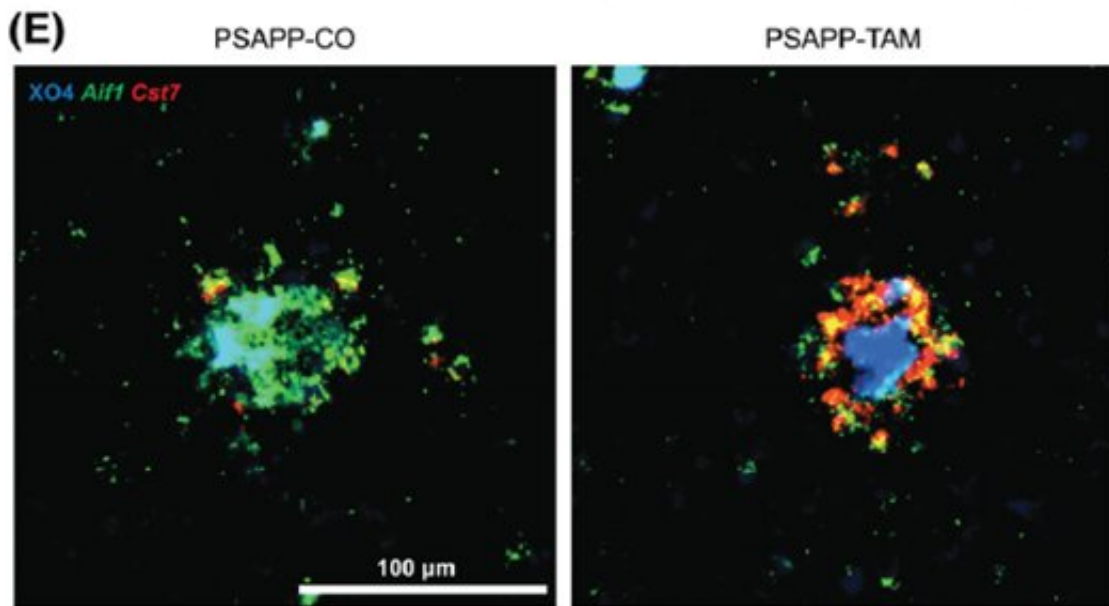


Researchers identify the role of an Alzheimer's disease risk gene in the brain

November 30 2022



Cst7 (yellow) is a Plaque Induced Gene (PIG) and its expression is specific to plaques in the INPP5D knockdown mouse (right), vs an AD mouse with normal levels of INPP5D (left). Credit: Emilie Castranio, PhD, postdoctoral fellow at the Icahn School of Medicine at Mount Sinai and co-first author of the paper.

A new study links a gene concentrated in the brain's cleanup cells, known as microglia, to the inflammation that has increasingly emerged as a key mechanism contributing to Alzheimer's disease. The findings may offer a new potential target for therapies for the intractable

condition.

The gene, known as inositol polyphosphate-5-phosphatase D (*INPP5D*), is the subject of a collaborative study conducted by researchers from the Icahn School of Medicine at Mount Sinai and the Grossman School of Medicine at NYU Langone Health that appears in the November 30 issue of *Alzheimer's and Dementia: The Journal of the Alzheimer's Association*.

Microglia are [immune cells](#) in the brain that act as scavengers to remove dying cells and amyloid plaques that are associated with the dementia of Alzheimer's [disease](#). Human genetics studies initially linked *INPP5D* to the risk for Alzheimer's disease. Other studies revealed elevated levels of *INPP5D* in the postmortem brain tissue of Alzheimer's disease patients, but the specific role(s) that the gene plays in both early or late disease and the mechanism contributing to these altered functions remains unknown.

Because *INPP5D* in the brain is concentrated in microglia, co-senior author Michelle E. Ehrlich, M.D., Professor of Neurology, Pediatrics, and Genetics and Genomic Sciences at Icahn Mount Sinai, used mice genetically engineered to "knock down" (turn off) the mouse *INPP5D* gene in their microglia at the onset of pathology. This process allowed them to better see the specific impact of the missing gene on brain tissue. They then measured [plaque buildup](#) and microglial behavior approximately three months later. Since *INPP5D* was known to be elevated in the brains of Alzheimer's patients, the scientists expected that the mice with that gene inactivated would be protected from the amyloid plaques that are hallmarks of Alzheimer's disease pathology.

"When I looked through the microscope, I was quite surprised to see that the mice lacking *INPP5D* in their microglia had more plaques than mice with normal microglia," said Emilie Castranio, Ph.D., a postdoctoral

fellow in Dr. Ehrlich's lab and co-first author on the new paper.

"Microglia frequently sit on the edges of the plaques but when *INPP5D* was knocked down, the plaques were completely covered with them."

"We are encountering unexpected results more and more with modulation of inflammation [genes](#) in Alzheimer's," said Dr. Ehrlich. "At this point in our understanding, we still do not know which of these genes to target for therapeutic intervention in humans, or whether to turn them up or turn them off depending on disease stage. Because these experiments are not possible in living humans, we rely on mouse models to show us the way. We also use these mice to help us predict whether a particular gene is more related to disease onset or disease progression, with the caveat that mouse and human microglia differ in important ways. Despite these differences, the plaque-associated gene signature we identified overlaps with human Alzheimer's disease gene networks."

When it became clear that the *INPP5D* knockdown moved microglia around the brain in unexpected ways, Dr. Ehrlich recognized that detailed spatial and quantitative gene expression information was required. Spatial transcriptomics is a molecular profiling method that allows scientists to measure all the gene expression in a tissue sample and map where the expression is occurring. Drs. Ehrlich and Castranio turned to Shane Liddelow, Ph.D., Assistant Professor of Neuroscience, Physiology, and Ophthalmology at NYU Langone, and co-senior author of the new study, who is a world leader in this approach.

The spatial transcriptomics findings emphasized the range of gene expression changes that [microglia](#) can display. Microglia near [amyloid plaques](#) are known to express genes designated as plaque-induced genes (PIGs). The *INPP5D* knockdown mice replicated the increases in PIGs that had been described in previous research, but the high quality of both the technical aspects and analysis of the spatial transcriptomics allowed for the identification of additional PIGs. The newly identified PIG with

the greatest increase in expression in these mice was *CST7*, a gene encoding the protein cystatin F that is known to be impacted in Alzheimer's and associated with prion diseases, a family of rare, progressive neurodegenerative disorders that affect both humans and animals. These findings suggest that both *INPP5D* and cystatin F should be considered as targets for development of novel interventions aimed at mitigating inflammation in the Alzheimer's brain.

More information: EL Castranio et al, *INPP5D* limits plaque formation and glial reactivity in the APP/PS1 mouse model of Alzheimer's disease, *bioRxiv* (2022). DOI: [10.1101/2022.04.29.490076](https://doi.org/10.1101/2022.04.29.490076)

Emilie L. Castranio et al, Microglial *INPP5D* limits plaque formation and glial reactivity in the PSAPP mouse model of Alzheimer's disease, *Alzheimer's & Dementia* (2022). DOI: [10.1002/alz.12821](https://doi.org/10.1002/alz.12821) , [dx.doi.org/10.1002/alz.12821](https://doi.org/10.1002/alz.12821)

Provided by The Mount Sinai Hospital

Citation: Researchers identify the role of an Alzheimer's disease risk gene in the brain (2022, November 30) retrieved 21 July 2024 from <https://medicalxpress.com/news/2022-11-role-alzheimer-disease-gene-brain.html>

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