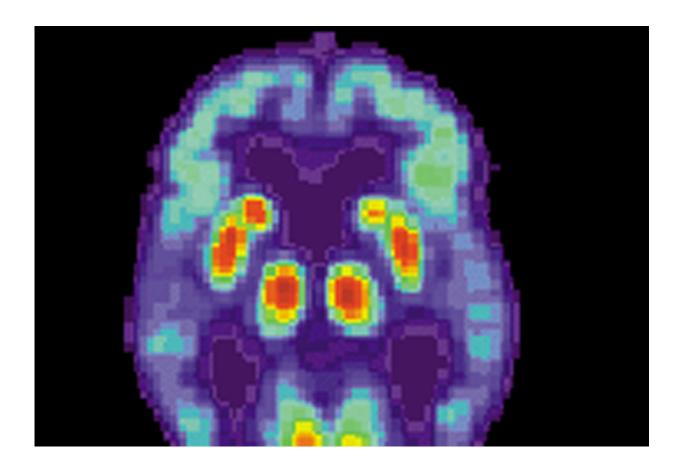


Researchers develop mouse model of human gene involved in Alzheimer's disease

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PET scan of a human brain with Alzheimer's disease. Credit: public domain

In research that helps scientists better understand and explore treatments for diseases like Alzheimer's, scientists have developed a line of mice in which the mouse version of the Alzheimer's-associated MAPT gene has



been fully replaced by the human version of the gene. In this new animal model, known as a full gene-replacement model, the MAPT gene will function the same way it does in humans, allowing researchers to more accurately develop and evaluate genetic therapies. The research was presented at the American Society of Human Genetics 2019 Annual Meeting in Houston, Texas.

Researchers have long studied <u>human genes</u> in mice and other animal models, usually by finding and manipulating the animal's version of the human gene being studied, explained Michael Koob, Ph.D., Associate Professor at the University of Minnesota, who presented the work.

"However, mice have different <u>genes</u> than people, and even if the gene's function is the same, its sequence is different," Dr. Koob said. For this reason, <u>animal model</u> work typically involves a great deal of trial and error, and it requires researchers to make assumptions about why and how a genetic change leads to the observed changes. In addition, drawing conclusions about the role of the human version of the gene in humans—and building on this knowledge by developing therapies—is difficult and prone to error, and the findings do not always translate.

In developing their approach to full gene-replacement models, Dr. Koob and colleagues decided to focus on MAPT, a gene that is known to play an important role in Alzheimer's <u>disease</u> but whose involvement is not well understood. They overcame several <u>technical challenges</u> to do so, such as identifying the boundaries of the gene's protein-coding segments and regulatory regions, inserting the human version into the mouse genome with precision, and working with the gene's relatively large size.

By replacing the mouse's version of MAPT with the human version, Dr. Koob said, "we will be able to better explain how MAPT would function in humans, learn how it contributes to the symptoms and progression of Alzheimer's disease, and do some early testing of the potential efficacy



of genetic therapies."

Longer term, the researchers plan to use the same approach to develop full gene-replacement models for other genes involved in Alzheimer's, eventually building up to a mouse model that expresses several genes involved in the disease.

"A mouse model that expresses multiple genes will help us understand the interactions among genes that may contribute to disease," Dr. Koob said.

They have also developed similar models for genes involved in other diseases, such as the neurological disorder spinocerebellar ataxia type 1 and the eye disease Fuchs endothelial corneal dystrophy.

More information: M Koob et al. (2019 Oct 16). Abstract: Moving human genetics into the mouse: Full human gene-replacement models. Presented at the American Society of Human Genetics 2019 Annual Meeting. Houston, Texas.

Provided by American Society of Human Genetics

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