

# Adult-onset neurodegeneration has roots in early development

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The disease mechanism for adult-onset progressive degenerative diseases begins much earlier than previously thought, according to a Northwestern Medicine study published in the *Journal of Clinical Investigation*.

Using a mouse model of spinocerebellar ataxia type 1 (SCA1) genetically engineered to precisely mirror the human disease, a team of investigators showed that there is an altered neural circuitry in the cerebellum that sets the stage for later disease vulnerability.

These findings are important since they raise the possibility that other adult-onset neurodegenerative diseases have their roots in early developmental defects, with implications for pathogenesis and treatment, said Puneet Opal, MD, PhD, professor in Ken & Ruth Davee Department of Neurology at Northwestern University Feinberg School of Medicine, and senior author on the study.

"This is the first discovery of alterations in an adult-onset spinocerebellar disorder that stem from such early developmental processes," said Opal, also a professor of Cell and Molecular Biology.

"This may well be generalizable to a whole host of other diseases, including Alzheimer's disease, Huntington's disease, Parkinson's disease and amyotrophic lateral sclerosis."

SCA1 is caused by a genetic defect in a protein involved in regulating gene expression called ATXN1. While ATXN1 is expressed throughout

the brain, when mutated it predominantly affects the cerebellum, leading to a loss of coordination and an abnormal gait in patients.

Previous studies have largely focused on neurons in the disease, but Opal and his colleagues theorized the cerebellar stem cell population might also behave differently since some of the gene expression changes in the brain in the disease occur relatively early.

Using a genetically engineered mouse model of the [disease](#), the scientists found that mutant ATXN1 had surprising effects on postnatal cerebellar stem cells.

"We were amazed to find that they multiplied excessively and tended to differentiate into [inhibitory neurons](#) called basket cells," said Opal. "We knew that cerebellar stem cells generate inhibitory neurons, but in this case the number of inhibitory neurons was so much more than normal that they generated an enhanced inhibitory effect on Purkinje neurons, the chief output neurons of the cerebellum. This results in a very different cerebellar network."

These changes began in the early postnatal weeks of life and may explain why the cerebellar neuronal network is particularly vulnerable to degeneration in patients with SCA1, according to Opal.

In addition, as more stem cells became inhibitory [neurons](#), fewer stem cells became brain [cells](#) called astrocytes, possibly further disrupting normal cerebellar function.

"This network dysfunction could be a constant stress, and that constant stress makes the neural network deteriorate over time," Opal said.

While it's still unclear exactly how the mutant protein is causing these changes, according to the study, the findings underscore the need to

study a relatively underappreciated phenomenon: that pathogenic processes in neurodegenerative diseases begin very early in life— well before signs or symptoms are noticed.

"Other diseases may have similar developmental defects, but we haven't looked for them," Opal said. "We found them in SCA1, but it could be true for the other diseases as well."

**More information:** Chandrakanth Reddy Edamakanti et al, Mutant ataxin1 disrupts cerebellar development in spinocerebellar ataxia type 1, *Journal of Clinical Investigation* (2018). [DOI: 10.1172/JCI96765](https://doi.org/10.1172/JCI96765)

Provided by Northwestern University

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