

New brain development disorder identified by scientists

May 22 2018



Credit: CC0 Public Domain

Researchers have identified a new inherited neurodevelopmental disease that causes slow growth, seizures and learning difficulties in humans.

Writing in the journal *eLife*, the team reveals that this disease is caused by a recessive mutation in *CAMK2A*—a gene that is well known for its

role in regulating learning and memory in animals. The findings suggest that dysfunctional CAMK2 genes may contribute to other neurological disorders, such as epilepsy and autism, opening up potential new avenues for treating these conditions.

"A significant number of children are born with growth delays, neurological defects and intellectual disabilities every year across the world," explains senior author Bruno Reversade, Research Director at the Institute of Medical Biology and Institute of Molecular and Cell Biology, A*STAR, Singapore, who supervised the study. "While specific genetic [mutations](#) have been identified for some patients, the cause remains unknown in many cases. Identifying novel mutations would not only advance our understanding of neurological diseases in general, but would also help clinicians diagnose children with similar symptoms and/or carry out genetic testing for expecting parents."

The team's research began when they identified a pair of siblings who demonstrated neurodevelopmental delay with frequent, unexplained seizures and convulsions. While the structure of their bodies developed normally, they did not gain the ability to walk or speak. "We believed that the children had novel mutations in CAMK2A, and we wanted to see if this were true," says Reversade.

The fully functional CAMK2A protein consists of multiple subunits. Using a genomic technique called exome sequencing, the team discovered a single coding error affecting a key residue in the CAMK2A gene that prevents its subunits from assembling correctly.

Moving their studies into the roundworm *Caenorhabditis elegans*, the scientists saw that this mutation disrupts the ability of CAMK2A to ensure proper neuronal communication and normal motor function. This suggests that the mutation is indeed the cause of the neurodevelopmental defects seen in the siblings.

To the best of the team's knowledge, this new disorder represents the first human disease caused by inherited mutations on both copies of the CAMK2A gene. In addition, another report* published recently identified single-copy mutations on both CAMK2A and CAMK2B that caused [intellectual disabilities](#) as soon as the mutations occurred. "We would like to bring these findings to the attention of those working in the area of paediatric genetics, such as clinicians and genetic counsellors, as there are likely more undiagnosed children with similar symptoms who have mutations in their CAMK2A gene," explains co-first author Franklin Zhong, Research Scientist in Reversade's lab at A*STAR.

"Neuroscientists working to understand childhood brain development, neuronal function and memory formation also need to consider this new disease, since CAMK2A is associated with these processes. In future, it would be interesting to test whether restoring CAMK2A activity can bring therapeutic benefit to patients with this condition, as well as those with related neurological disorders."

More information: Poh Hui Chia et al, A homozygous loss-of-function CAMK2A mutation causes growth delay, frequent seizures and severe intellectual disability, *eLife* (2018). [DOI: 10.7554/eLife.32451](https://doi.org/10.7554/eLife.32451)

*Kury, S., van Woerden, G.M., Besnard, T., Proietti Onori, M., Latypova, X., Towne, M.C., Cho, M.T., Prescott, T.E., Ploeg, M.A., Sanders, S., et al. (2017). De Novo Mutations in Protein Kinase Genes CAMK2A and CAMK2B Cause Intellectual Disability. *The American Journal of Human Genetics* 101, 768-788.

Provided by eLife

Citation: New brain development disorder identified by scientists (2018, May 22) retrieved 4

May 2024 from <https://medicalxpress.com/news/2018-05-brain-disorder-scientists.html>

This document is subject to copyright. Apart from any fair dealing for the purpose of private study or research, no part may be reproduced without the written permission. The content is provided for information purposes only.