CHRD 2024: Abstract Submission Form

Presenter Name Katelyn Panchoo

Role in the project Design Perform Experiments Analyze Data Write Abstract **Presenter Status** Undergraduate Students

Research Category Basic Science

Title

Investigating ELFN1 Variants in Relation to Neurodevelopmental Disorders

Background

Extracellular leucine-rich repeat and fibronectin type 3 domain containing 1, ELFN1, is an important synaptic adhesion molecule (SAM) that aids in neuronal communication. Previous studies have found that ELFN1 modulates the activity of a particular class of receptors known as group III metabotropic glutamate receptors (mGluRs), broadening our understanding of SAM function as more than structural components. Mouse studies have also found that downregulation of mGluRs at the synapse can result in excess amounts of glutamate being released, leading to symptoms of neurodevelopmental disorders such as hyperactivity, compulsivity, and anxiety. Interestingly, these symptoms have been found in children with different mutations of ELFN1. My research investigates an intracellular (ELFN1- Δ CT) and extracellular mutation (Δ NT-ELFN1).

Objective

The purpose of my experiment is to examine whether the expression of these variants is altered, affecting their interaction with mGluRs and leading to these neurodevelopmental disorders.

Methods

My experiments were done through western blot analysis, which separates proteins based on molecular weight or size via gel electrophoresis. After transferring to a solid support matrix, the western blot was imaged and densitometric analysis was conducted to quantify expression levels.

Results

Through quantitative analysis, we found ELFN1- Δ CT expressed significantly lower than Δ NT-ELFN1 and was trending towards a significant difference from wild-type ELFN1.

Conclusion

Lowered expression of ELFN1 may indicate a reduced number of synaptic ELFN1/mGluR interactions, resulting in the excess release of glutamate and behaviours found in neurodevelopmental disorders. Gaining a better understanding of the mechanistic consequences could help us identify better therapeutic strategies for patients harboring these mutations.

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Yes

Authors

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