

Seminar

Cellular and Developmental basis of neurodevelopmental defects In Lowe syndrome

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The activity of signalling pathways is required for coordinated cellular and physiological processes leading to normal development of brain structure and function. Mutations in *OCRL*, a phosphatidylinositol 4,5 biphosphate [PIP₂] 5-phosphatase leads to the neurodevelopmental disorder, Lowe Syndrome (LS). However, the mechanism by which mutations in *OCRL* leads to the brain phenotypes of LS is not understood. We find that on differentiation of LS patient derived iPSC, developing neural cultures show reduced excitability along with enhanced P levels of Glial Fibrillary Acidic Protein. Multiomic single-nucleus RNA and ATAC seq analysis of neural stem cells generated from LS patient iPSC revealed an enhanced number of cells with a gliogenic cell state. RNA seq analysis also revealed increased levels of *DLK1*, a non-canonical Notch ligand in LS patient NSC associated increased levels of cleaved Notch protein and elevation of its transcriptional target *HES5*, indicating upregulated Notch signalling. Treatment of iPSC derived brain organoids with an inhibitor of PIP5K, the lipid kinase that synthesises PIP₂, was able to restore neuronal excitability and rescue Notch signalling defects in LS patient derived organoid cultures. Overall, our results demonstrate a role for PIP₂ dependent regulation of Notch signalling, cell fate specification and development of neuronal excitability regulated by *OCRL* activity.

Tuesday, Aug 26th 2025

16:00 Hrs (Tea / Coffee 15:45 Hrs)

Auditorium, TIFRH