

ELUCIDATING NEURONAL CELL POPULATIONS UNDERLYING CIC HAPLOINSUFFICIENCY SYNDROME

Mark A Durham¹, Carolyn J Adamski², Matthew C Hill³, Zachary A Kadow³, James F Martin⁴, Huda Y Zoghbi⁵

¹ Baylor College of Medicine, Department of Molecular & Human Genetics

² Baylor College of Medicine, Human Genetics, NRI

³ Baylor College of Medicine, Developmental Biology, n/a

⁴ Baylor College of Medicine, Molecular Physiology, n/a

⁵ Baylor College of Medicine, Human Genetics, n/a

Keywords: cortical-development, hyperactivity, single-cell-RNA-sequencing

Background: Capicua (CIC) is a transcriptional repressor that reduces target gene expression by recruiting histone deacetylase complexes. Our lab discovered that heterozygous de-novo loss-of-function mutations in CIC lead to CIC Haploinsufficiency Syndrome (CHS), a neurodevelopmental disorder characterized by intellectual disability, ADHD, autism and seizures. To understand how CIC loss leads to these phenotypes, we deleted *Cic* conditionally in different brain regions in mice. Loss of *Cic* from the developing neocortex and hippocampus leads to a robust ADHD-like hyperactivity phenotype and impaired learning and memory. Additionally, ablation of *Cic* from inhibitory neurons recapitulated seizures observed in patients. The overlap in phenotypes between mice and humans suggests we can utilize these models to identify the biological processes driving CHS. Although *Cic* is ubiquitously expressed in neurons, its loss leads to a post-mitotic reduction in the number of upper layer cortical neurons but does not affect lower layers. Thus, it is possible that a population of upper layer cortical neurons is implicated in the behavioral phenotypes in *Cic* mutant mice, but a direct link is not yet known. We seek to determine if there is a subpopulation of upper layer cortical neurons responsible for hyperactivity and elucidate the role of *Cic* in this cell population.

Materials/Methods: To identify changes in cellular heterogeneity upon loss of *Cic* in the developing cortex we performed single-cell RNA sequencing (scRNA-seq) on postnatal day 5 cortices of *Emx1-Cre* *Cic* KO and control mice. We used CUT&RUN to identify CIC binding sites to distinguish primary vs secondary transcriptional changes. After identifying a *Cic* dependent cortical cell population, we used conditional mouse genetics to evaluate whether deletion of *Cic* in this population was sufficient to drive hyperactivity.

Results: scRNA-seq showed a 60% reduction in layer IV cortical neurons compared to other layers. Integration of CUT&RUN and scRNA-seq suggested that layer IV cortical neurons are preferentially vulnerable to *Cic* loss, rather than *Cic* having a unique function in this population. Finally, removal of *Cic* from layer IV cortical neurons caused a slight increase in hyperactivity.

Conclusions: Layer IV cortical neurons are preferentially affected upon loss of *Cic* from the developing cortex and likely contribute to hyperactivity in *Cic* mutant mice. Further investigation of the vulnerability of Layer IV neurons will help explain the processes underlying CHS.

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