

POSTERS

## DECODING THES PATHOGENESIS: HOW TTC37 SHAPES GUT HOMEOSTASIS AND PERISTALSIS

A. Tesoriere<sup>1</sup>, F. Sernesi<sup>1</sup>, A. Piersanti<sup>1</sup>, M. Gasparotto<sup>3</sup>, L. Dalla Valle<sup>1</sup>, M. Cananzi<sup>2</sup>, F. Argenton<sup>1</sup>

<sup>1</sup>University of Padova, Italy; <sup>2</sup>University Hospital of Padova, Italy; <sup>3</sup>Central Institute of Mental Health, Mannheim, Germany

Trichohepatoenteric Syndrome (THES) is an ultra-rare, multisystem disorder caused by biallelic mutations in the *TTC37* or *SKIV2L* genes, which encode essential subunits of the SKI complex, a cofactor for the RNA exosome. The hallmark of THES is intractable diarrhea, often necessitating parenteral nutrition for survival. THES is a life-threatening condition, with mortality mainly attributed to intestinal failure and infections. Despite its clinical implications, no curative treatment exists, and management is limited to supportive care.

The mechanisms through which *TTC37* mutations give rise to such profound gastrointestinal dysfunction and high mortality have remained entirely unclear. The existing body of literature consists largely of clinical descriptions and case reports and provides no mechanistic explanation linking *TTC37* deficiency to THES enteropathy. This critical gap in knowledge has hindered the identification of therapeutic targets for a condition in urgent need of effective interven-

tions.

To elucidate the complete unknown pathophysiological basis of THES, we established a *ttc37* knockout zebrafish line, the first animal model of the disease reported to date. This model recapitulates major aspects of the human condition and exhibits marked abnormalities in gut architecture and impaired peristalsis, all consistent with the clinical phenotype of THES.

Leveraging the strengths of our disease model, we dissected the impact of the *ttc37* mutation on gut cells. We identified marked structural defects associated with increased inflammation and apoptosis, alongside a reduction in stemness markers.

In conclusion, we provide the first insight into the molecular basis of THES symptomatology, enabled by the generation of a disease model. Our data support a pathogenic framework in which structural defects lead to impaired epithelial integrity, chronic inflammation, and reduced regenerative capacity, ultimately driving THES enteropathy.