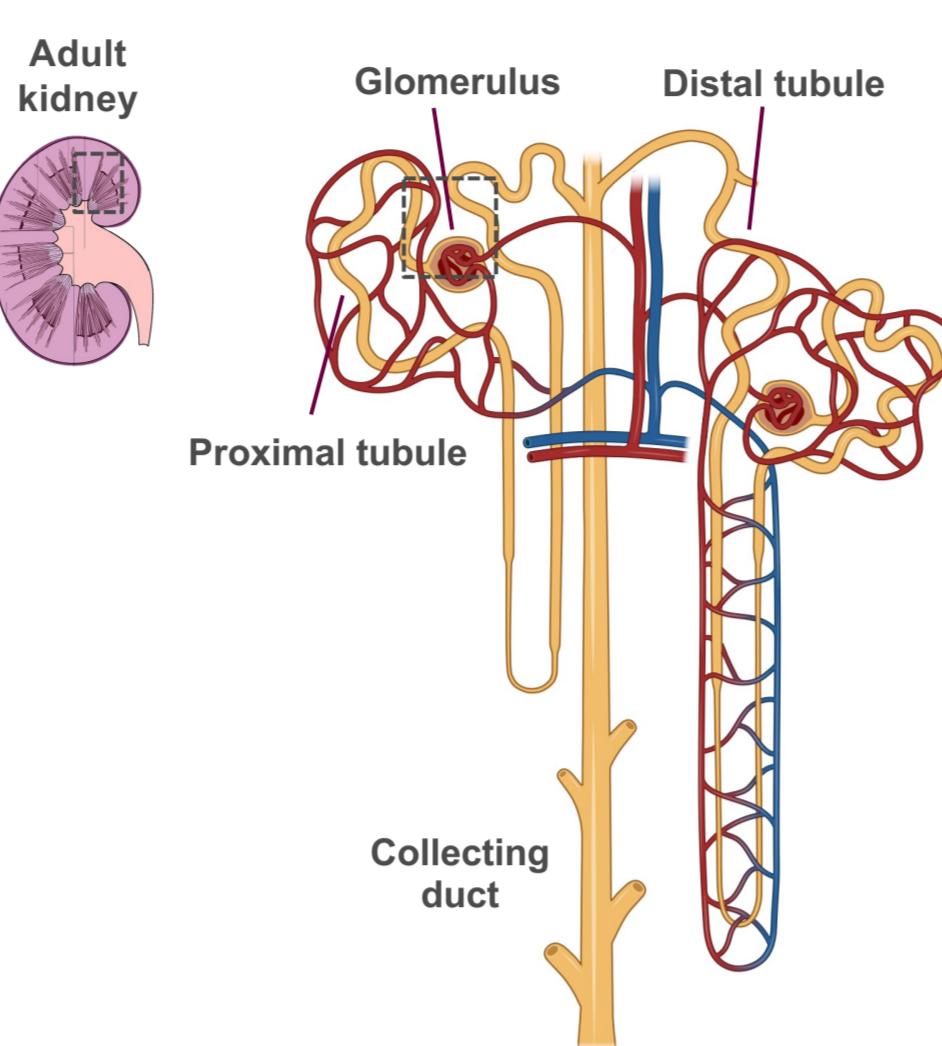


# Modeling X-linked Alport syndrome with deep-intronic variations in kidney organoids for antisense oligonucleotide-based therapy

Hassan Saei<sup>1</sup>, Bruno Estebe<sup>1</sup>, Nicolas Gaudin<sup>2</sup>, Mahsa Esmailpour<sup>1</sup>, Julie Haure<sup>1</sup>, Olivier Gribouval<sup>1</sup>, Christelle Arondel<sup>1</sup>, Vincent Moriniere<sup>3</sup>, Corinne Antignac<sup>1</sup>, Geraldine Mollet<sup>1\*</sup>, Guillaume Dorval<sup>1\*</sup>

## X-linked Alport Syndrome

**X-linked Alport syndrome (XLAS)** is a hereditary glomerulopathy arising from genetic mutations in the *COL4A5* gene, encoding the  $\alpha 5$  chain of the collagen IV [ $\alpha 5$ (IV)] in the glomerular basement membrane (GBM).

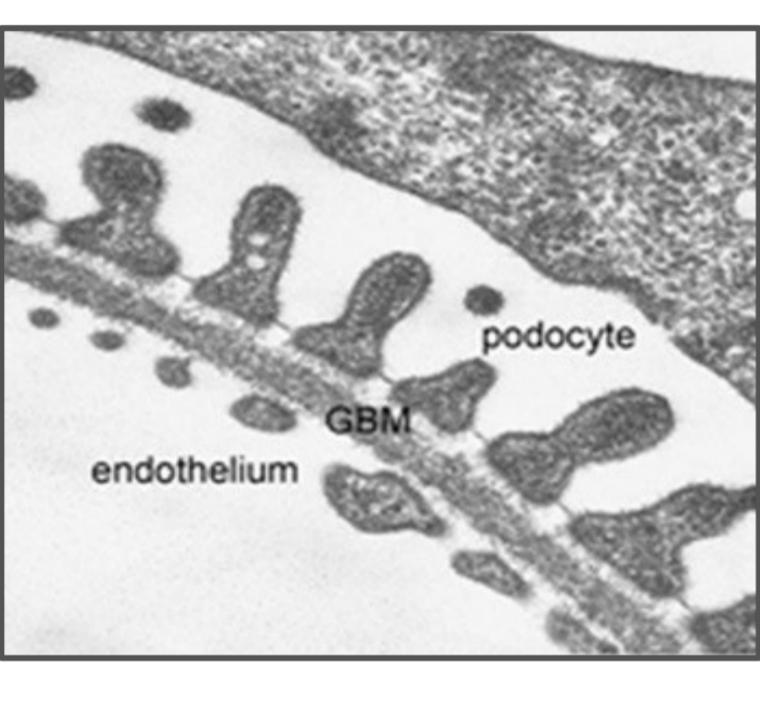


In a study on a cohort of **19 patients** with clinically proven XLAS, we identified deep-intronic variants responsible for the aberrant splicing events (17/19) using a **targeted RNA sequencing** approach.

The objective of this study is to develop a robust *in vitro* model for XLAS to characterize the disease and to test different therapeutic approaches including ASO therapy.



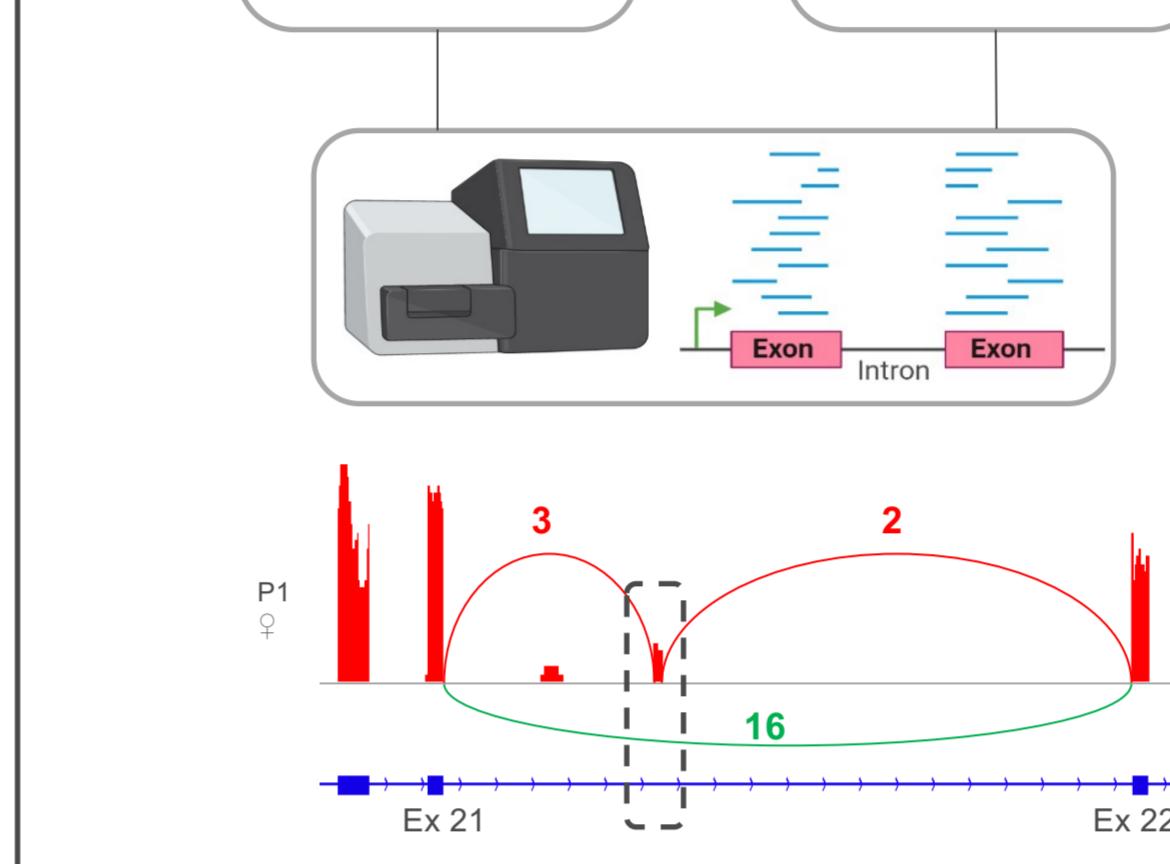
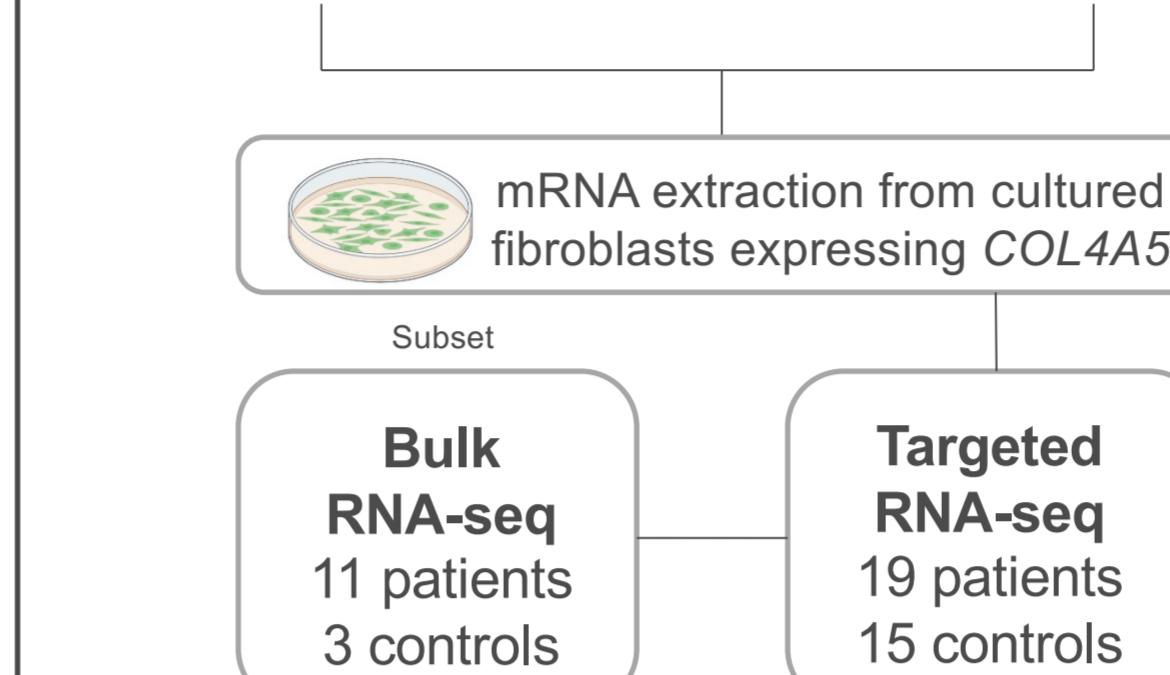
Andres et al., 2023; Naylor RW et al. Nat Rev Nephrol, 2021



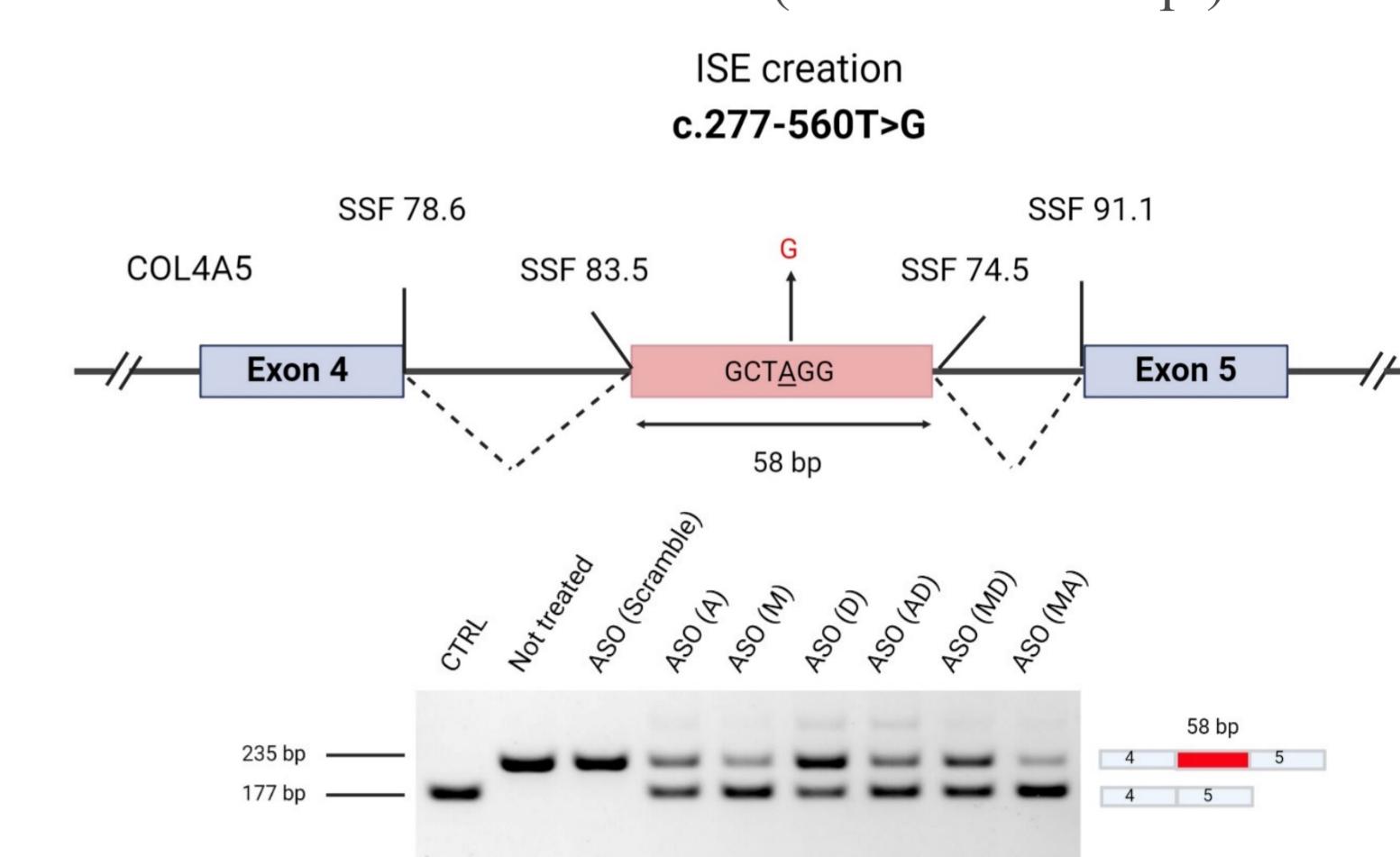
Boisson et al., Kidney International, 2023.

## Identification of Missing Variant Amenable to ASO Therapy

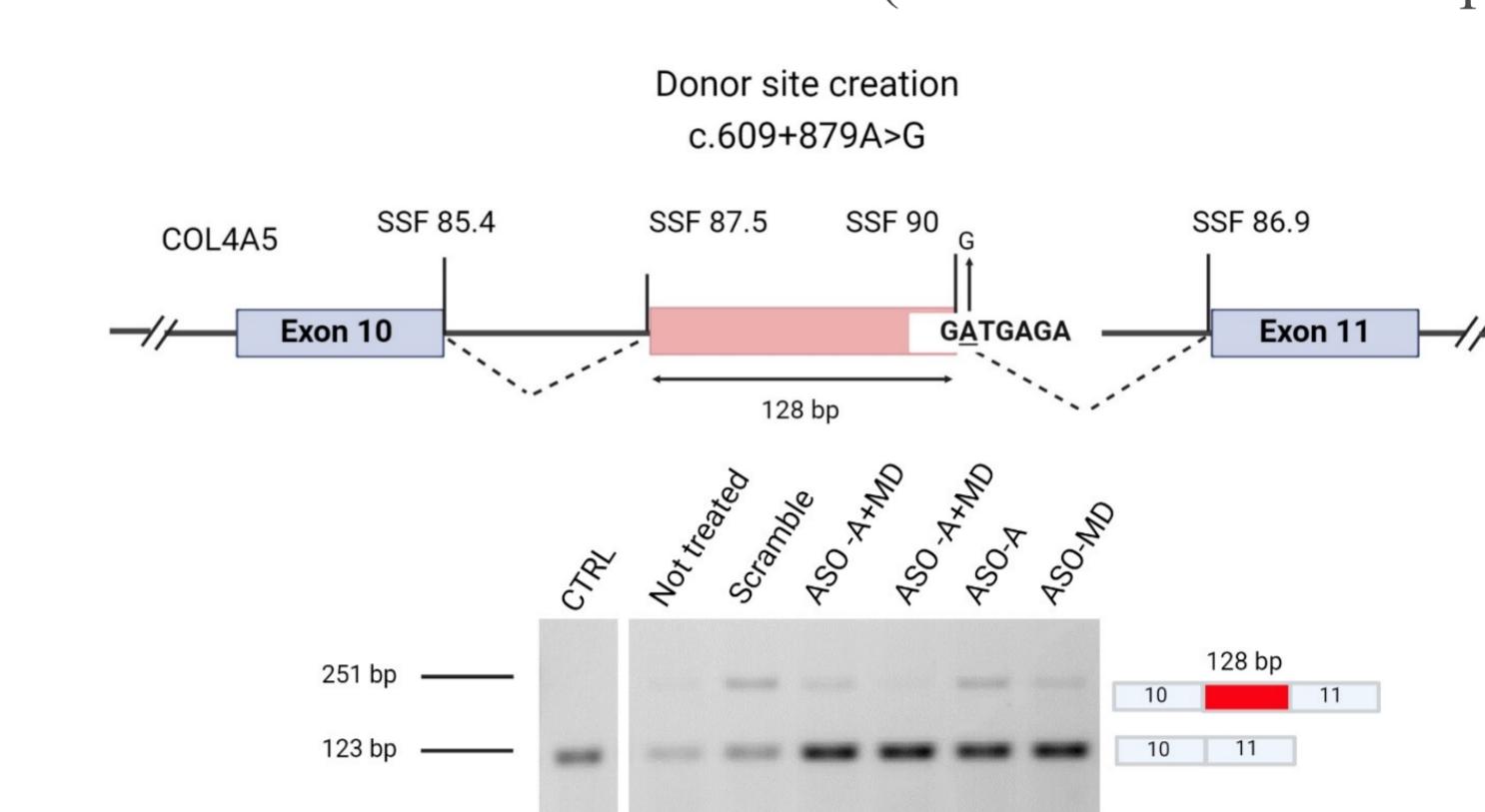
19 patients  
Clinically proven AS  
No pathogenic variant  
15 controls



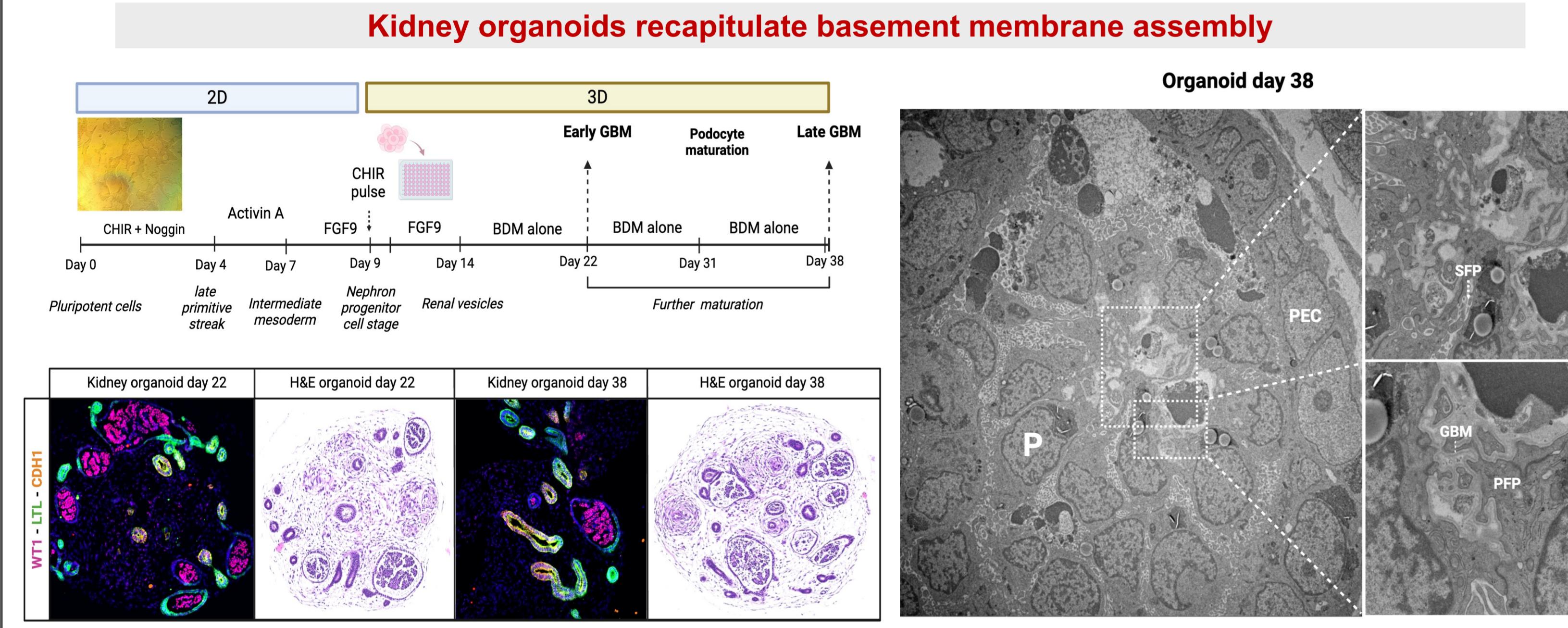
Patient 1: Severe XLAS (No WT transcript)



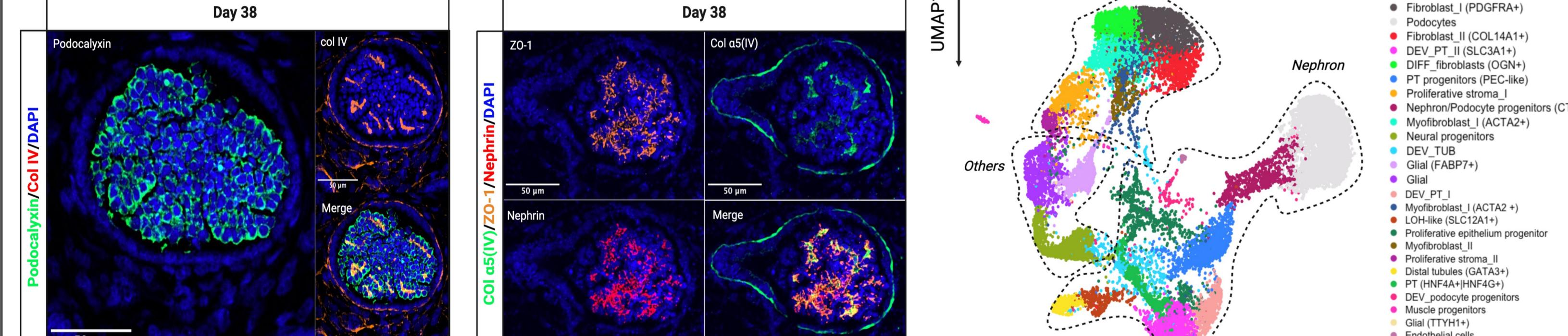
Patient 2: Moderate XLAS (Residual WT transcript)



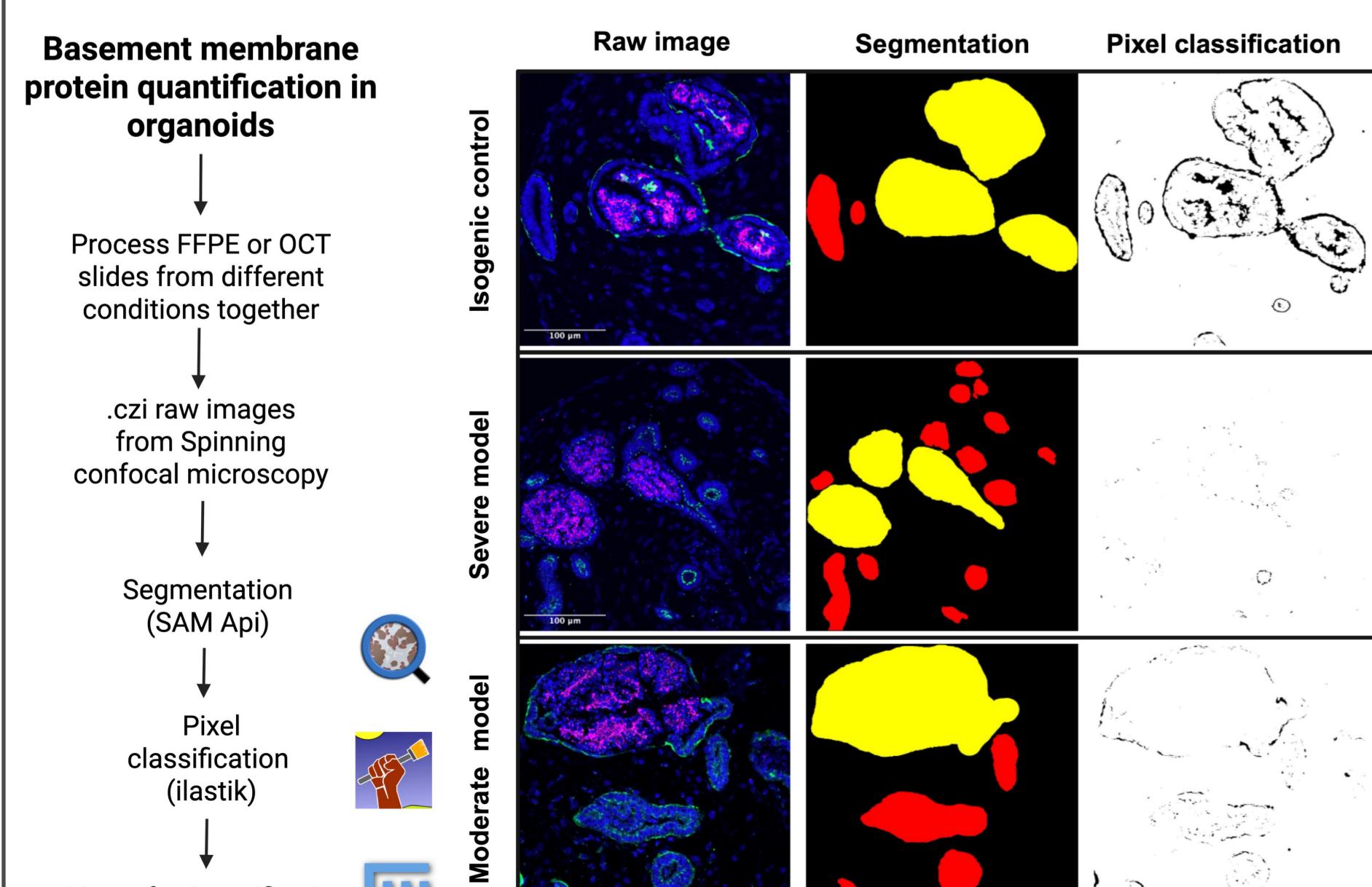
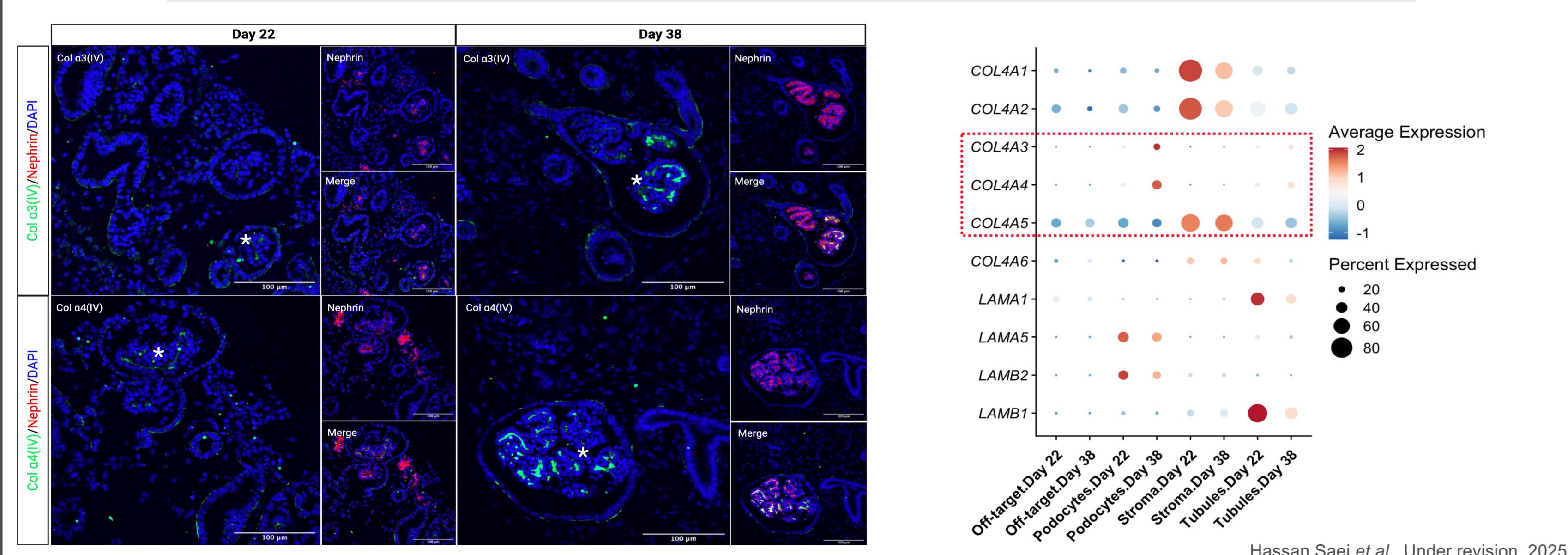
## Collagen IV Switch in Developing Organoids



### Podocytes in kidney organoids are polarized



### Prolonged organoid culture is essential for maturation of the GBM collagen network



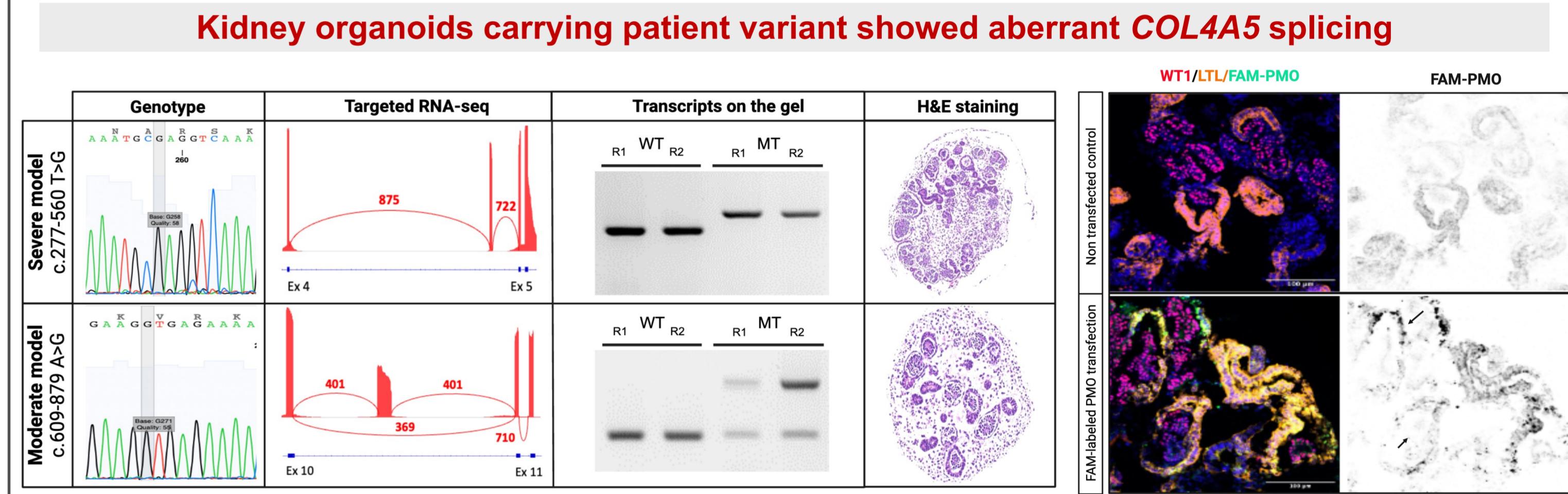
### BMQuant: A FIJI macro to quantify basement membrane proteins

This workflow begins with segmentation of glomeruli and tubule using QuPath, followed by pixel classification with ilastik. SAM API was used for accurate identification of object border and ROIs were saved. The channel of interest is being uploaded to ilastik for random forest model-based pixel classification. The FIJI macro generates csv files that contains measurements such as the full border area, mean channel intensity in the border and inside each object.

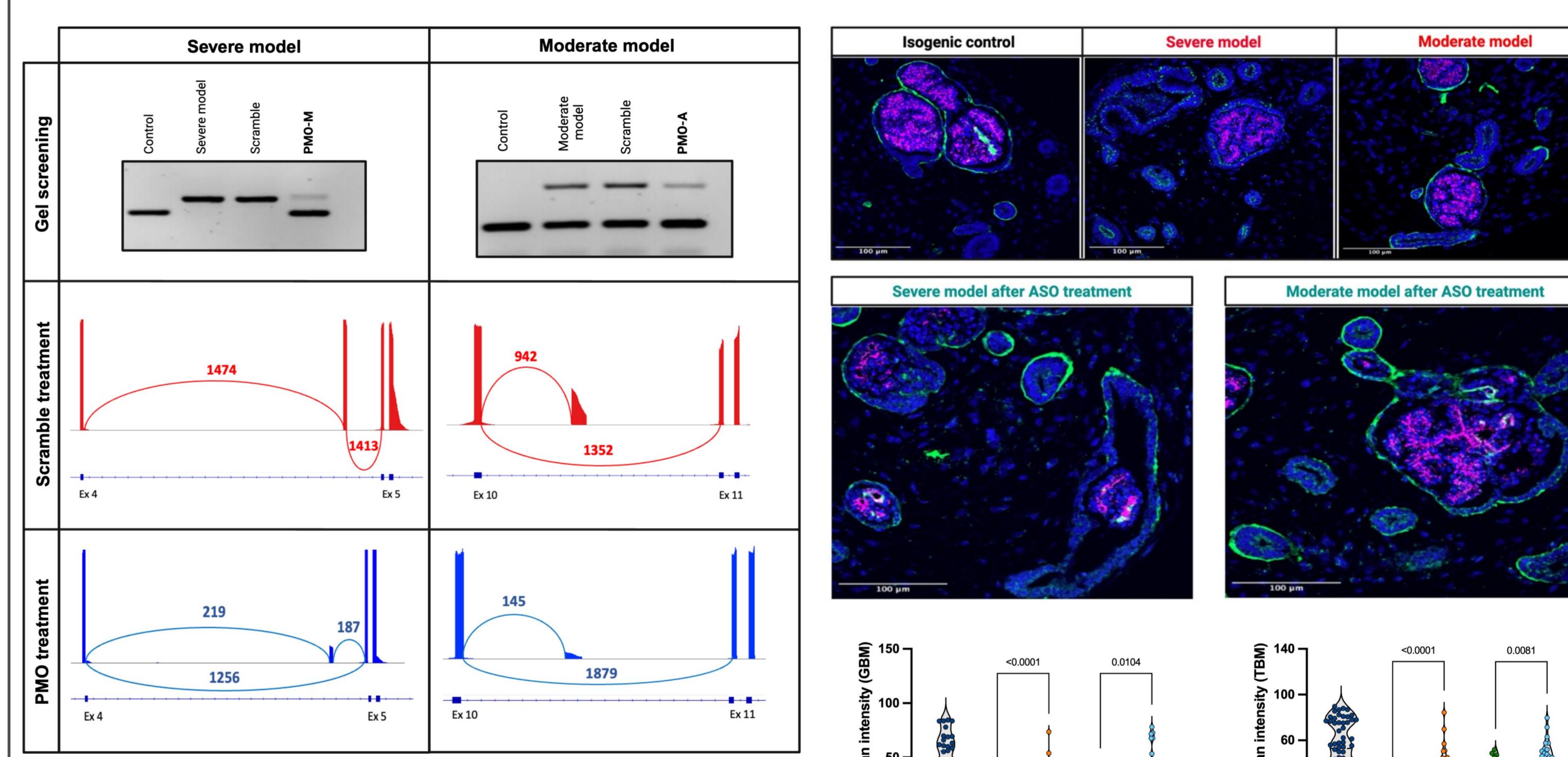


<https://github.com/hassansaei/BMQuant>

## ASO Treatment in XLAS Organoids with Deep Intronic Variants



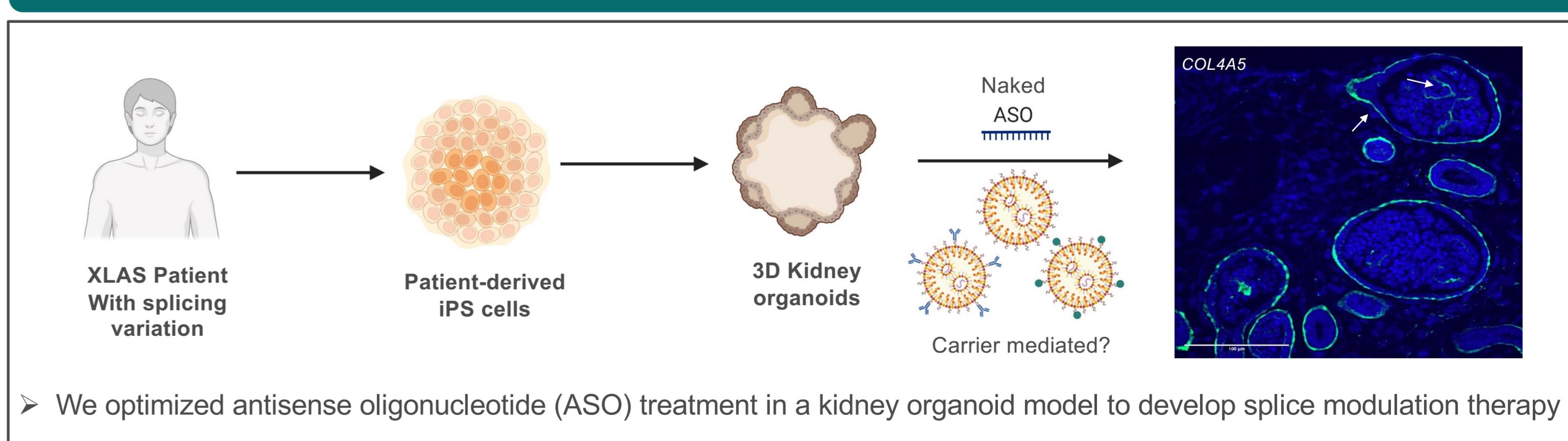
### ASO treatment reinstates collagen IV assembly in organoid models of XLAS



The severe model showed a complete absence of wild-type *COL4A5* transcript, which was significantly restored at both mRNA and protein levels following ASO transfection in organoids

Hassan Saei et al., Under revision, 2025

## Organoid Enable Scalable Development of Personalized Therapies



<sup>1</sup>Laboratory of hereditary kidney diseases, Inserm UMR 1163, Institut Imagine, Université Paris Cité, Paris, France <sup>2</sup>Necker Bioimage Analysis Core Facility of the Structure Fédérative de Recherche Necker, Paris, France <sup>3</sup>Medical genomics services for rare diseases, Hôpital Necker-Enfants Malades, Assistance Publique, Hôpitaux de Paris (AP-HP), Paris, France



Hassan.saei@inserm.fr



@hassansaei.bsky.social



<https://github.com/hassansaei>

