



MICHIGAN MEDICINE  
UNIVERSITY OF MICHIGAN

# Precision Medicine approach identifies patients with IgA nephropathy at risk for progression using endothelin activation signatures

Viji Nair<sup>1</sup>, Eric Olson<sup>2</sup>, Sean Eddy<sup>1</sup>, Wenjun Ju<sup>1</sup>, Marvin Gunawan<sup>2</sup>, Joyce Wu<sup>2</sup>, Jennifer Cox<sup>2</sup>, Andrew King<sup>2</sup>, Matthias Kretzler<sup>1</sup>.

<sup>1</sup>University of Michigan, Ann Arbor, MI USA,

<sup>2</sup> Chinook Therapeutics Inc, Vancouver, BC, Canada.



MICHIGAN KIDNEY  
TRANSLATIONAL MEDICINE  
CORE



KIDNEY  
WEEK 2021

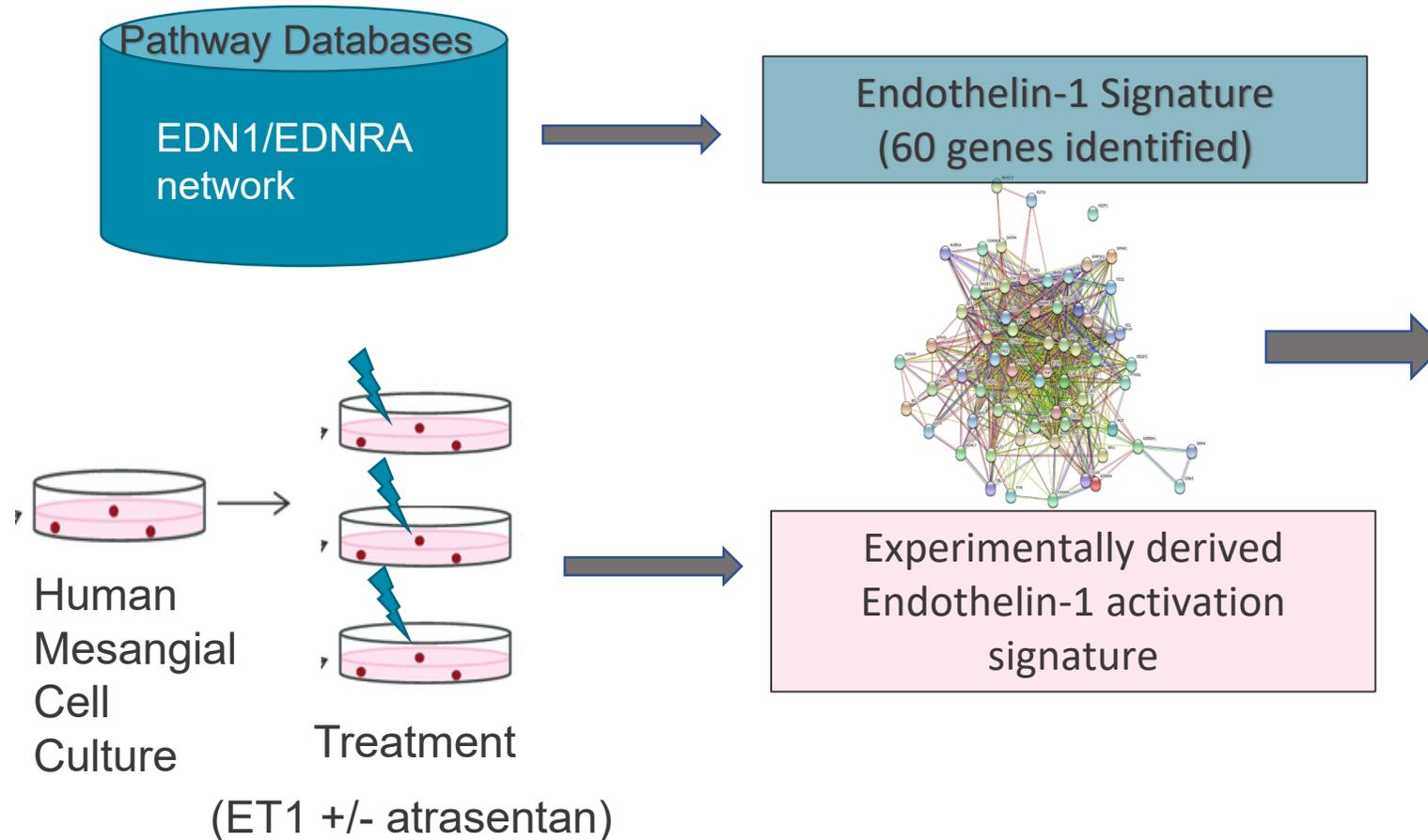
# Disclosures

- Funding from Chinook Therapeutics

# Background and Aims

- IgA nephropathy (IgAN) is the most common glomerulonephritis globally, with up to 40% of patients at risk of progressing to ESKD
- The aim of this study is to
  - Identify intra-renal transcriptional signatures of endothelin-1 (ET1)-activation network
  - Generate an ET1 activation score
  - Characterize the cellular location in the kidney of the signature genes
  - Relationship of ET1 activation and phenotype
  - Stratify patients at high risk of IgAN progression
  - Experimental evaluation of ETA receptor antagonist atrasentan treatment on the ET1 activation score in mesangial cells and animal models

# Methods overview

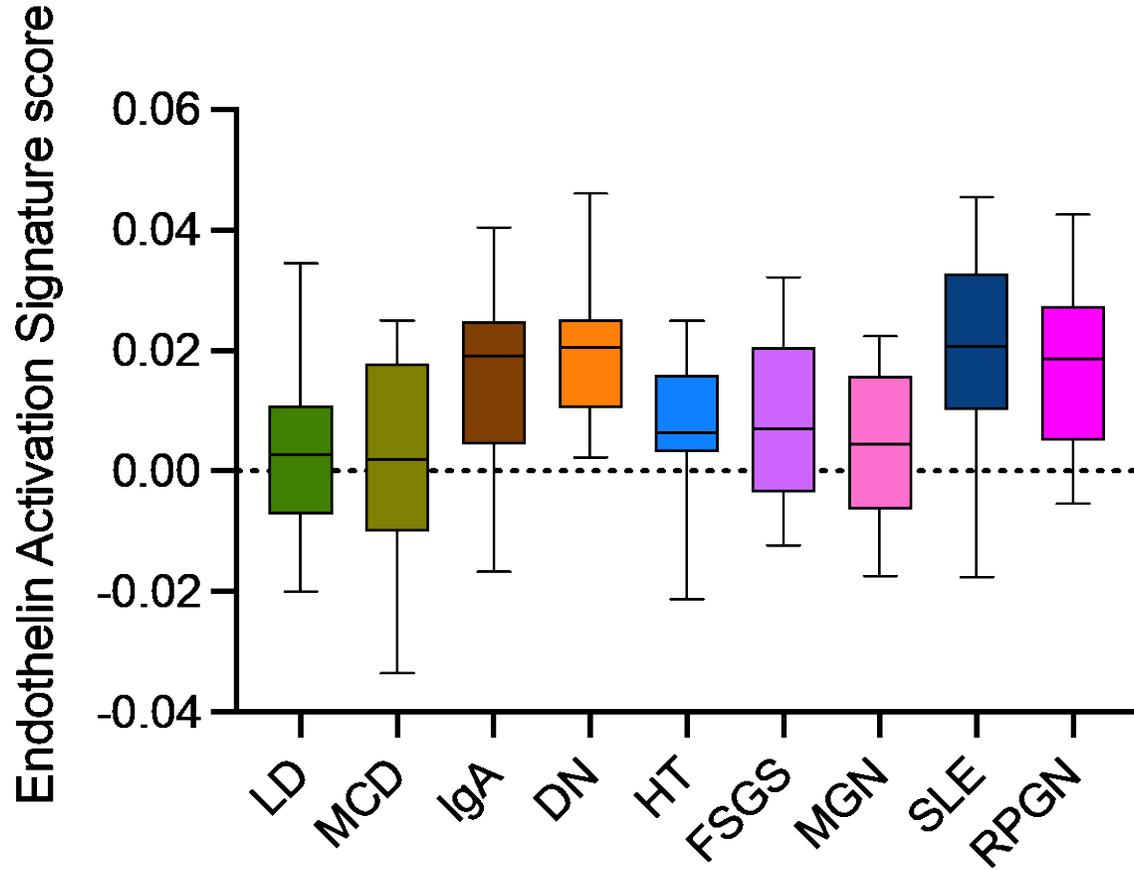


ET1 activation score with clinical measures and outcomes

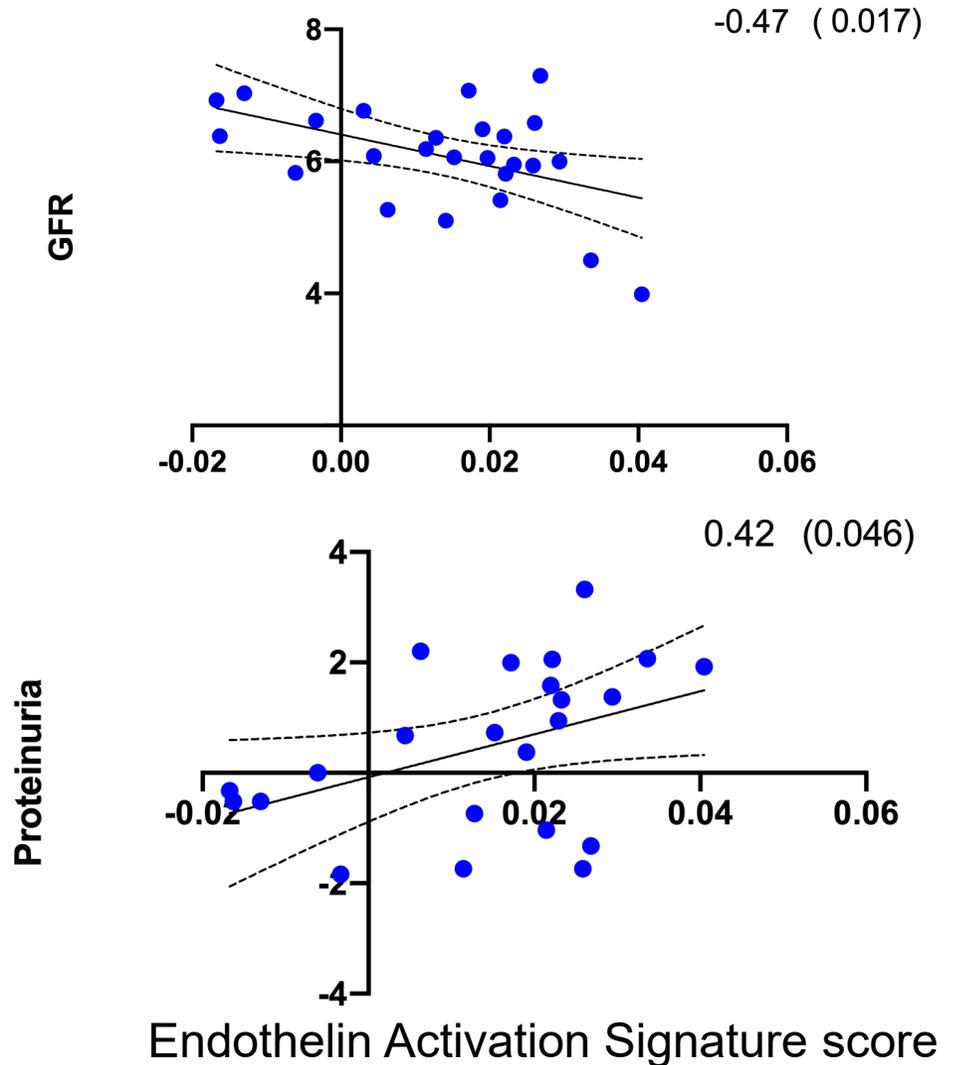
Cellular localization of signature in Single cell

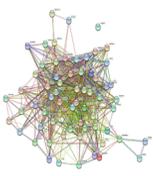


# ET1 activation in glomerular diseases (ERCBB cohort)



# ET1 activation scores strongly correlated with GFR/Proteinuria in IgAN patients

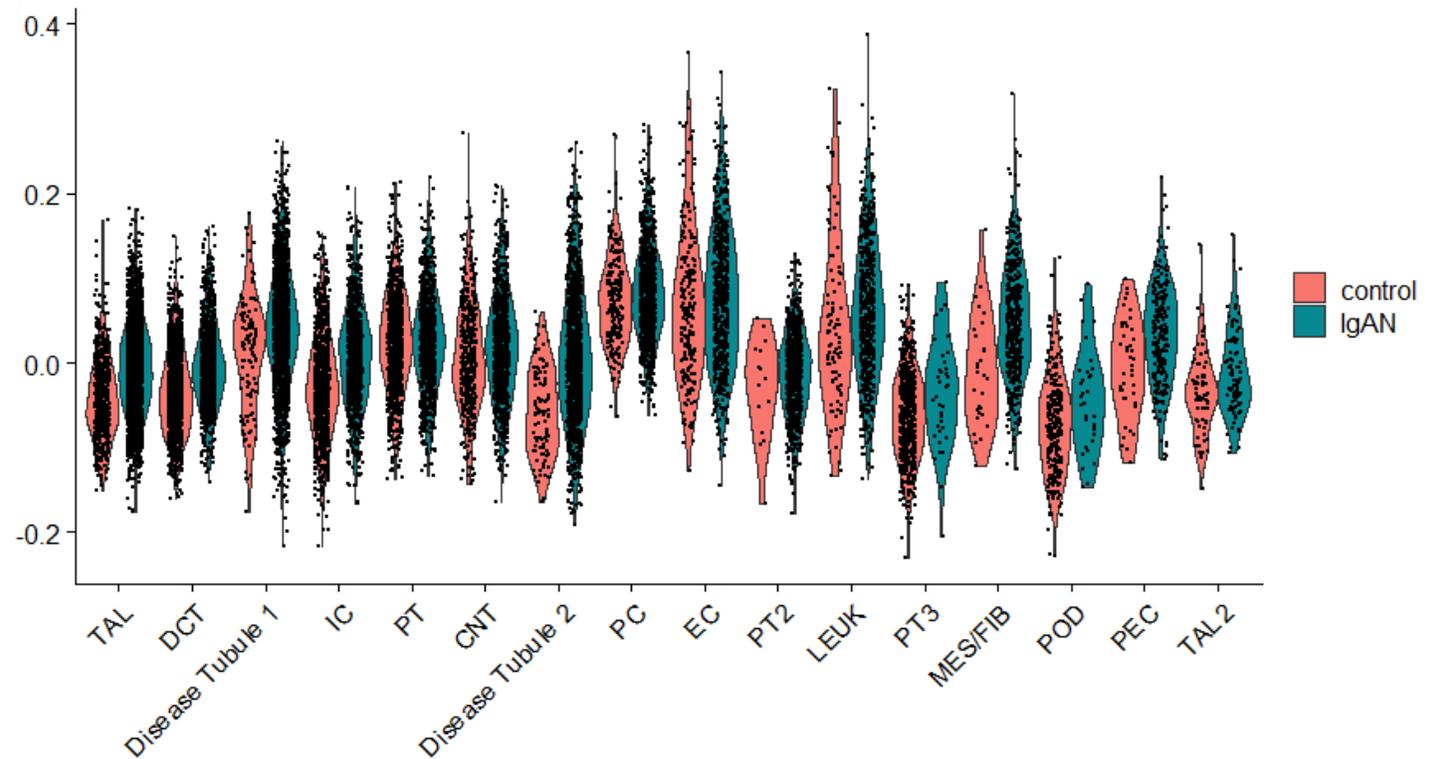




# ET-1 activation signatures in Single cell profiles derived from IgAN patients



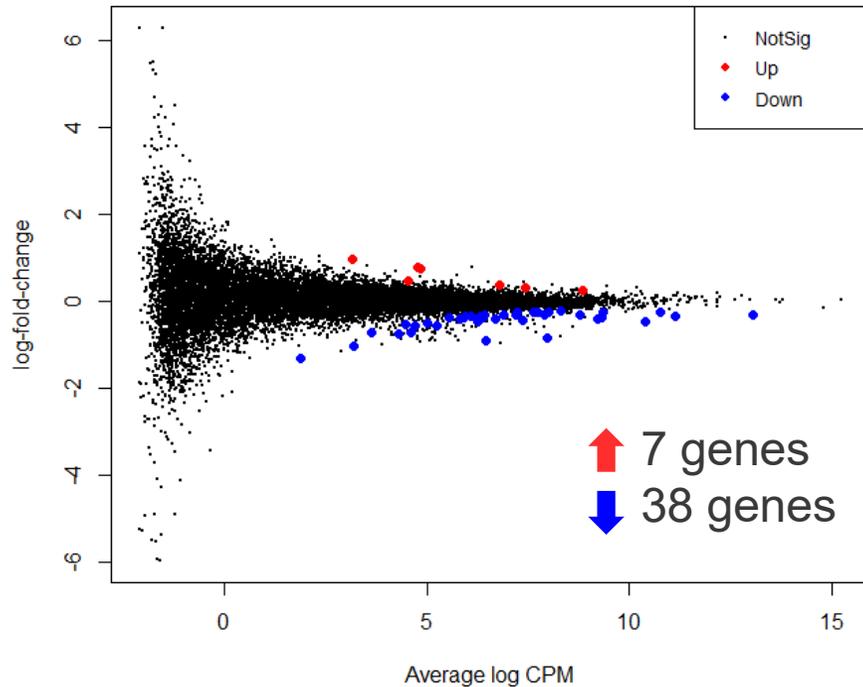
- ET1 activation scores in cell cluster profiles from IgAN patients\*
- The scores show higher enrichment in the mesangial cells, endothelial cells and Myeloid cells



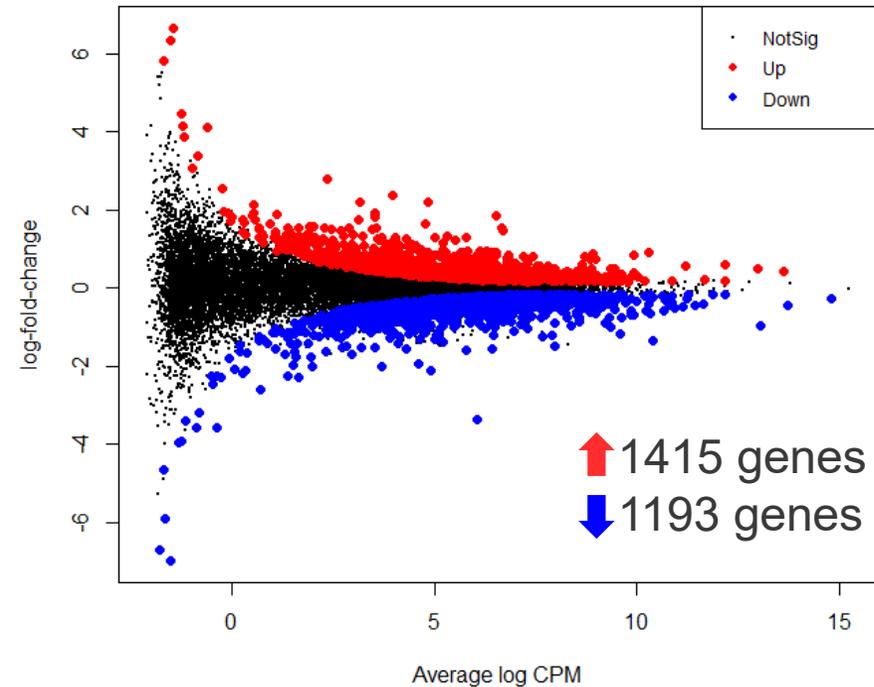
\*Tang R, Meng T, Lin W, Shen C, Ooi JD, Eggenhuizen PJ, Jin P, Ding X, Chen J, Tang Y, Xiao Z, Ao X, Peng W, Zhou Q, Xiao P, Zhong Y, Xiao X. A Partial Picture of the Single-Cell Transcriptomics of Human IgA Nephropathy. Front Immunol. 2021 Apr 16;12:645988.

# Atrasentan treatment shows dose-dependent blockade of ET-1 response in mesangial cells

Atra 1 nM (+ ET1) vs. ET1 (+DMSO)



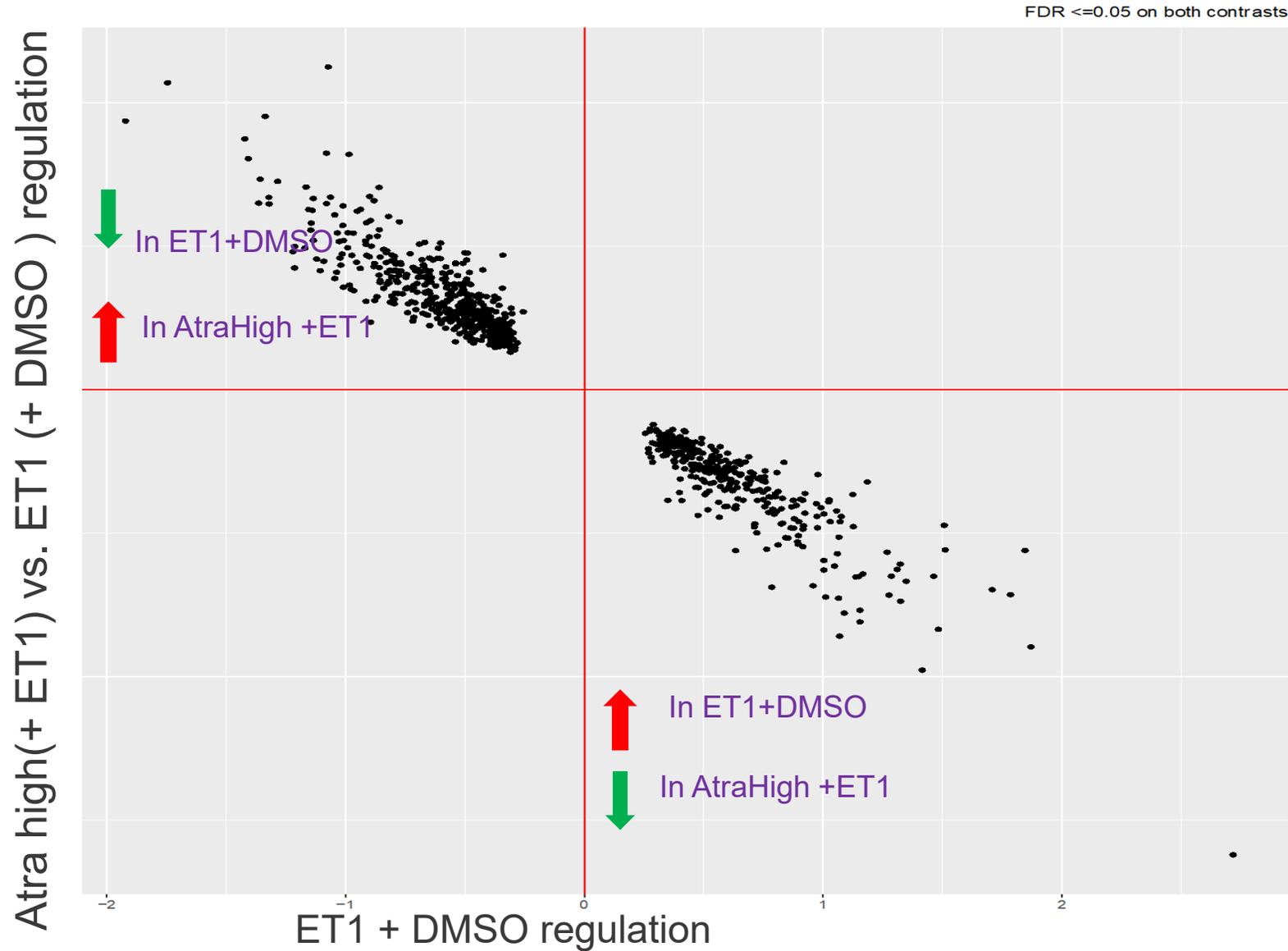
Atra 25 nM (+ ET1) vs. ET1 (+DMSO)



## Study design:

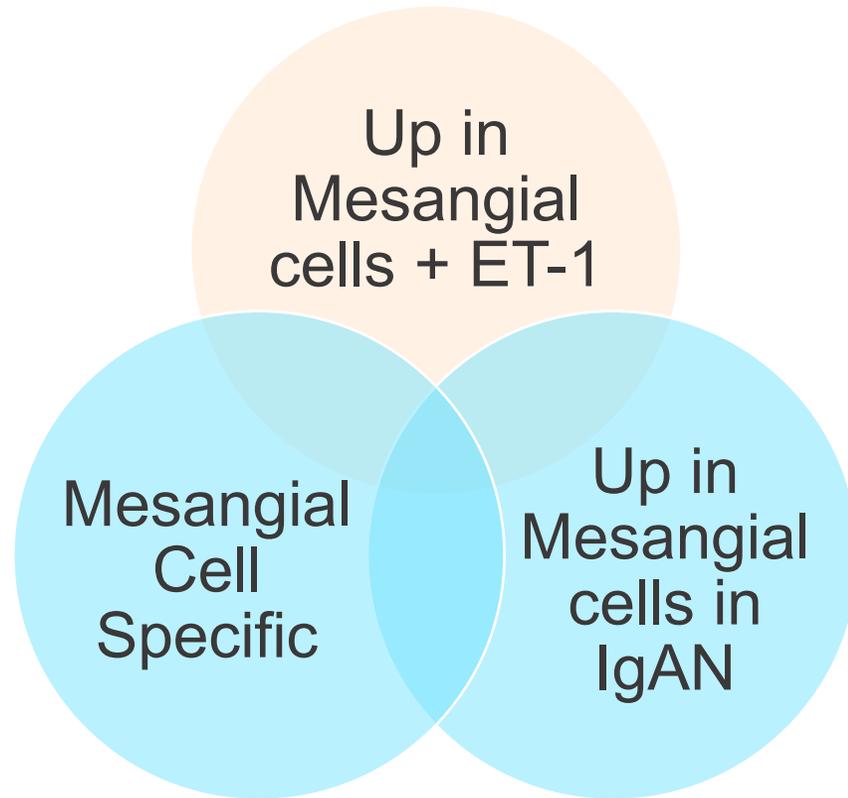
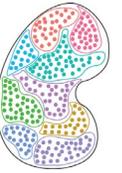
- In triplicates, cultured primary human renal mesangial cells from ScienCell were treated with ET-1 (4 nM) in the presence or absence of atrasentan (1 nM and 25 nM).
- 0.1% DMSO acted the vehicle control group.
- RNA-seq was performed 24 hours post-treatment.

# Reversal of ET-1 Gene Regulation by Atrasentan



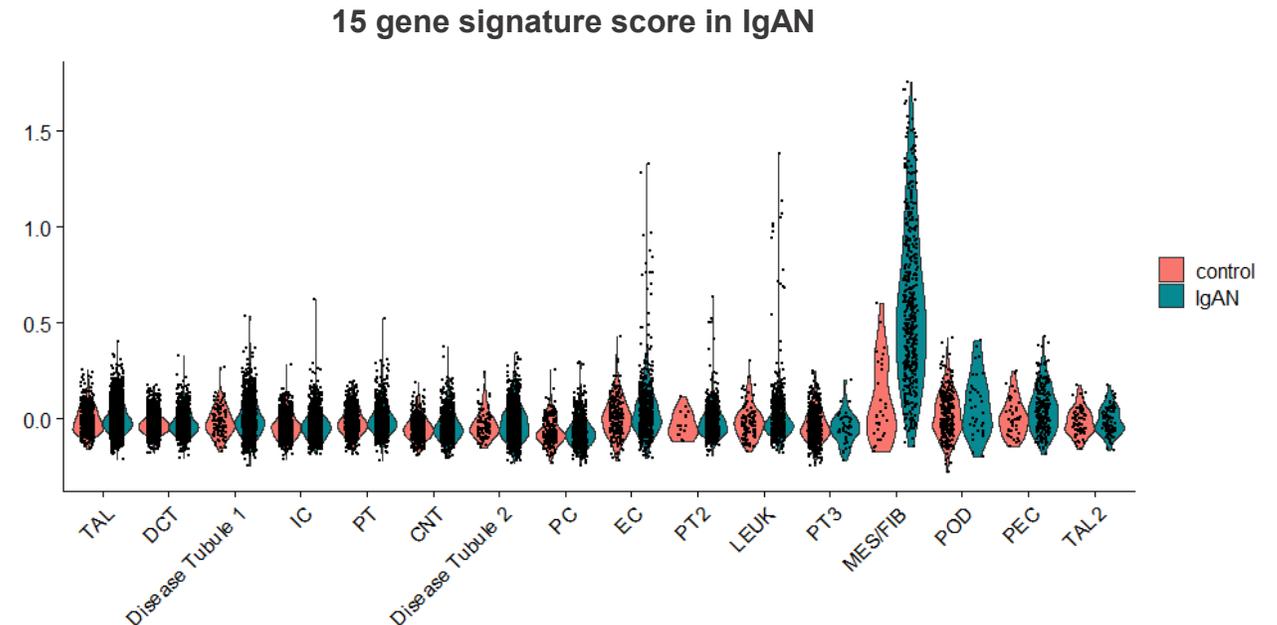
- Gene set enrichment analysis in MCs revealed up-regulation of cell proliferation, inflammatory and fibrotic networks, with ET1 treatment, which were blocked by atrasentan

# An approach to refine ET-1 activation signature for mesangial cell specificity



Other datasets –  
Bulk and snRNA-seq  
In vivo models +/- Atrasentan  
In vitro models +/- ET-1/Atrasentan

Strong activation of the ET1 signature in IgAN patient mesangial cells



# Conclusions

- Endothelin (ET) A receptor activation results in mesangial cell activation, proteinuria, inflammation, and fibrosis, all considered hallmarks of IgAN progression, suggesting the potential for therapeutic benefit of ETA antagonists
- ET1 transcriptional signatures show activation in IgAN patients, including in mesangial cells, and were significantly associated with clinical progression
- ET1 mediated transcript changes in mesangial cells, including hallmarks of mesangial cell activation (proliferation, inflammation and fibrosis) were dose-dependently inhibited by atrasentan