

Antibodies and Their Interactions with the Kidney Allograft Tissue in Antibody Mediated Rejection

Presented by Dr. Ana Konvalinka

Moderated by Dr. Lakshman Gunaratnam

The webinar will begin shortly



**Canadian Society of Nephrology/
Société canadienne de néphrologie**
CSN/SCN

LAND ACKNOWLEDGMENT

The Canadian Society of Nephrology Board of Directors has a commitment to diversity and inclusion.

We acknowledge that we are meeting on ancestral land that has been inhabited by Indigenous peoples from time immemorial. We are meeting virtually, so I would like to acknowledge that the Indigenous peoples are the traditional stewards of the lands and waters where each of us attends this meeting.



Ana Konvalinka
PhD, MD, FRCPC
Toronto, ON

Dr. Ana Konvalinka was recruited in 2015, as a transplant nephrologist and a Clinician Scientist at Toronto General Hospital, University Health Network. She is an Assistant Professor at the University of Toronto. Dr. Konvalinka completed medical studies at the University of Ottawa in 2003. She then completed internal medicine and nephrology training in Toronto in 2008. She subsequently embarked on a PhD in basic science at the University of Toronto. Her PhD thesis addressed the effect of angiotensin II on the proteome of primary human proximal tubular cells, and the relevance of this effect in vivo. Following completion of her PhD in 2013, she went on to complete the clinical kidney transplant fellowship at Toronto General Hospital. Her main clinical and research interests are in antibody-mediated rejection and kidney allograft fibrosis. She utilizes systems biology approaches and proteomics to enhance the understanding of the mechanisms, derive novel markers and to repurpose drugs for treatment of kidney disease. Dr. Konvalinka is the director of the Multi-Organ Transplant biobank for kidney, pancreas, and liver transplant programs. She is also the co-director of the Drug Discovery research group. She has received international research awards (the Human Proteome Project (2016), the American Society of Transplantation Faculty-Development Research Grant (2016) and the Advances in Organ Transplantation Award (2015)) and national research awards (Canadian Society of Nephrology New Investigator Lectureship (2017) and the KRESCENT New Investigator Award (2016)).



CSN Webinar

Antibodies and their Interactions with the Kidney Allograft Tissue in Antibody Mediated Rejection

Dr. Ana Konvalinka

Transplant Nephrologist, Clinician Scientist, University Health Network

Assistant Professor, University of Toronto

Senior Scientist, Toronto General Hospital Research Institute

Director, Ajmera Transplant Centre Multi-organ Transplant Biobank

Dec 13, 2022

OBJECTIVES

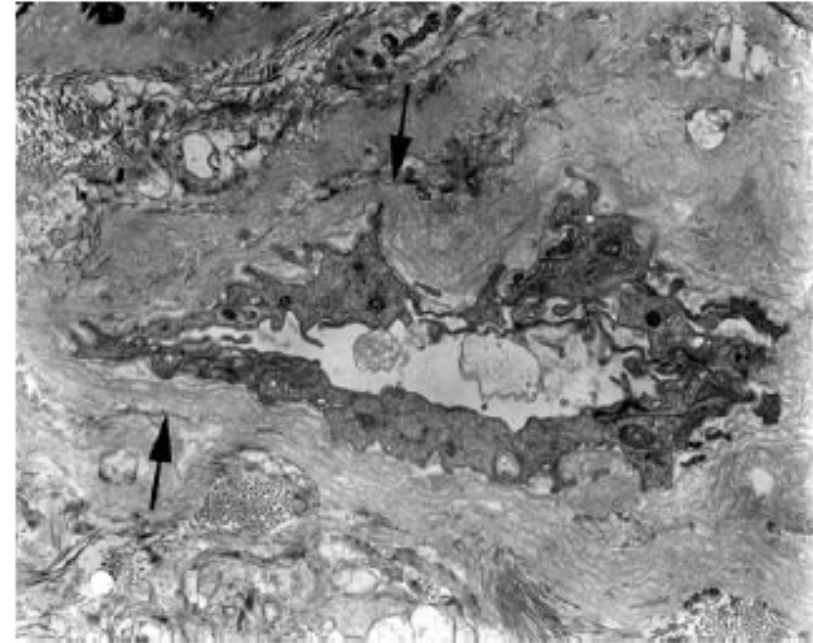
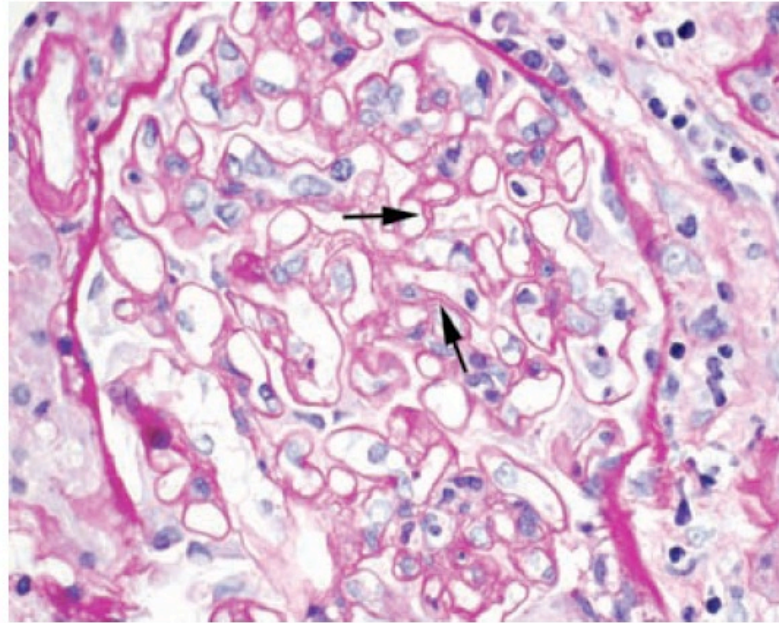
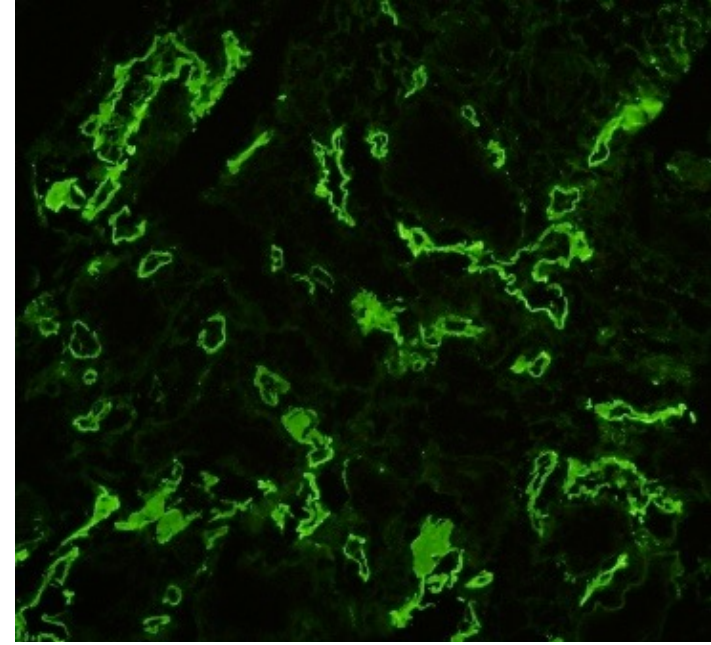
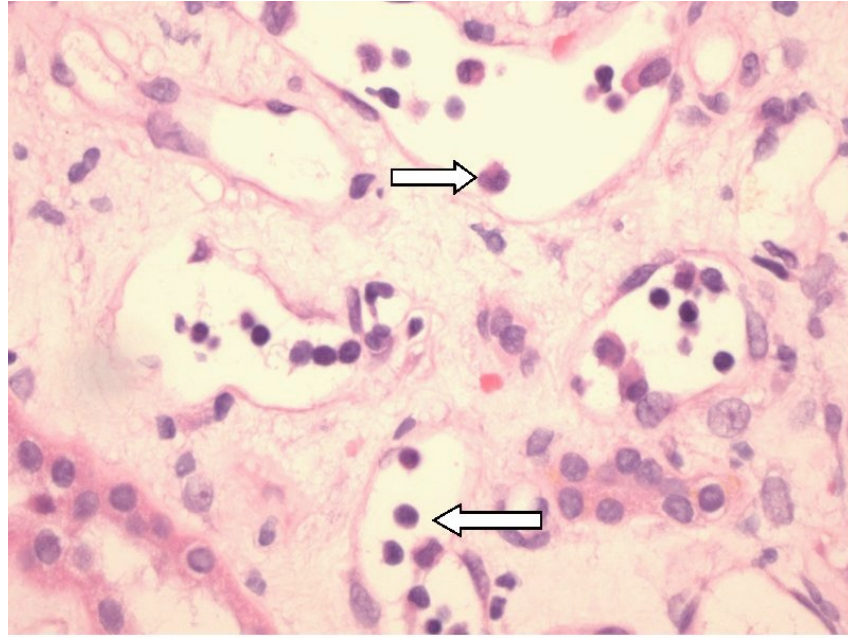
1. Provide an overview of antibody-mediated rejection in the kidney allograft
2. Describe the proteome changes in kidney glomeruli and tubulointerstitium in early antibody-mediated rejection
3. Review the importance of studying the extracellular matrix in the kidney
4. Describe novel approaches to delineate the biochemical features and roles of donor-specific antibodies in antibody-mediated rejection

DISCLOSURES

- Baxter (2022) - honoraria
- Promega - Academic access program

CASE 1

- 33F with ESKD due to IgA nephropathy
- Received a living donor kidney transplant 2 years previously
- Admits to non-adherence in the last few months; otherwise well
- Blood work reveals SCr of 210 μ mol/L from a base line of 120 μ mol/L
- US is normal
- New DSA against DQ6 is detected

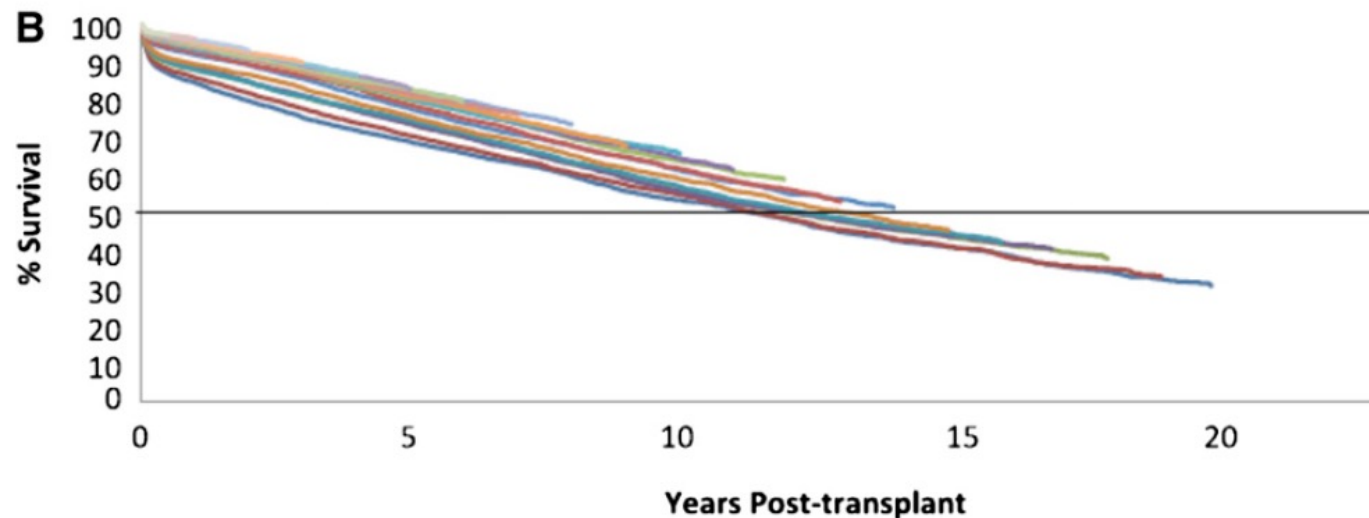


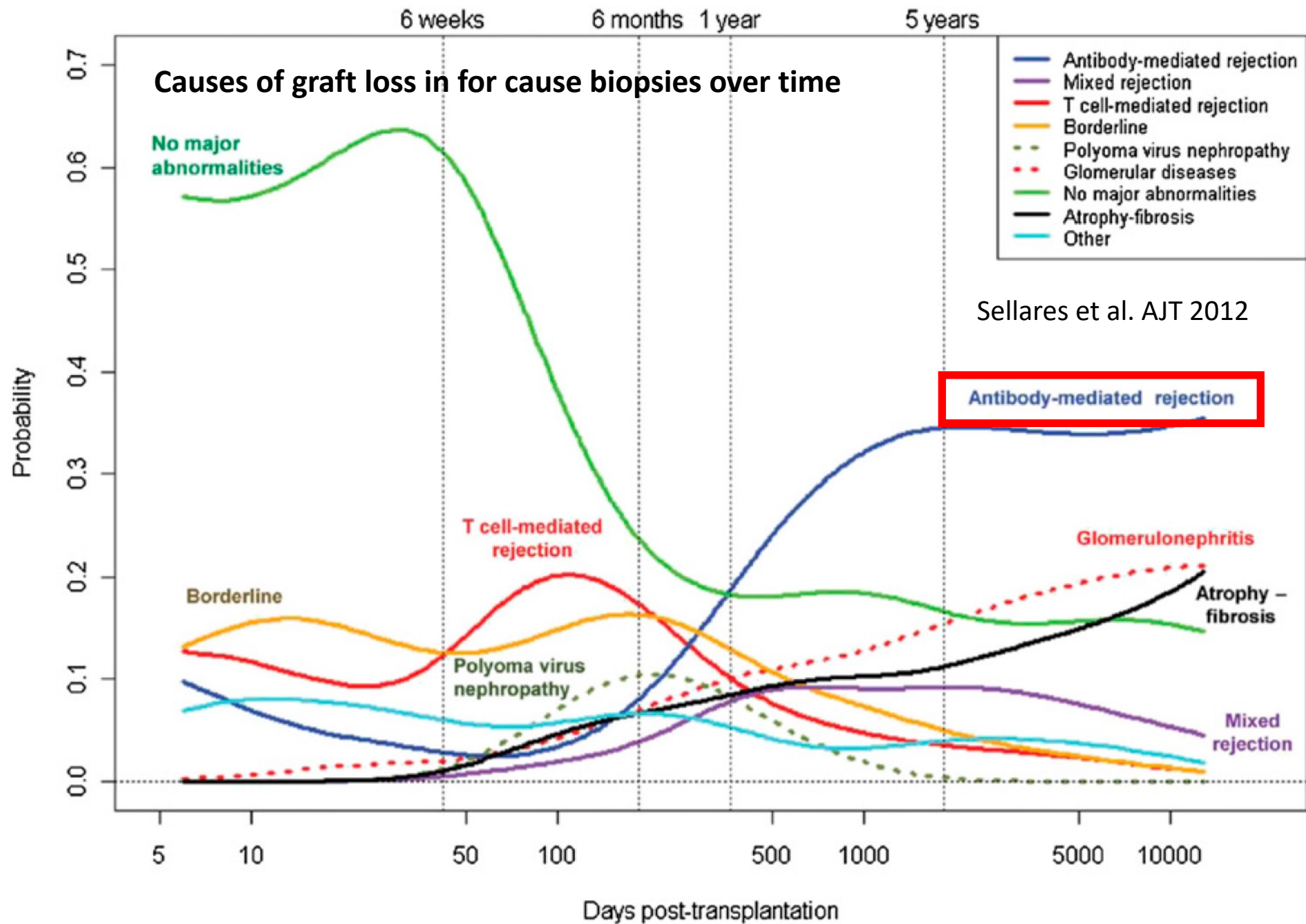
Overview

- Background
- Story #1 – Kidney glomerular and tubulointerstitial proteome in AMR
- Story #2 – Novel models to study AMR
- Story #3 – Assess circulating non-HLA antibodies in patients with AMR
- Future directions

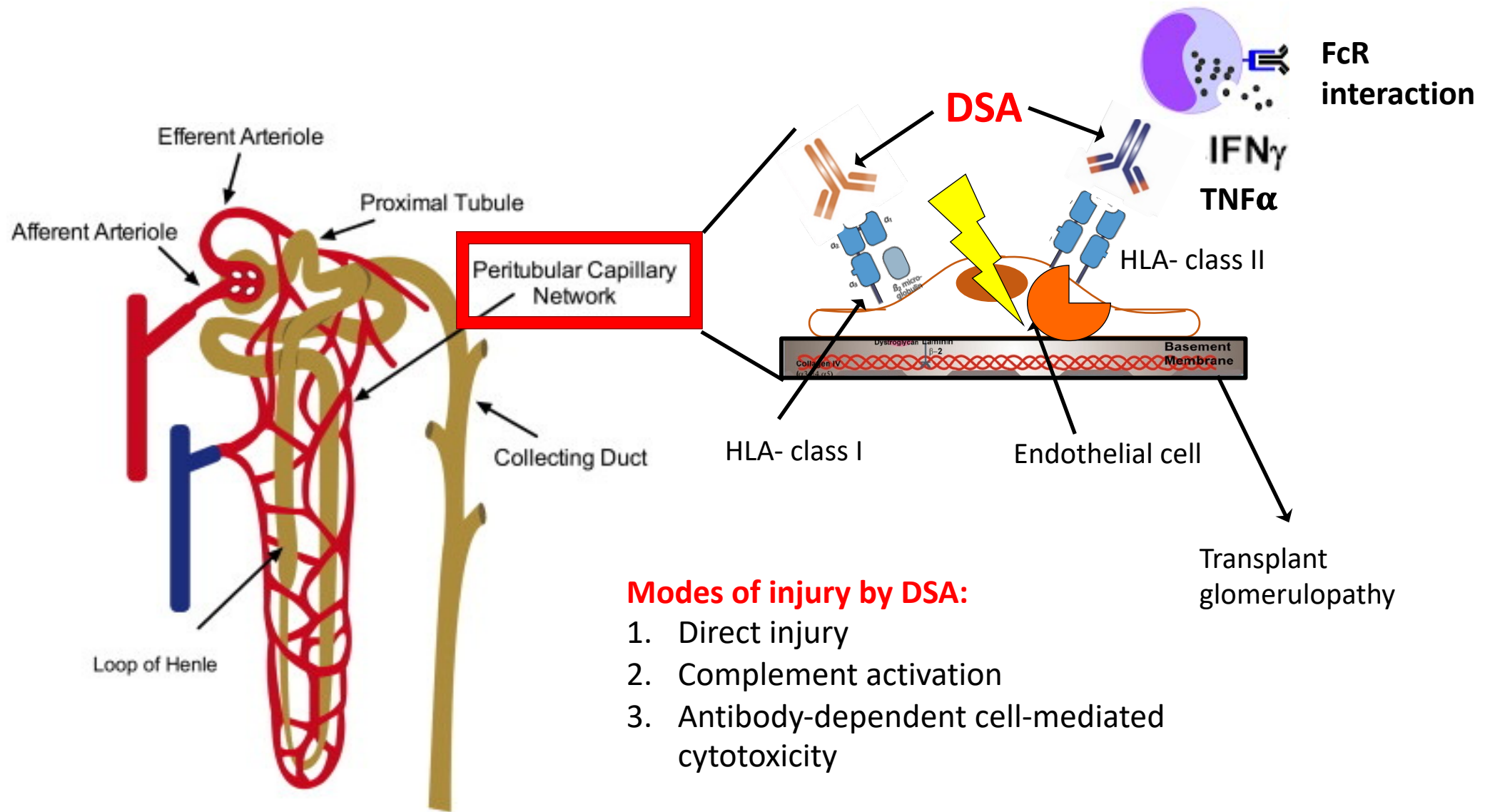
BACKGROUND

- Kidney transplantation is the best treatment for end stage kidney disease
- The rates of early kidney allograft rejection have diminished
- However, long-term allograft survival has not changed significantly (Lamb et al. AJT 2011)





Pathophysiology of AMR



Donor Specific Antibodies (DSA) – as a causal agent in AMR

- Class II HLA DSA appear more commonly after transplant and are associated with worse outcomes
- However, even in the presence of these antibodies, 30-60% of patients will do well
 - Yamamoto et al. *Transplantation* 2016
 - Matignon et al. *PLoS ONE* 2017
 - Heilman et al. *Transplantation* 2014
 - Lefacheur et al. *J Am Soc Nephrol* 2016
- Other DSA+ patients will experience AMR, TG and graft loss
 - Everly et al. *Am J Transplant* 2009, *Transplantation* 2010
 - DeVos et al. *Kidney Int* 2012
 - Willicombe et al. *Transplantation* 2012
 - Yabu et al. *Transplantation* 2011
- 40-60% of all AMR cases have no detectable DSA
 - Senev A, et al. *Am J Transplant* (2019) 19(3):763–80.
 - Bestard O, Grinyó J. *Am J Transplant* (2019) 19(3):952–3.
 - Koenig A, et al. *Nat Commun* (2019) 10(1):5350.

Gaps in knowledge

- The role of kidney tissue in antibody mediated rejection is poorly understood
- The most important non-HLA antibodies to be included in routine clinical monitoring are unknown
- Determination of pathogenic DSA is elusive
- Effective therapies are lacking

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Story # 1

Assess proteome of glomerular and tubular compartments of kidney biopsies with AMR compared to other forms of graft injury

Patient Characteristics

Group	Antibody-Mediated Rejection	Acute Cellular Rejection	Acute Tubular Necrosis
Number of patients	7	11	12
Sex - number of males (%)	4 (57.1)	9 (81.8)	9 (75)
Patient age at biopsy (years) median (IQR)	48 (43, 57.5)	42 (34, 48)	63.5 (57.5, 67.25)
Graft age (days post-transplant) median (IQR)	10 (8, 11)	16 (14.5, 57)	10 (8, 14)
Cause of ESRD (number)			
Diabetic Nephropathy	3	1	4
IgA Nephropathy	2	2	1
PCKD	0	2	3
FSGS	0	1	1
Vasculitis	0	2	0
Unknown	1	0	1
Other	1	3	2
Pre-existing autoimmune condition (number of patients)	1	1	0
Donor Type			
Deceased Donor	2	4	10
Living Donor	5	7	2
DSA pre-transplant (No of Patients)			
Class 1	3	0	4
Class 2	3	0	0
Prior desensitization- number (%)	2 (28.5)	0	0
Induction Agent - number (%)			
ATG	5 (71.4)	6 (54.54)	12 (100)
Basiliximab	2 (28.57)	5 (45.45)	0
Maintenance immunosuppression – number (%)			
Calcineurin Inhibitor	7 (100)	11 (100)	12 (100)
Anti-proliferative	7 (100)	11(100)	12 (100)
Prednisone	7 (100)	11(100)	12 (100)



Dr. Joseph Kim

Patient Characteristics

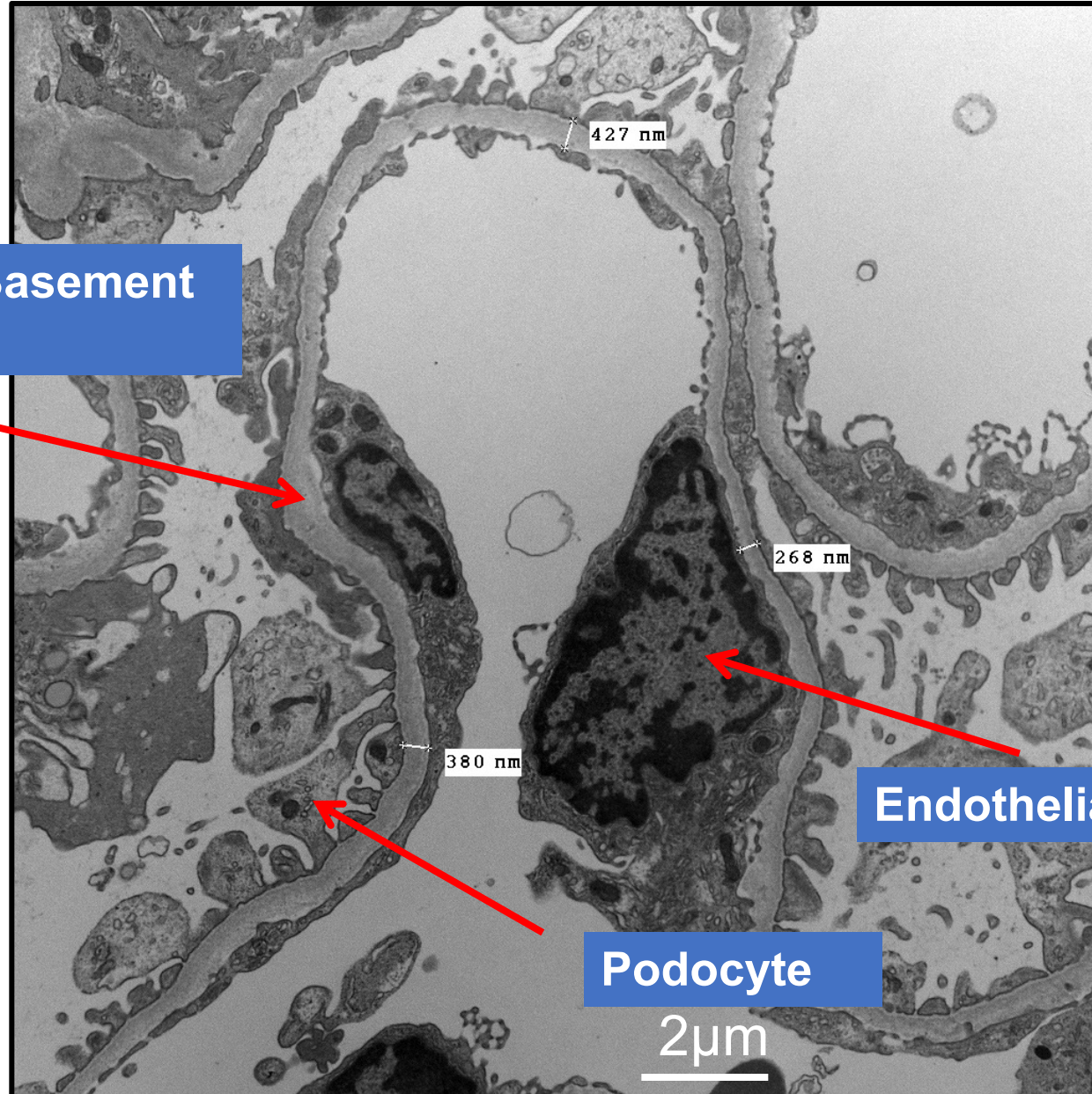
Group	Antibody-Mediated Rejection	Acute Cellular Rejection	Acute Tubular Necrosis
Biopsy Findings:			
Acute tubular necrosis (ATN)	5/7	3/11	12/12
Antibody-mediated rejection (AMR)	7	0	0
Acute cellular rejection (ACR)	0	11	0
Grade: 1A	NA	1	NA
1B	NA	7	NA
2A	NA	3	NA
Banff Scoring- median (IQR)			
% globally sclerosed glomeruli (gsg)	0 (0, 4.38)	3 (0, 5)	1 (0, 5.43)
Interstitial Inflammation (i)	1 (0, 1)	2 (2, 2)	0 (0, 0)
Tubulitis (t)	0 (0, 0)	3 (2.5, 3)	0 (0, 0)
Total inflammation (ti)	1 (0, 1)	2 (2, 3)	0 (0, 0)
Glomerulitis (g)	1 (0.5, 1.5)	0 (0, 0)	0 (0, 0)
Peritubular capillaritis (ptc)	1 (0, 2)	1 (0, 1)	0 (0, 1)
Intimal arteritis (v)	0 (0, 0)	0 (0, 0)	0 (0, 0)
Chronic Glomerulopathy (cg)	0 (0, 0)	0 (0, 0)	0 (0, 0)
Interstitial Fibrosis (ci)	0 (0, 0)	0 (0, 1)	0 (0, 0)
Tubular atrophy (ct)	0 (0, 0)	1 (0.5, 1)	1 (0, 1)
Arteriolar hyalinosis (ah)	0 (0, 0)	0 (0, 0.5)	0 (0, 0)
Vascular Fibrous Intimal Thickening (cv)	0 (0, 0)	1 (0, 1)	1 (0, 1.25)
C4D Staining	2 (1.5, 3)	0 (0, 0)	0 (0, 0)



Dr. Rohan John

Histopathology

Glomerular Basement Membrane



Endothelial cell

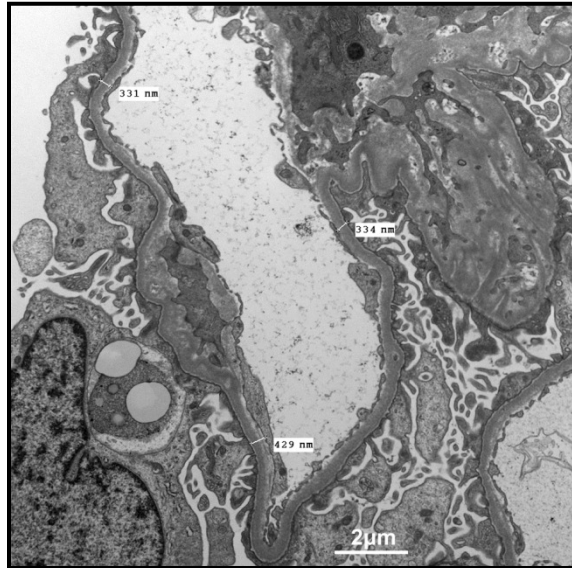
Podocyte

2 μm

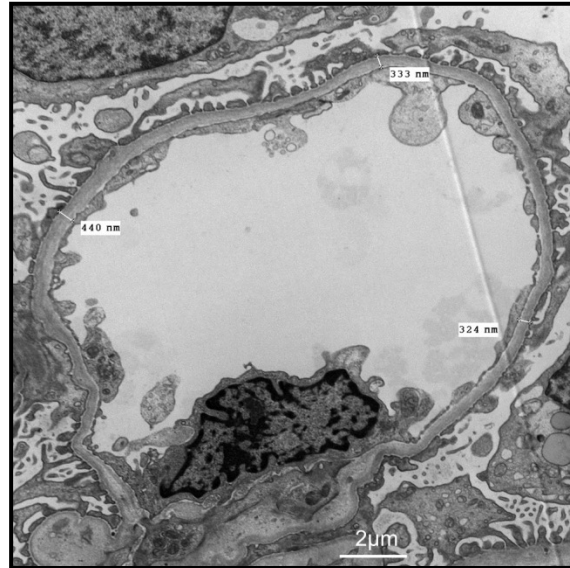
Histopathology

- No changes in Glomerular Basement Membrane Thickness

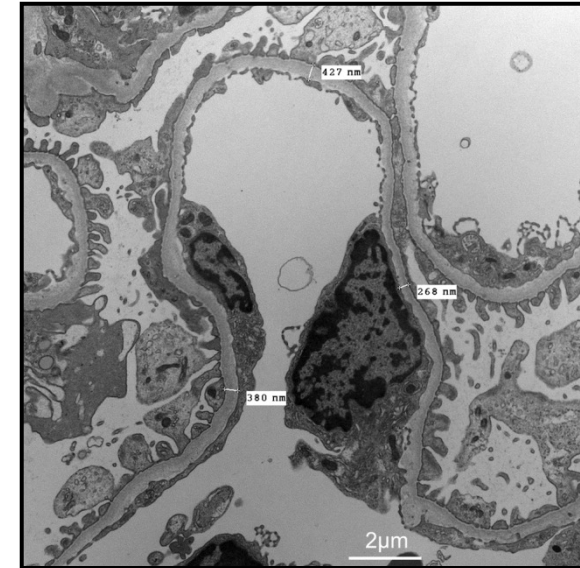
AMR



ACR



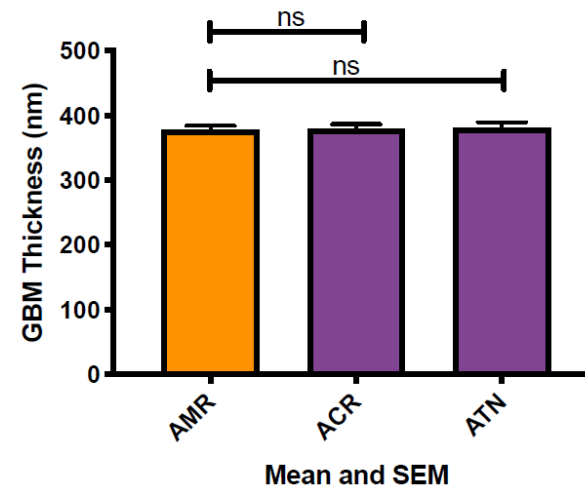
ATN



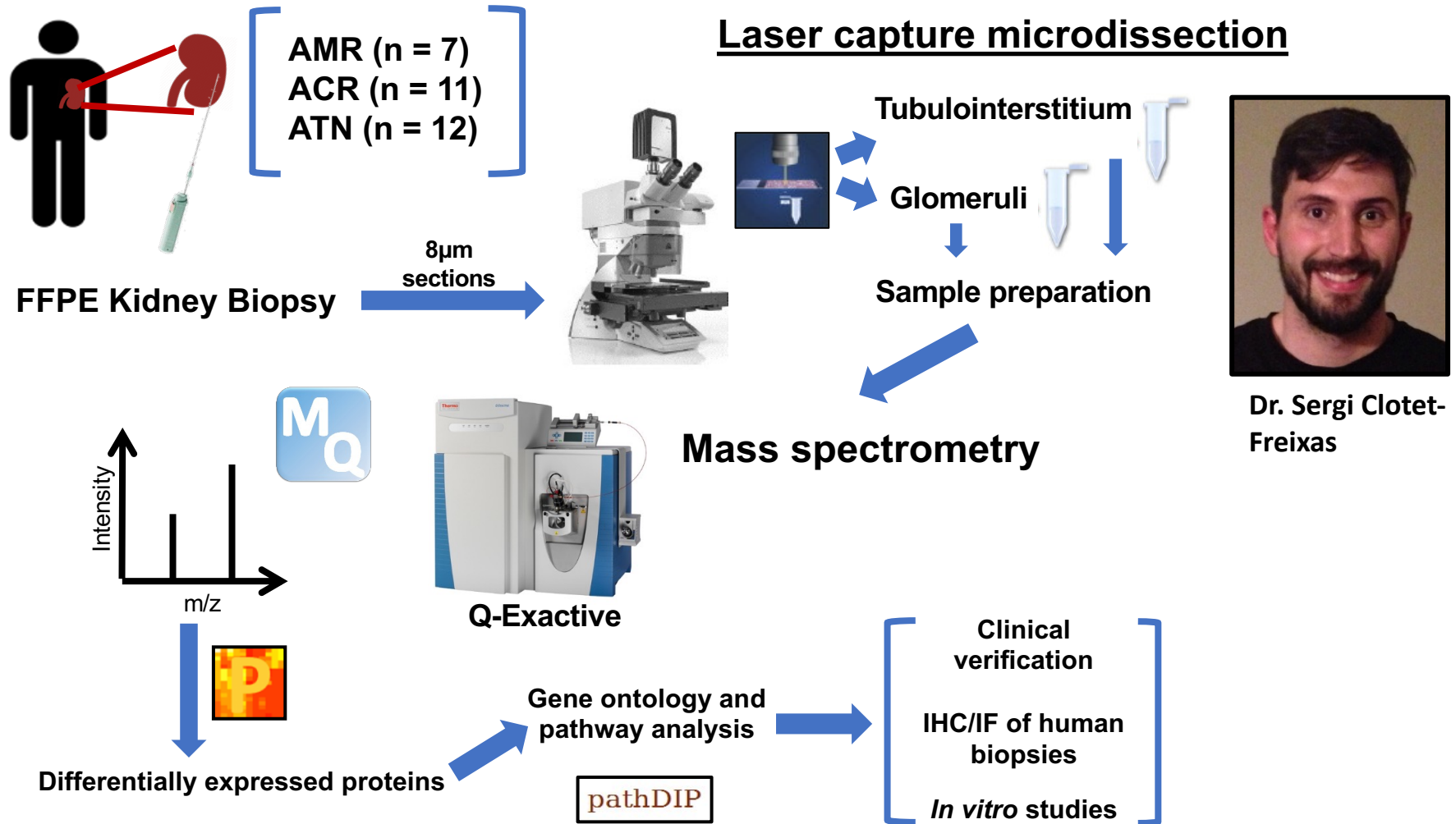
Dr. Rohan John



Dr. Caitriona McEvoy



METHODS



RESULTS

GLOMERULAR compartment

2026 proteins identified



1299 proteins quantified in
at least 50%
samples/group

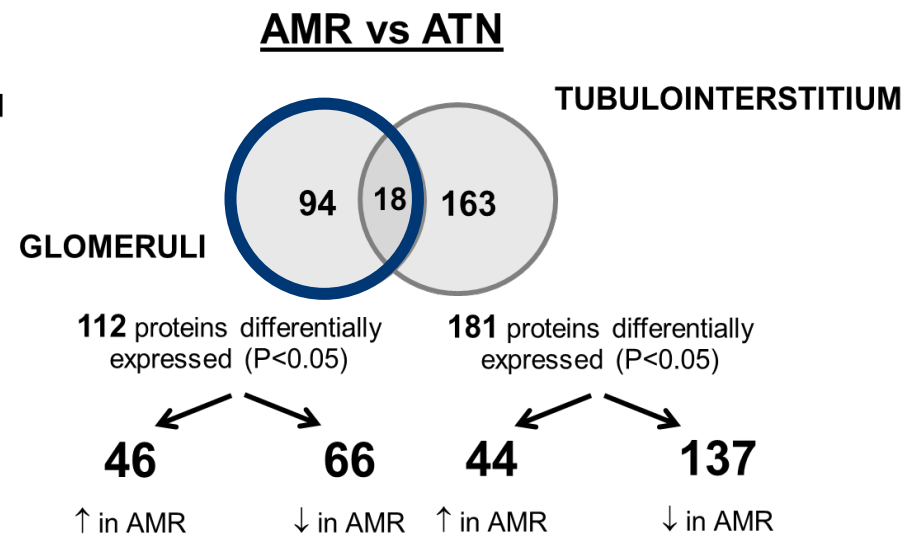
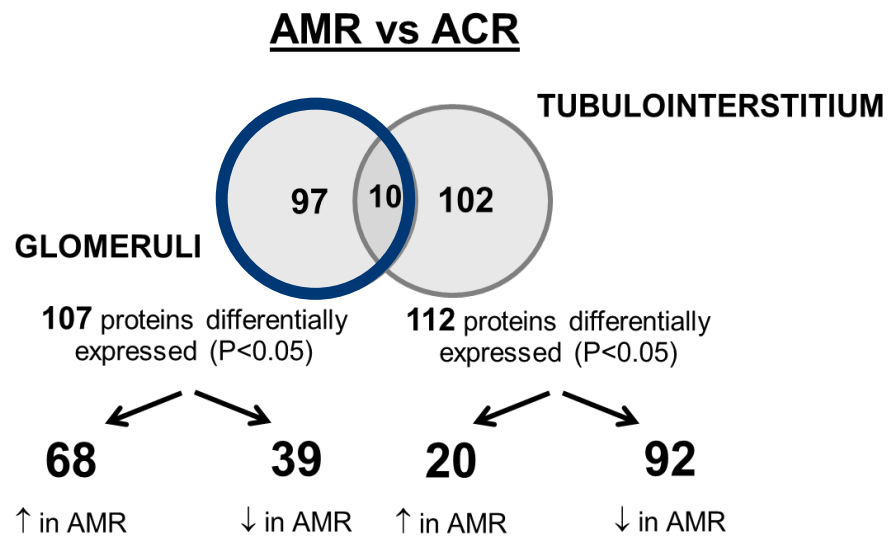
TUBULOINTERSTITIAL compartment

2399 proteins identified

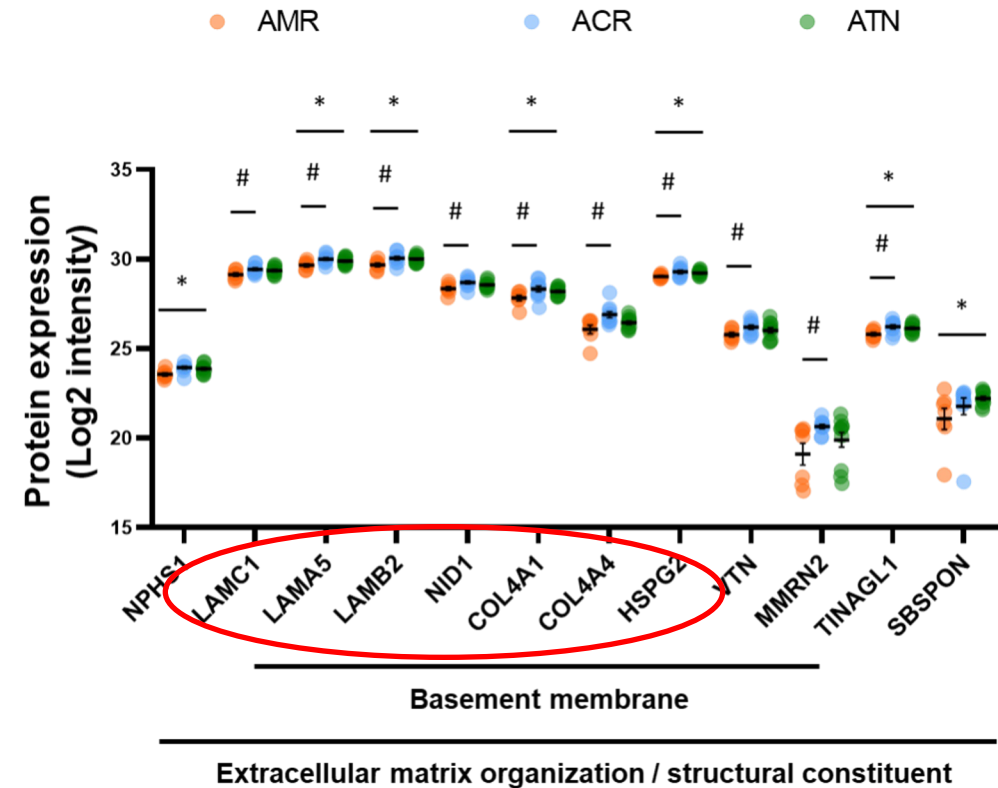


1842 proteins
quantified in at least
50% samples/group

DIFFERENTIALLY EXPRESSED PROTEINS

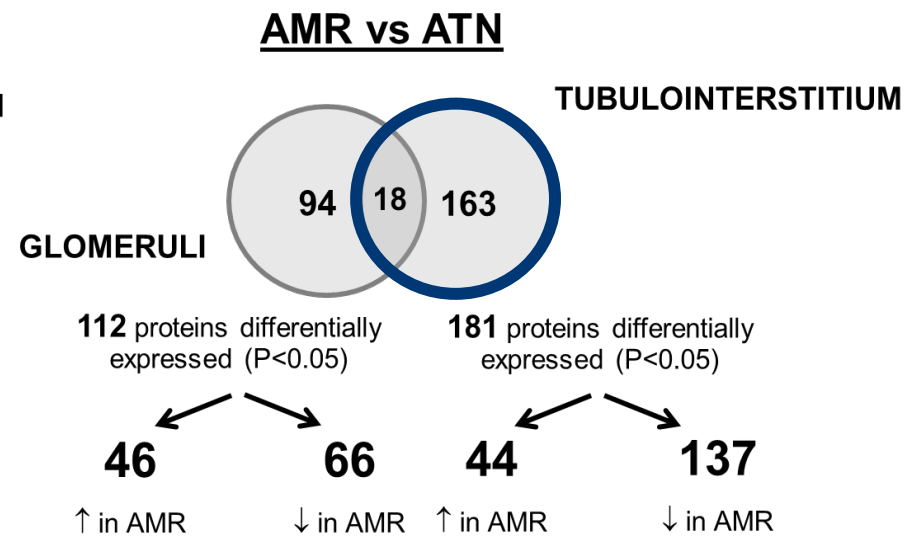
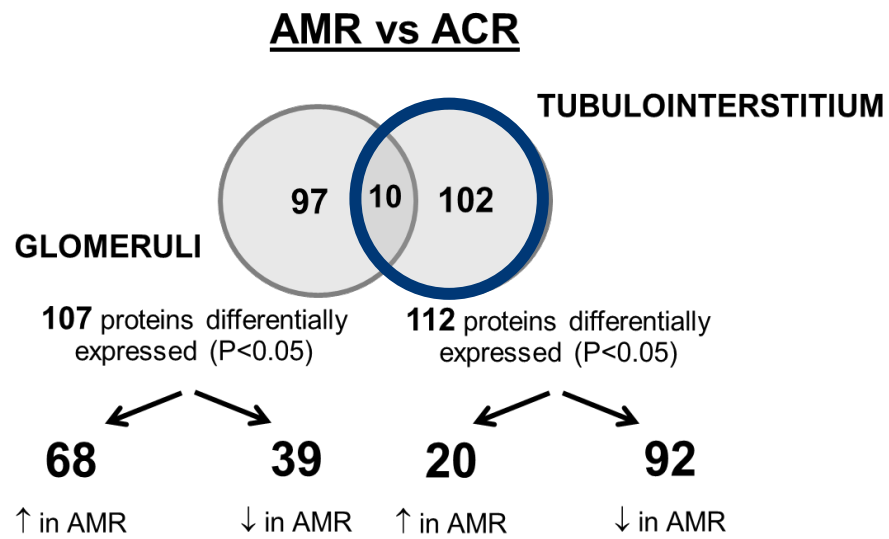


Glomerular AMR: Down-regulated proteins



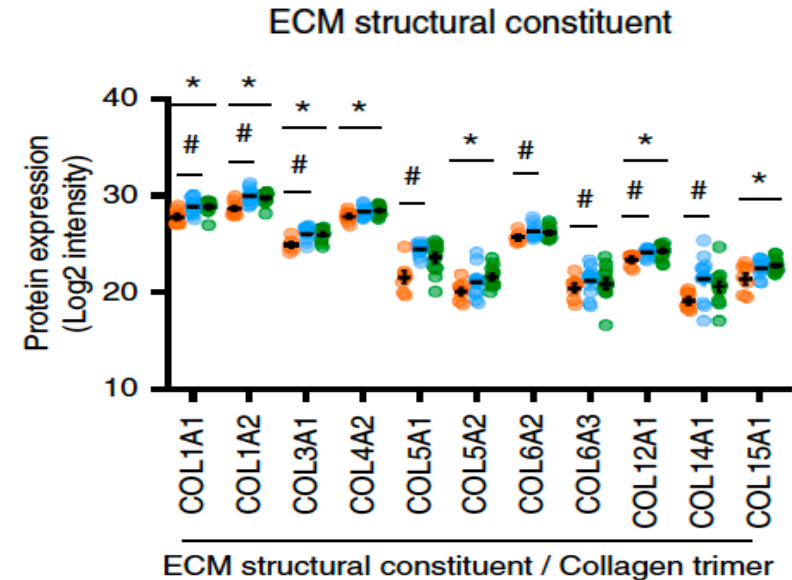
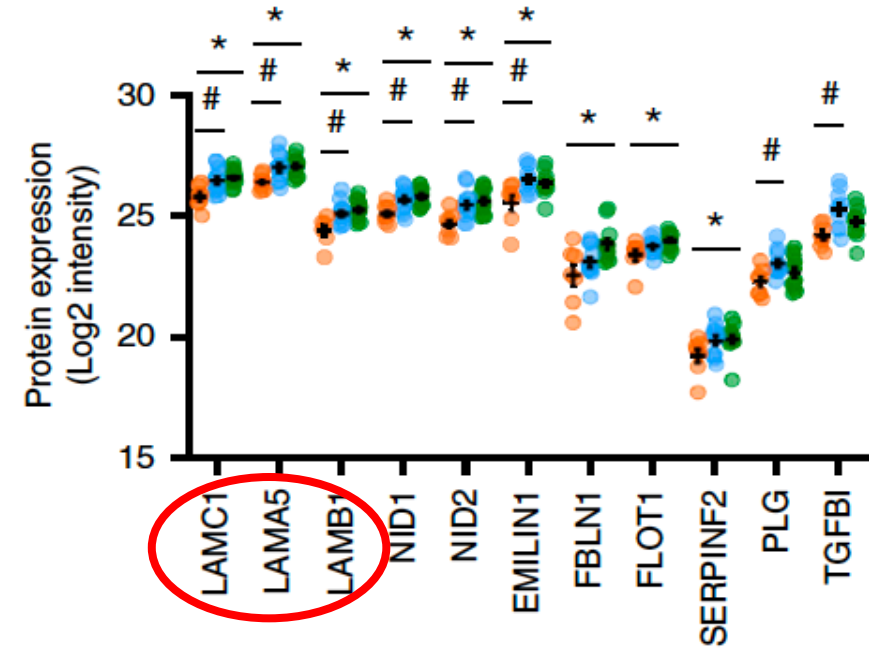
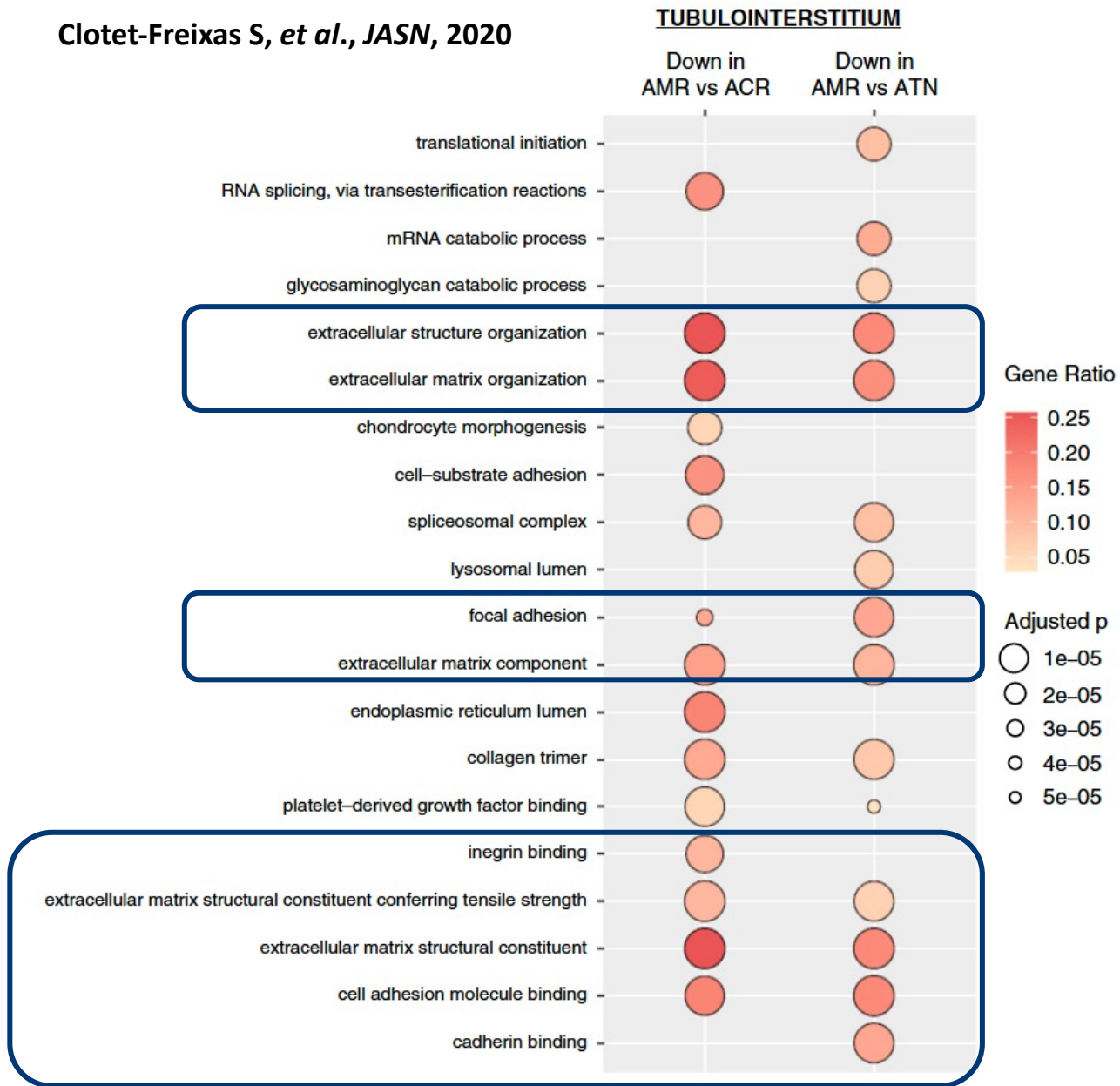
Clotet-Freixas S, et al., JASN, 2020

DIFFERENTIALLY EXPRESSED PROTEINS

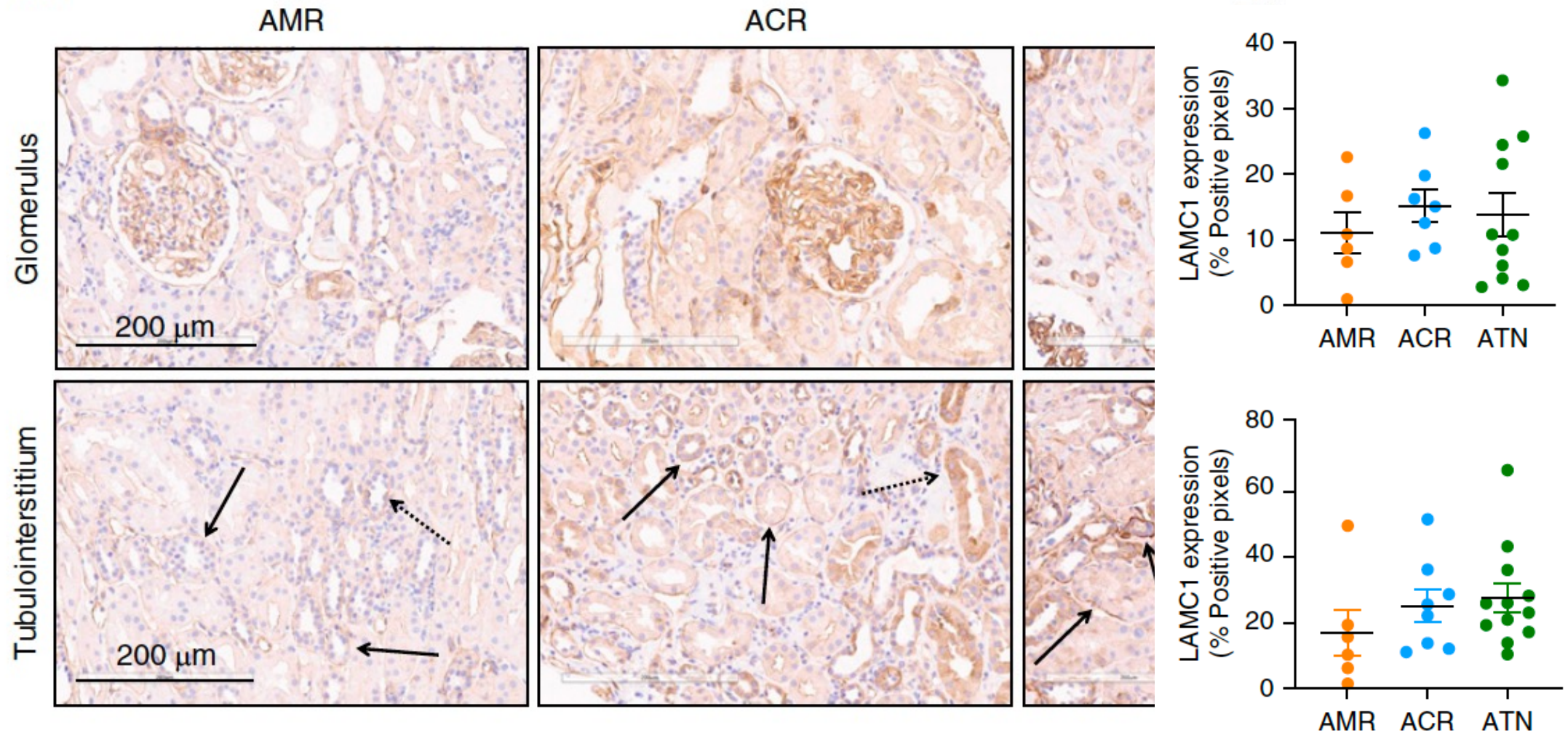


Tubulointerstitial AMR: Down-regulated proteins

Clotet-Freixas S, *et al.*, *JASN*, 2020



LAMC1 Immunohistochemistry



Clotet-Freixas S, et al., JASN, 2020

Glomerular and tubulointerstitial proteome in AMR

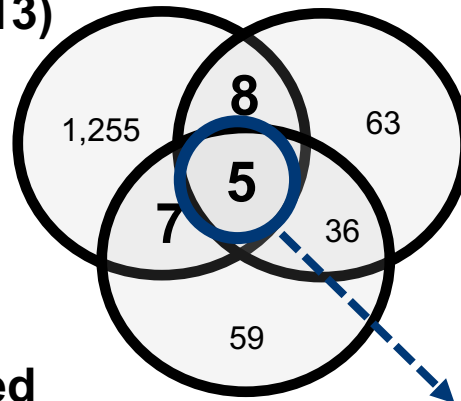
- Basement membrane and extracellular matrix proteins were significantly decreased in both compartments, even in the absence of histological lesions consistent with extracellular matrix remodeling
- Some proteins (e.g. LAMC1) were common to both compartments, while others were compartment specific

Glomerular AMR: External data set analysis

1,275 genes differentially expressed in AMR biopsies (Sellaes et al. *Am J Transplant* 2013)

112 proteins differentially expressed in AMR vs ATN Glomeruli

107 proteins differentially expressed in AMR vs ACR Glomeruli



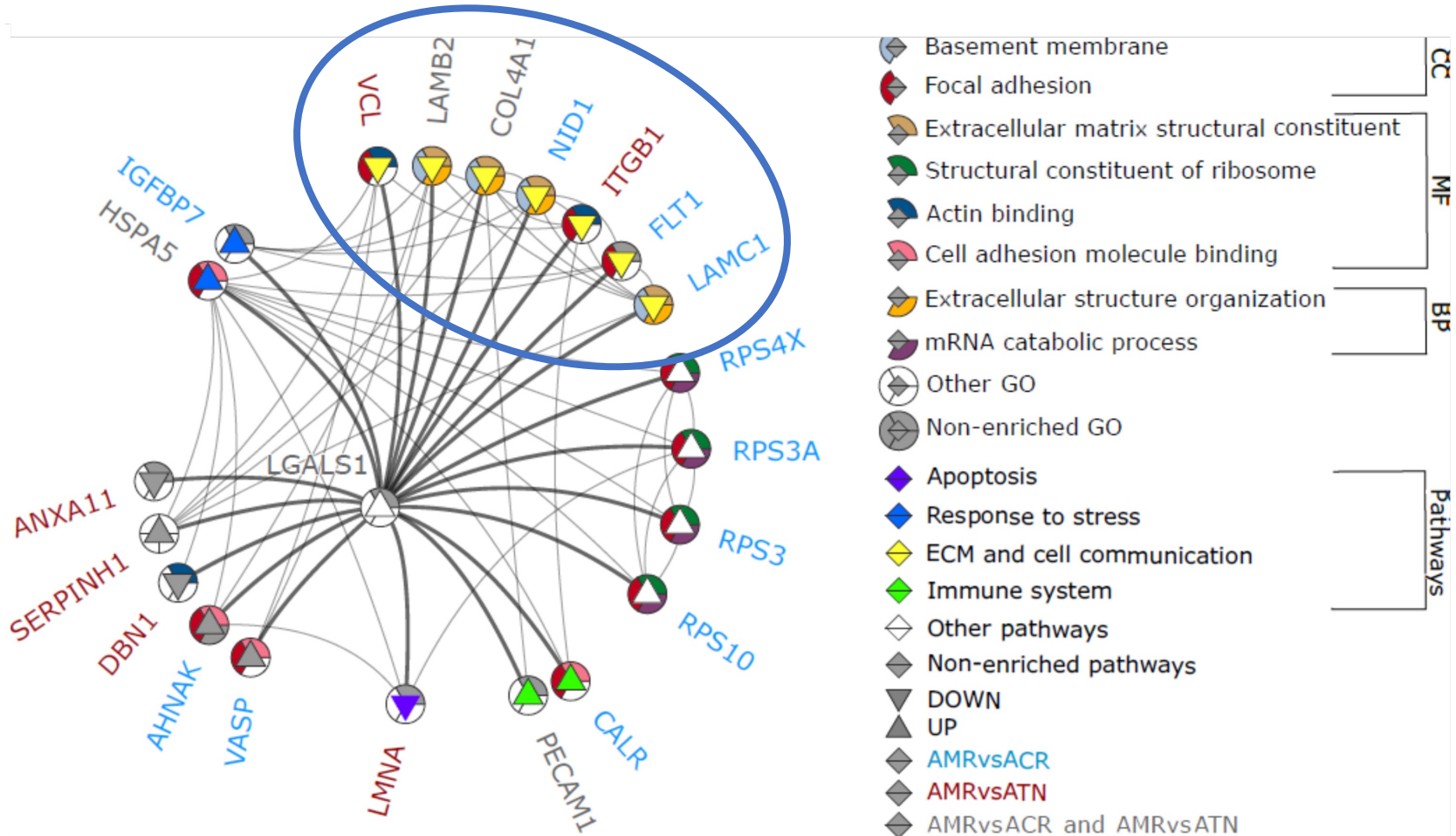
Galectin 1 (LGALS1) → UP in AMR

Clotet-Freixas S, et al., *JASN*, 2020

Galectin-1 (LGALS1) in glomerular AMR

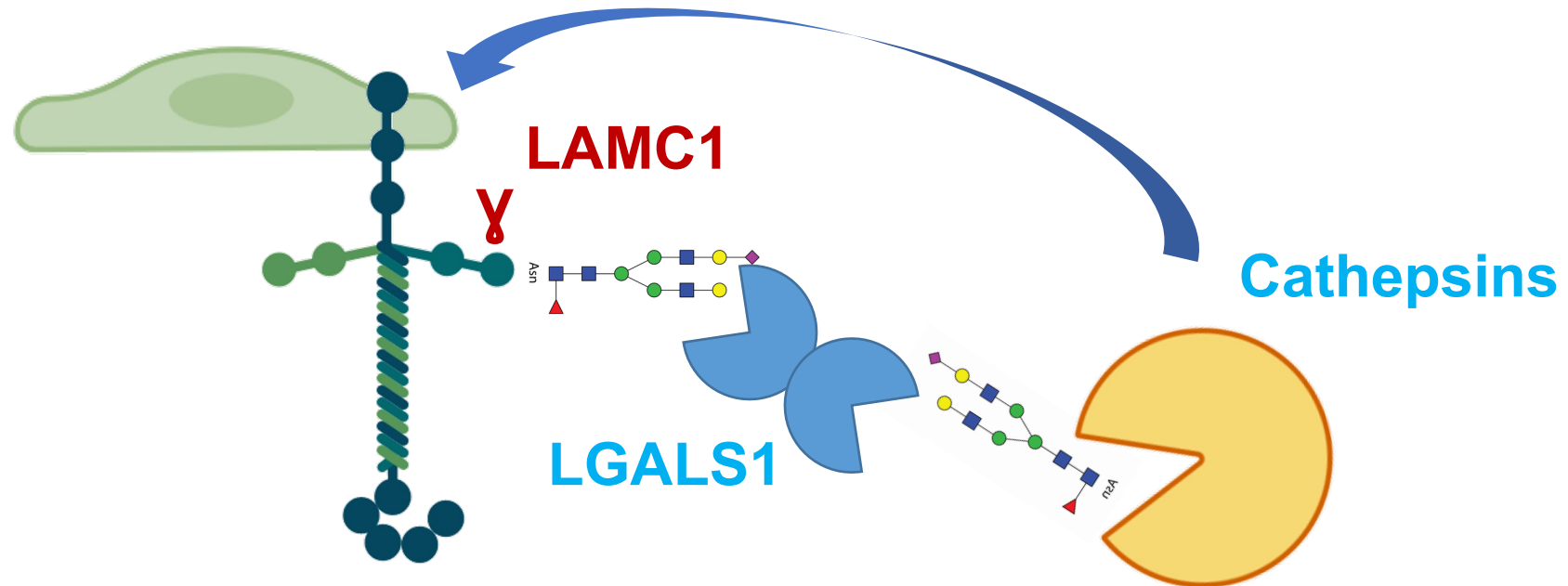


Dr. Igor Jurisica



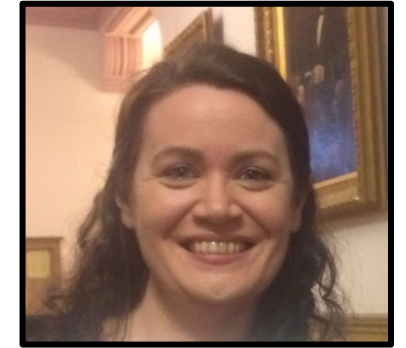
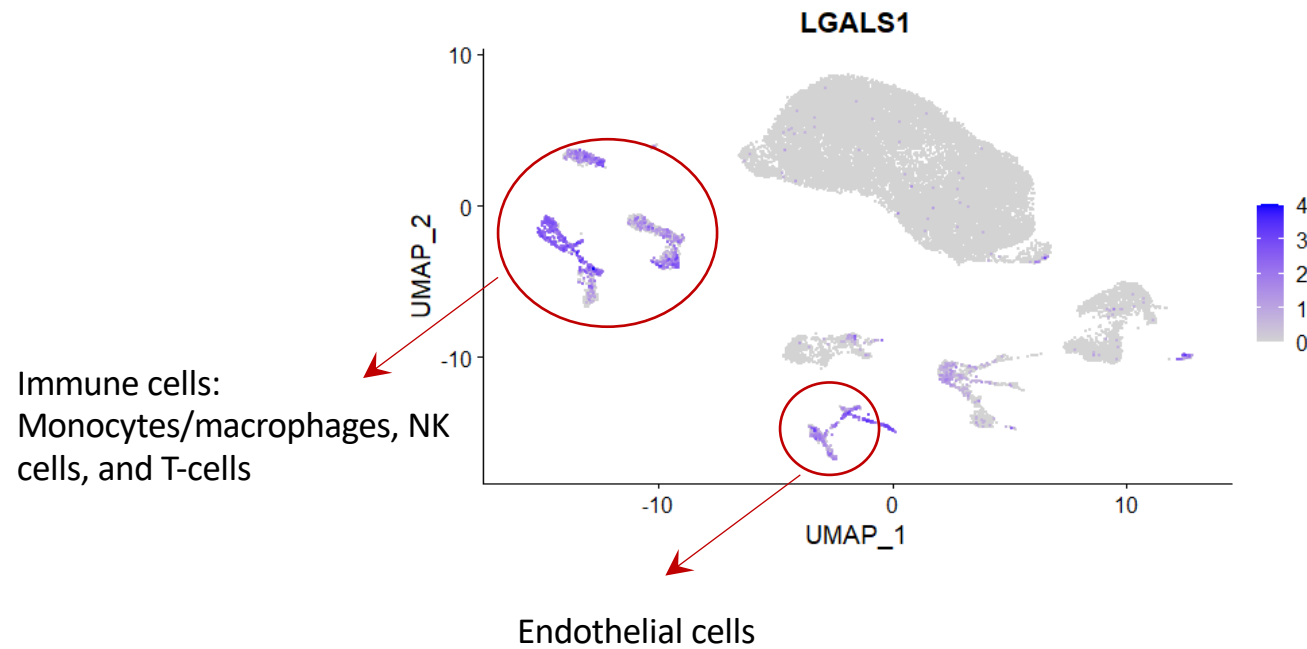
LGALS1: An interesting protein

- **Interacts with the ECM** by recognizing β -galactose sugar moieties of proteins such as laminin and fibronectin (Ilarregui, *et al.*, *Annals of the Rheumatic Diseases*, 2005).
- **Interacts with cathepsins**, which in turn induce its production in endothelial cells (Pranjol, *et al.*, *J Transl Med*, 2019).



- **Remodels endothelium** to prevent migration of T cells into the tumor microenvironment (Nambiar, *et al.*, *J Clin Invest*, 2019 and He *et al.* *Lab Invest* 2006).

Where is LGALS1 Expressed in the Healthy Human Kidney?



Dr. Caitriona McEvoy



Julia Murphy



Dr. Sarah Crome

Data from:

McEvoy, Murphy,....., Konvalinka*, Crome*, *Nature Communications* 2022

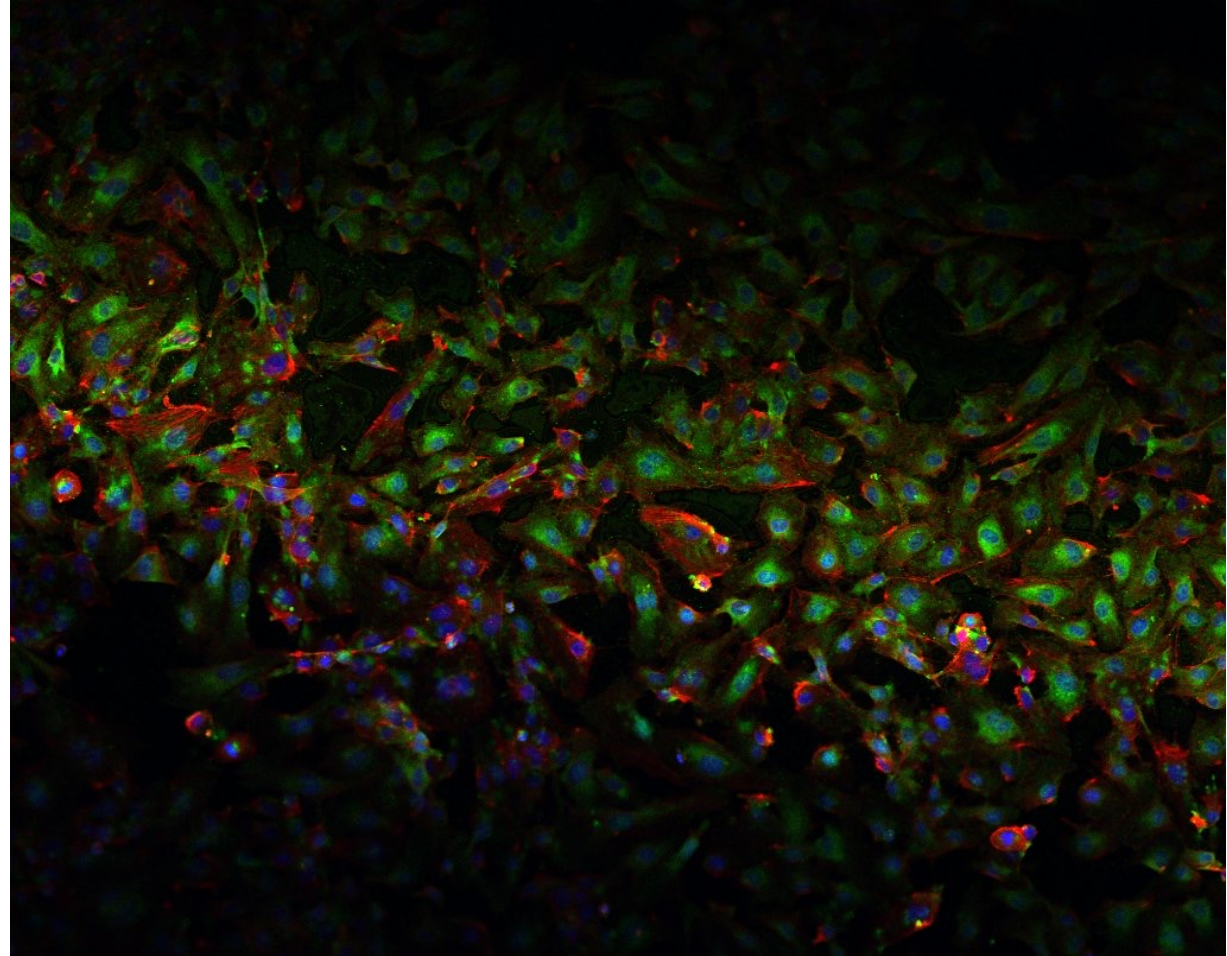
LGALS1 protein expression in kidney endothelial cells



Alex Boshart
PhD student

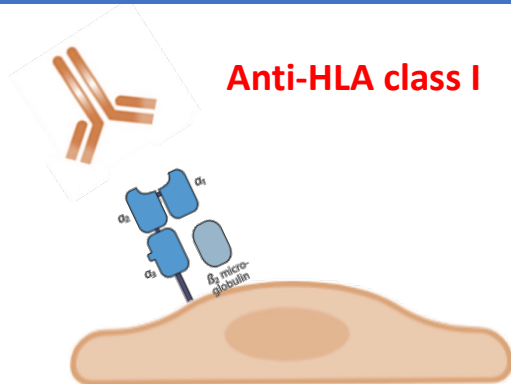
Legend:

-  DAPI
-  Actin
-  LGALS1

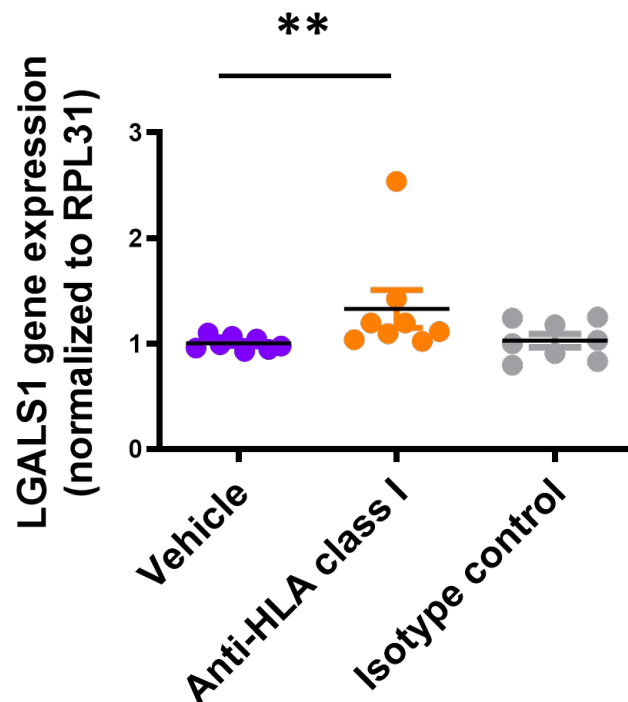
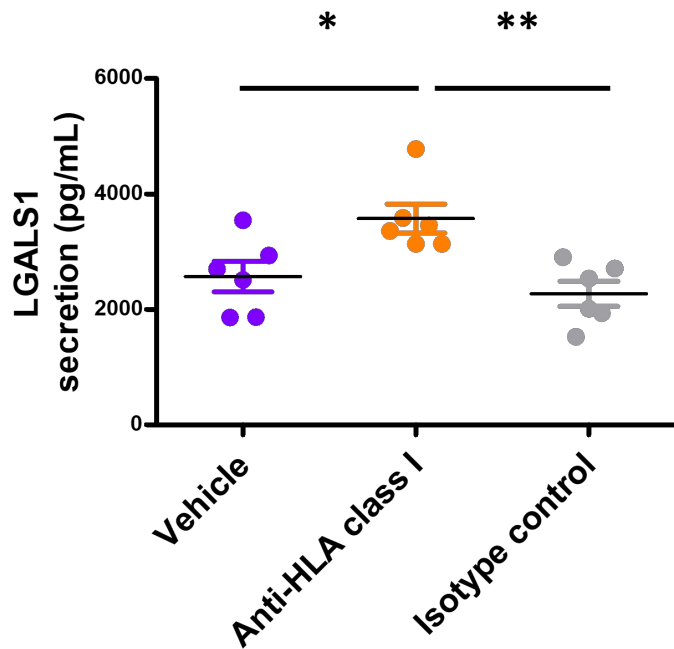


LGALS1 in glomerular AMR

Primary Human Glomerular Microvascular Endothelial Cells (HGMECs)



Alex is now knocking down LGALS1 in these cells and studying their phenotype



*p < 0.05 **p < 0.01

Clotet-Freixas S, et al., JASN, 2020

STORY #2

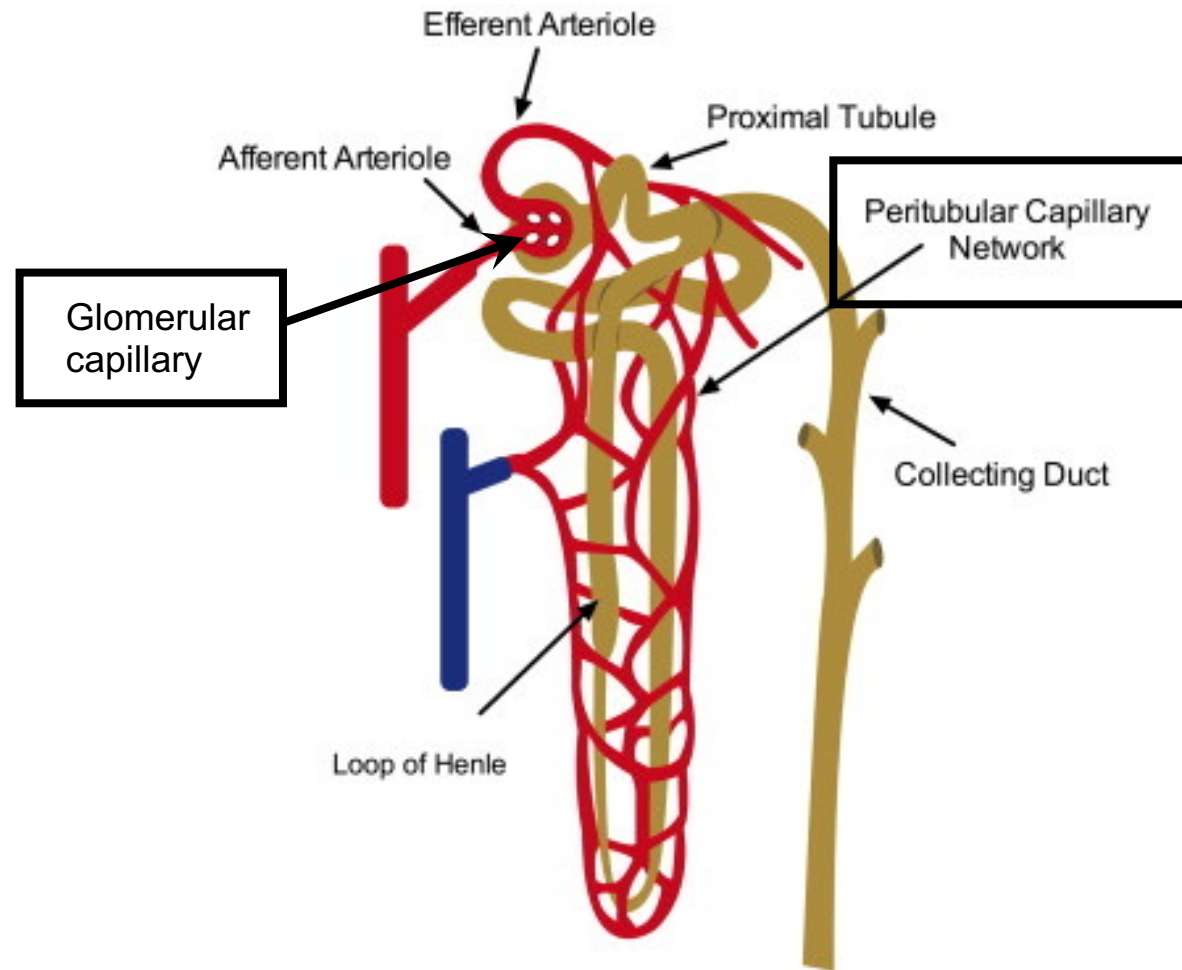
Novel Models to Study AMR in the Kidney Allograft



Dr. Boyang Zhang



Dr. Milica Radisic

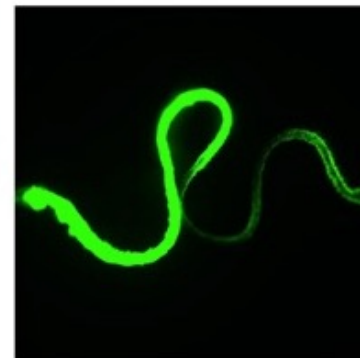
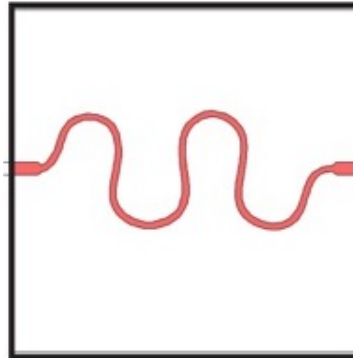
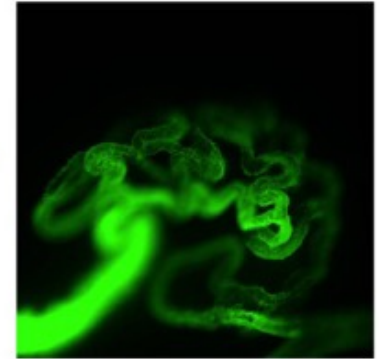
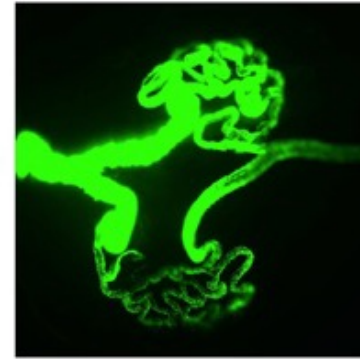
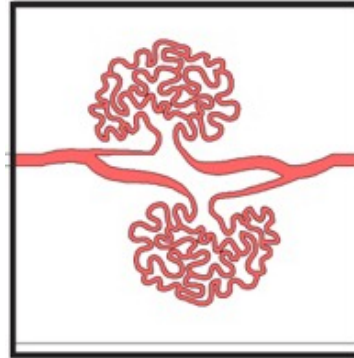
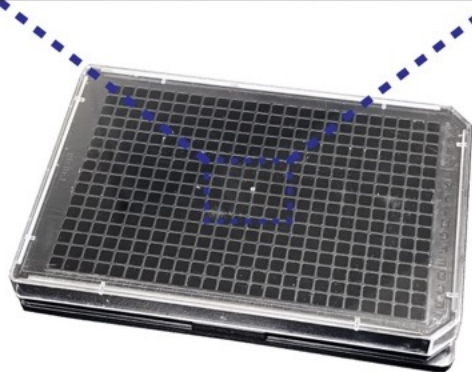
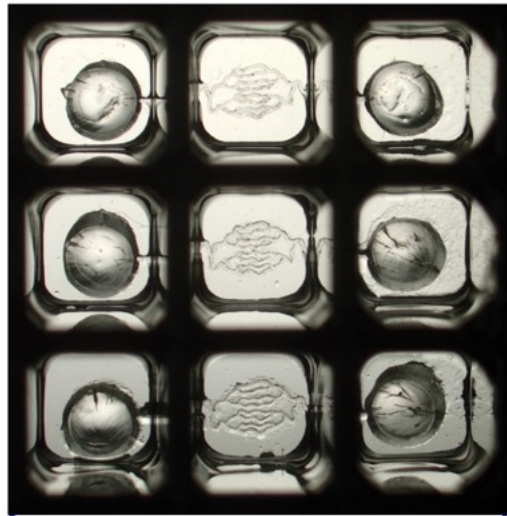


DEVELOPMENT OF NEW MODELS TO STUDY AMR

Kidney-on-a-chip



Dr. Boyang Zhang



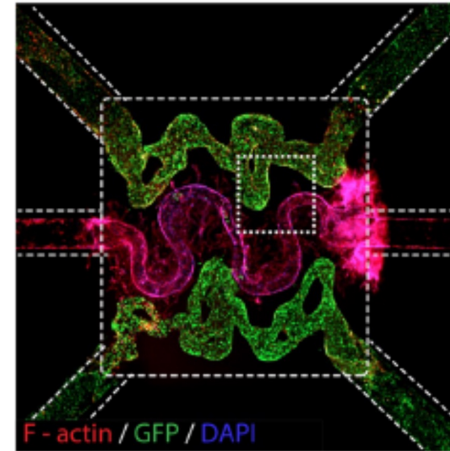
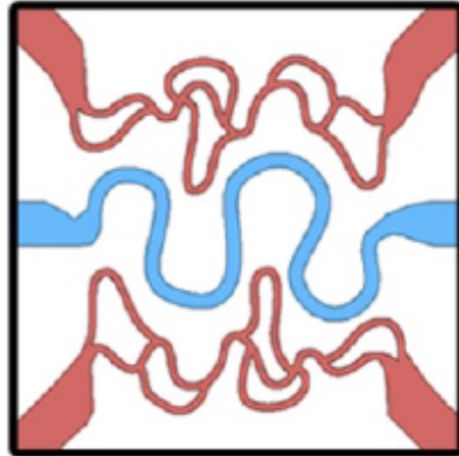
Rajasekar, et al., *Lab on a Chip*, 2022

PERITUBULAR CAPILLARY WITH PROXIMAL TUBULE

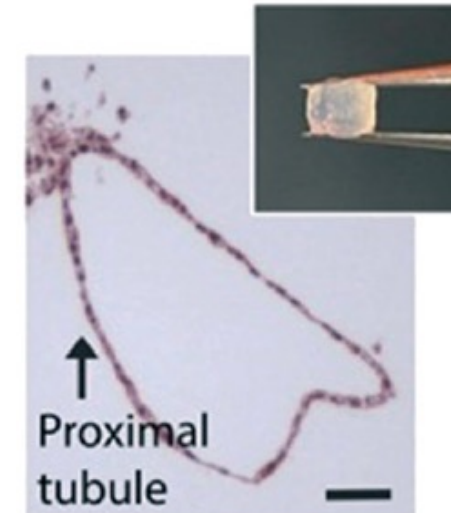
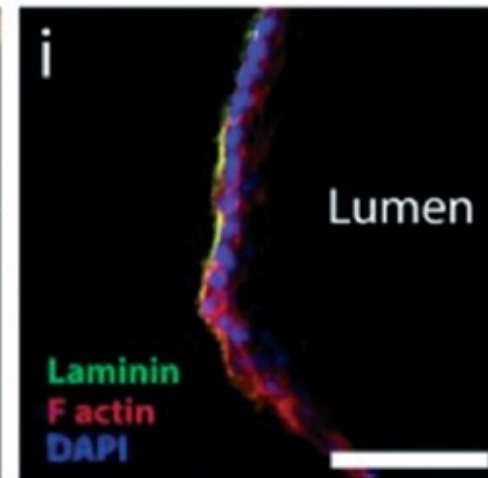
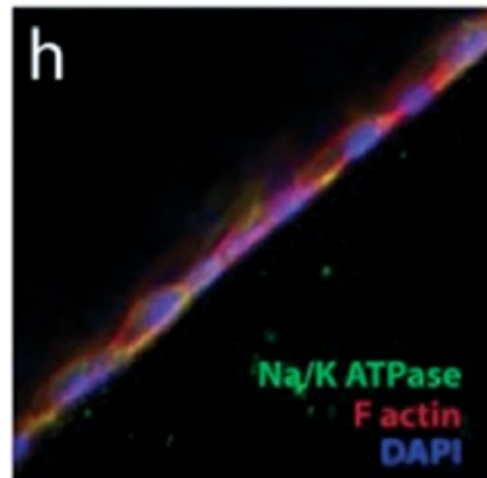
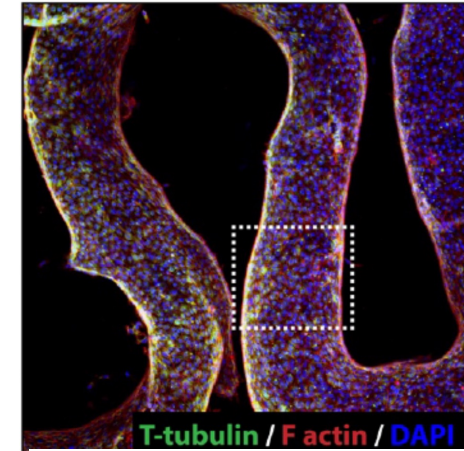


Dr. Boyang Zhang

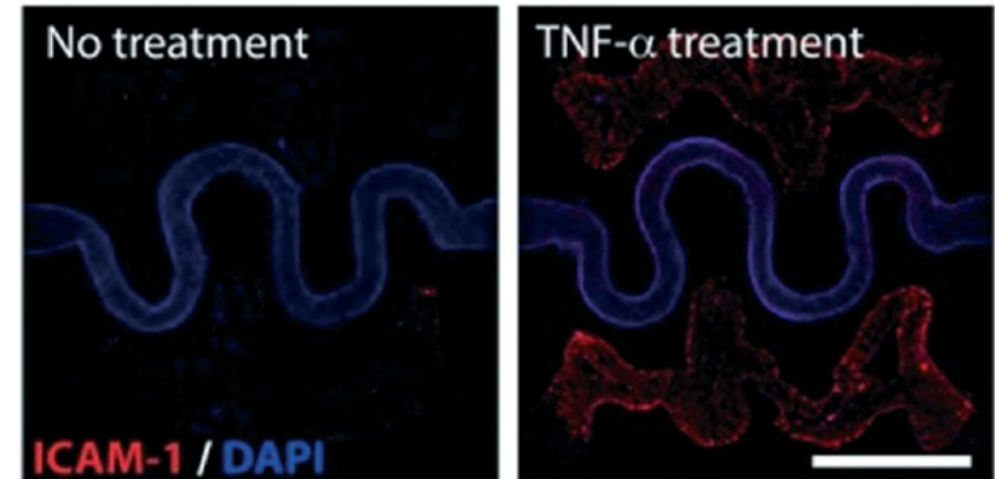
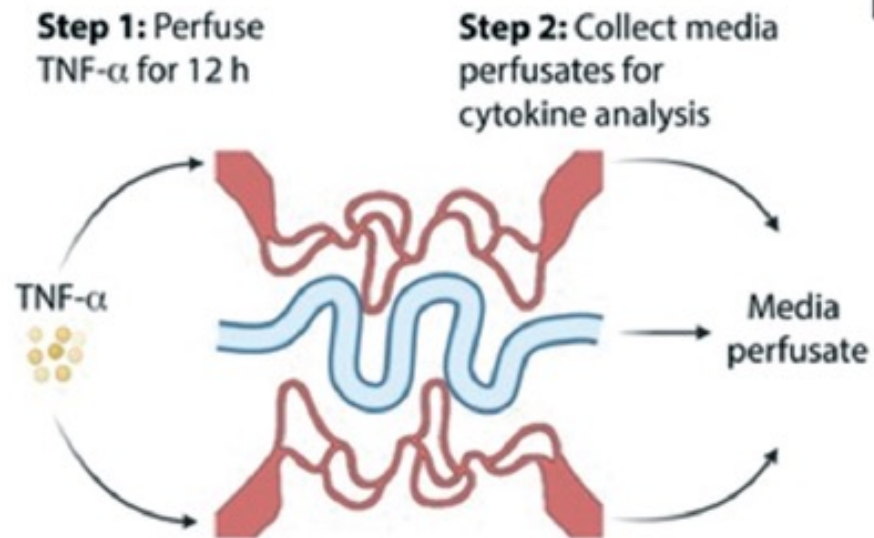
Kidney micro-capillary model



Rajasekar, *et al.*, *Lab on a Chip*, 2022



PERITUBULAR CAPILLARY WITH PROXIMAL TUBULE



Rajasekar, *et al.*, *Lab on a Chip*, 2022

CONCLUSIONS

- We have identified common and compartment-specific changes in the basement membrane proteins in the absence of transplant glomerulopathy
- LGALS1 may be a potential therapeutic target as it communicates with the basement membrane proteins and immune cells and can modify inflammation
- Novel models to study AMR in peritubular or glomerular capillary bed will enable us to integrate immune cells with endothelial cells and to perform drug screening

Gaps in knowledge

- The role of kidney tissue in antibody mediated rejection is poorly understood
- The most important non-HLA antibodies to be included in routine clinical monitoring are unknown
- Determination of pathogenic DSA is elusive
- Effective therapies are lacking

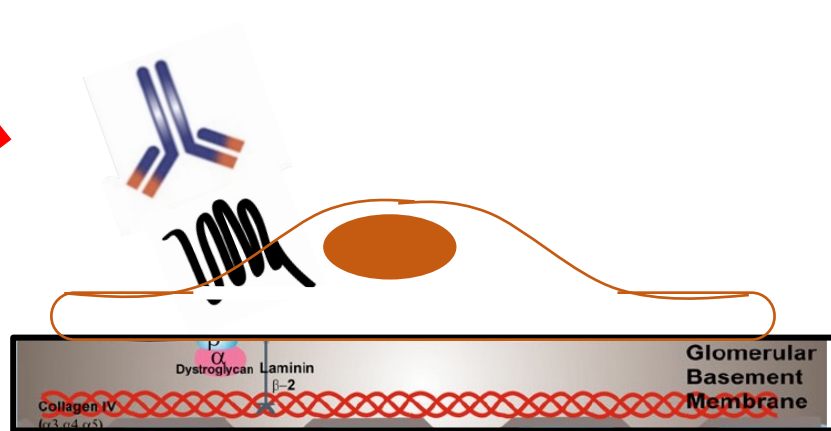
Non-HLA Antibodies

AT-1 receptor agonistic antibodies

Dragun et al. *NEJM* 2005

LG3 perlecan fragment antibodies

Cardinal et al. *AJT* 2013, Hebert *Circ Res* 2012



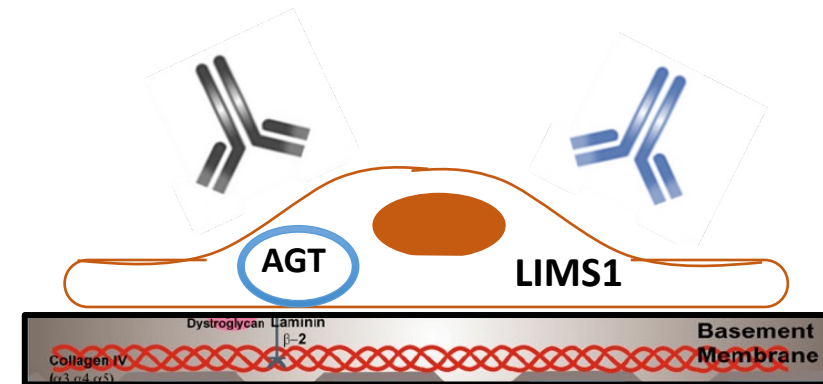
Donor **endothelial** cell

Anti-Angiotensinogen Abs

Li et al. *J Proteome Res* 2010

Anti-LIMS1 Abs in kidney allograft rejection

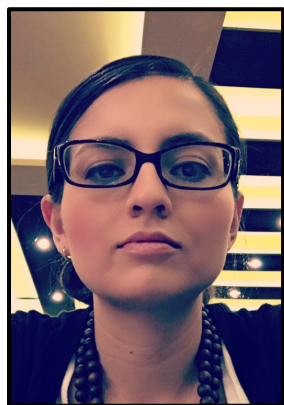
Steers et al. *N Engl J Med* 2019



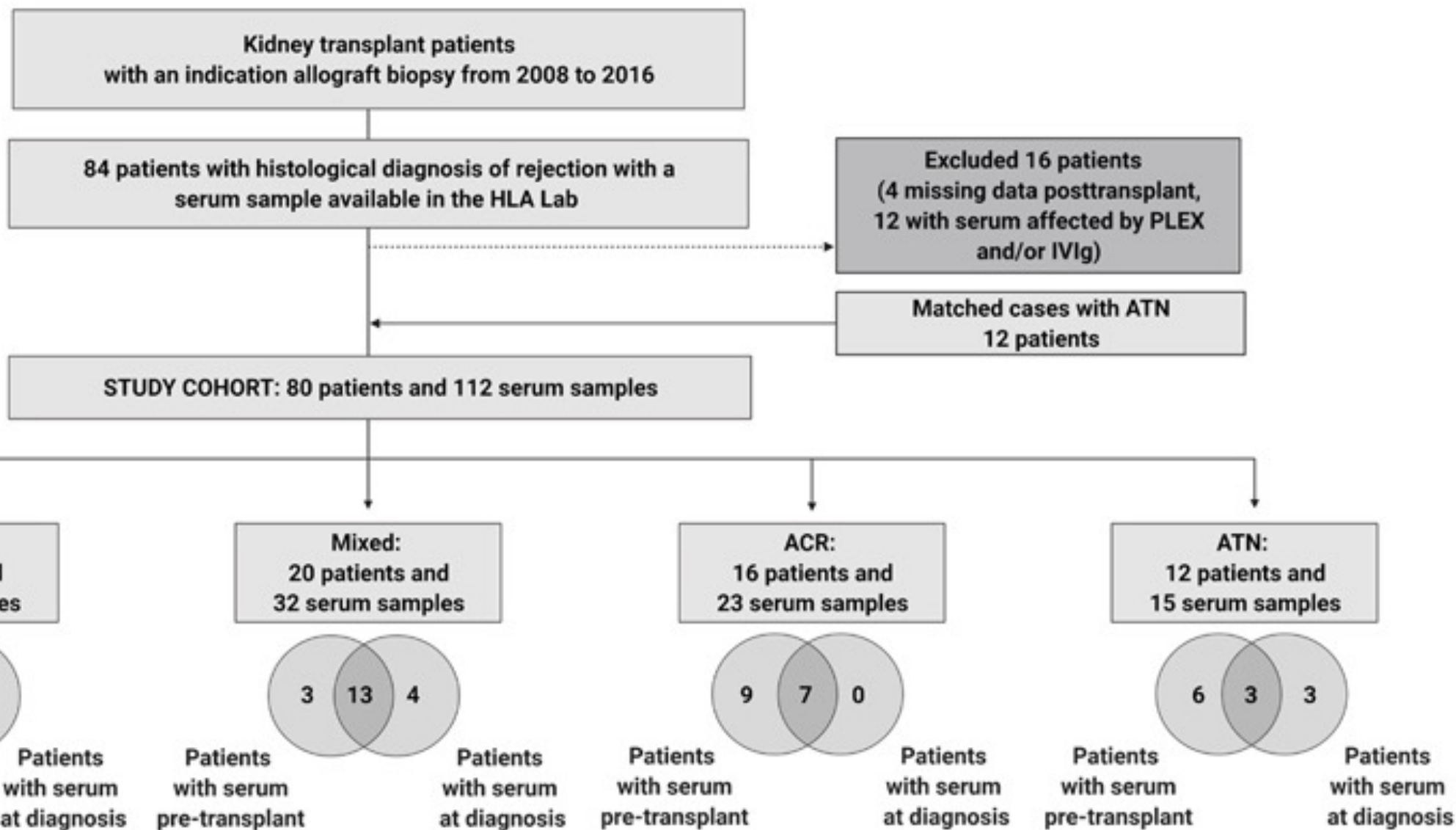
Donor **epithelial** cell

Story # 3

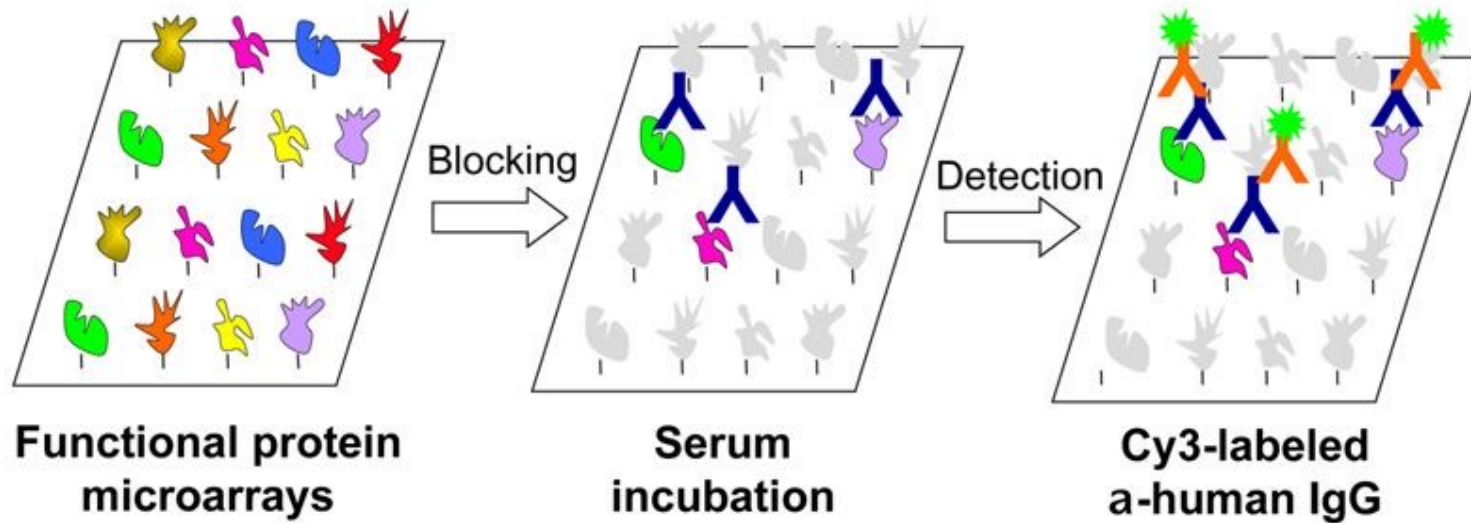
Assess circulating non-HLA antibodies in patients with AMR compared to other forms of graft injury



Dr. Sonia Rodriguez Ramirez
Post-doctoral fellow



Measuring non-HLA antibodies

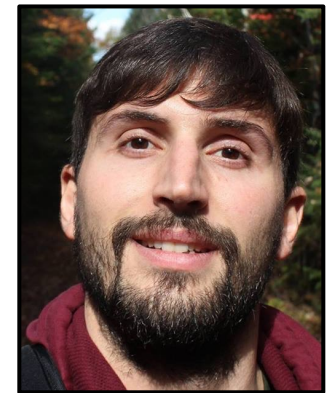


Zhu, H.; *et al.* Applications of functional protein microarrays in basic and clinical research. *Adv Genet.* 2012, 79: 123-155

Designed protein microarrays to measure antibodies against 134 antigens implicated in autoimmune disease or solid organ transplant rejection



Dr. Andrzej Chruscinski



Dr. Sergi Clotet-Freixas
Post-doctoral fellow

RESULTS



Dr. Max Kotlyar

Antibody name	Antigen specificity	Antibody levels		
		Before transplant		
		AMR/mixed, median (IQR)	ACR, median (IQR)	P
IgG M2	M2 (PDC-E2 + OGDC-E2 + BCOADC-E2)	267 (0–513)	0 (0–0)	0.0051
→ IgG CENP-B	Major centromere autoantigen B	312.5 (0–877.2)	0 (0–235.2)	0.0097
→ IgG Ro/SS-A (52 kDa)	Ro/SS-A (52 kDa)	315 (0–915.7)	0 (0–257.5)	0.0112
IgG gliadin	Gliadin	3209 (1485.8–5474)	1618.5 (793–3110.6)	0.0157
→ IgG PDH	Pyruvate dehydrogenase	238 (0–507)	0 (0–226)	0.0232
IgG smooth muscle	Smooth muscle actin	382.5 (304.5–532.8)	310.8 (230.5–374.1)	0.0457
		AMR/mixed, median (IQR)	ATN, median (IQR)	P
IgM PL-12	Alanyl-tRNA synthetase	1049.5 (544.2–1672.8)	352 (236.5–601)	0.0074
IgM HGMEC lysate	Glomerular endothelial cells	1393.5 (881–2234.5)	752 (522–1131.5)	0.0224
IgM PM/Sci-100	PM/Sci-100	317.5 (0–451.5)	0 (0–0)	0.0234
IgM OGDC-E2	M2 (OGDC)	426 (104.8–766)	0 (0–307.5)	0.0298
IgG HCEC cytoplasm	Cardiac endothelial cells	380.5 (229.2–532.7)	222 (0–294.5)	0.0309
IgM LG3	Basement membrane-specific heparan sulfate proteoglycan core protein	402.5 (238.3–607.8)	0 (0–401)	0.0365
IgM LKM 1 hp	LKM 1 hp antigen of cytochrome P450 2D6	210.5 (0–314)	0 (0–0)	0.0393
IgG Ro/SS-A (60 kDa, R)	Ro/SS-A (60 kDa)	0 (0–233.5)	296 (0–559.5)	0.0403
IgM Sm (NR, B)	Small nuclear ribonucleoprotein Sm	311 (0–495.7)	0 (0–299.5)	0.0470
→ IgG PDH	Pyruvate dehydrogenase	238 (0–507)	0 (0–0)	0.0494

RESULTS – Sera at the time of diagnosis

Antibody name	Antigen specificity	Antibody levels		
		At diagnosis		
		AMR/mixed, median (IQR)	ACR, median (IQR)	<i>P</i>
IgG CENP-B	Major centromere autoantigen B	510 (0–877.8)	0 (0–0)	0.0126
IgG Ro/SS-A (52 kDa)	Ro/SS-A (52 kDa)	353.5 (0–1494)	0 (0–0)	0.0325
IgM La/SS-B	La/SS-B	735.5 (390.8–1843)	296 (139.5–411)	0.0261
IgM CENP-B	Major centromere autoantigen B	666.5 (313.5–1889.5)	262 (100.2–429)	0.0447
IgM PDH	PDH	251 (0–684.5)	0 (0–0)	0.0472
		AMR/mixed, median (IQR)	ATN, median (IQR)	<i>P</i>
IgG M2	M2 (PDC-E2 + OGDC-E2 + BCOADC-E2)	219 (0–596.5)	0 (0–0)	0.0313
IgG human IgA	Human IgA	750.5 (229.5–1322)	0 (0–337.5)	0.0495

Validation of top non-HLA antibodies in an external cohort

60 kidney transplant recipients from Montreal previously described with serum samples analyzed using the same platform

Cardinal et al. Am J Transplant 2013

1. Antibody mediated or Mixed rejection
2. Acute cellular rejection
3. Controls without rejection



Dr. HELOÏSE CARDINAL



Dr. Marie-Josée Hébert



Dr. MÉLANIE DIEUDÉ

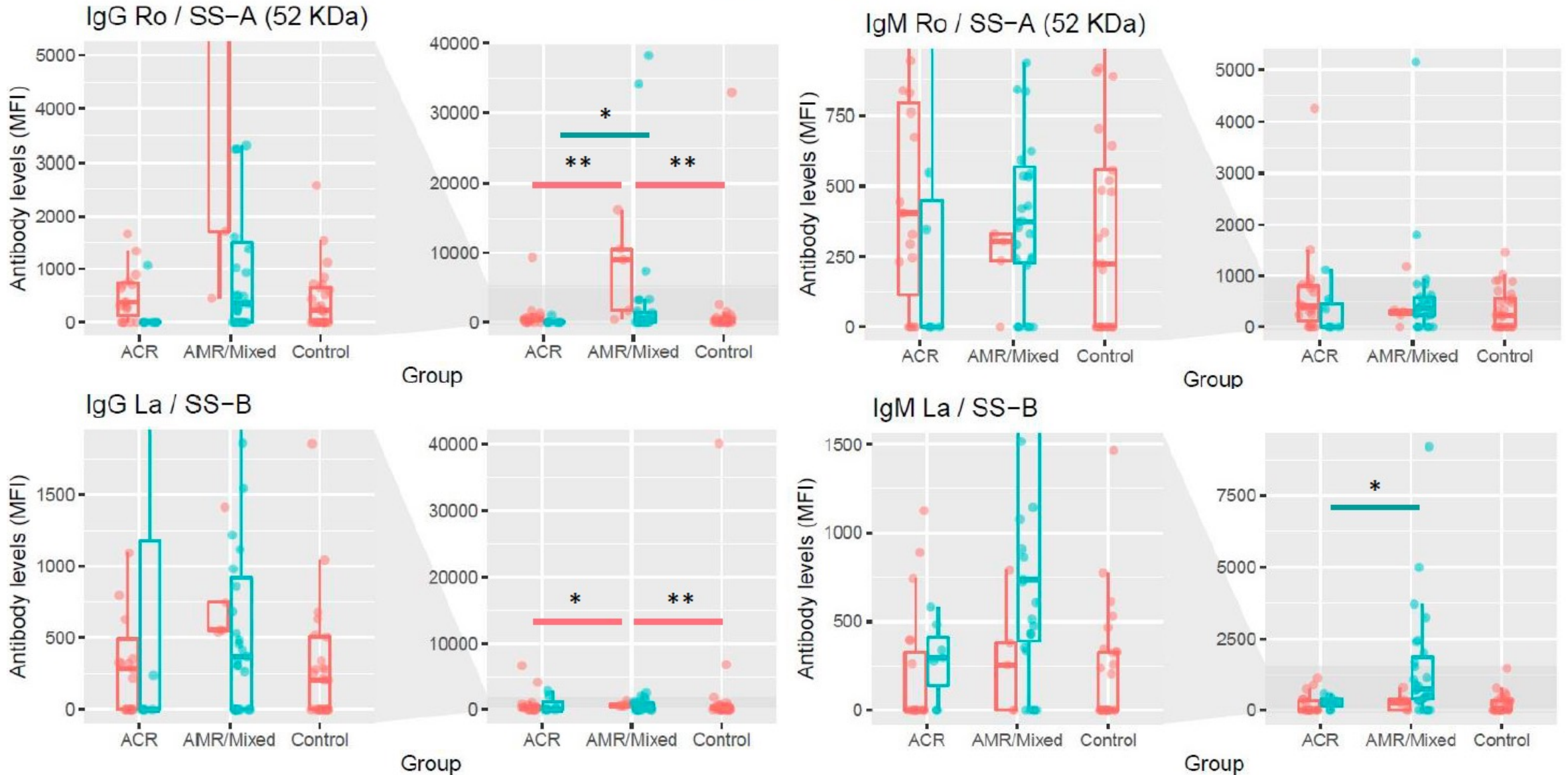
Applied the same method of non-HLA Ab measurement

Examine only antibodies most significant in our cohort:

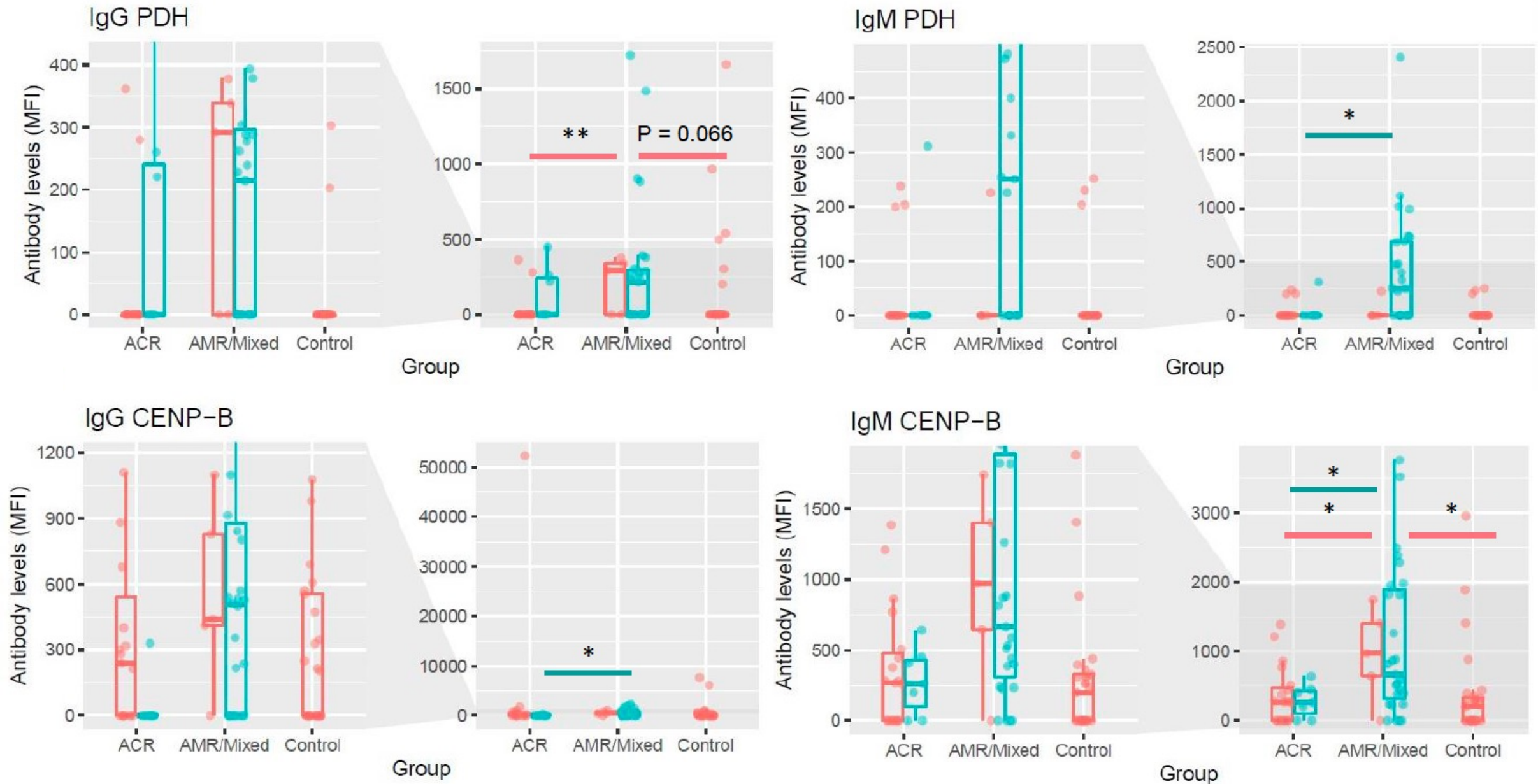
Anti-Ro, Anti-La, Anti-CENPB, Anti-PDH

Validation of Results

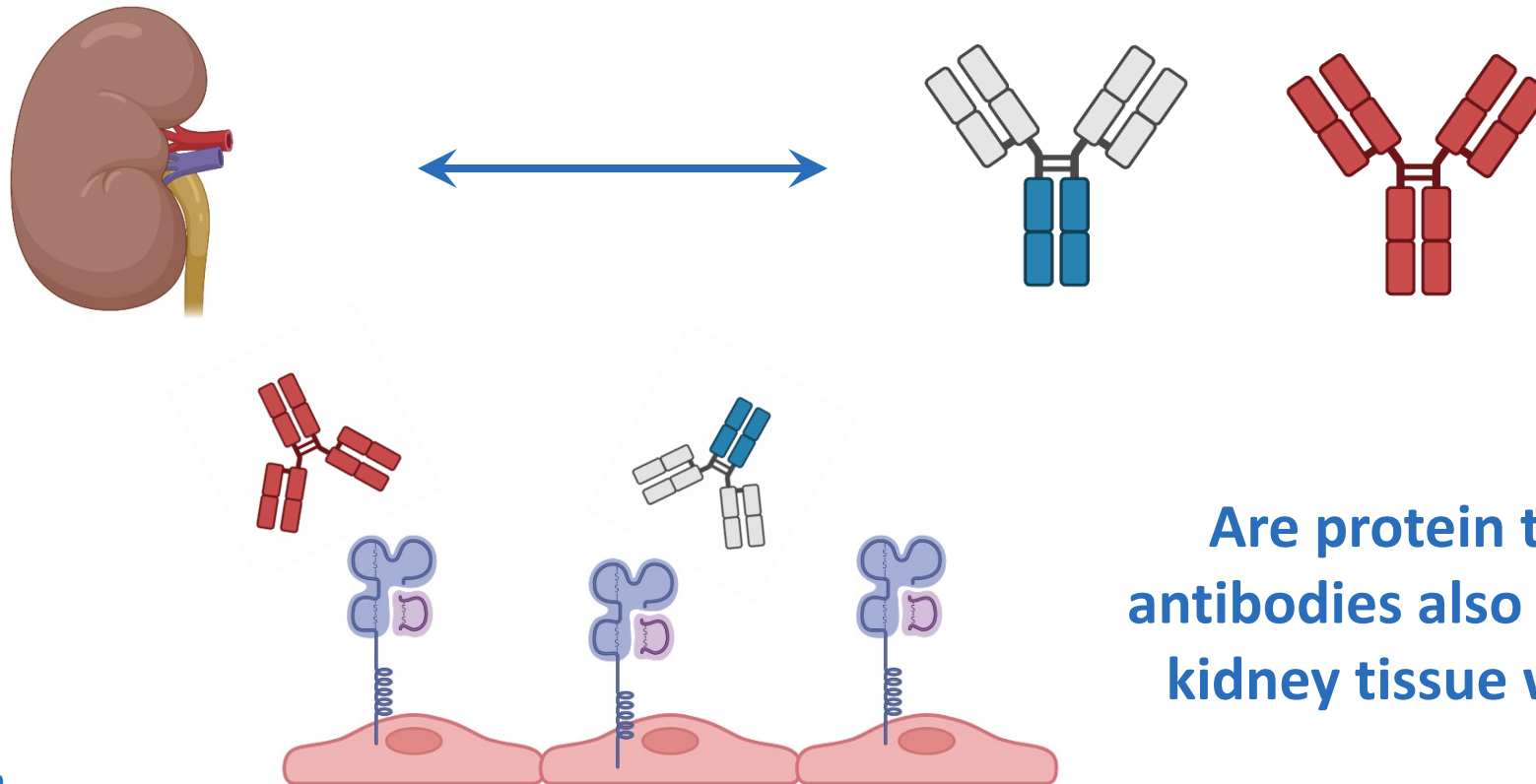
Cohort Montreal Toronto



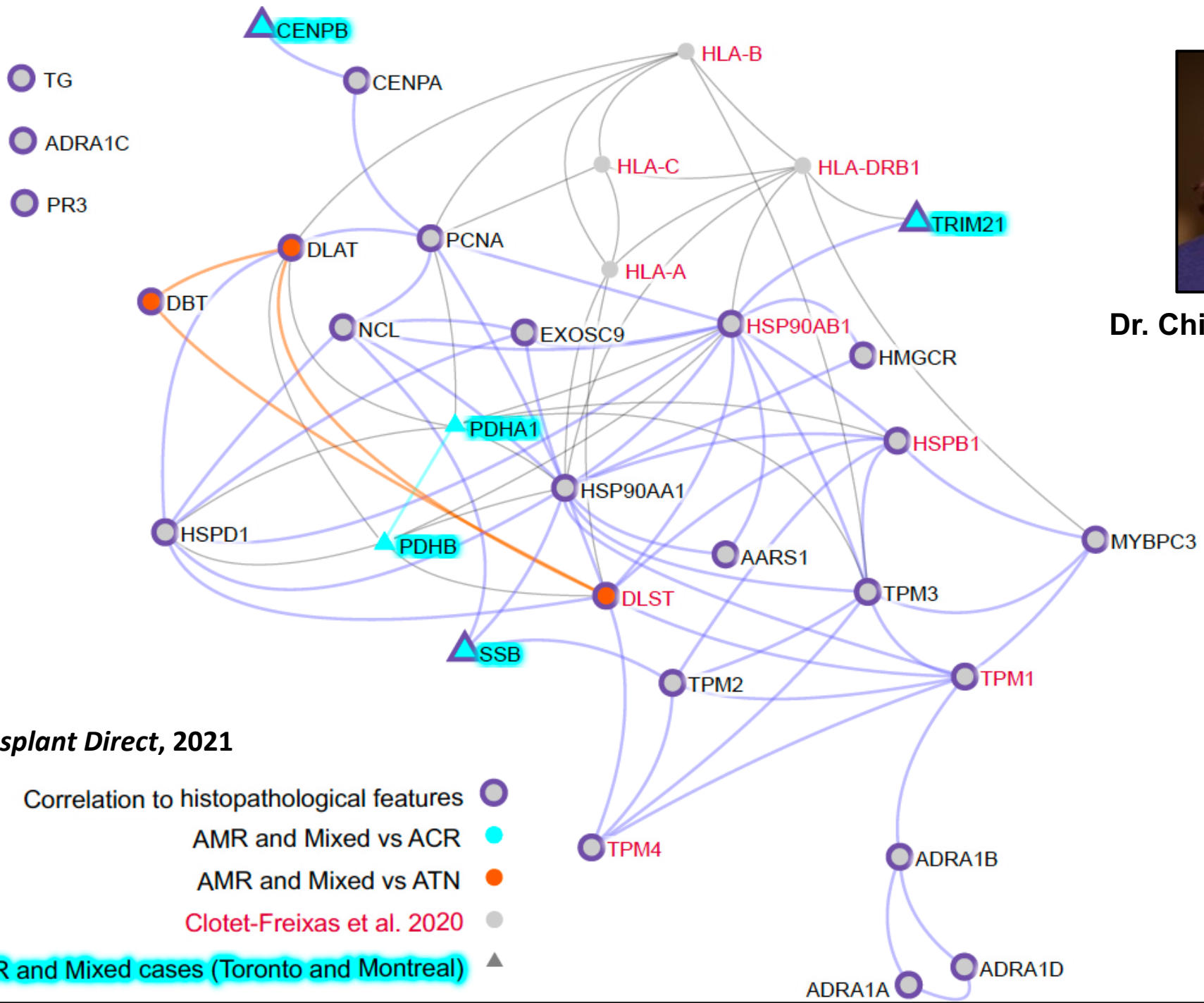
Validation of Results



Integration of circulating antibody specificity and kidney tissue protein expression



Are protein targets of antibodies also increased in kidney tissue with AMR?

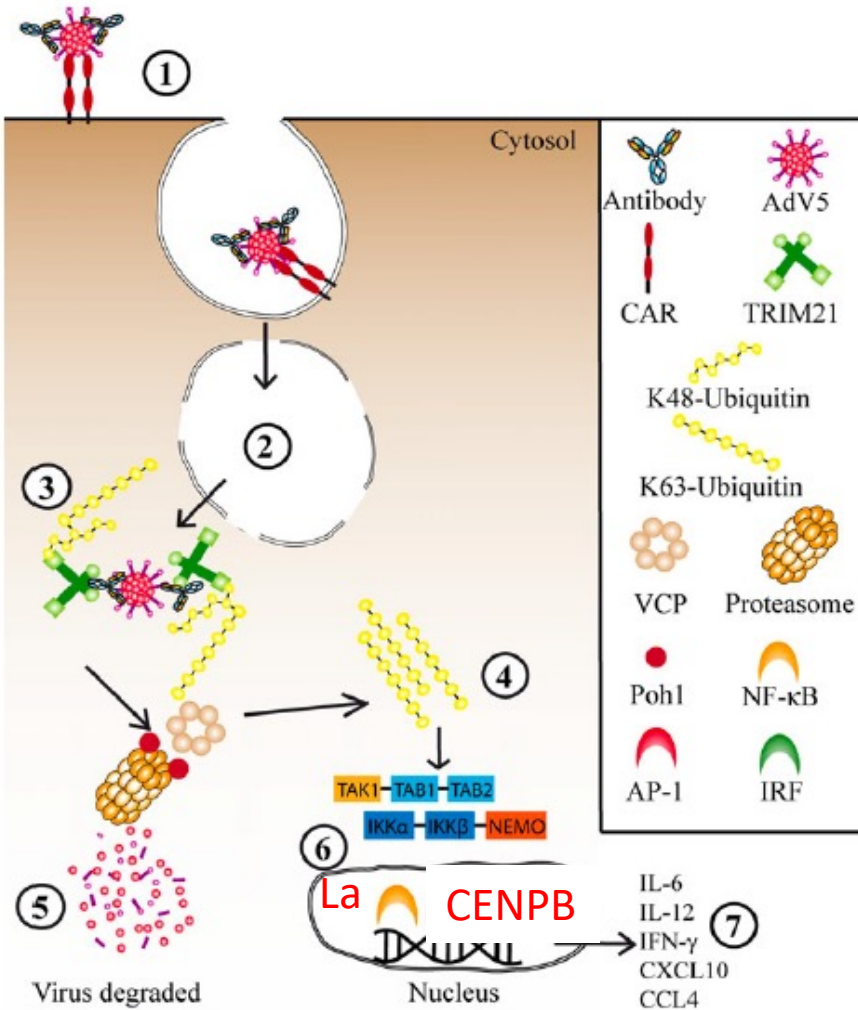


Dr. Chiara Pastrello

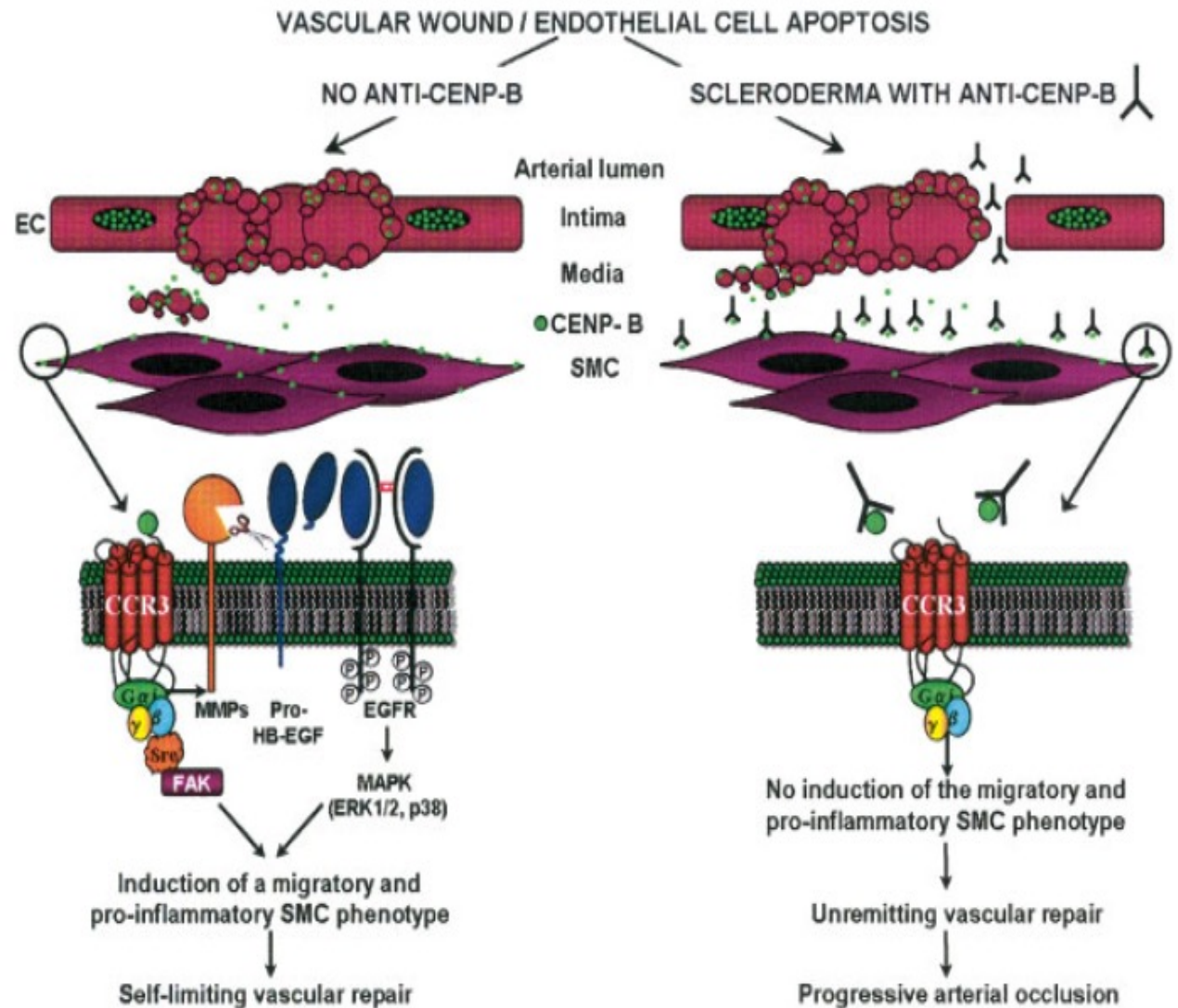
Clotet-Freixas, et al., *Transplant Direct*, 2021

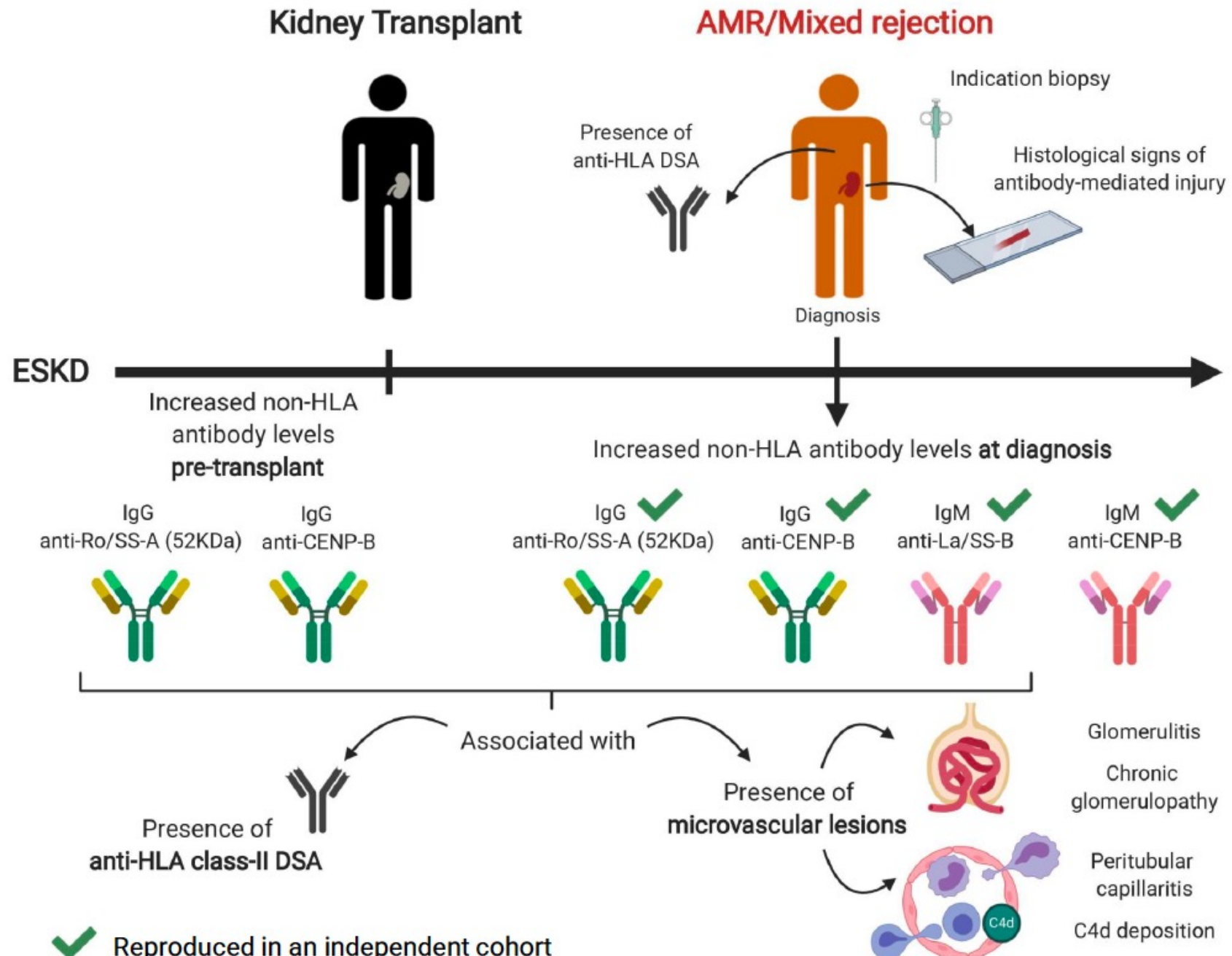
- Correlation to histopathological features
- AMR and Mixed vs ACR
- AMR and Mixed vs ATN
- Clotet-Freixas et al. 2020
- Increased in AMR and Mixed cases (Toronto and Montreal)

What do these antibodies target?



Immunological Reviews 2015
 Vol. 268: 328–339





Gaps in knowledge

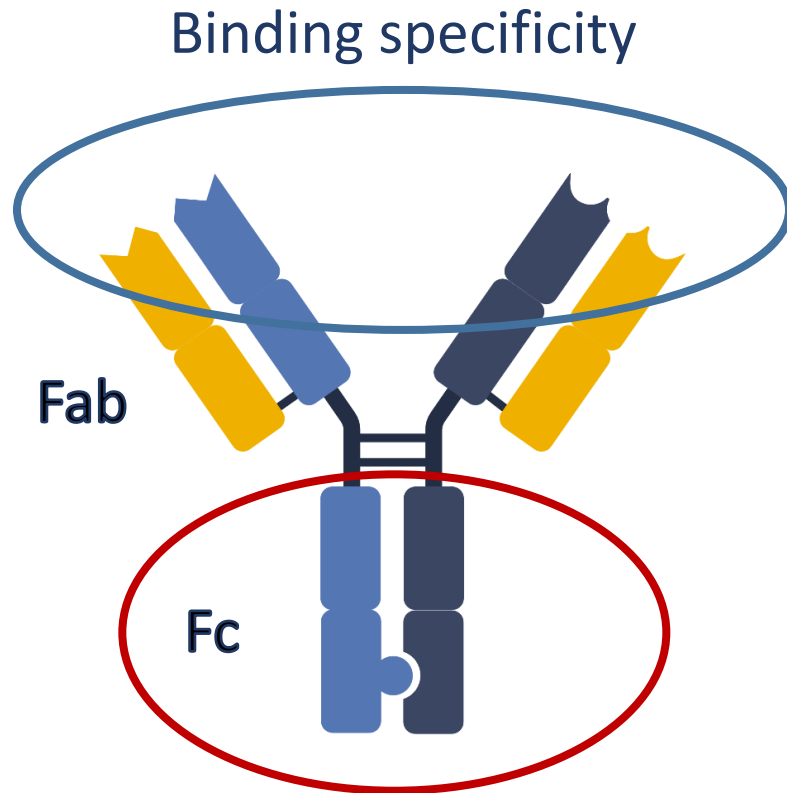
- The role of kidney tissue in antibody mediated rejection is poorly understood
- The most important non-HLA antibodies to be included in routine clinical monitoring are unknown
- **Determination of pathogenic DSA is elusive**
- Effective therapies are lacking

30-60% of patients with DSA do not develop rejection

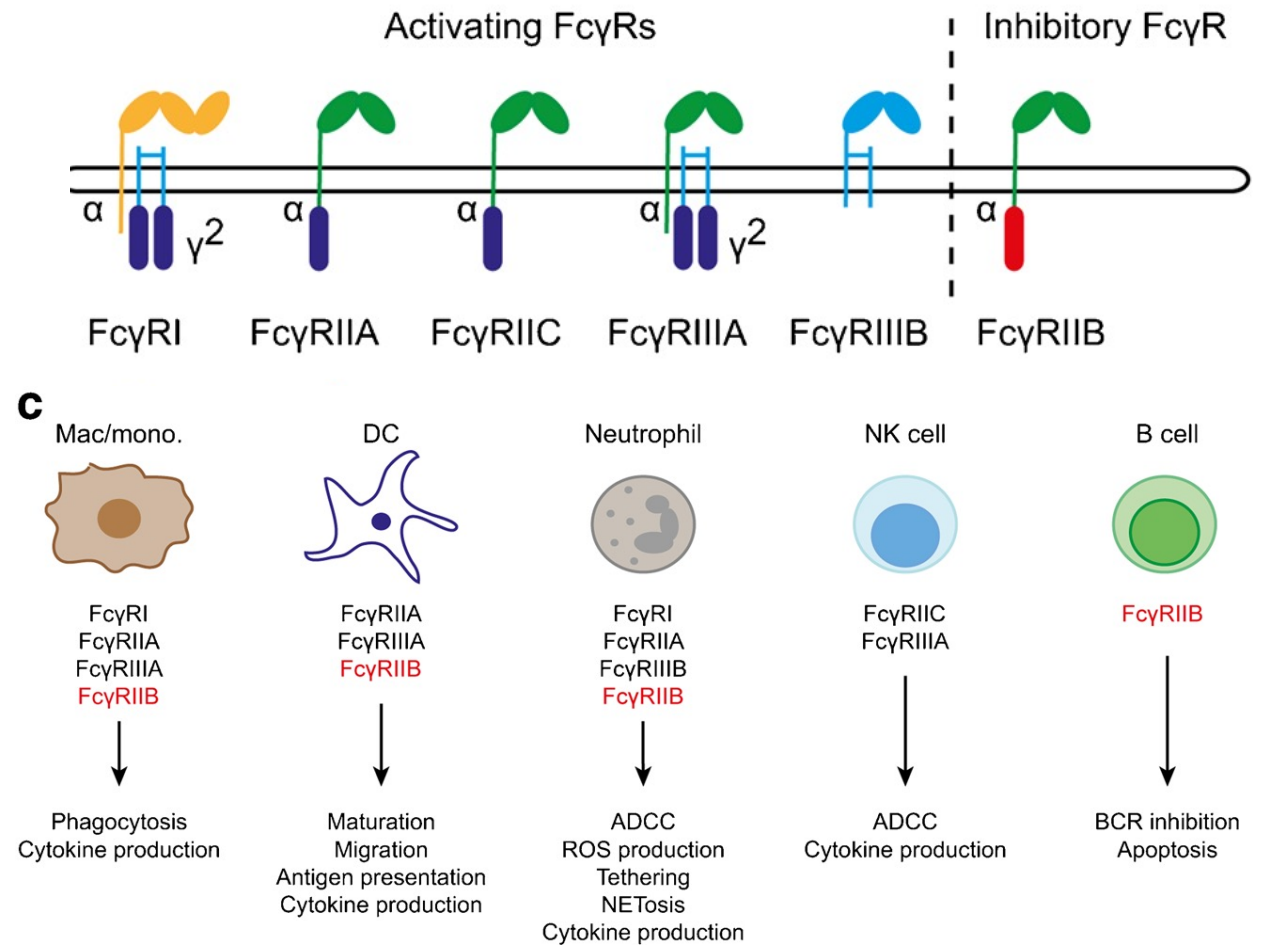
FUTURE DIRECTIONS

Decipher donor specific antibody biochemical and functional properties linked to its pathogenicity

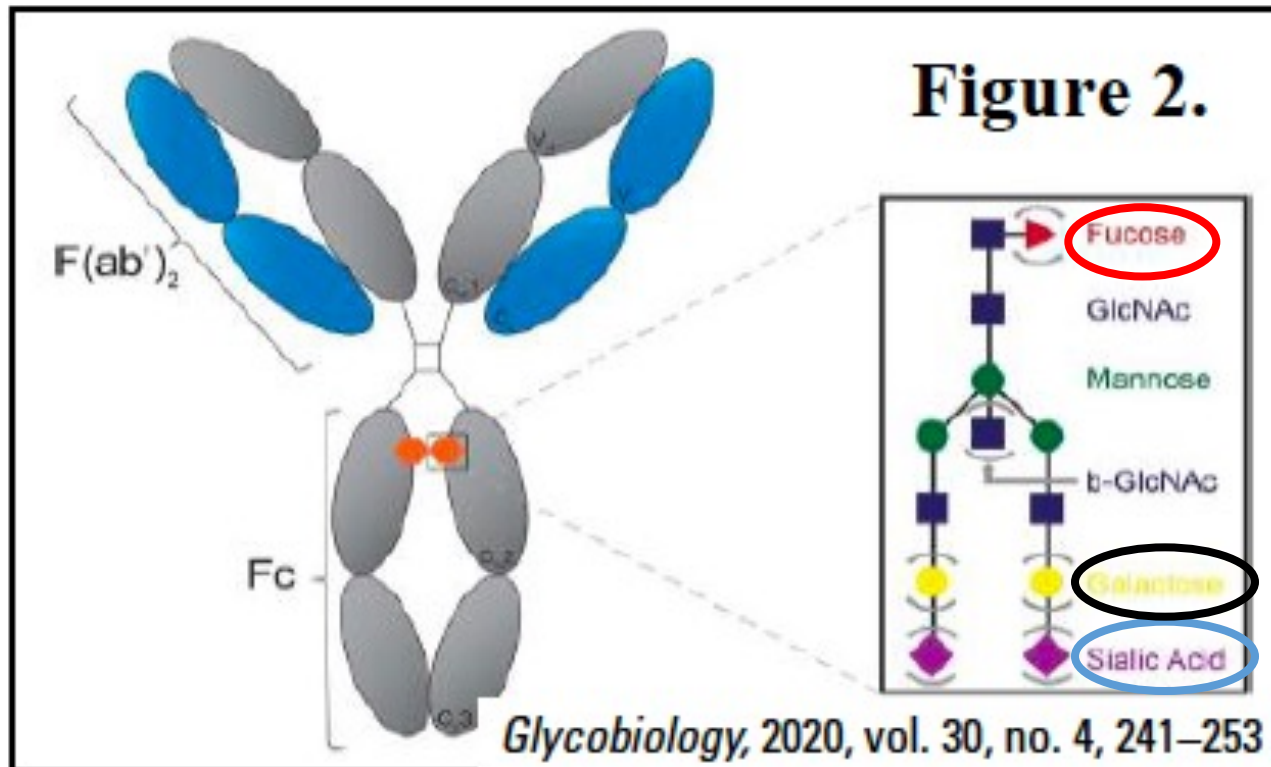
Donor Specific Antibodies (DSA) and Fc Receptors



**Functional interactions
with Fc receptors**



Fc Glycosylation



Lack of fucose activates FcGR3A on NK cells and monocytes/macrophages resulting in cytotoxicity

Lack of terminal galactose results in complement activation

Sialic acid activates inhibitory FcGR2B on monocytes, macrophages, B-cells

How can we decipher these glycans when DSA is different in every patient?



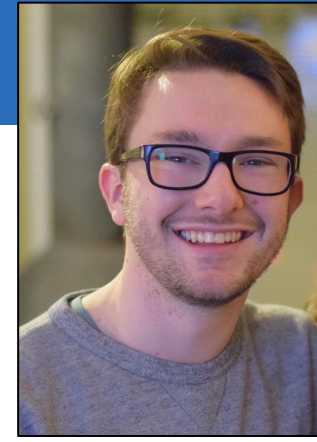
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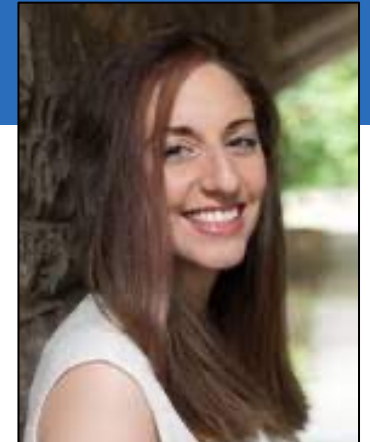
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KONVALINKA LAB



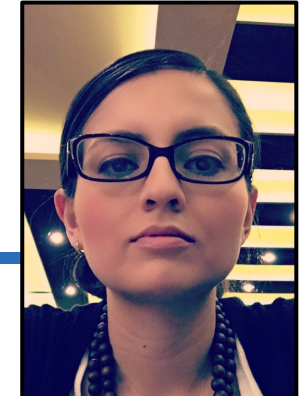
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Q & A

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