

SUPPLEMENTAL MATERIAL

Table of Contents

<i>Supplemental Table 1: Correlation coefficients between immunosuppressive medication use, UPCR, eGFR at biopsy and electron microscopy (EM) descriptors. Correlation coefficients with p-values < 0.05 are bolded.</i>	<i>2</i>
<i>Supplemental Table 2: Associations between clusters and clinical outcomes ($\geq 40\%$ decline in eGFR with eGFR < 60 or ESRD and complete proteinuria remission) from Cox models.</i>	<i>3</i>

Supplemental Table 1: Correlation coefficients between immunosuppressive medication use, UPCR, eGFR at biopsy and electron microscopy (EM) descriptors. Correlation coefficients with p-values < 0.05 are bolded.

	On Immunosuppressive Medications at Biopsy	UPCR at Biopsy	eGFR at Biopsy
Clinical measures			
On Immunosuppressive Medications at Biopsy	1.00		
UPCR at Biopsy	0.20	1.00	
eGFR at Biopsy	0.37	0.09	1.00
EM descriptors			
Foot Process Effacement	0.05	0.30	-0.01
Loss of Primary Processes	-0.07	0.08	-0.18
Condensation of Cytoskeleton	-0.03	0.19	-0.03
Podocyte Detachment	-0.10	-0.02	-0.16
Microvillous Transformation	0.25	0.30	0.05
Increased Thickening of GBM	-0.14	-0.06	-0.26
Increased Thinning of GBM	-0.13	0.04	-0.06
Tubuloreticular Inclusions	-0.05	0.02	-0.12
GBM Abnormal Texture	-0.11	-0.04	-0.05
Loss of Endothelial Cell Fenestration	-0.01	0.06	-0.02
Absence of Endothelial Cell Honeycombing	-0.10	-0.06	-0.06
Electron Densities/Hyaline Material	-0.12	-0.05	-0.17

UPCR, urinary protein/creatinine ratio; eGFR, estimated glomerular filtration rate

Supplemental Table 2: Associations between clusters and clinical outcomes ($\geq 40\%$ decline in eGFR with eGFR < 60 or ESRD and complete proteinuria remission) from Cox models

Composite ($\geq 40\%$ decline in eGFR with eGFR < 60 or ESRD)			Complete Proteinuria Remission			
Model 1 (unadjusted)			Model 1 (unadjusted)		Model 2 (adjusted)	
	HR (95% CI)	p-value	HR (95% CI)	p-value	HR (95% CI)	p-value
Clusters		0.0807		<.0001		0.1917
1: fewer changes (reference)						
2: thin GBM	1.39 (0.36, 5.39)	0.6307	1.43 (0.58, 3.51)	0.4334	1.11 (0.39, 3.15)	0.8514
3: classic MCD	0.16 (0.03, 0.76)	0.0215	2.87 (1.62, 5.09)	0.0003	1.99 (1.00, 3.93)	0.0491
4: abnormal GBM texture	0.51 (0.11, 2.46)	0.4015	1.42 (0.66, 3.03)	0.3683	1.45 (0.63, 3.34)	0.3869
5: thick GBM	1.25 (0.44, 3.56)	0.6787	1.26 (0.64, 2.47)	0.5102	1.21 (0.58, 2.55)	0.6092
6: endothelial cell changes	1.83 (0.53, 6.26)	0.3363	0.63 (0.23, 1.74)	0.3714	0.66 (0.23, 1.88)	0.4332

Model 2 adjusted for disease cohort (FSGS vs. MCD/MCD-Like), patient age, sex, black race, Hispanic ethnicity, eGFR and UPCR at biopsy, and immunosuppressant use at biopsy.

eGFR, estimated glomerular filtration rate; ESRD, end-stage renal disease; UPCR, urinary protein/creatinine ratio; GBM, glomerular basement membrane; MCD, minimal change disease