## Supplementary Table S2. Association of comorbidity with RCC subtype stratified by race and ethnicity and by sex.

## A. Papillary vs Clear Cell RCC

	OR and 95%CI								
	By race and ethnicity				By sex				
	Asian Pl	Latino	NH Black	NH White	Male	Female			
ESRD vs No kidney disease and no HTN	2.88	2.77	1.79	2.67	2.38	2.34			
	(1.65-5.01)	(1.95-3.94)	(1.31-2.46)	(2.14-3.33)	(2.00-2.84)	(1.72-3.19)			
CKD, not end-stage vs No kidney disease and no HTN	1.40	1.48	1.46	1.56	1.55	1.43			
	(0.80-2.46)	(1.03-2.12)	(1.07-2.00)	(1.33-1.83)	(1.34-1.79)	(1.09-1.87)			
HTN without kidney disease vs No kidney disease and no HTN	1.30	1.00	1.31	1.20	1.26	1.02			
	(0.93-1.81)	(0.80-1.25)	(1.05-1.64)	(1.09-1.32)	(1.16-1.38)	(0.87-1.19)			
Diabetes vs No diabetes	0.59	0.71	0.76	0.58	0.63	0.61			
	(0.41-0.85)	(0.57-0.89)	(0.61-0.93)	(0.51-0.65)	(0.57-0.70)	(0.51-0.74)			

## **B.** Chromophobe vs Clear Cell RCC

	OR and 95%CI								
	By race and ethnicity					By sex			
	Asian	Latino	NH Black	NH White		Male	Female		
	PI								
ESRD vs No kidney disease and no HTN	0.69	0.69	1.19	0.79		0.72	0.95		
	(0.24-1.96)	(0.40-1.19)	(0.67-2.10)	(0.52-1.20)		(0.50-1.04)	(0.64-1.42)		
CKD, not end-stage vs No kidney disease	1.26	0.52	0.94	0.81		0.80	0.73		
and no HTN	(0.60-2.62)	(0.31-0.87)	(0.53-1.67)	(0.63-1.03)		(0.63-1.03)	(0.52-1.01)		
HTN without kidney disease vs No kidney						0.97			
disease and no HTN	1.07	0.94	1.23	0.89		(0.85-	0.88		
	(0.74-1.54)	(0.76-1.16)	(0.85-1.79)	(0.79-1.01)		1.10)	(0.76-1.02)		
Diabetes vs No diabetes	0.38	0.66	0.75	0.60	•	0.67	0.55		
	(0.23-0.65)	(0.52-0.84)	(0.52-1.09)	(0.51-0.71)		(0.57-0.79)	(0.45-0.67)		

Estimates for race and ethnicity-stratified models computed from logistic regression model including terms for sex, age at diagnosis, comorbidity, neighborhood SES and year of diagnosis.

Estimates for sex-stratified models computed from logistic regression model including terms for race and ethnicity, age at diagnosis, comorbidity, neighborhood SES and year of diagnosis.