

Supplementary Material

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Item S1: Supplementary Methods

Study Data

CKD Progression Risk Factors

The set of demographic risk factors were: age (21-44, 45-64, and 65-75 years), gender, and race/ethnicity (non-Hispanic white, non-Hispanic black with low-risk APOL1 genotype, non-Hispanic black with high-risk APOL1 genotype, Hispanic, and other). APOL1 risk genotype was assessed using the number of G1 and G2 risk alleles characterizing individuals with black race as either low-risk (0 or 1 copies of an APOL1 risk allele) or high-risk (2 copies of a risk allele).¹ Kidney function measures included eGFR (<30, 30-44, 45-59, and ≥60 mL/min/1.73m²), and urine albumin:creatinine ratio (UACR; <30, 30-299, ≥300 mg/g). Blood pressure factors included systolic blood pressure (SBP; <120, 120-139, ≥140 mmHg) and self-reported use of an angiotensin converting enzyme inhibitor or angiotensin receptor blocker (ACE/ARB). Clinical risk factors included self-reported history of cardiovascular disease (CVD; myocardial infarction or revascularization, heart failure, stroke, or peripheral arterial disease) and serum uric acid level (<6.5, 6.5-8.1, ≥8.2 mg/dL). The socioeconomic factor was level of education (did not complete high school, high school diploma/some college, college graduate/graduate school) and behavioral factor was current smoking. Body composition measures included body mass index (<28, 28-34, ≥35 kg/m²), and fat-free mass in kilograms estimated using bioelectrical impedance analysis.^{2,3} Ankle-brachial index (<0.9, 0.9-1.0, 1.1-1.3, ≥1.4) was included as a peripheral vascular measure. Gender-specific categories of hemoglobin were used as a measure of anemia (<11, 11-12, ≥13 g/dL for women and <12, 12-13, ≥14 for men). High-sensitivity C-reactive protein (hsCRP) represented an inflammatory marker, and serum fractalkine (CX3CL1) and plasma CXCL12 were included as available inflammatory chemokines. Mineral metabolism markers included fibroblast growth factor-23 (FGF-23), serum phosphate, and intact parathyroid hormone (iPTH), and carbohydrate metabolism markers were hemoglobin A1c (HbA1c) and insulin resistance estimated by the Homeostatic Model Assessment (HOMA-IR) calculated using fasting plasma glucose and insulin.⁴ High-sensitivity Troponin T (hsTnT) and N-terminal pro-B-type natriuretic peptide (NTproBNP) were the considered cardiac markers. Serum bicarbonate served as the acidosis measure, urinary electrolytes included urine sodium and potassium, urine neutrophil gelatinase-associated lipocalin (NGAL) was used as a marker of kidney injury, and serum aldosterone was included as a marker related to the renin-angiotensin-aldosterone system. Continuous variables were assessed using quartiles unless otherwise noted.

Data Collection

SBP was measured in triplicate following a standardized protocol with the participant seated using a Tycos Classic hand cuff and aneroid sphygmomanometer. The baseline SBP value was the mean of all measurements. Self-reported current use of cigarettes along with at least 100 cigarettes smoked ever defined current smoking. Anthropometric measures were assessed using standard protocols.⁵ Participants were queried about any medication usage in the prior 30 days. Serum creatinine was measured at the CRIC Central Laboratory using an enzymatic method (www.orthoclinical.com) for samples collected through October 2008 and by the Jaffe method (www.beckmancoulter.com), thereafter, standardized to isotope dilution mass spectrometry-traceable values.^{6,7} Serum cystatin C was measured by particle-enhanced immunonephelometric assay on the Siemens BN™ II System (www.siemens.com). Estimated GFR was calculated from serum creatinine and cystatin C using a CRIC Study equation.⁸ Hemoglobin was obtained as part of a complete blood count at each of the CRIC clinical sites using local clinical laboratories. Hemoglobin A1c was analyzed in whole blood; CXCL12, glucose, insulin, hsCRP, iPTH, FGF-23, hsTnT, NTproBNP were measured in plasma; and uric acid, fractalkine, phosphate, bicarbonate, and aldosterone were evaluated using a serum sample. Urine albumin, creatinine, sodium, potassium, and NGAL were measured in 24-hour urine samples.

Outcomes and Censoring Events

KRT was defined as the initiation of maintenance dialysis or kidney transplantation and was ascertained through self-report and supplemented by data from the United States Renal Data System (USRDS). For participants who developed KRT, the eGFR from the Centers for Medicare and Medicaid Services Medical Evidence Form 2728 at onset of KRT was assigned as the last GFR value when available; for the remainder of participants reaching KRT, the mean eGFR reported by the USRDS for the age range in

CRIC (i.e., 10.8 mL/min/1.73m²) was assigned. Time to eGFR halving was imputed assuming a linear decline in kidney function between in-person annual visit measures.⁹

Deaths were determined from reports of next of kin, death certificates, obituaries, reviews of hospital records, the Social Security Death Master File and the National Death Index.

Table S1a. Intermediate models of chronic kidney disease progression adjusted only for demographic, kidney function and blood pressure risk factors among CRIC participants without diabetes

| | | Without Diabetes | | | |
|--|---------------------------|---|---------|------------------|---------|
| | | eGFR Slope, mL/min/1.73m ² /year | | KRT/eGFR Halving | |
| | | Beta Coefficient ^a (SE) | P value | HR (95% CI) | P value |
| Demographic Factors | | | | | |
| Age, years | <44 | -0.1 (0.1) | .4 | 1.3 (1.0, 1.7) | .1 |
| | 45-64 | Ref | | Ref | |
| | 65-75 | -0.1 (0.1) | | 1.0 (0.8, 1.3) | |
| Gender | Male | Ref | .003 | Ref | .002 |
| | Female | 0.3 (0.1) | | 0.7 (0.6, 0.9) | |
| Race/ethnicity | NH-White | Ref | <.001 | Ref | <.001 |
| | NH-Black, APOL1 low-risk | -0.2 (0.1) | | 1.8 (1.4, 2.3) | |
| | NH-Black, APOL1 high-risk | -0.6 (0.2) | | 2.3 (1.7, 3.1) | |
| | Hispanic | 0.1 (0.2) | | 1.2 (0.8, 1.8) | |
| | Other | -0.4 (0.2) | | 2.3 (1.4, 3.7) | |
| Kidney Function Measures | | | | | |
| Baseline eGFR, mL/min/1.73m ² | <30 | 0.2 (0.2) | .04 | 2.1 (1.7, 2.7) | <.001 |
| | 30-44.9 | Ref | | Ref | |
| | 45-59.9 | 0.0 (0.1) | | 0.5 (0.4, 0.7) | |
| | ≥60 | 0.3 (0.1) | | 0.2 (0.1, 0.3) | |
| UACR, mg/g | <30 | Ref | <.001 | Ref | <.001 |
| | 30-299 | -1.0 (0.1) | | 3.5 (2.6, 4.8) | |
| | ≥300 | -2.3 (0.1) | | 10.6 (7.8, 14.4) | |
| Blood Pressure Factors | | | | | |
| Systolic BP, mmHg | <120 | Ref | <.001 | Ref | .003 |
| | 120-139 | -0.2 (0.1) | | 1.3 (1.1, 1.7) | |
| | ≥140 | -0.5 (0.1) | | 1.5 (1.2, 2.0) | |
| ACE/ARB | No | Ref | .04 | Ref | .8 |
| | Yes | 0.2 (0.1) | | 1.0 (0.8, 1.3) | |
| Clinical Factors | | | | | |
| History of CVD | No | Ref | .6 | Ref | .8 |
| | Yes | -0.1 (0.1) | | 1.0 (0.8, 1.2) | |

| | | Without Diabetes | | | |
|---|--------------------------|--|---------|------------------|---------|
| | | eGFR Slope, mL/min/1.73m ² /year | | KRT/eGFR Halving | |
| | | Beta Coefficient ^a (SE) | P value | HR (95% CI) | P value |
| Uric acid, mg/dL | <6.5 | Ref | .9 | Ref | .9 |
| | 6.5-8.9 | 0.0 (0.1) | | 1.1 (0.8, 1.4) | |
| | ≥8.2 | 0.0 (0.1) | | 1.1 (0.8, 1.4) | |
| Socioeconomic Status Factor | | | | | |
| Education | Less than High School | 0.0 (0.1) | .5 | 1.1 (0.8, 1.4) | .6 |
| | High School/Some College | Ref | | Ref | |
| | Graduated College | -0.1 (0.1) | | 1.1 (0.9, 1.4) | |
| Behavioral Factor | | | | | |
| Current Smoker | No | Ref | .6 | Ref | .3 |
| | Yes | -0.1 (0.1) | | 1.2 (0.9, 1.5) | |
| Body Composition Measures | | | | | |
| BMI, kg/m ² | <28 | -0.1 (0.1) | .3 | 1.0 (0.8, 1.3) | .3 |
| | 28-34.9 | Ref | | Ref | |
| | ≥35 | 0.0 (0.1) | | 0.8 (0.6, 1.1) | |
| Fat-free mass, kg | Q1: [25.8, 49.1) | Ref | .1 | Ref | .02 |
| | Q2: [49.1, 59.1) | 0.3 (0.1) | | 0.7 (0.5, 0.9) | |
| | Q3: [59.1, 69.9) | 0.3 (0.1) | | 0.6 (0.5, 0.9) | |
| | Q4: [69.9, 167.1) | 0.2 (0.2) | | 0.8 (0.6, 1.1) | |
| Peripheral Vascular Measure | | | | | |
| Ankle- brachial index | <0.9 | -0.3 (0.2) | .2 | 1.2 (0.8, 1.6) | .2 |
| | 0.9-1.09 | Ref | | Ref | |
| | 1.1-1.39 | 0.0 (0.1) | | 0.9 (0.7, 1.1) | |
| | ≥1.4 | -0.4 (0.4) | | 1.6 (0.7, 3.7) | |
| Measure of Anemia | | | | | |
| Hemoglobin, mg/dL | M: <12, F: <11 | 0.1 (0.1) | .3 | 1.3 (1.0, 1.7) | .003 |
| | M: 12-13.9, F: 11-12.9 | Ref | | Ref | |
| | M: ≥14, F: ≥13 | 0.1 (0.1) | | 0.8 (0.6, 1.0) | |
| Inflammatory Marker and Chemokines | | | | | |
| hsCRP, mg/L | Q1: [0.1, 1.0) | Ref | .9 | Ref | .6 |
| | Q2: [1.0, 2.5) | 0.0 (0.1) | | 1.0 (0.8, 1.4) | |
| | Q3: [2.5, 6.3) | 0.1 (0.1) | | 0.9 (0.7, 1.2) | |

| | Without Diabetes | | | | |
|--|--|----------------|-------------------------|----------------|-----|
| | eGFR Slope, mL/min/1.73m²/year | | KRT/eGFR Halving | | |
| | Beta Coefficient^a (SE) | P value | HR (95% CI) | P value | |
| | Q4: [6.3, 187) | 0.0 (0.1) | | 1.0 (0.8, 1.4) | |
| Serum fractalkine (CX3CL1), pg/mL | Q1: [0.1, 0.6) | Ref | .9 | Ref | .3 |
| | Q2: [0.6, 0.8) | 0.0 (0.1) | | 1.3 (1.0, 1.8) | |
| | Q3: [0.8, 1.1) | 0.1 (0.1) | | 1.2 (0.8, 1.6) | |
| | Q4: [1.1, 3.6] | 0.0 (0.1) | | 1.3 (1.0, 1.8) | |
| Plasma CXCL12, pg/mL | Q1: [832.1, 2066.4) | Ref | .3 | Ref | .2 |
| | Q2: [2066.4, 2410.5) | -0.2 (0.1) | | 1.2 (0.9, 1.7) | |
| | Q3: [2410.5, 2797.8) | -0.1 (0.1) | | 1.3 (1.0, 1.7) | |
| | Q4: [2797.8, 6173.3] | -0.2 (0.1) | | 1.4 (1.0, 1.8) | |
| Mineral Metabolism Markers | | | | | |
| FGF23, RU/mL | Q1: [1.4, 94.1) | Ref | .1 | Ref | .1 |
| | Q2: [94.1, 139.1) | 0.0 (0.1) | | 1.4 (1.0, 1.9) | |
| | Q3: [139.1, 225.5) | 0.2 (0.1) | | 1.4 (1.0, 1.9) | |
| | Q4: [225.5, 14 318.9] | 0.3 (0.1) | | 1.4 (1.0, 1.9) | |
| Serum phosphate, mg/dL | Q1: [1.7, 3.3) | -0.3 (0.1) | .09 | 1.4 (1.0, 1.8) | .02 |
| | Q2: [3.3, 3.7) | Ref | | Ref | |
| | Q3: [3.7, 4.1) | 0.0 (0.1) | | 1.5 (1.1, 2.0) | |
| | Q4: [4.1, 9.3] | -0.1 (0.1) | | 1.4 (1.0, 1.9) | |
| Intact PTH pg/mL | Q1: [1.9, 34.1) | 0.2 (0.1) | .3 | 1.0 (0.7, 1.4) | .6 |
| | Q2: [34.1, 52.4) | Ref | | Ref | |
| | Q3: [52.4, 84.7) | 0.1 (0.1) | | 1.0 (0.8, 1.4) | |
| | Q4: [84.5, 1483] | 0.0 (0.1) | | 1.2 (0.9, 1.6) | |
| Carbohydrate Metabolism Markers | | | | | |
| HbA1c, % | Q1: [3.5, 5.6) | Ref | .8 | Ref | .6 |
| | Q2: [5.6, 6.2) | -0.1 (0.1) | | 0.9 (0.7, 1.1) | |
| | Q3: [6.2, 7.3) | 0.0 (0.1) | | 0.8 (0.6, 1.1) | |
| | Q4: [7.3, 15.2] | -0.3 (0.5) | | 0.9 (0.4, 2.3) | |
| HOMA-IR | Q1: [0, 2.5) | Ref | .9 | Ref | .3 |
| | Q2: [2.5, 4.1) | 0.0 (0.1) | | 0.9 (0.7, 1.2) | |
| | Q3: [4.1, 7.2) | 0.1 (0.1) | | 1.0 (0.8, 1.3) | |
| | Q4: [7.2, 224.8) | 0.1 (0.2) | | 0.7 (0.5, 1.0) | |

| | Without Diabetes | | | |
|---------------------------------|--|------------|------------------|----------------|
| | eGFR Slope, mL/min/1.73m ² /year | | KRT/eGFR Halving | |
| | Beta Coefficient ^a (SE) | P value | HR (95% CI) | P value |
| Cardiac Markers | | | | |
| hsTnT, pg/mL | Q1: [1.5, 5.5) | Ref | .7 | Ref |
| | Q2: [5.5, 11.4) | 0.1 (0.1) | | 1.2 (0.9, 1.5) |
| | Q3: (11.4, 22.2) | 0.0 (0.1) | | 0.9 (0.6, 1.2) |
| | Q4: (22.2, 738.7) | 0.2 (0.2) | | 1.0 (0.7, 1.4) |
| NTproBNP, pg/mL | Q1: [2.5, 60.2) | Ref | .02 | Ref |
| | Q2: [60.2, 139.4) | -0.2 (0.1) | | 1.5 (1.1, 2.1) |
| | Q3: [139.4, 366.6) | -0.4 (0.1) | | 1.7 (1.3, 2.4) |
| | Q4: [366.6, 33 742] | -0.2 (0.1) | | 1.7 (1.2, 2.4) |
| Acidosis Marker | | | | |
| Serum bicarbonate, mmol/L | ≤22 | -0.1 (0.1) | .8 | 1.5 (1.2, 2.0) |
| | (22, 24] | Ref | | Ref |
| | (24, 26] | 0.0 (0.1) | | 1.0 (0.7, 1.4) |
| | >26 | 0.0 (0.1) | | 1.1 (0.8, 1.5) |
| Urinary Electrolytes | | | | |
| Urine sodium, mEq/24h | Q1: [4.3, 108.6) | Ref | .2 | Ref |
| | Q2: [108.6, 151.3) | 0.2 (0.1) | | 0.8 (0.6, 1.1) |
| | Q3: [151.3, 203.1) | 0.1 (0.1) | | 0.9 (0.7, 1.2) |
| | Q4: [203.1, 699.8] | 0.2 (0.1) | | 0.9 (0.7, 1.2) |
| Urine potassium, mmol/24h | Q1: [3.0, 37.5) | Ref | .05 | Ref |
| | Q2: [37.5, 51.8) | 0.0 (0.1) | | 1.1 (0.9, 1.5) |
| | Q3: [51.8, 69.4) | -0.1 (0.1) | | 1.3 (0.9, 1.7) |
| | Q4: [69.4, 417.7] | 0.2 (0.1) | | 1.1 (0.8, 1.5) |
| Kidney Injury Marker | | | | |
| Urine NGAL, ng/ml | Q1: [0.4, 6.3) | Ref | .06 | Ref |
| | Q2: [6.3, 14.2) | -0.3 (0.1) | | 1.7 (1.2, 2.4) |
| | Q3: [14.2, 32.9) | -0.2 (0.1) | | 1.3 (0.9, 1.9) |
| | Q4: [32.9, 2743.8] | -0.3 (0.1) | | 1.7 (1.2, 2.5) |
| RAAS Marker | | | | |
| | Q1: [0.8, 71.2) | Ref | .03 | Ref |
| | Q2: [71.2, 101.4) | -0.3 (0.1) | | 1.4 (1.0, 1.9) |

| | Without Diabetes | | | |
|--------------------------------|--|------------|------------------|---------|
| | eGFR Slope, mL/min/1.73m ² /year | | KRT/eGFR Halving | |
| | Beta Coefficient ^a (SE) | P value | HR (95% CI) | P value |
| Serum aldosterone, pg/mL | Q3: [101.4, 152.8] | -0.2 (0.1) | 1.4 (1.1, 1.9) | |
| | Q4: [152.8, 15 630.9] | 0.0 (0.1) | | |

Abbreviations: ACE: angiotensin converting enzyme; ARB: angiotensin receptor blocker; BP: blood pressure; CI: confidence interval; CKD: chronic kidney disease; CVD: cardiovascular disease; eGFR: estimated glomerular filtration rate; KRT: kidney replacement therapy; FGF23: fibroblast growth factor-23; HR: hazard ratio; hsCRP: high-sensitivity C-reactive protein; hsTnT: high-sensitivity troponin T; NGAL: neutrophil gelatinase-associated lipocalin; NH: non-Hispanic; NTproBNP: N-terminal pro-B-type natriuretic peptide; PTH: parathyroid hormone; RAAS: renin-angiotensin-aldosterone system; Ref: reference; SE: standard error; UACR: urine albumin:creatinine ratio

SI conversion: To convert Uric acid to $\mu\text{mol/L}$, multiply by 59.485. To convert Hemoglobin to g/L, multiply by 0.01. To convert hsCRP to nmol/L, multiply by 9.524. To convert serum phosphate to mmol/L, multiply by 0.323. For serum bicarbonate, 1 mEq/L is equivalent to 1 mmol/L. For urine potassium, 1 mEq/24h is equivalent to 1 mmol/d. To convert serum aldosterone to pmol/L, multiply by 2.774.

^a Negative coefficients represent a faster decline in eGFR compared to the reference group.

Table S1b. Intermediate models of chronic kidney disease progression adjusted only for demographic, kidney function and blood pressure risk factors among CRIC participants with diabetes

| | | With Diabetes | | | |
|--|---------------------------|---|---------|------------------|---------|
| | | eGFR Slope, mL/min/1.73m ² /year | | KRT/eGFR Halving | |
| | | Beta Coefficient ^a (SE) | P value | HR (95% CI) | P value |
| Demographic Factors | | | | | |
| Age, years | <44 | -0.5 (0.2) | .1 | 1.3 (1.0, 1.7) | .001 |
| | 45-64 | Ref | | Ref | |
| | 65-75 | 0.0 (0.1) | | 0.8 (0.6, 0.9) | |
| Gender | Male | Ref | .6 | Ref | .01 |
| | Female | -0.1 (0.1) | | 0.8 (0.7, 1.0) | |
| Race/ethnicity | NH-White | Ref | <.001 | Ref | <.001 |
| | NH-Black, APOL1 low-risk | -0.7 (0.1) | | 1.5 (1.2, 1.8) | |
| | NH-Black, APOL1 high-risk | -1.1 (0.3) | | 1.7 (1.3, 2.3) | |
| | Hispanic | -0.5 (0.2) | | 1.4 (1.1, 1.9) | |
| | Other | -0.3 (0.3) | | 1.2 (0.8, 1.7) | |
| Kidney Function Measures | | | | | |
| Baseline eGFR, mL/min/1.73m ² | <30 | 0.4 (0.2) | <.001 | 1.9 (1.6, 2.3) | <.001 |
| | 30-44.9 | Ref | | Ref | |
| | 45-59.9 | -0.4 (0.1) | | 0.6 (0.5, 0.8) | |
| | ≥60 | -0.4 (0.2) | | 0.4 (0.3, 0.6) | |
| UACR, mg/g | <30 | Ref | <.001 | Ref | <.001 |
| | 30-299 | -1.1 (0.1) | | 2.8 (2.1, 3.7) | |
| | ≥300 | -2.7 (0.2) | | 7.1 (5.5, 9.3) | |
| Blood Pressure Factors | | | | | |
| Systolic BP, mmHg | <120 | Ref | <.001 | Ref | <.001 |
| | 120-139 | -0.2 (0.1) | | 1.2 (1.0, 1.5) | |
| | ≥140 | -0.8 (0.2) | | 1.9 (1.5, 2.3) | |
| ACE/ARB | No | Ref | .9 | Ref | .2 |
| | Yes | 0.0 (0.2) | | 0.9 (0.7, 1.1) | |
| Clinical Factors | | | | | |
| History of CVD | No | Ref | .3 | Ref | .02 |
| | Yes | -0.1 (0.1) | | 1.2 (1.0, 1.4) | |

| | | With Diabetes | | | |
|---|--------------------------|---|---------|------------------|---------|
| | | eGFR Slope, mL/min/1.73m ² /year | | KRT/eGFR Halving | |
| | | Beta Coefficient ^a (SE) | P value | HR (95% CI) | P value |
| Uric acid, mg/dL | <6.5 | Ref | .3 | Ref | .08 |
| | 6.5-8.9 | -0.2 (0.1) | | 1.2 (1.0, 1.5) | |
| | ≥8.2 | -0.2 (0.2) | | 1.3 (1.0, 1.6) | |
| Socioeconomic Status Factor | | | | | |
| Education | Less than High School | -0.2 (0.2) | .2 | 0.9 (0.8, 1.1) | .6 |
| | High School/Some College | Ref | | Ref | |
| | Graduated College | 0.1 (0.1) | | 0.9 (0.8, 1.1) | |
| Behavioral Factor | | | | | |
| Current Smoker | No | Ref | .07 | Ref | .06 |
| | Yes | -0.3 (0.2) | | 1.2 (1.0, 1.5) | |
| Body Composition Measures | | | | | |
| BMI, kg/m ² | <28 | -0.2 (0.2) | .5 | 1.1 (0.9, 1.4) | .4 |
| | 28-34.9 | Ref | | Ref | |
| | ≥35 | 0.0 (0.1) | | 1.0 (0.9, 1.2) | |
| Fat-free mass, kg | Q1: [25.8, 49.1) | Ref | .9 | Ref | .6 |
| | Q2: [49.1, 59.1) | 0.0 (0.2) | | 0.9 (0.7, 1.1) | |
| | Q3: [59.1, 69.9) | -0.1 (0.2) | | 0.9 (0.7, 1.2) | |
| | Q4: [69.9, 167.1) | 0.0 (0.2) | | 1.0 (0.7, 1.3) | |
| Peripheral Vascular Measure | | | | | |
| Ankle-brachial index | <0.9 | 0.0 (0.2) | .7 | 1.1 (0.9, 1.4) | .03 |
| | 0.9-1.09 | Ref | | Ref | |
| | 1.1-1.39 | 0.1 (0.1) | | 1.1 (0.9, 1.3) | |
| | ≥1.4 | -0.2 (0.3) | | 1.7 (1.2, 2.5) | |
| Measure of Anemia | | | | | |
| Hemoglobin, mg/dL | M: <12, F: <11 | -0.2 (0.1) | .2 | 1.1 (0.9, 1.3) | .1 |
| | M: 12-13.9, F: 11-12.9 | Ref | | Ref | |
| | M: ≥14, F: ≥13 | 0.2 (0.2) | | 0.9 (0.7, 1.1) | |
| Inflammatory Marker and Chemokines | | | | | |
| hsCRP, mg/L | Q1: [0.1, 1.0) | Ref | .3 | Ref | .4 |
| | Q2: [1.0, 2.5) | 0.1 (0.2) | | 0.9 (0.7, 1.1) | |
| | Q3: [2.5, 6.3) | 0.3 (0.2) | | 0.8 (0.7, 1.0) | |

| | With Diabetes | | | | |
|--|--|------------|------------------|----------------|-------|
| | eGFR Slope, mL/min/1.73m ² /year | | KRT/eGFR Halving | | |
| | Beta Coefficient ^a (SE) | P value | HR (95% CI) | P value | |
| Q4: [6.3, 187) | 0.2 (0.2) | | 0.9 (0.7, 1.1) | | |
| Serum fractalkine (CX3CL1), pg/mL | Q1: [0.1, 0.6) | Ref | .01 | Ref | .05 |
| | Q2: [0.6, 0.8) | 0.1 (0.2) | | 1.0 (0.7, 1.3) | |
| | Q3: [0.8, 1.1) | -0.4 (0.2) | | 1.2 (0.9, 1.6) | |
| | Q4: [1.1, 3.6] | -0.4 (0.2) | | 1.3 (1.0, 1.7) | |
| Plasma CXCL12, pg/mL | Q1: [832.1, 2066.4) | Ref | .3 | Ref | <.001 |
| | Q2: [2066.4, 2410.5) | -0.1 (0.2) | | 1.2 (0.9, 1.6) | |
| | Q3: [2410.5, 2797.8) | -0.3 (0.2) | | 1.5 (1.2, 2.0) | |
| | Q4: [2797.8, 6173.3] | -0.3 (0.2) | | 1.7 (1.4, 2.2) | |
| Mineral Metabolism Markers | | | | | |
| FGF23, RU/mL | Q1: [1.4, 94.1) | Ref | .6 | Ref | <.001 |
| | Q2: [94.1, 139.1) | -0.2 (0.2) | | 1.2 (0.9, 1.6) | |
| | Q3: [139.1, 225.5) | -0.1 (0.2) | | 1.6 (1.2, 2.1) | |
| | Q4: [225.5, 14 318.9] | -0.3 (0.2) | | 1.8 (1.3, 2.5) | |
| Serum phosphate, mg/dL | Q1: [1.7, 3.3) | 0.0 (0.2) | .9 | 0.9 (0.7, 1.1) | .2 |
| | Q2: [3.3, 3.7) | Ref | | Ref | |
| | Q3: [3.7, 4.1) | 0.0 (0.2) | | 1.0 (0.8, 1.2) | |
| | Q4: [4.1, 9.3] | 0.1 (0.2) | | 1.1 (0.9, 1.4) | |
| Intact PTH pg/mL | Q1: [1.9, 34.1) | 0.2 (0.2) | .2 | 1.1 (0.9, 1.5) | .06 |
| | Q2: [34.1, 52.4) | Ref | | Ref | |
| | Q3: [52.4, 84.7) | -0.1 (0.2) | | 1.1 (0.9, 1.4) | |
| | Q4: [84.5, 1483] | -.1 (0.2) | | 1.4 (1.1, 1.7) | |
| Carbohydrate Metabolism Markers | | | | | |
| HbA1c, % | Q1: [3.5, 5.6) | Ref | .3 | Ref | .1 |
| | Q2: [5.6, 6.2) | 0.2 (0.3) | | 1.3 (0.9, 2.1) | |
| | Q3: [6.2, 7.3) | 0.2 (0.3) | | 1.0 (0.6, 1.4) | |
| | Q4: [7.3, 15.2] | 0.0 (0.3) | | 1.0 (0.7, 1.6) | |
| HOMA-IR | Q1: [0, 2.5) | Ref | .8 | Ref | .3 |
| | Q2: [2.5, 4.1) | 0.1 (0.2) | | 0.8 (0.6, 1.0) | |
| | Q3: [4.1, 7.2) | 0.1 (0.2) | | 0.9 (0.6, 1.1) | |
| | Q4: [7.2, 224.8) | 0.2 (0.2) | | 0.8 (0.6, 1.0) | |

| | | With Diabetes | | | |
|---------------------------------|-----------------------|--|---------|------------------|---------|
| | | eGFR Slope, mL/min/1.73m ² /year | | KRT/eGFR Halving | |
| | | Beta Coefficient ^a (SE) | P value | HR (95% CI) | P value |
| hsTnT, pg/mL | Q1: [1.5, 5.5) | Ref | .2 | Ref | <.001 |
| | Q2: [5.5, 11.4) | -0.2 (0.2) | | 1.1 (0.8, 1.6) | |
| | Q3: (11.4, 22.2) | -0.2 (0.2) | | 1.3 (0.9, 1.8) | |
| | Q4: (22.2, 738.7) | -0.4 (0.2) | | 1.8 (1.3, 2.6) | |
| NTproBNP, pg/mL | Q1: [2.5, 60.2) | Ref | <.001 | Ref | <.001 |
| | Q2: [60.2, 139.4) | -0.5 (0.2) | | 1.5 (1.1, 2.0) | |
| | Q3: [139.4, 366.6) | -0.5 (0.2) | | 1.6 (1.2, 2.1) | |
| | Q4: [366.6, 33 742] | -0.8 (0.2) | | 2.1 (1.6, 2.8) | |
| Serum bicarbonate, mmol/L | ≤22 | 0.1 (0.2) | .5 | 0.9 (0.7, 1.1) | .7 |
| | (22, 24] | Ref | | Ref | |
| | (24, 26] | 0.1 (0.2) | | 0.9 (0.7, 1.1) | |
| | >26 | 0.3 (0.2) | | 0.9 (0.7, 1.1) | |
| Urine sodium, mEq/24h | Q1: [4.3, 108.6) | Ref | .2 | Ref | .7 |
| | Q2: [108.6, 151.3) | 0.0 (0.2) | | 1.0 (0.8, 1.2) | |
| | Q3: [151.3, 203.1) | 0.3 (0.2) | | 0.9 (0.8, 1.2) | |
| | Q4: [203.1, 699.8] | 0.2 (0.2) | | 0.9 (0.7, 1.1) | |
| Urine potassium, mmol/24h | Q1: [3.0, 37.5) | Ref | .2 | Ref | .1 |
| | Q2: [37.5, 51.8) | 0.0 (0.2) | | 0.8 (0.7, 1.0) | |
| | Q3: [51.8, 69.4) | 0.2 (0.2) | | 0.8 (0.6, 1.0) | |
| | Q4: [69.4, 417.7] | 0.3 (0.2) | | 0.8 (0.6, 1.0) | |
| Urine NGAL, ng/ml | Q1: [0.4, 6.3) | Ref | <.001 | Ref | <.001 |
| | Q2: [6.3, 14.2) | -0.4 (0.2) | | 1.2 (0.9, 1.6) | |
| | Q3: [14.2, 32.9) | -0.6 (0.2) | | 1.3 (1.0, 1.7) | |
| | Q4: [32.9, 2743.8] | -0.8 (0.2) | | 2.3 (1.7, 2.9) | |
| Serum aldosterone, pg/mL | Q1: [0.8, 71.2) | Ref | .8 | Ref | .2 |
| | Q2: [71.2, 101.4) | 0.1 (0.2) | | 0.9 (0.7, 1.1) | |
| | Q3: [101.4, 152.8) | 0.2 (0.2) | | 0.9 (0.7, 1.1) | |
| | Q4: [152.8, 15 630.9] | 0.1 (0.2) | | 1.1 (0.9, 1.4) | |

Abbreviations: ACE: angiotensin converting enzyme; ARB: angiotensin receptor blocker; BP: blood pressure; CI: confidence interval; CKD: chronic kidney disease; CVD: cardiovascular disease; eGFR: estimated glomerular filtration rate; KRT: kidney replacement therapy; FGF23: fibroblast growth factor-23; HR: hazard ratio; hsCRP: high-sensitivity C-reactive protein; hsTnT: high-sensitivity troponin T; NGAL: neutrophil gelatinase-associated lipocalin; NH: non-Hispanic; NTproBNP: N-terminal pro-B-type natriuretic peptide; PTH: parathyroid hormone; RAAS: renin-angiotensin-aldosterone system; Ref: reference; SE: standard error; UACR: urine albumin:creatinine ratio

SI conversion: To convert Uric acid to $\mu\text{mol/L}$, multiply by 59.485. To convert Hemoglobin to g/L , multiply by 0.01. To convert hsCRP to nmol/L , multiply by 9.524. To convert serum phosphate to mmol/L , multiply by 0.323. For serum bicarbonate, 1 mEq/L is equivalent to 1 mmol/L . For urine potassium, 1 mEq/24h is equivalent to 1 mmol/d . To convert serum aldosterone to pmol/L , multiply by 2.774.

^a Negative coefficients represent a faster decline in eGFR compared to the reference group.

Table S2a. Final multivariable-adjusted models of chronic kidney disease progression among CRIC participants without diabetes after exclusion of albuminuria

| | | Without Diabetes | | | |
|--|---------------------------|--|---------|--------------------------|-----------------------------------|
| | | eGFR Slope, mL/min/1.73m ² /year | | KRT/eGFR Halving | |
| | | Beta Coefficient ^a (SE) | P value | HR (95% CI) ^b | P value ^b |
| Demographic Factors | | | | | |
| Age, years | <45 | -0.7 (0.1) | <.001 | Y0-6: 2.4 (1.7, 3.4) | Y0-6: <.001 Y6+: <.001 |
| | 45-64 | Ref | | Y6+: 2.4 (1.4, 3.9) | |
| | 65-75 | 0.1 (0.1) | | Ref | |
| | | | | Y0-6: 0.7 (0.5, 1.0) | |
| Gender | Male | Ref | <.001 | Y6+: 0.8 (0.5, 1.3) | <.001 |
| | Female | 0.8 (0.1) | | Ref | |
| Race/ ethnicity | NH-White | Ref | <.001 | 0.4 (0.3, 0.5) | Y0-6: <.001 Y6+: .07 |
| | NH-Black, APOL1 low-risk | -0.2 (0.1) | | Y0-6: 1.8 (1.3, 2.5) | |
| | NH-Black, APOL1 high-risk | -0.8 (0.2) | | Y6+: 1.4 (0.9, 2.1) | |
| | Hispanic | 0.0 (0.2) | | Y0-6: 3.7 (2.5, 5.4) | |
| | Other | -0.7 (0.2) | | Y6+: 1.3 (0.6, 2.5) | |
| Kidney Function Measures | | | | | |
| Baseline eGFR, mL/min/ 1.73m ² | <30 | 0.1 (0.2) | <.001 | Y0-6: 1.3 (2.2) | <.001 |
| | 30-44.9 | Ref | | Ref | |
| | 45-59.9 | 0.2 (0.1) | | 0.5 (0.4, 0.6) | |
| | ≥60 | 0.7 (0.1) | | 0.2 (0.1, 0.3) | |
| Blood Pressure Factors | | | | | |
| Systolic BP, mmHg | <120 | Ref | <.001 | Ref | <.001 |
| | 120-139 | -0.3 (0.1) | | 1.4 (1.1, 1.8) | |
| | ≥140 | -0.8 (0.1) | | 2.0 (1.5, 2.7) | |
| ACE/ARB | No | - | - | Ref | .01 |

| | | Without Diabetes | | | | |
|-----------------------------------|------------------------|--|---------|--------------------------|----------------------|--|
| | | eGFR Slope, mL/min/1.73m ² /year | | KRT/eGFR Halving | | |
| | | Beta Coefficient ^a (SE) | P value | HR (95% CI) ^b | P value ^b | |
| Yes | | - | | 1.3 (1.1, 1.7) | | |
| Behavioral Factor | | | | | | |
| Current smoker | No | - | - | Ref | .08 | |
| | Yes | - | | 1.3 (1.0, 1.8) | | |
| Body Composition Measures | | | | | | |
| Fat-free mass, kg | Q1: [25.8, 49.1) | Ref | .04 | Ref | .01 | |
| | Q2: [49.1, 59.1) | 0.3 (0.1) | | 0.7 (0.5, 0.9) | | |
| | Q3: [59.1, 69.9) | 0.3 (0.2) | | 0.8 (0.5, 1.1) | | |
| | Q4: [69.9, 167.1) | 0.1 (0.2) | | 1.0 (0.7, 1.5) | | |
| Measure of Anemia | | | | | | |
| Hemoglobin, mg/dL | M: <12, F: <11 | - | - | Y0-6: 1.2 (0.9, 1.6) | Y0-6: .2 | |
| | M: 12-13.9, F: 11-12.9 | - | | Y6+: 0.9 (0.5, 1.4) | | |
| | M: ≥14, F: ≥13 | - | | Ref | | |
| | | | | Y0-6: 0.8 (0.6, 1.2) | Y6+: .008 | |
| | | | | Y6+: 0.5 (0.3, 0.8) | | |
| Mineral Metabolism Markers | | | | | | |
| FGF23, RU/mL | Q1: [1.4, 94.1) | - | - | Ref | .03 | |
| | Q2: [94.1, 139.1) | - | | 1.3 (1.0, 1.9) | | |
| | Q3: [139.1, 225.5) | - | | 1.6 (1.1, 2.2) | | |
| | Q4: [225.5, 14 318.9] | - | | 1.7 (1.2, 2.4) | | |
| Cardiac Markers | | | | | | |
| NTproBNP, pg/mL | Q1: [2.5, 60.2) | Ref | .009 | Ref | Y0-6: .002 | |
| | Q2: [60.2, 139.4) | -0.2 (0.1) | | Y0-6: 1.6 (1.1, 2.5) | | |
| | Q3: [139.4, 366.6) | -0.5 (0.1) | | Y6+: 0.8 (0.5, 1.4) | | |
| | Q4: [366.6, 33 742] | -0.4 (0.2) | | Y0-6: 2.2 (1.4, 3.4) | | |
| | | | | Y6+: 1.3 (0.8, 2.2) | | |
| | | | | Y0-6: 2.2 (1.4, 3.5) | Y6+: .3 | |
| | | | | Y6+: 1.2 (0.7, 2.1) | | |
| Acidosis Marker | | | | | | |
| | ≤22 | - | - | 1.7 (1.3, 2.3) | .003 | |
| | (22, 24] | - | | Ref | | |

| | | Without Diabetes | | | |
|---------------------------------|--------------------|--|---------|--------------------------|----------------------|
| | | eGFR Slope, mL/min/1.73m ² /year | | KRT/eGFR Halving | |
| | | Beta Coefficient ^a (SE) | P value | HR (95% CI) ^b | P value ^b |
| Serum bicarbonate, mmol/L | (24, 26] | - | | 1.3 (0.9, 1.8) | |
| | >26 | - | | 1.3 (0.9, 1.8) | |
| Urinary Electrolytes | | | | | |
| Urine potassium, mmol/24h | Q1: [3.0, 37.5) | Ref | .04 | - | - |
| | Q2: [37.5, 51.8) | -0.1 (0.1) | | - | |
| | Q3: [51.8, 69.4) | -0.3 (0.1) | | - | |
| | Q4: [69.4, 417.7] | 0.0 (0.1) | | - | |
| Kidney Injury Marker | | | | | |
| Urine NGAL, ng/ml | Q1: [0.4, 6.3) | Ref | <.001 | Ref | <.001 |
| | Q2: [6.3, 14.2) | -0.4 (0.1) | | 2.0 (1.4, 2.9) | |
| | Q3: [14.2, 32.9) | -0.5 (0.1) | | 2.0 (1.4, 2.9) | |
| | Q4: [32.9, 2743.8] | -0.7 (0.2) | | 2.7 (1.8, 4.0) | |

Abbreviations: ACE: angiotensin converting enzyme; ARB: angiotensin receptor blocker; BP: blood pressure; CI: confidence interval; CKD: chronic kidney disease; eGFR: estimated glomerular filtration rate; KRT: kidney replacement therapy; FGF23: fibroblast growth factor-23; HR: hazard ratio; NGAL: neutrophil gelatinase-associated lipocalin; NH: non-Hispanic; NTproBNP: N-terminal pro-B-type natriuretic peptide; Ref: reference; SE: standard error

SI conversion: To convert Uric acid to µmol/L, multiply by 59.485. To convert Hemoglobin to g/L, multiply by 0.01. To convert serum phosphate to mmol/L, multiply by 0.323. For serum bicarbonate, 1 mEq/L is equivalent to 1 mmol/L. For urine potassium, 1 mEq/24h is equivalent to 1 mmol/d.

^a Negative coefficients represent a faster decline in eGFR compared to the reference group.

^b HRs and p-values for factors with significant interactions with time are depicted separately for years 0 through 6 (Y0-6) and 6+ years (Y6+).

Table S2b. Final multivariable-adjusted models of chronic kidney disease progression among CRIC participants with diabetes after exclusion of albuminuria

| | | With Diabetes | | | |
|--|---------------------------|--|---------|--------------------------|------------------------------------|
| | | eGFR Slope, mL/min/1.73m ² /year | | KRT/eGFR Halving | |
| | | Beta Coefficient ^a (SE) | P value | HR (95% CI) ^b | P value ^b |
| Demographic Factors | | | | | |
| Age, years | <45 | -1.0 (0.3) | <.001 | Y0-6: 1.5 (1.1, 2.1) | <.001 |
| | 45-64 | Ref | | Y6+: 2.5 (1.5, 4.0) | |
| | 65-75 | 0.3 (0.2) | | Ref | |
| | | | | Y0-6: 0.6 (0.5, 0.8) | |
| | | | | Y6+: 0.7 (0.5, 1.1) | |
| Gender | Male | Ref | <.001 | Ref | <.001 |
| | Female | 0.7 (0.2) | | 0.6 (0.5, 0.7) | |
| Race/ ethnicity | NH-White | Ref | <.001 | Ref | <.001 |
| | NH-Black, APOL1 low-risk | -0.4 (0.2) | | 1.3 (1.0, 1.6) | |
| | NH-Black, APOL1 high-risk | -1.5 (0.3) | | 1.8 (1.3, 2.5) | |
| | Hispanic | -0.8 (0.2) | | 1.7 (1.2, 2.2) | |
| | Other | -0.5 (0.3) | | 1.5 (1.0, 2.2) | |
| Kidney Function Measure | | | | | |
| Baseline eGFR, mL/min/ 1.73m ² | <30 | 0.6 (0.2) | <.001 | Y0-6: 1.4 (1.1, 1.8) | Y0-6: <.001 Y6+: .004 |
| | 30-44.9 | Ref | | Y6+: 2.1 (1.3, 3.4) | |
| | 45-59.9 | -0.4 (0.2) | | Ref | |
| | | | | Y0-6: 0.7 (0.5, 0.9) | |
| | ≥60 | -0.6 (0.2) | | Y6+: 0.8 (0.5, 1.2) | |
| Blood Pressure Factor | | | | | |
| Systolic BP, mmHg | < 120 | Ref | <.001 | Ref | <.001 |
| | 120-139 | -0.5 (0.2) | | 1.6 (1.3, 2.0) | |
| | ≥ 140 | -1.2 (0.2) | | 2.4 (1.9, 3.0) | |
| Clinical Factors | | | | | |
| History of CVD | No | - | - | Ref | .04 |
| | Yes | - | | 1.2 (1.0, 1.4) | |

| | | With Diabetes | | | |
|---|----------------------|--|---------|--------------------------|----------------------|
| | | eGFR Slope, mL/min/1.73m ² /year | | KRT/eGFR Halving | |
| | | Beta Coefficient ^a (SE) | P value | HR (95% CI) ^b | P value ^b |
| Behavioral Factor | | | | | |
| Current smoker | No | Ref | .07 | Ref | .03 |
| | Yes | -0.4 (0.2) | | 1.3 (1.0, 1.7) | |
| Inflammatory Marker and Chemokines | | | | | |
| Serum fractalkine (CX3CL1), pg/mL | Q1: [0.1, 0.6) | Ref | <.001 | Ref | <.001 |
| | Q2: [0.6, 0.8) | -0.2 (0.2) | | 1.3 (1.0, 1.9) | |
| | Q3: [0.8, 1.1) | -0.7 (0.2) | | 1.8 (1.3, 2.4) | |
| | Q4: [1.1, 3.6] | -0.7 (0.2) | | 1.9 (1.4, 2.5) | |
| Plasma CXCL12, pg/mL | Q1: [832.1, 2066.4) | - | - | Ref | Y0-6: .02 Y6+: .3 |
| | Q2: [2066.4, 2410.5) | - | | Y0-6: 1.2 (0.9, 1.7) | |
| | Q3: [2410.5, 2797.8) | - | | Y6+: 1.2 (0.7, 2.0) | |
| | Q4: [2797.8, 6173.3] | - | | Y0-6: 1.6 (1.2, 2.2) | |
| | | | | Y6+: 0.9 (0.5, 1.5) | |
| Mineral Metabolism Markers | | | | | |
| Intact PTH pg/mL | Q1: [1.9, 34.1) | 0.3 (0.2) | .08. | 1.1 (0.8, 1.5) | .05 |
| | Q2: [34.1, 52.4) | Ref | | Ref | |
| | Q3: [52.4, 84.7) | -0.1 (0.2) | | 1.1 (0.9, 1.5) | |
| | Q4: [84.5, 1483] | -0.2 (0.2) | | 1.4 (1.1, 1.8) | |
| Carbohydrate Metabolism Markers | | | | | |
| HbA1c, % | Q1: [3.5, 5.6) | Ref | .003 | - | - |
| | Q2: [5.6, 6.2) | 0.5 (0.4) | | - | |
| | Q3: [6.2, 7.3) | 0.2 (0.3) | | - | |
| | Q4: [7.3, 15.2] | -0.2 (0.3) | | - | |
| Cardiac Markers | | | | | |
| hsTnT, pg/mL | Q1: [1.5, 5.5) | - | - | Ref | .009 |
| | Q2: [5.5, 11.4) | - | | 1.1 (0.7, 1.6) | |
| | Q3: (11.4, 22.2) | - | | 1.4 (0.9, 2.0) | |
| | Q4: (22.2, 738.7) | - | | 1.7 (1.1, 2.5) | |

| | | With Diabetes | | | | | |
|--|---------------------|--|---------|--------------------------|-----------------------------------|--|--|
| | | eGFR Slope, mL/min/1.73m ² /year | | KRT/eGFR Halving | | | |
| | | Beta Coefficient ^a (SE) | P value | HR (95% CI) ^b | P value ^b | | |
| NTproBNP, pg/mL | Q1: [2.5, 60.2] | Ref | <.001 | Ref | Y0-6: <.001 Y6+: .4 | | |
| | Q2: [60.2, 139.4] | -0.6 (0.2) | | Y0-6: 1.8 (1.2, 2.7) | | | |
| | Q3: [139.4, 366.6) | -0.7 (0.2) | | Y6+: 1.1 (0.6, 1.9) | | | |
| | Q4: [366.6, 33 742] | -1.0 (0.2) | | Y0-6: 1.7 (1.1, 2.6) | | | |
| | | | | Y6+: 1.4 (0.8, 2.3) | | | |
| | | | | Y0-6: 2.3 (1.5, 3.4) | | | |
| | | | | Y6+: 1.5 (0.9, 2.6) | | | |
| Acidosis Marker | | | | | | | |
| Serum bicarbonate, mmol/L | ≤22 | - | - | Y0-6: 0.8 (0.6, 1.1) | Y0-6: .05 Y6+: .2 | | |
| | (22, 24] | - | | Y6+: 0.6 (0.4, 1.0) | | | |
| | (24, 26] | - | | Ref | | | |
| | | - | | Y0-6: 1.0 (0.7, 1.3) | | | |
| | >26 | - | | Y6+: 0.9 (0.5, 1.4) | | | |
| Kidney Injury Marker | | | | | | | |
| Urine NGAL, ng/ml | Q1: [0.4, 6.3) | Ref | <.001 | Ref | Y0-6: <.001 Y6+: .02 | | |
| | Q2: [6.3, 14.2) | -0.5 (0.2) | | Y0-6: 1.3 (0.9, 1.8) | | | |
| | Q3: [14.2, 32.9) | -0.8 (0.2) | | Y6+: 1.0 (0.7, 1.7) | | | |
| | Q4: [32.9, 2743.8] | -1.3 (0.2) | | Y0-6: 1.6 (1.2, 2.3) | | | |
| | | | | Y6+: 1.2 (0.8, 1.9) | | | |
| Abbreviations: BP: blood pressure; CI: confidence interval; CKD: chronic kidney disease; CVD: cardiovascular disease; eGFR: estimated glomerular filtration rate; KRT: kidney replacement therapy; HbA1c: glycosylated hemoglobin; HR: hazard ratio; hsTnT: high-sensitivity troponin T; NGAL: neutrophil gelatinase-associated lipocalin; NH: non-Hispanic; NTproBNP: N-terminal pro-B-type natriuretic peptide; PTH: parathyroid hormone; Ref: reference; SE: standard error | | | | | | | |
| N/A indicates variable was excluded from analysis for indicated outcome due to multicollinearity with the outcome measure. | | | | | | | |
| SI conversion: To convert serum bicarbonate, 1 mEq/L is equivalent to 1 mmol/L. | | | | | | | |

^a Negative coefficients represent a faster decline in eGFR compared to the reference group.^b HRs and p-values for factors with significant interactions with time are depicted separately for years 0 through 6 (Y0-6) and 6+ years (Y6+).

Table S3a. Multivariable-adjusted models of chronic kidney disease progression with albuminuria interaction among CRIC participants without diabetes

| | | Without Diabetes | | | |
|---|---------------------------------------|--|---------|---------------------------------------|---------|
| | | eGFR Slope, mL/min/1.73m ² /year | | KRT/eGFR Halving | |
| | | Beta Coefficient ^a (SE) | P value | Beta Coefficient ^b (SE) | P value |
| Demographic Factors | | | | | |
| Gender | Male | Ref | <.001 | Ref | .004 |
| | Female | 0.3 (0.1) | | -1.3 (0.4) | |
| Gender* UACR | Male*<30 UACR | Ref | .2 | Ref | .4 |
| | Female*30-299 UACR | 0.0 (0.2) | | 0.7 (0.5) | |
| | Female*≥300 UACR | 0.6 (0.3) | | 0.6 (0.5) | |
| Race/ ethnicity | NH-White | Ref | <.001 | Ref | .1 |
| | NH-Black, APOL1 low-risk | 0.2 (0.1) | | 0.0 (0.4) | |
| | NH-Black, APOL1 high-risk | -0.1 (0.2) | | 0.9 (0.5) | |
| | Hispanic | 0.2 (0.3) | | 0.9 (0.5) | |
| | Other | -0.3 (0.3) | | 1.1 (0.8) | |
| Race/ ethnicity* UACR | NH-White*<30 UACR | Ref | <.001 | Ref | .4 |
| | NH-Black, APOL1 low-risk*30-299 UACR | -0.6 (0.3) | | 0.8 (0.4) | |
| | NH-Black, APOL1 high-risk*30-299 UACR | -0.8 (0.4) | | 0.7 (0.6) | |
| | Hispanic*30-299 UACR | 0.6 (0.4) | | -0.8 (0.7) | |
| | Other*30-299 UACR | -0.7 (0.5) | | 0.4 (0.9) | |
| | NH-Black, APOL1 low-risk*≥300 UACR | -1.1 (0.3) | | 0.6 (0.4) | |
| | NH-Black, APOL1 high-risk*≥300 UACR | -2.2 (0.5) | | 0.2 (0.6) | |
| | Hispanic*≥300 UACR | -0.5 (0.4) | | -0.6 (0.6) | |
| Kidney Function Measures | | | | | |
| Baseline eGFR, mL/min/ 1.73m ² | <30 | 0.1 (0.3) | .004 | 0.9 (0.4) | <.001 |
| | 30-44.9 | Ref | | Ref | |
| | 45-59.9 | 0.1 (0.2) | | -1.0 (0.4) | |
| | ≥60 | 0.3 (0.2) | | -1.5 (0.5) | |
| | 30-44.9 eGFR*<30 UACR | Ref | .1 | Ref | .6 |

| | | Without Diabetes | | | |
|----------------------------------|--------------------------|--|---------|---------------------------------------|---------|
| | | eGFR Slope, mL/min/1.73m ² /year | | KRT/eGFR Halving | |
| | | Beta Coefficient ^a (SE) | P value | Beta Coefficient ^b (SE) | P value |
| Baseline eGFR*UACR | <30 eGFR*30-299 UACR | 0.1 (0.4) | <.001 | -0.4 (0.5) | .2 |
| | 45-59.9 eGFR*30-299 UACR | -0.1 (0.3) | | 0.3 (0.5) | |
| | ≥60 eGFR*30-299 UACR | 0.1 (0.3) | | 0.1 (0.7) | |
| | <30 eGFR*≥300 UACR | 0.5 (0.4) | | -0.3 (0.4) | |
| | 45-59.9 eGFR*≥300 UACR | -0.8 (0.3) | | 0.6 (0.4) | |
| | ≥60 eGFR*≥300 UACR | 0.0 (0.5) | | -0.4 (0.8) | |
| UACR, mg/g | <30 | Ref | <.001 | Ref | .2 |
| | 30-299 | -0.3 (0.5) | | 1.7 (0.9) | |
| | ≥300 | 0.6 (0.6) | | 1.5 (0.9) | |
| Blood Pressure Factors | | | | | |
| Systolic BP, mmHg | <120 | Ref | <.001 | Ref | .2 |
| | 120-139 | -0.1 (0.1) | | 0.3 (0.3) | |
| | ≥140 | -0.4 (0.2) | | 0.7 (0.4) | |
| Systolic BP* UACR | <120 BP*<30 UACR | Ref | .06 | Ref | .04 |
| | 120-139 BP*30-299 UACR | -0.1 (0.2) | | -0.4 (0.4) | |
| | ≥140 BP*30-299 UACR | 0.3 (0.3) | | -0.9 (0.5) | |
| | 120-139 BP*≥300 UACR | -0.4 (0.3) | | 0.1 (0.4) | |
| | ≥140 BP*≥300 UACR | -0.9 (0.4) | | 0.1 (0.5) | |
| ACE/ARB | No | Ref | .6 | - | N/A |
| | Yes | 0.2 (0.1) | | - | |
| ACE/ARB* UACR | No*<30 UACR | Ref | .5 | - | N/A |
| | Yes*30-299 UACR | -0.2 (0.2) | | - | |
| | Yes*≥300 UACR | -0.2 (0.3) | | - | |
| Body Composition Measures | | | | | |
| Fat-free mass, kg | Q1: [25.8, 49.1) | - | N/A | Ref | .6 |
| | Q2: [49.1, 59.1) | - | | -0.4 (0.4) | |
| | Q3: [59.1, 69.9) | - | | -0.6 (0.5) | |
| | Q4: [69.9, 167.1) | - | | -0.7 (0.6) | |
| Fat-free mass*UACR | Q1 FFM*<30 UACR | - | N/A | Ref | .1 |
| | Q2 FFM*30-299 UACR | - | | 0.3 (0.5) | |

| | Without Diabetes | | | |
|-----------------------------------|--|------------|---------------------------------------|------------|
| | eGFR Slope, mL/min/1.73m ² /year | | KRT/eGFR Halving | |
| | Beta Coefficient ^a (SE) | P value | Beta Coefficient ^b (SE) | P value |
| Q3 FFM*30-299 UACR | - | | 0.5 (0.6) | |
| | - | | 0.5 (0.7) | |
| | - | | -0.2 (0.5) | |
| | - | | 0.2 (0.6) | |
| | - | | 1.0 (0.6) | |
| Measure of Anemia | | | | |
| Hemoglobin, mg/dL | M: <12, F: <11 | - | N/A | 0.3 (0.3) |
| | M: 12-13.9, F: 11-12.9 | - | | Ref |
| | M: ≥14, F: ≥13 | - | | -0.7 (0.4) |
| Hemoglobin* UACR | M: 12-13.9, F: 11-12.9 Hemoglobin*<30 UACR | - | N/A | Ref |
| | M: <12, F: <11 Hemoglobin *30-299 UACR | - | | -0.2 (0.4) |
| | M: ≥14, F: ≥13 Hemoglobin*30-299 UACR | - | | -0.2 (0.5) |
| | M: <12, F: <11 Hemoglobin*≥300 UACR | - | | -0.2 (0.4) |
| | M: ≥14, F: ≥13 Hemoglobin*≥300 UACR | - | | 0.8 (0.4) |
| Mineral Metabolism Markers | | | | |
| Serum phosphate, mg/dL | Q1: [1.7, 3.3) | -0.2 (0.1) | .05 | - |
| | Q2: [3.3, 3.7) | Ref | | - |
| | Q3: [3.7, 4.1) | 0.1 (0.2) | | - |
| | Q4: [4.1, 9.3] | -0.1 (0.2) | | - |
| Serum phosphate* UACR | Q2 phosphate*<30 UACR | Ref | .3 | - |
| | Q1 phosphate*30-299 UACR | -0.3 (0.3) | | - |
| | Q3 phosphate*30-299 UACR | -0.1 (0.3) | | - |
| | Q4 phosphate*30-299 UACR | 0.2 (0.3) | | - |
| | Q1 phosphate*≥300 UACR | -0.2 (0.3) | | - |

| | Without Diabetes | | | |
|-------------------------------|--|------------|---------------------------------------|------------|
| | eGFR Slope, mL/min/1.73m ² /year | | KRT/eGFR Halving | |
| | Beta Coefficient ^a (SE) | P value | Beta Coefficient ^b (SE) | P value |
| Q3 phosphate* \geq 300 UACR | -0.7 (0.4) | | - | |
| | -0.5 (0.4) | | - | |
| Cardiac Markers | | | | |
| hsTnT, pg/mL | Q1: [1.5, 5.5) | - | N/A | Ref |
| | Q2: [5.5, 11.4) | - | | 0.7 (0.4) |
| | Q3: (11.4, 22.2) | - | | -0.7 (0.5) |
| | Q4: (22.2, 738.7) | - | | -0.3 (0.6) |
| hsTnT*UACR | Q1 hsTNT*<30 UACR | - | N/A | Ref |
| | Q2 hsTNT*30-299 UACR | - | | -0.8 (0.5) |
| | Q3 hsTNT*30-299 UACR | - | | 0.2 (0.6) |
| | Q4 hsTNT*30-299 UACR | - | | 0.1 (0.7) |
| | Q2 hsTNT* \geq 300 UACR | - | | -0.5 (0.4) |
| | Q3 hsTNT* \geq 300 UACR | - | | 0.6 (0.5) |
| | Q4 hsTNT* \geq 300 UACR | - | | 0.3 (0.7) |
| NTproBNP, pg/mL | Q1: [2.5, 60.2) | Ref | .01 | Ref |
| | Q2: [60.2, 139.4) | 0.0 (0.1) | | 0.4 (0.5) |
| | Q3: [139.4, 366.6) | -0.2 (0.2) | | 0.9 (0.5) |
| | Q4: [366.6, 33 742] | -0.1 (0.2) | | 1.3 (0.5) |
| NTproBNP* UACR | Q1 NTproBNP*<30 UACR | Ref | .005 | Ref |
| | Q2 NTproBNP*30-299 UACR | -0.2 (0.3) | | -0.3 (0.5) |
| | Q3 NTproBNP*30-299 UACR | -0.6 (0.3) | | -0.4 (0.5) |
| | Q4 NTproBNP*30-299 UACR | -0.4 (0.4) | | -1.0 (0.6) |
| | Q2 NTproBNP* \geq 300 UACR | -1.0 (0.3) | | 0.4 (0.5) |
| | Q3 NTproBNP* \geq 300 UACR | -0.1 (0.4) | | -0.5 (0.5) |
| | Q4 NTproBNP* \geq 300 UACR | 0.3 (0.4) | | -1.1 (0.6) |
| | \leq 22 | - | N/A | 0.4 (0.4) |
| | | | | .03 |

| | | Without Diabetes | | | |
|---------------------------------|--------------------------------------|--|---------|---------------------------------------|---------|
| | | eGFR Slope, mL/min/1.73m ² /year | | KRT/eGFR Halving | |
| | | Beta Coefficient ^a (SE) | P value | Beta Coefficient ^b (SE) | P value |
| Serum bicarbonate, mmol/L | (22, 24] | - | | Ref | |
| | (24, 26] | - | | -0.9 (0.5) | |
| | >26 | - | | -0.2 (0.4) | |
| Serum bicarbonate* UACR | ≤22 bicarbonate*<30 UACR | - | N/A | Ref | .4 |
| | (22, 24] bicarbonate*30- 299 UACR | - | | 0.1 (0.4) | |
| | (24, 26] bicarbonate*30- 299 UACR | - | | 1.2 (0.6) | |
| | >26 bicarbonate*30-299 UACR | - | | 0.4 (0.5) | |
| | (22, 24] bicarbonate*≥300 UACR | - | | -0.2 (0.4) | |
| | (24, 26] bicarbonate*≥300 UACR | - | | 0.8 (0.5) | |
| | >26 bicarbonate*≥300 UACR | - | | 0.1 (0.5) | |
| Urinary Electrolytes | | | | | |
| Urine potassium, mmol/24h | Q1: [3.0, 37.5) | Ref | .4 | - | N/A |
| | Q2: [37.5, 51.8) | -0.1 (0.2) | | - | |
| | Q3: [51.8, 69.4) | -0.2 (0.2) | | - | |
| | Q4: [69.4, 417.7] | 0.2 (0.2) | | - | |
| Urine potassium* UACR | Q1 potassium*<30 UACR | Ref | .2 | - | N/A |
| | Q2 potassium*30-299 UACR | 0.4 (0.3) | | - | |
| | Q3 potassium*30-299 UACR | 0.2 (0.3) | | - | |
| | Q4 potassium*30-299 UACR | 0.2 (0.3) | | - | |
| | Q2 potassium*≥300 UACR | -0.5 (0.4) | | - | |
| | Q3 potassium*≥300 UACR | -0.4 (0.4) | | - | |
| | Q4 potassium*≥300 UACR | -0.9 (0.4) | | - | |

| | Without Diabetes | | | |
|--------------------------------|--|------------|---------------------------------------|------------|
| | eGFR Slope, mL/min/1.73m ² /year | | KRT/eGFR Halving | |
| | Beta Coefficient ^a (SE) | P value | Beta Coefficient ^b (SE) | P value |
| Kidney Injury Marker | | | | |
| Urine NGAL, ng/ml | Q1: [0.4, 6.3) | Ref | .02 | Ref |
| | Q2: [6.3, 14.2) | -0.2 (0.1) | | 0.7 (0.4) |
| | Q3: [14.2, 32.9) | -0.2 (0.2) | | 0.4 (0.5) |
| | Q4: [32.9, 2743.8] | -0.2 (0.2) | | 0.8 (0.5) |
| Urine NGAL*UACR | Q1 NGAL*<30 UACR | Ref | .09 | Ref |
| | Q2 NGAL*30-299 UACR | -0.1 (0.3) | | -0.4 (0.5) |
| | Q3 NGAL*30-299 UACR | 0.0 (0.3) | | -0.3 (0.6) |
| | Q4 NGAL*30-299 UACR | 0.3 (0.3) | | -0.9 (0.6) |
| | Q2 NGAL*≥300 UACR | -0.4 (0.4) | | 0.0 (0.5) |
| | Q3 NGAL*≥300 UACR | -0.6 (0.4) | | 0.2 (0.6) |
| | Q4 NGAL*≥300 UACR | -1.2 (0.4) | | 0.2 (0.6) |
| RAAS Marker | | | | |
| Serum aldosterone, pg/mL | Q1: [0.8, 71.2) | Ref | .03 | - |
| | Q2: [71.2, 101.4) | -0.1 (0.2) | | - |
| | Q3: [101.4, 152.8) | -0.2 (0.2) | | - |
| | Q4: [152.8, 15 630.9] | 0.2 (0.2) | | - |
| Serum aldosterone* UACR | Q1 aldosterone*<30 UACR | Ref | .3 | - |
| | Q2 aldosterone*30-299 UACR | -0.2 (0.3) | | - |
| | Q3 aldosterone*30-299 UACR | 0.0 (0.3) | | - |
| | Q4 aldosterone*30-299 UACR | -0.3 (0.3) | | - |
| | Q2 aldosterone*≥300 UACR | -0.8 (0.4) | | - |
| | Q3 aldosterone*≥300 UACR | -0.2 (0.4) | | - |
| | Q4 aldosterone*≥300 UACR | -0.6 (0.4) | | - |

Abbreviations: ACE: angiotensin converting enzyme; ARB: angiotensin receptor blocker; BP: blood pressure; CKD: chronic kidney disease; eGFR: estimated glomerular filtration rate; KRT: kidney replacement therapy; FFM: fat-free mass; hsTnT: high-sensitivity troponin T; NGAL: neutrophil gelatinase-associated lipocalin; NH: non-Hispanic; NTproBNP: N-terminal pro-B-type natriuretic peptide; RAAS: renin-angiotensin-aldosterone system; Ref: reference; SE: standard error; UACR: urine albumin:creatinine ratio

N/A indicates variable was excluded from analysis for indicated outcome as only variables selected by models described in Table 2 were tested for interaction.

SI conversion: To convert Hemoglobin to g/L, multiply by 0.01. To convert serum phosphate to mmol/L, multiply by 0.323. For serum bicarbonate, 1 mEq/L is equivalent to 1 mmol/L. For urine potassium, 1 mEq/24h is equivalent to 1 mmol/d. To convert serum aldosterone to pmol/L, multiply by 2.774.

^a Negative coefficients represent a faster decline in eGFR compared to the reference group.

^b Coefficients are the log hazard ratio to the reference level with positive coefficients indicating increased risk of KRT/eGFR Halving compared to the reference group.

* Indicates interaction between variables. Sum of main and interaction beta coefficients will provide estimate for given group.

Table S3b. Multivariable-adjusted models of chronic kidney disease progression with albuminuria interaction among CRIC participants with diabetes

| | | With Diabetes | | | |
|-----------------------------|---------------------------------------|--|---------|---------------------------------------|---------|
| | | eGFR Slope, mL/min/1.73m ² /year | | KRT/eGFR Halving | |
| | | Beta Coefficient ^a (SE) | P value | Beta Coefficient ^b (SE) | P value |
| Demographic Factors | | | | | |
| Age, years | <45 | -0.3 (0.4) | .03 | 0.1 (0.6) | .8 |
| | 45-64 | Ref | | Ref | |
| | 65-75 | -0.2 (0.2) | | -0.2 (0.3) | |
| Age*UACR | 45-64 age*<30 UACR | Ref | .2 | Ref | .9 |
| | <45 age*30-299 UACR | -0.5 (0.7) | | -0.2 (0.7) | |
| | 65-75 age*30-299 UACR | 0.0 (0.3) | | -0.1 (0.4) | |
| | <45 age*≥300 UACR | -0.3 (0.5) | | 0.2 (0.6) | |
| | 65-75*≥300 UACR | 0.8 (0.3) | | -0.1 (0.3) | |
| Gender | Male | - | N/A | Ref | .5 |
| | Female | - | | -0.2 (0.3) | |
| Gender* UACR | Male*<30 UACR | - | N/A | Ref | .8 |
| | Female*30-299 UACR | - | | -0.1 (0.4) | |
| | Female*≥300 UACR | - | | -0.2 (0.3) | |
| Race/ ethnicity | NH-White | Ref | <.001 | Ref | .4 |
| | NH-Black, APOL1 low-risk | -0.1 (0.2) | | -0.1 (0.3) | |
| | NH-Black, APOL1 high-risk | -0.2 (0.5) | | 1.0 (0.5) | |
| | Hispanic | -0.1 (0.3) | | 0.2 (0.5) | |
| | Other | -0.5 (0.5) | | 0.3 (0.6) | |
| Race/ ethnicity* UACR | NH-White*<30 UACR | Ref | .02 | Ref | .4 |
| | NH-Black, APOL1 low-risk*30-299 UACR | -0.8 (0.3) | | 0.7 (0.4) | |
| | NH-Black, APOL1 high-risk*30-299 UACR | -1.2 (0.7) | | -0.1 (0.6) | |
| | Hispanic*30-299 UACR | -1.3 (0.5) | | 0.7 (0.6) | |
| | Other*30-299 UACR | 0.4 (0.7) | | -0.3 (0.8) | |
| | NH-Black, APOL1 low-risk*≥300 UACR | -0.7 (0.3) | | 0.4 (0.4) | |
| | NH-Black, APOL1 high-risk*≥300 UACR | -1.1 (0.6) | | -0.5 (0.5) | |

| | | With Diabetes | | | |
|---|---------------------------------|--|---------|---------------------------------------|---------|
| | | eGFR Slope, mL/min/1.73m ² /year | | KRT/eGFR Halving | |
| | | Beta Coefficient ^a (SE) | P value | Beta Coefficient ^b (SE) | P value |
| Hispanic* \geq 300 UACR | <30 | -0.3 (0.4) | | 0.0 (0.6) | |
| | Other* \geq 300 UACR | 0.3 (0.7) | | 0.2 (0.7) | |
| Kidney Function Measures | | | | | |
| Baseline eGFR, mL/min/ 1.73m ² | <30 | 0.2 (0.3) | <.001 | 1.2 (0.4) | <.001 |
| | 30-44.9 | Ref | | Ref | |
| | 45-59.9 | -0.2 (0.2) | | -0.6 (0.4) | |
| | \geq 60 | -0.7 (0.3) | | -0.4 (0.5) | |
| Baseline eGFR*UACR | <30 eGFR*<30 UACR | Ref | .1 | Ref | .1 |
| | 30-44.9 eGFR*30-299 UACR | 0.3 (0.5) | | -0.6 (0.4) | |
| | 45-59.9 eGFR*30-299 UACR | -0.3 (0.3) | | 0.4 (0.4) | |
| | \geq 60 eGFR*30-299 UACR | 0.2 (0.5) | | 0.0 (0.6) | |
| | 30-44.9 eGFR* \geq 300 UACR | 0.9 (0.4) | | -0.9 (0.4) | |
| | 45-59.9 eGFR* \geq 300 UACR | -0.4 (0.4) | | 0.4 (0.4) | |
| | \geq 60 eGFR* \geq 300 UACR | -0.5 (0.5) | | -0.1 (0.6) | |
| UACR, mg/g | <30 | Ref | <.001 | Ref | .1 |
| | 30-299 | -1.2 (0.5) | | 1.7 (1.3) | |
| | \geq 300 | -1.5 (0.7) | | 2.4 (1.2) | |
| Blood Pressure Factors | | | | | |
| Systolic BP, mmHg | <120 | Ref | .002 | Ref | .005 |
| | 120-139 | -0.2 (0.2) | | 0.5 (0.3) | |
| | \geq 140 | -0.5 (0.3) | | 1.2 (0.4) | |
| Systolic BP* UACR | <120 BP*<30 UACR | Ref | .1 | Ref | .1 |
| | 120-139 BP*30-299 UACR | 0.3 (0.3) | | -0.6 (0.4) | |
| | \geq 140 BP*30-299 UACR | 0.4 (0.4) | | -1.2 (0.4) | |
| | 120-139 BP* \geq 300 UACR | -0.3 (0.4) | | -0.3 (0.4) | |
| | \geq 140 BP* \geq 300 UACR | -0.8 (0.4) | | -0.7 (0.4) | |
| Behavioral Factor | | | | | |
| Current smoker | No | Ref | .07 | - | N/A |
| | Yes | -0.3 (0.3) | | - | |

| | | With Diabetes | | | |
|--|-----------------------|--|---------|---------------------------------------|---------|
| | | eGFR Slope, mL/min/1.73m ² /year | | KRT/eGFR Halving | |
| | | Beta Coefficient ^a (SE) | P value | Beta Coefficient ^b (SE) | P value |
| Current smoker* UACR | No*<30 UACR | Ref | .9 | - | N/A |
| | Yes*30-299 UACR | -0.1 (0.5) | | - | |
| | Yes*≥300 UACR | 0.0 (0.5) | | - | |
| Inflammatory Marker and Chemokines | | | | | |
| Serum fractalkine (CX3CL1), pg/mL | Q1: [0.1, 0.6) | Ref | .01 | Ref | .2 |
| | Q2: [0.6, 0.8) | 0.0 (0.2) | | -0.3 (0.4) | |
| | Q3: [0.8, 1.1) | -0.4 (0.2) | | 0.4 (0.4) | |
| | Q4: [1.1, 3.6] | -0.4 (0.3) | | 0.5 (0.4) | |
| Serum fractalkine (CX3CL1)* UACR | Q1 CX3CL1*<30 UACR | Ref | .6 | Ref | .7 |
| | Q2 CX3CL1*30-299 UACR | 0.3 (0.4) | | 0.4 (0.5) | |
| | Q3 CX3CL1*30-299 UACR | 0.3 (0.4) | | 0.0 (0.5) | |
| | Q4 CX3CL1*30-299 UACR | 0.5 (0.4) | | -0.4 (0.5) | |
| | Q2 CX3CL1*≥300 UACR | 0.3 (0.5) | | 0.2 (0.5) | |
| | Q3 CX3CL1*≥300 UACR | -0.2 (0.5) | | -0.2 (0.5) | |
| | Q4 CX3CL1*≥300 UACR | 0.4 (0.5) | | -0.4 (0.4) | |
| Plasma CXCL12, pg/mL | Q1: [832.1, 2066.4) | - | N/A | Ref | .3 |
| | Q2: [2066.4, 2410.5) | - | | 0.1 (0.4) | |
| | Q3: [2410.5, 2797.8) | - | | 0.5 (0.5) | |
| | Q4: [2797.8, 6173.3] | - | | 0.7 (0.4) | |
| Plasma CXCL12* UACR | Q1 CXCL12*<30 UACR | - | N/A | Ref | .6 |
| | Q2 CXCL12*30-299 UACR | - | | -0.3 (0.5) | |
| | Q3 CXCL12*30-299 UACR | - | | -0.5 (0.5) | |
| | Q4 CXCL12*30-299 UACR | - | | -0.5 (0.5) | |
| | Q2 CXCL12*≥300 UACR | - | | 0.3 (0.5) | |
| | Q3 CXCL12*≥300 UACR | - | | 0.0 (0.5) | |
| | Q4 CXCL12*≥300 UACR | - | | -0.2 (0.5) | |
| Carbohydrate Metabolism Markers | | | | | |
| HbA1c, % | Q1: [3.5, 5.6) | - | N/A | Ref | .6 |
| | Q2: [5.6, 6.2) | - | | 0.7 (0.8) | |
| | Q3: [6.2, 7.3) | - | | 0.2 (0.8) | |
| | Q4: [7.3, 15.2] | - | | 0.6 (0.8) | |

| | | With Diabetes | | | |
|-----------------------------|----------------------------|--|---------|---------------------------------------|---------|
| | | eGFR Slope, mL/min/1.73m ² /year | | KRT/eGFR Halving | |
| | | Beta Coefficient ^a (SE) | P value | Beta Coefficient ^b (SE) | P value |
| HbA1c* UACR | Q1 HbA1c*<30 UACR | - | N/A | Ref | .9 |
| | Q2 HbA1c*30-299 UACR | - | | 0.0 (1.0) | |
| | Q3 HbA1c*30-299 UACR | - | | 0.1 (1.0) | |
| | Q4 HbA1c*30-299 UACR | - | | -0.1 (0.9) | |
| | Q2 HbA1c*≥300 UACR | - | | -0.3 (0.8) | |
| | Q3 HbA1c*≥300 UACR | - | | -0.3 (0.8) | |
| | Q4 HbA1c*≥300 UACR | - | | -0.6 (0.8) | |
| Cardiac Markers | | | | | |
| NTproBNP, pg/mL | Q1: [2.5, 60.2) | Ref | .01 | Ref | .04 |
| | Q2: [60.2, 139.4) | -0.5 (0.2) | | 0.9 (0.5) | |
| | Q3: [139.4, 366.6) | -0.6 (0.3) | | 0.8 (0.5) | |
| | Q4: [366.6, 33 742] | -0.7 (0.3) | | 1.4 (0.5) | |
| NTproBNP* UACR | Q1 NTproBNP*<30 UACR | Ref | .2 | Ref | .1 |
| | Q2 NTproBNP*30-299 UACR | -0.2 (0.4) | | -0.2 (0.6) | |
| | Q3 NTproBNP*30-299 UACR | 0.1 (0.4) | | -0.1 (0.6) | |
| | Q4 NTproBNP*30-299 UACR | 0.1 (0.4) | | -0.9 (0.6) | |
| | Q2 NTproBNP*≥300 UACR | 0.6 (0.4) | | -0.7 (0.5) | |
| | Q3 NTproBNP*≥300 UACR | 1.0 (0.5) | | -0.7 (0.5) | |
| | Q4 NTproBNP*≥300 UACR | 0.3 (0.5) | | -0.9 (0.6) | |
| Kidney Injury Marker | | | | | |
| Urine NGAL, ng/ml | Q1: [0.4, 6.3) | Ref | <.001 | Ref | .5 |
| | Q2: [6.3, 14.2) | -0.3 (0.2) | | 0.3 (0.4) | |
| | Q3: [14.2, 32.9) | -0.3 (0.2) | | -0.2 (0.4) | |
| | Q4: [32.9, 2743.8] | -0.2 (0.3) | | -0.1 (0.5) | |
| Urine NGAL* UACR | Q1 NGAL*<30 UACR | Ref | <.001 | Ref | .003 |
| | Q2 NGAL*30-299 UACR | 0.3 (0.4) | | -0.7 (0.4) | |
| | Q3 NGAL*30-299 UACR | 0.1 (0.4) | | 0.1 (0.5) | |
| | Q4 NGAL*30-299 UACR | 0.4 (0.4) | | 0.4 (0.6) | |
| | Q2 NGAL*≥300 UACR | -0.7 (0.4) | | 0.3 (0.4) | |

| | With Diabetes | | | |
|-------------------|--|---------|---------------------------------------|---------|
| | eGFR Slope, mL/min/1.73m ² /year | | KRT/eGFR Halving | |
| | Beta Coefficient ^a (SE) | P value | Beta Coefficient ^b (SE) | P value |
| Q3 NGAL*≥300 UACR | -1.3 (0.4) | | 1.0 (0.4) | |
| Q4 NGAL*≥300 UACR | -1.9 (0.4) | | 1.5 (0.5) | |

Abbreviations: BP: blood pressure; CKD: chronic kidney disease; eGFR: estimated glomerular filtration rate; KRT: kidney replacement therapy; HbA1c: glycosylated hemoglobin; NGAL: neutrophil gelatinase-associated lipocalin; NH: non-Hispanic; NTproBNP: N-terminal pro-B-type natriuretic peptide; Ref: reference; SE: standard error; UACR: urine albumin:creatinine ratio

N/A indicates variable was excluded from analysis for indicated outcome as only variables selected by models described in Table 3 were tested for interaction.

^a Negative coefficients represent a faster decline in eGFR compared to the reference group.

^b Coefficients are the log hazard ratio to the reference level with positive coefficients indicating increased risk of KRT/eGFR Halving compared to the reference group.

* Indicates interaction between variables. Sum of main and interaction beta coefficients will provide estimate for given group.

Table S4a. Multivariable-adjusted models of chronic kidney disease progression with race interaction among CRIC participants without diabetes

| | | Without Diabetes | | | |
|--|-------------------------------------|--|---------|---------------------------------------|---------|
| | | eGFR Slope, mL/min/1.73m ² /year | | KRT/eGFR Halving | |
| | | Beta Coefficient ^a (SE) | P value | Beta Coefficient ^b (SE) | P value |
| Demographic Factors | | | | | |
| Gender | Male | Ref | .2 | Ref | .05 |
| | Female | 0.3 (0.1) | | -0.5 (0.3) | |
| Race/ ethnicity | NH-White | Ref | <.001 | Ref | .5 |
| | NH-Black, APOL1 low-risk | 0.8 (0.5) | | 0.3 (0.8) | |
| | NH-Black, APOL1 high-risk | -0.9 (0.8) | | 2.3 (1.3) | |
| | Hispanic | 1.2 (1.0) | | 0.4 (1.3) | |
| | Other | -1.2 (1.6) | | -2.1 (3.7) | |
| Gender*Race /ethnicity | Male*NH-White | Ref | .7 | Ref | .8 |
| | Female*NH-Black, APOL1 low-risk | 0.2 (0.2) | | -0.3 (0.4) | |
| | Female*NH-Black, APOL1 high-risk | -0.1 (0.4) | | -0.6 (0.5) | |
| | Female*Hispanic | -0.4 (0.5) | | -0.2 (0.9) | |
| | Female*Other | 0.1 (0.8) | | -1.9 (2.6) | |
| Kidney Function Measures | | | | | |
| Baseline eGFR, mL/min/ 1.73m ² | <30 | 0.2 (0.2) | .005 | 0.4 (0.2) | <.001 |
| | 30-44.9 | Ref | | Ref | |
| | 45-59.9 | 0.0 (0.2) | | -0.7 (0.2) | |
| | ≥60 | 0.1 (0.2) | | -1.6 (0.4) | |

| | | Without Diabetes | | | |
|----------------------------------|---|--|---------|---------------------------------------|---------|
| | | eGFR Slope, mL/min/1.73m ² /year | | KRT/eGFR Halving | |
| | | Beta Coefficient ^a (SE) | P value | Beta Coefficient ^b (SE) | P value |
| Baseline eGFR*Race/ ethnicity | 30-44.9 eGFR*NH-White | Ref | .5 | Ref | .2 |
| | <30 eGFR*NH-Black, APOL1 low-risk | 0.4 (0.4) | | 0.4 (0.4) | |
| | 45-59.9 eGFR*NH-Black, APOL1 low-risk | 0.0 (0.3) | | 0.2 (0.4) | |
| | ≥60 eGFR*NH-Black, APOL1 low-risk | 0.2 (0.3) | | 0.6 (0.6) | |
| | <30 eGFR*NH-Black, APOL1 high-risk | 1.2 (0.7) | | -0.1 (0.5) | |
| | 45-59.9 eGFR*NH-Black, APOL1 high-risk | 0.9 (0.5) | | -1.1 (0.5) | |
| | ≥60 eGFR*NH-Black, APOL1 high-risk | 0.7 (0.5) | | -0.6 (0.9) | |
| | <30 eGFR*Hispanic | 0.2 (0.7) | | 0.5 (0.5) | |
| | 45-59.9 eGFR*Hispanic | 0.3 (0.5) | | 0.5 (0.6) | |
| | ≥60 eGFR*Hispanic | 1.1 (0.7) | | -11.9 (709.0) | |
| | <30 eGFR*Other | 1.2 (1.7) | | -1.4 (2.5) | |
| | 45-59.9 eGFR*Other | -0.2 (1.0) | | -5.5 (2.1) | |
| | ≥60 eGFR*Other | 1.2 (1.1) | | -12.5 (5.0) | |
| UACR, mg/g | <30 | Ref | <.001 | Ref | <.001 |
| | 30-299 | -0.8 (0.1) | | 1.1 (0.3) | |
| | ≥300 | -1.7 (0.2) | | 2.4 (0.3) | |
| UACR*Race/ ethnicity | <30 UACR*NH-White | Ref | <.001 | Ref | .05 |
| | 30-299 UACR*NH-Black, APOL1 low-risk | -0.5 (0.2) | | 0.4 (0.4) | |
| | ≥300 UACR*NH-Black, APOL1 low-risk | -1.0 (0.3) | | 0.4 (0.4) | |
| | 30-299 UACR*NH-Black, APOL1 high-risk | -0.7 (0.4) | | 1.3 (0.6) | |
| | ≥300 UACR*NH-Black, APOL1 high-risk | -2.4 (0.6) | | 1.1 (0.7) | |
| | 30-299 UACR*Hispanic | 0.8 (0.5) | | -1.7 (0.8) | |
| | ≥300 UACR*Hispanic | -0.1 (0.5) | | -1.0 (0.7) | |
| | 30-299 UACR*Other | 0.3 (0.7) | | 1.7 (1.6) | |

| | | Without Diabetes | | | |
|-----------------------------------|---|--|---------|---------------------------------------|---------|
| | | eGFR Slope, mL/min/1.73m ² /year | | KRT/eGFR Halving | |
| | | Beta Coefficient ^a (SE) | P value | Beta Coefficient ^b (SE) | P value |
| | ≥300 UACR*Other | 0.1 (0.9) | | 4.4 (2.3) | |
| Blood Pressure Factors | | | | | |
| Systolic BP, mmHg | <120 | Ref | .3 | Ref | .3 |
| | 120-139 | -0.1 (0.1) | | 0.2 (0.2) | |
| | ≥140 | -0.5 (0.2) | | 0.5 (0.3) | |
| Systolic BP*Race/ ethnicity | <120 BP*NH-White | Ref | .3 | Ref | .03 |
| | 120-139 BP*NH-Black, APOL1 low-risk | -0.3 (0.2) | | 0.0 (0.3) | |
| | ≥140 BP*NH-Black, APOL1 low-risk | -0.1 (0.3) | | 0.1 (0.4) | |
| | 120-139 BP*NH-Black, APOL1 high-risk | -0.2 (0.4) | | 0.5 (0.4) | |
| | ≥140 BP*NH-Black, APOL1 high-risk | 1.4 (0.6) | | -1.4 (0.6) | |
| | 120-139 BP*Hispanic | -0.3 (0.4) | | 0.1 (0.6) | |
| | ≥140 BP*Hispanic | 0.2 (0.6) | | 0.1 (0.7) | |
| | 120-139 BP*Other | -0.1 (0.7) | | -2.0 (1.5) | |
| | ≥140 BP*Other | 0.2 (0.8) | | 5.2 (2.6) | |
| ACE/ARB | No | Ref | .04 | - | N/A |
| | Yes | 0.2 (0.1) | | - | |
| ACE/ARB*Ra ce/ ethnicity | No*NH-White | Ref | .2 | - | N/A |
| | Yes*NH-Black, APOL1 low- risk | -0.2 (0.2) | | - | |
| | Yes*NH-Black, APOL1 high-risk | 0.3 (0.4) | | - | |
| | Yes*Hispanic | -0.3 (0.4) | | - | |
| | Yes*Other | 1.3 (0.7) | | - | |
| Body Composition Measures | | | | | |
| Fat-free mass, kg | Q1: [25.8, 49.1) | - | N/A | Ref | .6 |
| | Q2: [49.1, 59.1) | - | | -0.1 (0.3) | |
| | Q3: [59.1, 69.9) | - | | -0.3 (0.3) | |
| | Q4: [69.9, 167.1) | - | | 0.0 (0.3) | |
| | Q1 FFM*NH-White | - | N/A | Ref | <.001 |

| | | Without Diabetes | | | |
|-------------------------------------|---|--|---------|---------------------------------------|---------|
| | | eGFR Slope, mL/min/1.73m ² /year | | KRT/eGFR Halving | |
| | | Beta Coefficient ^a (SE) | P value | Beta Coefficient ^b (SE) | P value |
| Fat-free mass*Race/ ethnicity | Q2 FFM*NH-Black, APOL1 low-risk | - | | -0.5 (0.4) | |
| | Q3 FFM*NH-Black, APOL1 low-risk | - | | 0.4 (0.5) | |
| | Q4 FFM*NH-Black, APOL1 low-risk | - | | 0.0 (0.5) | |
| | Q2 FFM*NH-Black, APOL1 high-risk | - | | -1.4 (0.6) | |
| | Q3 FFM*NH-Black, APOL1 high-risk | - | | -2.5 (0.7) | |
| | Q4 FFM*NH-Black, APOL1 high-risk | - | | -0.5 (0.6) | |
| | Q2 FFM*Hispanic | - | | -1.4 (1.0) | |
| | Q3 FFM*Hispanic | - | | -1.0 (0.9) | |
| | Q4 FFM*Hispanic | - | | 0.0 (1.0) | |
| | Q2 FFM*Other | - | | 1.8 (2.4) | |
| | Q3 FFM*Other | - | | 8.7 (3.6) | |
| | Q4 FFM*Other | - | | 2.5 (2.4) | |
| Measure of Anemia | | | | | |
| Hemoglobin, mg/dL | M: <12, F: <11 | - | N/A | 0.2 (0.3) | .3 |
| | M: 12-13.9, F: 11-12.9 | - | | Ref | |
| | M: ≥14, F: ≥13 | - | | -0.3 (0.2) | |
| Hemoglobin* Race/ ethnicity | M: 12-13.9, F: 11-12.9 Hemoglobin*NH-White | - | N/A | Ref | .1 |
| | M: <12, F: <11 Hemoglobin*NH-Black, APOL1 low-risk | - | | -0.1 (0.4) | |
| | M: ≥14, F: ≥13 Hemoglobin*NH-Black, APOL1 low-risk | - | | -0.3 (0.4) | |
| | M: <12, F: <11 Hemoglobin*NH-Black, APOL1 high-risk | - | | -0.3 (0.5) | |
| | M: ≥14, F: ≥13 Hemoglobin*NH-Black, APOL1 high-risk | - | | -1.1 (0.5) | |

| | Without Diabetes | | | | |
|--|---|---|---------------------------------------|---|----|
| | eGFR Slope, mL/min/1.73m ² /year | | KRT/eGFR Halving | | |
| | Beta Coefficient ^a (SE) | P value | Beta Coefficient ^b (SE) | P value | |
| M: <12, F: <11 Hemoglobin*Hispanic | - | | -0.4 (0.6) | | |
| M: ≥14, F: ≥13 Hemoglobin*Hispanic | - | | 0.0 (0.7) | | |
| M: <12, F: <11 Hemoglobin*Other | - | | 4.7 (1.7) | | |
| M: ≥14, F: ≥13 Hemoglobin*Other | - | | 0.5 (0.9) | | |
| Mineral Metabolism Markers | | | | | |
| Serum phosphate, mg/dL | Q1: [1.7, 3.3) Q2: [3.3, 3.7) Q3: [3.7, 4.1) Q4: [4.1, 9.3] | -0.1 (0.1) Ref 0.2 (0.2) -0.3 (0.2) | .2 | - - - | |
| Serum phosphate* Race/ ethnicity | Q2 phosphate*NH-White Q1 phosphate*NH-Black, APOL1 low-risk Q3 phosphate*NH-Black, APOL1 low-risk Q4 phosphate*NH-Black, APOL1 low-risk Q1 phosphate*NH-Black, APOL1 high-risk Q3 phosphate*NH-Black, APOL1 high-risk Q4 phosphate*NH-Black, APOL1 high-risk Q1 phosphate*Hispanic Q3 phosphate*Hispanic Q4 phosphate*Hispanic Q1 phosphate*Other Q3 phosphate*Other Q4 phosphate*Other | Ref -0.6 (0.3) -0.3 (0.3) 0.2 (0.3) -1.4 (0.5) -1.3 (0.5) -0.5 (0.6) -0.7 (0.5) -0.4 (0.5) 0.4 (0.6) 0.8 (0.7) -0.2 (0.7) 0.2 (0.8) | .06 | - - - - - - - - - - - - - | |
| Cardiac Markers | | | | | |
| | Q1: [1.5, 5.5) | - | N/A | Ref | .2 |

| | | Without Diabetes | | | |
|---------------------------------|---|--|---------|---------------------------------------|---------|
| | | eGFR Slope, mL/min/1.73m ² /year | | KRT/eGFR Halving | |
| | | Beta Coefficient ^a (SE) | P value | Beta Coefficient ^b (SE) | P value |
| hsTnT, pg/mL | Q2: [5.5, 11.4) | - | | 0.3 (0.2) | |
| | Q3: (11.4, 22.2) | - | | -0.2 (0.3) | |
| | Q4: (22.2, 738.7) | - | | -0.3 (0.3) | |
| hsTnT*Race/ ethnicity | Q1 hsTNT*NH-White | - | N/A | Ref | .3 |
| | Q2 hsTNT*NH-Black, APOL1 low-risk | - | | 0.0 (0.4) | |
| | Q3 hsTNT*NH-Black, APOL1 low-risk | - | | 0.3 (0.4) | |
| | Q4 hsTNT*NH-Black, APOL1 low-risk | - | | 0.3 (0.5) | |
| | Q2 hsTNT*NH-Black, APOL1 high-risk | - | | 0.2 (0.5) | |
| | Q3 hsTNT*NH-Black, APOL1 high-risk | - | | -1.2 (0.7) | |
| | Q4 hsTNT*NH-Black, APOL1 high-risk | - | | 0.4 (0.7) | |
| | Q2 hsTNT*Hispanic | - | | -0.1 (0.6) | |
| | Q3 hsTNT*Hispanic | - | | -0.2 (0.6) | |
| | Q4 hsTNT*Hispanic | - | | -0.3 (0.8) | |
| | Q2 hsTNT*Other | - | | 3.9 (1.8) | |
| | Q3 hsTNT*Other | - | | -10.0 (4.5) | |
| | Q4 hsTNT*Other | - | | -4.3 (2.9) | |
| NTproBNP, pg/mL | Q1: [2.5, 60.2) | Ref | .08 | Ref | .03 |
| | Q2: [60.2, 139.4) | -0.3 (0.1) | | 0.7 (0.3) | |
| | Q3: [139.4, 366.6) | -0.2 (0.2) | | 0.7 (0.3) | |
| | Q4: [366.6, 33 742] | -0.3 (0.2) | | 0.8 (0.3) | |
| NTproBNP* Race/ ethnicity | Q1 NTproBNP*NH-White | Ref | .2 | Ref | .004 |
| | Q2 NTproBNP*NH-Black, APOL1 low-risk | 0.7 (0.3) | | -0.5 (0.4) | |
| | Q3 NTproBNP*NH-Black, APOL1 low-risk | -0.2 (0.3) | | -0.5 (0.4) | |
| | Q4 NTproBNP*NH-Black, APOL1 low-risk | 0.0 (0.3) | | -0.2 (0.5) | |

| | Without Diabetes | | | |
|---|--|---------|---------------------------------------|---------|
| | eGFR Slope, mL/min/1.73m ² /year | | KRT/eGFR Halving | |
| | Beta Coefficient ^a (SE) | P value | Beta Coefficient ^b (SE) | P value |
| Q2 NTproBNP*NH-Black, APOL1 high-risk | 0.1 (0.5) | | -2.0 (0.6) | |
| | -0.2 (0.5) | | -0.8 (0.6) | |
| | -0.2 (0.6) | | -2.2 (0.7) | |
| | -0.3 (0.6) | | 0.8 (0.8) | |
| | -0.8 (0.6) | | 0.8 (0.9) | |
| | 0.1 (0.7) | | 0.6 (1.0) | |
| | 0.1 (0.7) | | -3.8 (1.5) | |
| | -0.7 (0.9) | | 5.1 (2.4) | |
| | 0.6 (1.3) | | -6.8 (263.7) | |
| Acidosis Marker | | | | |
| Serum bicarbonate, mmol/L | ≤22 | N/A | 0.5 (0.3) | .1 |
| | (22, 24] | | Ref | |
| | (24, 26] | | 0.0 (0.3) | |
| | >26 | | 0.1 (0.3) | |
| Serum bicarbonate* Race/ ethnicity | (22, 24] bicarbonate*NH- White | N/A | Ref | .08 |
| | ≤22 bicarbonate*NH-Black, APOL1 low-risk | | -0.1 (0.4) | |
| | (24, 26] bicarbonate*NH- Black, APOL1 low-risk | | 0.0 (0.4) | |
| | >26 bicarbonate*NH-Black, APOL1 low-risk | | -0.2 (0.4) | |
| | ≤22 bicarbonate*NH-Black, APOL1 high-risk | | -1.5 (0.5) | |
| | (24, 26] bicarbonate*NH- Black, APOL1 high-risk | | -0.3 (0.5) | |
| | >26 bicarbonate*NH-Black, APOL1 high-risk | | 0.1 (0.6) | |
| | ≤22 bicarbonate*Hispanic | | -0.4 (0.6) | |
| | (24, 26] bicarbonate*Hispanic | | -1.0 (0.8) | |
| | >26 bicarbonate*Hispanic | | -0.5 (0.9) | |

| | Without Diabetes | | | |
|--|--|------------|---------------------------------------|------------|
| | eGFR Slope, mL/min/1.73m ² /year | | KRT/eGFR Halving | |
| | Beta Coefficient ^a (SE) | P value | Beta Coefficient ^b (SE) | P value |
| ≤22 bicarbonate*Other | - | | 5.9 (2.6) | |
| | - | | 4.8 (2.4) | |
| | - | | 0.0 (1.4) | |
| Urinary Electrolytes | | | | |
| Urine potassium, mmol/24h | Q1: [3.0, 37.5) | Ref | .5 | - |
| | Q2: [37.5, 51.8) | 0.2 (0.2) | | - |
| | Q3: [51.8, 69.4) | -0.1 (0.2) | | - |
| | Q4: [69.4, 417.7] | 0.1 (0.2) | | - |
| Urine potassium* Race/ ethnicity | Q1 potassium*NH-White | Ref | .05 | - |
| | Q2 potassium*NH-Black, APOL1 low-risk | -0.7 (0.3) | | - |
| | Q3 potassium*NH-Black, APOL1 low-risk | 0.1 (0.3) | | - |
| | Q4 potassium*NH-Black, APOL1 low-risk | -0.2 (0.4) | | - |
| | Q2 potassium*NH-Black, APOL1 high-risk | 0.3 (0.5) | | - |
| | Q3 potassium*NH-Black, APOL1 high-risk | 0.2 (0.5) | | - |
| | Q4 potassium*NH-Black, APOL1 high-risk | -0.3 (0.5) | | - |
| | Q2 potassium*Hispanic | 0.1 (0.6) | | - |
| | Q3 potassium*Hispanic | 0.7 (0.6) | | - |
| | Q4 potassium*Hispanic | 1.0 (0.5) | | - |
| | Q2 potassium*Other | -0.6 (1.3) | | - |
| | Q3 potassium*Other | -1.8 (1.3) | | - |
| | Q4 potassium*Other | -1.0 (1.3) | | - |
| Kidney Injury Marker | | | | |
| Urine NGAL, ng/ml | Q1: [0.4, 6.3) | Ref | .1 | Ref |
| | Q2: [6.3, 14.2) | -0.2 (0.1) | | 0.4 (0.3) |
| | Q3: [14.2, 32.9) | 0.0 (0.2) | | -0.1 (0.3) |
| | Q4: [32.9, 2743.8] | -0.1 (0.2) | | 0.5 (0.3) |
| | Q1 NGAL*NH-White | Ref | .5 | Ref |
| | | | | .2 |

| | | Without Diabetes | | | |
|---|---|--|---------|---------------------------------------|---------|
| | | eGFR Slope, mL/min/1.73m ² /year | | KRT/eGFR Halving | |
| | | Beta Coefficient ^a (SE) | P value | Beta Coefficient ^b (SE) | P value |
| Urine NGAL*Race/ ethnicity | Q2 NGAL*NH-Black, APOL1 low-risk | -0.3 (0.3) | .6 | 0.2 (0.4) | N/A |
| | Q3 NGAL*NH-Black, APOL1 low-risk | -0.5 (0.3) | | 0.5 (0.5) | |
| | Q4 NGAL*NH-Black, APOL1 low-risk | -0.5 (0.3) | | 0.0 (0.5) | |
| | Q2 NGAL*NH-Black, APOL1 high-risk | 0.6 (0.6) | | 1.6 (0.9) | |
| | Q3 NGAL*NH-Black, APOL1 high-risk | -0.4 (0.6) | | 1.7 (1.0) | |
| | Q4 NGAL*NH-Black, APOL1 high-risk | -0.5 (0.6) | | 1.8 (1.0) | |
| | Q2 NGAL*Hispanic | -0.6 (0.6) | | 0.9 (1.1) | |
| | Q3 NGAL*Hispanic | -1.1 (0.6) | | 1.3 (1.0) | |
| | Q4 NGAL*Hispanic | -0.7 (0.7) | | 1.1 (1.0) | |
| | Q2 NGAL*Other | 0.0 (0.8) | | -6.3 (2.1) | |
| | Q3 NGAL*Other | -0.1 (0.8) | | -0.4 (1.8) | |
| | Q4 NGAL*Other | -0.7 (1.0) | | -2.5 (1.8) | |
| RAAS Marker | | | | | |
| Serum aldosterone, pg/mL | Q1: [0.8, 71.2) | Ref | .6 | - | N/A |
| | Q2: [71.2, 101.4) | -0.1 (0.2) | | - | |
| | Q3: [101.4, 152.8) | -0.3 (0.2) | | - | |
| | Q4: [152.8, 15 630.9] | 0.0 (0.2) | | - | |
| Serum aldosterone* Race/ ethnicity | Q1 aldosterone*NH-White | Ref | .3 | - | N/A |
| | Q2 aldosterone*NH-Black, APOL1 low-risk | -0.3 (0.3) | | - | |
| | Q3 aldosterone*NH-Black, APOL1 low-risk | 0.2 (0.3) | | - | |
| | Q4 aldosterone*NH-Black, APOL1 low-risk | 0.3 (0.3) | | - | |
| | Q2 aldosterone*NH-Black, APOL1 high-risk | 0.5 (0.6) | | - | |
| | Q3 aldosterone*NH-Black, APOL1 high-risk | 0.9 (0.6) | | - | |

| | Without Diabetes | | | |
|---|--|---------|---------------------------------------|---------|
| | eGFR Slope, mL/min/1.73m ² /year | | KRT/eGFR Halving | |
| | Beta Coefficient ^a (SE) | P value | Beta Coefficient ^b (SE) | P value |
| Q4 aldosterone*NH-Black, APOL1 high-risk | 0.5 (0.5) | | - | |
| | -1.3 (0.5) | | - | |
| | -0.6 (0.5) | | - | |
| | -0.4 (0.7) | | - | |
| | 0.7 (0.8) | | - | |
| | 0.6 (0.9) | | - | |
| | 0.1 (0.8) | | - | |

Abbreviations: ACE: angiotensin converting enzyme; ARB: angiotensin receptor blocker; BP: blood pressure; CKD: chronic kidney disease; eGFR: estimated glomerular filtration rate; KRT: kidney replacement therapy; FFM: fat-free mass; hsTnT: high-sensitivity troponin T; NGAL: neutrophil gelatinase-associated lipocalin; NH: non-Hispanic; NTproBNP: N-terminal pro-B-type natriuretic peptide; RAAS: renin-angiotensin-aldosterone system; Ref: reference; SE: standard error; UACR: urine albumin:creatinine ratio

N/A indicates variable was excluded from analysis for indicated outcome as only variables selected by models described in Table 2 were tested for interaction.

SI conversion: To convert Hemoglobin to g/L, multiply by 0.01. To convert serum phosphate to mmol/L, multiply by 0.323. For serum bicarbonate, 1 mEq/L is equivalent to 1 mmol/L. For urine potassium, 1 mEq/24h is equivalent to 1 mmol/d. To convert serum aldosterone to pmol/L, multiply by 2.774.

^a Negative coefficients represent a faster decline in eGFR compared to the reference group.

^b Coefficients are the log hazard ratio to the reference level with positive coefficients indicating increased risk of KRT/eGFR Halving compared to the reference group.

* Indicates interaction between variables. Sum of main and interaction beta coefficients will provide estimate for given group.

Table S4b. Multivariable-adjusted models of chronic kidney disease progression with race interaction among CRIC participants with diabetes

| | | With Diabetes | | | |
|----------------------------|--|--|---------|---------------------------------------|---------|
| | | eGFR Slope, mL/min/1.73m ² /year | | KRT/eGFR Halving | |
| | | Beta Coefficient ^a (SE) | P value | Beta Coefficient ^b (SE) | P value |
| Demographic Factors | | | | | |
| Age, years | <45 | -0.5 (0.3) | .5 | 0.0 (0.3) | .03 |
| | 45-64 | Ref | | Ref | |
| | 65-75 | 0.0 (0.2) | | -0.5 (0.2) | |
| Race/ ethnicity | NH-White | Ref | .02 | Ref | .9 |
| | NH-Black, APOL1 low-risk | 0.1 (0.6) | | -0.3 (0.9) | |
| | NH-Black, APOL1 high-risk | -0.5 (1.4) | | 0.2 (1.7) | |
| | Hispanic | -0.8 (0.8) | | -1.0 (1.5) | |
| | Other | 0.5 (1.4) | | -0.1 (2.6) | |
| Age*Race/ ethnicity | 45-64 age*NH-White | Ref | .7 | Ref | .2 |
| | <45 age*NH-Black, APOL1 low-risk | -0.7 (0.6) | | 0.5 (0.4) | |
| | 65-75 age*NH-Black, APOL1 low-risk | 0.0 (0.3) | | 0.3 (0.3) | |
| | <45 age*NH-Black, APOL1 high-risk | 0.6 (1.1) | | -0.9 (0.7) | |
| | 65-75 age*NH-Black, APOL1 high-risk | -0.6 (0.7) | | 0.7 (0.5) | |
| | <45 age*Hispanic | 0.2 (0.7) | | 0.7 (0.4) | |
| | 65-75 age*Hispanic | 0.5 (0.4) | | 0.1 (0.3) | |
| | <45 age*Other | 0.1 (1.6) | | 1.3 (1.3) | |
| | 65-75 age*Other | -0.6 (0.8) | | 1.1 (0.9) | |
| Gender | Male | - | N/A | Ref | .2 |
| | Female | - | | -0.3 (0.2) | |
| Gender*Race/ ethnicity | Male*NH-White | - | N/A | Ref | .02 |
| | Female*NH-Black, APOL1 low-risk | - | | -0.1 (0.3) | |
| | Female*NH-Black, APOL1 high-risk | - | | -0.2 (0.5) | |
| | Female*Hispanic | - | | -0.2 (0.3) | |
| | Female*Other | - | | -3.6 (1.1) | |

| | | With Diabetes | | | |
|---|---|--|---------|---------------------------------------|---------|
| | | eGFR Slope, mL/min/1.73m ² /year | | KRT/eGFR Halving | |
| | | Beta Coefficient ^a (SE) | P value | Beta Coefficient ^b (SE) | P value |
| Kidney Function Measures | | | | | |
| Baseline eGFR, mL/min/ 1.73m ² | <30 | 0.2 (0.3) | <.001 | 0.7 (0.2) | <.001 |
| | 30-44.9 | Ref | | Ref | |
| | 45-59.9 | -0.4 (0.2) | | -0.3 (0.2) | |
| | ≥60 | -0.6 (0.3) | | -0.9 (0.4) | |
| Baseline eGFR*Race/ ethnicity | 30-44.9 eGFR*NH-White | Ref | .7 | Ref | .2 |
| | <30 eGFR*NH-Black, APOL1 low-risk | 0.6 (0.4) | | -0.4 (0.3) | |
| | 45-59.9 eGFR*NH-Black, APOL1 low-risk | 0.0 (0.3) | | 0.1 (0.3) | |
| | ≥60 eGFR*NH-Black, APOL1 low-risk | -0.1 (0.5) | | 1.0 (0.5) | |
| | <30 eGFR*NH-Black, APOL1 high-risk | 1.5 (0.9) | | -1.1 (0.5) | |
| | 45-59.9 eGFR*NH-Black, APOL1 high-risk | 0.2 (0.7) | | 0.0 (0.5) | |
| | ≥60 eGFR*NH-Black, APOL1 high-risk | -0.7 (1.1) | | 1.3 (0.9) | |
| | <30 eGFR*Hispanic | 0.7 (0.5) | | -0.2 (0.3) | |
| | 45-59.9 eGFR*Hispanic | 0.3 (0.4) | | -0.2 (0.4) | |
| | ≥60 eGFR*Hispanic | -0.3 (0.8) | | 0.5 (0.7) | |
| | <30 eGFR*Other | 0.5 (1.1) | | 1.2 (1.2) | |
| | 45-59.9 eGFR*Other | 0.1 (0.8) | | 0.0 (1.0) | |
| | ≥60 eGFR*Other | -1.9 (1.3) | | 1.0 (1.4) | |
| UACR, mg/g | <30 | Ref | <.001 | Ref | <.001 |
| | 30-299 | -0.7 (0.2) | | 0.7 (0.2) | |
| | ≥300 | -2.0 (0.3) | | 1.6 (0.2) | |
| UACR*Race/ ethnicity | <30 UACR*NH-White | Ref | .04 | Ref | .3 |
| | 30-299 UACR*NH-Black, APOL1 low-risk | -0.7 (0.3) | | 0.6 (0.4) | |
| | ≥300 UACR*NH-Black, APOL1 low-risk | -1.0 (0.4) | | 0.7 (0.4) | |
| | 30-299 UACR*NH-Black, APOL1 high-risk | -0.9 (0.7) | | 0.6 (0.7) | |

| | With Diabetes | | | |
|--|--|------------|---------------------------------------|------------|
| | eGFR Slope, mL/min/1.73m ² /year | | KRT/eGFR Halving | |
| | Beta Coefficient ^a (SE) | P value | Beta Coefficient ^b (SE) | P value |
| ≥300 UACR*NH-Black, APOL1 high-risk | -1.2 (0.8) | | 0.2 (0.7) | |
| 30-299 UACR*Hispanic | -1.4 (0.5) | | 0.9 (0.5) | |
| ≥300 UACR*Hispanic | -0.7 (0.5) | | 0.4 (0.5) | |
| 30-299 UACR*Other | 0.1 (0.9) | | -2.5 (1.9) | |
| ≥300 UACR*Other | 0.5 (0.9) | | -2.1 (1.6) | |
| Blood Pressure Factors | | | | |
| Systolic BP, mmHg | <120 | Ref | .006 | Ref |
| | 120-139 | -0.1 (0.2) | | 0.2 (0.2) |
| | ≥140 | -0.6 (0.3) | | 0.8 (0.2) |
| Systolic BP*Race/ ethnicity | <120 BP*NH-White | Ref | .6 | Ref |
| | 120-139 BP*NH-Black, APOL1 low-risk | -0.3 (0.3) | | 0.0 (0.3) |
| | ≥140 BP*NH-Black, APOL1 low-risk | 0.0 (0.4) | | -0.5 (0.3) |
| | 120-139 BP*NH-Black, APOL1 high-risk | 0.7 (0.8) | | -0.5 (0.6) |
| | ≥140 BP*NH-Black, APOL1 high-risk | -0.6 (0.8) | | -0.4 (0.6) |
| | 120-139 BP*Hispanic | 0.2 (0.5) | | -0.1 (0.4) |
| | ≥140 BP*Hispanic | 0.2 (0.5) | | -0.2 (0.4) |
| | 120-139 BP*Other | -0.6 (0.9) | | 1.2 (1.4) |
| | ≥140 BP*Other | -0.9 (1.2) | | 0.7 (1.6) |
| Behavioral Factor | | | | |
| Current smoker | No | Ref | .2 | - |
| | Yes | -0.6 (0.3) | | - |
| Current smoker*Race/ ethnicity | No*NH-White | Ref | .2 | - |
| | Yes*NH-Black, APOL1 low-risk | 0.4 (0.4) | | - |
| | Yes*NH-Black, APOL1 high-risk | -1.0 (0.8) | | - |
| | Yes*Hispanic | 1.4 (0.8) | | - |
| | Yes*Other | 0.0 (1.6) | | - |

| | | With Diabetes | | | |
|---|---------------------------------------|--|---------|---------------------------------------|---------|
| | | eGFR Slope, mL/min/1.73m ² /year | | KRT/eGFR Halving | |
| | | Beta Coefficient ^a (SE) | P value | Beta Coefficient ^b (SE) | P value |
| Inflammatory Marker and Chemokines | | | | | |
| Serum fractalkine (CX3CL1), pg/mL | Q1: [0.1, 0.6) | Ref | .007 | Ref | .9 |
| | Q2: [0.6, 0.8) | -0.1 (0.3) | | 0.0 (0.3) | |
| | Q3: [0.8, 1.1) | -0.1 (0.3) | | 0.1 (0.3) | |
| | Q4: [1.1, 3.6] | -0.2 (0.3) | | 0.1 (0.3) | |
| Serum fractalkine (CX3CL1)* Race/ ethnicity | Q1 CXCL1*NH-White | Ref | .05 | Ref | .2 |
| | Q2 CXCL1*NH-Black, APOL1 low-risk | -0.2 (0.4) | | 0.4 (0.4) | |
| | Q3 CXCL1*NH-Black, APOL1 low-risk | -0.7 (0.4) | | 0.6 (0.4) | |
| | Q4 CXCL1*NH-Black, APOL1 low-risk | -0.6 (0.4) | | 0.7 (0.4) | |
| | Q2 CXCL1*NH-Black, APOL1 high-risk | 1.3 (0.9) | | -1.0 (0.7) | |
| | Q3 CXCL1*NH-Black, APOL1 high-risk | -0.5 (0.9) | | 0.3 (0.7) | |
| | Q4 CXCL1*NH-Black, APOL1 high-risk | 1.9 (0.9) | | -0.3 (0.7) | |
| | Q2 CXCL1*Hispanic | 0.7 (0.6) | | -0.2 (0.5) | |
| | Q3 CXCL1*Hispanic | -0.1 (0.6) | | 0.2 (0.5) | |
| | Q4 CXCL1*Hispanic | -0.1 (0.6) | | 0.1 (0.5) | |
| | Q2 CXCL1*Other | 1.2 (1.0) | | 0.9 (1.2) | |
| | Q3 CXCL1*Other | 0.0 (1.1) | | -1.8 (1.2) | |
| | Q4 CXCL1*Other | -0.4 (1.2) | | 1.5 (1.5) | |
| Plasma CXCL12, pg/mL | Q1: [832.1, 2066.4) | - | N/A | Ref | .7 |
| | Q2: [2066.4, 2410.5) | - | | 0.3 (0.3) | |
| | Q3: [2410.5, 2797.8) | - | | 0.1 (0.3) | |
| | Q4: [2797.8, 6173.3] | - | | 0.2 (0.3) | |
| Plasma CXCL12* Race/ ethnicity | Q1 CXCL12*NH-White | - | N/A | Ref | .1 |
| | Q2 CXCL12*NH-Black, APOL1 low-risk | - | | -0.2 (0.4) | |
| | Q3 CXCL12*NH-Black, APOL1 low-risk | - | | 0.6 (0.3) | |

| | With Diabetes | | | | |
|--|--|---------|---------------------------------------|------------|----|
| | eGFR Slope, mL/min/1.73m ² /year | | KRT/eGFR Halving | | |
| | Beta Coefficient ^a (SE) | P value | Beta Coefficient ^b (SE) | P value | |
| Q4 CXCL12*NH-Black, APOL1 low-risk | - | | 0.5 (0.3) | | |
| Q2 CXCL12*NH-Black, APOL1 high-risk | - | | -0.2 (0.7) | | |
| Q3 CXCL12*NH-Black, APOL1 high-risk | - | | 0.4 (0.7) | | |
| Q4 CXCL12*NH-Black, APOL1 high-risk | - | | -0.1 (0.7) | | |
| Q2 CXCL12*Hispanic | - | | -0.6 (0.5) | | |
| Q3 CXCL12*Hispanic | - | | 0.4 (0.5) | | |
| Q4 CXCL12*Hispanic | - | | 0.0 (0.4) | | |
| Q2 CXCL12*Other | - | | 0.5 (1.0) | | |
| Q3 CXCL12*Other | - | | -1.1 (1.2) | | |
| Q4 CXCL12*Other | - | | -2.9 (1.4) | | |
| Carbohydrate Metabolism Markers | | | | | |
| HbA1c, % | Q1: [3.5, 5.6) | - | N/A | Ref | .4 |
| | Q2: [5.6, 6.2) | - | | 0.6 (0.5) | |
| | Q3: [6.2, 7.3) | - | | 0.2 (0.5) | |
| | Q4: [7.3, 15.2] | - | | 0.3 (0.5) | |
| HbA1c*Race/ ethnicity | Q1 HbA1c*NH-White | - | N/A | Ref | .6 |
| | Q2 HbA1c*NH-Black, APOL1 low-risk | - | | -0.7 (0.6) | |
| | Q3 HbA1c*NH-Black, APOL1 low-risk | - | | -0.3 (0.6) | |
| | Q4 HbA1c*NH-Black, APOL1 low-risk | - | | -0.5 (0.6) | |
| | Q2 HbA1c*NH-Black, APOL1 high-risk | - | | 1.3 (1.0) | |
| | Q3 HbA1c*NH-Black, APOL1 high-risk | - | | 0.0 (0.9) | |
| | Q4 HbA1c*NH-Black, APOL1 high-risk | - | | 0.5 (0.9) | |
| | Q2 HbA1c*Hispanic | - | | 0.3 (0.8) | |
| | Q3 HbA1c*Hispanic | - | | 0.1 (0.8) | |

| | With Diabetes | | | |
|---------------------------------|--|------------|---------------------------------------|------------|
| | eGFR Slope, mL/min/1.73m ² /year | | KRT/eGFR Halving | |
| | Beta Coefficient ^a (SE) | P value | Beta Coefficient ^b (SE) | P value |
| Q4 HbA1c*Hispanic | - | | 0.1 (0.8) | |
| | - | | 1.3 (1.9) | |
| | - | | 0.6 (1.5) | |
| | - | | 1.4 (1.5) | |
| Cardiac Markers | | | | |
| NTproBNP, pg/mL | Q1: [2.5, 60.2) | Ref | .03 | Ref |
| | Q2: [60.2, 139.4) | -0.4 (0.3) | | 0.5 (0.3) |
| | Q3: [139.4, 366.6) | -0.5 (0.3) | | 0.5 (0.3) |
| | Q4: [366.6, 33 742] | -0.7 (0.3) | | 0.6 (0.3) |
| NTproBNP* Race/ ethnicity | Q1 NTproBNP*NH-White | Ref | .9 | Ref |
| | Q2 NTproBNP*NH-Black, APOL1 low-risk | 0.0 (0.4) | | 0.0 (0.4) |
| | Q3 NTproBNP*NH-Black, APOL1 low-risk | 0.1 (0.4) | | 0.1 (0.4) |
| | Q4 NTproBNP*NH-Black, APOL1 low-risk | 0.1 (0.4) | | 0.3 (0.4) |
| | Q2 NTproBNP*NH-Black, APOL1 high-risk | -0.1 (0.8) | | -0.4 (0.7) |
| | Q3 NTproBNP*NH-Black, APOL1 high-risk | -0.2 (1.0) | | 0.0 (0.8) |
| | Q4 NTproBNP*NH-Black, APOL1 high-risk | 0.1 (1.0) | | 0.0 (0.8) |
| | Q2 NTproBNP*Hispanic | -0.2 (0.6) | | -0.1 (0.5) |
| | Q3 NTproBNP*Hispanic | 0.7 (0.6) | | -0.9 (0.5) |
| | Q4 NTproBNP*Hispanic | 0.1 (0.6) | | -0.3 (0.5) |
| | Q2 NTproBNP*Other | -0.9 (1.0) | | -2.3 (1.7) |
| | Q3 NTproBNP*Other | -0.4 (1.1) | | -1.6 (1.4) |
| | Q4 NTproBNP*Other | -0.7 (1.1) | | -2.9 (1.7) |
| Kidney Injury Marker | | | | |
| Urine NGAL, ng/ml | Q1: [0.4, 6.3) | Ref | .03 | Ref |
| | Q2: [6.3, 14.2) | -0.3 (0.2) | | 0.1 (0.2) |
| | Q3: [14.2, 32.9) | -0.7 (0.3) | | 0.4 (0.2) |
| | Q4: [32.9, 2743.8] | -0.2 (0.3) | | 0.8 (0.3) |

| | | With Diabetes | | | |
|-----------------------------------|--------------------------------------|--|---------|---------------------------------------|---------|
| | | eGFR Slope, mL/min/1.73m ² /year | | KRT/eGFR Halving | |
| | | Beta Coefficient ^a (SE) | P value | Beta Coefficient ^b (SE) | P value |
| Urine NGAL* Race/ ethnicity | Q1 NGAL*NH-White | Ref | .6 | Ref | .2 |
| | Q2 NGAL*NH-Black, APOL1 low-risk | 0.1 (0.4) | | -0.1 (0.3) | |
| | Q3 NGAL*NH-Black, APOL1 low-risk | 0.5 (0.4) | | -0.5 (0.3) | |
| | Q4 NGAL*NH-Black, APOL1 low-risk | -0.3 (0.4) | | -0.3 (0.4) | |
| | Q2 NGAL*NH-Black, APOL1 high-risk | -0.6 (0.9) | | 0.6 (0.8) | |
| | Q3 NGAL*NH-Black, APOL1 high-risk | -0.4 (0.9) | | 0.8 (0.8) | |
| | Q4 NGAL*NH-Black, APOL1 high-risk | -1.8 (1.0) | | 1.5 (0.9) | |
| | Q2 NGAL*Hispanic | 0.4 (0.6) | | -0.2 (0.6) | |
| | Q3 NGAL*Hispanic | 0.0 (0.5) | | -0.1 (0.5) | |
| | Q4 NGAL*Hispanic | -0.7 (0.6) | | 0.3 (0.5) | |
| | Q2 NGAL*Other | 0.4 (0.9) | | -0.1 (1.6) | |
| | Q3 NGAL*Other | 0.8 (1.0) | | 3.6 (1.6) | |
| | Q4 NGAL*Other | -1.0 (1.1) | | 3.5 (1.5) | |

Abbreviations: BP: blood pressure; CKD: chronic kidney disease; eGFR: estimated glomerular filtration rate; KRT: kidney replacement therapy; HbA1c: glycosylated hemoglobin; NGAL: neutrophil gelatinase-associated lipocalin; NH: non-Hispanic; NTproBNP: N-terminal pro-B-type natriuretic peptide; Ref: reference; SE: standard error; UACR: urine albumin:creatinine ratio

N/A indicates variable was excluded from analysis for indicated outcome as only variables selected by models described in Table 3 were tested for interaction.

SI conversion: To convert Uric acid to $\mu\text{mol/L}$, multiply by 59.485. For serum bicarbonate, 1 mEq/L is equivalent to 1 mmol/L.

^a Negative coefficients represent a faster decline in eGFR compared to the reference group.

^b Coefficients are the log hazard ratio to the reference level with positive coefficients indicating increased risk of KRT/eGFR Halving compared to the reference group.

* Indicates interaction between variables. Sum of main and interaction beta coefficients will provide estimate for given group.

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