Relationship of Race and Proteinuria to Renal Function Decline in Patients with CKD

Steven Rosansky1, Martin Durkin2, James Hardin3, Kirby Jackson3, Csaba Kovessy4, Jessica Sontrop5, Justin Reynolds1, Kathryn Haddock1, Frankie Richards1 and William Clark6

1WJBDA Hospital, Dorn Research Institute, University of South Carolina School of Public Health, Columbia, SC, USA
2Palmietto Health Hospital, Research Service, Columbia, SC, USA
3University of South Carolina School of Public Health, Columbia, SC, USA
4Memphis VA Medical Center, Nephrology, Memphis, TN, India
5Division of Nephrology London Health Sciences Centre London, Ontario, UK

Corresponding author: Steven Rosansky, WJBVA Hospital, Dorn Research Institute, University of South Carolina School of Public Health, Columbia, SC, USA, 526 N Trenholm Rd, Columbia SC 29206, USA, Tel: 8034225427; E-mail: sjrcra@yahoo.com

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Abstract

Background: To date, the reasons for the higher rates of treated end-stage renal failure in blacks versus whites are poorly understood. Proteinuria is the most important determinant of renal function loss (RFL). Blacks have a higher incidence of proteinuric CKD than whites. The current study evaluates the hypothesis that after adjustment for proteinuria, blacks have faster RFL, more prominent at lower levels of estimated glomerular filtration rates (eGFR).

Methods: In a non-referral outpatient CKD population of 1,935 blacks and 6,286 whites, the relationship of zero, <2 plus and ≥2 plus dipstick proteinuria to MDRD e GFR change per year (RFL) was analyzed. Next the relationship between race and RFL was examined in patients with higher versus lower eGFR (defined by never or ever having an eGFR <30 ml/min/1.73 m², respectively) during the study using a mixed effects model which includes longitudinal urinalysis (log converted), serum creatinine data points, age and whether a patient died during the study. Results: Versus whites, blacks had higher baseline eGFR (75.3 ml/min/1.73 m² versus 64.9 ml/min/1.73 m²), higher frequency of e GFR <30 ml/min/1.73 m² (30.8% versus 21%), higher dipstick proteinuria levels and faster RFL by proteinuria group (range -1.07 to -2.28 ml/min/m²/year in blacks and -0.68 to -1.80 ml/min/1.73 m²/year, in whites), p<0.01. In the mixed effects model, blacks had a 0.30 and 0.59 ml/min/1.73 m²/year, faster loss of renal function in the higher and lower eGFR groups, respectively, p<0.001.

Conclusion: Blacks with CKD appear to lose renal function faster than whites. This effect may be more pronounced at lower e GFR levels.

Keywords: Proteinuria, race, CKD, eGFR, RFL

Introduction

American blacks are four times more likely to initiate dialysis than American whites [1]. One possible explanation for this is a higher rate of progression to advanced chronic kidney disease (CKD) in blacks. Proteinuria has been reported to be the single strongest predictor of renal disease progression at all levels renal function [2]. Few studies to date have examined comparative rates of renal function loss (RFL) in blacks versus whites [1]. Most published studies included: patients with (eGFR) > 60 ml/min/1.73 m²; few longitudinal data points; short follow-up (<3 years), and few include the effect of albuminuria on RFL. Recently, Ericksen, et al. reported that in the in a non-referral Veterans Administration (VA) outpatient population of 1,935 blacks and 6,286 whites with an eGFR ≤ 60 ml/min/1.73 m², followed for an average of nine years with higher or lower eGFR levels defined by never or ever having an eGFR <30 ml/min/1.73 m² during the study interval, respectively.

Concise Methods

Patients from the Dorn VA Hospital, Columbia, SC who ever had a serum creatinine of 1.3 mg/dl (chosen to find patients with eGFR < 60 ml/min/1.73 m²) or greater during the interval 1989-2003 were included in the analysis. Patients were followed until December 31, 2008. eGFR was calculated utilizing the four variable equations from the modification of diet in renal disease (MDRD) study. The same automated serum creatinine methodology is used throughout the study (Beckman Coulter Synchron System Jaffe rate method). Only outpatient serum creatinine values were used to avoid the confounding effect of illness on eGFR (i.e., acute kidney injury). In order to enter the study a patient required a minimum of: three serum creatinine values over a minimum of three years of follow-up; at least one urinalysis with dipstick proteinuria measurement, and information on age and race. Subject specific measurements: whether died while under observation, whether the patient ever had an eGFR < 30 ml/min/1.73 m², the total number of and date of each serum creatinine measure, the total number and date of each urinalysis protein measure, and race were collected. Proteinuria was classified into three groups utilizing all urinalysis data, evaluated by automated colorimetric dipstick analysis, to define the following proteinuria
groupings: group 1, no proteinuria on any dipstick tests, group 2, <2 plus proteinuria (100 mg/dl) on all tests, group 3, ≥ 2 plus (100 mg/dl) on at least one urinalysis. The date of each urinalysis and creatinine measure was recorded. The relationship of RFL to level of urinalysis proteinuria was compared by race and by proteinuria group. Statistical significance by t test, and medians and interquartile ranges were reported. Proteinuria was included in the regression analysis by taking urine protein measures closest to longitudinal serum creatinine values. Given the units in which protein was recorded, dipstick proteinuria values of 10 mg/dl (trace), 1 plus proteinuria (30 mg/dl), 2 plus proteinuria (100 mg/dl), 3 plus proteinuria (300 mg/dl) and 4 plus proteinuria, 1000 mg/dl, were log10 transformed. We added 1 to the proteinuria measure to avoid numeric problems with measurement results of zero. Chi squared analysis for categorical and t test for continuous variables was used to test for statistical significance. We compared in blacks and whites.

Time is measured in years and interactions with time taking urine protein measures closest to longitudinal serum creatinine measure was recorded. The relationship of RFL to level of proteinuria was included in the regression analysis by using a mixed-effects linear regression model (Stata), which allows for multiple observations per patient. Predictor variables for regression model includes, whether died while under observation, whether the patient ever had an eGFR < 30 ml/min/1.73 m², all dated serum creatinine measures, all log converted urine protein values matching the urine protein closest to a dated serum creatinine value, and race for each person. Time is measured in years and interactions with time in years were examined by race, and ever had eGFR < 30 ml/min/1.73 m². Patients who ever had an eGFR of < 15 ml/min/1.73 m² on all tests, group 3, ≥ 2 plus (100 mg/dl) on at least one urinalysis. The date of each urinalysis and proteinuria measure to avoid numeric problems with measurement results of zero. Chi squared analysis for categorical and t test for continuous variables was used to test for statistical significance. We estimated the rate of change in eGFR (ml/min/1.73 m² per year) using a mixed-effects linear regression model (Stata), which allows for multiple observations per patient. Predictor variables for regression model includes, whether died while under observation, whether the patient ever had an eGFR < 30 ml/min/1.73 m², all dated serum creatinine measures, all log converted urine protein values matching the urine protein closest to a dated serum creatinine value, and race for each person. Time is measured in years and interactions with time in years were examined by race, and ever had eGFR < 30 ml/min/1.73 m². Patients who ever had an eGFR of < 15 ml/min/1.73 m² were compared in blacks and whites.

Main Results

Table 1 summarizes the subject characteristics and outcomes (medians and interquartile range) for the 8,221 subjects, 23.5% of who were black. The following parameters were significantly different in blacks versus whites at the p<.001 level, age 61.2 in blacks, 65.1 in whites, follow-up 9.9 years in blacks, 8.9 years in whites, mean serum creatinine measures/urine protein measures 25/8 in blacks, 19/6 in whites, median eGFR during the first 12 months 75.3 ml/min/1.73 m² in blacks vs. 64.9 ml/min/1.73 m² in whites, patients who ever had an eGFR < 30 ml/min/1.73 m² and an eGFR< 15 ml/min/1.73 m², in blacks versus whites was 30.8% versus 21%, and 13.6% versus 6.3%, respectively. There was no significant difference in number of black versus white patients who died during the study. Table 1 also presents the change in eGFR (ml/min/1.73 m² per year), utilizing the difference between the average of the first and average of the last years eGFR divided by time in years for three urine protein groups. Blacks had a higher frequency of urine protein group 2 compared with whites (Table 1). For blacks and whites, loss of eGFR was more rapid in those with higher levels of proteinuria. Median eGFR slope in ml/min/1.73 m²/year ranged from -0.74 for those with no proteinuria to -1.96 for those with at least one urinalysis with ≥ 2+ proteinuria, p< 0.01. Unadjusted eGFR slope (ml/min/1.73 m²/year) was significantly higher in blacks versus whites overall, -1.94 vs. -1.24 and after stratification by levels of proteinuria; group 0, -1.62 vs. -0.96; group 1,-1.07 vs. -0.68; group 2, -2.28 vs. -1.8, p<.01.

Table 1: Subject Characteristics

<table>
<thead>
<tr>
<th></th>
<th>All</th>
<th>Black</th>
<th>White</th>
</tr>
</thead>
<tbody>
<tr>
<td>Population (N)</td>
<td>8221</td>
<td>1935</td>
<td>6286</td>
</tr>
<tr>
<td>Age</td>
<td>64.4 (54.8-71.0) 1</td>
<td>61.2 (49.4-68.9)</td>
<td>65.1 (56.6-71.5)</td>
</tr>
<tr>
<td>Follow-up time in years</td>
<td>9.09 (6.43-13.12)</td>
<td>9.85 (6.8-14.3)</td>
<td>8.9 (6.3-12.7)</td>
</tr>
<tr>
<td># of Creatinine Measures</td>
<td>20 (12-31)</td>
<td>25 (15-40)</td>
<td>19 (12-29)</td>
</tr>
<tr>
<td># of Protein Measures</td>
<td>6 (3-12)</td>
<td>8 (4-14)</td>
<td>6 (3-11)</td>
</tr>
<tr>
<td>Average of first 12 mos. eGFR 3</td>
<td>66.9 (57.3-77.8)</td>
<td>75.3 (62.3-88.7)</td>
<td>64.9 (56.6-74.5)</td>
</tr>
<tr>
<td>Average of last 12 mos. eGFR 3</td>
<td>56.6 (43.8-67.6)</td>
<td>59.1 (42.2-72.7)</td>
<td>56.0 (44.1-66.3)</td>
</tr>
<tr>
<td>Protein Group 0 4</td>
<td>37.30%</td>
<td>22.40%</td>
<td>41.90%</td>
</tr>
<tr>
<td>Protein Group 1 5</td>
<td>32.70%</td>
<td>31.20%</td>
<td>33.20%</td>
</tr>
<tr>
<td>Protein Group 2 6</td>
<td>32.20%</td>
<td>46.40%</td>
<td>24.90%</td>
</tr>
<tr>
<td>RFL Group 0 7</td>
<td>-0.74 (-1.7, 0.27)</td>
<td>-1.07 (-2.5, 0.17)</td>
<td>-0.68 (-1.8, 0.28)</td>
</tr>
<tr>
<td>RFL Group 1 8</td>
<td>-0.95 (-2.10, 0.06)</td>
<td>-1.13 (-2.5, 0.7)</td>
<td>-0.91 (2.0, 0.50)</td>
</tr>
<tr>
<td>RFL Group 2 9</td>
<td>-1.96 (-3.8, -0.46)</td>
<td>-2.28 (-4.1, -0.75)</td>
<td>-1.8 (-3.6, -0.35)</td>
</tr>
<tr>
<td>Died during study</td>
<td>33.30%</td>
<td>35.30%</td>
<td>32.70%</td>
</tr>
<tr>
<td>Ever eGFR&lt;30 ml/min/1.73 m², 9</td>
<td>23.20%</td>
<td>30.80%</td>
<td>21.00%</td>
</tr>
<tr>
<td>Ever eGFR&lt;15 ml/min/1.73 m², 10</td>
<td>8.00%</td>
<td>13.60%</td>
<td>6.30%</td>
</tr>
</tbody>
</table>

Table 1: Subject Characteristics

1. Median (interquartile range)
2. Black versus white characteristics for all=P <.001, except died during study=P <.04
3. Estimated GRF (eGFR) using 4 variable MDRD equation in ml/min/1.73 m²
4. Protein Group 0: all urinalysis protein results negative
5. Protein Group 1: urinalysis results trace to 1+ protein (30 mg/dL)
6. Protein Group 2: urinalysis results had at least one urinalysis result ≥ 2 plus protein (100 mg/dL)
7. Renal function loss (RFL) change in eGFR per year
8. Median (interquartile range) for RFL in ml/min/1.73 m²/years, calculated as the difference in the average eGFR of the last 12 months minus the average of the first 12 months, divided by the years of follow up.
9. Had an eGFR < 30 ml/min/1.73 m² at some point of the study
10. Had an eGFR <15 ml/min/1.73 m² at some point in the study

The random effects model (Table 2) examines the differences between blacks and whites in the change in eGFR per year. Interaction terms not included in Table 2 were all not statistically significant (blacks died, blacks died years, blacks ever 30).

Compared to a white patient for each log protein unit increase in urine proteinuria, blacks had a -1.0 ml/min/1.73 m² lower eGFR data point. The other coefficients in the model all reported as ml/min/1.73 m²/year were, years-0.17, black years -0.30, ever 30 years -1.58, black ever 30 years, -0.29. For those whites who never had an eGFR <30, the RFL was -0.17 for whites; -0.17 + 0.30 = -0.47 for blacks. For those who ever had an eGFR < 30 for whites RFL was -0.17+1.58=-1.75; RFL for blacks was -0.17+0.30+1.58+0.29=-2.35 ml/min/1.73 m²/year. Thus the RFL difference between blacks and whites was -0.30 ml/min/1.73 m²/year if never had an eGFR < 30 and -0.60 ml/min/1.73 m² if ever had eGFR < 30.

### Table 2: Mixed Effects Model of Black versus White RFL Differences

<table>
<thead>
<tr>
<th></th>
<th>Black</th>
<th>White</th>
<th>Difference</th>
<th>P Value</th>
</tr>
</thead>
<tbody>
<tr>
<td>Constant</td>
<td>104.10</td>
<td>92.80</td>
<td>11.40</td>
<td>&lt;.001</td>
</tr>
<tr>
<td>eGFR 30³</td>
<td>-7.57</td>
<td>-7.80</td>
<td>0.23</td>
<td>NS</td>
</tr>
<tr>
<td>Died⁴</td>
<td>1.43</td>
<td>2.80</td>
<td>-0.75</td>
<td>NS</td>
</tr>
<tr>
<td>Log protein⁵</td>
<td>-3.00</td>
<td>-2.00</td>
<td>-1.00</td>
<td>&lt;.001</td>
</tr>
<tr>
<td>Age⁶</td>
<td>-0.49</td>
<td>-0.49</td>
<td>0</td>
<td>NS</td>
</tr>
<tr>
<td>Male</td>
<td>5.90</td>
<td>5.90</td>
<td>0</td>
<td>NS</td>
</tr>
<tr>
<td>No eGFR 30 yrs⁷</td>
<td>-0.47</td>
<td>-0.17</td>
<td>-0.30</td>
<td>&lt;.001</td>
</tr>
<tr>
<td>Yes eGFR 30 yrs⁸</td>
<td>-2.35</td>
<td>-1.75</td>
<td>-0.29</td>
<td>&lt;.001</td>
</tr>
</tbody>
</table>

### Implications/discussion

This is the first report in a non-referral black and white CKD population, followed for an average of nine years that compares rates of eGFR decline among blacks and whites after adjustment for level of proteinuria. Few studies have compared rates of eGFR decline in blacks versus whites [1]. Most included patients with an eGFR > 60 ml/min/1.73 m², and had only limited follow-up (<3 years) and few included the effect of albuminuria on decline in eGFR. Recently the need for long-term follow-up to adequately assess RFL has been emphasized [4]. Using reciprocal of serum creatinine as a measure of renal function, Brazy and Fitzwilliams found no difference in renal function loss in 112 blacks and 88 white VA patients followed an average of nine years [5]. However, these results did not account for differences in proteinuria. In a relatively small (289 blacks, 945 whites) proteinuric (approximately 800 mg/24 hour) referral VA renal clinic population, with advanced CKD (mean initial eGFR 37 ml/min/1.73 m²) followed for 2.8 years, Kovesdy, et al. found no difference in blacks versus white RFL after adjustment for proteinuria and other comorbidities [6].

In the current study, the higher frequency of 2+proteinuria (100 mg/dl) in blacks versus whites (46.4% versus 24.9%, Table 1) may relate to the finding that although blacks started out with higher first year average eGFR, they were approximately 50% more likely to develop CKD 4 and twice as likely to develop CKD 5 (Table 1). Gansevoort, et al. used a modification of the method of calculating RFL in Table 1 on the Chronic Kidney Disease Prognosis Consortium data [7]. In that report, the difference between initial and final eGFR divided by years of follow-up was used to calculate RFL and log proteinuria appears to influence the rate of change of eGFR per year more so in blacks than whites (Table 1). The mixed effects model (Table 2) utilizing all serum creatinine and urinalysis data points confirms this finding. Compared to whites, blacks had a -0.30 ml/min/1.73 m²/year greater difference in eGFR in the higher renal function group (never had an eGFR <30 ml/min/1.73 m²) and a -0.60 ml/min/1.73 m²/year greater difference in the lower eGFR subset (had at least one e GFR < 30 ml/min/1.73 m²). The failure of the two prior VA based studies to find a RFL difference in blacks versus whites may relate to their shorter follow-up, higher baseline serum creatinine (lower initial eGFR) or the lack of comorbidity adjustment in the current study [3,4].

The higher baseline e GFR in blacks versus whites in the current study may partly relate to the method of choosing patients for entry in to the study. To enroll in the study, patients need at least one serum creatinine of 1.3 mg/dL or greater. A serum creatinine of 1.3 mg/dl would translate in to a higher eGFR by the MDRD equation in blacks versus whites as a result of the MDRD formuale. It is possible that in the current study, blacks had a slower rate of loss of renal function at e GFR levels over 60 ml/min/1.73 m² followed by a slower rate of loss at e GFR <60 ml/min/1.73 m². A recent study by Peralta et al. found the...
reverse, that blacks had a faster RFL than whites at eGFR levels above 60 ml/min/1.73 m² [8].

Proteinuria is the most important factor that affects RFL. In a large Norwegian cohort followed 10.3 years, Hallan, et al. found that after adjusting for proteinuria and eGFR, none of the other variables examined including hypertension, diabetes, obesity, cardiovascular disease, smoking, and education predicted development of treated ESRD [9]. Similarly, Muntner, et al. found that after adjusting for eGFR and albuminuria other comorbidity and life style risk factors did not predict development of ESRD in blacks [10]. On the other hand, McClellan, et al. found that albuminuria prevalence and severity explained some of the excess risk of ESRD in blacks versus whites, but this relationship was attenuated by social and comorbidity factor adjustments [11]. VA care helps minimize some of the socioeconomic issues in black versus white medical care since it is assumed that both races have equal access to care in the VA system.

With regards to survival, similar fractions of blacks and whites in the current study died, 35.3% and 32.7% respectively and death during the study did not appear to be a significant factor in the mixed effects model (Table 2). The majority of the reported data on relative risk of death in blacks versus whites with similar levels of renal function found no difference or a higher death rate in blacks versus whites [12-14]. Thus, although we do not provide a formal survival analysis, a faster RFL in whites who died (censoring of data) is probably not the explanation for the black/white RFL difference.

The pathophysiology behind the putative faster loss of renal function in blacks versus whites may relate to the “genetic hypothesis” [15], which associate the faster decline in blacks to the presence of the APOL1 gene. In a recent report by Parsa et al. using data from the African American Study of Kidney Disease and Hypertension (AASK) and from the Chronic Renal Insufficiency Cohort (CRIC) study, patients with two copies of this gene were twice as likely to reach a renal end point as those with no or one copy of the gene [15]. This faster rate of loss did not appear to be related to degree of blood pressure control or presence of diabetes. Other studies found that the association of APOL1 genotype and faster RFL was not present in diabetics. Socioeconomic factors and other pathophysiological factors may also relate to the difference in black white RFL difference.

Others have reported dipstick proteinuria as a predictor of RFL with a similar finding of faster RFL with higher levels of dipstick proteinuria [16,17]. The utility of the urinalysis protein groups in the current study and the best predictor variable (initial value, all values, change in proteinuria) for log converted urinalysis proteinuria versus RFL needs further exploration. The faster RFL in blacks versus whites that appears to be increased at lower eGFR levels in the current study (Table 2) needs to be replicated in studies that control for social and comorbidity variables (including diagnosis diabetes), survival, as well as episodes of acute kidney injury and renal decline patterns [18].

References