

Association of Urine α 1-Microglobulin with Kidney Function Decline and Mortality in HIV-Infected Women

Vasantha Jotwani, Rebecca Scherzer, Alison Abraham, Michelle M. Estrella, Michael Bennett, Mardge H. Cohen, Marek Nowicki, Anjali Sharma, Mary Young, Phyllis C. Tien, Joachim H. Ix, Mark J. Sarnak, Chirag R. Parikh, and Michael G. Shlipak

Abstract

Background and objectives Despite advances in therapy, HIV-infected individuals remain at higher risk for kidney dysfunction than uninfected individuals. It was hypothesized that urine levels of α 1-microglobulin, a biomarker of proximal tubular dysfunction, would predict kidney function decline and mortality risk in HIV-infected and uninfected women.

Design, setting, participants, & measurements In the Women's Interagency HIV Study, urine α 1-microglobulin and creatinine concentrations were measured in 903 HIV-infected and 287 uninfected women using stored urine from 1999 to 2000, when prevalence of tenofovir use was <1%. Participants were categorized into three categories by level of α 1-microglobulin-to-creatinine ratio, and associations with kidney decline and all-cause mortality over 8 years were evaluated.

Results Urine α 1-microglobulin was detectable in 60% of HIV-infected and 40% of uninfected women ($P < 0.001$). Among HIV-infected women, there were 177 (22%), 61 (7%), and 128 (14%) patients with incident CKD, with 10% annual eGFR decline, and who died, respectively. Compared with HIV-infected women in the lowest α 1-microglobulin category, HIV-infected women in the highest α 1-microglobulin category had a 2.1-fold risk of incident CKD (95% confidence interval, 1.3 to 3.4), 2.7-fold risk of 10% annual eGFR decline (95% confidence interval, 1.2 to 5.9), and 1.6-fold mortality risk (95% confidence interval, 1.0 to 2.6) in models adjusting for kidney risk factors, baseline eGFR, and albuminuria. Among uninfected women, the highest α 1-microglobulin category was associated with 3% (relative risk, 2.2; 95% confidence interval, 1.4 to 3.5) and 5% (relative risk, 2.2; 95% confidence interval, 1.1 to 4.3) annual eGFR decline relative to the lowest α 1-microglobulin category.

Conclusions Proximal tubular dysfunction, indicated by urine α 1-microglobulin, was independently associated with kidney function decline in HIV-infected and uninfected women and mortality risk among HIV-infected women.

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Introduction

Kidney disease is an established complication of HIV infection. HIV-infected persons are at higher risk for proteinuria, CKD, and ESRD compared with uninfected individuals, even after controlling for traditional kidney disease risk factors (1–6). However, standard clinical measures of kidney disease, such as serum creatinine and dipstick proteinuria, are inadequate in their ability to detect early kidney damage (7). There is a strong clinical need for biomarkers that can identify kidney dysfunction at earlier stages and localize pathology within the nephron to provide prognostic and therapeutic guidance in HIV-infected individuals. Although serum cystatin C may be more sensitive than serum creatinine for the estimation of GFR in HIV-infected individuals (8,9), we hypothesize that tubular injury and dysfunction may occur earlier than clinically apparent reductions in glomerular filtration function.

We previously showed that two tubular injury markers, IL-18 and kidney injury molecule-1 (KIM-1), are present in higher levels in the urine of HIV-infected women compared with uninfected controls and that higher urine IL-18 and KIM-1 predict longitudinal eGFR decline and mortality, independent of traditional kidney disease risk factors and albuminuria (10–12). In contrast to IL-18 and KIM-1, which are believed to be produced by proximal tubular epithelial cells in the setting of injury (13,14), α 1-microglobulin (α 1m) is a 26-kD lipocalin that is filtered at the glomerulus but fully reabsorbed by proximal tubular epithelial cells, where it is degraded (15). The presence of α 1m in the urine is, therefore, indicative of proximal tubular dysfunction (16). A recent proteomic profiling study identified α 1m as one of the earliest biomarkers detectable in urine after ischemic injury (17), suggesting that urine α 1m may capture proximal tubular dysfunction at the earliest stages.

Due to the number of contributing authors, the affiliations are provided in the Supplemental Material.

Correspondence:

Dr. Michael G. Shlipak, Department of Medicine, San Francisco VA Medical Center, 4150 Clement Street, 94121, San Francisco, CA. Email: Michael.Shlipak@ucsf.edu

In this study of HIV-infected and uninfected women enrolled in the Women's Interagency HIV Study (WIHS), we tested the hypotheses that HIV infection is associated with a higher prevalence of detectable urine $\alpha 1m$, that urine $\alpha 1m$ predicts kidney function decline in HIV-infected and uninfected women, and that urine $\alpha 1m$ predicts mortality risk among HIV-infected women, independent of albuminuria and three previously studied tubular injury markers: IL-18, KIM-1, and neutrophil gelatinase-associated lipocalin (NGAL).

Materials and Methods

Study Population

The WIHS is a multicenter, prospective cohort study that enrolled 3067 HIV-infected and 1070 uninfected women from six United States locations: Bronx, Brooklyn, Chicago, Los Angeles, San Francisco, and Washington DC in 1994–1995, 2001–2002, and 2011–2012. Details of study design, data collection methods, and baseline characteristics are published elsewhere (18,19). Briefly, HIV-infected participants were recruited to be representative of the population of HIV-infected women in the community. Recruitment of uninfected participants targeted women who engaged in high-risk behaviors (defined by self-reported injection drug use or high-risk sexual behavior within the year before enrollment). Participants underwent semiannual visits that included an interviewer-administered questionnaire, a physical examination, and collection of laboratory specimens.

The WIHS Kidney Aging Study was designed as a nested cohort study to investigate the onset of kidney disease in the setting of HIV infection using stored urine and serum specimens. Baseline urine and serum samples were collected between October of 1999 and March of 2000. Serum was also collected at two visits occurring at approximately 4 and 8 years after the baseline of this ancillary study. For this study, we included all 903 HIV-infected and 287 uninfected women who had stored urine available and at least one follow-up serum cystatin C measurement. The institutional review boards of the participating institutions approved the study protocol at all WIHS study sites, and informed consent was obtained from all study participants. This study of kidney injury was also approved by the University of California, San Francisco, the San Francisco Veterans Affairs Medical Center, and the Yale Committees on Human Research.

Exposure Variables

Urine $\alpha 1m$ was measured at the Cincinnati Children's Hospital Medical Center Biomarker Laboratory by a commercially available assay (Siemens BNII nephelometer; Siemens, Munich, Germany). The detectable limit of the $\alpha 1m$ assay was 0.6 mg/dl. Intra- and interassay coefficients of variation (CVs) were 5.2% and 13.2%, respectively. Urine creatinine was measured by colorimetric enzyme assay using a Siemens Dimension Xp and plus HM clinical analyzer (Siemens). All urine specimens were in continuous storage at -80°C until biomarker measurement without prior freeze-thaw; mean urine storage time was 11 years. Laboratory personnel performing the biomarker assays were blinded to clinical information regarding WIHS participants, including their HIV status, and the samples were evaluated in random order. The associations of urine $\alpha 1m$

with kidney function decline were additionally compared with prior analyses of urine albumin-to-creatinine ratio (ACR), IL-18, KIM-1, and NGAL (10).

Outcomes

The primary outcomes of this study were longitudinal kidney function decline and all-cause mortality. Kidney function was estimated using the Chronic Kidney Disease Epidemiology Collaboration equation for serum cystatin C ($e\text{GFR}_{\text{Cys}}$) (20,21). We chose *a priori* to estimate GFR by cystatin C rather than creatinine, because it is less susceptible to bias by muscle mass and health status (8). Cystatin C was measured centrally at the University of California at Los Angeles Clinical Immunology Research Laboratory using a particle-enhanced immunoturbidimetric assay (Gentian, Moss, Norway), which has been calibrated against the new World Standard Reference material ERM-DA471/IFCC (20). The assay was run concurrently from all three clinic visits, minimizing concerns for drift in the assay over time. Intra-assay CVs, on the basis of 10 replicates, were $<2\%$ at serum concentrations of 0.7 and 1.1 mg/L. Interassay CVs were 4.4% and 3.9% at serum concentrations of 0.8 and 2.2 mg/L. As in our prior work, $e\text{GFR}$ was capped at 120 ml/min per 1.73 m^2 , because the equation has not been validated at higher values (22).

We analyzed $e\text{GFR}_{\text{Cys}}$ as a continuous outcome expressed as change in $e\text{GFR}_{\text{Cys}}$ in milliliters per minute per 1.73 m^2 per year over approximately 8 years of follow-up. Among HIV-infected women, two dichotomized kidney outcomes were analyzed: (1) incident CKD (defined as $e\text{GFR}_{\text{Cys}} < 60\text{ ml/min per } 1.73\text{ m}^2$ at either of two follow-up visits among HIV-infected women with baseline $e\text{GFR}_{\text{Cys}} \geq 60\text{ ml/min per } 1.73\text{ m}^2$) and (2) rapid decline (defined as 10% or greater annual decline in $e\text{GFR}_{\text{Cys}}$). Rapid decline was calculated as the relative change in $e\text{GFR}_{\text{Cys}}$ from baseline to each follow-up visit for each participant. As a sensitivity analysis, we required an $e\text{GFR}_{\text{Cys}}$ decline of $\geq 1\text{ ml/min per } 1.73\text{ m}^2$ per year of follow-up for all patients with incident CKD and found that this approach eliminated only one of 177 patients. Because of the small numbers of patients with incident CKD ($n=16$) and rapid decline ($n=8$) among uninfected women, we evaluated 3% and 5% annual $e\text{GFR}_{\text{Cys}}$ declines as dichotomous outcomes in these participants.

Vital status and date of death were determined using the National Death Index and data from medical records and providers. Detailed methods have been described in prior studies (23–25). Deaths were ascertained over 8 years of follow-up in parallel with the kidney decline analyses. As a sensitivity analysis, we updated vital status through April of 2013, the most recent possible update at the time of manuscript submission. Because there were only 13 deaths among the uninfected participants, we studied the associations of $\alpha 1m$ with mortality risk only in HIV-infected women.

Covariates

The following characteristics were tested as candidate covariates in all multivariate models: age and race/ethnicity, systolic and diastolic BP, antihypertensive use, diabetes (defined using confirmatory criteria for fasting glucose $\geq 126\text{ mg/dl}$, self-reported diabetes, self-reported diabetes medication use, or hemoglobin A1c $\geq 6.5\%$), cigarette smoking

status (current, former, or never), menopause status, LDL and HDL cholesterol, triglycerides, serum albumin, body mass index, waist circumference, hepatitis C virus (HCV) infection (confirmed by detectable HCV RNA after a positive HCV antibody result), and current heroin use. Candidate HIV-related characteristics included current CD4 lymphocyte count, nadir CD4 lymphocyte count, history of AIDS diagnosis, current HIV viral load, current highly active antiretroviral therapy use, current nucleoside reverse transcription inhibitor use, current non-nucleoside reverse transcription inhibitor use, and current protease inhibitor use. During the study period, the prevalence of tenofovir use increased from <1% at baseline to 49% at the end of the 8-year follow-up period. Multiple imputation with the Markov chain Monte Carlo method was used to impute missing covariates with five imputations to yield approximately 95% relative efficiency (26). The percentage of missing observations for each covariate ranged from <1% to 15%.

Statistical Analyses

Forty-five percent of participants had undetectable urine α 1m, and the distribution among those with detectable α 1m was right-skewed. Because of the left-censored nature of the data, we analyzed α 1m using three approaches: a dichotomized variable (detectable or undetectable), a log-transformed continuous variable using models that accommodate left-censored data, and an ordinal variable with three categories of urine α 1m-to-creatinine ratio (α 1m/cr). For analyses of HIV-infected and uninfected women modeling α 1m as an ordinal variable, category 1 of α 1m/cr included all participants with undetectable urine α 1m. The remaining participants were divided at the median value of urine α 1m/cr into categories 2 and 3 for HIV-infected and uninfected participants separately.

We compared baseline characteristics of HIV-infected women across the three categories of urine α 1m/cr using chi-squared and Kruskal–Wallis tests for categorical and continuous variables, respectively; these comparisons were replicated among the uninfected participants. We then used Poisson regression with a robust variance estimator (27) to assess the cross-sectional association of HIV infection with detectable urine α 1m and identify factors associated with detectable urine α 1m among HIV-infected and uninfected women. As an alternate approach, we used multivariable generalized γ -regression models to identify factors associated with higher urine α 1m concentration in HIV-infected and uninfected women separately. Similar to the Tobit regression method, generalized γ -regression models accommodate left-censored data but also allow log transformation of urine α 1m to normalize its right-skewed distribution. Results were back-transformed to produce estimated percentage differences in urine α 1m attributable to each factor.

Next, we used linear mixed models to evaluate the associations of urine α 1m/cr categories with the continuous outcome of eGFR_{Cys} with random intercepts and slopes across time to account for the correlation between repeated measures at baseline, year 4, and year 8. Multivariate models sequentially adjusted for demographics, traditional risk factors for kidney disease progression, baseline eGFR_{Cys}, ACR, and the previously studied biomarkers urine IL-18/cr, KIM-1/cr, and NGAL/cr. As a

sensitivity analysis, we additionally adjusted for tenofovir use during follow-up. To compare the effect size of α 1m/cr with those of the previously studied biomarkers, we modified our previously published WIHS analyses (10) assessing the associations of ACR, IL-18/cr, KIM-1/cr, and NGAL/cr with eGFR_{Cys} by incorporating α 1m/cr into the final multivariate-adjusted models for each biomarker.

We then used Poisson regression with a robust variance estimator to assess the associations of urine α 1m/cr with incident CKD and 10% annual eGFR_{Cys} decline among HIV-infected individuals and with 3% and 5% annual eGFR_{Cys} declines among uninfected individuals. We calculated risk ratios for each outcome using category 1 as the reference category. Women with eGFR_{Cys}<60 ml/min per 1.73 m² at baseline were excluded from the incident CKD analysis. To evaluate for possible effect modification by tenofovir use during follow-up, analyses of HIV-infected participants were stratified by tenofovir use during follow-up (ever used versus never used), and we evaluated an interaction term (α 1m \times tenofovir) for statistical significance in the overall model.

To account for losses to follow-up in all kidney function decline analyses, we adjusted estimates using an inverse probability weighting approach by modeling the participant's probability of having a nonmissing outcome, with separate weights calculated at each visit (28). The inverse of this probability was then used as a weight applied to persons with known outcomes in the multivariable regression analyses of kidney decline.

Finally, we used unadjusted generalized additive models to produce a spline plot depicting the probability of mortality over the detectable range of α 1m/cr in HIV-infected participants. We obtained *P* values for the association of α 1m/cr with mortality and the test of nonlinearity. We also evaluated the associations of urine α 1m/cr categories with all-cause mortality using multivariable Cox proportional hazards and the same series of adjusted models as for the kidney function decline outcomes.

Spearman coefficients were used to evaluate correlations between α 1m, IL-18, KIM-1, NGAL, ACR, and urine creatinine. Because of the moderately strong intercorrelations between all biomarkers and urine creatinine (Supplemental Table 1), primary analyses standardized the biomarkers to urine creatinine. In sensitivity analyses, we evaluated the associations of unstandardized biomarker levels with kidney decline and mortality using the methods described above.

All analyses were conducted using SAS version 9.3 (SAS Institute Inc., Cary, NC).

Results

Baseline Characteristics of WIHS Participants

The median age was 41 years among the 903 HIV-infected women and 40 years among the 287 uninfected women included in this study. At baseline, 9% of HIV-infected and 2% of uninfected women had an eGFR_{Cys}<60 ml/min per 1.73 m² (*P*<0.001); 13.8% of the participants were lost to follow-up after the second visit; these individuals provided a mean follow-up time of 3.7 years.

HIV-infected women in the highest category of α 1m/cr were older, were more often African American, and had higher prevalence of smoking, diabetes mellitus, hypertension,

HCV infection, and albuminuria as well as lower CD4 counts and higher HIV RNA levels (Table 1). Among uninfected participants, the highest $\alpha 1m/cr$ category was characterized by older age and higher prevalence of smoking, diabetes mellitus, hypertension, HCV infection, heroin use, and albuminuria (Table 2).

Prevalence of Detectable Urine $\alpha 1m$ in HIV-Infected and Uninfected Women

Urine $\alpha 1m$ was detectable in 545 (60%) HIV-infected and 114 (40%) uninfected women ($P<0.001$). HIV infection remained associated with a 51% higher prevalence of detectable $\alpha 1m$ after multivariable adjustment for age, race, and traditional kidney disease risk factors (Table 3). The association

was slightly attenuated with additional adjustment for ACR and $eGFR_{Cys}$ but remained statistically significant.

Factors Associated with Urinary $\alpha 1m$ in HIV-Infected and Uninfected Women

Among HIV-infected participants, African-American race (relative risk [RR], 1.32; 95% confidence interval [95% CI], 1.04 to 1.69; $P=0.02$), current smoking (RR, 1.24; 95% CI, 1.04 to 1.47; $P=0.01$), and CD4 lymphocyte count <200 cells/ mm^3 (RR, 1.27; 95% CI, 1.04 to 1.56; $P=0.02$) were independently associated with detectable $\alpha 1m$. When we modeled urine $\alpha 1m$ concentration as a continuous outcome, African-American race was associated with 52% higher $\alpha 1m$ (95% CI, 22% to 90%; $P<0.001$), HCV

Table 1. Baseline characteristics of HIV-infected women in the Women's Interagency HIV Study stratified by urine $\alpha 1$ -microglobulin category

Characteristic	1 (n=358)	2 (n=272)	3 (n=273)	P Value
Range of $\alpha 1m/cr$ (mg/g)	Below detectable	1.2–10.2	>10.2	
Baseline age (yr)	40 (35–44)	40 (36–45)	43 (38–48)	<0.001
Race				
African American	180 (50%)	151 (56%)	188 (69%)	<0.001
Caucasian	92 (26%)	46 (17%)	37 (14%)	
Other	86 (24%)	75 (28%)	48 (18%)	
Cigarette smoking				
Current	149 (42%)	144 (53%)	167 (61%)	<0.001
Past	100 (28%)	72 (26%)	52 (19%)	
Never	109 (30%)	56 (21%)	54 (20%)	
Diabetes mellitus	25 (7%)	23 (8%)	38 (14%)	0.01
Hypertension	72 (20%)	68 (25%)	85 (31%)	<0.01
Antihypertensive use	28 (8%)	28 (10%)	41 (15%)	0.02
Menopause	55 (16%)	45 (17%)	82 (31%)	<0.001
Hepatitis C	87 (25%)	77 (28%)	114 (42%)	<0.001
Current heroin use	11 (3%)	12 (4%)	20 (7%)	0.04
LDL (mg/dl)	107 (85–133)	106 (80–136)	95 (71–125)	<0.001
HDL (mg/dl)	44 (35–55)	45 (37–57)	42 (33–57)	0.16
Triglycerides (mg/dl)	135 (88–206)	117 (89–179)	142 (101–197)	0.004
Body mass index (kg/m ²)	26 (24–31)	27 (24–31)	25 (23–31)	0.02
Waist circumference (cm)	88 (81–98)	89 (80–100)	87 (78–98)	0.17
Current CD4 (cells/mm ³)	448 (282–647)	386 (234–534)	363 (195–548)	<0.001
Nadir CD4 (cells/mm ³)	235 (131–348)	214 (114–313)	186 (85–294)	0.001
History of AIDS	148 (41%)	135 (50%)	158 (58%)	<0.001
HIV viral load (copies/ml)				
≤ 80	126 (35%)	77 (29%)	71 (26%)	<0.01
81–1999	90 (25%)	60 (22%)	53 (19%)	
2000–9999	55 (15%)	48 (18%)	44 (16%)	
>10,000	84 (24%)	85 (31%)	104 (38%)	
Current HAART use	221 (62%)	153 (56%)	155 (57%)	0.30
Current NRTI use	254 (71%)	176 (65%)	172 (63%)	0.08
Current NNRTI use	97 (27%)	68 (25%)	79 (29%)	0.58
Current protease inhibitor use	158 (44%)	109 (40%)	110 (40%)	0.50
Albuminuria ^a	57 (16%)	46 (17%)	109 (40%)	<0.001
$eGFR_{Cys}$ (ml/min per 1.73 m ²)	97 (83–110)	91 (80–105)	76 (62–89)	<0.001
$eGFR_{Cys}<60$ ml/min per 1.73 m ²	15 (4%)	10 (4%)	59 (22%)	<0.001
Serum albumin (g/dl)	4.1 (3.8–4.4)	4.0 (3.8–4.3)	3.9 (3.6–4.2)	<0.001

Data are presented as medians (interquartile ranges) or numbers (percentages). $\alpha 1$ -Microglobulin is standardized to urine creatinine. Category 1 comprises all participants with undetectable urine $\alpha 1$ -microglobulin. $\alpha 1m/cr$, $\alpha 1$ -Microglobulin-to-creatinine ratio; HAART, highly active antiretroviral therapy; NRTI, nucleoside reverse transcription inhibitor; NNRTI, non-nucleoside reverse transcription inhibitor; $eGFR_{Cys}$, $eGFR$ by cystatin C.

^aDefined as a positive urine dipstick result ($\geq 1+$) or urine albumin-to-creatinine ratio >30 mg/g.

Table 2. Baseline characteristics of HIV-uninfected women in the Women's Interagency HIV Study stratified by urine α 1-microglobulin category

Characteristic	1 (n=173)	2 (n=57)	3 (n=57)	P Value
Range of α 1m/cr (mg/g)	Below detectable	1.3–7.5	>7.5	
Baseline age (yr)	38 (32–43)	42 (36–46)	42 (38–48)	<0.001
Race				
African American	99 (57%)	41 (72%)	37 (65%)	0.31
Caucasian	20 (12%)	6 (11%)	6 (11%)	
Other	54 (31%)	10 (18%)	14 (25%)	
Cigarette smoking				
Current	84 (49%)	39 (68%)	46 (81%)	<0.001
Past	49 (28%)	7 (12%)	6 (11%)	
Never	40 (23%)	11 (19%)	5 (9%)	
Diabetes mellitus	11 (6%)	5 (9%)	10 (18%)	0.04
Hypertension	39 (23%)	11 (19%)	30 (53%)	<0.001
Antihypertensive use	18 (10%)	6 (11%)	11 (19%)	0.19
Menopause	20 (12%)	6 (11%)	14 (25%)	0.03
Hepatitis C	23 (14%)	16 (28%)	23 (40%)	<0.001
Current heroin use	9 (5%)	2 (4%)	12 (21%)	<0.001
LDL (mg/dl)	107 (89–131)	104 (84–128)	100 (84–132)	0.71
HDL (mg/dl)	52 (44–63)	49 (41–61)	48 (39–59)	0.29
Triglycerides (mg/dl)	94 (71–141)	107 (69–154)	119 (87–156)	0.08
Body mass index (kg/m ²)	29 (24–35)	30 (26–34)	29 (25–32)	0.67
Waist circumference (cm)	92 (80–103)	95 (83–107)	94 (81–99)	0.55
Albuminuria ^a	15 (9%)	6 (11%)	12 (21%)	0.04
eGFR _{Cys} (ml/min per 1.73 m ²)	106 (96–117)	97 (87–112)	97 (79–112)	<0.001
eGFR _{Cys} <60 ml/min per 1.73 m ²	1 (2%)	5 (9%)	0	<0.001
Serum albumin (g/dl)	4.1 (3.9–4.3)	4.0 (3.8–4.2)	4.0 (3.8–4.3)	0.08

Data are presented as medians (interquartile ranges) or numbers (percentages). α 1-Microglobulin is standardized to urine creatinine. Category 1 comprises all participants with undetectable urine α 1-microglobulin.

^aDefined as a positive urine dipstick result ($\geq 1+$) or urine albumin-to-creatinine ratio >30 mg/g.

Table 3. Association of HIV infection with detectable urine α 1-microglobulin in HIV-infected (n=903) versus uninfected (n=287) women at baseline visit

Model	Prevalence Ratio ^a (95% Confidence Interval)	P Value
Demographic adjusted ^b	1.52 (1.24 to 1.86)	<0.001
Multivariate adjusted ^c	1.51 (1.22 to 1.85)	<0.001
Multivariate adjusted ^c +ACR	1.46 (1.19 to 1.81)	<0.001
Multivariate adjusted ^c +ACR+eGFR _{Cys}	1.34 (1.08 to 1.67)	<0.01

ACR, albumin-to-creatinine ratio.

^aAdjusted prevalence ratios of detectable α 1-microglobulin in HIV-infected versus uninfected women calculated using multivariable Poisson regression models.

^bAdjusted for age and race.

^cAdjusted for age, race, hypertension, diabetes mellitus, hepatitis C virus infection, smoking, LDL, triglycerides, and body mass index.

was associated with 39% higher α 1m (95% CI, 10% to 77%; $P<0.01$) in adjusted analyses. HIV-related factors independently associated with higher urine α 1m levels included CD4 lymphocyte count <200 cells/mm³ (65%; 95% CI, 34% to 103%; $P<0.001$) and history of AIDS (30%; 95% CI, 10% to 53%; $P=0.002$).

Among uninfected women, factors associated with higher levels of α 1m included diabetes mellitus (66%; 95% CI, 7% to 157%; $P=0.02$), heroin use (79%; 95% CI, 11 to 190; $P=0.02$), HCV infection (55%; 95% CI, 11 to 117; $P=0.01$), and current smoking (82%; 95% CI, 36% to 145%; $P<0.001$).

Association of Urine α 1m with Kidney Function Decline in HIV-Infected and Uninfected Women

Over the approximately 8-year follow-up period, the rate of annual eGFR_{Cys} decline was -1.18 (95% CI, -1.29 to -1.06) ml/min per 1.73 m² in HIV-infected women and -0.97 (95% CI, -1.16 to -0.79) ml/min per 1.73 m² in uninfected women.

Compared to HIV-infected women with undetectable α 1m, HIV-infected women in the highest category of α 1m/cr experienced faster eGFR_{Cys} decline over the follow-up period. The adjusted difference in annual eGFR_{Cys} change between categories 3 and 1 was -0.19 (95% CI, -0.22 to -0.16) ml/min per 1.73 m² per year in multivariate analyses and -0.16 (95% CI, -0.19 to -0.13) ml/min per 1.73 m² per year after additional adjustment for ACR, IL-18/cr,

infection was associated with 22% higher α 1m (95% CI, 2% to 46%; $P=0.03$), hypertension was associated with 29% higher α 1m (95% CI, 7% to 57%; $P=0.01$), and menopause

KIM-1/cr, and NGAL/cr. Adjustment for tenofovir use during follow-up did not alter the association of $\alpha 1m/cr$ with annual eGFR change. Compared with ACR, IL-18, KIM-1, and NGAL, urine $\alpha 1m$ was associated with the largest annual change in eGFR_{Cys} in multivariate models adjusting for all four biomarkers in creatinine-standardized (Figure 1) and unstandardized analyses (Supplemental Figure 1).

Among HIV-infected women, 177 (22%) and 61 (7%) cases of incident CKD and 10% annual eGFR_{Cys} decline occurred, respectively. Relative to those with undetectable $\alpha 1m$, the highest category of $\alpha 1m/cr$ was associated with 2- to 3-fold risks of incident CKD and rapid decline after adjustment for demographics, baseline eGFR_{Cys}, traditional kidney disease risk factors, and HIV-related factors (Table 4). Even after additional adjustment for ACR, IL-18, KIM-1, and NGAL, the highest category of $\alpha 1m/cr$ remained associated with a 1.9-fold risk of incident CKD and a 2.8-fold risk of rapid decline. Results were similar when urine $\alpha 1m$ was modeled without standardization to urine creatinine (Supplemental Table 2) and when we required an eGFR_{Cys} decline of ≥ 1 ml/min per 1.73 m² per year of follow-up for all patients with incident CKD. There were no statistically significant interactions with kidney decline outcomes when individuals were stratified by tenofovir use during follow-up.

Among the uninfected participants, we examined the associations of urine $\alpha 1m/cr$ with continuous eGFR_{Cys} decline and two dichotomous outcomes: 3% and 5% annual eGFR_{Cys} decline. Compared to individuals with undetectable $\alpha 1m$, the highest category of $\alpha 1m/cr$ was associated with -0.21 (95% CI, -0.28 to -0.13 ; $P < 0.001$) ml/min per 1.73 m² annual eGFR_{Cys} decline in multivariate analyses

and -0.27 (95% CI, -0.34 to -0.19 ; $P < 0.001$) ml/min per 1.73 m² annual eGFR_{Cys} decline after additional adjustment for ACR, IL-18/cr, KIM-1/cr, and NGAL/cr. Individuals in the highest $\alpha 1m/cr$ category had approximately 2-fold adjusted risks of 3% and 5% annual eGFR_{Cys} decline relative to those with undetectable $\alpha 1m$ (Table 5).

Association of Urine $\alpha 1m$ with Mortality in HIV-Infected Women

During the 8 years of follow-up in the WIHS Kidney Aging Study, there were 128 (14%) deaths among 903 HIV-infected women included in this study. When we used a spline plot to assess probability of mortality across the range of detectable urine $\alpha 1m/cr$ (Figure 2), higher urine $\alpha 1m/cr$ levels were incrementally associated with higher mortality ($P < 0.001$). Although the slope appeared to be steeper at higher values of $\alpha 1m/cr$, the test for nonlinearity did not reach statistical significance ($P = 0.37$).

Compared to women with undetectable $\alpha 1m$, women in the highest category of $\alpha 1m/cr$ had a 1.7-fold risk of all-cause mortality in multivariate analyses (Table 4). Results were similar when $\alpha 1m$ was not standardized to urine creatinine (hazard ratio [HR], 1.64; 95% CI, 0.94 to 2.87) (Supplemental Table 2). Updating vital status through April of 2013 resulted in a total of 224 deaths among 903 HIV-infected participants, with a median follow-up of 13 years. The highest category of $\alpha 1m/cr$ remained independently associated with mortality in the multivariate-adjusted model (HR, 1.49; 95% CI, 1.05 to 2.11; $P = 0.03$), but the association was no longer statistically significant after adjustment for albuminuria and the additional biomarkers (HR, 1.42; 95% CI, 0.98 to 2.05; $P = 0.06$).

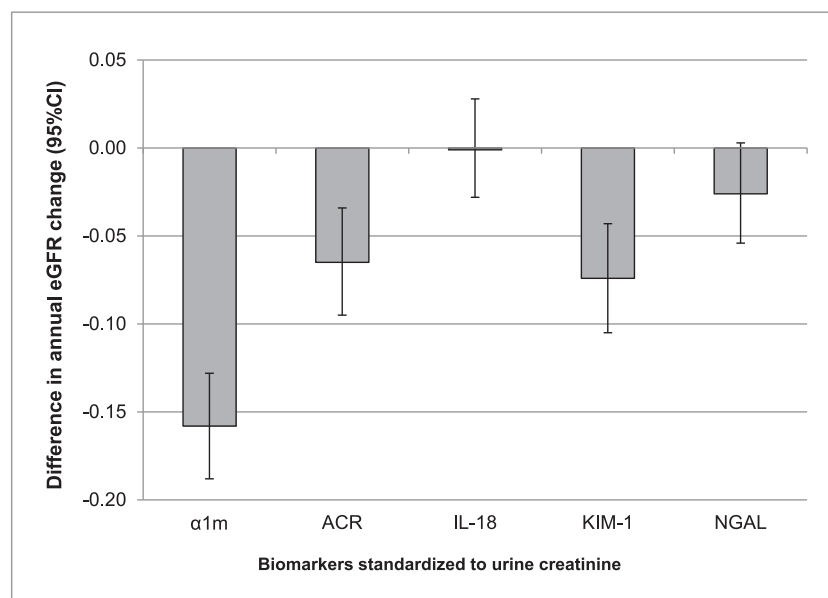


Figure 1. | Magnitudes of associations with annual eGFR change across creatinine-standardized urine biomarkers. All estimates were derived from multivariate models comparing category 3 with category 1 for $\alpha 1m$ -microglobulin ($\alpha 1m$) and comparing highest with lowest tertile for albumin-to-creatinine ratio (ACR), IL-18, kidney injury molecule-1 (KIM-1), and neutrophil gelatinase-associated lipocalin (NGAL). Multivariate models were adjusted for age, race, traditional kidney disease risk factors, HIV-related factors, and all displayed biomarkers. Results are reported as eGFR change in milliliters per minute per 1.73 m² per year (95% confidence interval [95% CI]).

Table 4. Associations of urine α1-microglobulin with kidney function decline and mortality in HIV-infected women.

	Incident CKD			10% Decline in eGFR _{Cys}			All-Cause Mortality		
	C1	C2	C3	C1	C2	C3	C1	C2	C3
Range of α1m/cr (mg/g)	—	1.2–10.2	>10.2	—	1.2–10.2	>10.2	—	1.2–10.2	>10.2
No. at risk	343	262	214	358	272	273	358	272	273
No. events	48	50	79	9	17	35	29	36	63
	Risk Ratio (95% CI)	Risk Ratio (95% CI)	Risk Ratio (95% CI)	Risk Ratio (95% CI)	Risk Ratio (95% CI)	Risk Ratio (95% CI)	Risk Ratio (95% CI)	Hazard Ratio (95% CI)	Hazard Ratio (95% CI)
Demographic adjusted ^a	Reference	1.35 (0.58 to 3.14)	3.56 (2.16 to 5.88)	Reference	2.53 (1.14 to 5.64)	4.53 (2.21 to 9.28)	Reference	1.60 (0.98 to 2.63)	2.66 (1.70 to 4.17)
Multivariate adjusted ^b	—	1.31 (0.62 to 2.74)	2.36 (1.42 to 3.92)	—	2.21 (1.01 to 4.84)	2.99 (1.42 to 6.29)	—	1.47 (0.89 to 2.44)	1.69 (1.06 to 2.69)
Adjusted+ACR ^c	—	1.34 (0.66 to 2.72)	2.08 (1.28 to 3.38)	—	2.38 (1.11 to 5.12)	2.70 (1.23 to 5.92)	—	1.50 (0.91 to 2.49)	1.61 (1.00 to 2.59)
Adjusted+ACR, IL-18, KIM-1, NGAL ^d	—	1.13 (0.65 to 1.98)	1.87 (1.23 to 2.84)	—	2.68 (1.26 to 5.72)	2.76 (1.28 to 5.98)	—	1.65 (0.98 to 2.76)	1.73 (1.07 to 2.82)

Standardized to urine creatinine and stratified by categories. Category 1 comprises all participants with undetectable urine α1-microglobulin. All analyses use category 1 as the reference category. eGFR_{Cys}, cystatin C-based eGFR; C1, category 1; C2, category 2; C3, category 3; α1m/cr, α1-microglobulin-to-creatinine ratio; 95% CI, 95% confidence interval; ACR, albumin-to-creatinine ratio; KIM-1, kidney injury molecule-1; NGAL, neutrophil gelatinase-associated lipocalin.

^a Adjusted for age and race.

^b Adjusted for age, race, baseline eGFR_{Cys}, hypertension, diabetes mellitus, hepatitis C virus infection, HIV viral load, CD4 lymphocyte count, antiretroviral therapy use, and serum albumin.

^c Adjusted for all covariates listed above with the addition of ACR.

^d Adjusted for all covariates listed above with the addition of ACR, IL-18, KIM-1, NGAL, and liver fatty acid binding protein.

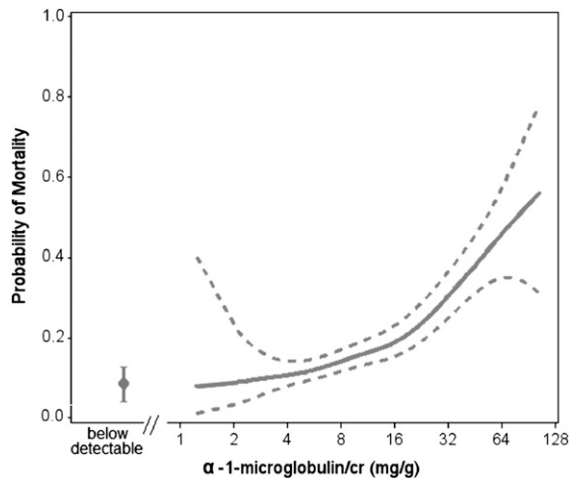


Figure 2. | Spline plot displaying unadjusted association of urine α 1-microglobulin with mortality over a median follow-up of 8 years. Solid line denotes predicted probability of mortality; dotted lines represent 95% confidence bounds. Below detectable estimate represents the proportion of deaths in individuals with undetectable α 1-microglobulin. The highest 2.5% of values were truncated. $P < 0.001$ for association of urine α 1-microglobulin-to-creatinine ratio with mortality. $P = 0.37$ for test of nonlinearity. cr, creatinine.

–0.22-ml/min per 1.73 m² annual eGFR change compared with systolic BP < 120, and hemoglobin A1c levels of 8–8.9 were associated with –0.23 ml/min per 1.73 m² annual eGFR decline compared with hemoglobin A1c levels of < 7. These observations suggest that associations of this magnitude are clinically important. Future studies must validate our findings in additional cohorts of individuals with varied risk factors for kidney disease progression.

In persons with HIV infection, proximal tubular dysfunction is an increasingly common presentation of renal toxicity (35), with one study reporting a 25% prevalence of tubular damage in individuals on antiretroviral therapy without proteinuria or significant impairment of glomerular function (36). Recent evidence also suggests that α 1m and other low molecular mass proteins can detect subclinical impairments in proximal tubular function among HIV-infected individuals receiving tenofovir, a nucleotide analog reverse transcription inhibitor associated with a particular form of proximal tubular dysfunction known as Fanconi's syndrome (37), AKI (38), and CKD (39). In a 48-week trial, Vrouenraets *et al.* (40) randomized 20 individuals with HIV infection to continue therapy with zidovudine/lamivudine or switch to tenofovir/emtricitabine and noted an approximately 50% rise ($P = 0.09$) in mean urine α 1m/cr in subjects randomized to tenofovir/emtricitabine compared with unchanged urine α 1m/cr levels in persons continued on zidovudine/lamivudine. Similarly, Post *et al.* reported higher urine levels at 48 weeks of retinol-binding protein and β 2-microglobulin by 50% ($P < 0.001$) and 24% ($P < 0.001$), respectively, in antiretroviral-naïve individuals randomized to tenofovir/emtricitabine ($n = 193$) compared with abacavir/lamivudine ($n = 192$) (41). Although urine α 1m was not measured in that study, retinol-binding protein and β 2-microglobulin undergo transport into proximal

tubular cells by the same endocytic receptor as urine α 1m (31), suggesting that these low molecular mass proteins may provide similar information when measured in urine. Notably, none of the subjects in the trial described above met the protocol definition for proximal tubular dysfunction, which relied on other criteria, such as hypophosphatemia, hypokalemia, nondiabetic glycosuria, metabolic acidosis, and rise in serum creatinine. This finding highlights the need for novel biomarkers that capture drug-related toxicity before the onset of clinical Fanconi's syndrome or reduction in glomerular filtration function.

This study offers several important implications for clinical care. First, the associations of urine α 1m with kidney function decline and mortality suggest that minor impairments of proximal tubular function are not benign. Additional studies should investigate the potential reversibility of proximal tubular dysfunction to determine whether tubular dysfunction is a modifiable risk factor for kidney decline. Larger studies of α 1m in diverse populations may also enable the identification of a threshold α 1m level beyond which kidney risk escalates. Second, our study utilized urine specimens collected prior to the widespread use of tenofovir. Although subsequent tenofovir use did not modify the observed associations of urine α 1m with kidney decline outcomes, our study was not designed or powered to determine whether urinary α 1m identifies a subset of individuals who are particularly susceptible to tubular toxicity from tenofovir or other drugs. Future clinical studies could specifically evaluate the potential applications of α 1m in risk stratification before initiation of tenofovir and early detection of tubular toxicity during therapy. Additionally, a growing number of uninfected individuals at high risk for HIV acquisition are now receiving pre-exposure prophylaxis with tenofovir. Early recognition of tubular toxicity from tenofovir will be particularly important in this population of predominantly young individuals, in whom the risks of therapy are incompletely delineated. Third, biomarkers sensitive for the detection of tubular dysfunction and injury could revolutionize our methods for detecting nephrotoxicity in drug trials, which often fail to identify drug-related nephrotoxicity in initial stages of development because of their sole reliance on serum creatinine and proteinuria.

There are several limitations to this study. First, because we exclusively studied women, we cannot generalize these results to men. Second, biomarker measurements were made using urine samples collected over a decade ago. However, with protein degradation over time, we would expect our results to shift toward null findings. Third, although it would have been desirable to confirm the diagnosis of CKD with at least two separate eGFR measurements at the 4- and 8-year time points, this was not feasible because of the availability of only one serum sample per patient at each of these visits. Fourth, to ensure comparability with the HIV-infected women, recruitment of the uninfected WIHS cohort targeted women who engage in high-risk behaviors, such as injection drug use and high-risk sexual behavior (19). Therefore, the uninfected women in this study had a high prevalence at baseline of established risk factors for kidney disease. This may limit the generalizability of our findings to healthier, uninfected populations. Fifth, because of the low rates of incident CKD, 10% annual eGFR_{Cys}

decline, and mortality in the uninfected participants, we lacked sufficient power to analyze the associations of $\alpha 1m$ with these outcomes. Sixth, we did not have access to serum concentrations of $\alpha 1m$, and therefore, we cannot exclude the possibility that higher serum levels in susceptible individuals contributed to our observations. Seventh, although we adjusted for multiple potential confounders, the possibility of residual confounding exists for our associations of urine $\alpha 1m$ with kidney function decline and mortality.

In conclusion, we found that urine $\alpha 1m$ independently predicted kidney function decline in HIV-infected and uninfected women as well as mortality risk among HIV-infected women, highlighting the importance of proximal tubular dysfunction to longitudinal outcomes. Future research should identify temporal patterns of $\alpha 1m$ excretion and their associations with clinical outcomes, and determine specific thresholds of risk in HIV-infected and uninfected individuals. In HIV-infected persons, studies should specifically investigate the role of urine $\alpha 1m$ in detection of drug toxicity before the onset of overt kidney disease. Although our findings must be validated in subsequent cohorts, urine $\alpha 1m$ seems to be a promising candidate for inclusion in a future kidney biomarker panel.

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Original Article: Association of urine α 1-microglobulin with kidney function decline and mortality in HIV-infected women

Authors: Vasantha Jotwani^{1,2}, Rebecca Scherzer^{1,2}, Alison Abraham³, Michelle M Estrella⁴, Michael Bennett⁵, Mardge H Cohen⁶, Marek Nowicki⁷, Anjali Sharma⁸, Mary Young⁹, Phyllis C Tien¹, Joachim H Ix^{10, 11}, Mark J Sarnak¹², Chirag R Parikh^{13,14}, Michael G Shlipak^{1,2}

¹Department of Medicine, San Francisco VA Medical Center, San Francisco, USA

²Department of Epidemiology and Biostatistics, University of California, San Francisco, USA

³Department of Epidemiology, Johns Hopkins Bloomberg School of Public Health, Baltimore, USA

⁴Department of Medicine, Johns Hopkins School of Medicine, Baltimore, USA

⁵Division of Nephrology and Hypertension, Cincinnati Children's Hospital Medical Center, Cincinnati, USA

⁶Departments of Medicine, Stroger Hospital and Rush University, Chicago, USA

⁷Department of Medicine, University of Southern California, Los Angeles, USA

⁸Division of Infectious Diseases, Department of Medicine, SUNY Downstate Medical Center, Brooklyn, USA

⁹Division of Infectious Diseases and Travel Medicine, Department of Medicine, Georgetown University Medical Center, Washington, D.C., USA

¹⁰Division of Nephrology-Hypertension, Department of Medicine, University of California, San Diego, USA

¹¹Nephrology Section, Veterans Affairs San Diego Healthcare System, USA

¹²Division of Nephrology, Tufts Medical Center, Boston, USA

¹³Section of Nephrology, Department of Medicine, Yale University, New Haven, USA

¹⁴Program of Applied Translational Research, Yale University, New Haven, USA

Appendix

Supplemental Tables 1-2

Supplemental Figure Legend

Supplemental Figure 1

Supplemental Table 1a. Correlation matrix of urine biomarker-creatinine ratios in HIV-infected women.

	α1m	IL-18	KIM-1	NGAL	ACR
α1m	1	0.225**	0.301**	0.209**	0.421**
IL-18		1	0.162**	0.232**	0.148**
KIM-1			1	0.124*	0.177**
NGAL				1	0.248**
ACR					1

All biomarkers are standardized to urine creatinine.

Note: *p<0.001 and **p<0.0001 denote statistical significance of Spearman correlations.

Abbreviations: α1m, α1-microglobulin; ACR, albumin-to-creatinine ratio; IL-18, interleukin-18; KIM-1, kidney injury molecule-1; NGAL, neutrophil gelatinase-associated lipocalin.

Supplemental Table 1b. Correlation matrix of urine biomarker concentrations in HIV-infected women.

	α1m	IL-18	KIM-1	NGAL	ACR	UCR
α1m	1	0.416**	0.396**	0.279**	0.211**	0.409**
IL-18		1	0.495**	0.438**	-0.018	0.642**
KIM-1			1	0.343**	0.014	0.577**
NGAL				1	0.119*	0.410**
ACR					1	-0.181*
UCR						1

Biomarkers are not standardized to urine creatinine, with exception of ACR.

Note: *p<0.001 and **p<0.0001 denote statistical significance of Spearman correlations.

Abbreviations: α1m, α1-microglobulin; ACR, albumin-to-creatinine ratio; IL-18, interleukin-18; KIM-1, kidney injury molecule-1; NGAL, neutrophil gelatinase-associated lipocalin; UCR, urine creatinine.

Supplemental Table 2. Associations of urine α 1m concentration¹ with kidney function decline and mortality in HIV-infected women.

Range of α 1m	Incident CKD			10% decline in eGFR _{Cys}			All-cause mortality		
	C1 -	C2 0.6-1.4 mg/dl	C3 >1.4 mg/dl	C1 -	C2 0.6-1.4 mg/dl	C3 >1.4 mg/dl	C1 -	C2 0.6-1.4 mg/dl	C3 >1.4 mg/dl
No. at risk	343	259	217	358	274	271	358	274	271
No. events	48	54	75	9	18	34	29	30	69
		Risk Ratio (95% CI)	Risk Ratio (95% CI)		Risk Ratio (95% CI)	Risk Ratio (95% CI)		Hazard Ratio (95% CI)	Hazard Ratio (95% CI)
Demographic Adjusted²	Ref	1.59 (0.78, 3.21)	3.09 (1.80, 5.32)	Ref	2.62 (1.20, 5.74)	4.58 (2.23, 9.41)	Ref	1.32 (0.79, 2.20)	3.01 (1.93, 4.68)
Multivariate Adjusted³	---	1.28 (0.71, 2.30)	2.21 (1.33, 3.69)	---	2.23 (1.03, 4.86)	3.01 (1.42, 6.38)	---	1.16 (0.69, 1.95)	1.95 (1.24, 3.08)
Adjusted +ACR⁴	---	1.27 (0.72, 2.24)	2.02 (1.23, 3.31)	---	2.31 (1.08, 4.93)	2.80 (1.29, 6.08)	---	1.14 (0.68, 1.93)	1.92 (1.21, 3.04)
Adjusted +ACR, IL-18, KIM-1, NGAL⁵	---	1.16 (0.68, 1.96)	1.70 (1.09, 2.64)	---	2.21 (1.02, 4.78)	2.28 (1.03, 5.03)	---	1.06 (0.61, 1.84)	1.64 (0.94, 2.87)

¹Stratified by categories. Category 1 comprises all participants with undetectable urine α 1m. All analyses use Category 1 as reference category.

²Adjusted for age and race.

³Adjusted for age, race, baseline eGFR_{Cys}, hypertension, diabetes mellitus, hepatitis C virus infection, HIV viral load, CD4 lymphocyte count, antiretroviral therapy use, and serum albumin.

⁴Adjusted for all covariates listed above with addition of albumin-creatinine ratio.

⁵Adjusted for all covariates listed above with addition of albumin-creatinine ratio, interleukin-18, kidney injury molecule-1, neutrophil gelatinase-associated lipocalin, and liver fatty acid binding protein.

Abbreviations: ACR, albumin-creatinine ratio; α 1m, α 1-microglobulin concentration; C1, Category 1; C2, Category 2; C3, Category 3; CI, confidence interval; CKD, chronic kidney disease; eGFR_{Cys}, cystatin C-based estimated glomerular filtration rate; IL-18, interleukin-18; KIM-1, kidney injury molecule-1; NGAL, neutrophil gelatinase-associated lipocalin; Ref, reference category.

Supplemental Figure Legend

Supplemental Figure 1. Magnitudes of associations of urinary biomarker concentrations with annual eGFR_{Cys} change. All estimates derived from multivariate models comparing Category 3 with Category 1 for α1m, and highest with lowest tertile for ACR, IL-18, KIM-1, and NGAL. Multivariate models adjust for age, race, traditional kidney disease risk factors, HIV-related factors, and all displayed biomarkers. Results reported as eGFR_{Cys} change in ml/min/1.73m² per year (95% confidence interval).

Supplemental Figure 1.

