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# Kidney function and multiple hemostatic markers: cross sectional associations in the multi-ethnic study of atherosclerosis

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## Abstract

**Background:** Defined as estimated glomerular filtration rate (eGFR) < 60 ml/min/1.73 m<sup>2</sup>, chronic kidney disease (CKD) is strongly and independently associated with cardiovascular and overall mortality. We hypothesized that reduced kidney function would be characterized by abnormalities of hemostasis.

**Methods:** We tested cross-sectional associations between (eGFR) and multiple hemostatic markers among 6751 participants representing a broad spectrum of kidney function in the Multi-Ethnic Study of Atherosclerosis (MESA). Kidney function was measured using cystatin C (eGFR<sub>cys</sub>) or creatinine, using CKD Epidemiology Collaboration (eGFR<sub>cr</sub>). Hemostatic markers included soluble thrombomodulin (sTM), soluble tissue factor (sTF), D-Dimer, von Willebrand factor (vWF), factor VIII, plasmin-antiplasmin complex (PAP), tissue factor pathway inhibitor (TFPI), plasminogen activator inhibitor-1 (PAI-1), and fibrinogen. Associations were tested using multivariable linear regression with adjustment for demographics and comorbidities.

**Results:** In comparison to persons with eGFR<sub>cys</sub> >90 ml/min/1.73 m<sup>2</sup>, subjects with eGFR<sub>cys</sub> < 60 ml/min/1.73 m<sup>2</sup> had adjusted levels of sTM, sTF, D-Dimer, PAP, Factor VIII, TFPI, vWF and fibrinogen that were respectively 86%, 68%, 44%, 22%, 17%, 15%, 12% and 6% higher. Subjects with eGFR<sub>cys</sub> 60-90 ml/min/1.73 m<sup>2</sup> had adjusted levels that were respectively 16%, 14%, 12%, 6%, 6%, 6%, 11% and 4% higher (p < 0.05 for all). Percent differences were not significantly different when groups were categorized by eGFR<sub>cr</sub>.

**Conclusions:** Throughout a broad spectrum of kidney function, lower eGFR was associated with higher levels of hemostatic markers. Dysregulation of hemostasis may be a mechanism by which reduced kidney function promotes higher cardiovascular risk.

## Background

Defined as estimated glomerular filtration rate (eGFR) <60 ml/min/1.73 m<sup>2</sup>, chronic kidney disease (CKD) stage 3 to 5 [1] is an established risk factor for cardiovascular morbidity and mortality [2,3]. This association begins in the pre-clinical stages of kidney disease, when kidney function may be better estimated by cystatin C (eGFR<sub>cys</sub>) [4-6]. The mechanisms underlying the association of CKD with cardiovascular morbidity and mortality are inadequately understood. In multiple cohorts

with and without CKD, the associations of lower eGFR with higher cardiovascular risk persist despite adjustment for traditional risk factors and inflammation [7-10], prompting investigation into novel pathways between renal and cardiovascular disease (CVD).

Atherosclerosis represents a pernicious interplay between systems of inflammation and coagulation [11]. Multiple investigators have reported an association between creatinine-based CKD and elevated levels of inflammatory and thrombotic markers [12,13]. In studies from the Health, Aging and Body Composition Study and the Multi-Ethnic Study of Atherosclerosis (MESA), subjects with mildly reduced kidney function measured with cystatin C had only modest elevations in inflammatory

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markers [13-16]. Furthermore, the positive association of cystatin C with cardiovascular and overall mortality is only partially attenuated by adjustment for C-reactive protein (CRP), interleukin-6 (IL-6) or TNF- $\alpha$  [5,6]. However, whether or not dysregulation of hemostasis contributes to the association of CKD and CVD has not been well evaluated.

We designed these analyses to explore the associations of reduced kidney function, as measured by eGFR<sub>cys</sub> and CKD Epidemiology Collaboration (eGFR<sub>cr</sub>), with hemostatic factors in an ethnically diverse cohort free of clinical cardiovascular disease.

## Methods

### Subjects

We conducted a cross-sectional analysis of 6751 participants in the Multi Ethnic Study of Atherosclerosis (MESA) at the baseline visit (July 2000 to July 2002). Soluble thrombomodulin (sTM), soluble tissue factor (sTF) and tissue factor pathway inhibitor (TFPI) were measured in a random sample of 1000 MESA participants; the other markers were measured in nearly the whole cohort. MESA is a cohort study sponsored by the National Heart, Lung, and Blood Institute aimed to improve understanding of subclinical cardiovascular disease and its progression in a multiethnic cohort. Details on recruitment and design have been published [17]. Briefly, MESA recruited 6,814 men and women between the ages of 45 and 84 years, who were free of cardiovascular disease and self-identified as white, African American, Hispanic, or Chinese-American. Subjects were recruited from Baltimore City and Baltimore County, Maryland; Chicago, Illinois; Forsyth County, North Carolina; Los Angeles County, California; northern Manhattan and the Bronx, New York; and St. Paul, Minnesota, between July 2000 and August 2002. Individuals were excluded if they had physician-diagnosed heart attack, angina, heart failure, stroke or transient ischemic attack, or atrial fibrillation; if they had undergone coronary artery bypass grafting, angioplasty, valve replacement, or pacemaker insertion; or if they weighed > 300 pounds (>136 kg). The institutional review boards at all participating centers approved the study, and all participants gave informed consent.

### Primary predictors

Cystatin C was measured in plasma using a BNII nephelometer on plasma specimens (Siemens Latex N-Latex cystatin C assay). Estimated glomerular filtration rate by cystatin C (eGFR<sub>cys</sub>) was calculated using the formula:  $76.7 \times \text{cys } C^{-1.19}$ . This formula was developed from pooled data from several cohorts with eGFR measured by iothalamate [18]. Serum creatinine was measured using colorimetry with a Johnson & Johnson

Vitros 950 analyzer (Johnson & Johnson Clinical Diagnostics Inc., Rochester, NY). Creatinine levels were calibrated to the Cleveland Clinic standard ( $0.9954 \times \text{Cr} + 0.0208$ ) [19]. The eGFR<sub>cr</sub> equation was used to calculate creatinine - based eGFR (eGFR<sub>cr</sub>) [20].

### Primary outcomes

Soluble thrombomodulin (sTM) was measured by enzyme immunoassay using a monoclonal antibody to TM as the capture antibody (Asserachrom Thrombomodulin, Diagnostica Stago; Asnières-sur-Seine, France, CV 12%). Soluble tissue factor (sTF) was measured by an enzyme-linked immunoassay that employs an anti-TF monoclonal capture antibody (Imubind Tissue Factor ELISA Kit, American Diagnostica, Inc.; Stamford, CT, CV 14.6%). Total tissue factor pathway inhibitor (TFPI) was measured by enzyme-linked sandwich ELISA using a polyclonal anti-TFPI antibody as the capture antibody (Imubind Total TFPI ELISA Kit, American Diagnostica, Inc.; Stamford, CT). Intra-assay and inter-assay analytical CVs range from 6.2-7.1% and 5.5-7.3%, respectively. Fibrin fragment D-dimer was measured using an immuno-turbidimetric assay (Liatest D-DI; Diagnostica Stago, Parsippany, NJ) on the Sta-R analyzer (Diagnostica Stago, Parsippany, NJ, CV 8%). Plasmin-antiplasmin complex (PAP) was measured in an assay that detects only plasmin in complex with  $\alpha_2$ -antiplasmin, and not free plasmin or  $\alpha_2$ -antiplasmin, CV 1.7%. von Willebrand factor was measured by an immunoturbidimetric assay on the Sta-R analyzer (Diagnostica Stago, Parsippany, NJ). The intra-assay and inter-assay analytical CV's are 3.7% and 4.5%, respectively. Factor VIII coagulant activity was determined by measuring the clotting time of a sample in factor VIII deficient plasma in the presence of activator (STA-Deficient VIII; Diagnostica Stago, Parsippany, NJ). Fibrinogen was measured using a BNII nephelometer (N Antiserum to Human Fibrinogen; Siemens) with intra-assay and inter-assay analytical CV's of 2.7% and 2.6%, respectively. Plasminogen activator inhibitor-1 (PAI-1) was measured by a two-site ELISA [21].

### Variables of interest

Blood pressure measurements were obtained by using an automated blood pressure device (DINAMAP PRO 100 monitor; General Electric Healthcare) following JNC guidelines [22]. Hypertension was defined as self-report of physician diagnosis, use of anti-hypertensive agents, or SBP  $\geq 140$ , or DBP  $\geq 90$ . Diabetes was defined by fasting glucose  $\geq 126$  or the use of hypoglycemic medication or insulin. Tobacco use was defined by current smoking. Height and weight were measured with participants wearing light clothing and no shoes. Body mass index was calculated as weight (kg)/height (m)<sup>2</sup>.

Fasting blood was collected and stored at -70 F (21.11 C) until analyzed for total and high density lipoprotein cholesterol, triglycerides and glucose. Low density lipoprotein cholesterol was calculated by using the Friedewald equation.

### Statistical analysis

We compared sociodemographic and anthropometric factors as well as comorbidities among subjects with eGFR<sub>cys</sub> > 90, 60-90 and < 60 ml/min/1.73 m<sup>2</sup> using ANOVA, Kruskal-Wallis test, or chi-square tests as appropriate. Using multivariable linear regression, we studied the associations between continuous eGFR and each of the seven biomarkers, using both eGFR<sub>cys</sub> and eGFR<sub>cr</sub>. Each biomarker was natural log transformed, and beta coefficients were transformed into percent differences using  $(e^{\beta}-1)*100$  to allow comparison across outcomes. In additional analyses we defined eGFR groups as >90, 60-90 and <60 ml/min/1.73 m<sup>2</sup>, and we compared the percent difference in each biomarker with eGFR > 90 as the reference. The fully adjusted models included age, race/ethnicity (white, black, Hispanic, Chinese), gender, annual income, study site, current smoking, current alcohol use, BMI, DM, HTN, statin use, ACE-I, LDL cholesterol, HDL cholesterol, triglycerides, fasting glucose, and ln(albumin/creatinine). All models assumptions for linear regression, including linearity, homoscedasticity and normality, were checked. Analyses were performed in SAS V 9.1.3 (SAS Institute, Cary, NC), and  $p < 0.05$  was considered as the threshold for statistical significance.

## Results

### Participants' characteristics

Among the 6751 participants, the mean age was 62 years; 53% were women; 38% were white, 27% black, 22% Hispanic, and 12% Chinese-American. The prevalences of diabetes and hypertension were 14% and 45%, respectively. Average eGFR<sub>cys</sub> was 92 ml/min/1.73 m<sup>2</sup> and average eGFR by eGFR<sub>cr</sub> was 79 ml/min/1.73 m<sup>2</sup>. Compared to those without CKD, subjects with CKD with eGFR<sub>cys</sub> <60 ml/min/1.73 m<sup>2</sup> were older, had a higher prevalence of diabetes and hypertension, higher rate of ACE-I and statin use, higher fasting glucose, triglycerides and urinary albumin to creatinine ratio, lower HDL and LDL, and lower rate of alcohol use. (Table 1)

### Variation by race/ethnicity and correlations among biomarkers

We analyzed levels of hemostatic markers in different groups of race/ethnicity. sTM showed significant heterogeneity by race/ethnicity: Caucasians and Hispanics had significantly higher sTM than African-Americans and Chinese. Levels of vWF, and fVIII were significantly

higher in African-Americans, and fibrinogen was higher in African Americans and Hispanics. (Table 2) As expected, many of the hemostatic biomarkers were highly correlated. vWF and factor VIII were positively correlated ( $r = 0.66$ ); PAP and PAI-1 were inversely correlated, ( $r = -0.48$ ). D-Dimer had moderate correlations ( $r = 0.23-0.34$ ) with VWF, factor VIII, PAP and fibrinogen. Fibrinogen had moderate correlations ( $r = 0.23-0.38$ ) with vWF, factor VIII, PAP, and TFPI ( $p < 0.05$  for all). No other strong correlations were observed. (Table 3)

### Associations between eGFR and biomarkers

We analyzed the associations between continuous measures of eGFR measured by both eGFR<sub>cys</sub> and eGFR<sub>cr</sub> equations and the seven biomarkers. In models adjusted for sociodemographics and comorbidities, sTM and sTF had strong correlations with eGFR by either method. For every 10 ml/min/1.73 m<sup>2</sup> lower eGFR<sub>cys</sub>, sTM was 7% higher ( $p < 0.001$ ) and sTF was 6% higher ( $p < 0.001$ ). D-Dimer, vWF, Factor VIII, PAP and TFPI had moderate associations with eGFR using either method. D-Dimer was 5% higher, vWF was 3% higher, and the other three were 2% higher for every 10 ml/min/1.73 m<sup>2</sup> lower eGFR<sub>cys</sub> ( $p < 0.001$  for all). PAI-1 was associated with eGFR<sub>cys</sub> but not eGFR<sub>cr</sub>; fibrinogen had a weak but statistically significant association with both GFR estimates. (Table 4)

In addition, we calculated percent differences for each marker between subjects grouped by eGFR, using eGFR > 90 ml/min/1.73 m<sup>2</sup> as the reference. The group with eGFR<sub>cys</sub> <60 ml/min/1.73 m<sup>2</sup> had an 86% higher sTM level, 68% higher sTF, 44% higher D-Dimer, 12% higher vWF, 17% higher Factor VIII, 22% higher PAP, 15% higher TFPI, 6.5% higher PAI-1, and 6.3% higher fibrinogen. Participants with eGFR 60-90 ml/min/1.73 m<sup>2</sup> had significantly higher levels of all biomarkers ranging from 16% for sTM to 3.8% for fibrinogen. (Figure 1)

## Discussion

In a large, multiethnic cohort free of cardiovascular disease and with a spectrum of kidney function ranging from normal to moderate CKD, even small decrements in eGFR were associated with significantly higher levels of sTM, sTF, D-Dimer, vWF, factor VIII, PAP and TFPI. Our results suggest that dysregulation of hemostasis could play an important pathologic role in CKD. In particular, these associations may help to explain the consistently strong and independent associations of cystatin C with cardiovascular and overall mortality, despite adjustment for known risk factors including CRP [5], IL-6 and TNF- $\alpha$  [6].

Soluble thrombomodulin and soluble tissue factor had the largest associations with eGFR<sub>cys</sub>. sTM is a receptor

**Table 1 Demographic and clinical characteristics of MESA participants by groups of eGFRcys, 2000-2002**

	eGFRcys <60 (n = 376)	eGFRcys 60-90 (n = 2746)	eGFRcys >90 (n = 3629)	P Value*
Age, years†	71 ± 9	66 ± 10	59 ± 9	<0.001
Female, n(%)	185 (49)	1131 (48)	2066 (57)	<0.001
Race/Ethnicity, n(%)				<0.001
Caucasian	165 (44)	1151 (42)	1286 (35)	
African-American	103 (27)	715 (26)	1048 (29)	
Hispanic	72 (19)	617 (23)	795 (22)	
Chinese-American	36 (10)	263 (10)	500 (14)	
Current Smoker, n(%)	50 (13)	386 (14)	440 (12)	0.08
Current Alcohol Use, n(%)	169 (45)	1457 (53)	2087 (58)	<0.001
BMI, kg/m <sup>2</sup>	30 ± 6	29 ± 6	28 ± 5	
Prevalent Diabetes, n(%)	94 (25)	401 (15)	464 (13)	<0.001
Prevalent Hypertension, n(%)	272 (72)	1451 (53)	1310 (36)	<0.001
ACE Inhibitor Use, n(%)	113 (30)	501 (18)	410 (11)	<0.001
Statin Use, n(%)	77 (21)	465 (17)	459 (13)	<0.001
Fasting Plasma Glucose, mg/dl	111 ± 34	105 ± 29	104 ± 32	<0.001
Triglycerides, mg/dl‡	131 (93, 180)	126 (88, 179)	107 (77, 157)	<0.001
HDL Cholesterol, mg/dl	47 ± 14	49 ± 14	53 ± 15	<0.001
LDL Cholesterol, mg/dl	114 ± 35	116 ± 6	118 ± 31	0.01
Urinary Creatinine, mg/dl‡	96 (65, 142)	111 (66, 163)	106 (55, 162)	0.001
Urine Albumin/Creatinine, mg/g	10 (5, 47)	6 (3, 12)	5 (3, 10)	<0.001
Serum Creatinine, mg/dl	1.4 ± 0.8	1.0 ± 0.2	0.9 ± 0.2	<0.001
eGFR Creatinine, ml/min/1.73 m <sup>2</sup>	52 ± 17	73 ± 14	87 ± 14	<0.001
eGFR Cystatin, ml/min/1.73 m <sup>2</sup>	50 ± 10	78 ± 8	108 ± 15	<0.001

\*P-values obtained using ANOVA.

†Unless otherwise noted, Mean (SD).

‡Median (Quartile 1, Quartile 3).

for thrombin found on vascular and lymphatic endothelial cells. Injured endothelial cells release sTM into the circulation, where it binds thrombin, inhibiting it from forming fibrin and activating platelets. sTM has additional anti-thrombotic activity as a cofactor for the thrombin-catalyzed activation of protein C [23]. Elevated sTM is associated with lower eGFR in CKD and end stage renal disease ESRD [24,25], and our results extend these findings to a broader spectrum of eGFR. The association of sTM and cardiovascular disease varies by patient

population: in healthy individuals, lower sTM predicts coronary heart disease (CHD), but in patients with established CHD, higher sTM levels are associated with recurrence of CHD and with higher mortality [26]. Whether higher levels of sTM are associated with CVD or mortality in our cohort would require further investigation.

Tissue factor (TF) initiates the extrinsic coagulation cascade. Smooth muscle cells, adventitial fibroblasts, and pericytes constitutively express TF, but endothelial cells express TF only after injury [27]. TF has a prominent

**Table 2 Levels of hemostatic markers by racial/ethnic group\***

	All	Caucasian	African-American	Chinese	Hispanic	P Value**
Soluble Thrombomodulin, ng/ml	35 (16, 45)	36 (27, 47)	32 (23, 42)	31 (23, 40)	37 (27, 46)	<0.001
Soluble Tissue Factor, pg/ml	106 (68, 156)	107 (66, 149)	106 (72, 175)	97 (65, 169)	108 (68, 155)	0.53
D-dimer, ug/ml	0.20 (0.13, 0.37)	0.20 (0.13, 0.35)	0.25 (0.15, 0.44)	0.18 (0.10, 0.30)	0.23 (0.13, 0.41)	<0.001
von Willebrand Factor, %	130 (99, 169)	124 (95, 166)	144 (113, 199)	130 (96, 168)	127 (96, 166)	<0.001
Factor VIII, %	93 (73, 120)	90 (70, 115)	100 (77, 131)	93 (73, 115)	91 (74, 117)	<0.001
Plasmin Antiplasmin, nM	4.4 (3.4, 5.6)	4.4 (3.4, 5.6)	4.8 (3.8, 6.2)	3.9 (3.1, 4.9)	4.3 (3.4, 5.4)	<0.001
Tissue Factor Pathway Inhibitor, ng/ml	48 (38, 57)	49 (38, 57)	48 (40, 58)	40 (33, 50)	49 (40, 58)	<0.001
Plasminogen Activator Inhibitor-1, ng/ml	19 (10, 36)	19 (9, 34)	17 (8, 33)	24 (12, 39)	21 (11, 40)	0.02
Fibrinogen, mg/dl	338 (295, 389)	328 (286, 373)	353 (304, 406)	323 (287, 366)	352 (308, 401)	<0.001

\*Levels displayed as median (Quartile 1, Quartile 3).

\*\*By Kruskal-Wallis test for difference among racial/ethnic groups.

**Table 3 Spearman correlations among hemostatic markers**

	sTM	STF	D-dimer	vWF	Factor VIII	PAP	TFPI	PAI-1	Fib
sTM	-								
STF	0.15†	-							
D-dimer	0.05	0.04	-						
vWF	0.15†	0.09**	0.25†	-					
Factor VIII	0.10**	0.02	0.23†	0.66†	-				
PAP	-0.03	0.15†	0.34†	0.25†	0.21†	-			
TFPI	0.21†	0.08*	0.14†	0.16†	0.16†	0.10**	-		
PAI-1	0.08*	-0.08*	0.02	0.05	0.07*	-0.48†	0.13†	-	
Fib	0.03	0.07*	0.29†	0.26†	0.24†	0.38†	0.23†	0.08*	-

\*0.01-<0.05.

\*\*0.001-<0.01.

†<0.001.

role in atherosclerotic plaque rupture and acute coronary syndrome (ACS). TF is concentrated in the core of the atherosclerotic plaque; when the plaque ruptures, TF comes into contact with blood, binds factor VII and initiates the extrinsic coagulation cascade [28]. Higher levels of sTF were reported previously in CKD as compared to healthy subjects [29]. Epidemiological studies of sTF and CVD occurrence in healthy individuals are lacking, but in patients with ACS, high levels of sTF on admission are associated with poor outcome [30]. To our knowledge this paper is the first to report an association between higher sTF and lower eGFR in patients

**Table 4 Cross-sectional associations of eGFR measures with hemostatic markers in MESA, 2000-2002: percent difference in hemostatic marker per 10 ml/min/1.73 m<sup>2</sup> decrement in eGFR\***

	eGFRcys, % (95% CI)	eGFRcr, % (95% CI)
Soluble Thrombomodulin	7.4 (5.9, 8.8)	8.2 (6.3, 10.1)
Soluble Tissue Factor	6.1 (3.6, 8.5)	7.6 (4.4, 10.6)
D-dimer	4.6 (3.5, 5.7)	4.7 (3.3, 6.1)
Von Willebrand Factor	2.8 (1.4, 4.2)	2.8 (0.93, 4.6)
Factor VIII	2.2 (2.7, 1.8)	1.8 (2.5, 1.2)
Plasmin-Antiplasmin Complex	2.2 (1.7, 2.7)	3.2 (2.6, 3.8)
Tissue Factor Pathway Inhibitor	2.1 (1.2, 3.1)	1.4 (0.2, 2.6)‡
Plasminogen Activator Inhibitor-1	4.7 (2.0, 7.4)	1.3 (-2.3, 5.1)§
Fibrinogen	1.0 (0.74, 1.2)	1.0 (0.64, 1.3)

\*Note: Models adjusted for age, race, gender, SES (income), site, current smoking, current alcohol, BMI, prevalent DM, prevalent HTN, statin use, ACE-I, LDL, HDL, TG, fasting glucose, and ln(albumin/creatinine). All p values are <0.001 unless otherwise noted.

‡p < 0.05.

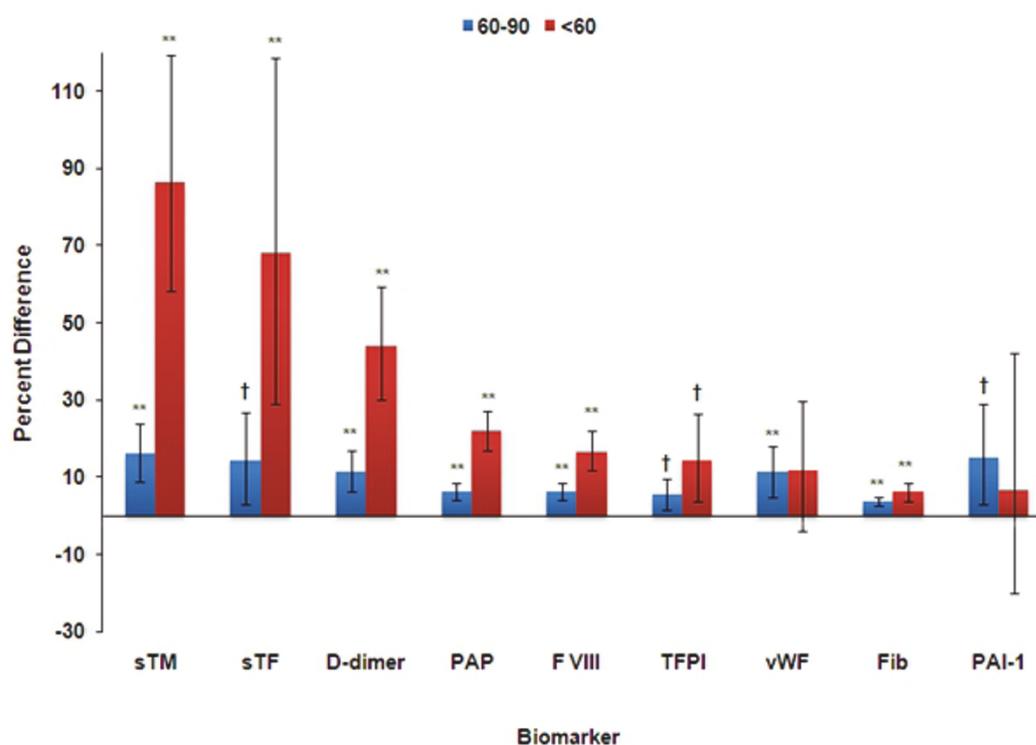
§Non-significant.

without CKD. Whether high levels of sTF contribute to the high cardiovascular mortality of persons with kidney disease is unknown.

D-Dimer, vWF, Factor VIII, and PAP showed moderate associations with lower eGFRcys. Elevated levels of D-Dimer are associated with lower eGFR [12] and CVD [31] across the spectrum of eGFR. vWF is increased in subjects with CKD [32], and it is considered to be a marker for endothelial dysfunction [33] and CVD risk [34,35]. Factor VIII is increased in subjects with CKD [12,32]. In the Atherosclerosis Risk in Communities (ARIC) Study, factor VIII was associated with progression of CKD and with fatal MI [36]. In MESA, Factor VIII was associated with overall but not cardiovascular mortality [37]. Plasmin, an enzyme that increases fibrinolysis, is inactivated when complexed with a2-antiplasmin to form PAP [11]. Subjects with CKD have higher levels of PAP than subjects without CKD [12]. In a cohort with CAD, PAP was associated with higher CVD risk [38]; in the Cardiovascular Health Study (CHS), PAP was associated with incident myocardial infarction [39].

Several of the hemostatic markers evaluated in this study are strong risk factors for venous thromboembolic disease, including D-dimer, factor VIII, and vWF [40]. Recent epidemiological studies have linked CKD with risk of venous thrombosis [41,42]. Thus, our findings may be applicable to both arterial and venous thrombosis, and these associations may extend into higher levels of kidney function than previously appreciated.

There are several possible mechanisms to explain the association of lower eGFR and higher levels of hemostatic factors. A direct effect of decreased renal clearance may explain an increase in levels of smaller molecular weight hemostatic markers such as sTF and sTM, as these may be filtered at the glomerulus. Fragments of sTM ranging from 28 to 108 kDa can be detected in blood and urine [43]. Soluble tissue factor has a MW of 29 kDa. Urinary tissue factor, with MW of 34 kDa, is elevated in glomerulonephritis and various malignancies, but it is thought to originate from the kidneys, rather than being blood borne and filtered at the glomerulus [44]. Whether sTF and sTM are filtered at the glomerulus would have to be tested in animal studies. Given the higher molecular weight of D-Dimer, vWF, factor VIII and PAP, it seems unlikely that these markers are elevated as a direct result of decreased renal clearance. It is possible that elevations in these markers relate to processes initiated by smaller molecules. Conversely, kidney dysfunction may generate a thrombotic milieu indirectly through electrolyte or acid-base abnormalities, which may alter activities of enzymes involved in coagulation. Inflammation associated with lower eGFR may cause elevations in markers that are acute phase reactants. Alternatively, microvascular disease may promote the development of CKD, and in



**Figure 1 Percent Difference in Biomarker in Groups of eGFR cystatin in MESA\*** \* Models adjusted for age, race, gender, SES (income), site, current smoking, current alcohol, BMI, prevalent DM, prevalent HTN, statin use, ACE-I, LDL, HDL, TG, fasting glucose, and ln(albumin/creatinine); eGFR cystatin > 90 is the reference group \*\* p<0.001 † p<0.05.

parallel, may cause elevations in these hemostatic biomarkers; in this scenario, the markers would be neither cause nor effect of lower eGFR.

A strength of this study is the unique setting, as MESA is a large, well-characterized cohort including four major race/ethnic groups. The use of cystatin C to measure eGFR increases our ability to detect associations of biomarkers across a broader spectrum of kidney function. However, there are important limitations to our study. As is typical for most large epidemiological studies, we did not use a gold standard measure of eGFR, such as iothalamate or iohexol clearance. Due to the cross-sectional design, we cannot ascertain the direction of association; it is equally possible that increased levels of thrombotic markers causes decreased eGFR, or that decreased eGFR causes the elevated markers. Finally, residual confounding may exist in our analysis, which may account for some of our findings, but would be less likely to account for the strong associations of eGFRcys with sTM and sTF.

## Conclusions

In summary, lower eGFR was significantly associated with higher levels of sTM, sTF, D-Dimer, PAP, Factor VIII, TFPI, vWF and fibrinogen in subjects with and without

CKD. Increased levels of smaller molecular weight molecules such as sTM and sTF may be a direct result of decreased renal clearance. Elevations of larger molecules may represent increased fibrinolytic activity and higher clot burden of a thrombotic milieu induced by declining kidney function. Alternatively, the atherosclerotic process may differ fundamentally in persons with lower eGFR. Further investigations may lead to a better understanding of the mechanisms underlying these associations, as well as the role of these associations in predicting cardiovascular disease in subjects with elevated cystatin C.

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#### Authors' contributions

RD drafted and revised the paper, and gave input into statistical analysis. MC helped to draft the manuscript and provided advice on hemostatic factors. AF helped draft the manuscript and provided advice on interpreting the data. LF helped draft the manuscript and provided advice on interpreting the data. WP helped draft the manuscript and provided advice on interpreting the data. CP helped draft the manuscript and provided advice on interpreting the data. CW performed statistical analysis. MS sponsored and conceived of the project, helped revise the paper, and gave input into statistical analysis. All authors read and approved the final manuscript.

#### Competing interests

The authors declare that they have no competing interests.

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