# Clinical, laboratory, and genetic risk factors for thrombosis in sickle cell disease

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#### **Key Points**

- Traditional (kidney disease) and SCD-specific (SS/Sβ0-thal, HU, low Hb F%, high WBC count, and Hb) risk factors are associated with thrombosis.
- Stroke recurrence was 24% despite being at transfusion goal and VTE recurrence was lower with direct oral anticoagulant vs other therapies.

Sickle cell disease (SCD) patients are at a four- to 100-fold increased risk for thrombosis compared with the general population, although the mechanisms and risk factors are not clear. We investigated the incidence and predictors for thrombosis in a retrospective, longitudinal cohort of 1193 pediatric and adult SCD patients treated at our institution between January 2008 and December 2017. SCD diagnosis and thrombotic complications were identified using International Classification of Diseases coding and verified through medical chart review. Clinical and laboratory data were extracted from the medical records. With a median follow-up of 6.4 years, 208 (17.4%) SCD patients experienced 352 thrombotic events (64 strokes, 288 venous thromboembolisms [VTE]). Risk factors for stroke included older age and HbSS/Sβ<sup>0</sup>-genotype and a lower hemoglobin (Hb) F% in the subset of HbSS/Sβ<sup>0</sup> -genotype patients (P < .05). VTE risk was independently associated with lower estimated glomerular filtration rate, hydroxyurea (HU) use, HbSS/Sβ<sup>0</sup> genotype, and higher white blood cell (WBC) counts and Hb ( $P \le .03$ ). Two thrombomodulin gene variants previously associated with thrombosis in the general African American population, THBD rs2567617 (minor allele frequency [MAF] 0.25; odds ratio [OR], 1.5; P = .049) and THBD rs1998081 (MAF, 0.24; OR, 1.5; P = .059), were associated with thrombosis in this cohort. In summary, thrombotic complications are common, and several traditional and SCD-specific risk factors are associated with thrombotic risk. Future studies integrating clinical, laboratory, and genetic risk factors may improve our understanding of thrombosis and guide intervention practices in SCD.

## Introduction

Sickle cell disease (SCD) is an inherited red blood cell disorder caused by homozygous or compound heterozygous inheritance of mutations in the  $\beta$ -globin gene that affects 1 in 365 African Americans.<sup>1</sup> Vasculopathy is a hallmark of SCD that contributes to the protean acute and chronic complications.<sup>2</sup> Thrombosis has been increasingly recognized as a complication of SCD. The hypercoagulable state may be due to increased exposure of phosphatidylserine on the outer surface of red blood cell membranes, increased tissue factor expression on endothelial cells and in circulation, depletion of protein C and S, endothelial dysfunction, and chronic platelet activation.<sup>3</sup>

Up to 30% of patients will have a cumulative lifetime risk of a clinically overt cerebrovascular accident (CVA), which is the most common arterial thrombotic event in SCD.<sup>4</sup> Risk factors for CVA include the hemoglobin (Hb) SS genotype, an elevated white blood cell count, and low Hb concentrations and

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Hb F%.<sup>4</sup> Chronic red blood cell transfusion therapy reduces the risk for subsequent CVA, but 18% of SCD children will have another overt CVA despite achieving transfusion goals.<sup>5</sup> Rates of CVA recurrence in SCD adults treated with transfusion therapy are less clear.

The reported cumulative risk for venous thromboembolism (VTE) in SCD ranges from 2.9% in children up to 25.0% in adults.<sup>6-9</sup> Although clinical and laboratory risk factors for VTE are not clear, its clinical importance in SCD is highlighted by a 2.3- to 2.9-fold increased risk of death in those with vs without a VTE event.<sup>7,9</sup> The genetic basis for VTE risk in SCD has not been previously reported, but variants implicated in other non-SCD African American cohorts may be relevant.<sup>10</sup> It is also important to understand how the type and duration of anticoagulant therapy affect the rate of VTE recurrence, which is also unclear.

In a cohort of 1193 SCD patients, we conducted a retrospective, longitudinal study to identify the (1) incidence, (2) clinical and laboratory risk factors, (3) association of genetic variants implicated in African American thrombosis risk, and (4) rates of recurrence based on type and duration of therapy for thrombotic events over a 10-year period.

# **Methods**

We analyzed SCD patients receiving medical care at the University of Illinois at Chicago (UIC) between January 2008 and December 2017. The protocol was approved by the UIC Institutional Review Board prior to extracting the data from the Cerner PowerChart electronic health records (EHRs).

Through a semiautomated algorithm using the UIC hospital enterprise data warehouse, the SCD cohort was identified through encounters with relevant 9th and 10th Clinical Modifications editions of the International Classification of Disease (ICD-9-CM and ICD-10-CM) diagnostic codes for SCD, Hb fractionations, and having either a minimum of 2 outpatient or 1 inpatient encounter between 1 January 2008 and 31 December 2017 (supplemental Table 1).<sup>11</sup> SCD genotype was determined by assessing the categorization of ICD coding and Hb fractionation results with manual confirmation by EHR review for those patients that could not be automatically classified. SCD patients were categorized into 4 distinct subgroups: Hb SS disease or Hb S $\beta^0$ -thalassemia, Hb S $\beta^+$  -thalassemia, Hb SC, or other sickle gene variant.

Baseline demographic and risk factors were gueried from our EHR database, including sex, race/ethnicity, past medical history, orthopedic and central venous catheterization procedure data, laboratory values, and medications prescribed. Baseline laboratory, blood pressure, and anthropometric results were determined for each patient using values from outpatient visits. The estimated glomerular filtration rate (eGFR) was calculated using the Chronic Kidney Disease-Epidemiology formula.<sup>12</sup> Hydroxyurea use was defined as a logical variable (ever vs never) based on if the patient was prescribed the medication during the study period. Pregnancy data were initially queried using relevant ICD-9/10-CM for pregnancy outcomes (including deliveries and abortions), and then manually confirmed by EHR review to determine number of pregnancy episodes. Orthopedic procedures and central venous catheterizations were queried using pertinent 9th and 10th Procedure Coding system editions of the International Classification of Disease (ICD-9-PCS and ICD-10-PCS). Thromboembolic event diagnoses were queried with pertinent ICD-9/10-CM coding and manually confirmed upon review of diagnostic imaging and/or medical records. Each event was characterized to determine the type of therapy initiated and the duration of anticoagulation. A recurrent thrombotic event was defined as having a second event, confirmed by manual review of the imaging and/or records, during the study time period. The ICD-9 and ICD-10 codes used for identifying pregnancy, procedures, and thrombotic events are provided in the supplemental Table 1.

## **Genetic risk variants**

Genotyping was carried out using Affymetrix Axiom genome-wide Pan-African GeneChip array at the Core Genomics Facility at UIC in DNA isolated from peripheral blood mononuclear cells in 327 SCD patients, as previously described.<sup>13</sup> All samples had genotype call rate >95%. Single-nucleotide polymorphisms (SNPs) deviating from Hardy-Weinberg equilibrium (P < .0001) or with a minor allele frequency (MAF) <0.01 were removed.

Candidate SNPs associated with thrombotic risk in the general African American population were tested.<sup>10</sup> Allele dosages were associated with thrombotic risk using an additive model, adjusting for the following covariates: age, sex, SCD genotype, hydroxyurea use, and population structure.

## **Statistical analysis**

For univariate analyses, the associations of linear and categorical variables with any thrombotic event, CVA, or VTE were assessed using the Kruskal-Wallis test and Pearson's  $\chi^2$  test, respectively. Variables with P < .002 were considered statistically significant after the Bonferroni correction. An initial multivariate Cox proportional hazards model was fit by including covariates with  $P \le .05$ from the univariate analyses. Patients who did not experience a thrombosis were censored at the date of last known contact. The final multivariate model was determined by stepwise variable selection, allowing for both backward and forward selection by sequentially selecting models that minimize the Akaike information criterion, adjusting for the following covariates: age, sex, sickle cell genotype, and hydroxyurea use.<sup>14</sup> The analyses were conducted using time updated variables for body mass index, systolic blood pressure, white blood cell count, Hb concentration, reticulocyte count, aspartate aminotransferase (AST), and eGFR.<sup>15,16</sup> The eGFR, white blood cell count, AST, and baseline Hb F% were logtransformed to achieve a normal distribution. We conducted the following subset analyses for the multivariate models: (1) risk factors for any thrombosis, CVA, and VTE in patients with the Hb  $SS/S\beta^{0}$ -thalassemia genotype; (2) risk factors for venous thrombosis in non-catheter-related VTE events. Patients who received an unknown therapy for CVA (n = 1) or VTE (n = 3) events were excluded from the analyses for recurrence based on initial therapy received. Data preprocessing and analysis were accomplished using custom-written scripts in R<sup>17</sup> using premade analytical packages, including tidyverse, lubridate, broom, survival, survminer, coin, and tidyforest.<sup>18-24</sup> Median values with interquartile range (IQR) are provided.

# **Results**

An initial ICD query identified 1350 potential SCD subjects. We were able to confirm the diagnosis of SCD in 1193 patients by review of Hb fractionations, transfusion history, and medical

Table 1. Baseline clinical and laboratory variables in 1193 SCD patients

Variable	Value
Age at first encounter, median (IQR), y	24 (17-35)
Sex, male/female, n (%)	515 (43.2)/678 (56.8)
Race, African American, n (%)	1123 (94.1)
Sickle cell genotype, n (%)	
HbSS/SB <sup>0</sup> -thalassemia	831 (69.7)
HbSC	290 (24.3)
HbSB <sup>+</sup> -thalassemia	69 (5.8)
Other	3 (0.3)
Hydroxyurea by sickle cell genotype, n/N (%)	
Total	659/1193 (55.2)
HbSS/SB <sup>0</sup> - thalassemia	560/831 (67.4)
HbSC	77/290 (26.6)
HbSB <sup>+</sup> -thalassemia	22/69 (31.9)
Other	0/3 (0)

records. The median age of the cohort was 24 years (IQR, 17-35 years); 57% were female, 55% were treated with hydroxyurea, and 70% were Hb SS/S $\beta^0$ -thalassemia genotype (Table 1). Based on ICD procedure coding, there were 83 orthopedic procedures in 52 patients, 1509 central venous catheterizations in 394 patients, and 209 pregnancies in 138 patients between January 2008 and December 2017.

#### All thrombotic episodes

With a median follow-up of 6.4 years (IQR, 2.7-9.8 years), 208 (17.4%) SCD patients experienced 352 thrombotic events (CVA: n = 64, VTE: n = 288). The median age at initial thrombosis was 30 years (IQR, 24-43 years) and median time to initial thrombosis from the first encounter was 3.2 years (IQR, 0.8-5.9 years). The distribution for the type of thrombotic event is provided in Figure 1. Differences in variables by thrombosis status are provided in Table 2. Independent risk factors for a thrombotic event on Cox regression analysis included a lower eGFR (P < .001), hydroxyurea use (P = .02), and Hb SS/S $\beta^0$ -thalassemia genotype (P = .05),



Figure 1. Types of thrombotic events in 1193 SCD patients between January 2008 and December 2017. DVT, deep vein thrombosis; SVC, superior vena cava.

Table 2. Clinical and laboratory values stratified by thrombotic events in patients with SCD

	n	No thrombosis	n	Thrombosis	P
Age at first encounter, y	985	23 (15-35)	208	27 (21-39)	<.0001
Male/female, %	985	45/55	208	36/64	.028
African American, n (%)	985	923 (94)	208	200 (96)	.2
Sickle cell genotype, %	985		208		<.0001
HbSS/SB <sup>0</sup> thalassemia		66.7	83.7		
HbSC, SB <sup>+</sup> -thalassemia, or other		33.3		16.3	
Hydroxyurea use, %	985	496 (50)	208	163 (78)	<.0001
Tobacco use, %	985	266 (27)	208	87 (42%)	<.0001
Body mass index, kg/m <sup>2</sup>	985	22 (18-26)	208	23 (20-27)	.0031
Systolic blood pressure, mm Hg	985	115 (104-127)	208	122 (110-132)	<.0001
White blood cell count, $ imes 10^3/\mu L$	985	9.7 (7.4-12.2)	208	10.1 (7.6-12.5)	.13
Platelet count, $\times 10^3/\mu L$	985	406.0 (287.0-525.0)	208	402.5 (312.5-532.2)	.4
Hb, g/dL	985	9.6 (8.3-11.1)	208	9.0 (8.2-10.2)	.002
Hb F, %	985	4.9 (2.1-11.2)	208	4.7 (2.0-9.2)	.12
$\begin{array}{c} \text{Reticulocyte count,} \\ \times 10^3 / \mu L \end{array}$	985	249 (158-384)	208	291 (204-423)	.00012
Lactate dehydrogenase, U/L	985	289 (222-408)	208	342 (247-443)	.018
AST, U/L	985	35 (25-47)	208	38 (28-54)	.0016
Serum creatinine, mg/dL	985	0.7 (0.5-0.8)	208	0.7 (0.6-0.9)	<.0001
eGFR, mL/min/1.73 m <sup>2</sup>	985	146 (119-171)	208	137 (101-154)	<.0001
Hospitalizations, no. per year	985	1 (0-3)	208	3 (1-7)	<.0001
Pregnancies, n (%) of patients	545	105 (19)	133	33 (25)	.15
Orthopedic procedures, n (%) of patients	985	29 (3)	208	23 (11)	<.0001
Central venous catheters, n (%) of patients	985	244 (25)	208	150 (72)	<.0001

Values are median (IQR) unless otherwise noted.

while trends for an association with tobacco use and higher AST were observed (Figure 2). In the subset of Hb SS/S $\beta^0$ -thalassemia patients (n = 831), independent risk factors for thrombosis were lower eGFR, higher AST, and hydroxyurea use ( $P \le .02$ ) (supplemental Figure 1).

#### **CVAs**

We observed 64 CVA events in 50 patients (4.2%). The median age at the first CVA event was 30 years (IQR, 20-46 years); 52 (81.3%) were ischemic events and 12 (18.8%) were ischemic events with hemorrhagic conversion. Differences in variables by CVA status are provided in Table 3. Independent risk factors for a CVA event on Cox regression analysis included older age and Hb SS/S $\beta^0$ -thalassemia genotype (P = .04) (Figure 3). In the subset of Hb SS/S $\beta^0$ -thalassemia genotype patients, a lower Hb F% was an independent risk factor for stroke (P = .046) (supplemental Figure 2).

We observed 20 recurrent CVA events in 9 (18%) of the 50 SCD patients that experienced an initial CVA. Rates of CVA recurrence

Figure 2. Independent risk factors for a thrombotic event in patients with SCD using a multivariate Cox proportional hazards model. CI, confidence interval.



did not differ based on initial CVA-related therapy (chronic transfusion, 8/36; anticoagulation, 0/5; aspirin, 0/4; no therapy, 1/4; P = .5). Stroke recurrence was observed in 24% (4/17) of SCD patients with Hb fractionations at goal for chronic transfusion (defined as >80% of Hb fractionations with Hb S  $\leq$ 30% within the first 2 years and Hb S  $\leq$ 50% after the first 2 years of a CVA event).<sup>5</sup>

## VTE

We observed 288 VTE events in 172 (14.4%) SCD patients; 194 VTE events occurred in females, and 94 VTE events occurred in males. The median age at the time of the first VTE event was 30 years (IQR, 25-43 years) and included 115 pulmonary embolism, 167 deep vein thrombosis (115 upper extremity, 51 lower extremity, and 1 superior vena cava), and 6 right atrial thrombus events. Onehundred and six (36.8%) of the VTEs occurred within 30 days of a previous hospital discharge. One-hundred and three (35.8%) VTE events were central venous catheter related, while 20 (6.9%) occurred in the perioperative period and 16 (8.2% of VTE events in females) during pregnancy. Differences in variables by VTE status are provided in Table 3. Independent risk factors for VTE risk on Cox regression analysis included lower eGFR (P < .001), hydroxyurea use (P = .003), Hb SS/S $\beta^{0}$ -thalassemia genotype (P = .01), higher Hb concentration (P = .03), and higher white blood cell count (P = .03), while a trend for an association with tobacco use was observed (P = .09) (Figure 4). A lower eGFR, hydroxyurea use, and a higher white blood cell count and Hb concentration were also independent risk factors for VTE in the subset of Hb SS/S $\beta^{0}$ -thalassemia genotype patients (supplemental Figure 3) and in the subset of non-catheter-related VTE events (n = 185) (supplemental Figure 4).

Fifty-seven of 172 (33.1%) patients with an initial VTE had recurrent VTE events. VTE recurrence was lower in SCD patients treated with an oral direct factor Xa inhibitor (3/29) compared with other therapies (warfarin, 44/113; low-molecular-weight heparin, 6/16; no therapy, 3/11; P = .008). For those patients started on warfarin, international normalized ratio (INR) results within the designated therapy duration were available for 104 of 113 patients. The

proportion of SCD patients on warfarin at goal (defined as >60% of INR results  $\ge 2.0$ )<sup>25</sup> was 17.3% (18/104). We did not observe an association between VTE recurrence and warfarin being at goal (VTE recurrence, 5/41 at goal; no VTE recurrence, 13/63 at goal; P = .4). We did not observe an association between VTE recurrence and duration of anticoagulation therapy (<3 months, 1/6; 3-6 months, 34/97; >6 months, 17/55, P = .8).

## African American gene risk variants for thrombosis

Next, we evaluated genetic risk variants that have been implicated in thrombosis risk in the general African American population in 327 SCD patients genotyped by the Affymetrix Pan-African Axiom GeneChip array. The median age of this cohort was 32 years (IQR, 23-43 years), 268 (82%) were Hb SS/S $\beta^{0}$ genotype, 196 (60%) were female, and 153 (47%) were on hydroxyurea. Two previously reported variants in the thrombomodulin gene (THBD) were associated with thrombosis risk (THBD rs2567617: MAF, 0.25; odds ratio [OR], 1.50; 95% confidence interval, 1.01-2.22; P = .049; and *THBD* rs1998081: MAF, 0.24; OR, 1.46; 95% confidence interval; 0.99-2.17; P = .059). The MAF for the factor V Leiden and prothrombin gene mutations was <0.01, so we were unable to impute and analyze these variants in this cohort. The factor V Leiden and the prothrombin gene mutations were tested as part of routine medical care in 56 and 42 SCD patients, respectively, and neither mutation was detected.

## Discussion

Thrombosis has been increasingly recognized as a complication of SCD that leads to substantial morbidity and early mortality.<sup>7-9,26</sup> In a longitudinal cohort of 1193 SCD patients with a median follow-up of 6.4 years, we demonstrate that thrombotic events are common, with a cumulative incidence of 17.4%. We observed high rates of recurrence for both CVA (18%) and venous thrombotic events (33%) and that the use of direct factor Xa inhibitors was associated with a lower rate of VTE recurrence compared with other anticoagulation therapies.

Table 3. Clinical and laborator	y values stratified by typ	e of thrombotic event in	patients with SCD

	n	No thrombosis	n	CVA	n	Venous thrombosis
Age at first encounter, y	985	23 (15-35)	50	26 (19-40)	172	27 (21-40)**
Male/female, %	985	45/55	50	40/60	172	35/65*
African American, n (%)	985	923 (94)	50	49 (98)	172	165 (96)
Sickle cell genotype, n (%)	985		50		172	
HbSS/SB0-thalassemia		657 (66.7)		44 (88.0)*		141 (82.0)**
HbSC or HbSB <sup>+</sup> -thalassemia or other		328 (33.3)		6 (12.0)		31 (18.0)
Hydroxyurea use, n (%)	985	496 (50)	50	34 (68)	172	141 (82)**
Tobacco use, n (%)	985	266 (27%)	50	15 (30%)	172	75 (44%)**
Body mass index, kg/m <sup>2</sup>	985	22 (18-26)	50	23 (20-26)	172	23 (20-28)**
Systolic blood pressure, mm Hg	985	115 (104-127)	50	122 (110-130)	172	122 (110-134)**
White blood cell count, $\times 10^3/\mu L$	985	9.7 (7.4-12.2)	50	9.1 (6.6-12.1)	172	10.2 (7.7-12.9)*
Platelet count, $\times 10^3/\mu L$	985	406 (287-525)	50	392 (335-569)	172	404 (311-529)
Hb, g/dL	985	9.6 (8.3-11.1)	50	8.8 (7.9-9.6)*	172	9.0 (8.3-10.3)*
Hb F, %	985	4.9 (2.1-11.2)	50	2.5 (1.1-7.8)*	172	5.2 (2.3-9.9)
Reticulocyte count, $\times 10^3/\mu L$	985	249 (158-384)	50	283 (249-488)*	172	291 (193-415)*
Lactate dehydrogenase, U/L	985	289 (222-408)	50	414 (296-481)**	172	323 (234-440)
AST, U/L	985	35 (25-47)	50	44 (30-58)*	172	37 (28-53)
Serum creatinine, mg/dL	985	0.7 (0.5-0.8)	50	0.7 (0.6-1.0)*	172	0.7 (0.6-0.9)**
eGFR, mL/min/1.73 m <sup>2</sup>	985	146 (119-171)	50	138 (95-154)*	172	136 (102-155)**
Hospitalizations, no. per year	985	1 (0-3)	50	1 (1-4)	172	4 (1-9)**
Pregnancies, n (%) of patients	545		30		112	
Before the first thrombotic event		105 (19)		1 (4)*		23 (21)
During the entire time period				3 (10)		31 (28)*
Orthopedic procedures, n (%) of patients	985		50		172	
Before the first thrombotic event		29 (3)		3 (6)		8 (5)
During the entire time period				5 (10)*		19 (11)**
Central venous catheters, n (%) of patients	985		50		172	
Before the first thrombotic event		244 (25)		16 (32)		89 (52)**
During the entire time period				36 (72)**		125 (72)**

Values are median (IQR) unless otherwise noted.

\*P < .05; \*\*P < .002 (statistically significant after Bonferroni correction).

Patients with SCD have a three- to 30-fold greater risk for CVA compared with the general African American population.<sup>27</sup> A CVA occurred in 4% of our SCD patients, which is consistent with what has been observed in other cohorts.<sup>28,29</sup> A lower Hb F% was associated with CVA risk in Hb SS/S $\beta^0$ -thalassemia patients, which has been observed in some SCD cohorts.<sup>4,30-32</sup> We observed CVA recurrence in 18% of SCD patients, including a rate of 24% for CVA recurrence in those SCD patients that maintained at Hb fractionation goal for at least 80% of their laboratory checks. These rates are similar to the 18% recurrence rate described in SCD children at goal<sup>5</sup> and highlights the need for alternative strategies to identify high-risk patients and for developing therapies to treat CVA in SCD.

SCD patients have a four- to 100-fold increased risk for VTE compared with the general population.<sup>33-35</sup> Many of these cohorts have relied solely on ICD coding and the risk factors for VTE in SCD patients are not clear. We applied a comprehensive approach integrating VTE events, identified by ICD coding and manually

confirmed through the medical records, with clinical and laboratory variables to identify risk factors for VTE. A VTE event occurred in 14% of our cohort, which is consistent with the 11% to 25% incidence reported in the literature.<sup>7-9</sup>

Decreasing eGFR was an independent risk factor for VTE in our cohort. In the general population, the risk for VTE is 1.3-fold higher in those with stage 1 CKD, increasing up to 2.1-fold higher in those with stage 3 or 4 CKD and 2.3-fold higher in those with stage 5 CKD.<sup>36</sup> Several pathways implicated in CKD-related thrombosis, including increased activation of procoagulant factors and platelets, endothelial injury, and reduced anticoagulant and fibrinolytic activity, overlap with the vasculopathy of SCD and may highlight common pathways for CKD and thrombosis in SCD patients. A high white blood cell count and Hb concentration were also risk factors for VTE in our cohort. Both a higher white blood cell count and a higher Hb concentration are also risk factors for thrombosis in polycythemia vera.<sup>37,38</sup> Common pathophysiologies, such as activated white blood cells promoting endothelial damage and a higher Hb

Figure 3. Risk factors for a cerebrovascular event in patients with SCD using a multivariate Cox proportional hazards model.



concentration leading to hyperviscosity, may be involved in thrombosis risk for both polycythemia vera and SCD. Patients with SCD who developed a VTE in our cohort had more frequent hospitalizations, orthopedic surgeries, and central venous catheter placements. These risk factors, also identified in the general population<sup>39</sup> and other SCD cohorts,<sup>6,8,9</sup> may promote thrombosis by reflecting the severity of SCD, increasing venous stasis, and/or causing direct vascular injury.

In our study, hydroxyurea was associated with a greater risk for VTE, but not CVA, in the entire cohort as well as in the subset of Hb SS/S $\beta^0$  patients and for non-catheter-related VTE. The use of hydroxyurea may be a marker for more severe SCD. Interestingly, hydroxyurea may have differential effects on the venous and arterial vasculature, as demonstrated in myeloproliferative neoplasms.<sup>40-42</sup> In a study of patients with polycythemia vera, hydroxyurea conferred

no protective benefit against VTE but led to a threefold reduction in arterial thrombotic events.<sup>40</sup> Hydroxyurea targets multiple processes implicated in arterial thrombosis, such as qualitative changes in white blood cells, reduced expression of endothelial adhesion molecules, and enhanced nitric oxide generation.<sup>43</sup> Reduced blood flow due to hyperviscosity is predominantly associated with venous thrombosis and may not be ameliorated by hydroxyurea.<sup>40</sup> The association of hydroxyurea with VTE risk in SCD patients will need to be investigated in future studies.

VTE recurrence was observed in 33% of SCD patients and was lower in those treated with an oral direct factor Xa inhibitor (10%) compared with alternative therapies (warfarin, 39%; low-molecular-weight heparin, 38%). The advantage of direct oral anticoagulants over warfarin for thrombotic risk has been observed in some,<sup>44,45</sup> but not all, non-SCD cohorts.<sup>46</sup> This advantage may be partially

Variable	Hazard Ratio (95% CI)		P value
Estimated GFR (log)	0.54 (0.39 - 0.74)	┝═┥	< 0.001
Hydroxyurea Use	2.42 (1.36 - 4.30)	⊢∎⊣	0.003
HbSS/HbS $\beta$ 0-thalassemia	2.47 (1.24 - 4.90)	∎	0.010
Hemoglobin	1.15 (1.01 — 1.31)	•	0.031
White Blood Cell Count (log)	1.85 (1.06 - 3.24)	⊢∎⊣	0.032
Tobacco Use	1.43 (0.95 - 2.14)	┠╼═╌┤	0.09
Age	1.01 (0.99 - 1.02)	•	0.4
Female Sex	1.01 (0.67 - 1.52)	<b>⊢</b> ∎-	1.0
		0.5 1.0	10.0

Figure 4. Risk factors for a venous thrombotic event in patients with SCD using a multivariate Cox proportional hazards model.

dependent on the time in therapeutic range for warfarin.<sup>44</sup> Interestingly, the proportion of INR results at goal was not associated with the rate of VTE recurrence. Future prospective studies comparing the safety and efficacy of direct oral anticoagulants vs warfarin to treat VTE in SCD are warranted.

Genetic variation may also contribute to the risk of thrombosis. Variants in *ABO*, *IGFBP3*, and *THBD* have been implicated in the thrombotic risk of African Americans.<sup>10</sup> We tested these variants and were able to identify a 1.5-fold increased risk for thrombosis in SCD patients with the *THBD* rs2567617 and rs1998081 variants. *THBD* encodes an endothelial cell glycoprotein, thrombomodulin, that plays a central role in anticoagulation by (1) reducing circulating thrombin levels, (2) increasing the activation of protein C by >2 orders of magnitude after complexing with thrombin, and (3) downregulating complement activation.<sup>47,48</sup> Variants in *THBD* have been implicated in both arterial<sup>49-53</sup> and venous<sup>54-56</sup> thrombotic complications as well as in the atypical hemolytic uremic syndrome.<sup>57</sup> Future investigation for the effects of *THBD* rs2567617 and rs1998081 on thrombomodulin function and thrombotic risk in SCD may provide insight into the mechanisms and risks of SCD vasculopathy.

Limitations of this study include that it was a single-center study and retrospective in nature. Fragmented care is common in SCD patients,<sup>58</sup> and our study was limited in the ability to detect thrombotic events that may have occurred at outside institutions, leading to an ascertainment bias. The median age at first encounter in SCD patients that developed a thrombotic event was older compared with those that did not have a thrombotic event, potentially leading to a selection bias. Orthopedic procedures, central venous catheter procedures, and pregnancies were captured regardless of timing to thrombosis, limiting our interpretation to associations, not causation. The definition of hydroxyurea was ever vs never in this study. The use and adherence of hydroxyurea among eligible patients is low,<sup>59</sup> and future prospective studies capturing the timing of procedures and pregnancies and confirming hydroxyurea use will help address these limitations.

In summary, in a cohort of nearly 1200 SCD patients, we observed a high rate of thrombotic events and identified several traditional (hospitalization rate, central venous catheter placement, pregnancy, and kidney disease) and SCD-specific (hydroxyurea therapy, Hb SS/S $\beta^0$ -thalassemia genotype, lower Hb F%, higher AST, higher white blood cell count, and higher Hb concentration) risk factors for thrombosis. In addition, we demonstrated a very high rate of recurrence for CVA and venous thrombotic events and that SCD patients treated with a direct oral factor Xa inhibitors had lower rates of VTE recurrence compared with other treatment strategies. Future prospective studies understanding the role of thrombomodulin in maintaining vascular flow and developing thrombotic risk profiles may guide the prophylactic and treatment strategies for SCD.

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# **Authorship**

Contribution: A.S., R.R., X.Z., and S.L.S. designed and performed research, analyzed the data, and wrote the paper; and B.N.S., J.H., M.G., R.E.M., and V.R.G. designed and performed research and wrote the paper.

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## References

- 1. Hassell KL. Population estimates of sickle cell disease in the U.S. Am J Prev Med. 2010;38(4 Suppl):S512-S521.
- 2. Kato GJ, Hebbel RP, Steinberg MH, Gladwin MT. Vasculopathy in sickle cell disease: Biology, pathophysiology, genetics, translational medicine, and new research directions. Am J Hematol. 2009;84(9):618-625.
- 3. Ataga KI, Key NS. Hypercoagulability in sickle cell disease: new approaches to an old problem. *Hematology Am Soc Hematol Educ Program*. 2007; 2007:91-96.
- 4. Wong WY, Powars DR. Overt and incomplete (silent) cerebral infarction in sickle cell anemia: diagnosis and management. *Hematol Oncol Clin North Am.* 2005;19(5):839-855, vi.
- 5. Hulbert ML, McKinstry RC, Lacey JL, et al. Silent cerebral infarcts occur despite regular blood transfusion therapy after first strokes in children with sickle cell disease. *Blood*. 2011;117(3):772-779.
- 6. Woods GM, Sharma R, Creary S, et al. Venous thromboembolism in children with sickle cell disease: a retrospective cohort study. *J Pediatr.* 2018;197: 186-190.e181.
- 7. Naik RP, Streiff MB, Haywood C Jr., Segal JB, Lanzkron S. Venous thromboembolism incidence in the Cooperative Study of Sickle Cell Disease. *J Thromb Haemost.* 2014;12(12):2010-2016.
- 8. Naik RP, Streiff MB, Haywood C Jr., Nelson JA, Lanzkron S. Venous thromboembolism in adults with sickle cell disease: a serious and under-recognized complication. *Am J Med.* 2013;126(5):443-449.

- 9. Brunson A, Lei A, Rosenberg AS, White RH, Keegan T, Wun T. Increased incidence of VTE in sickle cell disease patients: risk factors, recurrence and impact on mortality. Br J Haematol. 2017;178(2):319-326.
- 10. Hernandez W, Gamazon ER, Smithberger E, et al. Novel genetic predictors of venous thromboembolism risk in African Americans. *Blood*. 2016;127(15): 1923-1929.
- 11. Michalik DE, Taylor BW, Panepinto JA. Identification and validation of a sickle cell disease cohort within electronic health records. Acad Pediatr. 2017; 17(3):283-287.
- 12. Levey AS, Stevens LA, Schmid CH, et al; CKD-EPI (Chronic Kidney Disease Epidemiology Collaboration). A new equation to estimate glomerular filtration rate. *Ann Intern Med.* 2009;150(9):604-612.
- 13. Zhang X, Zhang W, Ma SF, et al. Hypoxic response contributes to altered gene expression and precapillary pulmonary hypertension in patients with sickle cell disease. *Circulation.* 2014;129(16):1650-1658.
- 14. Akaike H, Parzen E, Tanabe K, Kitagawa G. Selected papers of Hirotugu Akaike. New York: Springer; 1998.
- Therneau TM. A Package for Survival Analysis in R. R package version 2.38 (2015). Available at: https://CRAN.R-project.org/package=survival. Accessed 1 March 2018.
- 16. Therneau TM, Crowson C, Atkinson E. Using Time Dependent Covariates and Time Dependent Coefficients in the Cox Model. 2020. Available at: https://cran.r-project.org/web/packages/survival/vignettes/timedep.pdf. Accessed 1 February 2020.
- 17. R Core Team. R: A Language and Environment for Statistical Computing. Vienna, Austria: R Foundation for Statistical Computing; 2013.
- 18. Wickham H. tidyverse: Easily Install and Load the 'Tidyverse' (Version 1.2.1). 2017. Available at: https://CRAN.R-project.org/package=tidyverse. Accessed 1 October 2017.
- 19. Grolemund G, Wickham H. Dates and Times Made Easy with {lubridate}. J Stat Softw. 2011;40(3):1-25.
- Robinson D, Hayes A. broom: Convert Statistical Analysis Objects into Tidy Tibbles (Version 0.5.1). 2018. Available at: https://CRAN.R-project.org/ package=broom. Accessed 1 October 2017.
- 21. Therneau TM, Grambsch PM. Modeling Survival Data: Extending the Cox Model. New York, NY: Springer; 2000.
- 22. Kassambara A, Kosinski M. survminer: Drawing Survival Curves using 'ggplot2' (Version 0.4.3). 2018. Available at: https://CRAN.R-project.org/ package=survminer. Accessed 1 March 2018.
- 23. Hothorn T, Hornik K, van de Wiel MA, Zeileis A. Implementing a class of permutation tests: The {coin} package. J Stat Softw. 2008;28(8):1-23.
- healthyr.surgicalinformatics.org. tidyforest: A tidyverse workflow for running and presenting a logistic regression model in R. (Version 0.1.0). 2019. Available at:https://github.com/riinuots/tidyforest. Accessed 1 March 2019.
- 25. White HD. Comparison of outcomes among patients randomized to warfarin therapy according to anticoagulant control. Arch Intern Med. 2007;167(3): 239-245.
- Kumar R, Stanek J, Creary S, Dunn A, O'Brien SH. Prevalence and risk factors for venous thromboembolism in children with sickle cell disease: an administrative database study. Blood Adv. 2018;2(3):285-291.
- 27. Strouse JJ, Lanzkron S, Urrutia V. The epidemiology, evaluation and treatment of stroke in adults with sickle cell disease. *Expert Rev Hematol.* 2011;4(6): 597-606.
- 28. Ohene-Frempong K, Weiner SJ, Sleeper LA, et al. Cerebrovascular accidents in sickle cell disease: rates and risk factors. Blood. 1998;91(1):288-294.
- 29. Powars D, Wilson B, Imbus C, Pegelow C, Allen J. The natural history of stroke in sickle cell disease. Am J Med. 1978;65(3):461-471.
- Domingos IF, Falcão DA, Hatzlhofer BL, et al. Influence of the βs haplotype and α-thalassemia on stroke development in a Brazilian population with sickle cell anaemia. Ann Hematol. 2014;93(7):1123-1129.
- 31. Adams RJ, Kutlar A, McKie V, et al. Alpha thalassemia and stroke risk in sickle cell anemia. Am J Hematol. 1994;45(4):279-282.
- 32. Rees DC, Lambert C, Cooper E, et al. Glucose 6 phosphate dehydrogenase deficiency is not associated with cerebrovascular disease in children with sickle cell anemia. *Blood.* 2009;114(3):742-743, author reply 743-744.
- Novelli EM, Huynh C, Gladwin MT, Moore CG, Ragni MV. Pulmonary embolism in sickle cell disease: a case-control study. J Thromb Haemost. 2012; 10(5):760-766.
- 34. Noubiap JJ, Temgoua MN, Tankeu R, Tochie JN, Wonkam A, Bigna JJ. Sickle cell disease, sickle trait and the risk for venous thromboembolism: a systematic review and meta-analysis. *Thromb J.* 2018;16(1):27.
- 35. Stein PD, Beemath A, Meyers FA, Skaf E, Olson RE. Deep venous thrombosis and pulmonary embolism in hospitalized patients with sickle cell disease. *Am J Med.* 2006;119(10):897.e897-811.
- 36. Wattanakit K, Cushman M. Chronic kidney disease and venous thromboembolism: epidemiology and mechanisms. *Curr Opin Pulm Med.* 2009;15(5): 408-412.
- 37. Barbui T, Finazzi G, Falanga A. Myeloproliferative neoplasms and thrombosis. Blood. 2013;122(13):2176-2184.
- Marchioli R, Finazzi G, Specchia G, et al; CYTO-PV Collaborative Group. Cardiovascular events and intensity of treatment in polycythemia vera. N Engl J Med. 2013;368(1):22-33.
- Heit JA, Silverstein MD, Mohr DN, Petterson TM, O'Fallon WM, Melton LJ III. Risk factors for deep vein thrombosis and pulmonary embolism: a population-based case-control study. Arch Intern Med. 2000;160(6):809-815.

- 40. Barbui T, De Stefano V, Ghirardi A, Masciulli A, Finazzi G, Vannucchi AM. Different effect of hydroxyurea and phlebotomy on prevention of arterial and venous thrombosis in Polycythemia Vera. *Blood Cancer J.* 2018;8(12):124.
- 41. Stein BL, Martin K. From Budd-Chiari syndrome to acquired von Willebrand syndrome: thrombosis and bleeding complications in the myeloproliferative neoplasms. *Blood.* 2019;134(22):1902-1911.
- 42. De Stefano V, Rossi E, Carobbio A, et al. Hydroxyurea prevents arterial and late venous thrombotic recurrences in patients with myeloproliferative neoplasms but fails in the splanchnic venous district. Pooled analysis of 1500 cases. *Blood Cancer J*. 2018;8(11):112.
- 43. Maugeri N, Giordano G, Petrilli MP, et al. Inhibition of tissue factor expression by hydroxyurea in polymorphonuclear leukocytes from patients with myeloproliferative disorders: a new effect for an old drug? *J Thromb Haemost*. 2006;4(12):2593-2598.
- 44. Ruff CT, Giugliano RP, Braunwald E, et al. Comparison of the efficacy and safety of new oral anticoagulants with warfarin in patients with atrial fibrillation: a meta-analysis of randomised trials. *Lancet.* 2014;383(9921):955-962.
- 45. Shah S, Norby FL, Datta YH, et al. Comparative effectiveness of direct oral anticoagulants and warfarin in patients with cancer and atrial fibrillation. *Blood Adv.* 2018;2(3):200-209.
- 46. Burn J, Pirmohamed M. Direct oral anticoagulants versus warfarin: is new always better than the old [published correction appears in *Open Heart*. 2018]? *Open Heart*. 2018;5(1):e000712.
- 47. Weiler H, Isermann BH. Thrombomodulin. J Thromb Haemost. 2003;1(7):1515-1524.
- 48. Boffa MC, Karmochkine M. Thrombomodulin: an overview and potential implications in vascular disorders. Lupus. 1998;7(suppl 2):S120-S125.
- 49. Ireland H, Kunz G, Kyriakoulis K, Stubbs PJ, Lane DA. Thrombomodulin gene mutations associated with myocardial infarction. *Circulation*. 1997;96(1): 15-18.
- Wu KK, Aleksic N, Ahn C, Boerwinkle E, Folsom AR, Juneja H; Atherosclerosis Risk in Communities Study (ARIC) Investigators. Thrombomodulin Ala455Val polymorphism and risk of coronary heart disease. *Circulation*. 2001;103(10):1386-1389.
- 51. Doggen CJ, Kunz G, Rosendaal FR, et al. A mutation in the thrombomodulin gene, 127G to A coding for Ala25Thr, and the risk of myocardial infarction in men. *Thromb Haemost.* 1998;80(5):743-748.
- Kunz G, Ireland HA, Stubbs PJ, Kahan M, Coulton GC, Lane DA. Identification and characterization of a thrombomodulin gene mutation coding for an elongated protein with reduced expression in a kindred with myocardial infarction. *Blood.* 2000;95(2):569-576.
- 53. Norlund L, Holm J, Zöller B, Ohlin AK. A common thrombomodulin amino acid dimorphism is associated with myocardial infarction. *Thromb Haemost.* 1997;77(2):248-251.
- 54. Le Flem L, Picard V, Emmerich J, et al. Mutations in promoter region of thrombomodulin and venous thromboembolic disease. Arterioscler Thromb Vasc Biol. 1999;19(4):1098-1104.
- 55. Ohlin AK, Marlar RA. The first mutation identified in the thrombomodulin gene in a 45-year-old man presenting with thromboembolic disease. *Blood.* 1995;85(2):330-336.
- 56. Ohlin AK, Marlar RA. Thrombomodulin gene defects in families with thromboembolic disease-a report on four families. *Thromb Haemost*. 1999;81(3): 338-344.
- 57. Delvaeye M, Noris M, De Vriese A, et al. Thrombomodulin mutations in atypical hemolytic-uremic syndrome. N Engl J Med. 2009;361(4):345-357.
- Shatola A, Brunson A, Keegan T, Wun T, Mahajan A. Fragmentation of care for young adults with sickle cell disease in California [abstract]. Blood. 2019; 134(suppl 1). Abstract 4667.
- Candrilli SD, O'Brien SH, Ware RE, Nahata MC, Seiber EE, Balkrishnan R. Hydroxyurea adherence and associated outcomes among Medicaid enrollees with sickle cell disease. Am J Hematol. 2011;86(3):273-277.