

Effect of a Recombinant Human Soluble Thrombomodulin on Baseline Coagulation Biomarker Levels and Mortality Outcome in Patients With Sepsis-Associated Coagulopathy

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Objectives: To assess the effects of recombinant human soluble thrombomodulin treatment on 28-day all-cause mortality in subgroups categorized by baseline coagulation biomarker levels (prothrombin fragment 1.2, thrombin-antithrombin complex, D-dimer) in patients with sepsis-associated coagulopathy in the Sepsis Coagulopathy Asahi Recombinant LE Thrombomodulin trial (SCARLET) (NCT01598831).

Design: Post hoc, subgroup analysis of a randomized, double-blind, placebo-controlled, multinational, multicenter phase 3 study.

Setting: ICUs at 159 sites in 26 countries.

Patients: Eight-hundred adults with sepsis-associated coagulopathy defined as international normalized ratio greater than 1.40 and platelet count between $30 \times 10^9/L$ and $150 \times 10^9/L$ or greater than 30% decrease within 24 hours with concomitant cardiovascular and/or respiratory failure.

Interventions: Patients randomized and treated with recombinant human soluble thrombomodulin (0.06 mg/kg/d; $n = 395$) or equivalent placebo ($n = 405$) for 6 days.

Measurements and Main Results: Recombinant human soluble thrombomodulin did not significantly reduce 28-day all-cause mor-

tality in the Sepsis Coagulopathy Asahi Recombinant LE Thrombomodulin trial: absolute risk reduction was 2.55% ($p = 0.32$) in patients with sepsis-associated coagulopathy. In this post hoc analysis, mortality steadily increased with increasing baseline prothrombin fragment 1.2 and thrombin-antithrombin complex levels in the placebo group; for those values exceeding the upper limit of normal, the mortality increases in the recombinant human soluble thrombomodulin group were lower or negligible with increasing baseline prothrombin fragment 1.2 and thrombin-antithrombin complex. Consequently, absolute risk reductions were greater in subgroups with higher baseline prothrombin fragment 1.2 or thrombin-antithrombin complex. Absolute risk reductions were also greater in subgroups with baseline coagulation biomarker levels at or above median of the entire study population, ranging from 4.2% (95% CI, -5.0% to 13.4%) to 5.5% (95% CI, -4.0% to 14.9%).

Conclusions: Compared with patients receiving placebo, patients treated with recombinant human soluble thrombomodulin having higher baseline thrombin generation biomarker levels had lower mortality. Further research regarding the predictive role of coagulation biomarkers for recombinant human soluble thrombomodulin treatment response in sepsis-associated coagulopathy is warranted to evaluate clinical relevance. (*Crit Care Med* 2020; 48:1140–1147)

Key Words: international normalized ratio; platelet count; prothrombin fragment 1.2; recombinant human soluble thrombomodulin; sepsis; thrombin-antithrombin complex

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Sepsis can lead to life-threatening multiple organ dysfunction that is estimated to contribute to approximately 5 million deaths globally per year (1, 2). It is caused by a dysregulated response of the body to infection and is often complicated by coagulopathy and endothelial dysfunction (1, 3, 4). Disseminated intravascular coagulation (DIC) is a key

predictor of clinical outcomes in patients with sepsis and is associated with aberrantly enhanced thrombin generation (5, 6). Generated thrombin is primarily neutralized by antithrombin, thereby forming thrombin-antithrombin complexes (TAT). Prothrombin fragment 1.2 (F1.2) is an activation peptide generated when prothrombin is converted to thrombin (7). New criteria for DIC from the Japanese Society on Thrombosis and Hemostasis (JSTH) include these biomarkers of thrombin generation, TAT, and F1.2 to improve the sensitivity and specificity of the DIC criteria. If any of these values are greater than or equal to two-fold of the respective upper limit of normal, then a point is added to the JSTH diagnostic score (8).

There is an incremental increased risk of mortality with an increase in the severity of sepsis-associated coagulopathy (SAC), as defined by prolonged prothrombin time-international normalized ratio (PT-INR) and reduced platelet count (9). In SAC, a systemic inflammatory response to the infection is triggered, which leads to coagulation activation (10). Activating the coagulation cascade also leads to excess formation of fibrin, the primary component of a thrombus. A key enzymatic degradation product of this process is D-dimer (11). Endothelial expression of the pivotal receptor of the crucial regulatory protein C pathway, thrombomodulin, is severely reduced (10, 12), leading to reduced activated protein C (APC), which functions to proteolytically degrade coagulation factors Va and VIIIa, exerting an antithrombotic effect (13). In addition, the APC pathway has anti-inflammatory properties (10, 12, 14). An initial phase 3 study of recombinant human APC (rhAPC; Xigris, Eli Lilly, Indianapolis, IN) showed that rhAPC decreased mortality in patients with severe sepsis (15). However, subsequent studies showed less-favorable results and rhAPC has been withdrawn from the market (16).

Recombinant human soluble thrombomodulin (ART-123) exerts anticoagulant and anti-inflammatory effects (17). Unlike exogenously administered rhAPC, ART-123 binds circulating thrombin molecules to serve as an activation complex to convert endogenous protein C to endogenous APC, which is the underlying mechanism for the anticoagulant effect of ART-123 (17). Thus, the complex formed by the binding of ART-123 with thrombin is the initial step in exerting the anticoagulant and anti-inflammatory effects of ART-123. However, plasma APC activity is not increased in patients with sepsis-associated DIC who are treated with ART-123. It is possible that APC acts locally and does not systemically circulate (18). Thus, endogenously generated APC concentrations are very low and nearly undetectable. In addition, ART-123 inhibits inflammation and organ injury caused by damage-associated molecular patterns, such as high mobility group box protein 1 and histones (19–21).

ART-123 was evaluated in a global phase 2b clinical study in patients with sepsis and DIC (NCT00487656) (22). The 28-day mortality rates were 17.8% in the ART-123 treatment group and 21.6% in the placebo group ($p = 0.27$) (22). Biomarker analysis in this study showed that ART-123–treated patients exhibited a significant decrease in the levels of coagulation biomarkers F1.2, TAT, and D-dimer compared with patients who received placebo at day 7 ($p < 0.05$) (23).

Sepsis Coagulopathy Asahi Recombinant LE Thrombomodulin (SCARLET) (ClinicalTrials.gov identifier: NCT01598831) was a randomized, double-blind, placebo-controlled, multinational, multicenter, parallel-group phase 3 trial of patients with organ dysfunction and coagulopathy due to severe sepsis to assess the safety and efficacy of ART-123 treatment. ART-123 did not significantly reduce 28-day all-cause mortality (primary efficacy endpoint) versus placebo—the absolute risk reduction (ARR) of 28-day all-cause mortality in ART-123–treated patients was 2.55% ($p = 0.32$). However, in post hoc analyses, 28-day all-cause mortality rates in the subgroup of patients who maintained the protocol-specified coagulopathy criteria at baseline (after randomization and before the first dose of study drug; $n = 634$; PT-INR > 1.40 and platelet count $> 30 \times 10^9/L$) were 26.7% (82/307) for ART-123 and 32.1% (105/327) for placebo—a difference of 5.40% (95% CI, -1.7% to 12.5%). In the post hoc analysis, a significant change from baseline was observed in each biomarker during the 6-day treatment period among those receiving ART-123 versus those assigned to placebo ($p < 0.05$), despite the reduction in 28-day all-cause mortality not being significant in the ART-123 treatment group. Thrombin generation markers F1.2 and TAT appeared to undergo significantly more rapid reduction in the ART-123 group as compared with the placebo group. Similarly, plasma levels of D-dimer were lowered after treatment with ART-123 compared with placebo (24).

The objective of this post hoc analysis is to present the 28-day all-cause mortality data for patients treated with ART-123 categorized by baseline coagulation biomarker levels (F1.2, TAT, and D-dimer) in patients with SAC in the SCARLET trial.

MATERIALS AND METHODS

Study Design

A total of 816 patients who were critically ill with SAC were randomized in a 1:1 ratio to allocate to ART-123 ($n = 402$) or placebo ($n = 414$). Among these patients, 800 individuals in the full analysis set (FAS) were randomized and treated with either IV ART-123 ($n = 395$) or equivalent placebo ($n = 405$). ART-123 was dosed at 0.06 mg/kg/d up to a maximum dose of 6.0 mg/d for 6 days, as described previously (24). For each of the participating institutions, the institutional review board or local ethics committee reviewed and approved the study. Informed consent was obtained from all participants prior to performing any study-specific procedures to conform to all applicable local, regulatory, and ethical requirements. This study was conducted in compliance with the International Conference on Harmonisation Good Clinical Practice guideline (25).

Patient Population

Patients greater than or equal to 18 years old who were critically ill with SAC from a bacterial source and receiving treatment with IV antibiotics, with WBC counts greater than $12,000/mm^3$ or less than $4,000/mm^3$ or bandemia greater than 10%, temperature of less than $36^\circ C$ or fever greater than $38^\circ C$, with a concurrent diagnosis of cardiovascular dysfunction

(requiring both adequate fluid resuscitation and vasopressors) and/or respiratory dysfunction (requiring mechanical ventilation for hypoxemia), and coagulopathy were included in the study. Coagulopathy was characterized by 1) PT-INR greater than 1.40 without other known etiology (e.g., anticoagulant therapy, chronic liver disease) and 2) platelet count between $30 \times 10^9/L$ and $150 \times 10^9/L$ or greater than 30% decrease in platelet count within 24 hours. Subcutaneous heparin for thromboprophylaxis was permitted at doses per the manufacturer recommendations in the package inserts approved for each country. Wide variability was observed across countries in the heparin dose administered and percentage of enrolled patients receiving prophylactic heparin at baseline.

Approximately 80% of patients (634/800) maintained protocol-specified eligibility for coagulopathy of PT-INR greater than 1.40 and platelet count greater than $30 \times 10^9/L$ before initiating the first dose of study drug (baseline). In the remaining patients, PT-INR was less than or equal to 1.40 and/or platelet count less than $30 \times 10^9/L$ at baseline after initial confirmation of eligibility. The subset of patients with PT-INR greater

than 1.40 and platelet count greater than $30 \times 10^9/L$ at baseline will be referred to as having “confirmed coagulopathy” in this article.

Measurement and Analysis of Coagulation Biomarkers

Citrate- or ethylenediaminetetraacetic acid-anticoagulated whole blood was drawn from patients. Plasma was prepared from the blood and frozen at $-70^\circ C$ or $-20^\circ C$ and shipped to the central laboratory (PPD, Wilmington, NC). Plasma samples for F1.2 and TAT measurements were analyzed on a monthly basis using commercially available enzyme-linked immunosorbent assay (ELISA) kits (Siemens, Berlin, Germany) at Loyola University Chicago, Hemostasis and Thrombosis Research Laboratories. The coefficient of variation (CV) between assays for F1.2 was 4.8% and the CV for TAT was 4.1%. The plasma samples for D-dimer measurements were analyzed on a weekly basis using commercially available ELISA kits (Instrumentation Laboratory, Bedford, MA) at PPD. The CV for D-dimer intra-assay variation was 7.0%.

TABLE 1. Patient Characteristics and Baseline Coagulation Biomarker Levels by Subgroups Based on Prothrombin Fragment 1.2 Levels in the Full Analysis Set of the Sepsis Coagulopathy Asahi Recombinant LE Thrombomodulin Trial

Characteristic	Recombinant Human Soluble Thrombomodulin		Placebo	
	F1.2 < 358.9 (n = 187)	F1.2 ≥ 358.9 (n = 191)	F1.2 < 358.9 (n = 198)	F1.2 ≥ 358.9 (n = 194)
Age, mean (SD), yr				
Median	62	63	62	63
Q1–Q3	50–74	55–74	51–73	51–75
Male, n (%)	110 (58.8)	97 (50.8)	120 (60.6)	97 (50.0)
Baseline international normalized ratio				
Mean (SD)	1.7 (0.4)	1.8 (0.8)	1.7 (0.4)	1.8 (0.8)
Baseline platelets, $\times 10^9/L$				
Mean (SD)	134.2 (78.6)	114.1 (66.2)	129.2 (68.2)	114.5 (61.7)
Acute Physiology and Chronic Health Evaluation II score				
n	167	172	180	173
Mean (SD)	21.4 (8.4)	23.4 (7.8)	21.7 (8.3)	22.7 (7.8)
Cardiovascular dysfunction, n (%)	169 (90.4)	175 (91.6)	175 (88.4)	182 (93.8)
Respiratory dysfunction, n (%)	106 (56.7)	129 (67.5)	132 (66.7)	146 (75.3)
Renal dysfunction, n (%)	56 (29.9)	63 (33.0)	59 (29.8)	74 (38.1)
Hepatic dysfunction, n (%)	37 (19.8)	48 (25.1)	38 (19.2)	42 (21.6)
Arterial lactate, mmol/L				
n	148	152	169	158
Mean (SD)	3.5 (2.9)	4.9 (4.2)	3.3 (2.2)	4.4 (3.3)
Baseline heparin use, n (%)	107 (57.2)	86 (45.0)	110 (55.6)	97 (50.0)

F1.2 = prothrombin fragment 1.2.

Mortality Data for Baseline Coagulation Biomarker Subgroups

In this post hoc analysis, the placebo and ART-123 groups were categorized by baseline values of F1.2, TAT, and D-dimer (\leq upper limit of normal range, $>$ upper limit, ≥ 1.5 -fold upper limit, \geq two-fold upper limit, and ≥ 2.5 -fold upper limit; categories were based on normal range values from the package inserts of the assays for each of the three coagulation biomarkers), consistent with the new JSTH DIC criteria described previously (7). Mortality rates in the FAS population and the confirmed coagulopathy subgroup were evaluated according to baseline coagulation biomarker levels. According to the assay kit package inserts, the normal range for F1.2 was 69 to 229 pmol/L, and TAT was 1.0 to 4.1 μ g/L; for D-dimer, the upper limit was 243 ng/mL.

Statistical Analysis

CI_s were calculated using traditional 95% two-sided intervals and all calculations were performed using SAS Version 9.3 or 9.4 (SAS Institute, Cary, NC).

RESULTS

Patients

Baseline demographics and characteristics for all study patients were described previously (24). In summary, baseline characteristics were well balanced between the ART-123 and placebo groups in the FAS population. The mean age of the patients was 61.0 years in the ART-123 group and 60.5 years in the placebo group and approximately 55% of patients in each group were male. The demographics and baseline characteristics for patients with F1.2 and TAT biomarker levels below the median value and patients with biomarker levels at or above the median value were well balanced in the ART-123 and placebo groups (Tables 1 and 2). In patients with baseline thrombin generation markers at or above the median value, mean INR was slightly higher, platelet counts were slightly lower, and Acute Physiology and Chronic Health Evaluation II scores were slightly higher. In addition, standard-of-care prophylactic heparin use was lower at baseline in these patients.

TABLE 2. Patient Characteristics and Baseline Coagulation Biomarker Levels by Subgroups Based on Thrombin-Antithrombin Complex Levels in the Full Analysis Set of the Sepsis Coagulopathy Asahi Recombinant LE Thrombomodulin Trial

Characteristic	Recombinant Human Soluble Thrombomodulin		Placebo	
	TAT < 14.2 (n = 187)	TAT \geq 14.2 (n = 194)	TAT < 14.2 (n = 199)	TAT \geq 14.2 (n = 195)
Age, mean (SD), yr				
Median	61	64	64	61
Q1–Q3	49–72	56–75	52–74	50–72
Male, n (%)	119 (63.6)	91 (46.9)	120 (60.3)	97 (49.7)
Baseline international normalized ratio				
Mean (SD)	1.7 (0.6)	1.8 (0.7)	1.6 (0.3)	1.9 (0.8)
Baseline platelets, $\times 10^9/L$				
Mean (SD)	134.9 (72.7)	112.9 (72.6)	126.5 (65.6)	117.1 (65.1)
Acute Physiology and Chronic Health Evaluation II score				
n	166	176	176	179
Mean (SD)	20.7 (7.9)	24.0 (8.0)	21.1 (8.2)	23.2 (7.8)
Cardiovascular dysfunction, n (%)	166 (88.8)	181 (93.3)	179 (89.9)	181 (92.8)
Respiratory dysfunction, n (%)	107 (57.2)	132 (68.0)	137 (68.8)	143 (73.3)
Renal dysfunction, n (%)	52 (27.8)	68 (35.1)	57 (28.6)	75 (38.5)
Hepatic dysfunction, n (%)	36 (19.3)	50 (25.8)	43 (21.6)	38 (19.5)
Arterial lactate, mmol/L				
n	151	152	161	167
Mean (SD)	3.0 (2.1)	5.4 (4.4)	3.0 (1.8)	4.5 (3.5)
Baseline heparin use, n (%)	104 (55.6)	90 (46.4)	122 (61.3)	86 (44.1)

TAT = thrombin-antithrombin complex.

Mortality Data for Baseline Coagulation Biomarker Subgroups

As previously reported, the change from baseline of each coagulation biomarker (F1.2, TAT, D-dimer) during the treatment period (6 d) was statistically significantly greater in the ART-123 group than the placebo group (24). The baseline values of these coagulation biomarkers and 28-day all-cause mortality rates were also investigated in patients who were randomized and treated with ART-123 (*n* = 395) or placebo (*n* = 405; **Fig. 1**). These results were not statistically significant in this post hoc analysis. As the baseline plasma levels of F1.2 and TAT increased in the placebo group, the 28-day all-cause mortality rates increased from 20.0% and 19.2% in the group with biomarker levels less than or equal to the upper limit to 38.9% and 36.3% in the group with greater than or equal to the 2.5-fold upper limit, respectively. However, except for the group with baseline biomarker levels less than or equal to the upper limit, the 28-day all-cause mortality rates ranging from 28.8% to 31.8% and 27.0% to 29.4% were consistently lower or not apparent in patients treated with ART-123 versus those assigned to placebo, whose mortality rates ranged from 33.1% to 38.9% and 29.6% to 36.3%, across the range of increasing baseline F1.2 and TAT levels, respectively. Consequently, ARR were greater in the subgroups with higher baseline F1.2 or TAT levels. Analysis of D-dimer levels did not show a change in the 28-day all-cause mortality rates for either group with increasing baseline levels. ARR were also greater in the subgroups with baseline coagulation biomarker levels at or above median values of the entire study population: 4.7% (95% CI, -4.7% to 14.1%), 5.5% (95% CI, -4.0% to 14.9%), and 4.2% (95% CI, -5.0% to 13.4%) for F1.2, TAT, and D-dimer, respectively (**Table 3**). However, none of the ARRs indicate statistical significance.

A separate post hoc analyses was also performed in patients with confirmed coagulopathy (PT-INR > 1.40 and platelet count > 30 × 10⁹/L) at baseline who were treated with ART-123 (*n* = 308) and placebo (*n* = 326). In this subset of patients with confirmed coagulopathy and baseline coagulation biomarker levels at or above median values of the FAS population, ARRs were 10.3% (95% CI, -0.2% to 20.8%), 8.1% (95% CI, -2.3% to 18.6%), and 7.5% (95% CI, -2.8% to 17.8%) for F1.2, TAT,

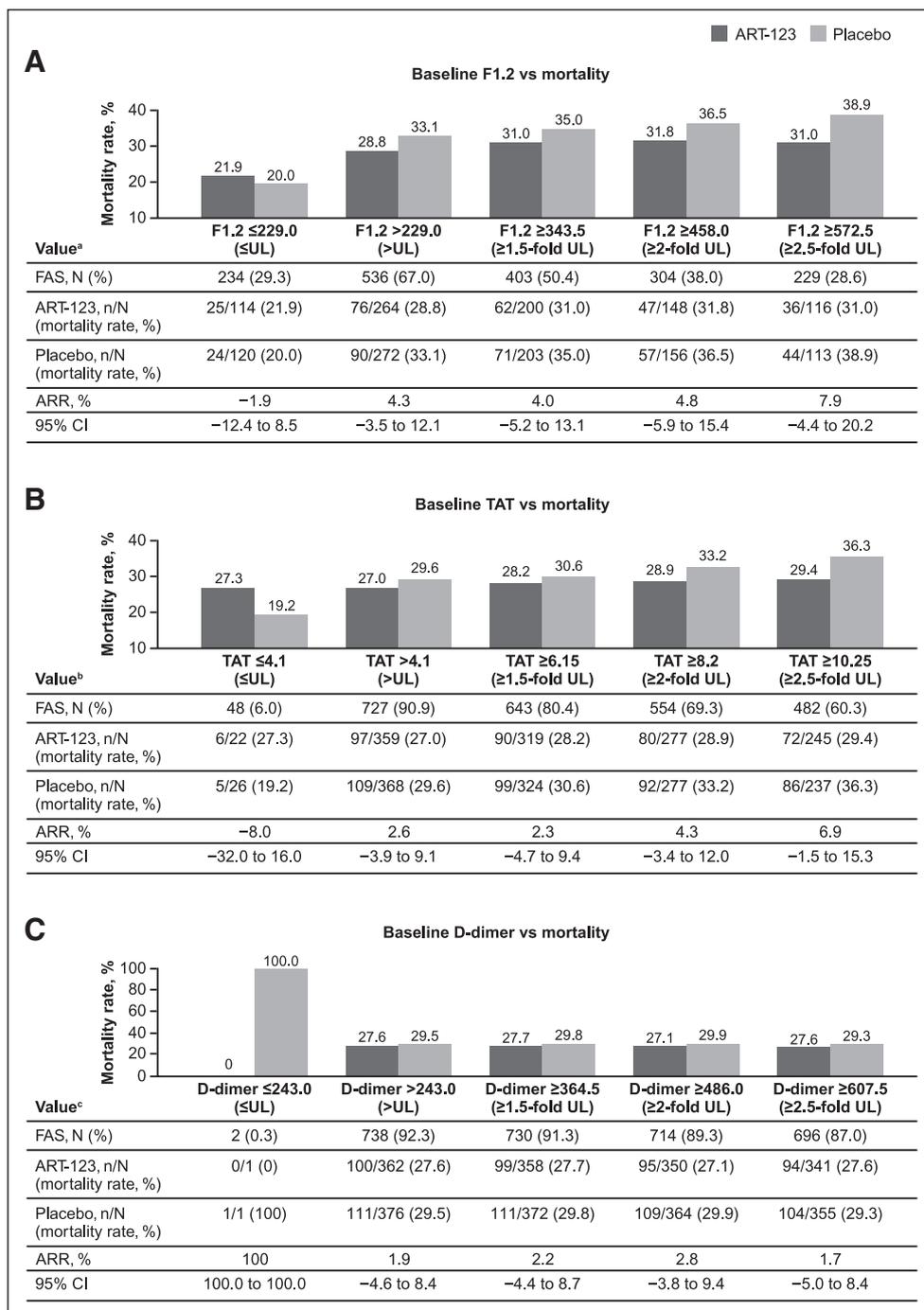


Figure 1. Mortality rates in coagulation biomarker subgroups. **A**, Prothrombin fragment 1.2 (F1.2), **B**, Thrombin-antithrombin complex (TAT), **C**, D-dimer, and 28-d all-cause mortality in patients treated with recombinant human soluble thrombomodulin (ART-123) or placebo. ^aData were unavailable for 30 patients. ^bData were unavailable for 25 patients. ^cData were unavailable for 60 patients. ARR = absolute risk reduction, FAS = full analysis set, UL = upper limit.

TABLE 3. Association of Baseline Coagulation Biomarkers With 28-Day All-Cause Mortality in the Post Hoc Analysis in the Full Analysis Set Population of the Sepsis Coagulopathy Asahi Recombinant LE Thrombomodulin Trial

Biomarker	Subgroup Concentration Relative to Median Value in Full Analysis Set ^a	Recombinant Human Soluble Thrombomodulin (n = 395) n (%) ^b		Placebo (n = 405) n (%) ^b		Absolute Risk Reduction (%) of 28-d All-Cause Mortality (95% CI)
Prothrombin fragment 1.2, pmol/L ^c	< 358.9	42/187 (22.5)	45/198 (22.7)	0.3	(-8.1 to 8.6)	
	≥ 358.9	59/191 (30.9)	69/194 (35.6)	4.7	(-4.7 to 14.1)	
Thrombin-antithrombin complex, ng/mL ^d	< 14.2	41/187 (21.9)	41/199 (20.6)	-1.3	(-9.5 to 6.9)	
	≥ 14.2	62/194 (32.0)	73/195 (37.4)	5.5	(-4.0 to 14.9)	
D-dimer, ng/mL ^e	< 3,030	50/173 (28.9)	57/197 (28.9)	0.0	(-9.2 to 9.3)	
	≥ 3,030	50/190 (26.3)	55/180 (30.6)	4.2	(-5.0 to 13.4)	

^aMedian values were calculated from baseline values in full analysis set.

^bRatio is number of patients with a particular biomarker result who were deceased by day 28 to number of patients with data for the particular biomarker.

^cData were unavailable for 30 patients.

^dData were unavailable for 25 patients.

^eData were unavailable for 60 patients.

and D-dimer, respectively, versus 2.55% for the FAS population and 5.40% for the patients with confirmed coagulopathy.

DISCUSSION

Coagulopathy is prevalent in patients with sepsis and likely plays a key role in multiple organ dysfunction (26, 27). SAC is correlated with an increased risk of mortality and is defined by a prolonged PT-INR and reduced platelet count. In this post hoc analysis, patients in the SCARLET trial were sorted by the baseline values of coagulation biomarkers (\leq upper limit, $>$ upper limit, \geq 1.5-fold upper limit, \geq two-fold upper limit, and \geq 2.5-fold upper limit), referring to the new JSTH DIC criteria, which include values of F1.2 or TAT (8). The upper limits of normal range were based on values from the package inserts for each of the three coagulation biomarkers. Findings from this post hoc analysis show that mortality rates steadily increased, with increasing levels of baseline thrombin generation markers, F1.2 and TAT, in the placebo group (Fig. 1). However, except for the group with F1.2 and TAT levels at or below the upper limit of normal, the increases in the 28-day all-cause mortality rates in the ART-123 group were lower or negligible with increasing baseline F1.2 and TAT levels. Consequently, ARRs were greater in the subgroups with higher baseline F1.2 or TAT levels. These results are consistent with the known anticoagulant effects of ART-123 exerted by binding with thrombin and modifying thrombin activity (17). The discrepancies between the observations with F1.2 or TAT levels and D-dimer levels are possibly due to F1.2 and TAT being pure markers of thrombin generation, whereas D-dimer is a marker of fibrin degradation (fibrinolysis). These D-dimer data may be a reflection of the impaired fibrinolysis known to occur in sepsis due to elevated levels of plasminogen activator inhibitor-1 (PAI-1), which inhibits plasmin formation and

subsequently the degradation of fibrin. High levels of PAI-1 are correlated with the severity of sepsis and are associated with increased mortality (28). The higher levels of PAI-1 in patients with severe sepsis could have affected the relationship between mortality and D-dimer levels in this analysis. In the previous post hoc analyses, 28-day all-cause mortality rates in the subgroup of patients who maintained the protocol-specified coagulopathy criteria at baseline (after randomization and before the first dose of study drug; $n = 634$, PT-INR > 1.40 and platelet count $> 30 \times 10^9/L$) were 26.7% (82/307) with ART-123 and 32.1% (105/327) with placebo—a difference of 5.40% (95% CI, -1.7% to 12.5%), exceeding the 2.55% ARR of the FAS ($p = 0.32$) (24). This post hoc analysis shows the mortality data for ART-123-treated patients with SAC and higher coagulation markers (F1.2, TAT, and D-dimer). The ARRs of these subgroups increased with higher baseline levels of thrombin generation biomarkers.

Results from a phase 1 study showed that the inhibition of thrombin generation was directly correlated with the concentration of ART-123 (29), and the pharmacokinetic profile of ART-123 showed that dosing at 0.06 mg/kg/d for 6 consecutive days caused a gradual increase in the plasma concentration of ART-123 up to 1,600 ng/mL at day 6, after which the levels slowly decreased (30). Similar outcomes were noted in the analysis of coagulation biomarker levels in patients in the SCARLET trial in which consistent reductions from baseline were observed in the plasma levels of each of the coagulation biomarkers F1.2, TAT, and D-dimer during the treatment period (6 d) and the reductions were greater with ART-123 than with placebo (24).

SAC is a heterogeneous condition, and there are limitations for applying the findings of this clinical study across all patients with this condition. Variations in standard of care for the treatment of SAC may exist across sites because

characteristics of ICUs vary among countries and regions (31). Such potential differences include use of prophylactic heparin. Data from this post hoc subgroup analysis should be reviewed with consideration that the sample size analyzed here limits the ability to draw robust conclusions. The findings from the post hoc analysis provide a solid rationale for pursuing additional studies with larger sample sizes to confirm these findings.

CONCLUSIONS

This post hoc analysis shows mortality data for patients with SAC according to baseline coagulation marker levels. Compared with patients treated with placebo, the subgroup of ART-123-treated patients with SAC who had higher baseline levels of thrombin generation biomarkers had lower mortality. Further research regarding the predictive role of coagulation biomarkers for ART-123 treatment response in patients with SAC is warranted to evaluate clinical relevance.

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Significance of present work: Sepsis is a host response to infection leading to activation of inflammation and coagulation and is associated with high mortality rates. Recombinant human soluble thrombomodulin (ART-123) has distinct anticoagulant and anti-inflammatory effects. Primary results from the randomized, double-blind, placebo-controlled, multinational phase 3 Sepsis Coagulopathy Asahi Recombinant LE Thrombomodulin trial (SCARLET) (NCT01598831) showed that ART-123 did not significantly reduce 28-day all-cause mortality (absolute risk reduction [ARR], 2.55% [$p = 0.32$]), in patients with sepsis-associated coagulopathy. Herein we report that the ARR values with ART-123 were higher in patients with higher baseline thrombin generation biomarker levels in the SCARLET trial.

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The work for the Sepsis Coagulopathy Asahi Recombinant LE Thrombomodulin trial, a multinational, multicenter study, was performed in 27 countries (320 sites), of which 26 countries enrolled patients (159 enrolling sites) spanning across Europe (including Israel), North and South Americas, Russia, and the Asia-Pacific region.

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