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Evaluation of high-sensitivity C-reactive protein and uric acid in vericiguat-treated patients with heart failure with reduced ejection fraction

Frank Kramer¹*, Sebastian Voss², Lothar Roessig³, Bernd-Wolfgang Igl⁴, Javed Butler⁵, Carolyn S.P. Lam^{6,7,8}, Aldo P. Maggioni⁹, Sanjiv J. Shah¹⁰, and Burkert Pieske^{11,12,13,14}

¹Medical Devices and eHealth Clinical, Bayer AG, Wuppertal, Germany; ²Chrestos Concept GmbH & Co. KG, Essen, Germany; ³Clinical Development, Bayer AG, Wuppertal, Germany; ⁴Research and Clinical Sciences Statistics, Bayer AG, Berlin, Germany; ⁵Department of Medicine, University of Mississippi Medical Center, Jackson, MS, USA; ⁶National Heart Centre and Duke University–National University of Singapore, Singapore, Singapore; ⁷University Medical Centre Groningen, Groningen, the Netherlands; ⁸The George Institute for Global Health, Sydney, NSW, Australia; ⁹Italian Association of Hospital Cardiologists (ANMCO) Research Centre, Florence, Italy; ¹⁰Division of Cardiology, Department of Medicine, Northwestern University Feinberg School of Medicine, Chicago, IL, USA; ¹¹Department of Internal Medicine and Cardiology, Campus Virchow Klinikum, Charité Universitätsmedizin Berlin, Germany; ¹²German Heart Centre Berlin (DHZB), Berlin, Germany; ¹³Berlin Institute of Health (BIH), Berlin, Germany; and ¹⁴German Centre for Cardiovascular Research (DZHK), partner site Berlin, Berlin, Germany

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Aims	The effects of vericiguat vs. placebo on high-sensitivity C-reactive protein (hsCRP) and serum uric acid (SUA) were assessed in patients with heart failure with reduced ejection fraction (HFrEF) in the Phase 2 SOCRATES-REDUCED study (NCT01951625).
Methods and results	Changes from baseline hsCRP and SUA values at 12 weeks with placebo and vericiguat (1.25 mg, 2.5 mg, 5.0 mg and 10.0 mg, respectively) were assessed. The probability of achieving an hsCRP value of \leq 3.0 mg/L or SUA value of $<$ 7.0 mg/dL at week 12 was tested. Median baseline hsCRP and SUA levels were 3.68 mg/L [interquartile range (IQR) 1.41–8.41; $n = 335$] and 7.80 mg/dL (IQR 6.40–9.33; $n = 348$), respectively. Baseline-adjusted mean percentage changes in hsCRP were 0.2%, -19.5%, -24.3%, -25.7% and -31.9% in the placebo and vericiguat 1.25 mg, 2.5 mg, 5.0 mg and 10.0 mg groups, respectively; significance vs. placebo was observed in the vericiguat 10.0 mg group ($P = 0.035$). Baseline-adjusted mean percentage changes in SUA were 5.0%, -1.3%, -1.1%, -3.5% and -5.3% in the placebo, and vericiguat 1.25 mg, 2.5 mg, 5.0 mg and 10.0 mg groups ($P = 0.0202$ and $P = 0.004$, respectively). Estimated probability for an end-of-treatment hsCRP value of \leq 3.0 mg/L and SUA value of $<$ 7.0 mg/dL was higher with vericiguat compared with placebo. The effect was dose-dependent, with the greatest effect observed in the 10.0 mg group.
Conclusions	Vericiguat treatment for 12 weeks was associated with reductions in hsCRP and SUA, and a higher likelihood of achieving an hsCRP value of \leq 3.0 mg/L and SUA value of $<$ 7.0 mg/dL.
Keywords	Heart failure • Ventricular ejection fraction • Biomarker • C-reactive protein • Uric acid • Vericiguat

*Corresponding author. Medical Devices and eHealth Clinical, Bayer AG, Aprather Weg 18a, 42096 Wuppertal, Germany. Tel: +49 202 365680, Email: frank.kramer@bayer.com

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Inflammation and oxidative stress play key roles in the development and progression of heart failure (HF).^{1,2} High-sensitivity C-reactive protein (hsCRP) is an established marker of systemic inflammation and is associated with HF severity and outcomes.^{1,3–7} Uric acid, the end product of purine metabolism, circulates at increased serum levels in conditions of high oxidative stress and is frequently elevated in patients with HF, in which it correlates with HF severity and outcomes.^{6,8–10} Under conditions of oxidative stress, endothelial dysfunction creates a deficiency in nitric oxide (NO) and leads to impaired NO-soluble guanylate cyclase-cyclic guanosine monophosphate (NO-sGC-cGMP) signalling, and contributes, amongst other pathways, to the development of HF.^{11,12}

Vericiguat is a stimulator of sGC under investigation as a first-in-class therapy for worsening chronic HF with reduced ejection fraction (HFrEF).¹³ Although vericiguat did not meet the primary endpoint [change from baseline in log-transformed N-terminal pro-B-type natriuretic peptide (NT-proBNP) compared with placebo at week 12] in the Phase 2 dose-finding SOCRATES-REDUCED study (NCT01951625) in patients with HFrEF, exploratory analyses suggested a dose-response association between higher vericiguat dose and greater reductions in NT-proBNP compared with placebo.¹⁴ As vericiguat may address inflammation and oxidative stress via its action on the NO-sGC-cGMP pathway, we aimed to evaluate the effects of vericiguat on hsCRP and serum uric acid (SUA) (markers of inflammation and oxidative stress) in patients with HFrEF using data from the Phase 2 SOCRATES-REDUCED study.

Methods

Study design, treatment and assessments

SOCRATES-REDUCED (n = 456; protocol BAY 1021189/15371) was a randomized, placebo-controlled, double-blind, dose-finding, Phase 2 study of vericiguat in patients with HFrEF.¹⁴ In brief, the study population comprised patients who were post-hospitalization for HF or had received outpatient treatment with i.v. diuretics for HF. Patients were eligible for inclusion in the SOCRATES-REDUCED study if they had a left ventricular ejection fraction (LVEF) of <45% within 4 weeks of a symptomatic HF event. Patients were randomized to 12 weeks of treatment with vericiguat (1.25 mg, 2.5 mg, 5.0 mg and 10.0 mg once daily) or placebo after clinical stabilization or within 4 weeks after discharge. Dose up-titration (dose doubling) or sham titration occurred at week 2 (visit 2) and week 4 (visit 3) after randomization, with dose titration predicated on safety assessments and systolic blood pressure criteria.¹⁵ The study design was published¹⁵ and is summarized in supplementary material online Figure S1. The study conformed with the principles outlined in the Declaration of Helsinki.¹⁶ Institutional review board or ethics committee approval was obtained at each study site. All patients provided written informed consent.

Blood samples were collected at all visits and centrifuged at 4°C. Serum and ethylenediaminetetraacetic acid plasma were aliquoted and stored immediately at -20° C or lower at study sites until they were transported on dry ice to the central laboratory (Covance Central Laboratory Services, Geneva, Switzerland), where samples were stored at -70° C or lower until analysis.

High-sensitivity CRP was measured using an immunoturbidimetry assay (Roche Diagnostics, Indianapolis, IN, USA). Assays for SUA (Roche Diagnostics) were performed with the Roche Modular and Cobas Analyzer. Baseline data were collected at visit 1, before randomization and before the first intake of study treatment.

Other biomarkers investigated included bone alkaline phosphatase (bAP), cGMP, C-terminal telopeptide of type 1 collagen (CTX), galectin-3 (Gal-3), growth differentiation factor 15 (GDF-15), high-sensitivity troponin T (hsTnT), osteopontin (OPN), pro-collagen III peptide (PIIINP), soluble suppression of tumorigenicity 2 (sST2) and tissue metallopeptidase inhibitor 4 (TIMP-4).

Statistical analysis and model generation

The impacts of treatment in terms of biomarker changes from baseline to end of treatment were evaluated with a linear mixed model in order to account for intra-patient variability and differences in biomarker baseline levels. All analyses were based on log-transformed biomarker levels and the per-protocol population. Baseline-adjusted mean hsCRP and SUA changes in each vericiguat treatment arm were statistically compared with those in the placebo arm using two-sided *t*-tests on the model contrasts. Dose dependency was assessed by a linear trend test. The analyses for hsCRP and SUA were performed with and without additional clinical covariates (age group, sex, New York Heart Association functional class, diabetes and atrial fibrillation).

Building on this analysis, risk cut-offs of hsCRP and SUA established from prior studies were examined: hsCRP concentrations of <1.0 mg/L, 1.0–3.0 mg/L and >3.0 mg/L, respectively, are associated with low, moderate and high categories of relative risk for cardiovascular disease.¹⁷ Based on the linear mixed model, the expected end of treatment value for a given baseline value and treatment was calculated. The estimated probabilities of biomarker decreases to levels below the risk cut-off (hsCRP \leq 3.0 mg/L, SUA <7.0 mg/dL) during the course of the study were calculated for all treatment groups and selected baseline values within the observed range. These calculations were performed with consideration of the observed inter-patient variability and on the assumption of a log-normal distribution of the biomarkers.

The associations of hsCRP and SUA changes from baseline to end of treatment with clinical outcomes (cardiovascular death, cardiovascular hospitalization and emergency presentation caused by worsening chronic HF) were assessed. The relative odds (odds ratio) of a clinical outcome in each group (segmented by the direction of biomarker change from baseline to end of treatment) were calculated and Fisher's exact test was conducted.

Results

Patients

The SOCRATES-REDUCED study was conducted across Europe, North America and Asia between November 2013 and January 2015, with follow-up ending in June 2015. Randomized patients with no major protocol deviations and biomarker values at baseline and week 12 were included in the analysis (n = 328 and n = 345 in the hsCRP and SUA analyses, respectively) (supplementary material online Figure S2). Baseline characteristics for the full analysis set are presented in supplementary material online Table S1.

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Baseline characteristic	Total	hsCRP <1.0 mg/L n = 53	hsCRP 1.0-3.0 mg/L n = 92	hsCRP >3.0 mg/L n = 183	P-value ^a
	N = 328	(16.2%)	(28.0%)	(55.8%)	
Age, years, median (IQR)	68.0 (58.0–76.0)	69.0 (60.0-78.0)	70.0 (60.8–79.2)	66.0 (57.0–74.5)	0.035
BMI, kg/m ² , median (IQR)	27.6 (24.0-31.4)	26.5 (23.7-29.6)	27.1 (24.0-30.9)	28.2 (24.5-31.9)	0.041
Female sex, n	69 (21.0%)	15 (28.3%)	21 (22.8%)	33 (18.0%)	0.231
Non-White race, <i>n</i>	78 (23.8%)	15 (28.3%)	21 (22.8%)	42 (23.0%)	0.716
NYHA class III/IV, n	139 (42.4%)	16 (30.2%)	40 (43.5%)	83 (45.4%)	0.140
Atrial fibrillation, <i>n</i>	157 (47.9%)	29 (54.7%)	46 (50.0%)	82 (44.8%)	0.402
Arterial hypertension, n	254 (77.4%)	44 (83.0%)	65 (70.7%)	145 (79.2%)	0.163
Coronary artery disease, n	167 (50.9%)	25 (47.2%)	48 (52.2%)	94 (51.4%)	0.844
Chronic kidney disease, n	121 (36.9%)	20 (37.7%)	25 (27.2%)	76 (41.5%)	0.063
Diabetes mellitus, n	155 (47.3%)	21 (39.6%)	43 (46.7%)	91 (49.7%)	0.421
DBP, mmHg, median (IQR)	75.2 (69.6–81.7)	75.3 (69.0–79.7)	75.3 (68.3-82.0)	75.0 (70.0-83.0)	0.814
SBP, mmHg, median (IQR)	121.7 (115.6–134.4)	120.0 (114.7-128.7)	125.3 (117.0-135.5)	121.0 (115.0-136.5)	0.104
Heart rate, b.p.m., median (IQR)	70.5 (62.0-81.0)	69.3 (61.0-78.7)	71.0 (60.7–81.5)	70.7 (63.1–81.2)	0.655
LVEF, %, median (IQR)	29.2 (23.3-36.0)	29.8 (24.5-33.9)	31.2 (24.9-36.8)	28.5 (23.0-37.1)	0.350
Serum sodium, mmol/L, median (IQR)	139.0 (137.0–141.0)	139.0 (137.0–141.0)	139.5 (137.0–141.0)	139.0 (136.0–140.0)	0.194
Serum creatinine, mg/dL, median (IQR)	1.2 (1.0–1.4)	1.1 (1.0–1.3)	1.2 (1.0–1.4)	1.2 (1.0–1.5)	0.397
eGFR, mL/min/1.73 m ² , median (IQR)	56.3 (44.4-72.6)	55.3 (47.9-64.6)	56.3 (41.3-75.0)	56.3 (44.5-73.8)	0.689

Table 1 Baseline characteristics by high-sensitivity C-reactive protein subgroup

BMI, body mass index; DBP, diastolic blood pressure; eGFR, estimated glomerular filtration rate; hsCRP, high-sensitivity C-reactive protein; IQR, interquartile range; LVEF, left ventricular ejection fraction; NYHA, New York Heart Association; SBP, systolic blood pressure.

^aP-values are based on Fisher's exact tests for sex, race, NYHA class, atrial fibrillation, arterial hypertension, coronary artery disease, chronic kidney disease and diabetes mellitus, and on Kruskal–Wallis tests for other baseline characteristics.

Biomarker evaluation at baseline

Overall median hsCRP was 3.68 mg/L [interquartile range (IQR) 1.41-8.41 mg/L] at baseline. Within each treatment arm, baseline median hsCRP values were 3.98 mg/L (IQR 1.54-8.46 mg/L), 5.45 mg/L (IQR 1.78-10.75 mg/L), 3.62 mg/L (IQR 1.29-7.58 mg/L), 3.88 mg/L (IQR 1.29-8.60 mg/L) and 2.81 mg/L (IQR 1.37-6.62 mg/L) in the placebo, and vericiguat 1.25 mg, 2.5 mg, 5.0 mg and 10.0 mg groups, respectively.

Median SUA was 7.80 mg/dL (IQR 6.40-9.33 mg/dL) at baseline. Within each treatment arm, baseline median SUA values were 7.50 mg/dL (IQR 6.30-9.90 mg/dL), 7.40 mg/dL (IQR 6.10-8.70 mg/dL), 7.90 mg/dL (IQR 6.88-9.13 mg/dL), 8.25 mg/dL (IQR 6.63-9.80 mg/dL) and 7.95 mg/dL (IQR 6.55-9.38 mg/dL) in the placebo, and vericiguat 1.25 mg, 2.5 mg, 5.0 mg and 10.0 mg groups, respectively.

Patient demographics by baseline hsCRP and SUA subgroups are shown in *Tables 1* and 2, respectively. There were several differences between patients with lower and higher hsCRP and SUA values at baseline: patients with higher hsCRP and SUA values at baseline were of a slightly younger age and had a higher body mass index; patients with higher SUA at baseline had a lower LVEF, systolic blood pressure, serum sodium and estimated glomerular filtration rate (eGFR), and a higher serum creatinine level.

Biomarker evaluation following treatment

Baseline-adjusted mean hsCRP and SUA changes (%) from baseline to week 12 across the placebo and vericiguat dose

groups are shown in Figure 1. Following 12 weeks of treatment, baseline-adjusted mean percentage changes from baseline in hsCRP were 0.2% [95% confidence interval (Cl)] -22.4 to 29.3], -19.5% (95% Cl -37.8 to 4.1), -24.3% (95% Cl -41.1 to -2.7), -25.7% (95% Cl -42.6 to -3.7) and -31.9% (95% Cl -47.2 to -12.3) in the placebo, and vericiguat 1.25 mg, 2.5 mg, 5.0 mg and 10.0 mg groups, respectively (Figure 1A). Dose-dependent reductions in hsCRP were observed; the analysis indicated a significantly greater reduction in hsCRP with the vericiguat target dose of 10.0 mg relative to placebo (-31.9% vs. 0.2%; P = 0.035) and a significant linear trend in the reduction of hsCRP from baseline to end of treatment from placebo up to the highest vericiguat dose of 10.0 mg (P = 0.039) (Table 3).

After 12 weeks of treatment, baseline-adjusted mean percentage changes from baseline in SUA were 5.0% (95% Cl -0.1 to 10.4), -1.3% (95% Cl -6.1 to 3.7), -1.1% (95% Cl -5.8 to 3.8), -3.5% (95% Cl -8.3 to 1.6) and -5.3% (95% Cl -9.8 to -0.6) in the placebo, and vericiguat 1.25 mg, 2.5 mg, 5.0 mg and 10.0 mg groups, respectively (*Figure 1B*). Significant reductions from baseline in SUA were observed in the vericiguat 5.0 mg and 10 mg groups relative to placebo [-3.5% vs. 5.0% (P = 0.02) and -5.3% vs. 5.0% (P = 0.004), respectively] and a significant linear trend in the reduction of SUA from baseline to end of treatment from placebo up to the highest vericiguat dose of 10.0 mg (P = 0.004) was apparent (*Table 3*), indicating a dose-dependent effect.

Additional consideration of clinical covariates in the analysis did not lead to noteworthy changes in the treatment effects (e.g. effect ratios of 0.66 vs. 0.68 for hsCRP and 0.90 vs. 0.90 for SUA in comparisons of 10.0 mg vericiguat and placebo in the model with and without clinical covariates).

Baseline characteristic	Total	SUA <7.0 mg/dL n = 116	SUA 7.0–10.0 mg/dL n = 159	SUA >10.0 mg/dL n = 70	P-value ^a
	N = 345	(33.6%)	(46.1%)	(20.3%)	
Age, years, median (IQR)	68.0 (59.0–77.0)	71.0 (62.0–79.0)	67.0 (59.0–75.0)	63.0 (54.0-73.0)	0.001
BMI, kg/m ² , median (IQR)	27.7 (24.3–31.5)	26.8 (23.2-30.2)	27.6 (24.8–31.4)	29.1 (26.2–33.6)	0.004
Female sex, n	76 (22.0%)	29 (25.0%)	32 (20.1%)	15 (21.4%)	0.637
Non-White race, <i>n</i>	76 (22.0%)	24 (20.7%)	32 (20.1%)	20 (28.6%)	0.344
NYHA class III/IV, n	152 (44.1%)	52 (44.8%)	67 (42.1%)	33 (47.1%)	0.761
Atrial fibrillation, <i>n</i>	167 (48.4%)	56 (48.3%)	78 (49.1%)	33 (47.1%)	0.970
Arterial hypertension, n	268 (77.7%)	89 (76.7%)	124 (78.0%)	55 (78.6%)	0.956
Coronary artery disease, n	174 (50.4%)	57 (49.1%)	81 (50.9%)	36 (51.4%)	0.950
Chronic kidney disease, n	128 (37.1%)	49 (42.2%)	52 (32.7%)	27 (38.6%)	0.267
Diabetes mellitus, n	168 (48.7%)	55 (47.4%)	80 (50.3%)	33 (47.1%)	0.874
DBP, mmHg, median (IQR)	75.0 (69.7–81.3)	74.8 (69.7–79.2)	75.3 (69.0-81.5)	76.3 (70.2–83.2)	0.480
SBP, mmHg, median (IQR)	121.3 (115.0–133.7)	125.3 (116.7–140.7)	120.7 (115.0–131.7)	119.3 (113.9–127.8)	0.010
Heart rate, b.p.m., median (IQR)	70.7 (62.3–81.0)	70.5 (61.7–76.8)	70.0 (62.5-81.0)	74.5 (64.7–85.3)	0.144
LVEF, %, median (IQR)	29.0 (23.2–35.8)	30.2 (25.6–37.6)	29.6 (23.0-36.0)	25.8 (21.8–33.1)	0.005
Serum sodium, mmol/L, median (IQR)	139.0 (137.0–141.0)	140.0 (138.0–141.0)	139.0 (136.0–140.5)	138.0 (136.0–140.0)	0.006
Serum creatinine, mg/dL, median (IQR)	1.2 (1.0–1.4)	1.1 (0.9–1.4)	1.2 (1.0–1.4)	1.4 (1.1–1.7)	<0.001
eGFR, mL/min/1.73 m ² , median (IQR)	56.3 (44.9–72.7)	60.2 (45.8-80.8)	59.2 (48.2–70.9)	47.2 (38.2–59.1)	<0.001

Table 2 Baseline characteristics by serum uric acid subgroup

BMI, body mass index; DBP, diastolic blood pressure; eGFR, estimated glomerular filtration rate; IQR, interquartile range; LVEF, left ventricular ejection fraction; NYHA, New York Heart Association; SBP, systolic blood pressure; SUA, serum uric acid.

^aP-values are based on Fisher's exact tests for sex, race, NYHA class, atrial fibrillation, arterial hypertension, coronary artery disease, chronic kidney disease and diabetes mellitus, and on Kruskal–Wallis tests for other baseline characteristics.

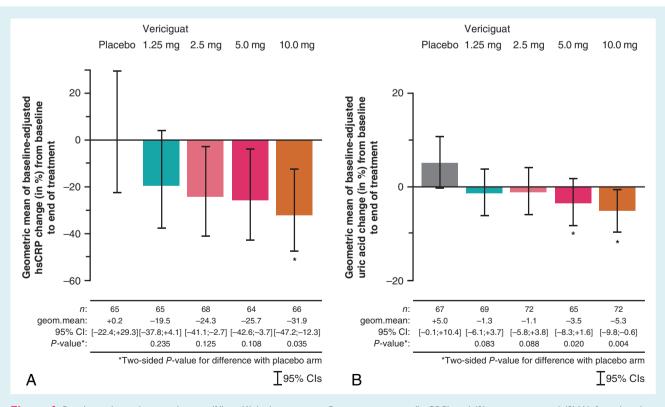


Figure 1 Baseline-adjusted mean changes (%) in (A) high-sensitivity C-reactive protein (hsCRP) and (B) serum uric acid (SUA) from baseline to end of treatment. Cl, confidence interval; geom.mean, geometric mean. *Statistical significance relative to placebo.

	hsCRP changes				SUA changes			
	Adjusted for hsCRP baseline level (log-transformed)	RP baseline rmed)	Adjusted for hsCRP baseline level (log-transformed) and clinical covariates	RP baseline med) and	Adjusted for SUA baseline level (log-transformed)	\ baseline rmed)	Adjusted for SUA baseline level (log-transformed) and clinical covariates	baseline med) and
	Effect ratio ^a (95% Cl)	P-value	Effect ratio ^a (95% CI)	P-value	Effect ratio ^a (95% CI)	P-value	Effect ratio ^a (95% CI)	P-value
Treatment groups								
Vericiguat 1.25 mg vs. placebo	0.80 (0.56–1.15)	0.235	0.84 (0.59–1.22)	0.367	0.94 (0.88–1.01)	0.083	0.94 (0.88–1.01)	0.081
Vericiguat 2.5 mg vs. placebo	0.76 (0.53–1.08)	0.125	0.71 (0.50–1.03)	0.069	0.94 (0.88–1.01)	0.088	0.95 (0.88–1.01)	0.114
Vericiguat 5.0 mg vs. placebo	0.74 (0.52-1.07)	0.108	0.69 (0.48–1.00)	0.049	0.92 (0.86–0.99)	0.020	0.91 (0.85–0.98)	0.010
Vericiguat 10.0 mg vs. placebo	0.68 (0.47–0.97)	0.035	0.66 (0.46–0.96)	0.024	0.90 (0.84–0.97)	0.004	0.90 (0.84–0.97)	0.004
Linear trend test		0.039		0.013		0.004		0.003

Estimated probability of achieving biomarker thresholds at end of treatment

The estimated probability for an end-of-treatment (week 12) hsCRP value of <3.0 mg/L and SUA value of <7.0 mg/dL was higher in all vericiguat treatment groups compared with placebo (Figure 2). The estimated probability for an end-of-treatment hsCRP value of \leq 3.0 mg/L and SUA value of <7.0 mg/dL varied by treatment and baseline biomarker level. The estimated probability differences between all vericiguat treatment groups and placebo for an end-of-treatment hsCRP value of \leq 3.0 mg/L were comparable across baseline hsCRP values (Figure 2A). The estimated probability increase (shift on y-axis) from placebo to vericiguat for an end-of-treatment hsCRP value of \leq 3.0 mg/L ranged from 8.3 to 14.6 percentage points in patients with a baseline hsCRP value of 4 mg/L (left dashed vertical line) and from 6.5 to 12.1 percentage points in patients with a baseline hsCRP value of 10 mg/L (right dashed vertical line) in the vericiguat 1.25 mg and 10.0 mg groups, respectively.

The SUA model indicated that the estimated probability difference between all vericiguat treatment groups and placebo for an end-of-treatment SUA value of <7.0 mg/dL was decreased in patients with higher baseline SUA levels (Figure 2B). The estimated probability increase from placebo to vericiguat for an end-of-treatment SUA value of <7.0 mg/L ranged from 9.1 to 16.5 percentage points in patients with a baseline SUA value of 8.0 mg/dL and from 1.9 to 4.0 percentage points in patients with a baseline SUA value of 11.0 mg/dL across the vericiguat treatment groups.

The distributions of the hsCRP and SUA subgroups in the placebo and 2.5-10.0 mg vericiguat treatment arms at baseline and end of treatment are shown in Figure 2C and 2D. At baseline, 11, 22 and 33 patients in the 10.0 mg vericiguat treatment arm had hsCRP values of <1.0 mg/L, 1.0-3.0 mg/L and >3.0 mg/L, respectively (Figure 2C). Following 12 weeks of treatment with vericiguat, 20, 19 and 27 patients had hsCRP values of <1.0 mg/L, 1.0-3.0 mg/L and >3.0 mg/L, respectively. Changes in the distribution of the hsCRP subgroups were also observed in the placebo treatment group, with fewer patients having an hsCRP value of <1.0 mg/L at week 12 than at baseline. These changes reflect a near doubling of patients (from 11 to 20) with hsCRP values of <1.0 mg/L by week 12 in the 10.0 mg vericiguat treatment group.

At baseline, 24, 32 and 16 patients in the 10.0 mg vericiguat treatment arm had SUA values of <7.0 mg/dL, 7.0-10.0 mg/dL and >10.0 mg/dL, respectively (Figure 2D). At the end of vericiguat treatment, 29, 30 and 13 patients had SUA values of <7.0 mg/dL, 7.0-10.0 mg/dL and >10.0 mg/dL, respectively. The distribution of SUA subgroups in the placebo treatment arm was largely unchanged throughout the study.

Association of biomarker changes from baseline with clinical outcomes

The proportion of clinical outcome events (cardiovascular death, cardiovascular hospitalization and emergency presentation caused

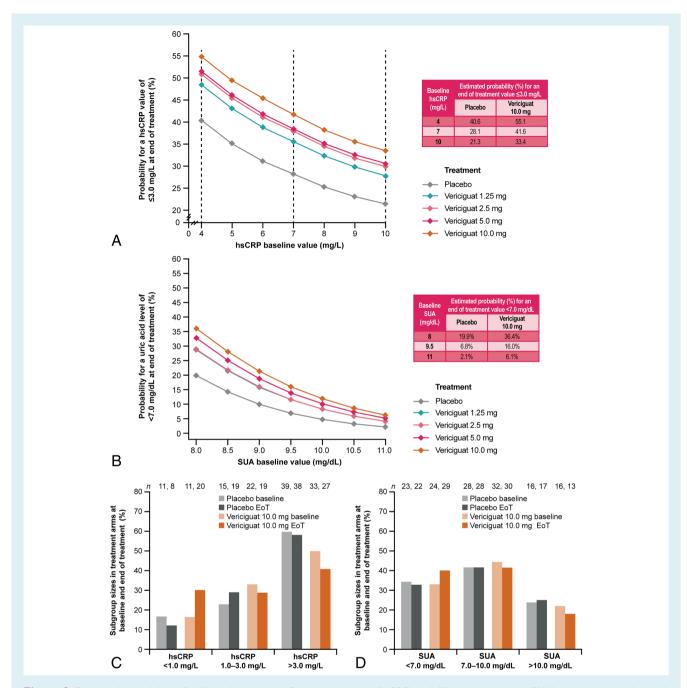


Figure 2 Estimated probabilities for (A) high-sensitivity C-reactive protein (hsCRP) and (B) serum uric acid (SUA) reduction to below risk level and (C, D) subgroup sizes per treatment arm at baseline and end of treatment (EoT). In (B), 1.25 mg and 2.5 mg vericiguat trend lines overlap SUA data. Numbers within the bar charts represent the numbers of patients with specified (C) hsCRP or (D) SUA values at baseline or EoT. The total numbers of patients per treatment arm for placebo and vericiguat 10.0 mg were 65 and 66, respectively, for hsCRP data, and 67 and 72, respectively, for SUA data.

by worsening chronic HF) was numerically smaller in patients with decreases in hsCRP and SUA than in those with increases from baseline (*Table 4*). Clinical outcome events occurred in 14.0% of patients who had a decrease in hsCRP compared with 21.8% who had an increase in hsCRP, relative to baseline, at the end of treatment (P = 0.08). Similarly, clinical outcome events occurred in 14.8% of patients who had a decrease in SUA compared with

19.9% who had an increase in SUA, relative to baseline, at the end of treatment (P = 0.26).

Other exploratory biomarkers

Geometric means at baseline and summary statistics of percentage changes from baseline in other exploratory biomarkers

Direction of biomarker change from baseline to end of treatment	n	Patients with an event, <i>n</i>	Event proportion, %	Odds ratio	95% CI	P-value
hsCRP						
Increase	142	31	21.8%	0.58	0.31-1.08	0.08
Decrease	186	26	14.0%			
SUA						
Increase	176	35	19.9%	0.70	0.38-1.27	0.26
Decrease	169	25	14.8%			

Table 4 Association of changes in high-sensitivity C-reactive protein and serum uric acid with clinical outcomes

CI, confidence interval; hsCRP, high-sensitivity C-reactive protein; SUA, serum uric acid.

(bAP, cGMP, CTX, Gal-3, GDF-15, hsTnT, OPN, PIIINP, sST2 and TIMP-4) are presented in supplementary material online *Table S2*. There was no effect of treatment with vericiguat compared with placebo on these biomarkers at week 12 (*P*-value of linear trend test >0.05).

Discussion

Vericiguat treatment for 12 weeks was associated with a decrease in hsCRP (significant in the vericiguat 10.0 mg group and a significant dose-dependent trend) and SUA concentrations (significant in the vericiguat 5.0 mg and 10.0 mg groups and a significant dose-dependent trend) from baseline in patients with HFrEF. Statistically significant dose-dependent reductions in hsCRP and SUA were observed in vericiguat-treated patients with HFrEF when data were adjusted for baseline levels and clinical covariates.

Vericiguat treatment was also associated with a higher probability of an hsCRP value of \leq 3.0 mg/L at the end of the study than placebo. The distribution of hsCRP subgroups in the 10.0 mg vericiguat treatment arm revealed that the increase in patients with an end-of-treatment hsCRP value of <1.0 mg/L corresponded with fewer patients having an hsCRP value of \geq 1.0 mg/L at week 12. These novel results point to a potential anti-inflammatory effect of sGC stimulation in patients with HFrEF after a recent worsening HF event who are post-hospitalization for HF or have required outpatient treatment with i.v. diuretics.

The mechanism underlying the lowering of hsCRP by vericiguat in patients with HFrEF is not directly addressed by these data. The anti-inflammatory capacity of sGC stimulation has been shown in a murine model of interleukin-1 β -induced leucocyte rolling and adhesion, in which sGC stimulation with BAY 41–2272 (a predecessor to the sGC activator) down-regulated P-selectin expression and inhibited leucocyte recruitment.¹⁸ However, studies using tumour necrosis factor inhibitors demonstrated no improvement in HF-related outcomes.⁴ Similarly, although statin treatment in the CORONA study was associated with a reduction in CRP, the study did not meet its primary endpoint.⁴ It remains to be determined whether hsCRP is a true prognostic biomarker in relation to clinical outcomes.

Systemic inflammation in HF has previously been postulated to result from an impaired intestinal mucosal barrier function secondary to mesenteric venous congestion in right ventricular (RV) HF. The intestinal damage leads to translocation of lipopolysaccharides from the intestinal lumen into the circulation, and endotoxaemia subsequently causes production of proinflammatory cytokines.¹⁹⁻²³ Thus, a decrease in systemic inflammatory markers could be hypothesized to reflect decreased peripheral venous congestion as a potential consequence of improved RV function and decreased right heart filling pressures. Reduced congestion may also result in less hepatic congestion,²⁴ which may, in turn, result in lower levels of inflammation and oxidative stress.²⁵ However, observed correlations between changes in hsCRP and SUA and changes in echocardiographic parameters following 12 weeks of treatment were only very small and not supportive of an imaging correlate reflecting lower RV filling pressures as an underlying mechanism (data not shown). In patients who experienced a decrease from baseline in hsCRP or SUA at week 12, the relative odds of a clinical outcome were non-significantly reduced compared with patients who had an increase from baseline in hsCRP or SUA at end of treatment. The collection of biomarkers from the Phase 3 vericiguat study VICTORIA (NCT02861534) in patients with HFrEF will provide an opportunity to assess clinical laboratory markers of haemodynamic right heart unloading in parallel with echocardiography in an imaging ancillary study as potential determinants of reduced systemic inflammation in response to treatment with vericiguat.¹³ In addition, the longer duration and increased number of clinical events in the VICTORIA study will enable a more robust assessment of the associations between inflammatory biomarkers and clinical outcomes.

Treatment with vericiguat in patients with HFrEF was associated with a higher likelihood of SUA levels falling below the threshold of hyperuricaemia (SUA \geq 7.0 mg/dL) at the end of the study. The magnitude of SUA reduction observed with vericiguat treatment in this analysis is comparable with that achieved by sacubitril/valsartan treatment in the PARADIGM-HF study.¹⁰ Hyperuricaemic levels are associated with increased risk for cardiovascular death, hospitalization for HF and all-cause mortality.^{26–28} The observed reduction in uric acid levels following treatment with vericiguat may also represent a potential protective mechanism to prevent the deterioration of kidney function in HF and ameliorate cardiorenal syndrome. Although they are not strongly correlated with changes in eGFR (Spearman correlation -0.363 (95% CI -0.452 to -0.267) (supplementary material online *Table* S3), the reductions from baseline in SUA reported here (geometric mean: <0.5 mg/dL) are smaller than those reported in two dedicated SUA-lowering trials in patients with HF (reductions of 2 mg/dL and 4.2 mg/dL), which found no improvements in survival, hospitalization for HF or LVEF.^{29,30} Thus, although effectively reducing SUA, the inconsistent clinical efficacy of xanthine oxidase inhibitors in these studies called into question the promise of SUA reduction to improve clinical outcomes. This contrasts with the broad evidence base in support of the pathophysiological relevance of an increase in SUA as a risk factor for adverse outcomes in both HF and chronic kidney disease. The SUA-reducing effect of vericiguat may support the hypothesis that mechanisms other than the inhibition of SUA generation via xanthine oxidase may have promise for tackling this risk factor.

The analysis of baseline characteristics by biomarker subgroup revealed that the baseline demographic characteristics of younger age and a higher body mass index were associated with higher hsCRP and SUA values. Higher SUA but not higher hsCRP were found at baseline in patients with worse renal function, more severely reduced LVEF and lower serum sodium. With the exception of serum sodium, these findings are consistent with those reported for SUA in the PARADIGM-HF study¹⁰ and suggest that the clinical phenotype associated with higher hsCRP and SUA is consistent with more advanced HF.

There was no effect of treatment with vericiguat compared with placebo on other exploratory biomarkers associated with HF (bAP, cGMP, CTX, Gal-3, GDF-15, hsTnT, OPN, PIIINP, sST2, TIMP-4) at week 12; these results are consistent with those reported in the full analysis set.¹⁴

One limitation of this analysis is that the mechanisms behind the effects of vericiguat on hsCRP and SUA are unknown and further studies are warranted to validate the prognostic value of reducing SUA as a treatment target in therapeutic approaches in HFrEF. Given the exploratory post hoc nature of the analysis, these results should be considered as hypothesis-generating. It should also be considered that elevated hsCRP levels at baseline may reflect the presence of aetiologies other than HF, such as metabolic conditions including insulin resistance.⁷ The analysis of SUA changes from baseline to week 12 did not take into consideration the concomitant use of SUA-altering therapies such as diuretics. Additionally, this study was not powered for outcomes and the reported associations between biomarkers and adverse clinical outcomes should be interpreted with caution.

In conclusion, dose-dependent decreases in hsCRP and SUA concentrations from baseline in patients with HFrEF were observed following 12 weeks of treatment with vericiguat. Whether the reductions in hsCRP and SUA with vericiguat are sustained with longer periods of treatment and are associated with improved clinical outcomes remains to be demonstrated in the Phase 3 VICTORIA study (NCT02861534).¹³

Supplementary Information

Additional supporting information may be found online in the Supporting Information section at the end of the article. **Table S1.** Baseline characteristics of patients in the full analysis set and biomarker analysis sets.

Table S2. Baseline levels and changes from baseline to end of treatment of additional biomarkers (per-protocol population).

Table S3. Correlation analysis of changes in estimated glomerular filtration rate and body mass index with changes in log-transformed high-sensitivity C-reactive protein and serum uric acid levels from baseline to end of treatment.

Figure S1. SOCRATES-REDUCED study design.

Figure S2. Patient disposition in the SOCRATES-REDUCED study.

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