

Plasminogen activator inhibitor-1 is elevated in patients with COPD independent of metabolic and cardiovascular function

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Introduction: Plasminogen activator inhibitor-1 (PAI-1), a major inhibitor of fibrinolysis, is associated with thrombosis, obesity, insulin resistance, dyslipidemia, and premature aging, which all are coexisting conditions of chronic obstructive pulmonary disease (COPD). The role of PAI-1 in COPD with respect to metabolic and cardiovascular functions is unclear.

Methods: In this study, which was nested within a prospective cohort study, the serum levels of PAI-1 were cross-sectionally measured in 74 stable COPD patients (Global Initiative for Chronic Obstructive Lung Disease [GOLD] Stages I–IV) and 18 controls without lung disease. In addition, triglycerides, high-density lipoprotein cholesterol, fasting plasma glucose, waist circumference, blood pressure, smoking status, high-sensitive C-reactive protein (hs-CRP), adiponectin, ankle-brachial index, N-terminal pro-B-type natriuretic peptide, and history of comorbidities were also determined.

Results: The serum levels of PAI-1 were significantly higher in COPD patients than in controls, independent of a broad spectrum of possible confounders including metabolic and cardiovascular dysfunction. A multivariate regression analysis revealed triglyceride and hs-CRP levels to be the best predictors of PAI-1 within COPD. GOLD Stages II and III remained independently associated with higher PAI-1 levels in a final regression analysis.

Conclusion: The data from the present study showed that the serum levels of PAI-1 are higher in patients with COPD and that moderate-to-severe airflow limitation, hypertriglyceridemia, and systemic inflammation are independent predictors of an elevated PAI-1 level. PAI-1 may be a potential biomarker candidate for COPD-specific and extra-pulmonary manifestations.

Keywords: plasminogen activator inhibitor-1, chronic obstructive pulmonary disease, cardiovascular disease, inflammation, hypertriglyceridemia, ankle-brachial index, N-terminal pro-B-type natriuretic peptide

Introduction

The coexistence of cardiovascular diseases (CVDs) and metabolic disorders is frequent in patients with chronic obstructive pulmonary disease (COPD).¹ The prevalence of both these conditions is highest in patients with moderate-to-severe COPD in particular and drops in patients with very severe COPD.² Accordingly, CVD is a leading cause of death in the earlier stage of the disease.³ Thrombosis, in turn, is the most frequent underlying mechanism of the three major CVDs: ischemic heart disease, stroke, and venous thromboembolism,⁴ which raises the question whether blood markers involved in the pathogenesis of thrombosis are linked to COPD.

Plasminogen activator inhibitor-1 (PAI-1) is a member of the superfamily of serine protease inhibitors and the principal inhibitor of fibrinolysis within the plasminogen

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activator system.^{5,6} High concentration of PAI-1 plays a pivotal role in the pathogenesis of arterial and venous thrombosis and, therefore, precedes the occurrence of thrombotic events.^{7,8} In addition, higher level of PAI-1 is associated with obesity, insulin resistance, diabetes, hyperlipidemia, and premature aging.^{5,9,10} All these conditions are prevalent in COPD,^{11,12} and a potential influence of PAI-1 on an enhanced thrombogenesis in COPD has been hypothesized.¹³ Indeed, a recent case-control study demonstrated that PAI-1 polymorphisms related to a higher PAI-1 expression are associated with COPD.¹⁴ Furthermore, a study conducted on patients with predominantly mild-to-moderate COPD found that higher serum PAI-1 levels are related to airflow limitation.¹⁵ Whether the observed elevation in PAI-1 levels in COPD is independent of metabolic and cardiovascular functions is unknown.

In order to elucidate the clinical value of PAI-1 as a potential biomarker in COPD, the serum levels of PAI-1 and a broad spectrum of potential confounding factors for PAI-1 were analyzed in patients with mild-to-very severe COPD and in control subjects.

Methods

This cross-sectional study is nested within a prospective COPD cohort study conducted at the Pulmonary Research Institute, the LungenClinic Grosshansdorf (Grosshansdorf, Germany). Details regarding the COPD population and the methodology of the study have been published elsewhere.^{16–18} In the present study, 74 stable outpatients with mild-to-very severe COPD (ie, $n=16, 20, 18,$ and 20 with Global Initiative for Chronic Obstructive Lung Disease [GOLD] Stages I, II, III, and IV, respectively) and 18 nonsmoking controls without any lung disease from the 3-year follow-up visit were included. The study was approved by the Ethics Committee of the Medical Association of Schleswig-Holstein, Bad Segeberg, Germany (III/EK 116/05[I]; 185/08[I]), and all the participants provided written informed consent. In this study, the serum levels of PAI-1 were analyzed by using enzyme-linked immunosorbent assay (ELISA; Human Serpin E1/PAI-1 DuoSet ELISA; R&D Systems, Wiesbaden-Nordenstadt, Germany; detection range =0.312–20 ng/mL); metabolic and cardiovascular functions as possible confounders of PAI-1, including triglyceride levels, high-density lipoprotein (HDL) cholesterol, fasting plasma glucose, waist circumference, blood pressure, the presence of the metabolic syndrome (based on the latter five variables according to the criteria of the International Diabetes Federation¹⁹), smoking status, and history of coronary artery disease and diabetes, were determined; high-sensitivity C-reactive protein (hs-CRP) was measured as an established marker of

systemic inflammation and adiponectin, which is involved in anti-inflammatory, anti-diabetic, and anti-atherogenic processes;²⁰ in addition, global cardiac function by serum levels of N-terminal pro-B-type natriuretic peptide (NT-proBNP) and the presence of atherosclerosis by the ankle-brachial index (ABI) were also determined as previously described.^{16,18}

First, the differences between groups were analyzed univariately by using unpaired *t*-test, χ^2 test, and analysis of variance (ANOVA) with least significant difference post hoc analysis. Then, a multivariate linear regression analysis was conducted with PAI-1 as the dependent variable and COPD as a predictor, and adjustments for possible confounders, such as age, sex, body mass index (BMI), impaired fasting glucose (ie, ≥ 100 mg/dL),¹⁹ dyslipidemia (ie, triglycerides > 150 mg/dL or HDL < 40 mg/dL in male and < 50 mg/dL in female),¹⁹ cardiac dysfunction (ie, NT-proBNP > 125 pg/mL),²¹ atherosclerosis (ie, ABI ≤ 0.90),²² as well as history of diabetes, hypertension, and coronary artery disease, were performed. A second multivariate regression analysis with PAI-1 as the dependent variable aimed at evaluating the best independent predictors for PAI-1 within COPD. Therefore, a stepwise approach was chosen with a backward elimination. Only the variables that showed at least a trend for statistical significance on the bivariate level, that is, $P < 0.10$, were included in the model. Next, whether COPD severity stages were still independently associated with PAI-1 within the final model was proved. Last, the interaction effects of GOLD stages with the remaining predictors of the final model were tested by using separate two-way ANOVAs.

Results

Impaired fasting glucose, dyslipidemia, cardiac dysfunction, hypertension, and history of diabetes ($P > 0.05$, respectively; Table 1) did not differ between the patients with COPD and controls. The frequency of atherosclerosis and history of coronary artery disease increased in patients with COPD compared with controls ($P < 0.05$). Serum levels of PAI-1 were significantly higher in COPD patients than in controls (Figure 1A). Stratified by severity stages, PAI-1 levels were highest in GOLD Stages II and III (Figure 1B). Smoking status and statin use had no effect on the level of PAI-1 in patients with COPD ($P=0.51$ and 0.50 , respectively). In a multivariate linear regression analysis, the presence of COPD remained an independent predictor for higher PAI-1 levels after adjustments for age, sex, BMI, impaired fasting glucose, dyslipidemia, cardiac dysfunction, hypertension, diabetes, atherosclerosis, and history of coronary artery disease (regression coefficient *B* for COPD patients versus

Table 1 Characteristics of patients with COPD and controls without any lung disease

	Controls (n=18)	COPD patients (n=74)	P-value
Age, years, mean (SD)	65.9 (5.8)	66.0 (6.6)	0.99
Men, n (%)	11 (61)	52 (70)	0.45
Current smokers, n (%)	0 (0)	24 (32)	0.005
Pack-years of smoking, mean (SD)	0.5 (1)	55 (26)	<0.001
Body mass index, kg/m ² , mean (SD)	25.7 (4.6)	26.0 (5.3)	0.82
FEV ₁ %pred, mean (SD)	116.2 (11.6)	55.2 (25.6)	<0.001
FEV ₁ /FVC, mean (SD)	78.6 (3.3)	51.0 (15.0)	<0.001
History of diabetes mellitus, n (%)	0 (0)	7 (10)	0.18
Impaired fasting glucose, n (%)	4 (22)	32 (43)	0.10
Dyslipidemia, n (%)	6 (33)	19 (26)	0.35
Waist circumference, >94/80 cm (M/F), n (%)	11 (61)	59 (80)	0.10
Cardiac dysfunction, n (%)	3 (17)	27 (37)	0.11
Atherosclerosis, n (%)	0 (0)	21 (29)	0.009
History of coronary artery disease, n (%)	0 (0)	14 (19)	0.045
Hypertension, n (%)	12 (67)	57 (77)	0.36
hs-CRP, mg/L, median (IQR)	1.1 (0.6–2.0)	3.0 (1.5–6.1)	<0.001
Adiponectin, ng/mL, median (IQR)	6,245 (4,219–8,147)	8,038 (6,219–11,651)	0.025

Notes: Dyslipidemia = triglycerides ≥ 150 mg/dL or high-density lipoprotein < 40 mg/dL in men and < 50 mg/dL in women; impaired fasting glucose = fasting plasma glucose ≥ 100 mg/dL or treatment; cardiac dysfunction = NT-proBNP > 125 pg/mL; atherosclerosis = ankle-brachial index ≤ 0.90 . Differences of pack-years, FEV₁, and FEV₁/FVC were tested by using rank test, as the homogeneity of variances was not given. hs-CRP and adiponectin were log-transformed to reveal a normal distribution.

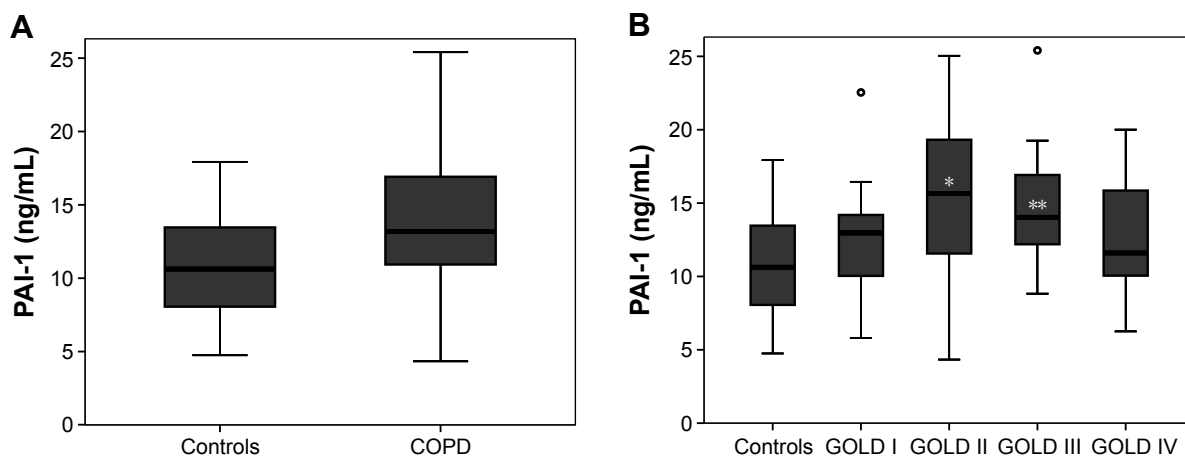
Abbreviations: COPD, chronic obstructive pulmonary disease; FEV₁, forced expiratory volume in 1 s; FVC, forced vital capacity; hs-CRP, high-sensitive C-reactive protein; IQR, interquartile range; M/F, male/female; NT-proBNP, N-terminal pro-B-type natriuretic peptide; SD, standard deviation.

controls =2.9, 95% confidence interval [CI] =0.34–5.41, $P=0.027$).

A next step was aimed at identifying the predictors for PAI-1 within COPD. Significant correlations of PAI-1 with HDL cholesterol, triglycerides, hs-CRP, and adiponectin ($P<0.05$, respectively; Table 2) were found on the bivariate level. The association between age and PAI-1 showed a trend for significance ($P=0.058$). In a multivariate regression analysis with a backward elimination, log-triglycerides (3.3, 95% CI =0.8–5.9, $P=0.012$) and log-hs-CRP (0.83,

95% CI =0.03–1.63, $P=0.043$) were found to be the best independent predictors for PAI-1 in COPD. Figure 2 visualizes the results from the multivariate regression analysis, that is, mean PAI-1 levels in COPD patients stratified according to triglycerides and hs-CRP in comparison with controls.

In a final model we studied whether COPD severity stages add additional information as predictors for PAI-1 independent of triglycerides and hs-CRP within the total study cohort. It was found that GOLD Stages II and III were independently associated with PAI-1 (Table 3). There were

**Figure 1** Serum levels of PAI-1 (A) in patients with COPD and controls without any lung disease ($P=0.015$) and (B) according to COPD severity.

Notes: After adjustment for a broad spectrum of possible confounders, PAI-1 still differed significantly between COPD patients and controls ($P=0.027$). After stratifying according to severity stages, PAI-1 was highest in GOLD II followed by GOLD III (* $P=0.003$ and ** $P=0.011$ for the comparison with controls). Lines represent the median, boxes represent the IQR, and circles represent outliers.

Abbreviations: COPD, chronic obstructive pulmonary disease; GOLD, Global Initiative for Chronic Obstructive Lung Disease; IQR, interquartile range; PAI-1, plasminogen activator inhibitor-1.

Table 2 Bivariate associations of PAI-1 with lung function, metabolic, cardiac, and inflammatory markers in patients with COPD

	R	P-value
Age, years	-0.22	0.058
Body mass index, kg/m ²	0.01	0.93
Waist circumference, cm	-0.01	0.96
FEV ₁ %predicted	-0.07	0.55
Log-triglycerides, mg/dL	0.31	0.007
High-density lipoprotein cholesterol, mg/dL	-0.23	0.044
Fasting plasma glucose, mg/dL	-0.16	0.19
Ankle-brachial index	-0.02	0.36
Log-NT-proBNP, pg/mL	0.05	0.68
Log-hs-CRP, mg/L	0.25	0.035
Log-adiponectin, ng/mL	-0.23	0.048

Notes: Bivariate associations were analyzed by using Pearson correlations. NT-proBNP, triglycerides, hs-CRP, and adiponectin were log-transformed to obtain a normal distribution.

Abbreviations: COPD, chronic obstructive pulmonary disease; FEV₁, forced expiratory volume in 1 s; hs-CRP, high-sensitive C-reactive protein; NT-proBNP, N-terminal pro-B-type natriuretic peptide; PAI-1, plasminogen activator inhibitor-1.

no significant interaction effects of GOLD stages either on triglycerides or on hs-CRP.

Discussion

The main finding of the present study was that serum levels of PAI-1 are higher in patients with COPD independent of metabolic and cardiovascular disorders. Furthermore, hypertriglyceridemia and systemic inflammation were found to be the best independent predictors for higher PAI-1 levels in COPD patients, whereas the presence of moderate-to-severe airflow limitation remained to be independently associated with PAI-1.

PAI-1 is a multifunctional protein associated with thrombosis as well as with metabolic conditions, such as obesity, dyslipidemia, insulin resistance, and diabetes,^{5,7,9,10} which all are important concomitant conditions of COPD.^{1,12} There are two very recent studies showing an association between elevated PAI-1 levels and COPD, but considered neither comorbidities of COPD nor the whole spectrum of airflow limitation, from mild to very severe stage.^{14,15} The present study is the first to demonstrate that higher serum PAI-1 levels in COPD are independent of a broad spectrum of possible confounders, in particular established variables indicating metabolic and cardiovascular dysfunction. Interestingly, this study found the highest PAI-1 levels in patients with GOLD Stages II and III. This observation is in good agreement with the finding of the previous study showing a significant association between higher levels of PAI-1 and airflow limitation in patients with predominantly mild-to-moderate COPD.¹⁵ Moreover, it was also found that triglycerides and hs-CRP are the best predictors for elevated PAI-1 levels in COPD patients. In general, an elevation in triglycerides and hs-CRP plays an important role for CVD.^{23,24} Moreover, an association of PAI-1 levels with serum lipids and inflammatory markers has been found,^{9,25} although the mechanistic link behind this remains not fully understood.^{5,26} In patients with COPD, hypertriglyceridemia and systemic inflammation are also found frequently and predict mortality.²⁷⁻²⁹ The present study demonstrated an independent association of high PAI-1 serum levels with moderate-to-severe airflow limitation, hypertriglyceridemia, and systemic inflammation.

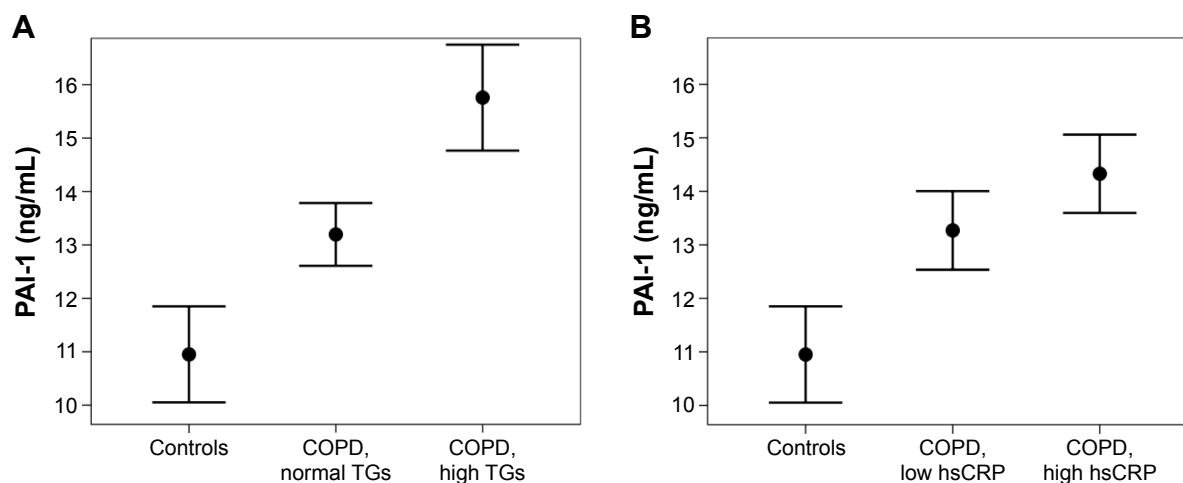


Figure 2 Mean serum PAI-1 levels in controls and COPD patients according to (A) normal or elevated triglyceride levels (cutoff: 150 mg/dL) and (B) low or high hs-CRP (cutoff: median hs-CRP, ie, 3.0 mg/L).

Notes: The control group also included five subjects (28%) with elevated triglycerides and four subjects (22%) with high hs-CRP levels. Points represent mean values and whiskers represent standard error.

Abbreviations: COPD, chronic obstructive pulmonary disease; hs-CRP, high-sensitive C-reactive protein; PAI-1, plasminogen activator inhibitor-1; TGs, triglycerides.

Table 3 Independent predictors for the level of PAI-1 – a multivariate linear regression analysis

	Coefficient B (95% CI)	Standardized β	P-value
Age, years	-0.10 (-0.24 to 0.04)	-0.14	0.14
Sex, male	0.14 (-1.75 to 2.03)	0.02	0.88
Log-triglycerides, mg/dL	2.46 (0.22 to 4.70)	0.22	0.032
Log-hs-CRP, mg/L	0.82 (0.06 to 1.58)	0.23	0.034
GOLD Stage I	0.99 (-1.84 to 3.82)	0.08	0.49
GOLD Stage II	3.02 (0.28 to 5.75)	0.28	0.031
GOLD Stage III	2.81 (0.02 to 5.59)	0.25	0.048
GOLD Stage IV	0.70 (-2.19 to 3.60)	0.07	0.63

Notes: Data from the multivariate linear regression analysis for the total study cohort (n=92) with PAI-1 as the dependent variable are shown. First, age and sex were included in the model, followed by both independent predictors for PAI-1 derived from the former model within COPD patients, that is, triglycerides and hs-CRP. Last, GOLD stages were included in the model after they were dummy-coded with controls as the reference category. Triglycerides and hs-CRP were log-transformed to obtain a normal distribution.

Abbreviations: COPD, chronic obstructive pulmonary disease; GOLD, Global Initiative for Chronic Obstructive Lung Disease; hs-CRP, high-sensitive C-reactive protein; PAI-1, plasminogen activator inhibitor-1.

COPD is characterized by airway and lung abnormalities and is frequently complicated by concomitant cardiovascular and metabolic disorders.¹ Local pathological changes of COPD include parenchymal tissue destruction, resulting in emphysema, and disrupted repair and defense mechanisms, resulting in small airway fibrosis.¹ PAI-1 is synthesized by a variety of cells including vascular endothelial cells, macrophages, and fibroblasts and not only is involved in the pathogenesis of thrombosis but also activates tissue repair.^{30–32} Transiently increased levels of PAI-1 may protect extracellular matrix proteins from proteolytic degradation, inducing matrix remodeling and enhancing wound healing.^{33,34} It is well documented that tissue homeostasis is maintained by the balance of extracellular matrix synthesis and degradation depending on the cellular proteolytic activities of urokinase-type/tissue-type plasminogen activator and plasmin, which mostly rely on the activity of PAI-1.^{33,35} According to the data from the present study, the independent elevation of PAI-1 in patients with COPD may indicate a disease-specific mechanism. We speculate that higher PAI-1 levels may to some extent reflect processes of airway remodeling in COPD. These processes start early in the course of the disease and are less present in advanced COPD,³⁶ which might be the reason why the PAI-1 level is highest in moderate COPD. This concept is in good agreement with the findings of Hogg et al showing that indicators of small airway remodeling are increased in moderate COPD in comparison with controls and declined in very severe COPD again.³⁶ In line with this result, the

plasminogen activator system has already been identified as a potential treatment target for chronic respiratory diseases including COPD.³⁷

On the other hand, the pivotal role of PAI-1 in the pathogenesis of thrombosis, in general, seems to be clear,^{5,7–10} which allows to propose that higher levels of PAI-1 in patients with moderate-to-severe COPD (GOLD Stages II and III) may be associated with an increased risk for cardiovascular events. This idea is supported by the observation that CVD and cardiovascular mortality are highest in patients with moderate-to-severe COPD and drops in very severe COPD.^{2,3} Overall, PAI-1 is recognized as one of the central molecules linking the metabolic syndrome/obesity to the increased CVD risk.^{26,38} Beyond its function as a tissue-remodeling activator, PAI-1 is also an adipocytokine, which is upregulated along with fat accumulation and, therefore, is associated with insulin resistance, dyslipidemia, and obesity.^{5,9,26,38} Metabolic comorbidities are, in turn, especially prevalent in COPD patients with GOLD Stages II and III.² Altogether, PAI-1 may serve as a cross-talk molecule between airway remodeling during the course of COPD, accompanied metabolic comorbidities, and a subsequent increased risk for cardiovascular events, especially in moderate COPD.

The present study has some limitations. First, this is a cross-sectional study and therefore does not allow any interpretation on causality. The elevation of PAI-1 in COPD could also represent a bystander effect. However, it was indicated that PAI-1 may play a plausible role in COPD, and this is the first study conducted in a well-characterized COPD cohort to demonstrate a clear association of PAI-1 with COPD independent of a broad range of possible confounders. Second, the sample size, especially in the control group, is small. Despite these limitations, a significant difference in PAI-1 between COPD patients and controls was found. Of note, the control subjects were free from lung diseases but not explicitly healthy and, therefore, were well matched in terms of metabolic comorbidities. Furthermore, for remaining differences in some cardiovascular conditions, statistical adjustments were performed. Nevertheless, further studies are needed to confirm these observations and to evaluate the role of PAI-1 for cardiovascular outcomes in COPD.

Conclusion

The results of this study showed that PAI-1 levels are higher in patients with COPD, independent of metabolic disorders and cardiovascular function. Hypertriglyceridemia and systemic inflammation may play a role for elevated levels of PAI-1 in COPD, even though they do

not explain this elevation solely, as moderate-to-severe airflow limitation is independently associated with higher levels of PAI-1. These findings suggest PAI-1 as a potential biomarker candidate that indicates a link between airway remodeling, metabolic comorbidities, and an increased cardiovascular risk.

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Disclosure

The authors report no conflicts of interest in this work.

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