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Effects of Pitavastatin on Coronary Artery Disease and Inflammatory Biomarkers in HIV Mechanistic Substudy of the REPRIEVE Randomized Clinical Trial

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IMPORTANCE Cardiovascular disease (CVD) is increased in people with HIV (PWH) and is characterized by premature noncalcified coronary plaque. In the Randomized Trial to Prevent Vascular Events in HIV (REPRIEVE), pitavastatin reduced major adverse cardiovascular events (MACE) by 35% over a median of 5.1 years.

OBJECTIVE To investigate the effects of pitavastatin on noncalcified coronary artery plaque by coronary computed tomography angiography (CTA) and on inflammatory biomarkers as potential mechanisms for MACE prevention.

DESIGN, SETTING, AND PARTICIPANTS This double-blind, placebo-controlled randomized clinical trial enrolled participants from April 2015 to February 2018 at 31 US clinical research sites. PWH without known CVD who were taking antiretroviral therapy and had low to moderate 10-year CVD risk were included. Data were analyzed from April to November 2023.

INTERVENTION Oral pitavastatin calcium, 4 mg per day.

MAIN OUTCOMES AND MEASURES Coronary CTA and inflammatory biomarkers at baseline and 24 months. The primary outcomes were change in noncalcified coronary plaque volume and progression of noncalcified plaque.

RESULTS Of 804 enrolled persons, 774 had at least 1 evaluable CTA. Plaque changes were assessed in 611 who completed both CT scans. Of 611 analyzed participants, 513 (84.0%) were male, the mean (SD) age was 51 (6) years, and the median (IQR) 10-year CVD risk was 4.5% (2.6-7.0). A total of 302 were included in the pitavastatin arm and 309 in the placebo arm. The mean noncalcified plaque volume decreased with pitavastatin compared with placebo (mean [SD] change, -1.7 [25.2] mm³ vs 2.6 [27.1] mm³; baseline adjusted difference, -4.3 mm³; 95% Cl, -8.6 to -0.1; P = .04; 7% [95% Cl, 1-12] greater reduction relative to placebo). A larger effect size was seen among the subgroup with plaque at baseline (-8.8 mm^3 [95% Cl, -17.9 to 0.4]). Progression of noncalcified plaque was 33% less likely with pitavastatin compared with placebo (relative risk, 0.67; 95% CI, 0.52-0.88; P = .003). Compared with placebo, the mean low-density lipoprotein cholesterol decreased with pitavastatin (mean change: pitavastatin, -28.5 mg/dL; 95% Cl, -31.9 to -25.1; placebo, -0.8; 95% Cl, -3.8 to 2.2). The pitavastatin arm had a reduction in both oxidized low-density lipoprotein (-29% [95% CI, -32 to -26] vs -13% [95% CI, -17 to -9]; P < .001) and lipoprotein-associated phospholipase A2 (-7% [95% CI, -11 to -4] vs 14% [95% CI, 10-18]; P < .001) compared with placebo at 24 months.

CONCLUSIONS AND RELEVANCE IN PWH at low to moderate CVD risk, 24 months of pitavastatin reduced noncalcified plaque volume and progression as well as markers of lipid oxidation and arterial inflammation. These changes may contribute to the observed MACE reduction in REPRIEVE.

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The Randomized Trial to Prevent Vascular Event in HIV (REPRIEVE) trial is a phase 3 multicenter randomized clinical trial of oral pitavastatin calcium, 4 mg per day, vs matched placebo for primary prevention of atherosclerotic CVD (ASCVD) events in PWH at low to moderate risk for ASCVD.⁸ In RE-PRIEVE, the pitavastatin arm had 35% less incident MACE compared with placebo over a median (IQR) of 5.1 (4.3-5.9) years, an effect beyond that anticipated from low-density lipoprotein (LDL) cholesterol reduction alone.⁹

The mechanistic substudy embedded within REPRIEVE was designed to determine potential statin effects to reduce noncalcified plaque volume and progression as assessed by entry and 2-year coronary computed tomography angiography (CTA). Moreover, the substudy sought to identify effects on lipid parameters and circulating biomarkers of inflammation.¹⁰

Methods

Trial Design and Oversight

The design of REPRIEVE and the mechanistic substudy have been previously described,^{8,10} and the trial protocol and statistical analysis plan are available in Supplement 1. In brief, 31 REPRIEVE sites, primarily from the AIDS Clinical Trial Group, were selected to participate in the mechanistic substudy based on capabilities for coronary CTA and blood biomarker acquisition. The trial was designed by study principal investigators in consultation with the National Heart, Lung, and Blood Institute, the National Institute of Allergy and Infectious Diseases, and the AIDS Clinical Trials Group. Institutional review board/ethics committee approval and any other applicable regulatory entity approvals were obtained from the Mass General Brigham Institutional Review Board and each clinical research site. Participants were provided with study information, including discussion of risks and benefits, and signed the approved declaration of informed consent. This study followed the Consolidated Standards of Reporting Trials (CONSORT) reporting guideline.

Trial Population

For the main REPRIEVE trial, eligible individuals were aged 40 to 75 years, had documented HIV, were taking antiretroviral therapy for 180 days or more, had a CD4 cell count greater than 100 cells/mm³, estimated glomerular filtration rate of 60 mL/min/1.73 m² or greater, had an alanine transaminase level 2.5-

Question Can pitavastatin reduce noncalcified coronary plaque and inflammatory biomarkers in people with HIV?

Findings In this substudy of the Randomized Trial to Prevent Vascular Events in HIV (REPRIEVE) including 611 people with HIV, compared with the placebo arm, the pitavastatin arm had a significant 7% relative reduction in noncalcified plaque. Pitavastatin use was associated with reductions in oxidized low-density lipoprotein cholesterol and lipoprotein-associated phospholipase A2.

Meaning In this study, among people with HIV at low to moderate cardiovascular risk, pitavastatin showed promise to reduce noncalcified coronary plaque as well as markers of lipid oxidation and arterial inflammation.

fold the upper limit of normal or lower, and low to moderate 10-year ASCVD risk as calculated by the 2013 Pooled Cohort Equation risk calculator, with LDL cholesterol less than threshold for a given risk score.⁸ Statin use in the 90 days before entry or prior history of CVD were exclusionary. Race and ethnicity were self-reported via a questionnaire; options included Asian, Black or African American, White, or other race, which included participants identifying as American Indian or Alaska Native, Native Hawaiian or Other Pacific Islander, or having more than 1 or unknown race. The REPRIEVE mechanistic substudy enrolled REPRIEVE participants without contraindications to coronary CTA. Detailed inclusion and exclusion criteria are available in the eMethods in Supplement 2.¹⁰

Randomization

REPRIEVE participants were randomized 1:1 to pitavastatin calcium, 4 mg per day, and to matched placebo. Randomization was accomplished using a centralized computer system, stratified by sex, CD4 cell count, and participation in the mechanistic substudy. Study investigators and participants were blinded to randomization. Participants received information on heart-healthy lifestyle, diet, and smoking cessation.

Coronary CTA and Plaque Assessment

Coronary CTA was performed on CT scanners using 64 slices or more according to Society of Cardiovascular CT guidelines.¹¹ Deidentified coronary CTA images were transmitted to the central REPRIEVE CT core laboratory, as previously described.^{10,12} Coronary plaque volume and composition were quantified using semi-automated plaque analysis software (QAngio CT; Medis Medical Imaging).^{13,14} The coronary tree was automatically extracted. Plaque length was established visually with markers at the proximal and distal edges of plaque, and each participant's CTs were read side-by-side blinded to scan order so that the same portion of the coronary artery was measured to minimize interscan variability. The inner and outer borders of the coronary wall were defined by the software and manually edited. Voxels between the outer wall of the coronary artery and the lumen were considered plaque; plaque composition was assigned based on voxel attenuation in Hounsfield units (HU; calcified plaque was defined as ≥350 HU, noncalcified plaque as <350 HU and low attenuation as <30 HU).^{15,16} Effects of Pitavastatin on Coronary Artery Disease and Inflammatory Biomarkers in HIV

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REPRIEVE indicates Randomized Trial to Prevent Vascular Events in HIV.

Participants were randomly assigned to 1 of 3 experienced CT core laboratory readers (B. F., J. K., and J. T.), who read CTs blinded to clinical information, order of CT acquisition, and randomization. Interobserver reproducibility for the primary outcome, noncalcified coronary plaque volume, was excellent (intraclass correlation coefficient, 0.97; 95% CI, 0.95-0.99) in a subset of 40 baseline and follow-up CTs.

The primary outcomes were change in noncalcified coronary plaque volume and progression of noncalcified coronary plaque, defined as any increase in noncalcified plaque volume or new noncalcified plaque. Supportive outcomes included change in total plaque volume (including both noncalcified and calcified components) and change in the composition of plaque (ratio of noncalcified to total plaque volume). An additional exploratory plaque outcome was change in low-attenuation plaque volume.^{17,18}

Lipids and Biomarkers

Blood was obtained fasting at entry, month 4, and month 24 for assessment of lipids (LDL and non-high-density lipoprotein cholesterol), inflammatory (high-sensitivity C-reactive protein [hsCRP], lipoprotein-associated phospholipase A2 [Lp-PLA2], and oxidized low-density lipoprotein [oxLDL]), and immune activation biomarkers (monocyte chemoattractant

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protein 1, IL-6, IL-10, IL-1 β , IL-18, soluble CD14, and soluble CD163). Lipids and hsCRP were measured from serum at Quest Diagnostics, Secaucus, New Jersey. Inflammatory and immune biomarkers were measured from plasma using enzyme-linked immunosorbent assay kits (R&D Systems) except for oxLDL (Mercodia) at Temple University, Philadelphia, Pennsylvania.

Statistical Analysis

The statistical analysis plan is provided in Supplement 1. While the main trial (N = 7769) was powered for a reduction in cardiovascular events, target enrollment for the mechanistic substudy was 800 participants to provide 90% power to detect an approximate 6% relative difference between the pitavastatin and placebo arms in noncalcified plaque volume over 2 years and a 27% relative change in the risk of plaque progression, assuming 50% had noncalcified plaque at entry and 15% dropout.¹⁹ Given the expected mixed population of participants with and without noncalcified plaque at entry, power was estimated via a simulation study including a 5% type error with no adjustment for the dual primary outcomes.

Participants completing a parent trial entry visit and a separate substudy registration were considered enrolled (**Figure 1**). The modified intention-to-treat population for plaque out-

Table 1. Baseline Characteristics by Treatment Arm

comes was limited to participants who had at least 1 evaluable CT. Primary analyses included participants with complete case outcomes (eg, paired entry and year 2); sensitivity analyses used multiple imputation in the population with at least 1 evaluable CT (Statistical Analysis Plan in Supplement 1). A per-protocol analysis was conducted including only those participants who completed their 2-year course of pitavastatin or placebo. A planned secondary analysis was restricted to participants with quantified coronary plaque at entry (ie, excluding those with no plaque).

Data are presented as counts with percentages, means with SDs, and medians with IQRs. Changes from baseline in continuous outcomes are presented as means with 95% CIs. Treatment group differences in plaque volumes were based on differences in means adjusted for entry plaque volume using linear regression. Analyses adjusting for race and ethnicity, sex, and ASCVD risk score did not change the findings and are not presented. Analyses were conducted on the measured scale and with a log transformation to reflect both absolute and relative changes. Probability of noncalcified and total plaque volume progression was compared by treatment group using a χ^2 test with the effect size presented as a relative risk (stratified on plaque presence at entry). Statin effects on lipids and markers of immune function and inflammation were assessed via Wilcoxon test for continuous variables.

Associations with changes in LDL cholesterol as well as immune and inflammatory biomarkers were assessed using linear regression for noncalcified coronary plaque volume and using log binomial regression for noncalcified plaque progression. These mechanistic analyses were limited to the perprotocol population with paired biomarkers and plaque outcomes (overall and the subpopulations with plaque at entry). The effects of biomarker changes on noncalcified coronary plaque volume at 4 and 24 months were examined to find the relationship of short-term and longer-term biomarkers changes to longer-term changes in noncalcified plaque volume. We also assessed whether adjusting for these biomarkers impacted the pitavastatin effect on plaque.

All inference used a 5% type I error with no adjustment for multiple comparisons, and all *P* values were 2-tailed. All analyses were conducted with SAS version 9.4M7 (SAS Institute).

Results

Study Population

A total of 804 participants from 31 US sites were enrolled from April 2015 to February 2018. Baseline characteristics of the enrolled population were previously reported and are compared with the parent REPRIEVE trial in eTable 1 in Supplement 2.¹² Of the 804 enrolled participants, 774 had at least 1 evaluable CT (eTable 2 in Supplement 2). Of these, 611 had complete data for evaluation of noncalcified plaque progression, and 587 had complete data for change in noncalcified plaque volume. Among the 611 complete case participants (**Table 1**), 513 (84.0%) were male, the mean (SD) age was 51 (6) years, and the median (IQR) 10-year CVD risk was 4.5% (2.6-7.0). A total of 302 were included in the pitavastatin arm

	No. (%)	
Characteristic	Pitavastatin (n = 302)	Placebo (n = 309)
Demographic characteristics		
Age, y		
Mean (SD)	51 (6)	51 (6)
40-49	117 (39)	141 (46)
50-59	163 (54)	142 (46)
≥60	22 (7)	26 (8)
Natal sex		
Female	51 (17)	47 (15)
Male	251 (83)	262 (85)
Gender identity		
Cisgender	293 (97)	303 (98)
Transgender spectrum	7 (2)	5 (2)
Not reported	2 (1)	1 (<0.5)
Self-reported race ^a		
Asian	4 (1)	5 (2)
Black or African American	105 (35)	115 (37)
White	159 (53)	168 (54)
Other race	34 (11)	21 (7)
Self-reported ethnicity ^b		
Hispanic or Latino	79 (26)	71 (23)
Not Hispanic or Latino	221 (73)	234 (76)
Unknown	2 (1)	4 (1)
Cardiovascular risk factors		
ASCVD risk score, %		
Median (IQR)	4.4 (2.7-6.7)	4.6 (2.5-7.0)
0-<2.5	67 (22)	75 (24)
2.5-<5	100 (33)	96 (31)
5-10	116 (38)	117 (38)
>10	19 (6)	21 (7)
Coronary artery calcium score		
in Agatston units, No./total No. (%)	104/200 (67)	101/202
0	194/290 (67)	(65)
1-100	67/290 (23)	74/293 (25)
101-400	25/290 (9)	20/293 (7)
>400	4/290 (1)	8/293 (3)
Smoking status, No./total No. (%)		
Current	74/301 (25)	71/308 (23)
Former	104/301 (35)	93/308 (30)
Never	123/301 (41)	144/308
		(47)
Substance use, No./total No. (%)	5 (2.04 (2))	= (2.0= (2))
Current	5/301 (2)	//30/(2)
Former	146/301 (49)	148/307 (48)
Never	150/301 (50)	152/307 (50)
Use of antihypertensive medication	61 (20)	63 (20)
Systolic blood pressure, mean (SD), mm Hg	123 (13)	122 (13)
Cholesterol level, median (IQR), mg/dL ^d		
Total	186 (164-205)	180 (159-207)
HDL	49 (40-61)	48 (39-59)
LDL	108 (89-126)	106 (86-128)
		(continued)

	No. (%)		
Characteristic	Pitavastatin (n = 302)	Placebo (n = 309)	
Triglycerides, median (IQR), mg/dL ^d	114 (78-163)	111 (76-173)	
Family history of premature CVD	77 (25)	58 (19)	
BMI			
Mean (SD)	27.3 (4.3)	27.3 (4.3)	
<25	97 (32)	107 (35)	
25-29.9	132 (44)	118 (38)	
≥30	73 (24)	84 (27)	
Prior statin use ^e	21 (7)	25 (8)	
Preexisting diabetes	1 (<0.5)	0	
Use of antidiabetic medication	1 (<0.5)	0	
Use of ACE inhibitors or ARBs	43 (14)	42 (14)	
Use of antiplatelet therapy (including aspirin) ^f	24 (8)	26 (8)	
Use of nonstatin lipid-lowering therapy	24 (8)	19 (6)	
HIV history			
Time since HIV diagnosis, median (IQR), y	15 (9-21)	15 (9-22)	
Nadir CD4 cell count, cells/mm ³			
<50	63 (21)	70 (23)	
50-199	88 (29)	90 (29)	
200-349	86 (28)	78 (25)	
≥350	55 (18)	63 (20)	
Unknown	10 (3)	8 (3)	
Total ART use, y			
<5	43 (14)	46 (15)	
5-10	75 (25)	83 (27)	
≥10	184 (61)	180 (58)	
HIV-related health			
CD4 cell count, cells/mm ³			
Mean (SD)	629 (269)	639 (286)	
<350	44 (15)	46 (15)	
350-499	56 (19)	59 (19)	
≥500	202 (67)	204 (66)	
HIV-1 RNA, No./total No. (%)			
<llq< td=""><td>263/300 (88)</td><td>264/303 (87)</td></llq<>	263/300 (88)	264/303 (87)	
LLQ -<400 copies/mL	29/300 (10)	32/303 (11)	
≥400 copies/mL	8/300 (3)	7/303 (2)	
Entry ART regimen class			
NRTI + INSTI	120 (40)	143 (46)	
NRTI + non-NRTI	85 (28)	81 (26)	
NRTI + PI	62 (21)	40 (13)	
NRTI sparing	7 (2)	12 (4)	
Other NRTI containing	28 (9)	33 (11)	
Entry NRTI			
TDF	148 (49)	162 (52)	
TAF	88 (29)	81 (26)	
Abacavir	52 (17)	47 (15)	
No NRTI	9 (3)	15 (5)	
Other	5 (2)	4(1)	

(continued)

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Table 1. Baseline Characteristics by Treatment Arm (continued)

	No. (%)			
Characteristic	Pitavastatin (n = 302)	Placebo (n = 309)		
Entry INSTI				
No INSTI	151 (50)	128 (41)		
Dolutegravir	71 (24)	83 (27)		
Elvitegravir	57 (19)	65 (21)		
Raltegravir	23 (8)	33 (11)		

Abbreviations: ACE, angiotensin-converting enzyme; ARB, angiotensin receptor blocker; ART, antiretroviral therapy; ASCVD, atherosclerotic cardiovascular disease; BMI, body mass index (calculated as weight in kilograms divided by height in meters squared); CVD, cardiovascular disease; HDL, high-density lipoprotein; INSTI, integrase strand inhibitor; LDL, low-density lipoprotein; LLQ, lower limit of quantification; NRTI, nucleoside reverse transcriptase inhibitor; PI, protease inhibitor; RNA, ribonucleic acid; TAF, tenofovir alafenamide; TDF, tenofovir disoproxil fumarate.

SI conversion factor: To convert cholesterol to mmol/L, multiply by 0.259; triglycerides to mmol/L, multiply by 0.0113.

- ^a The other race category included participants self-identifying as American Indian or Alaska Native, Native Hawaiian or Other Pacific Islander, or having more than 1 or unknown race.
- ^b Ethnicity is presented per National Institutes of Health definitions.
- $^{\rm c}$ Substance use includes use of cocaine, methamphetamine, and intravenous drugs.

^d Screening lipids presented.

- ^e Prior statin use based on site report at entry.
- ^f Aspirin use in antiplatelet therapy is limited to long-term aspirin use, defined as more than 60 days.

and 309 in the placebo arm. The median (IQR) HIV diagnosis duration at enrollment was 15 (9-22) years. The median (IQR) 10-year ASCVD risk score was 4.5% (2.6-7.0), with 338 (55.3%) having an ASCVD risk score less than 5%. The median (IQR) LDL cholesterol level was 108 (87-126) mg/dL (to convert to millimoles per liter, multiply by 0.0259). The median (IQR) body mass index (calculated as weight in kilograms divided by height in meters squared) was 26.9 (24.3-30.1). Participants had good virologic control, with a median (IQR) CD4 cell count of 609 (442-783) cells/mm³; 364 (59.6%) had been taking antiretroviral therapy for more than 10 years.

Primary Outcome: Noncalcified Plaque Volume

At baseline, the mean (SD) noncalcified plaque volume was 52.3 (193.5) mm^3 in the pitavastatin arm and 57.7 (109.2) mm^3 in the placebo arm (Table 2). Noncalcified plaque volume decreased in the pitavastatin arm compared with the placebo arm at month 24 (mean [SD] change, -1.7 [25.2] mm³ vs 2.6 [27.1] mm³; baseline adjusted difference, -4.3 mm³; 95% CI, -8.6 to -0.1; P = .04). Analyzed as a relative change from baseline, the treatment effect represented a 7% (95% CI, 1-12) greater decrease in noncalcified plaque volume compared with placebo (fold change, 0.93; 95% CI, 0.88-0.99; *P* = .02). Progression of noncalcified plaque was less likely in the pitavastatin arm (relative risk, 0.67; 95% CI, 0.52-0.88; P = .003). The sensitivity analysis using multiple imputation to account for missing data and the per-protocol analysis including only participants who completed treatment as randomized supported these findings (eTables 3 and 4 in Supplement 2).

	Treatment arm		Treatment effect ^b		
Outcome	Pitavastatin (n = 302)	Placebo (n = 309)	Estimated difference adjusted for baseline (95% CI)	P value	
Primary outcomes					
NCP volume, mean (SD), mm ^{3c}					
Baseline	52.3 (193.5)	57.7 (109.2)	NA	NA	
Month 24	50.6 (192.8)	60.4 (113.7)	NA	NA	
Change from baseline	-1.7 (25.2)	2.62 (27.1)	-4.3 (-8.6 to -0.1)	.04	
Fold change from baseline (95% CI)	0.95 (0.91-1.00)	1.02 (0.99-1.05)	0.93 (0.88-0.99)	.02	
Progression of NCP, No. (%)	53 (18)	85 (28)	0.67 (0.52-0.88)	.003	
Secondary outcomes					
Total plaque volume, mean (SD), mm ³					
Baseline	65.6 (211.7)	72.2 (135.9)	NA	NA	
Month 24	67.9 (218.4)	79.1 (149.1)	NA	NA	
Change from baseline	2.34 (28.7)	6.89 (32.3)	-4.3 (-9.2 to 0.62)	NA	
Progression of total plaque, No. (%)	80 (28)	100 (34)	0.89 (0.74-1.08)	NA	
Reduction in NCP percentage, mean (SD)	-6.6% (14.3)	-2.6% (9.21)	-4.1% (-7.0 to -1.3)	NA	
Exploratory outcomes					
Low-attenuation plaque volume, mean (SD), mm ³					
Baseline	3.32 (16.5)	4.25 (15.3)	NA	NA	
Month 24	2.43 (9.91)	4.17 (12.8)	NA	NA	
Change from baseline	-0.9 (9.97)	-0.1 (8.16)	-1.2 (-2.2 to -0.1)	NA	

Abbreviations: NA, not applicable; NCP, noncalcified plaque.

 ^a NCP defined as plaque voxels with attenuation <350 Hounsfield units. Total plaque includes all plaque voxels (noncalcified + calcified). Low-attenuation plaque defined as <30 Hounsfield units.

^b For continuous outcomes, the treatment effect is given as the mean difference or relative fold change between treatment groups; for binary outcomes, it is a relative risk. All treatment effects and associated *P* values are adjusted for baseline values. Intention-to-treat analyses were conducted with the exception of reduction in NCP percentage, which is calculated in the subgroup with plaque present at baseline.

^c Change in NCP volume could be assessed in a subset of 587 participants (297 in the pitavastatin arm and 290 in the placebo arm).

In a preplanned subgroup analysis, a larger effect size was seen among the subgroup with plaque at baseline (-8.8 mm³; 95% CI, -17.9 to 0.4; eTable 5 in Supplement 2), representing a 12% greater decrease in noncalcified plaque in pitavastatin compared with placebo arms. There was similar reduction in progression of noncalcified plaque compared with placebo (relative risk, 0.71; 95% CI, 0.55-0.93). Example images of plaque progression and regression are provided in eFigure 1 in Supplement 2.

Secondary and Exploratory Plaque Outcomes

There was a nonsignificantly smaller increase in total plaque volume (ie, noncalcified and calcified) with pitavastatin (mean [SD] change: pitavastatin, 2.3 [28.7] mm³; placebo, 6.9 [32.3] mm³; treatment effect, -4.3 mm³; 95% CI, -9.2 to 0.62). With respect to plaque composition, the percentage of noncalcified plaque decreased a mean (SD) of -6.6% (14.3) in the pitavastatin arm and -2.6% (9.2) in the placebo arm, for a treatment effect of -4.1% (95% CI, -7.0 to -1.3). Low-attenuation plaque volume also decreased on average in the pitavastatin arm compared with the placebo arm (mean [SD] change, -0.9 [10.0] mm³ vs -0.1[8.2] mm³; baseline adjusted difference, -1.2 mm³; 95% CI, -2.2 to -0.1).

At baseline, the mean (SD) calcified plaque volume was 10.4 (30.6) mm³ in the pitavastatin arm and 13.1 (42.4) mm³ in the placebo arm. Calcified plaque volume increased in both the pitavastatin and placebo arms at month 24 (mean [SD] change, 4.1 [13.2] mm³ vs 4.2 [15.4] mm³).

Effect on Lipids and Inflammatory Biomarkers and Relationship to Plaque

Entry and month 24 circulating lipids and measures of inflammation are provided in **Table 3** and eTable 6 in **Supplement 2**. Available month 4 values are provided in eTable 7 in **Supplement 2**. As in the parent REPRIEVE trial, LDL cholesterol levels were similar at baseline and changed from a median (IQR) of 105 (88-124) mg/dL to 75 (59-93) mg/dL in the pitavastatin group and from 107 (90-129) mg/dL to 108 (87-128) mg/dL in the placebo group at year 2. Similar effects were seen for non-high-density lipoprotein cholesterol.

The pitavastatin arm demonstrated a greater reduction in oxLDL at year 2 (pitavastatin, -29%; 95% CI, -32 to -26; placebo, -13%; 95% CI, -17 to -9; P < .001). Pitavastatin also reduced LpPLA2 by -7% (95% CI, -11 to -4), while there was a 14% (95% CI, 10-18) increase in LpPLA2 in the placebo arm (P < .001) at year 2. There was a nonsignificant decrease in hsCRP (Table 3). No significant effects were seen on markers of immune activation.

There was no significant association between change in LDL cholesterol and change in continuous noncalcified plaque volume (eFigure 2 in Supplement 2). Change in LDL cholesterol and an achieved LDL cholesterol level less than 70 mg/dL at 24 months were associated with lower risk of noncalcified plaque progression, though these relationships were not significant after adjustment for baseline plaque volume (Figure 2). No significant associations between

Table 3. Distributions of Inflammatory and Immune Biomarkers at Entry and Month 24

ImportImpor		Median (IQR)				
<table-container>HerePerformantPe</table-container>		Entry		Month 24		
set is a stateImplementation360360360360360360Implementation360360360360360360Implementation360360360360360360360Implementation360 <td< td=""><td>Measure^a</td><td>Pitavastatin (n = 402)</td><td>Placebo (n = 402)</td><td>Pitavastatin (n = 349)</td><td>Placebo (n = 350)</td><td>P value</td></td<>	Measure ^a	Pitavastatin (n = 402)	Placebo (n = 402)	Pitavastatin (n = 349)	Placebo (n = 350)	P value
Unit of the set of th	Inflammatory markers					
Form IP39839235533NALipelAceed.ng/md126(04.8-160)131(080-163)143(105-180)0.301Inchage from baselineNA-9.71-3.5 to 1.6.8.0139(-5.0 to 4.3.00.301IDNA0.302 080-0.001.14 (1.0-1.8)NAIDS1.04 (2.6.9.8.0)3.5 (3.0.9.0.90)1.42 (1.0-1.8)NAIDS1.04 (2.6.9.8.0)3.5 (3.0.042.9.0.30)A.20 (2.0.0.0.0.0.0.0.0.0.0.0.0.0.0.0.0.0.0.0	LpPLA2					
IpPLQ Interplay126 (94.8-169)131 (98.6-163)143 (0.6-160)143 (0.6-1000.001Man find knapp (SK)(D)NA0.33 (0.80-0.65)1.14 (1.10-1.18)NAMA0.33 (0.80-0.65)1.14 (1.10-1.18)NAMan find knapp (SK)(D)NA0.33 (0.80-0.65)1.14 (1.10-1.18)NAMain find knapp (SK)(D)S3.0 (42.2-69.8)S3.6 (41.7-69.9)37.6 (30.0-47.3)48.2 (38.9-58.4)<001	Total, No.	398	391	335	333	NA
And Index prom baselineNA-9.77 (-3.5 k to 16.5)19.9 (-5.6 to 43.0)-0.01Mana fold change (95% t)NANA0.93 (0.89-0.90)1.14 (1.01.10)NAMala398392335332NANAor.DLS3.0 (42.2-68.0)S3.6 (41.7-69.9)37.6 (30.0-7.3)42.2 (38.5-6.4)<001	LpPLA2 level, ng/mL	126 (94.8-169)	131 (89.6-163)	119 (84.0-160)	143 (105-186)	<.001
Mander damper dampe	Change from baseline	NA	NA	-9.77 (-33.6 to 16.8)	19.9 (-5.6 to 43.0)	<.001
netlectionnotal level363363363MAnotal level36.0 (42.2-69.30)53.6 (41.7-69.0)7.6 (30.0-47.3)4.8 (28.9-58.4)-001Manefold change (95%.0)NANA-14.9 (-28.9 to -3.9.3)6.45 (-20.2 to 5.2.0)-001Manefold change (95%.0)NANA-14.9 (-28.9 to -3.9.3)6.45 (-20.2 to 5.2.0)-001IncreationNANA-14.9 (-28.9 to -3.9.3)6.45 (-20.2 to 5.2.0)NAIncreationNANA-14.9 (-28.9 to -3.9.3)0.64 (-20.2 to 5.2.0)NAIncreationNANA0.71 (0.68-0.7.9.3)0.19 (0.10-0.000.01Scherel mojdtNANANA0.01 (-0.10 to 0.0.0)0.01 (-0.09 to 0.10)0.01Change fom baselineNANANANANANANAIncreationNANANANANANANANAIncreationNANA211 (-37.8 to 4.7.5)139 (16.4 '10.7.0.3).46MCP-1 level, pr/m189 (18-47.4)131 (14.9.2.00135 (14.2.47.50).47.48Schube CD14 level, pr/m189 (18-47.4)131 (14.9.2.00135 (14.2.47.50).54.48Schube CD14 level, pr/m189 (18-47.4)131 (14.9.4.5.8.4.5.	Mean fold change (95% CI)	NA	NA	0.93 (0.89-0.96)	1.14 (1.10-1.18)	NA
Forda, Ro.398392335335NAorda, Rove, M., Na37.6 (30.0.47.3)48.2 (38.9.58.4)6.001Change from baselineNANA0.419 (2.28.9.16.2.4)6.4.6 (2.02.16.2.520)6.010NaNA0.71 (0.68.0.74)0.87 (0.33.0.51)NANAhscRP*Total, Na397391431335NAhscRp0.18 (0.09-0.39)0.17 (0.07-0.36)0.19 (0.10-0.040)0.20Maen fold change (95% (1)NANA0.001 (0.10 to.00.8)0.10 (0.00 to.01.1)0.90Maen fold change (95% (1)NANA0.01 (0.10 to.00.8)0.10 (0.00 to.01.1)0.90Maen fold change (95% (1)NANA0.90 (0.79-1.01)1.06 (0.94-1.20)NAMaen fold change (95% (1)NANA0.90 (0.79-1.01)1.06 (0.94-1.20)NAMaen fold change (95% (1)NANA1.92 (1.57-235)1.93 (1.54-247).72Change from baselineNANA2.11 (-37.8 to.47.5).91 (-31.8 to.52.9.46Maen fold change (95% (1)NANA2.31 (-37.8 to.47.5).51 (-12.8 to.52.9.46Solube CD14 evel, ng/ml184 (1.64-21.69)1.54 (1.24-1.690.54 (1.24-1.690.54Solube CD14 evel, ng/mlNA1.91 (1.94-2.92).54 (1.24-1.690.54 (1.24-1.690.54 (1.24-1.690Solube CD14 evel, ng/mlNA1.91 (1.24-1.690.54 (1.24-1.690.54 (1.24-1.690.54 (1.24-1.690.54 (1.24-1.690.54 (1.24-1.690.54 (1.24-1.690<	oxLDL					
exclewed, U/L53.0 (42.e.69.8)53.6 (41.7-69.9)77.6 (30.0-73)48.2 (83.9-5.84)<010Change form baselineNANA-14.9 (-7.8.9 to -3.0.3)-6.45 (-7.0.2 to 5.2.0)<0.001	Total, No.	398	392	335	333	NA
Change from baselineNANA-1.4 9 (-28.8 to -3.3)-6.45 (-20.2 to 5.26)<.001Maan fold change (95% CI)NANA0.71 (0.66-0.7A)0.87 (0.83-0.91)NAISCRPPTotal, No.397391343335NAISCRPS0.18 (0.08-0.39)0.17 (0.07-0.36)0.10 (0.09-0.10.100.02Change from baselineNANA0.80 (0.07-1.01)1.06 (0.94-1.20)NAImmute markesNA0.80 (0.97-1.01)1.06 (0.94-1.20)NAImmute markesNA0.89 (0.79-1.01)1.06 (0.94-1.20)NAImmute markesNA1.02 (0.98-1.06)1.06 (0.94-1.20)NAImmute markesNA1.02 (0.98-1.06)1.06 (0.11)0.90Immute markesNA1.02 (0.98-1.06)1.06 (0.11)0.90Immute markesNA1.02 (0.98-1.06)1.05 (0.11)0.90 <t< td=""><td>oxLDL level, U/L</td><td>53.0 (42.2-69.8)</td><td>53.6 (41.7-69.9)</td><td>37.6 (30.0-47.3)</td><td>48.2 (38.9-58.4)</td><td><.001</td></t<>	oxLDL level, U/L	53.0 (42.2-69.8)	53.6 (41.7-69.9)	37.6 (30.0-47.3)	48.2 (38.9-58.4)	<.001
ManNA0.71 (0.68-0.74)0.87 (0.83-0.91)NAhcRPb*Total, No.397391343335NAhcRP (evel, mg/dL0.18 (0.08-0.39)0.18 (0.09-0.33)0.17 (0.07-0.36)0.19 (0.10-0.40).02Change from baselineNANA-0.01 (-0.01 to 0.08)0.01 (-0.09 to 0.11).09Maan fold change (95% CINANA0.90 (19-0.40)1.06 (0.94-1.20)NAMan fold change (95% CINANA0.90 (19-0.40).018 (0.08-0.31).01Man fold change (95% CINANA0.90 (19-0.40).016 (0.94-1.20)NAImmune markesNA1.92 (157-235)1.93 (154-247).77fotal, No.3983903353.91 (-33.3 to 53.2).46MCP-1 level, ng/mL1.89 (148-248)1.83 (144-238)1.92 (157-235)1.93 (154-247).77fotal, No.3983923553.31 (Na (196-200)NASoluble CD14NA2.11 (-37.81 to 47.5).39 (1-33.3 to 53.2).46for haselineNANA.280 (-588 to -5.08).72 (-52.858).52fotal, No.398392355.33.3NASoluble CD14Is40 (154-2176)I810 (1496-220)I541 (128-1809)I541 (132-1850).52for haselineNANA.03 (0.81-0.88).84.65.64.65Soluble CD15NA.84 (561-163).84 (561-163).84.65.64.65.64.65.64.65<	Change from baseline	NA	NA	-14.9 (-28.9 to -3.93)	-6.45 (-20.2 to 5.26)	<.001
InterfaceTotal, No.397391343335NAhaCRP level, mg/dt0.18 (0.08-0.39)0.17 (0.07-0.36)0.19 (0.10-0.40).02Change from baselineNANA0.89 (0.79-1.01)1.06 (0.94-1.20)NAImmune arkersNNA0.89 (0.79-1.01)1.06 (0.94-1.20)NAImmune arkersNNA0.89 (0.79-1.01)1.06 (0.94-1.20)NAICP-1NNA0.89 (0.79-1.01)1.06 (0.94-1.20)NAChange from baselineNA1.02 (0.98-1.02)1.91 (1.92, 1.92).05 (0.91, 1.92)Change from baselineNA1.02 (0.98-1.04).190 (1.01-1.00)NASoluble CD14NA1.02 (0.98-1.04).191 (1.92, 1.92).191 (1.92, 1.92).191 (1.92, 1.92)Soluble CD141.940 (1.94-2.17)1.910 (1.94-2.92).313NA.52Soluble CD14 level, ng/m1.940 (1.94-2.17)1.910 (1.94-2.92).314 (1.92, 1.92).52Soluble CD14 level, ng/m1.940 (1.94-2.17)1.910 (1.94-2.92).52.54Soluble CD14 level, ng/m1.940 (1.94-1.91)1.931 (1.94-1.92).54Soluble CD14 level, ng/m1.940 (1.94-1.91)1.931 (1.94-1.91).54Soluble CD163 level, ng/mAA.491 (-6.03.0.420).431 (-6.03.0.420).431 (-6.03.0.420).431 (-6.03.0.420).431 (-6.03.0.420).54Change from baselineNANA1.921 (1.91-7.91)1.931 (-6.03.0.420).431 (-6.03.0.420).45Change	Mean fold change (95% CI)	NA	NA	0.71 (0.68-0.74)	0.87 (0.83-0.91)	NA
For Lay No.397391343335NAIn ScR level, my/dL0.18 (0.09-0.39)0.18 (0.09-0.39)0.19 (0.10-04 (0.1)0.01Change from baselineNANA-0.01 (-0.10 (0.0)0.01 (-0.04 (0.1)NAImmerativesNA0.89 (0.79-1.01)1.06 (0.94-1.20)NAImmerativesNA0.89 (0.79-1.01)1.06 (0.94-1.20)NAImmerativesNA0.89 (0.79-1.01)1.06 (0.94-1.20)NAImmerativesNA0.89 (0.79-1.01)1.06 (0.94-1.20)NAImmerativesNA0.81 (1.42-23)1.92 (1.57-23)1.93 (1.64-24)NAImmerativesNA1.11 (2.73.51 4.72.51)1.93 (1.64-24).06 (0.11 (1.10))NAImmerativesNANA2.11 (2.73.51 4.72.51).93 (1.63.13 t.53.13).46ImmerativesNANA2.11 (2.73.51 4.72.51).93 (1.63.13 t.53.13).46ImmerativesNANA2.11 (2.73.51 4.53.13).54.12 (1.63.13 t.53.13).46ImmerativesNANA.281 (1.54.124.13).54 (1.24.136.0).54.54Soluble CD14NANA.289 (1.54.130).54 (1.24.136.0).54.54ImmerativesNA.33 (0.11 (0.11 A.73).33.84Soluble CD163NA.33.94.35.33.84ImmerativesNA.39.33.84.54.54ImmerativesNA.34.34 <td>hsCRP^b</td> <td></td> <td></td> <td></td> <td></td> <td></td>	hsCRP ^b					
bsCRP level, mg/dL 0.18 (0.08-0.39) 0.18 (0.09-0.33) 0.17 (0.07-0.36) 0.19 (0.10-0.40) .02 Change from baseline NA NA -0.01 (-0.10 to 0.08) 0.01 (-0.09 to 0.11) .09 Mean fold change (95% CI) NA NA 0.80 (0.79-1.01) 1.06 (0.94-1.20) NA MCP-1	Total, No.	397	391	343	335	NA
Change from baseline NA NA -0.01 (-0.10 to 0.08) 0.01 (-0.09 to 0.11) .09 Menn fold change (55%) NA NA 0.89 (0.79-1.01) 1.06 (0.94-1.20) NA Immune markers - - - - NA Immune markers - - - - - NA MCP-1 -	hsCRP level, mg/dL	0.18 (0.08-0.39)	0.18 (0.09-0.33)	0.17 (0.07-0.36)	0.19 (0.10-0.40)	.02
Mean fold change (95% CI) NA NA 0.89 (0.79-1.01) 1.06 (0.94-1.20) NA Immum markers	Change from baseline	NA	NA	-0.01 (-0.10 to 0.08)	0.01 (-0.09 to 0.11)	.09
Immune markers MCP-1 Total, No. 398 390 335 330 NA MCP-1 level, pg/mL 189 (148-245) 183 (144-238) 192 (157-235) 193 (154-247) .7.7 Change from baseline NA NA 2.11 (~37.8 to 47.5) 3.91 (~33.3 to 53.2) .46 Mean fold change (95% CI) NA NA 1.02 (0.98-1.06) 1.05 (1.01-1.09) NA Soluble CD14 Total, No. 398 392 335 333 NA Soluble CD14 level, ng/mL 1840 (1544-2176) 1810 (1496-2200) 1541 (1284-1809) 1534 (1324-1850) .52 Change from baseline NA NA -289 (~588 to ~5.08) ~237 (~528 to 36.1) .54 Mean fold change (95% CI) NA NA 0.83 (0.81-0.86) 0.86 (0.83-0.88) NA Soluble CD163 Total, No. 398 391 335 333 NA Soluble CD163 level, ng/mL 849 (651-1163) 845 (613-1074) 1013 (728-1419) 1003 (745-1384) .65 Cha	Mean fold change (95% CI)	NA	NA	0.89 (0.79-1.01)	1.06 (0.94-1.20)	NA
MCP-1Total, No.396390335330NATotal, No.398 (148-245)183 (144-283)192 (157-235)193 (154-277).7Change from baselineNANA1.10 (20.98-105).931 (-33.3 to 53.2).46Mean fol change (95%)NANA1.02 (0.98-105).931 (-33.3 to 53.2).46Soluble CD14Total, No.398 (39.2)395333NASoluble CD14 level, ng/m1840 (1544-2176)1810 (1496-2200)1541 (1284-1809).1534 (1324-1850).52Change from baselineNANA-289 (-588 to -5.08).237 (-528 to 36.1).54Mean fol change (95%)NANA.280 (-38.10.86).086 (0.83-0.88)NASoluble CD161Total, No.396 391.351 (33.2.333NASoluble CD163Colspan="2">Colspan="2"Cols	Immune markers					
Total, No.398390335330NAMCP-1 level, pg/mL189 (148-245)183 (144-238)192 (157-235)193 (154-247).77Mean fold change (95% CI)NANA2.11 (-37.8 to 47.5)3.91 (-33.3 to 53.2).46Mean fold change (95% CI)NANA1.02 (0.98-1.06)1.05 (0.10-1.09)NASoluble CD14Tatal, No.398392335333NASoluble CD14 level, ng/mL1840 (1544-2176)1810 (1496-2200)1541 (1284-1809)1534 (1324-1850).52Mean fold change (95% CI)NANA-289 (-588 to -5.08)-237 (-528 to 36.1).54Mean fold change (95% CI)NANA.083 (0.81-0.86).08 (0.83 - 0.88)NASoluble CD163VNA.083 (0.81-0.86).02 (149-1184).65Soluble CD163 level, ng/mL849 (651-1163)845 (613-1074)1013 (728-1419)1003 (745-1384).65Mean fold change (95% CI)NANA.121 (1.16-1.27).123 (1.18-1.29)NAIL-6Tatal, No.398391335.333NAIL-6Tatal, No.198.91.35.333NAIL-6Tatal, No.198.91.35.333.04IL-6Tatal, No.198.91.35.333.04IL-6Tatal, No98.91.35.333.04IL-6Tatal, No98.91.35.333.04IL-6Tatal, No.	MCP-1					
MCP-1 level, pg/mL189 (148-245)183 (144-238)192 (157-235)193 (154-247).77Change from baselineNANA2.11 (-37.8 to 47.5)3.91 (-33.3 to 53.2).46Man fold change (95% CI)NANA1.02 (0.98-1.06)1.05 (1.01-1.09)NASoluble CD14Total, No.398392335333NASoluble CD14 level, ng/mL1840 (1544-2176)1810 (1496-2200)1541 (1284-1809)1534 (1324-1850).52Change from baselineNANA-289 (-588 to -5.08)-237 (-528 to 36.1).54Mean fold change (95% CI)NANA0.83 (0.81-0.86)0.86 (0.83-0.88)NASoluble CD163TTotal, No.398391335333NASoluble CD163 level, ng/mL849 (651-1163)845 (613-1074)1013 (728-1419)1003 (745-1384).65Mean fold change (95% CI)NANA1.21 (1.16-1.27)1.23 (1.18-1.29)NASoluble CD163 level, ng/mL849 (651-1163)845 (613-1074)1013 (728-1419)1003 (745-1384).65IL-6TTTTotal, No.398391335333NAIL-6IL-6 level, ng/mL1.64 (0.99-2.76)1.64 (0.99-2.73)1.53 (0.93-2.62).24IL-18TTTTTTTotal, No.377375320320NAIL-19TTTTTTotal, No.377375320320	Total, No.	398	390	335	330	NA
Change from baseline NA NA 2.11 (-37.8 to 47.5) 3.91 (-33.3 to 53.2) .46 Mean fold change (95% CI) NA NA 1.02 (0.98-1.06) 1.05 (1.01-1.09) NA Soluble CD14 398 392 335 333 NA Soluble CD14 level, ng/mL 1840 (1544-2176) 1810 (1496-2200) 1541 (1284-1809) 1534 (1324-1850) 52 Change from baseline NA NA -289 (-588 to -5.08) -237 (-528 to 36.1) .54 Mean fold change (95% CI) NA NA 0.83 (0.81-0.86) 0.86 (0.83-0.88) NA Soluble CD163 335 333 NA Soluble CD163 494 (-60.180-86) 0.86 (0.83-0.88) NA Soluble CD163 335 333 NA Soluble CD163 494 (-60.181-0.86) 0.86 (0.83-0.88) NA Itals NA NA 1013 (728-1419) 1003 (745-1384) .65 Itals Soluble CD163 849 (651-1163)<	MCP-1 level, pg/mL	189 (148-245)	183 (144-238)	192 (157-235)	193 (154-247)	.77
Mean fold change (95% Cl) NA NA 1.02 (0.98-1.06) 1.05 (1.01-1.09) NA Soluble CD14 Valuation of the transmission of	Change from baseline	NA	NA	2.11 (-37.8 to 47.5)	3.91 (-33.3 to 53.2)	.46
Soluble CD14 Total, No. 398 392 335 333 NA Soluble CD14 level, ng/mL 1840 (1544-2176) 1810 (1496-2200) 1541 (1284-1809) 1534 (1324-1850) .52 Mean fold change (95% CI) NA NA -289 (-588 to -5.08) -237 (-528 to 36.1) .54 Mean fold change (95% CI) NA NA 0.83 (0.81-0.86) 0.86 (0.83-0.88) NA Soluble CD163 level, ng/mL 849 (651-1163) 845 (613-1074) 1013 (728-1419) 1003 (745-1384) .65 Change from baseline NA NA 149 (-60.8 to 420) 143 (-46.0 to 453) .88 Mean fold change (95% CI) NA NA 1.21 (1.16-1.27) 1.23 (1.18-1.29) NA IL-6 Total, No. 398 391 335 333 NA IL-6 Level, ng/mL 1.64 (0.99-2.76) 1.46 (0.99-2.73) 1.53 (0.93-2.62) .24 Mean fold change (95% CI) NA NA 0.90 (-0.95 to 0.77) -0.12 (-0.81 to 0.59) .46 Mean fold change (95% CI) NA	Mean fold change (95% CI)	NA	NA	1.02 (0.98-1.06)	1.05 (1.01-1.09)	NA
Total, No.398392335333NASoluble CD14 level, ng/mL1840 (1544-2176)1810 (1496-2200)1541 (1284-1809)1534 (1324-1850).52Man fold change (95% CI)NANA-289 (-588 to -5.08)-237 (-528 to 36.1).54Soluble CD163	Soluble CD14					
Soluble CD14 level, ng/mL 1840 (1544-2176) 1810 (1496-2200) 1541 (1284-1809) 1534 (1324-1850) .52 Change from baseline NA NA -289 (-588 to -5.08) -237 (-528 to 36.1) .54 Mean fold change (95% CI) NA NA 0.83 (0.81-0.86) 0.86 (0.83-0.88) NA Soluble CD163 Total, No. 398 391 335 333 NA Soluble CD163 level, ng/mL 849 (651-1163) 845 (613-1074) 1013 (728-1419) 1003 (745-1384) .65 Change from baseline NA NA 149 (-60.8 to 420) 143 (-46.0 to 453) .88 Mean fold change (95% CI) NA NA 1201 (1.16-1.27) 1.23 (1.18-1.29) NA IL-6 Total, No. 398 391 335 333 NA IL-6 Le-6 Total, No. 398 391 335 333 NA IL-16 Le-6 NA NA 0.90 (-0.95 to 0.77) -0.12 (-0.81 to 0.59) .66 Mean fold change (95% CI) NA NA	Total, No.	398	392	335	333	NA
Change from baseline NA NA -289 (-588 to -5.08) -237 (-528 to 3.6.1) .54 Mean fold change (95% CI) NA NA 0.83 (0.81-0.86) 0.86 (0.83-0.88) NA Soluble CD163 335 333 NA .65 Fotal, No. 398 391 335 333 NA Soluble CD163 level, ng/ml 849 (651-1163) 845 (613-1074) 1013 (728-1419) 1003 (745-1384) .65 Total, No. 398 391 335 333 NA Mean fold change (95% CI) NA NA 149 (-60.8 to 420) 143 (-46.0 to 453) .88 IL-6 NA NA 149 (-60.8 to 420) 1.23 (1.18-1.29) NA IL-6 IL-6 NA NA 1.49 (-60.8 to 420) 1.23 (1.18-1.29) NA IL-6 IL-6 NA NA 1.49 (-60.8 to 420) 1.23 (1.18-1.29) NA IL-6 IL-6 IL-6 IL-6 IL-6 IL-6 IL-6 IL-6 IL-8 IL-18 IL-18	Soluble CD14 level, ng/mL	1840 (1544-2176)	1810 (1496-2200)	1541 (1284-1809)	1534 (1324-1850)	.52
Mean fold change (95% CI) NA NA 0.83 (0.81-0.86) 0.86 (0.83-0.88) NA Soluble CD163 Total, No. 398 391 335 333 NA Soluble CD163 level, ng/mL 849 (651-1163) 845 (613-1074) 1013 (728-1419) 1003 (745-1384) .65 Change from baseline NA NA 149 (-60.8 to 420) 143 (-46.0 to 453) .88 Mean fold change (95% CI) NA NA 1.21 (1.16-1.27) 1.23 (1.18-1.29) NA IL-6 Total, No. 398 391 335 333 NA Mean fold change (95% CI) NA NA 1.21 (1.16-1.27) 1.23 (1.18-1.29) NA IL-6 Total, No. 398 391 335 333 NA Mean fold change (95% CI) NA NA 0.40.0.99-2.73) 1.53 (0.93-2.62) .24 Mean fold change (95% CI) NA NA 0.90 (0.92.108) 0.94 (0.86-1.03) NA IL-16 Total, No. 377 375 320 320 NA <td>Change from baseline</td> <td>NA</td> <td>NA</td> <td>-289 (-588 to -5.08)</td> <td>-237 (-528 to 36.1)</td> <td>.54</td>	Change from baseline	NA	NA	-289 (-588 to -5.08)	-237 (-528 to 36.1)	.54
Soluble CD163 Total, No. 398 391 335 333 NA Soluble CD163 level, ng/mL 849 (651-1163) 845 (613-1074) 1013 (728-1419) 1003 (745-1384) .65 Change from baseline NA NA 149 (-60.8 to 420) 143 (-46.0 to 453) .88 Mean fold change (95% CI) NA NA 1.21 (1.16-1.27) 1.23 (1.18-1.29) NA IL-6 NA 1.21 (1.16-1.27) 1.23 (0.18-1.29) NA IL-6 NA 1.21 (0.16-2.7) 1.23 (0.18-1.29) NA IL-6 335 333 NA IL-6 1.64 (0.99-2.73) 1.53 (0.93-2.62) .24 Change from baseline NA NA 0.98 (0.89-1.08) 0.94 (0.86-1.03) NA IL-1β 377 375 320 320 NA IL-1β <	Mean fold change (95% CI)	NA	NA	0.83 (0.81-0.86)	0.86 (0.83-0.88)	NA
Total, No. 398 391 335 333 NA Soluble CD163 level, ng/mL 849 (651-1163) 845 (613-1074) 1013 (728-1419) 1003 (745-1384) .65 Change from baseline NA NA 149 (-60.8 to 420) 143 (-46.0 to 453) .88 Mean fold change (95% CI) NA NA 1.21 (1.16-1.27) 1.23 (1.18-1.29) NA IL- Total, No. 398 391 335 333 NA IL-6 Level, pg/mL 1.64 (0.99-2.76) 1.46 (0.98-2.59) 1.64 (0.99-2.73) 1.53 (0.93-2.62) .24 Change from baseline NA NA 0 (-0.95 to 0.77) -0.12 (-0.81 to 0.59) .46 Mean fold change (95% CI) NA NA 0 (-0.95 to 0.77) -0.12 (-0.81 to 0.59) .46 IL-1β Total, No. 377 NA NA 0.94 (0.86-1.03) NA IL-1β Fotal, No. 377 375 320 320 NA IL-1β level, fg/mL 75.9 (43.9-151) 87.0 (44.9-168) 88.6 (50.2-164)	Soluble CD163					
Soluble CD163 level, ng/mL 849 (651-1163) 845 (613-1074) 1013 (728-1419) 1003 (745-1384) .65 Maan fold change (95% (1) NA NA 149 (-6.0.8 to 420) 143 (-46.0 to 453) .88 IL-6 NA NA 1.21 (1.16-1.27) 1.23 (1.18-1.29) NA IL-6 Soluble CD163 level, ng/mL 1.64 (0.99-2.76) 1.46 (0.98-2.59) 1.64 (0.99-2.73) 1.53 (0.93-2.62) .24 Change from baseline NA NA 0 (-0.95 to 0.77) -0.12 (-0.81 to 0.59) .46 Mean fold change (95% (1) NA NA 0.98 (0.89-1.08) 0.94 (0.86-1.03) NA IL-16 Visit Soluble CD164 level, ng/mL 377 375 320 320 NA IL-1β level, fg/mL 75.9 (43.9-151) 87.0 (44.9-168) 88.6 (50.2-164) 96.4 (52.6-191) .09 IL-1β level, fg/mL 75.9 (43.9-151) 87.0 (44.9-168) 88.6 (50.2-164) 96.4 (52.6-191) .09 IL-18 Visit NA NA 20.01 (-41.4 to 58.1) 11.4 (-28.9 to 69.2) .06	Total, No.	398	391	335	333	NA
Change from baseline NA NA 149 (-60.8 to 420) 143 (-46.0 to 453) .88 Mean fold change (95% CI) NA NA 1.21 (1.16-1.27) 1.23 (1.18-1.29) NA IL-6 Total, No. 398 391 335 333 NA IL-6 level, pg/mL 1.64 (0.99-2.76) 1.46 (0.98-2.59) 1.64 (0.99-2.73) 1.53 (0.93-2.62) .24 Change from baseline NA NA 0 (-0.95 to 0.77) -0.12 (-0.81 to 0.59) .46 Mean fold change (95% CI) NA NA 0.98 (0.89-1.08) 0.94 (0.86-1.03) NA IL-1p Total, No. 377 375 320 320 NA Mean fold change (95% CI) NA NA 2.01 (-41.4 to 58.1) 11.4 (-28.9 to 69.2) .06 IL-1p Vertical from baseline NA NA 2.01 (-41.4 to 58.1) 11.4 (-28.9 to 69.2) .06 Mean fold change (95% CI) NA NA 2.01 (-41.4 to 58.1) 11.4 (-28.9 to 69.2) .06 IL-1B Vertical from baseline NA NA	Soluble CD163 level, ng/mL	849 (651-1163)	845 (613-1074)	1013 (728-1419)	1003 (745-1384)	.65
Mean fold change (95% Cl) NA NA 1.21 (1.16-1.27) 1.23 (1.18-1.29) NA IL-6 Total, No. 398 391 335 333 NA IL-6 level, pg/mL 1.64 (0.99-2.76) 1.46 (0.98-2.59) 1.64 (0.99-2.73) 1.53 (0.93-2.62) .24 Change from baseline NA NA 0 (~0.95 to 0.77) ~0.12 (~0.81 to 0.59) .46 Mean fold change (95% Cl) NA NA 0.98 (0.89-1.08) 0.94 (0.86-1.03) NA IL-1β Total, No. 377 375 320 320 NA IL-1β level, fg/mL 75.9 (43.9-151) 87.0 (44.9-168) 88.6 (50.2-164) 96.4 (52.6-191) .09 Mean fold change (95% Cl) NA NA 2.01 (~41.4 to 58.1) 1.1.4 (~28.9 to 69.2) .06 Mean fold change (95% Cl) NA NA 2.05 (.03.1.17) 1.19 (1.07-1.33) NA IL-18 Total, No. 377 375 320 320 (.06 NA IL-18 Level, pg/mL 245 (179-330) 232 (169-310) 2	Change from baseline	NA	NA	149 (-60.8 to 420)	143 (-46.0 to 453)	.88
IL-6 Total, No. 398 391 335 333 NA IL-6 level, pg/mL 1.64 (0.99-2.76) 1.46 (0.98-2.59) 1.64 (0.99-2.73) 1.53 (0.93-2.62) .24 Change from baseline NA NA 0 (-0.95 to 0.77) -0.12 (-0.81 to 0.59) .46 Mean fold change (95% CI) NA NA 0.98 (0.89-1.08) 0.94 (0.86-1.03) NA IL-1β Total, No. 377 375 320 320 NA L1β level, fg/mL 75.9 (43.9-151) 87.0 (44.9-168) 88.6 (50.2-164) 96.4 (52.6-191) .09 Change from baseline NA NA 2.01 (-41.4 to 58.1) 11.4 (-28.9 to 69.2) .06 Mean fold change (95% CI) NA NA 2.01 (-41.4 to 58.1) 11.4 (-28.9 to 69.2) .06 IL-18 Total, No. 377 375 320 320 NA IL-18 IL-18 V NA 1.05 (0.93-1.17) 1.19 (1.07-1.33) NA IL-18 Jotal, No. 377 375 320 320	Mean fold change (95% CI)	NA	NA	1.21 (1.16-1.27)	1.23 (1.18-1.29)	NA
Total, No. 398 391 335 333 NA IL-6 level, pg/mL 1.64 (0.99-2.76) 1.46 (0.98-2.59) 1.64 (0.99-2.73) 1.53 (0.93-2.62) .24 Change from baseline NA NA 0 (-0.95 to 0.77) -0.12 (-0.81 to 0.59) .46 Mean fold change (95% Cl) NA NA 0.98 (0.89-1.08) 0.94 (0.86-1.03) NA IL-1β Total, No. 377 375 320 320 NA Change from baseline NA NA 2.01 (-41.4 to 58.1) 11.4 (-28.9 to 69.2) .06 Change from baseline NA NA 1.05 (0.93-1.17) 1.91 (1.07-1.33) NA IL-18 Total, No. 377 375 320 320 NA Mean fold change (95% Cl) NA NA 2.01 (-41.4 to 58.1) 11.4 (-28.9 to 69.2) .06 IL-18 Total, No. 377 375 320 320 NA IL-18 Total, No. 377 375 320 320 (176-308) .79	IL-6					
IL-6 level, pg/mL 1.64 (0.99-2.76) 1.46 (0.98-2.59) 1.64 (0.99-2.73) 1.53 (0.93-2.62) .24 Change from baseline NA NA 0 (-0.95 to 0.77) -0.12 (-0.81 to 0.59) .46 Mean fold change (95% CI) NA NA 0.98 (0.89-1.08) 0.94 (0.86-1.03) NA IL-1β Total, No. 377 375 320 320 NA Change from baseline NA NA 2.01 (-41.4 to 58.1) 11.4 (-28.9 to 69.2) .06 IL-1β level, fg/mL 75.9 (43.9-151) 87.0 (44.9-168) 88.6 (50.2-164) 96.4 (52.6-191) .09 Mean fold change (95% CI) NA NA 2.01 (-41.4 to 58.1) 11.4 (-28.9 to 69.2) .06 IL-18 Total, No. 377 375 320 320 NA IL-18 Jule Jule Jule Jule Jule Jule Jule Jule IL-18 Jule Jule Jule Jule Jule Jule Jule Jule IL-18 level, pg/mL 245 (179	Total, No.	398	391	335	333	NA
Change from baselineNANA0 (-0.95 to 0.77)-0.12 (-0.81 to 0.59).46Mean fold change (95% CI)NANA0.98 (0.89-1.08)0.94 (0.86-1.03)NAIL-1βTotal, No.377375320320NA[L-1β level, fg/mL75.9 (43.9-151)87.0 (44.9-168)88.6 (50.2-164)96.4 (52.6-191).09Change from baselineNANA2.01 (-41.4 to 58.1)11.4 (-28.9 to 69.2).06Mean fold change (95% CI)NANA2.01 (-41.4 to 58.1)1.19 (1.07-1.33)NAIL-18Total, No.377375320320NAIL-18IL-18IL-18 level, pg/mL245 (179-330)232 (169-310)236 (169-310)230 (176-308).79Change from baselineNANA-8.53 (-48.2 to 35.3)-3.77 (-43.3 to 44.1).26Mean fold change (95% CI)NANA0.96 (0.92-1.00)1.01 (0.97-1.04)NA	IL-6 level, pg/mL	1.64 (0.99-2.76)	1.46 (0.98-2.59)	1.64 (0.99-2.73)	1.53 (0.93-2.62)	.24
Mean fold change (95% CI) NA NA 0.98 (0.89-1.08) 0.94 (0.86-1.03) NA IL-1β	Change from baseline	NA	NA	0 (-0.95 to 0.77)	-0.12 (-0.81 to 0.59)	.46
IL-1β Total, No. 377 375 320 320 NA IL-1β level, fg/mL 75.9 (43.9-151) 87.0 (44.9-168) 88.6 (50.2-164) 96.4 (52.6-191) .09 Change from baseline NA NA 2.01 (-41.4 to 58.1) 11.4 (-28.9 to 69.2) .06 Mean fold change (95% CI) NA NA 1.05 (0.93-1.17) 1.19 (1.07-1.33) NA IL-18 Total, No. 377 375 320 320 (176-308) .79 IL-18 level, pg/mL 245 (179-330) 232 (169-310) 236 (169-310) 230 (176-308) .79 Change from baseline NA NA -8.53 (-48.2 to 35.3) -3.77 (-43.3 to 44.1) .26 Mean fold change (95% CI) NA NA 0.96 (0.92-1.00) 1.01 (0.97-1.04) NA	Mean fold change (95% CI)	NA	NA	0.98 (0.89-1.08)	0.94 (0.86-1.03)	NA
Total, No. 377 375 320 320 NA IL-1β level, fg/mL 75.9 (43.9-151) 87.0 (44.9-168) 88.6 (50.2-164) 96.4 (52.6-191) .09 Change from baseline NA NA 2.01 (-41.4 to 58.1) 11.4 (-28.9 to 69.2) .06 Mean fold change (95% CI) NA NA 1.05 (0.93-1.17) 1.19 (1.07-1.33) NA IL-18 Total, No. 377 375 320 320 (176-308) .79 Change from baseline NA NA 2.32 (169-310) 236 (169-310) 230 (176-308) .79 IL-18 level, pg/mL 245 (179-330) RA RA -8.53 (-48.2 to 35.3) -3.77 (-43.3 to 44.1) .26 Mean fold change (95% CI) NA NA 0.96 (0.92-1.00) 1.01 (0.97-1.04) NA	IL-1β					
IL-1β level, fg/mL 75.9 (43.9-151) 87.0 (44.9-168) 88.6 (50.2-164) 96.4 (52.6-191) .09 Change from baseline NA NA 2.01 (-41.4 to 58.1) 11.4 (-28.9 to 69.2) .06 Mean fold change (95% CI) NA NA 1.05 (0.93-1.17) 1.19 (1.07-1.33) NA IL-18 Total, No. 377 375 320 320 NA IL-18 level, pg/mL 245 (179-330) 232 (169-310) 236 (169-310) 230 (176-308) .79 Change from baseline NA NA -8.53 (-48.2 to 35.3) -3.77 (-43.3 to 44.1) .26 Mean fold change (95% CI) NA NA 0.96 (0.92-1.00) 1.01 (0.97-1.04) NA	Total, No.	377	375	320	320	NA
Change from baseline NA NA 2.01 (-41.4 to 58.1) 11.4 (-28.9 to 69.2) .06 Mean fold change (95% CI) NA NA 1.05 (0.93-1.17) 1.19 (1.07-1.33) NA IL-18 Total, No. 377 375 320 320 (176-308) .79 [L-18 level, pg/mL 245 (179-330) 232 (169-310) 236 (169-310) 230 (176-308) .79 Change from baseline NA NA -8.53 (-48.2 to 35.3) -3.77 (-43.3 to 44.1) .26 Mean fold change (95% CI) NA NA 0.96 (0.92-1.00) 1.01 (0.97-1.04) NA	IL-1β level, fg/mL	75.9 (43.9-151)	87.0 (44.9-168)	88.6 (50.2-164)	96.4 (52.6-191)	.09
Mean fold change (95% CI) NA NA 1.05 (0.93-1.17) 1.19 (1.07-1.33) NA IL-18	Change from baseline	NA	NA	2.01 (-41.4 to 58.1)	11.4 (-28.9 to 69.2)	.06
IL-18 Total, No. 377 375 320 320 NA IL-18 level, pg/mL 245 (179-330) 232 (169-310) 236 (169-310) 230 (176-308) .79 Change from baseline NA NA -8.53 (-48.2 to 35.3) -3.77 (-43.3 to 44.1) .26 Mean fold change (95% CI) NA NA 0.96 (0.92-1.00) 1.01 (0.97-1.04) NA	Mean fold change (95% CI)	NA	NA	1.05 (0.93-1.17)	1.19 (1.07-1.33)	NA
Total, No. 377 375 320 320 NA IL-18 level, pg/mL 245 (179-330) 232 (169-310) 236 (169-310) 230 (176-308) .79 Change from baseline NA NA -8.53 (-48.2 to 35.3) -3.77 (-43.3 to 44.1) .26 Mean fold change (95% CI) NA NA 0.96 (0.92-1.00) 1.01 (0.97-1.04) NA	IL-18					
IL-18 level, pg/mL 245 (179-330) 232 (169-310) 236 (169-310) 230 (176-308) .79 Change from baseline NA NA -8.53 (-48.2 to 35.3) -3.77 (-43.3 to 44.1) .26 Mean fold change (95% CI) NA NA 0.96 (0.92-1.00) 1.01 (0.97-1.04) NA	Total, No.	377	375	320	320	NA
Change from baseline NA NA -8.53 (-48.2 to 35.3) -3.77 (-43.3 to 44.1) .26 Mean fold change (95% CI) NA NA 0.96 (0.92-1.00) 1.01 (0.97-1.04) NA	IL-18 level, pg/mL	245 (179-330)	232 (169-310)	236 (169-310)	230 (176-308)	.79
Mean fold change (95% CI) NA NA 0.96 (0.92-1.00) 1.01 (0.97-1.04) NA	Change from baseline	NA	NA	-8.53 (-48.2 to 35.3)	-3.77 (-43.3 to 44.1)	.26
	Mean fold change (95% CI)	NA	NA	0.96 (0.92-1.00)	1.01 (0.97-1.04)	NA

(continued)

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Table 3. Distributions of Inflammatory and Immune Biomarkers at Entry and Month 24 (continued)

	Median (IQR)					
	Entry		Month 24			
Measure ^a	Pitavastatin (n = 402)	Placebo (n = 402)	Pitavastatin (n = 349)	Placebo (n = 350)	P value	
IL-10						
Total, No.	363	353	305	300	NA	
IL-10 level, pg/mL	0.24 (0.17-0.37)	0.22 (0.16-0.35)	0.21 (0.15-0.34)	0.22 (0.16-0.33)	.53	
Change from baseline	NA	NA	-0.02 (-0.12 to 0.06)	0 (-0.09 to 0.09)	.08	
Mean fold change (95% CI)	NA	NA	0.96 (0.87-1.05)	0.99 (0.91-1.07)	NA	
Caspase-1						
Total, No.	377	375	320	320	NA	
Caspase-1 level, pg/mL	66.6 (47.0-93.8)	69.6 (50.1-107)	69.6 (48.6-102)	74.2 (52.6-113)	.08	
Change from baseline	NA	NA	-3.41 (-22.2 to 26.2)	5.61 (-18.6 to 30.5)	.07	
Mean fold change (95% CI)	NA	NA	1.02 (0.94-1.10)	1.07 (0.98-1.16)	NA	

Abbreviations: hsCRP, high-sensitivity C-reactive protein; IL, interleukin; LpPLA2, lipoprotein-associated phospholipase A2; MCP-1, monocyte chemoattractant protein 1; NA, not applicable; oxLDL, oxidized low-density lipoprotein. ^a Data are reported for all enrolled participants with available biomarker results.

^b For hsCRP, censored values below the assay limit are randomly imputed based on a uniform distribution; otherwise, censored values below the assay limit are imputed as the result –0.01; values above the assay limit are imputed as result 0.1.

SI conversion factor: To convert hsCRP to mg/L, multiply by 10.

Figure 2. Effect of Biomarkers on Relative Risk of Progression of Noncalcified Plaque

A Unadjusted

A Unadjusted				
Measure	No.	Estimated difference (95% CI)	Lower Higher risk risk	P value
Change in hsCRP at month 4	499	0.98 (0.95-1.01)	-	.16
Change in hsCRP at month 24	524	0.99 (0.96-1.01)	- -	.32
Change in LpPLA-2 at month 4	505	0.99 (0.89-1.11)	- -	.92
Change in LpPLA-2 at month 24	515	1.01 (0.91-1.12)		.86
Change in oxidized LDL at month 4	505	1.01 (0.92-1.10)	- -	.88
Change in oxidized LDL at month 24	515	0.96 (0.89-1.04)	- 	.35
Change in LDL cholesterol at month 4	501	0.94 (0.85-1.04)		.25
Change in LDL cholesterol at month 24	519	0.91 (0.83-0.99)		.03
LDL cholesterol <70 at month 4	501	0.83 (0.58-1.18)		.30
LDL cholesterol <70 at month 24	519	0.62 (0.41-0.94)		.03
		(D.4 1	2
			Relative risk of progression (95)	% CI)

B Adjusted for NCP at baseline

Maarura	No	Estimated	Lower	Higher	Dyalua
Measure	INO.	uniference (95% CI)	TISK	115K	P value
Change in hsCRP at month 4	499	0.98 (0.96-1.00)	-		.04
Change in hsCRP at month 24	524	0.99 (0.97-1.02)			.65
Change in LpPLA-2 at month 4	505	1.00 (0.91-1.10)	-1		.99
Change in LpPLA-2 at month 24	515	1.02 (0.94-1.11)	-	-	.57
Change in oxidized LDL at month 4	505	1.00 (0.93-1.09)	-	-	.90
Change in oxidized LDL at month 24	515	0.99 (0.92-1.07)	-	-	.81
Change in LDL cholesterol at month 4	501	0.97 (0.89-1.06)	-	-	.50
Change in LDL cholesterol at month 24	519	0.94 (0.86-1.02)	-8-		.15
LDL cholesterol <70 at month 4	501	0.98 (0.73-1.33)			.91
LDL cholesterol <70 at month 24	519	0.79 (0.55-1.14)			.20
			· · · · · · · · ·		_
		0).4	1	2
			Relative risk of progr	ession (95% CI)	

Effect size is per 25% decrease in the biomarker. The *P* value tests for a linearly increasing log relative risk. Mechanistic analyses were limited to the per-protocol population with paired biomarkers and plaque outcomes. hsCRP indicates high-sensitivity C-reactive protein; LDL, low-density lipoprotein; LpPLA-2, lipoprotein-associated phospholipase A2; NCP, noncalcified plaque.

changes in oxLDL, hsCRP, or LpPLA2 and noncalcified plaque progression or change in noncalcified plaque volume were apparent. The association between pitavastatin and changes in noncalcified plaque did not change after adjustment for these biomarkers (eFigures 3 and 4 in Supplement 2).

Discussion

REPRIEVE demonstrated that primary cardiovascular prevention with pitavastatin reduced the risk of MACE by 35% over a median (IQR) of 5.1 (4.3-5.9) years in PWH at low to moderate traditional cardiovascular risk,⁹ an effect beyond that anticipated from LDL cholesterol reduction alone. Within the embedded REPRIEVE mechanistic substudy, the pitavastatin arm demonstrated a relative reduction (treatment effect, -4.3 mm³; 95% CI, -8.6 to -0.1; a 7% [95% CI, 1-12] greater reduction relative to placebo) in noncalcified coronary plaque volume over 2 years, with an estimated 33% lower risk of plaque progression. The pitavastatin arm also had a reduction in LDL cholesterol and biomarkers of lipid oxidation (oxLDL) and arterial inflammation (LpPLA2), although there was not clear association with changes in noncalcified plaque volume. These findings provide insight into how pitavastatin may prevent MACE.

In designing the mechanistic substudy, we sought to assess the effects of a statin intervention on coronary plaque and inflammation in PWH. Coronary CTA allows for noninvasive quantification of changes in plaque in response to therapy.²⁰⁻²² Here, we quantified noncalcified and lowattenuation coronary plaque, which are thought to be the biologically active component of plaque and are associated with MACE.^{18,23} Among PWH, increased noncalcified coronary plaque may be a mechanism of increased CVD.^{12,24-28} Noncalcified plaque was seen in a higher-than-expected percentage of participants at baseline in REPRIEVE, a relatively young primary prevention population with low to moderate traditional CVD risk.¹² The observed reduction in plaque volume may therefore represent an important statin benefit potentially contributing to the reduced MACE observed with statin in REPRIEVE. Studies have shown that relatively small effects on atheroma volume of 1% can lead to large reductions in MACE.²⁹ Other studies suggest decreases in noncalcified plaque with statin therapy may be due in part to conversion to calcified plaque, which may contribute to plaque stabilization.^{30,31} In this study, there were similar increases in calcified plaque volume between arms, while noncalcified plaque volume decreased more in the pitavastatin arm than the placebo arm. Indeed, there was a decrease in the proportion of noncalcified plaque by -4.1% (95% CI, -7.0 to -1.3) in the pitavastatin arm compared with the placebo arm, suggesting relatively greater effects of pitavastatin to reduce noncalcified plague than to increase calcified plaque in this population with low to moderate ASCVD risk.

The observation of a reduction in noncalcified coronary plaque in the pitavastatin arm compared with the placebo arm in REPRIEVE builds on an earlier trial of atorvastatin in PWH,¹⁹ which found that atorvastatin reduced noncalcified plaque volume in 40 PWH with preexisting plaque. The amount of baseline noncalcified plaque and change in noncalcified plaque were relatively small but occurred in a population very different from that investigated in other trials of high-risk cohorts^{20,21} with more risk factors^{32,33} or with established CAD.^{17,34} Consistent with this, the effects of our statin intervention on plaque were larger among those with plaque at baseline in RE-PRIEVE. The reduction in noncalcified plaque volume among those with preexisting plaque in REPRIEVE was similar to that seen in a metanalysis of 3 CTA statin studies.³⁵

While the focus was on the salutary effects of pitavastatin, the placebo arm also provides a valuable look at the natu-

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ral history of CAD in PWH. In this group with low to moderate traditional CVD risk who would not normally qualify for statin, there was on average progression of noncalcified coronary plaque over 2 years. These findings support previous observations of accelerated CAD in PWH,^{19,24,28,36} which, along with the 35% reduction in MACE seen in the parent trial,⁹ build the case for primary prevention in this population.

The pitavastatin arm had a reduction in oxLDL and Lp-PLA2 compared with the placebo arm, in addition to LDL cholesterol. These changes are notable, as specific inflammatory pathways have been hypothesized to increase the risk of CVD in PWH.⁴ Prior studies have shown increased arterial inflammation using fludeoxyglucose-18 positron emission tomography in PWH compared with controls without HIV in the general population,³⁷ linking arterial inflammation to noncalcified plaque among PWH.³⁸ In this regard, LpPLA2 is a plateletactivating factor that hydrolyzes oxidized lipids, thought to play a proinflammatory role in atherogenesis and is a marker of arterial inflammation in PWH.^{39,40} In contrast, markers of generalized inflammation, such as CRP, may not be increased in well-treated PWH taking antiretroviral therapy, potentially explaining the more modest changes in CRP observed in our study. oxLDL is a marker of oxidative stress, which increases the atherogenic properties of LDL cholesterol and induces foam cell formation through localized macrophage activation and endothelial activation, key processes in plaque formation.⁴¹ In contrast, no significant effects on immune pathways were found.

Among studies in the general population, there is an overall association between achieved LDL cholesterol and reductions in coronary plaque volume, though considerable heterogeneity exists at the level of the individual study.²² Similarly, in REPRIEVE, there was a suggestion that achieved LDL cholesterol less than 70 mg/dL at 24 months was associated with a lower risk of noncalcified plaque progression in unadjusted analyses. Our population was unusual, as it was composed of PWH at low to moderate CVD risk with lower entry LDL cholesterol levels in contrast to other higher-risk populations with higher LDL cholesterol levels, in which statin effects on LDL may relate more strongly to changes in plaque. Decreases in systemic markers of lipid oxidation and arterial inflammation were not significantly associated with the changes in noncalcified plaque. However, effects of statins may modify inflammatory pathways through lipid effects occurring at the local plaque level, leading to reductions in inflammation in the plaque microenvironment as a mechanism of their antiinflammatory effect.⁴² Moreover, changes in inflammatory pathways may be related to MACE, even if not related to changes in plaque volume, a hypothesis that will be further tested in the main REPRIEVE population. Further research is needed into the mechanism by which statins reduce noncalcified plaque among PWH, including potential effects related to atheroma formation, lipid uptake, and foam cell formation. Other effects of statins at the plaque surface to stabilize plaque, reduce inflammation within the plaque, and reduce plaque rupture, may be critical to preventing MACE in PWH. The data from REPRIEVE add to our growing understanding of the potential mechanisms of statin effects on CVD.⁴³

Limitations

Limitations of this study should be considered. Although our results are specific to pitavastatin, other statins may have similar effects. REPRIEVE was an international trial with 145 sites from 12 countries; the mechanistic substudy necessarily included a more limited population from 31 US sites and thus may not be representative of other regions. The substudy population was generally similar to the overall trial population in terms of ASCVD risk but included a higher proportion of male participants (eTable 1 in Supplement 2). A multiple imputation analysis addressed missing data and supported the overall conclusion of the study. The results may not be generalizable to

higher-risk populations, with higher baseline LDL cholesterol and more plaque.

Conclusions

In conclusion, in a primary prevention trial of PWH at low to moderate ASCVD risk, 2 years of pitavastatin reduced noncalcified coronary plaque volume and inflammatory biomarkers. These effects may contribute to the observed effects of pitavastatin on MACE in the parent REPRIEVE trial.

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