Relation of Coronary Artery Calcium to Flow-Mediated Dilation and C-Reactive Protein Levels in Asymptomatic Patients With Heterozygous Familial Hypercholesterolemia

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The extent of coronary artery calcium (CAC) is correlated with coronary artery disease prognosis. However, the relation of CAC to endothelial function and high-sensitivity C-reactive protein (hs-CRP) in patients with asymptomatic heterozygous familial hypercholesterolemia (FH) requires clarification. The study aim was to investigate the relations among CAC, endothelial function, and hs-CRP in patients with asymptomatic heterozygous FH. Thirty-two patients with asymptomatic heterozygous FH (mean age 42 years) and 34 healthy control subjects (mean age 36 years) were enrolled. We measured CAC by electron-beam computed tomography and endothelial function by flow-mediated dilation of the brachial artery. A higher percentage of patients with FH had a positive CAC score compared with the control group. Comparing the FH group with detectable CAC (CAC score >0) and undetectable CAC (CAC score of 0), we found higher hs-CRP levels (0.29 \pm 0.23 vs 0.07 \pm 0.08 mg/dl, p = 0.001) and reduced flow-mediated dilation (0.04 \pm 0.03 vs 0.08 ± 0.03 , p = 0.005) in the detectable CAC group. Multivariate analysis showed an independent correlation of hs-CRP with detectable CAC (relative risk 5.034, 95% confidence interval 1.525 to 16.613, p = 0.04). In conclusion, FH subjects with positive CAC scores have decreased flow-mediated dilation and increased hs-CRP levels. Furthermore, hs-CRP level is the only independent predictor of the presence of CAC. © 2007 Elsevier Inc. All rights reserved. (Am J Cardiol 2007;100:1119-1123)

Familial hypercholesterolemia (FH) is an inherited autosomal dominant disorder that affects approximately 1 in 500 persons worldwide and is the most common inborn error of cholesterol metabolism.¹ Coronary artery calcium (CAC), measured by electron-beam computed tomography, has been widely used to assess the extent of atherosclerosis and predict the prognosis of coronary artery disease.² Hoffmann et al³ have demonstrated that CAC is more extensive in cases of coronary artery disease in patients with FH, presumably resulting from a long history of increased lowdensity lipoprotein cholesterol (LDL-C) levels. However, the pathophysiology of vascular calcium deposition is unclear, and its relation to inflammation markers needs further clarification.^{4,5} In the present study, we tested the hypothesis that a high cholesterol burden would lead to increased high-sensitivity C-reactive protein (hs-CRP) and endothelial dysfunction, and could thereby contribute to CAC in asymptomatic patients with heterozygous FH.

Methods

Patients with LDL-C levels >190 mg/dl and positive family history of hypercholesterolemia were enrolled in a genetic screening program for FH in Taiwan. Methodology used for mutation detection has been described before.⁶ Family members who were genetically verified as not carrying an FH-related mutation allele and with LDL-C levels <130 mg/dl were enrolled as the control group. The exclusion criteria were age >55 years and history of medication for dyslipidemia, hypertension, or hyperglycemia. Patients with history of angina or coronary artery disease were also excluded. The study protocol was approved by the institutional review board, and informed consent was obtained from each patient.

Demographic and clinical parameters including age, height, weight, body mass index, waist circumference, systolic blood pressure, diastolic blood pressure, and smoking habits were collected. After 12 to 14 hours overnight fasting, total plasma cholesterol, high-density lipoprotein cholesterol, and triglyceride were measured using commercially available kits (Roche, Mannheim, Germany). LDL-C was calculated according to the formula of Friedewald et al.⁷ The concentrations of apolipoprotein A1 and apolipoprotein B were measured by nephelometry (Behring Diagnostic, Mar-

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Table 1
Comparison of subjects with or without heterozygous familial hypercholesterolemia (FH) mutation

Variable	Control Group (n = 34)	FH Group $(n = 32)$	p Value
Age (yrs)	42.1 ± 17.0	36.0 ± 17.8	0.165
Men/women	17/17	18/14	0.618
Body weight (kg)	58.3 ± 15.2	59.6 ± 17.6	0.792
Body mass index (kg/m ²)	23.4 ± 3.9	22.1 ± 4.9	0.243
Waist circumference (cm)	78.3 ± 8.5	78.6 ± 11.6	0.921
Systolic blood pressure (mm Hg)	109.5 ± 11.0	115.8 ± 14.4	0.106
Diastolic blood pressure (mm Hg)	65.0 ± 9.8	67.3 ± 10.8	0.448
Fasting blood glucose (mg/dl)	101.2 ± 30.7	93.0 ± 8.1	0.152
Total cholesterol (mg/dl)	196.6 ± 38.1	299.7 ± 73.7	< 0.001
Total triglyceride (mg/dl)	132.5 ± 65.0	132.9 ± 76.1	0.979
LDL-C (mg/dl)	120.0 ± 38.1	218.4 ± 66.1	< 0.001
HDL-C (mg/dl)	51.2 ± 18.3	56.5 ± 12.2	0.178
Total cholesterol/HDL-C	4.1 ± 0.9	5.5 ± 1.5	< 0.001
Cholesterol-year score	8,277.3 ± 3,391.1	$11,077.0 \pm 6,234.5$	0.026
Apolipoprotein B (mg/dl)	99.2 ± 30.3	151.5 ± 41.2	< 0.001
Apolipoprotein A1 (mg/dl)	132.0 ± 20.0	133.1 ± 30.8	0.897
hs-CRP (mg/dl)	0.11 ± 0.14	0.12 ± 0.17	0.175
Flow-mediated dilation	0.08 ± 0.03	0.06 ± 0.04	0.023
CAC score	0.5 ± 1.5	59.1 ± 19.4	0.048
Patients with positive CAC score	3 (8.8%)	10 (31.2%)	0.022

Values presented as means \pm SD where applicable.

HDL-C = high-density lipoprotein cholesterol.

burg, Germany). Blood glucose was measured using an enzymatic method. A highly sensitive latex-based immunoassay (Dade Behring, Milton Keynes, United Kingdom) was used for hs-CRP. Cholesterol–year score, which is dependent on cholesterol exposure duration and severity in people with FH, was selected to reflect the total burden of hypercholesterolemia.⁸ The score is computed by multiplying the initial serum cholesterol value (in milligrams per deciliter) by the age at diagnosis and then adding the annually determined total cholesterol values.

Endothelium dependent flow-mediated dilation was assessed using a 7.5-MHz linear array transducer (Sonos 5500; Hewlett-Packard; Andover, Massachusetts) to scan the brachial artery in longitudinal section. All subjects were asked to fast, refrain from smoking, and withhold all medications for 12 hours. The procedure was performed in a quiet, air-conditioned room (22°C to 25°C). The left arm was stabilized with a cushion, and a sphygmomanometric cuff was placed on the forearm. A baseline image was acquired, and blood flow was estimated by time-averaging the pulsed Doppler velocity signals obtained from a sample volume at the middle of the artery. The cuff was then inflated to \geq 50 mm Hg greater than systolic pressure to occlude the arteries for 5 minutes and released abruptly. Postocclusion diameters were obtained at 60, 80, 100, and 120 seconds after deflation. Flow-mediated dilation was calculated as the maximum postocclusion diameter relative to the average diameter before occlusion. A mid-artery pulsed Doppler signal was obtained immediately after cuff release and no later than 15 seconds after cuff deflation to assess hyperemic velocity.4

Electron-beam computed tomography was performed to measure CAC. A C-150 LXP scanner (Imatron, South San Francisco, California) calibrated daily with air and water phantoms and twice monthly with contrast medium and resolution phantoms was used. Images were obtained using a 40- to 50-slice (3-mm thickness) protocol with image acquisition triggered to 60% to 80% of the electrocardiographic RR interval while respirations were held. Scans were interpreted by an experienced radiologist blinded to the participant's group using the scoring method of Agatston et al.⁹ A focus of coronary calcium was defined as the presence of \geq 4 contiguous pixels with >130 Hounsfield units. Total CAC score was determined as the sum of individual scores of 4 major epicardial coronary arteries.¹⁰

All results were expressed as means with SDs for continuous variables and proportions for categorical variables. Comparisons between groups were performed using the Wilcoxon 2-sample test. Logarithmically transformed hs-CRP values were used in statistical analysis to achieve a normal distribution. Spearman correlation coefficients were determined to assess association among CAC score, flow-mediated dilation, plasma lipid profile, lipoprotein, blood pressure, and hs-CRP. To test whether variables had an independent effect on hs-CRP and CAC, stepwise logistic regression analysis was performed. All tests were 2-sided, and p values <0.05 were considered statistically significant.

Results

A total of 32 untreated participants genetically verified as being heterozygous for FH (mean age 36 ± 17.8 years, 56% male) and 34 family members (mean age 42.1 ± 17.0 years, 50% male) genetically verified as not carrying an FH-related allele were enrolled in this study. All were of

Table 2 Spearman correlation coefficients in 32 subjects with heterozygous familial hypercholesterolemia

Variable	CAC	Score
	R	p Value
Age	0.469	0.009
Systolic blood pressure	0.415	0.022
Diastolic blood pressure	0.375	0.041
Body weight	0.427	0.018
Body mass index	0.564	0.001
Waist circumference	0.490	0.006
Fasting blood glucose	0.130	0.495
Total cholesterol	0.206	0.275
Total triglyceride	0.493	0.006
LDL-C	0.171	0.350
HDL-C	-0.200	0.290
Total cholesterol/HDL-C	0.393	0.032
Cholesterol-year score	0.478	0.008
Apolipoprotein B	0.357	0.053
Apolipoprotein A1	-0.046	0.808
hs-CRP	0.572	0.001
Flow-mediated dilation	-0.458	0.001

Abbreviation as in Table 1.

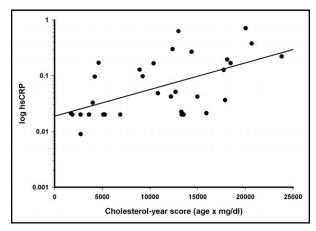


Figure 1. Semi-log scatterplot for correlation between log hs-CRP (y axis) and cholesterol–year score (x axis). Cholesterol–year scores are correlated with hs-CRP (r = 0.574, p = 0.001).

Chinese descent. The characteristics of the mutations of the LDL receptor have been described before.⁶ None of the participants had a history of cardiovascular disease, hypertension, or diabetes mellitus. Participant characteristics are listed in Table 1. Compared with healthy subjects, the FH subjects had significantly increased total cholesterol, LDL-C, and apolipoprotein B. The cholesterol–year scores were higher in the FH group. Subjects with FH had decreased flow-mediated dilation and remarkably higher CAC scores. A higher percentage of subjects with FH in this series had a positive CAC score compared with the control group. Serum levels of hs-CRP were higher in the FH group, but the difference did not reach statistical significance.

As the Spearman correlation analysis shows in Table 2, the CAC scores of patients with FH were correlated with age, systolic blood pressure, cholesterol-year score, body weight, body mass index, waist circumference, total triglyceride, hs-CRP, and flow-mediated dilation. As presented in Figure 1, cholesterol-year score was highly correlated with hs-CRP (r = 0.574, p = 0.001). Cholesterol-year score was also correlated with CAC score (r = 0.478, p = 0.008), even though only 10 subjects with FH had detectable CAC.

The 32 patients with FH-related mutations were further divided into 2 groups according to their CAC score: group 1 (n = 22) consisted of patients with FH without detectable CAC (CAC score of 0), and group 2 (n = 10) consisted of patients with FH with detectable CAC (CAC score >0). Table 3 compares baseline and clinical parameters between these 2 groups. The patients with FH with detectable CAC were older and had higher systolic/diastolic blood pressure and higher cholesterol-year scores than patients with undetectable CAC. The body weight, body mass index, and waist circumference were also increased in patients with detectable CAC. As Figure 2 shows, patients with FH with detectable CAC had significantly increased hs-CRP values (p < 0.001) and remarkably impaired flow-mediated dilation (p = 0.005). By stepwise logistic regression analysis adjusted for age, total cholesterol, LDL-C, systolic/diastolic blood pressure, body mass index, waist circumference, and other conventional risk factors, hs-CRP value was an independent predictor for the presence of CAC in subjects with FH (relative risk 5.034, 95% confidence interval 1.525 to 16.613, p = 0.04).

Discussion

In the present study, we demonstrated that subjects with asymptomatic heterozygous FH with positive CAC scores have decreased flow-mediated dilation and significantly increased levels of hs-CRP. On stepwise logistic regression analysis, hs-CRP is the only independent predictor for the presence of CAC in patients with FH.

Although CAC and hs-CRP are proven predictors of subclinical atherosclerosis,^{2,5} the correlation between CAC and hs-CRP results remains obscure.11-13 Huang et al4 reported that CAC was strongly related to endothelial dysfunction but was not related to hs-CRP in patients with suspected coronary artery disease. Compared with participants in other studies, our patients with FH were relatively healthy and untreated, and hypercholesterolemia was their only modifiable risk factor. Their atherosclerotic process, initiated by high cholesterol burden, was probably in the active inflammatory stage. In the absence of other confounding cardiovascular risk factors, our study clearly showed that, in patients with FH, the CAC score correlated not only with cholesterol burden but also with hs-CRP values. Previous reports have also demonstrated the relations between inflammatory markers and vascular injury in FH.14,15 In addition, stepwise logistic regression analysis showed that the hs-CRP level was an independent predictor for the presence of CAC in FH. Zouridakis et al¹⁶ demonstrated that increased C-reactive protein and other mediators of inflammation signal rapid progression of coronary artery disease in patients with stable angina pectoris. Therefore, our study supports the notion that the mechanism for initiating coronary artery calcification in

Comparisons of basic characteristics of patients with heterozygous familial hypercholesterolemia (FH) with versus without coronary artery calcium (CAC)

Variable	Group 1 (CAC score of 0; $n = 22$)	Group 2 (CAC score >0 ; n = 10)	p Value
Age (yrs)	32.0 ± 15.6	49.2 ± 18.3	0.014
Men/women	12/10	6/4	0.878
Systolic blood pressure (mm Hg)	112.9 ± 11.8	126.3 ± 15.7	0.015
Diastolic blood pressure (mm Hg)	65.1 ± 8.7	74.8 ± 12.5	0.022
Body weight (kg)	54.3 ± 14.2	73.2 ± 18.8	0.004
Body mass index (kg/m ²)	20.4 ± 3.7	26.4 ± 5.0	0.001
Waist circumference (cm)	75.0 ± 9.8	87.8 ± 11.0	0.003
Fasting blood glucose (mg/dl)	92.9 ± 7.0	94.6 ± 10.7	0.619
Total cholesterol (mg/dl)	288.8 ± 76.0	329.7 ± 70.9	0.180
Total triglyceride (mg/dl)	118.4 ± 77.3	171.7 ± 69.5	0.086
LDL-C (mg/dl)	210.2 ± 64.7	240.4 ± 67.6	0.270
HDL-C (mg/dl)	57.9 ± 14.1	53.2 ± 7.6	0.357
Total cholesterol/HDL-C	5.2 ± 1.5	6.3 ± 1.2	0.062
Cholesterol-year score	$9597.0 \pm 5,397.5$	$15,924.0 \pm 5,890.0$	0.008
Apolipoprotein B (mg/dl)	141.7 ± 40.3	178.2 ± 39.7	0.030
Apolipoprotein A1 (mg/dl)	136.0 ± 32.0	132.6 ± 29.4	0.786
hs-CRP (mg/dl)	0.07 ± 0.08	0.29 ± 0.23	< 0.001
Flow-mediated dilation	0.08 ± 0.03	0.04 ± 0.03	0.005

Values presented as means \pm SD where applicable.

Abbreviation as in Table 1.

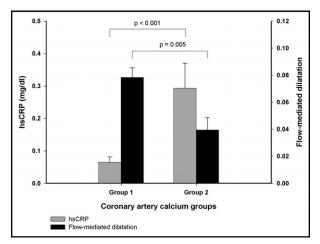


Figure 2. Comparisons of hs-CRP and flow-mediated dilation between groups 1 and 2 in patients with heterozygous FH. Patients with FH with detectable CAC have significantly increased hs-CRP values (p < 0.001) and remarkably impaired flow-mediated dilation (p = 0.005). Group 1 (n = 22) comprises patients with no detectable CAC (CAC score of 0); group 2 (n = 10) comprises those with detectable CAC (CAC score >0).

FH is vascular inflammation induced by a high cholesterol burden.¹⁷

Although it is difficult to infer a causal relation without a longitudinal follow-up study, our results clearly indicate the association between risk factors of hypercholesterolemia and increased hs-CRP level and the intermediate outcomes of CAC and flow-mediated dilation in heterozygous patients with FH. Despite a relatively small sample size, we still identified close correlations between these measurements in patients with heterozygous FH.

In conclusion, compared with patients with FH without detectable CAC scores, subjects with asymptomatic heterozygous FH with detectable CAC have increased hs-CRP and impaired endothelial function. Furthermore, the hs-CRP level is the only independent predictor of the presence of CAC. CAC is correlated with hs-CRP and flow-mediated dilation in patients with asymptomatic heterozygous FH.¹¹

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