# **ORIGINAL RESEARCH**

# Antiatherosclerotic Phenotype of Perivascular Adipose Tissue Surrounding the Saphenous Vein in Coronary Artery Bypass Grafting

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**BACKGROUND:** Perivascular adipose tissue (PVAT) is associated with metabolically driven chronic inflammation called metaflammation, which contributes to vascular function and the pathogenesis of vascular disease. The saphenous vein (SV) is commonly used as an essential conduit in coronary artery bypass grafting, but the long-term patency of SV grafts is a crucial issue. The use of the novel "no-touch" technique of SV harvesting together with its surrounding tissue has been reported to result in good long-term graft patency of SV grafts. Herein, we investigated whether PVAT surrounding the SV (SV-PVAT) has distinct phenotypes compared with other PVATs of vessels.

**METHODS AND RESULTS:** Fat pads were sampled from 48 patients (male/female, 32/16; age, 72±8 years) with coronary artery disease who underwent elective coronary artery bypass grafting. Adipocyte size in SV-PVAT was significantly larger than the sizes in PVATs surrounding the internal thoracic artery, coronary artery, and aorta. SV-PVAT and PVAT surrounding the internal thoracic artery had smaller extents of fibrosis, decreased gene expression levels of fibrosis-related markers, and less metaflammation, as indicated by a significantly smaller extent of cluster of differentiation 11c–positive M1 macrophage infiltration, higher gene expression levels of inflammatory cytokines, than did PVATs surrounding the coronary artery and aorta. Expression patterns of adipocyte developmental and pattern-forming genes were totally different among the PVATs of the vessels.

**CONCLUSIONS:** The phenotype of SV-PVAT, which may result from inherent differences in adipocytes, is closer to that of PVAT surrounding the internal thoracic artery than that of PVAT surrounding the coronary artery or that of PVAT surrounding the aorta. SV-PVAT has less metaflammation and consecutive adipose tissue remodeling, which may contribute to high long-term patency of grafting when the no-touch technique of SV harvesting is used.

Key Words: coronary artery bypass surgery 
epicardial fat 
perivascular fat 
saphenous vein

Perivascular adipose tissue (PVAT) plays a role in supporting vascular structure and metabolism, and it contributes to vascular function and the pathogenesis of vascular disease.<sup>1</sup> PVAT is implicated in the regulation of vascular tone by releasing adiposederived relaxing and contracting factors, which contribute to the protection and progression of atherosclerotic vascular disease.<sup>2–5</sup> It has been reported that epicardial

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## **CLINICAL PERSPECTIVE**

## What Is New?

- Perivascular adipose tissue surrounding the saphenous vein had less metabolically driven chronic inflammation called metaflammation, smaller extents of fibrosis, and decreased gene expression levels of inflammatory and fibrosisrelated markers than did perivascular adipose tissue surrounding the coronary artery and that surrounding the aorta.
- The phenotype of perivascular adipose tissue surrounding the saphenous vein was close to that of perivascular adipose tissue surrounding the internal thoracic artery, an atherosclerosisresistant vessel.

## What Are the Clinical Implications?

• The phenotype of perivascular adipose tissue surrounding the saphenous vein might be linked to the high rate of long-term patency of coronary artery bypass grafting using the "notouch" technique of saphenous vein harvesting together with its surrounding tissue and intact endothelium.

## Nonstandard Abbreviations and Acronyms

Ao-PVAT	perivascular adipose tissue surrounding the aorta				
ASC	apoptosis-associated speck-like protein containing caspase recruitment domain				
CA-PVAT	perivascular adipose tissue				
CHOP GRP78 ITA ITA-PVAT	surrounding the coronary artery C/EBP homologous protein glucose-regulated protein 78 internal thoracic artery perivascular adipose tissue surrounding the internal thoracic				
MCP-1 PVAT	monocyte chemoattractant protein-1 perivascular adipose tissue				
SV SV-PVAT	saphenous vein perivascular adipose tissue surrounding the saphenous vein				

adipose tissue is a source of inflammatory mediators<sup>6</sup> and that an elevated volume of PVAT surrounding the coronary artery (CA-PVAT) is strongly associated with coronary atherosclerosis<sup>7-9</sup> and vasospastic angina<sup>10</sup> in combination with metabolically driven chronic inflammation called metaflammation<sup>11</sup> and an imbalance of cytokines and adipokines.

The internal thoracic artery (ITA) is protected from the development of atherosclerosis<sup>12</sup> and is frequently used as the gold standard graft material for coronary artery bypass grafting (CABG) because of its excellent long-term patency.<sup>13</sup> It has been reported that the ITA has less concentric atheromatous intimal thickening involving the whole circumference than does the right gastroepiploic artery, a candidate of graft vessels for CABG, even in patients with several coronary risk factors.<sup>14</sup> Furthermore, excellent endothelial function, including shear stress sensing and generation of NO and antithrombotic factors, of the ITA has been reported to provide physiological and metabolic effects that are beneficial in both the graft and recipient coronary artery.<sup>12,13</sup> We previously proposed that PVAT surrounding the ITA (ITA-PVAT) is protected from metaflammation and consecutive adipose tissue remodeling, possibly contributing to the decreased atherosclerotic plaque burden in the ITA.<sup>15</sup>

The saphenous vein (SV) is still commonly used as an essential conduit in CABG. However, the long-term patency of an SV graft is one of the most crucial issues in cardiovascular surgery.<sup>13</sup> It has recently been proposed that the "no-touch" technique should be used to harvest a complete SV together with its surrounding tissue and intact endothelium.<sup>16</sup> The use of the novel no-touch technique of SV harvesting has been reported to result in superior short- and long-term graft patency rates than those achieved by using the conventional technique of SV harvesting, and the rates were comparable to those for ITA grafts.<sup>17–19</sup> Supply of nutrients and oxygen to the SV wall and its surrounding tissue, including the vasa vasorum,<sup>20</sup> and preservation of endothelial integrity and luminal NO synthase derived from the intact endothelium<sup>21,22</sup> have been proposed as possible mechanisms of the benefit of the no-touch technique.

PVAT can be a source of several vasoactive factors.<sup>23,24</sup> However, the difference between PVAT surrounding the SV (SV-PVAT) and other PVATs of vessels remains unclear. We hypothesized that SV-PVAT has less metaflammation and smaller extents of fibrosis than do other PVATs of vessels, possibly contributing to the protection against atherosclerotic plaque burden in the vessel when no-touch SV is used as a graft material in CABG. In the present study, we investigated whether SV-PVAT, ITA-PVAT, CA-PVAT, and PVAT surrounding the aorta (Ao-PVAT) have distinct histological and gene expression phenotypes and functional relevance to vascular remodeling in patients with coronary artery disease who have undergone elective CABG.

## **METHODS**

This study conformed to the principles outlined in the Declaration of Helsinki and was performed with the approval of the Ethical Committee of Sapporo Medical University (No. 312-34). Written informed consent was received from all of the study subjects. The data that support the findings of this study are available from the corresponding author on reasonable request.

## **Study Patients**

Patients diagnosed with coronary artery disease who underwent elective CABG surgery were consecutively recruited from Sapporo Medical University Hospital and the affiliated Hokkaido Ohno Memorial Hospital, Sapporo Central Hospital, Sapporo City General Hospital, Teine Keijinkai Hospital, and Otaru City General Hospital from March 2018 through March 2020. Patients treated with hemodialysis and those who underwent CABG without using the SV were excluded. The patients in the present study were not overlapped with the patients in our previous study in which PVAT was investigated.<sup>15</sup>

## **Collection of Fat Pads**

Fat pads of SV-PVAT from the area of the lower thigh SV, ITA-PVAT from the area of the left ITA, CA-PVAT from the area of the left coronary artery, and Ao-PVAT from the area of the ascending aorta were collected during surgery before anticoagulation and establishment of extracorporeal circulation. The samples were divided into 2 parts. The first specimen was stored in formalin and embedded in paraffin for histological staining, and the second specimen was frozen in liquid nitrogen for mRNA isolation and real-time polymerase chain reaction (PCR).

## **Histological Analysis**

Samples of fat pads were fixed in 10% formalin solution, dehydrated, and embedded in paraffin. Serial 5-µm-thick sections were prepared, mounted onto glass slides, deparaffinized, and rehydrated through degraded ethanol. Tissues were stained with hematoxylin-eosin and Masson trichrome reagents. Immunohistochemical 3,3'-diaminobenzidine staining using mouse anti–cluster of differentiation (CD) 68 (Dako, Santa Clara, CA), rabbit anit-CD11c (Abcam, Cambridge, UK), and mouse anti-CD206 (Abnova, Taipei, Taiwan) antibodies was performed as previously described.<sup>15,25,26</sup> Control experiments were performed by omitting the primary antibodies or adding goat anti-mouse or goat anti-rabbit immunoglobulins (Dako) for control primary antibodies (Figure S1).

## **Quantitative Image Analysis**

Images were captured with a microscope (BZ-X700; Keyence, Osaka, Japan). Image analyses were performed using ImageJ and Fiji software.

The cross-sectional area of an adipocyte in hematoxylin-eosin staining was determined by calculating the mean area of 100 randomly selected adipocytes per optical field at x200 magnification. The results were averaged, and they are expressed as adipocyte size  $(\mu m^2)$ . After checking sections of hematoxylin-eosin staining to avoid nonspecific fibrosis staining in the adventitia, sections were chosen to stain the fibrosis area in the inside, but not the edge, of adipose tissues. Fibrosis was assessed in Masson trichrome-stained tissue sections by quantifying the blue area representing collagen relative to the total tissue area at ×100 magnification. Furthermore, macrophage infiltration in tissue sections was assessed by quantifying the area stained by anti-CD68, CD11c, or CD206 antibodies relative to the total tissue area at ×200 magnification. For each quantitative analysis of fibrosis and macrophage infiltration, 3 randomly selected optical fields were examined per adipose tissue depot, and the results were averaged. Results are expressed as percentage positive area per total area. All of the measurements were performed in a double-blind manner by 2 different researchers.

## **Quantitative Real-Time PCR**

Total RNA was isolated from samples using the miRNeasy Micro Kit (QIAGEN, Hilden, Germany) and the RNase-Free DNase Set (QIAGEN). The amount and quality of isolated RNA were determined spectrophotometrically using Nanodrop (Thermo Fisher Scientific, Waltham, MA), and 500 ng of total RNA was reverse transcribed by using the high-capacity cDNA archive kit Reverse Transcription Kit (Applied Biosystems, Foster City, CA). Quantitative real-time PCR analysis was performed using SYBR Green in the real-time PCR system (Applied Biosystems, Warrington, UK). The thermal cycling program was 10 minutes at 95°C for enzyme activation and 40 cycles of denaturation for 15 seconds at 95°C, 30 seconds of annealing at 58°C, and 30 seconds of extension at 72°C. Primers used in the present study are listed in Table S1. The specificity of the PCR of genes was confirmed by one distinct band, which matched with the predicted size of the PCR amplicon (Figure S2). To normalize expression data, 18s rRNA was used as an internal control gene. Results are shown as relative expression of each target gene in SV-PVAT of each patient.

## **Statistical Analysis**

Numeric variables are expressed as means±SEM for quantitative real-time PCR data, means±SD for normal

distributions, and medians (interquartile ranges) for skewed variables. One-way repeated-measures ANOVA and paired *t*-test were used for detecting significant differences in data between SV-PVAT, ITA-PVAT, CA-PVAT, and Ao-PVAT, because a generalized linear mixed model analysis showed that there was no statistically significant interaction of any fat pad measure with age, sex, body mass index, or comorbidities, including diabetes mellitus, hypertension, dyslipidemia, and myocardial infarction, in the present study (data not shown). P<0.05 was considered statistically significant. All data were analyzed by using JMP15.2.1 for Macintosh (SAS Institute, Cary, NC).

## RESULTS

# Basal Characteristics of the Studied Patients

Basal characteristics of the 48 recruited patients (male/ female, 32/16) who underwent elective CABG surgery using the SV are shown in Table 1. Age and body mass index of the recruited patients were  $72\pm8$  years and  $23.8\pm3.7$  kg/m<sup>2</sup>, respectively. The numbers of patients with current and former smoking habits were 7 (14.6%) and 31 (64.6%), respectively. Most of the patients (91.7%) were diagnosed with multivessel coronary artery disease. The numbers of patients with diabetes mellitus, hypertension, dyslipidemia, and myocardial infarction were 21 (43.8%), 39 (81.3%), 42 (87.5%), and 9 (18.6%), respectively. The ejection fraction assessed by echocardiography was  $55\pm12\%$ . Data for laboratory measurements are shown in Table 2.

## Adipocyte Sizes in Fat Pads

Representative hematoxylin-eosin staining of SV-PVAT, ITA-PVAT, CA-PVAT, and Ao-PVAT is shown in Figure 1A. Adipocyte size in SV-PVAT was significantly larger by 47%≈57% than that in the other PVATs of vessels (Figure 1B). There was no significant difference in adipocyte size among ITA-PVAT, CA-PVAT, and Ao-PVAT. Crown-like structures, which have been reported to be composed of macrophages surrounding dead or dying adipocytes,<sup>27</sup> were observed in CA-PVAT and Ao-PVAT but not in SV-PVAT or ITA-PVAT.

## **Fibrosis in Fat Pads**

Representative Masson trichrome staining of PVAT is shown in Figure 2A. Samples for this fibrosis analysis were available from 28 of the 48 patients, and basal clinical characteristics in this subgroup of patients were similar to those in the entire study subjects (data not shown). The fibrosis areas in the inside, but not the edge, of adipose tissues were significantly larger in CA-PVAT and Ao-PVAT than in SV-PVAT and ITA-PVAT,

#### Table 1. Characteristics of the Recruited Patients

Characteristics	Values	
No. (men/women)	48 (32/16)	
Age, y	72±8	
Body mass index	23.8±3.7	
Waist circumference, cm	86.0±8.6	
Systolic blood pressure, mm Hg	128±17	
Diastolic blood pressure, mm Hg	67±15	
Pulse rate, beats/min	70±9	
Smoking habit, current/former/never	7 (14.6)/31 (64.6)/10 (20.8)	
Drinking habit	18 (37.5)	
Coronary artery disease		
1-Vessel disease	4 (8.3)	
2-Vessel disease	15 (31.3)	
3-Vessel disease	29 (60.4)	
Complications		
Diabetes mellitus	21 (43.8)	
Hypertension	39 (81.3)	
Dyslipidemia	42 (87.5)	
Myocardial infarction	9 (18.6)	
Medications		
Oral antidiabetic drugs	16 (33.3)	
Insulin	3 (6.3)	
ACEI or ARB	21 (45.7)	
β-Blocker	26 (54.2)	
Statin	38 (79.2)	
Antiplatelet drugs	38 (79.2)	
Echocardiography		
Ejection fraction, %	55±12	

Variables are expressed as number (percentage) or means±SD. ACEI indicates angiotensin-converting enzyme inhibitor; and ARB, angiotensin receptor blocker.

whereas the fibrosis area was significantly larger in Ao-PVAT than in CA-PVAT (Figure 2B). There was no significant difference in the fibrosis area between SV-PVAT and ITA-PVAT. CA-PVAT and Ao-PVAT had significantly higher gene expression levels of fibrosis-related molecules, transforming growth factor- $\beta$ , tissue inhibitor of metalloproteinase 1, and macrophage-inducible C-type lectin, than did SV-PVAT and ITA-PVAT (Figure 2C). The gene expression level of another fibrosis-related molecule, platelet-derived growth factor subunit B, in CA-PVAT and that in Ao-PVAT (Figure 2C).

### Macrophage Infiltration in Fat Pads

Representative immunohistological staining, including CD68, a marker of macrophages, CD11c, a marker of M1 macrophages, and CD206, a marker of M2 macrophages, of PVAT is shown in Figure 3A. The macrophage infiltration area quantified by the ....

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Table 2.	Laboratory Measurements					

Measures	Values		
Hemoglobin, g/dL	11.5±2.0		
AST, IU/L	26 (18–35)		
ALT, IU/L	21 (13–36)		
γGTP, IU/L	27 (17–40)		
Blood urea nitrogen, mg/dL	17 (14–23)		
Creatinine, mg/dL	0.96 (0.76–1.25)		
eGFR, mL/min per 1.73 m <sup>2</sup>	54.5±19.3		
Uric acid, mg/dL	5.6±1.6		
Total cholesterol, mg/dL	141±45		
LDL cholesterol, mg/dL	80±32		
HDL cholesterol, mg/dL	42±11		
Triglycerides, mg/dL	95 (59–126)		
Fasting glucose, mg/dL	106 (93–129)		
Insulin, µU/mL	5.1 (2.7–8.7)		
HOMA-R	1.22 (0.62–3.15)		
HbA1c, %	6.1±1.1		
hs-CRP, mg/dL	0.14 (0.06–0.31)		
NT-proBNP, pg/mL	288 (104–1004)		

Variables are expressed as means±SD or medians (interquartile ranges). ALT indicates alanine transaminase; AST, aspartate transaminase; eGFR, estimated glomerular filtration rate; yGTP, y-glutamyl transpeptidase; HbA1c, hemoglobin A1c; HDL, high-density lipoprotein; HOMA-R, homeostasis model assessment of insulin resistance; hs-CRP, high-sensitivity C-reactive protein; LDL, low-density lipoprotein; and NT-proBNP, N-terminal pro-B-type natriuretic peptide.

anti-CD68 antibody was significantly larger in CA-PVAT and Ao-PVAT than in SV-PVAT and ITA-PVAT. and there was no significant difference in the macrophage infiltration area between SV-PVAT and ITA-PVAT (Figure 3B). The inflammatory area of the M1 phenotype of macrophage infiltration quantified by the anti-CD11c antibody was significantly larger in CA-PVAT and Ao-PVAT than in SV-PVAT, and the area in CA-PVAT was larger than the areas in ITA-PVAT and Ao-PVAT (Figure 3C). The area of the M2 phenotype of macrophage infiltration quantified by the anti-CD206 antibody was significantly larger in CA-PVAT and Ao-PVAT than in SV-PVAT and ITA-PVAT (Figure 3D). The ratio of CD11c-positive/CD206positive macrophages was significantly higher in CA-PVAT than in the other PVATs of vessels (Figure 3E).

## **Metaflammation in Fat Pads**

Gene expression levels of CD68, a marker of macrophages, CD11c, a marker of M1 macrophages, CD163 and CD206, M2 macrophage markers, and interleukin-10, an anti-inflammatory cytokine as an M2 macrophage-related marker, in CA-PVAT and those in Ao-PVAT were significantly higher than those in SV-PVAT (Figure 4A). Gene expression levels of CD68 and CD11c in CA-PVAT were significantly higher than those in ITA-PVAT (Figure 4A). Gene expression levels of CD163, CD206, and interleukin-10 in ITA-PVAT were significantly higher than those in SV-PVAT (Figure 4A).

The gene expression level of adiponectin, an adipokine with antiatherogenic and anti-inflammatory features, in SV-PVAT was significantly higher than that in the other PVATs of vessels (Figure 4B). Gene expression levels of MCP-1 (monocyte chemoattractant protein-1), a chemokine, and inflammatory cytokines, including interleukin-1 $\beta$  and interleukin-6, were significantly higher in CA-PVAT and Ao-PVAT than in SV-PVAT and ITA-PVAT (Figure 4B). The gene expression level of tumor necrosis factor- $\alpha$ , an inflammatory cytokine, in SV-PVAT was significantly lower than that in the other PVATs of vessels (Figure 4B).

Gene expression levels of inflammasome-related molecules, including nucleotide-binding domain, leucinerich-containing family, pyrin domain-containing-3 and ASC (apoptosis-associated speck-like protein containing caspase recruitment domain), and endoplasmic reticulum stress-related molecules, including GRP78 (glucose-regulated protein 78) and CHOP (C/EBP homologous protein), were significantly higher in CA-PVAT and Ao-PVAT than in SV-PVAT (Figure 4C). Gene expression levels of nucleotide-binding domain, leucine-richcontaining family, pyrin domain-containing-3, GRP78, and CHOP in CA-PVAT were significantly higher than those in ITA-PVAT (Figure 4C).

## Adipocyte Developmental and Pattern-Forming Genes in Fat Pads

To further address phenotypic differences between adipose tissues in the 4 perivascular parts, we examined gene expression levels of adipocyte developmental and pattern-forming genes, including engrailed homeobox 1, empty spiracles homeobox 2, and homeobox A5, as in previous studies.<sup>15,28-30</sup> The gene expression level of engrailed homeobox 1 was significantly higher in SV-PVAT than in the other PVATs of vessels (Figure 5A). The expression level of empty spiracles homeobox 2 was significantly higher in SV-PVAT than in the other PVATs, and the empty spiracles homeobox 2 level in CA-PVAT was significantly higher than the levels in ITA-PVAT and Ao-PVAT (Figure 5B). The gene expression level of homeobox A5 was significantly higher in ITA-PVAT than in the other PVATs, and the homeobox A5 level in SV-PVAT was significantly higher than the levels in CA-PVAT and Ao-PVAT (Figure 5C).

## DISCUSSION

To the best of our knowledge, the present study is the first study to demonstrate a differential phenotype of metaflammation in fat pads surrounding the SV in





**A**, Representative hematoxylin-eosin (H-E) staining of perivascular adipose tissue surrounding the saphenous vein (SV-PVAT), that surrounding the internal thoracic artery (ITA-PVAT), that surrounding the coronary artery (CA-PVAT), and that surrounding the aorta (Ao-PVAT). Bar=100  $\mu$ m. **B**, Comparison of adipocyte sizes in SV-PVAT, ITA-PVAT, CA-PVAT, and Ao-PVAT of patients (n=48). Results are shown as means±SD. \**P*<0.05 vs SV-PVAT.

direct comparison to fat pads surrounding the coronary artery, ITA, and aorta. Interestingly, SV-PVAT was shown to have lower levels of metaflammation and consecutive adipose tissue remodeling (namely, augmented immune cell infiltration, induction of inflammatory cytokines, activation of inflammasomes and endoplasmic reticulum stress, and increased fibrosis) than those in CA-PVAT and Ao-PVAT. The





**A**, Representative Masson trichrome (MT) staining of perivascular adipose tissue surrounding the saphenous vein (SV-PVAT), that surrounding the internal thoracic artery (ITA-PVAT), that surrounding the coronary artery (CA-PVAT), and that surrounding the aorta (Ao-PVAT). Bar=200 µm. **B**, Fibrosis areas in SV-PVAT, ITA-PVAT, CA-PVAT, and Ao-PVAT of patients (n=28). Results are shown as means±SD. **C**, Gene expression levels of fibrosis-related molecules, including transforming growth factor- $\beta$  (TGF- $\beta$ ), tissue inhibitor of metalloproteinase 1 (TIMP1), platelet-derived growth factor subunit B (PDGFB), and macrophage-inducible C-type lectin (MINCLE), in SV-PVAT, ITA-PVAT, CA-PVAT, and Ao-PVAT. Results are shown as relative expression of each target gene in SV-PVAT of each patient (n=48) and as means±SEM for quantitative real-time polymerase chain reaction data. AU indicates arbitrary unit. \**P*<0.05 vs SV-PVAT, †*P*<0.05 vs CA-PVAT.



#### Figure 3. Macrophage infiltration in fat pads.

**A**, Representative immunohistological staining, including cluster of differentiation (CD) 68 (a marker of macrophages), CD11c (a marker of M1 macrophages), and CD206 (a marker of M2 macrophages), of perivascular adipose tissue surrounding the saphenous vein (SV-PVAT), that surrounding the internal thoracic artery (ITA-PVAT), that surrounding the coronary artery (CA-PVAT), and that surrounding the aorta (Ao-PVAT). Bar=100  $\mu$ m. **B** through **D**, Positive areas of immunohistological staining of CD68, CD11c, and CD206 in SV-PVAT, ITA-PVAT, CA-PVAT, and Ao-PVAT of patients (n=28). **E**, Ratios of the macrophage infiltration areas of CD11c/CD206 in SV-PVAT, ITA-PVAT, CA-PVAT, and Ao-PVAT. Results are shown as means $\pm$ SD. \**P*<0.05 vs SV-PVAT, †*P*<0.05 vs ITA-PVAT, †*P*<0.05 vs CA-PVAT.



#### Figure 4. Metaflammation in fat pads.

**A**, Gene expression levels of cluster of differentiation (CD) 68, a marker of macrophages, CD11c, a marker of M1 macrophages, and M2-polarized state-related molecules, including CD163, CD206, and interleukin-10 (IL-10), in perivascular adipose tissue surrounding the saphenous vein (SV-PVAT), that surrounding the internal thoracic artery (ITA-PVAT), that surrounding the coronary artery (CA-PVAT), and that surrounding the aorta (Ao-PVAT). **B**, Gene expression levels of adiponectin and inflammatory molecules, including MCP-1 (monocyte chemoattractant protein-1), interleukin-1 $\beta$  (IL-1 $\beta$ ), interleukin-6 (IL-6), and tumor necrosis factor- $\alpha$  (TNF- $\alpha$ ), in SV-PVAT, ITA-PVAT, CA-PVAT, and Ao-PVAT. **C**, Gene expression levels of inflammasome-related molecules, including nucleotide-binding domain, leucine-rich-containing family, pyrin domain-containing-3 (NLRP3) and ASC (apoptosis-associated speck-like protein containing caspase recruitment domain), and endoplasmic reticulum stress-related molecules, including GRP78 (glucose-regulated protein 78) and CHOP (C/EBP homologous protein), in SV-PVAT, ITA-PVAT, CA-PVAT, and Ao-PVAT. Results are shown as relative expression of each target gene in SV-PVAT of each patient (n=48) and as means±SEM for quantitative real-time polymerase chain reaction data. AU indicates arbitrary unit. \**P*<0.05 vs SV-PVAT, †*P*<0.05 vs ITA-PVAT.

level of metaflammation was higher in CA-PVAT than in Ao-PVAT, as indicated by a significantly larger extent of CD11c-positive M1 macrophage infiltration, and higher ratio of CD11c-positive M1/CD206-positive M2 macrophages. The phenotype

of SV-PVAT was closer to that of ITA-PVAT than the other PVATs. We previously reported that ITA-PVAT had less metaflammation, which may be related to the decreased atherosclerosis in the ITA used as the gold standard graft material in CABG.<sup>15</sup>





**A** through **C**, Gene expression levels of adipocyte developmental and pattern-forming factors, including engrailed homeobox 1 (EN1) (**A**), empty spiracles homeobox 2 (EMX2) (**B**), and homeobox A5 (HOXA5) (**C**), in perivascular adipose tissue surrounding the saphenous vein (SV-PVAT), that surrounding the internal thoracic artery (ITA-PVAT), that surrounding the coronary artery (CA-PVAT), and that surrounding the aorta (Ao-PVAT). Results are shown as relative expression of each target gene in SV-PVAT of each patient (n=48) and as means±SEM for quantitative real-time polymerase chain reaction data. AU indicates arbitrary unit. \**P*<0.05 vs SV-PVAT,  $^{+}P$ <0.05 vs ITA-PVAT.

The conventional SV harvesting removes all the vasa vasorum, resulting in ischemia of the wall in the SV.<sup>13</sup> Physical damage of the endothelial layer also occurs during the preparation of a graft by highpressure dilation of the vein, and degenerative and atherosclerotic processes can be promoted when a venous graft is implanted in the arterial circulation.<sup>13</sup> In contrast, in the "no-touch" technique, the SV is harvested together with its surrounding tissues.<sup>16</sup> The use of this novel technique has been reported to result in better short- and long-term graft patency rates than those achieved by using the conventional technique of SV harvesting, and the rates were comparable to those for ITA grafts.<sup>17-19</sup> Several mechanisms underlying the clinical benefit (or advantage) of no-touch vein grafts have been proposed. In the no-touch vein pedicle, the perivascular tissue is preserved and acts as a natural, external, and biological stent that consists of numerous collagen fibers, leading to prevention of the deleterious effects of aortic pressure to the vein wall and reduction of early thrombotic occlusion, neointimal and medial thickening, and kinking of the vein graft.<sup>17–19</sup> In addition, preservation of the vasa vasorum allows retrograde blood flow from the graft lumen to perfuse through the vein wall, leading to a decrease in transmural ischemic damage.<sup>31</sup> Furthermore, endothelial NO synthase is preserved in all 3 layers of the vein wall after the notouch technique,<sup>31,32</sup> whereas stripping and distention used during the conventional technique lead to a reduction of endothelial NO synthase.<sup>32,33</sup> It has also been proposed that SV-PVAT can be a source of several vasoactive factors.<sup>23,24</sup> In the present study, we demonstrated that SV-PVAT had an antiatherosclerotic phenotype, such as less metaflammation and proinflammatory cytokines. In addition, we found that such a phenotype of SV-PVAT is similar to that of ITA-PVAT. Taken together, the findings of the present study suggest that distinct phenotype of SV-PVAT is linked to a high rate of long-term patency of grafting in CABG using the no-touch technique of SV harvesting.

Anatomically separated regional fat pads have distinct gene expression patterns and functional characteristics.<sup>28-30</sup> It has been reported that in vitro differentiation of preadipocytes isolated from regional adipose depots produces adipocytes that phenotypically resemble in vivo counterparts of adipocvtes.<sup>28–30,34</sup> Phenotypic differences in adipocytes can stem from developmental divergence of precursor cells in distinct adipose tissue. In the present study, the expression patterns of adipocyte developmental and pattern-forming genes,<sup>15,28-30</sup> including engrailed homeobox 1, empty spiracles homeobox 2, and homeobox A5, were totally different among SV-PVAT, ITA-PVAT, CA-PVAT, and Ao-PVAT. These findings are consistent with the notion that adipocytes residing in these depots are derived from distinct precursor cells, which may underlie the phenotypic differences, although immune cells in adipose tissue may have some acquired effects.

The physiological development of PVAT has been reported to be reduced and impaired with increased inflammation in experimental models of atherosclerosis and endovascular injury.<sup>35–37</sup> Conversely, removal of normally functioning PVAT or implantation of PVAT with a proinflammatory phenotype promoted atherosclerosis, suggesting that the lack of normal PVAT is sufficient to drive increased atherosclerosis.<sup>38</sup> Autopsy analyses demonstrated that there was more macrophage infiltration in epicardial fat in vessels with large necrotic core plaques than in vessels without lipid core plaques.<sup>39,40</sup> In the present study, patients without coronary artery disease were not included as a control group. Therefore, it is difficult to determine whether the inflammatory response in PVAT is the cause or the result of atherosclerosis. Mechanistic studies are needed to clarify what underlies the relationship between the development of atherosclerosis and differential phenotypes of PVAT.

This study has several limitations. First, the number of patients was small, and there would have been selection bias of patients. However, the clinical profile of the patients enrolled in this study was not deviated from the average profiles in patients registered in Japan Cardiovascular Surgery Database. Second, because the recruited patients were only Japanese people, it is unclear whether the present findings can be generalized to other ethnicities. Third, because this study was a cross-sectional study using patients with coronary artery disease, the study does not prove a causal relationship between inflammatory changes in adipose tissue and the development and progression of atherosclerosis in the vein grafts. Interventional studies of CABG using the conventional technique as a control and the no-touch technique of SV harvesting are necessary for investigating the relationship between inflammatory changes in adipose tissue and graft patency in the future. Fourth, the recruited patients had several diseases, including diabetes mellites, hypertension, and dyslipidemia, and had been treated with several drugs, including angiotensin-converting enzyme inhibitors, angiotensin receptor blockers, and statins. Pretreatment with several drugs may have affected the extent of metaflammation and fibrosis in adipose tissue. Fifth, most of the recruited patients in the present study were adequately treated and were not obese, and risk factors in patients had been controlled well at the time of the operation of CABG. Such conditions may have affected the extent of inflammation and remodeling in adipose tissue. Finally, the gene expression of molecules was mainly investigated because of the small amount of fat pads in the present study, but the protein expression of molecules needs to be investigated in the future.

In conclusion, SV-PVAT has a distinct histological and gene expression phenotype, resulting from inherent differences in adipocytes, and is protected from metaflammation and consecutive adipose tissue remodeling in patients who have undergone elective CABG surgery. There are similarities in the gene expression phenotypes of SV-PVAT and PVAT surrounding the ITA, an atherosclerosis-resistant vessel. The phenotype of SV-PVAT might be linked to the high rate of long-term patency of grafting in CABG using the no-touch technique of SV harvesting.

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#### Disclosures

None.

#### **Supplementary Material**

Table S1 Figures S1-S2

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# **SUPPLEMENTAL MATERIAL**

Table S1. Prim	ers for humar	n genes in	quantitative	real-time	PCR.
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Genes	Accession #	Forward primer		Reverse primer		Amplicon size (bp)
18s	M10098	5'-GTAACCCGTTGAACCCCATT	-3'	5'-CCATCCAATCGGTAGTAGCG	-3'	151
Adiponectin	NM_004797	5'-GTCATGACCAGGAAACCAC	-3'	5'-TTCACCGATGTCTCCCTTAGG	-3'	203
Asc	NM_145182	5'-TGGATGCTCTGTACGGGAAG	-3'	5'-CCAGGCTGGTGTGAAACTGAA	-3'	110
Cd11c	NM_000887	5'-CTGCAAGGGTTTACATACACGG	-3'	5'-GAATTTTGGCGGCATCCCTAC	-3'	97
Cd68	NM_001040059	5'-CTTCTCTCATTCCCCTATGGACA	-3'	5'-GAAGGACACATTGTACTCCACC	-3'	105
Cd163	NM_004244	5'-GCGGGAGAGTGGAAGTGAAAG	-3'	5'-GTTACAAATCACAGAGACCGCT	-3'	89
Cd206	NM_002438	5'-GGGTTGCTATCACTCTCTATGC	-3'	5'-TTTCTTGTCTGTTGCCGTAGTT	-3'	126
Chop	NM_004083	5'-GGAGAACCAGGAAACGGAAAC	-3'	5'-TCTCCTTCATGCGCTGCTTT	-3'	69
Emx2	NM_004098	5'-CGGCACTCAGCTACGCTAAC	-3'	5'-CAAGTCCGGGTTGGAGTAGAC	-3'	107
En1	NM_001426	5'-GAGCGCAGGGCACCAAATA	-3'	5'-CGAGTCAGTTTTGACCACGG	-3'	92
Grp78	NM_005347	5'-CATCACGCCGTCCTATGTCG	-3'	5'-CGTCAAAGACCGTGTTCTCG	-3'	104
Hoxa5	NM_019102	5'-AACTCATTTTGCGGTCGCTAT	-3'	5'-TCCCTGAATTGCTCGCTCAC	-3'	89
ll1b	NM_000576	5'-CACGATGCACCTGTACGATCA	-3'	5'-GTTGCTCCATATCCTGTCCCT	-3'	121
116	NM_000600	5'-AAATTCGGTACATCCTCGACGG	-3'	5'-GGAAGGTTCAGGTTGTTTTCTGC	-3'	112
1110	NM_000572	5'-GACTTTAAGGGTTACCTGGGTTG	-3'	5'-TCACATGCGCCTTGATGTCTG	-3'	112
Мср1	NM_002982	5'-CAGCCAGATGCAATCAATGCC	-3'	5'-TGGAATCCTGAACCCACTTCT	-3'	190
Mincle	NM_014358	5'-CTGAAACACAATGCACAGAGAGA	-3'	5'-AAAGATGCGAAATGTCACAACAC	-3'	128
NIrp3	NM_001127462	5'-GATCTTCGCTGCGATCAACAG	-3'	5'-CGTGCATTATCTGAACCCCAC	-3'	81
Pdgfb	NM_033016.2	5'-CTCGATCCGCTCCTTTGATGA	-3'	5'-CGTTGGTGCGGTCTATGAG	-3'	239
Tgfb	NM_000660	5'-CAAGCAGAGTACACACAGCAT	-3'	5'-TGCTCCACTTTTAACTTGAGCC	-3'	126
Timp1	NM_003254	5'-CTTCTGCAATTCCGACCTCGT	-3'	5'-ACGCTGGTATAAGGTGGTCTG	-3'	79
Tnfa	NM_000594	5'-GAGGCCAAGCCCTGGTATG	-3'	5'-CGGGCCGATTGATCTCAGC	-3'	91



#### Figure S1. Control experiments of immunohistological staining.

**A-C.** Representative immunohistological staining using mouse anti-CD68 (A), rabbit anti-CD11c (B) and mouse anti-CD206 (C) primary antibodies in perivascular adipose tissue surrounding the aorta (Ao-PVAT). **D-F.** Representative immunohistological staining in the absence of a primary antibody (D) and using control primary antibodies of goat anti-mouse immunoglobulins (Ig) (E) and goat anti-rabbit Ig (F) in Ao-PVAT. Scale bar: 100 μm.



### Figure S2. Images of PCR amplicons in 2% agarose gels.

The specificity of the PCR reaction of genes was confirmed by one distinct band, which matched with the predicted size of the PCR amplicon.