

Fig. 1. Representative AAC on chest radiograph and inverse association of AAC grade with vascular endothelial function. (A) Representative AAC on postero-anterior chest radiograph and CT is shown. AAC extent was divided into four grades according to the categorization. (B) Endothelium-dependent vasodilation by reactive hyperemia (flow-mediated dilation, FMD) and endothelium-independent vasodilation by nitroglycerin (nitroglycerine-mediated dilation, NMD) in the brachial artery were measured. AAC grade was also negatively correlated with not only FMD but also NMD. The ratio of FMD to NTG was calculated, and compared with AAC grade. n.s.: not significant.

computer tomography (EB-CT) and multi-detector row CT (MD-CT), have been shown to be the gold standard for evaluating its calcification, with power to quantify its severity and progression [12]. Some reports demonstrated a positive correlation between coronary artery calcification and CV events [13]. A meta-analysis also showed that coronary artery calcification was associated with a more than 8-fold higher risk of CV events [14]. However, these high technical examinations are expensive and are not suitable for repeatable assessment in clinical practice.

Aortic calcification contributes to deterioration of Windkessel function and auto-regulation of hemodynamics. Recently, radiographically visible calcification has been suggested to be an independent risk factor for CV morbidity in large epidemiological studies [15,16]. However, most studies only evaluated whether aortic calcification is present or absent without considering its severity. In addition, although many reports show a correlation between abdominal aortic calcification and CV events, there are relatively few reports regarding aortic arch calcification (AAC).

We have recently reported the validity and usefulness of assessment of AAC grade, as determined by a simple chest X-ray [17]. AAC grade was significantly associated with clustering of traditional risk factors. However, it is still unknown whether AAC grade is a sensitive predictor of CV events beyond traditional risk factors. The purpose of the present study was to determine the validity of AAC grade as an independent predictor of CV events.

2. Methods

2.1. Study population

This study retrospectively evaluated the correlates and consequence of AAC detected by simple chest X-ray in 239 consecutive asymptomatic out-patients (male/female = 115/124, mean age 61.9 ± 10.8 years) who underwent measurement of endothelial function at the 1994–2000, as our previous report [17]. They comprised a non-selected sample of patients from the Department of Geriatric Medicine in University of Tokyo without a selection bias. All patients completed a detailed health history questionnaire just prior to follow-up of CV events, and this study was restricted to asymptomatic subjects with no history of CV events due to atherosclerotic diseases. Among 239 patients, the follow-up of CV events in 209 patients (male/female = 100/109, mean age 62.0 ± 11.0 years) was completed. Therefore, to evaluate the predictive value of each factor against CV events, we analyzed all data of these 209 patients in this study. Regarding ethical considerations, The Medical Ethics Committee of the University of Tokyo approved this study, and informed consent was obtained from all patients before the study.

2.2. Assessment of aortic arch calcification

We retrospectively reviewed postero-anterior chest X-rays, which were carried out simultaneously with measurement of endothelial function. The extent of AAC in each chest X-ray was assessed as shown in Fig. 1A. AAC extent was divided into four grades according to the following categorization, as our previous report [17]. Briefly, we scored the extent of calcification in the aortic arch as follows: grade 0, no visible calcification; grade 1, small spots or a single thin area of calcification; grade 2, one or more areas of thick calcification; grade 3, circumferential calcification. As our previous report, the accuracy and reproducibility of this technique for grading were employed 82%, 79%, 75% and 88% in grades 0, 1, 2 and 3, respectively [17].

2.3. Physical and laboratory measurements

BP was measured in the brachial artery with a random-zero sphygmomanometer. Traditional atherosclerotic risk factors, such as hypertension, dyslipidemia and diabetes mellitus, were defined as our previous report [17]. To evaluate renal function, we calculated the estimated glomerular filtration rate (eGFR). Excretion of urinary protein determined by qualitative analysis was simply scored from 0 to 4 (–: 0, +/-: 1, +: 2, ++: 3, +++: 4). Chronic kidney disease (CKD) was defined as eGFR of less than 60 ml/min/1.73 m² and/or marked proteinuria (+: 2 to +++: 4). Information on smoking obtained by interview was categorized as presence or absence of smoking.

2.4. Endothelial measurements

Endothelium-dependent vasodilation (presented as flow-mediated dilation: FMD) of brachial artery induced by reactive hyperemia was measured, according to the method described previously [18]. On the other hand, endothelium-independent

vasodilation by nitroglycerine (NTG) (presented as nitroglycerine-mediated dilation: NMD) was simultaneously measured. Briefly, the diameter of the brachial artery was measured in 2-dimensional ultrasound images, scanned over a longitudinal section 3–5 cm above the right elbow. The changes in diameter caused by FMD or NMD were expressed as the percent change relative to the diameter in the initial resting scan.

2.5. Clinical outcomes

Subjects in this study were continuously monitored for the occurrence of CV events. In the present study, the CV events according to our definition were specified clearly as coronary heart disease (CHD; angina pectoris (AP; criteria of AP were defined by presence of chest symptom and/or typical ST-T change in ECG), myocardial infarction (MI)), cerebrovascular disease (CVD; transient ischemic attack (TIA), stroke, cerebral hemorrhage), peripheral artery disease (PAD), heart failure (HF), and CV death. Individual diagnoses were classified according to the 9th International Classification of Disease (ICD-9) codes.

2.6. Statistical analysis

Differences between the groups were analyzed using ANOVA for continuous variables and χ^2 test for categorical variables. To determine the factors related to AAC grade, multivariate regression analysis was used. In the follow-up of CV events, event-free survival was analyzed using Kaplan–Meier curve and log-rank test. Hazard ratios (HRs) for CV events were analyzed using a multivariate Cox-proportional hazards regression. Data in the text, tables, and figures are expressed as mean \pm standard deviation (SD). A value of $p < 0.05$ was considered statistically significant.

3. Results

3.1. Patient characteristics

AAC grades of the subjects ($n = 209$) with completed follow-up of new CV events were distributed as follows: grade 0 (45%), grade 1 (22%), grade 2 (29%) and grade 3 (4%). Baseline characteristics of the total subjects and of the subjects with each grade are shown in Table 1. Because the number of subjects with grade 3 was small ($n = 8$), we combined grades 2 and 3 for the following analyses.

There were positive associations between AAC grade and the following variables: age, diastolic BP and pulse pressure, HbA1c, prevalence of diabetes and use of anti-diabetic agents (sulfonyl urea, alpha-glucosidase inhibitors and insulin). Regarding other traditional risk factors, no significant association of AAC grade with lipid parameters was found.

3.2. Association between AAC and renal dysfunction

The value of eGFR was declined according to increase in AAC grade (Table 1). Similarly, the score of urinary protein excretion was also positively associated with AAC grade. The level of serum phosphorus tended to increase according to AAC grade; however, the p -value for trend was not statistically significant (data not shown). Serum calcium showed no association with AAC grade.

3.3. Association between AAC and brachial artery measurements

The severity of AAC grade was significantly associated with a decline of FMD, suggesting the patients with high AAC grade had endothelial dysfunction (Fig. 1B). In addition, increasing AAC grade was also correlated with a decline of NMD. The ratio of FMD to NTG

Table 1
Baseline characteristics of patients overall and according to AAC grade.

	Total	Grade 0	Grade 1	Grades 2 + 3	<i>p</i> for trend
Number (male/female)	209 (100/109)	95 (47/48)	46 (22/24)	68 (31/37), G2(60), G3(8)	n.s.
Age (years)	62.0 ± 11.0	57.2 ± 9.5	64.0 ± 10.7	68.0 ± 9.5	<0.0001
Atherosclerotic risk factors (%)					
Hypertension	115 (55.0%)	46 (48%)	29 (63%)	40 (59%)	n.s.
Diabetes	61 (29.2%)	21 (22%)	11 (24%)	29 (43%)	<0.05
Dyslipidemia	141 (67.5%)	61 (64%)	32 (70%)	48 (71%)	n.s.
Smoking	74 (35.4%)	29 (31%)	18 (39%)	27 (39%)	n.s.
Anthropometric measurements					
Height (cm)	158.0 ± 9.3	159.4 ± 9.8	156.8 ± 9.1	156.3 ± 8.2	n.s.
Weight (kg)	60.4 ± 14.0	63.2 ± 16.4	59.4 ± 12.5	57.0 ± 10.0	n.s.
BMI (kg/m ²)	24.1 ± 4.3	24.7 ± 4.8	24.1 ± 4.1	23.3 ± 3.4	n.s.
Medical measurements					
sBP (mmHg)	133 ± 19	129 ± 18	138 ± 23	134 ± 17	n.s.
dBp (mmHg)	76 ± 12	79 ± 12	76 ± 13	73 ± 9	<0.05
mBP (mmHg)	95 ± 13	94 ± 13	98 ± 14	93 ± 11	n.s.
PP (mmHg)	56 ± 16	52 ± 12	58 ± 19	60 ± 14	<0.01
Laboratory measurements					
T-chol (mg/dl)	214 ± 42	223 ± 42	210 ± 42	205 ± 40	n.s.
LDL-C (mg/dl)	131 ± 43	135 ± 48	128 ± 44	127 ± 35	n.s.
HDL-C (mg/dl)	56.1 ± 16.2	59.3 ± 16.8	53.4 ± 16.2	53.4 ± 14.8	n.s.
TG (mg/dl)	135 ± 66	138 ± 96	140 ± 88	129 ± 94	n.s.
FPG (mg/dl)	107 ± 33	102 ± 23	106 ± 32	115 ± 42	n.s.
HbA1c (%)	6.1 ± 1.5	5.7 ± 1.1	6.0 ± 1.8	6.3 ± 1.6	<0.05
Cre (mg/dl)	0.8 ± 0.4	0.7 ± 0.2	0.9 ± 0.4	0.9 ± 0.6	n.s.
eGFR (ml/min/1.72 m ²)	67.7 ± 19.0	71.0 ± 17.0	64.5 ± 17.6	58.8 ± 21.9	<0.05
Proteinuria score (0–4)	0.6 ± 0.4	0.1 ± 0.1	0.3 ± 0.2	1.2 ± 0.6	<0.05
Anti-hypertensive drugs (%)					
ACE inhibitors/ARBs	23 (11.0%)	10 (11%)	6 (13%)	7 (10%)	n.s.
CCBs	79 (37.8%)	31 (33%)	18 (39%)	30 (44%)	n.s.
Alpha/beta-blockers	23 (11.0%)	7 (7%)	6 (13%)	10 (15%)	n.s.
Anti-diabetic drugs (%)					
Sulfonyl Urea	22 (10.5%)	5 (5%)	6 (13%)	11 (16%)	<0.05
Biguanides	1 (0.5%)	1 (1%)	0 (0%)	0 (0%)	n.s.
Alpha-glucosidase inhibitors	5 (2.4%)	0 (0%)	1 (2%)	4 (6%)	<0.05
Insulin	6 (2.9%)	0 (0%)	0 (0%)	6 (9%)	<0.05
Anti-hyperlipidemia drugs (%)					
HMG-CoA inhibitors	37 (17.7%)	15 (16%)	5 (11%)	17 (25%)	n.s.
Others	8 (3.8%)	2 (2%)	1 (2%)	5 (7%)	n.s.

G2, grade 2; G3, grade 3; BMI, body mass index; sBP, systolic blood pressure; dBp, diastolic blood pressure; mBP, mean blood pressure; PP, pulse pressure; T-chol, total cholesterol; LDL-C, low-density lipoprotein-cholesterol; HDL-C, high-density lipoprotein-cholesterol; TG, triglyceride; FPG, fasting plasma glucose; Cre, creatinine; eGFR, estimated glomerular filtration rate; ACE, angiotensin converting enzyme; ARBs, angiotensin II receptor blockers; CCBs, calcium channel blockers; n.s., not significant.

was calculated, and it was compared with AAC grade. No correlation between AAC grade and the ratio was found.

3.4. Correlates of AAC grade

As shown in Table 2, multiple regression analysis was performed to determine the factors related to AAC grade. Statistically significant correlates of AAC grade were age (coefficient $\beta=0.428$, $p<0.0001$), pulse pressure (coefficient $\beta=0.221$, $p=0.0004$), HbA1c (coefficient $\beta=0.197$, $p=0.048$), decreasing eGFR (coefficient $\beta=-0.148$, $p=0.033$) and proteinuria (coefficient $\beta=0.211$, $p=0.003$). In addition, both parameters for brachial artery vasodilation (FMD, coefficient $\beta=-0.222$, $p=0.001$; NMD; coefficient $\beta=-0.265$, $p<0.0001$) were also associated with AAC grade.

3.5. Association between AAC and CV events

New 57 CV events occurred during a mean follow-up period of 69 ± 45 months among 209 patients. These CV events included 18 stroke (13 cerebral infarction, 3 cerebral hemorrhage and 2 TIA), 31 CHD (23 AP and 8 MI including 1 CV death), 3 HF and 6 PAD. Fifty-eight percent of subjects who had CHD underwent coronary artery procedures (9 PCI and 9 CABG). We differentiated the CV out-

comes according to AAC grade. Kaplan–Meier analysis showed that the incidence of CV events was significantly higher in patients with higher AAC grade (grades 2 + 3) compared to those with AAC grade 0 or 1 (log-rank test; $p<0.01$) after adjustment for age, sex, and prevalence of diabetes, hypertension and dyslipidemia (Fig. 2A). Especially, patients with higher AAC grade had more cerebrovascular ischemic events. There was no sex-specific difference in each group. The HRs of each AAC grade were 1.69 (95%CI; 0.79–3.62, $p=0.176$) and 4.05 (95%CI; 2.16–7.58, $p<0.0001$) in AAC grade 1 and grades 2 + 3, respectively, compared to grade 0 as reference.

3.6. AAC is a predictor even in patients without CKD

AAC grade was significantly associated with renal dysfunction (Tables 1 and 2). Therefore, additional evaluation by Kaplan–Meier analysis was performed in each group with CKD or without CKD as based on the data of eGFR and proteinuria score (Fig. 2B). CKD group ($n=66$) was defined as an eGFR of less than 60 ml/min/1.73 m² and/or marked proteinuria (+: 2 to +++: 4). In addition, non-CKD group ($n=143$) was defined as an eGFR of over 60 ml/min/1.73 m² and no proteinuria (–: 0 to +/-: 1). In the CKD group, the patients with higher AAC grade (grades 2 + 3) had more CV events compared to grade 0 or 1; however, the statistical analysis (p -value for

Table 2
Multiple regression analysis of AAC grade and significantly associated variables.

	β	T-value	p-value
Age (years)	0.428	6.814	<0.0001
Sex (female)	-0.030	-0.426	0.671
Smoking	0.093	1.340	0.182
sBP (mmHg)	0.114	1.652	0.100
PP (mmHg)	0.221	3.258	0.001
LDL-C (mg/dl)	-0.063	-0.901	0.368
TG (mg/dl)	-0.037	-0.532	0.596
HbA1c (%)	0.197	2.788	0.048
eGFR (ml/min/1.72 m ²)	-0.148	-2.136	0.033
Proteinuria (grade)	0.211	2.967	0.003
FMD (%)	-0.222	-3.238	0.001

Abbreviations seen in Table 1.

trend) showed no significant (log-rank test; $p=0.06$). Of note, we found that AAC grade (grades 2 + 3) was a good predictor (log-rank test; $p<0.05$) even in the non-CKD group. When the analysis was stratified by age, we found similar results that higher AAC grades were also significantly associated with CV events in subjects aged 65 years and older (data not shown).

3.7. Multivariate Cox-proportional hazards analysis

As shown in Table 3, we evaluated the predictive value of each factor, such as age, atherosclerotic risk factors, endothelial dysfunction and AAC grade, against CV events using two kinds of analyses. In the first analysis, we divided the patients into two groups (less than 3.8% of mean value in FMD or over 3.8%) regarding endothelial function. The predictive value of endothelial dysfunction (HR = 1.29) was not statistically significant. Higher AAC grade (grades 2 + 3) was a significant predictor (HR; 2.49, 95%CI; 1.37–4.51, $p=0.01$) and the predictive power was superior to renal

dysfunction (low eGFR; HR = 1.97, proteinuria; HR = 2.25). Second, in multivariate analysis using the gross value of each laboratory parameter, statistical significance was found not only for increasing HbA1c but also for the presence of AAC grades 2 + 3 (HR; 2.56, 95%CI; 1.30–5.04, $p=0.007$). Unexpectedly, the predictive power of FMD was weaker than that of higher AAC grade in both analyses. In this connection, the increasing AAC grade (per 1-grade increase) had also statistical significance (HR; 1.6, $p=0.02$) using the same analysis (data not shown).

4. Discussion

Arterial calcification is a complication of advanced atherosclerosis. Especially, progressive arterial stiffness resulting from calcification makes the management of hemodynamics more difficult in the elderly. We have previously demonstrated the validity and usefulness of semi-quantitative estimation of AAC grade using chest X-ray examination, which is less expensive and is easy to

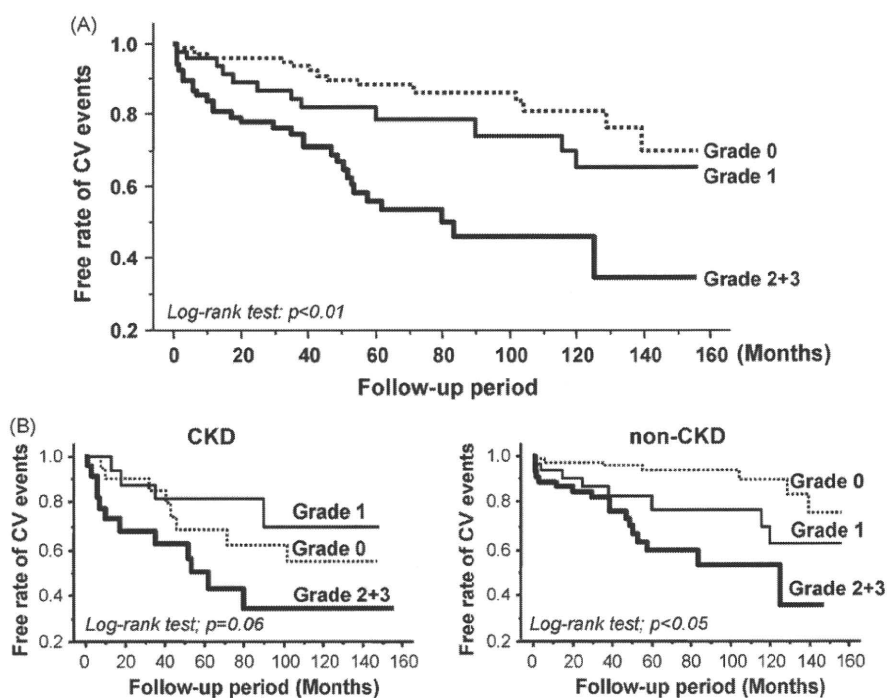


Fig. 2. Kaplan–Meier event-free survival analysis of all cardiovascular events according to AAC grade. (A) Kaplan–Meier analysis showed that the incidence of all new CV events was significantly higher in patients with higher AAC grade (grades 2 + 3) than in those with AAC grade 0 or 1 (p for trend; $p<0.01$) after adjustment for age, and prevalence of diabetes, hypertension and dyslipidemia. (B) The predictive value of AAC grade was also evaluated in each group with CKD or without CKD (non-CKD) using Kaplan–Meier analysis. Higher AAC grade (grades 2 + 3) was a good predictor in the non-CKD patients (p for trend; $p<0.01$). In the CKD patients, it seemed AAC grades 2 + 3 was strong compared to grade 0 or 1; however, the value was not statistically significant (p for trend; $p=0.06$).

Table 3

Multivariate analysis with Cox-proportional hazards models for cardiovascular events. Multivariate Cox-proportional hazards analysis showed each predictive value of several factors for CV events using the prevalence of several atherosclerotic risk factors (A) or using the gross value of each laboratory parameter (B).

Variables	Hazard ratio	95%CI	p-value
(A)			
Age (/year)	1.01	0.99–1.04	0.33
Gender (male)	1.24	0.61–2.55	0.55
Smoking	2.73	1.35–5.51	0.005
Dyslipidemia	0.89	0.51–1.58	0.69
Hypertension	1.26	0.69–2.31	0.44
Diabetes	1.28	0.67–2.32	0.44
Renal dysfunction (eGFR < 60)	1.97	1.11–3.50	0.02
Proteinuria (+, ++, +++)	2.25	1.05–4.76	0.03
Endothelial dysfunction (FMD < 3.8%)	1.29	0.72–2.31	0.39
AAC grade (grades 2 + 3)	2.49	1.37–4.51	0.01
(B)			
Age (/year)	1.01	0.98–1.04	0.62
Gender (male)	1.52	0.70–3.29	0.29
Smoking	2.01	0.92–4.41	0.08
LDL-C (/mg/dl)	1.00	0.99–1.01	0.82
HDL-C (/mg/dl)	0.99	0.97–1.01	0.31
Systolic BP (/mmHg)	1.00	0.99–1.02	0.75
HbA1c (%)	1.24	1.04–1.48	0.01
eGFR (/ml/min/1.73 m ²)	0.99	0.97–1.01	0.21
Proteinuria (per 1-score increase)	1.26	0.93–1.70	0.13
Endothelial function (%FMD)	1.05	0.94–1.19	0.39
AAC grade (grades 2 + 3)	2.56	1.30–5.04	0.007

follow-up routinely [17]. Among traditional risk factors, AAC grade was significantly correlated with the prevalence of diabetes and renal dysfunction (so-called CKD). In addition, AAC grade was associated with IMT and risk factor clustering. Therefore, we evaluated the effectiveness of AAC grade to predict new CV events using well-treated out-patients without any symptoms and past history. This study showed the usefulness of AAC grade as a strong independent predictor of new CV events. The result suggests an advantage of this simple non-invasive evaluation of AAC detectable on plain X-ray, as opposed to some recent high technical procedures like MD-CT.

If the level of calcification in the aorta predicts future CV events, it should be considered which area in the abdominal aorta or aortic arch is favorable and sensitive to evaluate the extent of calcification. First, there are much strong evidences that radiographic abdominal aortic calcification predicts CV events in huge populations during long-term follow-up. A report demonstrated that a significant correlation was found in 2467 Framingham Heart Study participants with plain abdominal X-ray during a period of 22 years [15,19]. Also regarding cerebrovascular disease, this correlation is supported by a very large prospective study of the Rotterdam Study cohort (HR of stroke; 1.89) [20]. These findings suggest valuable evaluation of abdominal aortic calcification using abdominal X-ray. On the other hand, as similar to the extent of calcification in the abdominal aorta, is the calcified level in the aortic arch also a good predictor? There are relatively few reports regarding the predictive value of AAC against CV events, although many previous studies have shown a positive association of CV events with abdominal aortic calcification. One possible explanation is that it is not easier to evaluate calcium deposition semi-quantitatively in the aortic arch as compared to that in the abdominal aorta. A report handling with population-based cohort has showed that the presence of AAC was independently associated with an increased risk of CHD (HR; 1.27 and 1.22 in male and female, respectively) [21], suggesting that our results are consistent with this evidence. However, that study evaluated whether AAC was present or absent using chest X-ray without considering the extent of calcification. There are several advantages and differences of our study as compared to the previous report. Our assessment by AAC grading certified a strong predictive value of higher AAC grade (grades 2 + 3) against inci-

dent CV events more precisely. However, in the analysis regarding HRs of each AAC grade, AAC grade 1 had no statistical significance of predictive value compared to grade 0. This result suggests that trivial calcium deposition in aortic arch only may not be a noticeable CV risk. In addition, we have confirmed that the AAC grade was significantly correlated with the calcified level in abdominal aorta using two independent non-invasive examinations [17]. In fact, huge differences in the hemodynamic state are found between patients with trivial AAC and those with severe AAC. Therefore, it is recommended at least to consider the AAC grade in routine clinical work.

Among our results obtained from this study, the predictive power of AAC grade was superior to that of FMD. In the early stage of atherosclerosis, endothelial dysfunction is found as an initial step, without organ damages, leading to a decline in release of nitric oxide (NO) and/or its bioavailability [22]. It is easier to decrease the value of FMD rather than that of NMD in patients with traditional risk factors; therefore, impaired brachial FMD has been shown to have a potent predictive value for subsequent clinical CV events [23]. Especially, a recent report also demonstrated the importance of prediction using FMD in a large population-based cohort [24]. However, there are few reports about the association between vascular function (FMD and NMD) and arterial calcification. Budoff et al. showed a fair correlation between decreasing brachial artery measurements and increasing calcification in the coronary artery, as determined by EB-CT [25]. However, the study evaluated vascular distensibility and resistance, but not FMD and NMD. In our study, patients with higher AAC grade (grades 2 + 3) showed a poor response not only in FMD but also in NMD. This result means that direct relaxation of smooth muscle cells themselves in the forearm by NTG was also impaired, suggesting that the artery may be stiffer. In other words, even if NO radical is smoothly released from endothelial cells, arterial dilation may be abrogated by the presence of microscopic calcification in the arterial media. However, microscopic calcification in the forearm is never assessed when vascular function is evaluated. Therefore, if we encounter patients with a decline in both endothelial-dependent and -independent vasodilation, we should assess whether they have a higher level of arterial calcification.

Patients with severe CKD, especially those on hemodialysis, show the strongest prognostic implication of arterial calcification [10]. In this study, AAC grade was significantly associated with a decline in eGFR and an increase in proteinuria. When we divided the patients into two groups, CKD or non-CKD, strong predictive power of higher AAC grade for CV events was found in patients without CKD. Many previous reports showed a good correlation between aortic calcification and CV events in hemodialysis patients; however, there are few reports regarding their correlation in the population with normal renal function. Our results suggest that prediction using AAC grade is valuable even in individuals without CKD.

It is still controversial how calcification in atheromatous plaque contributes to its stability. Although calcium deposition has been believed to play a protective role in plaque stability in the past [26], recent clinical reports using new techniques such as intravascular ultrasound have shown that it is a potent candidate for plaque rupture. One report showed that an important determinant is the anatomical distribution of calcification, but not its presence or absence, in the plaque [11]. In addition, calcification can be detected at two different anatomical sites in the arterial wall, Mönckeberg's medial and atherosclerotic intimal calcification. However, in the elderly and patients with advanced atherosclerosis, it is impossible to distinguish these two calcified lesions clearly using a plain radiographic approach only without a pathological approach, because the calcified lesions frequently present as a mixed pattern. In fact, when massive medial calcification appears as linear tram-tracks on radiographic examinations, it will probably not be possible to distinguish it from spotty intimal calcification. In the present study, we did not evaluate the extent of calcification in the coronary, carotid and cerebral arteries in this cohort; however, these may correlate with AAC grade. In addition, it has been generally shown that the important role of inflammation in arterial calcification in the patients with advanced atherosclerosis [27]. Unfortunately, in this study we do not have any data about detailed inflammatory cascades, such as high-sensitive CRP or any cytokines. Therefore, further investigation including the factors is needed.

Atherosclerosis and sclerosis have been shown to be epiphenomena associated with aging, diabetes and hypertension. In this study, AAC grade was strongly associated with diabetes and CKD. Among the medications used in the subjects, patients receiving anti-hyperglycemic drugs (insulin, sulfonyl urea and/or alpha-glucosidase inhibitors) had a higher AAC grade, although in cases of anti-hypertensive drugs or lipid-lowering drugs the significance was not found. Concretely, medications such as ACEI/ARB, CCBs and HMG-CoA inhibitors (statins) have recently been shown to improve arterial stiffness as a result of lowering BP and vasodilation. Unfortunately, we were not able to detect the precise change by drug intervention in this follow-up, because progressive change of calcium deposition in the aortic arch is very slow. However, if we could follow it up during more long-term, beneficial effects of intensive drug treatment might be found.

The principal limitation of this study is the semi-quantitative evaluation of AAC, because this method using four grades to evaluate AAC is relatively crude. Therefore, the true calcium deposition in aortic wall may be underestimated. Second, the largely retrospective design reduced the quality of the study, and the population size in this study was also small. Therefore, a largely prospective study using examinations for both AAC and abdominal aortic calcification is necessary.

In conclusion, the extent of aortic calcification and its significance are generally easily disregarded in routine clinical work. This study demonstrates that AAC is a strong independent predictor of increased risk of CV events. These findings support our hypothesis

that risk stratification by simple assessment of AAC using simple chest X-ray could play an important role in primary preventive management of atherosclerotic diseases. Beyond traditional risk factors, this assessment can provide additional beneficial information for the prediction of CV events.

Conflict of interest

We have no conflict of interest to disclose.

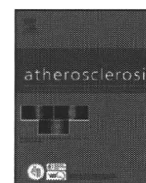
Acknowledgements

This study was supported by Grants-in-Aid for Scientific Research from the Ministry of Education, Science, Culture and Sports of Japan (Nos. 18590801, 19590854, 19590854, 20249041, and 21590947), Ono Medical Research Foundation, Kanzawa Medical Research Foundation, Novartis Foundation for Gerontological Research, Takeda Research Foundation, and Mitsui-Sumitomo Insurance Welfare Foundation.

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Low testosterone level as a predictor of cardiovascular events in Japanese men with coronary risk factors

Masahiro Akishita^{a,*}, Masayoshi Hashimoto^b, Yumiko Ohike^a, Sumito Ogawa^a, Katsuya Iijima^a, Masato Eto^a, Yasuyoshi Ouchi^a

^a Department of Geriatric Medicine, Graduate School of Medicine, University of Tokyo, 7-3-1 Hongo, Bunkyo-ku, Tokyo 113-8655, Japan

^b Department of General Internal Medicine, Kobe University School of Medicine, Kobe, Japan

ARTICLE INFO

Article history:

Received 25 August 2009
Received in revised form 5 October 2009
Accepted 22 October 2009
Available online 13 November 2009

Keywords:

Androgen
Sex hormone
Estrogen
Risk factor

ABSTRACT

Objective: Recent epidemiological studies have found that testosterone deficiency is associated with higher mortality largely due to cardiovascular (CV) disease in community-dwelling older men. We investigated whether a low plasma testosterone level could predict cardiovascular events in middle-aged Japanese men with coronary risk factors.

Methods: One hundred and seventy-one male outpatients (30–69 years old, mean \pm SD = 48 \pm 13 years) who had any coronary risk factor (hypertension, diabetes, dyslipidemia, smoking, and obesity) without a previous history of CV disease were followed up. At baseline, the subjects underwent examination of coronary risk factors, measurement of flow-mediated dilation (FMD) of the brachial artery as an indicator of vascular endothelial function and assays of plasma total testosterone, dehydroepiandrosterone-sulfate (DHEA-S), estradiol and cortisol.

Results: During the mean follow-up period of 77 months, a total of 20 CV events occurred. Kaplan–Meier survival analysis by tertile of plasma hormone levels revealed that the subjects with the lowest testosterone tertile were more likely to develop CV events than those with the highest tertile ($P < 0.01$ by log-rank test). Cox proportional hazards models showed that the subjects with the lowest tertile of plasma testosterone (< 14.2 nmol/L) had an approximately 4-fold higher CV event risk compared to those with the higher testosterone tertiles after adjustment for coronary risk factors including medication and FMD (unadjusted hazard ratio, 3.61; 95% CI, 1.47–8.86; multivariate-adjusted hazard ratio, 4.61; 95% CI, 1.02–21.04). Multivariate analysis did not show any significant association of DHEA-S, estradiol or cortisol with CV events.

Conclusions: A low plasma testosterone level is associated with CV events in middle-aged Japanese men, independent of coronary risk factors and endothelial function. This is the first report to show the relationship between endogenous testosterone and CV events in Asian population.

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1. Introduction

Plasma testosterone level declines with advancing age in men [1]. Testosterone deficiency is often associated with age-related diseases such as erectile dysfunction, osteoporosis, depressed mood, cognitive impairment and frailty [2,3]. Furthermore, a number of studies suggest that testosterone deficiency is related to cardiovascular (CV) disease and its risk factors in men. Inverse relations between testosterone level and coronary risk factors including obesity [4,5], hypertension [5,6], dyslipidemia [4,5], and diabetes [5,7] have been reported. In addition, we and others have

shown that a low testosterone level is associated with markers of atherosclerosis such as impaired endothelial vasomotor function [8], increased carotid intima-media thickness [9] and aortic calcification [4]. Although these data do not indicate a causal relationship between endogenous testosterone and CV disease, recent epidemiological studies have demonstrated that community-dwelling older men with a low testosterone level are more likely to die [10–12], largely due to CV disease [11,12]. However, this issue remains unknown in Asian population.

Based on these backgrounds, we tested the hypothesis that a low testosterone level is an independent risk factor for CV disease even in middle-aged Japanese men with coronary risk factors. For this purpose, we conducted a survey of 171 male patients by using baseline clinical information and by measuring sex hormone levels in stored plasma.

* Corresponding author. Tel.: +81 3 5800 8832; fax: +81 3 5800 8831.
E-mail address: akishita-tky@umin.ac.jp (M. Akishita).

2. Methods

2.1. Subjects

Male subjects aged 30–69 years at baseline, who were referred to our department to check for CV disease and undergo examination of vasomotor function of the brachial artery in 1996–2000, and had any of the classical coronary risk factors including hypertension, dyslipidemia, diabetes mellitus and current smoking, were eligible. Hypertension, dyslipidemia and diabetes mellitus were defined according to diagnostic criteria [13–15] or if the subject was taking any medication for these diseases. Subjects with a history of CV disease, including stroke, coronary heart disease, congestive heart failure and peripheral arterial disease, were excluded. Malignancy, overt endocrine disease and use of steroid hormones were also excluded, because these conditions may have a significant influence on both plasma sex hormones and clinical course.

Of the 188 eligible subjects whose plasma was stored, written informed consent was obtained from 171 subjects; 1 subject refused and 16 subjects were lost to follow-up. Then, plasma hormone levels were measured and follow-up data were obtained in 171 subjects. The study protocol was approved by the ethics committee of the Graduate School of Medicine, The University of Tokyo. Each subject or a family member, if the subject had died, gave written informed consent for enrollment in this study.

2.2. Clinical measurements

Clinical information was collected at baseline when each patient attended our department. Blood sampling and measurement of height, weight, blood pressure and vasomotor function were performed in the morning after a 14-h overnight fast. Blood pressure was measured at least twice using an automated, digital electrophygmomanometer (Omron Healthcare Co., Ltd., Kyoto, Japan) on the nondominant arm in a sitting position, and the average was used for analysis.

Serum total cholesterol and triglyceride concentrations were measured enzymatically, and serum high-density lipoprotein (HDL) cholesterol concentration was measured by the heparin- $\text{Ca}^{2+}\text{Ni}^{2+}$ precipitation method. Plasma glucose concentration was assayed by the glucose oxidase method, and hemoglobin A1c level was measured by high-performance liquid chromatography.

Plasma concentrations of total testosterone, dehydroepiandrosterone-sulfate (DHEA-S), estradiol and cortisol were determined using sensitive radioimmunoassays by a commercial laboratory (SRL, Inc., Tokyo, Japan). Because the plasma used for hormone assays was deep-frozen (-80°C) for up to 7 years, we checked the change in titers using the stored samples, which had been measured at sampling 5–7 years before. Pearson's correlation coefficient between the two measurements was 0.965 for estradiol ($n=34$), 0.976 for testosterone ($n=20$), 0.991 for DHEA-S ($n=15$) and 0.937 for cortisol ($n=16$), indicating that there was no significant change in plasma titers in our frozen samples. The intra-assay coefficients of variation for the measurements were less than 5%.

Vasomotor function of the brachial artery was evaluated using an ultrasound machine according to the method described previously [16]. Briefly, endothelium-dependent flow-mediated vasodilation (%FMD) was measured as the maximal percent change in the vessel diameter after reactive hyperemia. Subsequently, endothelium-independent nitroglycerin-induced vasodilation was measured as the maximal percent change in the vessel diameter after sublingual administration of nitroglycerin spray (0.3 mg; Toa Eiyō Co., Tokyo). The same examiner (M.H.) performed the measurements of FMD throughout this study.

2.3. Follow-up

The subjects were followed in 2006–2007 by mail and/or visits to our clinic. Each subject or a family member completed the questionnaire on CV disease and health status. CV events analyzed as the endpoints of this study included stroke, coronary artery disease, sudden cardiac death, and peripheral arterial disease. If CV events were reported on the questionnaire, we attempted to confirm the diagnosis of each event by medical records and/or interview by research doctors who were unaware of the patient's plasma hormone levels. Finally, after thorough examination, 20 cases were determined as CV events. Eighteen cases were ascertained by medical records which included clinical course, physical examination, laboratory tests and imagings. Because medical records were not available on other two cases of self-reported ischemic stroke, they were diagnosed according to the phone interview to each patient.

2.4. Data analysis

Values are expressed as mean \pm SD in the text unless otherwise stated. Differences between the groups were analyzed using ANOVA for continuous variables and Chi-squared test for categorical variables. Survival was analyzed using Kaplan–Meier plots and log-rank tests. Hazard ratios (HRs) for CV events were analyzed using Cox proportional hazards regression. A value of $P < 0.05$ was considered statistically significant. Data were analyzed using SPSS (Ver. 17.0, SPSS Inc., Chicago, IL).

3. Results

3.1. Characteristics of subjects according to plasma testosterone level

Table 1 shows the baseline characteristics of the subjects by tertile of plasma testosterone. As reported previously [4–8], subjects with the lowest testosterone tertile tended to be obese, hypertensive, dyslipidemic, diabetic, and to have impaired endothelial vasomotor function compared to those with higher testosterone tertiles. Age and smoking status were not different between the groups.

3.2. CV events and hormones

During the mean follow-up period of 77 ± 46 months (median = 54 months), a total of 20 CV events occurred (Table 2). Eleven cases of coronary artery disease included three of myocardial infarction, three of medically treated angina pectoris, four of percutaneous coronary intervention, and one of coronary artery bypass grafting. All of the five cases of stroke were due to cerebral infarction.

As shown in Fig. 1, Kaplan–Meier survival analysis by tertile of plasma testosterone level revealed that low testosterone was associated with CV events. Cox proportional hazards models showed that the subjects with the lowest tertile of plasma testosterone, but not those with the middle tertile, had significantly increased risk for CV events compared to those with the highest tertile (Table 2). Adjustment for age and body mass index did not attenuate the effect.

Then, HRs for the lowest tertile of plasma testosterone vs. the higher (middle and highest) tertiles were analyzed. The subjects with the lowest tertile (<14.2 nmol/L) showed an unadjusted HR of 3.61 (95% CI, 1.47–8.86), and an adjusted HR of 4.24 (95% CI, 1.67–10.78) for age, body mass index, and current smoking. The HR was 4.61 (95% CI, 1.02–21.04) after adjustment for age, body mass index, current smoking, systolic blood pressure, HDL cholesterol, non-HDL cholesterol, hemoglobin A1c, %FMD,

Table 1
Baseline characteristics of subjects by tertile group of plasma testosterone.

	Tertile 1 <14.2 nmol/L (n = 57)	Tertile 2 14.2–19.4 nmol/L (n = 57)	Tertile 3 >19.4 nmol/L (n = 57)	p for trend
Testosterone (nmol/L)	11.0 ± 3.0	17.0 ± 1.6	24.0 ± 3.0	<0.001
(ng/dL)	(318 ± 86)	(490 ± 45)	(693 ± 86)	
DHEA-S (μmol/L)	4.94 ± 2.68	4.55 ± 2.25	4.83 ± 2.64	0.81
Estradiol (pmol/L)	115 ± 30	116 ± 31	133 ± 30	0.004
Cortisol (nmol/L)	386 ± 138	378 ± 142	361 ± 120	0.67
Age (years)	47 ± 13	45 ± 13	50 ± 14	0.24
Body mass index (kg/m ²)	27.6 ± 5.5	25.6 ± 4.3	24.1 ± 3.6	<0.001
Systolic blood pressure (mmHg)	131 ± 18	125 ± 16	123 ± 12	0.01
Diastolic blood pressure (mmHg)	79 ± 15	74 ± 11	74 ± 9	0.04
Non-HDL cholesterol (mmol/L)	4.19 ± 1.27	3.91 ± 1.06	3.74 ± 1.01	0.10
HDL cholesterol (mmol/L)	1.20 ± 0.36	1.23 ± 0.41	1.44 ± 0.48	0.005
Triglycerides (mmol/L)	2.04 ± 2.12	1.91 ± 1.85	1.46 ± 1.28	0.18
Fasting plasma glucose (mmol/L)	6.00 ± 1.18	5.73 ± 0.92	5.73 ± 1.28	0.34
Hemoglobin A1c (%)	5.9 ± 1.7	5.2 ± 0.8	5.5 ± 1.2	0.03
%FMD	4.2 ± 2.7	5.7 ± 4.2	6.1 ± 3.8	0.01
%NTG	12.8 ± 4.3	14.2 ± 5.4	13.2 ± 5.0	0.30
Hypertension, n (%)	30 (53)	20 (35)	20 (35)	0.09
Dyslipidemia, n (%)	33 (58)	35 (61)	24 (42)	0.09
Diabetes mellitus, n (%)	15 (26)	7 (12)	9 (16)	0.13
Current smoker, n (%)	28 (49)	25 (44)	29 (51)	0.74

DHEA-S, dehydroepiandrosterone-sulfate; HDL, high-density lipoprotein; %FMD, percent flow-mediated dilation of brachial artery; %NTG, percent nitroglycerine-induced dilation of brachial artery.

Values are expressed as mean ± SD. Continuous variables were compared by ANOVA and categorical variables by Chi-squared test.

Table 2
Cardiovascular events by tertile of plasma testosterone.

	Tertile 1 <14.2 nmol/L (n = 57)	Tertile 2 14.2–19.4 nmol/L (n = 57)	Tertile 3 >19.4 nmol/L (n = 57)	Total (n = 57)
Number of events				
Stroke	2	3	0	5
Coronary artery disease	7	2	2	11
Sudden cardiac death	2	0	0	2
Peripheral arterial disease	1	0	1	2
Total cardiovascular events	12	5	3	20
HRs (95% CI) for total cardiovascular events				
Unadjusted	4.82 (1.36, 17.12)	1.67 (0.40, 6.99)	1(Ref)	
Adjusted for age	6.36 (1.78, 22.80)	1.82 (0.43, 7.71)	1(Ref)	
Adjusted for age and BMI	7.01 (1.94, 25.34)	1.86 (0.44, 7.86)	1(Ref)	

BMI, body mass index. HRs (Hazard ratios) were analyzed using Cox proportional hazards regression.

medications (antihypertensives, statins, hypoglycemic agents and antiplatelet agents), estradiol and DHEA-S. In addition to testosterone, age (HR per year, 1.12; 95% CI, 1.05–1.20), %FMD (HR per 1% increase, 0.80; 95% CI, 0.64–0.99) and HDL cholesterol (HR per 1 mg/dL, 0.88; 95% CI, 0.81–0.95) were independently asso-

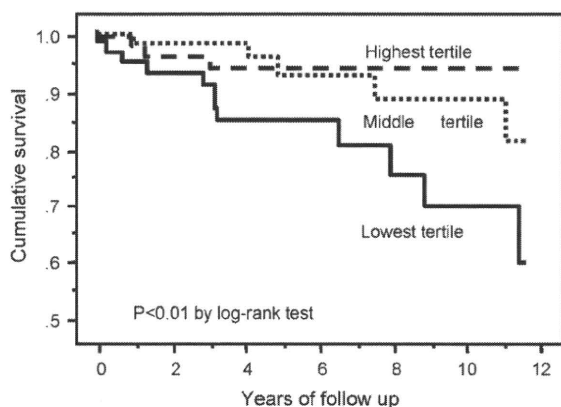


Fig. 1. Survival curves for cardiovascular events by tertile group of plasma concentration of testosterone. Cut-offs of the tertiles for testosterone were 14.2 and 19.4 nmol/L (410 and 560 ng/dL).

ciated with CV events, but other variables were not in this final model. Further inclusion of other hormones and nitroglycerin-induced endothelium-independent vasodilation into the model did not influence the statistical results (data not shown).

Two subjects with the lowest tertile of plasma testosterone suffered CV events within 6 months of follow-up; a case of sudden cardiac death and a case of coronary artery bypass grafting. Accordingly, similar statistical analyses were performed excluding these two cases. The results were essentially unchanged, although the HRs were slightly smaller (unadjusted HR, 3.06; 95% CI, 1.21–7.78; multivariate-adjusted HR, 3.80; 95% CI, 1.06–13.52).

Among other hormones examined, only DHEA-S was associated with increased risk for CV events, but was canceled by adjustment for age (data not shown). Further multivariate analysis did not show any significant association of DHEA-S, estradiol or cortisol with CV events.

4. Discussion

In this follow-up study of middle-aged Japanese men with coronary risk factors, a low plasma testosterone level was associated with CV events. Although the subjects with lower testosterone levels had worse profiles of coronary risk factors [4–7,11,12] and endothelial function [8] at baseline, as reported previously, adjustment for these confounders including age and cardiovascu-

lar medication indicated that low testosterone was an independent risk factor for CV events. In contrast, DHEA-S, estradiol and cortisol levels were not related to CV events.

A number of cross-sectional studies have shown an association between low testosterone level and CV disease [17,18], but have not provided evidence of a causal relationship between them. In recent years, longitudinal follow-up studies have demonstrated that community-dwelling older men (around 70 years on average) with lower testosterone levels are more likely to die from CV disease [11,12]. In contrast, a low testosterone level was not associated with CV deaths [19] or events [20] in community-dwelling middle-aged men (early 50s on average). These different findings might arise from the characteristics of the populations such as age and coronary risk factors, duration of follow-up and/or cut-off level of plasma testosterone at baseline. In any case, since all the above-mentioned studies were achieved in Caucasians, our study is the first to investigate the relationship between endogenous testosterone and CV events in Asians. Also, the present study showed a positive association between low testosterone level and CV events in middle-aged men with coronary risk factors, implying the clinical importance of measuring plasma testosterone in patients at risk, even if they are not old.

Unlike the previous reports showing an association of CV events with low levels of DHEA-S [21] and estradiol [22], and with a high cortisol:testosterone ratio [20], the present study did not show any significant association of CV events with estradiol, cortisol or cortisol:testosterone ratio (data not shown). The association between low DHEA-S and CV events was abolished by statistical adjustment for age, suggesting that the age-dependent decline of DHEA-S (Pearson's correlation coefficient between age and DHEA-S: -0.588 ; $P < 0.001$) might have eliminated the association with CV events if present. Taking together with the Cox regression model including all hormones, it is suggested that testosterone is the strongest among four steroid hormones that could be predictive of CV events in this population.

There could be several mechanisms by which endogenous testosterone protects men from CV disease. Consistent with the present study, observational studies [4–8,11,12] suggest that testosterone might prevent risk factors such as obesity, hypertension, dyslipidemia, diabetes and endothelial dysfunction. Supplementary studies support the beneficial effects of testosterone on adiposity [23] and endothelial vasomotor function [24]. Based on these findings, risk markers and endothelial vasomotor function were entered into the multivariate models. Although statistical adjustment may have been insufficient to exclude the interaction between testosterone and these risk factors, testosterone remained a significant predictor of CV events in the present study. Testosterone has been reported to inhibit vascular smooth muscle cell proliferation and neointima formation [25], suggesting the direct action of testosterone on the vasculature. Also, the effects of testosterone on inflammation, hemostasis and cardiac ischemia [26] might be involved in the final process leading to CV events. The precise mechanisms, including the role of the androgen receptor and aromatization to estrogen, should be addressed in the future.

The finding of this study should not be extended to men without coronary risk factors. Our preliminary data of 47 middle-aged men without coronary risk factors showed that no subject suffered CV events during the mean follow-up period of 102 months, although a quarter of them had plasma testosterone level below the cut-off of this study (<14.2 nmol/L). Thus, the relationship between plasma testosterone and CV outcomes might be totally different in middle-aged Japanese men without coronary risk factors.

This study has several limitations. First, the number of CV events was too small to reach a clear conclusion with strong statistical power, due primarily to the small sample size and secondarily to the low incidence of CV events (approximately 2%/year). Second,

the largely retrospective design (the protocol had been approved a few years before the final data collection) reduced the quality of the study and compelled us to lose many plasma samples and 16 subjects in the follow-up. Third, not all the CV events were confirmed by medical recordings. Two cases (a case in the lowest tertile and another in the middle tertile of plasma testosterone level) were determined according to the phone interview to each patient. Although the exclusion of these two cases did not significantly influence the statistical results (data not shown), self-reported outcomes limit the accuracy of this study. Fourth, the potential influence of medication on plasma testosterone level and on CV events cannot be excluded, although statistical adjustment for each class of drugs did not affect the results. For instance, beta-blockers have been reported to decrease plasma testosterone [27], but were taken by only nine subjects and were not related to testosterone level in our population (data not shown). Fifth, active forms of testosterone such as bioavailable and calculated free testosterone were not measured, because a direct assay of bioavailable testosterone or an assay of sex hormone binding globulin, which is necessary for free testosterone calculation, is not available in Japan. However, since previous longitudinal studies [11,12] have shown an association of total testosterone with CV mortality, the fundamental findings might not have differed if active forms of testosterone had been analyzed.

In summary, a low plasma testosterone level was associated with CV events in middle-aged Japanese men, independent of coronary risk factors and endothelial function. This study is the first to show the relationship between endogenous testosterone and CV events in Asian population, and provides evidence supporting the protective role of endogenous testosterone in the development of CV disease in men.

Acknowledgements

We thank Ms. Yuki Ito for her excellent technical assistance. This study was supported by a Health and Labor Sciences Research Grant (H17-Choju-046) from the Ministry of Health, Labour and Welfare of Japan, Grants-in-Aid for Scientific Research from the Ministry of Education, Science, Culture and Sports of Japan (21390220, 20249041) and grants from the NOVARTIS Foundation for Gerontological Research and the Yamaguchi Endocrine Research Association.

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Androgen Receptor-dependent Transactivation of Growth Arrest-specific Gene 6 Mediates Inhibitory Effects of Testosterone on Vascular Calcification^{*[S]}

Received for publication, August 17, 2009, and in revised form, December 16, 2009. Published, JBC Papers in Press, January 4, 2010, DOI 10.1074/jbc.M109.055087

Bo-Kyung Son[‡], Masahiro Akishita⁺¹, Katsuya Iijima[‡], Sumito Ogawa[‡], Koji Maemura[§], Jing Yu[¶], Kenichi Takeyama[¶], Shigeaki Kato[¶], Masato Eto[‡], and Yasuyoshi Ouchi[‡]

From the [‡]Department of Geriatric Medicine and the [¶]Department of Integrated Traditional Medicine, the Graduate School of Medicine and the [¶]Institute of Molecular and Cellular Biosciences, Graduate School of Agricultural and Life Sciences, University of Tokyo, Bunkyo-ku, Tokyo 113-8655 and the [§]Department of Cardiovascular Medicine Graduate School of Biomedical Sciences, Nagasaki University, Nagasaki 852-8501, Japan

Recent epidemiological studies have found that androgen deficiency is associated with a higher incidence of cardiovascular disease in men. However, little is known about the mechanism underlying the cardioprotective effects of androgens. Here we show the inhibitory effects of testosterone on vascular calcification and a critical role of androgen receptor (AR)-dependent transactivation of growth arrest-specific gene 6 (Gas6), a key regulator of inorganic phosphate (P_i)-induced calcification of vascular smooth muscle cells (VSMC). Testosterone and nonaromatizable androgen dihydrotestosterone inhibited P_i-induced calcification of human aortic VSMC in a concentration-dependent manner. Androgen inhibited P_i-induced VSMC apoptosis, an essential process for VSMC calcification. The effects on VSMC calcification were mediated by restoration of P_i-induced down-regulation of Gas6 expression and a subsequent reduction of Akt phosphorylation. These effects of androgen were blocked by an AR antagonist, flutamide, but not by an estrogen receptor antagonist, ICI 162,780. We then explored the mechanistic role of the AR in Gas6 expression and found an abundant expression of AR predominantly in the nucleus of VSMC and two consensus ARE sequences in the Gas6 promoter region. Dihydrotestosterone stimulated Gas6 promoter activity, and this effect was abrogated by flutamide and by AR siRNA. Site-specific mutation revealed that the proximal ARE was essential for androgen-dependent transactivation of Gas6. Furthermore, chromatin immunoprecipitation assays demonstrated ligand-dependent binding of the AR to the proximal ARE of Gas6. These results indicate that AR signaling directly regulates Gas6 transcription, which leads to inhibition of vascular calcification, and provides a mechanistic insight into the cardioprotective action of androgens.

Recent clinical studies have suggested that a low plasma testosterone level is associated with advanced atherosclerosis and is independently related to cardiovascular disease and death (1–5). Many but not all animal studies have also shown inhibitory effects of androgens on experimental atherosclerosis and vascular remodeling (6–8). Also, several clinical studies indicate that the testosterone level is inversely related to vascular calcification, a significant feature of vascular pathology (9). However, the mechanism underlying the vasoprotective effects of androgens is poorly understood.

Most of the actions of testosterone, particularly of nonaromatizable dihydrotestosterone (DHT),² are mediated by the androgen receptor (AR) (10, 11). In the nucleus the AR activates transcription by binding to androgen-response elements (AREs) in the promoter and enhancer regions of target genes (12). It further has been reported that AR is expressed in all layers of the arterial wall (13) and is involved in vascular disease (14, 15). However, the precise mechanism such as the signaling and molecular target of the AR has not been addressed.

We recently reported that growth arrest-specific gene 6 (Gas6) is a key molecule regulating calcification of vascular smooth muscle cells (VSMC) through the survival signal transduction mediated by phosphatidylinositol 3-OH kinase/Akt phosphorylation (16, 17). Gas6 is a member of the vitamin K-dependent protein family and is a secreted protein that harbors a γ -carboxylglutamic acid-rich domain and four epidermal growth factor-like repeats (18). In the present study we showed transcriptional activity of the AR in VSMC and an inhibitory effect of androgens on inorganic phosphate (P_i)-induced VSMC calcification. The inhibitory effect of androgens on VSMC calcification was attributable to restoration of the Gas6-mediated survival pathway. Furthermore, we found that the AR directly binds to the ARE in the Gas6 promoter region and transactivates the Gas6 gene.

* This work was supported by Health and Labor Sciences Research Grant H17-Choju-046 from the Ministry of Health, Labor, and Welfare of Japan and Grants-in-Aid for Scientific Research from the Ministry of Education, Science, Culture, and Sports of Japan 21390220 and 20249041.

[S] The on-line version of this article (available at <http://www.jbc.org>) contains supplemental Fig. 1.

¹ To whom correspondence should be addressed: Dept. of Geriatric Medicine, Graduate School of Medicine, The University of Tokyo, 7-3-1 Hongo, Bunkyo-ku, Tokyo 113-8655, Japan. Tel.: 81-3-5800-8832; Fax: 81-3-5800-8831; E-mail: akishita-tyk@umin.ac.jp.

² The abbreviations used are: DHT, dihydrotestosterone; AR, androgen receptor; ARE, androgen-response element; Gas6, growth arrest-specific gene 6; VSMC, vascular smooth muscle cells; HASMC, human aortic smooth muscle cells; DMEM, Dulbecco's modified Eagle's medium; siRNA, small interfering RNA; Act D, actinomycin D; ChIP, chromatin immunoprecipitation; luc, luciferase.

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EXPERIMENTAL PROCEDURES

Cell Culture—Human aortic smooth muscle cells (HASMC) derived from a 32-year-old man were purchased from Clonetics. HASMC were maintained in Dulbecco's modified Eagle's medium (DMEM) supplemented with 15% fetal bovine serum, 100 units/ml penicillin, and 100 mg/ml streptomycin at 37 °C in a humidified atmosphere with 5% CO₂. HASMC were used up to passage 8 for the experiments. In preliminary experiments HASMC were cultured in a calcifying condition of 2.6 mM P_i in DMEM without phenol red with 15% dextran-charcoal-stripped serum to remove steroids from the culture medium. This condition, however, induced marked apoptosis and an increase in calcification (4.7 ± 0.5-fold). Consequently, we performed all experiments in DMEM with 15% complete serum-supplemented medium. Human prostate cancer LNCaP and PC-3 cell lines were maintained in RPMI (Invitrogen) supplemented with 10% fetal bovine serum, 100 units/ml penicillin, and 100 mg/ml streptomycin.

Materials—Testosterone, DHT, 17 β -estradiol, and flutamide, an AR antagonist, were purchased from Sigma. ICI 182,780 was obtained from TOCRIS. These materials were dissolved in absolute ethanol and added to the cultures from a 1000-fold-concentrated stock. Control cultures received similar amounts of ethanol only. Final ethanol concentration did not exceed 0.1% (v/v).

Promoter Reporter Construct—The 1925-bp (−1827/+99) and 1070-bp (−971/+99) Gas6 promoter corresponding to the Gas6 promoter sequences were generated by PCR from human genomic DNA with the appropriate sets of primers. These inserts were cloned into a pGL3 basic vector (Promega). The pGL3-Gas6-ARE mutant construct was made by performing site-directed mutagenesis (Stratagene) with the appropriate primer pairs: AA82CC, 5'-CTGAGAATGGCAAGCCCTCC-ATTAActTCTC-3' (forward primer) and 5'-GAGAGTTA-ATGGAGGGCTTGCCATTCTCAG-3' (reverse primer); AA1281TT, 5'-CCAAGACAAGAGCCAGTTAGTCTTGGT-CTCTGAAG-3' (forward primer) and 5'-CTTCAGAGACCA-AGACTAACTGGCTCTTGCTTGG-3' (reverse primer); CT 1292 GA, 5'-GAGCCAGAAAGTCTTGGTGAAGAC-AAGACAATG-3' (forward primer) and 5'-CATTGTGCTTGTCTTCAGTCACCAAGACTTTCTGGCTC-3' (reverse primer). The constructs were verified by sequencing. The construct of ARE-luciferase (luc) was described previously (19).

Luciferase Assay—HASMC were seeded in 12-well plates at a density of 7 × 10⁴ cells/well and were transiently transfected with 0.8 μ g of ARE-luc construct or Gas6-luc construct using Lipofectamine 2000 (Invitrogen) according to the procedure recommended by the manufacturer. The next day the cells were treated with testosterone, DHT, or ethanol vehicle for an additional 24 h. Aliquots of 20 μ l of cleared lysate were assayed with a luciferase assay kit from Promega. Luciferase activity was normalized to that of vehicle-treated cells and adjusted to the cell protein content.

Small Interfering RNA—Two small interfering RNAs (siRNAs) were designed to target human Gas6 (GenBankTM accession no. NM_000820) using siRNA design software (Dharmacon). The sequences of Gas6 siRNA were 5'-GUGACGAGGGCUUUGCGUA-3' and 5'-GGAGAAGGCUUGCC-

GAGAU-3'. To evaluate the effect of Gas6 siRNA on calcium deposition, both of two siRNA were transfected when HASMC had reached 80~90% confluence and then transfected every time the medium was changed (every 2 days) up to 6 days. AR (GenBankTM accession no. NM_001011645) was knocked down with two siRNAs to evaluate the role of the AR in androgen-stimulated Gas6 transcription activity. The sequences of AR siRNA were 5'-GAGCGUGGACUUUCCGGAA-3' and 5'-UCAAGGAACUCGAUCGUAU-3' (Dharmacon). In HASMC, 6 h after transfection of the Gas6-luc construct, the two AR siRNAs or control siRNA (100 nM) was transfected using transfection reagent (Upstate Biotechnology). The next day DHT or ethanol vehicle was added for an additional 24 h, then luciferase assay was performed. The efficiency of siRNA was validated by immunoblotting the cell lysates at 48 h after transfection.

RNA Extraction, Real-time PCR, and mRNA Stability Analysis—Total RNA was prepared using an RNeasy RNA extraction kit (Qiagen); 3 μ g of total RNA from each of triplicate samples were reverse-transcribed into cDNA using an Omniscript first-strand synthesis system (Qiagen) according to the manufacturer's protocol. Assays for each sample were performed in triplicate using a 7300 real-time PCR system (Applied Biosystems). Then 5 μ l of the cDNA sample was amplified by PCR in a total reaction volume of 50 μ l using SYBR Green master mix (Applied Biosystems) and 500 nM concentrations of the forward 5'-GCCTTTCAGGTCTTCGAGGAG-3' and reverse 5'-GTCAGGCAGGTTTTGCACG-3' primers specific to Gas6. Amplification conditions were 10 min at 95 °C, 40 cycles of 15 s at 95 °C, and 1 min at 60 °C. Data were analyzed by 2^{− $\Delta\Delta$ Ct} method. The relative expression values of all mRNAs were normalized to the β -actin mRNA level (forward 5'-CTGGAACGGTGAAGGTGACA-3' and reverse 5'-AAGGGACTTCCTGTAACAATGC A-3').

To examine Gas6 mRNA stability, HASMC were incubated with actinomycin D (Act D, 5 μ g/ml) in the presence or absence of 2.6 mM P_i or DHT treatment (12 h). Total RNA was extracted at 0, 3, and 6 h after Act D treatment, and the decrease in mRNA expression was determined by real-time PCR analysis as described above. The RNA degradation curve was obtained by setting the maximum mRNA expression at 0 h before Act D treatment as 100%.

Chromatin Immunoprecipitation—Chromatin immunoprecipitation (ChIP) assays were performed using a chromatin immunoprecipitation assay kit (Upstate Biotechnology) according to the manufacturer's instructions. Briefly, HASMC were treated with DHT or ethanol vehicle for 12 h and cross-linked with 1% formaldehyde for 10 min at room temperature. After the cells were collected, nuclei were prepared by incubating the cells in SDS lysis buffer (50 mM Tris (pH 8.1), 10 mM EDTA, 1% SDS). Chromatin was sheared by sonication to an average size of 500~1000 base pairs and diluted 10-fold with dilution buffer. Immunoprecipitation was performed using a polyclonal AR antibody (Santa Cruz Biotechnology), polyclonal acetyl-histone H3 antibody (Millipore), monoclonal p300 antibody (Millipore), and polyclonal rabbit IgG antibody (Santa Cruz Biotechnology). PCR amplification of the Gas6 promoter region spanning the ARE was performed using the following

primers: proximal ARE (5'-GGATGCTGGGCTAACTGC-3') and 5'-GCAACATTGTGCTTGTCTTCA-3'); distal ARE (5'-CAGGCAGAGGCTAGAGATGC-3' and 5'-CAGCAGCC-ATGGATAAACT-3'). In all cases PCR was performed with serial dilutions of the input and various numbers of cycles (25~40 cycles) to ensure that amplification was maintained in the linear range.

Quantification of Calcification—For P_i -induced calcification, P_i (a mixed solution of Na_2HPO_4 and NaH_2PO_4 whose pH was adjusted to 7.4) was added to serum-supplemented DMEM to a final concentration of 2.6 mM. Calcium deposition was evaluated by the *o*-cresolphthalein complexone method (C-Test; WAKO) and von Kossa staining, as previously described (20).

Determination of Apoptosis—To examine the effect of androgens on P_i -induced apoptosis, androgens were added simultaneously to switch the medium of HASMC to medium containing 2.6 mM P_i . Apoptosis was detected by measuring DNA fragmentation with a cell-death detection ELISA^{plus} kit (Roche Applied Science) according to the manufacturer's instructions.

Immunoblotting and Immunofluorescent Analysis—To examine the location of the AR protein, HASMC were separated into cytoplasmic and nuclear fractions using a nuclear extract kit (Active Motif). Nuclear and cytoplasmic fractions (20~30 μ g) were applied to SDS-polyacrylamide gels under reducing conditions and transferred to a polyvinylidene difluoride membrane. Immunoblot analysis was performed using anti-AR polyclonal antibody (Santa Cruz Biotechnology). The effect of androgens on expression of Gas6, phospho-Akt and Akt were examined, as described previously (20). HASMC were grown in 15% fetal bovine serum in DMEM on 2-well chamber slides and fixed in 4% paraformaldehyde for 10 min, and for the AR assay they were incubated with rabbit anti-AR antibody at a 1:250 dilution. Detection of the AR was performed with a 1:100 dilution of fluorescein isothiocyanate-conjugated anti-rabbit antibody (Invitrogen). After several washes, the slides were counterstained with 4',6-diamidino-2-phenylindole.

Statistical Analysis—All values are presented as the mean \pm S.E. Statistical comparisons were made by analysis of variance followed by Fisher's test. A value of $p < 0.05$ was considered statistically significant.

RESULTS

Functional Androgen Receptor Expressed in the VSMC Nucleus—To investigate the action of androgens in VSMC, we first examined whether the AR is expressed in VSMC. In comparison with AR-positive (LNCaP) and AR-negative (PC-3) prostate cancer cells, we found that AR was endogenously expressed in HASMC (Fig. 1A). To determine the location of its expression, we separated the cytoplasmic and nuclear fractions of HASMC. AR was expressed mainly in the nucleus (Fig. 1B). These results were confirmed by immunofluorescence of the AR (Fig. 1C). Next, to examine whether the AR expressed in VSMC is functional, we transfected the ARE-luc construct into HASMC. Androgens (testosterone and DHT) increased luciferase activity by 2~2.5-fold, whereas 17 β -estradiol did not affect its activity. Furthermore, androgen-stimulated ARE activity was abrogated by flutamide, an AR antagonist (Fig. 1D). Taken together these results indicate that the AR expressed in

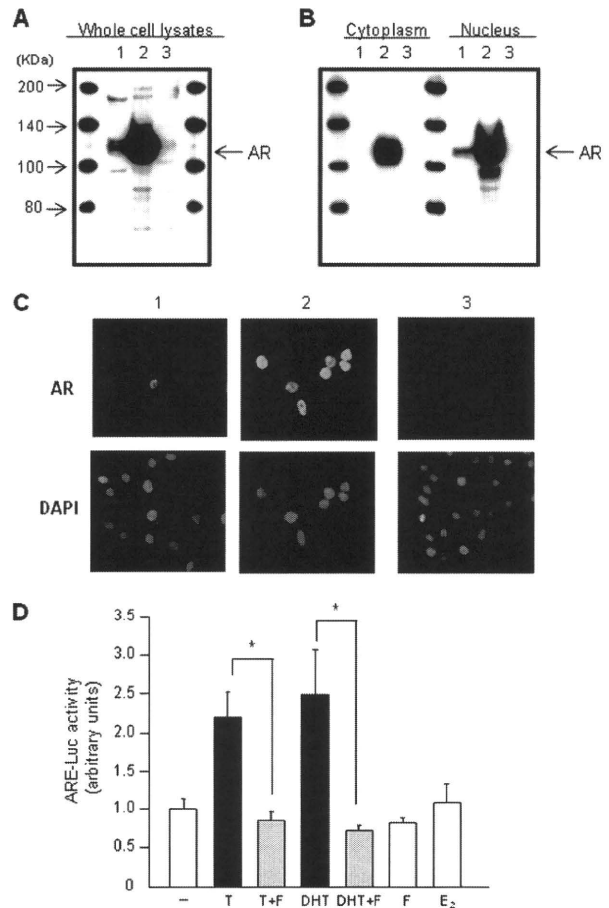


FIGURE 1. Expression of the functional the AR in HASMC. Endogenous expression of the AR in HASMC (lane 1) was examined in whole cell lysates (A) and cytoplasmic and nuclear fractions (B) compared with that in human prostate cancer cell lines, LNCaP (AR-positive; lane 2) and PC-3 (AR-negative; lane 3). C, AR expression was also detected by immunofluorescent staining (green). Nuclei were counterstained with 4',6-diamidino-2-phenylindole (DAPI, blue). D, HASMC were transiently transfected with 0.8 μ g of the ARE-luciferase construct. Twenty-four hours after transfection, androgens (testosterone (T) and DHT, 100 nM), 17 β -estradiol (E₂, 100 nM), and flutamide (F; 10 μ M) were added, and the cells were incubated for an additional 24 h. Relative promoter activities are expressed as the mean \pm S.E. of quadruplicate samples. Similar results were obtained from four independent experiments. *, $p < 0.05$ by Fisher's test.

the nucleus of VSMC participated in androgen-mediated regulation of the ARE.

Androgens Inhibit P_i -induced VSMC Calcification by Restoration of Gas6-mediated Survival Pathway—To investigate the role of the AR in VSMC, we examined the effects of androgens on vascular calcification, a critical and advanced phenotype of atherosclerosis. In the model of P_i -induced calcification (16), calcium deposition was significantly suppressed by both androgens in a concentration-dependent manner (Fig. 2, A and B). We then examined whether the effect of androgens was mediated by the AR. The effect of androgens was clearly abolished by flutamide but not by ICI 182,780, an estrogen receptor antagonist (Fig. 2C). Similar effects on calcification were confirmed by von Kossa staining (Fig. 2D).

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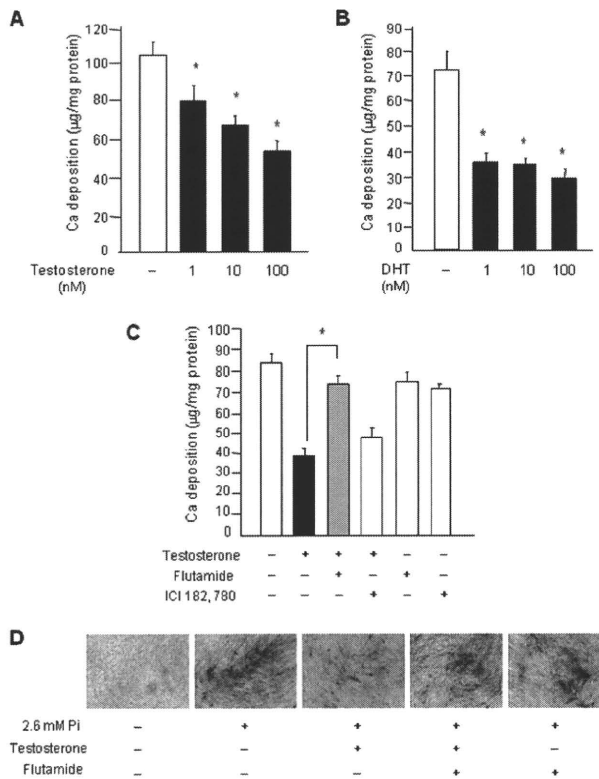


FIGURE 2. Androgens prevent P_i -induced calcification via the AR. HASMC were cultured with the indicated concentrations of androgens (testosterone (A) and DHT (B)) in the presence of 2.6 mM P_i for 6 days. Calcium deposition was measured by the *o*-cresolphthalein complexone method and normalized by cell protein content. *, $p < 0.05$ versus androgens (-) by Fisher's test. HASMC were cultured with flutamide (10 μ M) or ICI 182,780 (10 μ M) in the presence or absence of testosterone (100 nM) with 2.6 mM P_i treatment. On day 6 calcium deposition was measured (C) and was evaluated at the light microscopic level with von Kossa staining (D). All values of calcium deposition are presented as the mean \pm S.E. of quintuplicate samples. Similar results were obtained from three independent experiments. *, $p < 0.05$ by Fisher's test.

Because apoptosis is a crucial and initiating event in P_i -induced VSMC calcification (16, 17), we examined whether androgens inhibit P_i -induced apoptosis. Furthermore, in our recent study apoptosis induced by P_i has been shown to be associated with inhibition of Gas6 expression and secretion (16, 17). Androgens, at concentrations exerting an inhibitory effect on calcification, significantly reduced P_i -induced apoptosis, as quantified by analysis of cytoplasmic histone-associated DNA fragments (Fig. 3A). Flutamide significantly abrogated the inhibitory effect of androgens on apoptosis in HASMC (Fig. 3B). We further examined the effect of androgens on Gas6 expression. Both Gas6 mRNA and protein expression down-regulated by P_i were restored by the addition of testosterone. Moreover, flutamide abrogated the increase in Gas6 expression by testosterone in HASMC (Fig. 3, C and D).

The preventive effect of Gas6 on P_i -induced apoptosis and calcification is mediated by the phosphatidylinositol 3-OH kinase/Akt pathway, a well known anti-apoptotic signaling pathway, through Bcl2 family proteins (17). We found that testosterone restored the Akt phosphorylation down-regulated by

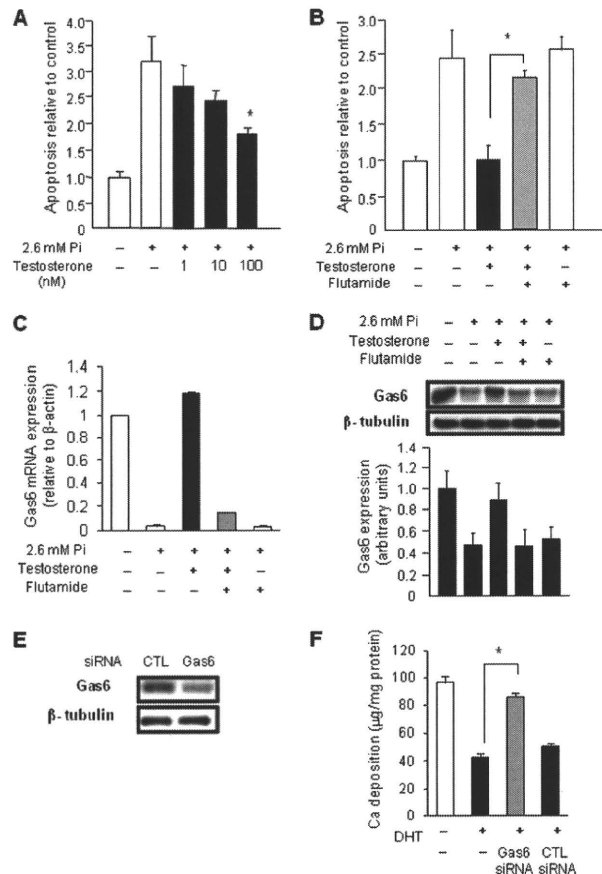


FIGURE 3. Androgens inhibit P_i -induced apoptosis and restore Gas6-mediated survival pathway. A, HASMC were cultured with the indicated concentrations of testosterone in the presence of 2.6 mM P_i for 6 days. A quantitative index of apoptosis, determined by DNA fragmentation enzyme-linked immunosorbent assay, is presented as the value relative to that without P_i treatment. *, $p < 0.05$ versus 2.6 mM P_i , testosterone (-) by Fisher's test. B, HASMC were treated with testosterone (100 nM), or flutamide (10 μ M) in the presence of 2.6 mM P_i for 6 days. C and D, on day 6, RNA and cell lysates were harvested and analyzed for Gas6 mRNA and protein levels by real-time PCR (C) and immunoblotting (D), respectively. β -Actin mRNA and β -tubulin protein levels were also measured as loading control. The average results of three separate measurements of mRNA are shown. The panel shows a representative blot, and bar graphs show quantitative analyses of three independent immunoblotting experiments. E, HASMC were transfected with two Gas6 or control siRNA (100 nM). Gas6 protein was efficiently decreased by two siRNAs targeting Gas6 at 48 h after transfection. CTL, control. F, for measurement of calcium deposition, HASMC were transfected with 100 nM Gas6 siRNA and nonspecific (CTL) siRNA and incubated with DHT (100 nM) and 2.6 mM P_i for 6 days. All values of apoptosis and calcium deposition are presented as the mean \pm S.E. of triplicate samples. Similar results were obtained from three independent experiments. *, $p < 0.05$ by Fisher's test.

P_i , and this increase in phosphorylation was blocked by flutamide (supplemental Fig. 1A). Furthermore, SH-5, an Akt inhibitor, abolished the effect of androgens on HASMC calcification (supplemental Fig. 1B).

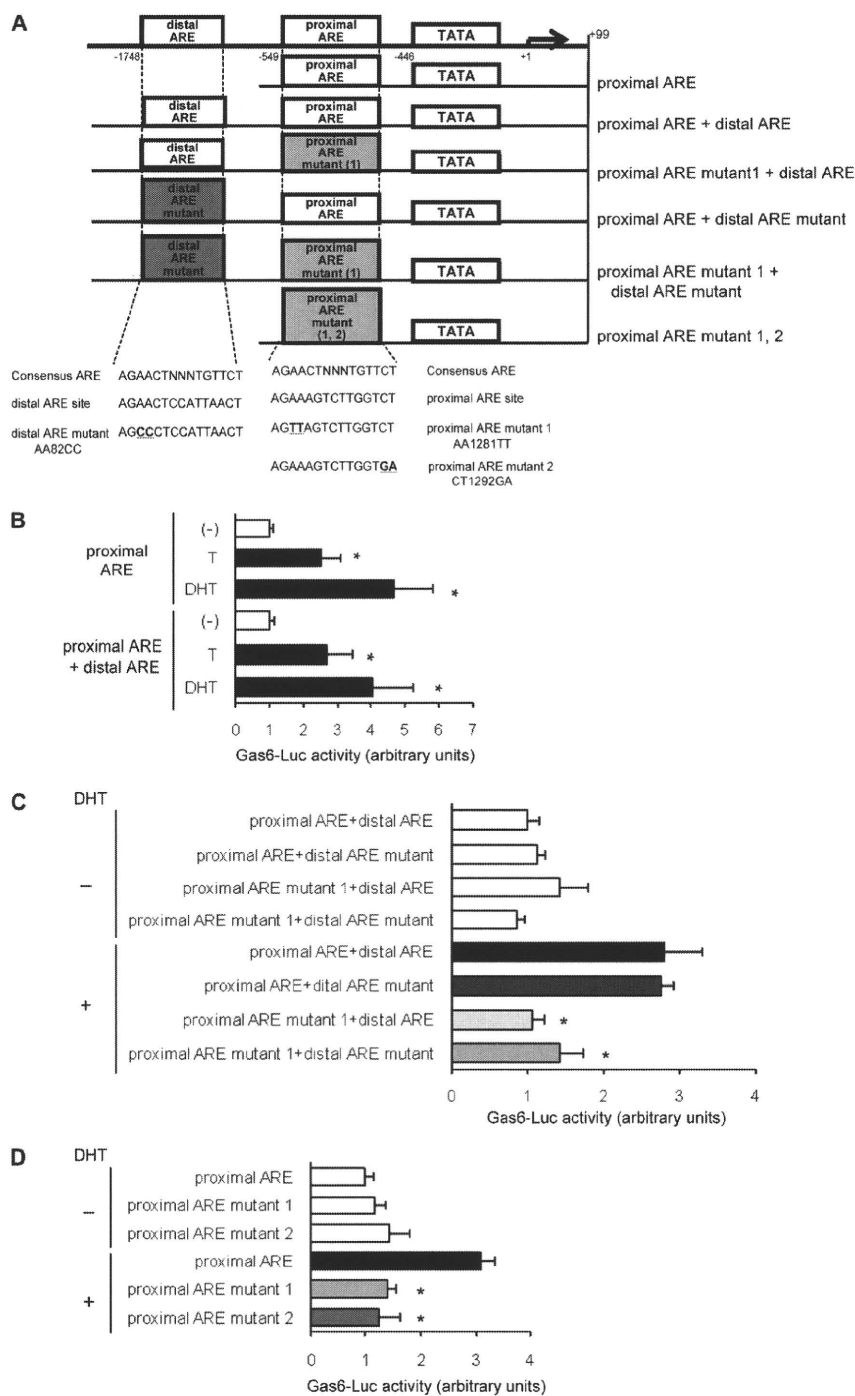
To determine whether Gas6 is required for androgen-mediated effects, we blocked the action of Gas6 using siRNA (Fig. 3E) and examined the effect of androgens on P_i -induced calcification. As shown in Fig. 3F, knockdown of the Gas6 gene significantly reversed the inhibitory effect of androgens on P_i -induced calcification.

The Proximal ARE in Gas6 Promoter Is Essential for Androgen-stimulated Gas6 Transcriptional Activation—To investigate the molecular mechanism involved in up-regulation of Gas6 expression by androgens, we explored the existence of ARE sites in the promoter region of the Gas6 gene (−1827 to +99 bp). We found that the Gas6 promoter contained two consensus ARE sites.

One ARE (−535 to −549 bp) was located close to the transcription start site, whereas the other was located at −1733 to −1748 bp (Fig. 4A). To examine whether AREs in Gas6 were functional, we made two constructs; one contained only the proximal ARE site of the Gas6 promoter, and the other contained both the proximal and distal ARE sites. With transient transfection, androgens significantly stimulated Gas6 promoter activity of the proximal ARE, whereas an additional increase in Gas6 promoter activity was not observed by transfection of the construct containing both the proximal ARE and the distal ARE (Fig. 4B). Then we performed site-directed mutagenesis to confirm whether the proximal ARE is critical. The distal and proximal ARE sites were mutated, as shown in Fig. 4A. Mutation of the proximal ARE completely abrogated DHT-stimulated Gas6 transcription activity. However, we did not observe a reduction in Gas6 transcription activity with the distal ARE mutation (Fig. 4C). To further verify the importance of the proximal ARE sequence in androgen-dependent activation of Gas6, we examined two mutants of the proximal ARE. As expected, both of the mutants abrogated DHT-stimulated Gas6 promoter activity, whereas they had no effect in the absence of DHT (Fig. 4D). Taking these results together, we identified two ARE sites in the Gas6 promoter and found that the proximal ARE is essential for androgen-induced activation of the Gas6 promoter.

Androgen-dependent Gas6 Promoter Activity Is Mediated by Binding of the AR to the ARE—To examine the role of the AR in androgen-dependent Gas6 promoter activation, we used flutamide and AR siRNA to block the function of the AR. First, we found that flut-

amide completely eliminated DHT-induced activation of the Gas6 promoter (Fig. 5A). However, P₁ did not affect Gas6 promoter activity. Next, AR siRNA clearly down-regulated AR protein expression, as shown in Fig. 5B. By transient transfection of AR siRNA, Gas6 promoter activity was significantly inhibited in the



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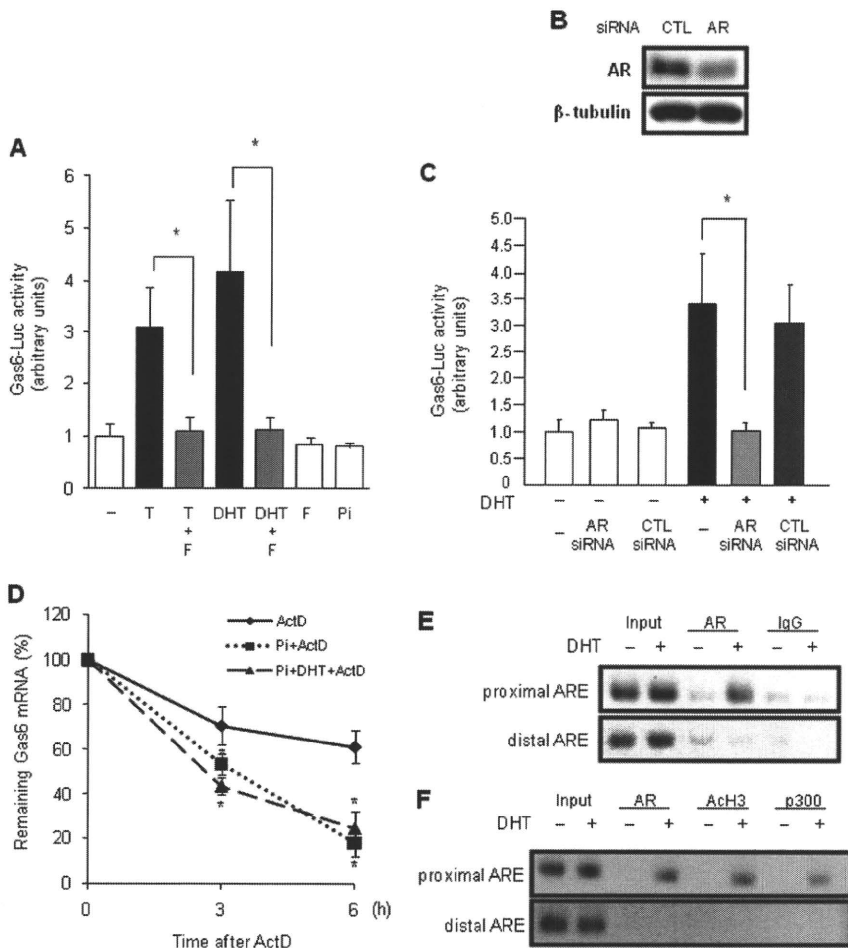


FIGURE 5. Interaction of the AR with the proximal ARE is essential for transactivation of Gas6 gene by androgen. *A*, HASMC were transfected with the Gas6-luc construct containing the proximal ARE. Twenty-four hours after transfection, testosterone (*T*, 100 nM), DHT (100 nM), P_i (P_i , 2.6 mM), or flutamide (*F*, 10 μ M) was added, and the cells were incubated for an additional 24 h. *, $p < 0.05$ by Fisher's test. *B*, HASMC were transfected with AR or control (*CTL*) siRNA (100 nM). The AR protein was efficiently decreased by AR siRNA at 48 h after transfection. *C*, HASMC were transfected with 0.8 μ g of Gas6 proximal ARE together with AR siRNA or nonspecific (*CTL*) siRNA (100 nM). Twenty-four hours later, DHT (100 nM) or vehicle was added. After a further 24 h, luciferase activity was assayed. *D*, serum-starved HASMC were incubated with Act D (5 μ g/ml) in the presence of 2.6 mM P_i after 12 h of DHT (100 nM) treatment. The remaining Gas6 mRNA was determined at 0, 3, and 6 h after Act D treatment by real-time PCR analysis. Values of Gas6 mRNA with P_i (dotted line with squares), with P_i and DHT (dashed line with triangles), or without P_i (solid line) in the presence of Act D were normalized to that of β -actin mRNA at each time point. Gas6 mRNA level at time 0 was expressed as a percentage of the maximum value. The results are the average of three separate experiments. *, $p < 0.05$ versus Act D by Fisher's test. *E*, chromatin extracts were obtained from HASMC after treatment with or without 100 nM DHT for 12 h, and the ChIP assay was performed using an antibody against AR or control IgG. DNA fragments were extracted from immunoprecipitates. The Gas6 promoter region containing proximal ARE was amplified, but distal ARE was not. *F*, a ChIP assay was performed using an antibody against AR, acetylhistone H3 (*AcH3*), or p300 with chromatin extracts with or without treatment with 100 nM DHT for 24 h. Relative promoter activities are expressed as the mean \pm S.E. of quadruplicate samples. Similar results were obtained from four independent experiments. *, $p < 0.05$ by Fisher's test.

FIGURE 4. Androgens stimulate Gas6 promoter activity in HASMC. *A*, shown is a schematic representation of the sequence for ARE sites in wild-type human Gas6 promoter and mutant construct. Site-directed mutagenesis was used to alter the ARE sites within the Gas6 construct. The sequences of the consensus ARE site, Gas6 ARE sites, and the mutated ARE sites with altered bases underlined are shown. *B*, 24 h after transfection of 0.8 μ g of Gas6-luc construct containing only the proximal ARE or the construct containing both the proximal and distal AREs, androgens (testosterone (*T*) and DHT, 100 nM) were added, and the cells were incubated for an additional 24 h. *, $p < 0.05$ versus androgens (-) by Fisher's test. *C*, HASMC were treated with DHT (100 nM) or vehicle for 24 h after transfection of the Gas6-luc constructs containing both proximal and distal AREs or mutants. *, $p < 0.05$ versus DHT (+) wild-type Gas6 by Fisher's test. *D*, HASMC were transfected with wild-type or two proximal ARE mutants. Twenty-four hours after transfection, DHT (100 nM) was added for an additional 24 h. Luciferase activity was normalized to that of the DHT-free wild-type Gas6 construct. *, $p < 0.05$ versus DHT (+) wild-type Gas6 by Fisher's test. Relative promoter activities are expressed as the mean \pm S.E. of quadruplicate samples. Similar results were obtained from five independent experiments.

presence of DHT (Fig. 5C). These findings suggest that Gas6 transactivation by androgens was dependent on the AR.

Because P_i did not affect Gas6 transcriptional activity, we further explored the effect of P_i on Gas6 regulation at the post-transcriptional level. The stability of Gas6 mRNA was examined in the presence or absence of Act D. We found that Gas6 mRNA was significantly more degraded in the presence of P_i than in its absence after Act D treatment (Fig. 5D). DHT did not have an effect on mRNA degradation (Fig. 5D). These findings suggest that P_i down-regulated Gas6 expression by increasing the mRNA degradation rate and not by decreasing transcriptional activity.

To confirm a direct association of the AR with the proximal ARE in the Gas6 gene, we performed a ChIP assay in HASMC. After 12 h of DHT treatment, a polyclonal antibody against the AR could efficiently precipitate the androgen-responsive region of Gas6, showing that the AR directly binds to the Gas6 gene promoter region containing the proximal ARE site in HASMC (Fig. 5E). We did not observe binding of the AR to the distal ARE site in the Gas6 gene (Fig. 5E). Furthermore, we attempted a characterization of the promoter interactions with an AR-containing transcriptional complex. Histone acetyltransferase, such as p300, is a well established coactivator of the AR, and acetylation of histone H3 is an important determinant of AR action, possibly mediated by p300 (19). We performed a ChIP assay with antibodies against acetylhistone H3 and p300. When the AR binds to the proximal ARE site of the Gas6 gene, acetylhistone H3 and p300 also bind to this site as coactivators (Fig. 5E). We did not

observe any binding of the AR, acetylhistone H3, or p300 to the distal ARE site in the *Gas6* gene (Fig. 5F).

DISCUSSION

The effect of testosterone replacement therapy on atherosclerosis is controversial (21–25), although testosterone deficiency is known to be associated with cardiovascular disease in men (26–30). We and others have shown that a low testosterone level is associated with markers of atherosclerosis such as impaired endothelial vasomotor function (27), increased carotid intima-media thickness (28), and aortic calcification (9). Recently, testosterone has also been reported to inhibit VSMC proliferation and neointima formation (7), suggesting a direct action of testosterone on the vasculature. In this *in vitro* study we examined the effect of androgens on P_1 -induced VSMC calcification and found that androgens at physiological concentrations exhibited inhibitory effects on VSMC calcification. In contrast to the present study, it has been reported that androgens induced vascular calcification in apolipoprotein E knock-out mice (31). This discrepancy may derive from the complex *in vivo* effects of testosterone. Further work is required to define the role of androgens in vascular calcification.

Androgens act mainly through transcriptional control of target genes mediated by the nuclear AR (11, 32). In the present study we found that the AR was expressed predominantly in the nucleus of VSMC and had transcriptional activity. Recently, it was demonstrated that the AR-dependent action of androgens protects against angiotensin II-induced vascular remodeling (33). Consistent with this, our results showed that the inhibitory effect of androgens on VSMC calcification was mediated by the AR and not by estrogen receptor.

Recently, we demonstrated that apoptosis plays a central role in the process of P_1 -induced VSMC calcification through downregulation of the *Gas6*-mediated survival pathway (16, 17). In the present study we found that androgens prevented VSMC apoptosis and restored *Gas6* expression and Akt survival signaling. These inhibitory effects of androgens on apoptosis and calcification were eliminated by flutamide and *Gas6* siRNA. Our findings indicate that AR-dependent restoration of *Gas6* by androgens contributes to the inhibition of apoptosis and VSMC calcification.

Although the involvement of other molecules such as protein kinase C δ (7) and endothelial nitric-oxide synthase (33) in the vasoprotective actions of androgens is unclear, our data showed that *Gas6* plays a pivotal role in the inhibitory effect of androgen on P_1 -induced calcification. Several genes containing AREs and having AR-mediated actions have been identified (34, 35). However, little is known about transcriptional regulation and the target genes of the actions of the AR in the vascular system. In this study we identified two AREs in the promoter region of the *Gas6* gene and characterized specific direct binding of the AR to the proximal ARE, in contrast to the nonfunctional distal ARE. Interestingly, Mo *et al.* (36) identified that an estrogen response (ER) element spanning –72 to –89 bp from the translation start site in *Gas6* and ER α is recruited by estrogen-mediated stimulation of *Gas6* gene expression in mouse mammary epithelial cells. In the human *Gas6* promoter domain, we also found the existence of an estrogen response element at –243 to

–251 bp. In clinical studies, a low serum estradiol level in women was correlated with increased arterial calcification (37), and estrogen replacement could reduce coronary calcification (38, 39). However, in experimental studies, estradiol treatment showed variable effects on vascular calcification with either inhibition (40, 41) or stimulation of calcification (42). Further studies are needed to elucidate the actions of estrogens in vascular calcification.

In summary, this study showed that *Gas6* is a novel target that is directly and transcriptionally regulated by the AR, and direct interaction of the AR and *Gas6* mediates the inhibitory effects of androgens on vascular calcification. This study provides a new mechanistic insight into the vascular protective action of androgens.

Acknowledgments—We thank Yuki Ito for technical assistance and Prof. Satoshi Inoue, Department of Geriatric Medicine, Graduate School of Medicine, University of Tokyo, for providing the LNCaP and PC3 cells.

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