

Original Article

Association of Polymorphism of Estrogen Receptor- α Gene with Circulating Levels of Adiponectin in Postmenopausal Women with Type 2 Diabetes

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Aim: Menopause is a risk factor for cardiovascular disease (CVD) in women because of the reduction in endogenous estrogen. Recently, single nucleotide polymorphisms (SNPs) of the estrogen receptor α (ESR-1) gene (c.454-397T>C) associated with the prognosis of myocardial infarction in postmenopausal women were identified; however, the mechanism by which genetic variation of ESR-1 contributes to the pathogenesis of CVD is unknown. Circulating levels of adipokines and inflammatory cytokines predict CVD risk; hence, this study aimed to investigate whether ESR-1 genotypes (c.454-397T>C) might influence circulating levels of adipokines and inflammatory cytokines in postmenopausal women with type 2 diabetes.

Methods: Sixty-three postmenopausal women with type 2 diabetes were recruited. Serum levels of adiponectin, resistin, interleukin-6 (IL-6), and high-sensitive C-reactive protein (hs-CRP) were determined.

Results: The genotype of ESR-1 was closely associated with serum adiponectin, which was decreased in subjects with the T allele and was lowest in those with the T/T genotype. Multiple logistic regression analysis revealed independent contribution of the homozygote for the T allele to low serum levels of adiponectin.

Conclusion: The T allele of the c.454-397T>C SNP of ESR-1 is associated with low serum levels of adiponectin, which may lead to a high risk of CVD in postmenopausal women.

J Atheroscler Thromb, 2009; 16:250-255.

Key words; Estrogen, Adiponectin, Diabetes, SNP, Atherosclerosis

Introduction

It is well known that the incidence of cardiovascular disease (CVD) is lower in women than men because of the protective effect of estrogen on atherosclerosis, and menopause is risk factor of CVD in

women. On the other hand, the increased relative risk of CVD associated with diabetes is greater in women than in men^{1, 2}), indicating that the role of estrogen in the pathogenesis of diabetic vascular disease is complicated. The effects of estradiol on vascular lineage cells are mediated via two different receptors, estrogen receptor (ER)- α and - β . Recently, single nucleotide polymorphisms (SNPs) of the estrogen receptor- α gene (ESR-1) (c.454-397T>C, also known as the PvuII polymorphism) associated with the prognosis of myocardial infarction in women were identified^{3, 4}); however, the underlying mechanism whereby these genetic variations in ESR-1 contribute to the risk of myocar-

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Received: September 7, 2008

Accepted for publication: December 10, 2008

dial infarction in women is unknown.

A series of recent investigations have revealed that adipokines and inflammatory mediators secreted by visceral adipose tissue profoundly contribute to the pathogenesis of insulin resistance and atherosclerosis, and that circulating levels of adipokines, especially adiponectin and resistin, predict the development of CVD^{5,6}. Circulating levels of these adipokines become more proatherogenic during menopause⁷, and low serum adiponectin is a risk of CVD in postmenopausal women with diabetes⁸. Hence, this study aimed to investigate whether ESR-1 genotypes might influence circulating levels of these adipokines and inflammatory mediators in postmenopausal women with type 2 diabetes.

Material and methods

Subjects and Determination of Genotype

Sixty-three postmenopausal women with type 2 diabetes but no diabetic renal complication, identified as having elevated serum creatinine or proteinuria, were recruited. Patients taking thiazolidinedione were excluded. Serum levels of the following parameters were determined: adiponectin by enzyme-linked immunosorbent assay (ELISA) (Otsuka Assay, Tokushima, Japan), resistin by ELISA (BioVender Laboratory Medicine, Brno, Czech Republic), interleukin-6 (IL-6) by chemiluminescent enzyme immunoassay (CLEIA) (Fujirebio, Tokyo, Japan), and high-sensitive C-reactive protein (hs-CRP) by ELISA (Date Behring, Marburg, Germany). DNA extracted from white blood cells was used to characterize ER- α genotypes. All subjects were genotyped for the c.454-397T>C (rs2234693) polymorphism using the sequence-specific primer cycle elongation and fluorescence correlation spectroscopy (SSPCE-FCS) method described elsewhere⁹. SSPCE primers were designed by NovusGene (for details, see <http://www.novusgene.co.jp>) as follows: GAGTTC-CAAATGTCCCAGCT for 5' labeling by TARM and GTTCCAATGTCCCAGCC for 5' labeling by Cy5. This study was approved by the Institutional Review Board of Jikei University School of Medicine and all subjects gave written informed consent.

Statistical Analysis

Correlation between parameters was evaluated by Spearman's rank test. To avoid a type 1 error due to multiple testing, genotype associations with other parameters (resistin, hs-CRP, IL-6, body mass index, adiponectin, hemoglobin A1c) were determined by a permutation test, where the distribution of maximum t statistics based on 100,000 random permutations

Table 1. Demographic characteristics of patients

	Mean	S.D.
Age (years)	63.2	6.1
Postmenopausal age (years)	49.8	5.1
BMI (kg/m ²)	23.9	3.4
HbA1c (%)	7.4	1.0
Adiponectin (μ g/mL)	11.5	6.8
hs-CRP (ng/mL)	705.5	737.2
IL-6 (pg/mL)	1.2	0.6
Resistin (ng/mL)	3.5	1.9
Treatment (n)		
No medication	3	
OHA	35	
Insulin therapy	25	

BMI: body mass index, hs-CRP: high-sensitive CRP, IL-6: interleukin-6, OHA: oral hypoglycemic agents

was compared with the observed values to determine the *p* value and its 95% confidence interval for each gene. Multiple logistic regression analysis was performed to determine factors independently contributing to serum levels of adiponectin. For all statistical analysis, Stata 8.0 (Stata Corp. College Station, TX) was used and *p*<0.05 was considered significant.

Results

Demographic characteristics of the patients are presented in **Table 1**. Serum levels of adiponectin showed a significant negative correlation with body mass index (BMI) (*p*=0.048, *r*=-0.30) (**Fig. 1**). Resistin showed a significant positive correlation with IL-6 (*p*=0.031, *r*=0.32) (**Fig. 2**), but not with BMI. There was no significant correlation between serum levels of adiponectin and resistin.

Changes in variables according to each genotype of ESR-1 are shown in **Table 2**. The incidence of each genotype of c.454-397T>C SNP was T/T 32%, T/C 55% and C/C 13%, respectively. There was no difference in BMI, hemoglobin A1c (HbA1c) or age among the 3 genotypes. Serum levels of adiponectin were decreased in the patients with T allele and were significantly lower in patients with T/T genotype than in those with other genotypes (T/C+C/C). These genotypes showed no association with serum levels of resistin, IL-6 or hs-CRP.

Consequently, the contribution of T/T genotype to serum levels of adiponectin was investigated by multiple logistic regression analysis. Because the median value of serum adiponectin of the patients in this study was 9.6 μ g/mL, we identified patients with serum

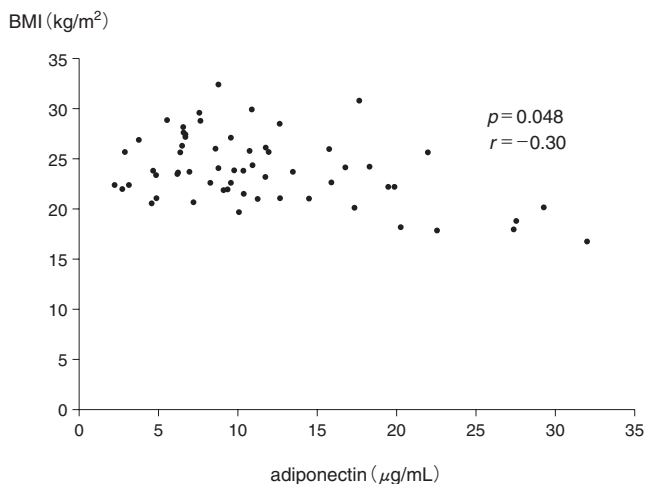


Fig. 1. Relationship between serum levels of adiponectin and BMI

BMI: body mass index

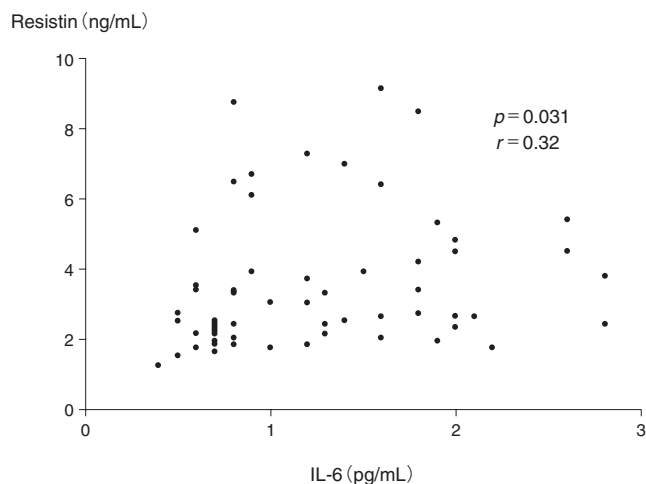


Fig. 2. Relationship between serum levels of resistin and IL-6
IL-6: interleukin-6

Table 2. Genotype of ESR-1 and clinical parameters

	T/T (<i>n</i> =20)	T/C (<i>n</i> =35)	C/C (<i>n</i> =8)	<i>p</i> value	95%CI of <i>p</i> value
Age (years)	63.7 ± 6.7	63.1 ± 5.8	63.0 ± 6.8	0.663	0.660–0.666
BMI (kg/m ²)	23.9 ± 3.6	23.9 ± 3.3	24.0 ± 4.0	0.951	0.950–0.952
HbA1c (%)	7.2 ± 1.0	7.4 ± 0.9	7.6 ± 1.6	0.443	0.440–0.446
Adiponectin (μg/mL)	9.1 ± 6.1	12.3 ± 7.2	14.5 ± 5.1	0.019	0.018–0.020
hs-CRP (ng/mL)	712 ± 781	726 ± 787	599 ± 372	0.667	0.664–0.670
IL-6 (pg/mL)	1.2 ± 0.6	1.3 ± 0.6	1.2 ± 0.7	0.716	0.713–0.718
Resistin (ng/mL)	3.1 ± 1.8	3.5 ± 1.9	4.8 ± 2.1	0.211	0.209–0.214

Figures are expressed as the mean ± S.D. *P* values denote comparison between T/T and T/C + C/C.
CI: confidence interval, BMI: body mass index, hs-CRP: high-sensitive CRP, IL-6: interleukin-6

Table 3. Multiple logistic regression analysis for independent contributors to low serum levels of adiponectin

	Odds Ratio	Standard Error	<i>p</i> value	95%CI
T/T genotype	5.80	4.10	0.013	1.446–23.245
BMI (kg/m ²)	1.28	0.13	0.013	1.054–1.560
Age (years)	0.96	0.05	0.417	0.870–1.059
Postmenopausal age (years)	0.92	0.06	0.172	0.870–1.059

BMI: body mass index, CI: confidence interval

adiponectin ≤ 10 μg/mL (*n*=33) as having low circulating levels of adiponectin, and examined the factors independently involved in low circulating levels of adiponectin as shown in **Table 3**. T/T genotype was an independent contributor to low circulating levels of adiponectin besides BMI.

Discussion

Various anti-atherosclerotic effects of estrogen on vascular cells are believed to be mediated mainly by estrogen receptor (ER)- α , which binds estradiol and regulates gene expression by acting as a transcription factor¹⁰. Menopause is a risk of CVD in women, exacerbating the known cardiovascular risk factors,

including hypertension, dyslipidemia and insulin resistance^{11, 12}). ESR-1 SNP, c.454-397T>C (rs2234693, or PvuII polymorphism), is recognized as a marker that identifies the risk of CVD in women^{3, 4}); however, the investigators were not able to explain this effect by changes in the known CVD risk factors³), and the mechanism whereby this SNP is involved in the progression of CVD is still unknown.

Visceral adipose tissue produces tissue-specific adipokines and inflammatory cytokines^{13, 14}). Both adipokines and inflammatory cytokines are implicated in obesity-associated inflammation, insulin resistance and, consequently, atherosclerosis. Of adipokines, adiponectin plays an important role in the regulation of insulin action via its own receptors in the liver and muscle¹⁵). Adiponectin has anti-atherosclerotic action on endothelial cells and vascular muscle cells^{16, 17}). Thus, low circulating levels of adiponectin could be a risk factor for developing type 2 diabetes^{18, 19}) and CVD^{5, 20}). In contrast, resistin causes insulin resistance²¹) and shows proatherogenic properties^{16, 22}). Circulating levels of resistin are closely associated with the severity of CVD^{23, 24}). In addition to these adipokines, obesity-associated induction of adipose IL-6 production stimulates the liver to secrete CRP, which is not only prognostic for the development of CVD, but also for the risk of type 2 diabetes^{25, 26}). The present study investigated a possible contribution of the specific ESR-1 SNP to circulating levels of these cytokines.

In this study, serum levels of adiponectin correlated significantly with BMI, as is already known (**Fig. 1**). In contrast, serum levels of resistin correlated significantly with IL-6 (**Fig. 2**), but not BMI, indicating that the circulating levels of these adipokines are regulated by a different mechanism, which was also suggested in a previous study²⁷). A novel finding of our study is that the c.454-397T>C SNP of ESR-1 is closely associated with serum levels of adiponectin that were decreased in the patients with the T allele and were lowest in those with the T/T genotype (**Table 2**). In contrast, this SNP showed no association with serum resistin, IL-6 or hs-CRP. Independent contribution of the T/T genotype to serum levels of adiponectin was confirmed by multiple logistic analysis (**Table 3**), where odds ratio of having the T/T genotype for low circulating levels of adiponectin reached 5.8. This study indicates that genetic variation in the action of estrogen plays a role in regulating circulating levels of adiponectin; hence, a homozygote for the T allele could be a risk of CVD in women, as previous investigations identified, because of low circulating levels of adiponectin.

The mechanism whereby this type of ESR-1 SNP

is involved in the regulation of circulating levels of adiponectin remains to be elucidated. This SNP reportedly affects the promoter activity of ER- α and expression of the ER- α promoter containing the T allele results in reduced ER- α expression compared to the C allele²⁸). Interestingly, recent studies suggested a close relationship between ESR-1 SNPs and obesity or metabolic syndrome. Okura *et al.* investigated the association of ESR-1 SNPs with body fat distribution and reported that XbaI polymorphism, which is in close linkage disequilibrium with PvuII polymorphism, was an independent contributor to fat mass and waist-to-hip ratio in middle-aged Japanese women²⁹). The Framingham Heart Study also investigated the sex-specific association of ESR-1 SNPs with adiposity³⁰) and described that waist circumference of homozygous men for C allele of PvuII polymorphism was significantly lower than in homozygous men for the T allele, while this association was not found in women. More recently, Gallagher *et al.* investigated the association of 17 SNPs of ESR-1 with metabolic syndrome and its components in African-American families and found a significant association of c.454-397T>C (rs2234693) with insulin sensitivity³¹). These studies indicate that ESR-1 polymorphisms are involved in the accumulation of visceral fat, which may lead to reduced adiponectin synthesis in visceral adipose tissue; however, it is uncertain whether estrogen affects circulating levels of adiponectin via a direct action.

An animal study using ovariectomized mice investigated the role of estrogen in the regulation of adiposity and lipolytic enzymes. In ovariectomized mice, replacement of estrogen decreased adiposity and adipocyte size, which was associated with the up-regulation of lipolytic enzymes in adipose tissue³²). Estrogen may regulate adiposity via its direct effect on lipolytic enzymes, which would be followed by a change in circulating levels of adiponectin. On the other hand, it is reported that 17- β estradiol enhances adipogenic differentiation of human mesenchymal stem cells and adipose-derived stromal cells^{33, 34}). ER- α plays a role in regulating nuclear factor of activated T-cell (NFAT) 3, which is involved in diverse cellular functions, including adipocyte differentiation³⁵). These findings suggest that estrogen may affect circulating levels of adiponectin, enhancing the differentiation of visceral adipocytes. More recently, estrogen-related receptor α (ERR α) was identified as an orphan nuclear receptor on the basis of its homology with ER- α ³⁶). ERR α function is linked to at least three pathways: estrogen signaling, bone formation and oxidative metabolism³⁷). Interestingly, ERR α and peroxisome proliferator-activated receptor γ (PPAR γ) coactivator (PGC)-1 α are

involved in the modulation of adipogenesis-related genes, such as the activation of PPAR γ during adipocyte differentiation³⁸). Collaboration between ER- α and ERR α in adipocyte differentiation is not well recognized; however, it is likely that ER- α modulates circulating levels of adiponectin in association with the upregulation of adipogenesis-related genes in visceral adipose tissue. The role of the genetic variation of ESR-1 as a contributor to serum adiponectin can be recognized from this view point. Nevertheless, further investigation is required to determine whether this speculation is reasonable.

There are several limitations of the present study. Firstly, the number of patients under investigation was small; however, it is noteworthy that the homozygote for T allele was present in over 30% of women enrolled in this study. We believe that contribution of the T allele of this SNP to circulating levels of adiponectin could be confirmed in a study with more subjects. Secondly, this study is cross sectional and the true risk of CVD in patients homozygous for the T allele should be evaluated by an observational study with prospective follow up.

In conclusion, this study demonstrated that the T allele of the c.454-397T>C SNP of ESR-1 is associated with low serum levels of adiponectin in postmenopausal women with type 2 diabetes, which may lead to a high risk of CVD.

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