

の中で、症例1, 2はFHヘテロ接合体のからの生体肝移植によって、HMG-CoA還元酵素阻害薬等の薬物療法の補助下で良好な脂質コントロールが得られている。移植手術は大きな侵襲を伴うが、2症例とも大きな問題なく成長している。またドナーである両親も術後問題なく経過している。以上から、わずか2例ではあるが、日本人のFHホモ接合体においても、肝臓移植は非常に有効な治療法であると考えられる。

LDL受容体変異のコンパウンドヘテロである症例3は、予想以上にスタチンの効果が認められ、約45%のLDL-C低下が得られている。早期にLDLアフェレシスの導入が望ましいことは明らかだが、遺伝子変異の部位によって良好な薬物治療効果が期待できることは、LDLアフェレシス導入に問題のある症例において薬物治療を考慮する判断材料になると考えられる。

#### E. 結論

FHホモ接合体は代表的な難治性の原発性高脂血症であるが、我が国でもLDLアフェレシスの普及によって生命予後が改善している。しかし希少疾患であるため国内からの報告は少数であり、LDLアフェレシス以外の治療法の有効性や問題点について、全国から症例を集積し検討していく必要がある。

#### F. 健康危険情報

なし

#### G. 研究発表

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- H. 知的財産権の出願・登録状況
1. 特許取得 なし
  2. 実用新案登録 なし
  3. その他 なし

研究成果の刊行に関する一覧表

書籍

著者氏名	論文タイトル名	書籍全体の編集者名	書 籍 名	出版社名	出版地	出版年	ページ
該当なし							

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# Association of glycated albumin with the presence of carotid plaque in patients with type 2 diabetes

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## ABSTRACT

**Aims/Introduction:** Postprandial hyperglycemia is a potent risk factor for cardiovascular disease. Serum glycated albumin (GA) has been reported to reflect postprandial blood glucose fluctuations. In the present study, we assessed the possible correlation of GA with the presence of carotid plaque to evaluate the potential clinical usefulness of GA for predicting atherosclerotic cardiovascular complications in patients with type 2 diabetes.

**Materials and Methods:** Patients with type 2 diabetes ( $n = 236$ ) admitted to Nippon Medical School Hospital (Tokyo, Japan) for glycemic control (aged 19–86 years, 81 females and 155 males) were examined. Clinical measurements were taken on admission. The presence of carotid plaque was assessed by ultrasonography.

**Results:** In patients with carotid plaque ( $n = 154$ ), GA ( $P = 0.023$ ) was higher than those without carotid plaque ( $n = 82$ ). In contrast, neither fasting plasma glucose ( $P = 0.48$ ) nor glycosylated hemoglobin ( $P = 0.41$ ) was significantly different between the groups. The results of logistic regression analysis showed that GA (age- and sex-adjusted odds ratio [95% confidence interval], 1.05 [1.01–1.09];  $P = 0.017$ ) and glycosylated hemoglobin (1.17 [1.01–1.37];  $P = 0.036$ ) were significantly associated with the presence of carotid plaque.

**Conclusions:** The positive correlation of serum GA with the presence of carotid plaque in type 2 diabetes suggests that GA will serve as a useful clinical marker for predicting diabetic cardiovascular complications. (*J Diabetes Invest*, doi: 10.1111/jdi.12085, 2013)

**KEY WORDS:** Carotid plaque, Glycated albumin, Type 2 diabetes mellitus

## INTRODUCTION

Glycated hemoglobin (HbA<sub>1c</sub>) is regarded as a gold standard for monitoring glycemic control. Most expert committees now recommend the use of HbA<sub>1c</sub> in the diagnosis of diabetes<sup>1</sup>. Although the overall usefulness of HbA<sub>1c</sub> is well accepted, epidemiological data suggest that HbA<sub>1c</sub> is not always versatile for predicting all types of diabetic complication; namely, whereas HbA<sub>1c</sub> is a good predictor of microvascular complications, it appears to be less so for macrovascular outcomes<sup>2</sup>. One interpretation is that the fluctuation of blood glucose contributes more to macrovascular diabetic complications than time-averaged blood glucose concentration represented by HbA<sub>1c</sub><sup>2</sup>. Postprandial acute glucose fluctuations are postulated to contribute to the pathogenesis of atherosclerotic cardiovascular complications through the induction of oxidative stress and consequent endothelial dysfunction<sup>3–5</sup>.

Whereas glycation of hemoglobin in erythrocytes (i.e., HbA<sub>1c</sub>) might not reflect blood glucose fluctuations, several reports have shown that glycation of serum albumin reflects the glycemic excursions<sup>6,7</sup>. The glycation of serum proteins has been long assessed by the measurement of fructosamine; however, the reduction reaction-based colorimetric assay is influenced by protein concentration and other coexisting substances in serum<sup>8</sup>. For assessing serum protein glycation more accurately, glycosylated albumin (GA) assay has been developed. GA was originally analyzed by high-performance liquid chromatography (HPLC), but it can currently be determined with automated clinical analyzers by a rapid and specific enzymatic method<sup>9</sup>. The measurement of GA is now available for monitoring glycemic control in patients with diabetes under public health insurance coverage in Japan. GA can be a potential index for predicting cardiovascular events, as it reflects blood glucose fluctuations better than HbA<sub>1c</sub><sup>6,7</sup>. Postprandial hyperglycemia is epidemiologically a more potent risk factor for diabetic cardiovascular complications than fasting plasma glucose (FPG)<sup>10,11</sup>.

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In the present study, we assessed the association of GA with the presence of carotid plaque, a surrogate marker for atherosclerotic disease, to evaluate the potential usefulness of the measurement of GA for predicting cardiovascular complications in type 2 diabetes.

## MATERIALS AND METHODS

### Participants

Patients with type 2 diabetes who were admitted to Nippon Medical School Hospital (Tokyo, Japan) for glycemic control during 2005–2012 were enrolled in the present study ( $n = 236$ , aged 19–86 years, 81 females and 155 males). Exclusion criteria included diabetic nephropathy stage 3 or higher (urinary albumin excretion  $\geq 300$  mg/g-Cr [spot], or  $\geq 300$  mg/day; Joint Committee of the Japan Diabetes Society and the Japanese Society of Nephrology<sup>12</sup>), uncontrolled endocrine disease, steroid treatment, ketoacidosis, estimated primary hyperlipidemia, malignant disease, anemia (hemoglobin  $< 10.0$  g/dL) and other systemic disorders. The present study was approved by the Institutional Review Board, and all participants were enrolled after giving informed consent.

### Clinical Measurements

All participants underwent a physical examination including height, bodyweight and blood pressure on the first morning of admission. Blood sample was taken after an overnight fast. FPG, HbA<sub>1c</sub> and serum GA were measured by the glucose oxidase method (ADAMS Glucose GA-1170; Arkray, Kyoto, Japan), HPLC (ADAMS A1c HA-8160; Arkray) and an enzymatic method using albumin-specific proteinase and ketoamine oxidase (Lucica GA-L; Asahi Kasei Pharma, Tokyo, Japan), respectively. HbA<sub>1c</sub> level was expressed as the percentage value of the National Glycohemoglobin Standardization Program according to the guideline of the Japan Diabetes Society<sup>13,14</sup>. GA was expressed as a percentage of total serum albumin. Serum total cholesterol, high-density lipoprotein (HDL)-cholesterol and triacylglycerol were measured enzymatically (Sekisui Medical, Tokyo, Japan). Low-density lipoprotein cholesterol concentration was calculated by the Friedewald formula<sup>15</sup>. Smoking habit (current, past or never) and type 2 diabetes duration were assessed by interview.

### Carotid Ultrasonography

Carotid artery status was examined with high-resolution B-mode ultrasound systems (SDU-2000; Shimadzu, Kyoto, Japan; iU22 and EnVisor; Philips Healthcare, Andover, MA, USA; LOGIQ 7; GE Healthcare, Little Chalfont, England, UK) equipped with linear transducers with a frequency of 3–12 MHz as previously described<sup>16</sup>. Carotid plaque was defined as a focal intima-media thickening  $\geq 1.0$  mm with marked protrusion into the lumen.

### Statistical Analysis

Continuous variables are expressed as means  $\pm$  standard deviation or median (interquartile range) for variables with normal

or skewed distribution, respectively. Differences in clinical data between participants with and without carotid plaque were assessed by Student's *t*-test, Mann–Whitney *U*-test or  $\chi^2$ -test as appropriate. Correlations between GA and other continuous variables were examined by Pearson's correlation analysis. A logistic regression model was applied to determine the odds ratio for the presence of carotid plaque. A *P*-value of  $< 0.05$  was considered as significant. All analyses were carried out with JMP software (version 9.0; SAS Institute, Cary, NC, USA).

## RESULTS

Of 236 participants enrolled in the present study, 154 (65%) had carotid plaque. The clinical characteristics for each group are shown in Table 1. In the participants with carotid plaque, age was higher and the duration of type 2 diabetes was longer than in those without carotid plaque. Body mass index (BMI) was lower in the participants with carotid plaque compared with those without. In glycemic control indices, GA and GA-to-HbA<sub>1c</sub> ratio (GA/HbA<sub>1c</sub>) were higher in the participants with carotid plaque than in those without, whereas neither FPG nor HbA<sub>1c</sub> was significantly different between the two groups. With regard to diabetic complications, the participants with carotid plaque had a higher incidence of retinopathy or abnormal Achilles tendon reflex than those without. Regarding prehospital medication, a higher population of the participants with carotid plaque had been treated with an antihypertensive agent or antiplatelet agent than those without.

Table 2 shows Pearson's correlation coefficients between GA and other continuous variables. GA was positively correlated with FPG, HbA<sub>1c</sub> and HDL-cholesterol, whereas inversely with systolic blood pressure, diastolic blood pressure and BMI.

Table 3 shows the results of logistic regression analysis for the presence of carotid plaque. Age, duration of type 2 diabetes, GA and GA/HbA<sub>1c</sub> were positively associated with the presence of carotid plaque in unadjusted univariate analysis, whereas BMI was inversely associated with that. After the analysis was adjusted for age and sex, GA and HbA<sub>1c</sub> were significant predictors of the carotid plaque presence. In contrast, no significant association was found between GA/HbA<sub>1c</sub> and the presence of carotid plaque after the adjustment. To address the discrepancies between these glycemic control indices (GA, HbA<sub>1c</sub> and GA/HbA<sub>1c</sub>) and the carotid plaque presence with or without the adjustment, we analyzed the correlations between these glycemic control indices and age. Whereas no correlation was found between GA and age (Figure 1a), HbA<sub>1c</sub> was inversely correlated with age (Figure 1b). As a consequence, GA/HbA<sub>1c</sub> was positively correlated with age (Figure 1c).

## DISCUSSION

HbA<sub>1c</sub> and GA basically provide similar clinical information on recent glycemic control. However, owing to the shorter half-life of serum albumin than erythrocyte hemoglobin, GA reflects shorter-term blood glucose concentration (over 2–3 weeks) compared with HbA<sub>1c</sub>, which reflects that over 2–3 months<sup>17</sup>.

**Table 1** | Clinical characteristics of the participants

Variable	All participants	Carotid plaque		P-value*
	(n = 236)	+	-	
Age (years)	56 ± 13	60 ± 11	50 ± 14	<0.0001
Sex (female/male)	81/155	52/102	29/53	0.81
Duration of type 2 diabetes (years)	5 [0–11]	6 [1–15]	3 [0–10]	0.0069
Systolic blood pressure (mmHg)	127 ± 15	128 ± 15	125 ± 15	0.073
Diastolic blood pressure (mmHg)	75 ± 11	75 ± 10	74 ± 11	0.58
BMI (kg/m <sup>2</sup> )	25.4 ± 4.9	24.8 ± 4.5	26.4 ± 5.6	0.014
Smoking habit, current or past (n [%])	148 [63]	102 [66]	46 [56]	0.13
Fasting plasma glucose (mmol/L)	9.79 ± 2.99	9.89 ± 3.04	9.60 ± 2.89	0.48
HbA <sub>1c</sub> (%)	10.3 ± 2.1	10.4 ± 2.1	10.1 ± 1.9	0.41
GA (%)	28.5 ± 8.4	29.4 ± 8.9	26.8 ± 7.0	0.023
GA/HbA <sub>1c</sub>	2.74 ± 0.45	2.80 ± 0.45	2.63 ± 0.43	0.0037
Total cholesterol (mmol/L)	5.31 ± 1.06	5.28 ± 1.06	5.37 ± 1.07	0.55
HDL-cholesterol (mmol/L)	1.27 ± 0.35	1.26 ± 0.32	1.27 ± 0.41	0.89
LDL-cholesterol (mmol/L)†	3.26 ± 0.89	3.26 ± 0.89	3.25 ± 0.88	0.93
Non HDL-cholesterol (mmol/L)	4.05 ± 1.05	4.02 ± 1.04	4.10 ± 1.08	0.58
Triacylglycerols (mmol/L)	1.72 ± 0.94	1.68 ± 0.88	1.81 ± 1.05	0.31
Retinopathy (n [%])	49 [21]	41 [27]	8 [10]	0.0014
Albuminuria, >30 mg/mg·Cr (n [%])	36 [15]	26 [17]	10 [12]	0.33
Abnormal Achilles tendon reflex (n [%])	98 [42]	75 [49]	23 [28]	0.0019
Prehospital medication				
Insulin (n [%])	29 [12]	23 [15]	6 [7]	0.078
Oral hypoglycemic agent (n [%])	120 [51]	81 [53]	39 [48]	0.46
Statin (n [%])	47 [20]	36 [23]	11 [13]	0.061
Antihypertensive agent (n [%])	75 [32]	57 [37]	18 [22]	0.016
Antiplatelet agent (n [%])	26 [11]	23 [15]	3 [4]	0.0043

Continuous variables are expressed as means ± SD or median [interquartile range]. \*For differences between the subjects with (+) and without (–) carotid plaque. †As low-density lipoprotein (LDL)-cholesterol was calculated by the Friedewald formula, three participants (one in + and two in –) with hypertriacylglycerolemia (≥4.5 mmol/L) were excluded from the statistical analysis of LDL-cholesterol. BMI, body mass index; GA, glycated albumin; GA/HbA<sub>1c</sub>, glycated albumin-to-glycated hemoglobin ratio; HbA<sub>1c</sub>, glycated hemoglobin; HDL, high-density lipoprotein.

GA should therefore be a more suitable index of glycemic control than HbA<sub>1c</sub> in cases where rapid changes might occur in blood glucose concentration; for example, when starting or changing diabetes treatments<sup>6,9,18</sup>. GA is also proposed as a useful measure of glycemic control in populations for whom HbA<sub>1c</sub> might not reflect glycemic status accurately; that is, anemics, those with hemoglobinopathies or neonates<sup>19</sup>. More intriguingly, GA has been reported to reflect blood glucose fluctuations<sup>7,20</sup>. A recent report clearly showed that GA, but not HbA<sub>1c</sub>, was positively associated with glycemic excursions, which were assessed with 48-h continuous glucose monitoring<sup>20</sup>.

Because GA has been reported to reflect blood glucose fluctuations and individuals with greater postprandial glycemic excursions are more likely to develop cardiovascular disease<sup>10,11</sup>, GA might serve as a useful marker for predicting diabetic cardiovascular complications. A recent study actually reported that GA, but not HbA<sub>1c</sub>, was associated with the increasing degree of coronary stenosis in type 2 diabetes<sup>21</sup>. The higher GA level in subjects with carotid plaque among the present participants (Table 1) also supports the clinical

usefulness of GA for evaluating the risk of cardiovascular complications.

The positive correlations of GA with other glycemic indices (FPG and HbA<sub>1c</sub>) suggest that GA increases with the deterioration of overall glycemic control (Table 2). Furthermore, GA was associated with the presence of carotid plaque, and the association remained significant after the logistic regression model was adjusted for age and sex (Table 3). As a recent meta-analysis of prospective cohorts<sup>22</sup> and an additional relevant report<sup>23</sup> showed, there is no doubt that higher HbA<sub>1c</sub> level is associated with the risk of cardiovascular events in the general population without diabetes. However, in subjects with type 2 diabetes, epidemiological evidence indicates a weaker association of HbA<sub>1c</sub> with macrovascular outcomes than that with microvascular complications<sup>24</sup>. Ishizaka *et al.*<sup>25</sup> actually demonstrated that HbA<sub>1c</sub> level was strongly associated with the presence of carotid plaque in normal subjects (normal FPG and normal glucose tolerance); however, the association was not observed in subjects with diabetes or prediabetes. Similar to that report, no significant difference was found in HbA<sub>1c</sub> between the subjects with

**Table 2** | Pearson's correlation coefficients between glycated albumin and other continuous variables

Variable	<i>r</i>	<i>P</i> -value
Age	0.0077	0.91
Duration of type 2 diabetes	-0.041	0.53
Systolic blood pressure	-0.19	0.0028
Diastolic blood pressure	-0.13	0.046
BMI	-0.39	<0.0001
Fasting plasma glucose	0.71	<0.0001
HbA <sub>1c</sub>	0.83	<0.0001
Total cholesterol	0.11	0.10
HDL-cholesterol	0.15	0.022
LDL-cholesterol*	0.083	0.20
Non HDL-cholesterol	0.056	0.39
Triacylglycerols	-0.044	0.50

\*Three participants were excluded from the statistical analysis of low-density lipoprotein (LDL)-cholesterol (see the footnote to Table 1). BMI, body mass index; HbA<sub>1c</sub>, glycated hemoglobin; HDL, high-density lipoprotein.

and without carotid plaque in the present study (Table 1). As patients with poor glycemic control were enrolled in the present study, their higher HbA<sub>1c</sub> levels within a narrow range might be one reason for the lack of association. However, age- and sex-adjusted logistic regression analysis showed the association between HbA<sub>1c</sub> and carotid plaque prevalence (Table 3).

Even though we excluded patients with anemia (hemoglobin <10.0 g/dL) and renal failure (stage 3 or higher), HbA<sub>1c</sub> showed a significant inverse correlation with age in the present participants (Figure 1b). Several reports suggested that HbA<sub>1c</sub>

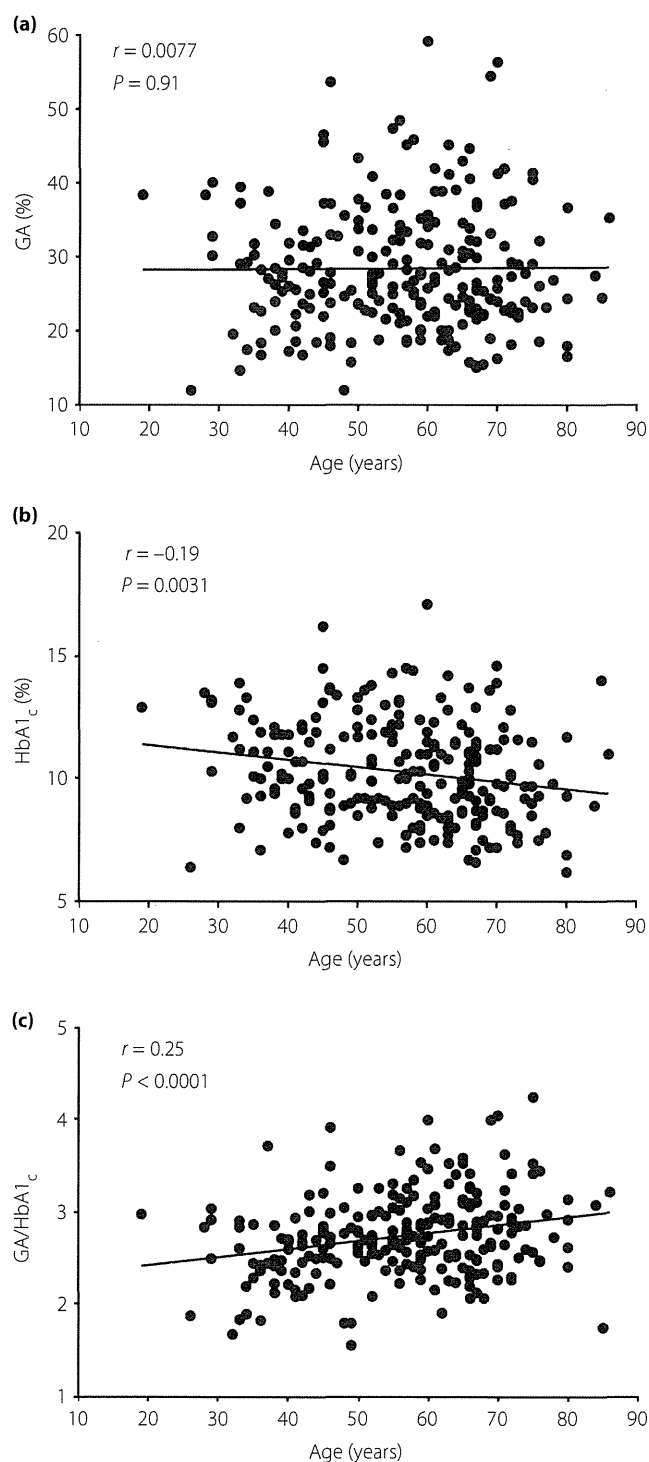
can be affected by age<sup>26,27</sup> and ethnic group<sup>28,29</sup>. Recently, Khoo *et al.*<sup>30</sup> reported a discrepancy between the oral glucose tolerance test (OGTT) and HbA<sub>1c</sub> to diagnose diabetes in elderly populations. They suggested a higher false negative rate when diagnosing diabetes by HbA<sub>1c</sub> alone as compared with by OGTT. These results suggest that HbA<sub>1c</sub> alone is not enough to evaluate glycemic control, especially in elderly patients with type 2 diabetes. Conversely, as GA was not affected by age (Figure 1a), the measurement of GA might have clinical value for evaluating glycemic control more accurately, and for predicting atherosclerotic cardiovascular outcomes, especially in patients with type 2 diabetes.

As GA reflects blood glucose fluctuations better than HbA<sub>1c</sub><sup>6,7</sup>, GA/HbA<sub>1c</sub> might reflect the postprandial glycemic response and could be useful for predicting diabetic complications<sup>31</sup>. Most recently, possible clinical use of both GA and GA/HbA<sub>1c</sub> for predicting the presence of carotid atherosclerosis<sup>32</sup> and the progression of intima-media thickness<sup>33</sup> were also reported in outpatients with type 2 diabetes. However, in the present study, the association between GA/HbA<sub>1c</sub> and the presence of carotid atherosclerosis was not significant after the adjustment for age and sex. Similarly, no significant correlation was found between GA/HbA<sub>1c</sub> and intima-media thickness after the adjustment (data not shown). One of the possible reasons for the differences from the previous reports<sup>32,33</sup> is the age distribution. Relative to the participants in those reports (aged 40–70 years and 53–68 years in the reports of Moon *et al.*<sup>32</sup> and Song *et al.*<sup>33</sup>, respectively), the present participants had a wider age distribution (19–86 years). As HbA<sub>1c</sub> was inversely correlated with age (Figure 1b), GA/HbA<sub>1c</sub> (the index defined by reciprocal

**Table 3** | Odds ratios of variables for the presence of plaque

Variable	Unadjusted		Age- and sex-adjusted	
	OR (95% CI)	<i>P</i> -value	OR (95% CI)	<i>P</i> -value
Age (per 1 year)	1.07 (1.04–1.09)	<0.0001		
Sex (male)*	1.07 (0.61–1.88)	0.81		
Duration of type 2 diabetes (per 1 year)	1.05 (1.01–1.09)	0.0050	1.01 (0.98–1.05)	0.51
BMI (per 1 kg/m <sup>2</sup> )	0.93 (0.88–0.99)	0.015	0.98 (0.92–1.04)	0.49
Smoking habit (current or past)†	1.54 (0.89–2.66)	0.13	1.80 (0.97–3.37)	0.062
Fasting plasma glucose (per 1 mmol/L)	1.03 (0.94–1.13)	0.48	1.06 (0.96–1.17)	0.24
HbA <sub>1c</sub> (per 1%)	1.06 (0.93–1.21)	0.41	1.17 (1.01–1.37)	0.036
GA (per 1%)	1.04 (1.01–1.08)	0.020	1.05 (1.01–1.09)	0.017
GA/HbA <sub>1c</sub> (per 1)	2.57 (1.36–5.05)	0.0032	1.74 (0.87–3.60)	0.12
Total cholesterol (per 1 mmol/L)	0.93 (0.72–1.19)	0.55	1.00 (0.76–1.32)	0.98
HDL-cholesterol (per 1 mmol/L)	0.95 (0.45–2.05)	0.89	0.42 (0.17–1.01)	0.052
Non HDL-cholesterol (per 1 mmol/L)	0.93 (0.72–1.20)	0.58	1.09 (0.82–1.44)	0.56
LDL-cholesterol (per 1 mmol/L)‡	1.01 (0.75–1.38)	0.93	1.08 (0.78–1.52)	0.63
Triacylglycerols (per 1 mmol/L)	0.87 (0.65–1.15)	0.31	1.15 (0.85–1.59)	0.37

BMI, body mass index; CI, confidence interval; GA, glycated albumin; GA/HbA<sub>1c</sub>, glycated albumin-to-glycated hemoglobin ratio; HbA<sub>1c</sub>, glycated hemoglobin; HDL, high-density lipoprotein; OR, odds ratio. \*Female as reference. †Never as reference. ‡Three participants were excluded from the statistical analysis of low-density lipoprotein (LDL)-cholesterol (see the footnote to Table 1).



**Figure 1** | Correlations of (a) glycated albumin (GA), (b) glycated hemoglobin (HbA<sub>1c</sub>) and (c) GA-to-HbA<sub>1c</sub> ratio (GA/HbA<sub>1c</sub>) with age in the participants. Pearson's correlation coefficient is shown in each panel.

value of HbA<sub>1c</sub>) showed a positive correlation with age in the present study (Figure 1c). These results suggest that age-related differences among the glycemic control indices

should be taken into account for clinical use of GA/HbA<sub>1c</sub> as a surrogate marker of diabetic complications.

In the present study, BMI was inversely correlated with GA (Table 2). Although the reasons remain unknown, several reports also showed that BMI was inversely correlated with GA<sup>34–36</sup>. Further investigation into the causes of the relationship between GA and BMI might provide new insights into the interpretation of glycemic control indices in the research and clinical practice of diabetes and its complications.

In conclusion, the present results showed that GA and GA/HbA<sub>1c</sub> were higher in subjects with carotid plaque among patients with type 2 diabetes. Logistic regression analysis showed the positive association of GA and HbA<sub>1c</sub> with the presence of carotid plaque when the models were adjusted for age and sex. Because this was a cross-sectional study, the present data provide just a snapshot of each patient at admission; this is a limitation of the present study. As atherosclerotic lesion formation is a longitudinal event, the longstanding history of vascular conditions should be taken into account. Nevertheless, the present cross-sectional data should warrant further prospective investigation to evaluate the clinical usefulness of GA for predicting cardiovascular outcomes in patients with type 2 diabetes.

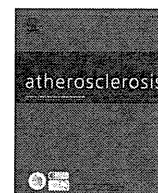
#### ACKNOWLEDGEMENTS

The authors declare that there is no duality of interest associated with this manuscript.

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## Effect of impaired glucose tolerance on atherosclerotic lesion formation: An evaluation in selectively bred mice with different susceptibilities to glucose intolerance



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### ABSTRACT

**Objective:** Impaired glucose tolerance (IGT) is an independent risk factor for atherosclerotic cardiovascular disease. However, due to the lack of appropriate animal models, the underlying mechanisms for IGT-induced atherosclerosis remain to be elucidated *in vivo*. We recently used selective breeding to establish 2 mouse lines with distinctively different susceptibilities to diet-induced glucose intolerance, designated selectively bred diet-induced glucose intolerance-resistant (SDG-R) and SDG-prone (SDG-P), respectively. Here, we assessed atherosclerotic lesion formation in these mice.

**Methods:** Female SDG-R and SDG-P mice were fed an atherogenic diet (AD; 1.25% cholesterol, 0.5% sodium cholate, and 36% energy as fat) for 20 weeks (8–28 weeks of age). Oral glucose tolerance tests were performed during the AD-feeding period. Atherosclerotic lesion formation was quantitatively analyzed in serial aortic sinus sections by oil red O staining. Plasma lipids were measured after the AD-feeding period.

**Results:** Glucose tolerance was impaired in SDG-P mice as compared to SDG-R mice over the 20-week AD-feeding period. No significant differences were observed in any plasma lipid measurement between the 2 mouse lines. Aortic sinus atherosclerotic lesion formation in SDG-P mice was approximately 4-fold greater than that in SDG-R mice.

**Conclusion:** In 2 mouse lines with different susceptibilities to diet-induced glucose intolerance, IGT accelerated atherosclerotic lesion formation. These mice may therefore serve as useful *in vivo* models for investigating the causal role of IGT in the pathogenesis of atherosclerosis.

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### 1. Introduction

Individuals with diabetes have an increased risk for atherosclerotic cardiovascular events, including myocardial infarction, stroke, and peripheral vascular disease [1,2]. In individuals with pre-diabetes as well, impaired glucose tolerance (IGT) is an independent risk factor for cardiovascular disease [3–9]. Although the benefit of intensive glycemic control for improved cardiovascular outcomes did not yield statistically significant results in 2 large prospective studies (the Diabetes Chronic Complications Trial [DCCT] in young subjects with type 1 diabetes [10] and the United Kingdom Prospective Diabetes Study [UKPDS] in subjects with newly diagnosed type 2 diabetes [11]) during the original study

periods, long-term follow-up of those subjects revealed that intensive glycemic control early in the course of diabetes reduced the occurrence subsequent cardiovascular events [12,13]. These intriguing observations of the so-called “legacy effect” or “metabolic memory” imply that glucose intolerance in pre-diabetes or early-stage diabetes may play a pivotal role in the initiation of the atherosclerotic process.

A number of animal models have been used to investigate the role of hyperglycemia in atherosclerosis [14,15]. The predominant models used have been genetically atherosclerosis-prone, hypercholesterolemic mice (apolipoprotein E [apoE]- or LDL receptor [LDLR]-deficient mice) in combination with chemical destruction of pancreatic  $\beta$ -cells (by streptozotocin) or crossbreeding with genetically obese type 2 diabetic models (*db/db* or *ob/ob*). However, severe hypercholesterolemia in apoE- or LDLR-deficient mice often masks the effect of hyperglycemia on the atherosclerotic process [16,17]. In addition, both streptozotocin-treated mice and

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genetically obese mice show severe hyperglycemia even under fasting conditions. Thus, these mice serve as appropriate models for established diabetes rather than for IGT early in the course of diabetes.

Recently, we established 2 mouse lines with distinctively different glucose tolerance by selective breeding. In brief, using C57BL/6, C3H, and AKR as background strains, mice exhibiting superior or inferior glucose tolerance after high-fat feeding were selectively bred and designated as selectively bred diet-induced glucose intolerance-resistant (SDG-R) and SDG-prone (SDG-P), respectively [18]. Since SDG-P mice show evident glucose intolerance as compared with SDG-R mice without any apparent difference in fasting blood glucose levels, these mice may serve as appropriate models for studying IGT-related disorders, including atherosclerotic cardiovascular complications. In this study, we assessed atherosclerotic lesion formation in these novel mouse lines.

## 2. Methods

### 2.1. Animals and diets

Female SDG-R and SDG-P mice (14th–19th generations [18]) bred at the Institute for Animal Reproduction (Kasumigaura, Japan) were used (see Supplementary Fig. S1 for detailed breeding history). Female mice are known to be more susceptible to atherosclerotic lesion formation than males [19,20]. The mice were fed a standard rodent chow (MF; Oriental Yeast, Tokyo, Japan) until 8 weeks of age. Subsequently, the diet was changed to an atherogenic diet (AD) containing 1.25% cholesterol, 0.5% sodium cholate, and 36% energy as fat (F2HFD1; Oriental Yeast; see Supplementary Table S1 for detailed composition). The mice were maintained on the AD for 20 weeks in standard housing (3 or 4 mice per cage) with a 14-h light (06:00–20:00 h)/10-h dark cycle. Food intake, body weight, and random-fed blood glucose levels were monitored every 4 weeks (at 16:00 h). Daily food intake per mouse was calculated from 1-week food consumption per cage. Circadian blood glucose levels were measured on 1 day in the 14th week of AD feeding; measurements were taken every 4 h. This study was conducted under approval from the institutional animal care and use committee of Nippon Medical School.

### 2.2. Oral glucose tolerance test (OGTT)

Glucose tolerance was evaluated by OGTT. At 1 week before (under standard rodent chow), 10 week after, and 19 week after the start of AD feeding, overnight-fasted mice were administered a 20% glucose solution (40, 50, and 60 mg glucose/mouse, respectively) by oral gavage. The dose was based on the average body weight at each time point (approximately 2 g/kg). Blood samples were obtained by tail bleeding. Blood glucose was measured with a glucose sensor (Glutest Neo Super; Sanwa Kagaku Kenkyusho, Nagoya, Japan). The insulin concentration of the plasma was measured by ELISA (Ultra Sensitive Mouse Insulin kit; Morinaga Institute of Biological Science, Yokohama, Japan).

### 2.3. Plasma lipid analysis

At the end of the 20-week AD-feeding period, blood was collected from the inferior vena cava of overnight-fasted mice under anesthesia. Total cholesterol, HDL-cholesterol (sodium phosphotungstate-magnesium chloride precipitation method), triacylglycerols, and non-esterified fatty acids in the blood plasma were measured using commercial kits (Wako Pure Chemical, Osaka, Japan).

### 2.4. Evaluation of atherosclerotic lesion formation

Atherosclerotic lesion formation in the aortic sinus was quantitatively analyzed based on the method of Paigen et al. [20] with modifications. After perfused *in situ* with saline followed by 4% formaldehyde in PBS from left ventricle, the heart was isolated and further fixed overnight with 4% formaldehyde in PBS. It was then cut at a plane parallel to atrial appendages and the upper part including aortic root was embedded in OCT compound (Sakura Finetek, Tokyo, Japan). Cryostat sections were cut from the left ventricular outflow tract and discarded until 3 valve cusps were shown. Then, 45 serial cross-sections (10- $\mu$ m thickness) of aortic sinus were prepared (i.e., covered a distance of 450  $\mu$ m). Of the 45 serial sections, every 5 sections (total of 9 sections each separated by 50  $\mu$ m) were stained with oil red O and counterstained with hematoxylin. The oil red O-stained sections were examined under a light microscope (AX80; Olympus, Tokyo, Japan) with cellSens imaging software (ver. 1.4.1; Olympus). The oil red O-stained area was determined manually from the photomicrograph images [21] on Photoshop Elements software (ver. 9.0.3; Adobe Systems, San Jose, CA). For each mouse, the oil red O-stained area of the 9 sections was averaged and expressed as the mean lesion size.

Immunohistochemical staining was performed to confirm macrophage infiltration into the atherosclerotic lesions. In brief, serial sections to the oil red O-stained ones were stained with MOMA-2 rat monoclonal antibody to mouse macrophages (AbD Serotec, Oxford, UK) using Vectastain Elite ABC kit (Vector, Burlingame, CA) followed by hematoxylin counter stain.

### 2.5. Statistical analysis

Values are presented as mean  $\pm$  SEM. Values of  $p < 0.05$  by Student's *t*-test were considered statistically different between the SDG-R and SDG-P groups.

## 3. Results

### 3.1. Body weight, food intake, and tissue weight

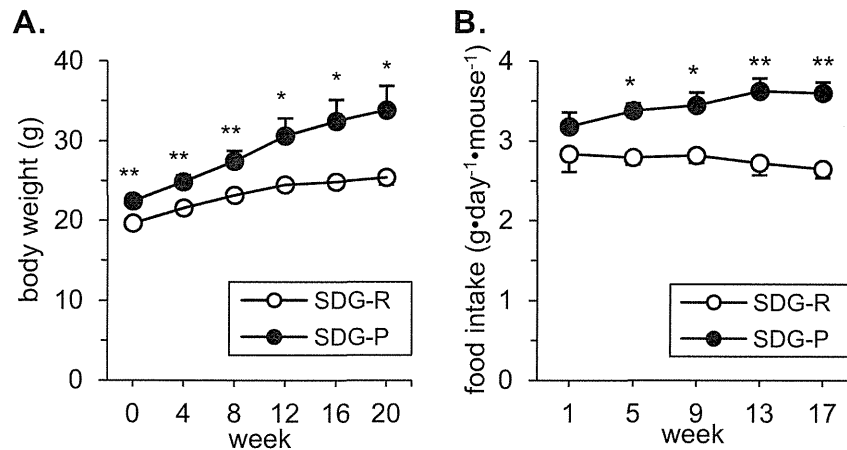
Over the 20-week AD-feeding period, SDG-P mice had a greater body weight than SDG-R mice (Fig. 1A). Food intake of SDG-P mice was also higher than that of SDG-R mice during most of the feeding period (Fig. 1B). At the end of the feeding period, SDG-P mice showed a greater gonadal fat mass as compared with SDG-R mice, whereas no differences were observed in liver weight (Table 1).

### 3.2. Glucose tolerance and random-fed blood glucose

In the OGTT, no significant differences were observed in fasting blood glucose levels between the SDG-R and the SDG-P mouse groups at 1 week before, 10 weeks after, and 19 weeks after the start of AD feeding (Fig. 2A–C; 0 min). However, SDG-P mice showed higher blood glucose levels at 30 min and subsequent times after glucose injection as compared with SDG-R mice during the 20-week AD-feeding period (Fig. 2A–C). Although a statistical difference was observed in the fasting insulin levels before the start of AD feeding, no such differences were observed in post-challenge insulin levels over the feeding period (Fig. 2E–G).

Random-fed blood glucose levels were higher in SDG-P mice than in SDG-R mice over the AD-feeding period (Fig. 3A). In addition, the circadian blood glucose profile revealed that SDG-P mice had higher blood glucose levels throughout the day with greater fluctuations than those seen in SDG-R mice (Fig. 3B). The standard deviations of individual circadian blood glucose levels (6 time





**Fig. 1.** Body weight gain (A) and food intake (B) of glucose intolerance-resistant (SDG-R) and prone (SDG-P) mice during atherogenic diet feeding. Data are expressed as mean  $\pm$  SEM (A,  $n = 10$ –13 mice; B,  $n = 4$  cages). \* $p < 0.05$ , \*\* $p < 0.01$ .

points) were  $0.60 \pm 0.09$  and  $1.01 \pm 0.12$  mmol/l in the SDG-R and SDG-P groups, respectively ( $p = 0.013$ ,  $n = 5$ –6).

### 3.3. Plasma lipid profile

Table 1 shows the plasma lipid profiles of the SDG-R and the SDG-P mouse groups at the end of the 20-week AD-feeding period. No significant differences were observed in total cholesterol, HDL-cholesterol, non-HDL-cholesterol, triacylglycerols, or non-esterified fatty acid levels between the 2 mouse lines.

### 3.4. Atherosclerotic lesion formation

At the end of the 20-week AD-feeding period, lipid-laden plaque formation was observed in the intimal area of the aortic sinus (Fig. 4A–D). The lesion formation was confirmed by the immunohistochemical detection of macrophage infiltration, another hallmark of atherosclerotic lesions (Fig. 4E–H). Atherosclerotic lesion formation in SDG-P mice was approximately 4-fold greater than that in SDG-R mice (Fig. 4I).

## 4. Discussion

IGT has been postulated to play an important role in the pathogenesis of atherosclerosis, but few animal models are available to investigate the causal mechanisms appropriately. Existing diabetic models display severe hyperglycemia, even in fasting conditions. In addition, the severe hyperglycemia is often accompanied by dyslipidemia [16,17]. The lipid abnormalities in diabetes make it

difficult to dissociate the effects of hyperglycemia *per se* from those of dyslipidemia.

In contrast to existing diabetic models, the present results demonstrate that SDG-P mice have the characteristics of a useful IGT model (modest hyperglycemia under post-challenge or random-fed conditions, but not fasting conditions), while SDG-R mice provide a normal glucose-tolerant control. Fasting blood glucose levels in SDG-P mice tended to be increased during the AD-feeding period most probably due to the insulin resistance arising from increased fat deposition (see Supplementary Fig. S2 for the results of insulin tolerance test). At 19 weeks after the start of AD feeding, fasting blood glucose levels of SDG-R and SDG-P were  $3.84 \pm 0.26$  ( $n = 6$ ) and  $8.26 \pm 1.51$  ( $n = 4$ ), respectively ( $p = 0.07$ ; Fig. 2C); however, the levels were still far lower than those of existing diabetes models such as streptozotocin-treated mice or *db/db* mice.

In addition, no significant differences in the lipid profiles were observed between the 2 mouse lines. Even under the AD feeding, these mice did not display overt hypercholesterolemia. Due to the overwhelming impact of cholesterol on atherosclerosis, extremely high levels of non-HDL-cholesterol in apoE- or LDLR-deficient mice can hamper any attempt to isolate the effect of hyperglycemia [16,17]. In contrast to the hypercholesterolemic mice in the abovementioned study, plasma lipid levels in our mice were almost comparable to the normal range in humans. Thus, our mouse lines will be appropriate models for defining the effect of glucose intolerance without the concomitant effect of overt dyslipidemia. However, more detailed lipoprotein fraction analysis may be needed to clarify the lipoprotein distributions. In patients with type 2 diabetes, lipoprotein abnormalities in the subclasses (in the size and function) have been discussed in relation to the increased cardiovascular risk [22]. Furthermore, a most recent report demonstrated that the subclass abnormalities already exist in individuals with IGT [23]. Hence, the subclass analysis may be useful in understanding the IGT-related lipoprotein abnormalities and their possible involvement in the accelerated lesion formation in SDG-P mice.

SDG-P mice with IGT were more susceptible to atherosclerotic lesion formation than glucose-tolerant SDG-R mice. Because our mice are not genetically engineered to form overt atherosclerotic lesions, the lesion formation was much less than that in apoE- or LDLR-deficient mice. Our mice are therefore not suitable for studying the advanced-stage of atherosclerotic disorders, but will be appropriate for investigating early stage atherogenesis. As

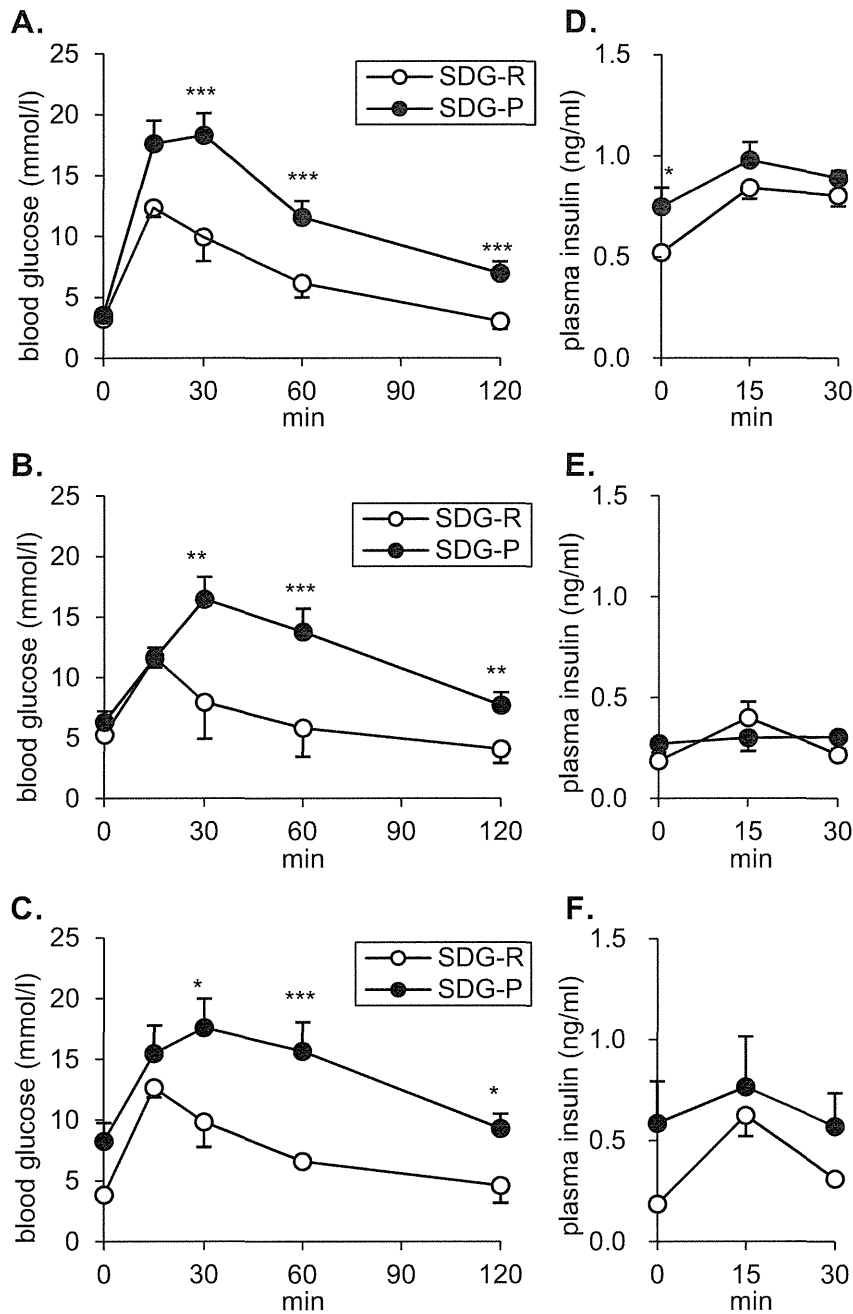
**Table 1**  
Relative tissue weights and plasma lipid profiles of glucose intolerance-resistant (SDG-R) and prone (SDG-P) mice after atherogenic diet feeding.

	SDG-R ( $n = 12$ )	SDG-P ( $n = 11$ )	$p$ value <sup>a</sup>
Tissue weight, mg/g body weight			
Liver	$78.7 \pm 5.0$	$67.1 \pm 3.4$	0.070
Gonadal fat	$13.2 \pm 1.7$	$27.1 \pm 4.2$	0.009
Plasma lipids, mmol/l			
Total-cholesterol	$4.36 \pm 0.33$	$4.36 \pm 0.73$	1.00
HDL-cholesterol	$1.32 \pm 0.34$	$0.74 \pm 0.11$	0.13
Non-HDL-cholesterol	$3.03 \pm 0.26$	$3.62 \pm 0.65$	0.42
Triacylglycerols	$0.39 \pm 0.04$	$0.43 \pm 0.05$	0.50
Non-esterified fatty acids	$0.77 \pm 0.07$	$0.72 \pm 0.07$	0.67

Values are expressed as mean  $\pm$  SEM.

<sup>a</sup> Student's *t*-test.



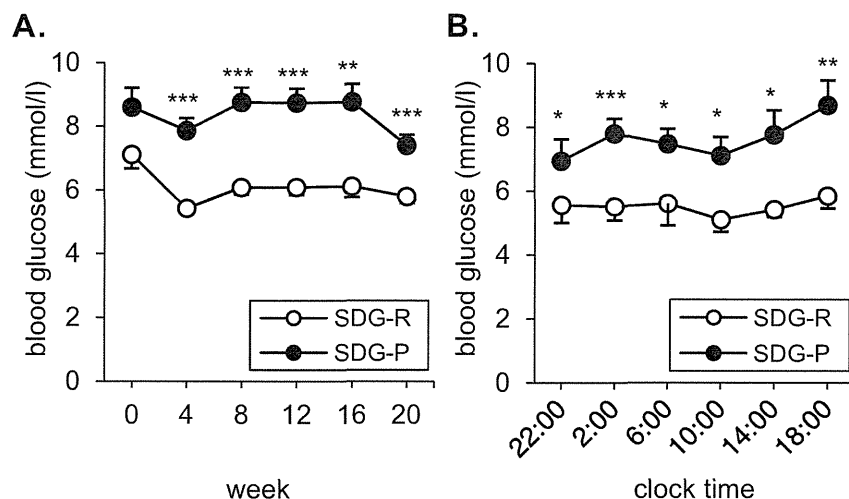


**Fig. 2.** Blood glucose (A–C) and plasma insulin (D–F) levels of glucose intolerance-resistant (SDG-R) and prone (SDG-P) mice in OGTT at 1 week before (A and D; 40 mg glucose/mouse), 10 weeks after (B and E; 50 mg glucose/mouse), and 19 weeks after (C and F; 60 mg glucose/mouse) the start of atherogenic diet feeding. Data are expressed as mean  $\pm$  SEM of 4–6 mice. \* $p < 0.05$ , \*\* $p < 0.01$ , \*\*\* $p < 0.001$ .

demonstrated in follow-up studies of the DCCT and UKPDS trials [12,13], glycemic control in early stage diabetes is important to prevent subsequent cardiovascular complications. This strongly suggests that IGT can modify early stage atherogenesis, which eventually leads to increased incidence of and mortality from cardiovascular events. Renard et al. [24] demonstrated in LDLR-deficient mice with viral  $\beta$ -cell destruction that hyperglycemia leads to the initiation of atherosclerotic lesions, whereas advanced lesion formation is largely dependent on diabetes-induced dyslipidemia. They also elucidated the distinct effect of hyperglycemia on early stage atherosclerosis; however, their diabetic mice had both severe hyperglycemia ( $\sim 20$  mmol/l) and hypercholesterolemia ( $\sim 9$  mmol/l), even under cholesterol-free diet feeding [24].

Moderately hyperglycemic SDG-P mice without apparent dyslipidemia will be preferable for studying the causal relationship between pre-diabetes or early stage diabetes and the pathogenesis of atherosclerosis. Recently, Bartels et al. [25] also reported that prolonged feeding (12 months) of high-fat diet (60% energy as fat) led to a high incidence (64%) of small atherosclerotic lesions in C57BL/6 mice in coincidence with glucose intolerance but without overt dyslipidemia.

Since considerable attention has been focused on the epidemiological impact of IGT on cardiovascular complications, a number of studies have reported the possible roles of glucose fluctuations, a hallmark of IGT, in atherogenesis. For example, intermittent high glucose exposures were demonstrated to induce oxidative stress

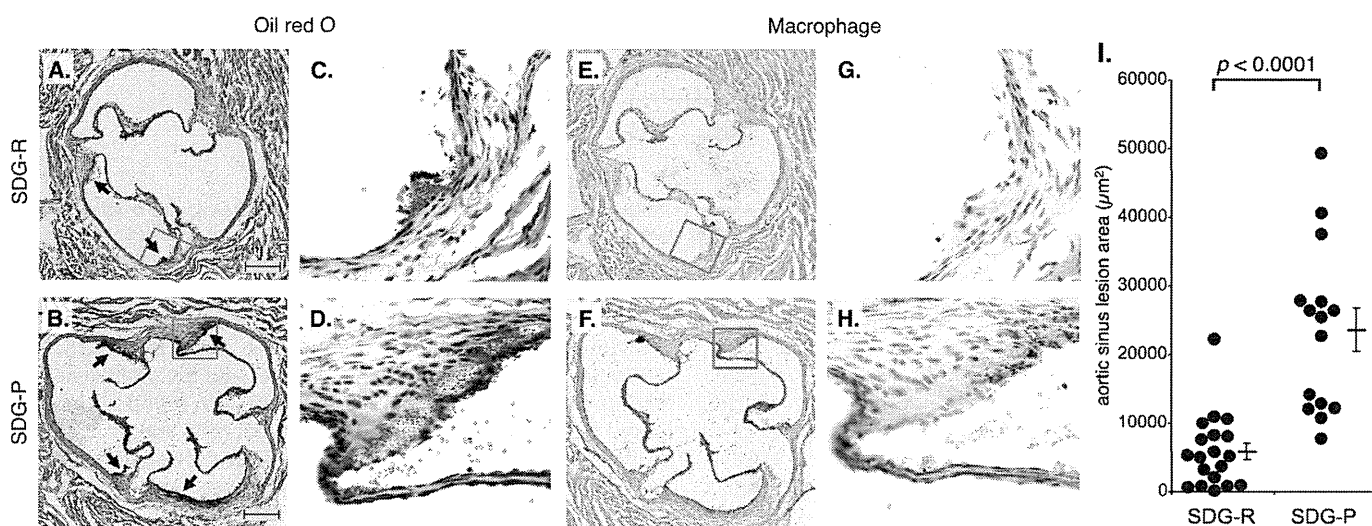


**Fig. 3.** Random-fed blood glucose levels and circadian blood glucose profiles in glucose intolerance-resistant (SDG-R) and prone (SDG-P) mice. (A) Random-fed blood glucose levels were measured every 4 weeks at 16:00 h during atherogenic diet feeding. (B) Circadian blood glucose profiles were obtained from measurements taken at 4-h intervals on 1 day in the 14th week of atherogenic diet feeding. Data are expressed as mean  $\pm$  SEM (A,  $n = 10$ –13; B,  $n = 5$ –6). \* $p < 0.05$ , \*\* $p < 0.01$ , \*\*\* $p < 0.001$ .

and subsequently produce various pro-atherogenic changes in cultured vascular endothelial cells, smooth muscle cells, and monocytes/macrophages (e.g., augmented expressions of adhesion molecules and inflammatory cytokines) [26]. Transient hyperglycemia-induced oxidative stress was also demonstrated in humans [27]. In the present study, SDG-P mice had greater circadian variations in random-fed blood glucose levels as compared with SDG-R mice, suggesting that the fluctuations of blood glucose, in addition to the mean levels, were likely to be involved in the accelerated lesion formation observed in SDG-P mice. Hence, these mice may contribute to a better understanding of the cellular and molecular mechanisms that are responsible *in vivo* for IGT-induced atherogenesis. For example, biochemical and histological analyses of the causal mechanisms (e.g., macrophage infiltration into and adhesion molecule expression on the aortic wall) at earlier time points are now of great interest.

Because SDG-R and SDG-P mice are outbred lines from 3 inbred strains (C57BL/6J, C3H/HeJ, and AKR/N) [18], our mice harbor

certain genetic variations. The genetic diversity was a key factor in the wide distribution of glucose tolerance, which enabled us to establish the 2 mouse lines. At the same time, although glucose tolerance was the sole criterion for the selective breeding, we cannot exclude the possibility that other genetic factors, which can affect atherogenesis independent of blood glucose levels, were concomitantly selected for in the SDG-R and SDG-P mouse lines. Among the 3 background strains, C3H and AKR are known to be highly resistant to atherosclerotic lesion formation, whereas C57BL/6 is relatively prone to that [19,28,29]. The genetic diversity in susceptibility to atherosclerosis may partially account for the different lesion size between SDG-R and SDG-P independent of blood glucose levels. For example, although it remains controversial [30], a defective allele of Toll-like receptor 4 (*Tlr4*) in C3H/HeJ has been implicated in the resistant phenotype to atherosclerosis [31]. Unexpectedly, however, the defective *Tlr4* allele was found only in atherosclerosis-prone SDG-P mice, but not in SDG-R (see Supplementary Fig. S3 for the results of genotyping). Hence, the



**Fig. 4.** Atherosclerotic lesion formation in the aortic sinus of glucose intolerance-resistant (SDG-R) and prone (SDG-P) mice after atherogenic diet feeding. (A, B) Representative images of oil red O-stained atherosclerotic lesions in the aortic sinus (arrows). Bar: 200  $\mu\text{m}$  (C, D) Higher magnification images from the fields indicated by squares in panels A and B. (E, F) Immunohistochemical staining for macrophages (brown) in the serial cross-sections to the corresponding oil-red O-stained images. (G, H) Higher magnification images from the fields indicated by squares in panels E and F. (I) Quantitative analysis of the lesion area. Each dot indicates mean lesion size in each mouse. Bars indicate mean  $\pm$  SEM of 15–19 mice.

atherosclerosis-resistant phenotype of SDG-R mice should not be the result of defective TLR4 signaling. A polymorphism in vascular cell adhesion molecule 1 (*Vcam1*) among the 3 background strains is also reported to affect the development of atherosclerosis [32,33]. The polymorphism of *Vcam1* was found in both mouse lines (Supplementary Fig. S3). In any case of *Tlr4* or *Vcam1* genotypes inherited in both mouse lines, however, SDG-P mice showed greater lesion formation than SDG-R mice carrying the same alleles. In addition, the hyperphagic behavior and consequent greater weight gain observed in the SDG-P as compared with the SDG-R mice might be determined by factors independent of glucose intolerance. Further research into such genetic variations will strengthen the value of the novel mouse lines as IGT-induced atherosclerosis model animals.

In this study, a cholate-containing AD was used to ensure the atherosclerotic lesion formation in wild-type mice. Besides the critical role in facilitating intestinal cholesterol absorption, bile acids can modulate diverse signaling pathways in metabolic homeostasis and inflammation [34–36]. The potential pleiotropic effects of exogenous cholate may therefore be taken into account in interpreting the present results.

In conclusion, the present results demonstrate that glucose intolerance-prone SDG-P mice are more susceptible to atherosclerotic lesion formation than glucose-intolerance resistant SDG-R mice. The novel mouse lines with different susceptibilities to glucose intolerance may therefore serve as useful animal models for studying the pathogenesis, prevention, and treatment of IGT-induced atherosclerotic disorders.

### Conflicts of interest

The authors declare that there is no duality of interest associated with this manuscript.

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### Appendix A. Supplementary data

Supplementary data related to this article can be found online at <http://dx.doi.org/10.1016/j.atherosclerosis.2013.10.009>.

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# Characterization of Pancreatic Islets in Two Selectively Bred Mouse Lines with Different Susceptibilities to High-Fat Diet-Induced Glucose Intolerance

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## Abstract

Hereditary predisposition to diet-induced type 2 diabetes has not yet been fully elucidated. We recently established 2 mouse lines with different susceptibilities (resistant and prone) to high-fat diet (HFD)-induced glucose intolerance by selective breeding (designated selectively bred diet-induced glucose intolerance-resistant [SDG-R] and -prone [SDG-P], respectively). To investigate the predisposition to HFD-induced glucose intolerance in pancreatic islets, we examined the islet morphological features and functions in these novel mouse lines. Male SDG-P and SDG-R mice were fed a HFD for 5 weeks. Before and after HFD feeding, glucose tolerance was evaluated by oral glucose tolerance test (OGTT). Morphometry and functional analyses of the pancreatic islets were also performed before and after the feeding period. Before HFD feeding, SDG-P mice showed modestly higher postchallenge blood glucose levels and lower insulin increments in OGTT than SDG-R mice. Although SDG-P mice showed greater  $\beta$  cell proliferation than SDG-R mice under HFD feeding, SDG-P mice developed overt glucose intolerance, whereas SDG-R mice maintained normal glucose tolerance. Regardless of whether it was before or after HFD feeding, the isolated islets from SDG-P mice showed impaired glucose- and KCl-stimulated insulin secretion relative to those from SDG-R mice; accordingly, the expression levels of the insulin secretion-related genes in SDG-P islets were significantly lower than those in SDG-R islets. These findings suggest that the innate predispositions in pancreatic islets may determine the susceptibility to diet-induced diabetes. SDG-R and SDG-P mice may therefore be useful polygenic animal models to study the gene–environment interactions in the development of type 2 diabetes.

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## Introduction

Gene–environment interactions play a crucial role in the development of type 2 diabetes. For analyzing genetic factors, the polygenic background of selectively bred animal models has been investigated [1]. For instance, the Goto-Kakizaki (GK) rat [2] and Nagoya-Shibata-Yasuda (NSY) mouse [3] are non-obese diabetic models produced by repetitive selective breeding for impaired glucose metabolism. Through the analyses of these polygenic rodent models, increasing numbers of candidate genes for the pathogenesis of type 2 diabetes have been identified, most of which well resemble the genetic basis of type 2 diabetes in humans [1,4,5].

Contemporary environmental factors (*e.g.*, nutritional excess and sedentary lifestyle) cause obesity, which leads to insulin resistance in peripheral tissue [6,7]. However, not all obese individuals with insulin resistance develop type 2 diabetes because the functional and morphological compensation capacities of  $\beta$  cells against insulin resistance vary between individuals [8]. In

rodent models, animals with high-fat diet (HFD)-induced obesity are chiefly used for assessing the impact of excess dietary fat as an environmental factor. However, the propensity for developing diet-induced diabetes varies widely even in a single strain [9]. Thus, individual differences in susceptibility to environmental factors are postulated to be determined by genetic factors. Existing polygenic models, which develop diabetes spontaneously [10], may therefore not be always appropriate to investigate the predisposition to lifestyle-related disorders because environmental factors had not been taken into account in their selective breeding.

To establish novel rodent models that can mimic the gene–environment interactions in the development of type 2 diabetes, we have performed a selective breeding of mice. In brief, using 3 inbred strains (C57BL/6, C3H, and AKR) as background, mice exhibiting superior and inferior glucose tolerance after HFD feeding have been bred repetitively to establish 2 distinct mouse lines with different susceptibilities (resistant and prone) to HFD-induced glucose intolerance, designated selectively bred diet-induced glucose intolerance-resistant (SDG-R) and -prone (SDG-