



# Insulin Resistance is Associated with Longitudinal Changes of Cardiac Repolarization Heterogeneity in Apparently Healthy Subjects

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## ABSTRACT

**Introduction:** Increased heterogeneity in ventricular repolarization is a risk factor of sudden cardiac death, but its natural history is unclear. Here we examined whether insulin resistance is associated with longitudinal change in ventricular repolarization heterogeneity in apparently healthy subjects.

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**Methods:** The study subjects were participants in health checkups in cohort 1 and cohort 2, which were followed up for 6 years and 5 years, respectively. Subjects with diabetes, cardiovascular disease, or renal disease at baseline were excluded from the analyses. As indices of insulin resistance, the homeostasis model assessment of insulin resistance (HOMA-IR) and triglyceride to HDL-cholesterol ratio (TG/HDL-C) were used in cohort 1 and cohort 2, respectively. Heterogeneity in ventricular repolarization was assessed by heart rate-corrected Tpeak–Tend interval in V5 (cTpTe), QT interval, and QT dispersion. In regression analyses, parameters with a skewed distribution were normalized by logarithmic transformation or by Box–Cox transformation.

**Results:** In longitudinal analyses, Box–Cox-transformed cTpTe at the end of follow-up was weakly correlated with log HOMA-IR at baseline in cohort 1 ( $n = 153$ ,  $r = -0.207$ , 95% CI  $-0.354$  to  $-0.050$ ,  $p = 0.010$ ) and with log TG/HDL-C at baseline in cohort 2 ( $n = 738$ ,  $r = -0.098$ , 95% CI  $-0.169$  to  $-0.026$ ,  $p = 0.008$ ). Multiple regression analysis showed that indices of insulin resistance, but not glycosylated hemoglobin (HbA1c) or plasma glucose, at baseline were significant explanatory variables for cTpTe at the end of follow-up. Neither QT interval nor QT dispersion was correlated with metabolic parameters.

**Conclusion:** Insulin resistance may be involved in the longitudinal increase of ventricular

repolarization heterogeneity in apparently healthy subjects.

**Keywords:** Electrocardiogram; Insulin resistance; Ventricular repolarization

## INTRODUCTION

The number of patients with diabetes mellitus has increased by fourfold worldwide in the past three decades [1], indicating expansion of the global risk of cardiovascular events. One of the phenotypes of diabetes-associated cardiovascular events is sudden cardiac death (SCD), and its mechanism is multifactorial, including acceleration of atherosclerosis, impairment of ventricular contractile function, and increased susceptibility to ventricular arrhythmias [2–5]. Pathological alterations in ventricular conduction and repolarization caused by diabetes are reflected by prolongation of the corrected QT (QTc) interval, increase in QT dispersion (QTd), and/or prolongation of the Tpeak–Tend interval (TpTe) in an electrocardiogram (ECG) [5–10]. Multiple factors are potentially involved in the abnormality of ventricular repolarization depending on the severity of diabetes and treatment [11–14]. However, the natural history of cardiac repolarization abnormality in patients with diabetes or other metabolic derangements is still unclear.

In the present study, we hypothesized that insulin resistance is associated with longitudinal increase in cardiac repolarization heterogeneity in an apparently healthy population. To examine this hypothesis, we determined relationships between indices of insulin resistance and longitudinal changes in indices of cardiac repolarization heterogeneity in apparently healthy subjects in two cohorts that we have prospectively followed.

## METHODS

### Compliance with Ethics Guidelines

This study was approved by the Ethics Committee of Sapporo Medical University and was

conducted in strict adherence with the principles of the Declaration of Helsinki. Written informed consent was obtained from all study subjects.

### Study Subjects

We retrieved data from two cohorts, the Tanno-Sobetsu cohort [15, 16] and Maruyama Clinic cohort [17], in both of which data of annual health checkups have been prospectively collected. Since storage periods of ECGs were 6 years and 5 years after the health checkups in the Tanno-Sobetsu cohort and Maruyama Clinic cohort, respectively, we selected those periods as follow-up periods in the present study.

Cohort 1 (Tanno-Sobetsu cohort): Subjects in this cohort were residents of Tanno and Sobetsu, rural towns in Japan. We retrieved data for 346 Sobetsu residents who received health checkups in both 2008 and 2014. To exclude effects of pre-existing morbidities on ventricular repolarization, we excluded subjects who met one or more of the following criteria: diabetes mellitus, regular use of medications for cardiovascular and/or metabolic diseases, hypo- or hyperkalemia (serum  $K^+$  < 3.5 or > 5.5 mEq/L), chronic kidney disease at stage 4 or 5, and electrocardiographic abnormalities that modify indices of ventricular repolarization (such as bundle branch block, WPW syndrome, and atrial fibrillation). Diabetes mellitus, which was defined as a history of diabetes mellitus, fasting plasma glucose (FPG) level  $\geq 126$  mg/dl (7 mmol/L), glycosylated hemoglobin (HbA1c) level  $\geq 6.5\%$ , and/or use of medication for diabetes mellitus, was diagnosed at baseline and at the end of the follow-up. In accordance with the exclusion criteria, 193 subjects were excluded, and data for 153 subjects in cohort 1 contributed to analyses.

Cohort 2 (Maruyama Clinic cohort): Subjects in this cohort were those who underwent medical health checkups at a clinic located in Sapporo, an urban area in Japan. We retrieved data for 1004 subjects who received medical checkups in both 2009 and 2014. The same exclusion criteria as those in cohort 1 were

applied in this cohort, and 266 subjects were excluded. Data for the remaining 738 subjects in this cohort were used for analyses.

### Demographic and Laboratory Examinations

Cohort 1: Medical checkups, including venous blood sampling, were conducted in the early morning after an overnight fast. Information regarding past history of diseases and medication was collected by public health nurses or clinic nurses on the basis of an interview form. Waist circumference (WC), body mass index (BMI), systolic blood pressure (SBP), and diastolic blood pressure (DBP) were measured by trained nurses. LDL-cholesterol (LDL-C) was calculated by the Friedewald equation. HbA1c was expressed on the National Glycohemoglobin Standardization Program (NGSP) scale. Serum insulin level was determined by an enzyme immunoassay, and the homeostasis model assessment of insulin resistance (HOMA-IR), an index of insulin sensitivity, was calculated by using following the equation:  $\text{HOMA-IR} = \text{fasting glucose (mg/dl)} \times \text{fasting insulin (mU/L)} / 405$  [18].

Cohort 2: Medical checkups including measurements of demographic parameters, blood pressure measurements, and analyses of venous blood samples were similar to those in cohort 1. Since serum insulin level was not determined in cohort 2, triglyceride (TG)/HDL-cholesterol (HDL-C) was calculated as an index of insulin resistance [19, 20].

### ECG Analysis

Standard 12-lead ECGs at rest were recorded using an electrocardiograph FCP-3610 (Fukuda Denshi Co., Ltd. Tokyo, Japan) in cohort 1 and by using CardioFaxV ECG-1550 (Nihon Kohden Co., Ltd. Tokyo, Japan) in cohort 2. Since both electrocardiographs had passed Japanese Industrial Standards, significant data variability by the electrocardiographs was unlikely. As indices of ventricular repolarization heterogeneity, three currently used indices were measured: QTc, QTd, and TpTe corrected by heart rate (cTpTe) [21]. The QT intervals were

measured in all 12 leads and corrected for heart rate by Bazett's formula. QTd was calculated as the difference between the longest and shortest QT intervals among the 12 leads. TpTe was measured in milliseconds (ms) from the peak of the T wave to the intersection between the tangent at the steepest point of the T wave downslope and the isoelectronic line at lead V5 [22]. Since correction of TpTe by heart rate using the Bazett or Fridericia formula has been shown to improve the predictive value of TpTe as a marker of sudden cardiac death [23], we calculated cTpTe by using Bazett's formula. ECG parameters in all study subjects were measured manually by a single investigator (T. Matsumoto), a board-certified cardiologist. Accuracy of her measurement of ECG indices had been confirmed by two co-authors using 10 randomly selected cases of ECG.

### Statistical Analysis

Numeric variables are expressed as mean  $\pm$  SD for data showing a normal distribution or medians and interquartile ranges for data showing a skewed distribution. Normality of data distribution was examined by the Shapiro–Wilk test. Differences in parameters between the two groups were examined by Student's *t* test, Mann–Whitney *U* test, or Chi-square test depending on the parameter and its distribution. The relationships between parameters were examined using simple and multiple linear regression analyses. Data with a skewed distribution were normalized by logarithmic transformation or by Box–Cox transformation [24] to make them usable for the regression analyses as in previous studies [25, 26]. Correlation of parameters was examined by use of Pearson's correlation coefficient. Multiple linear regression analyses were conducted by using the forced entry method to examine relationships between indices of insulin resistance at baseline and indices of ventricular repolarization at the end of follow-up. On the basis of current knowledge regarding insulin resistance and cardiac electrical remodeling [20, 27], we selected age, sex, BMI, and SBP as confounding factors of insulin resistance in the regression

analyses. For all tests,  $p < 0.05$  was considered statistically significant. All data were analyzed by using JMP11 for Windows (SAS Institute, Cary, NC, USA).

## RESULTS

### Cross-Sectional Analysis of Data in Cohort 1 (Tanno-Sobetsu Cohort)

Baseline characteristics of the study subjects (data in 2008) are shown in Table 1. The mean age of the subjects was 58.4 years and 46.4% of

them were men. BMI, WC, DBP, and percentage of subjects with a smoking habit were higher in men than in women. FPG, HOMA-IR, and serum TG were higher and HDL-C was lower in men than in women.

Mean QTc interval was shorter and SV1 + RV5 was larger in men than in women, whereas heart rate, QTd, TpTe, and cTpTe were comparable in men and women. HOMA-IR, cTpTe, and QTd were transformed into log HOMA-IR,  $1/\sqrt{cTpTe}$ , log QTd (2008), and  $\sqrt{QTd}$  (2014), respectively, for parametric analyses. SBP was correlated with  $1/\sqrt{cTpTe}$  ( $r = -0.162$ , 95% confidence interval [CI]  $-0.313$  to  $-0.004$ ,  $p = 0.045$ ), and DBP was

**Table 1** Baseline characteristics in cohort 1

	Total ( $n = 153$ )	Men ( $n = 71$ )	Women ( $n = 82$ )	$p$ value*
Age (years)	58.4 $\pm$ 12.5	56.6 $\pm$ 12.5	60.0 $\pm$ 12.4	0.102
BMI (kg/m <sup>2</sup> )	22.8 $\pm$ 3.4	24.0 $\pm$ 3.3	21.7 $\pm$ 3.2	< 0.001
WC (cm)	82.2 $\pm$ 10.5	84.9 $\pm$ 10.4	79.8 $\pm$ 10.0	0.002
SBP (mmHg)	130.9 $\pm$ 20.2	134.2 $\pm$ 20.1	128.0 $\pm$ 20.1	0.057
DBP (mmHg)	77.9 $\pm$ 11.3	82.1 $\pm$ 10.7	74.2 $\pm$ 10.7	< 0.001
FPG (mg/dl)	93.2 $\pm$ 8.4	95.4 $\pm$ 8.2	91.3 $\pm$ 8.1	0.002
HbA1c (%)	4.87 $\pm$ 0.30	4.88 $\pm$ 0.29	4.86 $\pm$ 0.30	0.574
HOMA-IR	1.04 (0.75, 1.51)	1.13 (0.84, 1.83)	0.97 (0.74, 1.42)	0.026
TG (mg/dl)	95 (65, 128)	110 (71, 156)	82.5 (62.8, 113)	0.008
HDL-C (mg/dl)	63.6 $\pm$ 15.4	59.1 $\pm$ 14.4	67.6 $\pm$ 15.1	0.001
LDL-C (mg/dl)	121.0 $\pm$ 28.3	122.7 $\pm$ 30.3	119.5 $\pm$ 26.5	0.490
Smoking ( $n$ , %)	29 (19.0)	23 (32.4)	6 (7.3)	< 0.001
HR (bpm)	62.5 $\pm$ 9.1	61.8 $\pm$ 8.8	63.2 $\pm$ 9.4	0.353
Mean QTc (ms)	403.1 $\pm$ 22.2	393.1 $\pm$ 22.2	411.7 $\pm$ 18.3	< 0.001
QTd (ms)	30 (20, 30)	30 (20, 30)	30 (20, 30)	0.945
TpTe (ms)	80 (70, 90)	80 (70, 80)	80 (70, 90)	0.640
cTpTe (ms)	79.6 (72.8, 87.2)	78.8 (73.0, 85.3)	79.8 (71.8, 87.6)	0.666
SV1 + RV5 (mV)	1.90 $\pm$ 0.54	2.07 $\pm$ 0.58	1.74 $\pm$ 0.45	< 0.001

Variables are expressed as number (%), mean  $\pm$  SD or medians (interquartile ranges)

BMI body mass index, WC waist circumference, SBP systolic blood pressure, DBP diastolic blood pressure, FPG fasting plasma glucose, HOMA-IR homeostasis model assessment of insulin resistance, TG triglycerides, HR heart rate, QTc corrected QT interval, QTd QT dispersion, TpTe Tpeak–Tend interval at V5 lead, cTpTe heart rate-corrected TpTe

\* $p$  values were calculated by Student's  $t$  test for normally distributed data, by Mann–Whitney  $U$  test for data in a skewed distribution, and by Chi-square test for categorical data for differences between men and women

**Table 2** Indices of ventricular heterogeneity at baseline: subjects with insulin resistance vs. subjects without insulin resistance in cohort 1

	HOMA-IR $\geq 1.73$ ( $n = 25$ )	HOMA-IR $< 1.73$ ( $n = 128$ )	<i>p</i> value*
Age (years)	60.8 $\pm$ 11.7	58.0 $\pm$ 12.7	0.306
Men (%)	18 (72.0)	53 (41.4)	0.005
HOMA-IR	2.43 (2.04, 2.81)	0.97 (0.67, 1.25)	$< 0.001$
TG/HDL-C ratio	1.55 (1.25, 2.94)	1.37 (0.94, 2.28)	0.094
Mean QTc (ms)	407.7 $\pm$ 26.6	402.2 $\pm$ 21.2	0.253
QTd (ms)	30 (20, 30)	30 (20, 30)	0.122
TpTe (ms)	80 (70, 90)	80 (70, 80)	0.326
cTpTe (ms)	80.4 (75.5, 90.8)	78.6 (71.5, 86.4)	0.112

Variables are expressed as mean  $\pm$  SD or medians (interquartile ranges)

\**p* values were calculated by Student's *t* test for normally distributed data, by Mann–Whitney *U* test for data in a skewed distribution, and by Chi-square test for categorical data

correlated with log QTd ( $r = -0.185$ , 95% CI  $-0.334$  to  $-0.027$ ,  $p = 0.022$ ) and  $1/\sqrt{cTpTe}$  ( $r = -0.199$ , 95% CI  $-0.346$  to  $-0.041$ ,  $p = 0.014$ ). There was a weak correlation between log HOMA-IR and  $1/\sqrt{cTpTe}$  ( $r = -0.159$ , 95% CI  $-0.319$  to  $-0.001$ ,  $p = 0.049$ ). However, neither FPG nor HbA1c was correlated with  $1/\sqrt{cTpTe}$ . Log QTd was not correlated with log HOMA-IR, FPG, HbA1c, or serum lipid levels (data not shown).

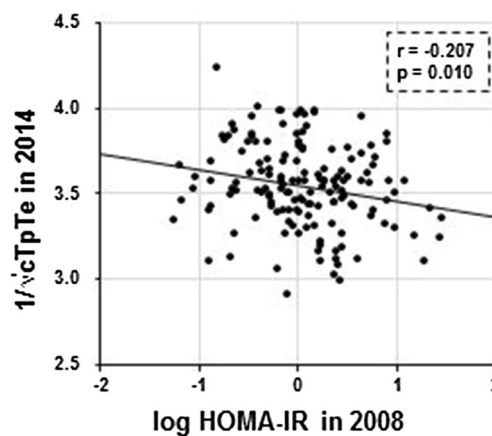
Insulin resistance defined by HOMA-IR  $\geq 1.73$  [28] was observed in 16.3% (25/153) of the study subjects at baseline, and their indices of ventricular repolarization heterogeneity are summarized in Table 2. Baseline cTpTe and QTd were not significantly different between subjects with HOMA-IR  $\geq 1.73$  and those with HOMA-IR  $< 1.73$ .

### Longitudinal Analysis of Data in Cohort 1

To determine factors that predict longitudinal changes in QTd and cTpTe, we examined correlations between clinical and laboratory variables in 2008 and  $\sqrt{QTd}$  and  $1/\sqrt{cTpTe}$  in 2014. There was a weak correlation between log HOMA-IR at baseline and  $1/\sqrt{cTpTe}$  in 2014 ( $r = -0.207$ , 95% CI  $-0.354$  to  $-0.050$ ,  $p = 0.010$ ) (Fig. 1), whereas such a correlation was not found between baseline FPG or HbA1c

and  $1/\sqrt{cTpTe}$  in 2014.  $\sqrt{QTd}$  in 2014 was not correlated with log HOMA-IR, HbA1c, or FPG at baseline (data not shown).

As shown in Table 3, multiple linear regression analysis for  $1/\sqrt{cTpTe}$  in 2014 revealed that log HOMA-IR at baseline was an independent explanatory variable of  $1/\sqrt{cTpTe}$  after adjustment for age and sex (model 1), and the relation was still significant after additional adjustment for SBP and BMI (model 2 and model 3). In



**Fig. 1** Correlation between HOMA-IR in 2008 and cTpTe in 2014: cohort 1. Because of the non-normalized distribution of data, HOMA-IR and cTpTe data were normalized to log HOMA-IR and  $1/\sqrt{cTpTe}$ , respectively. Regression line:  $y = 3.548 - 0.091x$ ,  $r = -0.207$  (95% CI  $-0.354$  to  $-0.050$ ),  $p = 0.010$

**Table 3** Multiple linear regression analysis for  $1/\sqrt{cTpTe}$  in 2014 in cohort 1

	Model 1				Model 2				Model 3			
	Estimate	SE	t	p value	Estimate	SE	t	p value	Estimate	SE	t	p value
08 log HOMA-IR	- 0.095	0.036	- 2.65	0.009	- 0.094	0.037	- 2.56	0.012	- 0.112	0.044	- 2.55	0.012
Age	- 0.001	0.002	- 0.89	0.377	- 0.001	0.002	- 0.75	0.457	- 0.001	0.002	- 0.75	0.454
Sex	- 0.020	0.020	- 0.98	0.326	- 0.020	0.025	- 0.99	0.326	- 0.016	0.021	- 0.75	0.453
08 SBP	-	-	-	-	- 0.001	0.001	- 0.12	0.906	- 0.001	0.001	- 0.19	0.847
08 BMI	-	-	-	-	-	-	-	-	0.006	0.007	0.75	0.454
08 HbA1c	0.022	0.072	0.30	0.765	0.030	0.073	0.41	0.680	0.036	0.074	0.48	0.630
Age	- 0.002	0.002	- 1.17	0.244	- 0.002	0.002	- 0.82	0.413	- 0.002	0.002	- 0.83	0.405
Sex	- 0.009	0.020	- 0.46	0.649	- 0.012	0.021	- 0.60	0.550	- 0.017	0.022	- 0.80	0.424
08 SBP	-	-	-	-	- 0.001	0.001	- 0.71	0.477	- 0.001	0.001	- 0.55	0.580
08 BMI	-	-	-	-	-	-	-	-	- 0.005	0.006	- 0.79	0.429
08 FPG	- 0.003	0.003	- 1.35	0.179	- 0.003	0.003	- 1.29	0.198	- 0.003	0.003	- 1.11	0.270
Age	- 0.001	0.002	- 0.87	0.386	- 0.001	0.002	- 0.56	0.576	- 0.001	0.002	- 0.57	0.568
Sex	- 0.018	0.021	- 0.84	0.402	- 0.020	0.021	- 0.93	0.354	- 0.022	0.022	- 0.99	0.326
08 SBP	-	-	-	-	- 0.001	0.001	- 0.54	0.593	- 0.001	0.001	- 0.46	0.647
08 BMI	-	-	-	-	-	-	-	-	- 0.002	0.007	- 0.37	0.715

SE standard error, 08 log HOMA-IR log homeostasis model assessment of insulin resistance in 2008, 08 HbA1c HbA1c in 2008, 08 FPG fasting plasma glucose in 2008, 08 SBP systolic blood pressure in 2008, 08 BMI body mass index in 2008, Model 1 adjusted for age and sex, Model 2 adjusted for age, sex, and SBP in 2008, Model 3 adjusted for age, sex, and SBP in 2008 and BMI in 2008



**Table 4** Baseline characteristics in cohort 2

	Total (n = 738)	Men (n = 421)	Women (n = 317)	p value*
Age (years)	46.0 ± 8.3	46.6 ± 8.2	45.2 ± 8.4	0.024
BMI (kg/m <sup>2</sup> )	22.5 ± 3.3	23.8 ± 3.0	20.9 ± 2.9	< 0.001
WC (cm)	81.8 ± 8.8	85.2 ± 7.9	77.3 ± 8.0	< 0.001
SBP (mmHg)	116.0 ± 14.2	119.8 ± 13.3	110.9 ± 13.8	< 0.001
DBP (mmHg)	74.1 ± 10.4	77.2 ± 9.9	70.0 ± 9.4	< 0.001
FPG (mg/dl)	86.8 ± 8.7	89.0 ± 8.8	83.9 ± 7.6	< 0.001
HbA1c (%)	4.80 ± 0.29	4.81 ± 0.30	4.79 ± 0.28	0.269
TG (mg/dl)	84 (60, 121.3)	104 (77, 156.5)	62 (46.5, 84.5)	< 0.001
HDL-C (mg/dl)	63.6 ± 16.8	56.9 ± 14.4	72.4 ± 15.8	< 0.001
LDL-C (mg/dl)	124.6 ± 31.2	126.9 ± 30.5	121.3 ± 31.9	0.021
TG/HDL-C ratio	1.32 (0.87, 2.30)	1.92 (1.26, 3.19)	0.90 (0.61, 1.25)	< 0.001
Smoking (n, %)	217 (29.4)	174 (41.3)	43 (13.6)	< 0.001
HR (bpm)	62.0 ± 8.5	61.2 ± 8.1	63.1 ± 8.8	0.003
Mean QTc (ms)	385.5 ± 21.0	378.2 ± 18.8	395.1 ± 19.9	< 0.001
QTd (ms)	30 (20, 30)	30 (20, 30)	30 (20, 40)	0.050
TpTe (ms)	80 (70, 80)	80 (70, 80)	80 (70, 90)	0.813
cTpTe (ms)	77.7 (70.9, 85.8)	77.7 (71.0, 85.3)	78.5 (70.7, 86.9)	0.271
SV1 + RV5 (mV)	2.55 ± 0.72	2.78 ± 0.75	2.24 ± 0.55	< 0.001

Variables are expressed as number (%), mean ± SD or medians (interquartile ranges)

*BMI* body mass index, *WC* waist circumference, *SBP* systolic blood pressure, *DBP* diastolic blood pressure, *FPG* fasting plasma glucose, *TG* triglycerides, *HR* heart rate, *QTc* corrected QT interval, *QTd* QT dispersion, *TpTe* Tpeak–Tend interval at V5 lead, *cTpTe* heart-rate corrected TpTe

\**p* values were calculated by Student's *t* test for normally distributed data, by Mann–Whitney *U* test for data in a skewed distribution, and by Chi-square test for categorical data for differences between men and women

contrast, FPG and HbA1c at baseline were not significant explanatory variables of  $1/\sqrt{cTpTe}$  in 2014 in any of the models.

### Cross-Sectional Analysis of Data in Cohort 2 (Maruyama Clinic Cohort)

As shown in Table 4, the mean age of the subjects was 46.0 years and 57.0% were men. Mean age, BMI, WC, SBP, DBP, FPG, TG, LDL-C, TG/HDL-C ratio, and percentage of subjects with a smoking habit were higher in men than in women, and HDL-C was lower in men.

In electrocardiographic variables, HR and QTd at baseline were slightly but significantly lower in men than in women in this cohort. Mean QTc interval was shorter and SV1 + RV5 was larger in men than in women and there was no statistical difference in TpTe and cTpTe between men and women at baseline, being consistent with data in cohort 1.

In cohort 2, TG/HDL-C, cTpTe, and QTd were transformed into log TG/HDL-C,  $1/cTpTe$ , and  $\sqrt{QTd}$ , respectively, for parametric analyses. SBP ( $r = -0.134$ , 95% CI  $-0.204$  to  $-0.062$ ,  $p = 0.0003$ ), DBP ( $r = -0.139$ , 95% CI  $-$

**Table 5** Indices of ventricular heterogeneity at baseline: subjects with insulin resistance vs. subjects without insulin resistance in cohort 2

	TG/HDL-C $\geq 3$ ( <i>n</i> = 121)	TG/HDL-C $< 3$ ( <i>n</i> = 617)	<i>p</i> value*
Age (years)	46.0 $\pm$ 7.9	46.0 $\pm$ 8.4	0.954
Men (%)	113 (93.4)	308 (49.9)	< 0.001
TG/HDL-C ratio	4.27 (3.63, 5.75)	1.15 (0.75, 1.70)	< 0.001
Mean QTc (ms)	383.1 $\pm$ 18.0	385.9 $\pm$ 21.5	0.184
QTd (ms)	30 (20, 30)	30 (20, 30)	0.996
TpTe (ms)	80 (70, 90)	80 (70, 80)	0.784
cTpTe (ms)	79.6 (71.4, 86.4)	77.3 (70.7, 85.4)	0.222

Variables are expressed as mean  $\pm$  SD or medians (interquartile ranges)

\**p* values were calculated by Student's *t* test for normally distributed data, by Mann–Whitney *U* test for data in a skewed distribution, and by Chi-square test for categorical data

0.209 to  $-0.068$ ,  $p = 0.0001$ ), log TG ( $r = -0.087$ , 95% CI  $-0.158$  to  $-0.015$ ,  $p = 0.019$ ), and log TG/HDL ( $r = -0.083$ , 95% CI  $-0.154$  to  $-0.011$ ,  $p = 0.025$ ), but not FPG or HbA1c, were weakly correlated with 1/cTpTe at baseline.  $\sqrt{\text{QTd}}$  was not correlated with any of the clinical or metabolic parameters.

Insulin resistance defined by TG/HDL-C  $\geq 3$  [19] was observed in 16.4% of the study subjects at baseline, and their indices of ventricular repolarization heterogeneity are summarized in Table 5. Baseline cTpTe and QTd were not significantly different between subjects with TG/HDL-C  $\geq 3$  and those with TG/HDL  $< 3$ .

## Longitudinal Analysis of Data in Cohort 2

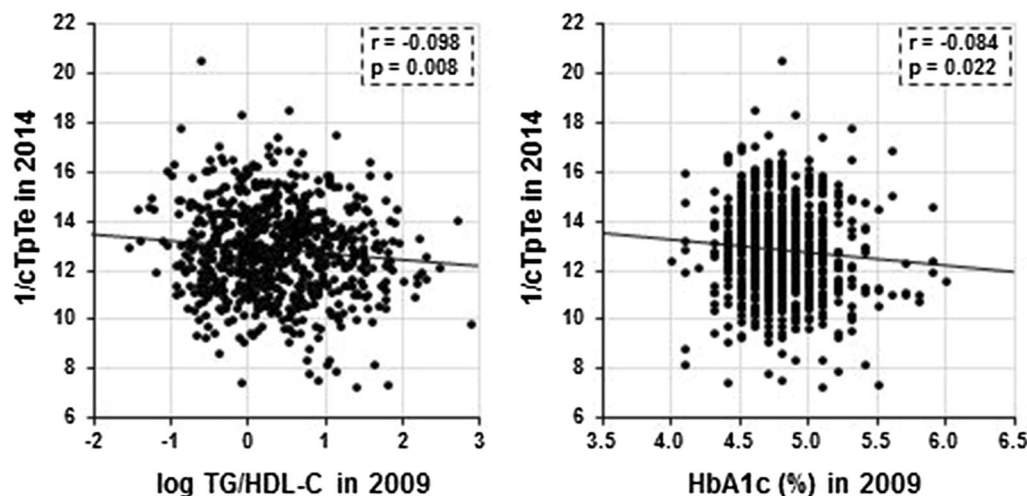
As shown in Fig. 2, 1/cTpTe in 2014 was very weakly correlated with log TG/HDL-C at baseline ( $r = -0.098$ , 95% CI  $-0.169$  to  $-0.026$ ,  $p = 0.008$ ) and with HbA1c at baseline ( $r = -0.084$ , 95% CI  $-0.156$  to  $-0.012$ ,  $p = 0.022$ ), whereas such a correlation was not found between baseline FPG and 1/cTpTe in 2014.  $\sqrt{\text{QTd}}$  in 2014 was not correlated with log TG/HDL-C, HbA1c, or FPG at baseline (data not shown).

In multiple linear regression analysis for 1/cTpTe in 2014 (Table 6), log TG/HDL-C at baseline was found to be an independent explanatory variable after adjustment for age and sex (model 1). The association of log TG/HDL-C at baseline with 1/cTpTe in 2014 was significant after additional adjustment for SBP and BMI (Table 6, model 2 and model 3). In contrast, HbA1c at baseline was not selected as an independent explanatory variable of 1/cTpTe in 2014 after adjustment for age and sex. There was no significant relation between  $\sqrt{\text{QTd}}$  in 2014 and TG/HDL-C ratio, HbA1c, or FPG at baseline (data not shown).

## DISCUSSION

In addition to weak correlations between indices of insulin resistance and cTpTe at baseline in both cohort 1 and cohort 2 in the cross-sectional analyses, a significant correlation was found between an index of insulin resistance at baseline and cTpTe in 2014. Furthermore, results of multiple linear regression analyses of data in the two cohorts (Tables 3 and 6) consistently indicated that insulin resistance assessed by HOMA-IR or TG/HDL-C was independently associated with increase in cTpTe after 5–6 years of follow-up. Correlation coefficients in the longitudinal relationship were small in both cohorts (Figs. 1 and 2), but it is not unexpected since we excluded subjects on regular medications, diabetic patients, and patients with chronic kidney disease from the study subjects for the analysis. Ranges of data for indices of insulin resistance and cTpTe in the study subjects were not largely deviated





**Fig. 2** Relationships between TG/HDL-C in 2009, HbA1c in 2009, and cTpTe in 2014: cohort 2. Because of the non-normalized distribution of data, TG/HDL-C and cTpTe data were normalized to log TG/HDL-C and to 1/cTpTe, respectively. **a**  $y = 12.937 - 0.246 x$ ,  $r = -$

$0.098$  (95% CI  $- 0.169$  to  $- 0.026$ ),  $p = 0.008$ ; **b**  $y = 15.403 - 0.532 x$ ,  $r = - 0.084$  (95% CI  $- 0.156$  to  $- 0.012$ ),  $p = 0.022$

from “normal” ranges both at baseline and during the follow-up, which may have made it difficult to detect an association of pathological changes in the two factors. Nevertheless, earlier studies have shown that abnormality of ventricular repolarization in obese subjects and patients with metabolic syndrome is significantly improved by long-term exercise and weight loss together with improvement of insulin sensitivity [29, 30]. Together with the earlier findings, the results of the present study suggest that insulin resistance plays a role in the increase in cardiac repolarization heterogeneity in apparently healthy subjects.

To use univariable and multivariable regression analyses, we normalized data for QTd and cTpTe by Box–Cox transformation and we normalized data for HOMA-IR and TG/HDL-C by logarithmic transformation as in earlier studies [25, 26]. Because of such transformations, the extent of change in cTpTe by the level of HOMA-IR or TG/HDL-C is not easily observable in Figs. 1 and 2. However, calculation using regression equations indicated that the median and 90th percentile values of HOMA-IR, 1.04 and 2.14, at baseline predict cTpTe of 78.2 and 82.6 ms, respectively, 6 years later in cohort 1. Similarly, the median and 90th percentile

values of TG/HDL-C, 1.32 and 3.93, predict cTpTe of 77.7 and 79.4 ms, respectively, 5 years later in cohort 2. cTpTe data for the subjects with 90th percentile HOMA-IR and TG/HDL-C are actually close to the average cTpTe in diabetic patients in earlier studies (i.e., 76.5–86.0 ms) [9, 10, 14, 31]. The modest impact of change in an index of insulin resistance on cTpTe prolongation (Figs. 1 and 2), as compared with the impact of diabetes, rather supports the notion that insulin resistance is the earliest causal factor of multiple factors involved in abnormalities of cardiac repolarization in diabetic patients.

In contrast to TpTe, QTd was not correlated with indices of insulin resistance in the two cohorts. QTd indicates the difference in action potential durations between two regions, and TpTe is an index of global dispersion of repolarization [20, 32]. Both QTd and TpTe have been shown to increase in patients with diabetes mellitus [6–10]. Obesity and metabolic syndrome are also reportedly associated with an increase in QTd [29, 30], though an increase in TpTe caused by obesity or metabolic syndrome, to our knowledge, has not been reported. Taken together, the findings suggest that dispersion of repolarization by insulin resistance precedes an

**Table 6** Multiple linear regression analysis for 1/cTnT<sub>Te</sub> in 2014 in cohort 2

	Model 1			Model 2			Model 3					
	Estimate	SE	t	p value	Estimate	SE	t	p value	Estimate	SE	t	p value
09 log TG/HDL-C	- 0.370	0.109	- 3.39	< 0.001	- 0.308	0.109	- 2.82	0.005	- 0.248	0.116	- 2.13	0.033
Age	- 0.016	0.008	- 2.01	0.045	- 0.009	0.008	- 1.14	0.255	- 0.010	0.008	- 1.17	0.242
Sex	- 0.184	0.080	- 2.30	0.022	- 0.240	0.080	- 2.98	0.003	- 0.262	0.082	- 3.20	0.001
09 SBP	-	-	-	-	- 0.019	0.005	- 3.82	<0.001	- 0.018	0.005	- 3.40	< 0.001
09 BMI	-	-	-	-	-	-	-	-	- 0.037	0.025	- 1.50	0.133
09 HbA1c	- 0.439	0.240	- 1.83	0.068	- 0.354	0.239	- 1.48	0.138	- 0.229	0.246	- 0.93	0.352
Age	- 0.013	0.008	- 1.55	0.122	- 0.006	0.008	- 0.73	0.464	- 0.008	0.008	- 0.90	0.368
Sex	- 0.040	0.068	- 0.58	0.560	- 0.126	0.070	- 1.79	0.073	- 0.185	0.076	- 2.45	0.015
09 SBP	-	-	-	-	- 0.021	0.005	- 4.13	< 0.001	- 0.018	0.005	- 3.5	<0.001
09 BMI	-	-	-	-	-	-	-	-	- 0.050	0.024	- 2.07	0.039
09 FPG	- 0.011	0.008	- 1.36	0.173	- 0.004	0.008	- 0.51	0.607	0.001	0.009	0.01	0.994
Age	- 0.014	0.008	- 1.71	0.088	- 0.008	0.008	- 1	0.317	- 0.010	0.008	- 1.16	0.264
Sex	- 0.061	0.071	- 0.86	0.390	- 0.132	0.072	- 1.84	0.066	- 0.189	0.076	- 2.50	0.013
09 SBP	-	-	-	-	- 0.021	0.005	- 4.16	< 0.001	- 0.019	0.005	- 3.57	<0.001
09 BMI	-	-	-	-	-	-	-	-	- 0.055	0.024	- 2.30	0.022

SE standard error, 09 log TG/HDL-C log triglycerides/HDL-C in 2009, 09 HbA1c HbA1c in 2009, 09 FPG fasting plasma glucose in 2009, 09 SBP systolic blood pressure in 2009, 09 BMI body mass index in 2009, Model 1 adjusted for age and sex, Model 2 adjusted for age, sex, and SBP in 2009, Model 3 adjusted for age, sex, and SBP in 2009 and BMI in 2009

increase in heterogeneity of action potential duration (APD) by hyperglycemia.

During the 6-year follow-up in cohort 1 and the 5-year follow-up in cohort 2, 3.3% ( $n = 5$ ) and 2.3% ( $n = 17$ ) of the subjects, respectively, newly developed diabetes mellitus. Since the number of patients with new-onset diabetes in each cohort was small, it was difficult to statistically determine the relationship between new-onset diabetes and change in QTd or cTpTe.

The mechanism by which insulin resistance induces abnormality of ventricular repolarization has been studied by animal experiments. Phosphoinositide 3-kinase (PI3K), which is coupled with the insulin receptor, has been shown to regulate multiple ion channels involved in repolarization of cardiomyocytes [33]. Knockout of the insulin receptors in cardiomyocytes suppressed protein expression of Kv4.2 and KChIP2, which reduced transient outward  $K^+$  current (Ito current), leading to prolongation of APD and QT on a surface ECG [34]. Similar reduction of Kv4.2 and KChIP2 and APD prolongation were observed in cardiomyocytes of a rat model of type 2 diabetes with hyperinsulinemia and insulin resistance [13]. However, since Ito current is smaller in human cardiomyocytes than in rodent cardiomyocytes, the findings in rat hearts may not be extrapolated to patients with insulin resistance. Increase in sympathetic tone might also be involved in increased heterogeneity of ventricular repolarization in association with insulin resistance [35, 36]. In addition, in diabetic patients, diabetes-related comorbidities such as coronary artery disease and cardiac hypertrophy are possibly involved in the abnormality of ventricular repolarization depending on the severity and duration of diabetes.

### Study Limitations

There are limitations in the present study. First, since different indices of insulin resistance and different equations for Box–Cox transformation were used in the two cohorts, we could not directly compare the changes in cTpTe caused by insulin resistance in cohort 1 and cohort 2. Additionally, we cannot exclude the possibility

that HDL-C [37] and/or LDL-C particle size, which is also reflected by TG/HDL-C [38], were responsible for the change in cTpTe during follow-up. Second, since follow-up periods were relatively short, the impact of a slight change in insulin sensitivity on a long natural history of ventricular repolarization remains unclear. Third, because a single investigator measured ECG parameters, ECG measurements might have been biased, though precision of the data appears high because the data are free from interobserver differences.

## CONCLUSIONS

Longitudinal analyses of two independent cohorts showed that indices of insulin resistance, HOMA-IR and TG/HDL-C, were independent explanatory variables for increase in cTpTe, an index of heterogeneity of ventricular repolarization, during 5- to 6-year follow-up in apparently healthy individuals. The findings support the notion that insulin resistance is the earliest causal factor in abnormal ventricular repolarization in patients with diabetes.

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**Data Availability.** The data sets generated and/or analyzed during the current study are available from the corresponding author on reasonable request.

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