ARTICLE

Association of glycaemia with lipids in adults with type 1 diabetes: modification by dyslipidaemia medication

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Abstract

Aims/hypothesis Hyperglycaemia and dyslipidaemia are common metabolic abnormalities in adults with type 1 diabetes and both increase cardiovascular disease (CVD) risk. The hypothesis of this study was that change in HbA_{1c} over 6 years would be associated with change in fasting lipids in adults with type 1 diabetes.

Methods The Coronary Artery Calcification in Type 1 Diabetes (CACTI) study examined 652 patients with type 1 diabetes (54% female); 559 and 543 had follow-up visits at 3 and 6 years. Baseline age (mean \pm SD) was $37\pm$ 9 years, diabetes duration $23\pm$ 9 years, and HbA_{1c} $8.0\pm$ 1.3%. Use of dyslipidaemia medication was 17%, 32%, and 46% at the three visits. Separate longitudinal mixed models were fitted to examine the relationship between change in HbA_{1c} and change in fasting total cholesterol (TC),

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Colorado School of Public Health, Department of Biostatistics and Informatics, University of Colorado Denver, Aurora, CO, USA HDL-cholesterol (HDL-c), LDL-cholesterol (LDL-c), log triacylglycerols (TG), and non-HDL-cholesterol (non-HDL-c). Because of an interaction between dyslipidaemia medication use and association of HbA_{1c} with lipids, results were stratified by dyslipidaemia medication use.

Results Among patients not using dyslipidaemia medication, a higher HbA_{1c} was associated with significantly worse levels of the lipids TC, LDL-c, TG and non-HDL-c (per 1% change in HbA1c, TC 0.101 mmol/l, 95% CI 0.050, 0.152; LDL-c 0.103 mmol/l, 95% CI 0.058, 0.148; TG 0.052 mmol/l, 95% CI 0.024, 0.081; and non-HDL-c 0.129 mmol/l, 95% CI 0.078, 0.180) but not HDL-c (-0.20 mmol/l, 95% CI -0.047, 0.007). The associations between HbA_{1c} and any lipid outcome among those on dyslipidaemia medication were in the same direction, but attenuated compared with persons not on medication.

Conclusions/interpretation Change in HbA_{1c} is significantly associated with change in fasting lipids, but dyslipidaemia medications may be required to optimise lipid and cardiovascular health.

Keywords CVD risk · Dyslipidaemia medications · Glycaemia · Lipids · Type 1 diabetes

Abbreviations

CACTI	Coronary Artery Calcification in Type 1 Diabetes
CAD	Coronary artery disease
CVD	Cardiovascular disease
DCCT	Diabetes Control and Complications Trial
EDC	Epidemiology of Diabetes Complications
EDIC	Epidemiology of Diabetes Control
HDL-c	HDL-cholesterol
LDL-c	LDL-cholesterol
TC	Total cholesterol
TG	Triacylglycerols

Introduction

Hyperglycaemia, as measured by HbA_{1c}, is the most important determinant of vascular complications of type 1 diabetes [1-4]. In addition to hyperglycaemia, dyslipidaemia is a well-established cardiovascular disease (CVD) risk factor [5, 6]. However, few data exist in people with type 1 diabetes on the relationship of glycaemia to other CVD risk factors such as dyslipidaemia, although aggressive control of glycaemia is recommended as first line treatment of dyslipidaemia in the care of people with type 1 diabetes, with addition of lipid-lowering medications if lipid goals are not achieved [7, 8]. Although hyperglycaemia and dyslipidaemia [9] both increase CVD risk and are metabolic abnormalities frequently found in individuals with type 1 diabetes, it is unclear how changes in HbA1c are associated with alterations in lipids and whether dyslipidaemia medications modify this association.

The Diabetes Control and Complications Trial (DCCT) demonstrated that intensive (vs conventional) insulin therapy was associated with significantly reduced total cholesterol (TC), LDL-cholesterol (LDL-c) and triacylglycerols (TG) (p < p0.01 for all) and decreased development of LDL-c >4.14 mmol/l in adults with type 1 diabetes, which was considered a treatment threshold at the time of the study [10], as well as a more favourable distribution of lipoprotein cholesterol [11]. Similarly, the DCCT/Epidemiology of Diabetes Control (EDIC) study demonstrated that intensive vs conventional insulin therapy was associated with a 24% reduced risk of developing incident hypertension over a median of 15.8 years of follow-up [12]. Thus, it is plausible that some of the deleterious effect of hyperglycaemia on cardiovascular health is mediated via the effects of hyperglycaemia on hypertension and dyslipidaemia [10, 12].

Therefore, to better describe the relationship of glycaemia to dyslipidaemia outside the setting of a clinical trial, we tested the hypothesis that change in HbA_{1c} over a follow-up interval of approximately 6 years resulted in a concordant change in the lipid profile in adults with type 1 diabetes participating in the Coronary Artery Calcification in Type 1 Diabetes (CACTI) study, independent of potential confounders, and accounting for dyslipidaemia medication usage.

Methods

Study participants The data presented in this report were collected as part of the baseline examination of 1,416 participants in the CACTI study who were 19–56 years of age and included 652 participants with type 1 diabetes [13]. All participants were asymptomatic for coronary artery disease (CAD) and had no history of coronary artery bypass graft, coronary angioplasty, or unstable angina. Participants

with diabetes generally had been diagnosed when <30 years of age, or had positive antibodies or a clinical course consistent with type 1 diabetes. Of the 652 persons enrolled at baseline, 559 (86%) participated in a second study visit ~3 years after baseline and 543 (83%) in a third study visit ~6 years after baseline, with 603 (92%) of participants having at least one follow-up study visit. Participants who did not have any follow-up data (n=49) had higher TC, LDL-c and non-HDL-cholesterol (non-HDL-c) at baseline than those with follow-up data. All participants provided informed consent and the study was approved by the Colorado Multiple Institutional Review Board.

Examination and laboratory measurements Participants completed a baseline examination between March 2000 and April 2002. A more detailed description of the study and baseline characteristics of this cohort has been published [14]. The second and third study visits followed the same protocol and occurred between 2003 and 2005 and between 2006 and 2008, respectively.

Laboratory analyses Blood was collected after an overnight fast and serum was stored at -70° C until assayed. Fasting plasma samples were obtained for measurement of TC, HDL-c and TG, in the University of Colorado Denver Clinical Translational Research Center core lab. Measurements of TC, HDL-c, and TG were performed enzymatically on a Hitachi 917 autoanalyser (Boehringer Mannheim Diagnostics, Indianapolis, IN, USA). LDL-c levels were calculated by the Friedewald equation for individuals with TG levels lower than 4.52 mmol/l [15] and by Lipid Research Clinics Beta Quantification [16] for those with TG≥4.52 mmol/l. Non-HDL-c was calculated as TC minus HDL-c. HbA_{1c} levels were measured by (DCCT-calibrated) ion-exchange high-performance liquid chromatography (Bio-Rad Laboratories, Hercules, CA, USA).

Variables Weight and height were measured using standard methods and BMI, defined as weight (kg) divided by height (m²), was calculated. Minimal waist circumference (measured at the smallest point between the tenth rib and the iliac crest over the bare skin) was measured. Intraabdominal fat and subcutaneous fat were assessed using abdominal computed tomography scan at the L4–L5 levels and volume in cubic centimeters was measured using the AccuAnalyzer software from AccuImage.

Dyslipidaemia medications Participants brought all current medications to each study visit for documentation by study staff. Dyslipidaemia treatment was defined as current use of lipid-lowering medication, such as hydroxymethylglutaryl-CoA reductase inhibitors, bile acid sequestrants, fibric acid derivatives, ezetimibe or niacin, and was recorded at each

study visit. Dyslipidaemia medication treatment decisions were made by the participants' physicians, not by the CACTI study team.

Statistical methods The mean and SE for each variable of interest at baseline and follow-up were estimated using the SAS Mixed procedure for linear mixed models. TG was log transformed due to non-normal distribution. Therefore, back transformed estimates were calculated as change in TG per 1% change in HbA_{1c} from a starting level of 0.93 mmol/l (median baseline TG level for cohort.) To determine the association of change in HbA1c with change in lipids, two a priori models (basic and saturated) were constructed with separate models for change in each lipid variable (TC, HDL-c, TG, LDL-c, and non-HDL-c) as an outcome. Time-varying change variables were created for HbA_{1c} and each lipid outcome for change from baseline to the first follow-up visit and from baseline to the second follow-up visit. The outcome of interest in all models was change in lipid levels over time. Baseline lipid levels were also included as a covariate to increase precision. In the basic model, demographic variables (sex, race/ethnicity, diabetes duration, and age as a time-varying variable) were adjusted for in addition to change in HbA_{1c}, visit, and a change in HbA_{1c} by visit interaction term to determine whether the association of change in HbA_{1c} with change in lipid outcome differed by visit. To account for differences in baseline levels of HbA1c, baseline HbA1c was also included. A baseline HbA_{1c} by time-varying change in HbA1c interaction term was tested for to determine whether the association of change in HbA1c with change in lipid was different according to baseline HbA1c, but this was not significant and was thus not included in the final models. Since the change in HbA_{1c} by visit interaction terms were not significant, ESTIMATE statements were used to determine the average association of change in HbA1c with change in each lipid outcome over time. In the saturated model, to account for the confounding role of adiposity on lipids, we additionally adjusted for BMI, waist circumference and visceral fat as time-varying variables. All three of these measures were included to fully capture the confounding effect of adiposity on lipids, as the individual role of each adiposity measure is not the focus of this analysis. Dyslipidaemia medication usage was considered as an effect modifier for the association of HbA1c with lipid levels. A time-varying dyslipidaemia medication usage (yes/no) by change in HbA_{1c} interaction term was included to determine whether the association of change in HbA_{1c} with change in the lipid outcome differed according to whether or not a participant was on a dyslipidaemia medication. As significant differences existed, models stratified by dyslipidaemia medication usage status are presented. In post hoc analyses, interactions for both timevarying age and sex by change in HbA_{1c} were explored and, since significant differences existed by sex, models stratified by sex are also presented. The Mixed procedure in SAS 9.2 (SAS Institute, Cary, NC, USA) was used for the analysis. A p value of <0.05 was considered statistically significant.

Results

Statistically significant improvements over 6 years (Table 1, p < 0.0001 for all) were noted for TC (mean decrease of 0.26 mmol/l), HDL-c (mean increase of 0.13 mmol/l), LDL-c (mean decrease of 0.33 mmol/l), TG (mean decrease of 0.14 mmol/l) and non-HDL-c (mean decrease of 0.39 mmol/l). HbA_{1c} also improved over time (mean decrease of 0.24% from baseline to first follow-up and mean decrease of 0.08% from baseline to second follow-up). Reported use of dyslipidaemia medications increased from 17% at baseline to 32% at the first follow-up and to 46% at the second follow-up visit. The improvements in lipid measures were noted despite increases in adiposity measures over the 6 year period.

The general pattern for reported dyslipidaemia medication use was that once participants started medications they continued on medications. Between the baseline and first follow-up visit, 85 additional participants were started on dyslipidaemia medications while 13 participants reported discontinuing dyslipidaemia medications. From the baseline to second follow-up visit 79 participants began and 13 discontinued dyslipidaemia medications. Of those on dyslipidaemia medications 97%, 95% and 93% were on statins at each visit.

In non-stratified models, there was a significantly different association of change in HbA_{1c} with change in lipid outcomes (TC, LDL-c, and non-HDL-c, p<0.0001 for each HbA1c by dyslipidaemia medication use interaction term) depending on whether a participant reported use of dyslipidaemia medication. We therefore report the results of stratified mixed model analysis of the association of change in HbA_{1c} to change in each lipid outcome (β -estimate \pm SE and p value) in the basic and saturated models as shown in Table 2. Among those not on dyslipidaemia medications, statistically significant associations existed for the association between change in HbA1c and change in TC, LDL-c, TG, and non-HDL-c, but not HDL-c. In the basic model, for every 1% change in HbA_{1c} the associated changes in lipids over 6 years were: TC +0.101 mmol/l (95% CI 0.050, 0.152, p=0.0001), HDL-c -0.20 mmol/l (95% CI -0.047, 0.007, p=0.15), LDL-c +0.103 mmol/l (95% CI 0.058, 0.148, p<0.0001), TG +0.052 mmol/l (95% CI 0.024, 0.081, p=0.0003), and non-HDL-c +0.129 mmol/l (95% CI 0.078, 0.180, p < 0.0001). Similar associations existed in the

Table 1 Characteristics of CACTI Type	1	Diabetes	Cohort
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Variable	Visit 0, <i>n</i> =652	Visit 1, <i>n</i> =559	Visit 2, <i>n</i> =543	p value
Age (years)	36.5±0.4	39.0±0.4	43.2±0.4	NA
Type 1 diabetes duration (years)	23.3 ± 0.3	25.8±0.3	29.6±0.3	NA
Time to follow-up (years)	NA	2.5 ± 0.4	$6.2 {\pm} 0.6$	NA
Sex (male/female), %	46%/54%	46%/54%	46%/54%	NA
Race/ethnicity (NHW), %	94%	94%	94%	NA
HbA _{1c} (%)	$7.98 {\pm} 0.05$	$7.74 {\pm} 0.05$	$7.90 {\pm} 0.05$	< 0.0001
TC (mmol/l)	$4.53 {\pm} 0.03$	$4.60 {\pm} 0.04$	$4.27 {\pm} 0.04$	< 0.0001
HDL (mmol/l)	$1.45 {\pm} 0.02$	$1.58 {\pm} 0.02$	$1.58 {\pm} 0.02$	< 0.0001
LDL (mmol/l)	2.61 ± 0.03	2.61 ± 0.03	$2.28 {\pm} 0.03$	< 0.0001
TG ^a (mmol/l)	0.93 (0.28-7.54)	0.75 (0.30-6.58)	0.79 (0.28-3.92)	< 0.0001
Non-HDL (mg/l)	$3.08 {\pm} 0.03$	$3.00 {\pm} 0.03$	2.69 ± 0.04	< 0.0001
Dyslipidaemia medications, yes/total, %	113/652 (17%)	180/559 (32%)	250/543 (46%)	< 0.0001
BMI (kg/m ²)	26.2 ± 0.2	26.5 ± 0.2	26.9 ± 0.2	< 0.0001
Waist circumference (cm)	85.1±0.5	85.2±0.5	88.7±0.5	< 0.0001
VSF (cm ³)	10.3 ± 0.03	10.5 ± 0.03	10.6 ± 0.02	< 0.0001

Results are presented as means±SE or count and frequency, except for 'Time to follow-up' (mean±SD)

p value, type 3 tests of fixed effects by visit in SAS Proc Mixed

n=603 with a baseline and either first or second visit

^a Geometric mean and range

NA, not applicable; NHW, non-Hispanic white; TC, total cholesterol; TG, triacylglycerols; VSF, visceral fat

saturated model (Table 2). In contrast, among those on dyslipidaemia medications, the association of change in HbA_{1c} with change in lipid level was reduced by approximately one-half and was no longer statistically significant, though the direction of the effect was similar to persons not on medication. Figure 1 displays the association of change in HbA_{1c} and change in lipid.

In participants not on dyslipidaemia medications, the following variables were also significantly associated with lipid outcomes in the basic model: male sex (HDL-c, LDL-c, non-HDL-c), older age (TC, HDL-c, LDL-c and non-HDL-c) and in the saturated models larger waist (HDL-c) and higher visceral fat (HDL-c, TG and non-HDL-c). Among participants on dyslipidaemia medications the following variables were significantly associated with lipid outcomes in the basic model: race/ethnicity other than non-Hispanic white (TG), higher baseline HbA_{1c} (all) and higher baseline lipid levels (all); and in the saturated models: male sex (HDL-c), older age (TC), higher baseline HbA_{1c} (all) and higher baseline lipid levels (all).

In post hoc analyses, significant differences existed in the association of HbA_{1c} with lipids, and models stratified by sex are presented in Tables 3 and 4. In men, there was a stronger association of HbA_{1c} with TC, LDL-c and non-HDL-c. HbA_{1c} was not associated with HDL-c in either sex. In men, HbA_{1c} was associated with TG in those not on dyslipidaemia medications in the basic but not the saturated model, and was

of borderline significance in those on dyslipidaemia medications, whereas in women HbA_{1c} was significantly associated with TG in those not on dyslipidaemia medications, but not in those on dyslipidaemia medications.

Discussion

We found that over 6 years of follow-up an increase in glycaemia was positively associated with change in TC, LDL-c, TG and non-HDL-c in adults with type 1 diabetes who were not on dyslipidaemia medications. This strong association was not observed in patients who were on dyslipidaemia medications. Although glycaemic control is important for reduction of vascular complications, we believe these data suggest that the effect of dyslipidaemia treatment on lipids is stronger than the effect of glycaemia on lipids and, therefore, may obscure the effect of glucose control on lipids. Therefore, this suggests that both glucose control and treatment with dyslipidaemia medications (if lipids exceed treatment goals) in adults with type 1 diabetes should be emphasised to achieve lipid and cardiovascular health. Furthermore, given the magnitude of the association of change in HbA_{1c} with change in lipids, for some patients with significantly elevated lipids it is likely that intensification of glycaemic control may be insufficient to achieve lipid treatment goals [7, 17].

	Basic model β (95% CI)	p value	Saturated model β (95% CI)	<i>p</i> value
NO lipid medications				
TC	0.101 (0.050,0.152)	0.0001	0.127 (0.072,0.182)	< 0.0001
HDL-c	-0.020 (-0.047,0.007)	0.15	-0.008 (-0.035,0.019)	0.57
LDL-c	0.103 (0.058,0.148)	< 0.0001	0.122 0.071,0.173)	< 0.0001
TG^{a}	0.052 (0.024,0.081)	0.0003	0.043 (0.014,0.073)	0.0004
Non-HDL-c	0.129 (0.078,0.180)	< 0.0001	0.140 (0.085,0.195)	< 0.0001
YES lipid medication	S			
TC	0.052 (-0.034,0.138)	0.23	0.036 (-0.055,0.127)	0.44
HDL-c	-0.018 (-0.053,0.017)	0.32	-0.022 (-0.059,0.015)	0.25
LDL-c	0.062 (-0.014,0.138)	0.12	0.054 (-0.26,0.134)	0.20
TG^{a}	0.025 (-0.018,0.070)	0.25	0.027 (-0.016,0.072)	0.22
Non-HDL-c	0.072 (-0.008,0.152)	0.09	0.065 (-0.021,0.151)	0.14

Table 2 Lo	ongitudinal	change in	lipid per	1% point	increase in	$h HbA_{1c}$ estimate
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Basic model includes sex, visit, race/ethnicity, diabetes duration, baseline HbA_{1c} and baseline lipid (model-specific) and time-varying age, change in HbA_{1c}, change in HbA_{1c} \times visit interaction

Saturated model includes variables included in the basic model plus time-varying BMI, waist circumference and visceral fat

^a Change in TG per 1% change in HbA_{1c} from a starting level of 0.93 mmol/l

In post hoc analyses, we found that the association of HbA_{1c} with TC, LDL-c and non-HDL-c was of reduced magnitude in women as compared with men. HbA_{1c} was not associated with HDL-c in either men or women, and the association of HbA_{1c} with TG was significant only in

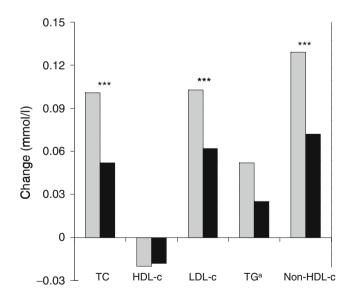


Fig. 1 Longitudinal association of change in lipid per 1% change in HbA_{1c} for the basic model (adjusted for sex, visit, race/ethnicity, diabetes duration, baseline HbA_{1c} and baseline lipid [model-specific] and time-varying age, change in HbA_{1c}, and change in HbA_{1c}×visit interaction). Grey bars, participants receiving no lipid medication; black bars, participants receiving lipid medication. ***p<0.001 for dyslipidaemia medication status for change per 1% increase in HbA_{1c}.^aEstimated change in TG from a starting level of 0.93 mmol/l

women not on dyslipidaemia medications, whereas men had a more variable pattern. Sex-based differences in CVD risk in type 1 diabetes have been previously reported [18–20] and we have reported that insulin resistance was associated with a proatherogenic lipoprotein subfraction cholesterol distribution in women [21]. Further investigation of sexbased differences in CVD risk in type 1 diabetes is needed.

CACTI is an observational cohort, and use of dyslipidaemia medications increased from 17% to 46% over the mean 6 years of follow-up, demonstrating increased use of dyslipidaemia medications for lipid and cardiovascular health in type 1 diabetes. Not surprisingly, the fasting lipid profile improved over the study period despite increases in adiposity measures. In contrast, there was minimal average improvement in HbA_{1c} over the study period. In the DCCT trial, TC, LDL-c and TG were significantly lower in the intensively treated arm than in the conventionally treated arm [10]. In addition, no participants were on statins at the end of the DCCT in 1993, while in 2004 in EDIC 33% and 34% in the intensive and conventional arms, respectively. were on statins [22]. The improved lipid profile in CACTI over time is likely due to increased use of dyslipidaemia medications, not improved glycaemic control.

Data clearly suggest that care for type 1 diabetes has improved in the past two decades, as evidenced by reduced rates of microvascular disease [23–25]; however, substantially less progress has been made in reduction of macrovascular disease rates [25–27], despite the DCCT/EDIC study showing that intensive glycaemic control over a mean of 6.5 years reduced CVD complications by 57% after a

Table 3 Longit Lipid	Longitudinal change in lipid per 1% point increase in HbA ₁ ^c estimate in men NO lipid medications	% point increase	e in HbA _{1c} estimate in men		YES lipid medications			
	Basic		Saturated		Basic		Saturated	
	β (95% CI)	p value	β (95% CI)	<i>p</i> value	β (95% CI)	<i>p</i> value	β (95% CI)	p value
TC HDI -c	0.220 (0.130, 0.310) 0.011 (-0.030_0.052)	<0.0001	0.232 (0.132, 0.332) 0.017 (-0.030_0.064)	<0.0001 0.49	0.141 (0.025, 0.257) 	0.02	0.121 (-0.002, 0.244) -0 037 (-0 080 0 006)	0.059 0.10
LDL-c	0.193 (0.109, 0.277)	<0.0001	0.214 (0.118, 0.310)	<0.0001	0.148 (0.047, 0.249)	0.005	0.136 (0.024, 0.248)	0.02
TG^{a}	0.058 (0.005, 0.113)	0.03	$0.024 \ (-0.030, \ 0.083)$	0.40	0.056 (-0.005, 0.120)	0.07	$0.064\ (0.002,\ 0.131)$	0.046
Non-HDL-c	0.217 (0.123, 0.311)	<0.0001	0.223 (0.117, 0.329)	<0.0001	0.172 (0.064, 0.280)	0.002	0.161 (0.047, 0.275)	0.007
	Basic Basic B (95% CD	n value	a	n value	Basic Basic	n value	Saturated 6 (95% CD)	n value
	β (95% CI)	<i>p</i> value	β (95% CI)	<i>p</i> value	β (95% CI)	<i>p</i> value	β (95% CI)	<i>p</i> value
TC	$0.066\ (0.001,\ 0.131)$	0.049	0.095 (0.024, 0.166)	0.00	-0.054 (-0.191, 0.083)	0.44	0.057 (-0.084, 0.198)	0.43
HDL-c	-0.025 (-0.060, 0.10)	0.16	-0.010 (-0.045, 0.025)	0.57	-0.010 (-0.075, 0.055)	0.77	0.12 (-0.057, 0.081)	0.73
LDL-c	0.075 (0.018, 0.132)	0.01	$0.090\ (0.031,\ 0.149)$	0.003	-0.035 (-0.155, 0.085)	0.57	-0.037 $(-0.159, 0.085)$	0.55
TG^{a}	$0.051 \ (0.018, \ 0.086)$	0.002	0.047 (0.013 , 0.084)	0.008	-0.020(-0.079, 0.043)	0.52	-0.006(-0.064, 0.057)	0.85
Non-HDL-c	0.101 (0.040, 0.162)	0.001	0.112 (0.049, 0.175)	0.0005	-0.057 (-0.188, 0.074)	0.40	-0.050 (-0.181, 0.081)	0.46
Basic model inc	Basic model includes sex, visit, race/ethnicity, diabetes duration,	y, diabetes dura	ttion, baseline HbA _{lc} and base	line lipid (mode	el specific) and time-varying a	ge, change in F	baseline HbA _{1c} and baseline lipid (model specific) and time-varying age, change in HbA _{1c} , change in HbA _{1c} × visit interaction	interaction
Saturated mode	I includes variables included 1	in the basic mod	Saturated model includes variables included in the basic model plus time-varying BMI, waist circumference and visceral fat β -coefficients, 95% CI and p value for 1% change in HbA1c estimate	st circumterenc	e and visceral fat (5-coefficient	s, 95% CI and	o value for 1% change in HbA	l c estimate

 $^{\rm a}$ Change in TG per 1% change in HbA $_{\rm lc}$ from a starting level of 0.93 mmol/l

mean of 17 years of follow-up [22]. People with type 1 diabetes continue to have excess morbidity and mortality, with a dramatic increase in CVD mortality risk compared with people without diabetes [27].

To describe the clinical course of type 1 diabetes in the post-DCCT age of intensive insulin therapy, data from 161 participants of the Pittsburgh Epidemiology of Diabetes Complications (EDC) Study (n=88 at the 2006 follow-up) were compared with those from the DCCT/EDIC Study [28]. The EDC reported a mean HbA_{1c} similar to that of the conventional arm of the DCCT (9%) until EDC year 8 (1994-1996), when HbA_{1c} fell by 0.5%, subsequent to publication of initial data from the DCCT demonstrating reduced microvascular complications with intensive control [28]. Subjects in the observational EDC reported a decrease in HbA_{1c} from 9.0% to 8.3%, an increase in BMI from 24 to 28 kg/m², and an increase in statin use from 1% to 38%from 1986–1988 to 2006, similar to the CACTI data. The authors emphasise that, based on these findings, current outcomes for people with type 1 diabetes should be much improved in comparison to cohorts who had onset of type 1 diabetes in the 1950s to 1970s.

In addition to glycaemic control, dyslipidaemia is another important CVD risk factor with extensive data to support its role as a target to improve cardiovascular health in people with type 1 diabetes [5]. Despite abundant data on the importance of controlling dyslipidaemia, adequate control of this CVD risk factor is frequently not achieved [9]. Our results add to those of the DCCT/EDIC and EDC studies by providing data on glycaemic and lipid control in a large observational cohort in the early 21st century. Moreover, we investigate the interplay of glycaemic control and dyslipidaemia medication use on the fasting lipid profile over a 6 year period in a large cohort of adults with type 1 diabetes. As the DCCT/EDIC study has shown that hyperglycaemia is a risk factor for hypertension [12], our data support a similar role for hyperglycaemia as a risk factor for a more atherogenic lipid profile; however, the association of glycaemia with lipids is attenuated by dyslipidaemia medication use in CACTI, whereas in the DCCT/EDIC study the association of intensive therapy with reduced incidence of hypertension remained significant in sensitivity analyses when accounting for use of antihypertensive medications [12].

While intensive treatment of glycaemia has been shown to reduce vascular complications in type 1 diabetes [22], there is also concern that intensive control is associated with weight gain, leading to adverse effects on blood pressure and lipids [29]. In the CACTI cohort, HbA_{1c} was relatively stable, adiposity measures increased and lipids improved over time. How this improved lipid profile translates into changes in health outcomes remains an important future research objective. Limited clinical trial data exist on the effect of dyslipidaemia medications on lipids and outcomes in type 1 diabetes, but the Heart Protection Study included 615 participants with type 1 diabetes. Though the study was underpowered to detect an effect in subgroup analysis, the magnitude of the risk reduction was similar to that in type 2 diabetes [30], consistent with a subsequent meta-analysis [31].

As noted, CACTI is an observational cohort study and not the result of a randomised clinical trial; therefore our data have limitations to consider when applying these findings to clinical scenarios. While 8% of participants did not have at least one follow-up visit, minimal differences exist between those who did and did not follow up. We also did not measure the effect of change in diet and exercise on change in lipid levels in these analyses, although these effects would likely be captured in the adiposity measures, which were adjusted for in the analyses. Nonetheless, CACTI participants represent one of the largest cohorts of persons with type 1 diabetes followed over 6 years who have had detailed and repeated measures of glycaemia, lipids, dyslipidaemia medications and adiposity. Our analyses stratified by sex were post hoc and should be further investigated. Finally, sensitivity analyses were performed in which participants with serum creatinine >2.0 mg/dl or microalbuminuria were removed from the analysis, and the associations of change in HbA_{1c} with change in lipids were similar.

In conclusion, over a 6 year period in the CACTI cohort, change in glycaemia was associated with a concomitant change in TC, LDL-c, TG and non-HDL-c, which was not observed in the participants who were on dyslipidaemia medications. These data suggest that there is a statistically strong association between change in glycaemia and change in lipid levels; however, this association may not be as clinically relevant compared with the stronger effect of treatment with dyslipidaemia medications on lipid levels. Therefore, both glycaemic control and treatment with dyslipidaemia medications, when indicated, in adults with type 1 diabetes may be required to optimise lipid and cardiovascular health and improve patient outcomes. Further data are needed to inform clinical decisions on lipid lowering in type 1 diabetes.

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