BMI is an important driver of β -cell loss in type 1 diabetes upon diagnosis in 10 to 18-year-old children

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Abstract

Objective: Body weight-related insulin resistance probably plays a role in progression to type 1 diabetes, but has an uncertain impact following diagnosis. In this study, we investigated whether BMI measured at diagnosis was an independent predictor of C-peptide decline 1-year post-diagnosis.

Design: Multicentre longitudinal study carried out at diagnosis and up to 1-year follow-up.

Methods: Data on C-peptide were collected from seven diabetes centres in Europe. Patients were grouped according to age at diagnosis (<5 years, $n=126$; >5 years <10 years, $n=295$; >10 years <18 years, $n=421$; >18 years, $n=410$). Linear regression was used to investigate whether BMI was an independent predictor of change in fasting C-peptide over 1 year. Models were additionally adjusted for baseline insulin dose and HbA1c.

Results: In individuals diagnosed between 0 and 5 years, 5 and 10 years and those diagnosed >18 years, we found no association between BMI and C-peptide decline. In patients aged 10–18 years, higher BMI at baseline was associated with a greater decline in fasting C-peptide over 1 year with a decrease (β 95% CI; P value) of 0.025 (0.010, 0.041) nM/kg per m^2 higher baseline BMI ($P=0.001$). This association remained significant after adjusting for gender and differences in HbA1c and insulin dose ($\beta=0.026$, 95% CI=0.0097, 0.042; $P=0.002$).

Conclusions: These observations indicate that increased body weight and increased insulin demand are associated with more rapid disease progression after diagnosis of type 1 diabetes in an age group 10–18 years. This should be considered in studies of β -cell function in type 1 diabetes.

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Introduction

Type 1 diabetes is considered an autoimmune disorder caused by inflammatory β -cell destruction, and presenting in childhood is characterised by loss of weight, ketoacidosis, acute onset and insulin dependency. In contrast, type 2 diabetes is considered a metabolic disorder usually presenting in adults, associated with obesity and insulin resistance (1).

However, insulin resistance may influence disease progression following diagnosis also in type 1 diabetes. Yki-Jarvinen & Koivisto (2) showed that newly diagnosed type 1 diabetes patients who went into remission of insulin therapy had greater insulin sensitivity than patients who continued to require insulin therapy. In addition, they found that more than a year post-diagnosis, insulin sensitivity in these patients was associated with body weight and was significantly reduced (2). Ludvigsson *et al.* (3) described in a national cohort of patients with type 1 diabetes that those with higher BMI had higher C-peptide at diagnosis, but then lost relatively more C-peptide during the first years after diagnosis.

The concept that insulin resistance may play a key role in the aetiology of type 1 diabetes is described by the accelerator hypothesis (4) and by the wider β -cell stress hypothesis (5).

The accelerator hypothesis argues that type 1 and type 2 diabetes are disorders that are aggravated by increased insulin resistance due to body weight gain, set against different genetic backgrounds, with the autoimmunity of type 1 diabetes being an expected response to upregulated β -cell response in individuals with genetic predisposition to heightened immunoreactivity.

Support for this hypothesis comes from a number of sources. In addition to the increase in incidence of type 2 diabetes that has paralleled increase in obesity, there has been an equally striking rise in the incidence of type 1 diabetes (6). Rapid weight gain during the first years of life has been found to increase the incidence of type 1 diabetes (7) and a number of studies have also shown a positive correlation between the year of onset of type 1 diabetes and BMI (8, 9, 10). More recently, we have shown that the decline in fasting C-peptide is more rapid in those patients with an earlier age of onset of type 1 diabetes, and that the factors influencing the decline in fasting C-peptide following diagnosis differ substantially according to the age of onset (11). Several studies have reported data showing that the age of onset of type 1 diabetes is inversely related with BMI, with children diagnosed at younger ages typically being heavier than those diagnosed

later (9, 12, 13, 14). Genetic studies have demonstrated that the proportion of high-risk HLA haplotypes has declined over time and been replaced by protective HLA haplotypes, arguing that the environmental contribution to disease risk (i.e. insulin resistance associated with increased obesity) has become more prominent in type 1 diabetes aetiology alongside the secular trends in obesity (15, 16). Finally, several studies indicate that in individuals at risk for the development of type 1 diabetes, insulin resistance is an independent risk factor for disease development (17, 18, 19, 20), and in studies of identical twins the co-twin who goes on to develop type 1 diabetes is more insulin resistant than the twin who does not (21). However it is important to add that other studies question these findings, and hence the theory is controversial (22).

Although some of the evidence summarised above suggests a potential key role for insulin resistance in type 1 diabetes aetiology, there has been no direct demonstration that insulin resistance in type 1 diabetes is detrimental to β -cell function even though, as mentioned above, it has been shown that C-peptide decline is more rapid in type 1 diabetic patients with higher BMI (3).

In order to investigate whether body weight influences loss of C-peptide in type 1 diabetes, we have investigated the association between BMI measured at age of onset and change in fasting C-peptide levels in the first year post-diagnosis using longitudinal data from a large data set of European type 1 diabetes patients recruited from seven different centres.

Subjects and methods

Subjects

From a cohort of 3929 Caucasian patients with type 1 diabetes of seven European registries (Belgian Diabetes Registry, Leuven Database, Hungary Database, Spain Database, Sweden Database, Germany Database, Italy Database IMDIAB), we included 1252 patients for whom data on fasting C-peptide, BMI, HbA1c levels and insulin dose at baseline and additionally for fasting C-peptide at 1 year follow-up were available. Eighty-one patients were placebo-treated subjects from immune intervention trials. In each study centre, type 1 diabetes was diagnosed according to ADA and WHO criteria. A detailed description of the cohort has been published elsewhere (11).

Measurements

Fasting, stimulated C-peptide and HbA1c were measured by standardised assays at each centre after an overnight pause in long-acting insulin; fasting glucose was required to be 70–200 mg/dl at the time of sampling. Baseline C-peptide was measured within the first 12 weeks since initiation of insulin therapy. Different methods were used for C-peptide determination, but all had similar detection limit (0.01 nM), and each centre covered patients of different age groups. Weight and height were measured using standardised procedures and used to calculate BMI (weight/height²).

Statistical methods

Statistical analysis was performed with Stata 11.2 (StataCorp. 2009, Stata Statistical Software: Release 11.2, TX, USA). Participants were stratified into four groups according to their age at onset of type 1 diabetes: very young (≤ 5 years), pre-pubertal (> 5 and ≤ 10 years), pubertal (> 10 and ≤ 18 years) and adult (18–65 years), and all analyses were performed for the whole study population and for each group.

To investigate the influence of BMI on change in fasting C-peptide over 1 year (Δ C-peptide_{0–1}), linear regression models were used with change in fasting C-peptide (C-peptide_{Year-1} – C-peptide_{baseline}) as an outcome variable and baseline BMI as the independent variable of interest. The models were adjusted for change in BMI over 1 year (Δ BMI – 1) for subjects with available data about BMI after 1 year from the diagnosis ($n=780$). The models were further adjusted for age of onset, sex, baseline HbA1c and baseline insulin dose to investigate whether differences in these variables explained any observed association between BMI and Δ C-peptide_{0–1}. Δ C-peptide_{0–1} was included in models as an untransformed variable since data were normally distributed. All models were also further adjusted for study centre to account for potential differences in the method of measurement of C-peptide and HbA1c.

In addition to analysing data using untransformed BMI variables, for children in the 0–5, 5–10 and 10–18 year age groups, we also performed regression analysis with BMI transformed to z-scores by the least mean squares (LMS) method (23) using the 1990 British Growth reference (zanthro command in Stata), which accounts for the fact that differences in BMI between children of different ages can be a result of differences in growth and development rather than solely adiposity. However, since

the use of untransformed BMI or BMI z-scores did not alter our results, data are reported for untransformed BMI variables given the ease of interpretation and comparison with results in adults. Finally, potential interactions between baseline variables were investigated by fitting continuous interactions terms in regression models.

To maximise the sample size, we report analyses that are not restricted to those with non-missing values for all parameters of interest (BMI, gender, HbA1c and insulin dose), and hence the sample size for each regression model is slightly different. However, the results were not different when we performed sensitivity analysis focusing solely on individuals with non-missing values for all traits.

Results

Baseline characteristics of study participants according to age of onset are given in Table 1.

Association of BMI with change in fasting C-peptide over 1 year (Δ C-peptide_{0–1}) in the total cohort

At baseline, individuals with a higher BMI had higher fasting C-peptide levels with a 5.2% (4.0, 6.5) increase per 1 kg/m² higher BMI ($P=3.3 \times 10^{-16}$).

Individuals with a higher BMI at baseline had a greater decline in Δ C-peptide_{0–1} with each 1 kg/m² higher BMI associated with a decrease (β 95% CI; P value) of -0.012 (-0.020 , -0.0046 ; 0.0002) nM in fasting C-peptide. This association remained significant after taking into account the differences in baseline HbA1c levels ($P=0.012$) and Δ BMI_{0–1} ($P=0.026$), but not when variation in baseline insulin dose ($P=0.060$) or both HbA1c levels and insulin dose ($P=0.11$) was controlled.

Association of BMI with change in fasting C-peptide over 1 year (Δ C-peptide_{0–1}) by age group

At baseline, differences in BMI were not associated with fasting C-peptide in those diagnosed at ≤ 5 years ($P=0.45$). In those diagnosed between 5 and 10 years, a larger BMI at baseline was significantly associated with greater fasting C-peptide levels, with each 1 kg/m² higher BMI associated with a 4.8% (2.1, 7.5) increase in fasting C-peptide ($P=0.00052$).

BMI was not a predictor of Δ C-peptide_{0–1} in those diagnosed at ≤ 5 years ($\beta=0.003$, 95% CI = -0.022 , 0.027; $P=NS$) or those diagnosed between 5 and 10 years of age ($\beta=0.011$, 95% CI = -0.0030 , 0.026; $P=NS$).

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Table 1 Characteristics of population by age of onset. Data are mean $\pm$ s.d.

Characteristic	Total cohort	Age at diagnosis (years)			
		≤ 5	> 5 and ≤ 10	> 10 and ≤ 18	> 18
<i>n</i> (m/f)	1252 (752/500)	126 (79/47)	295 (156/139)	421 (257/164)	410 (260/150)
Age of onset (years)	16.7 (10.6)	3.2 (1.5)	8.2 (1.4)	13.4 (2.1)	29.3 (7.4)
BMI (kg/m^2)	18.8 (3.9)	15.3 (2.3)	16.1 (2.7)	18.7 (3.1)	22.0 (3.4)
BMI z-score for age	-0.40 (1.3)	-1.1 (1.9)	-0.26 (1.3)	-0.31 (1.1)	-0.40 (1.3)
Insulin dose (IU/kg per day)	0.66 (0.37)	0.86 (0.50)	0.70 (0.34)	0.71 (0.35)	0.51 (0.30)
HbA1c (%)	10.3 (3.0)	9.9 (2.4)	9.9 (2.9)	10.0 (3.2)	11.0 (2.8)
Fasting C-peptide (nM)	0.31 (0.37)	0.16 (0.21)	0.21 (0.27)	0.33 (0.40)	0.41 (0.41)

At baseline, in individuals diagnosed between 10 and 18 years, a higher BMI was associated with higher fasting C-peptide levels with a 3.3% (0.98, 5.6) increase per 1 kg/m^2 difference in BMI ($P=0.0054$).

In individuals diagnosed between 10 and 18 years, a higher BMI at baseline was associated with a greater decline in fasting C-peptide over 1 year with a decrease of -0.025 (-0.010 , -0.041 ; 0.001) nM for each 1 kg/m^2 higher baseline BMI. This association remained significant even after differences in baseline HbA1c, and insulin dose were taken into account ($\beta = -0.026$, 95% CI = -0.0097 , -0.042 ; $P=0.002$). Data on BMI 1 year after the diagnosis were available for 264 subjects. In these subjects, the association between baseline BMI and the decline in C-peptide remained significant also after adjustment for ΔBMI_{0-1} ($\beta = -0.028$, 95% CI = -0.0049 , -0.006 ; $P=0.012$). The detrimental effect of increased BMI at baseline was reduced in those individuals with a larger baseline insulin dose ($p_{\text{interaction}}=0.004$).

In individuals diagnosed at > 18 years, each 1 kg/m^2 higher BMI was associated with a 5.4% (3.3, 7.4) increase in fasting C-peptide ($P=5.0 \times 10^{-7}$); however, baseline BMI was not a significant predictor of $\Delta \text{C-peptide}_{0-1}$.

The association between changes in BMI and $\Delta \text{C-peptide}_{0-1}$ in different age of onset groups is shown in Fig. 1.

Inclusion of BMI z-scores in regression models of individuals diagnosed at < 18 years of age (in place of untransformed BMI) did not alter the results.

Discussion

This study demonstrates that increased BMI is associated with an accelerated rate of fasting C-peptide decline over 1 year in newly diagnosed type 1 diabetes patients aged 10–18 years. It is assumed that insulin resistance may play a role in the pathophysiology or progression to type 1 diabetes, and the results reported here suggest that, in

those diagnosed with an age of onset of between 10 and 18 years, decreased insulin sensitivity may play an important role for residual C-peptide after diagnosis.

Although increasing insulin secretion strives to maintain normal glucose tolerance in response to reductions in insulin sensitivity, the increased secretory demand in insulin resistant states can potentially be detrimental to β -cell function, possibly due to β -cell dysfunction resulting from the excessive stimulation of β -cells necessary for the maintenance of normoglycaemia in insulin-resistant states (24, 25, 26).

It is well known that a greater BMI is associated with reduction in insulin sensitivity (26, 27), and hence the greater rate of decline of C-peptide in patients with a higher BMI could plausibly result from β -cell exhaustion by the mechanism described previously.

An important question to be addressed is why the detrimental association of BMI with fasting C-peptide decline is observed only in those individuals aged 10–18 years. A likely explanation is the pubertal transition that occurs during this age range. A previous longitudinal study has shown that during puberty there is a transient decrease in insulin sensitivity, and hence the pubertal transition is an example of an insulin-resistant state (28).

Figure 1

Association between 1 kg/m^2 change in BMI and $\Delta \text{C-peptide}^{0-1}$ in age of onset and sex-adjusted models by age of onset group.

The lack of association between BMI and fasting C-peptide decline over 1 year in younger and older age groups may be due to the fact that β -cell dysfunction is much more severe in the former and much less severe in the latter (29, 30, 31, 32, 33). Indeed individuals in our study diagnosed at ≥ 18 years of age were not obese according to typical BMI cut-offs.

The greater decline in C-peptide observed in patients with a higher BMI could also reflect greater low-grade inflammation associated with obesity (34, 35).

In the SEARCH for Diabetes in Youth study, the decline in β -cell function 12 months after the diagnosis was independent of BMI z-score and insulin resistance among autoantibodies-positive patients with diabetes diagnosed at the age of 10–20 years (36). Different to our data, the SEARCH study used a pathogenesis-based approach for the classification of diabetes; furthermore, the SEARCH study included a wide range of racial and ethnic groups and 23% of the population was not Non-Hispanic White.

Our work may have relevant clinical implications. Although we are not able to define whether insulin resistance or adiposity is the main determinant of the β -cell decline during the first year after diagnosis, the results presented here may suggest that, especially in individuals diagnosed between 10 and 18 years of age, intervention to reduce insulin resistance (e.g. weight loss) or drugs aimed at improving insulin sensitivity may be a possible component of therapy aimed at preventing fasting C-peptide decline, at least in adolescents. In addition, if the association between BMI and increased C-peptide decline is a result of pro-inflammatory processes associated with obesity, then treatments aimed to reduced inflammation would also be beneficial. Previous studies have shown that with increasing body weight, systemic immune mediators increase not only for type 2 diabetes but also for type 1 diabetes (37, 38). Although anti-IL1 therapy was not effective in a post hoc subanalysis of high BMI subjects (39), properly powered and designed studies also testing other anti-inflammatory interventions are needed to determine this question.

Limitations of our study include the reporting of results using untransformed BMI variables for the ease of interpretation and comparison with the data from adults in our cohorts. To overcome this problem, we repeated all analyses with BMI z-score variables and found no difference in results indicating that increased adiposity is likely to underlie this association.

We have used BMI as a proxy for insulin resistance in this study, and although the two variables are strongly related, it would have been preferable to have performed

the analysis by directly measuring insulin resistance; however, direct measurement of insulin resistance was not available for analysis. We quantified β -cell function by analysing C-peptide in the fasting state, which may only partially reflect β -cell function in the stimulated state (e.g. after a meal). Although we did have data on stimulated C-peptide, a limited number of patients had sufficient data for analysis; hence we decided to maximise our sample size by using fasting C-peptide alone as our outcome variable. Data are also limited by the use of different C-peptide assays though all assays had similar detection limits.

Moreover, we did not adjust linear regression models for baseline C-peptide. However, the inclusion of fasting C-peptide at baseline in models is actually a matter of debate in statistical analysis. Thus, we decided not to include it as it may introduce a bias (40).

Another limitation is that diabetes-associated autoimmunity was not evaluated in our study and this parameter appears to be a major variable associated with β -cell decline, at least in patients <20 years of age at diagnosis (36).

In conclusion, our work provides supportive evidence for a key role of body mass in type 1 diabetes disease progression after diagnosis. However, additional studies are needed to define whether insulin resistance or adiposity are key factors leading to β -cell decline soon after diagnosis of type 1 diabetes.

Declaration of interest

The authors declare that there is no conflict of interest that could be perceived as prejudicing the impartiality of the research reported.

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Author contribution statement

A Barker, A Lauria and P Pozzilli were responsible for the design of the study, acquisition of data, analysis and interpretation of data, writing the manuscript and revising it critically for important intellectual content. N Schloot, N Hosszofalusi, J Ludvigsson, C Mathieu, D Mauricio, M Nordwall, B Van der Schueren, T Mandrup-Poulsen, W A Scherbaum, I Weets and F K Gorus made substantial contributions to the acquisition of data and revised the manuscript critically for important intellectual content. N Wareham made a substantial contribution to the analysis of the data and critical revision of the manuscript for important intellectual content. R D Leslie contributed to discussion and reviewed/edited the manuscript. All authors approved the final version.

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