

Insulin action in adipose tissue in type 1 diabetes

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Background: Insulin action has been reported to be normal in type 1 diabetic patients. However, some studies have reported an insulin resistance state in these patients. The aim of this study was to investigate insulin resistance in a group of type 1 diabetic patients. We studied the insulin action in adipose tissue and analyzed the effects of duration of disease, body mass index (BMI), and glycosylated hemoglobin on insulin action at the receptor and postreceptor levels in adipocytes.

Methods: Nine female type 1 diabetic patients with different durations of disease and eight nondiabetic female patients of comparable age and BMI were studied. ¹²⁵I-insulin binding and U-[¹⁴C]-D-glucose transport was measured in a sample of subcutaneous gluteus adipose tissue obtained by open surgical biopsy from each subject.

Results: The duration of disease was negatively correlated with both ¹²⁵I-insulin binding capacity ($r = -0.70$, $P < 0.05$) and basal and maximum insulin-stimulated glucose transport ($r = -0.87$, $P < 0.01$, and $r = -0.88$, $P < 0.01$, respectively). Maximum specific ¹²⁵I-insulin binding to the receptors in adipocytes was higher in the group of patients with a shorter duration of disease ($P < 0.01$). Basal and maximum insulin-stimulated glucose transport was significantly higher in the group with less than 5 years of disease ($P < 0.01$). No correlation was found between BMI and insulin action.

Conclusion: Female type 1 diabetic patients have normal insulin action. There is a high glucose uptake in the early phase of the disease, although a longer duration of disease appears to be a contributing factor to a decrease in insulin action in these patients, and involving both receptor and postreceptor mechanisms.

Keywords: type 1 diabetes, insulin action, insulin resistance, adipose tissue

Introduction

Type 1 diabetes is an organ-specific autoimmune disease caused by altered immune tolerance to specific proteins, with selective destruction of insulin-producing beta cells in genetically predisposed individuals.¹ In contrast, insulin resistance is considered to be the main pathophysiological factor for type 2 diabetes.²

Type 1 diabetes patients have less cardiovascular disease than those who suffer from type 2 diabetes. However, a higher cardiovascular risk, as well as greater insulin resistance, has been reported in type 1 diabetes patients with a positive family history of type 2 diabetes compared with type 1 diabetic patients with no family history of type 2 diabetes.³ Several authors have shown that both hyperinsulinism and insulin resistance could explain why patients with type 2 diabetes generally have more macrovascular disease than those with type 1 diabetes.^{4,5}

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Normal insulin action has been reported in type 1 diabetes.³ In contrast, other studies have described insulin resistance in these individuals, with remission after a variable follow-up of disease. Prevention of insulin resistance in type 1 diabetes has recently been recommended. In overweight adult type 1 diabetes patients or in those with increased insulin resistance, insulin sensitizers could be adequate in addition to insulin therapy.⁶ This rationale would lead to a diminished risk of macrovascular disease and subsequently to a lower cardiovascular risk in these patients.

We have studied insulin action in type 1 diabetes patients and evaluated the effect of glycemic control, body mass index (BMI), and time of duration of disease on insulin action in these patients.

Materials and methods

Patients

Nine female patients with type 1 diabetes were recruited from endocrine clinics. A fasting and glucagon-stimulated C-peptide level <1 ng/dL, as well as positivity for islet-cell (IA2) and anti-GAD65 antibodies, were used as diagnostic criteria for type 1 diabetes. The duration of disease ranged from 4 months to 10 years. None of the patients were on medications that could alter insulin action, apart from exogenous insulin itself. A group of eight healthy women matched for age and BMI, and with a normal glucose response to an oral glucose tolerance test 2 hours after an overload of glucose 75 g, comprised the control group. Their BMIs ranged from 20 to 28 kg/m². The study was approved by the local ethics committee of the hospital and written informed consent was obtained from all the participants before entering the study.

Adipocyte isolation

An open surgical biopsy was performed in every patient to obtain subcutaneous gluteus adipose tissue for the measurement of ¹²⁵I-insulin binding, and U-[¹⁴C]-D-glucose transport (New Research Products, Dupont de Nemours, GmbH, Bad Homburg, Germany). Adipocytes were isolated by the method described by Rodbel and modified by Pedersen et al,⁷ and insulin binding was measured by the method described by Sinha et al.⁸ Fat tissue in the gluteus region was digested with collagenase 1.5 mg/dL in 10 mmol HEPES Krebs-Ringer bicarbonate (KRB) buffer, pH 7.4, containing 3% bovine serum albumin-KRB buffer, for 30–45 minutes at 37°C, under a 95% oxygen/5% CO₂ atmosphere with continuous shaking. Cells were filtered through a nylon mesh and washed with 1% bovine serum

albumin-KRB buffer. An aliquot of the cell suspension was fixed overnight in 10% formalin-buffered phosphate to determine cell size. Another aliquot was stained with trypan blue to determine viability of the adipose cells.

Insulin binding to adipocytes

Fresh adipocytes were incubated with 10 mol/L ¹²⁵I-insulin in the absence and presence, respectively, of increasing concentrations of unlabeled insulin (0–10⁻⁷ mol/L) at 22°C, in a final volume of 1 mL 3% bovine serum albumin-KRB buffer for 45 minutes. Aliquots of 250 µL were centrifuged through phthalate of bis-(2-ethyl-hexile), and cell-associated radioactivity was counted. Specific binding was determined after subtracting the nonspecific value obtained in the presence of 10⁻⁶ M of insulin.

Glucose transport

U-[¹⁴C]-D-glucose uptake by the cell as described by Kashiwagi et al was used for measuring glucose transport into the adipocyte.⁹ Isolated adipocytes were incubated with 500 nmol/L U-[¹⁴C]-D-glucose in the absence and presence of insulin (0–10⁻⁷ mol/L) in a final volume of 1 mL 3% bovine serum albumin-KRB buffer for 60 minutes, at 37°C and under a 95% O₂/5% CO atmosphere, with continuous shaking. Centrifugation was used to separate the cells and three 250 µL aliquots were used for the insulin binding assay. No specific glucose transport value was subtracted from each value.

Biochemical variables

Serum insulin and C-peptide levels were measured by a commercial radioimmunoanalysis kit (Inestar Corp, Stillwater, MN). Glycosylated hemoglobin (HbA_{1c}) levels were measured by high-performance liquid chromatography (Diamat, Bio-Rad, Vienna, Austria) using a Merck-Hitachi model L9100 autoanalyzer. Both intra-assay and interassay variability were below 10%.

Statistical analysis

Data were analyzed using nonparametric tests. For the comparison between both individual data and between groups, and also between patient subgroups segregated by duration of disease, we used the Mann-Whitney *U* and Wilcoxon tests as indicated. The Spearman bivariate correlation analysis was used for assessment of the influence of diabetes duration and HbA_{1c} levels on the different variables. Data are presented as means $\pm$ standard deviation, and a *P* value of <0.05 was considered to be statistically significant. Calculations were

performed using the Statistical Package for Social Science (SPSS), version 12.0.

Results

Nine type 1 diabetic women (mean age 33 ± 12 years, BMI 23.9 ± 5.0 kg/m², HbA_{1c} $6.5 \pm 0.9\%$), and eight healthy women of comparable age and BMI were studied. Clinical features of the study subjects are shown in Table 1.

Maximal insulin binding to adipocytes (5.8 ± 0.4 versus $6.4 \pm 1.3\%$ B/50,000 cells), basal glucose transport (71.7 ± 34 versus 54.38 ± 38 FI/cel/s), and maximum insulin-stimulated (10^{-7} mol/L insulin) glucose transport (135.3 ± 74 versus 102.5 ± 47 FI/cel/s) were similar in the type 1 diabetic and control groups (Table 1).

The whole group of type 1 diabetic patients was segregated into those with up to 5 years of diabetes ($n = 5$) and those with more than 5 years of diabetes ($n = 4$). Maximum specific ¹²⁵I-insulin binding to the receptor in adipocytes ($P < 0.01$), basal glucose transport ($P < 0.05$), and maximal insulin-stimulated glucose transport ($P < 0.01$) were higher in the former group (Figure 1, Table 2).

When we compared both type 1 diabetes subgroups with the control group, we only found significant differences in basal and maximal glucose transport, showing higher values in type 1 diabetic patients with less than 5 years of disease duration ($P < 0.05$, Table 2).

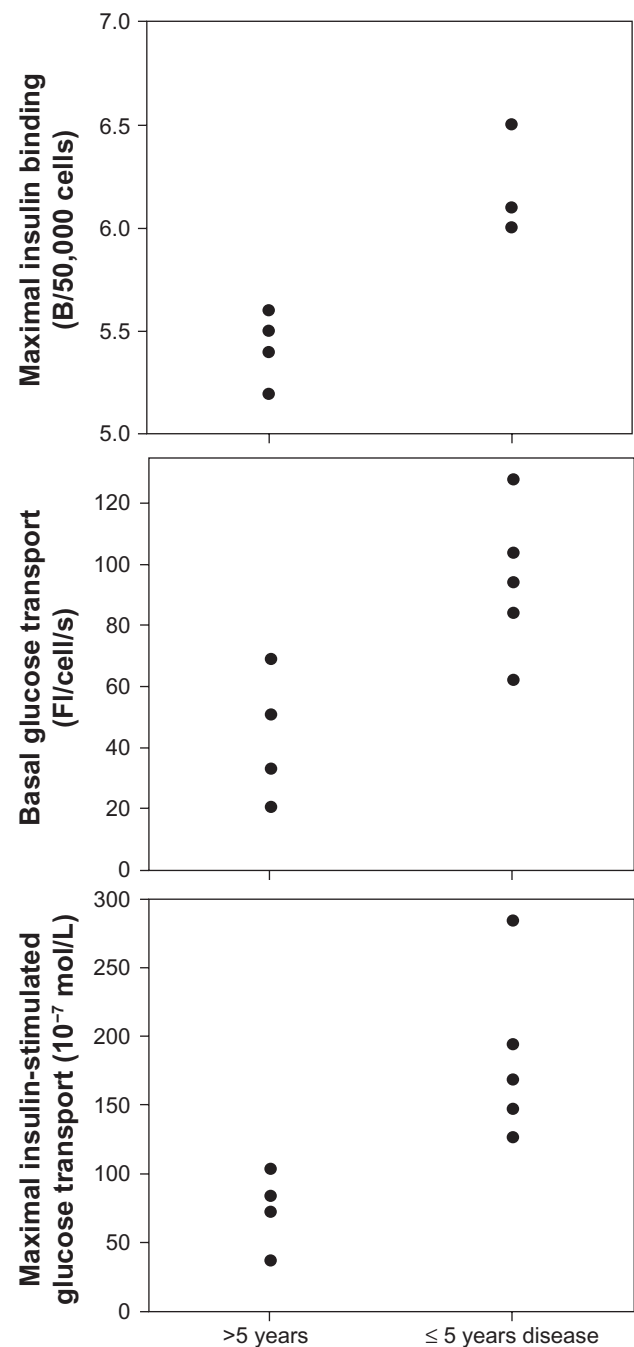
Glucose transport showed a significant negative correlation with duration of disease ($r = -0.84$, $P < 0.01$). There was a negative, but not significant, correlation between HbA_{1c} level

Table 1 Clinical and analytical data for the study groups

	T1D patients (n = 9)	Controls (n = 8)
Body mass index (kg/m ²)	23.9 ± 5.0	23.8 ± 5.2
HbA _{1c} (%)	6.5 ± 0.9^a	4.3 ± 1.0
Fasting plasma glucose (mg/dL)	249 ± 100^a	83 ± 7
Basal C-peptide (ng/mL)	0.5 ± 0.2^a	2.0 ± 1.0
C-peptide at 5 minutes after glucagon (I mg/IV) (ng/mL)	0.7 ± 0.3^a	5.0 ± 0.9
% maximum ¹²⁵ I-insulin binding	5.8 ± 0.4	6.4 ± 1.3
Basal U-(¹⁴ C)-D-glucose transport FI/cel/s	71.4 ± 34	54.3 ± 38
Maximal U-(¹⁴ C)-D-glucose transport after 10^{-7} mol/L insulin FI/cel/s	135.4 ± 74	102.5 ± 47

Notes: Data are means $\pm$ standard deviation. ^a $P < 0.01$.

Abbreviations: IV, intravenously; T1D, type 1 diabetes; HbA_{1c}, glycosylated hemoglobin.



	Time of evolution of diabetes
% Max insulin binding	$r = -0.70$ $P < 0.05$
Basal glucose transport FI/cel/s	$r = -0.87$ $P < 0.01$
Maximal glucose transport after Insulin-stimulated glucose transport FI/cel/s	$r = -0.88$ $P < 0.01$

Figure 1 Insulin action in the adipocyte: maximal insulin binding and basal and maximal glucose transport after stimulated insulin (10^{-7} mol/L) in type 1 diabetic patients with >5 years and $\leq$ 5 years of diabetes.

Table 2 Insulin binding, basal glucose transport, and insulin-stimulated glucose transport in adipocytes from type 1 diabetic patients according to duration of disease

	T1D ≤ 5 years group	T1D > 5 years group	Control group
% maximum ¹²⁵ I-insulin binding B/50,000 cells	6.2 ± 0.2 ^a	5.4 ± 0.1	6.4 ± 1.3
Basal U-(¹⁴ C)-D-glucose transport FI/cel/s	94.4 ± 24 ^{a,b}	43.5 ± 21	54.3 ± 31
Maximal U-(¹⁴ C)-D-glucose transport insulin-stimulated glucose transport (10 ⁻⁷ mol/L insulin) FI/cel/s	184.2 ± 61 ^{a,b}	74.2 ± 28	102.5 ± 47

Notes: Insulin binding is expressed as percentage of maximal ¹²⁵I-insulin binding; basal glucose transport is expressed as basal U-(¹⁴C)-D-glucose transport, and insulin-stimulated glucose transport as maximum insulin-stimulated (10⁻⁷ mol/L insulin) glucose transport. ^aP < 0.05 versus type 1 diabetes >5 years group; ^bP < 0.05 versus control group.

and insulin binding ($r = -0.33$, $P < 0.37$), as well as between HbA_{1c} and both basal and maximum glucose-stimulated transport, although neither of them were significant ($r = -0.42$, $P < 0.25$, and $r = -0.37$, $P < 0.32$ for basal and maximum glucose transport, respectively, Table 3).

Discussion

Glucose uptake is regulated by insulin in the case of high plasma glucose. Insulin is then secreted and stimulates the uptake and storage of glucose, as well as its metabolism in skeletal muscle and adipose tissue, in order to maintain a constant glucose level. In our study, the group with a longer disease duration had higher insulin requirements for metabolic control. Furthermore, a significant positive correlation was found between the time of disease and the dose of insulin required for treatment, which suggests that the appearance of insulin resistance occurs during the course of disease and becomes higher with longer duration of disease. The required insulin dose in patients with type 1

diabetes constitutes a marker of insulin resistance.^{10,11} Other factors which influence insulin action, such as suboptimal control, overweight, counter-regulatory hormone excess, or ketoacidosis, have been suggested as mechanisms for insulin resistance in patients with type 1 diabetes.¹² In our study, there were no differences in BMI and metabolic control as assessed by the level of HbA_{1c}, which could explain the difference in insulin action in type 1 diabetic patients with different durations of diabetes.

The first mechanism of insulin action is mediated by cell surface receptors which were described in 1971.¹³ An alteration in the binding of insulin to its receptor could explain an impairment of insulin action in diabetic patients. Studies of insulin action in type 1 diabetes have observed normal restored insulin action in these individuals. Although muscle tissue is the major peripheral target organ of insulin, adipose tissue is another important one, and previous studies have demonstrated that glucose uptake induced by insulin in this tissue correlates with overall glucose utilization.^{10,11} Moreover, in vitro experiments show that a reversal of hyperglycemia normalizes the insulin action in fat cells following insulin treatment.¹⁴ In addition, normal insulin sensitivity has also been reported in patients with type 1 diabetes in the preclinical stage of the disease. In subjects with newly diagnosed type 1 diabetes, a transient normalization of insulin action may occur after initial insulin treatment. In patients with type 1 diabetes, partial recovery of C-peptide secretion has been observed during the honeymoon period.^{15,16} We measured C-peptide levels, both at baseline and six minutes after stimulation with glucagon 1 mg intravenously, and an absence of insulin secretion was confirmed. Therefore, our results support the hypothesis that factors other than endogenous insulin secretion must contribute to the increase in daily insulin requirements over the duration of the disease.

High glucose transport in adipocytes as the cause of the honeymoon period, without changes in insulin secretion, has been reported.¹⁷ An increase in basal glucose transport and maximally stimulated glucose transport during the early phase of diabetes could explain the low insulin requirement needed to achieve metabolic control during the first year of the disease. In conceptual agreement, we now report high glucose transport, both basal and maximal, in type 1 diabetic patients with a shorter duration of disease. We did not observe insulin resistance in our type 1 diabetic patients, because both basal and maximum insulin-stimulated glucose uptake were not decreased in type 1 diabetic patients when compared with the control group. Furthermore, we found a negative

Table 3 Correlations between several clinical and analytical parameters and insulin action in adipocyte tissue in type 1 diabetes patients

	Length of diabetes (months)	HbA _{1c}	Body mass index
% maximum insulin binding	$r = -0.70$ $P < 0.05$	$r = -0.33$ $P < 0.34$	$r = -0.35$ $P < 0.35$
Basal glucose transport, FI/cel/s	$r = -0.87$ $P < 0.01$	$r = -0.42$ $P < 0.25$	$r = -0.51$ $P < 0.12$
Maximal glucose transport after insulin-stimulated glucose transport, FI/cel/s	$r = -0.88$ $P < 0.01$	$r = -0.37$ $P < 0.32$	$r = -0.38$ $P < 0.30$

correlation between insulin action and duration of disease in type 1 diabetic patients.

A limitation of our study is the low number of patients. Our results indicate that a longer duration of diabetes could lead to a decrease in insulin action in type 1 diabetic patients. In the early phase of the disease, an increase in glucose transport and maximal insulin glucose transport could explain the lower dose of insulin needed in type 1 diabetic patients in clinical practice. Moreover, a normal insulin action explains the frequent hypoglycemia seen in type 1 diabetes versus type 2 diabetes patients with high levels of insulin and low episodes of hypoglycemia.

In summary, type 1 diabetic patients are not insulin-resistant, and the early phase of the disease is characterized by an increase in glucose transport, both basal and insulin-stimulated. Further studies with larger number of patients are needed to confirm our results.

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Disclosure

The author reports no conflict of interest in this work.

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