

Demonstration of increased concentrations of circulating glycosylated insulin in human Type 2 diabetes using a novel and specific radioimmunoassay

J. R. Lindsay¹, A. M. McKillop², M. H. Mooney², F. P. M. O'Harte², P. M. Bell¹, P. R. Flatt²

¹ Regional Centre for Endocrinology, Royal Victoria Hospital, Belfast, Northern Ireland, UK

² School of Biomedical Sciences, University of Ulster, Coleraine, County Londonderry, Northern Ireland, UK

Abstract

Aims/hypothesis. Glycation of insulin, resulting in impaired bioactivity, has been shown within pancreatic beta cells. We have used a novel and specific radioimmunoassay to detect glycosylated insulin in plasma of Type 2 diabetic subjects.

Methods. Blood samples were collected from 102 Type 2 diabetic patients in three main categories: those with good glycaemic control with a HbA_{1c} less than 7%, moderate glycaemic control (HbA_{1c} 7–9%) and poor glycaemic control (HbA_{1c} greater than 9%). We used 75 age- and sex-matched non-diabetic subjects as controls. Samples were analysed for HbA_{1c}, glucose and plasma concentrations of glycosylated insulin and insulin.

Results. Glycosylated insulin was readily detected in control and Type 2 diabetic subjects. The mean circulating concentration of glycosylated insulin in control subjects was 12.6±0.9 pmol/l (*n*=75). Glycosylated insulin in the good, moderate and poorly controlled diabetic groups was increased 2.4-fold (*p*<0.001, *n*=44), 2.2-fold (*p*<0.001, *n*=41) and 1.1-fold (*n*=17) corresponding to 29.8±5.4, 27.3±5.7 and 13.5±2.9 pmol/l, respectively.

Conclusion/interpretation. Glycosylated insulin circulates at noticeably increased concentrations in Type 2 diabetic subjects. [Diabetologia (2003) 46:475–478]

Keywords Glycosylated insulin, glucose toxicity, Type 2 diabetes.

Formation of advanced glycation-end products (AGEs) plays an important role in long-term metabolic consequences of diabetes including ophthalmic, renal and atherosclerotic vascular complications. Glycation has been shown also to interfere with normal cellular functions including the activities of various enzymes such as Cu-Zn superoxide dismutase and several other peptide hormones. A growing body of evidence now exists to support the hypothesis that glycation of insulin in pancreatic beta cells occurs under conditions of hyperglycaemia. Glycosylated insulin has been identified in the pancreas of normal and diabetic animal models

[1]; the process of glycation in the beta cell is rapid and occurs in a time- and concentration-dependent manner [2]. Previous studies have shown that glycosylated insulin has a reduced ability to regulate plasma glucose homeostasis in vivo and to stimulate adipose tissue lipogenesis or glucose uptake and oxidation by isolated diaphragm and abdominal muscle in vitro [3, 4, 5]. Studies in healthy human volunteers using the hyperinsulinaemic-euglycaemic glucose clamp technique suggest that glycosylated insulin could contribute to insulin resistance in Type 2 diabetes mellitus [6].

The site of glycation of human insulin has now been identified by electrospray tandem mass spectrometry as the N-terminal Phe¹ of the B-chain [7], enabling the development of a sensitive and specific radioimmunoassay to measure concentrations of glycosylated insulin and to identify glycosylated insulin in pancreatic islets using immunohistochemistry [8]. High performance liquid chromatography techniques used previously were neither sensitive nor reliable

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Corresponding author: F. P. M. O'Harte, School of Biomedical Sciences, University of Ulster, Coleraine, County Londonderry, BT52 1SA, Northern Ireland, UK
E-mail: fpm.oharte@ulster.ac.uk

Table 1. Characteristics of controls subjects and Type 2 diabetic patients

	Control subjects	Patients with good control	Patients with moderate control	Patients with poor control
Number of subjects	75	44	41	17
Male/female ratio	31/44	21/23	24/17	9/8
Age (years)	59.5±2.1	64.6±1.4	63.4±1.7	61.0±3.7
Duration of diabetes (years)	–	4.8±0.6	7.1±0.9	8.5±1.3
BMI (kg/m ²)	25.6±0.5	29.6±1.0***	30.6±1.0***	28.9±1.3**
Serum Creatinine (µmol/l)	86.0±4.0	89.2±3.6	89.0±4.0	85.8±3.6
Diet (%)	–	40.9	26.8	17.6
Metformin (%)	–	18.2	14.6	11.8
Sulphonylureas (%)	–	29.5	26.8	17.6
Metformin/Sulphonylureas (%)	–	6.8	26.8	47.1
Other drug combinations (%) ^a	–	4.6	4.9	5.9

Values are means ± SEM. ** $p < 0.01$ *** $p < 0.001$ compared with control subjects. ^a Other treatment combinations included metformin and acarbose, gliclazide and acarbose and glibenclamide and acarbose

enough to measure circulating glycosylated insulin in normal or diabetic states. This paper describes the use of a sensitive and specific radioimmunoassay to measure increased concentrations of glycosylated insulin in the plasma of patients with Type 2 diabetes mellitus.

Subjects and methods

Subjects. Mid-morning blood samples were taken from Type 2 diabetic subjects ($n=102$). All subjects were controlled by diet alone or by oral hypoglycaemic agents. No subjects were receiving insulin. Type 2 diabetic subjects were divided into three groups depending upon their glycaemic control. Group 1 represented subjects under good glycaemic control with a HbA_{1c} less than 7%. Group 2 was comprised of subjects with moderate glycaemic control (HbA_{1c} 7–9%) and Group 3 represented subjects with a HbA_{1c} greater than 9% (poor glycaemic control). Age- and sex-matched normal healthy individuals served as controls. This study was approved by the ethics committee of The Queen's University of Belfast and all subjects gave their written informed consent.

Biochemical analyses. Development of a specific RIA for glycosylated insulin has been described in detail elsewhere [8]. In brief, an N-terminally glycosylated synthetic insulin peptide, closely related to the amino-terminal sequence of the insulin B-chain (Phe-Val-Asn-Gln-His-Leu-Tyr-Lys) was used to raise specific antibodies in rabbits. Antiserum R3/B/vi was used to establish a dextran-coated charcoal RIA with a glycosylated human insulin standard curve containing insulin free serum. Assay sensitivity was less than 3 pmol/l with an intra-assay coefficient of variation of 9%. The glycosylated insulin antibody cross-reacted 56% with glycosylated proinsulin but cross-reaction with non-glycosylated insulin, proinsulin and other pancreatic hormones was negligible [8]. Serum insulin was determined using a microparticle enzyme immunoassay. Assay sensitivity was less than 6 pmol/l with an intra-assay coefficient of variation of 4%. Cross-reactivity with proinsulin and split products was less than 0.01% with no detectable reaction with proinsulin C-peptide. Restricted cross-reaction was observed with glycosylated insulin corresponding to approximately 50%. Glucose concentrations, HbA_{1c} and creatinine were determined by conventional assays.

Statistical analysis. Data are expressed as means ± SEM. Significant differences between groups of data were assessed using the unpaired Student's *t* test and a *p* value of less than 0.05 was considered statistically significant.

Results

Control and diabetic groups were matched for age, sex, creatinine and BMI (Table 1). Duration of diabetes increased progressively from good, moderate to poorly controlled groups. A total of 70 diabetic subjects were taking oral hypoglycaemics of whom 16 were taking metformin alone, 27 sulphonylurea alone and 22 were taking a combination of metformin and sulphonylurea. The remaining patients were treated with dietary restriction alone, or a combination of other treatments, which included acarbose.

Control subjects had mean glycosylated haemoglobin values of 5.7±0.1% compared with 6.4±0.1 ($p < 0.001$), 7.9±0.1 ($p < 0.001$) and 10.4±0.4% ($p < 0.001$) for good, moderate and poorly controlled diabetic groups, respectively. Plasma glucose concentrations were similarly raised in the diabetic groups, with highest values observed in the poorly controlled group (Fig. 1). A positive correlation between glycosylated haemoglobin and glucose concentration was evident in the combined groups ($r=0.322$; $p < 0.01$). Plasma glycosylated insulin concentrations of good and moderately controlled diabetic groups were increased 2.4-fold ($p < 0.001$) and 2.2-fold ($p < 0.001$) compared with control subjects (Fig. 1). Plasma glycosylated insulin concentrations in the control group, well, moderately and poorly controlled diabetic groups were 12.8±1.1 (SD 9.2), 29.8±5.4 (SD 35.7), 27.3±5.7 (SD 36.5) and 13.5±2.9 (SD 11.9) pmol/l, respectively. The glycosylated insulin concentrations of the poorly controlled group were not noticeably increased and no correlation was found in the combined groups between glycosylated insulin and either glycosylated haemoglobin or plasma glucose

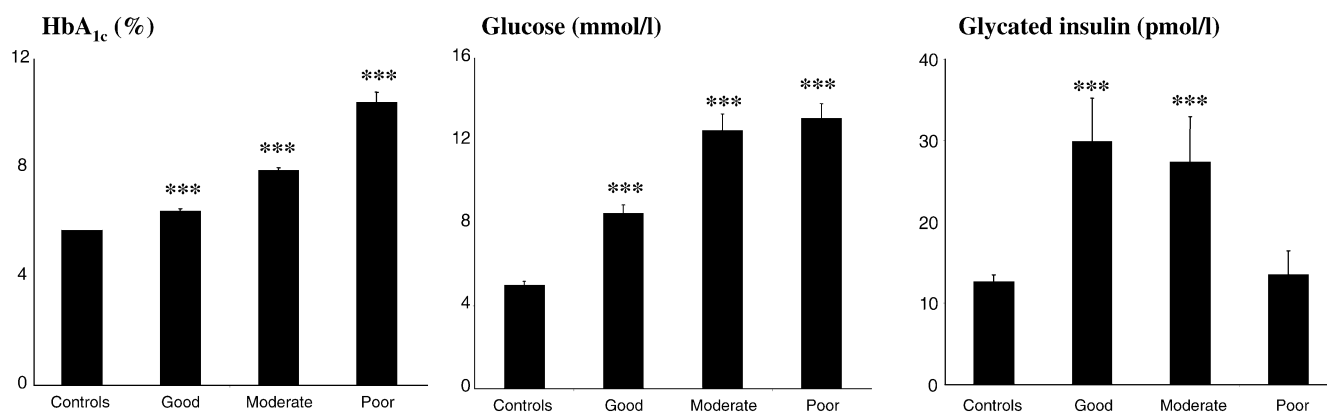


Fig. 1. HbA_{1c}, plasma glucose and glycated insulin concentrations of control subjects and diabetic patients with good, moderate or poor metabolic control. Values are means \pm SEM. *** $p < 0.001$ compared with control subjects

concentrations (data not shown). Glycated insulin was found to circulate at 26.1 ± 2.9 pmol/l in the combined groups ($p < 0.001$, compared with controls). Insulin concentrations measured using a specific insulin antiserum in the control, good, moderate and poorly controlled groups were 148 ± 19 , 191 ± 25 , 225 ± 40 and 159 ± 40 pmol/l, respectively.

Discussion

Disturbances of insulin secretion and peripheral insulin action are well-described features of Type 2 diabetes mellitus, and a growing body of data support the role of glycation of insulin as a feature of beta-cell dysfunction and insulin resistance. Evidence includes the demonstration of increased concentrations of glycated insulin in the pancreas of various animal models of diabetes [1, 8, 9] and the time- and concentration-dependent glycation of insulin in both isolated mouse islets or clonal insulin secreting cell lines under hyperglycaemic conditions [1, 2]. We have previously shown the impaired biological activity of glycated insulin using in vitro studies with isolated muscle preparations and in vivo studies using animal models or healthy human subjects subjected to hyperinsulinaemic euglycaemic glucose clamps [3, 4, 5, 6]. Having identified the site of glycation of insulin as the amino terminal Phe¹ residue of the insulin B-chain [7], a novel, sensitive and specific radioimmunoassay has recently been developed to detect glycated insulin in plasma and biological tissues [8].

In this study, we have examined circulating glycated insulin using this radioimmunoassay in 102 Type 2 diabetic patients, classified into good, moderate and poorly controlled groups, in comparison with age- and sex-matched control subjects. The results readily indicate that glycated insulin circulates at increased concentrations in Type 2 diabetic patients with increases

of 2.4-fold, 2.2-fold and 1.1-fold in the three diabetic subgroups, respectively. These results corroborate a recent study which determined raised concentrations of glycated insulin in the plasma of various animal models of Type 2 diabetes [8, 9]. Individual variations of glycated insulin in subjects from the various groups possibly reflect the collection of a single fasted sample together with possible individual differences in beta-cell secretory activity, diet history and duration of diabetes. However, it is clear from these observations that the concentration of insulin circulating in the glycated form in diabetes is significant compared with control subjects. Estimation of the precise proportion of glycated insulin relative to insulin is not possible at the moment due to differences in the cross-reactivity of the antisera with proinsulin or glycated proinsulin-derived peptides. The amounts of circulating proinsulin and split products are also likely to vary between the four observation groups.

We had initially anticipated increased concentrations of plasma glycated insulin in the most poorly controlled diabetic subjects in keeping with a time and concentration dependent glycation process in the pancreatic beta cell [2]. However this group showed relatively reduced concentrations of circulating plasma glycated insulin and insulin compared to the higher concentrations in the good and moderately controlled diabetic groups. Thus whereas a noticeable correlation existed between plasma glucose and glycated haemoglobin, concentrations of glycated insulin were not closely linked to either of these parameters. This indicates, perhaps not surprisingly, that the glycation and secretion of insulin from the pancreatic beta cells is a more dynamic process than glycation of haemoglobin which occurs progressively over a much longer period of weeks. However, circulating glycated insulin is much less susceptible to change than the prevailing plasma glucose concentration. Hence, it could be envisaged that circulating glycated insulin will depend not only on the glycaemic environment but also the ability of the beta cells to transport and metabolise glucose to reactive phosphorylated forms at the intracellular sites of (pro)insulin synthesis and storage [1, 2]. Additionally, the secretory activity and

competency of the beta cells together with metabolic clearance rate will also determine the rates of delivery and removal of glycosylated insulin from the circulation.

In conclusion this study describes increased concentrations of plasma glycosylated insulin in Type 2 diabetic subjects. We speculate that the combined effects of glucose toxicity and beta-cell failure [10], determine the total plasma concentration of glycosylated insulin in individual patients depending on the spectrum of their current disease.

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