

Glycated haemoglobin predicts progression to diabetes mellitus in Pima Indians with impaired glucose tolerance

R. R. Little^{1,2}, J. D. England², H. M. Wiedmeyer², R. W. Madsen³, D. J. Pettitt⁴, W. C. Knowler⁴, D. E. Goldstein^{1,2}

¹ Department of Pathology, University of Missouri, School of Medicine, Columbia, Missouri, USA

² Department of Child Health, University of Missouri, School of Medicine, Columbia, Missouri, USA

³ Department of Statistics, University of Missouri, School of Medicine, Columbia, Missouri, USA

⁴ Diabetes and Arthritis Epidemiology Section, National Institute of Diabetes and Digestive and Kidney Diseases, Phoenix, Arizona, USA

Summary Glycated haemoglobin could offer several practical advantages over the OGTT for assessing glucose metabolism. Initial cross-sectional studies (1983–1985) on 381 subjects (mostly Pima Indians) described the relationship between HbA_{1c} (a specific glycated Hb) and the OGTT. We performed follow-up OGTTs and HbA_{1c} measurements on 257 of these same subjects 1.6–6.1 years later. Subjects were again grouped according to both the result of the OGTT (normal, IGT or diabetes, by WHO criteria) and HbA_{1c} result (normal or elevated based on mean \pm 1.96 SD of normal). Of 66 subjects with IGT at baseline, 47 (71 %) had nor-

mal HbA_{1c} and 19 (29 %) had elevated HbA_{1c}. Twenty-six (39 %) of these subjects had diabetes at follow-up. Of these subjects with IGT, a significantly greater percentage of subjects with elevated HbA_{1c} at baseline (68 %) showed worsening to diabetes than those with a normal HbA_{1c} (28 %); (chi-square = 7.8, *df* = 1, *p* < 0.01). Thus, in subjects with IGT, glycated Hb may be a useful predictor of progression to diabetes. [Diabetologia (1994) 37: 252–256]

Key words Follow-up study, glycated Hb, HbA_{1c}, IGT, Pima Indians.

The OGTT is often used to diagnose diabetes mellitus as well as to detect more mildly impaired carbohydrate metabolism. Measurement of glycated Hb has been suggested as an alternative to the OGTT for assessing glucose metabolism [1–4]. The concentration of glycated Hb is increased within erythrocytes of patients in proportion to the degree of chronic hyperglycaemia; it is an indirect measure of the average blood glucose concentration over the previous 2 to 3 months [3]. Glycated Hb is widely used for monitoring long-term glycaemic control in known diabetic patients [5–8], and the prognostic value of glycated Hb has been investigated in a longitudinal study design of patients with normal glucose tolerance [9]. Measurement of glycated

Hb offers some practical advantages over the OGTT; it is not influenced by time of day, recent activity levels, metabolic stress, or food intake [7]. Only minimal patient cooperation is required before and during the test, and only one small blood sample is required.

Our initial cross-sectional studies [4] described the relationship between HbA_{1c}, a specific glycated Hb, and OGTT. In 381 subjects (mostly Pima Indians) from a population with a high prevalence of non-insulin-dependent diabetes mellitus, HbA_{1c} had high specificity and moderate sensitivity as a screening test for diabetes defined by the OGTT using WHO criteria [4]. Here we re-examine these same individuals 1 to 6 years later to investigate the prognostic value of glycated Hb. By using both OGTT and glycated Hb results, we attempted to identify, among a group of high risk subjects, those who are at greatest risk of developing diabetes.

Received: 8 February 1993
and in final revised form: 26 August 1993

Corresponding author: Dr. R. R. Little, Pathology Department M263, University of Missouri School of Medicine, One Hospital Drive, Columbia, MO 65212, USA

Abbreviations: OGTT, Oral glucose tolerance test; WHO, World Health Organisation; IGT, impaired glucose tolerance.

Subjects, materials, and methods

Subjects. Subjects included 257 individuals residing in the Gila River Indian Community of central Arizona (USA) who were participating in a longitudinal epidemiologic study of diabetes

Table 1. Results of OGTT and HbA_{1c} classification of 257 subjects at baseline and follow-up

Follow-up (n)	N-N (86)	N-Elev (6)	IGT-N (36)	IGT-Elev (16)	D-N (7)	D-Elev (106)
<i>Baseline (n)</i>						
N-N (93)	68	3	8	5	2	7
N-Elev (9)	5	2	1	0	0	1
IGT-N (47)	11	0	19	4	3	10
IGT-Elev (19)	1	0	1	4	0	13
D-N (14)	1	1	5	1	2	4
D-Elev (75)	0	0	2	2	0	71

N-N, Normal glucose tolerance and normal HbA_{1c};
 N-Elev, normal glucose tolerance and elevated HbA_{1c};
 IGT-N, IGT and normal HbA_{1c};

IGT-Elev, IGT and elevated HbA_{1c};
 D-N, diabetes and normal HbA_{1c};
 D-Elev, diabetes and elevated HbA_{1c}.

and obesity conducted by the National Institute of Diabetes and Digestive and Kidney Diseases since 1965.

Between 1983 and 1985, 381 subjects underwent a standard OGTT according to WHO recommendations in which plasma glucose concentrations were determined in the fasting state and 2 h after ingestion of a 75-g carbohydrate load (Glucola; Ames, Elkhart, Ind., USA). For purposes of the present study this was called the baseline visit. The sample was not representative of the total population but over represented people with IGT and diabetes. At the time of OGTT, a venous blood sample was obtained from each subject for measurement of HbA_{1c}. This blood sample was placed in an insulated container with wet ice and shipped overnight to the Diabetes Research Laboratory of the University of Missouri at Columbia (USA) [10]. Results of the baseline visits have been reported previously [4].

Between 1985 and 1990, 257 of these same individuals underwent at least one additional OGTT and HbA_{1c} determination 1 year or more after the baseline visit. This was designated the follow-up visit. For subjects with more than one follow-up visit, the most recent follow-up visit results were used in data analysis.

OGTT classification. Each subject was classified according to WHO criteria as having normal glucose tolerance, IGT, or diabetes mellitus [11]. Diagnosis of diabetes requires a fasting plasma glucose of 7.8 mmol/l or greater or a 2-h plasma glucose of 11.1 mmol/l or greater. Subjects are diagnosed as having IGT when fasting plasma glucose is less than 7.8 mmol/l and 2-h plasma glucose is 7.8 mmol/l or greater and less than 11.1 mmol/l. Individuals with a fasting and 2-h plasma glucose of less than 7.8 mmol/l were considered to have normal glucose tolerance. Results of glucose tolerance tests performed prior to the baseline visit were not considered for purposes of this classification. However, individuals who were receiving insulin or oral hypoglycaemic therapy at the baseline visit were excluded.

Preparation of blood samples and analysis of HbA_{1c}. Details of sample preparation and analysis have been described previously [4, 12]. Briefly, a haemolysate was prepared from washed and incubated erythrocytes and HbA_{1c} was measured by a component HPLC system which was calibrated at each run. Inter-assay coefficients of variation were 1.9 and 3.2% for quality control samples with mean values of 5.5 and 8.0%, respectively during the time period of this study.

HbA_{1c} classification. Each HbA_{1c} result was classified as either normal or elevated based on whether it was below or above the upper limit of the HbA_{1c} normal range which was 6.03% (mean + 1.96 SD; mean = 5.05%, SD = 0.50%). This range was determined for a mostly Caucasian population using 124 healthy volunteers at various centres throughout the United States, using the same assay method as described above. Although glucose tolerance tests were not performed on these subjects, all had

fasting plasma glucose values below 6.4 mmol/l. Mean HbA_{1c} for the Pima Indians with normal glucose tolerance was somewhat higher ($n = 159$, mean = 5.43%, SD = 0.40%).

OGTT-HbA_{1c} grouping. Each of the 257 study subjects was placed in one of six groups which describe both the OGTT and HbA_{1c} classification at baseline (Table 1).

Statistical analysis

A chi-squared test was used to test for the difference in the rate of progression to diabetes between two IGT groups; those with normal vs those with elevated HbA_{1c}. Comparison of the proportion of subjects whose diabetes persisted at follow-up among those having an OGTT in the diabetic range at baseline (diabetes/normal HbA_{1c} vs diabetes/elevated HbA_{1c}) was done using a two-tailed Fisher test. A multiple logistic regression model was used to investigate the relationship between several risk variables (e.g. HbA_{1c}, 2-h plasma glucose) and progression to diabetes in subjects with IGT.

Results

Of the 381 subjects studied at baseline (142 men, 239 women), 257 (85 men, 172 women) had follow-up examinations. The mean age of the 257 study subjects at baseline was 46.7 years (range 15–87 years). The mean time between baseline and follow-up exams was 3.3 years (range = 1.6–6.1 years). At the baseline exam, 39.7% had a normal OGTT, 25.7% had IGT, and 34.6% had diabetes (i.e. fasting glucose of 7.8 mmol/l or greater or a 2-h glucose of 11.1 mmol/l or greater). At follow-up, 35.8% had normal glucose tolerance, 20.2% had IGT, and 44.0% had diabetes. The most common change was from IGT to diabetes, although there were changes into and out of every classification. At baseline 60% of subjects had a normal HbA_{1c} while 40% had an elevated HbA_{1c} value. At follow-up only 50% had HbA_{1c} values in the normal range.

Table 1 shows the number of subjects in each of six OGTT-HbA_{1c} groups (as defined in Subjects, materials and methods) at baseline and at follow-up. Of those subjects with a normal OGTT at baseline; 93 had a normal and nine had an elevated HbA_{1c} (Table 1). At fol-

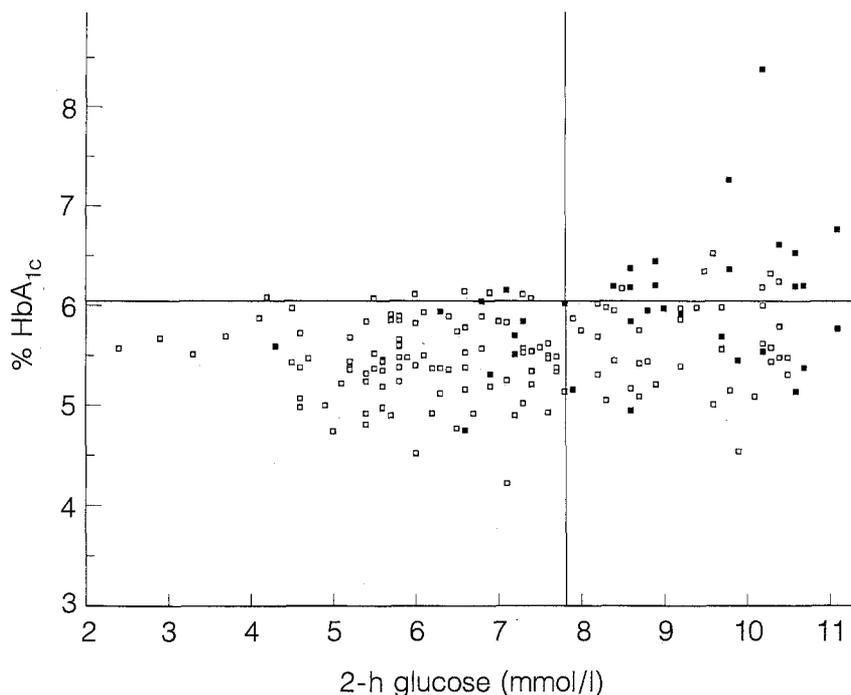


Fig. 1. Relationship between baseline 2-h glucose and HbA_{1c} in those subjects who were non-diabetic at baseline. The horizontal line indicates the upper limit of normal values for HbA_{1c} (6.05 %). The vertical line indicates the 2-h plasma glucose cutoff for IGT (7.8 mmol/l). □, subjects who were non-diabetic at follow-up; ■, subjects who were diabetic at follow-up

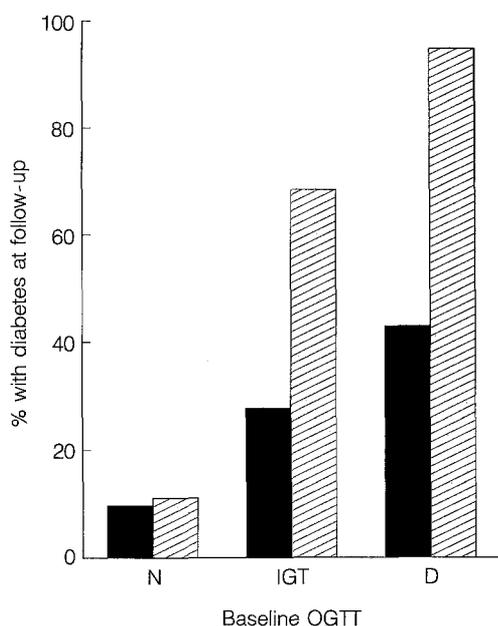


Fig. 2. Relationship between baseline OGTT/HbA_{1c} group and prevalence of diabetes by WHO criteria at follow-up. N, normal glucose tolerance; IGT, impaired glucose tolerance; D, diabetes; ■, normal HbA_{1c}; ▨, elevated HbA_{1c}

Table 2. Results of multiple logistic regression model in 66 subjects with IGT at baseline

Dependent variable: Diabetes		
Variable	Odds ratio	95 % confidence interval
HbA _{1c} ^a	6.76	1.77–25.8
Sex ^b	5.77	1.51–22.0

^a Odds ratio for a 1 % difference in HbA_{1c};

^b odds ratio for men to women

low-up, the two HbA_{1c} groups had similar rates of IGT (14.0 % in subjects with a normal HbA_{1c} and 11.1 % in those with an elevated HbA_{1c}) and of diabetes (9.7 % and 11.1 %, respectively). As shown graphically in Figure 1, it should be noted that all of the elevated HbA_{1c} values in this normal-OGTT group at baseline were only slightly elevated (6.07–6.15 % HbA_{1c}). At follow-up, those in the normal-OGTT/elevated HbA_{1c} group were also only slightly elevated (6.05–6.60 % HbA_{1c}).

Of subjects with IGT at baseline, 47 had a normal and 19 an elevated HbA_{1c} (Table 1). Among those with a normal HbA_{1c}, 27.7 % (13 of 47) progressed to diabetes during follow-up while among those with an elevated HbA_{1c}, 68.4 % (13 of 19) progressed to diabetes (chi-square = 7.8, *df* = 1, *p* < 0.01). Of subjects with IGT and elevated HbA_{1c}, in contrast to those with normal OGTT, HbA_{1c} values ranged widely from 6.17 to 8.37 % (Fig. 1). Prevalence of diabetes at follow-up is shown in Figure 2 by OGTT/HbA_{1c} group at baseline. In the 66 persons with IGT at baseline, the results of a multiple logistic regression analysis, with diabetes at follow-up as the binary outcome variable, are shown in Table 2. In this analysis, HbA_{1c} was used as a continuous variable. With both HbA_{1c} and 2-h post-load plasma glucose in the model, HbA_{1c} was a significant predictor of subsequent diabetes while 2-h glucose added little to the model. With sex and HbA_{1c} in the model, 2-h post-load plasma glucose concentration, age, body mass index, and the time between baseline and follow-up exams did not add to the predictive ability of the model. Controlled for sex, HbA_{1c} was still a significant predictor for the development of diabetes (odds ratio for having a HbA_{1c} concentration 1 % higher = 6.76, Table 2). Interestingly there was a significantly greater proportion of males than females that progressed to diabetes.

Of subjects with diabetes at baseline, 14 had a normal and 75 an elevated HbA_{1c} (Table 1). At follow-up, 57% (8 of 14) of those with a normal HbA_{1c} at baseline had a 2-h post-load plasma glucose concentration below 11.1 mmol/l including two (14.3%) with a glucose below 7.8 mmol/l. Less than one-half of these subjects would be classified as diabetic by OGTT at follow-up (43%, Figs. 1, 2). By contrast, only 5.3% (4 of 75) of those with an elevated baseline HbA_{1c} had a 2-h glucose below 11.1 mmol/l (Fig. 2); almost 95% had an OGTT in the diabetic range at follow-up ($p < 0.001$, two-tailed Fisher test).

Discussion

Previous studies have examined the relationship between glycosylated Hb and OGTT [1–3, 13–21]. Many of these studies have shown a good correlation between glycosylated Hb and either fasting or 2-h plasma glucose levels, but there has been considerable overlap of glycosylated Hb values in each OGTT category [3, 14, 19, 22, 23]. In our previous cross-sectional study of 381 subjects from this population of mostly Pima Indians, glycosylated Hb (HbA_{1c}) was highly specific and moderately sensitive in identifying subjects with diabetes (as diagnosed by OGTT). Diabetic subjects with normal HbA_{1c} had much lower fasting and 2-h post-load plasma glucose values than diabetic subjects with elevated HbA_{1c}. Subjects with elevated HbA_{1c} and IGT had significantly higher fasting glucose values than the normal HbA_{1c} group with IGT [4]. One explanation for the fact that a sub-group of diabetic subjects have normal glycosylated Hb levels is that large post-load (or post-meal) excursions (e.g. causing 2-h glucose values to fall in the diabetic range) have a minimal effect on the level of glycosylated Hb unless there is sustained hyperglycaemia.

In the present study, at follow-up there were fewer subjects who had either normal glucose tolerance and elevated HbA_{1c} or diabetes and normal HbA_{1c} than at baseline (15.1 vs 8.9% of subjects). These discrepant groups usually included borderline values; that is, normal glucose tolerance with HbA_{1c} values just outside the upper normal limit or glucose tolerance in the diabetic range with normal fasting glucose and 2-h glucose values only slightly above the 11.1 mmol/l cut-off for diagnosis of diabetes. As in our earlier study there were excellent correlations between HbA_{1c} and fasting and 2-h plasma glucose values both at baseline ($r = 0.91$, $r = 0.88$, respectively) and at follow-up ($r = 0.92$, $r = 0.91$, respectively). Because of the high prevalence of diabetes in this sample, this correlation may be higher than that observed in other studies.

Among subjects with a normal OGTT at the baseline examination, HbA_{1c} (when dichotomized into two groups) had no predictive value for the development of either IGT or diabetes. This finding is supported by the earlier findings of Modan et al. [9]. How-

ever, among subjects with IGT, those with a high HbA_{1c} were at a significantly higher risk of developing diabetes during the follow-up than were those with a low HbA_{1c}. Likewise, among subjects who already had diabetes at baseline, HbA_{1c} was significantly associated with the subsequent 2-h glucose. Screening characteristics might vary in populations with different plasma glucose frequency distributions. However, among subjects with IGT, glycosylated Hb may help determine who will progress to diabetes.

Acknowledgements. We thank the members of the Gila River Indian Community for cooperation and participation and the staff of the Diabetes and Arthritis Epidemiology Section, NIDDK for conducting the examinations.

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