

# The Classification of Diabetes by Clinical and C-Peptide Criteria

## A prospective population-based study

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**OBJECTIVE** — To evaluate both the concordance in the classification of diabetes by clinical and C-peptide criteria and, prospectively, the consistency of the classification by C-peptide.

**RESEARCH DESIGN AND METHODS** — Individuals with diabetes who were enlisted in the prospective epidemiological study of diabetic neuropathy (Rochester Diabetic Neuropathy Study [RDNS]) were classified clinically by National Diabetes Data Group (NDDG) criteria to IDDM and NIDDM at entry to the study. In addition, C-peptide response to 1 mg glucagon was measured at entry for the classification to IDDM (basal C-peptide,  $<0.17$  pmol/ml; increment above basal,  $<0.07$  pmol/ml) and NIDDM (all other responses) and for concordance with the clinical classification made. The consistency of the C-peptide response was assessed every 2 years for up to 8 years.

**RESULTS** — Among 346 individuals with diabetes, 84 were classified as IDDM and 262 as NIDDM by clinical algorithm. Concordance with the C-peptide response occurred in 89% of the patients and remained consistent during 8 years of follow-up. Among the 37 patients with discordant clinical and C-peptide classification, those considered clinically to have NIDDM had a consistent IDDM C-peptide response during follow-up, and most of those considered to have IDDM clinically eventually showed an IDDM C-peptide response during follow-up.

**CONCLUSIONS** — Clinical criteria for the classification of diabetes are highly correlated with the assessment of insulin secretory reserve. A small number of individuals considered to have NIDDM clinically or by C-peptide have or develop an IDDM peptide response.

The two major classes of diabetes, IDDM and NIDDM, were characterized by the National Diabetes Data Group (NDDG) primarily by clinical features (1). The qualitatively expressed discriminant relating to insulin secretory reserve (insulinopenia in IDDM or preserved insulin secretion in NIDDM) (1) has, over succeeding years, evolved to a quantitative corroboration of these classes on the basis of C-peptide concentrations. Various criteria for basal and stimulated C-peptide concentrations have been reported for the

classification of diabetes (2–21). There are scant epidemiological data regarding the concordance of classification by clinical and C-peptide criteria (13,15,19–21) or the prospective assessment of the consistency of C-peptide concentrations, especially in a population-based study (21). We had the opportunity to address these issues in the course of determining the epidemiological and demographic features of neuropathy in persons with diabetes under the aegis of the Rochester Diabetic Neuropathy Study (RDNS) (22).

### RESEARCH DESIGN AND METHODS

— Using the unique database that is available from the common medical record system of the Mayo Clinic and accessing the medical records of other medical care providers in the area who serve the local population, we identified all Rochester residents known to have diabetes and living within the geographic boundaries of the city on 1 January 1986. Residents who developed diabetes after this date were excluded.

Diabetes was confirmed by the satisfaction of NDDG criteria (1) and its classification was accomplished by the application of a clinical algorithm based on NDDG criteria (Fig. 1) (1).

The classification to IDDM and NIDDM by the clinical algorithm was compared with a classification based on basal and stimulated (6 minutes after 1 mg i.v. glucagon) C-peptide concentrations. C-peptide was measured by radioimmunoassay (23). Using previously published criteria for the characterization of IDDM and NIDDM as a guide, we arbitrarily segregated basal and the increment above basal C-peptide into three responses:  $<0.07$ ,  $0.07$ – $0.17$ , and  $>0.17$  pmol/ml. The various permutations and combinations of responses are shown in Table 1. Basal and stimulated C-peptide responses were assessed biannually over the succeeding years of follow-up.

### RESULTS

#### At entry

Of the total cohort of 381 patients enrolled in the RDNS, 346 had C-peptide responses to intravenous glucagon measured at entry to the study: 227 at year 2, 207 at year 4, 153 at year 6, and 35 at year 8. The clinical features of these patients classified by the clinical algorithm at entry to the study are shown in Table 2. Among the patients who had C-peptide measured at entry, 84 were classified by the clinical algorithm as IDDM and 262 as NIDDM. The concordance in classification between the clinical algorithm and C-peptide occurred in 67 IDDM and 242 NIDDM. Twenty patients

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DCCT, Diabetes Control and Complications Trial; NDDG, National Diabetes Data Group; RDNS, Rochester Diabetic Neuropathy Study.



Table 3—The classification by clinical algorithm

C-peptide pattern	IDDM (n = 84)		NIDDM (n = 262)	
	n	Percentage	n	Percentage
A	57	68	14	5
B	10	12	6	2
Subtotal	67	80	20	7
C	0	0	0	0
D	2	2	204	78
F	7	8	5	2
G	0	0	0	0
H	3	4	13	5
I	3	4	12	5
Subtotal	17	20	242	93
Total	84	100	262	100

Data are n or %. Responses A and B were considered to be indicative of IDDM; all other patterns of response were considered to be compatible with NIDDM.

**CONCLUSIONS**— Our selection of C-peptide criteria for the classification of diabetes, although arbitrary, was based on an interpretation of the literature extant about a decade ago when the present study was being designed. Our choice of basal C-peptide  $<0.17$  pmol/ml for IDDM is consistent with the Diabetes Control and Complications Trial (DCCT) criterion of  $<0.2$  pmol/ml (24) and that of the VA Cooperative Study of NIDDM, which used an exclusion criterion of  $<0.21$  pmol/ml (25). Unlike the DCCT, in which two absolute levels of stimulated C-peptide (primary prevention,  $<0.5$  pmol/ml; secondary intervention,  $<0.2$  pmol/ml) were used, and the VA Cooperative Study where the response to  $\beta$ -cell stimulus was not assessed, we reasoned that the ability to augment C-peptide concentrations would be a preferred way to assess  $\beta$ -cell reserve. Consequently, we examined the increment above baseline C-peptide as  $<0.07$ ,  $0.07$ – $0.17$ , and  $>0.17$  pmol/ml. Other investigators who have taken a diametrically opposite approach to ones in which C-peptide concentrations (whether basal or stimulated) were the independent variable and clinical characteristics the dependent variable observed  $\sim 90\%$  accuracy when cutoff values of C-peptide of  $0.16$  pmol/ml (13,20) and  $0.08$  pmol/ml (19) were used to classify IDDM and NIDDM. These criteria are remarkably close to the ones reported here.

At entry to the study, the concordance between the two systems of classification, clinical algorithm and C-peptide, occurred in 89% of the 346 patients. There was a

high degree of consistency of the C-peptide response over the subsequent years of follow-up, up to 8 years in some patients. No IDDM patient showed a sustained NIDDM-type C-peptide response, and only one NIDDM patient showed a sustained IDDM-type C-peptide response. Of the patients classified as NIDDM by the clinical algorithm, 14% showed an IDDM C-peptide response on at least one occasion, which suggests failing  $\beta$ -cell reserve but not to the IDDM level. Repeated fasting C-peptide concentrations after a 12-month interval in 215 insulin-treated individuals with diabetes on the island of Falster, Denmark, showed a high degree of consistency of response when the initial C-peptide concentration was  $<0.2$  pmol/ml (21).

The discordance in classification among 20 patients allocated to NIDDM by the clinical algorithm, all of whom had IDDM-type C-peptide responses at entry and almost consistently during follow-up, is not readily explained. At entry to the study, all were treated with insulin. Since four patients had a duration of diabetes  $<5$  years at entry to the study, the prolonged duration with a concomitant waning  $\beta$ -cell function cannot be the sole mechanism. It appears that one or more characteristics of the clinical algorithm misdirected the classification in these patients. The initial treatment with diet or sulfonylureas in 12 patients and obesity in 2 additional patients may have been misleading. Among the 17 patients classified as IDDM by clinical algorithm but NIDDM by C-peptide response, only 2 patients showed consistent NIDDM-type C-peptide responses during follow-up:

$>0.17$  pmol/ml basal and increment above basal. The remainder converted to consistent IDDM responses or showed variable responses during follow-up. The majority of these patients, therefore, could be considered to have IDDM in evolution. Although these patients had an NIDDM-type C-peptide response at entry, most had a blunted increment of C-peptide above basal at entry to the study. A limited response of C-peptide (e.g.,  $<0.17$  pmol/ml), regardless of the basal concentration of C-peptide, may be a harbinger of IDDM.

Are there clinical characteristics that suggest a measurement of insulin secretory reserve to classify a patient properly? This study does not identify any. One could argue that a  $\sim 90\%$  concordance between the clinical and C-peptide criteria for classification is better than might be expected.

We offer the following conclusions: 1) the clinical criteria for the classification of diabetes, although arbitrary and sometimes difficult to apply, nevertheless show a high degree of correlation with the assessment of insulin secretory reserve; 2) the clinical criteria for the classification of diabetes appear to be better at predicting IDDM (once classified as IDDM, it can be confirmed immediately or later by the assessment of insulin secretory reserve); and 3) the clinical criteria for NIDDM are good, but one should allow for the inclusion of some IDDM patients ( $\sim 10\%$ ), which is consistent with the studies that show  $\sim 10\%$  ICA-positivity in individuals who are considered clinically to be NIDDM.

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