

Originals

Evidence that the reduced number of natural killer cells in Type 1 (insulin-dependent) diabetes may be genetically determined

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Summary. Viruses may cause Type 1 (insulin-dependent) diabetes. We wondered whether the number and function of natural killer cells, which are important in anti-viral defense, are disturbed in diabetic patients. We studied 16 recently diagnosed Type 1 diabetic patients, 18 Type 1 diabetic patients diagnosed more than 15 years previously, 18 Type 2 (non-insulin-dependent) diabetic patients and 23 control subjects. We determined the number of natural killer cells (expressed as $\log_{10}\%$) using anti-Leu 11 monoclonal antibody and the function (in \log_{10} lytic units) concurrently using a ⁵¹Cr release assay with K562 as target cells. We found that the number of natural killer cells was reduced in Type 1 diabetes (1.01 ± 0.04) as compared with Type 2 diabetic patients (1.16 ± 0.04 , $p=0.004$) and normal control subjects (1.16 ± 0.04 , $p=0.006$). To establish whether the reduced natural killer

cell number is genetically determined we studied 19 identical twin pairs discordant for Type 1 diabetes; we found that even the non-diabetic co-twins had a reduced natural killer cell number (0.93 ± 0.05 , $p=0.0006$) as compared with normal control subjects. Natural killer cell function was similar in all groups while natural killer activity per cell was significantly increased in the recently diagnosed diabetic patients (1.63 ± 0.07) as compared with long-standing diabetic patients (1.26 ± 0.26 , $p=0.03$) and controls subjects (1.36 ± 0.07 , $p=0.006$). In conclusion the reduced number of natural killer cells in Type 1 diabetes appears to be genetically determined while their activity at diagnosis is increased.

Key words: Natural killer cells, identical twins.

The aim of the present study was to investigate the number and cytotoxic function of natural killer (NK) cells in Type 1 (insulin-dependent) diabetes. The immune system has been implicated in the pathogenesis of Type 1 diabetes [1] since recently diagnosed Type 1 diabetic patients have islet cell antibodies [2, 3], increased levels of activated T-lymphocytes [4, 5] and mononuclear cell infiltration of their pancreatic islets [6]. These immune changes are likely to result from environmental factors, possibly viruses [7, 8], acting on a genetically susceptible individual.

Natural killer cells are primary effectors of antiviral defense and lyse virus-infected cells [9, 10]. These cells can be accurately identified by monoclonal antibodies such as B73.1 and anti-Leu 11 which recognise phenotypes present on cells with NK function [11, 12]. The cytotoxic function of NK cells can be assessed by culturing NK cells with suitable target cells [13]. Previous studies which attempted to identify NK cells in Type 1 diabetes used techniques which are no longer considered valid [14]. Thus, neither anti-Leu 7 monoclonal antibody [15, 16] nor low affinity sheep erythrocyte ro-

setting [17] correctly characterise NK cells. Anti-Leu 7 antibody recognises only a terminal stage of NK cell maturation [18] while low affinity rosetting does not specifically recognise cells with NK function [19, 20].

Therefore, we studied NK cell number, using the monoclonal antibody anti-Leu 11, and NK cell function in Type 1 diabetic patients. In order to establish whether any abnormality in NK cell number or function in diabetic patients is influenced by hyperglycaemia or genetic factors we also investigated identical twin pairs discordant for diabetes.

Subjects and methods

We studied: (1) Thirty-four Type 1 diabetic patients (18 males, 16 females, median age 38 years, range 2–78); 16 had been diagnosed within one year (median 1 day, range 1 day–9 months) 11 of them were recently diagnosed and the remaining 5 were between 5–9 months after diagnosis; 18 had been diagnosed for more than 15 years (median 22 years, range 15–50); (2) Eighteen Type 2 diabetic patients (8 males, 10 females, median age 64 years, range 44–77); (3) Nineteen identical twin pairs discordant for Type 1 diabetes

(9 males, 10 females, median age 36 years, range 11–75); 9 pairs were short term discordant (median discordance time 2 years, range 6 months–3 years) and the remainder were long term discordant (median discordance time 15 years, range 15–19 years). Twins of 18 of the 19 pairs were living apart. (4) Twenty-three normal healthy control subjects (14 males, 9 females, median age 32 years, range 14–70).

Preparation of lymphocytes

Peripheral blood mononuclear cells (PBMC) were isolated from heparinised venous blood by sedimentation on a Ficoll-hypaque gradient (Pharmacia, Uppsala, Sweden) [21], washed in Hank's balanced salt solution (Gibco, Paisley, Scotland) and resuspended in RPMI 1640 medium (Gibco) containing 2 mmol/l glutamine, 200 U/ml penicillin, 100 µg/ml streptomycin and supplemented with 10% foetal calf serum.

Detection of NK cells

PBMC were stained by direct immunofluorescence with fluorescein labelled monoclonal anti-Leu 11 (Becton & Dickinson, Monoclonal Center, Mountain View, Calif, USA) [12]. One-hundred microlitres of PBMC at 4×10^6 /ml concentration were incubated with saturating amounts of anti-Leu 11 for 30 min at 4°C. After washing, the percentage of positive cells was determined using a U.V. microscope (Polyvar Reichert-Jung Fluorescence Microscope, Vienna, Austria). At least 400 cells were counted by a single observer unaware of the clinical details. The results are expressed as the percentage of NK cells in the PBMC population.

Natural killer cytotoxicity assay

The myeloid cell line K562 grown in suspension was used as a NK-sensitive target [22]. The assay used was a standard short-term (4 h) chromium-release assay. Briefly, K562 target cells were labelled by incubation in $\text{Na}_2^{51}\text{CrO}_4$ for 1 h at 37°C, using $100 \mu\text{Ci}/10^7$ cells in a volume of 0.5 ml. After washing and resuspending to 1×10^5 cells/ml, 100 µl of K562 target cells were mixed in the wells of a plastic plate (Nunc Plastic, Roskilde, Denmark) with 100 µl of PBMC at varying concentrations to give effector:target cells ratios of 40:1, 20:1, 10:1. Quadruplicates were performed at each dilution. After incubation for 4 h in 5% CO_2 at 37°C, 100 µl supernatants were harvested for counting in a gamma counter (LKB, 1260 MultiGamma, Bromma, Sweden). Spontaneous release of ^{51}Cr was measured in wells containing unlabelled K562 target cells as effector. Maximum release was measured by incubation of target cells in 10% Triton-100.

%NK activity =

$$\frac{\text{experimental } ^{51}\text{Cr release} - \text{spontaneous } ^{51}\text{Cr release}}{\text{maximum } ^{51}\text{Cr release} - \text{spontaneous } ^{51}\text{Cr release}} \times 100$$

Spontaneous release was between 7–13% of total labelled in all experiments.

Cytotoxic function was expressed as lytic units (LU) per 10^7 PBMC. A LU was defined as the number of PBMC needed to effect 33% cytotoxicity. LU were calculated from the cytotoxicity curve for each test [19].

The NK cell function was related to the frequencies of NK cells by dividing the $\text{LU}/10^6$ PBMC cells by the number of Leu 11 positive cells ($\text{LU}/10^6/\text{Leu 11}$) [23].

Interferon effect

A fraction of PBMC from each patient was preincubated with alpha interferon (INF) (Wellcome Diagnostics, Beckenham, England) at a concentration of 10^3 IU/ml for 1 h at 37°C. INF normally enhances the cytotoxic activity of NK cells [24]. These cells were used in the NK cytotoxicity assay. Another fraction of cells was incubated in the

same way in the absence of INF and served as control. Both the numbers of Leu 11 positive NK cells and their natural killing function were assessed from samples taken at the same time.

Statistical analysis

Results are expressed as mean \pm SEM. All data have been \log_{10} -transformed because of the marked variation in the level of NK cell number and function among individuals and to stabilise the variation around the \log_{10} -transformed mean. Following \log_{10} transformation the results corresponded to a normal distribution in that 66% of them were within 1SD of the mean. For comparison of NK cell number, function, and function per single cell among groups, the geometric means of each group were compared by subjecting the \log_{10} -transformed values of individual observations to two-tailed Student's t-test. To assess whether observations were correlated, we used the correlation coefficient r .

Results

NK cell number

Of the peripheral blood mononuclear cells the mean (\log_{10} -transformed) percentage of NK cells was significantly lower in Type 1 diabetic patients (1.01 ± 0.04) than in Type 2 diabetic patients (1.16 ± 0.04 , $p=0.004$) and normal control subjects (1.16 ± 0.04 , $p=0.006$) (Fig. 1). The mean levels were similar in recently diagnosed (0.97 ± 0.05) and long-standing diabetic patients (1.05 ± 0.04 , $p=0.13$). Of the six recently diagnosed diabetic patients tested repeatedly, four had levels less than the normal range (less than 0.89) which remained low on all 11 occasions they were tested, while two had

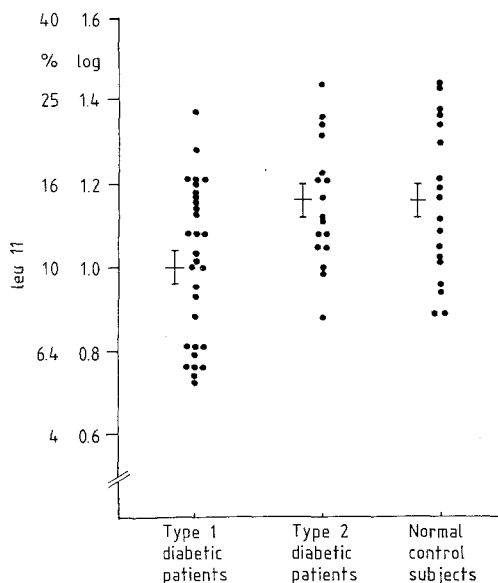


Fig. 1. Number of natural killer cells expressed as percentage and \log_{10} -transformed values in Type 1 diabetic patients, Type 2 diabetic patients and normal control subjects. Horizontal bars represent mean \pm SEM percent of the natural killer cell number. This is significantly lower in Type 1 patients when compared to Type 2 patients ($p=0.004$) and normal control subjects ($p=0.006$)

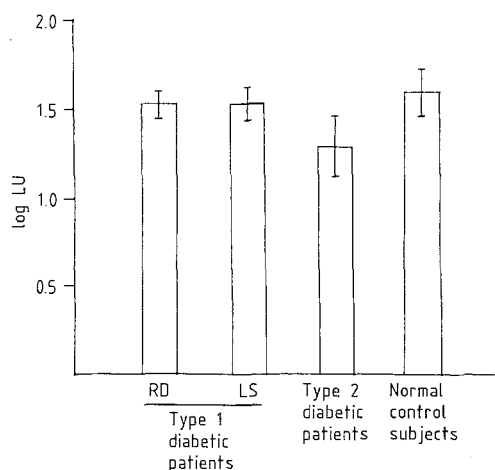


Fig. 2. Mean \pm SEM natural killer cell function expressed as \log_{10} -transformed lytic units (LU) in patients with recently diagnosed (RD) Type 1 diabetes, long-standing (LS) Type 1 diabetes, Type 2 diabetes and normal control subjects

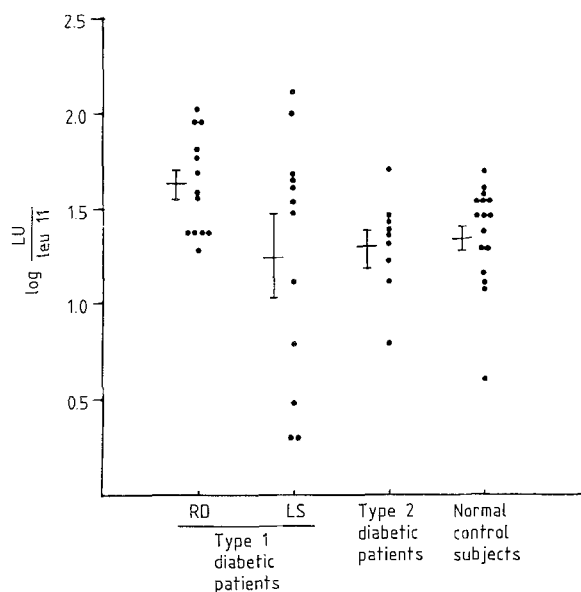


Fig. 3. Natural killer cell function per single cell expressed as a ratio between lytic units (LU) and the number of Leu 11 positive cells (\log_{10} -transformed) in patients with recently diagnosed (RD) Type 1 diabetes, long-standing (LS) Type 1 diabetes, Type 2 diabetes and normal control subjects. Horizontal bars represent mean \pm SEM natural killer cell function per single cell. This is significantly higher in recently diagnosed Type 1 patients than long-standing Type 1 patients ($p=0.03$), Type 2 patients ($p=0.005$) and normal control subjects ($p=0.006$)

normal levels which remained normal on all six occasions they were tested. The percentage of NK cell number did not correlate with the age of the patients ($r=0.15$).

NK cell function

Mean NK cell function of peripheral blood mononuclear cells (expressed as the \log_{10} -transformed lytic

units per 10^7 peripheral blood mononuclear cells) was not significantly different in recently diagnosed (1.54 ± 0.07 , $p=0.4$), long-standing Type 1 diabetic patients (1.54 ± 0.09 , $p=0.4$) and Type 2 diabetic patients (1.29 ± 0.16 , $p=0.08$) when compared to each other or normal control subjects (1.60 ± 0.13) (Fig. 2). The percentage change in NK cell function after alpha-interferon did not differ significantly from control subjects (20%) in recently diagnosed Type 1 diabetic patients (22%, $p=0.45$), long-standing Type 1 diabetic patients (26%, $p=0.23$) and Type 2 diabetic patients (28%, $p=0.13$).

NK function per cell

Mean NK function per single Leu 11 positive cell (\log_{10} -transformed) was significantly higher in recently diagnosed Type 1 diabetic patients (1.63 ± 0.07) than in long-standing Type 1 diabetic patients (1.26 ± 0.26 , $p=0.03$), Type 2 diabetic patients (1.31 ± 0.09 , $p=0.005$) and normal control subjects (1.36 ± 0.07 , $p=0.006$) (Fig. 3). Of the five recently diagnosed Type 1 diabetic patients tested repeatedly, one had levels above the normal range (greater than 1.7) on both occasions while the remainder had normal levels on 11 out of 12 occasions (one had a high level in one of four tests).

NK cell number and function

NK cell number and function of Leu 11 positive cells were significantly correlated in normal control subjects ($r=0.69$, $p=0.01$) but in neither Type 1 diabetic patients ($r=0.28$) nor in Type 2 diabetic patients ($r=0.02$).

Identical twin pairs

NK cell number

In the 19 identical twin pairs the percentage of NK cells was significantly reduced in both diabetic (0.91 ± 0.05 , $p=0.0005$) and non-diabetic twins (0.93 ± 0.05 , $p=0.0006$) when compared with normal control subjects (1.16 ± 0.04). The number of NK cells was correlated in the diabetic twin and non-diabetic co-twins ($r=0.66$, $p=0.004$) (Fig. 4).

NK cell function

NK cell function did not differ in the diabetic (1.49 ± 0.09 , $p=0.24$) and non-diabetic (1.43 ± 0.11 , $p=0.13$) twins from normal control subjects (1.60 ± 0.13). There was no significant correlation of NK function between the diabetic patients and their non-diabetic co-twins ($r=0.06$, $p=0.84$).

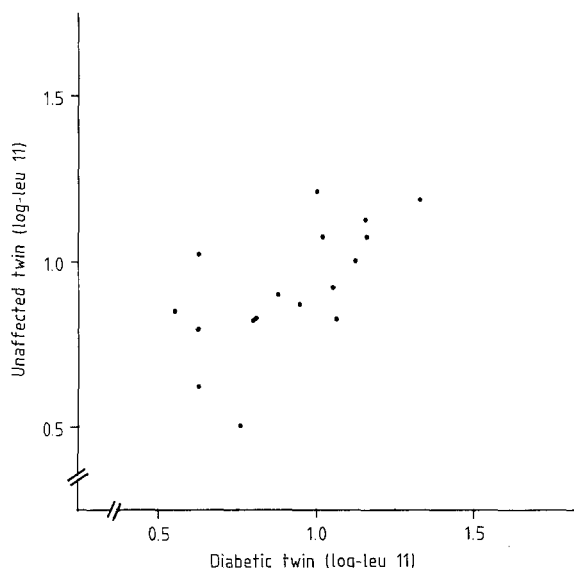


Fig. 4. Correlation for the number of natural killer cells, expressed as \log_{10} -transformed percentage of Leu 11 positive cells, between diabetic twin and non-diabetic co-twin ($r=0.66$, $p=0.004$). The 95% confidence interval for the correlation coefficient is 0.26–0.87

NK function per cell

NK cell function per single cell did not differ in the diabetic (1.41 ± 0.11 , $p=0.32$) and non-diabetic (1.49 ± 0.1 , $p=0.13$) twins when compared with normal control subjects (1.36 ± 0.07). The NK cell function per single cell in the diabetic twins correlated with that in their non-diabetic co-twins ($r=0.65$, $p=0.01$).

None of the diabetic twins was tested within 1 year of their diagnosis so we have no group of recently diagnosed diabetic twins to compare with the recently diagnosed diabetic singletons.

Discussion

This study demonstrates that Type 1 diabetic patients tend to have reduced numbers of natural killer cells. This reduction cannot be due solely to hyperglycaemia, since no change is found in Type 2 diabetic patients. We believe that the reduction in NK cell number is genetically determined because reduced numbers of these cells are found in both diabetic and non-diabetic identical twins, and, in them, the levels correlate significantly. In addition, the decrease in NK cells occurred irrespective of the duration of the disease.

In contrast to NK cell number, the overall NK cell function and response to interferon is normal in both groups of diabetic patients. It is likely that only large disturbances in NK cell function could be detected with the assay used in this study. These results are in apparent contrast to a recent report of decreased NK cell function in Type 1 diabetic patients [16]. However, in that study the population of cells used to assess NK function was depleted of T-lymphocytes which sub-

stantially contribute to NK function [19]. In our present study all cells with NK function were investigated. One might anticipate an increase in the NK cell function per single cell in Type 1 diabetes, since, in spite of the reduction in the NK cell number, the NK cell function was normal. However, we found that only the recently diagnosed Type 1 diabetic patients had an increase in NK cell function per single cell.

We do not know why an NK cell has increased function at, or soon after the diagnosis of Type 1 diabetes. Enhanced NK cell function can occur following viral infection or exposure to soluble factors released by activated T-lymphocytes [25, 26]. Recently diagnosed Type 1 diabetic patients show increased levels of activated T-lymphocytes [4, 5]. Thus, enhanced NK cell function in recently diagnosed Type 1 diabetes might be due either to a response to soluble factors released from activated T-lymphocytes or a factor which itself causes activation of T-lymphocytes.

In conclusion NK cell numbers tend to be reduced in Type 1 diabetes and this feature may be genetically determined. On the other hand, the NK function of each cell is increased in recently diagnosed Type 1 diabetes when there is also activation of T-lymphocytes.

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