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# Importance of Exercise in the Control of Metabolic and Inflammatory Parameters at the Moment of Onset in Type 1 Diabetic Subjects

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## Key words

- C-peptide
- $\beta$ -cell mass
- glycosylated haemoglobin
- inflammation

## Abstract

The onset of type 1 diabetes coincides with the final phase of  $\beta$ -cell destruction. In some cases, this period is characterized by the presence of a functional reserve of  $\beta$ -cells, favouring an adequate metabolic control (honeymoon phase). Therefore, the extension of this situation could have evident benefits in subsequent diabetes management. We aimed to study the influence of regular physical activity before and after the onset of the disease. We did an observational study of 2 groups of type 1 diabetic patients from onset to a 2-year period. One group (n=8) exercised regularly (5 or more hours/week) before onset and continued doing so with the same regularity. The second group (n=11) either did not perform physical activity or did so sporadically.

Circulating glycosylated haemoglobin (HbA<sub>1c</sub>), C-peptide, protein carbonyls and basal cytokine levels were determined at the beginning and at the end of the 1<sup>st</sup> and 2<sup>nd</sup> year. The more active group debuted with and maintained significantly lower HbA<sub>1c</sub> levels and insulin requirements compared to the more sedentary group. C-peptide levels were only significantly higher in the active group at the moment of onset compared to the sedentary group. In addition, determination of basal circulating cytokines revealed a large variability between individuals but no significant differences when comparing the groups. Altogether, the obtained results seem to indicate that physical activity allows a better control at the moment of onset regarding glycaemic control, residual endocrine pancreatic mass and subsequent insulin requirements.

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## Introduction

The clinical symptoms of type 1 diabetes coincide with the final phase of  $\beta$ -cell destruction. This stage presents islet inflammation or insulinitis, enhanced HLA (human leukocyte antigens) expression, presence of  $\beta$ -cell autoantibodies (antiGAD, antiIA-2, antiICA or anti-insulin among others) and  $\beta$ -cell dysfunction. Mononuclear cells infiltrate into the islet tissue, including CD4<sup>+</sup> and CD8<sup>+</sup> T-lymphocytes, macrophages and components of the complement system [1]. The high expression of HLA molecules and antigen transporter proteins indicates an enhanced antigen-presenting activity in the islet tissue, where the  $\beta$ -cell plays an instrumental role. Once the autoimmune process is activated by auto-reactive T-cells, the stimulation continues due to the presence of auto-antigens, leading to the expansion of effector cells and increasing the attack against  $\beta$ -cells [2].

Activated T-lymphocytes exist in circulation and in islet infiltrate. Under normal conditions, they are generally controlled by auto-tolerance mechanisms, however under diabetic conditions such control fails, leading to a cascade of inflammatory processes that culminate in insulinitis. The pathogenic immune response is mediated by Th1 lymphocytes and the protection by the Th2 subset. These populations are characterized by the production of different types of cytokines. Th1 subset secretes pro-inflammatory cytokines such as TNF- $\alpha$ , IFN- $\gamma$ , IL-1 $\beta$  and IL-12, while Th2 lymphocytes produce anti-inflammatory IL-4, IL-6 and IL-10 cytokines [3].

As previously mentioned, the destruction of pancreatic  $\beta$ -cells marks the initiation of the disease. However, in many cases a  $\beta$ -cell residual function exists at the moment of onset, allowing a good control of hyperglycaemia during a variable period of time. This transient phase is known as the honeymoon period. In this sense, several cohort studies and a retrospective analysis of the DCCT

(Diabetes Control and Complication Trial) demonstrated the benefits of preserving over years a residual  $\beta$ -cell function, monitored by C-peptide detection in circulation. These benefits are circumscribed to improved glycated haemoglobin ( $HbA_{1c}$ ) levels and reduction of severe hypoglycaemia crisis. Initial studies seem to indicate that this metabolic stability correlates with a preserved  $\beta$ -cell function [4]. This is of particular interest in certain moments of the patient's life, such as during adolescence, where the metabolic control of the disease is complicated due to the acquisition of new responsibilities and the lack of interest for certain topics, such as disease management. In addition, the DCCT indicates a low frequency to develop long-term complications, a reduced risk of ketoacidosis and better response to immunotherapy [4].

Therefore, the question is which strategies could help to debut with a significant residual  $\beta$ -cell mass at the moment of onset. Several pharmacological agents such as cyclosporine, azathioprine and prednisone were initially used to this end, but quickly discarded due to their secondary effects [5]. More recently, phase III trials using monoclonal antibodies against CD3 failed to retard the destruction of  $\beta$ -cells, although a single-dose regime yielded interesting results in young patients [6]. Other pharmacological compounds are still in the experimental phase. These include rituximab that acts by suppressing anti-insulin antibodies with no effect in other islet auto-antibodies [7]. The same rationale can be applied to anti-GAD antibodies, which act exclusively upon the GAD-expressing specific populations of  $\beta$ -cells [6]. Other experimental therapies include antibodies against CTLA4/CD152 which is a stimulatory molecule similar to CD28 that is expressed in T-helper cells [7]. Finally etanercept, a blocking agent of the TNF- $\alpha$  receptor, may affect other cellular responses in other organs where this cytokine is playing a signalling role [8]. Therefore, more research is necessary concerning the use of pharmacological agents.

On the other hand, several studies underline the metabolic and psychological advantages of physical exercise in the management of type 1 diabetes once the disease has been established and the onset period passed [9, 10]. However, there are no studies focused on the influence of physical activity at the moment of onset. With this objective, we have conducted an observational study in a group of patients at the moment of onset and during the subsequent 2 years. The group that regularly performed physical activity before, during and after the onset of diabetes presented low levels of  $HbA_{1c}$  as well as less exogenous insulin requirement. In addition, the more active group presented significant high circulating C-peptide levels only at the moment of onset. Values of protein carbonyl and circulating cytokines were not significantly different comparing both groups.

## Materials and Methods

### Subjects and study design

The key element of the pilot study design took in account that in the moment of onset, patients were asked about their habits on physical activity, being included in either Group A (for active) or Group S (for sedentary) (◉ Fig. 1). 19 subjects from 26 with recent onset of type 1 diabetes completed the 2 years study in the Endocrinology Service of the University Hospital Complex of A Coruña (Spain). The pilot study was registered in ClinicalTrials.gov (ID#: NCT01789190) and approved by the Autonomic Ethics Committee and met both local requirements as well as those stated in the Helsinki Declaration for research on human beings.

The inclusion criteria for group distribution took in account the principles of the American College of Sports Medicine, considering as active physical activity to practice a moderate-vigorous exercise during 1 h, 5 days or more/week. In this context, a sedentary or less active person should be a person that practices any or less than 5 h weekly [11]. Since this is an observational pilot study, we could not prescribe exercise nor intervene in the exercise routine of the volunteers. The information concerning the physical activity programmes was provided by each participant and recorded in the clinical story. Briefly, information concerning the type of exercise, training periods (hours/day), competition time (days/week), resting (days/week) and incidences (lesions, holidays, stages, trips, etc.) were recorded. Although we could not directly measure the intensity of exercise, nevertheless, each participant indicated its own perception about the intensity of work and this information was also included in the story. Therefore, group A (n=8) included patients that used to perform regular physical activity before onset and continued practicing afterwards with the same regularity. Group S (n=11) included patients that did not perform any physical activity at the moment of onset, continuing with the same habits during the subsequent observational period. 4 individuals of this group started to practice moderate exercise for a maximum of 3 days a week during sporadic moments of the observational period (less than 1 h each session). All individuals were informed at the beginning, in the presence of tutors in the case of minors, about the objective and justification of the study and signed a written consent. The informed consent included the following items in a total of 7 pages: Introduction (indicating the main idea of the study), Description (study design), Risks (concerning routine blood sample collection), Confidentiality (samples were codified to preserve anonymity), Benefits (there is no economic compensation to participate), Results (they will be submitted to scientific publication), Voluntary participation (participants can leave the study at any moment), Additional information (Spanish laws that regulate the study), Remaining samples (they will be custodied in the Hospital's Bank) and Letter of Consent (including data of each individual and signatures). The mean age of group A was  $21.9 \pm 4.2$  years and included 8 individuals, 5 men and 3 women diagnosed at the moment of onset by hyperglycaemia (n=2), hyperglycaemia+ketoacidosis (n=4) or ketoacidosis (n=2). The mean age of group S was  $23.3 \pm 2.2$  years and included 11 individuals, 7 men and 4 women diagnosed at the moment of onset by hyperglycaemia (n=1), hyperglycaemia+ketoacidosis (n=3) or ketoacidosis (n=7). A clinical story and physical exploration was also performed in the first consultation. Fasting blood samples were taken from the antecubital vein to determine  $HbA_{1c}$ , C-peptide, antiGAD, cytokines and protein carbonyls. Patients and tutors entered the hospital educational program in order to learn about diet management, glycaemia monitoring and insulin injections.

Consultations were initially performed on a weekly basis for the first month, then once a month for the remaining period. In each consultation, patients were asked about diet accomplishment, glycaemia monitoring, insulin injections and exercise performance. All individuals from group A declared to practice regularly intervallic routines, playing team sports (basketball for n=1 and soccer for n=6). Only one individual declared to perform aerobic exercise (long distance running and swimming). At the end of the first year a second blood sample was extracted in the same conditions from each individual to determine the same parameters. Exercise routine accomplishment and follow-up of diet

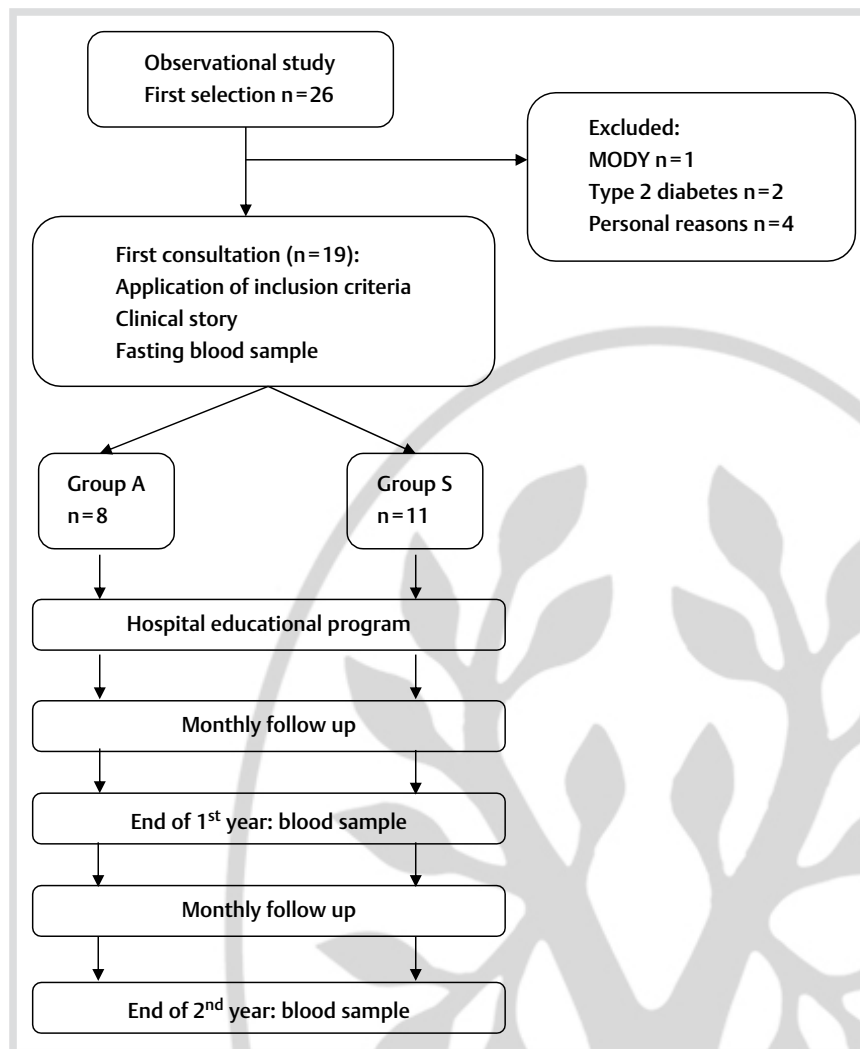


Fig. 1 Flow diagram for study design.

and insulin injections was followed monthly during one more year. At the end of this period, the pilot study was considered finished and the last blood extraction was performed in order to measure the same parameters. Other subjects were excluded because they were not capable to provide data required for study follow-up.

### Circulating HbA<sub>1c</sub>, C-peptide and protein carbonyl determinations

HbA<sub>1c</sub> was determined by immunoanalysis. C-peptide was determined by radioimmunoassay using the Coat-a-Count kit (Diagnosis Products Corporation, LA). Protein carbonyl derivatives were calculated by adapting the method developed by Levine (1994) [12].

### Circulating cytokine determinations

Cytokines were measured in plasma by Flow Cytometry (FACS-Calibur, BD Bioscience) through the Flowcytomix™ Multiplex Test (eBioscience) (n=3). The following cytokines were analysed for each patient: IL-1β, IL-2, IL-4, IL-5, IL-6, IL-9, IL-10, IL-12p70, IL-13, IL-17a, IL-22, TNF-α and INF-γ.

### Statistical analysis

Statistical analysis was performed using the SPSS-20 software for Windows. Normality testing was performed according to the Kolmogorov-Smirnov test (K-S test). Results in **Table 1** were

expressed as the mean ± sem and the values with a  $p < 0.05$  were considered statistically significant. The effect of physical activity on the changes induced in the different circulating parameters (HbA<sub>1c</sub>, C-peptide, protein carbonyls and cytokines) and injected insulin at each particular time was tested by Student t-test for independent samples. The evolution of the different cytokines into each particular group of individuals throughout time was analysed by Student t-test for paired samples. Although there is a limitation in the sample size (n=8 for group A and n=11 for group S), we have performed additional analyses such as a general lineal model (GLM) of repeated measures considering time as an intragroup factor and physical activity as an intergroup factor, and a GLM of repeated measures considering only the intragroup factor (time) and no intergroup factor. Correlation between different parameters was performed according to Pearson's correlation coefficient. We considered arbitrarily a low correlation at  $r = \pm 0-0.30$ , moderate at  $r = \pm 0.31-0.60$  and good at  $r = \pm 0.61-1$ .

### Results

**Table 1** shows the determinations at the moment of onset and after 1 and 2 years of the circulating parameters (HbA<sub>1c</sub>, C-peptide and protein carbonyls), insulin requirements and body mass index (BMI) of patients from both groups that completed the

**Table 1** Circulating values for HbA<sub>1c</sub>, insulin requirements, C-peptide, IDAA1C, protein carbonyls and BMI in the active (A) (n=8) and sedentary group (S) (n=11).

	Group A	Group S
<b>Time 0</b>		
HbA <sub>1c</sub> (%)	9.57 ± 3.47*	12.68 ± 2.83
injected insulin (IU/kg of weight)	0.48 ± 0.38	0.72 ± 0.27
total insulin injected (IU)	31.71 ± 9.74	44.67 ± 5.25
C-peptide (pmol/L)	1.09 ± 0.76*	0.37 ± 0.13
IDAA1C	11.50 ± 4.95	15.02 ± 3.03
protein carbonyls (μmol/L)	76.19 ± 40.62	75.64 ± 12.43
BMI (kg/m <sup>2</sup> )	20.80 ± 1.42	22.37 ± 0.71
<b>1<sup>st</sup> year</b>		
HbA <sub>1c</sub> (%)	6.12 ± 0.64*	7.96 ± 1.80
injected insulin (IU/kg of weight)	0.07 ± 0.04*	0.54 ± 0.25
total insulin injected (IU)	5.43 ± 1.29*	35.83 ± 5.15
C-peptide (pmol/L)	1.07 ± 0.68	0.67 ± 0.39
IDAA1C	6.18 ± 0.61*	10.23 ± 1.90
protein carbonyls (μmol/L)	80.73 ± 10.28	87.49 ± 19.25
BMI (kg/m <sup>2</sup> )	21.40 ± 1.33	24.41 ± 1.08
<b>2<sup>nd</sup> year</b>		
HbA <sub>1c</sub> (%)	6.20 ± 0.75*	8.50 ± 1.96
injected insulin (IU/kg of weight)	0.14 ± 0.07*	0.61 ± 0.29
total insulin injected (IU)	7.67 ± 1.94*	37.50 ± 6.53
C-peptide (pmol/L)	1.87 ± 1.58	0.67 ± 0.37
IDAA1C	6.72 ± 0.92*	10.96 ± 2.28
protein carbonyls (μmol/L)	88.48 ± 19.67	91.87 ± 12.56
BMI (kg/m <sup>2</sup> )	21.57 ± 1.25	24.06 ± 1.12

(\*) Significant values (p < 0.05) with respect to group S

study (n=19). Subjects that did not provide the necessary follow-up data were excluded from the study. HbA<sub>1c</sub> served as a marker for average glycaemia levels over long periods of time [13]. As it can be observed in Table 1, group A presented at the time of onset significantly lower values of HbA<sub>1c</sub> than group S. Interestingly, after the first year, group A presented HbA<sub>1c</sub> values similar to those recommended by the American Diabetes Association (ADA) (6.5%) for the reduction of vascular complications [14]. At the end of the first and second year, group S presented acceptable HbA<sub>1c</sub> values, but were significantly higher than those found in group A. However, the change observed in HbA<sub>1c</sub> levels in both groups was similar (around 4%).

Circulating C-peptide levels were determined in this study as an indicator of the existence of a residual pancreatic β-cell mass [15]. At the moment of onset, individuals from group A presented 3 times significantly higher values of circulating C-peptide than patients from group S (Table 1). Differences between both groups were not significant at the end of the first and second year, although the values of C-peptide were always higher in group A compared to group S. The concentration of C-peptide determined at the end of the study in both groups was approximately twice the ones determined at the moment of onset. However, C-peptide levels are insufficient for detecting functionality in residual β-cells. The IDAA1C (insulin dose-adjusted HbA<sub>1c</sub>) index has been proposed to define the partial remission period (honeymoon phase) and values of IDAA1C ≤ 9 strongly correlate with the existence of residual functional β-cells [16]. According to this, group A compared to group S displays IDAA1C values at the end of the first and second year that indicate the presence of functional endocrine pancreas (Table 1).

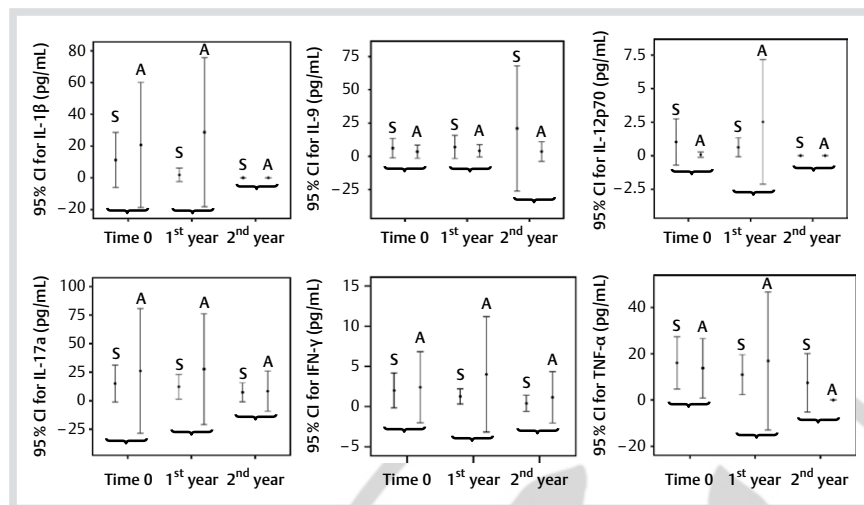
In the clinical practice, the insulin administered to each patient is strongly related, among other factors, to glycaemic control and to the presence of a residual pancreatic endocrine function.

According to data recorded in the clinical story of participants, we observed that individuals of group A required less insulin administration than subjects from group S, being significantly different at the end of the first and second year (Table 1). This is also reflected in the amount of total insulin units injected (Table 1). In addition, Pearson's correlation analysis revealed that the average amount of insulin injected measured in units/kg of body weight moderately correlated with HbA<sub>1c</sub> levels at the end of the first year (r=0.545) and very positively at the end of the second year (r=0.768) in group A. However, there was less correlation between insulin requirements and C-peptide levels, being r=-0.423 at the end of the first year and r=-0.334 at the end of the second year. Therefore, the adequate glycaemic control as reflected by low HbA<sub>1c</sub> levels seems to be the main determinant for reduced insulin requirements. Regular exercise could be instrumental to achieve this goal.

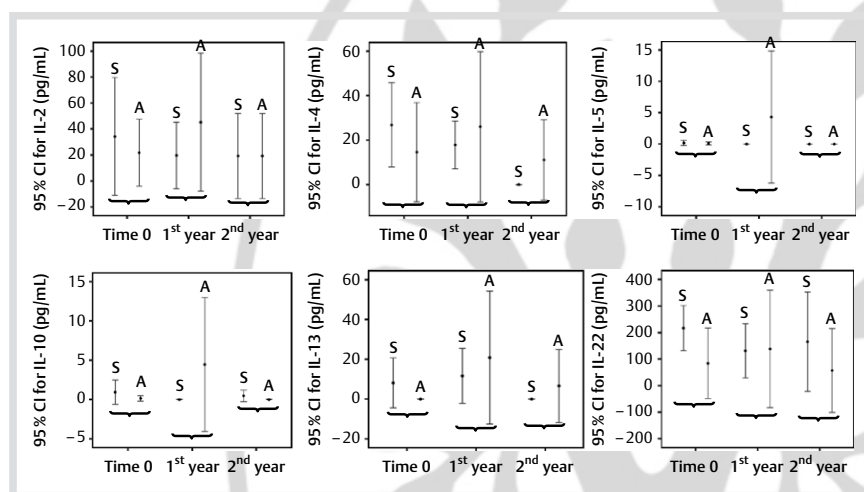
To verify this assumption, we applied the correlation analysis between exercise frequency and HbA<sub>1c</sub> levels, resulting in a moderate correlation at the moment of onset (r=-0.528) and at the end of the first year (r=-0.680), but good at the end of the second year (r=-0.798). As expected, correlation between exercise frequency and insulin requirements evolved positively, being r=-0.502 at the moment of onset, r=-0.810 at the end of the first year and r=-0.884 at the end of the second year. Nevertheless, correlation was moderate or low when comparing exercise frequency and C-peptide levels, being r=0.579 at the moment of onset, r=0.242 at the end of the first year and r=0.507 at the end of the second year.

In an attempt to better understand the evolution of the disease in both groups of patients, we determined the levels of plasmatic protein carbonyls. This election was based on the observation that diabetes is a disease that presents oxidative stress events, with sustained hyperglycaemia being one of the main pro-oxidant disruptors [17]. This provokes the glycation of haemoglobin and possibly of other body proteins, developing over time oxidative adducts, including protein carbonyls. Table 1 shows that there were no significant differences in the protein carbonyls of both groups, although group A tended to have slightly lower protein carbonyl values than group S. Finally, BMI did not present significant changes in both groups, although group A presented optimal values.

Altogether, these results seem to indicate that practicing regular physical activity before and after the onset gives rise to a positive evolution of different parameters, in particular HbA<sub>1c</sub> levels. The question that arises is if there are other parameters that could be modulated by regular exercise. To this end, we focused on the immunological and inflammatory response, since they play an instrumental role in the progression and severity of disease [18]. In particular, we determined the levels of GAD antibodies, one of the most frequent autoantibodies in type 1 diabetes [19]. Interestingly, not all individuals presented positive antiGAD in circulation. In group A, 3 of the 8 individuals presented antiGAD values lower than 1 U/mL, while in group S only one individual did not present antiGAD. For the rest of the individuals with positive antibody values, the antiGAD values varied greatly, ranging 2–34 U/mL in group A and 1.2–30 U/mL in group S. Individually, we observed that antiGAD values remained unchanged throughout the whole period studied. Only one individual of group A presented a decrease in circulating antiGAD ranging from 3.9 U/mL at the moment of onset to 2 U/mL at the end of the second year. In group S, 2 individuals presented a decrease, with 30 U/mL at the moment of onset and 6.4 U/mL at



**Fig. 2** Circulating cytokine ranges for sedentary (S) and active (A) individuals. Results are represented by error bars indicating 95% confidence interval (CI) based on Student t-test for paired data.



**Fig. 3** Circulating cytokine ranges for sedentary (S) and active (A) individuals. Results are represented by error bars indicating 95% confidence interval (CI) based on Student t-test for paired data.

the end of the second year and from 4.7 to 1.9U/mL for the second individual in the same period of time. Altogether, the obtained results do not allow us to establish a correlation between the frequency of exercise practice and the circulating levels of antiGAD. On the other hand, cytokines are involved in  $\beta$ -cell lesions in human and animal models of type 1 diabetes [3]. In this context, the second question that arises is if physical activity could modulate certain circulating cytokine levels. We measured a number of cytokines that have shown to be instrumental in the development of type 1 diabetes playing pro- as well anti-inflammatory roles [3,20]. When analyzed individually, we discovered that there was great variability among individuals, in other words, each individual presented their own cytokine profile. In addition, this profile changed in the same individual in response to a variety of external and internal stressors, being diabetes one of them.

Nevertheless, the analysis of the error bar representations revealed interesting tendencies (► **Fig. 2, 3**). At the moment of the onset of disease (Time 0), IL-4, IL-5, IL-9 and TNF- $\alpha$  presented a similar range of variability in individuals of group A and group S. At time 0, IL-1 $\beta$ , IL-17a, IL-22 and IFN- $\gamma$  levels were more varied in individuals of group A than group S. The opposite was observed with IL-2, IL-10, IL-13 and IL-12p70, which displayed wider variability in individuals of group S than group A. These patterns were maintained with more or less differences at the end of the first year for IL-1 $\beta$ , IL-9, IL-17a, IL-22 and IFN- $\gamma$ . However, at the end of the first year IL-2, IL-4, IL-5, IL-10, IL-13,

IL-12p70 and TNF- $\alpha$  levels were more varied in individuals of group A than group S. Finally at the end of the second year, IL-4, IL-13, IL-17a and IFN- $\gamma$  presented the same pattern observed at the end of the first year. At the end of the second year, IL-9, IL-10 and TNF- $\alpha$  presented a higher range of variability in individuals of group S compared with group A. A similar range of variability at the end of the second year was observed for IL-2 and IL-22, while IL-1 $\beta$ , IL-5 and IL-12p70 were not detected in either group. Basal IL-6 presented values under the detection limit of the technique in both groups during the study.

The analysis of evolution from time 0 to the first year revealed almost significant differences for specific cytokines, such as IL-4 in group A ( $p < 0.08$ ) and IL-22 in group S ( $p < 0.06$ ).

## Discussion

▼ The presented results seem to indicate that individuals who practice regular physical activity before the onset of type 1 diabetes initiate with a better metabolic profile than sedentary individuals, presenting significantly lower levels of HbA<sub>1c</sub> and higher C-peptide levels. In addition, individuals of group A present less ketoacidosis cases (2 from  $n=8$ ) than individuals of group S (7 from  $n=11$ ). We can speculate that exercise before onset helps to a better prognosis of the disease. In this context, it has been described that ketoacidosis seems to be associated

with a poor long-term metabolic regulation and residual  $\beta$ -cell mass [21]. In any case, additional experiments are necessary to verify this hypothesis.

The results also emphasize that the Hospital's educational program (which includes diet management and insulin injection monitoring) improved in a similar manner the management of the disease in both groups. In other words, during the 2 year period both groups presented a 4% decrease in HbA<sub>1c</sub> values and doubled circulating C-peptide levels. Since active individuals began the experimental period with a better circulating profile, this influenced at long term the functionality of residual  $\beta$ -cells as indicated by the values obtained for the IDAA1C index. These values indicate that active patients displayed an excellent metabolic control than sedentary, and that this event could be described as an extended honeymoon phase [16].

In this context, group A presented IDAA1C scores of 9 or less at the end of the first and second year. IDAA1C was used to define the honeymoon period, a moment that strongly correlates with the existence of residual functional  $\beta$ -cell mass [16]. Therefore we can hypothesize that the regular practice of physical activity before and after the onset of the disease allows individuals to enter in an optimal metabolic control period, similar in many aspects to the honeymoon phase mentioned before. This phase is characterised by presenting low levels of HbA<sub>1c</sub>, an indicator of correct glycaemia control, and therefore low requirements of exogenous insulin. The results of our study also emphasize that a sedentary lifestyle is not the best way to obtain certain metabolic advantages before the onset of the disease (compare groups A and S at time 0 in **Table 1**). In addition, diet and insulin injections are not sufficient to maintain an optimal metabolic control (i.e., individuals of group S) once the disease has been established, being the practice of regular physical activity the candidate factor to achieve both goals. Since this has been an observational study, we cannot provide data concerning frequency, intensity or type of activity that could be the most appropriate to prescribe to patients with type 1 diabetes. More studies are necessary to answer this particular point.

Since active individuals possessed better metabolic circulating parameters than the sedentary counterparts at the moment of onset, we could speculate that individuals of group A might be diagnosed later than sedentary ones. However, the average age of admission at the Endocrinology Unit was very similar in both groups. Concerning this point, it is important to mention that the active individuals of this study were practicing sports in a semi-professional level and for this reason submitted more often to routine blood and medical analysis, which was how the disease was detected. In any case, the severity of the disease seems to be a particular feature of each individual, as reflected by the large variability in values for antiGAD and the cytokines analysed. In other words, for a similar severity of the disease included in a similar environmental and genetic background, individuals with an active lifestyle would develop the disease later than sedentary ones. Unfortunately this did not occur neither in our study nor in the majority of the studies conducted at present. For this reason, this remains as a speculative point that needs very precise experimental designs for its study.

Several intervention and observational studies have analysed the impact of physical activity in diabetes control, but always once the pathology was established. In this respect, the changes in HbA<sub>1c</sub>, used as a marker of glycaemia management, observed in these studies were modest or not significant [9, 10]. In addition, we cannot compare these studies with ours due to several

key differences. First, in this study active individuals were performing exercise before the onset of the disease. Second, the exercise frequency to be considered active for an individual in our protocol was at least of 5 days/week and 1 h/day [11]. Regarding insulin requirements, both points seem to be instrumental in the subsequent management of the disease for individuals of group A (lower requirements) compared with those of group S, at least in a period of 2 years.

A similar tendency in individuals of group A was observed for protein carbonyls, however the differences with group S were not significant. We can hypothesize that hyperglycaemia in type 1 diabetes favoured the development of oxidative stress [17]. In this context, an adequate glycaemic control could minimize oxidative damage. Furthermore, it is known that physical activity allows the activation of genes coding for antioxidant enzymes, leading to a better antioxidant defence [22]. Therefore, the modulation of oxidative stress in diabetic patients that perform regular exercise remains to be studied in more detail.

On the other hand, exercise is capable of modulating cytokine production. However, these changes depend on intensity, type and duration of the exercise [23,24]. In this context, several studies have shown that specific routines can increase circulating levels of anti-inflammatory cytokines that can control the muscle inflammatory damage inherent to exercise [25]. However, this does not seem to be as simple in a chronic disorder such as type 1 diabetes (**Fig. 2, 3**). Furthermore, the complexity to interpret the results increases if we take in account that the majority of the cytokines are produced in response to a variety of environmental stressors, explaining the inter-individual variability. Otherwise said, it seems that each patient has its own particular cytokine circulating profile. Moreover, the majority of the data concerning the role of cytokines in the development of type 1 diabetes are from animal studies, where islet tissue can be directly analysed [26]. Therefore, our results must be interpreted with extreme caution.

Nevertheless and regarding this issue, some particular observations deserve further discussion. First of all, certain cytokines considered as pro-inflammatory in type 1 diabetes development were clearly detected in individuals of group A at the moment of onset, i.e., IL-1 $\beta$ , IL-17a, IFN- $\gamma$  and TNF- $\alpha$ , and even at the end of the first year (**Fig. 2**). These results could suggest that exercise has some influence in controlling the inflammatory response. However, at the end of the first year, when in addition to exercise the individuals control diet and insulin administration, we have observed that anti-inflammatory cytokines such as IL-2, IL-4, IL-5, IL-10 and IL-13 are largely detectable in individuals of group A compared with group S (**Fig. 3**). The pattern observed was not maintained at the end of the second year (**Fig. 2, 3**). This could be tentatively interpreted that over time the anti-inflammatory effects attributed to exercise practice is not sufficient to stall the pro-inflammatory progression of the disease.

IL-2 together with IL-21 is under the control of *Idd3*, a susceptible gene locus in diabetes development. Reduced levels of IL-2 in islets of NOD mice correlate with reduced function of Treg cells [27,28]. In this context, IL-2 helps to keep the ratio in favour of CD4<sup>+</sup>-Treg cells in swollen islets [29,30] and IL-4 avoids diabetes development by inducing Th2 cells in islet tissue [31]. In addition, IL-10 also plays a protective role in diabetes development [32], although some findings are discrepant [33].

On the other hand, IL-22, which is included in IL-10 family, is an interesting cytokine that deserves some attention. IL-22 upregulates the expression of *Reg* genes in islets. These genes code for

proteins such as Reg2 or Reg3 $\beta$  that are involved in  $\beta$ -cell regeneration [34] and are abundant in mouse pancreatic buds [35]. In addition, IL-22 receptor is expressed in pancreatic  $\beta$  and  $\alpha$ -cells, but not in exocrine acinar or ductal epithelium [36]. Moreover, IL-6, which is acutely secreted after exercise performance in diabetes patients and has a shorter half-life than IL-22 [37] is capable of also activating Reg proteins [38]. We can hypothesize that IL-22 is secreted when the endocrine pancreatic mass is reduced to a critical level in order to restore  $\beta$ -cell mass by activating Reg proteins. However, we can speculate that this objective is not reached due to the high presence of pro-inflammatory cytokines in islet tissue. Although, some anti-inflammatory cytokines are present to counteract the effect of the pro-inflammatory factors, at least during the first year, they cannot establish an optimal situation to restore  $\beta$ -cell mass. In any case, new experiments need to be designed in order to confirm these hypotheses. Finally, one limitation of our study concerns the low number of individuals studied. Nevertheless, the majority of the published studies present similar figures [9, 10]. In conclusion, this report suggests that adequate circulating parameters could be reached when physical activity is performed before the onset of type 1 diabetes, resulting in an optimal insulin administration in subsequent periods.

**Conflict of interest:** None.

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