



Interferon Gamma in Egyptian children with LN by Flow Cytometry.

KEYWORDS

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ABSTRACT Systemic lupus erythematosus (SLE) is a chronic, immune-mediated disease in which T and B cells cause systemic tissue injury. SLE is characterized by the production of high levels of auto-antibodies against nuclear antigens, resulting, at least in part, from a dysregulated T lymphocyte response to autoantigens. Results: the expression of $INF\gamma$ levels in T lymphocytes cells of blood samples obtained from LN group was significantly decreased as compared to healthy control (9.24 ± 3.6 and 14.05 ± 4.36 respectively; $P \leq 0.01$). In conclusion: the low expression of $INF\gamma$ in T lymphocyte cells of LN patients may be a good marker in prognosis and diagnosis of LN.

Introduction

Systemic lupus erythematosus (SLE) is a chronic, immune-mediated disease in which T and B cells cause systemic tissue injury [1]. T cell dysfunction has been described at the cellular and molecular level in both humans and animal models of lupus [2–4]. Impaired in vitro T cell proliferation and altered cytokine production (e.g., $INF\gamma$ production [5]) in response to foreign antigens, alloantigens (allogeneic major histocompatibility complex (MHC) molecules), and mitogens has been observed [4]. In general, it is thought that CD4 cell activity is exaggerated, while CD8 cell activity is diminished, resulting in an imbalanced T cell functional repertoire, a Th1/Th2 cytokine imbalance, and B cell hyperactivity [2]. However, it is not clear how functional and phenotypic aspects of antigen-specific CD8 cells compare in SLE. (B.R. Berner et al. / Cellular Immunology 235 (2005) 29–38)

A common and serious complication of SLE is lupus nephritis, which plays a key role in the prognosis of SLE and is a risk factor for the development of end-stage renal failure. It occurs in about 15% of patients at diagnosis and in approximately 40% during the course of the disease. Until now there are no general accepted methods to reliably identify patients who are at risk of flares. Disease activity is assessed using a combination of clinical history, physical examination, organ specific functional tests, and serologic studies. To date, determination of complement levels and anti-dsDNA antibodies are the serological parameters most commonly used to assess disease activity in patients with SLE. Numerous investigations of other immunological tests as indicators of disease activity have been reported (Cytokine 60 (2012) 410–416)

Systemic lupus erythematosus (SLE) is a disease characterized by the production of high levels of auto-antibodies against nuclear antigens, resulting, at least in part, from a dysregulated T lymphocyte response to autoantigens. (Kammer et al., 2002; Kong et al., 2003). In SLE, multiple serological factors get altered. These include cytokines, often representing altered ratio of Th1/Th2 and those released from different leukocytes; inflammatory peptides,

proteins and raised levels of circulating immune complexes. (V. Arora et al. / Molecular Immunology 44 (2007) 1722–1728)

Th1 type cytokines such as $INF\gamma$, IL-2 and $TNF\beta$ as well as Th2 type cytokines, such as IL-4, IL-6, IL-10 and IL-13 are known to be important regulators of the immune system. (Nagy et al., 2000)

The aim of our present work was to study the CD4, CD8 and $INF\gamma$ production of T lymphocytes by flow cytometric analysis using isolated lymphocytes from blood samples of pediatric patients with lupus nephritis and healthy donors.

2. Patients and methods

2.1. Patients and controls

The study cohort included 27 patients diagnosed as having lupus nephritis (% female and % male mean age 9.68 ± 2.89 years according to the American College of Rheumatology (ACR) criteria, Median (range) anti-dsDNA titers were 230 (3–1000 E/ml), median C3 and C4 were 0.55 g/l (0.05–1.03) and 0.14 g/l (0.04–0.30), and 11 healthy control subjects (% female and % male; mean age 9.98 ± 2.10 years) who had no history of autoimmune disease, infectious disease, malignancies or immunosuppressive therapy. The study was approved by Mansoura pediatric hospital, Mansoura University, and written informed consent was obtained from all participants.

Materials

Monoclonal antibodies (mAb) and flow cytometry.

The following mAb and reagents were used in this study: fluorescein isothiocyanate (FITC)-conjugated or peridinin chlorophyll A protein-conjugated-conjugated anti-CD8 mAb, FITC-conjugated anti-CD4 mAb, FITC-conjugated anti and anti $INF\gamma$ mAb,

EDTA-blood and fresh urine samples were collected from patients. Urine samples from patients with signs of urinary tract infection were excluded.

Sample preparation for flow cytometry

The fresh urine sample was collected from each patients and control. For 3 ml urine was centrifuged at 2000 rpm for 20 min. The supernatant was removed and the pellet which contains lymphocyte was shaken. The lymphocytes from urine sample was Fixed with ice cold absolute alcohol 1ml for each tube and was preserved in +4c forever until analysis.

Isolation of peripheral blood: Peripheral venous blood samples were collected into tubes containing EDTA, and peripheral blood mononuclear cells (PBMCs) were isolated by density-gradient centrifugation for 20 min at 1500 rpm without braking using Ficoll-Paque Plus solution. The bands of cell were pipette carefully into another centrifuge tube filled up with phosphate buffer saline (PBS) or hanks solution and mixed. After centrifugation for 10min at 1800 rpm, lymphocyte sediment were fixed with ice cold absolute alcohol 1ml and preserved in +4C° until analysis.

This technique was applicable where the fluorochrome was directly linked to the primary antibody e.g.: phycoerythrin (PE) or fluorescein isothicyanate (FITC)- conjugate. 200 µl of lymphocytes were added in test tubes. The lymphocytes were washed with 1ml PBS, and then were centrifuged at 200 rpm for 5 minutes, the supernatant was discarded. The lymphocytes pellet was washed with PBS and mixed well. 3µl of required marker was added and was mixed well. The tubes were incubated at room temperature for 20 minutes. The cells were washed with 1ml PBS; then the tubes were centrifuged at 2000 rpm for 5 minutes. The supernatant was removed then 200ml of 4% paraformaldehyde were added for analysis by flow cytometer.

Flow cytometric analysis was performed with FACCalibur flow cytometer using Cell Quest software (Becton Dickinson, San Jose, CA). A total o10,000 to 20,000events were collected for each analysis. Lymphocyte gates were set on live cells using forward (FSC) and side scatter (SSC).

Statistical analysis:

Data were explored, processed and analyzed using the statistical package for the social science, windows version 16, USA (SPSS PC+ version 16 software). Variable with normal distribution were expressed as mean± SD. In these variables, the T test was applied for group differences.

RESULTS

We studied a total of 38 subjects. Their characteristics and clinical data are summarized in Table (1).

Table 1.Characteristics data of LN patients and control subjects

	Control	Lupus Nephritis patients
Case no.	11	27
Sex (male\Female)	5\6	7\20
Age /years	9.98±2.10	9.68±2.89
Anti-dsDNA	7.5±5.5	230±168.23

T-cell count discriminates between LN patients and control

Table (3) represents the flow cytometry markers of T-cells% from blood samples. CD4⁺ T cells levels in blood samples obtained from LN patients were significantly elevated as compared to healthy control group (25.1± 8.43and 13.33± 3.97,respectively; p <0.001). The CD8⁺ T

cells levels in blood of samples obtained from LN group was significantly decreased as compared to healthy control (19.33±11.82and 30.02± 7.52respectively; P ≤0.01). Fig (2) showed the INF γ levels in blood of samples obtained from LN group was significantly decreased as compared to healthy control (9.24± 3.6and 14.05± 4.36 respectively; P ≤0.01).

Table (3): Flow cytometry markers for T-cells% from blood samples of studied subjects included in the study (n=38)

	Control (n=11)	LN (n=27)
CD4 ⁺	13.33± 3.97	25.1± 8.43***
CD8 ⁺	30.02± 7.52	19.33±11.82***
INF γ	14.05± 4.36	9.24± 3.6***

*** (P value <0.001)

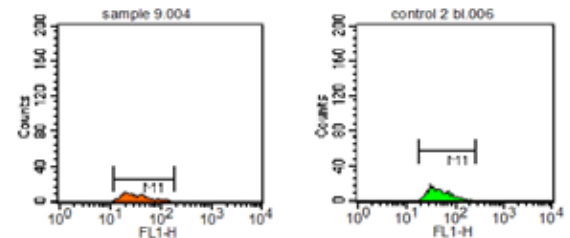


Fig (2) showed the% of blood cells stained with INF γ FITC conjugated fluorescence in LN patients and control.

Discussion

IFN- γ has been long associated with lupus pathology but its precise role in disease pathogenesis has been unclear. IFN- γ promotes T cell proliferation from the naive stage and augments Bcl-6 expression in Th cells and their precursors. These effects are likely to enhance the formation and possibly the maintenance of Th cells (Lin et al., 2011).

The role of IFN γ in the pathogenesis of SLE has been reported by Sau et al., (2012). However, there is contradictory data available about the dominance of Th1 and Th2 cytokines in SLE. Tucci et al. (2008) reported an increase level of serum IFN γ in patients with SLE.

The elevation of IFN γ level causes further inflammation and tissue injury in SLE patients. Increased level of IFN γ associates with nephritis in SLE patients. The result is similar to the study of Tucci et al. where they suggested the increased production of IFN γ in SLE patients. IFN γ are involved in cell mediated autoimmune disease, however, have been refuted by recent studies in several induced and spontaneous mouse models of SLE which showed that INF is a major effector molecule in this disease. These and additional findings reviewed here, suggested that these two cross-talking cytokines can exert autoimmune disease-promoting or disease-inhibiting effects without predictability or strict adherence to the Th1 versus Th2 dualism. (Tseiosopoulos et al., 2001).

In the present study, INF γ overexpression on T cells was significantly decreased as compared to the healthy control group (P ≤ 0.01). This finding disagrees with that of (Koenig et al., 2012). According to the low expression of INF γ in T lymphocyte cells of LN children, this is contrary to the high level of serum IFN γ in patients with SLE.

T cells with suppressor function were first described by

(Gershon and Kondo 1971). Thereafter, several observations supported their role in immune regulation processes. In fact, at least two lines of evidence suggest the existence of CD8⁺ T cells able to inhibit immune responses. First, the isolation of Ag-specific and alloantigen-specific T suppressor lymphocytes from mice and humans and, second, the identification and functional characterization of CD8⁺ T lymphocytes which suppress autoimmune responses in mice with experimental autoimmune encephalomyelitis. (Filaciet al., 2001)

In the current study, the CD8⁺ T cells (%) in blood samples obtained from LN patients was significantly decreased as compared to healthy control group ($p < 0.01$). This finding was in agreement with the study of Filaciet al., (2001) who demonstrated that, the lack of suppressor T cell activity may play a role in the pathogenesis of disease.

Dinesh et al., (2010) reported that, CD8⁺ Tregs in lupus patients have defective and/or reduced numbers, while Alvarado et al., (2006) study found no difference in the number of CD8⁺ Tregs in SLE patients. A multiplicity of different CD8⁺ Tregs subsets appears to have different mechanisms of suppression, markers, and modalities of induction. This diversity is one of the field's most salient characteristics because there is so far no single molecule that can specifically identify CD8⁺ suppressor or Tregs, either naturally occurring or induced.

In the present study, the surface markers CD4⁺ T cells (%) in blood samples obtained from LN patients was significantly elevated as compared to healthy control group ($p < 0.01$). This results are in agreement with those of Wanget al., 2013 and Tzifiet al., 2013. Tzifiet al., 2013 reported that increased numbers of CD4⁺ T cells were correlated with increased disease activity and the production of anti-self-antibodies.

The finding result was disagreement with the data obtained from Miyara et al., (2005). Who reported that all active patients studied presented not only with a decreased proportion of Tregs among CD4⁺ T cells, but also with decreased absolute number of the later cells, as compared with mean values obtained from healthy controls. These confictions may be attributing to the population and race differences.

In conclusion: the low expression of INF γ in T lymphocyte cells of LN patients may be a good marker in prognosis and diagnosis of LN .

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