Interferon α prevents spontaneous apoptosis of clonal Th2 cells associated with chronic hypereosinophilia

Liliane Schandené, Florence Roufosse, Aurore de Lavareille, Patrick Stordeur, André Efira, Bernard Kennès, Elie Cogan, and Michel Goldman

A recent study identified a clonal expansion of CD3⁺CD4⁺ cells secreting Th2-type cytokines in 4 patients with chronic hypereosinophilia. Because interferon α (IFN-α) is used in the therapy of the idiopathic hypereosinophilic syndrome, the effects of this cytokine on the survival of clonal Th2 cells isolated from the blood of 2 patients were determined. First, these cells displayed a high rate of spontaneous apoptosis on culture in cytokine-free medium and were also sensitive to Fas-mediated apoptosis induced by soluble Fas ligand. Addition of IFN-α or interleukin-2 (IL-2) to culture medium resulted in significant protection against spontaneous but not Fas-induced apoptosis. Although spontaneous apoptosis of the clonal Th2 cells was clearly associated with down-regulation of both bcl-2 and bcl-xL levels, IFN-α had no significant effect on the expression of these antiapoptotic proteins, whereas addition of IL-2 resulted in higher levels of bcl-2. On the other hand, IFN-α decreased the numbers of cells with disrupted mitochondrial transmembrane potential both during spontaneous apoptosis and after exposure to protoporphyrin IX. Thus, IFN-α might promote the survival of clonal Th2 cells, an effect that could be relevant to the therapeutic approach for patients with chronic hypereosinophilia caused by clonal expansion of Th2-type cells.

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Patients, materials, and methods

Patients

As previously described,15-17 patient 1 was a 20-year-old woman presenting with severe pruritus, eczema, and tenosynovitis of the right ankle. At presentation, the circulating leukocyte count was 16 900/µL, including 8923 eosinophils and 4630 lymphocytes. CD4⁺ T cells represented 87% of total lymphocytes and were composed of 88% CD3⁺CD4⁺ cells (3545 CD3⁺CD4⁺ cells/µL) and 12% CD3⁺CD4⁻ cells. Serum IgE and IgM levels reached 4285 BLOOD, 15 DECEMBER 2000 - VOLUME 96, NUMBER 13
levels were 340 U/mL (normal, < 20) and 310 mg/dL (normal, 40-250), respectively. Treatment with IFN-α was followed by regression of both clinical manifestations and decrease of eosinophil blood count, although the clonal CD3+CD4+ population remained stable. However, recurrence of hypereosinophilia was observed less than a year after initiation of IFN-α. Subsequent evolution has been characterized by expansion of the circulating CD3+CD4+ population (reaching 4800 µL) and appearance of chromosomal abnormalities in these cells. The patient has now developed peripheral T-cell lymphoma characterized by bilateral masses in the parotid regions and a submandibular mass composed of CD3+CD4+ cells. The patient has now developed peripheral T-cell lymphoma characterized by bilateral masses in the parotid regions and a submandibular mass composed of CD3+CD4+ cells.

Patient 21617 was a 21-year-old woman also presenting with severe pruritus and eczema, as well as cyclic angioedema. The circulating leukocyte count was 14 800 µL, including 9102 eosinophils and 3419 lymphocytes. CD4+ T cells represented 87% of total lymphocytes and were composed of 83% CD3+CD4+ cells (2469 CD3+CD4+ cells/µL) and 17% CD3+CD8+ cells. Serum IgE and IgM levels reached 15 640 U/mL and 1253 mg/dL, respectively. The clinical and biologic findings were consistent with the diagnosis of Gleich syndrome. Two-year follow-up has been characterized by good response to glucocorticoid treatment, as evidenced by normalization of eosinophil levels. Although the CD3+CD4+ cell population had decreased significantly (662µL) over this period, tapering of steroid dosage was quickly followed by recurrence of hypereosinophilia and clinical manifestations. In both patients, bone marrow aspiration showed abundant eosinophil precursors and absence of blastic cells. They were free of treatment at the time of blood sampling for the present study.

**Reagents**

Human recombinant IFN-α (rIFN-α) 2b with a specific activity of 3.10^6 IU/mL was obtained from Schering-Plough (Innishannon, Ireland). Human recombinant IL-2 (rIL-2) with a specific activity of 3.10^6 IU/mL was kindly provided by K. Willard-Gallo (Université Libre de Bruxelles, Rhode Saint-Gèneze, Belgium). The following monoclonal antibodies (mAbs) were purchased from Becton Dickinson (Mountain View, CA): fluorescein isothiocyanate (FITC)-labeled anti-CD45RA; peridinin chlorophyll protein (PerCP)-labeled anti-CD4; phycoerythrin (PE)-labeled anti-Fas (CD95), anti-CD3, and anti-CD45RO; purified anti-CD3; and anti-CD45RA. The neutralizing anti-Fas ligand (clone NOK-1) mAb was purchased from Pharmingen (San Diego, CA). The soluble human recombinant APO-1/Fas ligand (sFasL) and associated enhancer (antibody reacting with sFasL) were purchased from Alexis Corporation (Laufelfingen, Switzerland). Mouse monoclonal anti-bcl-xL (IgG1, H-5) antibody was obtained from Santa Cruz Biotechnology (Santa Cruz, CA) and mouse IgG1 isotype control from R & D Systems Europe (Abingdon, United Kingdom). FITC-labeled rabbit anti-mouse IgM used as second layers, FITC-conjugated anti-bcl-2 mAb (IgG1, clone 124), and IgG1 control mAb for intracellular staining of bcl-2 were obtained from Dako (Glostrup, Denmark).

**Isolation of CD3+CD4+ T cells from hypereosinophilic patients**

Blood leukocytes were obtained from both patients by cytopheresis, after informed consent was obtained. Peripheral blood mononuclear cells (PBMC) were isolated by density gradient centrifugation on Lymphoprep (Nycosmed, Oslo, Norway). After 3 washings with Hanks balanced salt solution (HBSS) (Gibco, Life Technologies, Paisley, Scotland), cells were resuspended in cold-conservation medium composed of 50% RPMI 1640, 10% dimethylsulfoxide (DMSO) (Sigma Chemical Co, St Louis, MO), and 40% heat-inactivated fetal calf serum (FCS) (BioWhittaker, Europe, Verviers, Belgium). Cells were quickly frozen in a −80°C freezer and then stored in liquid nitrogen until use. CD3+CD4+ T cells from both patients were prepared from CD4+ T cells (see below) by selective depletion of CD3+ T cells through incubation with anti-CD3 mAb for 30 minutes at 4°C, washing, and then incubation with immunomagnetic particles coated with sheep antimouse IgG (Dynabeads, Dynal, Oslo, Norway). The resulting cell preparations contained more than 90% viable CD3+CD4+ T cells as determined by flow cytometry.

**Isolation of CD3+CD4+CD45RO+ T cells from healthy volunteers**

The PBMC were isolated from buffy coats of healthy individuals by density gradient centrifugation on Lymphoprep. Cells were resuspended in cold-conservation medium, quickly frozen, and then stored in liquid nitrogen until use. After 3 washings with HBSS, PBMC were resuspended in complete medium (RPMI 1640, BioWhitaker, supplemented with 10% FCS and 40 µg/mL gentamycin). CD4+ lymphocytes were purified using the MACS CD4+ T cell isolation kit (Miltenyi Biotec GmbH, Bergisch Gladbach, Germany). The CD4+ T-cell suspensions were further incubated with an anti-CD45RA mAb for 30 minutes at 4°C, washed, and incubated with immunomagnetic particles coated with sheep antimouse IgG (Dynabeads, Dynal). Coated cells were removed with a magnet, leaving purified CD3+CD4+CD45RO+ cells in suspension. The resulting cell preparations routinely contained more than 90% viable CD3+CD4+CD45RO+ T cells.

**Lymphocyte phenotyping**

The expression of HLA-DR, CD45RO, and Fas (CD95) on CD4+ T cells was determined by flow cytometry. For that purpose, PBMC from patients and from healthy blood donors were washed in phosphate-buffered saline (PBS) and incubated for 30 minutes on ice with appropriate mAbs: FITC-, PE-, or PerCP-conjugated mAbs against CD4, CD3, CD45RO, HLA-DR, and CD95 antigens. After washing, 10 000 cells were acquired using a FACScan flow cytometer (Becton Dickinson) and analyzed with the Cellquest software (Becton Dickinson).

**Cell culture and determination of apoptosis**

Normal CD3+CD4+CD45RO+ and clonal CD3+CD4+ T cells were seeded at 2.10^5/well per 200 µL at 37°C and 5% CO2-humidified atmosphere in flat-bottomed 96-well plates (Nunc, Gibco) in complete medium in the presence or absence of IL-2 (100 U/mL) or IFN-α (10^5-1 U/mL). For induction of Fas-mediated apoptosis, CD4+ T cells were incubated in the presence of sFasL at concentrations ranging from 50 to 1.5 ng/mL and associated enhancer (1 µg/mL). After 24 to 96 hours of incubation, CD4+ T cells were collected and examined for apoptosis. In most experiments, recovered viable lymphocytes were detected by flow cytometry according to their light forward and side scatter properties (FSC/SSC). Indeed, apoptotic lymphocytes were smaller in size than resting viable lymphocytes but were distinct from debris by an increase in 90° scatter. After culture, numbers of viable CD3+CD4+CD45RO+ or CD3+CD4+ T lymphocytes (FSC<sup>hi</sup>/SSC<sup>low</sup>) were recovered and determined as expressed as a percentage of the original input viable cells. Based on these measurements, results were expressed as percentage of apoptotic cells. Measurement of apoptosis by the light scatter method was verified by 2 additional methods for apoptosis detection, namely annexin V binding and terminal deoxynucleotidyl transferase (TdT)–mediated deoxyuridine triphosphate (dUTP) nick end-labeling (TUNEL). FITC–annexin V binding detects phosphatidyl serine on the outer surface of cells undergoing apoptosis and was performed using a commercially available kit (Bender MedSystems, Vienna, Austria). TUNEL was performed using a kit according to the manufacturer’s protocol (Boehringer Mannheim, Mannheim, Germany). We confirmed that FSC<sup>low</sup>/SSC<sup>low</sup> cells were clearly enriched in apoptotic cells (98% annexin V positive [An+] and 97% TUNEL positive), whereas FSC<sup>hi</sup>/SSC<sup>low</sup> cells contained few apoptotic cells (9% An+ and 7% TUNEL positive). Double-labeling experiments with FITC–annexin V and propidium iodide (PI) revealed that most of the An+ cells were PI+ after 24 hours of culture (not shown).

**Expression of bcl-2 and bcl-xL**

Levels of bcl-2 and bcl-xL expression were determined by flow cytometry. Intraacellular staining for bcl-2 was performed as follows. Cells were washed in PBS supplemented with 10% FCS and 0.1% azide (PBS-FCS-Na<sub>3</sub>) fixed in FACS lysis solution (Becton Dickinson) for 10 minutes at room temperature, washed again in PBS, and then incubated for 15 minutes at room temperature in permeabilizing solution (Becton Dickinson) in PBS-FCS-Na<sub>3</sub>. Finally, cells were intracellularly stained with FITC-
To evaluate $Dc$ species generation cytometry using 0.25 $\text{m}$ that is oxidized into fluorescent ethidium (red) by superoxide anion. 30 $Dc$ with disrupted control Abs is represented by the broken line.

Representative of 3. The background staining using isotype-matched patients. The figure shows histograms from 1 experiment, representative of 2 patients with chronic hypereosinophilia. 15-17 As shown in Figure 1, by flow cytometry, their phenotype was further characterized by the expression of the CD45RO isoform and the activation antigen HLA-DR and by the absence of the CD7 and CD27 markers.16,17 (not shown). Moreover, the clonal T cells expressed high levels of CD95 as compared with CD3$^+$ CD4$^+$ CD45RO$^+$ T cells obtained from healthy individuals. The phenotype of the clonal Th2 cells is therefore consistent with that of activated memory T cells.

**Cyttofluorometric analysis of $\Delta \phi_m$ and reactive oxygen species generation**

To evaluate $\Delta \phi_m$, clonal Th2 cells (10$^4$/mL) from patient 1 were incubated in complete medium with 2.5 $\text{mmol/L}$ 3,3'-dihexyloxacarbocyanine (DiOC$_6$; Molecular Probes, Eugene, OR) for 15 minutes at 37°C followed by analysis on a FACScan flow cytometer, according to a published protocol.30 The fluorescence of DiOC$_6$, a potential-sensitive cationic lipophilic dye that incorporates into the mitochondrial matrix, correlates with the $\Delta \phi_m$ and is oxidation independent. Therefore, DiOC$_6$ cells correspond to cells with disrupted $\Delta \phi_m$. Superoxide generation was also measured by flow cytometry using 0.25 $\mu\text{mol/L}$ hydroethidine (HE; Molecular Probes), a dye that is oxidized into fluorescent ethidium (red) by superoxide anion.30 HE$^+$ cells therefore correspond to cells with increased production of superoxide anions. In some experiments, the clonal Th2 cells were cultured in the presence of 30 $\mu\text{mol/L}$ protoporphyrin IX (PPIX; Sigma), alone or in combination with IFN-$\alpha$ (10$^4$ IU/mL).

**Results**

**Clonal CD3$^+$ CD4$^+$ Th2 cells associated with chronic hypereosinophilia present an activated memory phenotype**

We described elsewhere the Th2-like cytokine profile of the monoclonal CD3$^+$ CD4$^+$ T lymphocytes present in the circulation of the 2 patients with chronic hypereosinophilia.15-17 As shown in Figure 1, by flow cytometry, their phenotype was further characterized by the expression of the CD45RO isoform and the activation antigen HLA-DR and by the absence of the CD7 and CD27 markers16,17 (not shown). Moreover, the clonal T cells expressed high levels of CD95 as compared with CD3$^+$ CD4$^+$ CD45RO$^+$ T cells obtained from healthy individuals. The phenotype of the clonal Th2 cells is therefore consistent with that of activated memory T cells.

**Interferon $\alpha$ protects clonal Th2 cells against spontaneous but not Fas-mediated apoptosis**

In a first series of experiments, we compared the rate of spontaneous apoptosis of clonal CD3$^+$ CD4$^+$ cells from patients with that of CD3$^+$ CD4$^+$ CD45RO$^+$ T cells from healthy individuals during a culture period of 1 to 4 days in the absence of any growth factor. The percentage of apoptotic cells was then quantified using flow cytometry analysis according FSC/SSC parameters, measuring DNA fragmentation using TUNEL assay and phosphatidylserine externalization using FITC–annexin V binding. The data presented in Figure 2 are a graphic representation of the flow cytometry data for FSC/SSC parameters. We found at all time points a significantly increased proportion of apoptotic cells within the clonal Th2 cells as compared with control CD3$^+$ CD4$^+$ CD45RO$^+$ T cells. Similar results were obtained using either FITC–annexin V binding or TUNEL assay, except that in every case more apoptotic cells were detected as defined by FSC/SSC or annexin V binding than as defined by the TUNEL assay. This difference was probably due to the fact that DNA degradation is a relatively late event in apoptosis.

The increased expression of the CD95 antigen on the clonal Th2 cells prompted us to examine whether the propensity of these cells to undergo apoptosis could be enhanced through ligation of Fas. As shown Figure 3, most of the clonal Th2 cells (> 90%) underwent apoptosis after 24 hours of incubation with 50 ng/mL sFasL, whereas less than 40% of apoptotic cells were found within control CD3$^+$ CD4$^+$ CD45RO$^+$ T cells under the same experimental conditions. The apoptosis induction by Fas engagement led us to...
determine whether this pathway was involved in the spontaneous apoptosis of those cells as they express mRNA encoding FasL (not shown). As shown in Table 1, addition of a blocking anti-FasL mAb did not affect the spontaneous apoptosis of the clonal Th2 cells, whereas it inhibited sFasL-induced apoptosis. These data establish that the spontaneous apoptosis of clonal Th2 cells is independent of the Fas/FasL pathway.

We then examined whether the clonal Th2 cells could be rescued from apoptosis by IL-2, which is known to exert antiapoptotic effects on activated T cells. As shown in Figure 4, addition of IL-2 (100 U/mL) to the culture medium rescued more than 80% of the clonal T cells from spontaneous apoptosis. Addition of rIFN-α (10^4 IU/mL) also exerted a clear-cut antiapoptotic action on clonal Th2 cells from both patients. As shown in Figure 5, the protective effect of rIFN-α on spontaneous apoptosis was dependent on the dose, a protective effect being already observed at a concentration of 1 IU/mL. The rescuing effect of IFN-α was also confirmed using FITC–annexin V binding and TUNEL assay (not shown). Contrasting with their inhibitory effects on spontaneous apoptosis, neither rIFN-α nor IL-2 affected Fas-induced apoptosis of the clonal T cells (Table 1). In a previous study, we demonstrated that IL-2 but not IFN-α induced proliferation of the Th2 cells. 16

**Table 1.** IFN-α and IL-2 protect the clonal Th2 cells against spontaneous but not Fas-mediated apoptosis

<table>
<thead>
<tr>
<th>Agent added</th>
<th>Patient 1</th>
<th>Patient 2</th>
<th>Patient 1</th>
<th>Patient 2</th>
</tr>
</thead>
<tbody>
<tr>
<td>None</td>
<td>28 ± 1</td>
<td>31 ± 3</td>
<td>89 ± 4</td>
<td>96 ± 2</td>
</tr>
<tr>
<td>Anti-Fas mAb</td>
<td>29 ± 2</td>
<td>28 ± 1</td>
<td>30 ± 3</td>
<td>28 ± 1</td>
</tr>
<tr>
<td>Control mAb</td>
<td>28 ± 1</td>
<td>28 ± 1</td>
<td>90 ± 4</td>
<td>97 ± 1</td>
</tr>
<tr>
<td>IL-2 (100 IU/mL)</td>
<td>11 ± 0.5*</td>
<td>10 ± 1*</td>
<td>78 ± 7</td>
<td>94 ± 1</td>
</tr>
<tr>
<td>IFN-α (10^4 IU/mL)</td>
<td>9 ± 1*</td>
<td>8 ± 0.5*</td>
<td>83 ± 5</td>
<td>89 ± 4</td>
</tr>
</tbody>
</table>

Purified CD3^+ CD4^+ CD45RO^+ cells from patient 1 and patient 2 were cultured for 24 hours in absence (spontaneous) or presence (Fas-mediated) of sFasL (25 ng/mL), Anti-Fas (1 μg/mL) or control mAb (1 μg/mL), IL-2 (100 IU/mL) or IFN-α (10^4 IU/mL) was added at the beginning of the culture. At the end of the culture, apoptotic lymphocytes were detected by flow cytometry according to their light scatter properties (FSC/SSC). Results are the mean percentages ± SD of 3 experiments.

*P < .05 as compared with culture without IFN-α or IL-2 (ANOVA test).
Interferon α does not prevent bcl-2 and bcl-xL down-regulation in clonal Th2 cells

To determine whether the susceptibility of the clonal Th2 cells to apoptosis was caused by changes in the constitutive expression of bcl-2 and bcl-xL, which are known to exert antiapoptotic effects, purified clonal Th2 cells and normal CD3⁺CD4⁺CD45RO⁺ T cells were stained for these proteins immediately after purification and analyzed by flow cytometry. As shown in Table 2, bcl-2 expression was clearly lower among the clonal Th2 cells, whereas no significant difference was observed as far as bcl-xL expression is concerned.

Interleukin-2 is known to protect activated T cells against death by increasing bcl-2 and, to a lesser extent, bcl-xL expression. We therefore decided to compare the effects of IL-2 and IFN-α on expression levels of both bcl proteins within the clonal Th2 cells of patient 1. For this purpose, bcl-2 and bcl-xL protein expressions were examined by flow cytometry after culture in the absence or presence of the cytokines. Apoptotic and live cells were analyzed separately on the basis of FSC/SSC parameters. As shown in Table 2, bcl-2 and bcl-xL expression by live clonal Th2 cells decreased by about 40% after 24 hours of culture in the absence of exogenous cytokines. The drop in bcl-xL protein expression was even more pronounced within apoptotic cells, resulting in a clear bimodal expression of bcl-xL by the clonal Th2 cells (not shown). On the other hand, normal CD3⁺CD4⁺CD45RO⁺ T cells decreased their bcl-2 and bcl-xL levels only by about 10%. Although addition of IL-2 (100 IU/mL) partially prevented the drop in bcl-2 expression in clonal Th2 cells, the antiapoptotic effect of IFN-α was not associated with any significant change in bcl-2 or bcl-xL down-regulation during the 24-hour culture period (similar data were obtained after 72 hours of culture).

**Interferon α prevents Δψm dissipation in clonal Th2 cells**

Loss of Δψm, generation of superoxide anions by the uncoupled respiratory chain,27,29 and phosphatidylserine exposure31 at the cell membrane are characteristic features of the apoptotic process. To get further inside the mechanism of action of IFN-α, we monitored these parameters within the clonal Th2 cells.

First, we observed that a fraction of Th2 cells spontaneously become DiOC6low and HEhigh after 24 hours in culture, indicating disrupted Δψm and increased superoxide production, respectively (Figure 6). Double-labeling experiments (not shown) revealed that most DiOC6low cells were HEhigh, FSCint/SSCint, and An−, confirming the association between Δψm disruption, superoxide generation, and apoptosis. Addition of IFN-α (10^4 IU/mL) to the culture medium resulted in a lower number of Th2 cells undergoing Δψm

Table 2. Expression of bcl-2 and bcl-xL in clonal Th2 cells

<table>
<thead>
<tr>
<th>Cells</th>
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<th>0</th>
<th>24 h</th>
<th>0</th>
<th>24 h</th>
</tr>
</thead>
<tbody>
<tr>
<td>CD3⁺CD45RO⁺ (controls)</td>
<td>—</td>
<td>83 021 ± 6427</td>
<td>71 524 ± 4752</td>
<td>64 986 ± 5483</td>
<td>58 933 ± 4982</td>
</tr>
<tr>
<td>CD3⁺CD4⁺ (patient 2)</td>
<td>—</td>
<td>85 432 ± 5561</td>
<td>—</td>
<td>26 048 ± 6843*</td>
<td>—</td>
</tr>
<tr>
<td>CD3⁺CD4⁺ (patient 1)</td>
<td>—</td>
<td>96 953 ± 11704</td>
<td>58 644 ± 10128</td>
<td>29 839 ± 5290*</td>
<td>19 973 ± 2213</td>
</tr>
<tr>
<td>+ IFN-α (10^5 IU/mL)</td>
<td>—</td>
<td>62 563 ± 13723</td>
<td>—</td>
<td>19 911 ± 2555</td>
<td></td>
</tr>
<tr>
<td>+ IL-2 (100 IU/mL)</td>
<td>—</td>
<td>66 238 ± 22834</td>
<td>—</td>
<td>23 647 ± 3371†</td>
<td></td>
</tr>
</tbody>
</table>

Intracytoplasmic bcl-xL and bcl-2 staining was performed on purified CD3⁺CD4⁺CD45RO⁺ cells from healthy individuals (n = 6) and CD3⁺CD4⁺ from hypereosinophilic patients (for each patient 6 determinations were performed). For patient 1, bcl-xL and bcl-2 levels were compared with those obtained after 24 hours of incubation in the absence or presence of IFN-α (10^4 IU/mL) or IL-2 (100 IU/mL). Results shown are the mean ± SD of bcl-xL and bcl-2 staining within live FSCint/SSCint cells expressed in MESF.

*P < .05 as compared with CD4⁺CD45RO⁺ cells from healthy individuals.
†P < .05 as compared with bcl-2 expression obtained in absence of IL-2 (ANOVA test).
loss and associated HE⁻/An⁺ conversion. Indeed, percentages of DiOC₆⁻/HE⁺/An⁺ cells after 24 hours of culture were 30.5% ± 3.3% and 16.4% ± 0.9% (mean ± SD of 5 determinations) in absence and presence of rIFN-α, respectively (P < .005 by Wilcoxon test). As shown in Figure 6, we also found that an important proportion of the clonal Th2 cells (70% mean) exhibited a drastic ΔΨm loss 4 hours after Fas cross-linking by sFasL (25 ng/mL). Most of these cells (90%) were HE⁺ and An⁺. Less than 5% of the clonal Th2 cells were rescued from Fas-mediated ΔΨm loss and HE⁺/An⁺ conversion by rIFN-α added at the beginning of the culture.

To further specify the mechanism of the antiapoptotic effect of IFN-α at the mitochondrial level, we studied the ΔΨm modification induced in Th2 cells by PPIX, an agent known to trigger permeability transition (PT) pore opening after binding to benzodiazepine receptors located in the outer mitochondrial membrane. After incubation with this agent for 24 hours, we found that more than 60% of the clonal Th2 cells were still alive with a FSClow/SSCmed/An⁺ phenotype. After gating on this population of viable cells, we observed 2 subpopulations of DiOC₆⁻⁰ Th2 cells (Figure 7), in agreement with a previous study on thymocytes, which demonstrated that the cell population with the lowest ΔΨm fluorescence was the one committed to apoptosis. As shown in Figure 7, addition of rIFN-α (10⁴ IU/mL) to PPIX significantly decreased the numbers of cells expressing the lowest ΔΨm fluorescence. Indeed, percentages of those cells after 24 hours of culture were 40 ± 8.9% and 22.6 ± 4.8% (mean ± SEM of 8 determinations) in the absence and presence of rIFN-α, respectively (P < .01 by the Wilcoxon test). Taken together, these data indicate that IFN-α inhibits ΔΨm decrease induced by PT pore opening in Th2 cells.

Discussion

In this study, we first demonstrated that the clonal Th2-type cells associated with chronic hypereosinophilia displayed a high rate of spontaneous apoptosis on culture in cytokine-free medium. This increased susceptibility to apoptosis was associated with a low level of bcl-2 expression, whereas bcl-xL expression was normal as compared to control cells.

It is well known that members of the IL-2 family that share the γc as part of their receptors can prevent the in vitro death of activated T cells. IL-2 and IL-4 (not shown) indeed prevented spontaneous apoptosis of the clonal Th2 cells. We observed that IFN-α represents another type of survival factor for the clonal Th2 cells. Autocrine production of IL-2 did not contribute to IFN-α-induced survival of the clonal Th2 cells, because this activity was inhibited neither by anti–IL-2R-α nor anti–IL-2R-γ neutralizing antibodies (data not shown). Several studies indicate that bcl-2 and bcl-xL repress cell death of cytokine-deprived activated T cells. In this report, when clonal Th2 cells were cultured alone, bcl-2 and bcl-xL protein expression decreased within live cells, suggesting that down-modulation of these antiapoptotic proteins could be involved in the onset of spontaneous apoptosis. Although addition of IL-2 or IFN-α resulted in an improved survival of the clonal Th2 cells, IFN-α had no significant effect on the expression of these antiapoptotic proteins, whereas addition of IL-2 resulted in higher levels of bcl-2, as expected. In contrast to IFN-α, IFN-γ did not influence the death rate of the clonal Th2 cells, although these cells bear receptors for IFN-γ because their expression of class I major histocompatibility complex molecules was upregulated by IFN-γ (data not shown). Because both IFN-γ and IFN-α activate Stat1, whereas only IFN-α protects Th2 cells from apoptosis, it seems evident that Stat1 activation might not be sufficient to obtain the antiapoptotic effect in those cells. However, it is also possible that in the case of IFN-γ, a proapoptotic pathway is simultaneously triggered.

To get further insight into the mechanism of action of IFN-α, we evaluated the possible role of mitochondria in the spontaneous apoptosis of the clonal cells. Indeed, recent studies establish that mitochondria play a critical role in the apoptotic cascade leading to T-cell death. Early after induction of apoptosis, PT-induced pore opening results in loss of ΔΨm and the generation of superoxide anions by the uncoupled respiratory chain. The hierarchical relationship between these events appears complex. Along this line, Hildeman and colleagues established that the production of superoxide anions associated with a loss of ΔΨm is a decisive contributor to the in vitro death of activated T cells obtained from mice injected with superantigen. Interestingly, it was reported in another work that type I IFNs can act as a survival factor for these cells. Moreover, Hyde and coworkers demonstrated that a fibroblast-derived factor, recently recognized as IFN-β, enhanced survival of activated human T cells by increasing their intracellular antioxidant glutathione levels.

Others works reported that PT pore opening induced in ΔΨm loss would constitute a sufficient event able to trigger apoptosis in thymocytes and T cells. In the present study, the capacity of the PT pore inducer PPIX to trigger ΔΨm disruption as well as apoptosis in the Th2 clonal cells supported this concept. The inhibitory effect of IFN-α on ΔΨm changes induced by PPIX strongly suggests that IFN-α promotes the survival of the clonal cells at least in part by preventing mitochondrial changes leading to apoptosis.

Our observations are also consistent with studies demonstrating that type I IFNs display antiapoptotic activity on in vivo activated T cells obtained from human inflamed rheumatoid synovium. Also, IFN-α has been reported to prevent spontaneous apoptosis of chronic lymphocytic leukemia cells and myeloma cells.
B-cell receptor-mediated apoptosis in B cells, as well as Fas-mediated apoptosis of myeloma cells, chronic myelogenous leukemia progenitors, and antigen-specific human T-cell clones generated in vitro. The fact that IFN-α did not affect Fas-mediated apoptosis in clonal Th2 cells from our patients with hypeeosinophilia indicates that the influence of IFN-α on the Fas-mediated death pathway might depend on the cell type considered.

The spontaneous apoptosis of the Th2-like cells contrasts with their persistence in vivo, in association with hypeeosinophilia and their contact with antigen-presenting cells. In the present study, we demonstrate that a type I IFN, IFN-α, represents another survival factor of Th2 cells that might be operative in vivo in a paracrine fashion. This line, recent data indicate that fibroblasts producing IFN-β keep alive activated T cells with low bcl-2 that have reverted to a quiescent memory state. Our findings might be clinically relevant because they suggest that the long-term effects of IFN-α therapy in HES might depend on the subtype of the syndrome considered. Initially, the rationale for use of IFN-α in the management of HES was based on its beneficial effects in patients with chronic myelogenous leukemia and the fact that some patients with HES presented several features suggestive of a chronic myeloproliferative disease. Although it is plausible that some patients with HES actually present a true myeloproliferative disorder, current techniques fail to provide evidence for this in most cases. On rare occasions, clonality of eosinophils has been demonstrated in patients fulfilling the diagnostic criteria of HES, leading to the more appropriate diagnosis of chronic eosinophilic leukemia. The beneficial effects observed with IFN-α in such patients may result from its suppressive action both on eosinophil progenitors and on differentiated eosinophils. The notion that clonal Th2 cells play a pathogenic role in a subgroup of patients with chronic hypeeosinophilia has emerged only recently, and most studies reporting favorable effects of IFN-α in the setting of HES have been conducted on patients with no evidence of an underlying T-cell disorder. In another study of 16 hypeeosinophilic patients presenting an aberrant population of IL-5–producing cells, T-cell lymphoma was ultimately diagnosed in 3 patients. One of these patients had previously been treated with IFN-α, and therapeutic measures were not specified for the 2 others. In view of the antiapoptotic effects of IFN-α reported in the present paper, we suggest carefully evaluating the long-term outcome of patients receiving isolated IFN-α therapy in the context of the Th2 variant of HES.

Acknowledgments

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