

Curcumin induces cell death without oligonucleosomal DNA fragmentation in quiescent and proliferating human CD8+ cells

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Cytotoxic CD8+ cells play an important role in determining host response to tumor, thus chemotherapy is potentially dangerous as it may lead to T cells depletion. The purpose of this study was to elucidate the propensity of quiescent and proliferating human CD8+ cells to undergo cell death upon treatment with curcumin, a natural dye in Phase I of clinical trials as a prospective chemopreventive agent. **Methods:** We treated human quiescent or proliferating CD8+ cells with 50 μ M curcumin or irradiated them with UVC. Cell death symptoms such as decreased cell viability, chromatin condensation, activation of caspase-3 and specific DFF40/CAD endonuclease and oligonucleosomal DNA fragmentation were analyzed using MTT test, microscopic observation, Western blotting and flow cytometry. **Results:** Curcumin decreased cell viability, activated caspase-3 and decreased the level of DFF45/ICAD, the inhibitor of the DFF40/CAD endonuclease. However, this did not lead to oligonucleosomal DNA degradation. In contrast, UVC-irradiated proliferating, but not quiescent CD8+ cells revealed molecular and morphological changes characteristic for apoptosis, including oligonucleosomal DNA fragmentation. Curcumin can induce cell death in normal human lymphocytes both quiescent and proliferating, without oligonucleosomal DNA degradation which is considered as a main hallmark of apoptotic cell death. Taking into account the role of CD8+ cells in tumor response, their depletion during chemotherapy could be particularly undesirable.

Keywords: CD8+, cell death, curcumin, DNA degradation

INTRODUCTION

Apoptosis, or programmed cell death, is a fundamental process essential for both development and maintenance of tissue homeostasis (reviewed in Jacobson *et al.*, 1997). Cells undergoing apoptosis exhibit specific morphological changes including membrane blebbing, cytoplasmic and chromatin condensation, DNA fragmentation, nuclear breakdown and assembly of membrane-enclosed vesicles termed apoptotic bodies, eventually subjected to phagocytosis (reviewed in Wyllie *et al.*, 1980). Multiple apoptotic stimuli trigger the activation of proteases called caspases, which in turn initiate and execute the apoptotic program (reviewed in Riedl

& Shi, 2004). One of the hallmarks of the terminal stages of apoptosis is oligonucleosomal DNA fragmentation (Wyllie *et al.*, 1980). Recent years have led to the discovery of two major apoptotic nucleases, termed DNA fragmentation factor (DFF) or caspase-activated DNase (CAD) and endonuclease G (Endo G). In non-apoptotic cells, DFF exists in the nucleus as a heterodimer, composed of a 45 kDa chaperone and inhibitory subunit (DFF45 or ICAD-L) and a 40 kDa latent nuclease subunit (DFF40/CAD). Apoptotic activation of caspase-3 or -7 results in the cleavage of DFF45/ICAD and release of the DFF40/CAD nuclease which then forms active homo-oligomers (reviewed in Widlak & Garrard, 2005).

Abbreviations: 7-AAD, 7-aminoactinomycin D; AICD, activation-induced cell death; APCs, antigen presenting cells; CAD, caspase-activated DNase; CFSE, carboxyfluorescein diacetate succinimidyl ester; DFF, DNA fragmentation factor; Endo G, endonuclease G; FCS, fetal calf serum; HA, host antigen; mAb, monoclonal antibody; MDR, multidrug resistance; MTT, 3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyl tetrazolium bromide; PBS, phosphate-buffered saline; PHA, phytohaemagglutinin; TCR, T-cell receptor.

Human lymphocytes undergo so called "activation-induced cell death" (AICD) when activated lymphocytes are induced to die during the down-phase of their immune response. This is caused by ligation of the Fas receptor (CD95) with Fas ligand (CD95-L) whose expression is up regulated by TCR activation (Krammer, 2000). AICD can be mimicked *in vitro* by stimulation of TCR with non-specific mitogen, phytohaemagglutinin (PHA), or by specific stimulation of TCR/CD28 with anti CD3/antiCD28. A second type of physiological cell death of T cells caused by deprivation of survival stimuli, such as cytokines, leading to down-regulation of anti-apoptotic proteins such as Bcl-2 can be called "passive cell death" (Akbar & Salmon, 1997).

Moreover, normal lymphocytes can undergo cell death induced by drugs as a side-effect of chemotherapy aimed at malignant cells. This type of lymphocyte cell death can be included in the category of damage-induced cell death (De Martinis *et al.*, 2005). Usually chemotherapy causes severe toxicity in normal tissues, leading to side-effects such as mucositis, hair loss, and myelosuppression. In addition, chemotherapy induces acute lymphopenia and chronic depletion of T cells, leading to increased susceptibility to opportunistic infections.

CD8+ cells play a very important role in the anti-tumor immune response. Antigens from peripheral tumor cells can enter the class I pathway for presentation by host antigen (HA) presenting cells (APCs) to CD8 cells, a process commonly known as "cross-presentation" (Heath & Carbone, 2001). It has recently been shown in mice that not only the tumor presence but also induction of tumor cell apoptosis *in vivo* increases tumor antigen cross-presentation of HA to CD8 cells which are not deleted but primed as was revealed by CFSE proliferation assay (Nowak *et al.*, 2003). Quantitation of lymphocyte populations in the peripheral blood of patients enrolled in multiagent dose-intensive regimens revealed significant lymphocyte depletion, but with a more profound effect on CD4+ than on CD8+ T-cells (Mackall, 2000). Accordingly, in this paper we checked interested whether CD8+ cells, both quiescent and activated, might be indeed resistant to apoptosis induced by a classical DNA damaging agent, UVC, and the natural dye curcumin.

Curcumin (diferuloyl methane) is a naturally occurring yellow pigment derived from the rhizome of *Curcuma longa*. Turmeric or curcuma, the powdered form of the rhizome, is widely used in Asian countries where this plant has been cultivated for centuries. Curcumin exhibits a variety of pharmacological effects including anti-inflammatory, anti-infectious and anticancer activities (Lin *et al.*, 2000). The exposure of populations worldwide to curcu-

min, and its many uses, has led to studies aimed at elucidating some of its activities with particular attention on the anticancer activity. These flourished with several Phase I human trials that have shown this compound to be well tolerated (Cheng *et al.*, 2001; Hsu *et al.*, 2002; Iqbal *et al.*, 2003; Sharma *et al.*, 2004).

It has been shown that curcumin can inhibit proliferation and/or induce cell death in *in vitro* experiments with different cancer cells. The most common cell death mode upon curcumin treatment seems to be apoptosis (reviewed in Lin *et al.*, 2000; Karunagaran *et al.*, 2005). There are also reports showing that unlike in cancer cells, curcumin does not induce apoptosis in normal cells (Jiang *et al.*, 1996). However, we showed that curcumin induced cell death not only in cancer but also in normal rat and human lymphocytes (Bielak-Zmijewska *et al.*, 2000). The cell death induced with curcumin in normal and transformed lymphocytes was not characterized oligonucleosomal DNA fragmentation that is typical for apoptosis. Moreover, we showed in Jurkat cells that curcumin induced caspase-3 activation and following DFF40/CAD activation but concomitantly blocked the active centre of the endonuclease thus precluding DNA fragmentation (Sikora *et al.*, 2006). Here we investigated whether also in normal human quiescent and proliferating lymphocytes curcumin gives the same symptoms, namely cell death with activation of caspase-3 and DFF40/CAD endonuclease but without oligonucleosomal DNA fragmentation. As the oligonucleosomal DNA degradation is considered a main hallmark of apoptotic cell death the lack of this sort of degradation can be improperly interpreted as an absence of apoptosis.

MATERIALS AND METHODS

T-cell preparation and treatment. Mononuclear cells were obtained by standard centrifugation over Ficoll-Paque from lymphocyte buffy coat (about 10^8 cells) of peripheral blood of eight healthy blood donors (aged 22–33 years). The buffy coats were obtained from the Regional Centre for Blood Donation and Blood Treatment in Warsaw (Poland). All cell cultures were conducted in RPMI 1640 supplemented with 10% FCS, 2 mM L-glutamine and antibiotics, at a starting density of 0.8×10^6 cells/ml. The cells were stimulated by PHA (5 $\mu\text{g}/\mu\text{l}$) added at the time of seeding (day 0). From day 3 on, 60 U/ml recombinant IL-2 (rIL-2; Peprotech EC LTD, UK) was added every 2–3 days. CD8+ cells from day 0 and day 10–14 were isolated by MACS-magnetic cell sorting (Miltenyi Biotec, Medianus, Poland) by positive selection. The purity was >97% and cell viability measured by trypan blue exclusion test was over 95%.

To induce apoptosis with curcumin, CD8+ cells were treated for 8 h with 50 μM dye (Merck, Warsaw, Poland). For UVC treatment cells were irradiated with a pulse of ultraviolet light of 254 nm and an energy output of 100 J/m² using a Stratalinker 2400 (Stratagene, La Jolla, CA, USA) and collected 6 h after irradiation.

Cell phenotyping. Phenotypes of cells were examined by standard flow cytometry procedures. This involved double immunofluorescence staining of blood samples using Simultest from Becton-Dickinson. Each experiment included cells incubated with isotype controls. Samples were analyzed on FACSCalibur (Becton-Dickinson, Warsaw, Poland) using Cell-Quest software (Becton-Dickinson).

Cell proliferation assay. Mononuclear lymphocytes were labelled with a tracker dye, carboxy-fluorescein diacetate succinimidyl ester (CFSE, Molecular Probes) according to Hasbold *et al.* (1999), before stimulation with PHA and then at the 1st and 7th day of cell culture as described previously (Brzezinska *et al.*, 2003; 2004; Brzezinska, 2005).

DNA content analysis by flow cytometry. Cells were analyzed for DNA content by flow cytometry. One million cells were collected, washed and suspended in Nicoletti buffer (0.1% sodium citrate, pH 7.4, 0.1% Triton X-100, and 50 $\mu\text{g}/\text{ml}$ propidium iodide). DNA content was determined on a flow cytometer (FACSCalibur, Becton Dickinson). The sub-G₁ fraction represents apoptotic cells; cellular debris was excluded from the analysis. The levels of apoptotic cells induced in specific experimental conditions were calculated according to the following formula: (percentage of induced apoptosis minus percentage of spontaneous apoptosis)/(100 minus percentage of spontaneous apoptosis) \times 100.

Cell viability measurement. Cell viability was measured by MTT (3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyl tetrazolium bromide; Sigma) assay as described by the manufacturer.

Spontaneous apoptosis was measured using 7-AAD (7-amino-actinomycin D; Calbiochem) which stains apoptotic and necrotic cells. The 7-AAD positive cells were analyzed by flow cytometry.

Cell morphology observation. Morphological observation was performed after Hoechst 33258 staining (Molecular Probes, Eugene, OR, USA). Cells (0.2–0.3 \times 10⁶) were centrifuged on cytospin, fixed with 70% ethanol, washed in PBS and stained for 10 min in 1 μM Hoechst 33258 dye. Samples were visualized by epifluorescence microscopy (Nikon) and images were acquired with a color CCD camera.

Caspase-3 activation measurement. The activation of caspase-3 was analyzed by flow cytometry using PE-conjugated anti-active caspase-3 mAb (BD Pharmingen) according to the manufacturer's protocol.

Western blotting. The protocol used for Western blotting has been reported (Bielak-Mijewska *et al.*, 2004).

Proteins (40 μg per lane) were separated on 12% PAGE/SDS and electrotransferred onto nitrocellulose membrane (Hybond-C, Amersham). Membranes were probed overnight at 4°C with rabbit polyclonal anti-DFF40 (1:500), anti-DFF45 (1:500), or mouse monoclonal anti-PARP-1 (1:500) (BD Biosciences). Specific proteins were visualized with horseradish peroxidase-conjugated anti-IgG antibodies and the enhanced chemiluminescence (ECL) reagent (Amersham Pharmacia).

Statistics. Statistical analysis was performed by using Student's *t*-test.

RESULTS

PHA induces cell to proliferation and cell death

The majority of human lymphocytes isolated from peripheral blood are in a quiescent state. They can be stimulated to proliferate in culture by mitogens and IL-2. Upon stimulation with PHA, which mimics pathogen activation, T cells undergo intense proliferation and then activation-induced cell death (AICD). The most intense proliferation and AICD of T cells were observed at the 7th day following PHA stimulation (Fig. 1A). After 10 days of culture, proliferation was still at a high level; almost 70% of live cells were proliferating as measured by CFSE assay. Also morphological observation indicated that the majority of cells on day 10–14 of culture were blasts (Fig. 2B). The percentage of dying cells measured by 7-AAD staining accounted for about 20% at the 5th day after stimulation and lasted at that level to the 14th day of culture, but afterwards it increased dramatically to about 80% in four-week-old cultures (Fig. 1B). As the 7-AAD assay does not discriminate between apoptotic and necrotic cell death, we checked cell DNA content by using propidium iodide to measure the sub-G₁ fraction corresponding apoptotic cells. The sub-G₁ fraction was relatively high (about 30%) at 5 days after PHA stimulation and practically absent in two-week-old cultures reaching again 20% at the end of culture (4 weeks) (Fig. 1B).

During the initial days of culture the proportion of cytotoxic and helper cells changed, namely the CD8+ population increased from about 20% on day 0 to about 60% on day 14 suggesting that CD8+ cells might be less prone to PHA-induced cell death than CD4+ cells (Fig. 1C).

Accordingly, for further experiments isolated CD8+ cells from day 10–14 of mononuclear lym-

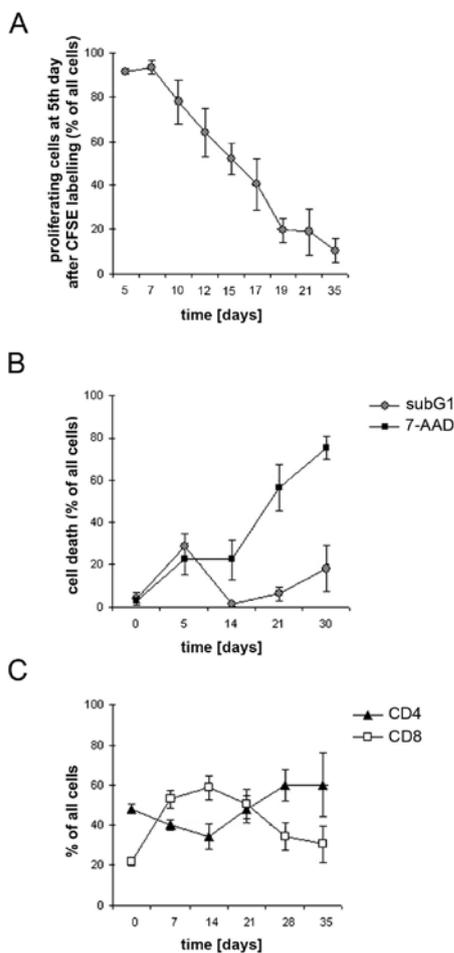


Figure 1. Proliferation, cell death and phenotype of PHA-stimulated T cells in long term culture of mononuclear lymphocytes.

A. Proliferation assay was performed by cell staining with CFSE and fluorescence was measured at 5th day after staining only in the gate of live cells (7-AAD negative). B. The amount of dying cells was assessed by 7-AAD staining and DNA content was measured with propidium iodide (sub-G₁ fraction). C. Cells were phenotyped using Simulstest. Data are expressed as mean \pm S.D. of eight independent experiments.

phocyte culture were chosen as they predominate in culture during the period of the more intense proliferation and they display the lowest ratio of the cell death. CD8+ cells activated stimulated after isolation from the mononuclear fraction did not respond with proliferation (not shown).

UVC but not curcumin induces DNA fragmentation in CD8+ cells

As previously we showed that rat and human quiescent T cells are less prone to undergo cell death induced by UVC than proliferating lymphocytes (Radziszewska *et al.*, 1999; 2000; Piwocka *et al.*, 1999) now we decided to study this difference in more detailed by focusing on UVC and curcumin's impact on non-activated and activated purified CD8+ cells.

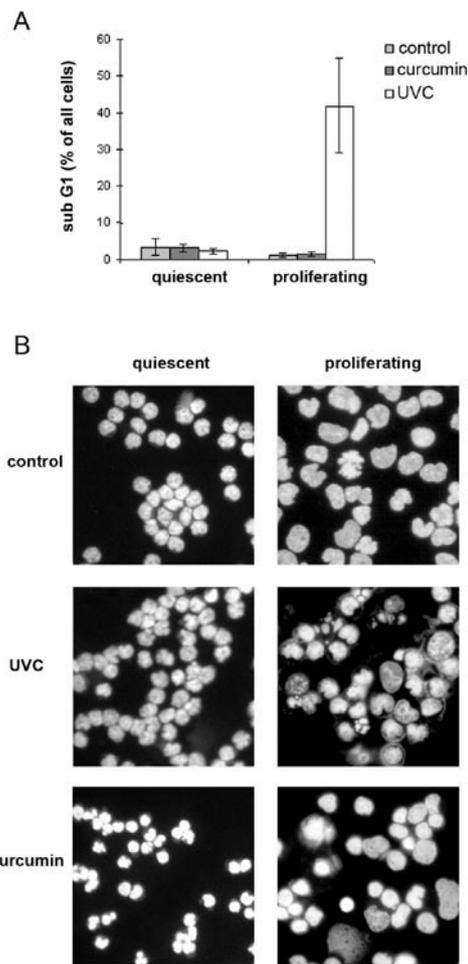


Figure 2. DNA fragmentation (A) and chromatin condensation (B) in quiescent and proliferating CD8+ cells treated with curcumin or UVC-irradiated.

DNA fragmentation was assessed by flow cytometry (sub-G₁). Data are expressed as mean \pm S.D. of three independent experiments. Chromatin condensation was observed microscopically after staining of cells with Hoechst 33258. Representative pictures of three independent experiments are shown.

To this end, CD8+ cells either isolated directly from non-stimulated peripheral lymphocytes (quiescent) or from cultured mononuclear human lymphocytes stimulated with PHA *in vitro* and cultured for 10–14 days (proliferating) were irradiated with UVC and analyzed 6 h later or treated with curcumin for 8 h. We chose 50 μ M concentration of curcumin which had earlier been found to induce in human T cells and Jurkat cells the mode of cell death without oligonucleosomal DNA degradation (Piwocka *et al.*, 1999; Bielak-Zmijewska *et al.*, 2000; Sikora *et al.*, 2006).

Data presented in Fig. 2A show that 6 h after UVC irradiation more than 40% of proliferating CD8+ cells revealed significant DNA fragmentation and were present in the sub-G₁ fraction. However, the sub-G₁ fraction of curcumin-treated cells did not exceed 5% in either quiescent or proliferating cells. Similarly only a few percent of quiescent UVC-treat-

ed cells were found in the sub-G₁ subpopulation (Fig. 2A).

Apoptotic chromatin condensation temporally correlates with DNA fragmentation and frequently its efficiency depends on the degree of internucleosomal cleavage (Widlak *et al.*, 2003). Thus, we have analyzed nuclear morphology of CD8+ cells treated with curcumin or UVC-irradiated (Fig. 2B). Only proliferating UVC-irradiated cells revealed characteristic chromatin condensation and fragmentation as well as formation of apoptotic bodies resembling typical apoptosis, which could be expected from DNA content analysis. Quiescent CD8+ cells after UVC irradiation seemed to be quite healthy. Curcumin treatment resulted in chromatin condensation in almost all cells, quiescent or proliferating. However, unlike in UVC-irradiated proliferating cells, no apoptotic bodies were seen (Fig. 2B).

Curcumin induces caspase-3 activation followed by cleavage of its substrates

In the majority of apoptotic cells oligonucleosomal DNA degradation results from the activity of endonuclease DFF40/CAD. The nuclease is activated upon caspase-3-catalyzed cleavage of its inhibitor DFF45/ICAD (Enari *et al.*, 1998; Halenbeck *et al.*, 1998; Liu *et al.*, 1998). Thus, one could conclude from the results shown above that curcumin, in contrast to UVC, did not induce caspase-3 activation. To verify this assumption the presence of activated caspase-3 was analyzed in curcumin-treated or UVC-irradiated CD8+ either quiescent or proliferating cells using

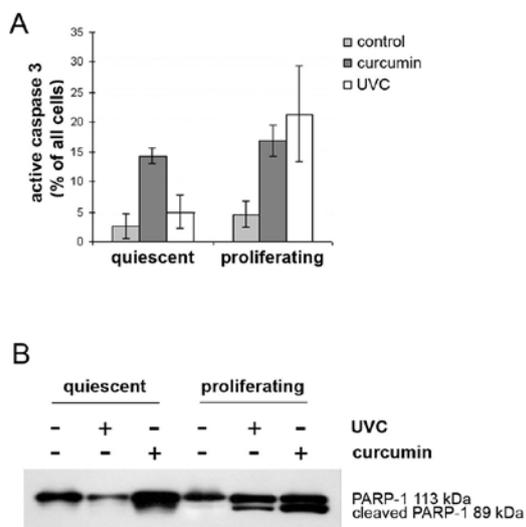


Figure 3. Active caspase-3 in quiescent and proliferating CD8+ cells at 8 h after treatment with 50 μ M curcumin.

(A) The number of cells that contained active form of caspase-3 was measured by flow cytometry and results are expressed as mean \pm S.D. of three independent experiments. (B) The truncated form of PARP-1 was assessed in whole cell lysates by Western blotting. A representative blot of three independent experiments is shown.

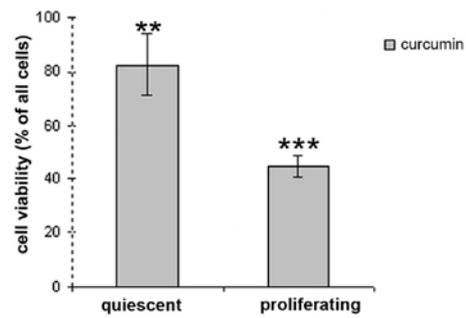


Figure 4. Viability of quiescent and proliferating CD8+ cells after treatment with 50 μ M curcumin.

Cell viability was measured after 8 h of curcumin treatment by the MTT assay (values are in the relation to untreated control). Data are expressed as mean \pm S.D. of three independent experiments. **, $P \leq 0.001$; ***, $P \leq 0.0001$.

flow cytometry (Fig. 3A). As expected the analysis revealed the presence of active caspase-3 in proliferating, but not quiescent CD8+ cells irradiated with UVC. Moreover, active caspase-3 was detected also in curcumin-treated cells, both quiescent and proliferating. The activity of caspase-3 was assessed in the same cells by detection of the 89 kDa fragment of PARP-1, which is a specific product of caspase-3 activity (Fig. 3B). The truncated form of PARP-1 could be detected in proliferating CD8+, both irradiated with UVC and treated with curcumin, as well as in quiescent curcumin-treated cells. Using MTT assay we showed that the percentage of survivals accounted for slightly more than 40% and 80% in curcumin-treated proliferating and quiescent cells, respectively (Fig. 4). These data show that in terms of survival, the impact of curcumin is stronger on proliferating than on quiescent CD8+ cells.

Curcumin induces degradation of DFF45/ICAD that leads to formation of the potentially active DFF40/CAD nuclease in CD8+ cells

In healthy non-apoptotic cells, the DFF40/CAD nuclease exists in the nucleus as a heterodim-

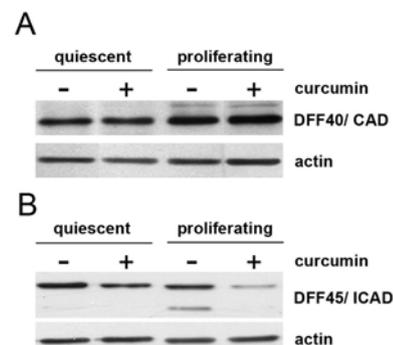


Figure 5. Levels of DFF subunits in CD8+ cells after 8 h of treatment with 50 μ M curcumin.

DFF40/CAD (A) and DFF45/ICAD (B) were assessed by Western blotting in whole cells lysates. The representative blots of three independent experiments are shown.

er with its inhibitor DFF45/ICAD (a 35-kDa splicing variant of DFF45-DFF35/ICAD-S-resides in the cytoplasm). Activation of caspase-3 results in the cleavage of DFF45/ICAD and release of DFF40/CAD, which forms active homo-oligomers (Liu *et al.*, 1999; Widlak *et al.*, 2003; Woo *et al.*, 2004). Here we show that DFF40/CAD is present in CD8+ cells, and its total level remains essentially unchanged in cells treated with UVC or curcumin as measured by Western blotting in whole cell lysates (Fig. 5A). In a marked contrast to the level of DFF40/CAD, the level of DFF45/ICAD decreased in curcumin-treated CD8+ cells, the decrease being less pronounced in quiescent cells than in proliferating ones (Fig. 5B). It is of note, that in quiescent CD8+ cells, the band of DFF35 was not present, but in proliferating cells, as expected, its disappearance was proportional to the decrease of the DFF45 band.

DISCUSSION

There are many facets to cancer prevention and one of them is using natural or synthetic compounds allowing suppression, retardation or inversion of carcinogenesis. Only a few such agents have been used to date in the clinic and these include non-steroidal anti-inflammatory drugs for colon, finasteride for prostate, and tamoxifen or reloxifene for breast tumors. An ideal chemopreventive agent should restore normal growth control to preneoplastic or cancerous cells by modifying aberrant signalling pathways or inducing cell death in cells beyond repair. The characteristics of such an agent include selectivity for transformed cells and more than one mechanism of action to foil the redundancy or crosstalk in signalling pathways (Manson *et al.*, 2005). It seems that curcumin fits this picture very well by affecting many signalling pathways, it is able to induce cell death in any tested cancer cells like leukaemia, melanoma, breast, lung, prostate, colon, renal, hepatocellular and ovarian carcinomas (reviewed in Karunakaran *et al.*, 2005), including those resistant to apoptosis due to the multidrug resistance phenotype (MDR) (Bielak-Mijewska *et al.*, 2004).

It appears from the available data that curcumin impact on cancer cells can be explained by its pleiotropic activity and many targets in the cell like COX-2 (Goel *et al.*, 2001; Shishodia *et al.*, 2003), HO-1 (Balogun *et al.*, 2003) or v-Src (Leu *et al.*, 2003). On the other hand, the multitude of curcumin's targets can also be explained by its influence on transcription regulation. Curcumin has been shown to inhibit the AP-1 transcription factor which is involved in apoptotic program and regulation of cell proliferation of many cells (Sikora *et al.*, 1997; Bharti *et al.*, 2004; Zheng *et al.*, 2004). Also NF- κ B involved in

pro-survival and apoptotic pathways is inhibited by curcumin (Deeb *et al.*, 2003; 2004). Recently curcumin has been shown to repress histone acetyltransferase-dependent chromatin transcription by inhibiting its p300/CREB-binding protein (Balasubramanyam *et al.*, 2004). p300 is a ubiquitously expressed global transcriptional coactivator that has a critical role in a wide variety of cellular phenomena including cell cycle control, differentiation and apoptosis (Giordano & Avantaggiati, 1999).

An open question is the curcumin's selectivity for transformed cells; alas, the data regarding curcumin impact on normal cells are rather scarce. Previously we showed that curcumin induced cell death in rat splenocytes as well as in human quiescent and proliferating T cells, but the cell death mode in terms of chromatin degradation and oligonucleosomal DNA fragmentation did not resemble typical apoptosis (Bielak-Zmijewska *et al.*, 2000). Also others described cell death in human V γ 9V Δ 2 T cells caused by curcumin, due to high molecular mass, but not oligonucleosomal DNA degradation (Cipriani *et al.*, 2001). Recently, the Gautam's group (Gao *et al.*, 2004) published data of an immunomodulatory activity of curcumin, namely its suppression of lymphocyte proliferation, development of cell-mediated cytotoxicity and cytokine production *in vitro*. Although curcumin at 30 μ M concentration irreversibly inhibited proliferation of splenocytes it did not affect cell viability measured by MTT test (Gao *et al.*, 2004). In our hands 50 μ M curcumin decreased proliferating cell viability to 40% already at 8 h as measured by MTT test. Also in a lower concentration, namely 25 μ M, curcumin was harmful to normal T cells (not shown).

It is believed that the side effects of anticancer therapy affect mainly proliferating cells. Indeed, previously (Radziszewska *et al.*, 1999) and in this paper we showed that UVC, which is a known DNA damaging agent, did not affect quiescent T cells, but induced apoptosis in proliferating ones. In contrast to UVC, curcumin induces cell death not only in proliferating but also in quiescent CD8+ cells, however, in a short-time treatment proliferating cells seem more sensitive to curcumin than quiescent ones. Moreover, the cell death mode induced by curcumin is distinct from that attributable to classical apoptosis whose main hallmark is oligonucleosomal DNA fragmentation resulting from caspase-3 activation.

Activation of caspase-3 led to proteolysis of its specific substrates including DFF45/ICAD, the inhibitor of the major apoptotic nuclease DFF40/CAD. One would assume that the release of the nuclease from its inhibitor should result in DNA cleavage. However, DNA fragmentation was not observed in curcumin-treated cells. Essentially the same effect was observed earlier in Jurkat cells. We have some

evidence that curcumin can inhibit DFF40/CAD by blocking magnesium binding in the active centre, thus preventing DNA fragmentation but not affecting cell death itself (Sikora *et al.*, 2006).

Curcumin is a very promising chemopreventive agent. It can be administered safely to patients at doses of up to 180 mg according to Sharma *et al.* (2001) and up to 8000 mg daily according to Cheng *et al.* (2001). However, curcumin has low oral bioavailability in humans and may undergo intestinal metabolism. At doses of 36–180 mg daily given for up to 4 months neither curcumin nor its metabolites were detected in blood or urine (Sharma *et al.*, 2001). Nonetheless, considering curcumin or curcumin derivatives' future use in the clinic and eventual intravenous application adverse side-effects could be expected such as death of T cells, including CD8+, and impairment of immune functions, e.g., suppression of proliferation, IL-2 production and decreased cytotoxicity of T cells (Gao *et al.*, 2004). Taking into account the role of CD8+ cells in tumor response, their depletion during chemotherapy could be particularly undesirable.

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