

Mycoplasma fermentans inhibits tumor necrosis factor α -induced apoptosis in the human myelomonocytic U937 cell line

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Abstract

Mycoplasma fermentans (*M. fermentans*) was shown to be involved in the alteration of several eukaryotic cell functions (i.e. cytokine production, gene expression), and was suggested as a causative agent in arthritic diseases involving impaired apoptosis. We investigated whether *M. fermentans* has a pathogenic potential by affecting tumor necrosis factor (TNF) α -induced apoptosis in the human myelomonocytic U937 cell line. A significant reduction in the TNF α -induced apoptosis (~60%) was demonstrated upon either infection with live *M. fermentans* or by stimulation with non-live *M. fermentans*. To investigate the mechanism of *M. fermentans* antiapoptotic effect, the reduction of mitochondrial transmembrane potential ($\Delta\Psi_m$) and the protease activity of caspase-8 were measured. In the infected cells, the reduction of $\Delta\Psi_m$ was inhibited (~75%), and a ~60% reduction of caspase-8 activity was measured. In conclusion, *M. fermentans* significantly inhibits TNF α -induced apoptosis in U937 cells, and its effect is upstream of the mitochondria and upstream of caspase-8.

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Abbreviations: AO, acridine orange; BSA, bovine serum albumin; CFU, colony-forming units; DiOC₆(3), 3,3'-dihexyloxa-carbocyanine iodide; EB, ethidium bromide; ELISA, enzyme-linked immunosorbent assay; FACS, FACSCalibur flow cytometer; FADD, fas-associated death domain; FCS, fetal calf serum; FITC, fluorescein isothiocyanate; mCiCCP, carbonyl cyanide *m*-chlorophenylhydrazone; MHC, major histocompatibility complex; *M. fermentans*, *Mycoplasma fermentans*; MOI, multiplicity of infection; NF- κ B, nuclear factor κ B; PBS, phosphate-buffered saline; PCR, polymerase chain reaction; PI, propidium iodide; PMSF, phenylmethylsulfonyl fluoride; PS,

phosphatidylserine; RA, rheumatoid arthritis; TNFR, tumor necrosis factor receptor; TNF α , tumor necrosis factor α ; $\Delta\Psi_m$, mitochondrial transmembrane potential

Introduction

Mycoplasmas, the smallest self-replicating wall-less bacteria, exist as parasites in humans, and were shown to interact with various cells of the immune system, either by activation or suppression, resulting in alteration of cell functions.¹ *Mycoplasma fermentans* (*M. fermentans*), a human pathogen, was shown to induce polyclonal activation of the immune cells, cytokine production, increased major histocompatibility complex (MHC) class II expression, increased cytotoxicity of T cells and expression of oncogenes.^{2–6} In the last decade, there have been increasing numbers of publications implicating *M. fermentans* in the development of rheumatoid arthritis (RA),^{7–9} a disease characterized by impaired apoptosis.^{8,10–12}

Apoptosis is induced in two different pathways: the intrinsic and extrinsic pathways. tumor necrosis factor α (TNF α), an inducer of the extrinsic pathway, crosslinks the TNF receptor (TNFR), which results in the activation of caspases, a class of cysteine proteases with specific functions in the execution of apoptosis.¹³ As a result of caspase activation, certain cellular substrates are cleaved and the cells undergo apoptosis.¹⁴

Several studies were published regarding mycoplasmas and apoptosis, but the results were contradictory. Mycoplasma proteins were shown to induce apoptosis in several cells,^{5,15} and it was shown that in Mycoplasma-contaminated cell cultures the internucleosomal DNA degradation is due to mycoplasmal endonuclease activity.¹⁶ Another study showed that mycoplasmal infections do not induce apoptosis, and even prevent it.¹⁷ In the present study, the indirect effect of mycoplasmal infection on apoptosis was investigated, by examining the effect of *M. fermentans* infection on apoptosis induced by TNF α , in the human myelomonocytic U937 cell line, as an experimental model.

Results

Infection of U937 cells with *M. fermentans* did not induce apoptosis in the cells

To investigate whether *M. fermentans* infection of U937 cells induces direct apoptosis of the cells, the viability of the cells was examined at 24 and 48 h post infection. The FACSCalibur flow cytometer (FACS) analysis of Annexin-V-fluorescein isothiocyanate (FITC)- and propidium iodide (PI)-stained cells (Figure 1a) and cell cycle (Figure 2) demonstrated no difference in the percent of apoptotic cells between non-infected and infected cells. Thus, under the conditions used,

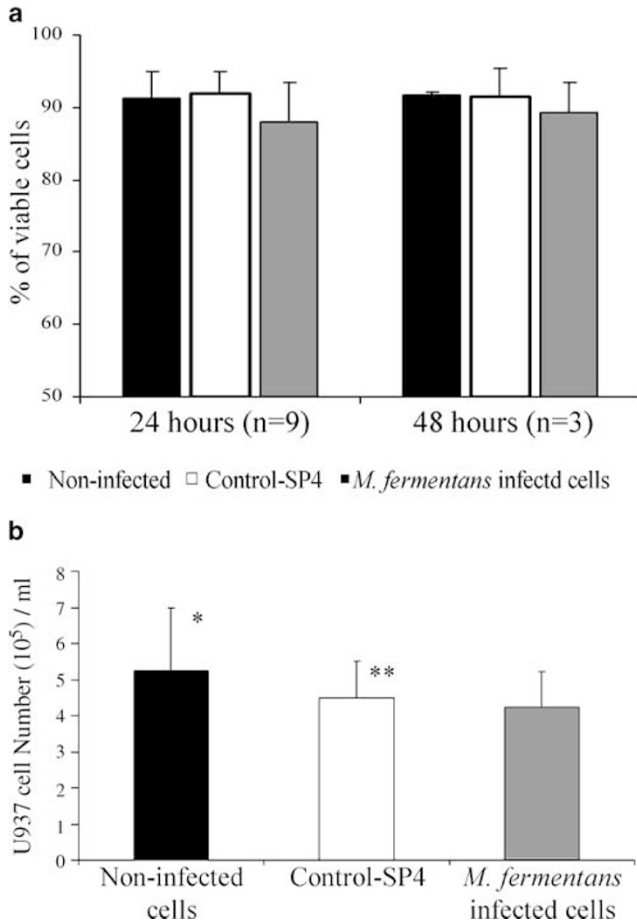


Figure 1 (a) Cell viability in *M. fermentans*-infected and noninfected U937 cells determined by Annexin-V-FITC-PI staining as analyzed by FACS. U937 cells were infected for 24 h ($n=9$) or 48 h ($n=3$) prior to cell staining. The mean percentage of live cells \pm standard deviation were as follows: $91 \pm 4\%$ (at 24 h) and $92 \pm 1\%$ (at 48 h), for noninfected cells (black); $92 \pm 3\%$ (at 24 h) and $91 \pm 4\%$ (at 48 h), for control culture treated with SP4 (*M. fermentans* medium) (white); and $88 \pm 6\%$ (at 24 h) and $89 \pm 4\%$ (at 48 h), for cells infected with *M. fermentans* (1000 CFU/cell) (gray). (b) Number of U937 cells in *M. fermentans*-infected and noninfected cultures, determined by trypan blue exclusion dye. U937 cells were infected for 24 h prior to cell staining (at a ratio of 1000 CFU/cells). The mean percentage of cell number $\times 10^5/\text{ml} \pm$ standard deviation was as follows: 5.2 ± 1.7 in noninfected cells; 4.5 ± 1 in SP4 control; 4.2 ± 1 in *M. fermentans*-infected cells (1000 CFU/cell). * $P < 0.05$, ** P – not significant

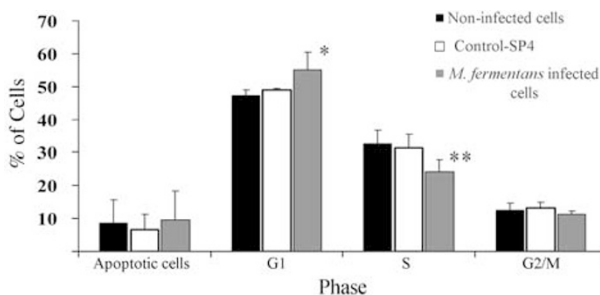


Figure 2 Cell cycle in *M. fermentans*-infected and noninfected U937 cells, determined by PI staining as analyzed by FACS. U937 cells were infected with *M. fermentans* (1000 CFU/cell) for 24 h prior to cell staining. The data are presented as the mean percentage \pm standard deviation of cells at different phases of the cell cycle, from three independent experiments. *M. fermentans* caused a G₁ arrest

infection of U937 cells with *M. fermentans* did not induce apoptosis.

Infection of U937 cells with *M. fermentans* causes a cell cycle arrest

To investigate whether *M. fermentans* infection of U937 cells induces proliferation of the cells, we determined, at 24 h post infection, the number and viability of the cells by trypan blue exclusion dye, and analyzed the cell cycle. As seen in Figure 1a, there was no significant difference in percentage of viable cells between the infected and noninfected cultures. By cell cycle analysis, a small but statistically significant decrease in the percentage of cells in S phase was observed ($P < 0.05$ for both controls), as well as a corresponding increase in the percentage of cells in G₁ phase ($P < 0.05$ for both controls) (Figure 2).

Confocal microscopy of U937 cells infected with *M. fermentans*

To determine the localization of *M. fermentans* with regard to U937 cells, we examined the cells by confocal microscopy. As seen in Figure 3, at 12 and 24 h post infection, most of *M. fermentans* were localized at the surface of U937 cells. Some *M. fermentans* were observed inside the cells.

TNF α -induced apoptosis is reduced in U937 cells infected with *M. fermentans*

Prior to investigating TNF α -induced apoptosis, the levels of TNF α secreted by U937 upon infection with *M. fermentans* were measured in the supernatants of the cultures, 24 h post infection. As shown in Figure 4, *M. fermentans* induced some secretion of TNF α , in a dose-response manner depending on the infection load ($P < 0.05$ for an infection ratio of 1000/1), but the amount secreted even at the highest colony-forming units (CFU)/cell ratio (the ratio we used in all experiments) is negligible (20 pg/ml) as compared to the amount we used to induce apoptosis (20 ng/ml). Indeed, as shown in Figure 1a, the infected cells, after 24 h, are not undergoing apoptosis.

The effect of *M. fermentans* on TNF α -induced apoptosis (20 ng/ml) in U937 cells was determined 24 h post infection. The rationale for choosing 24 h after infection emerged from the results obtained both in the confocal microscopy and in electron microscopy (data not shown). The percentage of apoptotic cells was examined by two techniques: (1) Acridine orange (AO)-ethidium bromide (EB) staining, which distinguishes between apoptotic and necrotic cells by the morphological changes in the nucleus (lack of DNA condensation in necrotic cells as opposed to apoptotic cells),¹⁸ was used to determine whether TNF α addition in our system induced apoptosis or necrosis. Firstly, TNF α addition induced apoptosis and not necrosis. Secondly, the apoptosis was reduced by ~60% in U937 cells infected with *M. fermentans* ($11.9 \pm 4.7\%$), in comparison to noninfected cells ($27.5 \pm 7.7\%$) and SP4-treated cells ($27.2 \pm 3.9\%$); $P < 0.05$ for both controls (Table 1); (2) A typical experiment of Annexin-V-FITC-PI staining is shown in Figure 5a. By this

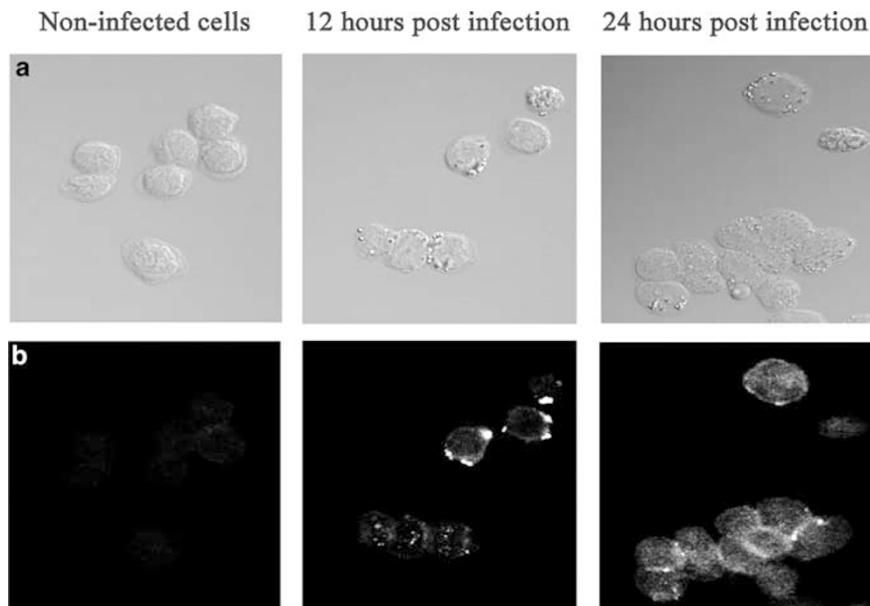


Figure 3 Confocal microscopy of *M. fermentans*-infected and noninfected U937 cells, at various infection times. Slides were prepared as described in Materials and methods. (a) Morphology of infected and noninfected cells. (b) The location of *M. fermentans* as detected with anti-*M. fermentans* antibodies

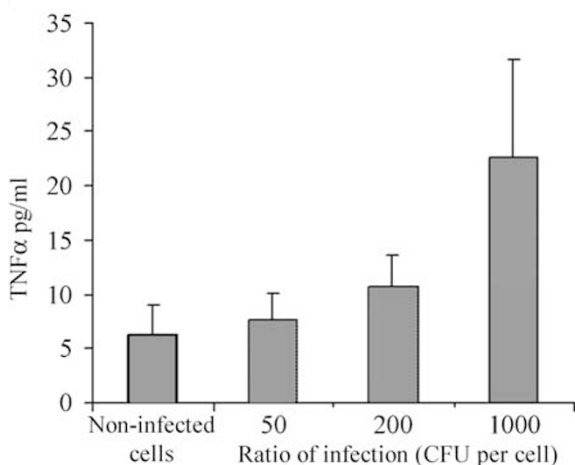


Figure 4 TNF α secretion by *M. fermentans*-infected and noninfected U937 cells, determined by ELISA. U937 cells were infected with *M. fermentans* for 24 h, and TNF α in the supernatants was measured. Mean percentage \pm standard deviation of TNF α levels in the different cell cultures, from four separate experiments, are depicted

technique, necrotic and late apoptotic cells cannot be distinguished from each other, but, relying on the AO-EB technique, we concluded that the double-positive cells (for Annexin-V-FITC and PI) represent late apoptotic cells and not necrotic ones. The results, obtained in nine independent experiments, showed approximately 60% reduction in the percentage of apoptotic U937 cells infected with *M. fermentans* at a ratio of 1000 CFU/cell ($6.5 \pm 2.8\%$), in comparison to noninfected cells ($19.8 \pm 4.3\%$) and SP4 control ($19.5 \pm 4.1\%$); $P < 0.01$ for both controls (Figure 5b). The specificity of the inhibitory effect of *M. fermentans* is

demonstrated by the magnitude of inhibition, which was dependent on the ratio of CFU/cell (Figure 5b).

Loss of mitochondrial inner transmembrane potential induced by TNF α is reduced in U937 cells infected with *M. fermentans*

In many apoptosis scenarios, including TNF α -mediated apoptosis, the mitochondrial inner transmembrane potential ($\Delta\Psi_m$) collapses.^{19,20} To investigate whether the antiapoptotic effect of *M. fermentans* in TNF α -induced apoptosis is upstream or downstream of the mitochondria, we measured the loss in $\Delta\Psi_m$, induced by TNF α (20 ng/ml), in infected and noninfected cells. At 24 h post infection, the cultures were stimulated with TNF α (20 ng/ml) for 2 h, and each culture was stained with 3,3'-dihexyloxycarbocyanine iodide (DiOC₆ (3)) and analyzed by FACS (a typical experiment is shown in Figure 6a).

In three independent experiments we found that, in U937 cells infected with *M. fermentans*, there was a fourfold reduction in $\Delta\Psi_m$ loss, $5.6 \pm 1.6\%$ cells with low DiOC₆(3) fluorescence dye, compared to noninfected cells ($23.7 \pm 7.2\%$) and SP4-treated cells ($22.6 \pm 2.2\%$); $P \leq 0.01$ for both controls (Figure 6b).

Protease activity of caspase-8 induced by TNF α is reduced in U937 cells infected with *M. fermentans*

Caspase-8 is at the apex of the caspase pathway and links death domain protein signaling (on TNF receptor) and caspase activation.¹⁴ To investigate whether the reduction of TNF α -induced apoptosis in infected cells is upstream or downstream of caspase-8, we measured the protease activity of caspase-8 induced by TNF α , in infected and noninfected

Table 1 Cell death induced by TNF α in *M. fermentans*-infected and noninfected U937 cells, as determined by fluorescence microscopy

Treatment	% of live cells	% of necrotic cells	% of apoptotic cells		
			Early	Late	Total
Noninfected U937	72.5 \pm 7.7	< 1	18.3 \pm 4.4	9.2 \pm 3.6	27.5 \pm 7.7
U937 treated with SP4	72.8 \pm 3.9	< 1	17.6 \pm 5.5	9.6 \pm 3.9	27.2 \pm 3.9
U937 infected with <i>M. fermentans</i>	88.1 \pm 4.7	< 1	9.0 \pm 3.1	2.9 \pm 2.9	11.9 \pm 4.7

Values, quantified by random counting (in duplicate) of acridine orange–ethidium bromide-stained cells, are mean \pm standard deviation of data from three independent experiments. Percentages represent the different stages of cell death. The decrease in TNF α -induced apoptosis, caused by *M. fermentans*, was \sim 60% ($P < 0.05$ for both controls as determined by ANOVA single factor)

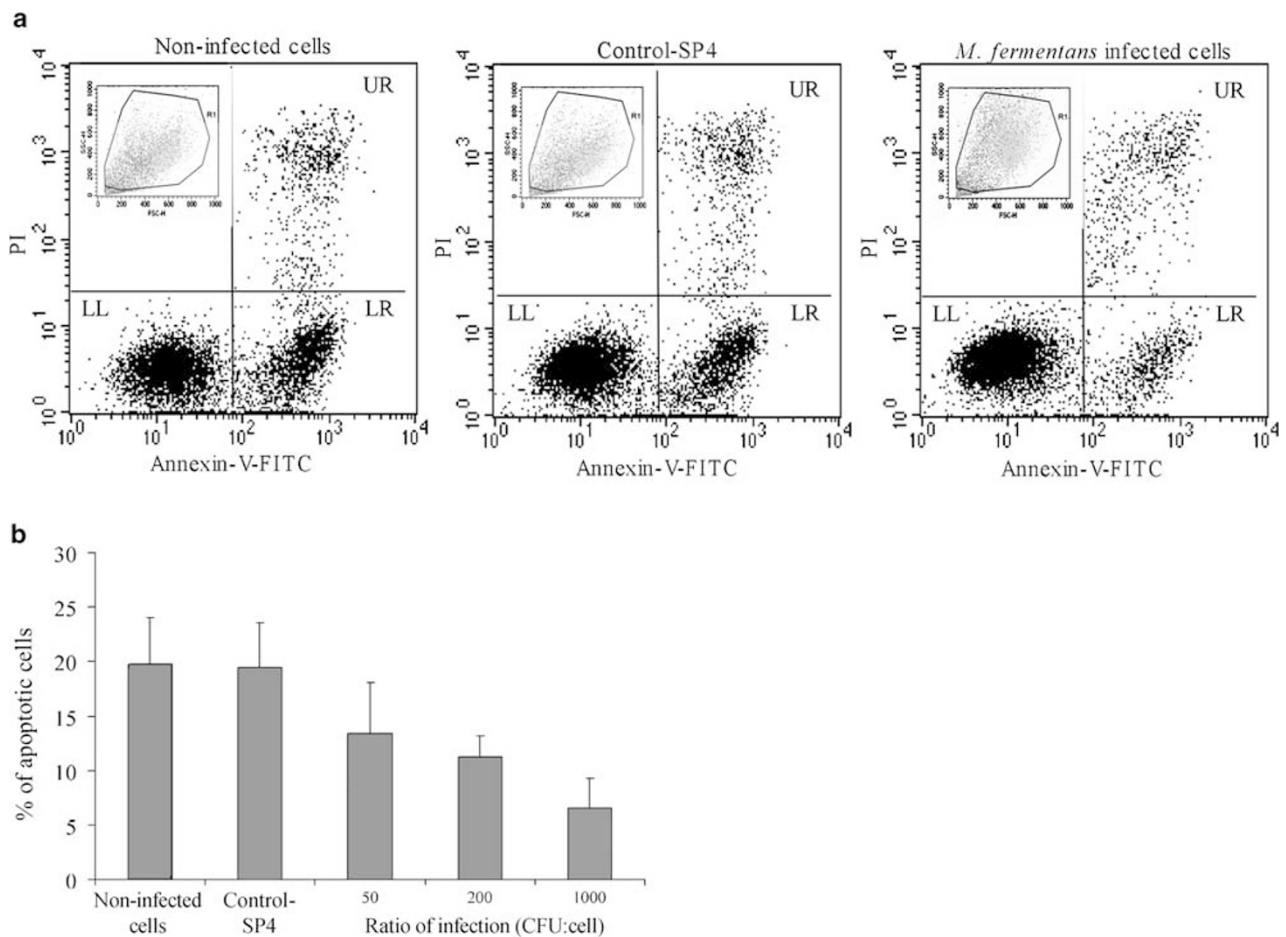


Figure 5 Measurement of TNF α -induced apoptosis by Annexin-V-FITC-PI staining in *M. fermentans*-infected and noninfected U937 cells. (a) A representative experiment: Cells were infected for 24 h (1000 CFU/cell), followed by addition of TNF α (20 ng/ml) for 4 h, stained with Annexin-V-FITC-PI and analyzed by FACS. Dot plots of forward and side scattered (with the gate used) are depicted on the upper-left corner of each treatment. Double negative staining represents live cells (LL), positive staining for Annexin-V-FITC and negative staining for PI represents early apoptotic stage (LR), and double-positive staining represents late apoptotic stage (UR). (b) Mean percentages of apoptotic cells \pm standard deviation were: 19.8 \pm 4.3% in noninfected cells, 19.5 \pm 4.1 in SP4 control, and 13.4 \pm 4.7, 11.2 \pm 2 and 6.5 \pm 2.8% in *M. fermentans*-infected cells (50, 200 and 1000 CFU/cell, respectively) ($P < 0.05$, $P < 0.01$ and $P < 0.01$ in *M. fermentans*-infected cells (50, 200 and 1000 CFU/cell, respectively) for both controls as determined by ANOVA single factor)

cells. At 24 h post infection, the cultures were stimulated with TNF α (20 ng/ml) for 4 h and cytosolic extracts were prepared. The assay for caspase-8 activity was performed as described in Materials and Methods.

A 60% reduction in protease activity of caspase-8 was measured in *M. fermentans*-infected U937 cells, compared to noninfected cells; $P < 0.05$ (Figure 7).

TNF α -induced apoptosis is reduced in U937 cells treated with non-live *M. fermentans*

To clarify whether the inhibitory effect of *M. fermentans* on TNF α -induced apoptosis was exerted by live or non-live mycoplasma, cells were stimulated with sonicated *M. fermentans* prior to induction of apoptosis by TNF α . The

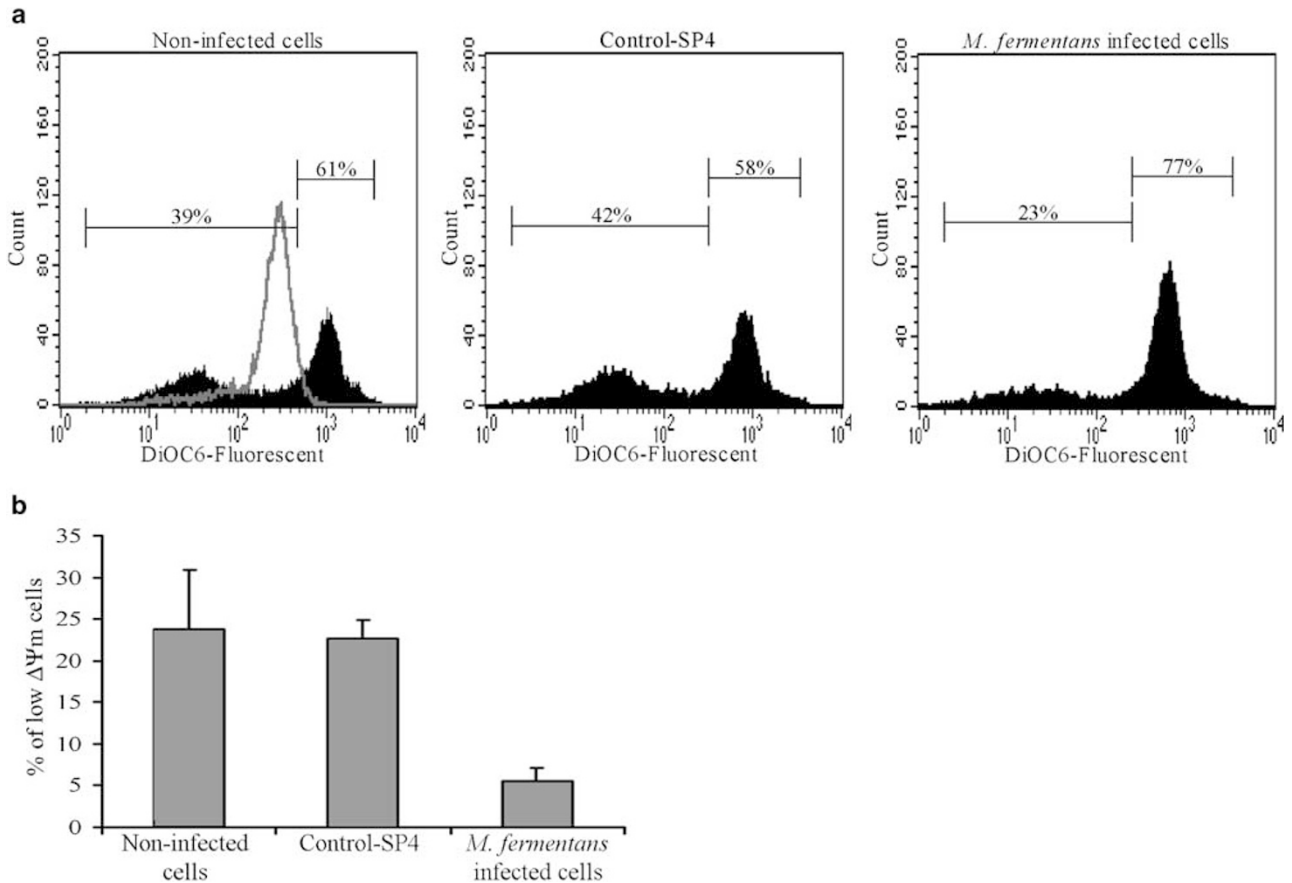


Figure 6 Determination of decrease in mitochondrial inner transmembrane potential induced by TNF α in *M. fermentans*-infected and noninfected U937 cells, by FACS of DiOC₆(3) fluorescence. (a) A representative experiment: Cells were infected for 24 h, followed by addition of TNF α (20 ng/ml) for 2 h, stained with DiOC₆(3) and analyzed by FACS. The percentage of cells with higher DiOC₆(3) staining (higher $\Delta\Psi_m$) is on the right of each histogram, while the percentage of cells with lower DiOC₆(3) staining (lower $\Delta\Psi_m$) is on the left. The line in the control histogram represents the positive control (mCiCCP). (b) Mean percentages of low DiOC₆(3) fluorescence dye \pm standard deviation ($n=3$) were: 23.7 \pm 7.2% in noninfected cells, 22.6 \pm 2.2% in SP4 control and 5.6 \pm 1.6% in *M. fermentans*-infected cells (1000 CFU/cell) ($P \leq 0.01$ for both controls, as determined by ANOVA single factor). Overall, a fourfold reduction of $\Delta\Psi_m$ loss was measured in *M. fermentans*-infected U937 cells

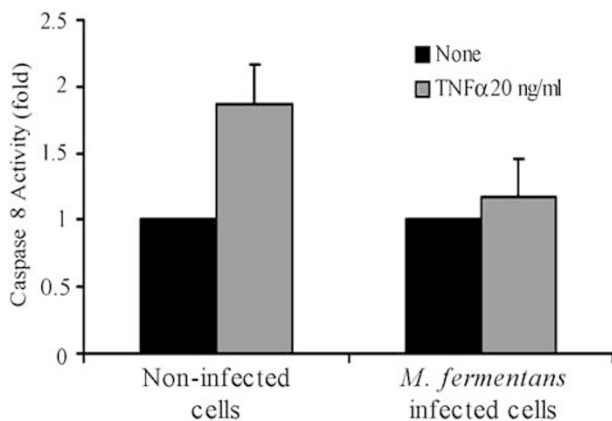


Figure 7 Protease activity of caspase-8 induced by TNF α in *M. fermentans*-infected and noninfected U937 cells. Cells were infected for 24 h, followed by addition of TNF α (20 ng/ml) for 4 h, and cytosolic extracts were prepared. Mean percentages of caspase-8 activity \pm standard deviation ($n=3$) were: 1.87 \pm 0.3-fold in TNF α -treated compared to untreated in noninfected cells, and 1.17 \pm 0.29-fold in TNF α -treated compared to untreated in *M. fermentans*-infected cells (1000 CFU/cell) ($P < 0.05$ as determined by ANOVA single factor). The decrease in protease activity of caspase-8 caused by *M. fermentans* was \sim 60%

percentage of apoptotic cells was examined by Annexin-V-FITC-PI staining. The results, obtained in four independent experiments, showed an \sim 60% reduction of apoptotic cells in U937 cells stimulated with 40 μ g/ml of sonicated *M. fermentans* (9.6 \pm 3.8%), in comparison to noninfected cells (23.7 \pm 4.7%) and phenylmethylsulfonyl fluoride (PMSF) control cells (26.3 \pm 3.8%); $P < 0.01$ for both controls (Figure 8). The inhibitory effect of sonicated *M. fermentans* on TNF α -induced apoptosis was dose dependent (Figure 8).

Discussion

In this article, we report that infection of human myelomonocytic U937 cell line with *M. fermentans* does not cause a direct cell death but, rather, inhibits TNF α -induced apoptosis. The block of apoptosis lies upstream of the mitochondria and upstream of caspase-8. It seems that the inhibitory effect of *M. fermentans* resides in constitutive components, since both live and non-live *M. fermentans* exert a similar effect.

M. fermentans has been shown to be both an extracellular and intracellular bacteria.^{7,21} We found, by confocal micro-

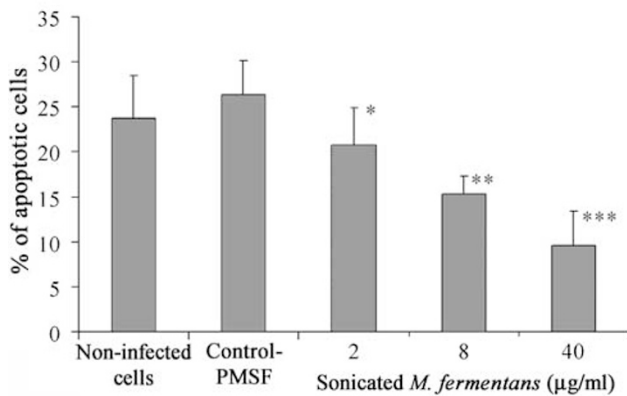


Figure 8 Measurement of TNF α -induced apoptosis by Annexin-V-FITC-PI staining in U937 cells treated with non-live sonicated *M. fermentans*. Cells were treated with non-live sonicated *M. fermentans* for 24 h, followed by addition of TNF α (20 ng/ml) for 4 h, stained with Annexin-V-FITC/PI and analyzed by FACS. Mean percentages of apoptotic cells \pm standard deviation were: 23.7 \pm 4.7% in noninfected cells, 26.3 \pm 3.8 in PMSF control, and 20.8 \pm 4.2%, 15.3 \pm 2 and 9.6 \pm 3.8% in cells treated with the sonicated *M. fermentans* (2, 8 and 40 μ g/ml respectively) (* P < 0.05, ** P < 0.01 and *** P < 0.01 for both controls as determined by ANOVA single factor)

scopy, that most *M. fermentans* cells were localized on the surface of U937 cells and that some of them became intracellular during the 8–24 h post infection (Figure 3). These findings suggest that *M. fermentans* might affect the cell either from the outer side of the membrane, or by stimulating the cell from its location within the cell.

The effect of mycoplasmas on apoptosis was previously examined in several studies.^{5,15–17} Most of the studies were engaged with the direct effect of Mycoplasma or mycoplasma components on the apoptotic processes in infected cells. *M. fermentans* proteins were shown to induce apoptosis in various cell lines.^{5,15} In contrast, it was shown that mycoplasma infections (under similar conditions, for example, multiplicity of infection (MOI)) do not induce apoptosis, and even prevent it.¹⁷ In the present study, we found that infection of U937 cells with *M. fermentans* for 24 or 48 h did not induce apoptosis (Figures 1a and 2), and induced an incomplete arrest in the cell cycle at G₁ (Figure 2), as was previously reported.⁶ The contradiction between the studies might be explained by the fact that non-live mycoplasma derivatives were used when apoptosis was demonstrated,^{5,15} whereas, when live mycoplasmas were employed, apoptosis was not induced.¹⁷ In another study, it was reported that Mycoplasma contamination of astrocytes (using similar MOI) could cause apoptosis, as a result of choline depletion in the medium, due to the mycoplasma nutritional requirements rather than a direct effect of Mycoplasma on the cells.²² Indeed, Rawadi *et al*⁶ reported that the proapoptotic effect resides exclusively in the nonlipid protein fraction of *M. fermentans* and not in the lipid fraction; this might explain why live Mycoplasma (enveloped only by a lipid membrane) do not cause apoptosis or cell death. In our system, U937 cells were infected with live *M. fermentans*. Although we were not able to grow *M. fermentans* from the culture 24 h post infection (data not shown), it is likely that U937 cells came into contact with *M.*

fermentans membrane lipoproteins. This might explain why, in our system, there was no apoptosis during the 24 or 48 h post infection.

To our knowledge, this is the first report presenting the ability of Mycoplasma to inhibit TNF α -induced apoptosis. Only a few papers have been published regarding inhibition of TNF α -induced apoptosis by *Chlamydia pneumoniae*^{23,24} and Hepatitis C Virus.²⁵ These support the hypothesis²⁶ that parasite bacteria might use a mechanism to prevent host cell apoptosis, in order to promote their survival and replication.

In TNF α -mediated apoptosis, the $\Delta\Psi_m$ collapses, resulting in cytochrome *c* release.^{19,20} A fourfold reduction in $\Delta\Psi_m$ loss was observed in infected cells as compared to noninfected cells (Figure 6). Therefore, we concluded that the effect of *M. fermentans* is upstream of the mitochondria. This result is similar to a recent observation regarding the effect of *Chlamydia pneumoniae* infection,^{23,24} which was reported to render epithelial and monocyte cell lines resistant to TNF α -induced apoptosis, via blockage of mitochondrial cytochrome *c* release.

It is well known that the Bcl-2 protein family regulates the $\Delta\Psi_m$ loss in apoptosis,^{27,28} but, as yet, we have no evidence that these proteins are upregulated or downregulated in our infected cells (data not shown).

Binding of TNF α to the TNFR1 causes fas-associated death domain (FADD) to bind the receptor. Procaspase-8 binds the receptor-bound FADD, leading to its proteolytic activation. This, in turn, can lead to an apoptotic process via two branches, either directly by activation of caspase-3 (extrinsic), or by release of cytochrome *c* from the mitochondria (intrinsic).¹³ By comparing caspase-8 activity (induced by TNF α) in infected and noninfected cells, we found that there was a 60% reduction of protease activity of caspase-8 in infected cells, indicating that the inhibitory effect of *M. fermentans* on TNF α -induced apoptosis is upstream of caspase-8 (Figure 7).

Our finding that *M. fermentans* inhibit TNF α -induced apoptosis might be explained by the effect of *M. fermentans* on the transcription factor, nuclear factor κ B (NF- κ B). NF- κ B activation is known to inhibit TNF α -induced apoptosis via suppression of caspase-8 activation.²⁹ It was previously reported that *M. fermentans* cause NF- κ B activation in human monocytic cell line and murine macrophages.^{5,17,30} In a preliminary study, we found that, in our system, NF- κ B was translocated to the nucleus in *M. fermentans*-infected cells (data not shown).

Alternatively, since most of *M. fermentans* were associated with the U937 cell membrane (Figure 3), the notion that the antiapoptotic effect of *M. fermentans* is exerted via modulation of TNF receptors has surfaced. This is currently under investigation in our laboratory, and preliminary results indicate that there is no significant difference in TNFR1 expression between infected and noninfected cells (data not shown).

The inhibition of TNF α -induced apoptosis was also evident when cells were stimulated by non-live *M. fermentans*, similar to the effect of infection with live *M. fermentans* (Figure 8). Although we infected cells with live *M. fermentans*, we cannot conclude whether the inhibition of TNF α -induced apoptosis was due to live or non-live Mycoplasma, since live *M. fermentans* could not be cultivated after 24 h. However, in

preliminary experiments, when live *M. fermentans* were recovered from the culture, 2 h post infection, the antiapoptotic effect was already demonstrated (data not shown). These results suggest that both live and non-live *M. fermentans* exerted inhibition of the TNF α -induced apoptosis.

In summary, the findings that *M. fermentans* causes a significant inhibition of TNF α -induced apoptosis in U937 cells strengthen the hypothesis that *M. fermentans* might play a pathogenic role in the development of diseases characterized by impaired apoptosis. These findings also imply that the presence of mycoplasmas in various cell lines might be affecting the results of apoptosis research.³¹

Materials and Methods

Reagents and antibodies

Recombinant TNF α and TNF α determination kit (enzyme-linked immunosorbent assay (ELISA)) were purchased from R&D Systems, Inc. (Minneapolis, MN, USA). Annexin-V-FITC and PI were purchased from Bender MedSystems (Vienna, Austria). The 3,3'-dihexyloxycarbocyanine iodide (DiOC₆(3)) and goat-anti-mouse conjugated to Alexa-Fluor 488 were purchased from Molecular Probes, Inc. (Junction City, OR, USA). Carbonyl cyanide *m*-chlorophenylhydrazine (mCICCP), EB, AO and PMSF were purchased from Sigma Chemical Co. (St. Louis, MO, USA). The caspase-8 Assay Kit was obtained from Calbiochem (Nottingham, UK). Cell culture media and supplements were purchased from Biological Industries Ltd (Beit Haemek, Israel).

Mycoplasma culture and mycoplasmal non-live preparation

M. fermentans K7 (originally obtained from JG Tully (National Institute of Allergy and Infectious Diseases, National Institutes of Health, Bethesda, MD, USA)) was cultured in SP4 broth containing 8.5% fetal calf serum (FCS).^{32,33} Aliquots (1 ml) (containing 4×10^8 CFU) were frozen at -70°C . For each experiment, one aliquot was thawed and cultivated in SP4 broth first, at a dilution of 1/10 for 24 h (37°C), and then stepped up at a 1/50 dilution for another 24 h, when log phase was observed. *M. fermentans* was quantified by counting of CFU on SP4 agar plates.^{32,33}

Non-live Mycoplasmas were prepared as follows: *M. fermentans* was cultivated as above and the 500 ml culture was pelleted ($13\,000 \times g$, 30 min at 4°C), and washed twice with phosphate-buffered saline (PBS). The pellet was then resuspended in 2 ml of PBS. CFU/ml was determined and then stored at -70°C for further experiments. Frozen pellets of *M. fermentans* were thawed and sonicated at 4°C for 4×30 s at 80% power (50% duty cycle; Heat Systems Ultrasonics, Inc.) in the presence of the protease inhibitor PMSF (10^{-4} M). These conditions of sonication resulted in a non-live mycoplasmal preparation (no growth was observed in repeated culturing procedures). Protein concentration of sonicated *M. fermentans* was determined by using the Bio-Rad protein assay kit (Richmond, CA, USA). An amount of 1×10^8 CFU corresponds to $10 \mu\text{g}$ of mycoplasmal proteins.

Cell culture

The human myelomonocytic U937 cell line was cultivated at 37°C and 5% CO₂, in RPMI 1640 culture medium containing 10% FCS, 1% HEPES, 1% penicillin–streptomycin and 1% glutamine. The cell line was tested every 4

weeks by a polymerase chain reaction (PCR)-based detection assay for *M. fermentans* contamination.⁹

Stimulation of U937 cells with live *M. fermentans* or sonicated *M. fermentans*

Prior to infection with live *M. fermentans*, Mycoplasmas (at a log phase) were washed once in PBS ($13\,000 \times g$, 30 min at 4°C), and resuspended in RPMI 1640 medium containing 10% FCS, 1% HEPES, 1% penicillin–streptomycin and 1% glutamine. Then, *M. fermentans* was added to U937 cells, pre-washed once with PBS ($500 \times g$, 10 min at 4°C), and resuspended in a new culture medium at a final concentration of 2×10^5 cells/ml. The ratio of Mycoplasmas/cell to be used was pre-determined by a series of dose–response experiments, at various ratios (50/1, 200/1, 1000/1). A ratio of 1000 CFU/cell was chosen since, under these conditions, the effect of *M. fermentans* on cells was more pronounced. The same procedure, with the same volume as in the *M. fermentans* culture, was performed with SP4 broth to be used as on the SP4 control. For experiments with non-live *M. fermentans*, the sonicated Mycoplasma (prepared as described above) was added to U937 (2×10^5 cells/ml) cells at concentrations of 2, 8 and $40 \mu\text{g}/\text{ml}$. PMSF control is a noninfected cell culture with the addition of PMSF, at the same concentration as in the sonicated *M. fermentans*.

Cell cycle analysis

At 24 h post infection, U937 cultures were washed and fixed with 1 ml of 70% cold ethanol. Cells were kept in the fixative for at least 24 h at -20°C . Followed by centrifugation and removal of the ethanol, pellets were resuspended in $900 \mu\text{l}$ of a solution containing 0.1% Triton X-100 and RNase ($10 \mu\text{g}/\text{ml}$) in PBS. The cells were kept for 40 min at room temperature, and stained with $100 \mu\text{l}$ of PBS containing PI ($150 \mu\text{g}/\text{ml}$). After incubation on ice for 10 min, the cells were analyzed by a FACS (Becton-Dickinson, Mansfield, MA, USA). The population of cells in each cell cycle phase was determined by using ModFit LT software (Becton Dickinson Immunocytometry Systems).

Confocal microscopy

U937 cells were infected with *M. fermentans*, as described above, for different time periods (0, 1, 2, 4, 12 and 24 h). Each culture was fixed with 3.7% formaldehyde (in Eppendorf tubes) for 30 min at room temperature and washed twice with PBS. Cells were then permeabilized by 0.2% Triton X-100 for 5 min at room temperature, washed twice with PBS, and blocked with 1% bovine serum albumin (BSA). Cells were then incubated with anti-*M. fermentans* hyperimmune mouse serum, prepared as described elsewhere,³⁴ for 40 min at room temperature. Then, the cells were washed three times with PBS, incubated with goat-anti-mouse conjugated Alexa-Fluor 488 for 40 min at room temperature and washed three times with PBS. The labeled cells were mounted on slides and examined by confocal laser-scanning microscopy (LSM410, Carl Zeiss, Jena, Germany).

Measurement of TNF α secreted from U937

At 24 h post infection of U937 cells with *M. fermentans*, as described above, the supernatants were collected by centrifugation ($500 \times g$ for 10 min) and stored at -20°C . TNF α ELISA was performed according to the manufacturer's instructions.

Induction of apoptosis

Infected U937 cells (24 h post infection) were counted, examined for viability by the trypan blue exclusion dye, centrifuged at $500 \times g$ for 10 min, and transferred to a new culture medium at a final concentration of 4×10^5 cells/ml. The cells were divided in 24-well sterile plates (Corning, Corning, NY), and apoptosis was induced by TNF α at a concentration of 20 ng/ml.

Measurement of apoptosis

Two different techniques were used: (1) AO-EB staining, an exclusion dye method, enables the differentiation between live, early-apoptotic, late-apoptotic and necrotic cells.¹⁸ At 8 h post induction of apoptosis by TNF α , infected and noninfected cells (4×10^5) were collected, centrifuged at $500 \times g$ for 10 min, and resuspended in 100 μ l PBS. Samples of 25 μ l from each culture were stained with AO-EB (final concentrations 1 μ g/ml for each AO and EB), and observed under a fluorescence microscope. At least 200 cells were randomly counted in each sample (in duplicates), and the percentage of apoptotic cells was calculated; (2) Annexin-V-FITC-PI staining,³⁵ which detects the exposure of phosphatidylserine (PS) to the external leaflet of the plasma membrane in early apoptosis, enables to differentiate between live, early apoptotic and dead cells (it does not differentiate between late apoptotic and necrotic cells).³⁶ At 4 h post induction of apoptosis by TNF α , infected and noninfected cells (2×10^5) were washed with PBS, centrifuged and resuspended in 400 μ l of a binding buffer (10 mM HEPES/NaOH, 140 mM NaCl, 2.5 mM CaCl₂, pH 7.4). Annexin-V-FITC (5 μ l; 10 μ g/ml) was added to a sample of 195 μ l of cell suspension, mixed, incubated for 10 min at room temperature in the dark, washed with PBS and resuspended in 190 μ l of binding buffer containing 10 μ l of PI (1 μ g/ml). The double-stained cells were analyzed by the FACS, within 10 min (10 000 cells/sample).

In all measurements of apoptosis, we determined the percent of cells that underwent apoptosis by TNF α , by subtracting the percent of spontaneous apoptosis (unstimulated cells) from the total apoptosis (stimulated cells).

Measurement of mitochondrial inner transmembrane potential ($\Delta\Psi_m$)

At 2 h post induction of apoptosis by TNF α , a total of 8×10^5 infected and noninfected cells were washed with PBS and resuspended in 990 μ l of PBS containing 10 μ l of DiOC₆(3) (final concentration 40 nM). Cells were mixed, incubated for 30 min at 37°C in the dark, and kept on ice until analyzed by the FACS within 10 min (10 000 cells per sample). Cells with low DiOC₆(3) fluorescence represent cells with permeabilized mitochondria. A positive control, comprising 8×10^5 cells with 5 mM of mCiCCP, a mitochondrial uncoupler and a known reducer of $\Delta\Psi_m$,³⁷ was incubated for 30 min at 37°C prior to the addition of DiOC₆(3).

Measurement of protease activity of caspase-8

At 4 h post induction of apoptosis by TNF α , a total of 1.6×10^6 infected and noninfected cells were washed with PBS, resuspended in 80 μ l of ice-cold cell lysis buffer (50 mM HEPES, 1 mM DTT, 0.1 mM EDTA, 0.1% CHAPS, 0.1% Tween20, pH 7.4), and incubated for 5 min on ice. The resulting cell lysates were centrifuged at $13\,000 \times g$ for 10 min at 4°C and the supernatant (cytosolic fraction) was kept at -70°C until analyzed. The test was performed according to the manufacturer's protocol (Calbiochem). Briefly, a sample of 15 μ l of each cell lysate was added to a well of

a microtiter plate (Corning), containing 75 μ l of assay buffer (100 mM NaCl, 50 mM HEPES, 10 mM DTT, 1 mM EDTA, 0.1% CHAPS, 10% glycerol, pH 7.4). The positive control in this assay was a recombinant human caspase-8. To validate the caspase-8 activity in the lysate, a negative control for each sample was analyzed. This control was an identical lysate sample to which 20 μ l of Granzyme B Inhibitor II (Ac-IETD-CHO), final concentration 50 ng/ml, was added. The plate was incubated at 37°C for 10 min. Then, the reaction was initiated by adding 10 μ l of Granzyme B Substrate I, Colorimetric (Ac-IETD-pNA), final concentration 200 mM, and the change in absorbance was monitored at 405 nm for 2 h, at 10-min intervals.

Protein concentration of all lysates was determined by using the Bio-Rad protein assay kit (Richmond, CA, USA), and protease activity of caspase-8 was calculated as a specific activity (pmol/min/ μ g protein).

Statistical analysis

The statistical significance of differences between groups was performed by analysis of variance ANOVA: single factor, given in *P*-value. *P*-values of <0.05 were considered statistically significant; *P*-values of <0.01 were considered highly significant.

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