

Neutrophil apoptosis mediated by nicotinic acid receptors (GPR109A)

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G protein-coupled receptor (GPR)109A (HM74A) is a G_i protein-coupled receptor, which is activated by nicotinic acid (NA), a lipid-lowering drug. Here, we demonstrate that mature human neutrophils, but not eosinophils, express functional GPR109A receptors. The induction of the GPR109A gene appears to occur late in the terminal differentiation process of neutrophils, since a mixed population of immature bone marrow neutrophils did not demonstrate evidence for its expression. NA accelerated apoptosis in cultured neutrophils in a concentration-dependent manner, as assessed by phosphatidylserine redistribution, caspase-3 activation, and DNA fragmentation assays. The pro-apoptotic effect of NA was abolished by pertussis toxin, which was used to block G_i proteins, suggesting a receptor-mediated mechanism. Activation of GPR109A by NA resulted in decreased levels of cyclic adenosine monophosphate (cAMP), most likely due to G_i-mediated inhibition of adenylyl cyclase activity. NA-induced apoptosis was reversed by the addition of cell-permeable cAMP, pointing to the possibility that reduced cAMP levels promote apoptosis in neutrophils. Distal mechanism involved in this process may include the post-translational modification of members of the Bcl-2 family, such as dephosphorylation of pro-apoptotic Bad and antiapoptotic Mcl-1 proteins. Taken together, following maturation in the bone marrow, neutrophils express functional GPR109A receptors, which might be involved in the regulation of neutrophil numbers. Moreover, this study identified a new cellular target of NA and future drugs activating GPR109A receptors, the mature neutrophil.

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The G_i protein-coupled receptor GPR109A (HM74A in humans; PUMA-G in mice) has been identified as a receptor for nicotinic acid (NA).^{1,2} NA is used to modify lipoprotein profiles in humans, resulting in a beneficial modification of multiple cardiovascular risk factors.³ This highly desirable therapeutic profile of NA is due to its direct activation of GPR109A on the surface of adipocytes.¹ However, GPR109A mRNA is also highly expressed in spleen and lung tissues,² but it remains unclear which cells indeed express the receptor and whether it is functional. Recently, it has been shown that interferon- γ is able to induce GPR109A receptors in macrophages, and that NA is subsequently able to stimulate prostaglandin E₂ and D₂ synthesis in these cells.⁴ The identification of functional GPR109A receptors in a cell outside adipose tissue raised the question whether additional cell types carry GPR109A receptors, which might also be targeted during therapy with NA.

Neutrophils are a critical component of the innate immune system with several effector and immunoregulatory functions. Their numbers are largely regulated by production in the bone marrow and death, which is commonly apoptosis.⁵ Delayed neutrophil apoptosis has been reported in association with several infectious diseases and is considered as a major mechanism of tissue neutrophilia. Overexpression of neutro-

phil survival factors, such as granulocyte colony-stimulating factor (G-CSF) and granulocyte-macrophage colony-stimulating factor (GM-CSF), play a major role in this process.^{6–9} These survival cytokines have been described to block the mitochondria-dependent death pathway in neutrophils.⁸ This is also reflected by the observation that these cytokines regulate the expression of Bcl-2 family members in these cells.⁵

In chronic inflammatory diseases, persistent accumulation of inflammatory cells may lead to increased release of toxic intracellular products, leading to tissue injury and subsequent remodeling with potential consequences on organ function. Therefore, it is important to study how inflammation normally resolves and under which conditions this process does not work. Although some surface receptors on neutrophils have been identified that transduce death signals, such as Fas¹⁰ and Siglec-9,¹¹ other death ligands may additionally contribute to the limitation of neutrophil numbers under inflammatory conditions. Additional knowledge in this field may provide novel approaches to anti-inflammatory pharmacotherapy.

In this article, we report the expression of functional GPR109A receptors on the surface of neutrophils, supporting the concept that they play a role in immunity and inflammation.

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Abbreviations: [Ca²⁺]_i, cytosolic-free calcium concentrations; cAMP, cyclic adenosine monophosphate; GAPDH, glyceraldehyde-3-phosphate dehydrogenase; G-CSF, granulocyte colony-stimulating factor; GM-CSF, granulocyte-macrophage colony-stimulating factor; GPR, G protein-coupled receptor; mAb, monoclonal antibody; NA, nicotinic acid; PKA, protein kinase A; PS, phosphatidylserine; PTX, pertussis toxin

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Moreover, GPR109A receptors are linked to a pro-apoptotic pathway in these cells, suggesting that NA may also mediate anti-inflammatory effects besides its inhibitory function on adipocyte lipolysis.

Results

Mature neutrophils, but not eosinophils or immature neutrophils, express GPR109A surface receptors. We investigated GPR109A surface expression on purified granulocyte population using indirect immunofluorescence and flow cytometric analysis. Freshly purified mature blood neutrophils, but not eosinophils or immature bone marrow neutrophils, expressed detectable GPR109A surface expression (Figure 1a). In addition, immature neutrophilic HL-60 cells were also GPR109A negative. Stimulation of mature neutrophils with GM-CSF slightly increased GPR109A expression (data not shown). Expression of GPR109A on mature neutrophils was confirmed using confocal microscopy. Both indirect immunofluorescence analysis and direct staining with FITC-conjugated NA demonstrated ring-like staining patterns, suggesting that GPR109A is predominantly expressed on the cell surface of these cells (Figure 1b).

Functional GPR109A receptors are present on mature neutrophils, but not on eosinophils or immature neutrophils. Expression of a functional GPR109A receptor

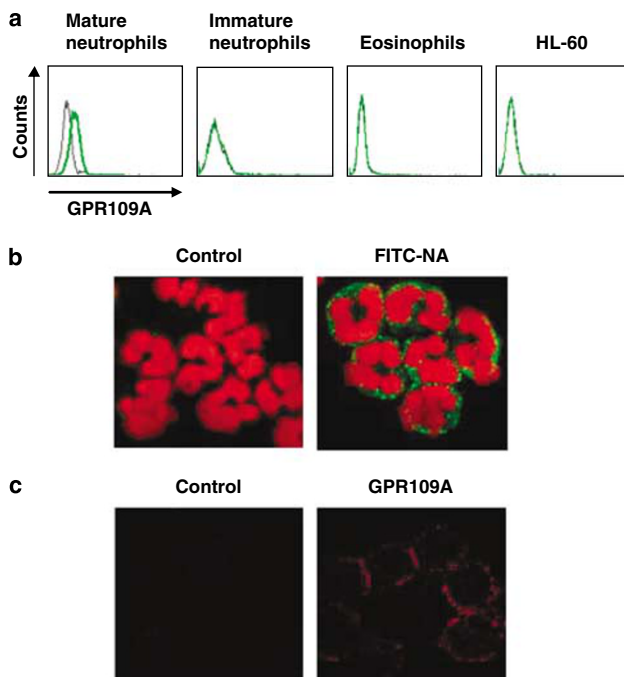


Figure 1 Mature blood neutrophils, but not immature bone marrow neutrophils or mature blood eosinophils, express the NA receptor GPR109A on their surface. (a) Flow cytometry. Cells were stained with control (black) or anti-GPR109A (green) mAbs. (b and c) Confocal microscopy. Neutrophils were stained with FITC-conjugated NA and anti-GPR109A mAb, respectively. Nuclei were stained with PE in (b). Results in each panel are representative of at least four independent experiments

was assessed by the ability of NA to specifically induce increases in cytosolic-free calcium concentrations ($[Ca^{2+}]_i$).¹ Addition of NA to fura-2-loaded mature neutrophils led to rapid and transient changes in $[Ca^{2+}]_i$ (Figure 2a). Peak calcium levels were observed within 1 min of addition of NA. Levels rapidly declined thereafter reaching baseline levels within 5 min. The increases in $[Ca^{2+}]_i$ were concentration dependent in the range of 10^{-7} – 10^{-3} M (Figure 2a). Furthermore, triggering of changes in $[Ca^{2+}]_i$ were caused by NA binding to its specific receptor because nicotinamide, a structurally related compound, which does not bind to GPR109A receptors,¹² had no effect on $[Ca^{2+}]_i$ at concentrations ranging from 10^{-7} to 10^{-3} M (data not shown). Moreover, immature bone marrow neutrophils, mature blood eosinophils, and HL-60 cells, which all did not express detectable levels of GPR109A surface protein, did not respond with increases in $[Ca^{2+}]_i$ following addition of NA. In contrast, addition of UTP, a ligand of P2Y₂ receptors expressed in hematopoietic cells including CD34⁺ stem cells,¹³ led to rapid and transient changes in $[Ca^{2+}]_i$ (Figure 2b), demonstrating that the cells were properly loaded and responsive, in principle.

To complement our studies of purified cell populations, we utilized a second functional approach analyzing $[Ca^{2+}]_i$ changes in single mature neutrophils. Interestingly, all cells responded similarly following activation with NA, regarding time and magnitude of the change in $[Ca^{2+}]_i$ (Figure 2c). These data suggested that all mature neutrophils express functional GPR109A receptors and also excluded the possibility that the NA-mediated effects on $[Ca^{2+}]_i$ seen using the bulk spectrofluorometric assay were due to GPR109A-expressing contaminating cells.

Pertussis toxin (PTX) is a known inhibitor of G_i proteins. Pre-incubation of mature neutrophils with PTX inhibited NA-induced increases in $[Ca^{2+}]_i$ in a concentration- and time-dependent manner (Figure 2d). In contrast, UTP-induced increases in $[Ca^{2+}]_i$ were not affected by PTX, consistent with the view that P2Y₂ receptors are able to stimulate, besides G_i, additional G proteins.¹⁴ Taken together, these data suggest that mature neutrophils express functional GPR109A receptors, which can be blocked by PTX. Moreover, increases in $[Ca^{2+}]_i$ induced by 10^{-5} M NA are an accurate and sensitive indicator for the presence of functional GPR109A receptors.

NA accelerates apoptosis in cultured mature neutrophils, but not in eosinophils or immature neutrophils. Mature neutrophils and eosinophils are known short-lived cells and undergo rapid apoptosis *in vitro*.⁵ In contrast, immature neutrophils exhibit a much longer life-span *in vitro*, although they also undergo cell death at later time points.^{15,16} Addition of NA to purified cell populations resulted in a detectable cytotoxic effect in mature neutrophils, but not in eosinophils or immature neutrophils (including HL-60 cells), in a concentration-dependent manner in 24-h cultures (Figure 3a). Longer culture times (48-h) revealed same results, except that spontaneous death of mature granulocytes was more advanced (data not shown). Concentrations of 10^{-5} and 10^{-4} M NA significantly accelerated the known spontaneous death in mature neutrophils.

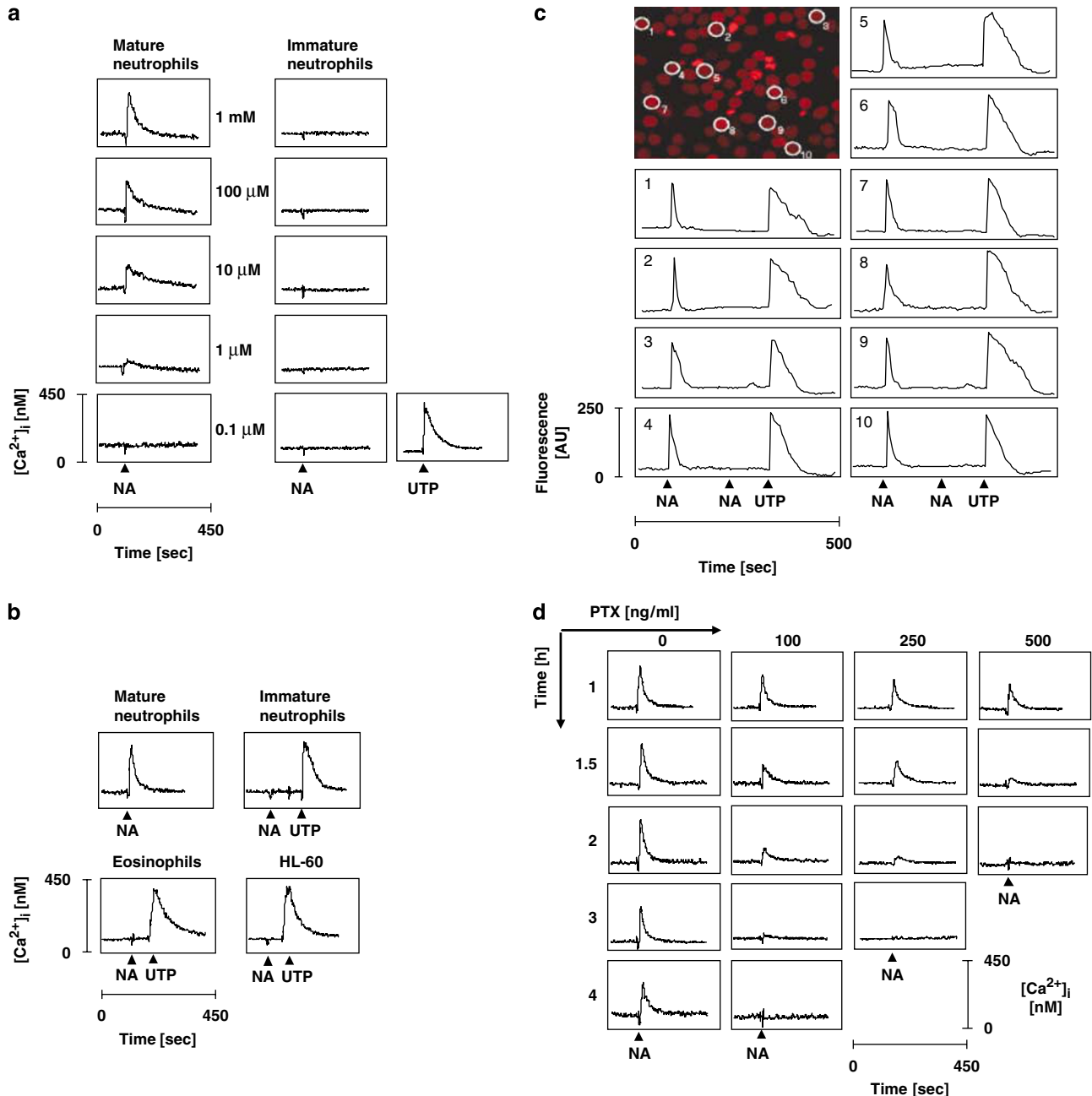


Figure 2 Mature blood neutrophils, but not immature bone marrow neutrophils or mature blood eosinophils, express functional GPR109A. **(a)** Concentration-dependent increases in [Ca²⁺]_i in mature neutrophils (left). NA had no effect on [Ca²⁺]_i in immature neutrophils (right). Positive control experiments were performed using UTP. **(b)** Functional GPR109A receptors were detectable in mature blood neutrophils, but not in their precursors in the bone marrow, HL-60 cells or in mature blood eosinophils. Positive control experiments were performed using UTP. NA was added at 100 μ M. **(c)** Single-cell analysis. Changes in [Ca²⁺]_i were analyzed in 10 single blood neutrophils. A second NA stimulation did not result in increases in [Ca²⁺]_i, suggesting homologous desensitization. **(d)** PTX inhibited increases in [Ca²⁺]_i in a concentration- and time-dependent manner. Positive control experiments were performed using UTP (data not shown). Representative examples of at least four independent experiments are shown in each panel. NA and UTP (20 μ M) were added where indicated (▲)

The death efficacy of NA was similar to that obtained with optimal concentrations of anti-Fas monoclonal antibody (mAb), a known trigger of neutrophil cell death.¹⁰ Moreover, since cytokine-mediated delayed apoptosis of mature neutrophils plays an important role under inflammatory conditions,⁵ we were interested in the effect of NA in the presence of GM-CSF. Interestingly, NA completely

abolished the cytoprotective effect of GM-CSF *in vitro* (Figure 3b).

The type of death, which was induced by NA was apoptosis, as assessed by phosphatidylserine (PS) redistribution (Figure 3c, upper panel), DNA fragmentation (Figure 3d), and morphological analysis (data not shown). The pro-apoptotic effect of NA was blocked by pre-incubation of

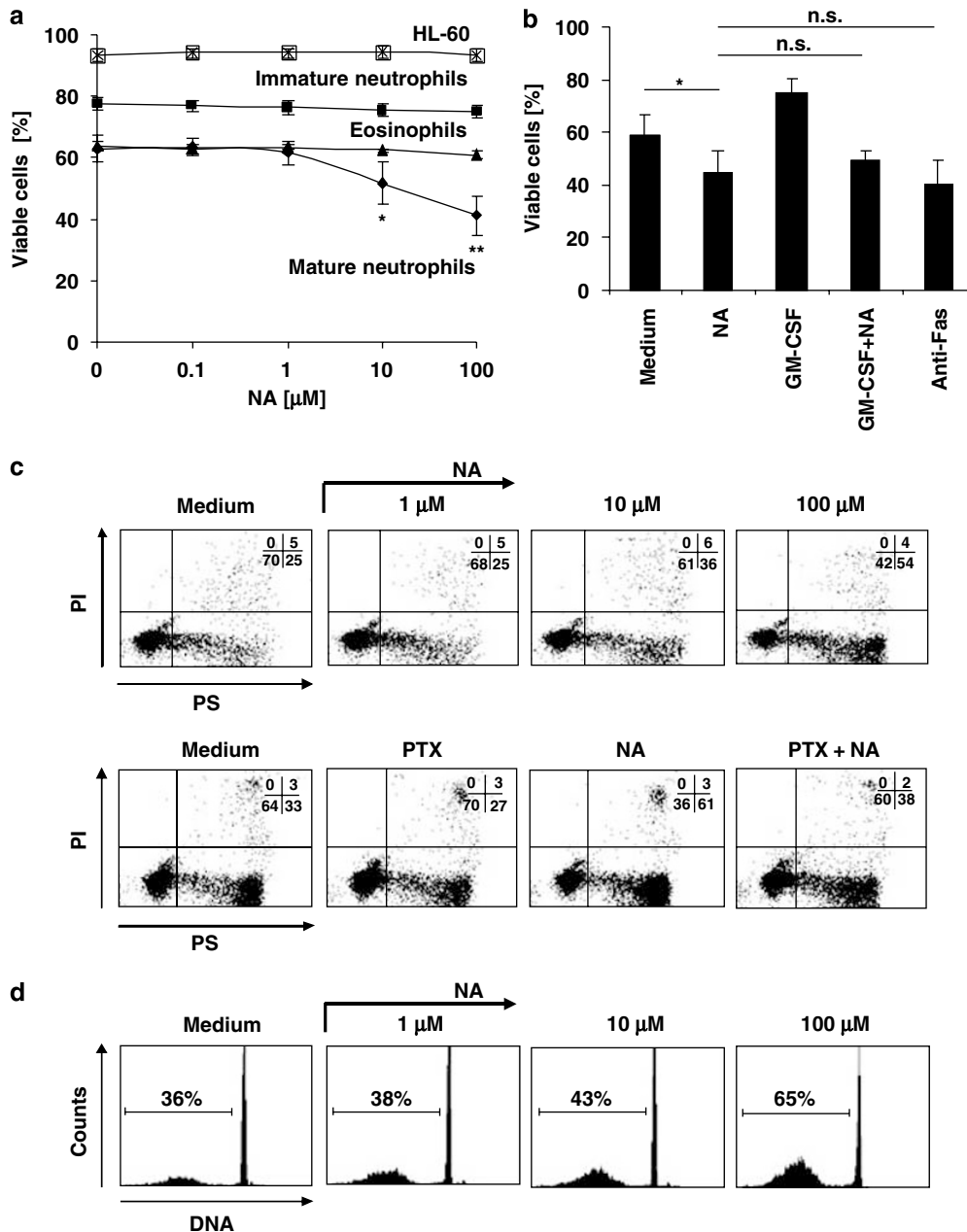


Figure 3 NA accelerates spontaneous neutrophil apoptosis. **(a)** NA induced death in mature neutrophils in a concentration-dependent manner. In contrast, NA had no cytotoxic effect on blood eosinophils, immature neutrophils, or HL-60 cells. Results of 24-h cultures are shown. Data are expressed as means \pm S.D. of 4–10 independent experiments. * $P < 0.05$; ** $P < 0.01$. Similar data were observed in 48-h cultures. **(b)** The death efficacy of 100 μ M NA and 1 μ g/ml anti-Fas mAb on blood neutrophils was similar. Moreover, NA completely abolished the anti-death effect of 25 ng/ml GM-CSF. Results of 24-h cultures are shown ($n = 3$). Data are expressed as means \pm S.D. * $P < 0.05$. **(c)** NA stimulation induced PS redistribution in blood neutrophils indicative of the induction of apoptosis in a concentration-dependent manner (upper panel). PTX (500 ng/ml for 2 h) abolished this effect. Quantitative analysis is shown in percent at the top right corner of each dot blot. The panels show one representative experiment out of three (8-h cultures). **(d)** NA induced DNA fragmentation in a concentration-dependent manner. The relative number of apoptotic blood neutrophils is indicated by the bracket in each subpanel. One representative experiment out of three is shown (14-h cultures)

neutrophils with PTX, suggesting that it is GPR109A-receptor-mediated (Figure 3c, lower panel). Moreover, caspase-3, a critical effector caspase in neutrophil apoptosis,¹⁷ is cleaved 4 h upon addition of NA into the apparent active enzyme (17-kDa fragment), whereas non-treated neutrophils do not demonstrate any detectable cleavage of caspase-3 at this time point (Figure 4a). We also measured increased enzy-

matic activity of caspase-3 in 12-h neutrophil cultures in the presence of NA compared to non-treated cells at this time point or freshly isolated neutrophils (Figure 4b).

The involvement of caspases was further supported by the observation that the pan-caspase inhibitor z-VAD-fmk as well as the caspase-3 inhibitor z-DEVD-fmk blocked both spontaneous and NA-accelerated neutrophil death (Figure 4c). The

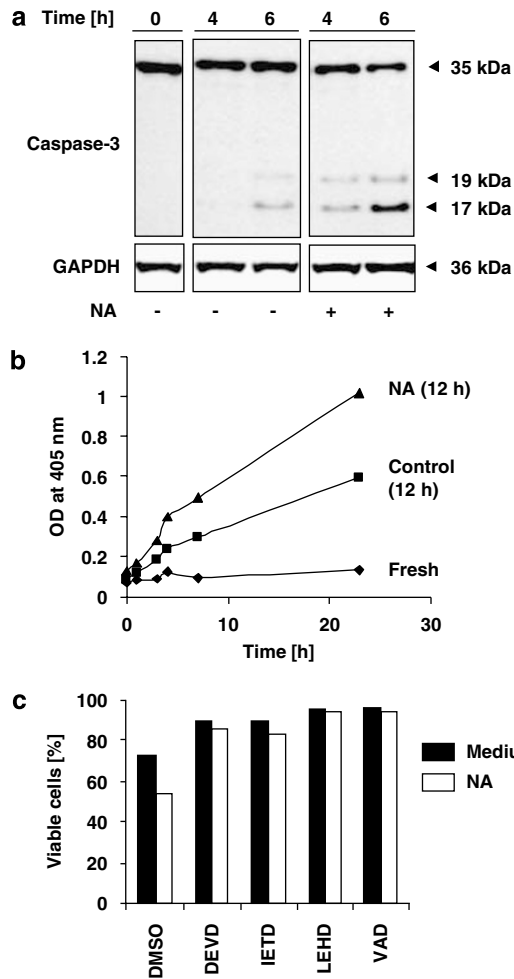


Figure 4 NA-mediated neutrophil apoptosis is caspase-dependent. (a) The induction of neutrophil apoptosis by NA was associated with the rapid occurrence of the 17-kDa caspase-3 cleavage product (detectable in 2-h cultures). (b) Spontaneous neutrophil apoptosis was associated with the induction of caspase-3-like enzymatic activity. NA accelerated caspase-3-like activity. Results of 12-h cultures were shown compared to freshly isolated blood neutrophils. One representative experiment out of three is shown. (c) All indicated caspase inhibitors (all used at 50 μ M) completely blocked NA-mediated neutrophil apoptosis. One representative experiment out of three is shown (24-h cultures)

caspase-8 inhibitor z-IETD-fmk and the caspase-9 inhibitor z-LEHD-fmk demonstrated similar blocking efficacies, consistent with previously published work suggesting that these caspases also play a role in neutrophil apoptosis.^{15,17} Taken together, NA accelerated spontaneous neutrophil apoptosis in a caspase-dependent manner *in vitro*.

NA-mediated reduction of cAMP levels associated with reduced phosphorylation of Bad at Ser 136. In initial experiments, we investigated whether pharmacological inhibition of calcineurin would block NA-mediated neutrophil death. Both FK506 (1–1000 ng/ml) and cyclosporin A (1–1000 ng/ml) had no effect on neutrophil viability in the presence and absence of NA (data not shown), excluding the possibility that calcium-mediated calcineurin activation is responsible for the pro-apoptotic effect of NA on neutrophils.

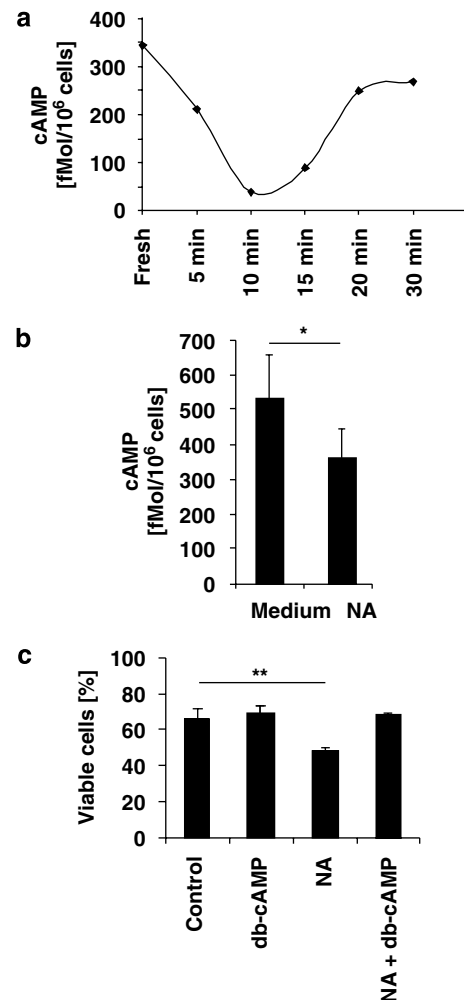


Figure 5 NA decreases intracellular cAMP levels in mature blood neutrophils. (a) NA stimulation of blood neutrophils resulted in a rapid reduction of intracellular cAMP levels. Maximal reduction was seen approximately 10 min upon NA stimulation. Concentration of cAMP increased thereafter, but did not reach the original levels. One representative experiment out of five is shown. (b) Reduced cAMP levels were also detected in neutrophils cultured for 6 h upon NA stimulation ($n=5$). Data are expressed as means \pm S.D. * $P<0.05$. (c) The NA-mediated cytotoxic effect on neutrophils was reversible by increasing the levels of cAMP (exogenous db-cAMP was added at 100 μ M). ** $P<0.01$

Besides intracellular calcium mobilization, GPR109A receptors have also been reported to inhibit adenylate cyclase activity.¹⁸ Therefore, we analyzed intracellular cyclic adenosine monophosphate (cAMP) levels in NA-stimulated mature neutrophils. NA dramatically reduced cAMP levels within 10 min (Figure 5a) and remained at this reduced level at least for 6 h (Figure 5b).

Previous reports suggested that cAMP levels regulate neutrophil apoptosis.^{19–21} We therefore asked the question whether the cell-permeable analogue of cAMP, dibutyl cyclic adenosine monophosphate (db-cAMP), can reverse the pro-apoptotic effect of NA. Although db-cAMP had no significant effect on spontaneous neutrophil death, it completely abolished the NA-mediated cytotoxic effect (Figure 5c). This suggested that the NA-mediated reduction

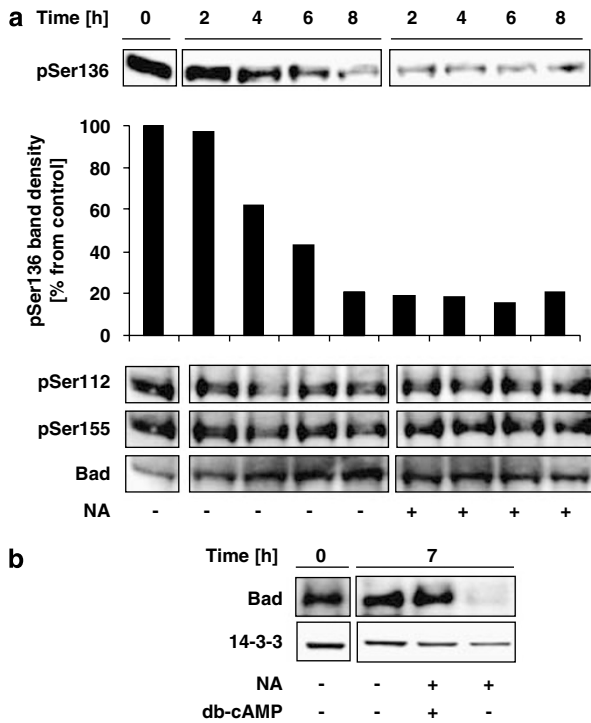


Figure 6 NA reduces phosphorylation of Bad at Ser 136 in mature blood neutrophils. **(a)** Neutrophils were cultured in the presence and absence of NA for the indicated time periods. Phospho Ser 112, Ser 136 and Ser 155 Bad, and total Bad were analyzed by immunoblotting. NA reduced Bad phosphorylation at Ser 136 that was quantified by densitometry. **(b)** 14-3-3 proteins in neutrophils were immunoprecipitated. Bad and 14-3-3 proteins were subsequently detected by immunoblotting. NA reduced the association between Bad and 14-3-3. This effect was not seen in the presence of exogenous db-cAMP. In both panels, representative immunoblots of at least two independent experiments are shown

in cAMP levels is a critical event for the subsequent induction of apoptosis.

Most effects of cAMP are mediated through activation of cAMP-dependent protein kinase A (PKA). One potential target of PKA is Bad.^{22,23} We, therefore, analyzed the phosphorylation pattern of Bad in neutrophils in response to NA stimulation. We observed that treatment of mature neutrophils with NA resulted in dephosphorylation of Bad at Ser 136, but not at Ser 112 or Ser 155, within 2 h (Figure 6a). Densitometry analysis indicated that 2-h NA treatment reduced Bad phosphorylation at Ser 136 approximately fivefold. These data suggested that Ser 136 phosphorylation is under the control of PKA in cultured neutrophils.

To demonstrate that attenuated Bad phosphorylation levels in NA-treated cells are functionally relevant, we investigated the physical association between Bad and 14-3-3 proteins.²⁴ Neutrophil lysates were immunoprecipitated with anti-14-3-3 and blotted with anti-Bad and anti-14-3-3 antibodies. In freshly isolated as well as cultured neutrophils, Bad associated with 14-3-3 proteins, suggesting that the pro-apoptotic function of Bad was largely blocked.²⁵ However, the interaction of Bad and 14-3-3 proteins decreased 7 h upon NA stimulation (Figure 6b). Importantly, db-cAMP completely prevented the NA-mediated disassociation of Bad and 14-3-3 proteins, in correlation with the prevention of NA-induced cell death (Figure 5c). Since the pharmacological inhibition of calcineurin

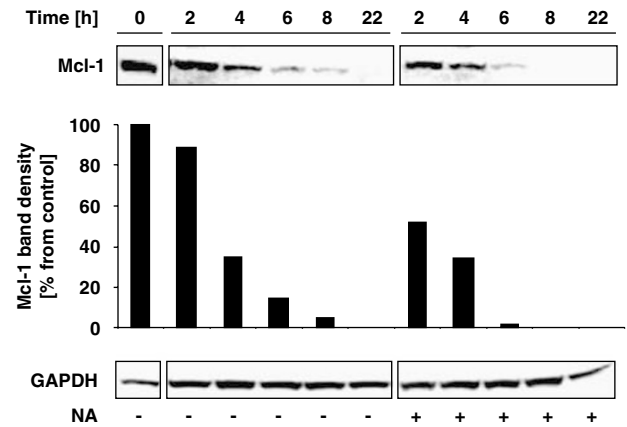


Figure 7 NA accelerates Mcl-1 degradation in mature blood neutrophils. Neutrophils were cultured in the presence and absence of NA for the indicated time periods. Mcl-1 expression was analyzed by immunoblotting. Mcl-1 declined over time, but more rapidly in the NA-stimulated neutrophils. Mcl-1 levels were quantified by densitometry. Representative immunoblots of three independent experiments are shown

had no effect on NA-mediated neutrophil death, these data suggested that NA-mediated reduced cAMP levels cause Bad dephosphorylation at Ser 136, resulting in its redistribution to mitochondria and consequently in increased apoptosis.

Mcl-1 is rapidly degraded during neutrophil apoptosis and cAMP was shown to stabilize Mcl-1 protein.²¹ We, therefore, analyzed Mcl-1 protein expression during spontaneous and NA-triggered neutrophil apoptosis. In NA-treated neutrophils, we observed markedly reduced Mcl-1 levels following culturing cells for 2 h as well as at later time points (6 and 8 h). In contrast, in non-treated neutrophils, Mcl-1 levels declined less rapidly (Figure 7). Taken together, lowering cAMP levels by NA stimulation is associated with reduced Bad phosphorylation and accelerated Mcl-1 degradation in mature neutrophils.

Discussion

The results of the current study suggest that mature neutrophils express functional GPR109A receptors, which have been known to inhibit lipolysis in adipocytes.¹ Receptor expression in the neutrophilic lineage appears to be induced during the terminal differentiation phase, since a mixed population of immature neutrophils containing myeloblasts, promyelocytes, and metamyelocytes were GPR109A negative. Moreover, GPR109A expression was not seen in mature eosinophils, suggesting that it is not a general feature of all granulocytes. Recently, it has been demonstrated that the receptor is also expressed on macrophages and dendritic cells, but not on monocytes.⁴ Therefore, GPR109A receptors are expressed on some, but not all, leukocytes and may play a role in defense mechanisms against pathogens, at least within the innate immune system.

NA binds to both low- (GPR109B; HM74) and high-affinity (GPR109A; HM74A) receptors. Activation of GPR109A receptors occurs in an NA concentration range between 10^{-7} and 10^{-4} M, whereas GPR109B receptors require 10^{-3} M NA for activation.² Therefore, mature neutrophils express GPR109A receptors, since significant signal transduction

(changes in $[Ca^{2+}]_i$) and apoptosis induction was seen at 10^{-5} M NA. However, we cannot exclude the possibility that these cells additionally express GPR109B receptors, in particular, since the antibody used to measure protein expression may recognize both receptors. The two receptors are highly homologous, displaying 96% identity at the protein level.²

Neutrophil apoptosis has previously been reported to be regulated by cAMP. For instance, agents and inflammatory stimuli that increase cAMP levels delayed spontaneous neutrophil apoptosis.¹⁹ The delay of apoptosis is also observed when the cell-permeable analogue of cAMP, db-cAMP, is added to neutrophils, directly demonstrating that cAMP mediates an antiapoptotic effect in these cells.^{20,21} Interestingly, elevation of cAMP promotes apoptosis in T cells,^{26,27} suggesting cell-type-specific differences. In agreement to previously published work performed in cell lines,^{28,29} we also observed reduced cAMP levels upon NA-mediated triggering of GPR109A receptors in mature neutrophils. This suggested a potential link to the NA-mediated pro-apoptotic effect observed in these cells. The fact that increasing the levels of cAMP by adding db-cAMP completely blocked the pro-apoptotic effect of NA supports this assumption.

PKA is a major target of cAMP³⁰ and may therefore be involved in cAMP-mediated antiapoptosis in neutrophils. However, the targets of PKA responsible for this effect remain to be determined. PKA is known to phosphorylate multiple targets, including Bad, a known pro-apoptotic member of the Bcl-2 family.^{22,23} Phosphorylated Bad is considered as being antiapoptotic, since it is bound in the cytosol to 14-3-3 proteins.²⁵ We, therefore, hypothesized that NA-mediated decreased cAMP levels and subsequent decreased PKA activity may reduce Bad phosphorylation levels. Indeed, in the presence of NA, Bad demonstrated decreased phosphorylation at Ser 136. Moreover, we obtained evidence that less Bad is sequestered in the cytosol by 14-3-3 proteins as the consequence of NA treatment. Interestingly, among the different phosphorylation sites of Bad, Ser 136 appears dominant in 14-3-3 binding.^{31,32} Taken together, our data suggest that less phosphorylated Bad as a consequence of NA stimulation is indeed pro-apoptotic and largely contributes to accelerated neutrophil apoptosis.

It has also been suggested that PKA may phosphorylate specific residues of the antiapoptotic Bcl-2 family member Mcl-1, leading to its stabilization.²¹ Mcl-1 plays an important role in neutrophil apoptosis.⁵ The more rapid degradation of Mcl-1 in neutrophils in the presence of NA compared with non-treated cells could therefore be explained by reduced Mcl-1 phosphorylation, leading to more rapid degradation. However, we cannot exclude the possibility that Mcl-1 degradation just correlated with the rate of apoptosis. Although the exact mechanisms of how NA induces neutrophil apoptosis remain to be determined, our initial data suggest that signal transduction events are involved in this pathway.

NA has been used as a lipid-lowering drug for decades.³³ The primary action of NA is the inhibition of lipolysis in adipocytes, leading to an increase in HDL and a decrease in VLDL and LDL levels in blood.³³ However, obesity is also considered as being a chronic inflammatory disorder.³⁴ Limiting neutrophil numbers in chronic inflammatory re-

sponses is an important anti-inflammatory mechanism.⁵ Therefore, the pro-apoptotic effect of NA on neutrophils may contribute to its overall beneficial effect in obese patients and GPR109A receptors continue to be promising targets for future pharmacotherapy.

Materials and Methods

Reagents. Anti-GPR109A (HM74a) rat mAb was from R&D Systems Europe Ltd. (Abingdon, UK). NA, FK-506, and cyclosporin A were obtained from Sigma-Aldrich (Buchs, Switzerland). Fluorescein isothiocyanate (FITC)-conjugated NA was synthesized by Molecular Probes Europe BV (Leiden, The Netherlands). Anti-CD7 and anti-CD36 mAbs for immature neutrophil isolations, anti-Mcl-1 mAb, the Annexin V apoptosis detection kit and all caspase inhibitors (z-VAD-fmk, z-DEVD-fmk, z-IETD-fmk, z-LEHD-fmk) were from BD Biosciences (Basel, Switzerland). Anti-CD16 mAb microbeads for eosinophil isolations were from Miltenyi Biotec GmbH (Bergisch Gladbach, Germany). Polyclonal anti-caspase-3 and all anti-Bad Abs were from Cell Signaling Technology Inc. (BioConcept, Allschwil, Switzerland). Polyclonal anti-14-3-3 β (K-19) Ab was from Santa Cruz Biotechnology Inc. (Heidelberg, Germany). Anti-glyceraldehyde-3-phosphate dehydrogenase (GAPDH) mAb was from Chemicon International Inc. (Juro, Lucerne, Switzerland). Horseradish peroxidase (HRP)-conjugated secondary Abs were from Amersham Bioscience Europe GmbH (Freiburg, Germany). Phycoerythrin (PE) and tetramethylrhodamine isothiocyanate (TRITC)-conjugated donkey anti-rat secondary Abs were purchased from Jackson ImmunoResearch Laboratories (Milan Analytica, La Roche, Switzerland). Anti-Fas mAb (clone CH-11) was obtained from MBL International Corporation (LabForce AG, Nunningen, Switzerland). PTX and fura-2-acetoxymethyl ester (fura-2/AM) were from Alexis (Lausen, Switzerland). Human GM-CSF was purchased from Novartis Pharma GmbH (Nürnberg, Germany). db-cAMP was provided by Biomol Research Laboratories Inc. (Plymouth Meeting, PA, USA). Fluo-3/AM was from Molecular Probes (Juro, Lucerne, Switzerland). UTP was from Boehringer Mannheim (Mannheim, Germany). All other chemicals were, unless stated otherwise, from Sigma-Aldrich.

Cells. Mature blood neutrophils were isolated from peripheral blood of healthy donors by Ficoll-Hypaque centrifugation.^{11,15,35,36} Briefly, peripheral blood mononuclear cells were separated by centrifugation on Ficoll-Hypaque (Seromed-Fakola AG, Basel, Switzerland). The lower phase, mainly granulocytes and erythrocytes, was treated with erythrocyte lysis solution (155 mmol/l NH_4Cl , 10 mmol/l $KHCO_3$, and 0.1 mmol/l EDTA, pH 7.3). The resulting cell populations contained greater than 95% mature neutrophils as assessed by staining with Diff-Quik (Medion GmbH, Düringen, Switzerland) and light microscopy analysis.

Immature neutrophils were isolated from bone marrow aspirates with normal cellular morphology and distribution as described previously.^{15,16} Briefly, following centrifugation on a two-step discontinuous percoll density gradient, cells were negatively isolated using anti-CD7 and anti-CD36 mAbs to eliminate contaminating lymphoid and erythroid precursors. The resulting cell population contained >97% cells of the neutrophil lineage as determined by myeloperoxidase staining, flow cytometric analysis of lineage-associated surface proteins,¹⁵ as well as with Diff-Quik and light microscopy. The proportion of mature neutrophils was less than 5%.

Peripheral blood eosinophils from control individuals were purified as described previously.^{37,38} The resulting cell populations contained more than 99% eosinophils as controlled by staining with Diff-Quik and light microscopy.

All individuals who donated blood or bone marrow aspirates signed a written informed consent agreement in accordance with the Declaration of Helsinki. The study was approved by the local medical ethics committee (the Kantonale Ethikkommission Bern).

Cell cultures. Neutrophils were cultured at 1×10^6 /ml in RPMI-1640 containing 1% fetal calf serum (FCS) and 200 IU/ml penicillin/100 μ g/ml streptomycin (all from Life Technologies, Basel, Switzerland) in the absence or presence of the indicated concentrations of NA, FK-506, cyclosporine A, PTX, GM-CSF, caspase inhibitors, and db-cAMP for the indicated time periods. If not indicated, NA was used at 100 μ M. HL-60 cells were cultured in Iscove's medium containing 10% FCS.

Cell-surface expression analysis

Method 1. Neutrophils were stained with 2.5 μ g/ml antihuman GPR109A mAb or isotype-matched control mAb. Binding of the primary Abs was visualized with

PE-conjugated donkey anti-rat secondary Ab (1/100) using flow cytometric analysis (FACS-Calibur, BD Biosciences).

Methods 2 and 3. Cytospins of fresh or cultured neutrophils were made on non-coated slides. Cells were fixed in 4% paraformaldehyde at room temperature for 10 min and washed three times in PBS, pH 7.4. To prevent non-specific binding, slides were incubated in blocking buffer (25% human immunoglobulins, 25% normal goat serum, 25% normal donkey serum, and 25% BSA) at room temperature for 1 h. Indirect immunostaining of GPR109A was performed at room temperature for 1 h with 2.5 μ g/ml anti-GPR109A or isotype-matched control mAb. Cells were then incubated with TRITC-conjugated donkey anti-rat secondary Ab (1/100) in the dark at room temperature for 1 h. In other experiments, cells were incubated with FITC-conjugated NA for 30 min. Specificity was controlled by pre-incubation of the cells with unlabeled 1 mM NA for 1 min. In the latter experiments, nuclei were counterstained with propidium iodide.³⁹ Slides were covered by coverslips and analyzed by confocal laser scanning microscopy (LSM 510, Carl Zeiss, Heidelberg, Germany) equipped with Ar and HeNe lasers.

Intracellular calcium measurements

Method 1. Intracellular ionized calcium concentrations ($[Ca^{2+}]_i$) were assayed with a bulk spectrofluorometric assay using a Perkin Elmer LS50 spectrofluorimeter as described previously.^{40,41} In the PTX experiments, cells were pre-incubated with the inhibitor at the indicated concentrations and for the indicated times.

Method 2. $[Ca^{2+}]_i$ were also assayed in single cells. Cells (3×10^5) were immobilized on glass cover slides for live-cell microscopy¹⁴ and loaded with 10 μ M fluo-3/AM for 30 min. The attached cells were carefully washed and 200 μ l PBS supplemented with glucose (2 mM) was added. Calcium measurements were performed in a 37°C-chamber using a confocal laser scanning microscopy (LSM 510). Fluo-3- Ca^{2+} was excited at 514 nm and fluorescence emission was detected using a 560-nm longpass filter. Images were collected with a $\times 40$, 1.3 NA Zeiss oil immersion objective. Single plane eight-bit images of 256×256 pixels representing 325.7μ m in each dimension were recorded at intervals of 1.5 s. After starting the recording process, agonists were applied at the indicated concentrations. From the resulting image series, signal intensities were calculated using the LSM510's built-in quantification software (Zeiss 510, version 2.5).

cAMP measurements. To rise cAMP to detectable levels, the phosphodiesterase inhibitor 3-isobutyl-1-methylxanthine (100 μ M) was added for 10 min. For the time-dependent experiments, mature neutrophils (1×10^6 /ml) were placed into 48-well plates and cultured in the presence and absence of 100 μ M NA for the indicated times. Intracellular levels of cAMP were analyzed using a commercial kit (Amersham) according to the recommendations of the manufacturer.

Determination of cell death and apoptosis. Cell death was assessed by uptake of 1 μ M ethidium bromide and flow cytometric analysis (FACS-Calibur).^{11,15,35,36} To determine whether cell death was apoptosis, DNA fragmentation and redistribution of PS were measured.^{11,15,35,36}

Immunoprecipitation. Neutrophils (5×10^6) were lysed in a buffer containing 0.2% Nonidet P-40, and protein samples were immunoprecipitated with 2 μ g of anti-14-3-3 Ab followed by precipitation with A-Sepharose.

Immunoblotting. Gel electrophoresis and immunoblotting were performed as described previously.^{11,15,16,35,36} Briefly, after electrotransfer of the separated proteins, the filters were incubated overnight with anti-caspase-3 Ab, anti-Mcl-1 mAb, or anti-phospho-Bad Abs (all 1/1000) at 4°C in TBS/0.1% Tween 20/5% non-fat dry milk. For loading controls, stripped filters were incubated with anti-GAPDH mAb (1/2000) or anti-Bad Ab (1/1000). Filters were washed, incubated with the appropriate HRP-conjugated secondary Ab (Amersham), and developed by an ECL-technique (ECL-Kit, Amersham) according to the manufacturer's instructions.

Enzymatic caspase assay. Enzymatic caspase-3-like activity in cell-free extracts was measured as enzymatic conversion of the colorimetric substrate Ac-DEVD-pNA at 405 nm according to the manufacturer's instructions (QuantiZyme caspase-3 cellular activity assay kit, Biomol).

Statistical analysis. Statistical analysis was performed by using the Mann-Whitney U-test. If mean levels are presented, the standard deviation of the mean (S.D.) and the number (n) of independent experiments are indicated in each case. A probability value of <0.05 was considered statistically significant.

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