

**Glycogen Synthase Kinase-3 Mediates Acetaminophen-induced Apoptosis in Human
Hepatoma Cells**

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ABSTRACT

The mild analgesic drug acetaminophen (AAP) induces severe hepatic injury when taken at excessive doses. Recent evidence shows that the initial form of damage is through apoptosis but this fails to go to completion and degenerates into necrosis. The aim of this study was to elucidate the mechanism through which AAP induces apoptosis using human HuH7 hepatoma cells as an *in vitro* model system to investigate the initial phase of AAP-induced hepatic injury. AAP-induced apoptosis in HuH7 cells as evidenced by chromatin condensation was preceded by the translocation of Bax to mitochondria and the cytoplasmic release of the proapoptotic factors cytochrome *c* and Smac/DIABLO. A concomitant loss of mitochondrial membrane potential occurred. Activation of the mitochondrial pathway of apoptosis led to the activation of execution caspases-3 and -7. AAP-induced apoptosis and cell death was blocked by inhibitors of caspases but not by inhibitors of calpains, cathepsins and serine proteases. Apoptosis was unaffected by inhibitors of the mitochondrial permeability transition pore and by inhibitors of Jun N-terminal kinases, p38 MAP kinase or MEK1/2. However, pharmacological inhibition of glycogen synthase kinase-3 (GSK-3) delayed and decreased the extent of AAP-induced apoptosis. In comparison, endoplasmic reticulum stress-induced but not prooxidant-induced apoptosis of HuH7 cells was sensitive to GSK-3 inhibition. It is concluded that AAP-induced apoptosis involves the mitochondrial pathway of apoptosis that is mediated by GSK-3 and most likely initiated through an endoplasmic reticulum stress response.

INTRODUCTION

The mild analgesic acetaminophen (paracetamol, AAP) remains the commonest cause of acute liver failure in the USA and other parts of the world as a result of accidental or deliberate overdose (Dargan and Jones, 2003). At therapeutic doses, AAP is primarily eliminated as its glucuronide and sulfate conjugate although a minor portion becomes bioactivated in the liver by CYP2E1 (Tonge *et al.*, 1998; Zaher *et al.*, 1998) to the reactive metabolite, *N*-acetyl-*p*-benzoquinoneimine (NAPQI). However, following an AAP overdose the rate and quantity of NAPQI production exceeds its detoxification through conjugation with glutathione, and hepatotoxicity ensues as a consequence of the binding of NAPQI to cellular proteins (Cohen *et al.*, 1997; Qiu *et al.*, 1998) and their oxidative damage (Tirmenstein and Nelson, 1990). A contribution to AAP hepatotoxicity also comes from the release of soluble pro-inflammatory and pro-apoptotic molecules such as tumor necrosis factor (TNF) (Blazka *et al.*, 1996).

Many of the histochemical and biochemical features of the late stages of AAP toxicity, particularly following high doses, support the conclusion that AAP induces hepatocellular necrosis (Adams *et al.*, 2001; Pierce *et al.*, 2002; Knight and Jaeschke, 2002; Gujral *et al.*, 2002; Jaeschke *et al.*, 2004). However, several reports have presented evidence for the occurrence of apoptosis in AAP-induced hepatic damage. For instance, AAP toxicity in rats is accompanied by an increase albeit small in the number of isolated parenchymal cells with apoptotic morphology (Dixon *et al.*, 1975; Gujral *et al.*, 2002), the formation of oligonucleosomal-length DNA fragmentation (Ray *et al.*, 1990) and the appearance of TdT-catalyzed dUTP-fluorescein nick end labeling (TUNEL) positive nuclei that were highly condensed (El-Hassan *et al.*, 2003). Several early mitochondrial events linked to apoptosis have been documented, including the truncation of the BH3-only proapoptotic Bcl-2 family

member Bid and its relocation to mitochondria (El-Hassan *et al.*, 2003), Bax translocation to mitochondria (Adams *et al.*, 2001; El-Hassan *et al.*, 2003) and the release of mitochondrial cytochrome *c* (Ferret *et al.*, 2001; Adams *et al.*, 2001; Knight and Jaeschke, 2002; El-Hassan *et al.*, 2003). However, there is little or no activation of the execution caspases-3 or -7 in liver tissues (Ferret *et al.*, 2001; Pierce *et al.*, 2002; Knight and Jaeschke, 2002; El-Hassan *et al.*, 2003), and it can be concluded that AAP induces an hepatocellular apoptotic response that fails to go to full completion and instead degenerates into necrosis (Adams *et al.*, 2001; Pierce *et al.*, 2002; El-Hassan *et al.*, 2003; Jaeschke *et al.*, 2004). However, the critical role of apoptosis in eliciting AAP-induced liver injury was demonstrated by a report from our laboratory showing that inhibitors of caspases protect mice from AAP-induced hepatic injury and ultimately necrosis *in vivo* by blocking the activation of the mitochondrial pathway of apoptosis (El-Hassan *et al.*, 2003).

The cytotoxicity of AAP in cultured hepatocytes requires concentrations of the drug that are higher than those required to induce hepatotoxicity *in vivo* (Prescott, 1996) and is much slower to develop than in mice. More importantly, cultured hepatocytes exposed to AAP die by necrosis in a mechanism that, unlike the situation *in vivo* (El-Hassan *et al.*, 2003), does not involve cytochrome *c* release from mitochondria and that does not involve nor is dependent on caspases (Neuman *et al.*, 1999; Nagai *et al.*, 2002). In this study, HuH7 hepatoma cells were used as an *in vitro* model to specifically study the early events that lead to the induction of apoptosis by the hepatotoxic drug. We report here that AAP activated the mitochondrial pathway of apoptosis in a mechanism that was mediated by the release of pro-apoptotic mitochondrial proteins. The apoptotic response required glycogen synthase kinase-3 (GSK-3) activity through a mechanism that appeared to involve an endoplasmic reticular (ER) stress response.

METHODS

Reagents and antibodies. Acetaminophen (99% pure), digitonin, N-tosyl-L-phenylalanine chloromethyl ketone (TPCK), saponin, 3-(2,4-dichlorophenyl)-4-(1-methyl-1H-indol-3-yl)-1H-pyrrole-2,5-dione (SB-216763), 3-[(3-chloro-4-hydroxy phenyl)amino]-4-(2-nitrophenyl)-1H-pyrrol-2,5-dione (SB-415286) and propidium iodide were purchased from Sigma (UK). Duroquinone was bought from Aldrich. Benzyloxycarbonyl-Val-Ala-DL-Asp-fluoromethylketone (Z-VAD-fmk) and N-acetyl-Asp-Glu-Val-Asp-aldehyde (Ac-DEVD-CHO) were obtained from Bachem (Bubendorf, Switzerland). Z-Phe-Ala-fmk (Z-FA-fmk) was obtained from Enzyme Systems Products (Livermore, CA, USA). Z-Leu-Leu-Leu-CHO (MG-132), thapsigargin, N-Acetyl-Leu-Leu-Nle-CHO (ALLN), 3-(4-iodophenyl)-2-mercapto-(Z)-2-propenoic acid (PD 150606), 2'-amino-3'-methoxyflavone (PD 98059), 4-(4-fluorophenyl)-2-(4-methylsulfinylphenyl)-5-(4-pyridyl) 1H-imidazole (SB-203580), bongkreikic acid, cyclosporin A and 1,9-pyrazoloanthrone (SP600125) were purchased from Calbiochem-Novabiochem (UK) and staurosporine and puromycin were from Alexis Corporation Ltd, UK. Bis-benzimide (Hoechst 33258) and tetramethylrhodamine ethyl ester (TMRE) were from Molecular Probes. Pefabloc, phenylmethylsulfonylfluoride (PMSF), aprotinin, pepstatin, leupeptin and lactate dehydrogenase (LDH) cytotoxicity detection kit were purchased from Roche. Anti-cytochrome *c* antibodies (clones 7H8.2C12 and 6H2.B4 for Western blot analysis and immunocytochemistry, respectively) and phycoerythrin-conjugated anti-caspase-3 (active form) (clone C92-605) were purchased from BD Pharmingen (San Diego, CA, USA). Anti-cytochrome *c* oxidase (subunit IV) was obtained from Molecular Probes (clone 20E8-C12) (Eugene, Oregon, USA). Anti- α -fodrin and anti-cytokeratin-18 (cleaved, M30) were obtained from Affiniti Research Products and Roche, respectively. Antisera against caspase-3 and Bax (B-9) were obtained from Calbiochem-Novabiochem (UK) and Santa Cruz, respectively. Anti-phospho-SAPK/JNK (Thr183/Tyr185) antibodies,

anti-Hsc70 antibodies and anti-SAPK1b/JNK3 rabbit antibodies were purchased from Cell Signalling Inc., Stressgen Biotechnology and Upstate Biotechnology, respectively. Anti-caspase-7 and anti-Smac/DIABLO antibodies were a gift from Prof Gerald M. Cohen (MRC Toxicology Unit, Leicester, UK). All secondary HRP-conjugated and fluorescein isothiocyanate (FITC)-conjugated antibodies were from Dako Ltd (Cambridge, UK). All other reagents were of analytical grade.

Cell culture and treatments. Human hepatoma HuH7 cells [a gift from Prof R. Bartenschlager (University of Mainz, Germany)] were grown at 37°C in DMEM supplemented with 10% fetal bovine serum, penicillin (100 IU/ml) and streptomycin (100 µg/ml), non-essential amino acids and 2 mM L-glutamine in an humidified incubator with 5% carbon dioxide/95% air. The cells were exposed in medium supplemented with 2% fetal bovine serum to AAP in the absence or presence of modulators of apoptosis (or a corresponding volume of solvent) as described in Results. The final concentration of solvent never exceeded 0.1% (v/v).

Analysis of apoptosis and cell death. Changes in nuclear morphology were assessed by fixing the cells grown on poly-L-lysine coated glass coverslips in 4 % (v/v) formaldehyde at room temperature for 30 minutes. The fixed cells were stained with Hoechst 33258 (2 µg/ml) and nuclear morphology analyzed by fluorescence microscopy. The nuclei were scored as either normal, in which the chromatin was uncondensed, or apoptotic, in which the chromatin was highly condensed or fragmented into discrete apoptotic bodies (Jones *et al.*, 1998). The assessment of cell viability was performed using the LDH cytotoxicity detection kit according to the manufacturer's instructions. DNA fragmentation was analyzed by flow cytometric detection of hypodiploid DNA. In brief, cells were detached by trypsinization, combined with medium containing floating cells and centrifuged at 100 x g for 5 minutes. The pellets were fixed in ice-cold 70% (v/v) ethanol in PBS overnight at 4°C by gradual addition whilst vortex

mixing. The cells were subsequently stained with propidium iodide (10 μ g/ml) and treated with RNase (1mg/ml) for 30 minutes at 37°C before analysis using a Beckman Coulter Epics XL flow cytometer (argon laser, excitation wavelength 488 nm). A minimum of 20,000 events were acquired in list mode while gating the forward and side scatters to exclude propidium iodide-positive cell debris and analyzed in FL-3 for the appearance of the sub-G₁ peak. Cellular GSH was measured using the Oxis Research Bioxytech GSH-400 kit on an clinical analyser (SpACE) and ATP content were assayed by luciferin/luciferase.

Analysis of pro-caspase processing and caspase substrate cleavage. Western blot analysis was performed as previously reported (Jones *et al.*, 1998). Cell extracts were prepared in cell lysis buffer (50mM Tris and 150mM NaCl, pH 7.5 supplemented with 1% (v/v) NP-40, 0.2 % (w/v) SDS, 1mM pefabloc, 2 μ g/ml aprotinin, 2 μ g/ml leupeptin, 1 mM PMSF and 1 mM Na₃VO₄). To separate mitochondrial proteins from cytosolic proteins, the cells were washed, resuspended in intracellular medium buffer (20 mM NaCl, 100 mM KCl, 2 mM MgCl₂, 1 mM EGTA, 23 mM Hepes, pH 7.1) supplemented with 1 μ M cyclosporin A and permeabilized on ice for 10 min by the addition of saponin (1 mg/ml). The cells were then centrifuged (2000 x g for 5 min at 4 °C) to separate cytosolic proteins from membrane (mitochondria)-associated proteins. Protein concentration in the cell samples was assessed using the BioRad protein assay (BioRad Laboratories). The cellular proteins were resolved by SDS-PAGE using 7.5% or 15% gels, blotted onto Hybond-C nitrocellulose membranes and probed with antibodies raised against caspase-3 (1:1000), caspase-7 (1:2000), Bax (1:500), cytochrome *c* (1:500), cytochrome *c* oxidase (1:1000), fodrin (1:2000), Smac/DIABLO (1:1000), Hsc70 (1:1000), phospho-SAPK/JNK (1:1000) or SAPK1b/JNK3 (1:1000). Incubation with primary antibodies was carried out overnight at 4°C in an orbital roller. After washing and incubation with peroxidase-conjugated secondary antibody (1:1000), the blots were visualized using the SuperSignal West Pico chemiluminescent detection system (Pierce). The reprobing for

loading controls was carried out as described (Kaufmann, 2001). In brief, the Hybond-C nitrocellulose membranes were incubated in 15ml PBS supplemented with 1mM sodium azide for 2 hours. The membranes were then rinsed in 1 x TBS-T buffer for 5 minutes and re-probed with primary antibody. Caspase-3 processing and caspase-mediated cytokeratin-18 cleavage were measured by flow cytometric detection of the corresponding neo-epitopes as described by the manufacturers (BD Pharmingen and Roche, respectively). A minimum of 10,000 events were acquired in list mode while gating the forward and side scatters to exclude cell debris and analyzed in FL-2 and FL-1, respectively for the appearance of the neo-epitopes.

Immunolocalization of cytochrome c. HuH7 cells grown on poly-L-lysine coated glass coverslips were fixed with 4% paraformaldehyde buffer for 30 minutes at room temperature. Following washes with blocking buffer (PBS supplemented with 3% bovine serum albumin and 0.05% saponin), the cells were incubated with anti-cytochrome c antibody (1:100 in blocking buffer) overnight at 4°C followed by FITC-linked secondary antibody (1:100) for 1 hour at room temperature and stained with Hoechst 33258 (2 µg/ml).

Detection of mitochondrial membrane potential with TMRE. HuH7 cells grown on poly-L-lysine coated round glass coverslips were treated with AAP, washed with pre-warmed PBS and incubated at 37°C for 15 minutes with medium supplemented with 2% fetal bovine serum, 0.5µM TMRE and Hoechst 33258 (2 µg/ml). The cells were then immediately viewed under a Zeiss Axiovert 135 fluorescence microscope.

Statistical analysis of the data. All data are given as means \pm SD of at least three independent experiments. Comparison of treatments against controls was made using one-way ANOVA followed by Fisher's LSD post hoc test with the SPSS statistical package. The significance level chosen for the statistical analysis was $p < 0.05$.

RESULTS

AAP induces apoptosis in HuH7 cells. A human hepatoma cell line, HuH7, was used to investigate the mechanism of AAP-induced apoptosis induction *in vitro*. AAP induced a concentration and time-dependent loss in cell viability in HuH7 cells. The losses of viability at 24h (as measured by the MTT assay) after AAP dosing were 21% (1 mM AAP), 24% (5 mM AAP) and 28% (10 mM AAP) and at 48h were 19% (1 mM AAP), 46% (5 mM AAP) and 68% (10 mM AAP). The loss of cell viability after exposure to 10 mM AAP (Fig. 1) was characterized by the appearance of a large number of cells displaying nuclear apoptotic morphology (chromatin condensation and fragmentation) and DNA fragmentation. Chromatin condensation also occurred after exposure to 5 mM AAP although in a reduced number of cells and only a few isolated apoptotic cells were observed after a 48h-treatment with 1 mM AAP (data not shown). The time course of the development of the cytotoxicity in HuH7 cells followed that occurring in humans where typically 24-48 h are required before transaminases, bilirubin levels and prothrombin time begin to increase (Salgia and Kosnik, 1999). The analysis of biochemical events preceding the loss of HuH7 cell viability and apoptosis showed a decrease in cellular GSH content ($53 \pm 12\%$ and $39 \pm 4\%$ decrease at 30h and 48h, respectively) and ATP content ($65 \pm 5\%$ and $35 \pm 1\%$ decrease at 30h and 48h, respectively) in response to AAP (10 mM) exposure. These findings support a mechanism whereby AAP requires bioactivation to NAPQI for toxicity and apoptosis to occur and that the apoptotic response is concentration-dependent.

Activation and role of caspases in AAP-induced apoptosis. Extracts from cells treated with AAP were analyzed by Western blotting for the processing and activation of the execution caspases-3 and -7. Solvent control-treated cells showed the distinct presence of pro-caspase-3 (p32) and a second band possibly corresponding to caspase-3 minus the pro-domain (p29) (Fig. 2A). Treatment of HuH7 cells with the apoptosis inducing drugs staurosporine (1 μ M

for 24h) or puromycin (20 μ M for 24h) generated a band that corresponded to the active fragment p17. This band was absent from control cells but was detected in HuH7 cells treated with AAP (10 mM) for 24 or 48h (Fig. 2A). This supports the conclusion that AAP-induced apoptosis correlated with the proteolytic processing and activation of caspase-3 in HuH7 cells. A fourth immunoreactive (p20) band was present in all samples. The latter band could have corresponded to the large subunit plus prodomain that in a cellular environment as p20-p12 complex is inactive (Li *et al.*, 2002). To address this possibility we investigated the processing of caspase-3 by flow cytometry using an antibody specifically recognizing the p20/p17 large fragments of caspase-3 but not the inactive proform. The lack of immunoreactivity in control cells demonstrates that the 20-kDa band detected by Western blot was unrelated to the p20 large subunit of caspase-3. Most importantly, Figure 2B shows that the time course of the appearance of the active caspase-3 correlated with the induction of apoptosis. Similarly, immunoblotting for caspase-7 revealed that this executioner caspase was similarly activated by AAP as evidenced by the appearance of the large fragment, p19 (Fig. 2A).

The appearance of the cleaved fragments does not necessarily imply that the caspases were catalytically active. However, the latter was demonstrated by examining the *in situ* cleavage of the substrate α -fodrin and the formation of a band corresponding to the p120 fragment of α -fodrin. Likewise, the detection of the M30 immunoreactive neo-epitope formed as a result of cytokeratin-18 cleavage confirmed the *in situ* activity of the execution caspases (Fig. 2C). The pivotal role of caspases in AAP-induced HuH7 cell apoptosis was established by the ability of the pan-caspase inhibitor Z-VAD-fmk to block cell death (Fig 2D) and caspase-mediated proteolysis *in situ* (Fig. 2C). The cytoprotective effect of the caspase inhibitor was dose-dependent with 50 μ M causing a 50% decrease in loss of cell viability and near complete protection was obtained with 100 μ M. A significant albeit incomplete protection

from AAP-induced cytotoxicity was also obtained with the caspase-specific inhibitor Ac-DEVD-CHO [cytotoxicity (%) at 54 h: AAP, 51.6 ± 10.8 ; AAP + 200 μ M Ac-DEVD-CHO, 39.5 ± 1.9 ; $p < 0.05$]. However, no protection occurred with the analogue Z-FA-fmk (50 μ M) that was used as a negative control to rule out possible non-specific effects of the fmk moiety [cytotoxicity (%) at 54 h: AAP, 53.5 ± 5.6 ; AAP + 50 μ M Z-FA-fmk, 58.4 ± 1.4 (n=3)]. Moreover, ALLN (range tested, 1-10 μ M), PD 150606 (range tested, 1-10 μ M), MG-132 (range tested, 0.5-10 μ M) or TPCK (range tested, 1-10 μ M) failed to protect from AAP-induced cytotoxicity (data not shown). These findings demonstrate that cytoprotection was achieved through caspase inhibition rather than interference with cathepsins or calpain activity by the reactive fmk moiety.

Role of mitochondria in AAP-induced apoptosis. AAP-induced apoptosis involved the release of the pro-apoptotic proteins, including cytochrome *c* from mitochondria in HuH7 cells (Fig. 3). At the cellular level, the relocation of cytochrome *c* from mitochondria always correlated with the onset of chromatin condensation (Fig. 3A). Furthermore, the release of mitochondrial intermembrane space proteins was not restricted to cytochrome *c* but was also observed with Smac/DIABLO (Fig. 3B). The mobilization of the pro-apoptotic mitochondrial proteins correlated with the translocation of Bax to the mitochondrial membranes (Fig. 3C) as well as a loss of mitochondrial membrane potential ($\Delta\Psi_m$) (Fig. 3D). As shown in Figure 3D, the loss of $\Delta\Psi_m$ in response to AAP coincided with a corresponding increase in nuclear staining by Hoechst 33258. The onset of apoptosis is characterized by an enhanced fluorescence of the nucleus that precedes the loss of cell viability (Ormerod *et al.*, 1993). Two stages of $\Delta\Psi_m$ loss as evidenced by TMRE staining were observed (Fig. 3D). In the first stage, near complete loss of TMRE uptake occurred in conjunction with enhanced nuclear staining. However, at this stage no chromatin condensation had taken place. Subsequently, TMRE fluorescence was undetectable but chromatin condensation was observed.

Both the loss of mitochondrial membrane potential and the release of pro-apoptotic proteins could have resulted from the induction of the mitochondrial permeability transition through pore opening (PTP). Indeed, the induction of PTP in isolated rat liver mitochondria by NAPQI and its dimethylated analogues is well established (Weis *et al.*, 1992). However, in the present study, neither cyclosporin A nor bongkreic acid was able to prevent or delay the loss of cell viability caused by AAP (Fig. 4).

GSK-3 but not stress kinases are involved in AAP-induced apoptosis. Given that the interaction between AAP and mitochondria seemed to be indirect and not mediated by NAPQI-induced mitochondrial PTP opening suggests the involvement of upstream signal(s) that co-operate(s) with Bax to initiate apoptosis. The primary site of AAP metabolism and its bioactivation to NAPQI is the ER. Hence, localized damage within this organelle may play a crucial role in the initiation of a cellular stress response that triggers the mitochondrial pathway of apoptosis. We therefore investigated the role of the MAPKinase pathway members, JNK, p38 MAP kinase and ERK in AAP-induced apoptosis. Figure 5A shows that two weak bands corresponding to phospho-JNK were detected in control cells. The intensity of the upper band was markedly enhanced in AAP-treated cells which is in contrast to the preferential phosphorylation of the lower band by the redox-cycling agent duroquinone (500 μ M). Taken together, phosphorylation of JNK occurs in response to AAP treatment although the pattern is distinct from that elicited by reactive oxygen species. The role of JNK activation in the induction of apoptosis was probed by using the JNK inhibitor SP 600125, and our results show that inhibition of JNK activity did not prevent the loss of cell viability in response to AAP (Fig. 5B). The pharmacological inhibition of p38 MAP kinase activity or MEK1/2 activity by SB-203580 (10 μ M) and PB98059 (50 μ M), respectively, also failed to protect or modify AAP cytotoxicity (Fig. 5B). These findings support the conclusion that the members of the MAP kinase pathways studied here are not involved in AAP-induced

apoptosis in HuH7 cells. In contrast, pretreatment with LiCl significantly protected the cells by delaying both the onset and extent of loss of cell viability induced by AAP (Fig. 6A). As GSK-3 is the primary pharmacological target of LiCl, these data suggest that GSK-3 plays a critical role in the events leading to the cytotoxicity of AAP. This was further confirmed by using two additional and specific inhibitors of this enzyme, SB-216763 and SB-415286 (Coghlan *et al.*, 2000) (Fig. 6B). To rule out the possibility that the inhibitors of GSK-3 were cytoprotective by delaying the onset of secondary necrosis, we tested both SB-415286 and SB-216763 for their ability to prevent caspase-3 activation and hence the execution of apoptosis. Fig. 6C demonstrates that the processing of pro-caspase-3 to its active fragment was significantly inhibited by SB-415286 (48h: control, $2.8 \pm 1.0\%$; AAP, $29.3 \pm 3.5\%$; AAP + SB-415286, $20.0 \pm 2.1\%$, $p < 0.05$) to a similar extent as the loss of cell viability. Similar results were obtained with SB-216763 (data not shown).

A role for ER stress in AAP-induced apoptosis? GSK-3 has recently been shown to represent a link between ER stress and the activation of the mitochondrial pathway of apoptosis (Song *et al.*, 2002; Ghribi *et al.*, 2003). We therefore investigated the ability of GSK-3 inhibitors to modulate ER-stress-induced apoptosis in HuH7 cells. Using the SERCA ATPase inhibitor thapsigargin that is well known to induce a condition of ER stress (Kass and Orrenius, 1999) we found that GSK-3 inhibitors blocked thapsigargin-induced apoptosis to a similar degree as their inhibition of AAP-induced apoptosis (Fig. 7A). In contrast, the induction of apoptosis by duroquinone-generated reactive oxygen species was essentially insensitive to GSK-3 inhibition (Fig. 7B).

DISCUSSION

Apoptosis plays a critical role in the mouse in initiating the events that lead to the destruction of the liver following an hepatotoxic dose of AAP (El-Hassan *et al.*, 2003). However, the ability to study the mechanism of AAP-induced apoptosis under *in vitro* conditions has been hampered by the lack of appropriate *in vitro* cellular model. Primary cultures of rat hepatocytes respond to cytotoxic doses of AAP by necrosis (Neuman *et al.*, 1999; Nagai *et al.*, 2002) and are therefore an unsuitable model system for the investigation of the early apoptotic events that occur following *in vivo* administration of AAP. Likewise, the differentiated transgenic mouse hepatocyte cell line (TAMH cells) displays only incomplete biochemical evidence for apoptosis although some degree of chromatin condensation and caspase activation was reported (Pierce *et al.*, 2002). Inhibition of the latter with z-VAD-fmk failed to prevent AAP-induced TAMH cell death (Pierce *et al.*, 2002). In contrast, AAP induced apoptosis in HuH7 cells as determined by chromatin condensation and fragmentation, the release of pro-apoptotic factors from mitochondria and the activation of caspases. The observation that HuH7 cell death by AAP was critically dependent on caspase activity, and that z-VAD-fmk nearly completely blocked AAP-induced cell death suggest that apoptosis rather than necrosis was the major form of cell death elicited by AAP. This conclusion is based on reports that caspase inhibitors are unable to block cell death by necrosis (Denecker *et al.*, 2001; Nagai *et al.*, 2002). It should be stressed, however, that the initial apoptotic response elicited in HuH7 cells did not prematurely degenerate into the necrosis typically observed *in vivo*. The mechanism underlying this transition in cell death mode is unclear. Therefore, HuH7 cells were found to be a good model for the investigation of the mechanism of AAP-induced apoptosis but the model is not intended to represent the later stages of liver injury. While this work was in progress, a report was published showing that AAP induced a caspase-mediated apoptotic response also in SK-Hep1 cells (Boulares *et al.*, 2002).

AAP induced the release of pro-apoptotic factors such as cytochrome *c* and Smac/DIABLO into the cytosol. Two different mechanisms have been postulated for the activation of this pathway and were investigated here. The release of mitochondrial pro-apoptotic factors could have been the result of mitochondrial PTP opening. The mitochondrial PTP is believed to be a protein complex formed primarily by outer membrane porin, the adenine nucleotide translocators and cyclophilin D. Indeed, the pharmacological inhibition of mitochondrial PTP opening has been demonstrated to prevent apoptosis and cell death in liver cells following exposure to a variety of apoptosis-inducing triggers (Pastorino and Hoek, 2000; Yerushalmi *et al.*, 2001; Zhao *et al.*, 2003). However, neither cyclosporin A nor bongkreikic acid prevented or attenuated AAP-induced HuH7 cell apoptosis. This suggests that the mitochondrial pathway of apoptosis was not initiated by mitochondrial PTP opening.

The execution of AAP-induced apoptosis was dependent on caspases as demonstrated by the protection afforded by caspase inhibitors, Z-VAD-fmk and Ac-DEVD-CHO. The pan-caspase inhibitor Z-VAD-fmk was a more potent inhibitor due to its superior ability to penetrate cells and to irreversibly inactivate most cellular caspases as compared to the tetrapeptide inhibitor. Non-specific effects of the reactive fmk moiety leading to the inhibition of cathepsins and calpains were ruled out by the findings that Z-FA-fmk and calpain inhibitors did not affect the cytotoxicity of AAP. Moreover, the cytoprotection provided by Z-VAD-fmk compared well with the prevention of caspase-3 processing and its *in situ* proteolytic activity. The pharmacological inhibition of calpains or cathepsins did not affect AAP-induced apoptosis ruling out their role in AAP-induced apoptosis. A recent report (Pierce *et al.*, 2002) showed that inhibition of proteasomal activity in TAMH cells with a low concentration of MG-132 promoted chromatin condensation and nuclear fragmentation induced by AAP whereas in the absence of the proteasome inhibitor only chromatin margination to the nuclear envelope was observed. Here, no modulation of AAP-induced apoptosis by MG-132 was observed, possibly

because a full apoptotic response was already elicited by AAP in HuH7 cells. Taken together, our results suggest that non-caspase proteases did not independently contribute to the initiation or execution of apoptosis in HuH7 cells by the hepatotoxic drug. However, a striking difference between the response of HuH7 cells to AAP and the *in vivo* situation is that following administration of AAP in mice, the caspase cascade, although required for the initiation of the events leading to the destruction of the liver, failed to fully activate the downstream execution caspases-3 and -7 (El-Hassan *et al.*, 2003). This leaves open the possibility that non-caspase proteases may contribute to liver cell death downstream of caspases under *in vivo* conditions and be responsible for some of the necrotic features observed after an hepatotoxic dose of AAP (Adams *et al.*, 2001; Pierce *et al.*, 2002; Jaeschke *et al.*, 2004).

If the trigger for activation of the mitochondrial pathway is not through PTP opening, then an alternative mechanism that most likely participates with Bax needs to be considered. AAP is bioactivated in the ER to the reactive metabolite NAPQI and AAP-induced injury also leads to cellular oxidative stress. Hence, a cellular stress response may be involved in AAP-induced apoptosis. Indeed, recent evidence has shown that AAP activates JNK in a glioma cell line (Bae *et al.*, 2001). Our results demonstrate that JNK is activated prior to apoptosis, as evidenced by the detection of the corresponding phospho-JNK bands. Interestingly, the pattern of activation was distinct from that elicited by the redox-cycling quinone duroquinone, suggesting that oxidative damage to cells in AAP-induced cytotoxicity (Tirmenstein and Nelson, 1990) is not the primary mechanism of JNK activation in HuH7 cells. However, in contrast to the pro-apoptotic role of JNK in glioma cell apoptosis, the pharmacological inhibition of JNK activity did not alter AAP-induced apoptosis of HuH7 cells. Likewise, the inhibition of p38 MAP kinase or MEK1/2 failed to modulate the ability of AAP to trigger apoptosis in the hepatoma cells. Although a cellular specificity intrinsic to HuH7 cells cannot

be excluded, these findings lead to the conclusion that neither the stress kinase pathways nor the ERK pathway of the MAP kinase family of signaling pathways are involved in AAP-induced liver cell apoptosis. This is in contrast to inducers of liver cell apoptosis such as menadione and TNF whose mechanisms of apoptosis action have been causally linked to the activation of one or more MAP kinase pathways (Czaja *et al.*, 2003; Pastorino *et al.*, 2003).

GSK-3 β is now recognized as being a central regulator of apoptosis and cell survival (Jope and Johnson, 2004). In cells this enzyme is maintained in an inactive state among others by AKT-, p90^{RSK}- and protein kinase A-mediated phosphorylation of N-terminal serine-9 and other mechanisms (Jope and Johnson, 2004). Activation of GSK-3 results in the upregulation of a number of pro-apoptotic genes and downregulation of anti-apoptotic genes through its effect on a wide number of transcription factors. Hence, its silencing is important in survival signals against a wide range of apoptosis-inducing conditions. Much of our current knowledge on the role of GSK-3 β in apoptosis stems from work in the area of neurodegeneration, and relatively little is known about its potential role in liver cells. Yet, GSK-3 β deficiency in mice is embryolethal as a result of massive liver degeneration in mid-gestation. This is caused by a failure of protection from tumor necrosis factor-induced apoptosis by GSK-3 mediated p65 phosphorylation and NF- κ B transactivation (Schwabe and Brenner, 2002). We therefore investigated the potential role of GSK-3 in AAP-induced apoptosis, and as reported here, a pro-apoptotic function was uncovered. A comparison with other inducers of apoptosis in HuH7 cells showed that the apoptotic effect of reactive oxygen species was insensitive to GSK-3 inhibition but that thapsigargin-mediated apoptosis was decreased by a similar degree as AAP-induced apoptosis in the presence of LiCl, SB-216763 or SB-415286. These findings show that GSK-3 plays a fundamental role in ER stress-induced apoptosis in HuH7 liver cells that is similar to its role in neuronal cells (Song *et al.*, 2002; Ghribi *et al.*, 2003). Moreover, our data support the hypothesis that AAP triggers an ER

stress response in these cells. This response is most likely to occur during the bioactivation of AAP to NAPQI which takes place in the ER by cytochrome P450-mediated metabolism (Tonge *et al.*, 1998; Zaher *et al.*, 1998). Indeed, it was recently suggested AAP-mediated renal tubular injury involved ER stress (Lorz *et al.*, 2004). Taken together, our findings suggest that AAP induces an ER stress response and that GSK-3 provides a mechanistic link between the ER and the ensuing activation of the mitochondrial pathway of apoptosis. However, the molecular mechanisms involved remain to be elucidated.

In conclusion, AAP induces apoptosis in HuH7 liver cells by activating the mitochondrial pathway of apoptosis through the release of pro-apoptotic factors such as cytochrome *c* and Smac/DIABLO and downstream caspase activation. GSK-3 was identified as a major regulator of the apoptotic response and is proposed to functionally link AAP-induced ER stress to the mitochondrial pathway of apoptosis.

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FOOTNOTES

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FIGURE LEGENDS

Figure 1. Acetaminophen induces apoptosis in HuH7 cells. A. HuH7 cells were treated in the absence (a) or presence of AAP (10 mM) for 24h (b) or 48h (c) before fixing, staining with Hoechst 33258 and visualization by fluorescence microscopy. B. HuH7 cells were treated in the absence (closed circles) or presence of AAP (10 mM) (open circles) and at the indicated time points the cells were harvested, stained with propidium iodide and analyzed for their DNA content by flow cytometry as described under *Methods*. Each point is the mean \pm SD of three independent experiments. ***, $p < 0.001$.

Figure 2. Processing of execution pro-caspases-3 and -7 in acetaminophen-induced apoptosis. A. HuH7 cells were exposed to AAP (10 mM) for 24 or 48h, or staurosporine (1 μ M for 24h) or puromycin (20 μ M for 24h), harvested and probed for the processing of pro-caspase-3 by Western blot analysis. Similarly, HuH7 cells were treated with 0.5, 5 or 10 mM AAP for 48h, staurosporine (1 μ M for 24h) or puromycin (20 μ M for 24h) and examined for the activation of caspase-7. B. Immunodetection of active (processed) caspase-3 by flow cytometry. Top, histogram showing the presence of active caspase-3 in control cells (.....) or following exposure to AAP (10 mM for 48h) (____). Bottom, time course of pro-caspase-3 processing following exposure to AAP (10 mM). C. Cleavage of cytokeratin-18 and α -fodrin in response to AAP. Control HuH7 cells (.....) and cells treated with AAP (10 mM) (____) were harvested after 48h, immunostained for cleaved cytokeratin-18 and analyzed by flow cytometry. The time course of cytokeratin-18 cleavage is shown in the middle panel [control cells, circles; cells treated with AAP in the absence (triangles) or presence (squares) of Z-VAD-fmk (100 μ M)]. Bottom panel, cleavage of α -fodrin in AAP-induced apoptosis. HuH7 cells were treated with AAP (10 mM for 48h), staurosporine or puromycin (as above) before immunoblotting for α -fodrin. D. AAP-induced apoptosis requires caspases. HuH7 cells, untreated (closed circles) or treated with AAP (10 mM) in the absence (open triangles) or

presence of Z-VAD-fmk (50 μ M, squares; 150 μ M, closed triangles) were assayed for loss of cell viability as measured by the release of cellular lactate dehydrogenase activity. Each point is the mean \pm SD of three independent experiments. *, $p < 0.05$; **, $p < 0.01$; ***, $p < 0.001$ (AAP + Z-VAD-fmk versus AAP alone); #, $p < 0.05$; ###, $p < 0.001$ (AAP versus control).

Figure 3. Release of apoptogenic factors from mitochondria, Bax translocation and loss of mitochondrial $\Delta\Psi_m$. HuH7 cells were exposed to AAP (10 mM) for 24h before (A) immunostaining for cytochrome *c* (a, cytochrome *c*; b, Hoechst 33258-counterstained cells), (B) Western blot analysis of mitochondrial (P) or cytosolic (S) content of cytochrome *c* and Smac/DIABLO and (C) Western blot analysis of mitochondrial (P) or cytosolic (S) content of Bax were performed as described under *Methods*. Loading controls were done by reprobating the blots for COX IV and Hsc70. D. HuH7 cells grown on glass coverslips were treated with AAP (10 mM) for 24h before staining with TMRE (500 nM) and Hoechst 33258 (2 μ g/ml) and transferring to a Zeiss POC chamber. The cells were viewed on a Zeiss Axiovert 135 fluorescence microscope for TMRE fluorescence (a) and Hoechst 33258 fluorescence (b). Cells undergoing apoptosis are indicated by arrows. Magnification, 400X.

Figure 4. Inhibitors of the mitochondrial PTP do not protect from acetaminophen-induced cytotoxicity. HuH7 cells were treated with AAP (10 mM) in the absence (black shaded) or presence (grey shaded) of (a) cyclosporin A (1 μ M) or (b) bongkreikic acid (50 μ M) before assaying for the loss of cell viability as described under *Methods*. Data are the mean \pm SD of three independent experiments.

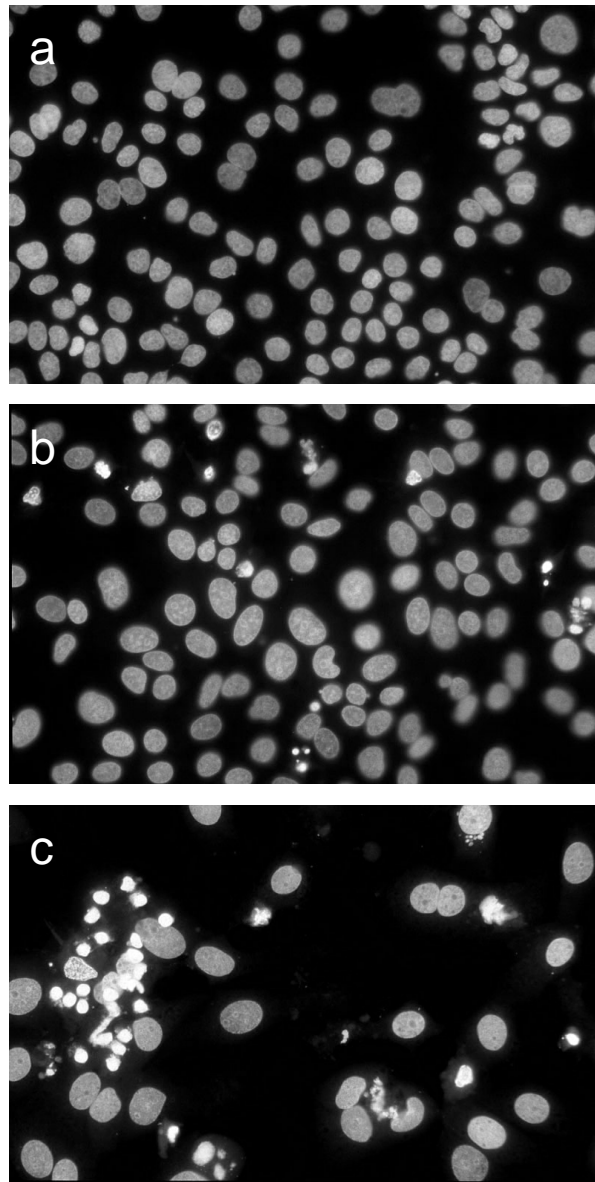
Figure 5. Acetaminophen induces the phosphorylation of JNK but its toxicity is insensitive to inhibitors of MAP kinase pathways. A. HuH7 cells, controls or treated with AAP (10 mM for 48h) or duroquinone (DQ, 500 μ M for 24h) were harvested and probed for phospho-JNK by Western blot analysis. Loading controls were carried out by reprobating the blots for total cellular JNK. B. HuH7 cells, untreated (closed circles) or treated with AAP (10 mM) in the

absence (open triangles) or presence of SP600125 (30 μ M, closed triangles), SB-203580 (10 μ M, diamonds) or PB98059 (50 μ M, squares) were assayed for loss of cell viability as measured by the release of cellular lactate dehydrogenase activity. Each point is the mean \pm SD of three independent experiments. #, $p < 0.05$; ##, $p < 0.01$; ###, $p < 0.001$ (AAP versus control).

Figure 6. GSK-3 mediates acetaminophen-induced apoptosis. A. HuH7 cells, untreated (closed circles) or treated with AAP (10 mM) in the absence (open triangles) or presence of LiCl (20 mM, squares) were assayed for loss of cell viability as measured by the release of cellular lactate dehydrogenase activity. B. Effect of the GSK-3 inhibitors SB-216763 (25 μ M, squares) and SB-415286 (3 μ M, diamonds) on AAP-induced cell death. C. The protection from AAP-induced cell death was caused by a decrease in caspase-3 activation. The histogram shows the presence of active caspase-3 as detected by flow cytometry in control cells (thin line) or following exposure to AAP (10 mM for 48h) in the absence (thick line) or presence (shaded) of SB-415286 (3 μ M). The cells were pre-incubated with the GSK-3 inhibitors for 1h prior to the addition of AAP. Each point is the mean \pm SD of three independent experiments. *, $p < 0.05$; **, $p < 0.01$ (AAP + inhibitor versus AAP alone); #, $p < 0.05$; ###, $p < 0.001$ (AAP versus control).

Figure 7. Role of GSK-3 in ER stress-mediated apoptosis. HuH7 cells were treated with thapsigargin (1 μ M, A) or duroquinone (500 μ M, B) in the absence (triangles) or presence (squares) of SB-415286 (3 μ M, A) or LiCl (20 mM, B). Control cells are represented by the circles. Cytotoxicity was assayed at the indicated time points as described under *Methods*. Data are the mean \pm SD of three independent experiments. *, $p < 0.05$; **, $p < 0.01$ (Thapsigargin or duroquinone + inhibitor versus thapsigargin or duroquinone alone); #, $p < 0.05$; ###, $p < 0.001$ (Thapsigargin or duroquinone versus control).

A



B

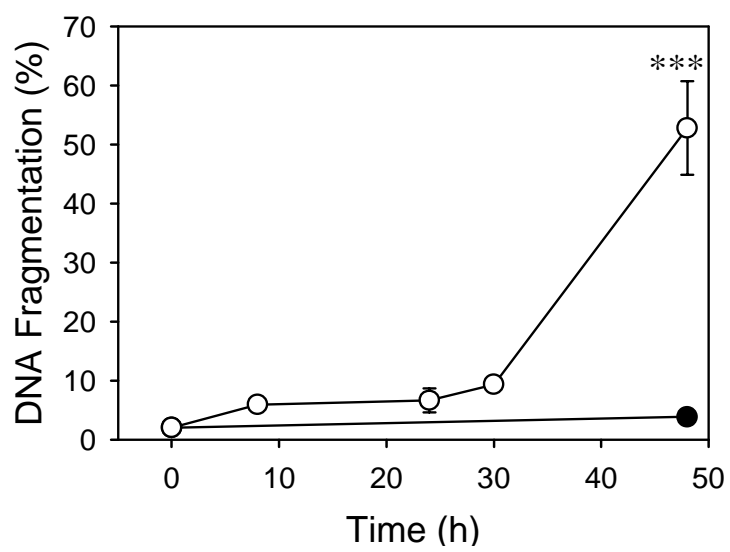


Figure 1

A

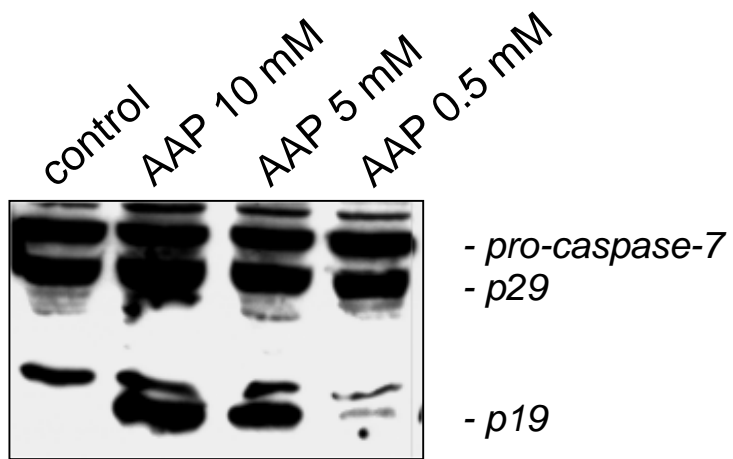
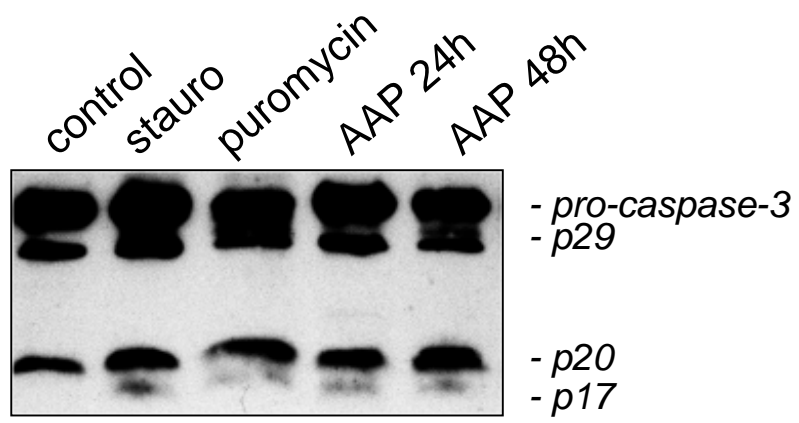


Figure 2A

B

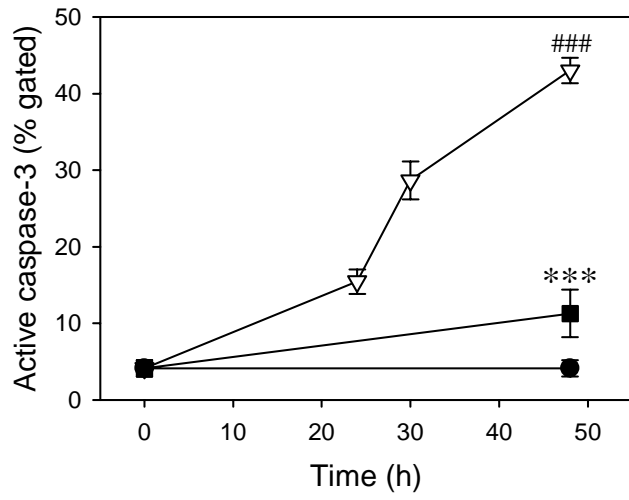
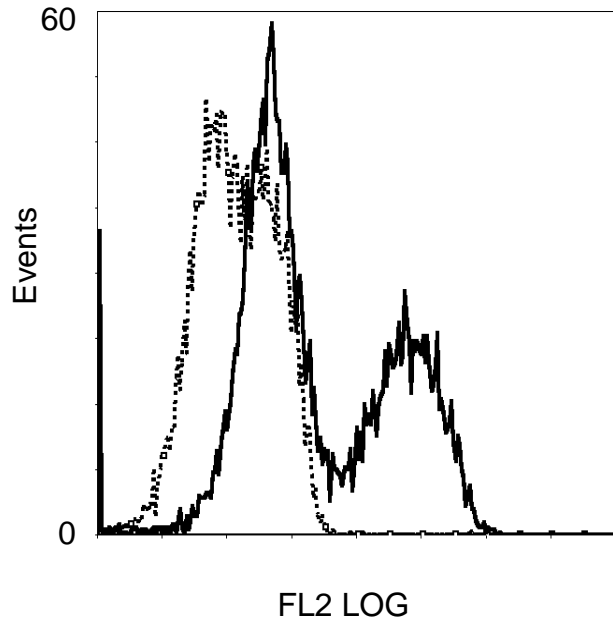


Figure 2B

C

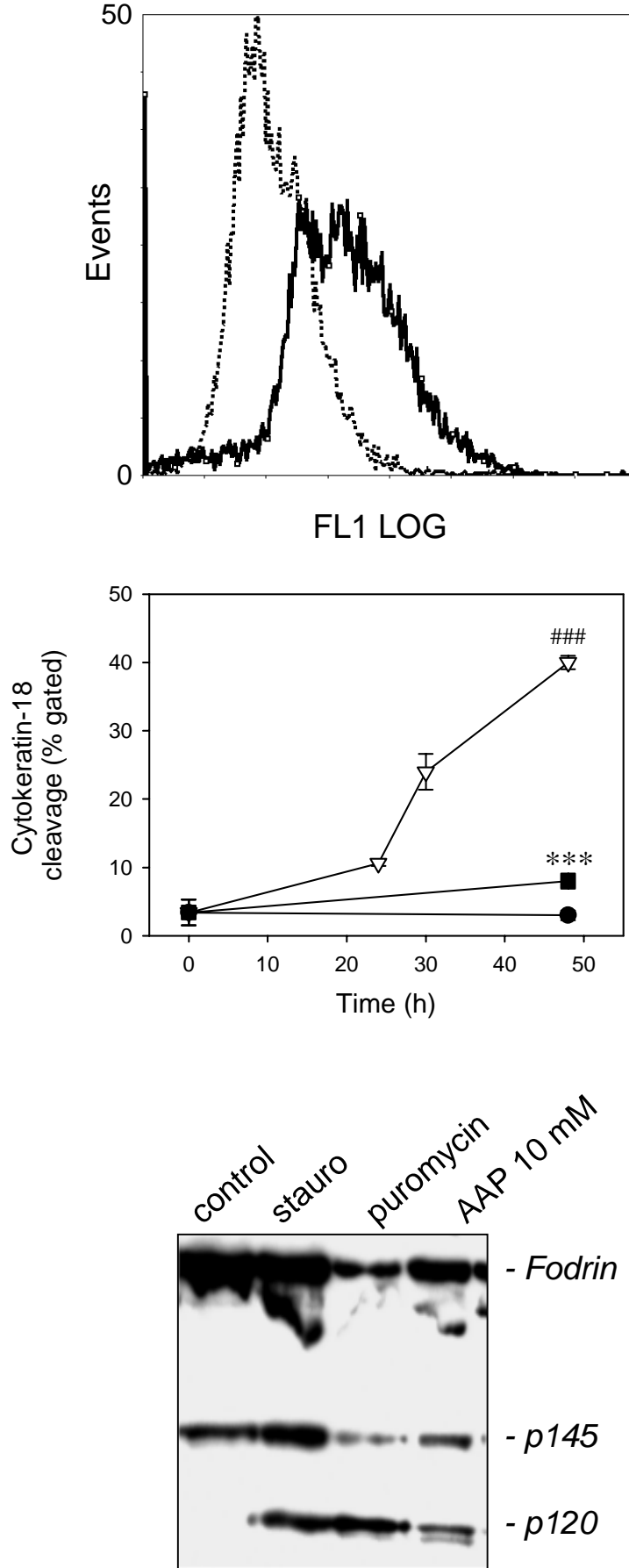


Figure 2C

D

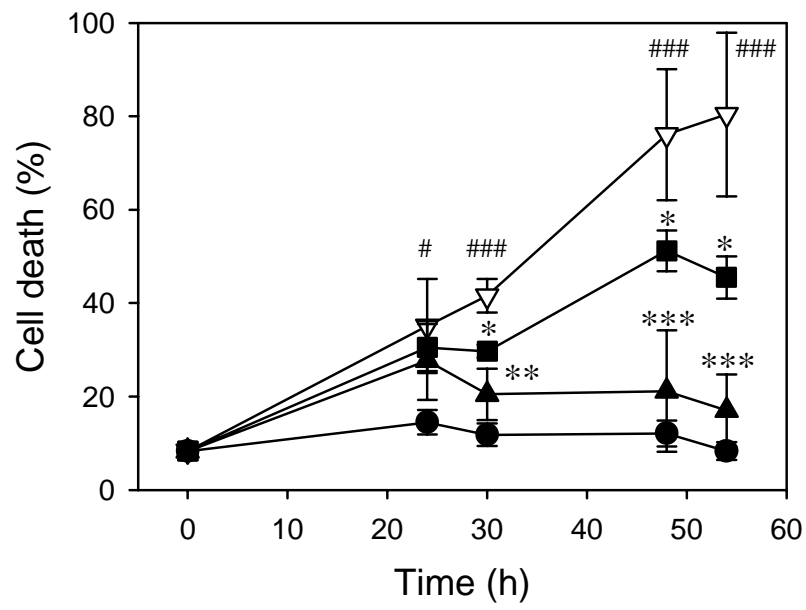
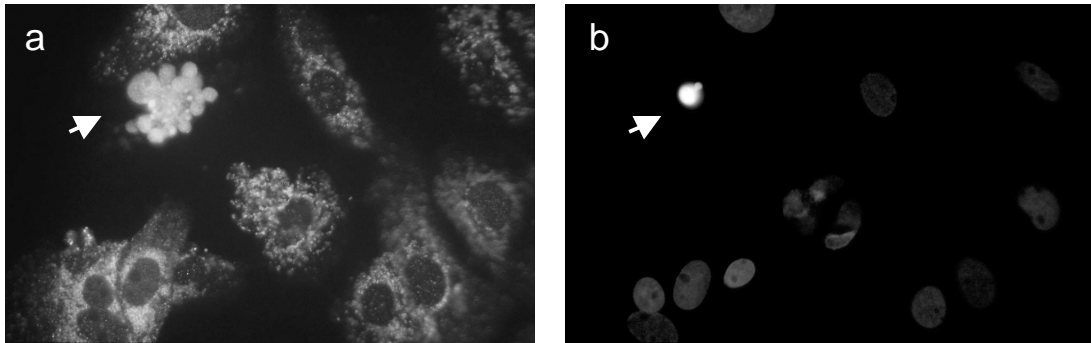


Figure 2D

A



B

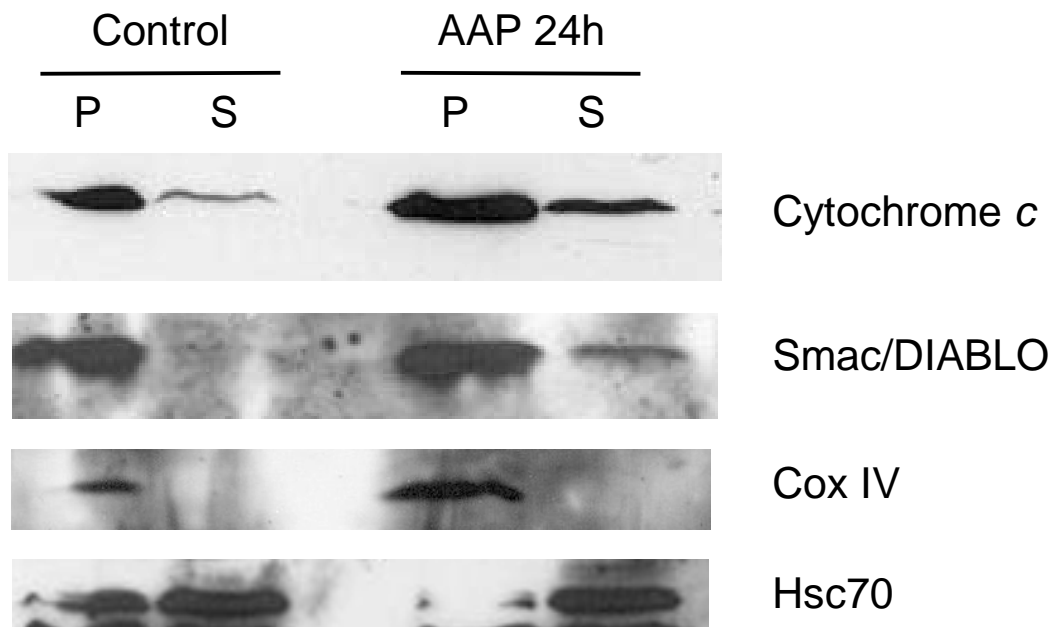
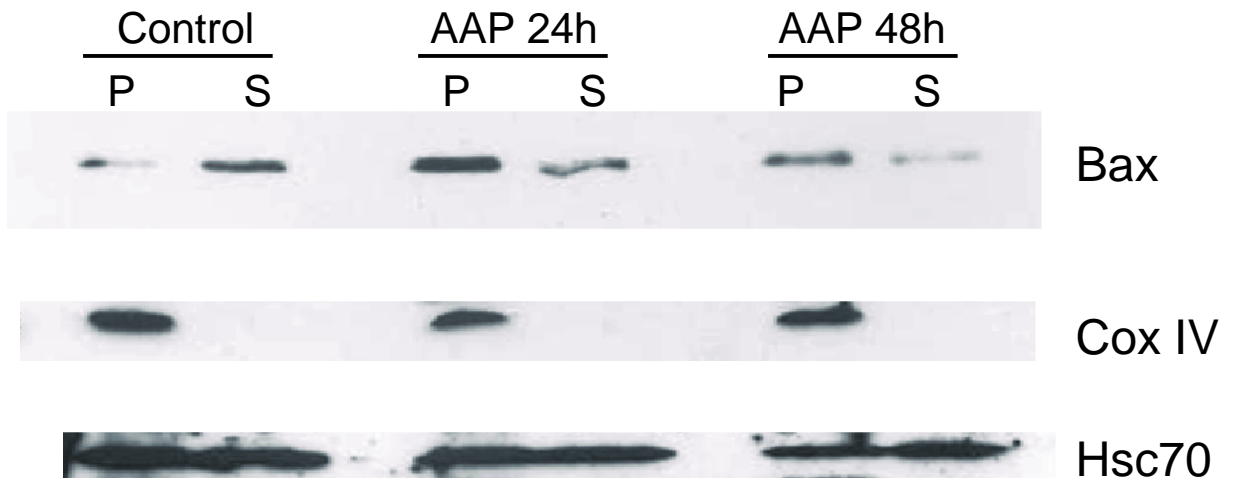


Figure 3AB

C



D

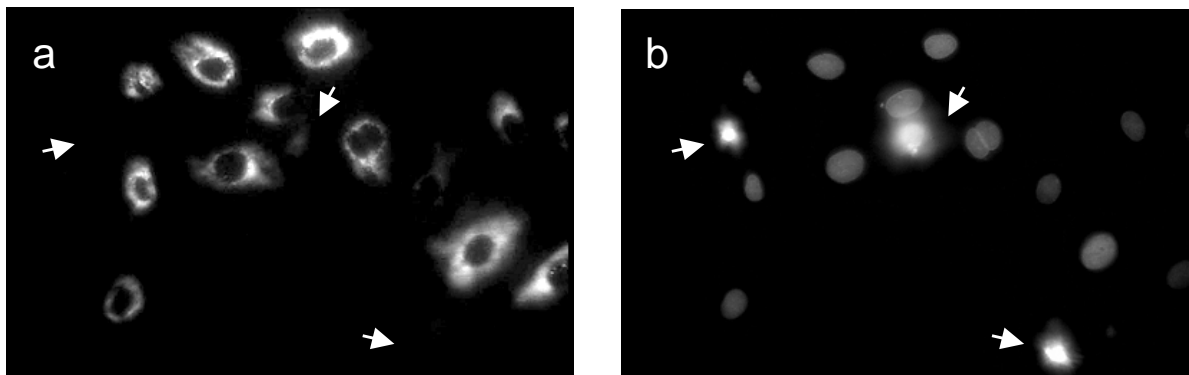


Figure 3CD

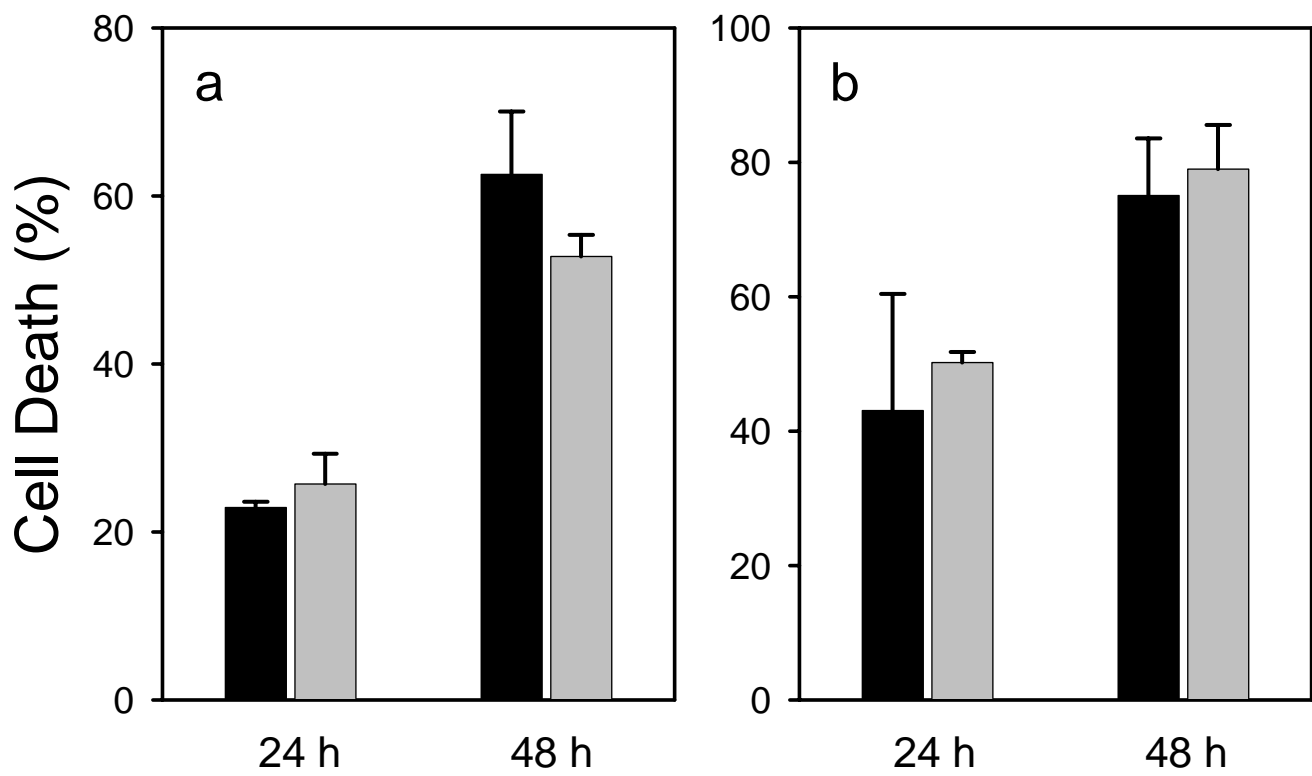
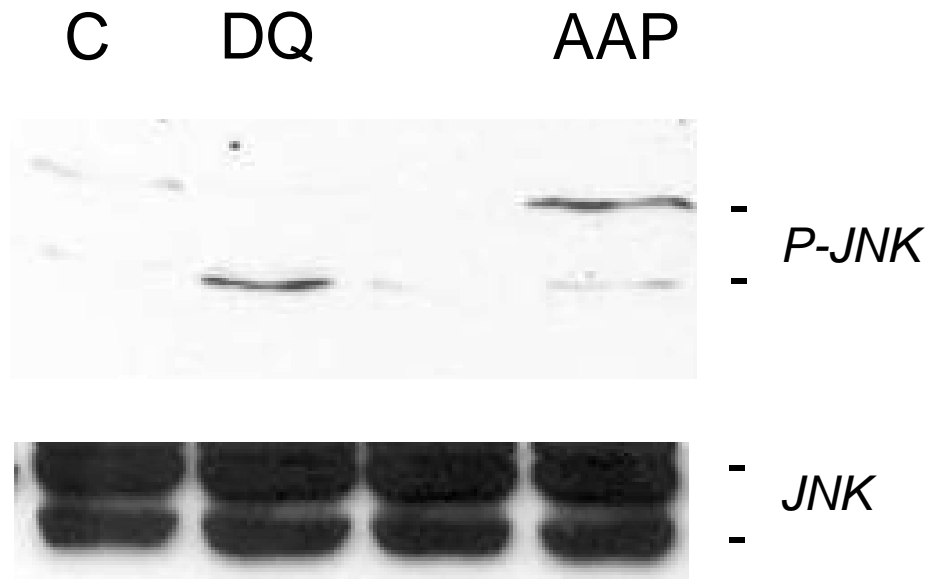


Figure 4

A



B

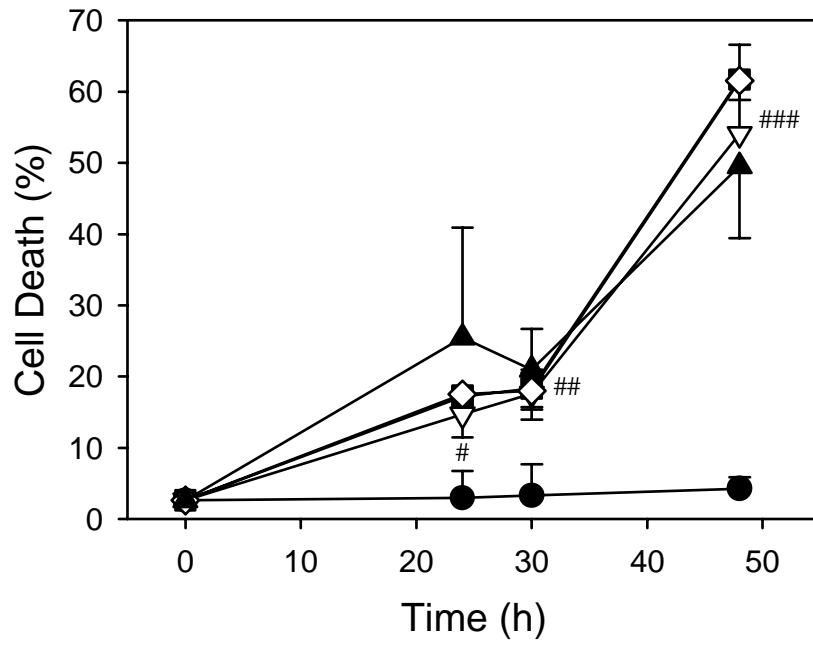


Figure 5

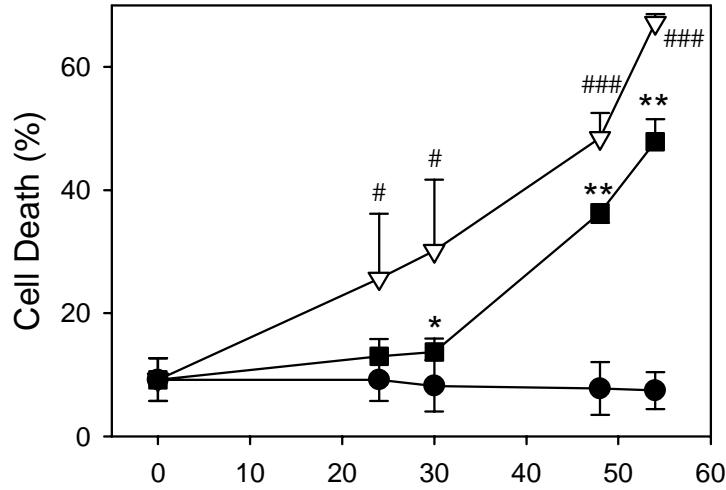
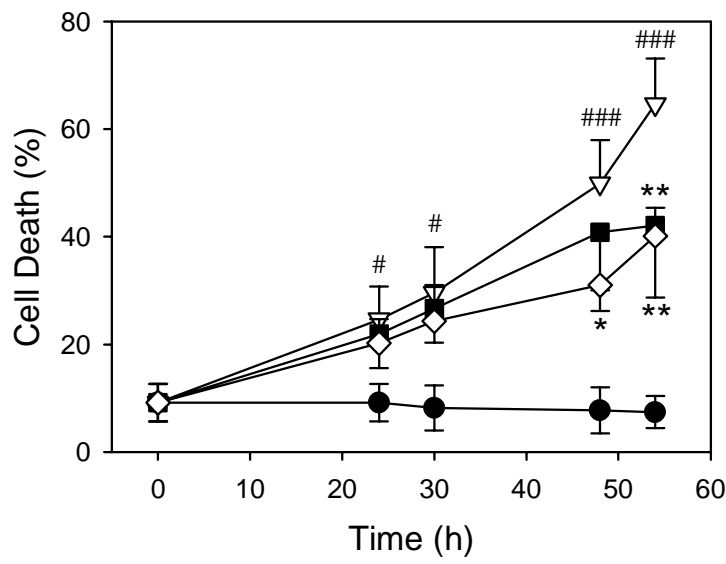
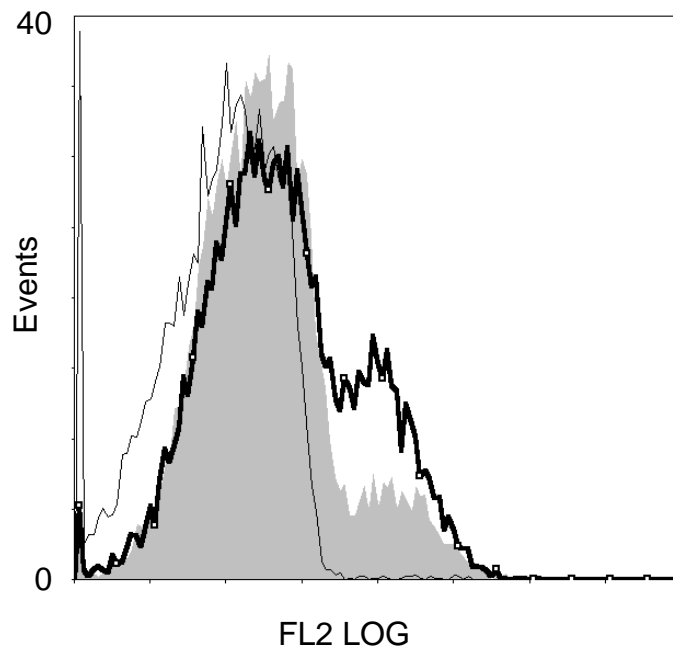
A**B****C**

Figure 6

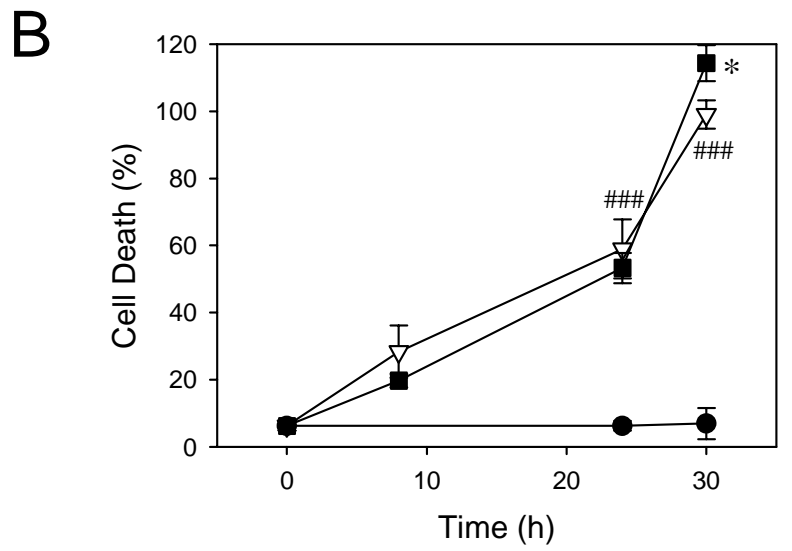
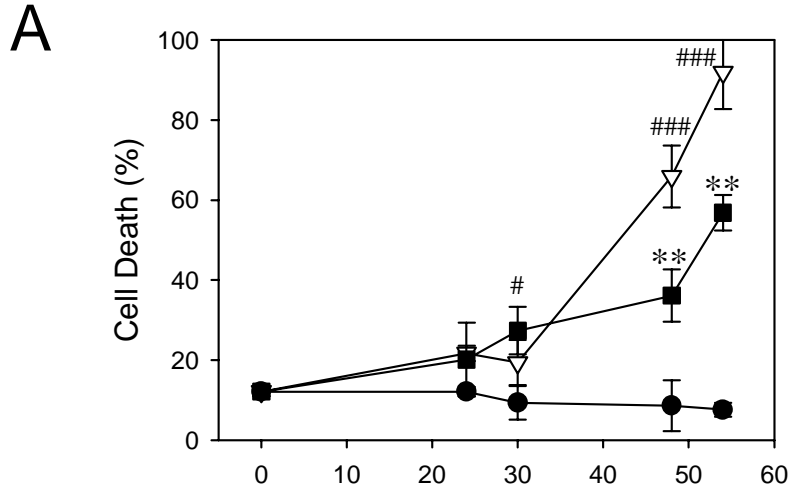


Figure 7