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Glycogen Synthase Kinase-3 (GSK-3) Inhibition Attenuates

Hepatocyte Lipoapoptosis

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Abstract

BACKGROUNDS AND AIMS—Saturated free fatty acids induce hepatocyte lipoapoptosis, a key pathologic feature of nonalcoholic steatohepatitis. The saturated free fatty acid palmitate induces hepatocyte lipoapoptosis via an endoplasmic reticulum stress pathway resulting in c-Jun-N-terminal (JNK) activation. Glycogen synthase kinase (GSK)-3 is a serine/threonine kinase which may also promote JNK activation. Thus our aim was to determine if GSK-3 inhibition suppresses palmitate induced JNK activation and lipoapoptosis.

METHODS—We employed mouse primary hepatocytes, Huh-7 and Hep3B cell lines for these studies.

RESULTS—Palmitate-induced GSK-3 activation was identified by phosphorylation of its substrate glycogen synthase. GSK-3 pharmacologic inhibition, by GSK-3 inhibitor IX and enzastaurin, significantly reduced PA-mediated lipoapoptosis. More importantly, Huh-7 cells in which either GSK-3 α or GSK-3 β isoforms were stably and selectively knocked down by shRNA displayed resistance to palmitate-induced cytotoxicity. GSK-3 pharmacological inhibitors and shRNA-targeted knockdown of GSK-3 α or GSK-3 β also suppressed JNK activation by palmitate. JNK activation, in part, promotes lipoapotosis by inducing expression of the pro-apoptotic effector p53-upregulated modulator of apoptosis (PUMA). Consistent with this concept, GSK-3 pharmacologic inhibition also reduced PUMA cellular protein levels during exposure to palmitate. On the other hand, the GSK-3 inhibitors did not prevent PA induction of ER stress.

In **CONCLUSION**, our results suggest GSK-3 activation promotes a JNK-dependent cytotoxic signaling cascade culminating in lipoapoptosis.

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Keywords

endoplasmic reticulum stress; glycogen synthase kinase; JNK; lipoapoptosis; nonalcoholic steatohepatitis; PUMA

INTRODUCTION

Nonalcoholic fatty liver disease (NAFLD) is an emerging public health problem linked to the increased incidence of obesity, insulin resistance and overt diabetes [1]. Up to 30% of the population is afflicted by NAFLD [2] and a subset of these patients develops nonalcoholic steatohepatitis (NASH). Individuals with NASH can progress to cirrhosis with its sequelae including the development of portal hypertension, end-stage liver disease and hepatocellular carcinoma [3]. Thus, the cellular and molecular mechanisms promoting liver injury in NASH are of biomedical and public health interest.

Obesity and insulin resistance are associated with increased circulating levels of nonesterified or free fatty acids (FFA), which are disproportionately increased in NASH as compared to NAFLD [4]. FFAs are taken up in the liver where they are esterified into neutral triglycerides. However, an excess of FFA is deleterious for the liver. Indeed, saturated FFAs are directly hepatotoxic, in part, by promoting apoptosis, which in this context is referred to as lipoapoptosis [5,6]. The importance of this process in human disease is highlighted by observations that hepatocyte lipoapoptosis and serum biomarkers of this pathologic process correlate with NASH severity [7,8]. Saturated FFA are more toxic than unsaturated FFA, presumably given their ability to disrupt endoplasmic reticulum (ER) homeostasis causing an ER stress response [9]. Biosensors of ER stress include protein kinase RNA-like ER kinase (PERK) and inositol-requiring protein-1a (IRE-1a). These two resident transmembrane-ER proteins sense and transduce the ER stress response [10]. PERK activation drives the expression of the transcription factor C/EBP-homologous protein (CHOP) [11,12]. CHOP in turn increases expression of Growth Arrest and DNA Damage Protein 34 (GADD34). PERK dimerization also results in inactivating phosphorylation of elongation initiation factor 2α (eIF2 α), thereby limiting protein synthesis. On the other hand GADD34 promotes protein synthesis by dephosphorylating $eIF2\alpha$, shutting down the unfolded protein response and exacerbating the ER stress [11]. IRE-1a possesses endonuclease activity for XBP-1 mRNA, creating a spliced form of XBP-1, the resultant spliced XBP-1 mRNA via transcriptional activity promotes degradation of misfolded ER glycoproteins [10,13].

We and others have reported that FFA-induced ER stress is associated with c-Jun-Nterminal kinase (JNK) activation, which has been well documented in both rodent and human steatohepatitis [14–17]. ER stress-associated JNK activation promotes apoptosis by modifying expression and function of pro-apoptotic members of the Bcl-2 family, especially the Bcl-2 homology 3 (BH3) only protein Bcl-2-interacting mediator of cell death (Bim) and p53-upregulated modulator of apoptosis (PUMA) [18,19]. During lipotoxicity, JNK appears to promote apoptosis predominantly by inducing expression of PUMA. As we have previously reported, JNK inhibition reduces PA-induced increases of PUMA expression [18], and protects against PA induced apoptosis [14]; also PUMA knockdown by shRNA markedly reduces lipoapoptosis [18]. On the other hand, JNK inhibition does not prevent PA-induced Bim protein induction/activation during lipoapoptosis [20]. However, the precise cellular and molecular mechanisms resulting in JNK activation have not been fully elucidated, and mechanistic insight into this process may identify therapeutic targets to treat human NASH. Glycogen synthase kinase (GSK)- 3α and GSK- 3β are serine/threonine kinases which can participate in pro-apoptotic signaling [21–24]. These kinases share 84% overall identity and 98% homology in their catalytic domains [25]. The mechanisms governing their respective and unique activation remain unclear. However, several studies have identified GSK-3activation during ER stress [26–28]. For example, pharmacologic inhibition of GSK-3attenuates ER stress-induced apoptosis in neuroblastoma cells, neurons and fibroblasts [29,30]. Moreover, GSK-3 is capable of activating JNK directly and GSK-3 activation of JNK contributes to an acute model of liver injury by acetaminophen [24]. Given this information, a potential role for GSK-3 in FFA-associated ER stress mediated JNK activation warrants exploration.

MATERIALS AND METHODS

Cell lines, cell isolation, and transfection

The human hepatocellular carcinoma cell lines, Huh-7, and Hep3B cells were cultured in Dulbecco's modified Eagle's medium (DMEM) as previously described [31]. Mouse hepatocytes were isolated from C57/Bl6 wild type mice by collagenase perfusion, purified by Percoll gradient, and cultured in Waymouth Medium [32]. Huh-7 cells were transfected separately with 1 μ g/ml DNA plasmid (GSK-3 α or GSK-3 β MISSION short hairpin (sh) RNA lentiviral plasmid; Sigma Aldrich) using Lipofectamine (Invitrogen, Carlsbad, CA). Stably transfected GSK-3 α and GSK-3 β clones were selected in medium containing 1200 mg/liter G418 and screened by immunoblot analysis.

Fatty acid treatment

Palmitate (PA) (# P5585) was obtained from Sigma-Aldrich (St. Louis, MO). PA was dissolved in isopropanol at a concentration of 160 mM. PA was added to DMEM containing 1% bovine serum albumin to obtain a physiologic ratio between bound and unbound FFA in the media [33]. The concentration of PA used in the experiments varied between 400–800 μ M and are similar to the fasting total FFA plasma concentrations observed in human with non-alcoholic steatohepatitis [4,34].

Quantification of apoptosis, Bax activation and mitochondrial membrane potential (MMP) Assay

Apoptosis was quantified by evaluating the characteristic nuclear changes of apoptosis using the nuclear binding dye DAPI (Molecular Probes, Eugene, OR) and fluorescence microscopy (Zeiss LSM 510, Carl Zeiss, Jena, Germany)[14]. Caspase 3/7 activation in cell lines was measured using Apo-ONE homogeneous caspase 3/7 kit (Promega, Madison, WI) according to the manufacturer's instructions [14]. Immunocytochemistry for active confirmation of Bax was performed using mouse anti-universal Bax (clone 6A7,1:100 dilution, Santa Cruz) as previously described by us in detail [18]. Huh 7 cells were cultured on glass-bottom plates (MatTek, Ashland, MA) and treated with reagents for 18 hours. The mitochondrial membrane potential was quantified as previously described by us in detail [35].

Real time polymerase chain reaction (PCR)

Real time PCR was performed as previously discussed by us [20] using the following primers: human PUMA, forward 5'-GACGACCTCAACGCACAGTA-3' and reverse 5'-AGGAGTCCCATGATGAGATTGT-3', size 101 bp; human CHOP, forward 5'-ATGGCAGCTGAGTCAATGCCTTTC-3' and reverse 5'-AGAAGCAGGGTCAAGAGTGGTGAA-3', size 177 bp; and human GADD34 forward 5'-CGACTGCAAAGGCGGC-3' and reverse 5'-CAGGAAATGGACAGTGACCTTC-3', size

107 bp. As internal control, primers for 18S ribosomal RNA (rRNA) were used: forward 5'-CGTTCTTAGTTGGTGGAGCG-3' reverse 5'-CGCTGAGCCAGTCAGTGTAG-3', size 212 bp

Detection of XBP-1 splicing

Human XBP-1 cDNA encompassing the region of PstI restriction site was amplified using the following primers: forward 5'-AAACAGAGTAGCAGCTCAGACTGC-3' and reverse 5'-TCCTTCTGGGTAGACCTCTGGGAG-3'. The PCR products were then digested with PstI restriction enzyme (Biolabs, catalog #R0140) for 1 hour at 37°C, and subjected to 1.5% agarose gel electrophoresis; the gels were photographed under UV transillumination as previously described by us in detail [31]. Thapsigargin (TG) was used as an established inducer of XBP-1 splicing [9].

Immunoblot analysis

Whole cell lysates were prepared and subjected to immunoblot analysis as previously described [20]. Samples containing 50 µg proteins were resolved by 10–15% SDS-PAGE, transferred to nitrocellulose membranes, and incubated with primary antibodies at a dilution of 1:1000. Membranes were incubated with appropriate horseradish peroxidase-conjugated secondary antibodies (Biosource International, Camarillo, CA) at a dilution of 1:3000. Bound antibody was visualized using chemiluminescent substrate (ECL, Amersham, Arlington Heights, IL) and was exposed to Kodak X-OMAT film (Eastman Kodak, Rochester, NY).

Antibodies and Reagents

Rabbit anti-JNK (#9252), rabbit anti-phospho-JNK (Thr183/Tyr185) (# 9251), rabbit anti-Phospho-eIF2 α (Ser51) (#9721), rabbit anti-eIF2 α (#9722), rabbit anti-GSK-3 α (#9338), rabbit anti-GSK-3 β (#9315) and rabbit anti-phospho-glycogen synthase antibody (Ser649) (#3891) were from Cell Signaling Technology (Beverly, MA). Rabbit anti-PUMA was from Abcam (Cambridge, MA). Goat anti- β -actin was from Santa Cruz Biotechnology (Inc., Santa Cruz, CA). GSK-3 inhibitor IX (GSK IX) (#361550) and thapsigargin (#586005) were from Calbiochem (San Diego, CA). Z-VD(OMe)VAD(OMe)-FMK (ZVAD) was from MP Biomedicals (Aurora, Ohio). Enzastaurin (Enz) was provided by Lilly (Indianapolis, Indiana). BSA, Bradford reagent, and all other chemicals were from Sigma Aldrich.

Statistical analysis

All data represent at least three independent experiments and are expressed as means \pm SE. Differences between groups were compared using Student's *t* tests and one-way analysis of variance with *post hoc* Dunnett test, and *p* values <0.05 were considered statistically significant.

RESULTS

GSK-3 inhibitor attenuates Palmitate-mediated lipoapoptosis

Initially we examined the effect of GSK- 3α and GSK- 3β inhibition on PA-mediated lipoapoptosis in Huh-7 cells. The GSK IX [36] and enzastaurin (a bisindolylmaleimide serine threonine kinase inhibitor that inhibits protein kinase C and GSK-3 [37]) were employed to inhibit both GSK- 3α and GSK- 3β . We confirmed that PA stimulated GSK-3, and the inhibitors effectively reduced PA-induced GSK-3 activity by assessing phosphorylation of glycogen synthase, a specific GSK-3 substrate. Following PA treatment, glycogen synthase phosphorylation was identified by phospho immunoblot analysis which was reduced by both inhibitors (Fig. 1A). The inhibitors also reduced PA-mediated

cytotoxicity in a concentration-dependent manner, each with a maximal inhibition at 2 μ M (Fig. 1B). Because caspase 3/7 activation mediates the apoptotic phenotype, we next confirmed that the observed apoptosis was caspase-mediated. Indeed, PA-induced apoptosis was caspase-dependent, as it was reduced by the caspase inhibitor ZVAD (Fig. 1C, 1D). Both GSK-3 inhibitors also reduced apoptosis as assessed by this biochemical parameter (Fig. 1E). The GSK-3 α and GSK-3 β inhibitor IX also reduced caspase 3/7 activity in Hep3B cells and primary murine hepatocytes (Fig. 1F, 1G). Finally to exclude off-target effect of the pharmacologic inhibitor, apoptosis was examined in Huh-7 cells, in which GSK-3 α or GSK-3 β had been reduced by shRNA technology (Fig. 2A). Importantly, the knockdown of either GSK-3 α or GSK-3 β reduced PA-induced lipoapoptosis as assessed by both morphological and biochemical criteria (Fig. 2B, 2C).

Activation of Bax, a known mediator of mitochondrial dysfunction, is required for induction of hepatocyte lipoapoptosis downstream of JNK and BH3-only proteins activation [14,18]. Activated Bax was identified by immunofluorescence using the 6A7 monoclonal antibody which detects an active Bax conformation [38]. As previously reported, Bax activation was observed in PA-treated cells [14], but this activation was decreased when cells were treated with PA plus GSK IX (Fig. 3A). Likewise, selective shRNA targeted knockdown of either GSK-3α or GSK-3β, also reduced PA-induced Bax activation (Fig. 3B). Along with Bax activation, mitochondrial dysfunction with loss of mitochondrial membrane permeability (MMP) is a prominent feature of the mitochondrial pathway of apoptosis. The MMP was reduced to $39 \pm 3\%$ of the control value following treatment with PA; in contrast the MMP was maintained with the addition of the GSK-3 inhibitor (Fig 3C). Taken together, these data indicate that GSK-3α and GSK-3β pharmacologic or genetic inhibition potently suppresses saturated FFA cytotoxicity. Both isoforms appear to contribute to cytotoxicity by PA suggesting cooperation between the isoforms during lipoapoptosis. Given that GSK-3 inhibitor cytoprotection was similar in cell lines and primary mouse hepatocytes, we employed the Huh-7 cell line for subsequent studies to further elucidate the role of GSK-3 in lipotoxicity.

GSK-3 activation is downstream of ER stress-induced Lipoapoptosis

Lipoapoptosis can be attenuated downstream or upstream of FFA-induced ER stress [14,18,20,31,39]; therefore, we sought to determine if GSK-3 inhibition modulated PAmediated ER stress. However, the pharmacologic inhibitors GSK-3 IX and enzastaurin did not prevent *CHOP* mRNA expression (Fig. 4A). GSK IX also did not prevent *GADD34* mRNA expression (Fig. 4B), or eIF2 α phosphorylation (Fig. 4C) by PA, all indices of PERK activation. Likewise, GSK IX did not block *XBP-1* mRNA splicing (Fig. 4D), an indicator of IRE-1 α activation. Collectively these data suggest that GSK-3 inhibition attenuates PAinduced lipoapoptosis downstream of ER Stress.

GSK-3 inhibition reduces JNK activation and PUMA upregulation during treatment with PA

Interestingly, activating JNK phosphorylation by PA was significantly reduced in the presence of the GSK IX (Fig. 5A) or enzastaurin (data not shown). Consistent with the pharmacologic inhibition, knockdown of GSK-3 α or GSK-3 β also attenuated JNK activation by palmitate (Fig. 5B). Consistent with this observation, GSK IX or enzastaurin also effectively attenuate upregulation of *PUMA* mRNA expression by PA (Fig. 6A). GSK IX was effective in reducing PA-induced PUMA expression at the protein level (Fig. 6B). However GSK IX did not reduce JNK activation by thapsigargin (Fig. 5C). These results suggest that GSK-3 inhibition may inhibit PA- mediated cell death through inhibition of JNK and reduction of the associated increase of PUMA.

DISCUSSION

Results of the present study provide mechanistic insights regarding the pro-apoptotic effects of GSK-3 during FFA-induced lipoapoptosis. The principal findings of this study indicate that during PA-mediated lipoapotosis in vitro: i) either pharmacological or genetic inhibition of GSK-3 attenuates apoptosis; ii) GSK-3 inhibition does not reduce the ER stress response; and iii) GSK-3 inhibition attenuates JNK activation, and subsequent PUMA induction. Each of these results is discussed in greater details below.

GSK-3 is a serine/threenine protein kinase with two isoforms (α and β). Both isoforms are ubiquitously expressed in cells and tissues, and have similar, although, not identical biochemical properties [25]. However, they are not interchangeable functionally, as demonstrated by the embryonic-lethal phenotype observed when the gene that encodes GSK-3 β was knocked out [40]. Although GSK-3 has multiple functions in health, in the context of cytotoxic stimuli, GSK-3 (especially GSK-3β) appears to function as a proapoptotic kinase [41]. Previous studies have suggested that GSK-3ß promotes the mitochondrial pathway of apoptosis through phosphorylation of Bax thereby facilitating its mitochondrial translocation [21]. More recently GSK-3β was also reported to activate JNK in acetaminophen-induced liver injury [24]. These prior studies have focused on GSK- 3β rendering the role of GSK-3 α in apoptosis ambiguous [42]. Our current study extends these observations by suggesting that both GSK-3 isoforms contribute to lipoapoptosis. Indeed, shRNA targeted knockdown of either GSK-3 isoform was protective against lipoapoptosis. The two isoforms apparently have complementary, non redundant roles in this model of apoptosis. The mechanisms by which these two isoforms cooperate in lipoapoptosis will require further detailed study.

Lipoapoptosis can be inhibited either upstream or downstream of ER stress. Although GSK-3 inhibition has been reported to reduce ER stress [28], in our current study the PA-induced ER stress response was not repressed by GSK-3 inhibition. These observations indicate that GSK-3 inhibition likely attenuates lipoapoptosis downstream of ER stress; however, we can not exclude a non ER stress-dependent pathway of GSK-3 activation. How GSK-3 is activated is not clear. Several studies suggest GSK-3 is activated by protein phosphatase 2A (PP2A) [43,44], GSK-3 β is also known to be activated by PP2A during ceramide induced apoptosis [45]. Moreover, we have previously reported that PP2A activity was stimulated by saturated free fatty acids in hepatocytes during lipoapotosis [20]. Thus, PP2A activation is a potential mechanism contributing to GSK-3 stimulation during lipoapoptosis.

Activated JNK is an important downstream mediator of ER stress-associated lipoapoptosis and it has been implicated in human and murine steatohepatitis [15–18]. Several studies have suggested that GSK-3β is involved in JNK activation through interaction with upstream kinases such as mitogen-activated protein kinase/extracellular signal-regulated kinase kinase (MAPKKK) and mixed lineage kinase (MLK) in neuronal and kidney cell lines [46,47]. GSK-3 was also reported to cooperate with JNK for the execution of the c-Jun stress response and neuronal death in response to trophic deprivation [22]. We note that in our present study, GSK-3 inhibition reduced PA-induced JNK phosphorylation, but it did not inhibit JNK phosphorylation in response to thapsigargin-induced ER stress. These data indicate that GSK-3 inhibition is not a general inhibitor of ER stress-associated JNK activation, but rather specifically antagonizes JNK activation by cytotoxic FFA.

The current study demonstrates that GSK-3 inhibition attenuates JNK-dependent dysregulation of PUMA, a key pro-apoptotic protein. PUMA promotes Bax activation and the mitochondrial pathway of cell death (Fig. 7). Activated Bax induces mitochondrial outer

membrane permeabilization (MOMP) which leads to the egress of these pro-apoptotic mediators into the cytosol. Subsequently, these mediators help activate downstream effectors caspases culminating in the apoptotic phenotype [48]. Collectively, our study suggests that GSK-3 genetic or pharmacologic inhibition attenuates saturated FFA-induced cell death in hepatocytes by inhibiting JNK-mediated PUMA upregulation. Given this mechanistic insight, we speculate that GSK-3 inhibition could have a potential therapeutic role in human NAFLD.

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Abbreviations

nonalcoholic fatty liver disease
nonalcoholic steatohepatitis
free fatty acids
endoplasmic reticulum
protein kinase RNA-like ER kinase
inositol-requiring protein-1α
C/EBP-homologous protein
growth arrest and DNA damage protein 34
elongation initiation factor 2α
c-Jun-N-terminal kinase
Bcl-2 homology 3
Bcl-2-interacting mediator of cell death
p53-upregulated modulator of apoptosis
glycogen synthase kinase
palmitate
mitochondrial membrane potential
short hairpin
polymerase chain reaction
thapsigargin
GSK-3 inhibitor IX
Z-VD(OMe)VAD(OMe)-FMK
enzastaurin
protein phosphatase 2A
vehicle
wild type

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Figure 1. GSK-3 inhibition attenuates PA mediated apoptosis

(A) Whole cell lysates were prepared from Huh-7 cells treated with vehicle (Veh) or PA at 800 μ M in the presence of the GSK-3 inhibitors, GSK IX or enzastaurin (Enz) (10 μ M) for 4, 8 and 16 hours, or ZVAD (25 μ M) for 16 hrs. Immunoblot analyses were performed for phosphorylated glycogen synthase (Phospho-GS) and β -actin was used as a control for protein loading. (B, C) Huh-7 cells were treated for 24 hours with Veh or PA at 800 μ M in the presence of either an increasing concentrations of GSK IX or Enz up to 2 μ M, or ZVAD (25 μ M). Apoptosis was assessed by morphological criteria after DAPI staining. Data represents the mean ± SEM for three experiments. (D, E) Huh-7 cells were treated for 24 hours with Veh or PA at 800 μ M in the presence of either GSK IX, Enz at 2 μ M, or ZVAD (25 μ M). (F) Hep3B cells (G) or mouse primary hepatocytes were treated for 16 hours with Veh or PA at 400 μ M in the presence of GSK IX at 2 μ M. (D, E, F and G) Caspase 3/7 catalytic activity was measured by a fluorogenic assay. Fold-increase was determined over control value (vehicle-treated cells), arbitrarily set to 1. Data represent the mean ± SEM for three experiments. *p < 0.05, Veh-treated cells vs PA-treated cells; **p < 0.05, PA-treated cells vs. PA plus GSK IX-treated cells or PA plus Enz-treated cells or PA plus ZVAD.

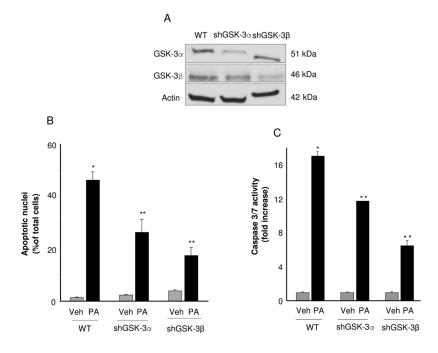
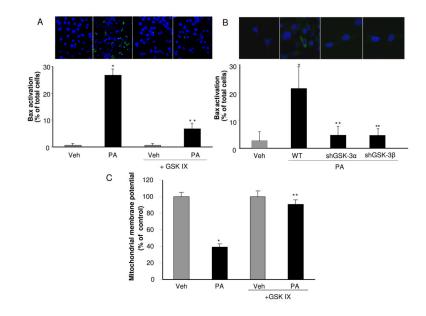
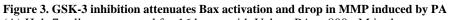


Figure 2. GSK-3a and GSK-3β targeted shRNA reduce PA-mediated lipotoxicity

Huh-7 Wild type (WT) or Huh-7 cells stably expressing short hairpin RNA targeting GSK-3 α (shGSK-3 α) or GSK-3 β (shGSK-3 β) were treated for 16 hours with Veh, or PA at 400 μ M. (A) Effective and selective downregulation of GSK-3 α or GSK-3 β protein levels in shGSK-3 α or shGSK-3 β Huh-7 cells, respectively, compared to WT Huh-7 cells was verified by immunoblot analysis on whole cell lysates. (B) Apoptosis was assessed by morphological criteria after DAPI staining. (C) Caspase 3/7 catalytic activity was measured by the fluorogenic assay. Fold-increase was determined over control value (vehicle-treated cells), arbitrarily set to 1. *p < 0.05, Veh-treated cells vs PA-treated cells; **p < 0.05, PA-treated WT cells vs PA-treated shGSK-3 α or shGSK-3 β .





(A) Huh-7 cells were treated for 16 hours with Veh or PA at 800 μ M in the presence of the GSK-3 inhibitor GSK IX at 2 μ M. (B) WT Huh-7, shGSK-3 α Huh-7 and shGSK-3 β Huh-7 were treated for 16 hours with Veh, or PA at 400 μ M. (A and B) Cells were fixed and Bax activation was assessed using conformation specific antisera (6A7) and immunofluorescence microscopy. Representative images of three independent experiments are depicted. Bax activation was quantified in 5 random 40 X objective field for each condition with automated software. (C) Huh-7 cells were treated for 16 hours with Veh or PA at 400 μ M in the presence of the GSK-3 inhibitor GSK IX at 10 μ M. Mitochondrial depolarization was measured using tetramethylrhodamine methylester. A minimum of 15 randomly selected cells were analyzed per condition from multiple microscopic fields. Data represents the mean \pm SEM for three experiments. *p < 0.05, Veh-treated cells vs PA-treated cells; **p < 0.01, PA-treated cells vs PA plus GSK IX-treated cells or PA-treated WT cells vs. PA-treated shGSK-3 α or shGSK-3 β .

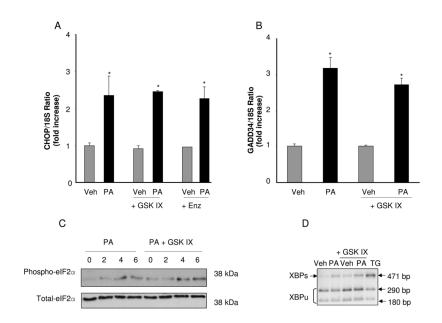


Figure 4. GSK-3 inhibition does not decrease PA-induced CHOP and GADD34 up-regulation, eIF2a phosphorylation or XBP-1 splicing

Huh-7 cells were treated with Veh or PA at 800 μ M in the presence of the GSK-3 inhibitors GSK IX or Enz at 2 μ M. (A and B) Total RNA was extracted 8 hours after treatments, and *CHOP* and *GADD34* mRNA were quantified by real-time PCR. Fold induction was determined after normalization to 18S. Data represent the mean ± SEM, of three independent experiments. *p < 0.05, Veh-treated cells vs PA-treated cells. (C) Whole cell lysates were prepared 2, 4 and 6 hours after treatments. Immunoblot analyses were performed for phosphorylated eIF2 α and total eIF2 α was used as a control for protein loading. (D) *XBP-1* cDNA was amplified by PCR, followed by incubation with PstI. In Veh-treated cells, most XBP-1 PCR products were cut by PstI, producing 290 bp and 180 bp amplification products, indicative of native and unspliced form of *XBP-1* mRNA (XBP1u). A 471 bp amplification product, indicative of spliced *XBP-1* mRNA (XBP1s) was mainly observed in PA-treated cells. Thapsigargin (TG)-treated cells at 450 nM were used as a positive control for *XBP-1* mRNA splicing.



Figure 5. Pharmacological or genetic inhibition of GSK-3 reduces PA-induced activating JNK phosphorylation

Whole cell lysates were prepared from Huh-7 cells treated for 4 hours (A) with Veh or PA at 800 μ M in presence of the pharmacological GSK-3 inhibitor GSK IX (2 μ M), and (C) with Veh or thapsigargin (TG) (450 nM) in the presence of GSK IX (2 μ M). (B) Whole cell lysate were prepared from WT Huh-7, shGSK-3 α Huh-7 and shGSK-3 β Huh-7 treated for 4 and 8 hours with Veh or PA at 800 μ M. (A, B and C) Immunoblot analyses were performed for phosphorylated JNK; total JNK was used as a control for protein loading.

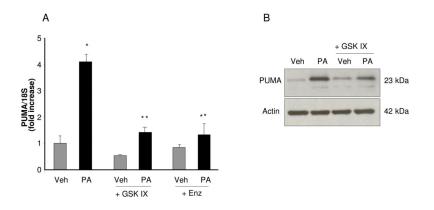


Figure 6. GSK-3 inhibitor suppresses PA-mediated PUMA upregulation

(A and B) Huh-7 cells were treated with Veh or PA at 800 μ M in the presence of the GSK-3 inhibitors GSK IX or Enz at 2 μ M. (A) Total RNA was extracted 8 hours after treatments and *PUMA* mRNA was quantified by real-time PCR. Fold induction was determined after normalization to 18S. Data represent the mean ± SEM, of three independent experiments. **p* < 0.05, Veh-treated cells vs PA-treated cells; ***p* <0.05, PA-treated cells vs. PA plus GSK IX or PA plus Enz-treated cells. (B) Whole cell lysates were prepared 16 hours after treatments and immunoblot analyses were performed for PUMA; β-actin was used as a control for protein loading.



Figure 7. Proposed model for the effects of GSK-3 inhibition on PA-induced apoptosis

GSK-3 inhibition blocks JNK phosphorylation thereby preventing PUMA upregulation, Bax activation, and attenuates PA-induced cell death. ¹ reference ⁹, ² reference ¹⁸, ³ reference ¹⁴.