

doi: 10.1093/toxsci/kfv091 Advance Access Publication Date: May 7, 2015 Research Article

Cytotoxic Synergy Between Cytokines and NSAIDs Associated With Idiosyncratic Hepatotoxicity Is Driven by Mitogen-Activated Protein Kinases

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ABSTRACT

Nonsteroidal anti-inflammatory drugs (NSAIDs) are among the most frequent causes of idiosyncratic, drug-induced liver injury (IDILI). Mechanisms of IDILI are unknown, but immune responses are suspected to underlie them. In animal models of IDILI, the cytokines tumor necrosis factor-alpha (TNFα) and interferon-gamma (IFNγ) are essential to the pathogenesis. Some drugs associated with IDILI interact with cytokines to kill hepatocytes in vitro, and mitogen-activated protein kinases (MAPKs) might play a role. We tested the hypothesis that caspases and MAPKs are involved in NSAID/cytokine-induced cytotoxicity. NSAIDs that are acetic acid (AA) derivatives and associated with IDILI synergized with TNFα in causing cytotoxicity in HepG2 cells, and IFNγ enhanced this interaction. NSAIDs that are propionic acid (PA) derivatives and cause IDILI that is of less clinical concern also synergized with TNFα, but IFNγ was without effect. Caspase inhibition prevented cytotoxicity from AA and PA derivative/cytokine treatment. Treatment with a representative AA or PA derivative induced activation of the MAPKs c-Jun N-terminal kinase (JNK), extracellular signal-regulated kinase (ERK), and p38. Inhibition of either JNK or ERK reduced cytotoxicity from cytokine interactions with AA derivatives. In contrast, an ERK inhibitor potentiated cytotoxicity from cytokine interactions with PA derivatives. An AA derivative but not a PA derivative enhanced IFNγ-mediated activation of STAT-1, and this enhancement was ERK-dependent. These findings raise the possibility that some IDILI reactions result from drug/cytokine synergy involving caspases and MAPKs and suggest that, even for drugs within the same pharmacologic class, synergy with cytokines occurs by different kinase signaling mechanisms.

Key words: NSAID; cytokine; idiosyncratic drug-induced liver injury; JNK; ERK

Drug-induced liver injury (DILI) is the leading cause of acute liver failure in the USA and remains the most common adverse effect associated with failure to obtain U.S. Food and Drug Administration approval for new drugs (Aithal *et al.*, 2011). Idiosyncratic DILI (IDILI), a subset of DILI, occurs in a small fraction of patients taking a drug and can result in severe liver injury or death. These reactions have resulted in removal of many drugs from the market that are efficacious and safe in the majority of patients.

Mechanisms underlying IDILI remain unproven, and the reactions are not predicted by typical preclinical toxicity testing. The infrequency of most IDILI responses suggests that individual susceptibility as well as characteristics of the offending drug are needed to elicit a response. A longstanding hypothesis is that IDILI-associated drugs activate a damaging adaptive immune response (Uetrecht, 1999). Specific human leukocyte antigen polymorphisms are associated with liver injury induced by some drugs, suggesting an important role for adaptive immune responses in the pathogenesis of IDILI (Tujios and Fontana, 2011). Another hypothesis suggests that activation of the innate immune system during drug therapy can render an individual susceptible to injury from an otherwise nontoxic dose of the

© The Author 2015. Published by Oxford University Press on behalf of the Society of Toxicology. All rights reserved. For Permissions, please e-mail: journals.permissions@oup.com drug (Roth and Ganey, 2011). Importantly, inflammatory cytokines are expressed and mediate critical events in both adaptive and innate immune responses. Indeed, in several rodent models of IDILI based on interaction of drugs with an immune response, the cytokines tumor necrosis factor-alpha ($TNF\alpha$) and interferon-gamma ($IFN\gamma$) proved to be critical to the pathogenesis of hepatocellular injury (Dugan *et al.*, 2011; Shaw *et al.*, 2009a, b; Zou *et al.*, 2009).

Recently published studies suggest that toxic cytokine/drug synergy can be recapitulated in vitro. For example, some drugs associated with IDILI synergize with $TNF\alpha$ to kill hepatocytes in vitro, and a role for aberrant mitogen-activated protein kinase (MAPK) signaling has been implicated in this response (Beggs et al., 2014; Cosgrove et al., 2009, 2010; Fredriksson et al., 2011). TNFa is known to activate the MAPKs c-Jun N-terminal kinase (JNK) and p38 transiently (Wullaert et al., 2007). MAPKs are commonly activated in response to cellular stress, and if their activation is prolonged cell death can ensue (Anderson, 1997). IFNy activates the Janus kinase-signal transducer and activator of transcription (JAK-STAT) pathway, which can also mediate cell death (Stephanou and Latchman, 2003). Exactly how drugs associated with IDILI synergize with cytokines to cause hepatocellular damage remains incompletely understood, although it is likely that activation of caspases and MAPKs plays a role (Beggs et al., 2014; Fredriksson et al., 2011).

Nonsteroidal anti-inflammatory drugs (NSAIDs) and antibiotics are the most frequent causes of IDILI. The observation that rheumatoid arthritis increases the risk of NSAID-induced liver injury more than 10-fold in human patients suggests that immune mediators contribute to IDILI pathogenesis from drugs in this pharmacologic class (Garcia Rodriguez et al., 1994). This suggestion is supported by results in animal models. In one such model, rodents treated with diclofenac (DCLF) in combination with lipopolysaccharide (LPS), an activator of the innate immune system, developed hepatocellular injury which did not occur after treatment with either DCLF or LPS alone (Deng et al., 2006). In addition, DCLF potentiated LPS-mediated expression of TNF α and IFN γ genes in rats (Ramm and Mally, 2013). Similarly, sulindac (SLD), another NSAID associated with IDILI, produced $TNF\alpha$ -dependent liver injury in rats cotreated with LPS (Zou et al., 2009).

Nearly all NSAIDs have been implicated in causing IDILI; however, the severity and lesion morphology of NSAID-induced hepatotoxicity varies substantially, likely due at least in part to the diversity of chemical structures within this drug class (O'Connor *et al.*, 2003; Teoh and Farrell., 2003). In this study, we tested the hypothesis that NSAIDs with idiosyncrasy liability synergize with TNF α and IFN γ to cause hepatocellular toxicity in vitro. To gain insight into the mechanism of the NSAID/cytokine-induced cytotoxic interaction, the involvement of caspases and MAPKs was examined.

MATERIALS AND METHODS

Materials. NSAIDs and MAPK inhibitors were purchased from Sigma-Aldrich (St. Louis, Missouri). Z-VAD-FMK and recombinant human TNF α and IFN γ were purchased from R&D Systems (Minneapolis, Minnesota). Phosphate-buffered saline (PBS), Dulbecco's Modified Eagles Medium (DMEM), Williams' Medium E, L-glutamine, fetal bovine serum (FBS), Antibiotic-Antimycotic (ABAM), and 0.25% Trypsin-EDTA were purchased from Life Technologies (Carlsbad, California). All antibodies were

purchased from Cell Signaling Technology (Beverly, Massachusetts).

Animals. Male C57Bl/6J mice purchased from Jackson Laboratory (Bar Harbor, Maine) were allowed to acclimate for at least 1 week upon arrival. They were housed in a 12-hour light/dark cycle, were fed a standard chow (8640 Teklad 22/5 Rodent Diet, Harlan Laboratories, Madison, WI, USA), and had continual access to bottled spring water. All procedures were conducted with the approval of the Michigan State University Institutional Animal Care and Use Committee.

Cell culture. Human hepatoma HepG2 cells (American Type Culture Collection, Manassas, Virginia) were chosen because they are insensitive to the harmful effects of cytokines yet express both $TNF\alpha$ and $IFN\gamma$ receptors (Hershey and Schreiber, 1989; Stonans et al., 1999). Moreover, HepG2 cells respond similarly to primary mouse hepatocytes and primary human hepatocytes with regard to the cytotoxic interaction between IDILIassociated drugs and cytokines (Beggs et al., 2014; Cosgrove et al., 2009). It is known that HepG2 cells have low expression of phase 1 drug metabolizing enzymes compared with primary human hepatocytes. However, compared with primary human hepatocytes, they have similar expression of phase 2 drug metabolizing enzymes (Westerink and Schoonen, 2007a, b). The only NSAID used in this study for which it is suspected that phase 1 metabolism is required for liver injury is SLD. In this study, the active metabolite of SLD, SLD sulfide, was used. We previously demonstrated that SLD sulfide, but not the parent compound, synergized with cytokines to cause cytotoxicity in HepG2 cells and primary rat hepatocytes (Zou et al., 2009). With regard to the remaining NSAIDs used in this study, there is not convincing evidence that bioactivation is required for liver injury in humans. Cells were grown in 25-cm² tissue culturetreated flasks and maintained in DMEM supplemented with 10% FBS and 1% ABAM. They were cultured at 37°C in 95% air and 5% CO₂ in a humidified incubator. They were passaged when they reached \sim 80% confluence.

Primary murine hepatocytes were isolated as described previously by Klaunig *et al.* (1981). Hepatocytes were isolated using a 2-step collagenase perfusion method. Viability of isolated hepatocytes was assessed by trypan blue exclusion. Only cells with >85% viability were used for experiments. Hepatocytes were plated with Williams' Medium E supplemented with 10% FBS, 1% ABAM, 2 mM L-glutamine, and 100 nM insulin. After plating, they were cultured at 37°C in 95% air and 5% CO₂ in a humidified incubator and allowed 3 h to attach prior to treatment.

IDILI classification. NSAIDs were classified according to their ability to cause IDILI (Table 1). The criteria used to classify the drugs in this study was established by Xu *et al.* (2008) and takes into consideration postmarketing label information as well as numbers of published clinical case reports.

Cytotoxicity assessment. HepG2 cells were plated at a density of 4×10^4 cells per well in black-walled, 96-well tissue culture plates and were allowed to attach overnight before treatment with compounds. DCLF, bromfenac (BRM), ibuprofen (IBU), and naproxen (NAP) were reconstituted in sterile water. SLD sulfide and aspirin were reconstituted in DMSO. Cells were treated with various concentrations of each NSAID or its vehicle, and simultaneously with TNF α (10 ng/ml) and/or IFN γ (10 ng/ml) or their vehicles (PBS). Cells were treated with NSAID concentrations ranging from 0 to 100 times the maximal plasma concentration

NSAID	Structural Sub-class	IDILI Potential	C_{\max} (µg/ml)	Molecular Weight (g/mol)	C _{max} Reference
DCLF	AA derivative	Yes	2.4	296.15	Xu et al. (2008)
SLD sulfide	AA derivative	Yes	1.6	340.41	Reid et al. (2008)
BRM	AA derivative	Yes	4.8	356.15	Gumbhir-Shah et al. (1997)
IBU	PA derivative	Yes	32.9	206.29	Bramlage and Goldis (2008)
NAP	PA derivative	Yes	75.9	252.23	Setiawati et al. (2009)
Aspirin	Salicylic acid derivative	No	7.6	180.16	Brandon et al. (1986)

Table 1. NSAID Subclass and Maximal Plasma Concentration (C_{max}) From Therapeutic Doses in Human Patients

 (C_{max}) observed in human patients. The C_{max} value for each NSAID is presented in Table 1. One hundred times the C_{max} was considered a pharmacologically relevant dosing limit for this in vitro study based on scaling factors described in Xu et al. (2008). Preliminary concentration-response studies were conducted with each cytokine to find a concentration that produced a robust cytotoxic interaction in the presence of DCLF (Supplementary Figs. 1A and 1B). The cytokine concentrations chosen for this study are within 10-fold of the concentrations found in serum of human patients undergoing an inflammatory response (Pinsky et al., 1993; Taudorf et al., 2007). Cells were exposed to drug/cytokine combinations for 24 h, and cytotoxicity was evaluated by measuring lactate dehydrogenase (LDH) release from the cells into culture medium using the Homogeneous Membrane Integrity Assay kit from Promega (Madison, Wisconsin). For drugs that interfered with the fluorescence-based assay (IBU and NAP), a spectrophotometric method was used to measure LDH release (Vanderlinde, 1985).

To investigate the roles of caspases and the MAPKs, pharmacological inhibitors of these pathways were used (40 μ M Z-VAD-FMK for caspases, 20 μ M SP600125 for JNK, 20 μ M U0126 for extracellular signal-regulated kinase (ERK), and 20 μ M SB203580 for p38). Inhibitors were reconstituted in DMSO, resulting in a maximal final concentration of 0.4% DMSO in experiments involving SLD sulfide/Z-VAD-FMK or 0.2% DMSO in all other experiments. In brief, cells were treated with an inhibitor alone or simultaneously in combination with TNF α and/or IFN γ and with one concentration of NSAID that produced strong cytotoxic synergy in the presence of cytokines. LDH release was measured 24 h after treatment.

Primary mouse hepatocytes were plated at a density of 1.25×10^5 cells per well in collagen-coated 24-well tissue culture-treated plates. Cells were allowed 3 h to attach followed by 2 washes with warm PBS then were treated with BRM alone or in combination with TNF α and/or IFN γ prepared in serum-free Williams' Medium E supplemented with 1% ABAM and 2 mM L-glutamine. After 24 h of exposure to drugs and/or cytokines, cell supernatant was collected, and attached cells were lysed with 1% triton-X. The supernatant and lysate were transferred to 96-well plates and analyzed for alanine aminotransferase activity as described by Luyendyk *et al.* (2005).

Caspase-3 activity. Caspase-3 activity was measured using the Caspase-3 Fluorometric Assay Kit purchased from R&D Systems. HepG2 cells were plated at 1.2×10^6 cells per well in 6-well tissue culture plates. Cells were treated with a NSAID alone or in combination with TNF α and/or IFN γ . Cells were lysed and centrifuged after 24 h of exposure. 50 μl of lysate were added to black-walled, 96-well plates and incubated with assay reaction buffer and fluorogenic substrate for 1 h. The plate was then read on a fluorescent plate reader at an excitation wavelength of 400 nm and an emission wavelength of 505 nm.

Protein isolation. Cells (1.2×10^6 per well) were plated in 6-well tissue culture plates and allowed to adhere overnight. They were exposed to one concentration of NSAID and its vehicle alone or in combination with TNF α and/or IFN γ for 12 or 18 h. Cells were rinsed with cold PBS followed by an addition of 150 μ l of radio-immunoprecipitation assay buffer containing HALT protease and phosphatase inhibitor cocktails (Thermo Scientific, Rockford, Illinois). Cells were scraped, collected, placed in microcentrifuge tubes, and incubated on ice for 10 min. During the 10-min incubation, the tubes were vortexed intermittently. Lysates were centrifuged for 25 min at 20000× g. The supernatants containing whole cell extracts were collected, placed in fresh, chilled tubes, and stored at -80° C until use. Protein concentrations were quantified using the bicinchoninic acid assay (Thermo Scientific).

Western analysis. For detection of MAPKs and phosphorylated STAT-1 (pSTAT-1) in whole cell lysates, protein (30 µg for JNK and $15\,\mu g$ for ERK, p38 and STAT-1) was loaded onto precast NuPAGE 12% Bis-Tris gels (Life Technologies) and subjected to electrophoresis. Proteins were transferred onto polyvinylidene fluoride membranes (Millipore, Billerica, Massachusetts). Membranes were blocked for 1 h with 5% bovine serum albumin (BSA) reconstituted in 1% tris-buffered saline (TBS) containing 0.1% tween-20 (TBSt) for detection of JNK, p38 and STAT-1 or blocked with TBS-based LI-COR blocking buffer (Lincoln, Nebraska) for detection of ERK. Membranes were probed with antibodies directed against phosphorylated JNK (pJNK), total JNK, phosphorylated ERK (pERK), total ERK, phosphorylated p38 (pp38), pSTAT-1 (Tyrosine 701), pSTAT-1 (Serine 727), and α -Tubulin. Primary antibodies were diluted in appropriate buffers to 1:1000. Membranes were incubated with primary antibodies overnight at 4°C, after which they were washed with TBSt followed by an addition of secondary antibodies. Goat anti-rabbit horseradish peroxidase (HRP)-conjugated secondary antibody was diluted in 5% BSA in TBSt at a concentration of 1:2500 for pJNK, 1:5000 for total JNK, 1:5000 for pp38, 1:5000 for total p38, 1:5000 for pSTAT-1 (Tyrosine 701), 1:5000 for pSTAT-1 (Serine 727), and 1:5000 for α -Tubulin. Clarity Western ECL substrate (Bio-Rad, Hercules, California) was used to visualize HRP and the substrate was developed on HyBlot CL film (Denville Scientific, Metuchen, New Jersey). For the detection of ERK, donkey anti-mouse or goat anti-rabbit infrared (IR) dye-conjugated secondary antibodies were diluted in LI-COR blocking buffer to 1:3000, and IR fluorescence was detected using the LI-COR Odyssey IR Imaging System. All images were quantified by performing densitometry using image J software.

Statistical analysis. All results are expressed as mean \pm standard error of the mean (SEM). Data were subjected to log transformation as necessary to achieve normality and equal variance. Data were analyzed via a 1-way or 2-way analysis of variance (ANOVA). Tukey's post-hoc test was used to perform multiple,



FIG. 1. Interaction of NSAIDs with cytokines leads to synergistic cytotoxicity. HepG2 cells were treated with NSAIDs that have (A, B) or do not have (C) IDILI liability alone or in combination with $TNF\alpha$ (10 ng/ml) and/or IFN γ (10 ng/ml). The NSAIDs that have IDILI liability are grouped according to structural subclass: (A) AA derivatives; (B) PA derivatives; (C) salicylic acid derivative. a, significantly different from VEH (no cytokine) within an NSAID concentration; b, significantly different from TNF α within an NSAID concentration; c, significantly different from IFN γ within an NSAID concentration; d, significantly different from γ within a cytokine group. Data are represented as mean ± S.E.M of at least 5 separate experiments. VEH, vehicle; TNF, tumor necrosis factor-alpha; IFN, interferon-gamma; LDH, lactate dehydrogenase.

pairwise comparisons between treatment groups. The criterion for significance was set at $\alpha = 0.05$.

RESULTS

NSAID/Cytokine-Induced Cytotoxicity: Concentration-Response The NSAIDs chosen for this study are DCLF, BRM, SLD sulfide (the active metabolite of SLD), NAP, IBU, and aspirin. Aspirin is the only one of these drugs that has not been associated with IDILI. Within the NSAID class of pharmaceuticals, there are a variety of subclasses based on chemical structure. There are 3 NSAID subclasses represented among the drugs used in our study. DCLF, BRM, and SLD sulfide are acetic acid (AA) derivatives, IBU and NAP are propionic acid (PA) derivatives, and aspirin is a salicylic acid derivative. It is worth noting that AA derivatives and PA derivatives are among the most hepatotoxic NSAIDs (Teoh and Farrell, 2003). Moreover, among the



FIG. 2. Caspase activation in response to DCLF/cytokine and IBU/cytokine treatment. HepG2 cells were treated with (A) a representative AA derivative (DCLF: 250 μ M) or (B) a representative PA derivative (IBU: 6 mM) alone or in combination with TNF α and/or IFN γ , and cell lysates were collected 24h after treatment for measurement of caspase 3 activity. a, significantly different from VEH; b, significantly different from TNF α ; c, significantly different from IFN γ ; d, significantly different from control. Data are represented as mean \pm S.E.M of at least 3 separate experiments.VEH, vehicle; TNF, tumor necrosis factor-alpha; IFN, interferon-gamma; DCLF, diclofenac; IBU, ibuprofen.

IDILI-associated drugs used in this study, the AA derivatives are of greater clinical concern than the PA derivatives (Unzueta and Vargas 2013).

As expected, treatment of cells with TNF α and/or IFN γ in the absence of drug did not result in release of LDH (Fig. 1). With the exception of SLD sulfide, treatment with NSAID alone did not result in cytotoxicity. All 5 IDILI-associated NSAIDs synergized with TNF α in a concentration-dependent manner to cause cytotoxicity (Figs. 1A and 1B). IFN γ by itself did not influence drug-induced cytotoxicity; however, in the presence of TNF α it enhanced the cytotoxicity mediated by DCLF, BRM, and SLD sulfide (Fig. 1A) but had no effect on the toxicity of IBU and NAP (Fig. 1B). Aspirin did not synergize with TNF α or IFN γ alone or in combination to kill HepG2 cells (Fig. 1C).

Some of the NSAIDs used in this study, in addition to dozens of other IDILI-associated drugs, synergize with cytokines to cause death of primary human hepatocytes (Cosgrove *et al.*, 2009). Drug/cytokine-induced cytotoxic synergy observed in primary human hepatocytes was recapitulated in HepG2 cells in spite of the low phase 1 metabolism observed in this cell line (Cosgrove *et al.*, 2009). The cytotoxic interaction observed here in HepG2 cells between BRM and TNF α , as well as the IFN γ -mediated enhancement of BRM/TNF α -induced cytotoxicity

(Fig. 1A), was observed in primary mouse hepatocytes as well (Supplementary Fig. 2).

Cytotoxic Synergy Between Cytokines and NSAIDs Requires Caspases

Fredriksson *et al.* (2011) reported that DCLF/TNFα-mediated cytotoxicity in HepG2 cells depends on caspases. In addition, Zou *et al.* (2009) demonstrated that SLD sulfide synergizes with TNFα to cause caspase activation that led to cell death. We tested the hypothesis that this holds true for other NSAIDs and for the IFN_γ-mediated enhancement of NSAID/TNFα-induced cytotoxicity.

Of the 3 NSAID subclasses used in this study, 2 of them (the AA derivatives and PA derivatives) interacted with cytokines to kill cells. Although these 2 subclasses differed in the manner in which they synergized with the cytokines, within subclass they responded similarly to each other. Consequently, we selected a representative AA derivative (DCLF) and PA derivative (IBU) to evaluate caspase involvement in the NSAID/cytokine-induced cytotoxic interaction. For this experiment and all subsequent ones, we selected an NSAID concentration that resulted in strong cytotoxic synergy with $TNF\alpha$ and/or IFN $\gamma.$ Both DCLF and IBU induced caspase 3 activation within 24h of treatment (Fig. 2). Addition of TNFa potentiated caspase 3 activation by the drugs. Consistent with the cytotoxicity data presented in Figure 1, IFN γ enhanced caspase 3 activation induced by DCLF/TNFa but not by IBU/TNFa (Fig. 2). Treatment with the pan-caspase inhibitor Z-VAD-FMK completely protected cells from NSAID/TNFα-induced cytotoxicity (Fig. 3). In addition, the IFN_γ-mediated enhancement of cytotoxicity in cells treated with AA derivatives/TNFa was eliminated by Z-VAD-FMK treatment.

Cytotoxic Synergy Between Cytokines and NSAIDs Involves Activation of JNK

As described earlier, a representative AA derivative (DCLF) and PA derivative (IBU) were selected to examine the expression of phosphorylated (activated) JNK (pJNK). Based on the timecourse of cytotoxicity after treatment with DCLF and cytokines (Supplementary Fig. 3), 2 times were selected: a time at which there was no cytotoxicity (12 h) and a time at which cytotoxicity was observed (18h). Treatment with TNFa in the absence of drug (Control) caused expression of pJNK at 12 h (Fig. 4A). In the absence of drug, expression of pJNK was unchanged by treatment with TNF α , IFN γ , or TNF α /IFN γ at 18 h (Figs. 4A and 4B). At 12 h, treatment with DCLF caused phosphorylation of JNK which was enhanced in the presence of TNF α . Treatment with IFN γ did not alter DCLF/TNFα-induced JNK phosphorylation. By 18h, DCLF caused JNK activation in the absence of TNF α . TNF α increased this response, but $\text{IFN}\gamma$ did not alter JNK activation either alone or in the presence of DCLF/TNFa. In contrast, IBU significantly activated JNK by itself at both 12 and 18 h (Fig. 4B). IBU-mediated JNK activation was enhanced in the presence of TNF α at 12 h, whereas IFN γ was without effect.

We next examined the involvement of JNK in NSAID/cytokine-induced cytotoxicity. Treatment with SP600125, an inhibitor of JNK activation, significantly reduced cytotoxicity mediated by cytokines in combination with NSAIDs containing an AA moiety (Fig. 5A). In contrast, SP600125 treatment did not alter the cytotoxic interaction for NSAIDs containing a PA moiety (Fig. 5B). Interestingly, treatment with SP600125 eliminated VEH- and DCLF-induced JNK activation but was ineffective at eliminating IBU-mediated JNK activation (Fig. 5C, Supplementary Fig. 4A).



FIG. 3. Caspases are involved in the NSAID/cytokine-induced cytotoxic interaction. HepG2 cells were treated with (A) AA derivatives (DCLF: 250μ M, BRM: 750μ M or SLD sulfide: 200μ M), or (B) PA derivatives (IBU: 6μ M or NAP: 10μ M) alone or in combination with TNF α and/or IFN γ . Cells were also incubated in the presence and absence of the pan-caspase inhibitor ZVAD-FMK (40μ M). Cytotoxicity was measured 24 h later. a, significantly different from VEH within NSAID/inhibitor treatment; b, significantly different from TNF α within NSAID/inhibitor treatment; c, significantly different from Control within a cytokine group; d, significantly different from NSAID without inhibitor within a cytokine group. Data are represented as mean \pm S.E.M of at least 5 separate experiments. VEH, vehicle; TNF, tumor necrosis factor-alpha; IFN, interferon-gamma; LDH, lactate dehydrogenase; DCLF, diclofenac; BRM, bromfenac; SLD, sulindac; IBU, ibuprofen; NAP, naproxen.

Cytotoxic Synergy Between Cytokines and NSAIDs Involves Activation of ERK

Treatment with cytokines in the absence of drug (Control) did not result in ERK activation (Fig. 6). Treatment with DCLF caused activation of ERK at 12 h that was still evident at 18 h (Fig. 6A). Neither TNF α nor IFN γ alone or in combination affected DCLFmediated activation of ERK. Similarly, treatment with IBU caused persistent ERK activation that was unaltered by the presence of cytokines (Fig. 6B). U0126 prevents ERK phosphorylation by inhibiting the MAPK kinase (MEK) that directly phosphorylates ERK. Treatment with U0126 did not affect cytotoxicity mediated by AA derivatives in combination with TNF α ; however, it significantly reduced the IFN γ -mediated enhancement of AA derivative/TNF α -mediated cytotoxicity (Fig. 7A). In contrast, treatment with U0126 potentiated cytotoxicity caused by PA derivatives in combination with cytokines (Fig. 7B). The MEK inhibitor U0126 was effective at eliminating ERK



FIG. 4. HepG2 cells were treated with (A) a representative AA derivative (DCLF: 250μ M) or (B) a representative PA derivative (IBU: 6 mM) alone or in combination with TNF α and/or IFN γ , and protein extracts were collected 12 or 18 hours after treatment. p-JNK and total JNK protein were detected via western analysis. Representive blots are shown. Densitometry was performed using image J software. a, significantly different from VEH within an NSAID treatment. b, significantly different from IFN γ within an NSAID treatment. c, significantly different from Control within a cytokine treatment. d, significantly different from TNF α within an NSAID treatment. Data are represented as mean \pm S.E.M of 3 separate experiments. Abbreviations: VEH, vehicle; TNF, tumor necrosis factor-alpha; IFN, interferon gamma; TI, tumor necrosis factor-alpha and interferon-gamma; DCLF, diclofenac; IBU, ibuprofen.

activation induced by DCLF and IBU (Fig. 7C, Supplementary Fig. 4B).

p38 Attenuates NSAID/Cytokine-Induced Cytotoxic Synergy

Treatment with either TNF α alone or DCLF alone induced phosphorylation of p38 at 12 h but not at 18 h (Fig. 8A). There was no change in DCLF-induced p38 activation in the presence of TNF α at 12 h, and treatment with DCLF/TNF α /IFN γ increased p38

phosphorylation relative to DCLF/IFN γ or TNF α /IFN γ treatment (Fig. 8A). IBU strongly induced p38 activation relative to VEH at 12 and 18 h (Fig. 8B). Cytokine treatment did not significantly alter IBU-mediated p38 activation (Fig. 8B).

Activation of p38 is typically associated with activation of pathways leading to cell death. Surprisingly, with the exception of DCLF/TNF α exposure, treatment with the p38 inhibitor SB203580 enhanced cytotoxicity mediated by AA derivative/



FIG. 5. JNK is involved in the NSAID/cytokine-induced cytotoxic interaction. HepG2 cells were treated with (A) AA derivatives (DCLF: 250μ M, BRM: 750μ M or SLD sulfide: 200 μ M), or (B) PA derivatives (IBU: 6 mM or NAP: 10 mM) alone or in combination with TNF α and/or IFN γ . Cells were also incubated in the presence and absence of the JNK inhibitor SP600125 (20 μ M). Cytotoxicity was measured 24 h later. (C) Cells were treated with a representative AA derivative (DCLF: 250μ M) or a representative PA derivative (IBU: 6 mM) in the presence or absence of SP60125 for 12 h, and p-JNK and total JNK protein were detected via western analysis. a, significantly different from VEH within NSAID/inhibitor treatment; b, significantly different from TNF α within NSAID/inhibitor treatment; c, significantly different from Control within a cytokine group. Data are represented as mean ± S.E.M of at least 5 separate experiments. VEH, vehicle; TNF, tumor necrosis factor-alpha; IFN, interferon-gamma; LDH, lactate dehydrogenase; DCLF, diclofenac; BRM, bromfenac; SLD, sulindac; IBU, ibuprofen; NAP, naproxen; SP, SP600125.

TNF α exposure in the presence and absence of IFN γ (Fig. 9A). Treatment with SB203580 potentiated cytotoxicity from PA derivative/TNF α exposure as well, irrespective of IFN γ exposure (Fig. 9B). These data suggest that p38 plays a protective role in NSAID/cytokine-induced cytotoxicity.

DCLF But Not IBU Promotes Dual Phosphorylation of STAT-1 in an ERK-Dependent Manner

Upon binding to its receptor, IFN γ is well known to activate the JAK-STAT pathway. As expected, treatment with IFN γ resulted in phosphorylation of STAT-1 at tyrosine (Tyr) 701, irrespective of TNF α exposure (Fig. 10). DCLF treatment did not influence Tyr 701 phosphorylation. It has been reported that phosphorylation of STAT-1 at serine (Ser) 727 in addition to Tyr 701 is required for maximal activation (Varinou *et al.*, 2003). There was a modest increase in phosphorylation of STAT-1 at Ser 727 in response to IFN γ treatment, irrespective of TNF α treatment. Interestingly, treatment with DCLF markedly enhanced the IFN γ -mediated

phosphorylation at Ser 727 (Fig. 10). JAK is responsible for phosphorylating Tyr 701 on STAT-1. It is unclear which specific kinases are responsible for phosphorylating Ser 727, but it has been suggested that MAPKs, specifically ERK, can perform this phosphorylation (Zhang *et al.*, 2004). Treatment with the ERK inhibitor U0126 did not alter phosphorylation of Tyr 701 but completely prevented phosphorylation of Ser 727 (Fig. 10). Treatment with IBU prevented IFN γ -mediated phosphorylation of STAT-1 at both Tyr 701 and Ser 727 (Fig. 11).

DISCUSSION

In this study, we showed that NSAIDs associated with IDILI synergize with TNF α to kill HepG2 cells in vitro and that treatment with an additional cytokine, IFN γ , enhanced the NSAID/TNF α -induced cytotoxic interaction. These results are consistent with what has been reported previously in studies involving animal models of drug/inflammatory stress-induced liver injury



FIG. 6. DCLF and IBU treatment induce prolonged activation of ERK. HepG2 cells were treated with (A) a representative AA derivative (DCLF: 250μ M) or (B) a representative PA derivative (IBU: 6 mM) alone or in combination with TNF α and/or IFN γ , and protein extracts were collected 12 or 18 h after treatment. p-ERK and total ERK protein were detected via western analysis. Representative blots are shown. Densitometry was performed using image J software. c, significantly different from Control within a cytokine treatment. Data are represented as mean \pm S.E.M of 3 separate experiments. VEH, vehicle; TNF, tumor necrosis factor-alpha, IFN, interferon-gamma; TI, tumor necrosis factor-alpha and interferon-gamma; DCLF, diclofenac; IBU, ibuprofen.

(Dugan et al., 2011; Shaw et al., 2009a, b; Zou et al., 2009). The IFN γ -mediated enhancement of NSAID/TNF α -induced cytotoxicity was observed with the AA derivatives (DCLF, BRM, and SLD sulfide) but not with the PA derivatives (IBU or NAP), suggesting that this is a phenomenon related to chemical structure.

To gain insight into the mechanism underlying NSAID/cytokine-induced cytotoxic synergy, the roles of caspases and MAPKs were examined. Caspase enzymes are crucial to the initiation of apoptosis (Porter and Jänicke, 1999), and results of previous studies suggested a role for caspases in drug/ cytokine-induced cytotoxic synergy (Beggs *et al.*, 2014; Fredriksson et al., 2011; Zou et al., 2009). Treatment with a representative AA derivative or PA derivative resulted in activation of caspase 3, which was increased further in the presence of TNFα (Fig. 2). Treatment with IFNγ enhanced caspase 3 activation in the presence of DCLF/TNFα but not IBU/TNFα (Fig. 2), consistent with its effect on cytotoxicity (Fig. 1). Treatment with the caspase inhibitor Z-VAD-FMK eliminated the NSAID/TNFα-mediated cytotoxic interaction and also eliminated the IFNγ-mediated enhancement of NSAID/TNFα cytotoxicity (Fig. 3). These results suggest that caspase-mediated apoptosis is the mode of cell death in NSAID/cytokine cytotoxic synergy.



FIG. 7. ERK is involved in the NSAID/cytokine-induced cytotoxic interaction. HepG2 cells were treated with (A) AA derivatives (DCLF: 250 μ M, BRM: 750 μ M or SLD sulfide: 200 μ M), or (B) PA derivatives (IBU: 6 mM or NAP: 10 mM) alone or in combination with TNF α and/or IFN γ . NSAID/cytokine combinations were also incubated in the presence and absence of the ERK inhibitor U0126 (20 μ M). Cytotoxicity was measured 24 h later. (C) Cells were treated with a representative AA derivative (DCLF: 250 μ M) or a representative PA derivative (IBU: 6 mM) in the presence or absence of 100126 for 12 h and p-ERK and total ERK protein was detected via western analysis. a, significantly different from TNF α within NSAID/inhibitor treatment; c, significantly different from TNF α within a cytokine group; d, significantly different from NSAID without inhibitor within a cytokine group; d, significantly different from SAID without inhibitor within a cytokine group; d, significantly different from SAID without inhibitor within a cytokine group; d, significantly different from SAID without inhibitor within a cytokine group; d, significantly different from SAID without inhibitor within a cytokine group; d, significantly different from SAID without inhibitor within a cytokine group. Data are represented as mean ± S.E.M of at least 5 separate experiments. VEH, vehicle; TNF, tumor necrosis factor-alpha; IFN, interferon-gamma; LDH, lactate dehydrogenase; DCLF, diclofenac; BRM, bromfenac; SLD, sulindac; IBU, ibuprofen; NAP, naproxen; U0, U0126.

Prolonged activation of JNK is associated with signaling through pathways leading to cell death (Wullaert *et al.*, 2007), and JNK activation contributes to cytotoxicity mediated by TNF α in combination with trovafloxacin, another IDILI-associated drug, and, as mentioned previously, also by DCLF/TNF α (Beggs *et al.*, 2014; Fredriksson *et al.*, 2011). In this study, DCLF and IBU caused similar patterns of JNK activation (Fig. 4). That is, both drugs caused persistent JNK activation that was enhanced in the presence of TNF α but unaffected by IFN γ exposure. A JNK inhibitor eliminated the activation of JNK and completely prevented the cytotoxic synergy caused by DCLF/TNF α in the absence and presence of IFN γ (Fig. 5). This is consistent with IFN γ interacting somehow with the cell death pathway initiated by TNF α . Inhibition of JNK similarly prevented cytotoxicity from other AA derivative NSAID/cytokine combinations (Fig. 5A).

In contrast to the AA derivatives, JNK inhibition did not affect cytotoxicity mediated by PA derivative/cytokine combinations (Fig. 5B). At first glance, these results suggest that JNK plays a role in cytotoxic synergy mediated by AA derivatives but not PA derivatives. However, treatment with the JNK inhibitor did not eliminate phosphorylation of JNK induced by IBU treatment, which might explain the lack of effect on cytotoxicity (Fig. 5C). Moreover, since SP600125 acts as a reversible, ATPcompetitive inhibitor (Bennett *et al.*, 2001), it is possible that IBU physically interacts with SP600125 or with JNK, thereby preventing SP600125 from completely inhibiting ATP binding to JNK. Treatments with larger concentrations of SP600125 were



FIG. 8. Treatment with TNFα, DCLF or IBU induces activation of p38. HepG2 cells were treated with (A) a representative AA derivative (DCLF: 250 μM) or (B) a representative PA derivative (IBU: 6 mM) alone or in combination with TNFα and/or IFNγ, and protein extracts were collected 12 or 18 h after treatment. pp38 and total p38 protein were detected via western analysis. Representative blots are shown. Densitometry was performed using image J software. a, significantly different from VEH; b, significantly different from IFNγ; c, significantly different from Control. Data are represented as mean ± S.E.M of 3 separate experiments. VEH, vehicle; TNF, tumor necrosis factor-alpha, IFN, interferon-gamma; TI, tumor necrosis factor-alpha and interferon-gamma; DCLF, diclofenac; IBU, ibuprofen.

attempted but were unsuccessful due to solubility limitations at concentrations ${>}30\,\mu\text{M}.$

ERK phosphorylation is typically associated with activating cell survival signaling pathways; however, it has become clear that under some conditions, ERK activates cell death pathways (Cagnol and Chambard, 2010). The duration of ERK activation can be an important factor in determining cellular fate. It has been suggested that prolonged ERK signaling can lead to cell death (Cagnol and Chambard, 2010). The involvement of ERK in

the cytotoxic synergy mediated by IDILI-associated drugs in combination with cytokines has not been reported previously. The representative AA and PA derivatives induced very similar patterns of ERK activation (Fig. 6), and these were not affected by cytokine treatment. U0126 treatment effectively inhibited both DCLF- and IBU-mediated ERK activation (Fig. 7C) but had opposite effects on the synergistic cytotoxicity caused by AA and PA derivatives. U0126 eliminated the IFN γ -mediated enhancement of AA derivative/TNF α -induced cytotoxicity,



FIG. 9. P38 plays a protective role in NSAID/cytokine-induced cytotoxicity. HepG2 cells were treated with (A) AA derivatives (DCLF: 250 μ M, BRM: 750 μ M or SLD sulfide: 200 μ M), or (B) PA derivatives (IBU: 6 mM or NAP: 10 mM) alone or in combination with TNF α and/or IFN γ . NSAID/cytokine combinations were also incubated in the presence and absence of the p38 inhibitor SB203580 (20 μ M). Cytotoxicity was measured 24 h later. a, significantly different from VEH within NSAID/inhibitor treatment; b, significantly different from TNF α within NSAID/inhibitor treatment; c, significantly different from Control within a cytokine group; d, significantly different from NSAID without inhibitor within a cytokine group. Data are represented as mean ± S.E.M of at least 4 separate experiments. VEH, vehicle; TNF, tumor necrosis factor-alpha; IFN, interferon-gamma; LDH, lactate dehydrogenase; DCLF, diclofenac; BRM, bromfenac; SLD, sulindac; IBU, ibuprofen; NAP, naproxen.

suggesting that the cytotoxic effect of IFN γ required ERK activity. In contrast, U0126 treatment potentiated cytotoxicity mediated by the PA derivative/cytokine combinations (Fig. 7B). These results suggest that ERK signaling plays a protective role in this case. Together, the findings indicate that persistent ERK activation can promote either cell survival or cell death, depending on the particular NSAID involved in its activation.

The involvement of p38 in drug/cytokine-induced cytotoxic synergy has not been reported. TNF α transiently activates p38 in a variety of cell types (Anderson, 1997), as was seen here in HepG2 cells (Fig. 8). DCLF also caused a transient activation of p38 that was modestly affected by the addition of cytokines (Fig. 8A). In contrast, IBU caused activation of p38 that was longer lived but unaffected by cytokine treatment (Fig. 8B). The observation that inhibition of p38 enhanced the NSAID/ cytokine-induced cytotoxic interaction (Fig. 9) suggests that p38

dampens this toxic response. Although activation of p38 is commonly associated with activation of cell death pathways (Anderson, 1997), it can promote cell survival under certain conditions. For example, transient activation of p38 by $TNF\alpha$ is essential to mediating signals that protect cells from apoptosis (Roulston *et al.* 1998).

As mentioned earlier, AA and PA derivatives responded differently in terms of their interaction with cytokines to kill cells. Although both subclasses interacted with TNF α to cause cytotoxicity, IFN γ enhanced the synergy from TNF α and AA derivatives but not PA derivatives. This observation raises the questions of how AA derivatives sensitize cells to the harmful effects of IFN γ and why are cells treated with PA derivatives not sensitive to IFN γ . To answer these questions, we evaluated the phosphorylation status of STAT-1, a critical component of the IFN γ signaling pathway. The IFN γ receptor is a heterodimer-associated



FIG. 10. DCLF promotes ERK-dependent phosphorylation of STAT-1 in the presence of IFN γ . HepG2 cells were treated with (A) a representative AA derivative (DCLF: 250 μ M) alone or in combination with TNF α and/or IFN γ and incubated in the presence or absence of U0126. Protein extracts were collected 18 h after treatment. pSTAT-1 (Tyrosine 701), pSTAT-1 (Serine 727) and α -Tubulin protein levels were detected via western analysis. Representative blots are shown. (B) Densitometry was performed using image J software. a, significantly different from VEH; b, significantly different from TNF α ; c, significantly different from Control; d, significantly different from DCLF (without inhibitor). Data are represented as mean ± S.E.M of 3 separate experiments. VEH, vehicle; TNF, tumor necrosis factor-alpha, IFN, interferon-gamma; TI, tumor necrosis factor-alpha and interferon-gamma; DCLF, diclofenac; Tyr, tyrosine; Ser, serine.

intracellularly with JAK. When bound to IFNy, the receptor becomes activated, leading to activation of JAK which phosphorylates STAT-1 at Tyr 701. Upon phosphorylation, STAT-1 dimerizes and translocates to the nucleus, where it binds to specific DNA sequences (Farrar and Schreiber, 1993). Phosphorylation at Ser 727 is required for maximal STAT-1 activation (Varinou et al., 2003). The kinases responsible for phosphorylation at Ser 727 include MAPKs, specifically ERK (Li et al., 2010). IFNγ caused pronounced phosphorylation of STAT-1 at Tyr 701 but had only a modest effect at Ser 727 (Figs. 10 and 11). Conversely, DCLF was without effect on Tyr 701 phosphorylation but in the presence of IFN γ caused a pronounced increase in phosphorylation at Ser 727 which depended on ERK (Fig. 10). These findings might explain why inhibition of ERK prevented the IFN_γ-mediated enhancement of DCLF/TNFα-induced cytotoxicity. Interestingly, phosphorylation of STAT-1 at Tyr 701 in response to IFN γ treatment was necessary for robust DCLF-induced phosphorylation of Ser 727. Consistent with this observation, in several cell types phosphorylation at Tyr 701 by JAK was required for Ser 727 phosphorylation (Sadzak et al., 2008).

In stark contrast to DCLF, treatment with IBU prevented IFN γ -mediated phosphorylation of STAT-1 at both Tyr 701 and Ser 727 (Fig. 11). These results are consistent with the observation that IFN γ failed to enhance cytotoxicity mediated by PA derivatives in combination with TNF α . Given that phosphorylation of Ser 727 was dependent on ERK, it is puzzling that both

DCLF and IBU treatment induced the same pattern of ERK activation, yet only DCLF led to phosphorylation of Ser 727. Our findings suggest that treatment with DCLF and IFN γ unmasks a substrate for ERK at Ser 727 of STAT-1, which is not available in cells treated with IBU. In addition, the observation that inhibition of ERK increased cytotoxicity from PA derivatives/TNF α treatment suggests that ERK activated by IBU treatment interacts with a cytoprotective substrate rather than one that leads to enhanced cytotoxicity.

In summary, NSAIDs associated with IDILI synergize with TNF α to cause death of HepG2 cells. IFN γ treatment enhances the cytotoxicity mediated by some NSAIDs in the presence of TNF α . Aspirin, an NSAID that is not associated with IDILI, did not synergize with any combination of cytokines to kill cells. These findings raise the possibility that drug/cytokine cytotoxic synergy contributes to human IDILI from NSAIDs.

With regard to mechanism, NSAID/cytokine-induced cytotoxicity requires caspases, suggesting an apoptotic mode of cell death. Persistent JNK activation plays an important role in the cytotoxic synergy. Prolonged ERK activation plays either a cytotoxic or a protective role, depending on NSAID chemical structure, whereas p38 is cytoprotective. Cosgrove *et al.* (2010) evaluated the signaling pathways involved in drug/cytokineinduced cytotoxic synergy in primary human hepatocytes. They found that various drugs associated with IDILI (including NSAIDs) synergized with cytokines to cause MAPK signaling



FIG. 11. IBU treatment prevents IFN γ -mediated phosphorylation of STAT-1. HepG2 cells were treated with (A) a representative PA derivative (IBU: 6 mM) alone or in combination with TNF α and/or IFN γ . Protein extracts were collected 18 h after treatment. pSTAT 1 (Tyrosine 701), pSTAT-1 (Serine 727), and α -Tubulin protein levels, were detected via western analysis. Representative blots are shown. (B) Densitometry was performed using image J software. a, significantly different from VEH; b, significantly different from TNF α ; c, significantly different from Control. Data are represented as mean \pm S.E.M of 3 separate experiments. VEH, vehicle; TNF, tumor necrosis factor-alpha, IFN, interferon-gamma; TI, tumor necrosis factor-alpha and interferon-gamma; IBU, ibuprofen; Tyr, tyrosine; Ser, serine.

dysregulation and consequently death of primary human hepatocytes, which lends support to our findings concerning involvement of MAPKs in NSAID/cytokine-induced cytotoxic synergy in HepG2 cells. NSAIDs from different structural classes differentially modify the phosphorylation status of STAT-1, and this appears to explain why IFN γ potentiates the cytotoxic interaction with TNF for some NSAIDs but not others.

These findings suggest that cytotoxic synergy of drugs with cytokines occurs through different kinase signaling mechanisms, even for drugs within the same pharmacologic class, and that these differences are related to chemical structure and IDILI liability. Knowledge generated from this study could be useful in developing an in vitro approach to classify drugs according to their potential to cause IDILI in humans.

SUPPLEMENTARY DATA

Supplementary data are available online at http://toxsci. oxfordjournals.org/.

FUNDING

National Institutes of Health (Grant R01 DK087886) and a Colgate Palmolive Award for Research in Alternative Methods.

ACKNOWLEDGMENTS

We thank Jonathan Turkus for technical assistance.

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