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Acetaminophen - Dose-dependent Drug Hepatotoxicity and Acute Liver Failure in Patients

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Abstract

Background—Drug-induced liver injury (DILI) is a rare but serious clinical problem. A number of drugs can cause severe liver injury and acute liver failure (ALF) at therapeutic doses in a very limited number of patients (<1:10,000). This idiosyncratic DILI (iDILI), which is currently not predictable in preclinical safety studies, appears to depend on individual susceptibility and the inability to adapt to the cellular stress caused by a particular drug. In striking contrast to iDILI, drugs with dose-dependent hepatotoxicity are mostly detected during preclinical studies and do not reach the market. One notable exception is acetaminophen (APAP, paracetamol), which is a safe drug at therapeutic doses but can cause severe liver injury and ALF after intentional and unintentional overdoses.

Key Messages—APAP overdose is responsible for more ALF cases in the US or UK than all other etiologies combined. Because APAP overdose in the mouse represents a model for the human pathophysiology, substantial progress has been made during the last decade in understanding the mechanisms of cell death, liver injury and recovery. More recently, emerging evidence based on mechanistic biomarker analysis in patients and studies of cell death in human hepatocytes suggests that most of the mechanisms discovered in mice also apply to patients. The rapid development of N-acetylcysteine as antidote against APAP overdose was based on the early understanding of APAP toxicity in mice. However, despite the efficacy of N-acetylcysteine in patients who present early after APAP overdose, there is a need to develop intervention strategies for late presenting patients.

Conclusions—The challenges related to APAP toxicity are to better understand the mechanisms of cell death in order to limit liver injury and prevent ALF and to develop biomarkers that better predict as early as possible who is at risk for developing acute liver failure with poor outcome.

Keywords

Acetaminophen hepatotoxicity; acute liver failure; biomarker; innate immune response; mitochondrial dysfunction

Disclosure Statement

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INTRODUCTION

Drug hepatotoxicity, although a relatively rare event, is the leading cause of acute liver failure in the clinic accounting for more than 60% of all cases in the US [1]. A general classification considers drug toxicity either idiosyncratic or dose-dependent. Idiosyncratic drug-induced liver injury (iDILI) is one of the biggest concerns with development of new drugs as neither predictable animal models nor in vitro tests are available and the limited number of patients in clinical trials does generally not allow to detect the delayed, mostly immune-based toxicity that occurs in less than 1 in 10,000 patients [2]. On the other hand, a considerable effort is being made to detect dose-dependent drug toxicity in preclinical testing and during clinical trials. Because of the use of numerous in vitro and in vivo approaches, dose-dependent direct toxicity is reliably detectable and generally, dosedependent hepatotoxins are eliminated early in the drug development process. However, one of the most used pain medication worldwide, acetaminophen (APAP; paracetamol), is a classical dose-dependent hepatotoxin that is responsible for almost 50% of all acute liver failure cases in the US, the UK and many Western countries [1]. Therefore, this review will focus on the recent progress made in the understanding of the pathophysiology of APAP hepatotoxicity in patients and the development of biomarkers that can predict whether a patient will spontaneously recover or will need a liver transplant to survive.

MECHANISMS OF APAP HEPATOTOXICTY

Clinical Aspects of APAP Overdose

APAP is a safe drug at therapeutic doses of 4 g per day for an adult. Extensive literature reviews suggest that even susceptible people such as alcoholics are unlikely to suffer adverse effects from therapeutic doses of APAP [3]. However, an overdose can cause severe liver injury and even acute liver failure [4,5]. Suicide attempts are a frequent cause of exposure to a single, high overdose of APAP but unintentional, cumulative overdosing is an increasing problem. APAP is not only in specific pain medications but is also present in sleeping aids, cold medicine and numerous other over-the-counter drugs. Thus, patients taking multiple drugs may inadvertently exceed recommended doses of APAP. Unintentional overdosing has frequently a poorer outcome because these patients generally present later than patients who attempted suicide [4].

The only clinically approved antidote against APAP-induced liver injury is N-acetylcysteine (NAC), which is most effective when administered within 8 h of the overdose [4]. Although NAC is still beneficial even after 24 h, the efficacy is substantially diminished [6]. NAC is a precursor for GSH synthesis and acts through facilitating scavenging of the reactive metabolite NAPQI during the metabolism phase [7]. At later time points, GSH assists in scavenging reactive oxygen in mitochondria and excess NAC is converted to Krebs cycle intermediates and supports mitochondrial energy metabolism [8]. Delayed treatment with NAC increases the risk for acute liver failure. However, in contrast to idiosyncratic drug toxicity, APAP-induced liver injury and liver failure has a relatively high survival rate. Nevertheless, due to the wide-spread use of the drug, APAP overdose is by far the most frequent cause of acute liver failure of any etiology in the US and the UK [1]. Thus, the major challenges related to APAP overdose are: 1. Improve the understanding of the

Mechanisms of APAP-induced liver injury in preclinical studies

Shortly after the first report on APAP hepatotoxicity in humans, a mouse model was developed that showed a similar degree of liver injury as observed in patients [9]. The early experiments demonstrated the formation of a reactive metabolite, later suggested to be *N*-acetyl-*p*benzoquinone imine (NAPQI), which can be scavenged by glutathione (GSH) [9]. However, the excess formation of NAPQI after an overdose causes reactions with protein sulfhydryl groups (protein adduct formation) and toxicity [9]. This early insight into the mechanism led to the development of NAC.

Subsequent studies demonstrated that APAP protein adducts formation *per se* do not directly cause cell death because no critical targets could be identified [10]. Instead, protein adducts, especially in mitochondria, seem to disturb the electron transport chain and generate an oxidant stress and peroxynitrite in mitochondria [11], which eventually triggers the membrane permeability transition (MPT) pore opening and collapse of the membrane potential [12] (**Figure 1**). In addition, formation of a Bax pore and later mitochondrial matrix swelling leads to release of intermembrane proteins, especially endonuclease G and apoptosis inducing factor (AIF), both of which translocate to the nucleus and cause extensive DNA fragmentation [10,11]. The combination of massive mitochondrial dysfunction and nuclear disintegration leads to cellular necrosis [13].

More recently, it was recognized that the initial oxidant stress was insufficient to trigger the MPT. Instead, an amplification loop is necessary that involves the activation of various redox-sensitive MAP kinases including apoptosis signal regulating kinase 1 (ASK1) and mixed lineage kinase 3 (MLK3), which ultimately causes the activation of c-jun N-terminal kinase 1/2 (JNK) and translocation of p-JNK to mitochondria [14] (Figure 1). Upon interaction of p-JNK with an anchor protein in the outer mitochondrial membrane [14], the mitochondrial oxidant stress is amplified [15]. Together with lysosomal iron, which enters mitochondria through the Ca^{2+} uniporter, the oxidant stress triggers the MPT [16]. Thus, in the mouse model, the JNK loop is critical for the pathophysiology. Recently, the importance of additional kinases such as receptor interacting protein kinase 1 (RIP1) and RIP3 has been shown in APAP hepatotoxicity [17, 18]. Whether these kinases act through JNK or other pathways remains unclear. RIP3 has been shown to promote mitochondrial fission and oxidant stress [17]. Ultimately, all signaling pathways lead to cell necrosis (necroptosis) [13, 17]. Apoptosis is not a relevant cell death mechanism of APAP-induced liver injury. There is no evidence for caspase activation and highly potent caspase inhibitors have no effect on the injury [13, 19, 20].

In addition to the pro-cell death signaling pathways, adaptive defense mechanisms also affect the pathophysiology. Autophagy is a process where damaged organelles are being enveloped with lipid membranes (autophagosomes), which then fuse with lysosomes for degradation [21]. APAP overdose activates autophagy of damaged mitochondria

(mitophagy), which can limit liver injury [22] (**Figure 1**). Mitophagy can affect cell death around the outer area of necrosis where the insult is not too overwhelming [23]. Beyond autophagy, the unfolded protein response (ER stress) can be activated [24] but the pathophysiological relevance remains controversial. Furthermore, adaptive responses can be beneficial such as induction of heat shock proteins [25], metallothionein [26] or antioxidant enzymes, e.g., glutamate cysteine ligase (GCL), which is the rate-limiting enzyme for hepatic GSH synthesis. In fact, the lower susceptibility of female mice to APAP overdose is thought to be mainly caused by the more robust induction of GCL and faster recovery of GSH levels [27]. Overall, for all experiments a number of pro-cell death pathways and adaptive defense mechanisms need to be considered.

Mechanisms of APAP-induced liver injury in human liver

Although most of the known mechanisms of APAP-induced liver injury were discovered in a rodent model, the important question is how relevant are these mechanisms for patients. Similar to mouse hepatocytes, the initial response of primary human hepatocytes or metabolically competent hepatoma cells (HepaRG cells) to high APAP levels is rapid GSH depletion and protein adduct formation, including mitochondrial adducts [28, 29]. In contrast to the original assumption that GSH levels need to be first depleted by 70% before adducts are being formed [9], it is now evident that APAP protein adducts form immediately [28, 30]. In fact, even after therapeutic doses, which cause only very minor, temporary GSH loss, protein adducts can be detected in the liver and in plasma in mice [30] and humans [31]. A difference between rodents and humans appears to be a higher overall protein adduct generation in human hepatocytes and a more delayed formation of mitochondrial adducts [28] compared to rodents [30, 32]. Oxidant stress and mitochondrial dysfunction, detected as breakdown of the mitochondrial membrane potential, is followed by cell contents release or propidium iodide uptake indicating necrotic cell death [28, 29]. APAP also caused JNK activation in the cytosol of primary human hepatocytes by 6 h; p-JNK translocation to mitochondria was observed by 15 h [28]. In contrast, JNK activation and mitochondrial p-JNK translocation occurs in mice within 1-2 h [14, 32]. However, APAP did not cause JNK activation in HepaRG cells [28] despite that mitochondrial dysfunction and oxidant stress are central to the mechanism of cell death in this cell type [29]. A JNK inhibitor only partially reduced cell death in primary human hepatocytes suggesting that human cells are less dependent on the JNK amplification loop than mouse hepatocytes [28].

In support of the findings with primary cells, evidence for mitochondrial dysfunction was also observed in APAP overdose patients where release of biomarkers for mitochondrial damage such as mitochondrial DNA (mtDNA), the mitochondrial matrix enzyme glutamate dehydrogenase (GDH) and the indirect mitochondrial damage biomarker nuclear DNA fragments (nDNA) could be measured in plasma [5] (**Figure 2**). The fact that these biomarkers are not released after furosemide overdose in mice despite the release of ALT and development of extensive necrosis suggests these biomarkers are specific for mitochondrial damage and not just cell death [5]. Patients with more mitochondrial damage are less likely to survive, demonstrating that mitochondrial dysfunction is central to the mechanisms of APAP hepatotoxicity in humans [33].

Mechanistic and predictive biomarkers of APAP hepatotoxicity and acute liver failure

In addition to the above mentioned mitochondria damage biomarkers mtDNA, GDH and nDNA, acylcarnitines were found to be elevated after APAP overdose in mice [34, 35]. Carnitine is used to shuttle long chain fatty acids into mitochondria for degradation. Thus, an increase in plasma acylcarnitines, even in the absence of ALT release, reflects mitochondrial dysfunction in hepatocytes [35]. However, no significant increase in plasma acylcarnitine levels was observed in human APAP overdose patients with injury [35]. The reason seems to be at least in part that all patients are treated very early with NAC, which can suppress acylcarnitine release [35]. The likely explanation for this effect is that any NAC not used for GSH synthesis is degraded to form Krebs cycle intermediates, which support mitochondrial energetics [8]. Other biomarkers such as miR-122 [36], full-length cytokeratin-18 [37], high mobility group box-1 (HMGB1) protein [37], and argininosuccinate synthetase [38] have all been shown to be highly sensitive biomarker for cell damage and necrosis in both animals and APAP overdose patients. These biomarkers are superior to ALT as indicator of cell injury, i.e. these biomarkers can be detected earlier and, due to shorter half-life in plasma, are better indicators of acute cell injury [39]. In addition, a panel of miRNAs was shown to be specific for APAP-induced cell death compared to liver injury caused by ischemic hepatitis [40]. In contrast to ALT, most of these biomarkers allow to predict whether a patient is likely to survive or will develop ALF and may need a transplant to survive [33, 37, 41]. A limitation of virtually all biomarkers evaluated so far is that although within a large group the biomarker can predict outcome, the substantial overlap in the individual values between groups makes it difficult to rely on the absolute value of any single biomarker [39]. However, several biomarkers could be combined as part of a panel of biomarkers to predict patient outcome as was used for a mitochondrial damage biomarker [33].

Innate immune response and APAP hepatotoxicity in rodents and humans

Necrotic cell death as observed after APAP overdose in mice and humans causes an innate immune response by release of damage associated molecular patterns (DAMPs) such as HMGB1, mtDNA, nDNA fragments, etc. [42, 43] (Figure 2). DAMPs bind to various Tolllike receptors, especially on macrophages, and trigger pro-inflammatory cytokine and chemokine formation. Interleukin-1 β (IL-1 β) received special attention because its formation is not only dependent on the transcriptional activation of the gene but also on a posttranslational cleavage by caspase-1 [43]. For its activation, caspase-1 requires the assistance of several scaffolding proteins of the nod-like receptor family, pyrin domain containing-3 (NLRP3) inflammasome. The purinergic receptor P2X7, when stimulated by DAMPs like ATP, can activate the NLRP3 inflammasome [43]. Although there is clear evidence for inflammasome activation and IL-1 β formation during APAP overdose in mice [20, 44], the pathophysiological relevance of this mediator and the inflammasome is controversial [20, 45]. The main reason why IL-1 β and the inflammasome is likely of very limited importance for the pathophysiology is the fact that the amount of IL-1 β being formed is orders of magnitude below levels that would be required to activate immune cells [20]. In addition, mice which lack the IL-1 receptor or any component of the NLRP3 inflammasome are not protected against APAP hepatotoxicity [20, 45]. Furthermore, caspase-1 gene

knockout mice or pancaspase inhibitors, which effectively prevented the processing of pro-IL-1 β to the active cytokine, did not affect APAP toxicity [20, 45]. Because IL-1 receptors cannot trigger cell death directly, the main mechanism by which IL-1 β could affect APAPinduced liver injury would be to activate inflammatory cells such as neutrophils. Although neutrophils are recruited during the injury phase of APAP hepatotoxicity, a large number of different interventions against neutrophils failed to reduce liver injury [reviewed in 42]. The conclusions of a few studies that suggested an aggravation of APAP-induced liver injury by neutrophils have to be questioned due to off-target effects [42]. In fact, some of the most convincing evidence against a role of neutrophils is the observation that circulating and liver-derived neutrophils are not activated during the initial phase of liver injury [46]. In addition, mice deficient in NADPH oxidase, the enzyme responsible for reactive oxygen formation by phagocytes, are not protected against APAP toxicity [42, 47]. Thus, neither neutrophils nor the resident macrophages of the liver (Kupffer cells) contribute to the oxidant stress or injury induced by APAP overdose [42, 47]. In contrast, Kupffer cells actually attenuate the injury through formation of IL-10 [48]. IL-10 limits the generation of pro-inflammatory cytokines, which may aggravate liver injury by inducing the inducible form of nitric oxide synthase and promoting the mitochondrial oxidant stress [48]. On the other hand, recruitment of neutrophils and macrophages into the area of necrosis is essential for removing cellular debris and promoting regeneration [49]. Taken together, the release of DAMPs by necrotic cell triggers an innate immune response with late activation and hepatic recruitment of neutrophils and macrophages (Figure 2). The majority of the experimental evidence supports the hypothesis that the inflammatory cells are essential for regeneration rather than contribute to the injury in mice [42].

An innate immune response is also observed in humans during APAP-induced liver injury and acute liver failure [47, 50]. However, there is little evidence for neutrophil activation before and during the peak of liver injury [47]. In contrast, during periods of declining serum ALT levels, i.e. after the active cell death is over, neutrophil activation as indicated by priming for reactive oxygen formation and the enhanced capacity for phagocytosis, could be observed [47]. As all these patients recovered, the enhanced activation of neutrophils correlated with the regeneration phase [47]. These data support the hypothesis that similar to the mouse model, neutrophils contribute to the recovery after APAP hepatotoxicity in humans. In addition, hepatic macrophages in livers of APAP-induced liver injury patients are derived from resident macrophages and from circulating monocytes [50]. The expression of the chemokine receptor CCR2 on these cells and the formation of CCL2 indicate that similar to mice [49], the chemokine CCL2 is responsible for hepatic monocyte recruitment in patients [50]. However, the environment in the liver in the presence of hepatic macrophages is characterized by anti-inflammatory cytokine formation and by expression of proliferation markers [50]. Together these findings support the hypothesis that hepatic macrophages contribute to the regeneration of the damaged tissue in patients [50] (Figure 2).

SUMMARY AND CONCLUSIONS

Acetaminophen (paracetamol) overdose is the most frequent cause of drug-induced liver injury and acute liver failure in Western countries. The mechanisms, which apply to the

mouse model as well as to the human pathophysiology, include the formation of a reactive metabolite, which depletes glutathione and triggers mitochondrial dysfunction with oxidant stress, JNK activation, nuclear DNA fragmentation and the MPT resulting in cellular necrosis. The clinical antidote NAC was developed based on these mechanisms. Although NAC is highly effective when given within 8 h of the overdose, the efficacy is diminished when treatment is delayed. Further studies of the later phase of the injury and the transition to regeneration are needed to identify therapeutic targets that would benefit late presenting patients. In addition, an increasing number of biomarkers are being identified which help in understanding the mechanisms of liver injury in patients but also will eventually allow prediction at the time of admission of whether a patient is going to recover or will need a liver transplant to survive.

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Jaeschke

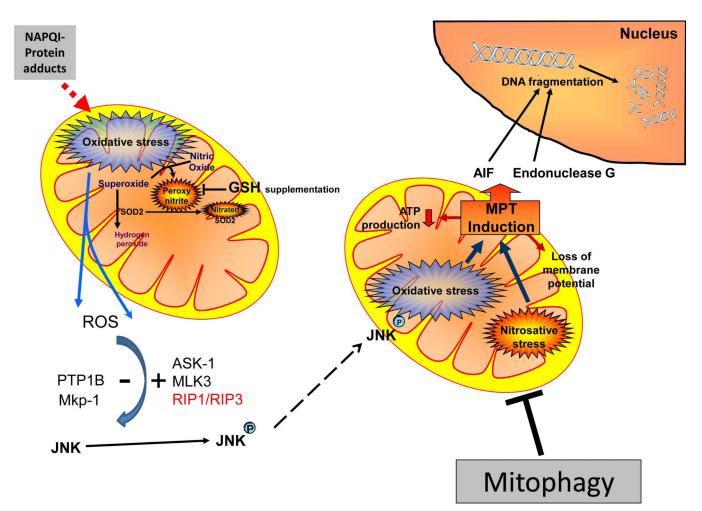


Figure 1. Mechanisms of Acetaminophen-induced Necrotic Cell Death

The P450-dependent metabolism of acetaminophen results in formation of the reactive intermediate N-acetyl-p-benzoquinone imine (NAPQI), which forms mitochondrial protein adducts and induces a mitochondrial oxidative stress. Superoxide derived from the mitochondrial electron transport chain reacts with nitric oxide resulting in production of peroxynitrite. Superoxide can be metabolized by superoxide dismutase 2 (SOD2) and converted to hydrogen peroxide, though peroxynitrite can inactivate SOD2 by nitration. The initial mitochondrial oxidative stress can also activate the MAP kinase c-jun-N-terminal kinase (JNK) by multiple pathways (redox-sensitive kinases), resulting in its phosphorylation and translocation to the mitochondria. Mitochondrial JNK translocation amplifies the mitochondrial oxidant stress, which subsequently leads to activation of the mitochondrial permeability transition (MPT) pore opening, translocation of mitochondrial proteins such as apoptosis inducing factor (AIF) and endonuclease G to the nucleus. This then results in DNA fragmentation and finally oncotic necrosis. Mitophagy can counteract the process by removing damaged mitochondria. Abbreviations: ASK-1, apoptosis signal regulating kinase-1; Mkp-1, mitogen-activated protein kinase (MAPK) phosphatase (Mkp)-1; MLK-3, mixed-lineage kinase 3; PTP1B, protein tyrosine phosphatase 1B; RIP,

receptor interacting protein kinase. (Adapted from Jaeschke et al., Drug Metab Rev 2012;44:88-106).

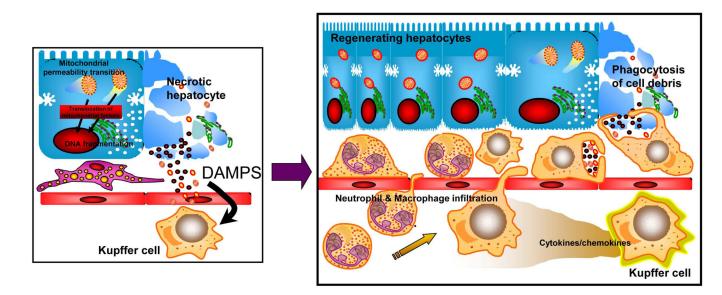


Figure 2. Acetaminophen-induced Sterile Inflammatory Response Promotes Regeneration

During acetaminophen-induced necrotic cell death a number of cellular components are being released including nuclear DNA fragments, formyl peptides and HMGB1, which can act as damage associated molecular patterns (DAMPs). DAMPs activate resident liver macrophages (Kupffer cells) via toll-like receptors. In the regeneration and repair phase, cytokine and chemokine formation by activated Kupffer cell results in activation, hepatic recruitment and transmigration of neutrophils and macrophages into the damaged tissue to facilitate removal of dead cells and activation of regenerative pathways. (Adapted from Jaeschke et al., Liver Int 2012;32:8-20).