



DAMPs and sterile inflammation in drug hepatotoxicity

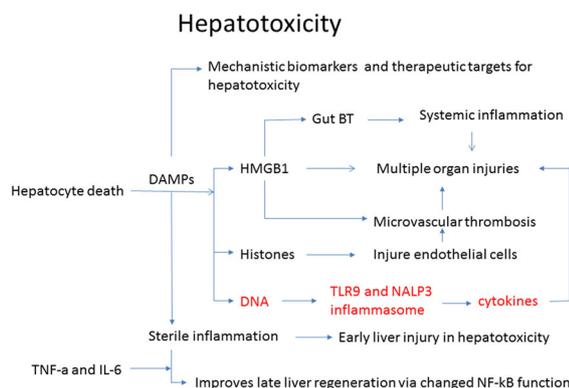
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Abstract

Drug hepatotoxicity is the leading cause of acute liver failure (ALF) in the developed countries. The early diagnosis and treatment are still problematic, and one important reason is the lack of reliable mechanistic biomarkers and therapeutic targets; therefore, searching for new biomarkers and therapeutic targets is urgent. Drug hepatotoxicity induces severe liver cells damage and death. Dead and damaged cells release endogenous damage-associated molecular patterns (DAMPs). Increased circulating levels of DAMPs (HMGB1, histones and DNA) can reflect the severity of drug hepatotoxicity. Elevated plasma HMGB1 concentrations can serve as early and sensitive mechanistic biomarker for clinical acetaminophen hepatotoxicity. DAMPs significantly contribute to liver injury and inhibiting the release of DAMPs ameliorates experimental hepatotoxicity. In addition, HMGB1 mediates 80% of gut bacterial translocation (BT) during acetaminophen toxicity. Gut BT triggers systemic inflammation, leading to multiple organ injury and mortality. Moreover, DAMPs can trigger and extend sterile inflammation, which contributes to early phase liver injury but improves liver regeneration at the late phase of acetaminophen overdose, because anti-inflammatory treatment reduces liver injury at early phase but impairs liver regeneration at late phase of acetaminophen toxicity, whereas pro-inflammatory therapy improves late phase liver regeneration. DAMPs are promising mechanistic biomarkers and could also be the potential therapeutic targets for drug hepatotoxicity. DAMPs-triggered sterile inflammation contributes to liver injury at early phase but improves liver regeneration at later phase of acetaminophen hepatotoxicity; therefore, anti-inflammatory therapy would be beneficial at early phase but should be avoided at the late phase of acetaminophen overdose.

Graphical abstract



Keywords Hepatotoxicity · DAMPs · Sterile inflammation · Regeneration

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Abbreviations

ALF	Acute liver failure
MOF	Multiple organ failure
SIRS	Systemic inflammatory response syndrome
BT	Bacterial translocation
HMGB1	High mobility group box 1
NETs	Neutrophil extracellular traps
DAMP	Damage-associated molecular pattern
TLR	Toll-like receptor
APAP	Acetaminophen
MODS	Multiple organ dysfunction syndrome
LPS	Lipopolysaccharide

Introduction

Drug hepatotoxicity is the main cause of acute liver failure (ALF) in the west industrialized countries [1]. Acetaminophen hepatotoxicity is a global issue; it is responsible for nearly half of the ALF cases in the US and remains the leading cause of liver transplantation [1]. Drug hepatotoxicity induces severe liver cells damage and results in hepatocyte cell death and inflammation [2, 3]. Dead, dying and damaged cells release DAMPs [4, 5], leading to increased circulating DAMPs concentrations in drug-induced acute fatal liver injury [5]. Once rapidly released extracellularly, these DAMPs can activate the innate and adaptive immune system [6] and resultantly initiate and extend sterile inflammation [4], which also significantly contribute to hepatotoxicity [5]. Therefore, endogenous DAMPs, also known as alarmins, can signal cellular damage [6]. Elevated circulating levels of DAMPs in patients and animals with severe acute pancreatitis (SAP) positively correlate with disease severity [4]. Inhibiting DAMPs (for example, HMGB1, histones and DNA) release can significantly ameliorate experimental SAP and multiple organ injuries [4]. Similarly, the increased plasma DAMPs concentrations can accurately reflect the severity of liver injury and blockade of these alarmins can attenuate experimental drug hepatotoxicity [5]. These evidences indicate that DAMPs could be the promising mechanistic biomarkers and the potential therapeutic targets for drug hepatotoxicity. Currently, the early diagnosis of drug hepatotoxicity and the treatment are still problematic, one important reason is lack of reliable biomarkers and therapeutic targets. Anti-inflammatory therapy is clinically available; however, when and how to use the anti-inflammation treatment is still difficult in practice, because it is increasingly clear that inflammation contributes to liver injury at early phase but improves hepatocyte regeneration at the late phase of drug hepatotoxicity. In this manuscript,

we review the current understanding of DAMPs and DAMPs-initiated sterile inflammation in drug hepatotoxicity.

HMGB1 in drug hepatotoxicity

HMGB1 is a non-histone nuclear protein that binds DNA [7]. Under resting conditions, HMGB1 is localized in the nucleus, where HMGB1 is involved in chromosomal DNA repair and contributes to nucleosome mobility by promoting histone sliding along the DNA strand [7]. Intracellular HMGB1 also plays an important role in modulating gene expression [7]. Following cell activation, HMGB1 undergoes post-translational modifications (such as acetylation) and the modified HMGB1 can translocate to cytoplasm. Extracellularly HMGB1 acts as DAMPs to interact with receptors that include RAGE (receptor for advanced glycation endproducts) as well as TLR2 and TLR4; therefore, extracellular HMGB1 can activate the innate immune system and mediate a wide range of pathological responses including sterile inflammation [7]. Among the DAMPs molecules, HMGB1 is a typical alarmin [5, 7] and plays an important role in the pathogenesis of drug hepatotoxicity [5].

HMGB1 is a critical mediator of lethal systemic inflammation

Drug hepatotoxicity is featured with severe liver cell damage and death [2]. Dead and damaged cells can rapidly release HMGB1 [8–10], which can also be actively secreted by immunocompetent cells [10]. Extracellular HMGB1 can signal inflammation [9] and modulate inflammatory cascade [11, 12] in which HMGB1 can stimulate LPS-stimulated macrophages to secrete early inflammatory cytokines TNF- α and IL-6 [11, 12]. HMGB1 is a potent late inflammatory mediator [8]; this well-studied alarmin molecule plays a critical role in lethal systemic inflammation [13, 14] in which HMGB1 may damage vascular endothelial cells and cause microvascular thrombosis, and this detrimental effect can lead to multiple organ dysfunction [15]. In addition, HMGB1 has a longer therapeutic window as compared to those early inflammatory cytokines: serum TNF- α and IL-6 are elevated for 5 days after the onset of sepsis, whereas circulating HMGB1 levels are increased for at least 3 weeks [16]. Therefore, compared to early inflammatory cytokines, HMGB1 is potentially a better therapeutic target for treating drug hepatotoxicity and sepsis, because drug-induced acute fatal liver injury has a potential to develop sepsis [5, 17, 18].

Circulating HMGB1 level is increased in drug hepatotoxicity

The clinical presentation of acetaminophen hepatotoxicity may vary; this makes the early diagnosis more difficult. Therefore, it is necessary to find new mechanistic biomarkers. Since HMGB1 can be rapidly released by damaged/dead hepatocyte and extracellular HMGB1 can signal cellular damage [6], HMGB1 might be a good biomarker candidate for the early diagnosis of drug-induced liver injury, and this notion is supported by the following evidences. In multiple acetaminophen-induced hepatotoxicity animal models, the significantly elevated circulating HMGB1 concentrations can signal liver cells injury [6] and reflect the severity of drug hepatotoxicity [17, 18]. In an acetaminophen-induced hepatotoxicity clinical trial, the circulating HMGB1 concentration represents hepatocyte necrosis, and serum hyper-acetylated HMGB1 levels indicate pyroptosis and immune cell activation [19, 20]. The increased plasma HMGB1 levels are able to sensitively reveal early acetaminophen-induced acute liver injury at first presentation to hospital [19]. In addition, the markedly increased total and acetylated HMGB1 concentrations indicate worse prognosis in the same acetaminophen clinical trial [20]. Moreover, ALF patients also have high levels of circulating HMGB1, which are positively correlated with disease severity [21, 22]. These data indicate that HMGB1 is a promising mechanistic biomarker of clinical drug hepatotoxicity (Fig. 1).

HMGB1 significantly contributes to drug hepatotoxicity and other vital organ injuries

Exogenous HMGB1 injection to mice can induce liver injury in which the serum ALT level is even higher than alcoholic steatohepatitis [23]. In a lethal acetaminophen overdose animal model, HMGB1 markedly impairs hepatocyte regeneration and blockade of HMGB1 significantly improves the late phase liver structure recovery [17]. Neutralization of HMGB1 can also ameliorate the early phase liver injury in two different experimental hepatotoxicity models, which are induced by acetaminophen overdose and D-galactosamine/LPS [24, 25]. In addition, inhibition of HMGB1 cytoplasmic translocation in liver cells can reduce acute fatal liver injury induced by D-galactosamine/LPS [26]. These data indicate that HMGB1 significantly contributes to drug hepatotoxicity at both the early injury phase and the late regeneration phase.

Moreover, circulating HMGB1 concentration has been reported as a sensitive disease biomarker [27], which can be used to reliably predict the severity of severe acute

pancreatitis (SAP) in clinic [28, 29]. The elevated levels of circulating HMGB1 can accurately reflect organ dysfunction, infection and fatality in patients with SAP [28, 29]. Inhibition of HMGB1 release can reduce pancreatic injury in an animal model of SAP [30]. HMGB1 has also been proved to promote acute kidney injury in multiple animal models, which are induced by renal ischemia–reperfusion [31], sepsis [32] and SAP [33]; inhibition of HMGB1 can attenuate kidney injury in these animal models. HMGB1 also significantly contributes to acute lung injury (ALI) in multiple animal models, which are induced by hemorrhagic shock [34], hyperoxia [35] and SAP [36]; blockade of HMGB1 protects against lung injury in these models.

HMGB1 mediates gut BT that triggers systemic inflammation during acetaminophen hepatotoxicity

Gut mucosal injury and BT play an important role in the development of multiple organ dysfunction during drug-induced hepatotoxicity, because acetaminophen overdose can induce evident mucosal injury [18], which can facilitate the leakage of gut luminal bacteria/LPS into the systemic circulation to trigger systemic inflammatory response syndrome (SIRS) [37]. The latter can lead to multiple organ dysfunctions (MOD), which can culminate in multiple organ failure (MOF) and fatality [38]. Therefore, the digestive tract is considered as the origin of systemic inflammation [38]. The following evidences support the notion that HMGB1 plays an important role in gut BT in drug hepatotoxicity. Exogenous HMGB1 injection to mice can induce gut mucosal injury and evident intestinal BT [23]. In an acetaminophen-induced hepatotoxicity murine model and a severe acute pancreatitis animal model, serum HMGB1 levels are elevated and the circulating HMGB1 levels are correlated with the amount of gut BT [17, 18, 39]; neutralization of HMGB1 decreases 70–80% of gut BT in both models; however, gut mucosal injury is not significantly improved in these two models [18, 39]. These data indicate that gut bacteria need adhere to the damaged gut mucosa to cross the intestinal mucosal barrier; however, this step is necessary but insufficient to complete the complicated gut BT procedure [18, 39]; HMGB1 is one of the important factors that are also needed to mediate intestinal BT; gut bacteria cross the mucosal barrier likely via an active “transcellular” pathway instead of a passive “paracellular” pathway through the damaged tight junction between the mucosal epithelial cells. It is becoming increasingly clear that gut BT can induce systemic inflammation in critical illness such as severe acute pancreatitis and trauma [40, 41]. Systemic inflammation can lead to MOD and mortality [42]. Therefore, HMGB1

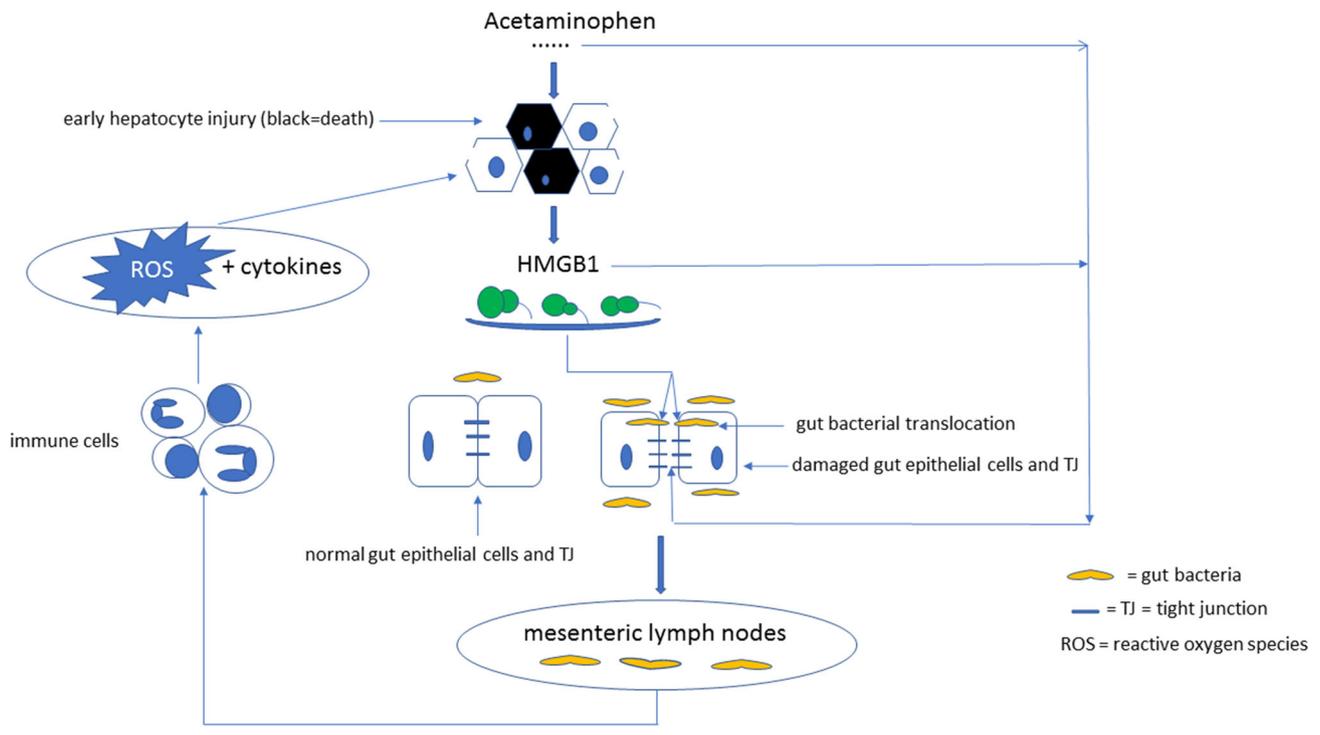
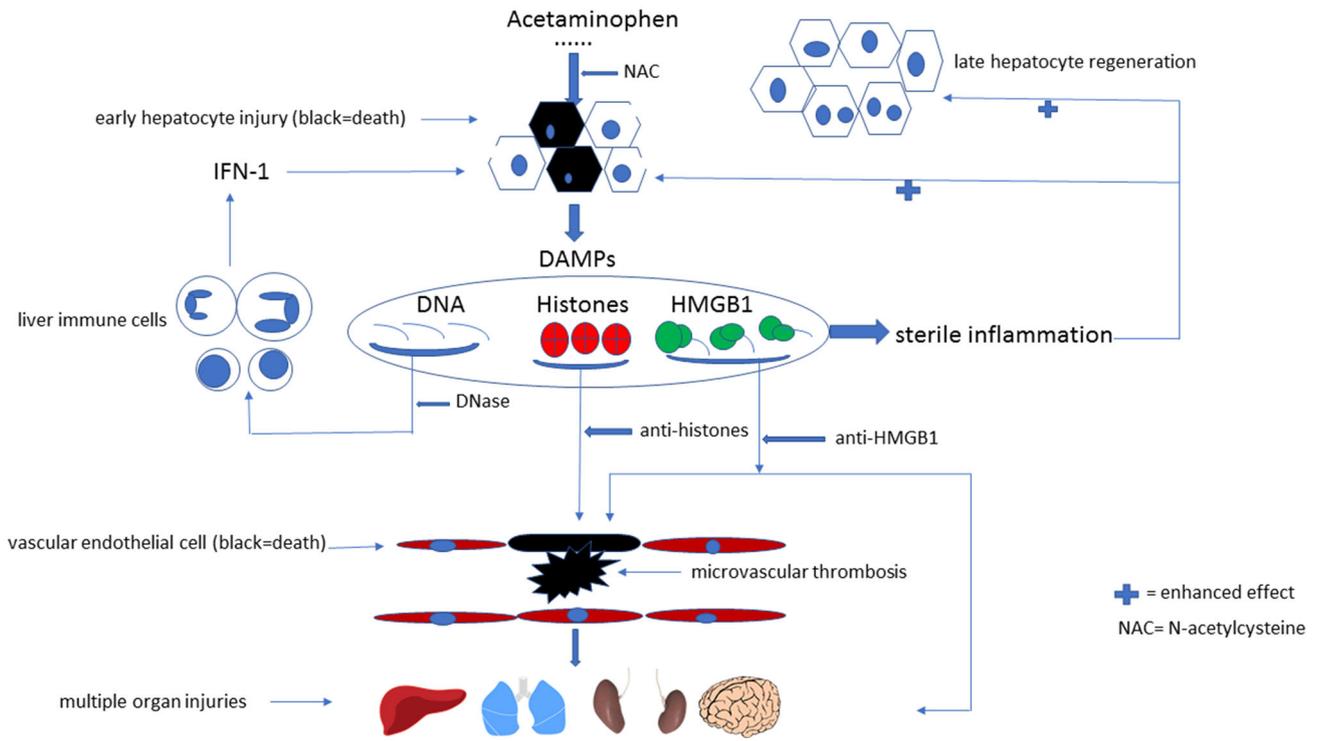


Fig. 1 In drug hepatotoxicity, the dying and dead hepatocyte releases damage-associated molecular patterns (DAMPs) such as HMGB1 and histones. HMGB1 significantly contributes to acetaminophen-induced hepatotoxicity and mediates gut bacterial translocation (BT), which induces systemic inflammation, leading to multiple organ dysfunction. Extracellular histones damage vascular endothelial cells and activate platelets to induce vascular thrombosis; this detrimental effect can lead to acute liver, lung, kidney and brain injuries. DNA released from dying hepatocytes is capable of activating liver immune cells via TLR9 and NALP3 inflammasome complexes resulting in cytokine production; the latter can amplify liver injury. Therefore, HMGB1, extracellular histones and DNA can signal cellular injury and are promising mechanistic biomarkers, and these molecules could also be the potential therapeutic targets for drug hepatotoxicity. In addition, DAMPs trigger sterile inflammation; the latter contributes to liver injury at early time point but improves liver regeneration at the late phase of drug hepatotoxicity. Pro-regenerative cytokines TNF- α and IL-6 can prime the quiescent hepatocyte to facilitate liver regeneration at the late time point of drug hepatotoxicity

appears to be the important factor that links BT and systemic inflammation in drug hepatotoxicity.

Extracellular histones in hepatotoxicity

Histones are important nuclear proteins that facilitate packaging of DNA into nucleosomes [43]. Upon cell injury or cell signaling process, histones can be passively or actively released [43–45]. Extracellular histones act as DAMPs to interact with receptors that include TLR2 as well as TLR4 to induce sterile inflammatory organ injury [43–45].

Circulating histones concentrations are increased in patients and animals with hepatotoxicity

The necrotic/damaged liver cells can passively release histones during drug hepatotoxicity, leading to elevated levels of circulating histones [44, 45]. The increased circulating histones levels can accurately reveal the severity of hepatotoxicity in two different animal models, which are induced by lethal doses of concanavalin A and acetaminophen [45]. Except the animal models, patients with acute-on-chronic liver failure and ALF are also reported to have elevated circulating histones concentrations, which can reflect disease severity and fatality [46, 47].

Extracellular histones significantly contribute to hepatotoxicity

Extracellular histones are cell toxic to host cells; therefore, the noxious extracellular histones can promote drug-induced liver injury by damaging tissue cells and killing

endothelial cells [43, 45]. In addition, extracellular histones can also elicit immuno-stimulatory effect to exaggerate multiple organ injuries [43–45, 48, 49]. In the following two different animal models, lethal doses of concanavalin A and acetaminophen are used to induce hepatotoxicity, and blockade of histones can attenuate acute fatal liver injury in both animal models, suggesting that extracellular histones are the major mediators of death in acute fatal liver injury [45]. In another animal model, the acute liver failure is induced by D-galactosamine/LPS, circulating histones can exacerbate systemic inflammation and neutralization of histones can ameliorate the acute liver failure in this model [50]. Except animal models, ALF patients also have increased serum histones levels, which can induce hepatocyte death and stimulate monocytes to release inflammatory mediators; these detrimental effects can be reversed by heparin treatment that can bind histones, suggesting that circulating histones are crucial mediators of systemic inflammation in ALF patients [46]. Histone H4 together with increased circulating NETs can activate platelets to facilitate the formation of microvascular thrombosis in sepsis [39, 45, 49, 51]. Extracellular histones can damage microvascular endothelial cells and induce the formation of microvascular thrombosis, these detrimental effects can lead to acute kidney [51, 52], lung [53] and brain injury in multiple animal models, and neutralization of histones can ameliorate these organ injuries and reduce the brain infarct size [54]. These data suggest that histones could be the potential novel biomarkers and therapeutic targets in drug-induced acute fatal liver injury.

Extracellular DNA contributes to acetaminophen-induced liver injury

After a drug overdose, hepatocytes may rupture and release a large amount of DNA, this can lead to the increased levels of circulating DNA [44, 45]. ALF patients also have increased plasma DNA levels that are closely associated with extracellular histones [44]. As part of the DAMPs, DNA together with histones/HMGB1 can stimulate the formation of neutrophil extracellular traps (NETs), and NETs can augment liver injury induced by sterile inflammation [48]. In addition, DNA together with histones/HMGB1 can mediate sterile inflammatory liver injury during drug hepatotoxicity, and DNase treatment can attenuate liver injury by reducing DNA levels [48]. Moreover, necrosis-derived DNA can stimulate liver immune cells (such as sinusoidal endothelial cells) via TLR9 and NALP3 inflammasome to release cytokines (such as IFN-1); the latter can amplify acetaminophen-induced liver necrosis [55, 56].

Inflammation contributes to liver injury at early phase but improves liver regeneration at late phase of acetaminophen hepatotoxicity

Drug hepatotoxicity results in cell death and sterile inflammation, and the latter also contributes to drug-induced liver injury [2].

Early inflammatory cytokines in drug hepatotoxicity

Early inflammatory mediators such as TNF- α , IL-6 and IL-1 β contribute to early phase acetaminophen hepatotoxicity via NF- κ B signal pathway [57, 58], and these detrimental effects can be reduced by anti-inflammatory therapies at the early phase of acetaminophen overdose [57, 58]. TNF- α acts synergistically with HMGB1 to promote the acute fatal liver injury induced by D-galactosamine/LPS, and blockade of TNF- α and HMGB1 can synergistically ameliorate liver injury [59]. These data indicate that early inflammatory mediators promote liver injury at the early phase of drug hepatotoxicity.

TNF- α is a well-known early pro-inflammatory mediator that promotes hepatic tissue damage at the early phase of acetaminophen overdose [57–59]. However, TNF- α has also been reported as an important pro-regenerative cytokine that can facilitate the late phase liver regeneration during acetaminophen toxicity, because TNF- α can make the quiescent hepatocyte become more responsive to growth factors [60–62]. Acetaminophen overdose induces hepatocyte necrosis; liver regeneration is, therefore, vital for survival after the drug overdose insult [60, 63]. Prior to liver repair, the pro-regenerative cytokines such as TNF- α , IL-6 and IL-8 are needed to sensitize the hepatocyte, this process is crucial for the late phase liver regeneration [60, 64, 65], and this notion is supported by the following evidences. In an acetaminophen-induced acute liver injury animal model, Ringer's lactate solution (RLS) is used to improve the late phase hepatocyte regeneration, because this frequently used resuscitative fluid has pro-inflammatory effect, which can elevate the circulating levels of IL-6, IL-8 [66, 67] and TNF- α in patients and animals [67, 68]. As compared to the saline treatment, RLS therapy increases hepatic tissue TNF- α level at the early phase and improves liver repair at the late phase of acetaminophen hepatotoxicity; the improved liver regeneration is associated with elevated serum TNF- α concentration [69]. In another acetaminophen overdose animal model, the human well-tolerated novel anti-inflammatory agent ethyl pyruvate (EP) is used to ameliorate liver injury. Compared to the saline therapy, EP treatment attenuates liver injury at

early phase but impairs hepatocyte regeneration at the late phase of acetaminophen overdose, and the impaired late phase liver repair is associated with decreased serum TNF- α level [70]. In addition, neutralization of HMGB1 enhances hepatocyte regeneration at the late phase of acetaminophen overdose, and this beneficial effect is associated with increased hepatic TNF- α level [17]. These data indicate that TNF- α might contribute to early liver injury, which possibly acts synergistically with the other early inflammatory mediators. However, increased TNF- α level at early phase might prime hepatocyte and facilitate the late phase liver regeneration in acetaminophen hepatotoxicity.

Kupffer cells in acetaminophen hepatotoxicity

Kupffer cells (KCs) are hepatic resident phagocytes that serve as sentinels for liver homeostasis [71]. Once activated, KCs can release a large amount of inflammatory mediators that can promote liver injury [69]. Therefore, KCs play an important role in acetaminophen hepatotoxicity [72, 73]. In an acetaminophen overdose animal model, the depletion of KCs protects against liver injury at the early time point [73] but leads to more severe liver injury at the late phase of hepatotoxicity [72]. Macrophages contain two different subsets [74]. The M1 macrophages appear to promote liver injury at the early time point but the M2 macrophages show evidence to protect against liver injury at the late time point of acetaminophen overdose [74]. Except that KCs can produce inflammatory cytokines, new evidence shows that KCs also express multiple angiogenic factors, which can induce liver sinusoidal endothelial cell (LSEC) proliferation and migration [75]. When KCs are depleted in acetaminophen-induced liver injury animal models, this can significantly affect the late phase liver regeneration because of the lack of KCs-related angiogenic factors. This theory is confirmed in the following acetaminophen overdose animal model in which the depletion of KCs significantly delays liver structure recovery at the late time point; this detrimental effect is caused by acetaminophen-induced liver sinusoidal endothelial cell injury, which cannot rapidly recover without KCs-related angiogenic factors, leading to the prolonged vascular leakage. The delayed liver regeneration is not caused by impaired hepatocyte proliferation in this acetaminophen overdose model [75]. These evidences suggest that KCs play a complicated role in acetaminophen hepatotoxicity. KCs promote liver injury at early time point by producing pro-inflammatory mediators; these hepatic resident macrophages also have a significant impact on the late phase liver regeneration by affecting liver blood vessel repair via the angiogenic factors during hepatotoxicity.

Conclusions

DAMPs significantly contribute to liver injury and link local tissue damage to systemic inflammation during drug hepatotoxicity. DAMPs are promising mechanistic biomarkers and could also be the potential therapeutic targets for drug hepatotoxicity. DAMPs-triggered sterile inflammation contributes to liver injury at early phase but improves liver regeneration at later phase of acetaminophen hepatotoxicity; therefore, anti-inflammatory therapy would be beneficial at early phase but should be avoided at the late phase of acetaminophen toxicity.

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Conflict of interest The authors declare that they have no conflicts of interest.

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