

NIH Public Access

Author Manuscript

Alcohol Clin Exp Res. Author manuscript; available in PMC 2013 August 13.

Published in final edited form as:

Alcohol Clin Exp Res. 2011 May ; 35(5): 782–786. doi:10.1111/j.1530-0277.2010.01398.x.

The unfolding web of innate immune dysregulation in alcoholic liver injury

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Summary

Inflammatory cell and cytokine cascade activation is present in humans with alcoholic liver disease as well as in animal models of alcohol-induced liver damage. Gut-derived lipopolysaccahride (LPS), a ligand of the Toll-like receptor 4 (TLR4), plays a central role in triggering and maintaining activation of Kupffer cells in alcoholic hepatitis. In this mini-review, we describe molecular mechanisms that lead to increased inflammatory cell activation by alcohol and LPS and discuss the mechanism for activation in alcohol-exposed macrophages. In alcohol-induced liver disease we discuss the role of MyD88-independent but IRF3-mediated TLR4 signaling in alcohol-related liver inflammation and liver damage.

Introduction

The history of alcohol consumption traces back to Mesopotamia but the ancient Egyptians were also drinkers as they invented the first straws for drinking beer that still contained wheat-husks. Some of Egyptian texts even refer to the social problems associated with drunkenness. Throughout human civilization to date alcohol use remained the most commonly used substance of abuse in all societies and remains to be one of the most common etiologies for liver disease. There are multiple factors contributing to alcoholic liver disease including genetic factors, environment, gender and co-existing diseases. The pioneering research of the late Charlie Lieber opened the stage for recent discoveries in alcoholic liver disease after showing that alcohol has direct biological effects on the liver as opposed to only causing nutritional impairment (Lieber, 1975). While the direct effects of alcohol and its metabolites have received significant attention in causing direct damage to hepatocytes in the liver, the pathomechanism of alcoholic liver disease is more complex and it involves interactions between the gut, the site of alcohol absorption, the portal circulation and the different cell types in the liver. Crosstalk between the various cell types in the liver including parenchymal cells such as hepatocytes, biliary epithelium, stellate cells, liver sinusoidal endothelial cells, and Kupffer cells adds another level of regulation disrupted during liver injury.

Liver-gut axis in alcoholic liver disease

The unique anatomical position, blood supply, and architecture of the liver expose hepatocytes and cells in the liver sinusoids not only to gut-derived nutrients required for metabolism but also to gut-derived microbial products. In health, a normal balance of gut barrier function, gut permeability and a balance of commensal and pathogenic microbes in the gut lumen is maintained. This prevents microbial translocation from the gut (Rao, 2009). In normal homeostasis, the liver participates by detoxification of gut-derived toxins and

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microbial products through uptake by hepatocytes and Kupffer cells in a manner that prevents cell damage or inflammation.

There is increasing evidence for a dysbalance of the gut-liver axis in various liver diseases including alcoholic liver disease, non-alcoholic fatty liver disease and steatohepatitis (Keshavarzian, 1999; Keshavarzian, 2009; Miele, 2009). It is believed that gut microbial translocation can be a result of gut barrier dysfunction, increased gut permeability, and/or microbial overgrowth. Hepatocyte dysfunction and Kupffer cell activation also contribute to the dysbalance of the gut-liver axis when increased microbial products enter the liver to activate immune cells, particularly Kupffer cells, the liver resident macrophages, and other immune cells to include production of oxygen radicals and pro-inflammatory mediators (Mandrekar, 2009; Nagy, 2003; Wu, 2009:). Liver-derived inflammatory cytokines, particularly TNFa by entering the systemic circulation, can further increase gut permeability by disrupting tight junctions of gut epithelial cells (Yajima, 2009). Alcohol consumption can directly impair the gut barrier function by disruption of epithelial cell tight junctions, bacterial overgrowth and potentially through other, yet to be explored mechanisms (Bode, 2003). Several studies indicate that bacterial translocation as well as Kupffer cell activation play a central role in the pathomechanism of alcoholic liver disease. Sterilization of the gut with antibiotics or elimination of Kupffer cells, respectively, attenuated alcoholic fatty liver in animal models (Adachi, 1994; Adachi, 1995).

LPS sensing and its modulation by alcohol use

Lipopolysaccharide, a component of Gram negative bacteria, is a potent activator of innate immune responses through its binding to the Toll-like receptor 4 (TLR4) complex. The lipid A component of LPS is sensed as a danger signal by the mammalian hosts via the TLR4 receptor complexes. While TLR4 cannot directly bind LPS, the co-receptors, CD14 or MD-2 bind LPS and upon LPS binding activate TLR4. TLR4 dimerization leads to downstream signaling of two pathways: the MyD88-dependent and MyD88-independent pathways. Association of the intracellular TIR domain of TLR4 with the adapter molecule MyD88 through TRAM, results in downstream activation of the IRAK1/4/TRAF6 complex and further activation of the IKK kinase complex that phosphorylates IrB to allow nuclear translocation of NF-κB (Verstak, 2009). NF-κB binding to the NF-κB responsive element in the promoter region of pro-inflammatory cytokine genes results in production of TNFa, and other pro-inflammatory cytokines and chemokines (Akira, 2006). The MyD88-independent signaling pathway is activated by TLR4 after recruitment of the TRIF adapter, TRAF6 and TBK/IKKe phosphorylation leading to phosphorylation of the interleukin regulatory factor 3 (IRF3) (Schafer, 1998). This leads to IRF3 nuclear translocation and induction of Type I IFNs.

There is increasing evidence that moderate/acute and heavy/chronic alcohol use have different effects on human health (Sazbo and Mandrekar, 2009). Moderate alcohol use is associated with benefits on overall mortality, cardiovascular diseases and diabetes, while chronic excessive alcohol intake can lead to damage of the liver, lung, pancreas and other organs. Many of these processes involve activation of inflammatory cells as a critical component of the disease pathomechanism. At the cellular and molecular level, acute alcohol administration inhibited while chronic alcohol use increased production of pro-inflammatory mediators particularly when LPS was used as an inflammatory insult (Messingham, 2002).

Previous studies demonstrated that consumption of a single dose of alcohol in vivo significantly attenuated LPS-induced production of TNFa and IL-1ß and NF- κ B activation in human monocytes (Szabo, 1996). This was in contrast with the increased TNFa

production and NF-xB activation found in monocytes of patients with alcoholic steatohepatitis or after prolonged in vitro alcohol treatment (McClain, 1989). The molecular mechanisms involved in the effects of acute and prolonged alcohol exposure were tested in normal human monocytes. It was shown that acute and prolonged alcohol use had opposite effects on TNFa production in monocytes (Mandrekar, 2009). Acute alcohol (25 mM) resulted in significant attenuation of LPS-induced TNFa production at the mRNA and protein levels. Assessment of the TLR4 signaling pathway revealed that acute alcohol treatment inhibited IRAK1 phosphorylation, IKK kinase activity and NF-rB phosphorylation and activation in response to a subsequent LPS stimulation (Mandrekar, 2009). Such attenuation of TLR4 signaling by acute alcohol was similar to LPS tolerance (Medvedev, 2002). Recent studies found that TLR/LPS tolerance is induced via upregulation of TLR inhibitor molecules. In monocytes/macrophages the negative regulator, IRAKmonocytes (IRAK-M) is upregulated after a single LPS dose resulting in tolerance to the second LPS dose (Escoll, 2003). In human monocytes after acute alcohol treatment, IRAK-M was upregulated both at the RNA and protein levels, and siRNA knock-down of IRAK-M prevented the alcohol-induced inhibition of TNFa production in response to LPS (Mandrekar, 2009). In contrast to acute alcohol, prolonged alcohol exposure of monocytes for 4 days or longer in vitro, augmented LPS-induced TNFa production compared to alcohol-naïve cells (Mandrekar, 2009). The involvement of the TLR4 signaling pathway was suggested by increased IRAK-1 phosphorylation, increased IKK kinase activity, increased NF-kB nuclear translocation and DNA transactivation (Mandrekar, 2009). Upregulation of TLR4 signaling occurred in the presence of diminished expression of IRAK-M in monocytes after prolonged alcohol treatment. Over-expression of IRAK-M prevented the increased LPS-induced TNFa production in chronic alcohol-treated cells suggesting that loss of IRAK-M is likely to contribute to the loss of TLR4 tolerance in monocytes after prolonged alcohol exposure (Mandrekar, 2009).

TLR4 signaling in alcoholic liver disease

The importance of LPS-induced cell signaling has been demonstrated in animal models where mutation in the LPS receptor or the LPS adapter, CD14, showed protection from early alcoholic liver steatosis (Uesugi, 2001; Yin, 2001). However, the involvement of the specific components of TLR4 signaling pathway have not been investigated. In a recent study we found that mice deficient in the expression of TLR4 were protected from development of ALD on a Lieber-DeCarli diet (Hritz, 2008). There were decreased steatosis and inflammation and significantly reduced liver triglyceride, TNFa and IL-6 serum levels in TLR4-deficient mice after chronic alcohol feeding, suggesting a key role for TLR4mediated signals in alcoholic liver disease. Engagement of TLR4 with its ligand, LPS, leads to activation of NF- κ B and pro-inflammatory cytokine induction via recruitment of the MyD88 adapter molecule to the TLR4 complex while recruitment of the TRIF adapter activates the TBK/IKKe kinase and IRF3 to induce Type I IFN production as well as NF-xB activation (Covert, 2005). To evaluate the hypothesis that MyD88-dependent and MyD88independent TLR4 signaling pathways may have different involvement in alcoholic liver disease, first we tested MyD88-deficient mice because the role of NF-rcB activation and TNFa induction has been identified as a major component in alcoholic liver disease (Szabo, 2009). Alcohol feeding with the Lieber-DeCarli diet (Lieber, 1975; Lieber, 1982) resulted in significant steatosis and liver damage indicated by ALT elevation in MyD88 deficient mice compared to pair-fed diet and the extent of alcohol-induced changes were comparable between alcohol-fed MyD88-deficient and wild-type mice. This observation suggests that TLR4-mediated signaling via MyD88-independent pathways is important in induction of alcoholic liver disease. The TLR4-induced MyD88-independent, TRIF-mediated pathway activation results in Type I IFN induction. Indeed, we found evidence for activation of the

Type I IFN pathway in livers with alcoholic liver disease where increased mRNA expression of ISG56 was found (Szabo unpublished data).

Furthermore, cell-specific activation of the IFN-inducible gene was reported; IRF7 in Kupffer cells of alcohol-fed mice was increased compared to control diet-fed animals (Hritz, 2008). In a different study it was also reported that TRIF-deficient mice were protected against alcohol-induced liver disease and it is likely that IRF3, a transcription factor downstream to TLR4/TRIF, may bind to the TNFa promoter region resulting in induction of TNFa (Zhao, 2008). The role of TLR4 in alcoholic liver disease also extends to stellate cell activation (Seki, 2007). However, the role of IRF3 in stellate cell activation is yet to be evaluated.

Conclusion

Both steatosis and inflammatory cascade activation are induced by TLR4-mediated intracellular signaling pathways in alcoholic liver disease. Studies with acute and prolonged alcohol administration suggest that in monocytes/macrophages acute alcohol results in attenuated while prolonged alcohol use sensitizes to LPS and TLR4 activation.. Recent data also suggests that the MyD88-independent TLR4 signaling cascade that induces Type I IFNs is important in the development of alcoholic liver injury. Further studies on cell-specific regulation and cross-cellular communication and determination of the cell-specific role of IRF3 in ALD await investigation.

Acknowledgments

This work was supported by NIAAA grants # AA017729 and AA011576

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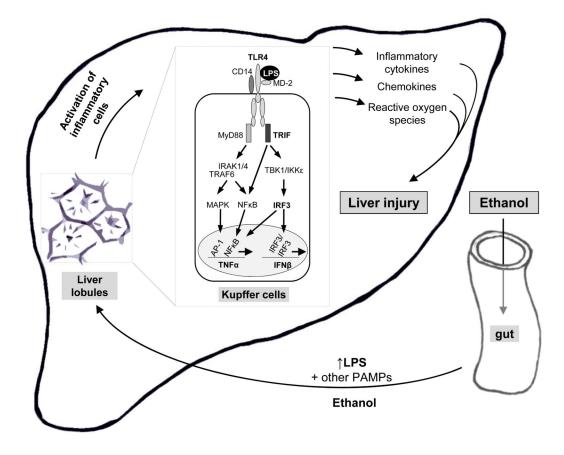


Fig. 1. New concepts in the pathophysiology of alcohol-induced liver injury

Ethanol increases translocation of bacterial endotoxin (lipopolysaccharide, LPS) and other pathogen-associated molecular patterns (PAMPs) from the gut to the liver. In the liver, LPS activates inflammatory cells, in particular Kupffer cells. In Kupffer cells, TLR4 recognizes LPS in cooperation with its co-receptors, CD14 and MD-2. The signal is passed preferentially through the TRIF-dependent intracellular pathways, which activate various transcription factors, including IRF3 and NF κ B, and induces pro-inflammatory cytokine and Type I interferon genes. Activated Kupffer cells are crucial in alcohol-induced liver injury by induction of inflammatory cytokines, chemokines and reactive oxygen species.

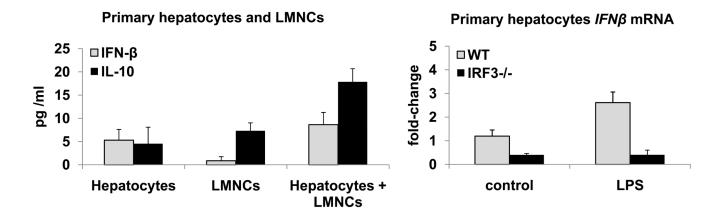


Fig. 2.

A. Cell-specific induction of IFN-ß and IL-10 in co-cultures of hepatocytes and liver mononuclear cells (LMNCs)

B. IRF3-dependent induction of IFN- β in hepatocytes

- TLR4 signaling via IRF3
- Cell-specific effects of IRF3 signaling

