

COMMENTARY

TLR4-MyD88 signalling: a molecular target for alcohol actions

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Several studies implicate Toll-like receptors (TLRs) in alcohol-induced neuroinflammatory processes. The work reported by Wu *et al.*, in this issue of the *British Journal of Pharmacology*, indicates that TLR4 along with its intracellular adaptor protein, MyD88, may play crucial roles in the acute actions of alcohol. The deletions of TLR4 or MyD88 gene or pharmacological inhibition of TLR4 by (+)-naloxone were able to attenuate alcohol-induced sedation, motor impairment and acute alcohol-induced increases in I κ B α protein levels in the hippocampus of mice. These results clearly suggest that TLR4-MyD88 signalling may play a causal role in the mediation of the behavioural effects of acute alcohol.

LINKED ARTICLE

This article is a commentary on Wu *et al.*, pp. 1319–1329 of this issue. To view this paper visit <http://dx.doi.org/10.1111/j.1476-5381.2011.01572.x>

Abbreviations

CeA, central nucleus of amygdale; Erk1/2, extracellular signal regulated kinase 1/2; I κ B α , inhibitor of κ B; JNK, c-Jun N-terminal kinase; LPS, Lipopolysaccharide; MeA, medial nucleus of amygdale; MyD88, myeloid differentiation primary response gene 88; TLR4, Toll-like receptor 4

Understanding the mechanisms underlying the acute actions of alcohol is crucial in order to identify, in more detail, the behavioural processes of alcohol tolerance and dependence and thus to advance the treatment of alcohol abuse disorders (Koob, 2003; Pandey *et al.*, 2008; Crews *et al.*, 2011). Alcohol addictive processes develop from a series of neuroadaptive changes leading to tolerance, withdrawal and relapse (Koob, 2003). Studies indicate that GABA, NMDA and other neurotransmitter systems may be involved in the acute and chronic effects of ethanol (Koob, 2003). Recently, a role for the innate immune system has been implicated in ethanol-induced neuroinflammatory processes leading to brain damage (Crews *et al.*, 2011). Toll-like receptors (TLRs) have been shown to be involved in these neuroinflammatory responses to chronic and acute ethanol exposure (Alfonso-Loeches *et al.*, 2010; Crews *et al.*, 2011; Pascual *et al.*, 2011). One of the TLRs, TLR4, has been shown to initiate the innate immune response by activating MyD88-dependent and -independent signalling pathways. MyD88 is an intracellular adaptor protein that interacts with TLR4 in response to LPS

exposure and activates protein kinases and gene transcription factors, such as NF- κ B, that can lead to increased expression of genes related to inflammation (Laird *et al.*, 2009; Alfonso-Loeches *et al.*, 2010; Crews *et al.*, 2011).

Acute ethanol exposure produces a variety of behavioural modifications, such as anxiolytic effects, sedation and motor impairment (Pandey *et al.*, 2008; Wu *et al.*, 2011; 2012). The studies published in this issue of the *British Journal of Pharmacology* by Wu *et al.* (2012) on TLR4-MyD88 signalling in relation to the behavioural responses of ethanol, clearly establish causal roles for TLR4 and MyD88 genes in alcohol-induced sedation and motor impairment in mice. They found that acute ethanol produced a sedative response in a dose-dependent manner and motor impairment, as measured by the rotor rod procedure in wild-type (+/+) mice, which was attenuated in TLR4 and MyD88 deficient (-/-) mice. Interestingly, these behavioural responses in wild-type mice were significantly attenuated by treatment with a TLR4 antagonist (+)-naloxone (Figure 1). They did not observe any effects of acute ethanol on p38, JNK or Erk 1/2 phosphorylation. NF- κ B

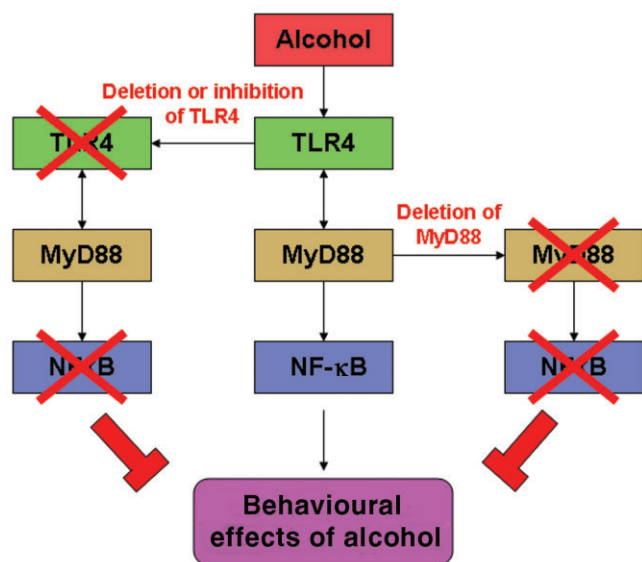


Figure 1

Molecular model depicting the actions of acute ethanol on TLR4-MyD88 signalling. As shown by Wu *et al.* (2012), activation of TLR4-MyD88 signalling by alcohol may mediate sedative and motor impairment effects, as they were attenuated by the deletion of TLR4 and MyD88 genes, or by the pharmacological blocking of TLR4 with (+)-naloxone.

is considered an important pro-inflammatory transcription factor and one of the genes that is activated by NF- κ B codes for I κ B α (Ferreiro and Komives, 2010). Wu *et al.* (2012) found that I κ B α protein levels were increased due to acute ethanol exposure in the hippocampus of wild-type (+/+) mice but not in TLR4 and MyD88 deficient (-/-) mice. These results suggest that TLR4-MyD88 signalling may be important in the molecular mechanisms underlying the actions of alcohol.

Several studies have supported this suggestion. A deficiency of TLR4 has been shown to protect against alcohol-induced glial activation and prevent induction of inflammatory processes including activation of NF- κ B pathways and apoptosis in mice (Alfonso-Loeches *et al.*, 2010). Interestingly, the motor ataxic effects of ethanol were enhanced in rats after 24 h, but not after 2 h of LPS exposure (Drugan *et al.*, 2007). Also, a single LPS injection increased innate immune gene expression in the brain (Crews *et al.*, 2011) and alcohol consumption in mice (Blednov *et al.*, 2011). Another interesting study pointed out the interactions between GABA $_A$ - α 2 and TLR4 in relation to binge drinking in alcohol-preferring (P) rats. Knocking down the expression of GABA $_A$ - α 2 by small interfering RNA (siRNA) in the central nucleus of amygdala (CeA) also decreased TLR4 expression and decreased binge drinking in P rats. Furthermore, siRNA knock down of TLR4 expression directly decreased binge drinking in P rats (Liu *et al.*, 2011). Deletion of certain genes related to chemokine networks resulted in altered alcohol-related behaviours in mice (Blednov *et al.*, 2005) and treatment with minocycline (blocker of pro-inflammatory microglial activation) and IL-1 receptor antagonists reduced the sedative effects of acute ethanol in mice (Wu *et al.*, 2011).

Studies conducted in mice and human post-mortem brains suggest that levels of a key innate immune chemokine, CCL2 (MCP-1), were increased in mice chronically treated with ethanol and also in several brain regions of human alcoholics (Crews *et al.*, 2011). A dysregulation in the NF- κ B system has also been observed in the prefrontal cortex of human alcoholics (Okvist *et al.*, 2007). Taken together, these studies suggest that altered innate immune gene expression along with its interaction with TLR4-MyD88 signalling and NF- κ B may play a neuroadaptive role in the development of alcoholism.

Anxiety that occurs during withdrawal from chronic ethanol exposure plays a fundamental role in relapse and may therefore maintain the continued abuse of alcohol (Koob, 2003). One recent study (Pascual *et al.*, 2011) demonstrated a role for TLR4 in anxiety-like behaviours developed in wild-type mice during abstinence after chronic ethanol exposure, as they were abolished in TLR4 deficient (-/-) mice. Alcohol also activated TLR4 receptors in glial cells and caused the induction of pro-inflammatory cytokines in the brain (Alfonso-Loeches *et al.*, 2010). In the absence of TLR4, these responses may not be initiated by alcohol; therefore, one can assume that anxiety-like behaviours were not seen in TLR4 deficient (-/-) mice, during withdrawal after chronic exposure.

Another interesting observation reported was that epigenetic modifications, such as changes in histone acetylation, were not observed in brain regions of TLR4 deficient (-/-) mice. Activity of histone acetyltransferases (HAT) and acetylation of histones H4 and H3 were decreased in several brain regions of wild-type mice during abstinence (15 days after a 5 month period of ethanol drinking). Reduced HAT activity and histone acetylation were not observed in the brains of TLR4 deficient (-/-) mice during abstinence after chronic ethanol exposure in mice (Pascual *et al.*, 2011). We reported that decreased histone (H3 and H4) acetylation in the CeA and medial nucleus of amygdala (MeA) may be involved in the development of anxiety-like behaviours during withdrawal after chronic ethanol exposure in rats. Acute ethanol exposure increased histone acetylation in the CeA and MeA and produced anxiolytic effects in rats (Pandey *et al.*, 2008). As mentioned previously, a deficiency in TLR4 in mice prevented changes in histone acetylation in the brain during the abstinence phase after chronic ethanol exposure (Pascual *et al.*, 2011), suggesting the possibility that TLR4 may interact with the chromatin architecture and may also be involved in the regulation of innate immune gene expression and the neurobiology of alcohol addiction.

In summary, the work conducted by Wu *et al.* (2012) along with other studies (Crews *et al.*, 2011; Pascual *et al.*, 2011) of the interactions between TLR4 and ethanol, suggests that TLR4 is an important molecular target in the brain that may mediate the behavioural responses related to acute and chronic ethanol exposure. Alcohol-induced activation of TLR4-MyD88 signalling may lead to activation of the NF- κ B gene transcription factor and, along with chromatin remodelling, may modify the expression of chemokines and cytokines, which in turn can regulate neuroinflammatory processes. Further work is needed to explore the links between TLR4 signalling and chromatin remodelling and interactions with the NF- κ B gene transcription factor and innate immune gene expression in brain regions implicated in cognition, reward and anxiety during alcoholism.

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Conflict of interest

None.

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