The anxious amygdala: CREB signaling and predisposition to anxiety and alcoholism

Gary Wand


Commentary

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The anxious amygdala: CREB signaling and predisposition to anxiety and alcoholism

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The amygdala is believed to play a key role in assigning emotional significance to specific sensory input, and conditions such as anxiety, autism, stress, and phobias are thought to be linked to its abnormal function. Growing evidence has also implicated the amygdala in mediation of the stress-dampening properties of alcohol. In this issue of the JCI, Pandey and colleagues identify a central amygdaloid signaling pathway involved in anxiety-like and alcohol-drinking behaviors in rats (see the related article beginning on page 2762). They report that decreased phosphorylation of cAMP responsive element–binding protein (CREB) resulted in decreased neuropeptide Y (NPY) expression in the central amygdala of alcohol-prefering rats, causing high anxiety-like behavior. Alcohol intake by these animals was shown to increase PKA-dependent CREB phosphorylation and thereby NPY expression, subsequently ameliorating anxiety-like behavior. These provocative data suggest that a CREB-dependent neuromechanism underlies high anxiety-like and excessive alcohol-drinking behavior.

Alcohol-abuse disorders

Alcohol-use disorders have been placed into 4 diagnostic categories of increasing severity: risky use, problem drinking, alcohol abuse, and alcohol dependence (1). This last and most severe form of alcohol disorders is characterized by loss of control over alcohol intake, tolerance, and physical dependence. It is estimated that approximately 14% of men and 5% of women in the US will experience the symptoms of alcohol abuse or dependence over their lifetimes (2). The economic cost of alcohol-use disorders is approximately $185 billion per year (3). Many people try alcohol, but few develop alcohol dependence. This selective vulnerability is, in part, because alcohol dependence has strong genetic determinants as evidenced by family, twin, and adoption studies (4). The genetic determinants for alcoholism create a vulnerable neural substrate that interacts with alcohol to create abuse potential.

Alcohol and the nucleus accumbens

Alcohol and alcoholism have both positive and negative reinforcing properties. The positive reinforcing properties of alcohol are linked to the hedonic aspects of alcohol intoxication. Considerable evidence shows that these positive reinforcing effects act through signal transduction systems affecting mesocorticolimbic dopamine (DA) pathways. The nucleus accumbens, a region at the base of the striatum, appears to be the key zone assigning importance to the alcohol exposure experience. Psychostimulants, opioids, and alcohol all increase synaptic DA accumulation within this important brain region (5). With PET imaging, we and others (6–8) have shown that mesolimbic DA release following drug administration is correlated with positive subjective effects. Preclinical studies have shown that drug reward can be attenuated by pharmacological or genetic manipulations that alter mesolimbic DA neurotransmission (9).

Alcohol and the amygdala

Alcohol-use disorders are also associated with negative reinforcing states. For example, individuals may drink alcohol to reduce anxiety or symptoms related to alcohol withdrawal. A large international study found strong associations between alcohol dependence and anxiety (10). Of particular interest, anxiety disorders were highly likely to predate the onset of alcoholism, lending support for a causal relationship. These data support the self-medicating hypothesis for alcohol-use disorders. This theory posits that a subset of individuals self-medicate with alcohol to reduce anxiety and/or stress (11). Additionally, an anxiety-like emotional state is often present during alcohol withdrawal and is thought to contribute to alcohol relapse in alcoholics (12).

In this issue of the JCI, Pandey and colleagues suggest that the neural substrate underlying the stress-dampening properties of alcohol is not the nucleus accumbens, but is probably the extended amygdala (13). The extended amygdala plays a crucial role not only in anxiety behaviors, but also in promoting alcohol intake (14). The amygdala receives limbic and olfactory afferents and projects fibers that innervate the hypothalamus and midbrain. Thus, the extended amygdala links the basal forebrain to the mesolimbic reward systems. This circuitry may mediate the anxiolytic effects of ethanol (14).

Selectively bred rodent lines

Rodent lines selectively bred to have a preference for alcohol over water have been an invaluable resource in the field of alcohol studies. Particularly useful lines are the alcohol-prefering (P) and alcohol-nonpreferring (NP) rats derived from a heterogeneous stock of Wistar rats at Indiana University (15). These rats are products of inbreeding high-alcohol drinking and low–alcohol drinking rats, respectively. The P rats generally drink greater than 5 grams of alcohol per kilogram body weight per day, whereas the NP line drinks less that 1.5 g/kg/d. In this issue of the JCI, Pandey and coworkers used these rats to investigate the role of cAMP/PKA signaling in the central amygdala (CeA) as it relates to alcohol consumption and anxiety (13). The CeA is a major component of the extended amygdala. It has been identified as a site of action for negative reinforcement associated with drug abstinence (14).

The gene transcription factor cAMP responsive element–binding protein (CREB) — a component of many signaling cascades...
activated by neurotransmitter receptor engagement — is regulated by cAMP-dependent PKA. PKA-mediated phosphorylation of CREB regulates the downstream expression of cAMP-inducible genes including NPY (16). Many studies have shown that a relationship exists between this G protein-mediated signaling cascade and responses to alcohol (17). Our group initially showed that alcohol can modulate the phosphorylation of CREB (18), and several studies subsequently demonstrated that CREB plays a role in alcohol dependence and preference (19–21). For example, CREB haplodeficient mice have a higher preference for alcohol than wild-type mice (21). NPY has been shown to play a role in anxiety and alcohol abuse (22). NPY-null mice display more anxiety-like behaviors and consume more alcohol than wild-type mice (23). The targeted disruption of specific NPY receptors alters ethanol consumption, and selective NPY receptor antagonists suppress ethanol intake in P C57BL/6j mice (24). NPY expression levels, similar to phosphorylated CREB levels, have been reported to be lower in the CeA of P rats compared to NP rats (25).

In this issue of the JCI, Pandey and coworkers show that P rats displayed higher baseline anxiety-like behaviors and consumed greater amounts of alcohol compared with NP rats (13), supporting prior observations (26). This finding prompted the investigators to posit that P rats drink excessive amounts of ethanol in order to reduce anxiety levels. Both investigator- and self-administered ethanol stimulated cAMP/PKA signaling in the CeA and medial amygdala (but not the basolateral amygdala), as evidenced by increased expression levels of the α-catalytic subunit of PKA (PKA-Cα), phosphorylated CREB, and NPY. These changes were accompanied by a reduction in anxiety-like behavior in the P rats (Figure 1). Impressively, the neurochemical and behavioral effects of alcohol were then mimicked by infusion of the PKA activator Sp-cAMP, or of NPY, into the CeA of these rats. In contrast, these manipulations did not produce any changes in anxiety levels or in phosphorylated CREB–induced NPY expression in the amygdaloid structures of NP rats. However, infusion of the PKA inhibitor Rp-cAMP into the CeA provoked anxiety-like behaviors and increased alcohol intake in NP rats.

A provocative implication of these findings (13) is that decreased CREB function in the CeA may be operative in maintaining high anxiety levels and alcohol-drinking behaviors of P rats. These data need to be replicated and then placed in context with 2 other very important neurotransmitter systems in the amygdala: corticotropin-releasing factor–mediated (CRF-mediated) and GABA-mediated signaling. CRF is an anxiogenic neuropeptide. Microdialysis studies have measured increased extracellular levels of CRF in the CeA during acute ethanol withdrawal in rats (27). CRF antagonists have been shown to reverse the excessive drinking of ethanol associated with ethanol withdrawal (28). Likewise, behavioral and neurochemical studies have implicated GABAergic transmission in the CeA in regulating alcohol intake (29). Chronic administration of ethanol is associated with increased GABA release in the CeA, which is mediated in part through CRF (11).

Although the findings reported by Pandey and coworkers (13) are important, cautious interpretation is appropriate. While animal models employed to study alcohol-use disorders have been extremely helpful, they have limitations. Alcoholism is a complex behavioral disorder, the product of environmental and polygenic interactions. Equating the human experience of anxiety with a rodent’s “choice” in an open field test may be stretching the limits of extrapolation. Furthermore, while rodent studies have demonstrated powerful efficacy for several classes of medications in reducing alcohol consumption, the effect of these drugs in treatment of alcohol-use disorders in humans has been, at best, weak (30). There is a clear need to develop more reliably predictive animal models for alcoholism.

With these caveats, the findings of Pandey and coworkers (13) are noteworthy. Their data inform us that we cannot limit our investigational focus to the mesolimbic DA system; the amygdala must be included when formulating models for alcoholism.

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Cholesterol efflux from macrophages, the first step in reverse cholesterol transport (RCT), is assumed to play a critical role in the pathogenesis of atherosclerosis. However, in vivo proof supporting this hypothesis is lacking, due to difficulties in determining the activity of this first step in RCT. In this issue of the JCI, Zhang et al. apply their recently developed method for measuring RCT in vivo to estimate RCT in mouse models with varying levels of HDL turnover. A surprisingly efficient clearance of cholesterol to feces is observed in mice overexpressing hepatic scavenger receptor class B type 1 (SR-BI), whereas in SR-BI–knockout mice, cholesterol clearance is diminished (see the related article beginning on page 2870). The study demonstrates that hepatic SR-BI is a positive regulator of macrophage RCT in vivo.

The transport of excess cholesterol from the periphery into the liver and bile, followed by excretion in the feces, is defined as reverse cholesterol transport (RCT) (Figure 1). Since the original definition of RCT by Glomset and Norum in 1973 (1), this pathway has become increasingly popular as a target for therapeutic strategies aimed at achieving the regression of atherosclerosis. Theoretically, a lipid-laden macrophage can release its contents by excretion in the feces, but conclusive evidence for this activity has been lacking until now. It is not an easy task to estimate net cholesterol flux from peripheral tissues to feces. Neutral sterols found in feces are derived from several sources. The major source is the liver, where the bulk of body cholesterol and bile salts are synthesized. After their secretion from the liver in bile, both components are secreted into the intestine, where up to 95% of bile salts and 30–60% of cholesterol is reabsorbed. The nonabsorbed cholesterol partly undergoes bacte-