Behavioral/Systems/Cognitive

Corticotropin Releasing Factor Signaling in the Central Amygdala is Recruited during Binge-Like Ethanol Consumption in C57BL/6J Mice

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A well established body of work indicates a crucial role for corticotropin-releasing factor (CRF) in neurobiological responses associated with excessive dependence-like ethanol drinking in ethanol-vapor-exposed rodents. Recent evidence demonstrates a role for CRF in the modulation of binge-like ethanol consumption by nondependent mice, a behavior that can precede ethanol dependence. The CRF circuitry that is engaged by binge-like ethanol exposure, however, is unknown. Using converging approaches, we provide evidence that, similar to ethanol-vapor-induced increases in ethanol intake, CRF signaling in the central nucleus of the amygdala (CeA) is engaged during binge-like ethanol consumption by C57BL/6J mice. Specifically, we found that binge-like consumption of an ethanol solution (20% ethanol v/v) was attenuated by pretreatment with the CRF1R antagonists antalarmin, 4-ethyl-[2,5,6-trimethyl-7-(2,4,6-trimethyl-phenyl)-7H-pyrrolo[2,3-d]pyrimidin-4-yl]amino-1-butanol, and NBI-27914 at doses (30 mg/kg, i.p.) that did not alter nonbinge-like ethanol consumption. Binge-like ethanol consumption resulted in significant increases of CRF immunoreactivity in the CeA immediately following ethanol drinking and 18-24 h following ethanol removal and also blocked the ability of CRF to enhance GABAergic transmission in the CeA 18-24 h following ethanol removal. Pretreatment with bilateral injections of antalarmin ($1 \mu g/0.5 \mu$ l per side) into the CeA, but not the adjacent basolateral amygdala, significantly attenuated binge-like ethanol consumption. These findings suggest that CRF signaling in the CeA is recruited during excessive ethanol intake, before the development of dependence. We hypothesize that plastic changes in CRF signaling develop with repeated binge-like drinking episodes, contributing to the transition to dependence.

Introduction

Binge drinking defines an episode in which enough alcohol is consumed to generate blood alcohol concentrations (BECs) of 80 mg/dl or greater (National Institute on Alcohol Abuse and Alcoholism, 2004). Binge drinking is associated with numerous adverse consequences (Read et al., 2008; Stahre et al., 2009), including an increased risk of developing alcohol dependence (Jennison, 2004; McCarty et al., 2004). As such, recruitment of mechanisms thought to underlie alcohol dependence may occur during bouts of high alcohol drinking, including alcohol binges.

Corticotropin-releasing factor (CRF) plays a key role in excessive dependence-like ethanol drinking in rats exposed to high levels of ethanol via vapor exposure, through activation of the

G-protein-coupled CRF-1 receptors (CRF1R) in the central nucleus of the amygdala (CeA) (Roberto et al., 2010). The CRF system does not significantly modulate ethanol consumption in nondependent rats (Roberts et al., 2000), thus leading to the hypothesis that neuroadaptations within the CRF system develop over an extended course of excessive ethanol exposure, culminating in ethanol dependence (Koob, 2003, 2009). A defining feature of this hypothesis is that CRF signaling is not crucial to maintain the initial stages of ethanol consumption that precede a dependent state (Heilig et al., 2010).

An extension of this model was revealed by several recent studies from our group that established a role for CRF signaling in maintaining binge-like ethanol consumption by nondependent animals (Sparta et al., 2008; Lowery et al., 2010). Specifically, systemic administration of a selective CRF1R antagonist, CP-154,526, attenuated binge-like ethanol consumption by male C57BL/6J mice that achieved BECs of at least 80 mg/dl under control conditions. Interestingly, CRF1R signaling appeared to be exclusively recruited by excessive levels of ethanol intake, as pretreatment with CP-154,526 did not alter ethanol consumption by mice that drank moderate levels of ethanol with associated BECs of 40 mg/dl or less (Sparta et al., 2008). While these findings indicate that the CRF system modulates excessive bingelike ethanol drinking, the central site of action and the overlap

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with the CRF pathways that modulate dependence-like drinking are unknown.

The current studies were designed to test the hypothesis that the CRF neurocircuitry that underlies binge-like ethanol consumption by nondependent animals overlaps with the CRF neurocircuitry that has been implicated in dependence-like ethanol drinking. We show that three different selective CRF1R antagonists attenuate binge-like ethanol consumption, but do not affect low-level, nonbinge-like ethanol consumption. Furthermore, binge-like ethanol consumption alters CRF immunoreactivity (IR) in extrahypothalamic brain regions and alters the neuromodulatory actions of CRF in the CeA of nondependent C57BL/6J mice. We also show that binge-like ethanol consumption is attenuated by intra-CeA microinjections of a selective CRF1R antagonist, while microinjections of the antagonist into the basolateral amygdala (BLA) do not alter binge-like ethanol consumption. Together, these results provide support for the hypothesis that high levels of ethanol intake by nondependent animals, such as those observed during binge-like drinking, recruit CRF signaling in the CeA, a region that is critical to the development of ethanol dependence.

Materials and Methods

Animals

Male C57BL/6J mice (Jackson Laboratories) were 6–8 weeks of age and weighed 20–30 g upon arrival. Mice were housed individually in plastic cages and allowed to habituate to the environment for at least 1 week before experimental procedures were initiated. The animal colony room was maintained at \sim 22°C with a 12 h/12 h light/dark cycle (lights on at 0700 or 1300 h). Mice had *ad libitum* access to food throughout all experiments and *ad libitum* access to water except during ethanol access, as noted. All procedures used are in accordance with the National Institute of Health guidelines and were approved by the University of North Carolina Institutional Animal Care and Use Committee.

Drugs

Ethanol (20% v/v) solutions were prepared using tap water and 95% ethyl alcohol; sucrose (10%, w/v) solutions were prepared using tap water and D-sucrose. The selective CRF1R nonpeptidic antagonists NBI27914 (10-60 mg/kg; Tocris Bioscience), antalarmin (30 mg/kg; Sigma-Aldrich), and 4-ethyl-[2,5,6-trimethyl-7-(2,4,6-trimethylphenyl)-7*H*-pyrrolo[2,3-*d*]pyrimidin-4-yl]amino-1-butanol (LWH-63; 10-30 mg/kg; Sigma-Aldrich) were dissolved in a 0.9% saline and emulphor (also known as Cremphor EL; Sigma-Aldrich) solution (10% emulphor v/v) and delivered intraperitoneally in a 10 ml/kg injection volume ~30 min before ethanol access. Doses were extrapolated from previous studies (Robison et al., 2004; Sabino et al., 2006; Ji et al., 2007; Kim and Han, 2009). For the intra-CeA and intra-BLA microinjections, antalarmin (1.0 μ g/ 0.5 μ l per side) was dissolved in 0.9% saline and emulphor solution (5% emulphor, v/v) and delivered $\sim\!1\,h$ before ethanol access. Previous studies were used to select the dose (Robison et al., 2004; Wang et al., 2006) and injection volume (Finn et al., 2007). CRF (Tocris Bioscience) was dissolved in dH₂O to a concentration of 0.1 mm, diluted to 200 nm in artificial CSF (ACSF), and bath applied.

Behavioral procedures

Drinking-in-the-dark experiments. All experiments used a 4 d drinking-in-the-dark (DID) procedure that has been used previously to generate high levels of ethanol intake that are associated with BECs in excess of 80 mg/dl (Rhodes et al., 2005, 2007; Sparta et al., 2008; Lowery et al., 2010). On days 1–3, beginning 3 h into the dark cycle, water bottles were removed from all cages and replaced with bottles containing a 20% (v/v) ethanol solution. Mice had 2 h of access to ethanol, after which ethanol bottles were removed from cages and water bottles were replaced. The same procedure was followed on day 4 except that ethanol access was extended to 4 h. Immediately following 4 h ethanol access on day 4, \sim 10 μ l of blood were collected from the tail vein of each animal. Blood samples were centrifuged, and 5 μ l of plasma were analyzed for BECs using

the Analox Alcohol Analyzer (Analox Instruments). For all experiments involving pretreatment with drug on day 4, drug treatment groups were equated based on ethanol consumption on days 1–3. For all experiments involving multiple cycles of DID procedures, each cycle was separated by 3 d. To assess the specificity of experimental manipulations to binge-like ethanol consumption, sucrose control studies and nonbinge-like ethanol consumption control studies were conducted. Briefly, sucrose control studies were conducted following the DID procedure similar to ethanol DID procedures except that 10% sucrose (w/v) solution was presented in place of 20% ethanol solution. Nonbinge-like ethanol control studies were conducted using similar procedures to DID, except that access to 20% ethanol began 3 h into the light portion of the light/dark cycle.

Perfusion, brain preparation, and immunohistochemistry procedures. Immunohistochemistry procedures were based on those routinely used in our laboratory (Thiele et al., 1996, 1997, 2000; Hayes et al., 2005). Mice were perfused transcardially with 0.1 mm PBS (pH 7.4) followed by 4% paraformaldehyde in phosphate buffer immediately following ethanol access on day 4 of the final DID cycle or 18-24 h following ethanol access on day 4 of the final DID cycle. The brains were collected and postfixed in paraformaldehyde for 48 h at 4°C and then transferred to PBS. The brains were cut using a vibratome into 40 μ m sections and stored in cryopreserve until the immunohistochemistry assay. Sections were then transferred to PBS for 24 h before processing with CRF antibody. After rinsing in fresh PBS four times (10 min each), tissue sections were blocked in 10% goat serum and 0.1% Triton X-100 in PBS for 1 h. Sections were then transferred to fresh PBS containing primary antibody for 72 h at 5°C. CRF expression was detected using primary rabbit anti-CRF (T-4037; 1:5000, 1:10,000, or 1:20,000 as determined by pilot experiments; Peninsula Laboratories). As a control to determine whether staining required the presence of the primary antibodies, some sections were run through the assay without primary antibody. After the 72 h of incubation, the sections were rinsed four times and then processed with Vectastain Elite kits (Vector Laboratories) as per the manufacturer's instructions for standard ABC/HRP/diaminobenzidine-based immunohistochemistry. The sections processed for CRF were visualized by reacting the sections with a 3,3'-diamino-benzidine tetrahydrochloride (DAB; Polysciences) reaction solution containing 0.05% DAB, 0.005% cobalt, 0.007% nickel ammonium sulfate, and 0.006% hydrogen peroxide. All sections were mounted on glass slides, air-dried overnight, and coverslipped for viewing. Digital images of each candidate brain region were obtained on an E400 microscope equipped with a Digital Sight DS-U1 digital camera run with manufacturer-provided software (all from Nikon Instruments). CRF-IR was quantified in each region at the level that showed the most robust staining and great care was taken to match sections within each brain region at the same level using anatomic landmarks with the aid of a mouse stereotaxic atlas (Paxinos and Franklin, 2004). Densitometric procedures were used to assess levels of CRF-IR in each brain region. Flat-field corrected digital pictures (8-bit grayscale) were taken using the Digital Sight DS-U1 camera. The density of staining was analyzed using ImageJ software (NIH) by calculating the percent of the total area examined that showed signal (cell bodies and processes) relative to a subthreshold background. To account for differences in background between pictures within the same region, only the darkest area(s) of staining in each image was defined as CRF-IR. Specifically, the darkest area of staining was defined as the first complete cell body to be highlighted by ImageJ's threshold function. This cell body and all other areas of staining (including both cell bodies and processes) that were highlighted at this threshold were identified as CRF-IR. Anatomically matched pictures of the left and right sides of the brain were used to produce an average density for each brain region from each slice. In all cases, quantification of immunohistochemistry data was conducted by an experimenter who was blinded to group identity.

Brain slice preparation and electrophysiology. Mice were killed $18-24\,h$ following ethanol access on day 4 of the final DID cycle and their brains were placed in ice-cold sucrose ACSF containing the following (in mm): 194 sucrose, 20 NaCl, 4.4 KCl, 2 CaCl₂, 1 Mg Cl₂, 1.2 NaH₂PO₄, 10.0 glucose, and 26.0 NaHCO₃ saturated with 95% O₂/5%CO₂. The brains were cut into $300~\mu m$ sections using a vibratome and slices containing the CeA were identified using well characterized landmarks such as the BLA.

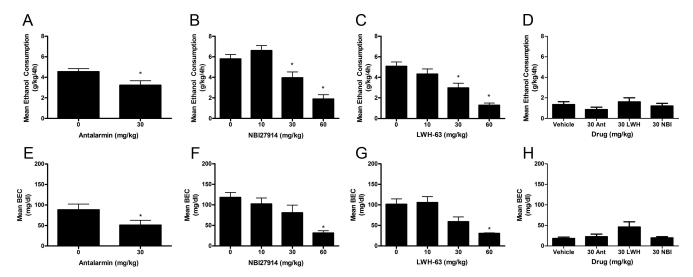


Figure 1. Pretreatment with CRF1R antagonists attenuate binge-like ethanol consumption and BECs but does not alter nonbinge-like ethanol consumption. Antalarmin (30 mg/kg dose) attenuated mean binge-like ethanol consumption (*A*) and mean BECs (*E*). NBI27914 reduced mean binge-like ethanol consumption at the 30 and 60 mg/kg doses (*B*) and the associated mean BECs at the 60 mg/kg dose (*F*). LWH-63 (30 and 60 mg/kg doses) attenuated mean binge-like ethanol consumption (*C*) and the associated mean BECs at the 60 mg/kg dose (*G*). *D*, Mean nonbinge-like ethanol consumption was not altered by 30 mg/kg doses of antalarmin, NBI27914, or LWH-63. *H*, Associated mean BECs were not altered by 30 mg/kg doses of antalarmin (Ant), NBI27914 (NBI), or LWH-63 (LWH). All data are presented as means ± SEM, significance was accepted at the p < 0.05 level, and n = 8-13 animals per group. *, Significant differences from the vehicle group.

Slices were then stored in a heated (28°C), oxygenated (95% O₂/5%CO₂) holding chamber containing ACSF [(in mm) 124 NaCl, 4.4 KCl, 2 CaCl₂, 1.2 MgSO₄, 1 NaH₂PO₄, 10.0 glucose, and 26.0 NaHCO₃₁ or transferred to a submerged recording chamber (Warner Instruments) where they were perfused with heated, oxygenated ACSF at a rate of 2 ml/min. Slices were allowed to equilibrate in ACSF for 1 h before the start of experiments. Whole-cell voltage-clamp recordings were conducted from slices in the submerged recording chamber. Neurons of the CeA were visualized using infrared video microscopy (Olympus America). Recording electrodes (3–6 M Ω) were pulled on a micropipette puller (Sutter Instruments) using thin-walled borosilicate glass capillaries. To analyze evoked IPSCs (eIPSCs), electrodes were filled with the following (in mm): 70 K⁺-gluconate, 80 KCl, 10 HEPES, 1 EGTA, 4 ATP, 0.4 GTP, pH 7.2, 290-295 mOsmol. Twisted nichrome wire stimulating electrodes were placed in the CeA, $100-500 \mu m$ from the recorded neuron. After entering a whole-cell configuration, cells were held at -70 mV and GABA type A receptor-mediated IPSCs were evoked at 0.2 Hz by local fiber stimulation with bipolar electrodes. GABA_A-IPSCs were pharmacologically isolated by adding 10 μ M NBQX. Signals were acquired via a Multiclamp 700B amplifier (Molecular Devices), digitized, and analyzed via pClamp 10.2 software (Molecular Devices). Input resistance and access resistance were continuously monitored during experiments. Experiments in which changes in access resistance were >20% were not included in the data analysis. eIPSC experiments were analyzed by measuring the peak amplitude of the synaptic response, which was normalized to the baseline period. The baseline period was defined as the 2 min period immediately preceding the application of the drug. CRF (200 nm) was bath applied at minutes 6-20.

Surgery and infusion procedures. Mice were surgically implanted with bilateral 26-gauge guide cannulae (Plastics One) aimed at the CeA (0.94 mm posterior to bregma; 2.55 mm lateral to the midline; 2.60 mm ventral to the skull surface) or the BLA (1.22 mm posterior to bregma; 3.01 mm lateral to the midline; 4.74 ventral to the skull surface) using the Angle II stereotax (Leica Instruments) with the aid of a mouse stereotaxic atlas (Paxinos and Franklin, 2004). Following surgery, DID procedures were initiated. On day 4, animals were injected bilaterally with antalarmin (1 μ g/ 0.5 μ l per side) or vehicle (5% emulphor in saline; 0.5 μ l per side) over a period of 60 s approximately 1 h before ethanol access using a 33-gauge injector designed to extend 2 mm beyond the cannula tip. A sucrose control study was conducted to evaluate the effects of intra-CeA antalarmin or vehicle on consumption of a 10% sucrose solution over a 4 h period beginning 3 h into the dark portion of the light/dark cycle. For

this control study, mice were assigned to drug treatment groups based on their body weights. Cannulae placements were determined for each animal using injections of blue dye and thionin staining of mounted sections of brain tissue. Only the data from mice in which both cannulae were correctly aimed at the CeA or the BLA were included. All data from mice in which both cannulae (n=1 for the CeA; n=3 for the BLA) or one cannula (n=8 for the CeA; n=4 for the BLA) did not target the CeA or the BLA were excluded.

Statistical analyses

One-way ANOVAs were used to assess between-group differences in binge-like ethanol consumption and BECs, CRF-IR, and basal GABAergic signaling. Repeated-measures ANOVAs were used to assess eIPSC peak amplitudes before CRF application (minutes 1–5) and while CRF was bath applied (minutes 6–20). A Student's t test was used to assess the main effect of DID treatment on CRF's modulation of GABAergic signaling. As the aim of the immunohistochemistry experiments were to assess the specific effects of binge-like ethanol consumption on CRF-IR, data from animals that did not achieve BECs of 80 mg/dl or greater are not presented (n=4). Because unquantifiable tissue varied by animal and region in these experiments, the degrees of freedom may differ for each statistical analysis. Tukey's post hoc tests were used to assess specific between-group differences where appropriate. Significance was accepted at the p<0.05 (two-tailed). All data are presented as mean \pm SEM.

Results

Systemic administration of CRF1R antagonists attenuates binge-like ethanol consumption

To test the hypothesis that high levels of ethanol intake require CRF signaling, the effects of several small molecule, nonpeptide CRF1R antagonists on binge-like ethanol consumption were assessed. Average ethanol consumption and BEC data from day 4 are presented in Figure 1. An ANOVA revealed a significant main effect of antalarmin on binge-like ethanol consumption ($F_{(1,25)} = 6.548$, p = 0.017), confirming that the 30 mg/kg dose of antalarmin significantly blunted binge-like ethanol drinking (Fig. 1A). An ANOVA performed on associated BEC data was also significant ($F_{(1,25)} = 4.296$, p = 0.049), as antalarmin-induced reduction of ethanol drinking was associated with a significant reduction of BECs (Fig. 1 E). An ANOVA revealed a significant

main effect of NBI27914 on binge-like ethanol consumption $(F_{(3.37)} = 18.512, p < 0.001)$, and post hoc analyses revealed that pretreatment with the 30 or 60, but not the 10, mg/kg doses of NBI27914 significantly attenuated binge-like ethanol consumption relative to vehicle (Fig. 1B). Similar significant effects were observed on BEC data ($F_{(3,37)} = 8.048$, p < 0.001) and post hoc analyses indicated that pretreatment with 60 mg/kg resulted in significantly reduced BECs relative to vehicle (Fig. 1F). An ANOVA revealed a significant main effect of LWH-63 on bingelike ethanol consumption ($F_{(3,39)} = 13.591$, p < 0.001) and post hoc analyses showed that pretreatment with both 30 and 60 mg/kg doses of LWH-63 significantly attenuated binge-like ethanol consumption relative to vehicle treatment (Fig. 1C). Similarly, an ANOVA performed on associated BECs was statistically significant ($F_{(3,39)} = 9.402$, p < 0.001) and post hoc analyses indicated that pretreatment with the 60 mg/kg dose significantly reduced BECs (Fig. 1G). Overall results showed that 30 mg/kg antalarmin, 30 mg/kg LWH-63, and 30 mg/kg NBI27914 selectively altered binge-like ethanol consumption. A 30 mg/kg dose of antalarmin did not significantly alter sucrose intake (220.76 \pm 20.49 ml/kg for vehicle-treated mice vs 214.43 \pm 39.67 ml/kg for antalarmin-treated mice). Furthermore, pretreatment with 30 mg/kg doses of LWH-63 or NBI27914 did not significantly alter sucrose drinking (234.05 \pm 29.30 ml/kg for vehicle-treated mice vs 267.02 \pm 17.07 ml/kg for LWH-63-treated mice and 173.48 \pm 26.02 ml/kg for NBI27914-treated mice). These results indicate that CRF1R antagonists selectively modulate binge-like ethanol consumption.

Systemic administration of CRF1R antagonists does not alter nonbinge-like ethanol consumption

To test the hypothesis that low or moderate levels of ethanol intake do not require CRF signaling, the effects of doses of each of the CRF1R antagonists that altered binge-like ethanol consumption were assessed on nonbinge-like ethanol consumption. An ANOVA assessing the effects of CRF1R antagonists on nonbingelike ethanol consumption data was not statistically significant $(F_{(3,34)} = 1.068, p = 0.377; Fig. 1D)$. An ANOVA performed on associated BECs was statistically significant ($F_{(3,34)} = 3.190, p =$ 0.037), though post hoc analyses did not indicate any significant differences in BECs between the vehicle group and any of the groups pretreated with a CRF1R antagonist (Fig. 1H). These results are in agreement with previous investigations showing that CRF1R antagonists have no effect on ethanol consumption that is not preceded by procedures to induce dependence-like drinking (Sabino et al., 2006; Chu et al., 2007; Gehlert et al., 2007) or to induce stress (Marinelli et al., 2007; Lowery et al., 2008), as well as a previous report comparing the effects of the CRF1R antagonist CP-154,526 on binge-like ethanol consumption and nonbingelike ethanol consumption (Sparta et al., 2008; Lowery et al., 2010). Because the Sparta et al. (2008) study also found that a CRF1R antagonist did not alter nonbinge-like ethanol consumption during the dark portion of the light/dark cycle, the lack of effect observed in the current study is unlikely to be due to diurnal effects on CRF (Owens et al., 1990).

CRF-IR is increased immediately following binge-like ethanol drinking and 18–24 h following the termination of binge-like ethanol drinking

To verify the specificity of our antibody, brain tissue from a CRF wild-type (WT) mouse (Fig. 2*A*; CeA shown) and a CRF knock-out (KO) mouse (Fig. 2*B*; CeA shown) was processed with CRF antibody. Importantly, CRF-IR was not observed in the paraven-

tricular nucleus of the hypothalamus (PVN), the bed nucleus of the stria terminalis (BNST), the ventral tegmental area (VTA), or the CeA of a CRF KO mouse, but was evident in each of these regions of a CRF WT mouse. The effects of binge-like ethanol consumption on CRF-IR immediately following one (1-cycle) or six (6-cycle) binge-like drinking exposures were investigated to assess the effects of a single episode and multiple episodes of high levels of ethanol intake on central CRF signaling. All comparisons were made with an ethanol-naive group of mice that consumed water for the duration of the experiment (Water). Consumption of ethanol during the 4 h access period for the duration of the experiment is shown in Table 1. Immediately following the final day of the final DID cycle, the 1-cycle group had mean BECs of 123.28 ± 12.69 mg/dl and the 6-cycle group had mean BECs of 161.71 ± 17.15 mg/dl. The results of a one-way ANOVA confirmed that ethanol consumption and BECs on the final day of the experiment did not differ between DID treatment groups $(F_{(1,15)} = 0.430, p = 0.523 \text{ for ethanol consumption; } F_{(1,14)} =$ 2.657, p = 0.127 for BECs). Figure 2 shows CRF-IR in the CeA and VTA in DID treatment groups. Binge-like ethanol consumption significantly altered CRF-IR in the CeA ($F_{(2,21)}=6.398, p=$ 0.008) and post hoc analyses revealed that the 1-cycle and 6-cycle groups had significantly greater CRF-IR than the water group in this region (Fig. 2C-E). A significant effect of binge-like ethanol consumption on CRF-IR in the VTA was also observed ($F_{(2,18)} =$ 8.931, p = 0.002; Fig. 2). Post hoc analyses revealed that the 1-cycle group displayed significantly greater CRF-IR than the water group and the 6-cycle group (Fig. 2F-H). CRF-IR was not significantly altered following sucrose consumption in the CeA $(0.10 \pm 0.01\%$ area for the water group vs $0.15 \pm 0.01\%$ area for the 1-cycle group vs $0.15 \pm 0.02\%$ area for the 6-cycle group) or in the VTA (0.05 \pm 0.01% area for the water group vs 0.06 \pm 0.01% area for the 1-cycle group vs 0.04 \pm 0.01% area for the 6-cycle group). CRF-IR was not significantly altered in any other brain region assessed following binge-like ethanol consumption (Table 2) or following sucrose consumption (data not shown). These results indicate that binge-like ethanol consumption significantly alters CRF-IR of the CeA and the VTA. Because previous studies have primarily assessed the effects of ethanol on CRF expression after ethanol removal (Lowery and Thiele, 2010), we assessed CRF-IR in mice that had undergone three exposures to DID procedures 18–24 h after their final ethanol exposure (3cycle). Ethanol consumption over the course of this experiment is shown in Table 1. Animals of the 3-cycle group achieved mean BECs of 115.8 \pm 20.72 mg/dl during the final 4 h ethanol access. A one-way ANOVA revealed that the 3-cycle group had significantly greater CRF-IR in the CeA than the water group did $(F_{(1.19)} = 7.586, p = 0.013; Fig. 2 I-K)$. The resolution of the stain did not allow us to determine whether increases of CRF-IR were due to increases of CRF-IR within cells or an increase in the number of cells exhibiting CRF-IR (or both possibilities). Together, these findings extend those of previous reports showing upregulations of CRF markers in the CeA immediately following chronic ethanol ingestion (Läck et al., 2005) and during acute ethanol withdrawal from chronic intermittent ethanol vapor exposure (Sommer et al., 2008) by demonstrating that similar alterations in CRF-IR are induced by voluntary ethanol consumption that is associated with intoxicating BECs. Additionally, a lack of effect of binge-like ethanol consumption on CRF expression in the PVN corroborates the results of a previous investigation from our group that suggested that binge-like ethanol consumption was modulated by CRF activity outside of the hypothalamic-pituitary-adrenal axis (Lowery et al., 2008).

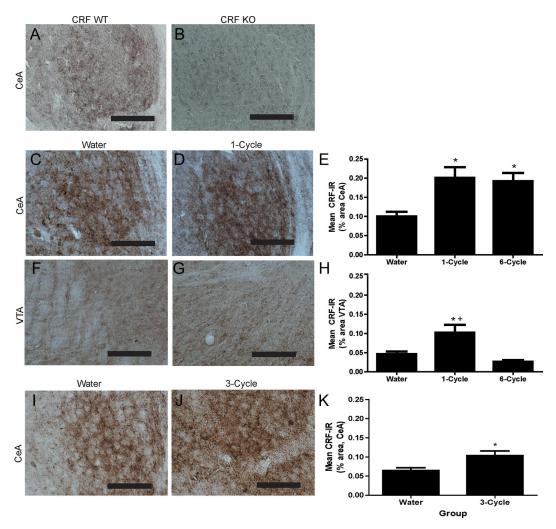


Figure 2. CRF-IR is significantly altered immediately following binge-like ethanol consumption and 24 h following ethanol removal. Representative photomicrographs demonstrate that the CRF antibody (antibody concentration: 1:20,000) is specific for CRF protein, as CRF-IR was evident in the CeA of a CRF WT mouse (A), but was not observed in the CeA of a CRF KO mouse (B). Representative photomicrographs of the CeA (C, D) and the VTA (F, G) show enhanced CRF-IR (antibody concentration: 1:5000) immediately following ethanol removal in animals exposed to one DID cycle (D, G) compared with the water-drinking group (C, F). E, The 1-cycle and 6-cycle groups had significantly greater mean CRF-IR immediately following ethanol removal in the CeA (percentage of the total area) than the water-drinking group. H, The 1-cycle group had significantly greater mean CRF-IR in the VTA (percentage of the total area) immediately following ethanol removal than the water group and the 6-cycle group. I, J, Representative photomicrographs of the CeA show CRF-IR (antibody concentration: 1:10,000) in animals of the water group (I) and the 3-cycle group (J) 18–24 h following ethanol removal. K, The 3-cycle group had significantly greater mean CRF-IR in the CeA (percentage of the total area) than the water group 18–24 h following their final exposure to ethanol. All data are presented as means ± SEM, significance was accepted at the p < 0.05 level, and n = 5–10 animals per group. *, Significant differences from the water group; +, significant differences from the 6-cycle group. Scale bars, 50 µm.

Table 1. Mean binge-like ethanol (g/kg/4 h \pm SEM) consumption on day 4 of each DID cycle for the duration of each experiment

CRF-IR immed binge-like dri	liately following nking	CRF-IR 18 —24 h following	GABAergic transmission 18 –24 h following binge-like drinking		
1-cycle	6-cycle	binge-like drinking			
Cycle 1 —	5.62 ± 0.38	5.37 ± 0.31	5.64 ± 0.41		
Cycle 2 —	6.02 ± 0.40	5.39 ± 0.23	6.07 ± 0.31		
Cycle 3 —	6.01 ± 0.31	5.27 ± 0.38	5.27 ± 0.34		
Cycle 4 —	6.70 ± 0.19	_	_		
Cycle 5 —	6.45 ± 0.31	_	_		
Cycle 6 5.81 ± 0.35	6.63 ± 0.34	_	_		

The modulatory effects of CRF on GABAergic transmission in the CeA are disrupted following binge-like ethanol drinking

Given that alterations of CRF-IR in the CeA were observed 18–24 h following ethanol removal in mice that had undergone three binge cycles, we hypothesized that multiple episodes of high ethanol intake would result in functional alterations in the CRF system. To maxi-

mize the effects of binge history on CRF function and minimize the likelihood of inducing dependence, we assessed the effects of three cycles of binge-like ethanol consumption on the neuromodulatory effects of CRF on GABAergic signaling in the CeA. As we were interested in the effects of a binge history on CRF function regardless of the effects of acute ethanol, this experiment was conducted 18-24 h following the final binge episode to ensure that ethanol was not on board at the time of testing. Binge-like ethanol consumption on day 4 of each DID cycle is shown in Table 1. Mice of the 3-cycle group achieved mean BECs of 133.48 \pm 10.5 mg/dl during the final 4 h ethanol access. Representative traces showing the effects of CRF on GABAergic transmission are shown in Figure 3A. The results of a repeated-measures ANOVA revealed that a history of binge-like ethanol exposure lead to a significant reduction in the ability of CRF to enhance GABAergic transmission in the CeA ($F_{(1,9)} = 9.163$, p = 0.014; Fig. 3B). The results of a t test revealed a significant main effect of ethanol history on the average modulation of eIPSC peak amplitude by

Table 2. Average CRF-IR (percentage area ± SEM) immediately following binge-like ethanol consumption

	BLA	MeA	NAccCore	NAccSh	PVN	LH	dIBNST	dmBNST	vBNST	LS
Water	0.02 ± 0.01	0.07 ± 0.03	0.06 ± 0.02	0.05 ± 0.02	0.11 ± 0.04	0.07 ± 0.03	0.15 ± 0.05	0.08 ± 0.03	0.06 ± 0.02	0.07 ± 0.03
1-cycle	0.02 ± 0.01	0.08 ± 0.03	0.06 ± 0.02	0.09 ± 0.03	0.09 ± 0.03	0.07 ± 0.02	0.13 ± 0.05	0.10 ± 0.04	0.06 ± 0.02	0.08 ± 0.03
6-cycle	0.02 ± 0.01	0.09 ± 0.03	0.10 ± 0.04	0.15 ± 0.05	0.14 ± 0.04	0.10 ± 0.04	0.21 ± 0.08	0.12 ± 0.04	0.06 ± 0.02	0.13 ± 0.05
F-statistic	0.220	0.624	2.483	2.848	1.589	1.677	1.306	1.434	0.154	2.876
p value	0.804	0.545	0.113	0.082	0.228	0.215	0.294	0.264	0.859	0.082

MeA, Medial amygdala; NAccCore, nucleus accumbens core; NAccSh, nucleus accumbens shell; LH, lateral hypothalamus; dIBNST, dorsolateral BNST; dmBNST, dorsomedial BNST; vBNST, ventral BNST; LS, lateral septum.

CRF ($t_{(9)} = 2.445$, p = 0.0370), as the average peak amplitude of IPSCs evoked in the presence of CRF was enhanced by $29 \pm 11\%$ in water-drinking control mice but was not increased ($-7 \pm 9\%$) in mice with a history of binge-like ethanol consumption (Fig. 3C). In addition, to determine whether there were any differences in the probability of basal GABA release, the paired-pulse ratio of the evoked IPSC in both water- and alcohol-exposed mice were assessed, and no significant differences were found ($t_{(9)} = 0.3392$, p = 0.74; Fig. 3D). Finally, we evaluated the ability of a CRFR1 antagonist, NBI27914 (1 µM), to alter GABAergic transmission in both control and alcohol-exposed mice. This experiment was done to determine whether there was an endogenous CRF tone in the slice related to the ethanol exposure. We found that bath application of NBI27914 had no effect on the peak amplitude of eIPSC in either controls (93 ± 8% of baseline, n = 4) or alcohol-exposed mice (108 \pm 4% of baseline, n = 3), suggesting that there is no alteration in endogenous CRF tone in the CeA following binge-like ethanol drinking. Together, these results suggest that, despite concomitant upregulations of CRF-IR in this region at the same time point, the modulatory effects of CRF on GABAergic transmission in the CeA are blunted in mice with a history of binge-like ethanol consumption.

Intra-CeA microinjections of a CRF1R antagonist attenuate binge-like ethanol consumption

Our IR results suggest that CRF signaling may be engaged in the CeA and motivate binge-like drinking. To test this hypothesis, the effects of microinjections of antalarmin into the CeA on binge-like ethanol consumption were assessed. Average ethanol consumption and BEC data from day 4 are presented in Figure 4. The results of a one-way ANOVA revealed a significant effect of drug treatment ($F_{(1,14)} = 8.921, p = 0.011$), confirming that intra-CeA administration of antalarmin significantly blunted binge-like ethanol consumption (Fig. 4A). Significant between-group differences in BECs were also observed ($F_{(1,14)} = 4.847, p = 0.046$), as mice pretreated with antalarmin achieved significantly lower BECs than mice that received vehicle (Fig. 4B). Conversely, pretreatment with antalarmin into the CeA did not have effects on 4 h sucrose consumption ($F_{(1,16)} = 1.458, p = 0.246$; Fig. 4C). Cannula placements are shown in Figure 4D. These results dem-

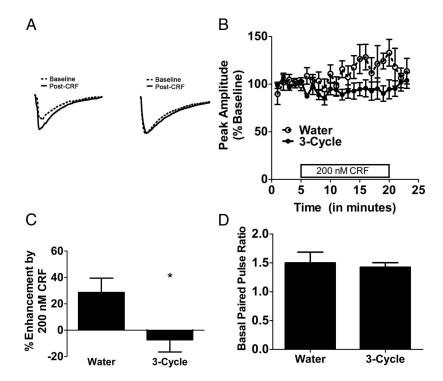
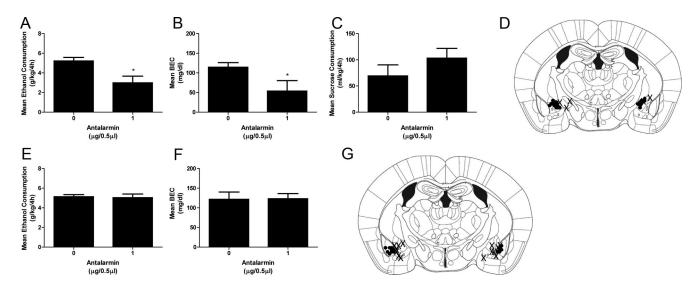


Figure 3. The modulation of GABAergic transmission by CRF in the CeA is disrupted 18-24 h following binge-like ethanol consumption. $\textbf{\textit{A}}$, Representative traces show that CRF significantly enhanced elPSCs relative to baseline IPSCs in a slice from the water group (left), an effect which was abolished in a slice from the 3-cycle group (right). $\textbf{\textit{B}}$, Bath application of CRF to slices containing the CeA enhanced mean elPSC peak amplitudes (percentage of baseline) of the water group. This effect was abolished in the 3-cycle group. $\textbf{\textit{C}}$, Bath application of CRF enhanced elPSC peak amplitudes of the water group by >20%, but did not alter the elPSC peak amplitudes of the 3-cycle group. $\textbf{\textit{D}}$, The water and 3-cycle groups did not differ in basal paired-pulse ratio. All data are presented as means \pm SEM, significance was accepted at the p < 0.05 level, and n = 5-6 cells per group (recordings taken from a single cell from each animal). *, Significant differences from the water group.

onstrate that binge-like ethanol consumption, like dependence-like ethanol consumption (Funk et al., 2006), requires CRF signaling in the CeA and are the first to demonstrate a direct role for CRF1Rs in this region in ethanol consumption (Lowery and Thiele, 2010).

Intra-BLA microinjections of a CRF1R antagonist do not alter binge-like ethanol consumption

To determine whether the effects of microinjections of antalarmin into the CeA on binge-like ethanol consumption could have resulted from diffusion of the drug to other regions of the amygdala, we also assessed the effects of microinjections of antalarmin into the BLA, a region proximal to the CeA and dense in CRF1R expression (Hauger et al., 2006), on binge-like ethanol consumption. Average ethanol consumption and BEC data from day 4 are presented in Figure 4. The results of a one-way ANOVA indicated no significant main effect of drug treatment on binge-like ethanol consumption ($F_{(1,16)} = 1.124$, p = 0.306; Fig. 4E) or BECs ($F_{(1,16)} = 0.082$, p = 0.778; Fig. 4F). Cannula placements are shown in Figure 4G. These results suggest that the attenuation



of binge-like ethanol consumption by intra-CeA administration of antalarmin was not likely to be due to diffusion of the drug into the BLA.

Discussion

The results of the current study provide novel evidence that endogenous CRF signaling is recruited in the CeA during bouts of binge-like ethanol consumption. First, three selective CRF1R antagonists reduced binge-like ethanol consumption at doses that did not alter nonbinge-like ethanol intake. Second, a history of binge-like ethanol drinking was associated with a significant increase of CRF-IR in the CeA and in the VTA. Additionally, a history of binge-like ethanol drinking abolished the ability of exogenous CRF to increase GABAergic transmission in the CeA at a time point when endogenous CRF-IR was upregulated in the same region. Finally, antagonism of CRF1R in the CeA, but not in the BLA, attenuated binge-like ethanol consumption. These converging results show that binge-like ethanol consumption recruits CRF1R signaling in the CeA, which is necessary for maintaining excessive binge-like ethanol intake.

CRF1Rs and ethanol consumption

Previous investigations comparing the effects of CRF1R antagonists on binge-like ethanol consumption to their effects on nonbinge-like ethanol consumption (Sparta et al., 2008; Lowery et al., 2010) have shown that CRF1R antagonists reduce high levels of ethanol consumption without effects on moderate or low levels of ethanol consumption (for review, see Lowery and Thiele, 2010). Similarly, CRF1R antagonists attenuate elevated ethanol consumption stemming from procedures to induce dependence-like drinking (for review, see Lowery and Thiele, 2010), stress (Marinelli et al., 2007; Lowery et al., 2008; Sommer et al., 2008), and deprivation-induced drinking (Sparta et al., 2009). Additionally, CRF1R antagonists attenuate ethanol consumption by the selectively bred Marchigian Sardinian alcohol-preferring rats (Ciccocioppo et al., 2006; Gehlert et al., 2007). Thus, while these compounds are effective under conditions that

elicit high levels of ethanol consumption, CRF1R antagonists do not alter moderate levels of ethanol consumption by nonstressed, nonselected animals in which dependence-like drinking has not been established (for review, see Lowery and Thiele, 2010). Together, these data may suggest that high levels of ethanol consumption that promote pharmacologically relevant BECs (i.e., ≥80 mg/dl) recruit CRF signaling, while moderate or low levels of ethanol consumption (associated with low BECs) do not. Interestingly, the role of CRF in cocaine self-administration also appears to be intake-dependent, as escalated cocaine selfadministration by animals with a history of high cocaine intake was attenuated by CRF1R antagonists at doses that did not alter cocaine self-administration by animals with a history of low cocaine intake (Specio et al., 2008). Thus, the hypothesis that the recruitment of CRF is dependent on the amount of ethanol consumed during a given period of access may also extend to other drugs of abuse.

Ethanol's effects on CRF-IR

Ethanol-induced alterations in CRF signaling in regions such as the CeA have previously been noted. Specifically, previous investigations have demonstrated significantly augmented CRF mRNA in the CeA of animals immediately following chronic ethanol ingestion (Läck et al., 2005) and in ethanol-dependent animals during acute withdrawal (Sommer et al., 2008). Additionally, increases in extracellular CRF were suggested in the CeA of ethanol-dependent animals during acute withdrawal, as decreases in total CRF-IR, but no change in the total number of CRF-immunoreactive cell bodies, were observed relative to nondependent animals (Funk et al., 2006). Our data extend the current literature by demonstrating that CRF-IR in the CeA is altered by high levels of ethanol consumption in animals with little (i.e., one binge-like drinking cycle) or moderate (i.e., six binge-like drinking cycles) ethanol exposure. This is an important observation because it suggests that recruitment of CRF signaling in the CeA during early binge-like drinking episodes, before dependence, may trigger initial neuroplastic changes in this system.

Theoretically, these changes may become more robust and rigid with repeated binge-like drinking experiences, contributing to the transition to dependence. Though increases of CRF-IR may reflect increased CRF activity (via increased CRF synthesis and/or release) or decreased CRF activity (via decreased CRF release), our converging pharmacological and electrophysiological results strongly suggest that CRF activity is elevated during binge-like ethanol consumption.

We also observed changes in CRF-IR in the VTA following a single binge-like drinking cycle. Previous studies have implicated CRF signaling in the VTA in the modulation of stress-induced reinstatement to cocaine seeking (Wang et al., 2005, 2007), which may be due to CRF's neuromodulatory effects on dopaminergic and glutamatergic transmission within this region (Wise and Morales, 2010). Thus, administration of CRF or a CRF1R agonist directly into the VTA significantly reinstated cocaine-seeking by animals that had undergone long access to cocaine selfadministration (Blacktop et al., 2011). Notably, intra-VTA administration of CRF did not alter stress-induced reinstatement to cocaine-seeking by animals that had undergone short access to cocaine self-administration, suggesting that CRF of the VTA modulates drug-taking behavior only in animals with a history of high cocaine intake. As ethanol has known actions on dopaminergic and glutamatergic systems of the VTA (Stuber et al., 2008), in addition to the CeA, binge-like ethanol drinking may also be modulated by CRF in the VTA.

Ethanol's effects on CRF-induced GABAergic transmission in the CeA

In the CeA, CRF exerts modulatory effects on GABAergic signaling via CRF1Rs in a way that mirrors ethanol's effects on GABAergic transmission (Nie et al., 2009; Roberto et al., 2010). The current data and previous work (Roberto et al., 2010) show that CRF enhances GABAergic transmission by neurons of the CeA in ethanol-naive animals. Previous investigations have shown that relative to ethanol-naive controls, this effect is augmented by chronic intermittent exposure to alcohol vapor (Roberto et al., 2010). However, we found that a history of binge-like ethanol consumption abolished the ability of CRF to enhance GABAergic transmission. Relative to water-drinking controls, we observed a blunted response to bath-applied CRF in brain slices of mice that experienced three binge-like drinking cycles at a time point when they also had increased CRF-IR in the CeA. This may reflect homeostatic regulation of the CRF-GABA system in response to high synaptic CRF concentrations (e.g., internalization of CRF receptors). Alternatively, upregulated tonic activity of CRF may have occluded the effect of subsequently applied CRF. However, this is unlikely given that there was no difference in basal pairedpulse ratio between water-drinking mice and mice that experienced three cycles of binge-like ethanol drinking (which would be expected if CRF was already active) and the lack of an effect of the CRFR1 antagonist on evoked GABAergic transmission.

The divergent results of the current study and previous studies of animals exposed to ethanol vapor suggest that binge-like ethanol consumption engenders different patterns of GABAergic transmission than does ethanol dependence, evidenced by differences in responses to exogenous CRF in the CeA between animals that experienced procedures to induce dependence-like drinking and nondependent animals with a history of binge-like ethanol intake, and by the observation that a history of binge-like drinking did not alter baseline GABAergic transmission as has been described in vapor-exposed rats (Roberto et al., 2010). It is important to consider factors that may have contributed to the dif-

ferences between dependence-like and binge-like drinking models noted above. Few exposures to high levels of ethanol (as in the current study) may cause transient perturbations of the CRF system that return to a homeostatic set point, while many exposures to high levels of ethanol (as in studies of ethanoldependent animals) may cause long-lasting adaptations in the CRF system that reflect the establishment of an allostatic set point. Interestingly, the disruption of CRF signaling found in the present report is similar to what was seen in the closely related structure, the BNST, following chronic cocaine exposure (Nobis et al., 2011), as repeated cocaine exposure led to a transient disruption in CRFR1 modulation of excitatory transmission with no changes in basal synaptic properties. Also consistent with our data is a study demonstrating that stress leads to an internalization of CRFR1 receptor in the dorsal raphe (Waselus et al., 2009). It is important to note that while binge-like ethanol consumption and dependence-induced ethanol consumption appear to fundamentally differ in some aspects, behaviors require CRF signaling in the CeA.

Conclusions and future directions

The results of the current study suggest that CRF signaling in the CeA is critical for the expression of binge-like drinking behavior. CRF signaling in the CeA is recruited during excessive ethanol intake, before the development of dependence, and we speculate that plastic changes in CRF signaling within the CeA develop over the course of repeated binge-like drinking episodes, contributing to alterations in critical downstream targets like the BNST (Li et al., 2012), and ultimately to the transition to ethanol dependence. Future investigations will examine potential differences in the CRF systems of binge-like drinking relative to nonbinge-like drinking mice and will elucidate the effects of binge-induced modifications of CRF activity in the CeA on downstream targets. Our observations suggest that compounds aimed at CRF1R may have therapeutic value for treating excessive binge drinking, perhaps protecting vulnerable individuals from progressing to dependence.

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