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CRF modulates glutamate transmission in the central amygdala of naïve and ethanol-dependent rats

Florence P. Varodayan1,* , **Diego Correia**1,2,* , **Dean Kirson**1,* , **Sophia Khom**1, **Christopher S. Oleata**1, **George Luu**1, **Paul Schweitzer**1, and **Marisa Roberto**¹

¹Department of Neuroscience, The Scripps Research Institute, La Jolla, California, USA

²Department of Pharmacology, Universidade Federal do Paraná, Jardim das Américas, Curitiba, Paraná, CEP 81531-990, Brazil

Abstract

Corticotropin-releasing factor (CRF) signaling in the central nucleus of the amygdala (CeA) is hypothesized to drive the development of alcohol dependence, as it regulates ethanol intake and several anxiogenic behaviors linked to withdrawal. Excitatory glutamatergic neurotransmission contributes to alcohol reinforcement, tolerance and dependence. Therefore, in this study we used in vitro slice electrophysiology to investigate the effects of CRF and its receptor subtype $(CRF₁)$ and CRF2) antagonists on both evoked and spontaneous action potential-independent glutamatergic transmission in the CeA of naive and ethanol-dependent Sprague-Dawley rats. We found that CRF (25–200 nM) concentration-dependently diminished evoked compound excitatory postsynaptic potentials (EPSPs), but increased miniature excitatory postsynaptic current (mEPSC) frequencies similarly in CeA neurons of both naïve and ethanol-dependent rats, indicating reduced evoked glutamatergic responses and enhanced vesicular glutamate release, respectively. This CRFinduced vesicular glutamate release was prevented by the $CRF_{1/2}$ antagonist (Astressin B) and the $CRF₁$ antagonist (R121919), but not by the $CRF₂$ antagonist (Astressin 2B). Similarly, $CRF₃$ effects on evoked glutamatergic responses were completely blocked by CRF₁ antagonism, but only slightly decreased in the presence of the CRF₂ antagonist. Moreover, CRF_1 antagonism reveals a tonic facilitation of vesicular glutamate, whereas the $CRF₂$ antagonism revealed a tonic inhibition of vesicular glutamate release. Collectively our data show that CRF primarily acts at presynaptic CRF1 to produce opposite effects on CeA evoked and spontaneous glutamatergic release and that the CRF system modulates CeA glutamatergic synapses throughout the development of alcohol dependence.

Keywords

amygdala; glutamate; alcohol/ethanol; CRF; CRF receptor; electrophysiology

Corresponding author: Marisa Roberto, The Scripps Research Institute, Department of Neuroscience, SP30-1150, 10550 North Torrey Pines Road; La Jolla, California 92037; USA, Phone: 1-858-784-7262; Fax: 1-858-784-7405, mroberto@scripps.edu. *These Authors Contributed Equally

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1. INTRODUCTION

Alcohol dependence is a chronic relapsing disorder, defined by the emergence of a negative emotional state that is mediated by the recruitment of brain stress systems, including the central nucleus of the amygdala (CeA) (Koob, 2008; Koob and Le Moal, 2008; Roberto et al., 2012). The CeA is the major output relay of the amygdala complex, and serves as a neuropeptidergic hub of anxiety, stress, and addiction-related functioning. In particular, the stress peptide corticotropin-releasing factor (CRF) plays a critical role in several CeA-driven addiction-related processes, including ethanol consumption and the anxiogenic effects of drug withdrawal (Breese et al., 2005; Eckardt et al., 1998; Heinrichs et al., 1995; Koob, 1998; Koob and Le Moal, 2001; Pich et al., 1995; Rassnick et al., 1993; Roberts et al., 1996).

CRF and its receptors $(CRF_1$ and CRF_2) are widely distributed in several stress-responsive brains regions of many species (e.g., human (Charlton et al., 1997), rat (Fischman and Moldow, 1982), and mouse (Justice et al., 2008)), playing a critical role in integrating the body's overall response to stress (Koob, 1999). The peptide can be synthesized and stored at specific GABAergic and glutamatergic synapses (Cain et al., 1991; Valentino et al., 2001a; Valentino et al., 2001b), and is usually co-released with classical neurotransmitters (Partridge et al., 2016) in response to neuronal firing to modulate their synaptic effects (Rainnie et al., 1992; Yu and Shinnick-Gallagher, 1998). The CRF system can produce a variety of region-, cell type- and synapse-specific effects depending on the distribution of its signaling elements within local and regional circuits (Henckens et al., 2016a). Both $CRF₁$ and $CRF₂$ mRNA expression have been detected in the CeA (Van Pett et al., 2000), and their pre- vs. postsynaptic distribution is critical for the CRF-induced inhibition and facilitation of glutamatergic transmission within this nucleus (Liu et al., 2004; Liu et al., 2005). As CRF signaling is critical to the role of the CeA in anxiety and alcohol use disorders (Gilpin et al., 2015), understanding its regulation of CeA glutamatergic transmission and its potential neuroadaptation with ethanol dependence are important.

Glutamate is the major excitatory neurotransmitter in the central nervous system and its ethanol-induced dysregulation contributes to several addiction-related behaviors, including drug reinforcement, tolerance and dependence (Lovinger and Roberto, 2013). Although the CeA mainly (95%) comprises γ-aminobutyric acid (GABA) neurons, it receives numerous glutamatergic afferents from other important addiction-related regions, including the basolateral amygdala, thalamus and cortex (Krettek and Price, 1978; Pitkanen et al., 1995; Savander et al., 1995). We recently reported on the effects of CRF and a $CRF₁$ antagonist on spontaneous and evoked glutamatergic transmission in CeA neurons of Wistar rats and Marchigian Sardinian Preferring (msP) rats, a line genetically selected for excessive ethanol drinking and characterized by heightened activity of the $CRF₁$ system that mimics the postdependent phenotype (Ciccocioppo et al., 2006; Herman et al., 2016; Natividad et al., 2017). We found that CRF had mixed effects on spontaneous action potential-dependent and independent glutamate release in the CeA of Wistar rats (with mainly a CRF-induced increase in action potential-independent glutamate release). However, in the msP rats, CRF had mixed effects on action potential-independent glutamate release, but only increased action potential-dependent glutamate release. CRF also decreased evoked CeA glutamatergic

transmission in Wistar, but not in msP rats, though its site(s) of action $(CRF₁$ vs. $CRF₂$) remain unknown. We and others have also previously shown that ethanol dependence is associated with increased CRF influence over GABA release in the CeA (Roberto et al., 2010), and that systemic and intra-CeA CRF_1 blockade both prevent the development of excessive ethanol consumption (Roberto et al., 2010; Varodayan et al., 2017) and withdrawal-associated anxiety-like behavior (Rassnick et al., 1993). However, to our knowledge, CRF's regulation of CeA glutamatergic synapses has yet to be investigated after ethanol dependence.

In this study, we used an *in vitro* slice preparation to assess the effects of CRF and its receptor antagonists on both evoked and spontaneous action potential-independent glutamatergic transmission in the CeA of naive and ethanol-dependent Sprague-Dawley rats. We found that CRF acts predominantly at $CRF₁$ to differentially modulate both forms of glutamatergic transmission at CeA synapses, and that ethanol dependence does not alter this regulation by the CRF system.

2. MATERIALS AND METHODS

2.1 Animals and slice preparation

We used 107 adult male Sprague-Dawley rats (average weight 346 ± 10 g) from Charles River (Raleigh, NC), housed 2–3 per cage in a temperature- and humidity-controlled room on a 12-h light/dark cycle (lights on at 8:00 am) with food and water available ad libitum. We conducted all care procedures in accordance with the National Institutes of Health Guide for the Care and Use of Laboratory Animals and with the Institutional Animal Care and Use Committee (IACUC) policies of The Scripps Research Institute (TSRI).

2.2 Chronic Ethanol Treatment

We used the standard ethanol inhalation method of the TSRI Alcohol Research Center to induce ethanol dependence (Cruz et al., 2013; Kallupi et al., 2014; Roberto et al., 2010; Roberto et al., 2004a; Roberto et al., 2004b; Roberto and Siggins, 2006; Rogers et al., 1979). Briefly, ethanol-dependent rats (n=36) had their home cages placed inside ethanol vapor chambers, and were intermittently exposed (14 h on, 10 h off) to ethanol vapor for 5–7 weeks. We determined the blood alcohol levels (BALs) of the chronic ethanol-treated animals using weekly tail-blood samples. The target BAL range was 150–200 mg/dL and the mean BAL was 177±10 mg/dL. Naïve/control rats (n=71) were treated similarly, but exposed to air 24 h/day. On experiment days, chronic ethanol-exposed rats were maintained in the vapor chamber until preparation of the CeA slices in ethanol-free solutions. Thus, all electrophysiology experiments were performed in brain slices undergoing acute in vitro withdrawal (2–8 hours), as previously described (Kallupi et al., 2014; Roberto et al., 2010; Roberto et al., 2004a; Roberto et al., 2004b; Varodayan et al., 2017).

2.3 Slice preparation

We prepared CeA slices as previously described (Cruz et al., 2012; Herman et al., 2016; Roberto et al., 2003; Roberto et al., 2004a; Roberto et al., 2004b; Roberto and Siggins, 2006). Briefly, the animals were anesthetized with isoflurane (3%) prior to decapitation, and

300–400 μm coronal slices were sectioned on a Leica 1000S vibratome (Leica Microsystems, Buffalo Grove, IL).

2.4 Intracellular recording of evoked responses

CeA slices were incubated in an interface configuration for 15–20 min, and then completely submerged and continuously superfused (flow rate of $2-4$ ml/min) with 95% O₂/5% CO₂ equilibrated artificial cerebrospinal fluid (aCSF) of the following composition in mM: NaCl, 130; KCl, 3.5; NaH₂PO₄, 1.25; MgSO₄•7H₂O, 1.5; CaCl₂, 2.0; NaHCO₃, 24; glucose, 10. We recorded from a total of 150 CeA neurons (from the medial subdivision of the CeA) with sharp micropipettes filled with 3M KCl using discontinuous current-clamp mode (Kallupi et al., 2014). Data were acquired with an Axoclamp-2A amplifier (Axon Instruments, Foster City, CA) and stored for later analysis using pClamp software (Axon Instruments). We held most neurons near their resting membrane potential (RMP) and applied hyperpolarizing and depolarizing current steps (200 pA increments, 750 msec duration) to generate I–V curves. We evoked pharmacologically-isolated excitatory postsynaptic potentials (EPSPs) by stimulating locally within the CeA through a bipolar stimulating electrode and superfusing the slices with aCSF containing the GABA receptor blockers: 30 μM bicuculline (to block $GABA_A$ receptors) and 1 μM [1-(S)-3,4-dichlorophenyl)ethyl]amino-2-(S)-hydroxypropyl p - benzylphosphonic acid (CGP 55845A) (to block GABA_B receptors). At the end of the recording session we often added 30 μM 6,7-Dinitroquinoxaline-2,3-dione (DNQX) and 30 μM DL-2-Amino-5-phosphonopentanoic acid (DL-AP5) to the aCSF to confirm the glutamatergic nature of the EPSP. To determine the synaptic response parameters for each cell, we performed an input-output (I-O) protocol (Cruz et al., 2012; Kallupi et al., 2014; Roberto et al., 2010; Roberto et al., 2003; Roberto et al., 2004a) consisting of a range of five current stimulations (50–250 mA; 0.125 Hz), starting at the minimum current required to elicit an EPSP up to the strength required to elicit the maximum subthreshold amplitude. These stimulus strengths were maintained throughout the entire duration of the experiment.

We examined paired-pulse facilitation (PPF) in each neuron using paired stimuli at 50 and 100 ms inter-stimulus interval (Andreasen and Hablitz, 1994; Logrip M., 2017; Roberto et al., 2004b). PPF was calculated as the amplitude of the second EPSP over that of the first EPSP. The stimulus strength was adjusted such that the amplitude of the first EPSP was ~50% of the maximal amplitude determined by the I-O protocol. A drug-induced change in PPF reflects presynaptic effects such that an increase in PPF suggests a decrease in neurotransmitter (glutamate) release. All measures were taken before drug superfusion (control) and during drug superfusion (10–20 min).

2.5 Whole-cell patch-clamp recording of glutamate currents

CeA slices were sectioned and then incubated in oxygenated aCSF for 30 min at 37°C and then 30 min at room temperature. We recorded from 87 medial CeA neurons that were visualized with infrared differential interference contrast (IR-DIC) optics, a x60 water immersion objective (Olympus BX51WI, Tokyo, Japan) and a CCD camera (EXi Aqua, QImaging, Surrey, BC, Canada). Whole-cell voltage-clamp recordings were performed in gap-free acquisition mode with a sampling rate of 10 kHz and low-pass filtered at 10 kHz, using a Multiclamp 700B amplifier, Digidata 1440A and pClamp 10 software (Molecular

Devices, Sunnyvale, CA). Pipettes (3–7 MΩ; King Precision, Claremont, CA) were filled with internal solution (in mM): 145 Kgluconate; 0.5 EGTA; 2 MgCl₂; 10 HEPES; 2 Na-ATP; 0.2 Na-GTP. Spontaneous miniature excitatory postsynaptic currents (mEPSCs) were isolated with the GABA receptor blockers, 30 μM bicuculline and 1 μM CGP 55845A, as well as 0.5 μM tetrodotoxin (TTX). Neurons were clamped at −60 mV and the series resistance was monitored with a 10 mV pulse (experiments with a series resistance >15 M Ω or a >20% change were excluded).

2.6 Drugs

We purchased CGP 55845A and bicuculline from Sigma (St. Louis, MO, USA), DNQX and DL-AP5 from Tocris (Ellisville, MO), and tetrodotoxin (TTX) from Biotum (Hayward, CA). CRF, Astressin B (antagonist for both CRF_1 and CRF_2) and Astressin 2B (CRF₂ antagonist) were synthesized and provided by Dr. Jean Rivier at the Salk Institute for Biological Studies. $R121919$ (CRF₁ antagonist) was synthesized by Dr. Kenner Rice at the Drug Design and Synthesis Section, Chemical Biology Research Branch, National Institute on Drug Abuse, National Institutes of Health, Bethesda, MD. Bicuculline was first dissolved in dimethylsulfoxide (DMSO) before being added to aCSF superfusate (final concentration of 0.05–0.1% DMSO). Drugs were added to the aCSF from stock solutions to obtain known concentrations in the superfusate.

2.7 Data analysis and statistics

To analyze data acquired from intracellular recordings, we quantified synaptic responses by calculating the EPSP amplitude with Clampfit 10 software. For whole cell recordings, the mEPSC frequency, amplitude (only mEPSCs >5 pA were included), and kinetics were analyzed using Mini Analysis software (Synaptosoft Inc., Fort Lee, NJ) and visually confirmed. Average mEPSC characteristics were determined over 3–5 min of recording trace and a minimum of 50 events. To control for cell-to-cell variation in all baseline electrophysiology properties, drug effects were normalized to their own baseline prior to group analyses. We used GraphPad Prism 6 software (GraphPad Software, La Jolla, CA) for all statistical analysis of results.

Pooled data for each experimental condition were analyzed by one-sample t-test for individual means comparisons to baseline to evaluate single drug (e.g. CRF) effects and paired t-test for comparison of successive drug effects (e.g. CRF antagonist vs. CRF antagonist + CRF) within the same animal treatment group. To assess differences resulting from ethanol exposure (i.e. naive vs. ethanol-dependent rats) and drug effects between these groups, we used a t-test or ANOVA as appropriate. Additionally, the Bonferroni post hoc test was used to assess significance between treatments as appropriate. We accepted statistical significance at the $p<0.05$ level. All data are presented as mean \pm standard error of the mean (SEM). n is reported in the figure legends, and represents the cell number from a minimum of four animals (1 to 5 cells were recorded per animal).

3. RESULTS

3.1 Chronic ethanol exposure alters glutamatergic transmission at CeA synapses

To investigate whether chronic ethanol treatment altered baseline glutamatergic signaling in the rat CeA, we recorded both evoked (intracellular configuration with sharp pipettes) and spontaneous (whole-cell voltage-clamp configuration) glutamate transmission.

We recorded intracellularly from a total of 157 neurons in the medial subdivision of the CeA. The mean resting membrane potential (RMP) and input resistance for the 84 CeA neurons from naïve rats were −81±0.7 mV and 169±7 MΩ, and for the 73 CeA neurons from ethanol-dependent rats (cells recorded in ethanol-free conditions equivalent to early withdrawal) were -80 ± 0.6 mV and 172 ± 8 M Ω , respectively. We generated compound excitatory postsynaptic potential (EPSP) input-output (I-O) curves using 5 normalized stimulation intensities (Fig. 1A). A two-way ANOVA of stimulus intensity and ethanol treatment did not show a significant difference between the compound EPSP I-O curves in slices from naive and ethanol-dependent animals $(F(1,155)=1.42, p=0.2, n=157$ cells), indicating that chronic ethanol treatment did not alter evoked glutamate transmission in the CeA. There was a main effect of stimulus intensity $(F(4,620)=483.4, p<0.001)$, with a Bonferroni *post-hoc* test indicating EPSP amplitudes were significantly different from each other $(p<0.001$ per comparison) among the five intensities. Finally, there was no interaction effect between the stimulus intensity and ethanol treatment ($F(4,620)=0.96$, $p=0.4$).

We also examined paired-pulse facilitation (PPF), a phenomenon whereby a secondary synaptic response is influenced by a preceding primary stimulus of equal intensity (Andreasen and Hablitz, 1994; Manabe et al., 1993). Generally, changes in the PPF ratio (second EPSP/first EPSP) are inversely related to transmitter release, such that a reduction of the PPF ratio is associated with an increased probability of transmitter release (Roberto et al., 2004b). We examined the PPF at the half-maximal intensity determined from the I-O curve using inter-stimulus intervals (ISI) of 50 ms and 100 ms (Fig. 1B). In slices from naïve animals, the PPF was 1.45 ± 0.08 and 1.23 ± 0.04 for 50 ms and 100 ms ISI, respectively (n=46 cells). In slices from dependent animals, the PPF was 1.27 ± 0.07 and 1.20 ± 0.06 for 50 ms and 100 ms ISI (n=30 cells). An unpaired t-test between naïve and dependent animals found no differences between groups for both the 50 ms $(t(74)=1.64, p=0.1)$ and 100 ms $(t(74)=0.37, p=0.7)$ ISI.

Next, we determined the effects of chronic ethanol treatment on spontaneous glutamate transmission by studying miniature excitatory postsynaptic currents (mEPSCs). This vesicular form of glutamate release is recorded in the presence of TTX to pharmacologically block action potentials, and so is distinct from the evoked glutamatergic transmission (EPSP) that is produced by electrically stimulating the entire synaptic network (Atasoy et al., 2008; Kavalali, 2015). We found that ethanol-dependent rats had a significantly lower mEPSC frequency (t(85)=3.22, $p<0.01$, n=42 naïve cells, n=45 dependent cells) and amplitude (t(85)=2.64, p <0.01) than that of naïve rats, with no change in the rise or decay times (Fig. 1C). Specifically, in naïve rats the mean frequency was 0.90±0.13 Hz, amplitude was 26.1±1.0 pA, rise time was 1.23±0.06 ms and decay time was 0.90±0.11 ms, while in the ethanol-dependent rats, the mean frequency was 0.46 ± 0.06 Hz, amplitude was 22.9 ± 0.68

pA, rise time was 1.18±0.05 ms and decay time was 0.97±0.07 ms. As mEPSCs are action potential-independent, increased frequencies reflect higher glutamate release probabilities, whereas altered amplitudes/kinetics indicate changed glutamate receptor sensitivity (De Koninck and Mody, 1994; Otis et al., 1994). Therefore, chronic ethanol decreased both spontaneous glutamate release and postsynaptic glutamate receptor function at CeA synapses.

3.2 CRF decreases CeA evoked glutamatergic transmission similarly in naïve and ethanoldependent rats

We first assessed the effects of 4 concentrations of CRF (25, 50, 100, and 200 nM) on evoked glutamate transmission in the CeA of naïve rats. The histograms in Figure 2A express the peak CRF effects as a percent of baseline evoked EPSP amplitudes at half-max stimulus intensity obtained from the I-O relationship during CRF superfusion. The lowest concentration of CRF (25 nM) caused a non-significant decrease in EPSP amplitude $(90.0 \pm 5.2\%$ of control; t(7)=1.91, p=0.1, n=8 cells), while 50, 100 and 200 nM CRF significantly decreased the EPSP amplitude to $89.3\pm3.4\%$ (t(6)=3.19, p<0.05, n=7 cells), 89.6 \pm 2.7% (t(15)=3.91, p<0.01, n=16 cells) and 83.4 \pm 3.0% (t(6)=5.48, p<0.01, n=7 cells) of control, respectively. Overall, a one-way ANOVA demonstrated that there was no significant difference in the inhibitory effect of CRF among the different concentrations after 15 min of application $(F(3,34)=0.75, p=0.5)$. In this experiment, we also found that CRF, at all concentrations applied, did not significantly alter the neuronal RMP or input resistance.

We then tested whether chronic ethanol exposure altered the effects of CRF (25–200 nM) on CeA evoked glutamate transmission. All 4 concentrations of CRF significantly decreased the EPSP amplitude at the middle stimulus intensity in the CeA of ethanol-dependent rats, similar to naïve animals (Fig. 2A). CRF reduced EPSP amplitudes to $88.7\pm4.0\%$ (t(6)=2.85, p<0.05, n=7 cells), 76.4±8.3% (t(5)=2.85, p<0.05, n=6 cells), 80.9±4.9% (t(8)=3.95, p<0.01, n=9 cells), and 87.1 \pm 4.5% (t(5)=2.86, p<0.05, n=6 cells) of control for 25, 50, 100, and 200 nM CRF respectively. A two-way ANOVA of CRF concentration and ethanol treatment showed no difference in the effect of CRF concentration $(F(3,58)=0.36; p=0.8)$ or ethanol treatment (F(1,58)=1.00; p=0.3), with no interaction effect (F(3,58)=2.08, p=0.1), suggesting that ethanol dependence does not alter the CRF-induced decrease in glutamatergic transmission in the CeA.

We also examined the effects of CRF on PPF in CeA neurons of naïve and ethanoldependent rats, and observed mixed responses (Fig. 2B and C). In the CeA of naïve rats, only 100 nM CRF at the 100 ms ISI produced a significant (t(12)=2.56, p<0.05, n=13 cells) change in PPF ratio, increasing it to $128.2 \pm 11.0\%$ of baseline (Fig. 2B). Additionally, oneway ANOVAs at 50 ms (F(3,29)=0.92, p=0.4, n=33 cells) and 100 ms (F(3,30)=0.61, p=0.6, n=34 cells) ISI found no significant effects of CRF concentration on PPF ratios.

Similarly, in the CeA of ethanol-dependent rats, only 50 nM CRF at 100 ms ISI caused a significant (t(4)=2.86, p<0.05, n=5 cells) increase (116.2 \pm 5.6% of baseline) in PPF ratio (Fig. 2C). Also, one-way ANOVAs of CRF concentrations at the 50 ms (F(3,22)=1.77, $p=0.2$, $n=26$ cells) and 100 ms (F(3,19)=0.80, $p=0.5$, $n=22$ cells) ISI revealed no effects on PPF ratios. Finally, we used a two-way RM ANOVA of CRF treatment and ethanol exposure

to examine whether the effects of CRF differed between naïve and ethanol-dependent animals. A two-way ANOVA of CRF concentration and ethanol treatment found no significant effects of CRF (F(3,51)=0.72, p=0.55, n=53 cells) or ethanol treatment $(F(1,51)=0.20, p=0.7)$ with no interaction effect $(F(3,51)=1.45, p=0.2)$ at the 50 ms ISI, and no significant effects of CRF ($F(3,55)=0.61$, $p=0.6$, $n=57$ cells) or ethanol treatment $(F(1,55)=0.75, p=0.4)$, and no interaction effect $(F(3,55)=0.46, p=0.7)$ at the 100 ms ISI. Therefore, CRF decreased evoked glutamate release at CeA synapses similarly in naïve and ethanol-dependent rats. Because 50, 100, and 200 nM CRF induced similar inhibitions of the evoked EPSPs, we used 100 nM CRF throughout the rest of the study.

3.3 CRF1 receptors mediate the CRF-induced decrease of evoked glutamatergic transmission

To assess whether endogenous CRF acts at CRF receptors to regulate evoked glutamate transmission, and to identify the specific CRF receptor(s) that mediates this effect, we used several pharmacological tools. We found that 200 nM Astressin B, a non-selective $CRF_{1/2}$ receptor antagonist (Liu et al., 2004), significantly decreased evoked EPSP amplitudes (to 76.8 \pm 4.2% of control baseline; t(7)=5.57, p <0.001, n=8 cells) in the CeA of naïve rats (Fig. 3A). In 7 of these neurons, we co-applied 100 nM CRF in the continued presence of Astressin B and found that CRF did not further change the evoked EPSPs (72.9±6.6% of control baseline; t(6)=0.43, p=0.7 by paired t-test between AB and AB+CRF data) (Fig. 3A). These data indicate that the $CRF_{1/2}$ system promotes excitatory transmission in the CeA by increasing glutamate release under basal conditions.

We next tested R121919, a selective antagonist for $CRF₁$, to probe for a potential role of this receptor in CeA evoked glutamate transmission. R121919 (1 μM) (Herman et al., 2016) significantly decreased evoked EPSPs $(86.4 \pm 2.7\%)$ of control at half maximal stimulus intensity; t(19)=4.96, p<0.001, n=20 cells; Fig. 3B and D). To investigate whether CRF decreased EPSPs in the CeA by activating $CRF₁$, we applied CRF in the presence of R121919 in 9 cells. We found that CRF no longer decreased the EPSP amplitude (R121919: 88.0±3.1% of control; R121919+CRF: 85.7±4.4% of control; t(8)=0.57, p=0.6; Fig. 3B), indicating that the inhibitory effect of CRF occurs mainly through CRF_1 .

We also tested whether selective $CRF₂$ blockade affects CRF-induced decreases in EPSPs. Application of the selective CRF2 antagonist Astressin 2B (200 nM) (Silberman and Winder, 2013) did not significantly affect evoked EPSPs $(92.3\pm3.7\%)$ of control; t(9)=2.08, p=0.1, n=10 cells) and the addition of CRF decreased EPSPs (to $84.7\pm4.7\%$ of control; t(9)=3.25, p<0.05; Fig. 3C and D). However, a paired t-test between the effects of Astressin 2B alone and Astressin 2B+CRF showed no significant difference in their effects on EPSPs (t(9)=1.65, p=0.1), indicating that CRF_2 may also partially contribute to CRF's inhibitory effects.

We next examined the effects of R121919 and Astressin 2B on PPF in naïve rats. R121919 increased the PPF ratio to $115.3\pm9.2\%$ of baseline for the 50 ms ISI and to $122.3\pm8.9\%$ of baseline for the 100 ms ISI (Fig. 3E). This increase in PPF ratio was significant at the 100 ms ISI (t(13)=2.50, p<0.05, n=14 cells) indicating a decrease of evoked glutamate release,

but not at the 50 ms ISI $(t(14)=1.66, p=0.1, n=15$ cells). Astressin 2B did not alter PPF ratios (Fig. 3E).

To determine whether chronic ethanol exposure altered the CRF system's regulation of evoked glutamatergic transmission in CeA neurons, we tested 1 μM R121919 and 200 nM Astressin 2B. The CRF₁ antagonist significantly $(t(8)=3.12, p<0.05)$ decreased the EPSP amplitude to $90.8\pm3.0\%$ of control (at half maximal stimulus intensity; n=9 cells; Fig. 3D), suggesting that endogenous CRF/CRF_1 signaling regulates CeA activity in ethanoldependent rats. This effect of R121919 was not significantly different between naïve and ethanol-dependent rats $(t(27)=0.96, p=0.3, n=29$ cells). Also similar to naïve rats, R121919 increased PPF ratios in the dependent rats to 149.7±32.1% of baseline for the 50 ms ISI and 125.2±16.5% of baseline for the 100 ms ISI, however, neither of these effects were significantly different from baseline (t(8)=1.55, p=0.2, n=9 cells and t(8)=1.53, p=0.2, n=9 cells for 50 and 100 ms ISI respectively; Fig. 3E). However, there was no significant difference between the effect of R121919 on PPF ratios between naïve and dependent rats for either the 50 ms (t(22)=1.26, p=0.1, n=24 cells) or the 100 ms (t(21)=0.17, p=0.9, n=23 cells) ISI. The CRF₂ antagonist did not significantly alter (t(6)=0.87, p=0.4, n=7 cells) the EPSP amplitude in the ethanol-dependent rats $(94.14 \pm 6.7\%)$ of baseline), and an unpaired ttest found no differences in the effects of Astressin 2B between naïve and dependent animals $(t(15)=0.27, p=0.8, n=17$ cells) (Fig 3D). Similar to CeA neurons of naive rats, Astressin 2B did not alter PPF ratios in 7 neurons of ethanol-dependent rats (Fig. 3E). Therefore, the CRF/CRF1 system continues to modulate evoked glutamatergic transmission in the CeA after ethanol dependence.

3.4 CRF increases action potential-independent glutamate release similarly in the CeA of naïve and ethanol-dependent rats

To further pursue the site of CRF's action at CeA glutamatergic synapses, we examined mEPSCs in neurons of naive and ethanol-dependent rats. We found that CRF (100 nM) increased the mEPSC frequency in CeA cells from naïve rats $(155.7\pm16.9\%; t(8)=3.29)$, $p<0.05$, n=9 cells; Fig. 4A); however, in 2/9 cells CRF had no effect or slightly decreased (<5% change from baseline) mEPSC frequencies, indicating that CRF does not enhance the glutamatergic input to all CeA neurons (similar to (Herman et al., 2016)). There were no CRF-induced changes in mEPSC amplitude or kinetics (Fig. 4A). Therefore, our data suggest that CRF acts presynaptically to increase vesicular (action potential-independent) glutamate release in the CeA of naïve rats. Similarly, CRF significantly increased the mEPSC frequency in cells from ethanol-dependent rats $(141.2 \pm 17.1\%; t(9)=2.41, p<0.05,$ n=10 cells), though it had no effect or a slight decrease on mEPSC frequencies in 4/10 cells and did not alter the overall mEPSC amplitude or kinetics (Fig. 4A). Therefore, CRF enhanced action potential-independent glutamate release at a subset of CeA synapses similarly in naïve and ethanol-dependent rats, without significantly affecting the region's glutamate receptor composition or expression.

3.5 CRF1 activation promotes action potential-independent glutamate release

We next assessed the effects of the $CRF_{1/2}$ antagonist Astressin B (200 nM) and found that it significantly decreased the mEPSC frequency (to $82.4 \pm 6.1\%$ of control; t(9)=2.89, $p \le 0.05$,

n=10 cells), but had no effect on the amplitude or kinetics (Fig. 4B). In 7 of these neurons, we co-applied CRF in the presence of Astressin B and found that there was no significant difference in mEPSC frequency between the antagonist alone and Astressin B+CRF (Astressin B: $80.5 \pm 8.4\%$ of control; Astressin B+CRF: $88.9 \pm 10.5\%$ of control; t(6)=1.61, $p=0.16$; Fig. 4C). There were also no drug-induced changes in the mEPSC amplitudes and kinetics. Overall, these data indicate that the CRF system tonically promotes excitatory transmission in the CeA by increasing glutamate release.

Similar to the effects of Astressin B on vesicular glutamate release, the $CRF₁$ antagonist R121919 (1 μ M) significantly decreased the mEPSC frequency (to 76.2 \pm 4.8% of control; t(16)=4.94, $p<0.05$, n=17 cells), with no effect on the amplitude or kinetics (Fig. 4D). In 7 of these neurons, we applied CRF in the presence of R121919 and there was no further change in mEPSC frequency (R121919: 85.1±4.7% of control; R121919+CRF: 80.0±5.6% of control; $t(6)=1.09$, $p=0.32$; Fig. 4E). However, CRF decreased the mEPSC amplitude $(t(6)=3.29, p<0.05)$ and decay time $(t(6)=3.65, p<0.05)$ in this R121919+CRF experiment (data not shown), suggesting that CRF may have additional postsynaptic effects that are not mediated by CRF_1 . Overall, the R121919 data indicate that CRF_1 mediates the tonic effects of endogenous CRF to promote CeA glutamate release, as well as the exogenous CRFinduced facilitation of glutamate release we observed in this region.

In a third set of experiments, we used the $CRF₂$ antagonist Astressin 2B. Notably, Astressin 2B alone significantly increased the mEPSC frequency (to $133.3\pm9.8\%$ of control; t(6)=3.39, $p<0.05$, n=7 cells), with no effect on the amplitude or kinetics (Fig. 4F), indicating that the basal activation of $CRF₂$ in the CeA limits glutamate release. Additionally, co-application of Astressin 2B and CRF in 6 of these cells led to a further significant increase in mEPSC frequency (Astressin 2B: $136.5\pm11.0\%$ of control; Astressin 2B+CRF: $173.0\pm20.6\%$; t(5)=2.82, p <0.05; Fig. 4G), with no change in the mEPSC amplitude or kinetics. Thus, CRF appears to activate both CRF_1 and CRF_2 to regulate vesicular glutamate transmission in the CeA, but its facilitation of glutamate release occurs predominantly via CRF1.

4. DISCUSSION

The amygdalar CRF/CRF_1 system plays a critical role in ethanol dependence (Gilpin et al., 2015; Koob, 1999; Koob and Volkow, 2010) and chronic ethanol treatment upregulates the expression of CRF and $CRF₁$ in the CeA (Merlo Pich et al., 1995; Roberto et al., 2010). Systemic CRF₁ antagonism prevented the escalated ethanol drinking of dependent rats (Roberto et al., 2010), and intra-CeA administration of a $CRF₁$ antagonist reduced both the ethanol consumption (Funk et al., 2006; Varodayan et al., 2017) and anxiety-like behavior (Rassnick et al., 1993) of these rats. Additionally, CRF acting at $CRF₁$ increased CeA GABAergic transmission in naïve rats, with a greater effect observed in ethanol-dependent animals (Roberto et al., 2010). Despite the key implications of this neuropeptide in the CeA with regard to alcohol-related behaviors, the present study is the first to our knowledge to determine whether ethanol dependence produces neuroadaptation within the CRF system to impact its neuromodulation of CeA glutamatergic synapse function.

Here we showed that CRF increases vesicular glutamate release at CeA synapses in both naive and ethanol-dependent animals, and that tonic $CRF₁$ activity enhances vesicular glutamate release, whereas tonic $CRF₂$ activity inhibits it. This CRF-induced vesicular glutamate release was prevented by $CRF_{1/2}$ and CRF_1 antagonists, but not by the CRF_2 antagonist. CRF decreases evoked glutamatergic responses, and this decrease was smaller in the presence of the CRF_2 antagonist, but blocked by CRF_1 antagonism. Therefore, despite the fact that CRF produces opposite effects on evoked and spontaneous glutamatergic transmission in the CeA of both naïve and ethanol-dependent rats, it acts predominantly via $CRF₁$ at these synapses.

We and others have previously shown that CRF modulates glutamate transmission in the CeA (Gallagher et al., 2008; Herman et al., 2016; Liu et al., 2004; Orozco-Cabal et al., 2006; Silberman and Winder, 2013). Consistent with our previous work in Wistar rats (Herman et al., 2016), we found here that CRF decreases evoked glutamatergic transmission in CeA neurons of naïve Sprague Dawley rats. This effect most likely results from decreases in evoked glutamate release, though we cannot rule out the possibility of CRF having postsynaptic effects across the synaptic network. Moreover, previous work by Liu et al. (Liu et al., 2004) reported a similar CRF-induced inhibition of basolateral amygdala glutamatergic afferents into the CeA (Liu et al., 2004), suggesting that CRF may modulate these projections in the present study. In contrast to these evoked glutamatergic responses, CRF increased the frequency of mEPSCs in the majority of CeA neurons both in the present study and our previous work using Wistar rats (Herman et al., 2016), indicating that it increases spontaneous action potential-independent glutamate release onto a subset of CeA neurons similarly in both naïve and ethanol-dependent rats. Notably, the two forms of glutamate transmission examined here (evoked and spontaneous action potentialindependent) are differentially regulated by CRF. Evoked neurotransmission results from the stimulation of the entire synaptic network to elicit widespread classical neural communication, whereas spontaneous action potential-independent neurotransmission reflects isolated synaptic communication that can maintain homeostasis and mediate plasticity at mature synapses (Kavalali, 2015). Thus, the differences we observed in CRF's modulation of spontaneous and evoked glutamatergic release are likely the result of distinct presynaptic neurotransmitter release mechanisms (e.g. vesicle fusion machinery, spatial segregation of the vesicles and/or vesicle populations) (Atasoy et al., 2008; Kavalali, 2015).

To better understand the mechanisms underlying the opposing actions of CRF at CeA glutamatergic synapses, we performed a series of pharmacology studies using its receptor antagonists. CRF and its G protein-coupled receptors $CRF₁$ and $CRF₂$ are widely expressed in the extended amygdala (Chalmers et al., 1996; Justice et al., 2008; Van Pett et al., 2000). In the CeA, $CRF₁$ activation increases anxiety-like responses (Muller et al., 2003; Smith et al., 1998; Zorrilla et al., 2002a; Zorrilla et al., 2002b), whereas CRF₂ activity has mixed behavioral effects that appear to be circuit- and synapse-specific (Radulovic et al., 1999). We identified dual effects of the CRF system on basal action potential-independent glutamatergic transmission; activation of $CRF₁$ promotes glutamate release under basal conditions, while activation of CRF_2 limits it. As the $CRF_{1/2}$ antagonist and the CRF_1 specific antagonist produce similar decreases in mEPSC frequency, endogenous CRF acts primarily at $CRF₁$ to increase glutamate release. $CRF₁$ also mediates the increased action

potential-independent glutamatergic release that we observed upon exogenous CRF application.

Regarding evoked glutamatergic responses, exogenous CRF acts primarily through CRF1, although CRF₂ may also be involved. Notably, the CRF₁ antagonist and CRF each inhibited evoked glutamate responses, which may seem counter-intuitive. We speculate that the heterogeneous expression of CRF_1 and CRF_2 at both pre- and postsynaptic sites (Gallagher et al., 2008; Liu et al., 2004; Orozco-Cabal et al., 2006), as well as the CRF system's selective modulation of specific CeA glutamatergic inputs as determined by the distribution of $CRF₁$ and $CRF₂$ (Gallagher et al., 2008; Liu et al., 2004), results in a predominant upstream network inhibition of evoked glutamate release that acts on distinct sites on our pool of stimulated presynaptic terminals. Accordingly, the Gallagher group (Gallagher et al., 2008; Liu et al., 2004; Orozco-Cabal et al., 2006) reported that CRF and urocortin I (a CRFrelated peptide that has a higher affinity for $CRF₂$ than $CRF₁$) have opposing effects on CeA glutamate transmission, potentially due to their different binding affinities for each CRF receptor subtype and the presynaptic $CRF₁$ vs. pre- and postsynaptic $CRF₂$ distributions of their main targets (but see also (Henckens et al., 2016a; Henckens et al., 2016b)). Thus, similar to how the CRF system's neuromodulation of CeA glutamate release varies based on its pre- and/or postsynaptic receptor expression, it likely also varies in terms of the origin of glutamatergic afferents and CeA target cell types.

Glutamatergic dysfunction is a prime molecular mechanism for many of the long-term behavioral effects resulting from chronic ethanol consumption (Lovinger and Roberto, 2013; Roberto et al., 2006; Roberto et al., 2004b), and the CRF system within the CeA is positioned to play a critical role in the emergence of the negative affect associated with dependence (Koob and Zorrilla, 2010). We have previously observed increased CRF and CRF1 expression in the CeA of ethanol-dependent rats (Roberto et al., 2010) and an increased CRF/CRF1 influence over CeA GABAergic transmission in ethanol-dependent rats, suggesting that these synapses may play an important role in the transition to alcohol dependence. However, a recent study reported decreased amygdalar $CRF₂$ density in ethanol-naïve rats selectively bred for their alcohol preference (P rats) compared to their non-preferring counterparts (NP rats) (Yong et al., 2014). Although we have studied the effects of acute ethanol on CeA glutamatergic signaling in naïve and ethanol-dependent rats (Roberto et al., 2006; Roberto et al., 2004b), to our knowledge, this is the first study exploring whether the effects of CRF on CeA glutamatergic transmission are altered by chronic ethanol exposure. Surprisingly, we found no differences in the CRF-induced glutamatergic responses in the CeA of naïve versus ethanol-dependent rats, perhaps due to compensatory mechanisms at these synapses. Therefore, our present work highlights the critical functional role of the CeA glutamatergic system in the maintenance of homeostasis during chronic ethanol exposure. Interestingly, CRF has been reported to potentiate CeA glutamate transmission after 2 weeks, but not 24 hours, of withdrawal from repeated cocaine exposure (Pollandt et al., 2006), highlighting the important contribution of both stress history and current stress levels in shaping the CRF system's neuroadaptation at CeA glutamatergic synapses (Henckens et al., 2016a; Henckens et al., 2016b).

CONCLUSION

The CeA CRF system has been hypothesized to drive the escalation of ethanol intake and the kindling of anxiety-like behaviors in dependent individuals (Gilpin et al., 2015; Koob and Volkow, 2010). We have previously shown that alcohol dependence is associated with increased CRF/CRF_1 influence over GABA transmission in the CeA, leading to greater local inhibition (Roberto et al., 2010). Here we found that CRF primarily acts at presynaptic $CRF₁$ to produce opposite effects on CeA evoked and spontaneous glutamatergic release in both naïve and ethanol-dependent animals. In addition, CRF also appears to have modest effects at CRF2, highlighting the complexity of the CRF system's neuromodulation of CeA glutamatergic synapses. This diversity in receptor type, localization and action allows greater flexibility for the CRF system to respond to different levels of stress, to both promote stress responses and to ensure stress recovery and the maintenance of homeostasis (Henckens et al., 2016a; Henckens et al., 2016b). As we observed no alteration in the CRF system's ability to regulate CeA glutamatergic synapses in ethanol-dependent animals, our data identify the CRF system as a sustained regulator of CeA glutamatergic synapses throughout the development of alcohol dependence.

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- CRF acts primarily via CRF₁ at central amygdala glutamatergic synapses.
- **•** CRF decreases evoked glutamate transmission at central amygdala synapses.
- **•** CRF increases spontaneous glutamate release at central amygdala synapses.
- **•** CRF affects glutamate transmission similarly in naïve and ethanol-dependent rats.

Figure 1. Chronic ethanol exposure alters CeA glutamatergic synapse function

A) Top: Representative traces of evoked CeA EPSPs at the five normalized stimulus intensities from a naïve and an ethanol-dependent rat (Dep). Bottom: I-O relationship curves are similar for naïve and dependent rats (84 cells from 61 naïve rats, 73 cells from 31 dependent rats). **B)** Top: Representative traces of PPF of EPSPs at the 50 ms ISI from naïve and dependent rat CeA neurons. Bottom: Histograms representing group PPF ratio for naïve and dependent rats at 50 ms and 100 ms ISI (46 cells from 34 naïve rats, 30 cells from 17 dependent rats). **C)** Left: Representative mEPSC traces from CeA neurons of naive and ethanol-dependent rats in baseline conditions. Middle and Right: Group comparison showed significantly lower baseline mEPSC frequencies and amplitudes in dependent rats compared to naïve rats (42 cells from 19 naïve rats, 45 cells from 10 dependent rats). ** $p \le 0.01$ by unpaired t-test.

Figure 2. CRF decreases evoked EPSP amplitude in CeA neurons of naïve and ethanoldependent rats

A) Top: Representative recordings of evoked EPSPs in CeA neurons from naïve rats before and during CRF application. Bottom: Histograms representing percent peak decrease in evoked (at half max stimulus intensity) EPSP amplitudes during superfusion of different concentrations (25, 50, 100 and 200 nM) of CRF on the CeA of naïve and ethanol-dependent rats (6–16 cells from at least 4 naïve or dependent rats per group). **B)** Top: Representative recordings of PPF at both 50 ms (top traces) and 100 ms (bottom traces) ISI in a CeA neuron from a naïve rat before and during superfusion of 100 nM CRF. Bottom: Histograms representing PPF ratio as a percentage of baseline after superfusion of 4 concentrations of CRF at 50 ms and 100 ms ISI in naïve rats. CRF significantly increased the PPF ratio of evoked EPSPs at the 100 ms ISI with 100 nM CRF **C)** Histograms representing PPF ratio as a percentage of baseline after superfusion of 4 concentrations of CRF at 50 ms and 100 ms ISI in ethanol-dependent rats. CRF significantly increased the PPF ratio of evoked EPSPs at the 100 ms ISI with 50 nM CRF. $p < 0.05$ and $* \cdot p < 0.01$ by one-sample t-test.

Figure 3. The CRF-induced decrease in evoked EPSP amplitude is mediated by CRF1

A) Top: Representative recordings of evoked EPSPs in a CeA neuron from a naïve rat recorded in control conditions, during Astressin B (AB) application, and during AB+CRF application. Bottom: Plot representing the percent peak decrease in evoked EPSP amplitudes for the middle three stimulus intensities during superfusion of AB and AB+CRF (7–8 cells from 7–8 naïve rats). CRF receptor antagonism decreases the evoked EPSP amplitude in naïve rats and blocks any further CRF-induced decrease. **B)** Top: Representative evoked EPSPs in a CeA neuron from a naïve rat recorded in control conditions, during R121919 (R12) application, and during R12+CRF application. Bottom: Plot representing the percent peak decrease in evoked EPSP amplitudes for the middle three stimulus intensities during superfusion of R12 and R12+CRF (9–20 cells from 9–17 naïve rats). CRF₁ antagonism decreases the evoked EPSP amplitude in naïve rats and blocks any further CRF-induced decrease. **C)** Top: Representative evoked EPSPs in a CeA neuron from a naïve rat in control conditions, during Astressin 2B (A2B) application, and during A2B+CRF application. Bottom: Plot representing the percent peak decrease in evoked EPSP amplitudes for the middle three stimulus intensities during superfusion of A2B and A2B+CRF (10 cells from 9 naïve rats). $CRF₂$ antagonism does not alter the evoked EPSP amplitude from baseline at the middle stimulus intensity in naïve rats, while co-application of CRF with A2B decreases the evoked EPSP. **D)** Histograms representing the percent peak decrease in evoked EPSP amplitudes for the middle stimulus intensity during superfusion of R12 and A2B show no difference between naïve and ethanol-dependent rats (Dep). **E)** Left: Representative PPF of EPSPs at 50 ms (top) and 100 ms (bottom) ISI in CeA neurons from an ethanol-dependent rat before and during superfusion of R12. Right: Histograms representing PPF ratio as a percentage of baseline after superfusion of R12 and A2B at 50 ms and 100 ms ISI in naïve and ethanol-dependent rats. R12 significantly increased the PPF ratio of evoked EPSPs for naïve rats at the 100 ms ISI. $*\infty$ 0.05 from baseline by one-sample t-test.

Figure 4. CRF1 activation promotes action potential-independent glutamate release A) Left: Representative mEPSC traces from CeA neurons of naive and ethanol-dependent (Dep) rats in baseline conditions and during 100 nM CRF superfusion. Right: CRF significantly increased mEPSC frequencies in the CeA of naïve and ethanol-dependent rats (9 cells from 4 naïve rats, 10 cells from 5 dependent rats). **B)** Left: Representative mEPSCs from CeA neurons of naive rats in baseline conditions and during 200 nM Astressin B (AB) superfusion. Right: AB significantly reduced the mEPSC frequency, but had no effect on amplitudes or kinetics (10 cells from 6 naïve rats). **C)** Co-application of CRF and AB did not significantly alter the mEPSC frequency compared to AB alone (7 cells from 5 naïve rats). **D)** Left: Representative mEPSCs from CeA neurons of naive rats in baseline conditions and during 1 μ M R121919 (R12) superfusion. *Right:* R12 significantly reduced the mEPSC frequency, but had no effect on amplitudes or kinetics (17 cells from 10 naïve rats). **E)** CRF+R12 did not significantly alter the mEPSC frequency compared to R12 alone (7 cells from 4 naïve rats). **F)** Left: Representative mEPSCs from CeA neurons of naive rats

in baseline conditions and during 1 μM Astressin 2B (A2B) superfusion. Right: A2B significantly increased the mEPSC frequency, but had no effect on amplitudes or kinetics (7 cells from 5 naïve rats). **G)** CRF+A2B significantly increased the mEPSC frequency compared to A2B alone (6 cells from 5 naïve rats). *p<0.05 by one-sample or paired t-test, as appropriate.