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## **OPIATE EXPOSURE AND WITHDRAWAL DYNAMICALLY REGULATE mRNA EXPRESSION IN THE SEROTONERGIC DORSAL RAPHE NUCLEUS**

**Jason Lunden**1,2 and **Lynn G. Kirby**1,\*

<sup>1</sup>Dept. Anatomy and Cell Biology and Center for Substance Abuse Research, Temple University School of Medicine, Philadelphia, PA, 19140, USA

## **Abstract**

Previous results from our lab suggest that hypofunctioning of the serotonergic (5-HT) dorsal raphe nucleus (DRN) is involved in stress-induced opiate reinstatement. To further investigate the effects of morphine dependence and withdrawal on the 5-HT DRN system, we measured gene expression at the level of mRNA in the DRN during a model of morphine dependence, withdrawal and post withdrawal stress exposure in rats. Morphine pellets were implanted for 72h and then either removed or animals were injected with naloxone to produce spontaneous or precipitated withdrawal, respectively. Animals exposed to these conditions exhibited withdrawal symptoms including weight loss, wet dog shakes and jumping behavior. Gene expression for brain-derived neurotrophic factor (BDNF), TrkB, corticotrophin releasing-factor (CRF)-R1, CRF-R2, GABAA- $\alpha$ 1, μ-opioid receptor (MOR), 5-HT<sub>1A</sub>, tryptophan hydroxylase2 and the 5-HT transporter was then measured using quantitative real-time PCR at multiple time-points across the model of morphine exposure, withdrawal and post withdrawal stress. Expression levels of BDNF, TrkB and CRF-R1 mRNA were decreased during both morphine exposure and following seven days of withdrawal. CRF-R2 mRNA expression was elevated after seven days of withdrawal.  $5-HT_{1A}$ receptor mRNA expression was decreased following 3 hours of morphine exposure, while TPH2 mRNA expression was decreased after seven days of withdrawal with swim stress. There were no changes in the expression of  $GABA_A-a1$ , MOR or 5-HT transporter mRNA. Collectively these results suggest that alterations in neurotrophin support, CRF-dependent stress signaling, 5-HT synthesis and release may underlie 5-HT DRN hypofunction that can potentially lead to stressinduced opiate relapse.

## **1.0 Introduction**

The serotonin (5-hydroxytryptamine, 5-HT) system plays an important role in stress-related psychiatric disorders and substance abuse (Charney et al., 1990; Meltzer, 1990; Valentino et al., 2010; Kirby et al., 2011; Waselus et al., 2011). Previous studies from our laboratory and others have shown that the serotonergic dorsal raphe nucleus (DRN) expresses receptors for the stress neuropeptide corticotrophin-releasing factor (CRF) (Swanson et al., 1983;

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<sup>\*</sup>Corresponding author: Lynn Kirby, MERB 857, Center for Substance Abuse Research, Temple University School of Medicine, 3500 N. Broad St, Philadelphia, PA 19140, Phone: 215-707-8556, Fax: 215-707-6661, lkirby@temple.edu.<br><sup>2</sup>Present address: Dept. Neuroscience and Cell Biology, Rutgers University - Robert Wood Johnson Medical School, Piscataway, 08854, USA

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Sakanaka et al., 1987; Chalmers et al., 1995) and is densely innervated by CRF terminals (Kirby et al., 2000; Valentino et al., 2001; Waselus et al., 2003). Both CRF and forced swim stress (FS) have the capacity to inhibit 5-HT output to specific forebrain regions (Price et al., 1998; Kirby et al., 2000; Kirby et al., 2008; Waselus et al., 2009; Valentino et al., 2010). Furthermore, CRF release within the DRN mediates the inhibitory effects of FS on 5-HT release (Price et al., 2002). Electrophysiology studies have demonstrated that intra-DRN CRF inhibits 5-HT neuronal activity (Price et al., 1998; Kirby et al., 2000), potentially via stimulation of GABAergic synaptic activity at 5-HT DRN neurons (Kirby et al., 2008).

In addition to an abundance of CRF receptors, the DRN expresses receptors for a number of other potential regulatory neuropeptides associated with drug abuse including opiates (Mansour et al., 1994; Neal, Jr. et al., 1999) and brain-derived neurotrophic factor (BDNF) (Merlio et al., 1992; Madhav et al., 2001). Opiate exposure has been shown to increase 5-HT output from the DRN through disinhibition of inhibitory GABA afferents caused by the activation of the  $\mu$ -opioid receptor (MOR) (Jolas and Aghajanian, 1997; Tao and Auerbach, 2002a; Tao and Auerbach, 2002b). Interestingly, during opiate withdrawal, 5-HT levels decrease below baseline (Tao et al., 1998), an effect caused by MOR-mediated stimulation of GABA synaptic activity at 5-HT DRN neurons (Jolas et al., 2000).

There is also a high degree of comorbidity of affective disorders including major depression with opioid dependence (Woody et al., 1975; Rounsaville et al., 1982; Brooner et al., 1997; Pani et al., 1997; Mason et al., 1998). Additionally, stress has been shown to be a potential trigger of opioid relapse in both the clinic and in animal models of drug abuse (de Wit H. and Stewart, 1981; de Wit H. and Stewart, 1983; Self and Nestler, 1998; De Vries and Shippenberg, 2002; Shaham et al., 2003; Brown and Lawrence, 2009). Thus the 5-HT system may contribute to negative affective states which underlie vulnerability to both mood disorders as well as drug abuse, potentially via overlapping neural mechanisms.

Recent work from our laboratory shows that CRF increases GABA synaptic activity in 5-HT cells within the DRN (Kirby et al., 2008). FS has been shown to inhibit 5-HT DRN activity by increasing CRF within the DRN (Price and Lucki, 2001). Furthermore FS can trigger reinstatement of previously extinguished morphine conditioned place-preference (Staub et al., 2012). Additionally 5-HT DRN neurons from animals exposed to both morphine and forced swim showed an increase in GABAergic synaptic activity (Staub et al., 2012). A more recent study from our lab shows that intra-DRN injections of GABA agonists can produce reinstatement of CPP without swim stress, while GABA antagonists can block swim stress-induced reinstatement of CPP (Li et al., 2013). These results underscore the importance of FS, CRF, and GABA in behavioral models of opiate relapse.

In the current study we hypothesize that the dynamic regulation of the 5-HT system during opiate exposure, withdrawal and stress-induced relapse is mediated by adaptations in neuropeptides and neurotransmitters within the DRN and that opiate history induces long term changes to the plasticity of the 5-HT DRN system. To test this hypothesis we measured the expression of genes that potentially regulate 5-HT DRN transmission at multiple timepoints in a model for opiate dependence, withdrawal and post withdrawal stress exposure. These time points included baseline, acute and chronic opiate exposure, precipitated withdrawal, spontaneous withdrawal, seven days post withdrawal and seven days post withdrawal with forced swim. While post withdrawal swim stress is not a model of stressinduced relapse, inclusion of this group in our model allows us to test the hypothesis that stress and opiate history interact, producing neuroadaptations within the DRN that could contribute to stress induced relapse.

Nine genes were chosen to study under the conditions of our model: BDNF, TrkB, CRF-R1, CRF-R2,  $GABA_A-a1$ , MOR, 5-HT<sub>1A</sub>, tryptophan hydroxylase2 (TPH2) and 5-HT transporter (SERT). BDNF (Boyarskikh et al., 2013) and its receptor TrkB (Merlio et al., 1992; Madhav et al., 2001) are known to be expressed within the DRN. Additionally intracerebroventricular BDNF administration has been shown to increase 5-HT activity (Siuciak et al., 1996). The receptors CRF-R1, CRF-R2,  $GABA_A$ , and MOR are known to affect 5-HT DRN activity under our experimental conditions (Jolas and Aghajanian, 1997; Kirby et al., 2008). The  $\alpha$ 1 subunit of the GABA<sub>A</sub> receptor was specifically chosen because it is expressed at high levels in the DRN (Sieghart and Sperk, 2002) and is the most abundantly expressed subunit in 5-HT DRN neurons (Lemos et al., 2011). Finally,  $5-HT<sub>1A</sub>$ , TPH2 and SERT are widely known to have more global effects on 5-HT regulation (Jacobs and Azmitia, 1992; Shishkina et al., 2007; Donner et al., 2012).

## **2.0 Materials and Methods**

## **2.1 Animals**

Male Sprague-Dawley rats (Taconic Farms, Germantown, NY) between 8–9 weeks of age were housed 2 per cage under standard temperature (20 °C) and humidity (40%). Rats were kept under a 12 h light/dark cycle (lights on at 7:00 AM). Food and water were provided ad libitum. Animal protocols were approved by the Temple University Institutional Animal Care and Use Committee and were conducted in accordance to the National Research Council Guide for the Care and Use of Laboratory Animals.

## **2.2 Surgery and post-operative care**

Prior to the start of experiments, rats were handled for 5 min over 2 days after which one rat per cage was assigned to either the placebo or morphine group. Before surgery two 2 cellulose placebo pellets or 2 morphine pellets (75 mg each pellet) (The National Institute on Drug Abuse (NIDA), Rockville, MD) were encased in a  $1 \times 1$ -cm square nylon mesh bag for each animal undergoing surgery. Rats were anesthetized with isoflurane after which nylon mesh bags containing either the placebo or morphine pellets were implanted subcutaneously in the scruff of the neck. In some groups, pellets were surgically removed 72 after implantation under isoflurane anesthesia. Rats were weighed and monitored daily for the duration of the experiment.

## **2.3 Animal behavior and experimental groups**

One baseline and six experimental groups were generated by euthanasia at different timepoints during a model of opiate exposure, withdrawal and post-withdrawal stress exposure (Figure 1) followed by tissue collection for PCR analysis. In addition to the baseline (untreated) group, the experimental group was implanted with placebo or morphine pellets and then sacrificed 3 hr post-pellet implantation (group 1, acute morphine), 72 hr post-pellet implantation (group 2, chronic morphine), 72 hr post-pellet implantation with naloxone injection (1mg/kg, subcutaneous injection) (group 3, precipitated withdrawal), 18 hr postpellet removal (group 4, spontaneous withdrawal), 7 days following pellet removal (group 5, abstinence) or 7 days following pellet removal with subsequent exposure to swim stress (animals placed in a swim tank that is 20 cm in diameter and filled with 21–22°C water to depth of 30 cm for 6-min) (group 6, abstinence + stress) to examine the effects of opiate history x stress interactions.

Animals undergoing withdrawal were recorded on video camera in a behavioral chamber and assessed for quantified behavioral signs of withdrawal including wet dog shakes, jumping, teeth chattering and writhing as well as the presence or absence of

chromodacryorrhea, eye-twitching, ptosis, rhinorrhea, salivation and diarrhea over a 15 min period, then euthanized 30 minutes later (Cerletti et al., 1975).

## **2.4 RNA processing**

Rats were sacrificed by decapitation and their brains were rapidly frozen on dry ice and stored at −80 °C for RNA extraction and real time PCR processing. Rat brains were sliced (200  $\mu$ m) in a cryostat at −20 °C and mounted on slides while RNA Later Ice (Invitrogen, Grand Island, NY) was added to maintain RNA stability. Following at least 24 hours of incubation with RNA Later Ice, 15 punches were collected from 5 sections spanning the DR using a 0.5 mm circular punch (Ted Pella, Inc., Redding, CA) (see Figure 2), pooled together and placed in TRIzol (Invitrogen) utilizing the small tissue extraction method supplied by Invitrogen. Briefly, tissues were added to  $800 \mu$  of TRIzol, and homogenized using an electric tissue homogenizer. Samples were spun down to remove cellular debris and genomic DNA was sheared by passing TRIzol solution through a 26 gauge needle before adding chloroform. RNA was then precipitated by addition of isopropanol and glycogen (GlycoBlue, Invitrogen). Samples were further cleaned through ethanol precipitation. Following RNA purification integrity of the samples was measured by spectrophotometry. All samples had A260/280 ratios between 1.8–2.0. Samples were run on a formaldehyde gel to confirm RNA integrity.

Following resuspension, 1 μg of total RNA went through a DNase I digestion protocol (Invitrogen) and was reverse transcribed using Affinity Script QPCR cDNA Synthesis Kit (Agilent Technologies, Santa Clara, CA). The cDNA samples were then diluted to a concentration of 2 ng/μl and stored for real time PCR.

## **2.5 Housekeeping gene selection**

Three potential housekeeping genes: β actin (actb, Rn00667869\_m1), GAPDH (Gapdh, Rn01775763\_g1), HPRT (Hprt1, Rn01527840\_m1) were tested at sample timepoints including 3 hours, 72 hours, precipitated withdrawal, and 7 days abstinence to assess consistency under the experimental conditions. Both Data Assist Software (Applied Biosystems, Foster City, CA) and REST Software (Qiagen, Germantown, MD) indicated that β actin showed the least differences between the treatment groups.

## **2.6 Real time quantitative PCR**

Real time PCR was performed using TaqMan gene expression assays with TaqMan universal PCR master mix (both Applied Biosystems). Assays utilized were: BDNF (Bdnf, Rn02531967\_s1), TrkB (Ntrk2, Rn01441749\_m1), CRF-R1 (Crhr1, Rn00578611\_m1), CRF-R2 (Crhr2, Rn00575617\_m1), 5-HT1A (Htr1a, Rn00561409\_s1), TPH2 (Tph2, Rn00598017\_m1), GABAA-α1 (Gabra1, Rn00788315\_m1), MOR (Oprm1, Rn01430371\_m1), SERT, (Slc6a4, Rn00564737\_m1) and β actin (Actb, Rn00667869\_m1). Each well consisted of 20 μl reaction mixture containing 10 μl master mix, 1 μl enzyme, 5 μl cDNA and 4 μl double autoclaved deionized water. Reactions were performed in triplicate using an ABI 7500 Real-Time PCR System (Applied Biosystems) under manufacturers recommended settings.

## **2.7 Data analysis**

Gene expression analysis was performed using the comparative CT (cycle threshold) method as described by Schmittgen and Livak (Schmittgen and Livak, 2008). A one tailed Student's T test was used for morphine to placebo comparisons as well as baseline to morphine comparisons, as all comparisons were a priori planned. Outliers were excluded based on the result of the Grubbs' test for outliers (Grubbs, 1969; Stefansky, 1972).

## **3.0 Results**

The hypothesis of the study is that the regulation of the 5-HT system across the opiate addiction cycle is mediated by adaptations in regulatory neurotransmitter and neuropeptide systems within the DRN. To test the hypothesis we quantified mRNA for genes that could potentially regulate 5-HT DRN neurotransmission within a model for opiate dependence, withdrawal, and stress exposure. All mRNA levels were normalized to a naïve baseline group or to placebo controls (Figs. 5–9; Table 1). The genes examined were BDNF, TrkB, CRF-R1, CRF-R2, 5-HT<sub>1A</sub>, TPH2,  $GABA_A$ -a1, MOR, and SERT.

## **3.1 Withdrawal syndrome**

For precipitated withdrawal, following 72 hours of placebo or morphine exposure the animals were weighed, given a single injection of naloxone (1 mg/kg), and observed for 15 minutes for withdrawal symptoms. All other animals had the placebo or morphine pellets surgically removed and 18 hours later were observed for withdrawal symptoms for 15 minutes. Animal experiencing both precipitated and spontaneous withdrawal experienced significant weight loss (precipitated:  $p < 0.01$ ; spontaneous:  $p < 0.01$ ) (Figure 3A). Both groups of animals were monitored for withdrawal behavior as well. In the precipitated and spontaneous withdrawal group, morphine-treated subjects showed a significant elevation of wet dog shakes compared to placebo controls (precipitated:  $p < 0.01$ ; spontaneous:  $p < 0.01$ ) (Figure 3B). In contrast, jumping behavior was seen in morphine treated animals undergoing precipitated but not spontaneous withdrawal ( $p < 0.05$  vs. placebo controls) (Figure 3C). Additionally, we observed the presence of diarrhea in both withdrawal groups as well as writhing within the precipitated withdrawal group. These data indicate somatic and behavioral signs of withdrawal in both withdrawal groups, though in precipitated withdrawal the behavioral signs are most prominent whereas in spontaneous withdrawal, somatic elements (i.e. weight loss) predominate.

#### **3.2 mRNA expression**

Utilizing all baseline groups, the mRNA of each gene was compared to CRF-R1, the lowest expressed gene. Generally we found that CRF-R1, CRF-R2, MOR and BDNF mRNA had low levels of expression. GABA<sub>A</sub>-α1, 5-HT<sub>1A</sub> and TrkB were expressed at middle levels and SERT and TPH2 mRNA were expressed most abundantly.

mRNA levels were measured at baseline, 3 hours and 72 hours after pellet implantation, following precipitated and spontaneous withdrawal, and 7 days following pellet removal with and without forced swim. Expression of BDNF mRNA was significantly decreased after 72 hours of morphine exposure compared to baseline (Figure 4D,  $p<0.05$ ). In contrast precipitated withdrawal increased BDNF mRNA compared to placebo (Figure 4B,  $p<0.05$ ) and then by 7 days following pellet removal, BDNF mRNA was downregulated in the morphine group compared to placebo (Figure 4C, p< 0.05). The BDNF receptor TrkB was significantly decreased compared to placebo after 3 hours of morphine exposure (Figure 5A, p<0.05) and again 7 days following morphine pellet removal compared to placebo (Figure 5C, p<0.05). TrkB mRNA was also elevated in the placebo group at 7 days post-pellet removal compared to baseline (Figure 5E, p< 0.05). CRF-R1 mRNA was decreased after 3 hours of morphine exposure (Figure 6A, p<0.05) and again 7 days following pellet removal compared to placebo (Figure 6C,  $p<0.05$ ). Seven days after morphine pellet removal, CRF-R2 mRNA was significantly increased compared to baseline (Figure 7D, p<0.01). The 5- $HT<sub>1A</sub>$  receptor mRNA was significantly decreased after 3 hours of morphine exposure compared to placebo (Figure 8A, p<0.01) and baseline (Figure 8D, p<0.01). 5-HT<sub>1A</sub> receptor mRNA was also elevated in the placebo group exposed to forced swim stress at 7 days post-pellet removal compared to baseline (Figure 8E, p<0.05). The TPH2 mRNA was

significantly decreased compared to placebo 7 days following morphine pellet removal in the forced swim group (Figure 9C,  $p<0.05$ ). These changes in mRNA expression are summarized in Table 1. All other mRNA expression data  $(GABA_A-1, MOR, and SERT)$  that were unaffected by the experimental conditions can be found in Table 2.

## **4.0 Discussion**

The main results of this study show that morphine dependence, withdrawal, and post withdrawal stress bring about differential regulation of mRNA expression for BDNF, TrkB, CRF receptors, and other genes regulating 5-HT in the rat DRN.

## **4.1 Baseline DRN mRNA expression**

We ranked mRNA levels in three distinct groups based on relative expression. Our results are consistent with published literature in many cases. CRF-R2 mRNA has previously been shown to be expressed at higher levels than CRF-R1 mRNA within the DRN (Van Pett et al., 2000; Day et al., 2004). BDNF mRNA has only recently been shown to be expressed in the DRN by quantitative PCR (Boyarskikh et al., 2013) as early in situ hybridization studies failed to measure expression in the raphe (Hofer et al., 1990; Wetmore et al., 1990; Gall et al., 1992), indicating that the high sensitivity of quantitative PCR is necessary to detect BDNF mRNA in the DRN. Genes for the 5-HT synthetic enzyme TPH2, the 5-HT transporter and the  $5-HT<sub>1A</sub>$  autoreceptor were, as expected, abundant in the serotonergic DRN.

## **4.2 Opiate regulation of BDNF and TrkB mRNA expression**

The neurotrophin BDNF and its receptor TrkB have been implicated in addiction-related neural plasticity in several monoaminergic nuclei (Koo et al., 2012; Mashayekhi et al., 2012) and reward-related brain regions (Berglind et al., 2007; Graham et al., 2007). More generally, BDNF activation of TrkB has been proposed to play a role in blocking the default pro-apoptotic pathway that is theoretically present in all neurons (Ichim et al., 2012). Additionally BDNF signaling is involved in neural differentiation and maturation (Waterhouse et al., 2012), regulating dendrite structure, modulation of synapse strength and circuit formation (English et al., 2012). Decreases in BDNF and TrkB activity have been associated with neuronal atrophy in both the hippocampus and frontal cortex as well as depression (for a review see (Gray et al., 2013) and (Duman, 2009) respectively). Our data show that following 3 hours of morphine exposure TrkB mRNA was decreased in the DRN relative to placebo, while a decrease in BDNF mRNA relative to baseline was seen after 72 hours. Interestingly a transient upregulation of DRN BDNF mRNA following precipitated opiate withdrawal was seen as well. While withdrawal-induced upregulation of BDNF and TrkB have been observed in other brain regions (Numan et al., 1998; Grimm et al., 2003; McGinty et al., 2010; Lu et al., 2010), the effect that we see in the DRN is less persistent. More experiments are needed to determine the functional implications of such a transient BDNF mRNA response.

Both chronic and repeated morphine exposure has been shown to reduce the number and complexity of dendritic spines in several reward-related brain regions including nucleus accumbens medium spiny neurons, medial prefrontal cortex and hippocampus pyramidal neurons (for a review see (Russo et al., 2009)). Opioid exposure has also been shown to impact morphology of neurons within the mesocorticolimbic dopamine system. Ventral tegmental area (VTA) dopamine neurons in rats treated with chronic morphine were shown to have a decreased area ( $\approx$ 20–25%) as well as reduced process length ( $\approx$ 30%), results that were prevented by BDNF pretreatment followed by administration of a mixture of BDNF and morphine (Sklair-Tavron et al., 1996). Both chronic morphine and heroin self-

administration in rats led to significant decreases in the surface area of VTA dopamine neurons (Russo et al., 2007). These morphological effects in VTA dopamine neurons persist when opioids are removed, lasting beyond the withdrawal phase. Fourteen days following opiate withdrawal, VTA neurons are reduced in size (Chu et al., 2007; Russo et al., 2007), an effect that is accompanied by a significant reduction in BDNF positive cells and VTA BDNF content (Chu et al., 2007). Collectively these other studies show that opiate exposure leads to long term changes in plasticity of limbic brain regions that are in part BDNF dependent. In a similar fashion, our data indicate that both BDNF mRNA and TrkB mRNA were downregulated compared to placebo seven days after the removal of morphine pellets. It is interesting to speculate that the decrease in neurotrophic support in the DRN following opiate exposure could potentially lead to a decrease in cellular surface area, process length or overall synaptic activity within DRN neurons. If so, these effects could potentially contribute to hypofunction of the 5-HT system that has been observed during protracted abstinence from chronic morphine (Goeldner et al., 2011) as well as susceptibility to stress induced reinstatement that we have observed in our previous studies (Staub et al., 2012; Li et al., 2013).

## **4.3 Opiate regulation of CRF receptor mRNA expression**

Anatomical data indicate that CRF axonal projections are present throughout the DRN and make contact with both GABA and 5-HT neurons (Kirby et al., 2000; Roche et al., 2003; Waselus et al., 2005). Electrophysiology studies further show that CRF stimulates inhibitory GABA synaptic activity at 5-HT DRN neurons at the pre- and postsynaptic levels with contributions from both CRF-R1 and –R2 receptor subtypes (Kirby et al., 2008). In the DRN of naïve animals CRF-R2 is expressed at higher levels than CRF-R1 (Van Pett et al., 2000; Day et al., 2004) and most of the CRF-R2 protein is intracellular leaving a higher functional ratio of CRF-R1/CRF-R2 on the cell surface (Waselus et al., 2009). Stress has been shown to induce a cellular redistribution of CRF receptor subtypes in DRN neurons, the majority of CRF-R1 receptors become internalized while CRF-R2 receptors are recruited to the membrane, effectively reversing the functional ratio of CRF-R1/CRF-R2 receptors on 5-HT DRN neurons (Waselus et al., 2009). Waselus et al (2009) further hypothesize that this functional CRF receptor redistribution underlies the switch from active to passive behavioral strategies to cope with the initiating stressor. Our results show that following 7 days of opiate abstinence CRF-R1 mRNA expression is decreased compared to placebo, while CRF-R2 mRNA expression is increased compared to baseline. These results indicate that this functional reversal of CRF receptors in the DRN may occur more ubiquitously, in this case produced by opiate history rather than by stress, and may occur at the level of the genes that encode CRF receptors as well as at the level of cell surface expression of the receptor protein (Waselus et al., 2009). It is also possible that this shift in CRF receptor mRNA expression in the DRN has similar behavioral consequences, predisposing subjects to passive, depression-like behaviors that might contribute to opiate relapse vulnerability. Previous studies from our lab (Staub et al., 2012) show that animals with an opiate history show sensitization of CRF-mediated inward current in 5-HT DRN neurons (a CRF-R2 mediated response; (Kirby et al., 2008) but CRF-R1-mediated stimulation of presynaptic GABA release that is normally seen in naïve subjects (Kirby et al., 2008) is absent in these animals. This opiate-induced shift in CRF receptor sensitivity of 5-HT DRN neurons is consistent with, and may be the result of the upregulation of the CRF-R2 mRNA and downregulation of the CRF-R1 mRNA that we observe in animals with opiate history in the current study. These alterations within the CRF stress system further support the hypothesis that opiate history produces an enduring change in plasticity of the 5-HT DRN system, potentially contributing to relapse susceptibility.

## **4.4 Opiate regulation of serotonin-related genes**

In the DRN the  $5-HT<sub>1A</sub>$  receptor is located on the soma and dendrites of  $5-HT$  neurons and effectively functions as an autoreceptor to maintain 5-HT homeostasis by regulating DRN electrical activity as well as 5-HT synthesis and release (Sprouse and Aghajanian, 1987; Hamon et al., 1988; Sharp et al., 1989). Expression of  $5-HT<sub>1A</sub>$  receptor mRNA was significantly decreased following 3 hours of morphine exposure compared to both placebo and baseline, yet remained at baseline for the duration of the experiment. Given that acute opiate exposure increases 5-HT output (Jolas and Aghajanian, 1997),  $5-HT<sub>1A</sub>$  mRNA downregulation may be an acute compensatory adaptation to elevated 5-HT levels. The net effect of this downregulation would be to further increase 5-HT levels. Therefore this mechanism may contribute to the overall stimulation of 5-HT in response to acute morphine exposure. The fact that the expression returns to baseline at 72 hours and remains steady for the rest of the experiment indicates that this  $5-HT<sub>1A</sub>$  mRNA response is transient and not be recruited beyond the acute response phase of the 5-HT system to morphine exposure.

Expression of TPH2 mRNA in contrast, was significantly decreased at seven days of abstinence with swim stress, but remained constant across all other time points. We have previously demonstrated that in animals with a morphine history, there is an interaction with swim stress leading to 5-HT hypofunction, an adaptation that may contribute to stressinduced opiate relapse in the CPP model (Staub et al., 2012). This adaptation was seen as an increased sensitivity of 5-HT DRN neurons (but not non 5-HT neurons) to GABAergic inhibition (Staub et al., 2012). Furthermore we have shown that intra-DRN injections of GABA agonists can produce reinstatement of CPP without swim stress, while GABA antagonists can block reinstatement of CPP following swim stress (Li et al., 2013). TPH2 is the main enzyme that drives 5-HT synthesis, and TPH2 mRNA is expressed almost 150 fold higher in the rodent brain stem compared to TPH1 (Walther and Bader, 2003). It is remarkable that TPH2 is steady across all non-stress time points examined suggesting that 5- HT synthesis within the DRN could be steady as well. Exposure to stress in subjects with an opioid history, in contrast, may decrease 5-HT synthesis. Furthermore the decrease in TPH2 mRNA is the opposite response seen in animals without exposure to opiates administered forced swim, as it has been demonstrated that forced swim increases TPH2 mRNA levels (Shishkina et al., 2008). These studies are consistent with our model that opiate history and stress interactions drive relapse through a decrease in 5-HT DRN activity. Results from the current study show that the interaction is not limited to GABAergic activity (Staub et al., 2012), as 5-HT synthesis may be altered as well. Finally this is a further example of the enduring plasticity in the 5-HT DRN system that results from an opiate history.

## **4.5 Gene expression changes in the placebo control group**

It is interesting to note that at two time points the placebo group significantly differed from baseline values (see elevated TrkB mRNA in 7-day group, Fig. 5 and elevated  $5-HT<sub>1A</sub>$ mRNA in the 7-day group exposed to FS, Fig. 8). The baseline group was included for comparison of all experimental groups to a naïve control group that had not experienced the stress of the surgery, anesthesia, etc. It is possible that in the placebo group, gene expression in these cases was upregulated in response to stress (related to surgery, anesthesia, and/or FS) but this effect was mitigated in the corresponding morphine group because the analgesic effects of the chronic morphine treatment may have reduced the stressfulness of these procedures (Buckingham and Cooper, 1984; Houshyar et al., 2001).

#### **4.6 Conclusions**

In this study we have shown that morphine exposure, withdrawal and post withdrawal stress regulate mRNA expression for BDNF, TrkB, CRF receptors, and other genes regulating 5- HT in the rat DRN. Following seven days of abstinence, we found decreases in BDNF, TrkB and CRF-R1, with an increase in CRF-R2. Additionally swim stress following abstinence decreased TPH2 mRNA. Our previous results indicate possible hypofunction of the 5-HT system in a model of stress-induced opiate relapse (Staub et al., 2012; Li et al., 2013). In this study we have identified differential mRNA expression patterns in neurotrophins, the CRF system and other aspects of 5-HT regulation that may contribute to this hypofunction and confer susceptibility to relapse.

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## **Highlights**

Morphine decreased BDNF, TrkB and 5-HT1A mRNA expression in the serotonergic DRN

Morphine decreased CRF-R1 and increased CRF-R2 mRNA expression in the DRN

Opiate history and swim stress interact to reduce DRN TPH2 mRNA expression

Opiate modulation of DRN circuits may promote 5-HT hypofunction

5-HT hypofunction may contribute to vulnerability to stress-induced opiate relapse



## **Figure 1. Experimental groups**

Gene expression at the mRNA level was assessed at the following seven time points: baseline, 3 hours of morphine exposure, 72 hours of morphine exposure, precipitated withdrawal (naloxone injection following 72 hours of morphine exposure), spontaneous withdrawal (18 hours following surgical removal of morphine pellets), abstinence (seven days following surgical removal of morphine pellets), and abstinence with forced swim (6 minutes of swim stress following seven days of abstinence). Placebo pellet controls were included for each experimental group.

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## **Figure 2. DRN tissue collection**

Brains were sliced into 200 μm sections on a cryostat and incubated for 24 hours in RNA Later ICE solution to preserve RNA integrity. Fifteen DRN tissue punches (500 μm diameter) from five brain slices (7.32–8.26 mm posterior to Bregma) were collected as indicated by grey circles (Paxinos and Watson, 2007). Tissue punches were pooled together placed immediately into TRIzol solution for RNA extraction.  $Aq = aqueduct; LPAG =$ lateral periaqueductal gray; VLPAG = ventrolateral periaqueductal gray; LDTg = lateral dorsal tegmental nucleus; DRD = dorsal raphe nucleus, dorsal part; DRV = dorsal raphe nucleus, ventral part; DRL = dorsal raphe nucleus, lateral part; PDR = posterodorsal dorsal raphe nucleus; Rbd = rhabdoid nucleus; mlf = medial longitudinal fasciculus; 4N = trochlear nucleus.

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#### **Figure 3. Withdrawal syndrome**

Physiological and behavioral withdrawal symptoms were measured in animals undergoing both precipitated withdrawal (PWD) and spontaneous withdrawal (SWD). For PWD, weights were recorded before and 30 minutes after subcutaneous naloxone injection and behavior was scored over 15 min following naloxone injection (n=4). For SWD animals, weights were recorded immediately and 18 hours after surgical pellet removal (n=23–24) and behavior was scored over 15 min at the 18-hour time-point (n=5). Data indicate mean  $\pm$ SEM. \*\* P < 0.01 vs. placebo; # P < 0.05 vs. PWD, ## P < 0.01 vs. PWD by Student's t-test. Lunden and Kirby Page 18



#### **Figure 4. BDNF mRNA expression**

BDNF mRNA was quantified relative to placebo (A–C), and baseline (D and E). No differences were seen at 3 hours and 72 hours of morphine exposure (A). Precipitated withdrawal elevated BDNF mRNA expression (B), while seven days of abstinence reduced BDNF mRNA expression (C). BDNF mRNA expression was also decreased following 72 hours of morphine exposure compared to baseline (D). There were no differences in the placebo groups compared to baseline (E). Data indicate mean  $\pm$  SEM. \* P < 0.05 vs. placebo;  $\# P$  < 0.05 vs. baseline by Student's t-test. Baseline data: BL (n=8); placebo: 3 H (n=6), 72 H (n=6), PWD (n=5–6), SWD (n=5–6), 7D (n=6), 7D+FS (n=6); 3 H (n=6); morphine: 3 H (n=6), 72 H (n=7), PWD (n=7), SWD (n=5), 7D (n=5), 7D+FS (n=6).



#### **Figure 5. TrkB mRNA expression**

TrkB mRNA was quantified relative to placebo (AC), and baseline (D and E). TrkB mRNA expression was decreased after 3 hours of morphine exposure; no differences were seen at 72 hours of morphine exposure (A). No differences were seen in either withdrawal group (B). TrkB mRNA expression was also decreased following seven days of abstinence (C). There were no differences in the morphine groups compared to baseline (D), but there was a significant increase in placebo at 7 days (E). Data indicate mean  $\pm$  SEM. \* P < 0.05 vs. placebo; # P < 0.05 vs. baseline by Student's t-test. Baseline data: BL (n=8); placebo: 3 H  $(n=6)$ , 72 H  $(n=6)$ , PWD  $(n=6)$ , SWD  $(n=5)$ , 7D  $(n=6)$ , 7D+FS  $(n=6)$ ; morphine: 3 H  $(n=6)$ , 72 H (n=7), PWD (n=7), SWD (n=5), 7D (n=5), 7D+FS (n=6).



## **Figure 6. CRF-R1 mRNA expression**

CRF-R1 mRNA was quantified relative to placebo (A–C), and baseline (D and E). CRF-R1 mRNA expression was decreased after 3 hours of morphine exposure; no differences were seen at 72 hours of morphine exposure (A). No differences were seen in either withdrawal group (B). CRF-R1 mRNA expression was also decreased following seven days of abstinence (C). There were no differences in the morphine or placebo groups compared to baseline (D and E). Data indicate mean  $\pm$  SEM.  $*$  P < 0.05 vs. placebo by Student's t-test. Baseline data: BL (n=8); placebo: 3 H (n=5), 72 H (n=5), PWD (n=6), SWD (n=5), 7D  $(n=6)$ ,  $7D+FS$   $(n=6)$ ; morphine: 3 H  $(n=5)$ ,  $72$  H  $(n=6)$ ,  $PWD$   $(n=7)$ ,  $SWD$   $(n=5)$ ,  $7D$   $(n=5)$ , 7D+FS (n=6).

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## **Figure 7. CRF-R2 mRNA expression**

CRF-R2 mRNA was quantified relative to placebo (A–C), and baseline (D and E). No differences were seen in any group with respect to placebo (A–C). CRF-R2 mRNA expression was significantly higher than baseline following seven days of abstinence (D). There were no differences in the placebo groups compared to baseline (E). Data indicate mean  $\pm$  SEM. ## P < 0.01 vs. baseline by Student's t-test. Baseline data: BL (n=8); placebo: 3 H (n=6), 72 H (n=6), PWD (n=6), SWD (n=5), 7D (n=6), 7D+FS (n=6); morphine: 3 H  $(n=6)$ , 72 H (n=6–7), PWD (n=7), SWD (n=5), 7D (n=6), 7D+FS (n=5–6).



## **Figure 8. 5-HT1A mRNA expression**

5-HT<sub>1A</sub> mRNA was quantified relative to placebo (A–C), and baseline (D and E). 5-HT<sub>1A</sub> mRNA expression was decreased after 3 hours of morphine exposure; no differences were seen at 72 hours of morphine exposure (A). No differences were seen in either withdrawal group or in abstinence with or without stress (B and C).  $5-HT_{1A}$  mRNA expression was also decreased following 3 hours of morphine exposure compared to baseline (D) and there was a significant increase in placebo at 7 days + FS compared to baseline (E). Data indicate mean  $\pm$  SEM. \*\* P < 0.01 vs. placebo; # P < 0.05 and ## P < 0.01 vs. baseline by Student's t-test. BL (n=8); placebo: 3 H (n=6), 72 H (n=6), PWD (n=6), SWD (n=4), 7D (n=6), 7D+FS (n=6); morphine: 3 H (n=5), 72 H (n=7), PWD (n=7), SWD (n=5), 7D (n=6), 7D+FS (n=6).



#### **Figure 9. TPH2 mRNA expression**

TPH2 mRNA was quantified relative to placebo (A–C), and baseline (D and E). TPH2 mRNA expression was not changed after 3 hours or 72 hours of morphine exposure (A). No differences were seen in either withdrawal group (B). TPH2 mRNA expression was decreased following seven days of abstinence with swim stress (C). There were no differences in the morphine or placebo groups compared to baseline (D and E). Data indicate mean ± SEM. \* P < 0.05 vs. placebo by Student's t-test. Baseline data: BL (n=8); placebo: 3 H (n=5–6), 72 H (n=5–6), PWD (n=6), SWD (n=5), 7D (n=6), 7D+FS (n=6); morphine: 3 H  $(n=5-6)$ , 72 H (n=6–7), PWD (n=7), SWD (n=5), 7D (n=5–6), 7D+FS (n=6).

#### **Table 1**

#### Summary of mRNA expression changes

#### **A. Direction of mRNA expression changes with respect to placebo**



#### **B. Direction of mRNA expression changes with respect to baseline**



Data represent the direction of mRNA changes compared to placebo (panel A) or baseline (panel B). Treatment groups include 3 hours of morphine exposure, 72 hours of morphine exposure, morphine precipitated withdrawal, seven days abstinence from morphine and seven days abstinence from morphine with forced swim.

#### **Table 2**

## GABAA-α1, MOR and SERT mRNA Expression



Data represent mean ± SEM. BL = baseline, FS = forced swim, MOR = μ-opioid receptor, PWD = precipitated withdrawal, SERT = 5-HT transporter, SWD = spontaneous withdrawal.