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Genetic and Pharmacologic Studies Towards Prevention of opioid use disorder

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Genetic and Pharmacologic Studies Towards Prevention of opioid use disorder

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Dissertation submitted
to the School of Medicine
at West Virginia University

In partial fulfillment of the requirements for the degree of

Doctor of Philosophy in
Cellular and Integrative Physiology

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ABSTRACT

Genetic and Pharmacologic Studies Towards Prevention of opioid use disorder Shane West Kaski

Endogenous opioids mediate both analgesic and affective responses to stress. While the mu opioid receptor (MOR) produces the reinforcing euphoric effects responsible for the high abuse potential of traditional opioid analgesics, the kappa opioid receptor (KOR) is hypothesized to mediate the dysphoric component of stress. With increasing rates of opioid addiction, extensive research efforts are focused on better understanding each of these systems and how they interact, with the goal of developing less addictive pain management strategies and better approaches to addiction treatment. With this in mind, we conducted both clinical and pre-clinical studies aimed at developing better treatment strategies for opioid use and the management of patients with opioid use disorder (OUD). First, we assessed a local cohort of patients with (OUD) for a collection of single nucleotide polymorphisms (SNPs) related to reward processing, with the goal of furthering efforts to develop a gene-based screening mechanism for OUD risk assessment. Second, we conducted preclinical assessments of the G protein-biased KOR agonist nalfurafine as a potential adjuvant medication for increasing the therapeutic efficacy of opioid-based analgesia. Genetic analysis of OUD patients identified two SNPs within the gene encoding the mu opioid receptor, one SNP within the gene encoding the serotonin 2B (5-HT_{2B}) receptor, and one SNP within the gene encoding Regulator of G protein Signaling 2 (RGS2), with the frequency of each SNP varying significantly from that observed in reference populations of European descent. Preclinical investigations with nalfurafine use in mice demonstrated a greater analgesic synergy when co-administered with morphine than morphine co-administered with the unbiased KOR agonist U50,488. As G protein-biased KOR agonists are hypothesized to produce less dysphoria (a therapeutically limiting side effect of KOR agonists), they may present a viable method for reducing the dose of MOR-targeting analgesic necessary for adequate pain relief, thereby reducing the likelihood of developing OUD. Further research is necessary to identify any anti-therapeutic effects of co-administering these two classes of drugs, as well as the range of pain modalities for which this approach is efficacious.

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Dedication

To my brother, whose kind heart and self-less nature drives my focus and dedication to helping those struggling with addiction.

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Abbreviations

5HT	Serotonin
ACC	Anterior Cingulate Cortex
ANOVA	Analysis of Variance
cAMP	Cyclic Adenosine Monophosphate
COAT	Comprehensive Opioid Addiction Treatment
CPA	Conditioned Place Aversion
CPP	Conditioned Place Preference
CREB	Cyclic-AMP Response Element Binding Protein
CRF	Corticotropin Releasing Factor
DMSO	Dimethylsulfoxide
DNA	Deoxyribonucleic Acid
DOR	Delta Opioid Receptor
EPM	Elevated Plus Maze
ERK	Extracellular Signal-Regulated Kinase
FDA	Food and Drug Administration
FST	Forced Swim Test
GABA	Gamma-Aminobutyric Acid
GDP	Guanosine Diphosphate
GPCR	G Protein-Coupled Receptor
GRK	G Protein-Receptor Kinase
GTP	Guanosine Triphosphate
icv	Intra-Cerebroventricular
it	Intra-Thecal
ICSS	Intra-Cranial Self-Stimulation
JNK	c-Jun N-Terminal Kinase
KO	Knockout
KOR	kappa Opioid Receptor
MAPK	Mitogen-Activated Protein Kinase
MAT	Medication Assisted Treatment
MDMA	Methylenedioxymethamphetamine
MOR	mu Opioid Receptor
NAc	Nucleus Accumbens
NIDA	National Institute on Drug Abuse
norBNI	nor-Binaltorphimine
ODD	opioid use disorder
PAG	Periaqueductal Gray
PAS	Photobeam Activated System
PDYN	Prodynorphin
RVM	Rostral Ventromedial Medulla
SERT	Serotonin Transporter
SNP	Single Nucleotide Polymorphism
SUD	Substance Use Disorder
VTA	Ventral Tegmental Area

Chapter One

1 - Introduction

1.1 - A Brief History of the Opioid Epidemic

Opium, an extract of the poppy plant, has left its mark on human culture since the plant's first recorded cultivation by the people of Sumeria in 3400 BCE (Rosenblum et al., 2008). As a powerful analgesic producing feelings of euphoria, it was used both medicinally and recreationally (Rosenblum et al., 2008). In the early 1800's Friedrich Serturmer isolated morphine, one of the active chemicals in opium (Wilkerson et al., 2016). Morphine saw extensive use during the American Civil War, with subsequent morphine addiction developing consistently enough that it earned the name "the army disease" (Courtwright, 1978). Not long afterward began the pursuit of a non-addictive substitute for morphine, with Charles Wright's synthesis of diacetylmorphine (Wilkerson et al., 2016). Today, diacetylmorphine is an illicit drug more commonly known as heroin and used by nearly 1 million people as reported in the 2016 National Survey on Drug Use and Health. Without a non-addictive replacement, opioid analgesics were largely avoided outside of acute and cancer-related pain until the 1980s, when a pair of publications claimed to demonstrate little risk of addiction when prescribed to pain patients without a history of drug abuse (Porter and Jick, 1980; Portenoy and Foley, 1986; Wilkerson et al., 2016). As a result, prescribing practices loosened and by the end of the 1990s pain was seen as "the fifth vital sign," with the management of chronic pain even becoming part of the accreditation process for hospitals (Phillips, 2000). Released in 1996, OxyContin® (extended release oxycodone) constituted a major portion of the increase in opioid prescriptions. The extended release formulation, aggressively marketed as less likely to produce addiction, was easily circumvented allowing for large, instant release doses of oxycodone (Remillard et al., 2019) and many consider OxyContin to be a central player in the rise of opioid addiction (Dyer, 2014; Jones et al., 2017). In 2017, the number of opioid prescriptions neared 200 million (CDC, 2017), while the number of opioid use disorder (OUD) diagnoses exceeded 2.1 million (SAMHSA, 2018).

Opioid use disorder is defined in the 5th edition of the Diagnostic and Statistical Manual of Mental Disorders (DSM-V) as the "...*compulsive, prolonged self-administration...*" of opioid drugs without a

legitimate medical purpose, or at higher doses than therapeutically required, despite “...*clinically significant impairment or distress.*” While this diagnosis is commonly thought of as addiction to the illicit drug heroin, roughly 1.67 million of the 2.1 million Americans diagnosed with OUD were using prescription opioids (SAMHSA, 2018). While the use of prescription opioids for the treatment of moderate to severe acute and cancer-related pain is regarded as standard and necessary (WHO, 1996; Christo and Mazloomdoost, 2008), their use in the management of chronic pain is contentious (Rosenblum et al., 2008; Salsitz, 2016).

Ironically, these long term outpatient prescriptions of opioids for chronic pain constitute the majority of the increase in opioid prescriptions (Boudreau et al., 2009; Gomes et al., 2011) and carry with them an increased potential for abuse (Shah et al., 2017; Barry et al., 2018; Brat et al., 2018). It is estimated that approximately 8-12% of patients on long term opioid therapy for chronic pain will develop OUD (Martell et al., 2007; Vowles et al., 2015; Volkow and McLellan, 2016) while closer to 25% will engage in “aberrant medication-taking behaviors” such as diversion of the medication to other individuals (Martell et al., 2007; Vowles et al., 2015; Volkow and McLellan, 2016). Addiction to and misuse of these prescription opioids is also linked to the use of heroin, with 80% of new heroin addicts reporting first abusing prescription opioids (Jones, 2013b). With such a strong correlation to prescription rates, it is hoped that alterations in prescribing practices will help curb the rise of OUD (Volkow and McLellan, 2016). A better understanding, however, of why some prove more susceptible than others to opioid abuse will be needed to inform truly responsible and effective prescribing guidelines.

1.2 - The Genetics of Opioid Addiction

Extensive investigation is underway to identify single-nucleotide polymorphisms (SNPs) that predict an individual’s likelihood of developing a substance use disorder (SUD) of any kind, including OUD. Currently, physicians do no better than chance at predicting which patients abuse opioids (Blum et al., 2018a) and current prescribing guidelines focus on generally reducing exposure and screening for a history of substance abuse (CDC, 2018). A better understanding of the genetics underlying OUD may lead to a standardized method of

screening patients before initiating opioid therapy to assess their likelihood of abuse and to tailor treatment for greatest efficacy. The genetics of SUD and psychiatric disorders in general, however, is complicated and now accepted to involve many genes, environmental factors, and gene x environment interactions. The Harvard Twin Study of Substance Abuse found that 34% of the variance in all substance abuse was attributable to additive genetic effects and that 43% of the variance in opioid abuse specifically was accounted for by additive genetic effects (Tsuang et al., 1998). This result is supported by at least one other twin study showing 48% genetic association of opioid addiction (Kendler et al., 2003). Though still in its nascency, ongoing research has identified several SNPs, found largely within genes expressed in limbic circuits, that correlate with rates of OUD and (in some combination) may account for this relationship.

1.2.1 - mu opioid receptor (MOR)

Of the predominant opioid receptors (mu, delta and kappa), MOR, encoded by the gene *OPRM1* is the primary site of action for opioid analgesics (Basbaum and Fields, 1984a; Pasternak, 1993) and is thus a likely site of variability in response to this class of drugs. Bond *et al.* identified the A118G SNP of the *OPRM1* gene (rs1799971) as having decreased expression and a greater affinity for its endogenous ligand, β -endorphin (Bond et al., 1998). In addition, Zhang et al. showed a reduced abundance of G allele mRNA in eight postmortem samples from subjects heterozygous for the A118G SNP, as well as reduced mRNA and protein levels in transfected Chinese Hamster Ovarian (CHO) (Zhang et al., 2005b). Human subjects carrying rs1799971 have shown increased odds for heroin addiction (Bart et al., 2004), increased nociceptive sensitivity (Tan et al., 2009), and decreased sensitivity to the analgesic effects of opioids in a post-operative setting (Chou et al., 2006). A meta-analysis by Haerian and Haerian of 13 studies with more than 9,000 subjects, however, identified no increased risk of opioid dependence among Caucasian or African-American subjects but an increased risk in Asian subjects with this specific SNP (Haerian and Haerian, 2013). While the A118G SNP remains the most well studied, other polymorphisms of MOR demonstrate significant associations with various parameters of OUD. The G-to-A transition of *OPRM1* SNP rs9479757 correlated with decreased severity of heroin addiction

in a Han Chinese population (Xu et al., 2014), while SNPs rs510769 and rs3778151 in Caucasians (Levrant et al., 2008), along with rs62638690 in people of European ancestry (Clarke et al., 2013), all affect susceptibility to heroin addiction. These results support further investigation of MOR polymorphisms as drivers of susceptibility to OUD

1.2.2 - kappa opioid receptor (KOR)

The kappa opioid receptor, encoded by the gene *OPRK1*, plays a strong role in the regulation of negative affective states (see section on *MOR, KOR and Addiction* below) and its endogenous ligand, dynorphin, is increased after exposure to opioid drugs (Wang et al., 1999). The *OPRK1* SNP rs6473797 demonstrated a reduced risk of heroin addiction in Caucasians (Levrant et al., 2008) while Yuferov and Kreek showed that the G36T transition of rs6473797 correlates with a higher susceptibility to heroin addiction in a mixed population (Yuferov et al., 2004). Evidence exists that at least one polymorphism in *OPRK1*, rs16918875, may increase the risk of heroin addiction in patients who also possess the A118G SNP of *OPRM1* beyond that conferred by either SNP alone (Kumar et al., 2012), hinting at a complex interplay of genetic susceptibilities underlying OUD. Various SNPs have also been identified in the gene encoding prodynorphin, the precursor of the endogenous KOR agonist dynorphin, *PDYN*. The majority of these *PDYN* SNPs, such as rs35286281 (Wei et al., 2011) and rs1022563 (Clarke et al., 2012), result in decreased expression of *PDYN* mRNA, as assessed both *in vitro* and in post-mortem analysis of human samples (Yuferov et al., 2004). These genetic variations in both *OPRK1* and the gene encoding its cognate agonist, *PDYN*, demonstrate the potential importance of this KOR signaling system in the development of addiction and thus is examined further in later sections.

1.2.3 - Delta Opioid Receptor (DOR)

Encoded by the gene *OPRD1*, the delta opioid receptor has been the subject of fewer studies identifying polymorphisms in human subjects with OUD, although preclinical data do support a role for DOR signaling in altering the rewarding properties of MOR targeting analgesics (Suzuki et al., 1996; Shippenberg et al., 2009). Two independent groups demonstrated correlations between *OPRD1* SNPs rs3766951 and rs2236857 and

heroin addiction in Caucasians (Levrán et al., 2008; Nelson et al., 2014) while a third group found a correlation between *OPRD1* SNP rs508448 and earlier age of heroin use in a population of Han Chinese (Gao et al., 2017). These initial population studies, along with the available preclinical evidence, support further investigation into variations in *OPRD1* and their relationships to the development of OUD.

1.2.4 - Serotonin Receptors (5HTR)

The serotonin (5-hydroxytryptamine) receptors are a large group, with 14 distinct receptors from seven different families (Barnes and Sharp, 1999); variations in the genes *HTR1B*, *HTR2A* and *HTR3A/B* (encoding the 5-HT_{1B}, 5-HT_{2A}, and 5-HT₃ receptors respectively) have shown correlations with heroin addiction. The G allele of the G861C variant of *HTR1B* (rs6296) correlated with heroin addiction, as well as amount of heroin-injection, in a study of a Han Chinese population (Gao et al., 2011), while a meta-analysis of studies assessing *HTR1B* SNPs for relationships with substance abuse found that rs130058 significantly correlated with heroin dependence in a mixed population (Cao et al., 2013). The A allele of the A1438G polymorphism (rs6311) of *HTR2A* correlates with heroin dependence in one study of a population in Spain (Saiz et al., 2008), while SNPs within *HTR3B* (rs3758987) (Levrán et al., 2008) and *HTR3A* (rs1176724 and rs897687) (Levrán et al., 2009) show correlations with heroin addiction in Caucasian and African-American populations, respectively. Although all these reports involve serotonin receptors, their individual biological functions vary significantly, and thus further research is needed to elucidate the role of each of these polymorphisms in increasing susceptibility to OUD.

Taken together, the studies outlined above support the genetic association seen in OUD, while at the same time make obvious the nascency of this line of research. Such factors as patient stratification, statistical analysis and lack of replication make it difficult to draw strong conclusions from the present reports. These findings, however, along with future advances in both genetic and epidemiologic methods, should encourage further efforts to identify predictive genetic factors for OUD.

1.3 - MOR, KOR, and Addiction

The opioid family of neuron signaling-inhibitory G protein-coupled receptors (GPCRs) not only includes the “classical” mu, kappa, and delta opioid receptors, but also the “non-classical” Nociceptin receptor. While all have been shown to modulate certain aspects of opioid-induced behaviors in rodent studies, MOR and KOR are thought to exhibit the strongest influence on the development of addiction (Koob and Le Moal, 1997; Crowley and Kash, 2015; Chavkin and Koob, 2016; Koob and Volkow, 2016). Opioid painkillers target primarily the mu opioid receptor, agonism of which produces both analgesia and euphoria (Koob and Le Moal, 1997; Koob and Volkow, 2016). Activation of the kappa opioid receptor produces analgesia as well (albeit more weakly), and carries with it clinically limiting effects, such as dysphoria and psychotomimesis (Pfeiffer et al., 1986; Siebert, 1994; Greenwald and Stitzer, 1998; Kivell and Prisinzano, 2010). The following sections outline the neurobiology of each of these opioid receptors as they relate to analgesia as well as the development and maintenance of opioid addiction.

1.3.1 - Analgesia

The clinical success of MOR-targeting analgesics arose from their activity at both spinal and supraspinal sites. At the level of the spinal cord, MOR localizes to both the presynaptic terminal of primary nociceptive afferents transmitting pain signals into the spinal cord (Fields and Basbaum, 1978; Pasternak, 1981; Moriwaki et al., 1996; Kline and Wiley, 2008; Todd, 2010b), as well as the post-synaptic cell bodies of secondary nociceptive efferents transmitting pain signals to higher brain centers (Moriwaki et al., 1996; Kline and Wiley, 2008; Todd, 2010b). MOR activation also modulates pain signals through its activity in the periaqueductal gray (PAG) matter (Basbaum and Fields, 1978; Yaksh et al., 1980; Basbaum and Fields, 1984b; Ghelardini et al., 2015b), a midbrain region that controls numerous functions, from fear and anxiety responses to blood pressure (Basbaum and Fields, 1978; Yaksh et al., 1980; Basbaum and Fields, 1984b; Behbehani, 1995; Ghelardini et al., 2015b).

The PAG modulates pain by controlling projections from midbrain nuclei to the dorsal horn of the spinal cord, a descending pathway that inhibits nociception (Basbaum and Fields, 1978; Yaksh et al., 1980; Basbaum and Fields, 1984b; Behbehani, 1995; Ghelardini et al., 2015b). This pathway begins in the periaqueductal gray, where glutamatergic neurons projecting to serotonergic and noradrenergic cell bodies in the rostral ventromedial medulla (RVM) (Basbaum and Fields, 1978; Yaksh et al., 1980; Basbaum and Fields, 1984b; Behbehani, 1995; Ghelardini et al., 2015b) remain under tonic inhibition by a local GABAergic circuit (Basbaum and Fields, 1978; Yaksh et al., 1980; Basbaum and Fields, 1984b; Behbehani, 1995; Ghelardini et al., 2015b). Activation of MORs located on the GABAergic neurons of the PAG leads to disinhibition of the glutamatergic projection neurons, in turn stimulating the serotonergic and noradrenergic neurons of the RVM (Pasternak, 2005b; Ghelardini et al., 2015b). These RVM neurons then project to the dorsal horn of the spinal cord, wherein release of serotonin and norepinephrine inhibit transmission of nociceptive signals (Pasternak, 2005b; Ghelardini et al., 2015b). In this way, MOR-targeting analgesics prevent the transmission of nociceptive signals directly, through actions at receptors in the dorsal horn, and indirectly through the modulation of regulatory projections.

The role of KOR signaling in producing spinal analgesia is similar to MOR activation outcomes, as the distribution of KOR resembles that of MOR (Besse et al., 1990; Todd, 2010b; Stein, 2016b) and KOR activation serves to inhibit the transmission of nociceptive signals through the dorsal horn (Todd, 2010a; Stein, 2016a). In the PAG and descending pain-inhibitory circuit, however, the role of KOR remains unclear. For example, one study found that microinjection of pentazocine, a KOR agonist, into the PAG produced analgesia against noxious heat but not against pressure, while fentanyl, a MOR-selective agonist, produced analgesia against both pain modalities (Llewelyn et al., 1983). A second group, however, used a thermal stimulus to show that morphine (a MOR-selective agonist), but not U50,488 (a KOR-selective agonist), produced an analgesic effect when microinjected directly into the PAG (Llewelyn et al., 1983). In addition to being a KOR agonist, pentazocine also acts as a partial agonist of MOR, while U50,488 acts more selectively at KOR and not MOR

(Clark and Pasternak, 1988; Hoskin and Hanks, 1991). This difference likely explains the disparate results between these two studies.

As a further complication, the distribution of MOR and KOR in these descending circuits remains unclear. While overlap may exist in their distribution, as reported in some histological studies (Gutstein et al., 1998), others demonstrate that MOR and KOR act in specific subpopulations of neurons in the RVM (Pan et al., 1997; Winkler et al., 2006), and that this distribution produces opposing analgesic effects (Pan et al., 1997; Tershner et al., 2000; Bie and Pan, 2003) (discussed further below). These discrepancies may also contribute to reported differences regarding receptor signaling effects on particular pain modalities. While less potent for somatic pain, KOR agonists display a greater potency for visceral pain in both preclinical studies (Su et al., 2002; Kamp et al., 2003) and in a clinical trial using a peripherally restricted KOR agonist (Arendt-Nielsen et al., 2009). As the underlying mechanisms and circuitry of somatic and visceral pain differ (Cervero and Laird, 1999), it is possible that these ambiguities in the distribution of each receptor play a role in their differing effects.

Finally, the variable of sex plays a large role in opioid analgesia. While it was long held that opioid analgesics produced greater analgesia in human females (Miaskowski and Levine, 1999), more recent studies complicate the issue, showing not only a greater degree of opioid-induced analgesia in males than in females, but also that these results vary depending on the opioid analgesic used, the type of pain, and the method used to assess pain/opioid efficacy (Gourlay et al., 1988; Cepeda and Carr, 2003; Miller and Ernst, 2004; Fillingim et al., 2005; Sibille et al., 2011; Lee and Ho, 2013). A 2010 meta-analysis of fifty studies found no association between sex and analgesia, although sub-group analysis found greater morphine-induced analgesia in women (Niesters et al., 2010). Studies of KOR-agonist induced analgesia produce the most reliable findings, repeatedly demonstrating greater analgesic efficacy in human females compared with males (Gear et al., 1996; Fillingim and Gear, 2004; Miller and Ernst, 2004). As a further complication, results from preclinical experiments on rats and mice show that both MOR-targeting and KOR-targeting opioid analgesics prove more efficacious in males than females (Lee and Ho, 2013; Abraham et al., 2018).

While both MOR and KOR produce analgesia through their activities in the spinal cord and descending pain-modulatory pathways, MOR agonists provide greater analgesia than KOR agonists due to their functional divergence in higher brain centers. Compared with KOR agonists, MOR-targeting analgesics retain broad applicability to nearly all types of pain, in part because of their ability to produce euphoria, thereby reducing the affective component of pain and preventing “agony” even when the conscious perception of a nociceptive stimulus remains (Zubieta et al., 2005; Porreca and Navratilova, 2017a). This ability, however, also leads to the dramatic abuse potential of MOR-targeting analgesics. In contrast, KOR agonism produces anxiety and dysphoria, limiting the clinical utility of KOR agonists and playing a major role in the cycle of addiction through its role in the production of negative affect. The following sections explore the effects of MOR and KOR agonism on affective state and motivation with respect to the development of addiction.

1.3.2 - Euphoria and Dysphoria

Both mu and kappa opioid receptors play central roles in the cycle of addiction. While MOR activation figures prominently in the initial development of drug preference through its euphorogenic properties (Koob and Le Moal, 1997; Koob and Volkow, 2016), KOR activation reinforces habitual drug taking through its dysphoric properties, producing a state of negative affect and “emotional pain” (Koob and Le Moal, 1997; Bruchas et al., 2010; Koob, 2013). In the CNS, these two receptors normally function in opposition to produce an affective homeostasis. When perturbed, however, this system may spiral to lower and lower set-points, leading to the destructive cycle of drug seeking known as addiction.

1.3.2a - Euphoria, reward and the mu opioid receptor

While MOR expression is distributed widely throughout the brain, the highest levels of MOR expression are found in regions involved in pain processing (notably the thalamus and periaqueductal gray matter) and motivated behavior (the mesolimbic system) (Le Merrer et al., 2009), with the latter playing the strongest role in the development of addiction. The endogenous ligand of the mu opioid receptor, β -endorphin, is produced in

the hypothalamus as well as the nucleus tractus solitarius (Froehlich, 1997), and is distributed throughout the brain as a homeostatic mechanism of reducing stress and producing reward, being released in response to novel stimuli as well as in times of stress and exercise (Rubinstein et al., 1996; Goldfarb and Jamurtas, 1997; Veening and Barendregt, 2015). As such, the mu opioid receptor system largely produces a positive affective response and motivates behavior through its effects on the mesolimbic system.

The mesolimbic pathway comprises dopaminergic projections from the ventral tegmental area (VTA) to the nucleus accumbens (NAc) and plays a key role in reward prediction, in addition to encoding the incentive salience of natural rewarding stimuli (Fields and Margolis, 2015). When encountering a rewarding stimulus, ranging from basic rewards like food to complex social rewards such as praise, burst firing of the VTA releases dopamine in the NAc and produces a positive reinforcement of the behavioral pattern that produced the rewarding stimulus (Bromberg-Martin et al., 2010; Salamone and Correa, 2012). VTA dopamine neurons remain under tonic inhibition by local GABAergic interneurons that express mu opioid receptors (Leone et al., 1991; Johnson and North, 1992). Endogenous opioid release in this area serves to inhibit these GABAergic interneurons, leading to disinhibition of VTA dopaminergic projections to the NAc, producing positive reinforcement (Leone et al., 1991; Johnson and North, 1992). Strong evidence supports this action on VTA mu opioid receptors as the primary mechanism of opioid analgesic induced reward (Leone et al., 1991; Fields and Margolis, 2015; Valentino and Volkow, 2018). Place preference for MOR-targeting analgesics is blocked in rats by administration of a MOR antagonist directly into the VTA (Olmstead and Franklin, 1997); conversely, MOR agonists are self-administered by rats when applied directly to the VTA (Bozarth and Wise, 1981; Zangen et al., 2002). While animals will self-administer MOR agonists applied to several other brain regions as well (David and Cazala, 1994), their actions in the VTA are widely regarded as necessary and sufficient for the reinforcing properties of opioid analgesics (Fields and Margolis, 2015). Furthermore, disinhibition of dopamine release likely plays a role in the ability of MOR-targeting opioid analgesics to reduce the affective experience of pain, as decreased function of midbrain dopaminergic neurons leads to increased pain (Saadé et al., 1997; Sophie and Ford, 2012), while dopaminergic agents can produce analgesia themselves. Mice lacking tyrosine hydroxylase

(dopamine deficient mice) show reduced thermal nociception in the tail withdrawal assay in response to morphine, supporting the role of dopaminergic signaling in the analgesic response to traditional opioid analgesics (Hnasko et al., 2005).

This canonical model, however, is not without its complications. Destruction of dopaminergic projections from the VTA to the anterior cingulate cortex (ACC), for example, blocks the acquisition of place preference to morphine administered to rodents systemically as well as directly into the VTA, implicating mesocortical circuitry in opioid reinforcement (Narita et al., 2010). In addition, while dopamine antagonists may block acquisition of morphine-induced place preference, the potency of this effect varies with the specificity of the antagonist for the various subtypes of dopamine receptor (Manzanedo et al., 2001). More confounding still are findings dopamine deficient mice (lacking functional tyrosine hydroxylase) still display a place preference to morphine (Hnasko et al., 2005), implicating factors beyond dopaminergic signaling in producing opioid reward.

While the intricacies of dopamine signaling in addiction require further inquiry, the initial phase of the development of opioid addiction -- acute reward and euphoria -- remain best explained by disinhibition of mesolimbic circuitry. The ability of opioid analgesics to potently disinhibit this circuitry underpins both their profound analgesic potential and their devastating propensity to incentivize compulsive use. As such a powerful circuit, the brain employs potent means of regulation, none linked more tightly to the cycle of addiction than the kappa opioid receptor.

1.3.2b - Dysphoria, altered set-points and the kappa opioid receptor

While the distribution of kappa opioid receptors mirrors that of MOR in many regions, oppositional localization in specific regions produces profound differences in the effect of KOR stimulation (Le Merrer et al., 2009). While stimulation of MORs produces euphoria, activation of KORs produces dysphoria, a negative affective state that includes components of anxiety, anhedonia, and depressive symptoms (Knoll and Carlezon, 2010; Lalanne et al., 2014; Crowley and Kash, 2015).

Dynorphin, a neuropeptide cleaved to six different lengths from the precursor prodynorphin and the endogenous ligand of the kappa opioid receptor, is produced in the hypothalamus (Froehlich, 1997), the amygdala (Marchant et al., 2007), and direct pathway GABAergic medium spiny neurons of the striatum (Steiner and Gerfen, 1998) (among other regions), and is released in response to stress (Knoll and Carlezon, 2010). Unlike β -endorphin, however, dynorphin plays the role of a negative homeostatic regulator. Dynorphin produces the negative experiences of stress (Land et al., 2008) and plays a role in mediating the central effects of corticotropin releasing factor (CRF), a major hormonal regulator of the stress response (Land et al., 2008). The areas most heavily implicated in the production of this negative state include the VTA-to-NAc circuit, along with the amygdala and dorsal raphe nucleus (DRN), the latter being the major serotonergic nucleus of the brain (Luo et al., 2015). In each of these regions, activation of KOR produces effects largely in opposition to those of MOR, culminating in the overall negative affective state of dysphoria.

In the mesolimbic system, KOR activation functions to reduce dopamine release, with KOR expression found on both the cell bodies of dopaminergic VTA neurons, as well as the presynaptic terminals of these

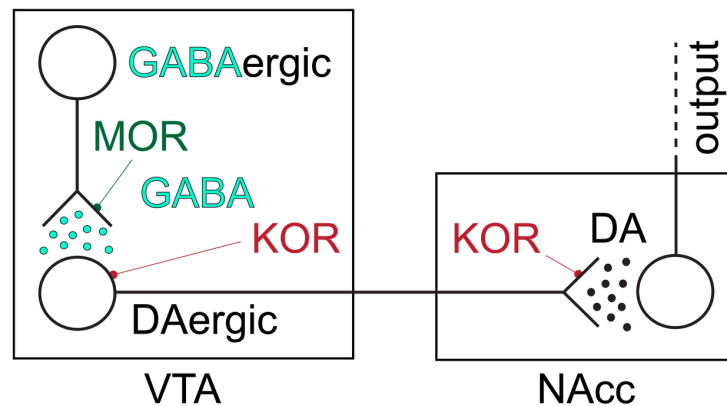


Figure 1.3.2b.1. Schematic of KOR and MOR receptors in mesolimbic pathway. Activation of MORs on GABAergic interneurons of the VTA disinhibits dopamine release, producing reward. Activation of KORs on dopaminergic neurons inhibits dopamine release, producing dysphoria.

neurons within the NAc (Figure 1) (Svingos et al., 2001; Chefer et al., 2013; Ehrich et al., 2015a). In both these

regions, KOR activation acts as negative feedback regulation on mesolimbic dopamine release, serving to restore homeostatic function within this circuit. When dopamine D1 receptor (D1R)-expressing neurons within the NAc are stimulated by dopamine, they release both dynorphin and GABA from axonal projections terminating in the VTA to reduce the firing rate of the dopaminergic neurons located therein (Nestler and Carlezon, 2006; Mysels and Sullivan, 2009). As disinhibition of this circuitry by MOR agonists produces a rewarding stimulus, inhibition by KOR agonists produces the opposite (*i.e.*, aversion, as measured in rodents in the conditioned place aversion (CPA) paradigm (Chefer et al., 2013; Ehrich et al., 2015a)); as a result, KOR agonists do not support self-administration in rats or rhesus monkeys (Tang and Collins, 1985; Serra et al., 2015; Ruan et al., 2016). Recent studies have strongly linked these aversive properties of KOR agonists to actions on mesolimbic dopamine neurons (Chefer et al., 2013; Ehrich et al., 2015a).

KOR actions in the amygdala appear to play a role in modulation of fear and anxiety related behaviors, possibly through modulation of local GABAergic neurotransmission and long-term potentiation (Huge et al., 2009; Kang-Park et al., 2013; Gilpin et al., 2014). Fear conditioning in rats is reported to increase *OPRK1* mRNA in this region, and local inhibition of KORs reduces this response (Knoll et al., 2011). In addition, KOR antagonism in the amygdala blocks the anxiogenic effect of a natural stressor (repeated forced swim stress) or intra-amygdalar injection of CRF in the elevated plus maze (EPM) assay (Bruchas et al., 2009). The DRN also plays a central role in mediating the affective response to KOR agonists. Stimulation of KORs in this region reduces extracellular serotonin concentration (Tao and Auerbach, 2005), likely through increasing surface levels of the serotonin transporter (SERT) via a p38 α MAPK dependent mechanism (Bruchas et al., 2011), an effect seen in mice after both local administration of a KOR agonist and social defeat stress (Bruchas et al., 2011). Furthermore, while the KOR agonist U50,488H did not produce a place aversion in mice lacking the kappa opioid receptor, viral re-expression of KORs on DRN neurons projecting to the NAc restored the effect in these KOR knockout mice (Land et al., 2009a).

The effects of KORs within these specific brain regions support the view that the dynorphin/KOR system plays a central role in mediating the aversive effects of stress (Land et al., 2009a; Bruchas et al., 2010;

Chavkin and Koob, 2016). Dynorphin levels increase due to stress (Nabeshima et al., 1992), and blockade or genetic deletion of *OPRK1* in mice prevents the pro-depressive and anxiogenic effects of forced swim, social defeat, and foot-shock stressors (Mague et al., 2003; Land et al., 2008; Bruchas et al., 2009; Bruchas et al., 2011). Systemic administration to rodents of KOR agonists produces a place aversion (Zhang et al., 2005a; Chefer et al., 2013) and depressive-like phenotypes in the forced swim and intra-cranial self-stimulation (ICSS) assays (Todtenkopf et al., 2004; Carlezon et al., 2006; Brust et al., 2016b). By producing these negative affective states, the dynorphin/KOR system also figures heavily into the addictive cycle. Stressful stimuli reinstate extinguished drug seeking behaviors in both place preference and self-administration paradigms (Mantsch et al., 2016). Pre-treatment of mice with the KOR antagonist nor-binaltorphimine (norBNI), as well as genetic deletion of either *OPRK1* or *PDYN*, prevented the reinstatement of cocaine conditioned place preference (CPP) by foot shock or forced swim stressors, as well as to a single administration of U50,488H (Redila and Chavkin, 2008). In addition, CPP for cocaine increased in mice exposed to forced swim stress prior to conditioning, an effect blocked by pre-treatment with norBNI (McLaughlin et al., 2003). While KOR antagonism also reduced ethanol self-administration by rats (Walker and Koob, 2008), the applicability to other drugs of abuse requires further investigation.

In light of its role in regulating limbic function and mediating stress responses, the kappa opioid receptor should be viewed as a key component of the “Allostatic Model” of addiction (Koob and Le Moal, 2001; Koob and Schulkin, 2018). This model contextualizes the state of addiction as one of a pathologically altered homeostatic set point, or “allostasis” (McEwen, 1998; Koob, 2008). The dynorphin/KOR system contributes to this allostatic response largely through modulation of dopamine tone within the mesolimbic system (Bruchas et al., 2010; Koob, 2013). As described above, MOR-targeting analgesics increase dopaminergic tone in the NAc, a property common to nearly all drugs of abuse (Wise, 2009; Koob and Volkow, 2016). Repeated use results in increased production of cAMP response element binding protein (CREB) within the NAc (Carlezon et al., 2005; Nestler and Carlezon, 2006) and subsequent upregulation of dynorphin and increased dynorphin tone (Nestler and Carlezon, 2006; Mysels and Sullivan, 2009; Bruchas et al., 2010). With cessation of opioid use, this

elevated dynorphin tone leads to a state of dysphoria, increasing the hedonic valence of the drug and increasing the susceptibility of relapse in an attempt to attain the now pathologically altered homeostatic set point (Bruchas et al., 2010; Koob, 2013; Koob and Schulkin, 2018).

In summary, while activation of both mu and kappa opioid receptors produces analgesia in response to stressful stimuli, the psychological functions of the receptors diverge dramatically and represent the opposing faces of opioid addiction. The desire of clinicians to alleviate pain placed the profound effects of MOR stimulation at the forefront of medicine in the early 2000s. Today, KOR antagonists sit among the National Institute of Drug Abuse (NIDA)'s ten most wanted drugs to help end the opioid crisis (Rasmussen et al., 2019), with the thought that the clinical application of KOR antagonism might reduce the contribution of stress, and its molecular and circuitry sequelae, in drug seeking and addiction reinstatement.

1.4 - Pharmacological approaches for improving opioid analgesics

The search for a non-addictive opioid analgesic began nearly the moment Friedrich Serturmer first isolated morphine in the early 1800's (Wilkerson et al., 2016), and the search regrettably continues today, over 200 years later. Many opioids are sold today as combination drugs, compounding opioids with non-steroidal anti-inflammatories like ibuprofen and acetaminophen as adjuvants meant to reduce the dose of opioids necessary to effectively treat pain, otherwise known as "dose sparing" (Remy, 2005; Maund et al., 2011). Other adjuvant therapies, such as suboxone, focus on preventing improper use of opioid medications by compounding them with opioid antagonists active only when taken through improper routes (Chen et al., 2014), but the effect of these measures are unclear. While many approaches are currently being investigated in labs across the world, from targeting little-understood splice-variants of the mu opioid receptor (Xu et al., 2017) to designing opioids that function only in the presence of inflammation (Spahn et al., 2017), the following sections focus on two widely studied and promising avenues: combination therapies activating multiple opioid receptors, and the development of "biased agonists."

1.4.1 - Combining MOR and KOR Activation

In the 1990s, researchers began investigating the possibility of utilizing the analgesic and non-reinforcing properties of KOR agonists to augment the effects of traditional MOR agonists. While KOR agonists alone proved less potent in producing analgesia (Pizziketti et al., 1985; Briggs et al., 1998), the combination of fentanyl (MOR agonist) with U69593 (KOR agonist) showed additive analgesic effects in rhesus monkeys while also reducing self-administration levels (Negus et al., 2008a). Further studies demonstrated that combined administration of MOR and KOR agonists prevented development of the clinical liabilities of opioid tolerance, hyperalgesia, and respiratory depression (He and Lee, 1997; Verborgh et al., 1997; Black and Trevethick, 1998; Field et al., 1999; Khotib et al., 2004)

While pre-clinical experiments demonstrated the utility of combined agonism of each of these opioid receptors, enthusiasm for this approach waned in the face of early clinical experiments utilizing mixed MOR/KOR active compounds. For example, while nalbuphine (a KOR agonist and MOR partial agonist) demonstrated reduced morphine-induced CPP and enhanced analgesia in pre-clinical studies (Tao et al., 2006), human studies demonstrated dose-dependent increases in dysphoria (Jasinski and Mansky, 1972; Preston et al., 1989a; Walsh and Babalonis, 2017). Also, pentazocine (similar to nalbuphine) and butorphanol (a KOR agonist and MOR antagonist) demonstrated the classic dysphoric effects of traditional KOR-targeting agonists in humans at higher doses (Preston et al., 1987; Zacny et al., 1994; Greenwald and Stitzer, 1998; Walsh et al., 2008; Walsh and Babalonis, 2017), limiting their therapeutic potential.

Of note, however, is one report (Preston and Bigelow, 1993) that co-administration of the opioid receptor antagonist naltrexone with pentazocine in human subjects “unmasks” the KOR-related dysphoric effects of pentazocine at lower doses, but blocks both MOR-dependent and KOR-dependent effects at higher doses. As the antagonist naltrexone possesses greater affinity for mu than kappa opioid receptors, these results of Preston & Bigelow support the premise that MOR agonism can mitigate KOR-related anti-therapeutic effects to some degree. While a single compound is generally considered a more attractive drug candidate than a dual-drug

formulation, these studies demonstrate that relative activity at each receptor plays a critical role in the interaction between these two receptor systems. Studies in patients receiving methadone maintenance therapy demonstrate that administration of nalbuphine, pentazocine, or butorphanol can precipitate a withdrawal syndrome (Preston et al., 1988; Preston et al., 1989b; Strain et al., 1993), indicating that none of these three, mixed-action drugs achieves activity at MOR comparable to methadone. While new mixed MOR/KOR drugs under development may possess more favorable activity profiles, such as the orvinols and 6 β -naltrexamines (Cami-Kobeci et al., 2009; Greedy et al., 2013), a co-administration paradigm may allow for greater control over relative activity at each receptor.

With these clinical studies dampening the potential therapeutic implications for KOR agonist development, research efforts dwindled. The rise of opioid addiction rates, however, has reinvigorated interest in the kappa opioid receptor. Cara Therapeutics invested heavily in developing a peripherally restricted KOR agonist, and its novel compound, CR845/difelikefalin is currently in Phase III clinical trials (Cara, 2018). In addition to peripheral restriction of KOR agonism (*i.e.*, to gain analgesic effect without concomitant CNS-centered dysphoric effects), major preclinical research efforts focus on utilizing a relatively new concept in pharmacology known as “biased agonism.” This approach utilizes variations in the efficacy of different compounds to engage various aspects of intracellular machinery, designing drugs that differ in their ability to produce canonical effects associated with receptor stimulation.

1.4.2 - Biased Agonism: All the good and none of the bad?

G protein-coupled receptors regulate an astoundingly diverse array of physiological processes and represent the target of approximately 37% of all drugs approved by the FDA as of 2017 (Hauser et al., 2017). Structurally, GPCRs contain seven transmembrane domains that couple to a heterotrimeric complex of G α , β and γ subunits on the intracellular face (McCudden et al., 2005). The G α subunit of this heterotrimeric complex associates with guanine nucleotide diphosphate (GDP) in the resting (inactive) conformation. When bound by agonist, the activated GPCR catalyzes the exchange of GDP for guanine nucleotide triphosphate (GTP), at

which point the $G\alpha$ subunit bound to GTP dissociates from the $G\beta\gamma$ obligate heterodimer (McCudden et al., 2005). Both the $G\alpha$ -GTP and $G\beta\gamma$ complexes initiate signaling cascades that vary depending upon the receptor and cellular context and are considered “G protein dependent” as a whole (McCudden et al., 2005). All opioid receptors couple to the $G\alpha i/o$ class of $G\alpha$ proteins. (Connor and Christie, 1999; Stein, 2016a). The $G i/o$ heterotrimeric complex, when activated (and separated) by GTP binding, generally functions to inhibit the production of cyclic adenosine monophosphate (cAMP), as well as increase potassium conductance (hyperpolarizing the cell) and inhibit calcium conductance (preventing vesicular release of neurotransmitters) (Pasternak, 2005a; Al-Hasani and Bruchas, 2011; Ghelardini et al., 2015a). Activation of opioid receptors inhibits neuronal function through these G protein-dependent mechanisms.

In addition to G protein-dependent signaling, work in the 1990’s demonstrated that GPCRs also initiate signaling cascades through a class of proteins known as β -arrestins (Wisler et al., 2014). Canonically, when a $G\beta\gamma$ subunit dissociates from $G\alpha$ -GTP upon agonist binding and receptor-catalyzed nucleotide exchange, it may activate a class of proteins known as G protein-receptor kinases (GRKs) (McCudden et al., 2005). This class of proteins serves to phosphorylate activated GPCRs, subsequently leading to the recruitment of β -arrestin (McCudden et al., 2005; Wisler et al., 2014). One of the functions of β -arrestin proteins is to act as a molecular brake on GPCR signaling, leading to the cessation of further G protein heterotrimer activation and subsequent receptor internalization. The internalized β -arrestin/GPCR complex then activates signaling pathways distinct from those activated by G protein subunits (Raehal, 2005; Wisler et al., 2014).

When a GPCR agonist compound activates either the G protein-dependent or β -arrestin dependent pathway to a greater extent than the other, this agonist is considered “biased” towards that signaling pathway (Wisler et al., 2014). Research in recent years demonstrates that the physiological and behavioral effects classically associated with stimulation of specific GPCRs may occur to a greater or lesser extent upon the application of biased compounds (Raehal, 2005; Ehrich et al., 2015b; Ewald et al., 2017b; Schmid et al., 2017b). The

possibility of discovering opioid drugs that lack traditional adverse effects has led to a surge in research focused on biased MOR and KOR compounds.

1.4.2a - G protein-biased MOR agonists

At the mu opioid receptor, G protein-dependent signaling produces the intended analgesic response (Todd, 2010a; Stein, 2016a), while β -arrestin signaling governs many of the unwanted side-effects. As might be predicted by the function of β -arrestin, mice lacking *BARR2*, the gene encoding β -arrestin2, displayed a decrease in the development of opioid analgesic tolerance and prolonged analgesic responses to morphine (Bohn, 1999; Bohn et al., 2000). In addition, *BARR2* knockout (KO) mice also display reduced respiratory depression and constipation after morphine administration (Raehal, 2005). Loss of β -arrestin2 did not, however, affect the development of naloxone-precipitated withdrawal symptoms (Bohn et al., 2000) or of morphine-induced hypothermia (Bohn, 1999). Pursuit of a G protein-biased MOR agonist led to the development of TRV130 (oliceridine), which shows potent analgesic effects and reduced tolerance development in rodents, yet retains the clinical liability of inhibitory effects on GI motility (Altarifi et al., 2017). Importantly, TRV130 also displays robust reinforcing effects, linking abuse liability to G protein-dependent signaling (Altarifi et al., 2017; Negus and Freeman, 2018). Despite these pre-clinical findings, TRV130 has proceeded to clinical trials as a potent opioid with reduced likelihood of requiring escalating doses due to a decreased propensity to induce tolerance.

A Phase I, double-blind, placebo-controlled study comparing morphine with TRV130 reported that the latter drug exhibited analgesic efficacy with reduced respiratory depression at equianalgesic doses, as well as a lower incidence of severe nausea (Soergel et al., 2014). While the FDA Advisory Committee stated that oliceridine demonstrated a “favorable benefit/risk profile” and proved superior to placebo in both APOLLO-1 and APOLLO-2 Phase III clinical trials (Research, 2018), the Anesthetic and Analgesic Drug Products Advisory Committee did not grant FDA approval to oliceridine, stating that “...oliceridine is a mu opioid agonist with an abuse potential, overdose potential and ability to produce physical dependence that is similar to other mu

opioid agonists” (FDA, 2018). As stated by this same report, “...it does not appear that the biased agonism of *oliceridine* with regard to preferential recruitment of G-protein over β -arrestin2 translates into a human safety advantage...” (FDA, 2018). Although the G protein signaling-biased TRV130 is only the first drug of its kind to undergo FDA review, these statements demonstrate not only the significant focus of the FDA on abuse potential with regard to opioid drugs, but also the difficulties of translating preclinical findings into human subjects.

1.4.2b - G protein-biased KOR agonists

At the kappa opioid receptor, G protein-dependent signaling also produces analgesic effects through inhibition of pain transmission through the dorsal horn of the spinal cord (Todd, 2010a; Stein, 2016a). In contrast, a strong arc of research implicates β -arrestin2 signaling in the therapeutically limiting dysphoric effects of KOR agonists (Bruchas et al., 2006; Bruchas et al., 2007a; Ehrich et al., 2015b). As discussed above, natural stressors such as forced swim, activate kappa opioid receptors in mice. It has been reported that pharmacological inhibition of p38, a mitogen activated protein kinase (MAPK) downstream of β -arrestin2 signaling at KOR (Bruchas et al., 2006), with SB203580, or preventing β -arrestin2 recruitment to activated KOR through gene deletion of the gene encoding the upstream GRK3 (*GRK3*), prevents the dysphoric component of this stress (Bruchas et al., 2007a). Furthermore, deletion of the gene encoding p38- α (MAPK14) from dopamine neurons prevents conditioned place aversion (CPA) in mice to the unbiased KOR agonist U50,488 (Ehrich et al., 2015b). These results and others have spurred investigation into a “non-dysphoric” replacement of traditional MOR-targeting opioid analgesics with a single KOR agonist biased toward G protein signal transduction.

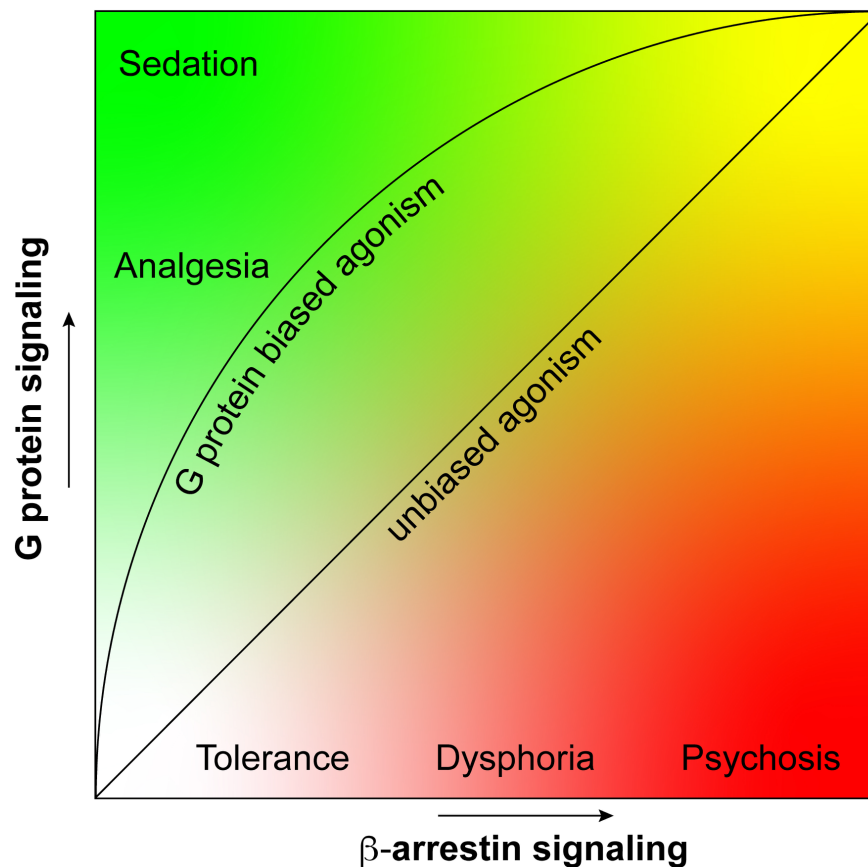


Figure 1.4.2b.1. Therapeutic potential of G protein-biased KOR agonists. Reducing signaling through β -arrestin may allow for the use of KOR agonists by reducing the incidence of therapeutically limiting side-effects.

The G protein-biased KOR agonist triazole 1.1 did not suppress ICSS thresholds, reduce NAc dopamine release, or inhibit novelty-induced locomotion at doses producing analgesia equivalent to U50,488 (Brust et al., 2016a). These results support the hypothesis that a G protein-biased KOR agonist would not produce the sedative and dysphoric effects that have limited traditional, unbiased agonist. Another G protein-biased KOR agonist, RB-64, however, produced CPA in both wild-type and β -arrestin2 KO mice without inhibiting novelty induced locomotion in either genotype (White et al., 2014). These conflicting results demonstrate the nascency of this line of research. First, more work is needed to determine if traditional KOR-related behavioral outcomes segregate to specific signaling pathways as cleanly as hypothesized. Second, two G protein-biased ligands (as assessed *in vitro*) may produce disparate results *in vivo*. While the current body of work focuses on G protein

and β -arrestin pathways, KOR ligands may possess biases in other signaling pathways such as extracellular signal-regulated kinase (ERK) and c-Jun N-terminal kinase (JNK) (Jamshidi et al., 2015) that could explain differences in their behavioral effects *in vivo*. In addition, differential expression of intracellular effector proteins within different brain regions has been shown to affect the regiospecific signaling efficacy of a biased agonist of the D2 dopamine receptor (Urs et al., 2016b), a process likely at work with KOR ligands.

Taken together, these studies reveal the complexities, and potential promise, of developing such signaling-selective ligands. Specific G protein-biased KOR agonists, such as triazole 1.1, may produce analgesia without producing sedative or dysphoric side effects, while others such as RB-64, may lack sedation but not dysphoria. A great deal more research into this field of behavioral pharmacology is required to disentangle the complexities of such biased ligands and their potential utility as replacements for traditional MOR-targeting analgesics.

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Chapter Two

Four single nucleotide polymorphisms in genes involved in neuronal signaling are associated with opioid use disorder in West Virginia

Original Article -in press *Journal of Opioid Management*

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2.1 - ABSTRACT

Objective: Pilot study to assess utility in opioid use disorder (OUD) of a panel of single nucleotide polymorphisms in genes previously related to substance use disorder (SUD) and/or phenotypes that predispose individuals to OUD/SUD.

Design: Genetic association study.

Setting: West Virginia University's Chestnut Ridge Center Comprehensive Opioid Abuse Treatment (COAT) clinic for individuals diagnosed with OUD.

Patients: Sixty patients 18 years of age or older with OUD undergoing medication (buprenorphine/naloxone)-assisted treatment (MAT); all sixty patients recruited contributed samples for genetic analysis.

Outcome Measure(s): Minor allele frequencies for single nucleotide polymorphisms.

Results: Four of the fourteen single nucleotide polymorphisms examined were present at frequencies that are statistically significantly different than in a demographically-matched general population.

Conclusions: For the purposes of testing WV individuals via genetic means for predisposition to OUD, at least four single nucleotide polymorphisms in three genes are likely to have utility in predicting susceptibility. Additional studies with larger populations will need to be conducted to confirm these results before use in a clinical setting.

Keywords: Opioid use disorder (OUD), substance use disorder (SUD), single nucleotide polymorphisms (SNPs), genetic testing

2.2 - INTRODUCTION

Opioid use disorder (OUD) has reached epidemic proportions in America (Rudd et al., 2016; McBain et al., 2018). Non-medical use of opioid analgesics and heroin is on the rise (Fischer et al., 2014; Jordan et al., 2017; Pergolizzi et al., 2018). Opioid overdose is now the leading cause of death of people under 50 years old (Dowell et al., 2017). Mindful of these developments, many addiction researchers are focusing combatting the epidemic with prevention and treatment (Marshalek and Sullivan, 2010; Schmid et al., 2017a; Vashishtha et al., 2017; Ratycz et al., 2018). A shared aspiration is to obtain an individual's genetic "fingerprint" and identify whether they are at increased risk for developing OUD (*e.g.*, ref. (Blum et al., 2018b)). Such testing is likely to interrogate several genes, as a multitude of factors (Lu et al., 2003; Badiani and Robinson, 2004; Scherbaum and Specka, 2008; Valentine and Fraser, 2008; Ducci and Goldman, 2012; Kreek et al., 2012; Nielsen and Kreek, 2012; Mistry et al., 2014; Bawor et al., 2015; Curtis et al., 2017; Crist et al., 2018; Fonseca and Torrens, 2018) contribute to OUD predisposition. Several gene variations are associated with predisposition to OUD and/or OUD treatment response (Kreek et al., 2012; Bawor et al., 2015; Curtis et al., 2017; Crist et al., 2018; Fonseca and Torrens, 2018). Environmental factors also contribute to OUD (Lu et al., 2003; Badiani and Robinson, 2004; Scherbaum and Specka, 2008; Valentine and Fraser, 2008), and some environmental factors can affect the influence of genetic factors on predisposition to developing addiction (Ducci and Goldman, 2012). The ability to identify *a priori* individuals at increased OUD risk would help greatly in prevention. For instance, using opioid analgesics with these individuals could be avoided or strictly monitored.

West Virginia is the epicenter of this opioid crisis, leading in opioid-related overdose deaths (Seth et al., 2018). The state is also demographically homogenous (Gall et al., 2016)*, being predominantly whites (94%) of European Caucasian ancestry (2010 US Census Data; <http://censusviewer.com/state/WV>). These factors make West Virginia ideal for genetic studies aimed at curbing the opioid epidemic. West Virginia University's Chestnut Ridge Center is home to a successful medication-assisted treatment program, known as the Comprehensive Opioid Addiction Treatment (COAT) clinic (Marshalek and Sullivan, 2010; Lander et al., 2013; Zheng et al., 2016; Zheng et al., 2017). Given access to this WV patient population, we sought to address

whether there are single nucleotide polymorphisms (SNPs) more or less prevalent in WV persons with OUD than in the general population.

2.3 – METHODS

2.3.1 - Recruitment

60 volunteers, diagnosed with OUD using DSM-V criteria (Association, 2013), were recruited from WVU's COAT clinic under IRB 1506733605. All participants were patients of the COAT clinic in Morgantown, West Virginia. The COAT clinic's service area (and hence our research subject area) is geographically diverse, as it draws from the entire state.

2.3.2 - Tissue Collection

Consenting volunteers provided buccal swabs, which were placed in 15-ml conical centrifuge tubes (Corning) and stored at -80°C.

2.3.3 - Genomic DNA Extraction

DNA was extracted using QIAamp DNA Mini Kits according to manufacturer's protocol. Yield was assessed by measuring 260-nm absorbance of DNA extracts on a QIAxpert microfluidic reader. Purity was assessed by measuring 260-nm and 280-nm absorbance; all samples had a 260-nm/280-nm quotient greater than 1.8 units.

2.3.4 - Genotyping

Genotyping was performed using TaqMan (FisherSci #4351379) primer-probe sets for each SNP (Table 1) and Type-it Fast SNP Probe PCR Kits (Qiagen) according to manufacturer's protocol. Samples were run on a Rotor-Gene Q (Qiagen) in duplicate, and no-DNA negative controls were also performed for each TaqMan primer-probe set.

2.3.5 - Statistical Analysis

Allele frequencies were compared to those of 1006 individuals of European descent (1000 Genomes Project (Genomes Project et al., 2015)); if no information was available from the 1000 Genomes Project, allele

frequencies were compared to ExAc database(Song et al., 2016) of ~60,000 genomes. Statistical analyses for differences of SNP variant frequency between groups were performed using Pearson's chi-square statistical test. The null hypothesis was that the allele frequencies from WV COAT clinic participants were the same as those from public databases. Therefore, $p\text{-value} < 0.05$ implies that the allele frequencies from WV COAT clinic participants are significantly different from the general public.

2.4 - RESULTS

A literature search for SNPs associated with OUD/SUD and/or related traits (*e.g.*, anxiety) identified fourteen of known or suspected relevance to the biological actions of opioids (*i.e.*, variations within genes encoding G protein-coupled receptors [the GPCRs dopamine D₂ receptor (Blomqvist et al., 2000; Ikeda et al.,

2008), serotonin 5-HT_{2B} receptor (Lin et al., 2004; Doly et al., 2008; Doly et al., 2009), and μ -opioid receptor (Oertel et al., 2012; Beer et al., 2013; Carpentier et al., 2013; Gong et al., 2013; Haerian and Haerian, 2013; Kolesnikov et al., 2013; Manini et al., 2013; Song et al., 2013; Cajanus et al., 2014; Solak et al., 2014; Xu et al., 2014; Adrian et al., 2015; Bartosova et al., 2015; Hancock et al., 2015; Laugsand et al., 2015; Olesen et al., 2015; Pfeifer et al., 2015; Wachman et al., 2015; Woodcock et al., 2015; Matic et al., 2016; Schwantes-An et al., 2016; Choi et al., 2017; Gao et al., 2017; Levran et al., 2017)]; Regulators of G protein Signaling [RGS2, RGS17] (Smoller et al., 2008; Amstadter et al., 2009b; Amstadter et al., 2009a; Koenen et al., 2009; Zhang et al., 2012; Dunn et al., 2014; Stein et al., 2014; Hohoff et al., 2015); a nerve growth factor [brain-derived neurotrophic factor] (Egan et al., 2003; de Cid et al., 2008); a voltage-gated potassium channel [KCNC1](Nascimento and Andrade, 2016); and, a second messenger-regulated transcription factor [cyclic-AMP responsive element binding protein-1] (Pal et al., 2014); Table 1). We isolated buccal swabs from 60 WV COAT clinic participants, prepared genomic DNA, and performed genotyping of the following SNPs.

2.4.1 - *OPRM1* rs9479757 (mu opioid receptor gene variant) (Beer et al., 2013; Xu et al., 2014; Olesen et al., 2015)): Xu *et al.* (Xu et al., 2014) identified rs9479757 as linked to heroin addiction severity among Han Chinese males; the G-to-A transition facilitates exon 2 skipping, leading to altered *OPRM1* splice-variant mRNAs and encoded mu opioid receptor (MOR) isoforms. All sixty participants were heterozygous for *OPRM1* rs9479757; in contrast, the minor allele frequency in the general European population is only 0.097 (Table 1).

2.4.2 - *OPRM1* rs1799971 (mu opioid receptor gene variant) (Oertel et al., 2012; Carpentier et al., 2013; Gong et al., 2013; Haerian and Haerian, 2013; Kolesnikov et al., 2013; Manini et al., 2013; Song et al., 2013; Cajanus et al., 2014; Solak et al., 2014; Adrian et al., 2015; Bartosova et al., 2015; Hancock et al., 2015; Laugsand et al., 2015; Olesen et al., 2015; Pfeifer et al., 2015; Wachman et al., 2015; Woodcock et al., 2015; Matic et al., 2016; Schwantes-An et al., 2016; Choi et al., 2017; Gao et al., 2017; Levran et al., 2017)): Woodcock *et al.*(Woodcock et al., 2015) published that Caucasian male carriers of the non-synonymous

OPRM1 rs1799971 minor allele (G) reported significantly more heroin use-related consequences and quit attempts, and were more likely to seek OUD treatment, than individuals homozygous for the ancestral A allele. Schwantes-An *et al.* (Schwantes-An et al., 2016), in a meta-analysis of ~28,000 Europeans, investigated non-specific risk for SUD and compared cases dependent on any substance to controls not dependent on all assessed substances. The rs1799971 G allele showed a modest protective effect on general substance dependence (Schwantes-An et al., 2016). The rs1799971 G allele was significantly less frequent in WV COAT clinic participants than in the population at large, by a factor of ~2 (Table 1; 0.083 vs 0.162, $p = 0.018$).

[Location of Table 1]

2.4.3 - *RGS2* rs4606 (Regulator of G protein Signaling-2 gene variant) (Smoller et al., 2008; Amstadter et al., 2009b; Amstadter et al., 2009a; Koenen et al., 2009; Dunn et al., 2014; Stein et al., 2014; Hohoff et al., 2015)): One of us (Oliveira-Dos-Santos et al., 2000) previously reported that loss of *RGS2* in mice engenders heightened anxiety and diminished aggression. Subsequently, Smoller *et al.* (Smoller et al., 2008) found that *RGS2* variations, including rs4606, were associated with introversion (a core personality trait in social anxiety disorder) (Smoller et al., 2008). Associations between the *RGS2* SNP rs4606 and panic disorder (Hohoff et al., 2015), suicidal ideation (Amstadter et al., 2009b), post-traumatic stress disorder (Amstadter et al., 2009a), and generalized anxiety disorder (Koenen et al., 2009) have also been reported. We found that rs4606 was more frequent in COAT clinic volunteers than in the general population by a factor of ~2 (Table 1; 0.533 vs 0.276, $p < 0.001$).

2.4.4 - *HTR2B* rs6736017 (serotonin [5-HT] receptor-2B gene) (Lin et al., 2004; Doly et al., 2008; Doly et al., 2009) **variant**): There are presently no published details on the *HTR2B* SNP rs6736017. However, *HTR2B* variations likely play a role in drug abuse. Doly *et al.* (Doly et al., 2008; Doly et al., 2009) showed that the serotonin 5-HT_{2B} receptor, encoded by the *HTR2B* gene, is required for 3,4-methylene-dioxymethamphetamine

(MDMA)-induced hyperlocomotion, serotonin and dopamine release, and conditioned place preference, the latter a rodent-based metric of reinforcing properties. Furthermore, Lin *et al.* (Lin et al., 2004) conducted a genome-scan identifying the human *HTR2B* gene as a candidate for drug abuse liability. The *HTR2B* SNP rs6736017 was markedly more frequent in COAT clinic participants than in the population at large, when comparing to the 60,000 genomes of the ExAc database (Table 1; 0.500 vs 0.0078, $p < 0.01$).

None of the other 10 SNPs investigated exhibited a significantly different frequency in the COAT clinic volunteers *versus* the population at large (Table 1).

2.5 - DISCUSSION

We identified four SNPs present at different frequencies in persons with OUD compared with a demographically-matched general population. It is currently difficult to quantify the degree of significance of each of these altered frequencies on OUD in the COAT population. One reason for this difficulty is that many gene variants are likely to play an integrated role in predisposition to OUD, and these variants may have additive or synergistic effects; here, we have only interrogated 14 SNPs in 8 candidate genes. Also, possible gene x environment effects were not assessed in the present study.

Two SNPs were in the *OPRM1* gene encoding MOR -- the main molecular target for the euphoric effects of opioids, as well as for the therapeutic effects of buprenorphine (Marshalek and Sullivan, 2010; Zheng et al., 2017) and methadone in OUD treatment. Therefore, it is not surprising that *OPRM1* variations would be associated with OUD.

We also identified an *RGS2* variant associated with OUD. This finding is novel, but it is also not surprising. Opioids like oxycodone and buprenorphine generate neuronal signals by activating opioid-binding GPCRs, altering receptor conformation to cause nucleotide exchange (Siderovski and Willard, 2005) by the associated G protein heterotrimer (G α ·GDP/G β /G γ). GTP-loaded G- α then dissociates from the G β / γ heterodimer; both subunits become free to modulate enzymes that generate intracellular second messengers (e.g., cyclic AMP, calcium) or stimulate potassium channels, leading to neuronal membrane hyperpolarization and thus action potential inhibition. G- α proteins have intrinsic GTPase activity, leading to self-inactivation and reassociation with G $\beta\gamma$ subunits (Siderovski and Willard, 2005). RGS proteins like RGS2 greatly enhance the GTP hydrolysis rate of G α subunits (Ingi et al., 1998; Siderovski and Willard, 2005; Kimple et al., 2009), thereby inhibiting GPCR signaling. Thus, one might indeed expect that a polymorphism in an *RGS* gene – encoding a negative regulator of GPCR signaling -- would be implicated in OUD. Moreover, *RGS2* function is particularly germane to multiple anxiety disorders (Smoller et al., 2008; Amstadter et al., 2009b; Amstadter et al., 2009a; Koenen et al., 2009; Dunn et al., 2014; Stein et al., 2014; Hohoff et al., 2015) that could predispose an individual toward OUD.

HTR2B encodes the serotonin 5-HT_{2B} receptor, implicated in the rewarding actions of MDMA (Doly et al., 2008; Doly et al., 2009) and cocaine (Doly et al., 2017), and thought to be involved in impulsivity (Bevilacqua et al., 2010) and resistance to antidepressants (Diaz et al., 2016), conditions that could predispose to OUD. Other polymorphisms in *HTR2B* have also been associated with vulnerability to illegal drug use (Lin et al., 2004). Therefore, our identification of a *HTR2B* polymorphism significantly more frequent in persons with OUD compared with a demographically-matched general population is not surprising.

There are several goals for identifying SNPs associated with OUD. One of these goals is to identify markers that, in genetic screens, could identify people at risk for developing OUD (*e.g.*, before prescribing opioid analgesics). Another goal is to identify potential novel “druggable” targets for preventing and/or treating OUD. For instance, the published role of the 5-HT_{2B} receptor in the actions of illicit drugs, coupled with our finding that rs6736017 is so frequent in OUD individuals, suggests that 5-HT_{2B} receptor-active pharmaceuticals (of which many are already approved for human use) might have efficacy in preventing/treating OUD. Future studies with larger cohorts are required to validate our findings, both in West Virginia and in other, more demographically heterogeneous regions. Such studies should identify genes to interrogate in order to predict individuals with susceptibility to OUD/SUD and in doing so provide an opportunity for the prevention of the disease.

Footnote:

* Demographic homogeneity in West Virginia in no way reflects any heightened degree of consanguinity. Tincher (Tincher, 1980) examined 140 years' worth of marriage records in a remote four-county Appalachian region. While cousin marriage was found to occur in Appalachia, it was not conspicuously more prevalent than in other geographical locales: by 1970, the cousin marriage rate was no higher than in the general U.S. population.

2.6 – TABLES

2.6.1 Table 1. Genetic markers and minor allele frequency (MAF) data obtained from public databases (1000 Genomes Project and/or ExAc) and WV COAT clinic patients (N=60)

Table 1. Genetic markers and minor allele frequency (MAF) data obtained from public databases (1000 Genomes Project and/or ExAc) and WV COAT clinic patients (N = 60).						
Gene	dbSNP ID	Minor Allele	TaqMan Probe	MAF via 1000 Genomes*	MAF via ExAc†	MAF from COAT clinic‡
<i>OPRM1</i>	rs9479757	A	C_25472011_10	0.097	0.082	0.500 p < 0.001
<i>OPRM1</i>	rs1799971	G	C_8950074_1	0.162	0.185	0.083 p = 0.018
<i>RGS2</i>	rs4606	G	C_2498717_10	0.276	n.a.	0.533 p < 0.001
<i>HTR2B</i>	rs6736017	C	C_25614588_20	n.a.	0.0078	0.500 p < 0.001
<i>OPRM1</i>	rs2075572	G	C_1691815_1	0.413	n.a.	0.483 p = 0.12
<i>RGS17</i>	rs596359	T	C_7830523_10	0.517	n.a.	0.467 p = 0.27
<i>DRD2 / ANKK1</i>	rs1800497	A	C_7486676_10	0.188	0.256	0.217 p = 0.41
<i>DRD2</i>	rs1799978	C	C_7486599_20	0.060	n.a.	0.075 p = 0.44
<i>BDNF</i>	rs6265	T	C_11592758_10	0.197	0.194	0.167 p = 0.49
<i>KCNC1</i>	rs60349741	C	C_89088414_10	0.001	n.a.	0.000
<i>HTR2B</i>	rs77982984	A	C_99996051_10	0.003	0.0012	0.000
<i>HTR2B</i>	rs78484969	C	C_99996081_10	0.001 (N = 1030 Finns)	n.a.	0.000
<i>HTR2B</i>	rs77570025	G	C_99996068_10	0.006	0.002	0.000
<i>CREB1</i>	rs35349697	A	C_25636228_10	n.a.	0.000008	0.000

* Minor allele frequency reported on N = 1006 Europeans via 1000 Genomes (except as otherwise noted); n.a. = not available.

† Minor allele frequency reported on N ~ 60,000 people via ExAc database of the Broad Institute; n.a. = not available.

‡ p-value, denoting significance of difference between MAFs, is derived by comparison between WV COAT clinic and European (if not, then ExAc) MAF using binomial test (null hypothesis is "no difference between WV COAT patients and Europeans"; alternative is "there is difference between WV COAT patients and Europeans")

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Chapter Three

Preclinical testing of nalfurafine as an opioid-sparing adjuvant that potentiates analgesia by the mu opioid receptor-targeting agonist morphine

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3.1 - Abstract:

mu opioid receptor (MOR)-targeting analgesics are efficacious pain treatments, but notorious for their abuse potential. In animal models, co-administration of traditional kappa opioid receptor (KOR)-targeting agonists with MOR-targeting analgesics can decrease reward and potentiate analgesia. However, traditional KOR-targeting agonists are well known for inducing anti-therapeutic side effects (psychotomimesis, depression, anxiety, dysphoria). Recent data suggest that some functionally selective or “biased” KOR-targeting agonists might retain the therapeutic effects of KOR activation without inducing undesirable side effects. Nalfurafine, used safely in Japan since 2009 for uremic pruritus, is one such functionally selective KOR-targeting agonist. Here we quantify the bias of nalfurafine and several other KOR agonists against the unbiased reference standard U50,488 and show that nalfurafine demonstrates a significant G protein-signaling bias. While nalfurafine (0.015 mg/kg) produces spinal analgesia (*via* a tail-immersion test) equivalent to 5 mg/kg of U50,488, it significantly enhances the analgesic effect of 5 mg/kg morphine in the hot plate assay. In addition, 0.015 mg/kg nalfurafine does not produce conditioned place aversion (CPA) yet retains the ability to reduce morphine-induced conditioned place preference (CPP) in C57BL/6J mice. Nalfurafine did produce robust inhibition of both spontaneous and morphine-stimulated locomotor behavior, however, suggesting a persistence of sedative effects when co-administered with morphine. Taken together, these findings suggest that G protein-biased KOR agonists may produce greater analgesic augmentation than unbiased agonists when combined with traditional opioid analgesics, while also reducing opioid-induced reward with less risk of dysphoria. Thus, adjuvant administration of G protein-biased KOR agonists may be beneficial in enhancing the therapeutic potential of MOR-targeting drugs, such as morphine.

3.2 - Introduction

Opioid use disorder (OUD) is an American health crisis (Manchikanti et al., 2012; Koh, 2015; Volkow and Collins, 2017). Use of prescription opioids, such as morphine and oxycodone, has contributed significantly to the OUD problem (Jones, 2013a). Morphine, the prototypical opioid analgesic, is a mainstay for treating moderate to severe pain as may occur post-operatively or in chronic illness. While prescription opioids are effective painkillers, there currently exists no facile way to use them for extended periods of time without the prospect of inducing tolerance, dependence and, in many cases, addiction. One plausible solution is to use morphine (and other MOR-targeting analgesics) in a way that increases its analgesic effects, thereby requiring lower doses (an approach known as “dose sparing”), or in a way that reduces addictive potential. In this vein, if the anti-nociceptive potential of MOR-targeting analgesics could be enhanced *via* an adjuvant that does not induce therapeutically limiting side effects, then their continued therapeutic use in treating pain could be extended.

kappa opioid receptor (KOR)-targeting agonists, when given with MOR-targeting analgesics, not only produce additive analgesia (Sutters et al., 1990; Negus et al., 2008b; Briggs and Rech, 2009), reduced tolerance (He and Lee, 1997; Khotib et al., 2004), and reduced respiratory depression (Verborgh et al., 1997) but can also reduce the rewarding properties of MOR-targeting analgesics (Kuzmin et al., 1997; Tsuji et al., 2001; Tao et al., 2006). However, clinical use of some KOR agonists, such as ketocyclazocine (Kumor et al., 1986) or salvinorin A (MacLean et al., 2013), has largely failed due to poor tolerability. In rodents, traditional KOR agonists produce anxiety- and depressive-like behaviors (Carlezon et al., 2006; Bruchas et al., 2009; Ehrich et al., 2015a), as well as locomotor suppression (Kunihara et al., 1993; Narita et al., 1993; Ehrich et al., 2015a; Brust et al., 2016b). In humans, the highly selective KOR agonist salvinorin A produces similar therapeutically limiting effects (MacLean et al., 2013; Maqueda et al., 2015; Johnson et al., 2016).

Thus, it is not surprising that initial enthusiasm for KOR agonists as adjuvants to MOR-targeting analgesics was reduced by early preclinical and clinical findings using single applications of mixed MOR/KOR agonists. For example, while nalbuphine studies in rodents demonstrated reduced conditioned place preference (CPP) to

morphine and enhanced morphine-induced analgesia (Tao et al., 2006), human studies demonstrated dose-dependent increases in dysphoria (Jasinski and Mansky, 1972; Preston et al., 1989a; Walsh and Babalonis, 2017). Similarly, pentazocine and butorphanol also induce classic dysphoric effects of traditional KOR-targeting agonists in humans, especially at higher doses (Preston et al., 1987; Zacny et al., 1994; Greenwald and Stitzer, 1998; Walsh et al., 2008; Walsh and Babalonis, 2017), limiting their therapeutic potential.

Recent efforts to develop non-addictive opioids however, have renewed interest in KOR ligands with specific emphasis on developing functionally selective (or “biased”) KOR agonists (Urban et al., 2007; Rankovic et al., 2016). G protein-biased KOR agonists demonstrate reduced potencies for signaling through the GRK/ β -arrestin pathway (Bohn and Aube, 2017). While G protein-dependent signaling through KOR is well established as producing the primary therapeutic outcome of analgesia (Chavkin, 2011; Brust et al., 2016b; Schattauer et al., 2017), the role of GRK/ β -arrestin signaling in producing dysphoria downstream of KOR activation is less clear. Some have shown that aversive behavior toward KOR agonists requires GRK3 (and p38 MAPK) activation (Bruchas et al., 2007b; Chavkin et al., 2014; Ehrich et al., 2015a), while others have observed little effect of *Arreb2* (β -arrestin-2) genetic ablation on the aversion produced by a variety of KOR agonists (White et al., 2015). Recent work with triazole 1.1 (Brust et al., 2016b), however, and nalfurafine (Liu et al., 2018) suggests that these G protein-biased KOR agonists may lack dysphoric effects at doses that produce analgesia, supporting the role of β -arrestin signaling in the dysphoric effects of traditional KOR agonists. G protein-biased KOR agonists may, thus, present a strategy for reducing the rewarding properties of MOR-targeting analgesics while avoiding the pitfalls of traditional KOR agonists. This notion is supported by a recent preclinical study in rats (Townsend et al., 2017b) showing that nalfurafine reduces intravenous self-administration of the MOR-targeting analgesic oxycodone while also potentiating its analgesic effect.

Nalfurafine is a G protein functionally selective KOR-targeting agonist with high translational potential given its safe usage in Japan since 2009 to treat uremic pruritis and, importantly, does not produce psychotomimesis (Inui, 2015). Therefore, we aimed to assess the therapeutic efficacy of nalfurafine in both

potentiating the analgesic effect and reducing the rewarding properties and of the MOR-targeting analgesic morphine, while also assessing the biased signaling properties of nalfurafine both *in vitro* and *in vivo*.

3.3 - Methods:

3.3.1 - Animal Subjects

All experiments were conducted using male and female C57BL/6J mice (RRID: IMSR_JAX:000664) between 8-14 weeks of age and 17-35 g; the sexes were used in similar proportions across all experiments. Original breeding pairs were obtained from The Jackson Laboratory (Bar Harbor, ME). All procedures were carried out in accordance with the National Research Council's *Guide to the Care and Use of Laboratory Animals (Eighth Edition)*, and were approved by West Virginia University's Institutional Animal Care and Use Committee.

3.3.2 - Study Design

Mice were randomly assigned to treatment conditions, after being balanced for sex. Mice were acclimated to the test room for at least 30 min prior to any study. All mice were used for a single experimental assay. Sample size for each experiment was determined by power analysis using effect sizes derived from literature, power value of 0.8, and an alpha value of 0.05. All experiments were done by personnel who were blinded to treatment conditions.

3.3.3 - Drugs

(±) U50,488, GR 89696, and ICI 199,441 were all purchased from Tocris Biosciences (Minneapolis, MN). Nalfurafine hydrochloride was purchased from Medchem Express (Monmouth Junction, NJ). Morphine sulfate salt pentahydrate was purchased from Sigma Aldrich (St. Louis, MO). Salvinorin A, ethoxymethyl ether Sal B (EOM Sal B), and triazole 1.1 were prepared as previously described (Zhou et al., 2013; Ewald et al., 2017a). All drugs were dissolved in 100% DMSO and diluted in saline to the desired concentration (no DMSO concentration exceeded 5% v/v final). In drug combination experiments, nalfurafine was injected subcutaneously (sc) as described in the limited literature available regarding this compound's use in mice (Endoh et al., 1999; Tsuji et al., 2001; Liu et al., 2018), while U50,488 and morphine were delivered by intraperitoneal (ip) injection as commonly described (Rada et al., 1996; McLaughlin et al., 2006; Koo et al., 2012; Muschamp et al., 2012; Laman-Maharg et al., 2017; Robinson et al., 2017). Dose range for nalfurafine

and U50,488 were chosen based upon the place preference data in the literature (Tsuji et al., 2001; Land et al., 2009b; Ehrich et al., 2015b). Dose range for morphine was chosen to produce consistent place preference while allowing for high sensitivity to detect potentiation in the hot plate assay of analgesia (Mueller et al., 2002; Raehal and Bohn, 2011)

3.3.4 - Measurement of Signaling Bias

Quantification of G protein signaling dependent changes in intracellular cyclic AMP (cAMP) levels was conducted using a GloSensor luciferase-based assay in HEK293T cells transiently transfected with 5 μ g of KOR expression vector DNA (3xHA-hKOR; www.cDNA.org) and 5 μ g of pGloSensor-22F cAMP biosensor expression vector DNA (Promega, Madison, WI). Quantification of β -arrestin recruitment was conducted using the luciferase-based TANGO assay (Barnea et al., 2008; Allen et al., 2011) in HTLA cells (generously provided by Dr. Gilad Barnea) transiently transfected with 5 μ g of KOR-V2-TEV-tTA expression vector DNA (Addgene plasmid #66462 donated by Dr. Bryan Roth). The appropriate masses of expression vector DNAs transfected was empirically determined prior to acquisition of the data contained herein. Transfections were performed in 10-cm dishes using the calcium phosphate-based method (Jordan et al., 1996). Culture media was replaced 16 hrs after transfection, and cells then plated on 96-well plates approximately 24 hrs after transfection. All cells were serum-starved with 1% dialyzed serum-containing medium for 8-16 hrs before assaying. For assays of cAMP inhibition, medium was replaced with GloSensor detection reagent dissolved in HEPES (20 mM)-buffered Hank's balanced saline solution, and cells were incubated for 2 hrs at room temperature. KOR-selective agonists were applied for 15 min prior to a 15-min stimulation with 100 nM isoproterenol (to induce endogenous β -adrenergic receptor-mediated and $G\alpha_s$ -dependent production of cAMP) and subsequent quantification of luminescence. For TANGO assays of β -arrestin recruitment, drug stimulation was performed overnight in 1% dialyzed serum-containing medium. Medium was removed the following morning, the cells lysed with BrightGlo reagent (Promega), and luminescence quantified using a FlexStation 3 multi-mode plate reader (Molecular Devices).

3.3.5 - Calculation of Bias Factor

Bias factors were calculated using the method previously described (Kenakin, 2017). Briefly, $\text{Log}(E_{\text{max}}/EC_{50})$ values for both GloSensor and TANGO assays were separately generated for each ligand using potency (EC_{50}) and maximal efficacy (E_{max}) values obtained from these assays. $\Delta\text{Log}(E_{\text{max}}/EC_{50})$ was calculated in each assay as the difference in $\text{Log}(E_{\text{max}}/EC_{50})$ for each ligand from the reference ligand (the unbiased KOR agonist U50,488). $\Delta\Delta\text{Log}(E_{\text{max}}/EC_{50})$ was then calculated as the difference in $\Delta\text{Log}(E_{\text{max}}/EC_{50})$ between GloSensor and TANGO assays for each ligand. The 95% confidence interval (CI) for each bias factor was calculated using the E_{max} and EC_{50} values generated in individual experiments for each ligand in each assay.

3.3.6 - Warm Water Tail Withdrawal

The warm water tail withdrawal test was conducted as described previously (Schattauer et al., 2017). Briefly, mice were gently wrapped in cloth before immersing the distal 2 cm of the tail in 55°C water and recording the latency for the mouse to withdraw its tail from the water. Baseline latencies were recorded 30 min after administration of vehicle. Mice were then injected with drug and latencies were recorded after 30 min. A maximum cutoff of 10 s was used to avoid tissue damage. Data are presented as latency to tail-withdrawal in seconds as well as percent maximum possible effect (%MPE) following the equation: $\%MPE = (\text{latency} - \text{baseline}) / (10 \text{ sec} - \text{baseline}) * 100\%$.

3.3.7 - Hot Plate Analgesia Test

To assess integrated pain perception, the hot plate test of nociception was performed as described previously (Raehal and Bohn, 2011) with minor modification. Nociceptive latency after vehicle injection was measured on Day 1. On Day 2, mice were injected with drug and response latencies were measured after 30 min. Individual mice were placed onto a 53°C hot plate (IITC Life Science Inc.; Woodland Hills, CA) inside of a clear, open-bottom acrylic cylinder. A maximum cutoff of 30 s was used to avoid tissue damage. Time until the first nociceptive sign (*i.e.*, licking of the hind paw, flicking of the hind paw, or jumping) was recorded by a trained observer to the nearest 0.1 sec and reported as percent maximum possible effect ($\%MPE =$

(latency - baseline) / (30 sec - baseline) * 100%). Data were analyzed by one way ANOVA with a Tukey's *post hoc* test, using Prism 7 software.

3.3.8 - Locomotion

Locomotor activity was assessed using a 16 X 16 open-field infrared photobeam activity system (PAS) from San Diego Instruments (San Diego, CA), as described previously (Gross et al., 2018). Briefly, locomotion was quantified as total beam breaks in the *x* and *y* coordinates in five minute bins. Spontaneous locomotion was assessed by administering drug or vehicle immediately before placing the mouse in the locomotor chamber for 60 min. Morphine-induced locomotion was assessed in a three-day procedure. On Day 1, mice were habituated to the locomotor chamber for 60 min. On Day 2, mice were habituated to the locomotor chamber for 30 min before receiving an injection of vehicle and returning to the chamber for an additional 60 min. On Day 3, mice were habituated to the locomotor chamber for 30 min before receiving an injection of either morphine with vehicle or morphine with KOR agonist, and returning to the chamber for an additional 60 min. Data plotted over time were normalized to the average vehicle locomotion in the first 5 minute bin for novelty induced locomotion and to the average locomotion of each subject during the 30 min prior to injection for morphine suppressed locomotion. For total novelty locomotion, data are normalized to the average total locomotor activity of the vehicle condition over the full 60 minute trial. For total morphine stimulated locomotion, data are normalized to the average total locomotor activity during the 60 minutes post vehicle injection on day 2.

3.3.9 - Conditioned Place Preference/Aversion

CPP and CPA studies were conducted as described previously (Redila and Chavkin, 2008) with the following modifications. A two-chambered place preference apparatus with differential floor grating in the white and black chambers was used. Baseline preference was established on Day 1 by allowing the mouse free access to both chambers for 30 min (for U50,488 studies) or 50 min (for nalfurafine studies, given prior evidence of time-of-effect for this compound (Tsuji et al., 2001)) and recording the amount of time spent in each chamber. The conditioning phase occurred on days 2 and 3 in 30 min (U50,488) or 50 min (nalfurafine) sessions. Vehicle was administered in the AM session and drug in the PM session with a 4-hr interval between

sessions. For CPP, drug was administered in the least-preferred chamber to override the initial preference of the mouse. Drug was administered in the most-preferred chamber for CPA. Preference was assessed on Day 4 by allowing the mouse free access to both chambers for 30 min (U50,488) or 50 (nalfurafine) min, and recording the amount of time spent in each chamber. ANY-maze video tracking software (Stoelting Co., Wood Dale, IL) was used to track movement and time spent in each chamber. Data are presented as the difference in time spent in the drug-paired chamber for each mouse. Data were analyzed by two-way ANOVA with Dunnett's *post hoc* test to compare to vehicle controls and Sidak's *post hoc* test to compare across genotypes, each run using Prism 7 software (GraphPad).

3.4 - Results

3.4.1 - Selection of nalfurafine as KOR biased agonist under consideration as a possible morphine adjuvant

Before launching our assessment of the efficacy of G protein-biased KOR agonists in reducing the addictive properties of morphine, we characterized the signaling properties of a variety of KOR-specific compounds to identify a suitable biased KOR agonist. Utilizing a GloSensor-based cAMP assay to determine G protein signaling, and the TANGO assay to determine β -arrestin recruitment, we identified EOM-salvinorin B (Bias Factor = 15.26, +95%CI = 29.95, -95%CI = 7.78) and nalfurafine (Bias Factor = 7.73, +95%CI = 15.16, -95%CI = 3.94) as strongly G protein-biased KOR agonists (Figure 1), the latter consistent with prior findings (Schattauer et al., 2017). As nalfurafine has been used successfully for nearly a decade in Japanese patients for the treatment of uremic pruritus (Ueno et al., 2013; Inui, 2015), we chose it as our biased KOR agonist given its greater translational potential as a viable pharmaceutical adjuvant.

3.4.2 - Spinal anti-nociception by nalfurafine

To establish the level of analgesia to be expected from these doses of nalfurafine, we assessed their abilities in the mouse tail immersion test (relative to the unbiased KOR agonist U50,488) to produce spinal anti-nociception, an outcome of KOR agonism known to be G protein-dependent. Tail withdrawal latencies were analyzed by one-way ANOVA: $F(6,152)=29.97$, $p<0.0001$. All doses of nalfurafine produced significant increases in withdrawal latency compared to vehicle (Figure 2; $p<0.0001$ for each; Dunnett's *post hoc* test). Administration of 1.25 mg/kg and 5 mg/kg U50,488 also increased tail withdrawal latency compared to vehicle (Figure 4; $p<0.001$ and $P<0.0001$ respectively), while no significant difference was observed between 2.5 mg/kg U50,488 and vehicle (Holm-sidak *post hoc* test, $p>0.05$), though 5 mg/kg was significantly higher than 2.5 mg/kg U50,488 (Holm-sidak *post hoc* test, $p<0.05$). No significant difference was observed between 1.25 mg/kg and 5 mg/kg U50,488 (Holm-sidak *post hoc* test, $p>0.05$). Both 0.015 mg/kg and 0.03 mg/kg doses of nalfurafine elicited comparable anti-nociception to 5 mg/kg U50,488 (Holm-Sidak *post hoc* test, $p=0.613$ for both).

3.4.3 - Enhancement of morphine-induced supraspinal analgesia by nalfurafine co-administration

To assess the effect of KOR agonism on the integrated nociceptive response, we assessed the effect of nalfurafine co-administration on morphine-induced analgesia *via* the hot plate test. Nociceptive latency with

administration of U50,488 was analyzed by one-way ANOVA: $F(6,57)=4.798$, $p=0.0005$. Nociceptive latency with administration of nalfurafine was also analyzed by one-way ANOVA: $F(6,68)=31.80$, $p<0.0001$. We found that no dose of U50,488 nor nalfurafine produced a significant analgesic response on its own in the hot plate assay of nociception (Figure 5; Sidak's *post hoc* test, $p > 0.05$). Co-administration of 2.5 mg/kg U50,488 with 5 mg/kg morphine significantly increased analgesia compared to U50,488 alone (Holm-Sidak's *post hoc* test, $p < 0.01$; Figure 5A), but this level of analgesia was not significantly different from 5 mg/kg morphine alone (Holm-Sidak's *post hoc* test, $p > 0.05$; Figure 5A).

Co-administration of all doses of nalfurafine tested with 5 mg/kg morphine significantly increased analgesia compared to nalfurafine alone (Holm-Sidak's *post hoc* test, $p < 0.0001$; Figure 5B) and 5 mg/kg morphine alone (Holm-Sidak's *post hoc* test, $p<0.0001$ for 0.015 and 0.06 mg/kg nalfurafine co-admin; $p<0.001$ for 0.03 mg/kg nalfurafine co-admin, Figure 5B).

3.4.4 - Locomotor suppression by nalfurafine administration and co-administration

Open-field locomotion represents a behavioral outcome of mesolimbic dopamine release and is increased by movement of mice into a novel environment, as well as by administration of many drugs of abuse to mice (Bardo et al., 1990; Sellings and Clarke, 2003; Bromberg-Martin et al., 2010; Fields and Margolis, 2015). Traditional KOR agonists suppress both spontaneous as well as opioid analgesic-induced locomotion (Narita et al., 1993; Chefer et al., 2005; Bruijnzeel, 2009). Suppression of total novelty-induced locomotion was analyzed by one-way ANOVA: $F(6,104)=18.01$; $p<0.0001$. We found that total spontaneous locomotion was significantly reduced upon administration of 2.5 and 5 mg/kg U50,488 ($p<0.05$ and $p<0.01$ respectively), as well as 0.015, 0.03 and 0.06 mg/kg nalfurafine ($p<0.001$, and $p<0.0001$ respectively) compared to vehicle (Dunnett's *post hoc* test; Figure 6A and inset). The dose of 0.06 mg/kg nalfurafine elicited a significantly greater reduction in spontaneous locomotion, as compared to 1.25, 2.5 or 5 mg/kg U50,488 ($p<0.0001$, $p<0.001$, and $p<0.0001$ respectively), or 0.015 and 0.03 mg/kg nalfurafine ($p<0.05$, $p<0.01$ respectively); Tukey's *post hoc* test; Figure 6A and inset). No significant differences in locomotor suppression were seen between doses of

U50,488 (Tukey's *post hoc* test, $p \geq 0.4698$), nor between 5 mg/kg U50,488 and 0.015 or 0.03 mg/kg nalfurafine (Tukey's *post hoc* test, $p=0.9336$ and $p=0.9480$, respectively).

Modulation of total morphine-stimulated locomotion by U50,488 was analyzed by one-way ANOVA: $F(3,32)=6.142$, $p=0.0020$. Modulation by nalfurafine was also analyzed by one-way ANOVA: $F(3,27)=30.11$, $p<0.0001$. Suppression of locomotion was seen with co-administration of 5 mg/kg morphine with 5 mg/kg U50,488 (Dunnett's *post hoc* test, $p<0.01$; Figure 6D and inset), as well as with all doses of nalfurafine (Dunnett's *post hoc* test, $p<0.0001$; Figure 6E and inset).

3.4.5 - Dose-dependent conditioned place aversion by nalfurafine

To help assess the level of anti-therapeutic liability presented by nalfurafine, we measured its ability to produce conditioned place aversion in mice (Figure 3), a behavior linked to β -arrestin signaling downstream of KOR activation (Bruchas et al., 2007b; Ehrich et al., 2015a) and a therapeutically-limiting effect of traditional KOR agonists like U50,488 (Jasinski and Mansky, 1972; Preston and Bigelow, 1993; Siebert, 1994; Greenwald and Stitzer, 1998). U50,488-induced CPA was analyzed by two-way ANOVA: test day effect, $F(1,39)=20.41$, $p<0.0001$; treatment effect, $F(3,39)=8.940$, $p=0.0001$; interaction effect, $F(3,39)=4.684$, $p=0.0072$; subject effect, $F(39,39)=0.8332$, $p=0.7143$. Nalfurafine-induced CPA was also analyzed by two-way ANOVA: test day effect, $F(1,50)=13.04$, $p=0.0007$; treatment effect, $F(3,50)=2.302$, $p=0.0884$; interaction effect, $F(3,50)=2.572$, $p=0.0645$; subjects effect, $F(50,50)=0.9840$, $p=0.5227$. We found that all doses of U50,488 produced aversion compared to vehicle (Dunnett's *post hoc* test, 1.25 mg/kg = $p<0.05$, 2.5 mg/kg = $p<0.001$, and 5 mg/kg = $p<0.01$ Figure 3A) with no significant difference between doses (Holm-Sidak *post hoc* test, $p>0.05$) while only the 0.03 mg/kg dose of nalfurafine produced statistically significant aversion compared to vehicle (Dunnett's *post hoc* test, $p<0.01$; Figure 3B) with no significant difference between doses (Holm-sidak *post hoc* test, $p>0.05$).

3.4.6 - Modulation of morphine-induced conditioned place preference by nalfurafine

As unbiased KOR agonists have been observed, when co-administered, to reduce the rewarding properties of various drugs of abuse, including morphine (Kuzmin et al., 1997; Tao et al., 2006), we evaluated the ability

of nalfurafine co-administration in mice to reduce the conditioned place preference produced by morphine. U50,488 results were analyzed by two-way ANOVA: test day effect, $F(1,78)=218.2$, $p<0.0001$; treatment effect, $F(3,78)=0.8699$, $p=0.4604$; interaction effect, $F(3,78)=3.482$, $p=0.0197$; subject effect, $F(78,78)=1.032$, $p=0.4441$. Morphine alone (5 mg/kg) produced place preference compared to preconditioning (Sidak *post hoc test*, $p<0.0001$; Figure 2). U50,488 co-administration (at 1.25 mg/kg and 5 mg/kg) reduced preference for morphine (Dunnett *post hoc test*, $p < 0.05$; Figure 2A), though there was no significant difference between doses (Holm-Sidak *post hoc test*, $p>0.05$). Nalfurafine results were also analyzed by two-way ANOVA: test day effect, $F(1,85)=169$, $p<0.0001$; treatment effect, $F(3,85)=5.049$, $p=0.0029$; interaction effect, $F(3,85)=1.209$, $p=0.3116$; subjects effect, $F(85,85)=0.9096$, $p=0.6683$. Nalfurafine co-administration showed no effect on morphine-induced preference at 0.03 mg/kg, but produced a significant reduction in morphine preference at 0.015 mg/kg and 0.06 mg/kg, with 0.06 mg/kg being significantly different from 0.03 mg/kg nalfurafine co-administration. Dunnett *post hoc test*, $p<0.05$, $p<0.01$ and Holm-Sidak *post hoc test*, $p<0.05$, respectively; Figure 2B).

3.5 - Discussion

Our results support nalfurafine as a G protein-biased KOR agonist that can beneficially modulate both the analgesic and rewarding properties of morphine *in vivo* upon co-administration; however, nalfurafine also produced aversion and locomotor suppression in mice. KOR agonists reduce reward associated with drugs of abuse (Lindholm et al., 2001; Morani et al., 2009a; Townsend et al., 2017b); however, their clinical viability has been hampered by dysphoria in patients (Jasinski and Mansky, 1972; Preston and Bigelow, 1993; Siebert, 1994; Greenwald and Stitzer, 1998). The dysphoria produced by KOR agonism has been linked to signaling

downstream of β -arrestin mobilization (Bruchas et al., 2007b; Ehrich et al., 2015a) and, thus, G protein-biased KOR agonists may lack dysphoric properties (Brust et al., 2016b; Liu et al., 2018). Our data demonstrate that while nalfurafine retains the ability to produce conditioned place aversion, the augmentation of morphine's analgesic effect persists at lower doses, providing evidence that G protein-biased KOR agonists may have increased clinical utility as dose-sparing agents compared to unbiased KOR agonists. Proper dose combinations may provide augmented analgesia while avoiding dysphoria and minimizing sedation.

Initial *in vitro* screening identified both nalfurafine and EOM Sal-B as potent and significantly G-protein biased KOR agonists. We chose to focus our initial efforts on nalfurafine due to its greater translational potential, being used safely in Japan since 2009 (Inui, 2015).

We first assessed spinal nociception utilizing the warm water tail withdrawal assay. As previous studies demonstrate that KOR agonist-induced increases in withdrawal latency are G protein-signaling dependent (Hernandez et al., 1995; Berg et al., 2011; White et al., 2015; Abraham et al., 2018), this assay also allowed for comparison of *in vivo* G protein-signaling between nalfurafine and U50 at the selected doses. A dose of 5 mg/kg U50,488 produced analgesia that was statistically equivalent to both 0.015 and 0.03 mg/kg nalfurafine. This finding suggests that KOR-mediated G protein signaling is equivalent *in vivo* between the higher dose of U50,488 and the lowest tested doses of nalfurafine.

As the supraspinal analgesia produced by MOR-targeting analgesics represents a major factor in their clinical success (Kanjhan, 1995; Jensen, 1997), we next assessed the effect of co-administration in a non-reflexive assay of nociception. While no significant analgesic effect of U50,488 or nalfurafine alone was observed in the hot plate assay compared to morphine alone, there was a significant effect of nalfurafine upon co-administration with morphine (Figure 5B). This increased analgesic effect was not seen with co-administration of U50,488. One potential explanation of this effect is that the locomotor-suppressing property of nalfurafine (Figure 6) hinders the expression of nociceptive behaviors (*e.g.*, jumping or paw flicking/licking), thereby falsely increasing nociceptive latency measurements. In the assay of morphine stimulated locomotion however, 5 mg/kg U50,488 suppresses locomotion to a similar extent as both 0.015 and 0.03 mg/kg (Figure 6),

yet only nalfurafine co-administration is seen to elicit augmented analgesia (Figure 5B). These findings make a locomotor confound to nociception measurement less likely. In this light, future studies are planned to investigate the effects of nalfurafine and MOR agonist co-administration on pain-suppressed behaviors to further investigate the potential confound and better characterize the analgesic interactions of these two drugs.

As 5 mg/kg U50,488 and 0.15 mg/kg nalfurafine produced equivalent increases in tail withdrawal latencies, the increased analgesic effect seen in the hot plate with co-administration of nalfurafine may indicate a beneficial supra-spinal or spinal interaction between morphine and nalfurafine downstream of G protein-signaling at KOR. Early work with site specific administration of MOR and KOR agonists demonstrate analgesic synergism when given intrathecally (i.t.) but not intracerebroventricularly (i.c.v.) (Ren et al., 1985; Sutters et al., 1990; Miaskowski et al., 1992; Miaskowski et al., 1993), suggesting that while KOR-dependent spinal analgesia is equivalent between 0.015 nalfurafine and 5 mg/kg U50,488, the biased signaling profile of nalfurafine produces a greater analgesic interaction between MOR and KOR signaling.

To assess the potential clinical viability of this co-administration paradigm, we evaluated these doses of nalfurafine and U50,488 for sedation and dysphoria. In the present study, significant reductions in spontaneous locomotion were seen with all KOR agonists tested except 1.25 mg/kg U50,488. This observation of sedation persisted when these KOR agonists were co-administered with morphine. However, post-marketing surveillance of nalfurafine reports a low incidence of somnolence (1%), at doses safe and efficacious in human patients (Kozono et al., 2018), suggesting that sedation may not necessarily prove to be an anti-therapeutic effect of nalfurafine co-administration with a MOR-targeting analgesic.

To assess dysphoria, we utilized the CPA assay. Aversion to KOR agonists has been linked to signaling downstream of β -arrestin (Bruchas et al., 2007b; Ehrich et al., 2015a) and previous work has demonstrated a lack of CPA to nalfurafine (Tsuji et al., 2001; Liu et al., 2018). In our CPA assays with nalfurafine, only 0.03 mg/kg produced significant aversion, as compared to U50,488 which produced significant aversion at all doses tested. The observed decrease in aversion with the highest dose of KOR agonist has been demonstrated in a

previous study (Robles et al., 2014) and could reflect the appearance of cognitive deficits (Castellano et al., 1988; Carey et al., 2009).

Of note is the finding that 0.03 mg/kg nalfurafine produced significant aversion. This result is supported by recent work demonstrating that the G protein biased KOR agonist RB-64 produces aversion in both wild-type and β -arrestin-2 knockout mice (in addition to the unbiased KOR agonists U69,593 and salvinorin A (White et al., 2015)), suggesting that aversion to KOR agonists may be produced in part through mechanisms independent of β -arrestin-dependent signaling. G protein-biased KOR agonists might thus retain the ability to produce aversion, though less potently than unbiased agonists.

Lastly, KOR agonists are known to reduce the rewarding properties of a variety of drugs of abuse (Kuzmin et al., 1997; Lindholm et al., 2001; Tsuji et al., 2001; Tao et al., 2006; Morani et al., 2009a; Ewald et al., 2017a; Townsend et al., 2017a), presenting a second possible benefit of combination therapy. Using the CPP assay, we found that nalfurafine reduces the rewarding properties of 5 mg/kg morphine at 0.015 and 0.06 mg/kg, while 0.03 mg/kg produced no effect on morphine place preference. Though a dose of 0.03 mg/kg reduced morphine CPP in a previous study, a potential source of this difference could be our use of inbred C57BL/6J mice, as opposed to the use of outbred ddY male mice (Tsuji et al., 2001). Indeed, strain differences in response to KOR agonists, as well as to psychostimulant drugs of abuse, have previously been demonstrated (e.g., (Castellano et al., 1988; Mouri et al., 2012)).

Administration of U50,488 displayed a similar pattern, with doses of 1.25 and 5 mg/kg suppressing morphine place preference, while 2.5 mg/kg was not a statistically significant suppression. Similar to results seen in CPA, the reduction in morphine induced CPP at the highest doses of each KOR agonist may be explained in part by cognitive disruption preventing the effective pairing of stimulus with context. Effects seen with the low and middle dose of each drug were opposite of expected. Interestingly, the doses of both drugs producing the greatest level of aversion produced the least suppression of morphine induced CPP. While extensive work focuses on KOR agonists effects on cocaine CPP (Suzuki et al., 1992; Redila and Chavkin, 2008; Morani et al., 2009b), less work investigates their effects on opioid-analgesic induced CPP, with few if

any using multiple doses of KOR agonist. Outside of the work by Tsuji et al. referenced earlier, a related lab demonstrated that 1 mg/kg U50,488 reduced CPP to 5 mg/kg morphine, though other doses of U50,488 were not assessed (Masahiko et al., 1993). A drawback of the CPP procedure is its relatively low sensitivity to dose magnitude (Napier et al., 2013), making dose-response relationships difficult to properly establish, however the effects reported here indicate an interaction between the dose of KOR agonist and the rewarding properties of morphine. As pretreatment with U50,488 has been shown to increase CPP to cocaine, it is possible that differences in the timing of KOR agonist effects at different doses in relation to the rewarding effects of morphine play a role in the observed response pattern. A more detailed interrogation of the effects seen in CPP with an expanded dose range and varied pretreatment intervals could aid in determining the pharmacodynamic and pharmacokinetic factors involved.

Taken together, these data suggest that G protein-biased KOR agonists may be effective dose-sparing adjuvants for traditional MOR-targeting analgesics. As all doses of nalfurafine provided similar augmentation of morphine's analgesic effect, low doses may provide adequate analgesic synergism while avoiding dysphoria and sedation. Nalfurafine is used in the treatment of uremic pruritis at doses that produce low rates of sedation, insomnia, constipation, psychosis and other common KOR related side-effects (Kozono et al., 2018). The anti-pruritic dose in rodents extends as low as 5 µg/kg (Liu et al., 2018). As a dose of 0.015 mg/kg effectively reduces morphine CPP without significant aversion, it is possible that doses already used in human subjects to treat pruritis may produce the benefits of co-administration described herein while avoiding significant anti-therapeutic effects. Though the signaling pathways involved in both the intended and unintended effects of KOR agonism requires further investigation, the reduced side-effect profile of nalfurafine as compared with unbiased KOR agonists suggests that G protein-biased KOR agonists may be clinically relevant adjuvants for opioid-based pain therapies, particularly in acute in-patient settings. Additional studies are planned to investigate the effects of chronic co-administration on analgesic tolerance, withdrawal and reward related behaviors.

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3.6 – Figures

3.6.1 Nalfurafine is a potent G protein biased KOR agonist

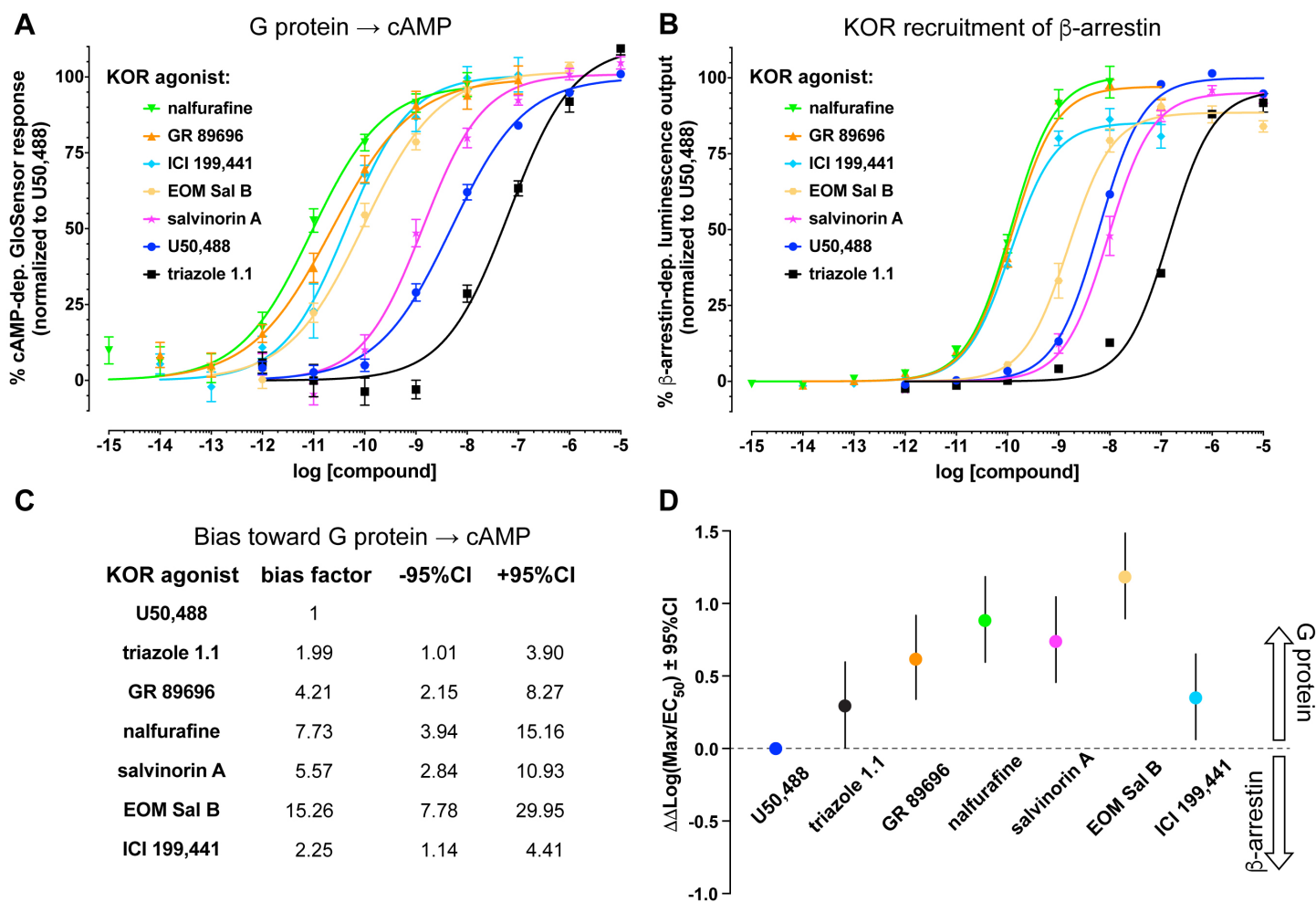


Figure 1. Nalfurafine is a potent G protein biased KOR agonist. Comparison of G protein and β -arrestin signaling outcomes with a variety of KOR ligands (at indicated concentrations) as compared to the reference standard U50,488. **(A)** G protein signaling was assessed with a GloSensor assay of cyclic AMP inhibition. **(B)** β -arrestin recruitment to the activated KOR was assessed via the TANGO assay. **(C,D)** Results from GloSensor and TANGO assays used to calculate Bias Factors for each compound from observed maximal efficacy (E_{\max} or “Max”) and potency (EC_{50}) values. \square \square $\text{Log}(\text{Max}/EC_{50})$ values, with their 95% confidence intervals (CIs), are plotted in panel D for each compound to indicate relative bias towards G protein signaling. All compounds were tested in triplicate alongside the reference standard U50,488. $N \geq 3$ for all compounds.

3.6.2 Spinal anti-nociception in C57BL/6J mice is equivalent at 5 mg/kg U50,488 and 0.03 mg/kg nalfurafine

tail immersion assay of spinal anti-nociception

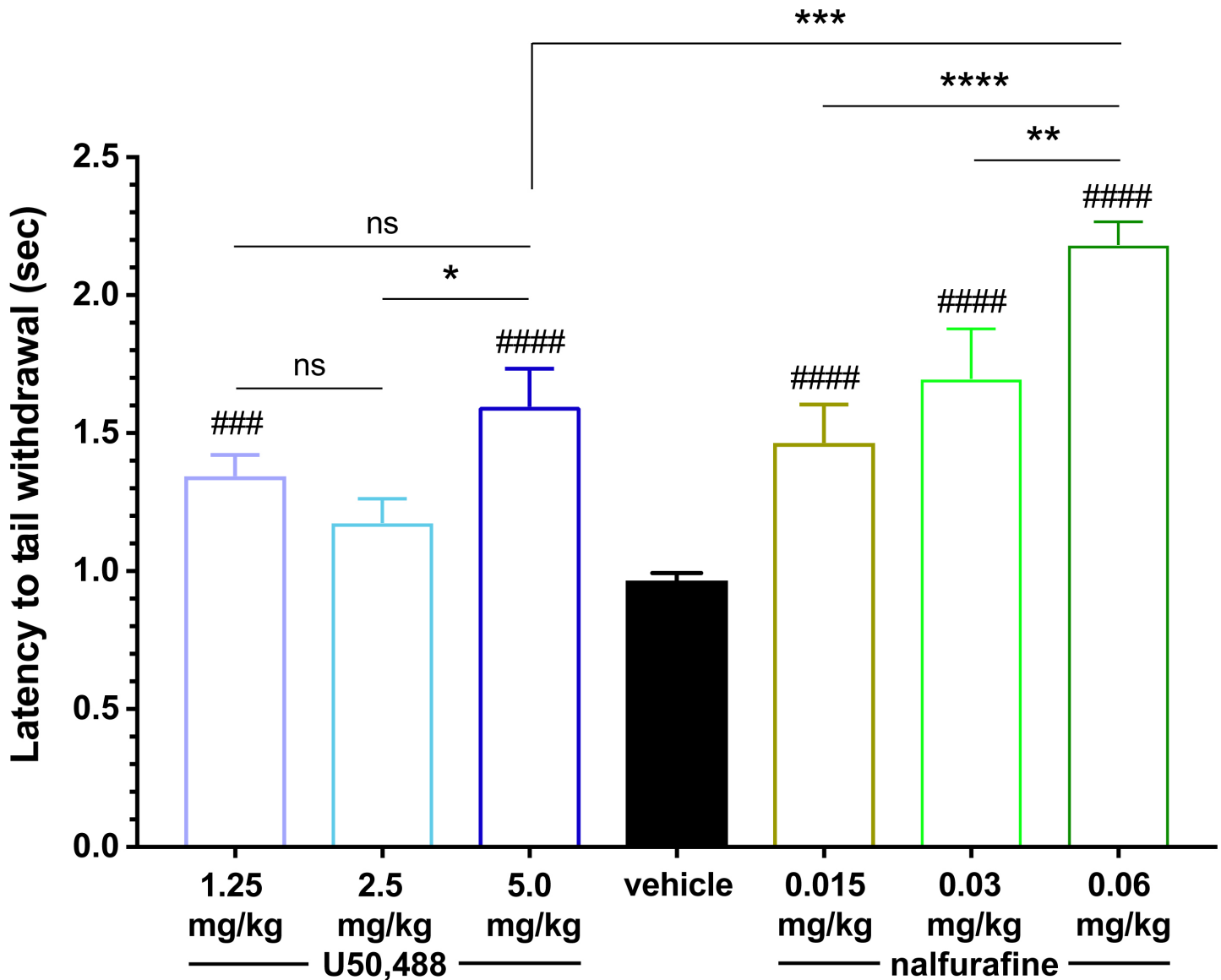


Figure 2. Spinal anti-nociception in C57BL/6J mice is equivalent at 5 mg/kg U50,488 and 0.03 mg/kg nalfurafine. Mice were treated with vehicle 30 min prior to establishing baseline latencies, then treated with indicated doses of nalfurafine or U50,488 30 min prior to testing. Data were analyzed by one-way ANOVA with Dunnett's multiple comparison *post hoc* test to assess for differences from the pooled vehicle. Holm-sidak multiple comparison *post hoc* test was used to assess for differences between treatment groups (n= 10-18 for all treatment conditions; ns, non-significant; *, $p < 0.05$; **, $p < 0.01$; ****, $p < 0.0001$; ###, $p < 0.001$; ####, $p < 0.0001$.)

3.6.3 Nalfurafine potentiates morphine-induced supraspinal analgesia in C57BL/6J mice

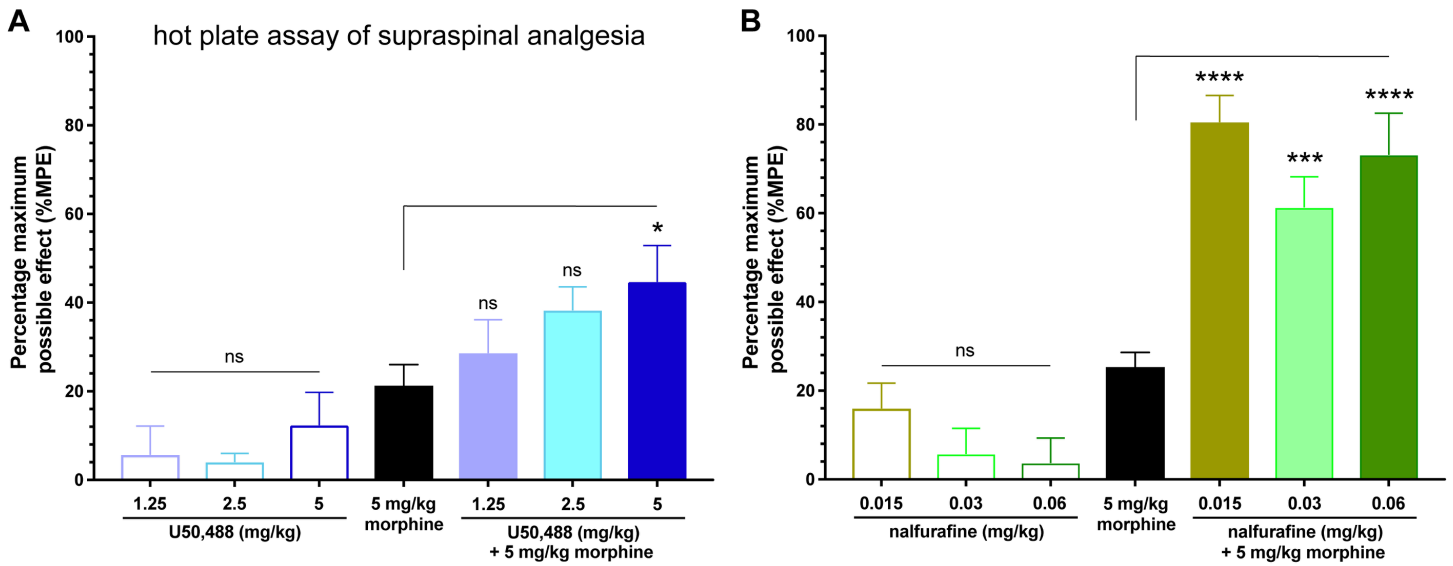


Figure 3. Nalfurafine potentiates morphine-induced supraspinal analgesia in C57BL/6J mice. Mice were treated with indicated doses of morphine, nalfurafine, U50,488, or a combination of morphine (5 mg/kg) with either nalfurafine or U50,488. Latency to nociceptive response was recorded and compared to the response to saline [%MPE = (test response - baseline)/(30 sec - baseline); thus, 100% MPE = 30 sec with no sign of nociception]. Data were analyzed by one-way ANOVA with Dunnett' multiple comparison *post hoc* test to assess differences from 5 mg/kg Morphine and Holm-sidak multiple comparison *post hoc* test to assess differences between treatment groups. ns, non-significant; *, $p < 0.05$; **, $p < 0.01$; ***, $p < 0.001$; **** or #####, $p < 0.0001$. **A.** 5 mg/kg morphine $n=16$; all other conditions $n=7-9$. **B.** 5 mg/kg morphine $n=17$; all other conditions $n=8-13$.

3.6.4 Nalfurafine treatment suppresses both spontaneous and morphine-induced locomotion

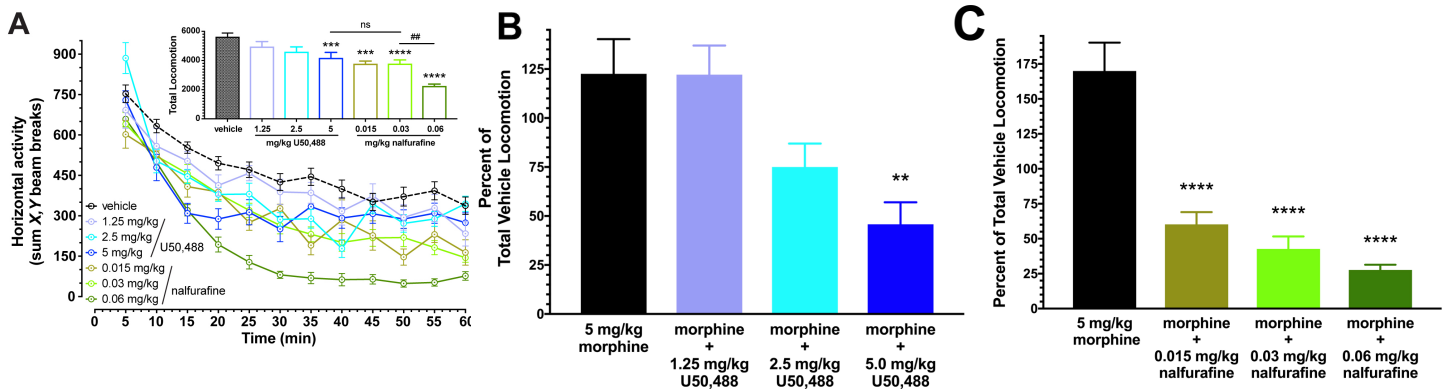


Figure 4. Nalfurafine treatment suppresses both spontaneous and morphine-induced locomotion. Assessment of open field locomotion elicited by environmental novelty or morphine administration. **(A)** For assessment of spontaneous locomotion, mice were administered vehicle or indicated single drug and immediately placed into the open field chamber for 60 min. **(B, C)** For assessment of morphine-induced locomotion, mice were again habituated to the chamber for 30 min (on Day 3) before receiving 5 mg/kg morphine with vehicle, or 5 mg/kg morphine with either 1.25, 2.5 or 5 mg/kg U50,488 (panel B), or 5 mg/kg morphine with either 0.015, 0.03 or 0.06 mg/kg nalfurafine (panel C) and assessed for locomotor behavior for an additional 60 min. Locomotion was measured by number of IR-beam breaks per 1 min. Data are presented as percent of pooled vehicle locomotion over the 60 min post injection for suppression of novelty induced locomotion. Total locomotion data were analyzed by one-way ANOVA with a Dunnett's *post hoc* analysis to compare treatment groups to the vehicle alone or vehicle + morphine condition and Tukey's *post hoc* analysis to assess difference between treatment conditions for novelty locomotion. Novelty locomotion n: vehicle= 41, all other conditions 7-16. Morphine suppressed locomotion n= 7-11 for all conditions. *, $p < 0.05$; **, $p < 0.01$; ***, $p < 0.001$; ****, $p < 0.0001$. ##, $p < 0.01$; #, $p < 0.05$.

3.6.5 Nalfurafine produces conditioned place aversion (CPA) in C57BL/6J mice at doses that do not reduce morphine CPP

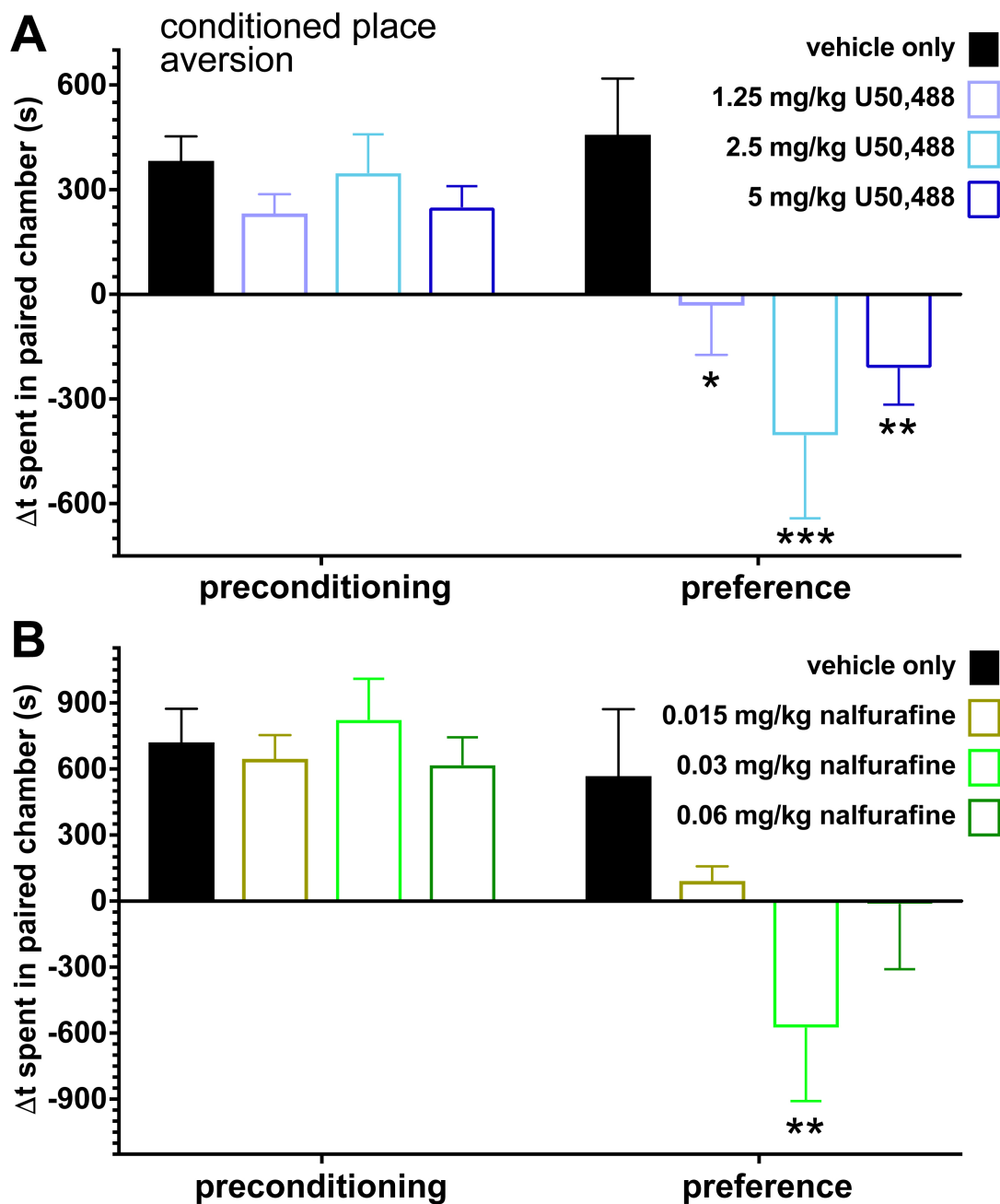


Figure 5. Nalfurafine produces conditioned place aversion (CPA) in C57BL/6J mice at doses that do not reduce morphine CPP. Mice were acclimated to the two-chambered apparatus and allowed to display a side preference. Then, mice were treated with saline (in their non-preferred chamber) or drug (in their preferred chamber) and confined to that chamber. Vehicle and drug were given on alternate days for four days. On test day, the mice were allowed free access to either chamber. The preference for the drug-paired chamber was measured as the time spent in the drug-paired chamber minus the time spent in the vehicle-paired chamber. (A)

Mice treated with U50,488 displayed a significantly reduced preference for the drug-paired chamber (**, $p < 0.01$; ****, $p < 0.0001$, ***, $p < 0.0004$). Vehicle, $n=8$; 1.25 mg/kg, $n=15$; 2.5 mg/kg U50, $n=9$; 5 mg/kg U50,488, $n= 11$. **(B)** Mice treated with 0.03 mg/kg nalfurafine, but not 0.015 or 0.06 mg/kg nalfurafine, displayed reduced preference for the drug-paired chamber. Vehicle, $n=20$; 0.015 mg/kg, $n=8$; 0.03 mg/kg nalfurafine, $n= 12$; 0.06 mg/kg nalfurafine $n=14$. Data were analyzed by one-way ANOVA with a Dunnett's multiple comparisons test comparing to vehicle (*) or a Holm-Sidak test to compare between doses of U50,488 or nalfurafine. **, $p < 0.01$; ***, $p < 0.001$; ****, $p < 0.0001$.

3.6.6 Nalfurafine, like the conventional KOR agonist U50,488, reduces morphine-induced conditioned place preference (CPP) in C57BL/6J mice

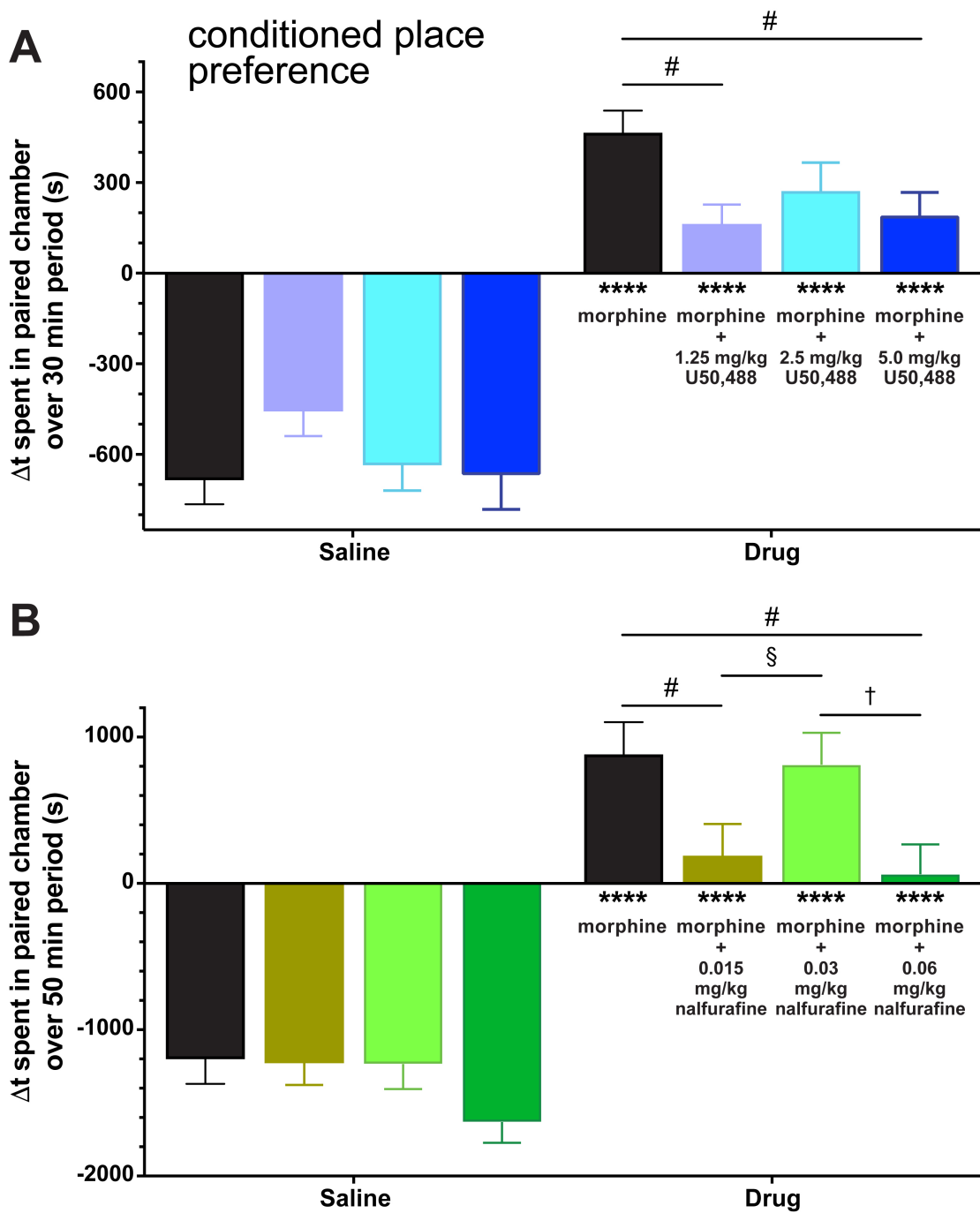


Figure 6. Nalfurafine, like the conventional KOR agonist U50,488, reduces morphine-induced conditioned place preference (CPP) in C57BL/6J mice. Mice were acclimated to the two-chambered apparatus and allowed to display a side preference. Then, mice were treated with vehicle (in their preferred chamber) or drug (in their non-preferred chamber) and confined to that chamber for 30 or 50 min (for U50,488 or nalfurafine, respectively). Vehicle or morphine was given on alternate days for four days. On test day, the

mice were allowed free access to both chambers. The preference for the drug-paired chamber was measured as the time spent in the drug-paired chamber minus the time spent in the vehicle-paired chamber. **(A)** Mice co-administered morphine (5 mg/kg) and U50,488 (1.25 or 5 mg/kg) displayed reduced preference for the drug-paired chamber. Morphine alone (+ vehicle), $n = 25$; Morphine plus U50,488, $n = 18-21$. **(B)** Mice co-administered morphine (5 mg/kg) and nalfurafine (0.015 or 0.06 mg/kg) displayed reduced preference for the drug-paired chamber. Morphine alone (+ vehicle), $n = 23$; Morphine plus nalfurafine, $n = 18-26$. Data were analyzed by two-way ANOVA with a Sidak's multiple comparisons test comparing to preconditioning (*), a Dunnett's test comparing to morphine alone (#), or a Holm-Sidak test to compare between doses of U50,488 or nalfurafine co-administration (††). #, $p < 0.05$; ††, $p < 0.01$; ****, $p < 0.0001$.

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Chapter Four

CONCLUSIONS AND FUTURE DIRECTIONS

4.1 – SNPs identified in a West Virginian population of patients with OUD

4.1.1 – Significant Results

Our analysis of a local cohort of patients in treatment for OUD identified several SNPs, out of the fourteen tested, with significantly different minor allele frequencies (MAFs) from the reference population. Of particular interest to the Siderovski lab is the finding that the rs4606 SNP within *RGS2* displayed increased prevalence of the minor allele in the test population, given Dr. Siderovski's role in the first cloning of the human *RGS2* cDNA in the 1990s (Siderovski et al., 1990; Siderovski et al., 1994; Wu et al., 1995) and first reporting the anxiety phenotype of mice lacking *Rgs2* expression (Oliveira-Dos-Santos et al., 2000). This SNP

within *RGS2* has since been associated with anxiety disorders (Smoller et al., 2008; Stein et al., 2014) and suicidality (Cui et al., 2008) in humans. Studies of reduced *RGS2* expression in mice have also demonstrated increased depressive and anxiety phenotypes (Lifschytz et al., 2012). As discussed previously, stress is a motivator of drug seeking behavior in rodents and anxiety disorders are commonly associated with substance use disorder (SUD) diagnoses (Brady et al., 2013) in humans, providing a plausible rationale for the association of the identified *RGS2* SNP with the WV OUD patient population.

The presence of the minor allele of the rs1799971 SNP within *OPRM1*, has previously been associated with increased odds for heroin addiction and decreased analgesic sensitivity (Bart et al., 2004; Tan et al., 2009). Assessment of this polymorphism in cell culture shows decreased mu opioid receptor expression yet increased affinity for β -endorphin (Bond et al., 1998) as well as engendering increased heroin self-administration in a transgenic mouse model (Zhang et al., 2014). Interestingly, our WV OUD patient population displayed a decreased prevalence of this polymorphism compared to the control population. As our patient cohort represents a population that is not in active addiction and has begun medication assisted therapy (MAT), this finding may indicate an interaction between the rs179971 SNP of *OPRM1* and the persistence of active addiction compared to the willingness to enter treatment. Comparisons of SNP frequency within a population of active users would inform this hypothesis. The Milan Puskar Health Right free clinic in Morgantown runs one of the only needle exchange programs in the state of WV. With the goal of reducing the harm related to drug use, the clinic provides clean needles and offers counseling to those in active addiction. The needle exchange could be used to gain access to a population of active heroin users from the same geographic region as those used for the genetics studies presented in this dissertation. Should the minor allele of the rs179971 SNP within *OPRM1* appear with greater frequency in this population of active heroin users than those assessed receiving MAT, it may indicate an association of the minor allele with reduced severity of addiction.

4.1.2 – Future Directions

The foremost limitations of the present study are the low sample size and restricted patient population. Recruitment of treatment centers outside of Morgantown and including additional border states within the Appalachian region would allow for both greater statistical power and patient stratification based on ethnicity as well as medical history (*e.g.*, first diagnosis of OUD vs. history of OUD). Considering the relationship of the kappa opioid receptor to stress (Massaly et al., 2016; Tejada and Bonci, 2018), inclusion of KOR (*OPRK1*) polymorphisms could provide important insight into the interactions between these two systems. At least one study (Kumar et al., 2012) has shown that the rs16918875 polymorphism of KOR increases addiction risk when patients also express the A118G (rs1799971) SNP of *OPRM1*, supporting the need to assess multiple components of addiction-related circuitry when assessing addiction risk. In addition, buprenorphine, a MOR partial agonist and KOR antagonist, is commonly used in MAT for opioid addiction (Marshalek and Sullivan, 2010; Zheng et al., 2016). Polymorphisms at KOR may potentially confer variations in the efficacy of buprenorphine for this purpose, providing a possible mechanism for screening patients into high and low responder groups before beginning therapy.

4.2 – Preclinical assessment of the G protein-biased KOR agonists

4.2.1 – Significant Results

The findings of these studies support the further investigation of G protein-biased KOR agonists as dose-sparing adjuvants to traditional MOR-targeting analgesics. Nalfurafine was shown *in vitro* to possess significant bias towards G protein-signaling at KOR and contrary to the unbiased agonist KOR U50,488, augmented the analgesic effect of morphine in the hot plate assay. This augmentation was observed at doses of KOR agonist that produced equivalent results in the tail-immersion assay of spinal nociception. As the analgesia provided by KOR agonism in the tail immersion assay is G protein dependent (Hernandez et al., 1995; Berg et al., 2011; White et al., 2015; Abraham et al., 2018), the augmented analgesia seen in the hot plate assay when co-administering nalfurafine with morphine may be a result of decreased β -arrestin signaling emanating from KOR agonism at either spinal or supraspinal sites.

Sedation (as measured in mice by hypolocomotion) and dysphoria (as measured in mice by conditioned place aversion) are two effects of KOR agonists that have limited their therapeutic use in humans (Kumor et al., 1986; Zacny et al., 1994; Walsh et al., 2008). While others have reported a lack of CPA to nalfurafine (Liu et al., 2018), our results demonstrate that aversion was present at 0.03 mg/kg nalfurafine, and that locomotion was reduced with all tested doses of nalfurafine when given alone or co-administered with morphine. A dose of 0.015 mg/kg nalfurafine, however, did not produce significant CPA, a result supported by previous work (Liu et al., 2018). As all doses of nalfurafine tested produced equivalent augmentation of morphine analgesia in the hot plate assay, the possibility remains that further dose optimization to avoid both the sedative and aversive effects of nalfurafine may be possible. While the unbiased KOR agonist U50,488 did not produce locomotor sedation to as great an extent as nalfurafine, all doses of U50,488 tested produced significant CPA. Taken together, these findings suggest that the sedative effects of KOR agonists occur via G protein-mediated signaling, while the aversive effects arise downstream of β -arrestin signaling. This interpretation, however, is complicated by work utilizing *BARR2* knockout mice that display both locomotor sedation and aversion when treated with unbiased KOR agonists (White et al., 2015). Further work is needed to determine if these effects truly arise via β -arrestin independent mechanisms or if there is compensation via β -arrestin1 in these mice.

The effects of nalfurafine on morphine-induced CPP are notable. KOR agonists reduce CPP for a variety of drugs of abuse (Kuzmin et al., 1997; Lindholm et al., 2001; Tsuji et al., 2001; Tao et al., 2006; Morani et al., 2009a; Ewald et al., 2017a; Townsend et al., 2017a), providing the possible additional benefit of reducing addiction liability with a co-administration paradigm. While both U50,488 and nalfurafine reduced morphine-induced place preference in our studies, each KOR agonist failed to do so with the middle dose of the ranges tested; *i.e.*, the dose of each drug that produced the greatest CPA. A reduction in place preference when co-administering the highest dose of either KOR agonist may be attributable to the induction of cognitive deficits known to occur with high doses of KOR agonists (Castellano et al., 1988; Carey et al., 2009), also potentially explaining the lack of CPA seen with the highest dose of nalfurafine. The persistence of morphine's rewarding effects, however, when co-administered with the most aversive dose of either drug is intriguing. A common

interpretation of the anti-rewarding properties of KOR agonists is that of positive punishment (Shippenberg et al., 1998; Wee and Koob, 2010), or an addition of aversive effects rather than the diminishment of positive effects produced by the rewarding drug (negative punishment). Rats pretreated with morphine for five consecutive days display an increased place preference for morphine, an effect that is not reduced with the addition of the KOR agonist U69593 during pretreatment (Shippenberg et al., 1998). This result supports the notion that the rewarding properties of morphine and the aversive properties of U69593 function through separate circuits when co-administered, producing a reduced preference at the level of the whole organism without altering the underlying biochemical effects of each drug.

This theory, however, would predict that doses of KOR agonist producing greater dysphoria (as measured by CPA) would produce increased levels of punishment and therefore suppress morphine-induced CPP to a greater extent. Our present results indicate the opposite effect, that doses of nalfurafine and U50,488 producing the greatest aversion reduce place preference for morphine the least. In addition, the lowest dose of both drugs (that show reduced levels of aversion) successfully reduced morphine-induced CPP. If KOR agonists do indeed reduce CPP through engagement of a punishment circuit rather than inhibition of a rewarding circuit engaged by morphine, then timing and intensity of engagement of these circuits may play an integral role in the behavioral outcomes observed. Extensive manipulation of dose of both KOR agonist and morphine, as well as alterations of pretreatment intervals will be needed as an initial step in investigating the complex interaction among the affective responses to each of these drugs.

4.2.2 – Future Directions

The most clinically relevant issue arising from the results of these studies centers on identifying appropriate dose combinations of nalfurafine and co-administered opioid analgesic (Ueno et al., 2013; Kozono et al., 2018; Siderovski and Setola, 2018). As nalfurafine is currently used in Japan with a low incidence of therapeutically limiting side-effects, clinical investigations into the viability of its use as a dose-sparing adjuvant face fewer hurdles than any other G protein-biased KOR agonist used in pre-clinical studies. While our

work supports the possibility that reduced dosing of both morphine and nalfurafine may achieve analgesia equivalent to a higher dose of morphine alone, this result needs to be confirmed experimentally by further adjusting the dose range of nalfurafine in addition to the dose range of morphine. Other nociceptive assays should be included as well to assess for therapeutic relevancy beyond thermal pain. MOR-targeting analgesics show reduced efficacy in treating inflammatory and neuropathic pain (Przewlocki and Przewlocka, 2005; McAlindon et al., 2014), while these conditions constitute the major reason for chronic out-patient opioid prescriptions (Boudreau et al., 2009; Gomes et al., 2011). An increase in the ability to manage these types of pain while reducing the likelihood of addiction would be of major therapeutic benefit to the medical community. Other therapeutically limiting effects must also be investigated, such as constipation, a side effect of MOR agonists prevalent enough to prompt the development of naloxegol (a peripherally restricted MOR antagonist) and diuresis, a side effect of KOR agonism (Leander et al., 1985; Webster, 2015). In addition, while nalfurafine was shown to reduce acetic acid-induced writhing in mice (a pain-stimulated behavior), it did not restore ICSS thresholds depressed by acetic acid administration (a pain-suppressed behavior) (Lazenka et al., 2018). Furthermore, nalfurafine alone suppressed ICSS thresholds in the absence of a painful stimulus (Lazenka et al., 2018). These results seem to indicate that nalfurafine alone does not alleviate the negative affect produced by pain, unlike traditional opioid analgesics (Altarifi et al., 2014; Porreca and Navratilova, 2017b), and nalfurafine may prevent this positive effect of traditional opioid analgesics when co-administered.

While these proposed pre-clinical and clinical studies with nalfurafine will better elucidate the viability of this drug as a dose-sparing adjuvant, additional experiments are necessary to elucidate the true relevance of G protein and β -arrestin signaling in producing the observed effects. Conducting parallel studies with other G protein-biased KOR agonists with greater bias factors will inform this question. In addition, future studies using *BARR2* knockout mice will aid in determining which effects of G protein-biased KOR agonists are dependent upon this particular signaling pathway, as well as if any traditionally β -arrestin-dependent effects seen with putatively G protein-biased KOR agonists (*e.g.*, CPA to 0.03 mg/kg nalfurafine; Figure 3.6.5 above) are simply a result of a dose range that engages the β -arrestin pathway *in vivo* despite the ligand possessing a demonstrable

bias toward G protein signaling *in vitro*. Such studies would be greatly enhanced with the use of β -arrestin cell-type and region-specific knockout mice (*e.g.*, (Urs et al., 2016a; Zhu et al., 2017), allowing for more specific interrogation of the cell populations and neuronal circuits involved. A deeper understanding of just how MOR and KOR produce their euphoric and dysphoric effects, respectively, will hopefully allow for the design of targeted therapeutics that alleviate pain without addiction or dysphoria.

4.3 – References

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