Inspiratory rhythmogenic activity is burst-independent and opioid-sensitive

Xiaolu Sun[1*], Carolina Thörn Pérez[1*], Nagaraj Halemani D[2], Xuesi M. Shao[1], Morgan Greenwood[3], Sarah Heath[1], Jack L. Feldman[1], and Kaiwen Kam[2,†]

[1]Department of Neurobiology, David Geffen School of Medicine at UCLA, Los Angeles, CA 90095, USA
[2]Department of Cell Biology and Anatomy, Chicago Medical School, Rosalind Franklin University of Medicine and Science, North Chicago, IL 60064, USA
[3]RFUMS/DePaul Research Internship Program, Rosalind Franklin University of Medicine and Science, North Chicago, IL 60064, USA

*Equal contribution

[†]Corresponding author and Lead Contact:
    Kaiwen Kam, Ph.D.
    Department of Cell Biology and Anatomy
    Chicago Medical School
    Rosalind Franklin University of Medicine and Science
    3333 Green Bay Rd.
    North Chicago, IL 60064-3095, USA
    Email: kaiwen.kam@rosalindfranklin.edu
    Phone: (847) 578-8630
    Fax: (847) 578-3253

Summary

How mammalian neural circuits generate rhythmic activity in motor behaviors, such as

breathing, walking, and chewing, remains elusive. For breathing, rhythm generation can be

localized to a brainstem nucleus called the preBötzinger Complex (preBötC). Rhythmic preBötC

population activity consists of small amplitude burstlets, which we hypothesize are

rhythmogenic, and larger inspiratory bursts, which drive motoneuronal activity. If burstlets are

rhythmogenic, opioids, analgesics that can cause profound respiratory depression, should

similarly reduce burstlet frequency. In conditions where burstlets were separated from bursts

in medullary slices from neonatal mice, the μ-opioid receptor (μOR) agonist DAMGO decreased

burstlet frequency. DAMGO-mediated depression was abolished by genetic deletion of μORs in

a glutamatergic preBötC subpopulation and was reduced by Substance P, but not blockade of

inhibitory synaptic transmission. Our findings suggest that rhythmogenesis need not rely on

strong bursts of activity associated with motor output and point to strategies for ameliorating

opioid-induced depression of breathing.

Keywords: opioids, breathing, rhythm generation, central pattern generator

Introduction

Rhythmic motor behaviors in mammals, such as breathing, walking, and chewing, are controlled

by neural circuits that determine the period of the movement and shape the pattern of motor

activity. Despite the apparent simplicity of the computation, the mechanisms underlying

rhythmogenesis in these circuits are unknown. Generation of breathing rhythm is localized to a

compact nucleus in the ventrolateral medulla, the preBötzinger Complex (preBötC; Smith et al.,

1991). In particular, two overlapping subpopulations of glutamatergic preBötC neurons, those

expressing the neurokinin-1 receptor (NK1R) and those derived from progenitors that express

the transcription factor Dbx1 (Dbx1$^+$), are necessary for inspiratory rhythmogenesis (Bouvier et

al., 2010; Gray et al., 2010; Gray et al., 2001).

Until recently, rhythmic bursts of preBötC population activity were considered unitary events

that determined both the timing and pattern of periodic inspiratory movements. However, we

recently showed that these preBötC bursts consist of two separable components: i) large

amplitude inspiratory bursts that are transmitted via premotoneurons to inspiratory

motoneurons to activate muscles that produce inspiratory airflow; and ii) low amplitude

burstlets that appear as preinspiratory activity when preceding these bursts and are not

normally seen in motor nerve or muscle activity (Kam et al., 2013a). Because preBötC rhythmic

activity can consist solely of burstlets that do not produce motoneuronal output, we

hypothesize that burstlets are rhythmogenic (Kam et al., 2013a). A strong prediction of this

hypothesis is that manipulations that affect inspiratory burst frequency should have similar

effects on preBötC burstlet frequency.

To test this prediction, we examined the effects of opioids, analgesics that depress breathing at

high doses, on bursts and burstlets and tested opioidergic interactions with other

neurotransmitters and neuromodulators that affect preBötC neuronal excitability. We bath-

applied [D-Ala^2,N-MePhe^4,Gly-ol^5]-enkephalin (DAMGO), a potent synthetic μ-opioid receptor

(μOR) agonist, in conditions of low excitability *in vitro* when burstlets appeared at times bursts

would have been expected or when preBötC rhythmic activity consisted solely of burstlets. We

measured the frequency of inspiratory-related preBötC population activity, i.e., the intervals

between successive bursts or burstlets, to assay opioidergic effects on rhythmogenesis and

calculated the fraction of rhythmic preBötC events that were burstlets to assay changes in the

pattern-generating threshold mechanism that converts burstlets into bursts. We conclude that

DAMGO depresses inspiratory frequency by acting on burstlet producing preBötC Dbx1$^+$

neurons. These data are consistent with our hypothesis of distinct rhythm and pattern

generating mechanisms within the preBötC, with rhythmogenesis being mediated by burstlets

and a separable burst-generating mechanism governing patterned preBötC output. Our results

also inform potential approaches for combating opioid-induced depression of breathing.

Results

DAMGO slows preBötC rhythmic activity without affecting burstlet fraction.

Systemic administration of opioids slows breathing *in vivo* and *in vitro* (Boom et al., 2012; Gray

et al., 1999). We investigated how preBötC burstlets and bursts mediate these opioidergic

effects in neonatal mouse tissue slices that contain the preBötC and the hypoglossal motor

nucleus and nerve (XII) and generate a physiologically relevant motor output, i.e., inspiratory-

related rhythmic activity. Under baseline conditions in artificial cerebrospinal fluid (ACSF)

containing 9 mM K$^+$ and 1.5 mM Ca^{2+} ("9/1.5"), preBötC population activity and XII activity

consisted primarily of coincident, large amplitude rhythmic bursts (Fig. 1A; Smith et al., 1991).

Subsaturating concentrations of DAMGO caused a dose-dependent decrease in the frequency

(*f*) of rhythmic preBötC and XII bursts (Fig. 1A-C). 10 nM DAMGO significantly decreased

preBötC *f* from 0.22 ± 0.05 Hz to 0.13 ± 0.04 Hz (p=0.01, n=6; Fig. 1C). 30 nM DAMGO

application further decreased preBötC *f* to 0.07 ± 0.04 Hz (p=0.0001, n=6; Fig. 1C), and 100 nM

DAMGO led to complete cessation of XII activity and, in some cases, also blocked all preBötC

activity (Fig. 1B).

We previously showed that changes in XII *f* induced by decreases in extracellular K$^+$ could be

caused by a reversion of preBötC bursts to preBötC burstlets (Kam et al., 2013a). This was not

the case for the effects of DAMGO on *f*, as the fraction of total preBötC events that were

burstlets (burstlet fraction), which was very low in control conditions (0.14 ± 0.14), was not

significantly different from that in the presence of 30 or 100 nM DAMGO (p=0.8, n=6; Fig. 1A,

C). Moreover, the amplitude of preBötC bursts (0.60 ± 0.10 a.u.) was not significantly different

between control conditions without DAMGO and when 10 nM (0.61 ± 0.13 a.u.) or 30 nM

DAMGO (0.56 ± 0.19 a.u.) was added to the bath (p=0.8, n=6).

To determine whether DAMGO could affect the conversion of burstlets into bursts when the

preBötC was operating in a different dynamic mode, we decreased ACSF K^+ and Ca^{2+}

concentrations (Kam et al., 2013a). In 3 mM K^+ and 1 mM Ca^{2+} ("3/1"), preBötC activity was a

mixed pattern of burstlets and bursts (Fig. 1D, upper traces). While the burstlet fraction under

these conditions (0.58 ± 0.26) was significantly higher than that in 9/1.5 ACSF (0.14 ± 0.14;

p=0.003, n= 6 for 9/1.5, n=7 for 3/1), preBötC f was similar (0.23 ± 0.10 Hz in 3/1 vs. 0.22 ± 0.05

Hz in 9/1.5; p=0.8, n= 6 for 9/1.5, n=7 for 3/1; Fig. 1C, F). XII f in 3/1 ACSF (0.08 ± 0.05 Hz), on

the other hand, was significantly lower than XII f in 9/1.5 ACSF (0.2 ± 0.06 Hz; p=0.002, n= 6 for

9/1.5, n=7 for 3/1; Fig. 1B, E), a consequence of the increased fraction of preBötC burstlets that

were not transmitted to XII output.

In 3/1 ACSF, DAMGO decreased preBötC f in a dose-dependent manner, similar to its effects in

9/1.5 ACSF. Addition of 30 nM DAMGO significantly reduced preBötC f from 0.22 ± 0.1 Hz to

0.04 ± 0.04 Hz (p=0.001, n=7; Fig. 1D-F). This reduction was not significantly different from the

effects of 30 nM DAMGO in 9/1.5 ACSF (p=0.2, n=6 for 9/1.5, n=7 for 3/1; Fig. 1C, F). Higher

concentrations of DAMGO (≥100 nM) frequently blocked all preBötC population activity.

Burstlet and burst amplitudes in 3/1 ACSF (burstlet: 0.38 ± 0.05 a.u.; burst: 0.73 ± 0.08 a.u.) did

not change significantly when 10 nM DAMGO (burstlet: 0.36 ± 0.04 a.u.; burst: 0.78 ± 0.10 a.u.)

or 30 nM DAMGO (burstlet: 0.34 ± 0.11 a.u.; burst: 0.65 ± 0.21 a.u.) was bath-applied (burstlet:

p=0.7, burst: p=0.3, n=7), and the burstlet fraction was again unaffected by DAMGO (p=1.0,

n=7; Fig. 1F).

To determine whether opioids specifically affected rhythmogenic activity independent of burst

production, we isolated burstlets using low concentrations of Cd^{2+} (6-25 µM; Kam et al., 2013a),

which significantly increased the burstlet fraction from 0.42 ± 0.30 to 0.78 ± 0.20 (p=0.02, n=8)

without significantly changing preBötC f or amplitude (f: p=0.7, amplitude: p=0.06, n=8; Fig. 2A,

B). The decreases in excitability and synaptic efficacy caused by Cd^{2+} (Kam et al., 2013a; Lu et al., 2007) resulted in an increased sensitivity to DAMGO. Bath application of 1 nM DAMGO on this burstlet only rhythm was sufficient to abolish preBötC rhythmic activity completely in 7 of 8 slices. Concentrations of DAMGO below that causing rhythmic cessation (0.01-10 nM) slowed preBötC f significantly from 0.28 ± 0.09 Hz to 0.18 ± 0.07 Hz (p=0.001, n=8; Fig. 2A, B). This relative decrease in preBötC f (0.66 ± 0.16) was not significantly different from the effect of 10 nM DAMGO on preBötC f in 3/1 ACSF with no Cd^{2+} (0.63 ± 0.16; p=0.8, n=5 for 3/1, n=8 for Cd^{2+}). However, burstlet amplitudes in the presence of both Cd^{2+} and DAMGO (0.29 ± 0.08) were significantly reduced compared to Cd^{2+} alone (0.35 ± 0.09; p=0.003, n=8; Fig. 2A, B).

Depression of preBötC burstlet f by μOR activation points to DAMGO modulating preBötC rhythmogenic mechanisms. We previously hypothesized that the emergent rhythmogenic processes that produce burstlets manifested as the latency in ectopic burst initiation following targeted excitation of 4-9 preBötC inspiratory neurons, whose molecular identity was unspecified (Kam et al., 2013b). Here, in slices from Dbx1^{+} reporter (*Dbx1^{cre};Rosa26^{tdTomato}*) mice, we targeted preBötC Dbx1^{+} neurons for holographic photostimulation to determine if opioids also modulate this rhythmogenesis-related process (Fig. 2C-F). In control 9/1.5 ACSF, ectopic bursts were reliably elicited when 5-10 Dbx1^{+} neurons were targeted (probability of success: 0.86 ± 0.08, n=6; Fig. 2D). When a set of neurons that reliably resulted in bursts in response to photostimulation in control conditions were then excited in 30 nM DAMGO, the probability of eliciting a burst was reduced to 0.48 ± 0.25 (n=4; Fig. 2D, E). In 100 nM DAMGO, the success rate decreased significantly as stimulation of the threshold set that had previously reliably triggered bursts was further compromised and, in some cases, failed to trigger a single burst (p=0.006, n=6; Fig. 2D, E). In 200 nM DAMGO, stimulation was unsuccessful at eliciting bursts (p=0.0003, n=6; Fig. 2E). Importantly, the latency between laser stimulation and burst initiation in the cases when bursts were triggered in DAMGO (157 ± 22 ms) was significantly increased compared to that in baseline conditions (105 ± 24 ms; p=0.002, n=5; Fig. 2F).

μOR-expressing Dbx1^{+} neurons mediate the rhythmogenic effects of opioids.

Which preBötC neurons does DAMGO target to depress inspiratory rhythm? We hypothesized

that DAMGO was acting on preBötC Dbx1$^+$ neurons, which are essential for respiratory rhythm

generation (Bouvier et al., 2010; Gray et al., 2010; Wang et al., 2014) and contain µOR mRNA

(Hayes et al., 2017). We first sought to determine whether these neurons express µOR protein

using immunohistochemistry. In *Dbx1^{Cre};Rosa26^{tdTomato}* mice, µOR protein expression

overlapped with the spatial distribution of tdTomato-expressing Dbx1$^+$ neurons in preBötC and

the adjacent intermediate band of the reticular formation (IRt; Fig. 3A) and was localized to

individual preBötC Dbx1$^+$ neurons (Fig. 3B-C).

To test whether µORs in Dbx1$^+$ neurons mediated DAMGO inhibition, we used a

Dbx1^{Cre};Oprm1$^{fl/fl}$ mouse line (Weibel et al., 2013), in which µORs are genetically deleted in

Dbx1$^+$ neurons. We first examined whether preBötC µOR expression was knocked down in P2

Dbx1^{Cre};Oprm1$^{fl/fl}$ mice compared to their littermate controls (Fig. 3D-I). *Dbx1* is not expressed

postnatally (Bouvier et al., 2010; Gray et al., 2010), and NK1R protein is predominantly found in

neuronal processes (Tan et al., 2010) while NK1R mRNA is sparsely expressed in somata (Fig.

S1). We therefore localized preBötC using tachykinin precursor peptide (TAC1) expression,

which colocalizes with NK1R, ventral to nucleus ambiguus (NA) as a marker for preBötC (Fig. S1;

Hayes et al., 2017). To measure µOR and TAC1 expression, we performed fluorescence *in situ*

hybridization, which avoids the non-specific labeling that may occur with some µOR antibodies

(Schmidt et al., 2013) and allows localization of TAC1 expression to somata (Fig. S1). Abundant

(>15 puncta) TAC1 mRNA was found in larger neurons confined to the preBötC (somatic

diameter: 27.08 ± 4.3 µm, n=121). Choline acetyl transferase (ChAT) mRNA probes were used to

mark NA. In *Dbx1$^{+/+}$; Oprm1$^{fl/fl}$* control mice, µOR mRNA puncta colocalized with TAC1 and NK1R

mRNA in the preBötC (Figs. 3D-F, S1), recapitulating the immunohistochemical distribution of

Dbx1$^+$ neurons in the Dbx1$^+$ reporter mice (Fig. 3A-C). Compared with µOR expression in

Dbx1$^{+/+}$; Oprm1$^{fl/fl}$ mice (Fig. 3D-F), µOR expression in *Dbx1^{Cre}; Oprm1$^{fl/fl}$* mice was reduced in

regions where Dbx1$^+$ neurons are normally found (Fig. 3D, G) and significantly reduced in TAC1

positive preBötC neurons (p=0.01, *Dbx1$^{+/+}$; Oprm1$^{fl/fl}$*: H-score = 139.07 ± 39.33, n = 5; *Dbx1^{Cre};*

Oprm1$^{fl/fl}$: H-score= 47.98 ± 19.77, n=3; Fig. 3G-I).

Having confirmed a decrease in µOR expression in preBötC in *Dbx1^{Cre}; Oprm1$^{fl/fl}$* mice, we

determined whether the functional effects of DAMGO on preBötC rhythm were similarly

affected. In rhythmic tissue slices from $Dbx1^{Cre}$; $Oprm1^{fl/fl}$ mice bathed in 9/1.5 ACSF, 100 nM

DAMGO had little effect on preBötC f (p=0.2, n=9; Figs. 3J, S2). In contrast, bath-applied

DAMGO in $Dbx1^{+/+}$; $Oprm1^{fl/fl}$ cre-negative littermates reduced preBötC f significantly (p=0.01,

n=6; Fig. 3J).

DAMGO-mediated depression is diminished by Substance P, but not blockade of inhibitory

synaptic transmission.

The coexpression of μOR and NK1R in preBötC Dbx1$^+$ neurons raises the possibility that NK1R

activation, which increases respiratory frequency in wild type rodents (Gray et al., 1999; Yeh et

al., 2017), could be effective in reducing the depressant effects of DAMGO. When applied

together in 3/1 ACSF, 30 nM DAMGO was much less effective in reducing rhythmic activity in

the presence of the NK1R agonist Substance P (SP; 500 nM), and preBötC f was not significantly

different from that in control 3/1 ACSF conditions, i.e., in the absence of SP and DAMGO

(p=0.09, n=6; Fig. 4A,C). We next tested whether altering preBötC excitability by blockade of

inhibitory synaptic transmission using a cocktail of the GABA$_A$ receptor antagonist picrotoxin

and the glycine receptor antagonist strychnine (picrotoxin/strychnine; Del Negro et al., 2009;

Ren and Greer, 2006; Shao and Feldman, 1997) would similarly block DAMGO-mediated

depression. In contrast to SP, picrotoxin/strychnine (100 μM/1 μM) did not reduce the effects

on preBötC f of 30 nM DAMGO compared to control conditions in either 9/1.5 or 3/1 ACSF

(p=0.2, n=6 for 9/1.5, n=7 for 3/1, n=13 for picrotoxin/strychnine; Fig. 4B, C). Indeed, preBötC f

when SP was applied with 30 nM DAMGO was greater than that in 9/1.5 ACSF, with or without

picrotoxin/strychnine, or 3/1 ACSF in the presence of DAMGO (p=0.001, n=6 for 9/1.5, n=7 for

3/1, n=12 for picrotoxin/strychnine, n=6 for SP; Fig. 4C).

Are the differential efficacies of SP and blockade of inhibitory transmission expressed in their

effects on baseline rhythmic activity? To determine how SP and blockade of inhibitory

transmission affect rhythm and pattern generation, we tested the effects of SP and

picrotoxin/strychnine on preBötC activity in 3/1 ACSF without DAMGO. Under these conditions,

bath-applied SP (500 nM) significantly reduced the burstlet fraction from 0.47 ± 0.3 to

0.10 ± 0.12 (p=0.01, n=6), but, surprisingly, did not affect preBötC f (p=0.4, n=6; Fig. 4D, E). In

slices from *Dbx1^{Cre}; Oprm1$^{fl/fl}$* mice bathed in 9/1.5 ACSF, we confirmed that 500 nM SP

significantly increased preBötC f (Fig. S2) and modulated preBötC f independent of μORs, which

were deleted in Dbx1$^+$ neurons in these slices. In 3/1 ACSF, bath-applied picrotoxin/strychnine

(100 μM/1 μM) also significantly decreased burstlet fraction (p=0.0001, n=13) and did not alter

preBötC f significantly (p=0.9, n=13; Fig. 4F, G).

Discussion

Inspiratory rhythm generation must be both robust, to operate almost continuously throughout

the lifetime of the animal, and labile, to adapt to changing metabolic and environmental

conditions and coordinate with other behaviors. How the preBötC generates rhythm to meet

these challenges has thus far defied a thorough understanding. We put forth a hypothesis that

respiratory rhythmogenesis is driven not by strong bursts, but by preBötC burstlets (Fig. 4H;

Kam et al., 2013a). Burstlets occur when spontaneous firing among preBötC neurons reaches a

threshold for synchrony, and the time required for preBötC neurons to assemble and achieve

this synchrony (percolation) determines the period of the rhythm (Kam et al., 2013a; S.A. and

J.L.F., in preparation).

A strong prediction of this hypothesis is that manipulations that depress burst frequency should

also slow burstlet frequency. Indeed, we found that DAMGO specifically slowed preBötC f

without affecting burstlet fraction when the rhythm: i) was composed of mostly bursts; ii) was a

mixed pattern of bursts and burstlets; or iii) was mostly (or all) burstlets (Fig. 4H). Additionally,

DAMGO prolonged the latency between photostimulation and burst generation when threshold

sets of ≤10 neurons were excited. The congruent effects of DAMGO on both burstlet f and

photostimulation latency: i) is consistent with our hypothesis that both are products of

percolation of activity within the preBötC microcircuit, a critical process for rhythmogenesis

(Kam et al., 2013a; Kam et al., 2013b), and ii) suggests that DAMGO directly affects

rhythmogenic mechanisms.

Consistent with our functional studies, we found abundant μOR expression in preBötC Dbx1$^+$

neurons that form the core circuitry for inspiratory rhythmogenesis (Wang et al., 2014). We

also showed colocalization of μOR with NK1R in some preBötC neurons as well as with a

partially overlapping preBötC subpopulation expressing TAC1 that may be one endogenous

source of SP (Liu et al., 2004). When μORs were genetically deleted in Dbx1$^+$ neurons in *Dbx1^{Cre};*

Oprm1$^{fl/fl}$ mice, the potent rhythmic slowing effect of DAMGO in slices was no longer present.

These data suggest that μORs in preBötC Dbx1$^+$ neurons mediate DAMGO inhibition of

rhythmicity.

How could μOR activation affect the percolation of activity in the preBötC that we hypothesize

underlies rhythmogenesis and transform, i.e., slow down, the dynamics of the Dbx1$^+$ preBötC

network? μORs can be localized postsynaptically on dendrites and cell bodies, where they

regulate neuronal excitability and transduce receptor activation to downstream signal

transduction pathways; μORs are also found presynaptically on axon terminals where they

inhibit neurotransmitter release via activation of K$^+$ conductances and/or inhibition of Ca^{2+}

conductances (Le Merrer et al., 2009; Mansour et al., 1988; Montandon et al., 2016; Williams et

al., 2001). We suggest that DAMGO-mediated hyperpolarization and/or decreases in synaptic

probability of release in and among μOR-expressing preBötC Dbx1$^+$ neurons reduce

spontaneous firing rate, weaken the propagation of action potentials to postsynaptic neurons,

and increase the number of inputs required to reach action potential threshold. These effects

reduce the moment-to-moment probability that a critical number of neurons will be

simultaneously active and prolong the time required for preBötC neurons to increase their

firing, achieve synchrony, and generate burstlets (S.A. and J.L.F., in preparation).

The pattern-generating mechanism that converts burstlets to bursts may, by analogy, require a

higher threshold of activity and synchrony be reached via independently regulated, distinct

mechanisms, such as the activation of persistent inward conductances (Picardo et al., 2019).

Through this threshold mechanism, SP and blockade of inhibitory synaptic transmission

increase, whereas Cd^{2+} decreases, the fraction of preBötC events that are burstlets, without

altering frequency of these events (Fig. 4H). Interestingly, the depressant effects of DAMGO

were reduced by SP, but not inhibitory blockade. Inhibitory synaptic activity in preBötC is

dispensable for generation of breathing rhythm *in vivo* and *in vitro* (Janczewski et al., 2013;

Shao and Feldman, 1997) and has variable effects on excitability (Ren and Greer, 2006), which

may account for the inability of picrotoxin and strychnine to mitigate DAMGO-induced

depression. Alternatively, inhibition may act elsewhere in the microcircuit or upstream of

opioidergic signaling pathways within Dbx1^{+} neurons.

While inhibitory blockade specifically affected pattern generation and was unable to rescue

DAMGO-mediated depression, NK1R activation seems capable of modulating both rhythm and

pattern in a state-dependent manner that can offset the effects of DAMGO (Fig. 4H). A ceiling

effect, i.e., preBötC f being near or at maximal, may have limited the rhythmogenic effect of SP

in 3/1 ACSF alone, but did reveal a pattern generating effect on burstlet fraction. Consistent

with SP effects in 9/1.5 ACSF (Gray et al., 1999; Yeh et al., 2017), co-application of opioids and

SP demonstrated a role for NK1R activation in modulating rhythm that is likely mediated by

intracellular interactions in preBötC Dbx1^{+} neurons that coexpress µOR and NK1R. The µOR

expression and DAMGO sensitivity of NK1R-expressing neurons are well-documented (Gray et

al., 1999; Montandon et al., 2011), and G_s-protein coupled signaling of NK1R opposes the $G_{i/o}$-

coupled signaling of µORs (Gray et al., 1999; Johnson et al., 1996). However, the interaction of

NK1R and µOR can be complicated. Opioids may inhibit SP release, and SP potentiates the

antinociceptive effects of morphine through release of endogenous opioids (Fukazawa et al.,

2007; Kream et al., 1993). SP may also act directly on µORs as a weak agonist since naloxone,

the µOR antagonist, blocks SP excitation in some neurons (Davies and Dray, 1977).

Nonetheless, we predict that only neuropeptides and neuromodulators, whose receptors are

coexpressed with µORs in rhythmogenic preBötC neurons, like NK1Rs, are able to modulate

µOR signaling intracellularly to mitigate DAMGO-mediated depression.

The interaction between excitatory neuromodulatory systems with opiates in respiratory neural

circuits is relevant for addressing the depressive effects of opiates on breathing. Opioids are the

most commonly prescribed drugs for severe acute and chronic pain and play an important role

in palliative care. However, with a high potential for addiction, opiate overprescription and

abuse has created a pressing public health crisis that affects millions (CBHSQ, 2018). Morbidity

and mortality from opioid addiction and overdose is largely a result of opioid-induced

depression of breathing (Jaffe & Martin, 1990). Opiates given systemically will act on opioid

receptors throughout the nervous system (Kibaly et al., 2019), and µORs are expressed in

various brain structures regulating breathing (Gray et al., 1999; Manzke et al., 2003; Phillips et

al., 2012; Pokorski and Lahiri, 1981; Prkic et al., 2012; Zhang et al., 2007; Zhang et al., 2011).

While depression of breathing is unlikely to be dependent on actions at a single site (Lalley et

al., 2014; Montandon and Horner, 2014; Stucke et al., 2015), our data is consistent with

compelling *in vitro* and *in vivo* evidence that the preBötC is the most sensitive site to µOR

agonists and that it mediates respiratory frequency depression by opioids (Ballanyi and

Ruangkittisakul, 2009; Janczewski et al., 2002; Montandon and Horner, 2014; Montandon et al.,

2011).

In summary, the depressive effect of the µOR agonist DAMGO on preBötC burstlet frequency, in

the absence of a change in burstlet fraction, supports a primary role for burstlets in respiratory

rhythm generation and is consistent with the hypothesis that rhythm and pattern generating

mechanisms within the preBötC are separable. We propose that having distinct mechanisms

generating burstlets and bursts provides functional substrates for targeted modulation of the

rhythmic timing and/or pattern of breathing by other brain areas and neuromodulators. This

organization enables the dynamic temporal and pattern lability characteristic of breathing

essential for adaptation to changing metabolic demands and for coordination with other

respiratory-related and orofacial behaviors. A better understanding of the mechanisms of µOR

actions in respiratory neural circuits has important implications for developing stimulants that

excite breathing via non-opioidergic pathways (van der Schier et al., 2014) and drugs that

reverse opioid actions on preBötC Dbx1$^+$ neurons to maintain breathing without affecting

analgesia.

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and K.K. wrote the paper.

Declaration of Interests

The authors declare no competing interests.

Figure legends

Figure 1. μOR activation depresses inspiratory frequency without affecting burstlet fraction.

(A) Representative traces of ∫preBötC and ∫XII population activity in 9/1.5 ACSF control (ctrl) and 30 nM DAMGO. Scale bar, 10 s.

(B) Representative time course of an experiment where increasing concentrations of DAMGO were bath-applied in 9/1.5 ACSF. Interevent intervals (int.) in preBötC (top) and XII (bottom) are plotted. In this experiment, 100 nM DAMGO led to cessation of rhythm.

(C) Average burstlet fraction and preBötC f in 9/1.5 ACSF (ctrl) and 10 nM and 30 nM DAMGO. Increasing concentrations of DAMGO did not significantly affect burstlet fraction whereas preBötC f was significantly lower in 10 nM and 30 nM DAMGO. *, $p < 0.05$, One-way ANOVA, post-hoc Tukey test, n=6.

(D) Representative traces of ∫preBötC and ∫XII population activity in 3/1 ACSF control (ctrl) and 30 nM DAMGO. Both small amplitude burstlets (•), which did not produce XII activity, and large amplitude bursts, which generated XII bursts, were observed. Scale bar, 10 s.

(E) Representative time course of an experiment, where increasing concentrations of DAMGO were bath-applied in 3/1 ACSF. Interevent intervals in preBötC (top) and XII (bottom) are plotted. In this experiment, 100 nM DAMGO led to cessation of XII activity while preBötC rhythm persisted.

(F) Average burstlet fraction and f in 3/1 ACSF (ctrl) and 10 nM and 30 nM DAMGO. Increasing concentrations of DAMGO did not significantly affect burstlet fraction whereas preBötC f was significantly lower in 30 nM DAMGO. *, $p < 0.05$, One-way ANOVA, post-hoc Tukey test, n=7.

Figure 2. DAMGO specifically modulates preBötC rhythmogenesis.

(A) Representative traces of ∫preBötC and ∫XII population activity in 3/1 ACSF control (ctrl), 8 μM Cd^{2+}, and 8 μM Cd^{2+} + 10 nM DAMGO. In 3/1 ACSF (top), rhythmic activity consisted of both burstlets and bursts. After Cd^{2+} application (middle), preBötC bursts were no longer observed, and XII activity was abolished, leaving only a preBötC burstlet rhythm with a f similar to control. With addition of DAMGO (bottom), burstlet f and amplitude decreased. Scale bar, 5 s.

(B) Average burstlet fraction, preBötC f, and burstlet amplitude (amp.) in 3/1 ACSF control (ctrl),

6-25 µM Cd^{2+}, and 6-25 µM Cd^{2+} + 0.1-10 nM DAMGO. Addition of Cd^{2+} significantly increased

the burstlet fraction while preBötC f and burstlet amplitudes were unchanged. Bath application

of DAMGO significantly decreased preBötC f (consisting almost solely of burstlets) and burstlet

amplitude. *, $p < 0.05$, One-way ANOVA, post-hoc Tukey test, n=8.

(C) Representative image of tdTomato fluorescence in the preBötC of a medullary slice from a

$Dbx1^{cre};Rosa26^{tdTomato}$ mouse. Dashed circles represent holographic spots over targeted $Dbx1^+$

neurons. Scale bar, 20 µm.

(D) Overlays of representative traces of ∫preBötC from successive laser stimulations of the 8

$Dbx1^+$ neurons shown in (C) in control 9/1.5 ACSF (ctrl), 30 nM DAMGO, and 100 nM DAMGO.

Arrow represents time of laser stimulation. In 9/1.5 ACSF, a failure was seen, but burst initiation

was otherwise reliably successful. Bursts were consistently triggered after a ~100 ms latency. In

30 nM DAMGO, failures were more frequent when the same stimulation parameters were

used, and the latency between laser stimulation and burst initiation was longer and more

variable. In 100 nM DAMGO, bursts were no longer triggered with the same stimulation

parameters. Scale bar, 200 ms.

(E) Average success rate (stim→burst) for burst initiation in control 9/1.5 ACSF (ctrl) conditions

and in increasing concentrations of DAMGO using entraining stimuli that elicits >80% success in

9/1.5 ACSF. To entrain the rhythm, 5-10 $Dbx1^+$ neurons were stimulated 8 - 20 times every 3-6

s. DAMGO produced a dose-dependent decrease in the success rate for triggering bursts with a

significant reduction in success between control and at 100 nM and 200 nM DAMGO. *, $p <$

0.05, One-way ANOVA, post-hoc Tukey test, n=6.

(F) Average latency between laser stimulation and burst initiation when bursts were

successfully triggered during entraining stimuli was increased in 30 nM DAMGO compared to

control 9/1.5 ACSF (ctrl). *, $p<0.05$, Student's t-test, n=5.

Figure 3. µORs expressed in Dbx1$^+$ neurons mediate DAMGO inhibition of preBötC rhythmic

activities.

(A) Confocal images of brainstem sections of a $Dbx1^{Cre};Rosa26^{tdTomato}$ mouse pup (P2) double-

stained with antibodies to ChAT (blue) and µOR (green). Coronal sections showing µOR

expression in the preBötC and the Intermediate band of the Reticular Formation (IRt) and

colocalization with Dbx1$^+$ neurons in the preBötC. Scale bar, 500 μm.

(B) Higher magnification of nucleus ambiguus (NA) and preBötC from the boxed region in A.

Scale bar, 50 μm.

(C) Expanded images of the boxed preBötC region highlighted in B. The arrow shows a preBötC

Dbx1$^+$ neuron with abundant punctate μOR expression on the cell body. Scale bar, 10 μm.

(D) Confocal image of a medullary brainstem section at the level of the preBötC from a control

P2 *Dbx1$^{+/+}$; Oprm1$^{fl/fl}$* mouse processed with RNAScope. Probes for μOR (magenta), TAC1

(white), and acetylcholine esterase (ChAT; green) were used, and the tissue was counterstained

with DAPI (blue). A box marks the location of preBötC. Scale bar, 1 mm.

(E) Higher magnification of the boxed preBötC region in D. μOR and TAC1 were highly expressed

in this region. Scale bar, 100 μm.

(F) Higher magnification images of μOR-expressing neurons with (arrows) and without

(arrowhead) TAC1 expression from E. Colocalization of TAC1 and multiple μOR puncta was

observed in several neurons as well as abundant μOR puncta in neurons that did not express

TAC1. Scale bar, 40 μm.

(G) Confocal image of a medullary brainstem section at the level of the preBötC from a P2

Dbx1$^{cre/cre}$; Oprm1$^{fl/fl}$ mouse processed with RNAScope. Probes for μOR (magenta), TAC1

(white), and acetylcholine esterase (ChAT; green) were used, and the tissue was counterstained

with DAPI (blue). In the preBötC of these mice, the density of μOR puncta was low. A box marks

the location of preBötC. Scale bar, 1 mm.

(H) Higher magnification of the boxed preBötC region in G. The density of μOR puncta was low,

and μOR expression was not associated with TAC1 expression. Scale bar, 100 μm.

(I) Higher magnification images of TAC1-expressing neurons from H with few to no μOR puncta

(arrows) as well as μOR-expressing neurons without TAC1 expression (arrowhead). Scale bar, 40

μm.

(J) Average *f* of preBötC activity in control 9/1.5 conditions (ctrl) and 100 nM DAMGO in

Dbx1^{Cre}; Oprm1$^{fl/fl}$ mice (right), where μORs are selectively deleted in Dbx1$^+$ neurons, and their

Dbx1$^{+/+}$;Oprm1$^{fl/fl}$ littermate controls with preserved μOR expression (left). Whereas 100 nM

DAMGO reduced preBötC f significantly in *Dbx1^+/+^;Oprm1^fl/fl^* control mice, the same

concentration had no effect on f when μORs were deleted. *, p<0.05, Student's *t*-test, n=6 for

*Dbx1^+/+^;Oprm1^fl/fl^*, n=9 for *Dbx1^Cre^;Oprm1^fl/fl^*.

Figure 4. Substance P, but not blockade of inhibitory synaptic transmission, relieves DAMGO-

mediated depression.

(A) Representative traces of ∫preBötC and ∫XII population activity in 3/1 ACSF control (ctrl) and

500 nM SP and 30 nM DAMGO. SP mitigated the effects of DAMGO, and preBötC activity

resembled that in control conditions. Scale bar, 10 s.

(B) Representative traces of ∫preBötC and ∫XII population activity in 3/1 ACSF control (ctrl) and

100 μM picrotoxin/1 μM strychnine and 30 nM DAMGO. The addition of picrotoxin/strychnine

following DAMGO application did not recover preBötC f to control levels. Scale bar, 10 s.

(C) Average normalized f of preBötC activity when 30 nM DAMGO was added alone (9/1.5, 3/1)

or following application of either 500 nM SP or 100 μM picrotoxin/1 μM strychnine with 30 nM

DAMGO. For each experiment, values were normalized to preBötC f during baseline 9/1.5 or

3/1 ACSF in that experiment. When picrotoxin/strychnine were added with DAMGO, the

relative change in f did not differ from the normalized f when DAMGO was applied alone in

either 9/1.5 ACSF or 3/1 ACSF. The normalized f when SP was added with DAMGO was

significantly higher than all other conditions. *, p < 0.05, One-way ANOVA, post-hoc Tukey test,

n=6 for 9/1.5, n=7 for 3/1, n=12 for picrotoxin/strychnine, n=6 for SP.

(D) Representative traces of ∫preBötC and ∫XII population activity in 3/1 ACSF control (ctrl) and

500 nM SP. In 3/1 ACSF, rhythmic activity consisted of both burstlets and bursts. After SP, f was

unchanged, but more preBötC bursts were observed. Scale bar, 10 s.

(E) Average burstlet fraction and f in 3/1 ACSF control condition (ctrl) and 500 nM SP. SP

significantly decreased the burstlet fraction, but did not change preBötC f. *, p < 0.05, Student's

t-test, n=13.

(F) Representative traces of ∫preBötC and ∫XII population activity in 3/1 ACSF control (ctrl) and

100 μM picrotoxin/1 μM strychnine. In 3/1 ACSF, rhythmic activity consisted of both burstlets

and bursts. After picrotoxin/strychnine, f was unchanged; however, more preBötC bursts were

observed, and XII bursts were larger. Scale bar, 10 s.

(G) Average burstlet fraction and f in 3/1 ACSF control condition (ctrl) and 100 µM picrotoxin/1

µM strychnine. Picrotoxin/strychnine significantly decreased the burstlet fraction, but did not

change preBötC f. *, $p < 0.05$, Student's t-test, n=13.

(H) Model for preBötC rhythm and pattern generation. In the absence of neuromodulation

(center), rhythm generation is the alternation between burstlet activity and the silent

interburstlet interval (IBI). Pattern generation in the preBötC occurs when burstlet activity

reaches a threshold (dashed line) and triggers a burst that is then transmitted to downstream

premotoneuronal and motoneuronal populations. These processes can be regulated

independently by a variety of factors (double-headed arrows). Activation of µORs expressed in

Dbx1^{+} neurons (left) slows the rhythm by decreasing excitability in Dbx1^{+} neurons, reducing

synchrony, prolonging IBI, and reducing burstlet amplitude. SP (right) may mitigate the effects

of µOR activation by increasing f or reducing the threshold for burst generation.

Experimental Procedures

Experimental procedures were carried out in accordance with the United States Public Health Service and Institute for Laboratory Animal Research Guide for the Care and Use of Laboratory Animals. All protocols were approved by University of California Animal Research Committee and the Rosalind Franklin University of Medicine and Science Institutional Animal Care and Use Committee. Every effort was made to minimize pain and discomfort, as well as the number of animals.

Transgenic mice

$Dbx1^{Cre}$ mice (Bielle et al., 2005) were crossed with either $Rosa26^{tdTomato}$ (JAX Stock No. 007908) or $Oprm1^{fl/fl}$ (Weibel et al., 2013) mouse lines. $Dbx1^{Cre};Rosa26^{tdTomato}$ were used to visualize Dbx1$^+$ neurons, and $Dbx1^{Cre};Oprm1^{fl/fl}$ mice were used to delete μORs selectively in Dbx1$^+$ neurons using cre-lox recombination. Animal genotypes were verified via real time polymerase chain reaction using primers specific for the $Oprm1^{fl/fl}$ conditional allele or cre recombinase (Pierani et al., 2001; Rose et al., 2009; Weibel et al., 2013) or by direct visualization of fluorescent reporter.

Immunohistochemistry and confocal imaging

$Dbx1^{Cre};Rosa26^{tdTomato}$ and $Dbx1^{Cre};Oprm1^{fl/fl}$ mice were anesthetized at P0-4 by inhalation of isoflurane. The brainstem was isolated from the pons to the rostral cervical spinal cord, fixed in 4% paraformaldehyde, and cryoprotected in 30% sucrose before sectioned at 70mm thickness using CRYOSTAR NX70 (Thermo Scientific). Tissue sections were washed in PBS, and incubated in primary antibody at least overnight, followed by incubation in secondary antibody for 2 hrs at room temperature, and cover-slipped in Vectashield with or without DAPI. Primary antibodies used included rabbit anti-μOR (1:400), goat anti-ChAT (1:400, SCBT); Secondary antibodies were species specific (1:250) and conjugated to Alexa 488, Rhodamine or Cy5 (Invitrogen or Jackson ImmunoResearch). Sections were imaged using confocal laser scanning microscopy. Confocal images were captured on a Zeiss LSM 710 Meta confocal microscope implemented on an upright Axioplan 2 microscope. Zen software was used to capture images. Confocal images

were analyzed with ImageJ, and the final figures were composed in Adobe Photoshop and/or

Microsoft Powerpoint.

In situ hybridization

As described above, mice were anesthetized at P0-4 by inhalation of isoflurane. The isolated

brainstem was rapidly frozen in dry ice and stored at -80°C until sectioning. Transverse sections

(14-20 μm) were cut on a cryostat (CryoStar NX70, Thermo Scientific) and mounted on

SuperFrost Plus (Fisher Scientific) slides. Slides were processed for fluorescence *in situ*

hybridization of three target RNAs simultaneously according to manufacturer protocols for

RNAScope® (Advanced Cell Diagnostics). Briefly, fresh frozen tissue samples were postfixed in

10% neutral buffered formalin, washed, and dehydrated in sequential concentrations of

ethanol (50, 70 and 100%) at RT. Samples were treated with protease IV, then incubated for

two hours at 40°C in the presence of target probes to allow for hybridization. The following

target probes were used: Mm-Oprm1, Mm-Tac1, and Mm-ChAT. A series of three amplification

steps was necessary to provide substrates for target fluorophore labeling (ChAT conjugated to

fluorescein (1:1200), Oprm1 to Cy5, (1:1200) and Tac1 to Cy3 (1:1500) (Jackson

ImmunoResearch). After labeling, samples were counterstained with DAPI. ProLong® Gold

(Invitrogen), was applied to each slide prior to cover slipping. Images were acquired on a

confocal laser scanning microscope (LSM710 META, Zeiss). High-resolution z-stack confocal

images were taken at 0.3 μm intervals. Cell-by-cell μOR expression profiles were quantified in

selected regions of interest (TAC1 positive neurons in preBötC) according to a five-grade scoring

system recommended by the manufacturer (ACD score; 0, no staining; 1, 1-3 dots/cell; 2, 4-9

dots/cell; 3, 10-15 dots/cell; 4, >15 dots/cell), where a dot (>0.60 μm) represented a single RNA

transcript. A Histo score (H-score) was then calculated from the data, using image-based

software analysis (FIJI):

$$\text{H} - \text{score} = \sum_{\substack{bin \\ 0-4}}^{n} (\text{ACD score x percentage of cells per bin})$$

Slice preparation and electrophysiology

Neonatal C57BL/6 or mutant mice (P0-5) of either sex were used for cutting brainstem

medullary transverse slices (550-600 µm thick) containing the rhythm-generating preBötC, XII

inspiratory motoneurons, and premotoneurons linking these populations (Koizumi et al., 2008;

Smith et al., 1991). To obtain slices with the preBötC at the surface, the rostral cut was made

above the first set of XII nerve rootlets at the level of the dorsomedial cell column and principal

lateral loop of the inferior olive, and the caudal cut captured the obex (Ruangkittisakul et al.,

2011). Slice cutting artificial cerebrospinal fluid (ACSF) solution was composed of (in mM): 124

NaCl, 3 KCl, 1.5 $CaCl_2$, 1 $MgSO_4$, 25 $NaHCO_3$, 0.5 NaH_2PO_4, and 30 D-glucose, equilibrated with

95% O_2 and 5% CO_2, 27°C, pH 7.4. Slices were placed rostral side up in a chamber and perfused

at 2-3 ml/min with 28-30°C recording ACSF. Baseline recording ACSF consisted of either the

cutting ACSF solution with K^+ raised to 9 mM and Ca^{2+} maintained at 1.5 mM (9/1.5 ACSF) for a

robust burst rhythm or the cutting ACSF solution with K^+ maintained at 3 mM and Ca^{2+} lowered

to 1 mM (3/1 ACSF) to study burstlets and bursts together (Kam et al., 2013a). Drugs, including

DAMGO, SP, picrotoxin, strychnine, and CdCl, were obtained from Sigma-Aldrich and bath-

applied at the specified concentrations. In all experiments, slices were allowed to equilibrate

for 30 min to ensure that the frequency and magnitude of XII and preBötC population activities

reached steady-state. Respiratory activity reflecting suprathreshold action potential firing from

populations of neurons was recorded from XII motor nucleus or nerve roots and as population

activity directly from the preBötC using suction electrodes (tip size ~50 µm) and an Axiopatch

200A (Molecular Devices), Multiclamp 700B (Molecular Devices), and/or a differential AC

amplifier (AM systems), filtered at 2–4 kHz, integrated, and digitized at 10 kHz. Integration was

performed on custom built Paynter filter with a 20-100 ms time constant. Digitized data were

analyzed off-line using custom procedures written for IgorPro (Wavemetrics).

Holographic photostimulation

Holographic photostimulation was performed on a Phasor SLM system (Intelligent Imaging

Innovations, Inc.) mounted around an epifluorescence upright microscope (Axioscope; Zeiss).

We used a 405 nm diode laser (CUBE 405-100; Coherent) to uncage MNI-glutamate (0.5 mM)

and depolarize targeted neurons. An iterative Fourier transform algorithm, implemented in

Slidebook 5 (Intelligent Imaging Innovations, Inc.), computed the phase pattern on the SLM

corresponding to the desired distribution of light intensity at the focal plane of the objective. A

blocker was inserted in the intermediate Fourier plane to block the unmodulated light

component (zero order spot) and replicate patterns (Golan et al., 2009). Neurons were targeted

by centering 10 um circular spots over the somata of the selected neurons. The laser intensity

per spot was set at 1-3 mW, which was previously determined to elicit bursts of action

potentials in the targeted neurons without the spread of glutamate to neighboring neurons

(Kam et al., 2013b). Each laser stimulation consisted of 5 x 0.8 ms pulses, delivered at 200 Hz. A

threshold stimulus was determined as the minimum number of neurons and minimum laser

power required to entrain preBötC bursts at slightly faster than the endogenous frequency with

a success rate > 80%. The latency between laser stimulation and burst initiation was calculated

as the time between the start of the next preBötC burst and the beginning of laser stimulation.

preBötC activity was used here, rather than XII activity, to isolate the effects of DAMGO on

rhythmogenic mechanisms and avoid additional delays or failures due to depression of XII

premotoneurons and motoneurons. Successful triggering of a burst by photostimulation was

defined as burst detection within 0.5 s of laser stimulation.

Data analysis and statistics

Semiautomated event detection of respiratory-related activity recorded in XII output or

preBötC population recordings was performed using custom procedures written in IgorPro.

Multiple criteria, including slope and amplitude thresholds, were used to select events

automatically, which were then confirmed visually. Event duration, amplitude, shape, and

synchrony between XII and preBötC activity were criteria used to categorize detected events as

burstlets, bursts, and doublets. Burstlets were events in preBötC that did not temporally

overlap a XII burst. Doublets were distinguished from bursts by their longer duration and the

presence of multiple peaks of activity. Two closely spaced bursts were considered a doublet

based on the distribution of the period of XII output. A small peak at <2 s in the distribution of

periods of XII bursts was usually observed. A Gaussian was fit to this small peak and a threshold

time interval was set. Two bursts separated by less than this threshold were considered a

doublet. As doublets were in phase with bursts and burstlets, these events were grouped with

bursts and not separately analyzed.

Rhythmogenic processes were captured by measuring the frequency of preBötC rhythmic

activity, which included small amplitude burstlets that were not transmitted to XII motor

output, larger amplitude bursts that produced XII bursts, and longer duration doublets that

were also transmitted to XII motoneurons. The average frequency was calculated as the mean

of the instantaneous frequencies (1/interevent interval) across all preBötC activity (burstlets,

bursts, and doublets) in that condition. Pattern generating processes that included the

production and transmission of preBötC activity to XII output were characterized by measuring

the fraction of preBötC events that were burstlets. The burstlet fraction was calculated as the

ratio of the number of burstlets to the total number of rhythmic preBötC events (burstlets,

bursts, and doublets). Amplitudes were normalized to the amplitude of the largest burst in the

control condition.

Data are expressed as mean ± SD. Comparisons between two groups were performed with

paired or unpaired Student's *t*-tests. Multiple comparisons were evaluated using parametric

one-way or two-way repeated measures ANOVA. Post-hoc Tukey or paired Student's *t*-tests

with the Bonferroni-Holm comparison were then used to determine significance for within

group comparisons. Statistical significance was set at $p < 0.05$.

    Supplemental Figure Legends

Figure S1. µOR, TAC1, and NK1R expression in preBötC. Related to Figure 3.

(A) Confocal image of a medullary brainstem section at the level of the preBötC from a P2 wild

type mouse triple processed with RNAScope. Probes for NK1R (green), µOR (magenta), and

TAC1 (white) were used, and the tissue was counterstained with DAPI (blue). Scale bar, 20 µm.

(B) Higher magnification images showing colocalization of TAC1, NK1R, and µOR puncta in

several preBötC neurons (arrows) as well as a neuron that expressed NK1R and µOR, but not

TAC1 (arrowhead). Scale bar, 25 µm.

Figure S2. Substance P increases preBötC *f* in *Dbx1^{Cre};Oprm1^{fl/fl}* mice. Related to Figure 4.

(A) Representative traces showing ∫preBötC and ∫XII activity from a *Dbx1^{Cre};Oprm1^{fl/fl}* slice in

control (ctrl, left), 100 nM DAMGO (middle), and 500 nM Substance P (SP, right). Scale bar, 10 s.

(B) SP (500 nM) significantly increased preBötC *f* in slices from *Dbx1^{Cre};Oprm1^{fl/fl}* mice, while 100

nM DAMGO had no effect. *, $p < 0.05$, One-way ANOVA post-hoc Tukey test, n=4.

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Figure 1

Figure 2

Figure 3

Figure 4