**Title:** Dopaminergic control of postnatal mammalian spinal network function is predominantly inhibitory

**Abbreviated title:** Dopaminergic control of dedicated spinal network components

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**Abstract**

Dopamine is well known to regulate movement through the differential control of direct and indirect pathways in the striatum that express $D_1$ and $D_2$ receptors respectively. The spinal cord also expresses all dopamine receptors however; how the specific receptors regulate spinal network output in mammals is poorly understood. We explore the receptor-specific mechanisms that underlie dopaminergic control of spinal network output of neonatal mice during changes in spinal network excitability. During spontaneous activity, which is a characteristic of developing spinal networks operating in a low excitability state, we found that dopamine is primarily inhibitory. We uncover an excitatory $D_1$-mediated effect of dopamine on motoneurons and network output that also involves co-activation with $D_2$ receptors. Critically, these excitatory actions require higher concentrations of dopamine; however, analysis of dopamine concentrations of neonates indicates that endogenous levels of spinal dopamine are low. Because endogenous levels of spinal dopamine are low, this excitatory dopaminergic pathway is likely physiologically-silent at this stage in development. In contrast, the inhibitory effect of dopamine, at low physiological concentrations is mediated by parallel activation of $D_2$, $D_3$, $D_4$ and $\alpha_2$ receptors which is reproduced when endogenous dopamine levels are increased by blocking dopamine reuptake and metabolism. We provide evidence in support of dedicated spinal network components that are controlled by excitatory $D_1$ and inhibitory $D_2$ receptors that is reminiscent of the classic dopaminergic indirect and direct pathway within the striatum. These results indicate that network state is an important factor that dictates receptor-specific and therefore dose-dependent control of neuromodulators on spinal network output and advances our understanding of how neuromodulators regulate neural networks under dynamically changing excitability.

**Keywords:** dopamine, dopamine receptors, spinal cord, rhythmicity, motor systems

**Significance statement:** Monoaminergic neuromodulation of neural networks is dependent not only on target receptors but also on network state. We studied the concentration-dependent control of spinal networks of the neonatal mouse, in vitro, during a low excitability state characterized by spontaneous network activity. Spontaneous activity is an essential element for the development of networks. Under these conditions, we defined converging receptor and cellular mechanisms that contribute to the diverse, concentration-dependent control of spinal motor networks by dopamine, in vitro. These experiments advance understanding of how monoamines modulate neuronal networks under dynamically changing excitability conditions and provide evidence of dedicated $D_1$ and $D_2$ regulated network components in the spinal cord that are consistent with those reported in the striatum.

## 1 Introduction

Neuromodulators are critical for central nervous system function and diversify circuit outputs by altering

synaptic and intrinsic properties [1–3]. Dopamine is a monoamine neuromodulator that is well known for

action selection in vertebrates through the control of direct and indirect circuits of the basal ganglia that

express excitatory $D_1$ and inhibitory $D_2$ receptors respectively (for review, see [4] and [5–7] for

examples). Dopamine is also important for the regulation of spinal motor networks that control rhythmic

movements including but not limited to locomotion (for review, see [8]). In larval zebrafish, phasic and

tonic firing patterns of descending neurons that provide the primary source of spinal dopamine correlates

with locomotor episodes and quiescence, respectively [9]. These different firing patterns can impact the

cellular release of dopamine and as a consequence, the receptor subtypes it activates [10]. For example,

high levels of dopamine released during phasic cell firing activate lower affinity excitatory $D_1$ receptors

and promote locomotor activity, and lower levels of dopamine released during tonic activity activate

higher affinity inhibitory $D_2$ receptors and suppress motor output [11]. Similarly, dopamine has dose-

dependent effects on locomotor circuits in precocial species with functional swim networks [12].

We have recently demonstrated that neuromodulation of developing mammalian spinal circuits is state-

dependent [13] which is consistent with work in invertebrates [14,15]. This is important in developing

spinal motor networks which produce a wide repertoire of patterned outputs at birth, including locomotor

activity, as a consequence of dynamically fluctuating network excitability. That being said, neonatal

rodents rarely produce coordinated bouts of locomotion and in vitro preparations of isolated spinal cord

require pharmacological or electrical stimulation to drive the network into a high excitability state to

produce fictive walking patterns. Instead, most of the movements observed in neonatal mice are ataxic.

(See Supplementary Video 1). These movements correlate with spontaneous network activity, which can

be observed in vitro [16–19]. Nevertheless, the vast majority of what we know about neuromodulation of

developing mammalian spinal networks has been derived from studies on fictive locomotor activities with

dopamine being predominantly excitatory when spinal networks are operating in this state [20–31].

Based on our previous work, we hypothesized that receptor actions, and therefore concentration-

dependent control of network output by dopamine, is linked to the underlying network excitability state

[13]. As a result, more complex receptor-dependent actions may have been masked during high

excitability states of fictive locomotion. We therefore examined how dopamine modulates spinal output at

the network and cellular level of neonatal mouse spinal cords in vitro during a low excitability state

characterized by spontaneous activity  [16,17]. We found that the receptor-specific effects of dopamine

are fundamentally different during a low excitability state compared to what has been previously reported

during high excitability states. Specifically, during a low excitability state, dopamine has primarily

inhibitory effects on network output acting in parallel through activation of $D_2$, $D_3$, $D_4$, and $\alpha_2$ receptors.

We uncover an excitatory effect of dopamine that is likely physiologically silent. This is because

endogenous dopamine levels in the spinal cord are low at this age and the excitatory effects, involving

coactivation of both $D_1$ and $D_2$ receptors, require higher concentrations of dopamine. We also found that

excitatory and inhibitory dopaminergic pathways act through the control of dedicated network

components. Specifically, the $D_1$ pathway acts through excitation of motoneurons, likely recruiting

recurrent excitatory circuits [32], with the $D_2$ pathway acting through hyperpolarization of multiple

ventral interneuron subclasses. Portions of these data were presented in abstract form  [33,34].

**Results**

*Dopaminergic modulation of spinal motor networks is dose-dependent and bidirectional*

We focused on the modulation of perinatal spontaneous activity patterns to investigate how dopamine

modulates spinal network output during a physiologically relevant low excitability state. Using the same

experimental set-up as Sharples et al. [23], we recorded spontaneous motor activity with extracellular

suction electrodes from single ventral roots of the second or fifth lumbar (L2/L5) segments

simultaneously from two or four spinal cord preparations sharing the same chamber. Each preparation in

this series of experiments was naïve to dopamine exposure and received only a single dose of its

respective concentration. This configuration ensured consistent experimental conditions across several

preparations exposed to different concentrations of dopamine. Low concentrations of dopamine (1–30

µM) consistently suppressed spontaneous motor activity, whereas higher concentrations (100–300 µM)

excited spontaneous motor activity, evoking episodic and continuous rhythmic patterns (Fig. 1; n = 42,

$F_{(5,36)} = 10.5.$ p < 0.001).

*Parallel actions of $D_2$, $D_3$, $D_4$, dopamine and alpha-2 adrenergic receptors mediate dopaminergic*

*inhibition of spontaneous activity*

To delineate receptor contributions to dopamine's bidirectional effects on endogenous spontaneous

activity in isolated spinal cord preparations, we used antagonists selective for the family of dopamine

receptors. At low concentrations of dopamine, we observed a negative response ratio (Fig. 2A, Di), which

was due to a reduction in the number (Fig. 2Dii), but not the amplitude, of spontaneous episodes (Fig.

2Diii). In contrast to our hypotheses, the inhibitory effect of dopamine at low concentrations (10 µM) was

not altered by antagonists targeting $D_2$ (L-741,626; n = 3, response ratio = −0.57 ± 0.26 ), $D_2/D_3$

(sulpiride; 20 µM; n = 10, response ratio = −0.36 ± 0.2), $D_3$ (SB-27701-A, 5 µM; n = 3; response ratio =

−0.36 ± 0.3), $D_4$ (L-745,870; n = 3, response ratio = −0.32 ± 0.1) or $D_1/D_5$ receptors (SCH 23390; n = 4,

response ratio = −0.43 ± 0.2). However, when all $D_2$-like receptors were blocked with a cocktail of

sulpiride ($D_2/D_3$, 20 µM) and L-745,870 ($D_4$, 5 µM), response ratios indicated that the inhibitory effect of

10 µM dopamine was attenuated, compared with dopamine alone (Fig. 2C; n = 6; one-way analysis of

variance (ANOVA), $F_{(4,32)}$ = 7.3, p < 0.001; Tukey post hoc p = 0.09), to a level where it was not

significantly different from a time-matched vehicle control (Fig. 2C, Di.; n = 6, $F_{(4,32)}$ = 7.3, p < 0.001;

Tukey post hoc, p = 0.98). Burst analysis revealed no difference in the number or amplitude of episodes

when antagonists were present, compared with baseline (Fig. 2Dii, Diii; two-way ANOVAs for number,

$F_{(2,28)}$ = 9.5, p < 0.0001; two-way ANOVA for amplitude: $F_{(4,28)}$ = 3.4, p = 0.023). Together these data

suggest that the inhibitory actions of dopamine depend on parallel action of $D_2$, $D_3$ and $D_4$ receptors.

We considered the possibility that the inhibitory influence of low dopamine concentrations was partly due

to the activation of non-dopamine receptors. Previous work conducted in our laboratory showed that

dopamine inhibits cauda equina-evoked locomotion, partially via $\alpha_2$-adrenergic receptors [30]. The

remaining minor inhibitory effect of dopamine at 10 µM was blocked by antagonizing the $\alpha_2$-adrenergic

receptors with yohimbine (4 µM) in the presence of $D_2/D_3$ and $D_4$ receptor antagonists. We observed

significantly different response ratios, compared with 10 µM dopamine alone (Fig. 2Di; n = 6, $F_{(4,32)}$ =

7.3, p < 0.001; Tukey post hoc, p = 0.006). Low dopamine concentrations did not change the number or

amplitude of episodes in the presence of these antagonists, compared with baseline (Fig. 2Dii, Diii)

suggesting that non-dopamine receptors also contribute to dopamine's inhibitory effects.

***$D_1$ - $D_2$ receptor coactivation contributes to dopaminergic excitation of spinal network activity***

Dopamine binding to $D_1$ and $D_2$ heteromers can lead to depolarization via an increase in intracellular

calcium levels, mediated by the enzyme phospholipase C (PLC) [35–38]. Thus, we first tested the role of

the lower affinity $D_1$ receptor system and then examined whether the $D_2$ receptor system has a

cooperative role in the control of spontaneous activity. With the addition of dopamine at high

concentrations (i.e., 50 & 100 µM), spontaneous activity patterns became rhythmic, often producing a

slow rhythm with episodes of high frequency rhythmic activity. Moreover, the presence of the $D_1$-like

receptor antagonist, SCH-23390, reduced this effect (Fig. 3; 100 µM dopamine with 10 µM SCH-23390;

n = 5; response ratio, $F_{(5,35)}$ = 11.4, p < 0.001); fast rhythm power, $F_{(4,27)}$ = 12.6, p < 0.001; slow rhythm

power, $H_{(4)}$ = 12.8, p = 0.013; Fig. 3 B & C, 50 µM dopamine with 10 µM SCH-23390; response ratio,

$F_{(5,35)}$ = 11.4, p < 0.001; fast rhythm power, $F_{(4,27)}$ = 12.6, p < 0.001; slow rhythm power, $H_{(4)}$ = 12.8, p =

0.013, Dunn's post hoc: p = 0.03; one way ANOVA used in tests). These results suggest that the

excitatory effects of dopamine are primarily mediated by the $D_1$-like receptor family.

If $D_2$ receptors form heteromers with $D_1$ receptors, we would expect that antagonists would inhibit, rather

than excite responses. We tested this idea by administering $D_2$-like antagonists (sulpiride + L745,870) and

found that the power of the fast rhythm elicited by dopamine at 100 µM was reduced (Fig. 3 Aiii, Biii,

Cii; n = 4; $F_{(4,27)}$ = 12.6, p < 0.001) to the same extent as the $D_1$-antagonist, with no effect on the power of

the slow rhythm (Fig. 3Ciii; $H_{(4)}$ = 12.8, p = 0.013; Dunn's post hoc, p = 1.0). This suggests that $D_2$

receptors may contribute to the excitatory effects of dopamine.

To explore the interaction and co-activation profile of $D_1$ and $D_2$ receptors in isolated neonatal spinal

cords, we performed co-immunoprecipitation for $D_1$ and $D_2$ receptors and used agonists to activate both

receptor subtypes. After immunoprecipitating $D_2$ receptors from neonatal spinal cord lysates, we used an

antibody to probe for $D_1$ receptors. We detected $D_1$ receptor protein within the $D_2$ receptor

immunoprecipitates (Fig. 4Ciii) and these bands were blocked when pre-incubated with an antigen-

blocking peptide. This result indicates that $D_1$ and $D_2$ receptors form a protein complex in neonatal mouse

spinal cords.

In support of this idea, co-application of the $D_1$ agonist SKF 81297 (50 µM) and the $D_2$ agonist quinpirole

(50 µM) elicited a more robust depolarization of the ventral root DC potential, compared with 50 µM of

the $D_1$ agonist alone (Fig. 4A, Ci; $D_1$, n = 8; $D_1/D_2$, n = 8; one-way ANOVA $F_{(2,21)}$ = 5.2, p = 0.01; Tukey

post hoc: p = 0.02). We observed no difference in the amount of spontaneous network activity evoked

with co-application of a $D_2$ agonist, compared with application of the $D_1$ agonist alone, as indicated by the

response ratio (Fig. 4B, Cii; one-way ANOVA, $F_{(3,29)} = 12.0$, $p < 0.001$; Tukey post hoc, $p = 0.5$). In

contrast, lower concentrations of the same agonists (10 µM) produced no effects (n = 8 for each

condition; DC potential, $t_{(6)} = 0.73$, $p = 0.24$; response ratio, $t_{(6)} = 0.9$, $p = 0.19$). Thus, consistent with

previous reports for striatal neurons [36], we found a dose-dependent effect of dopamine agonists wherein

co-applying high doses, but not low doses, of $D_1$ and $D_2$ receptor agonists, produced more robust

depolarization than a $D_1$ agonist alone.

In addition to co-applying separate $D_1$ and $D_2$ agonists, we tested co-activating $D_1$ and $D_2$-like receptors

with the $D_1/D_2$ co-agonist SKF 83959 (50 µM) [38,39]. As predicted, the co-agonist elicited a more

robust depolarization of the ventral root DC potential, compared with the $D_1$ agonist, when applied alone

(Fig. 4A, Ci; D1, n = 8; $D_1/D_2$, n = 8; one-way ANOVA, $F_{(2,21)} = 5.2$, $p = 0.01$; Tukey post hoc, $p = 0.03$).

Interestingly, the co-agonist also robustly facilitated superimposed spontaneous activity, as indicated by a

larger response ratio than co-application of the $D_1$ and $D_2$ agonists produced (Fig. 4B, Cii; one-way

ANOVA, $F_{(3,29)} = 12.0$, $p < 0.001$; Tukey post hoc, $p = 0.004$). These data suggest that under certain

conditions, $D_2$ receptors that are typically inhibitory can play an excitatory role via $D_1/D_2$ co-activation

and contribute to lumbar motor network excitation in the neonatal mouse spinal cord.

***Low levels of endogenous spinal dopamine inhibit spontaneous activity***

We next examined how the endogenous dopamine system regulates perinatal spinal network function.

Endogenous levels of dopamine were increased by blocking the dopamine transporter (DAT) with an

antagonist GBR 12909 (10 µM). Blocking DAT (n = 8 preparations) produced a modest but significant

reduction in spontaneous activity compared to time-matched vehicle controls, reflected by a reduced

response ratio (Fig. 5A, Bi; one-way ANOVA, $F_{(2,19)} = 18.0$, $p < 0.001$) and a reduced number of

spontaneous episodes (Fig. 5Bii; n = 8, two-way ANOVA, $F_{(2,20)}$ = 11.8, p = 0.0004), with no change in

amplitude (Fig. 5Biii; n = 6, two-way ANOVA, $F_{(2,20)}$ = 1.3, p = 0.3). We questioned whether

extracellular dopamine metabolism may have dampened the effect of the reuptake blocker, thus

diminishing the predicted increase in endogenous dopamine levels. Therefore, we repeated this

experiment in the presence of bifemelane, a monoamine oxidase A & B inhibitor. Under these conditions,

we found further reductions in spontaneous activity when DAT was blocked, as indicated by a

significantly reduced response ratio **(**Fig. 5Bi; n = 6, one-way ANOVA, $F_{(2,19)}$ = 18.0, p < 0.001). Burst

analysis revealed significantly fewer episodes (Fig. 5Bii; n = 6, two-way ANOVA, $F_{(2,20)}$ = 11.8, p =

0.0004) with no change in amplitude (Fig. 5Biii; n = 6, two-way ANOVA, $F_{(2,20)}$ = 1.3, p = 0.3).

We followed these experiments up with high-performance liquid chromatography (HPLC) to verify

endogenous levels of dopamine. In P3 lumbar spinal cords (n = 11) we detected low levels of dopamine;

in P60 adults (n = 17) we detected a threefold increase in dopamine levels (Fig. 5Biv; Mann–Whitney U =

10.0, T = 76, p < 0.001). Thus, our in vitro experiments indicated that low levels of endogenous dopamine

play a role in $D_2$-mediated inhibition.

***$D_1$ receptor activation increases motoneuron excitability by reducing afterhyperpolarization properties***

In the next set of experiments, we were interested in determining the cellular mechanisms that mediate

dopamine's complex modulatory effects on spinal network output. As integrators of premotor network

activity that generate many of the rhythmic outputs of the spinal cord, motoneurons are ideally suited to

amplify spontaneous activity and respond to dopaminergic modulation. Given that they not only serve as

the final output for spinal networks, but they also participate in the generation of rhythmic activity [40],

we initially selected motoneurons as a locus for determining the cellular mechanisms for dopaminergic

excitation and inhibition. We made whole-cell patch-clamp recordings from 75 motoneurons (across 42

animals). Some of these motoneurons (n = 18 across seven animals) were filled with fluorescein-

conjugated dextran amine (FITC; Molecular Probes, Inc., Eugene Oregon) and verified post hoc using

immunohistochemistry for the presence of choline acetyltransferase (ChAT; Fig. 6A). One hundred

percent of filled cells were ChAT positive indicating that we were indeed recording from motoneurons.

Electrical properties measured in a group of these cells (n = 8 cells across five animals) were comparable

to cells from putative motoneurons recorded in the remaining experiments (Table 1) indicating that these

cells are also likely motoneurons. Motoneurons also display characteristic time-dependent changes in

repetitive firing frequency during sustained depolarizing current injections (Fig. 6 Cii) [41–43]. We found

that all but two cells demonstrated a time-dependent reduction in firing rate throughout the duration of a

0.5 s depolarizing current injection, as indicated by the ratio between maximum steady-state and first

spike interval frequencies. This phenomenon is known as spike frequency adaptation (SFA; mean SFA

ratio = $0.47 \pm 0.19$). In addition, doublet firing occurred at the onset of repetitive firing in 5 cells (8%).

Consistent with previous reports from our laboratory [44,45] and with our network recordings in this

report, 100 µM of dopamine increased motoneuron excitability (n = 5 cells across four animals); we

reproduced this effect with the $D_1$ agonist SKF 81297 (20 µM; n = 8 cells across five animals). Both 100

µM dopamine and the $D_1$ agonist depolarized the membrane potential (Fig. 6Bi; $H_{(2)} = 18.9$, $p < 0.001$),

increased negative holding current (vehicle, $-7.1 \pm 33$ pA; DA, $-247 \pm 78$ pA; $D_1$, $-174 \pm 49$ pA; $H_{(2)} =$

18.9, p = 0.001) and input resistance (Fig. 6Bii; $H_{(2)} = 16.0$, $p < 0.001$), and decreased rheobase (Fig. 6Bii;

$F_{(2,22)} = 5.0$, $p = 0.016$) beyond that of the time-matched vehicle control. We found no change in spike rise

time ($F_{(2,22)} = 1.0$, $p = 0.4$) or half width ($F_{(2,22)} = 0.8$, $p = 0.5$). The $D_1$ agonist reduced the amplitude of

the afterhyperpolarization (AHP); however, 100 µM dopamine did not reduce AHP amplitude beyond

that of a time-matched vehicle control (Fig 6Biv; $F_{(2,22)} = 7.7$, $p = 0.003$). Frequency–current (FI)

relationships were measured for steady state and first spike interval during a series of depolarizing current

pulses (Fig. 6Ci). Both 100 µM dopamine and the $D_1$ agonist reduced the latency to first spike beyond

that of the time-matched vehicle control (Fig. 6Cii; $F_{(2,17)} = 9.6$, $p = 0.002$). Dopamine (100 µM) and the

$D_1$ agonist increased the slope of the exponential region of the FI relationship for the first spike interval

(Fig. 6Di–Dii; $F_{(2,22)}$ = 8.4, p = 0.002) and reduced the slope of the steady-state FI relationship (Fig.

6Diii–Div; $H_{(2)}$ = 9.4, p = 0.009). The reduction in steady-state slope was due to a leftward shift in steady-

state FI relationship (Fig. 6Diii) characterized by a reduction in the threshold for repetitive firing (Fig.

6Ciii; $H_{(2)}$ = 14, p < 0.001) with no change in the maximum steady-state firing rate ($H_{(2)}$ = 1.4, p = 0.5).

These results indicate that activation of $D_1$ receptors elicit consistent effects as high concentrations of

dopamine on motoneuron excitability and is a likely mechanism contributing to dopaminergic excitation

of motor output.

***Dopaminergic inhibition through $D_2$ - receptor hyperpolarization of distributed populations of ventral***

***interneurons***

We next set out to determine the cellular mechanisms that mediate the inhibitory effects of dopamine on

spinal network output with motoneurons as our first target. In contrast to our network recordings, low

concentrations of dopamine (1–10 µM; n = 20 cells across 14 animals; Table 1) and the $D_2$ agonist

quinpirole (20 µM; n = 12 cells across seven animals; Table 1) did not alter any passive, spike or

repetitive firing properties of motoneurons beyond that of the time-matched vehicle control (n = 12 cells

across six animals). A lack of responsiveness was not due to the diversity of motoneurons as there was no

correlation between cell capacitance and changes in membrane potential or holding current elicited by

low concentrations of dopamine (r = -0.3, p = 0.13) or quinpirole (r = -0.1, p = 0.72). These results

suggest that the inhibitory actions of dopamine on spinal network output are not due to $D_2$-receptor

inhibition of intrinsic motoneuron excitability; instead, dopamine may be acting on premotor

interneurons.

Many of the premotor interneurons that produce the rhythmic activities generated by the spinal cord are

distributed across lamina's VII - X of the ventral lumbar spinal cord. We next recorded from ventral

interneurons located in lamina VII–X to determine a cellular locus for $D_2$-mediated inhibition of spinal

network output (n = 30 cells across 17 animals; Table 2). Quinpirole produced a sustained

hyperpolarization of the resting membrane potential in 33% of interneurons (Fig. 7B, C, Di; n = 10; dVm

= −4.8 ± 2.4 mV; two-way ANOVA, $F_{(2,37)}$ = 26.1, p < 0.0001) and a transient hyperpolarization of

membrane potential in 10% of interneurons (n = 3; dVm = −5.16 ± 1.9 mV) that persisted for 209 ± 108 s

before returning to baseline levels. The change in resting membrane potential was greater in the

responders compared to non-responders (dVm responders = - 4.8 ± 2.4 mV; dVm non-responders = 1.6 ±

1.9 mV; unpaired t-test $t_{(28)}$ = 7.8, p < 0.0001). Quinpirole reduced the input resistance (Fig. 7Dii; two-

way ANOVA, $F_{(2,37)}$ = 4.1, p = 0.025) and increased the spike rise time (Fig. 7Div; two-way ANOVA,

$F_{(2,35)}$ = 5.2, p = 0.01; Dunn post hoc test, p=0.02) but not the half width ($F_{(2,35)}$ = 2.6, p = 0.086) in the

group of responding interneurons ("responders"). For both responders and non - responders, quinpirole

did not alter rheobase (Fig. 7Diii; two-way ANOVA, $F_{(2,37)}$ = 0.5, p = 0.6), action potential threshold

(two-way ANOVA, $F_{(2,37)}$ = 0.3, p = 0.7), AHP amplitude (two-way ANOVA, $F_{(2,37)}$ = 0.04, p = 0.96),

duration (two-way ANOVA, $F_{(2,37)}$ = 0.8, p = 0.45), threshold for repetitive firing (two-way ANOVA,

$F_{(2,37)}$ = 0.01, p = 0.98), maximum steady-state firing rate (Fig. 7Dvi; two-way ANOVA, $F_{(3,37)}$ = 0.8, p =

0.5 ) or steady-state FI slope (Fig. 7Dv: two-way ANOVA, $F_{(2,37)}$ = 0.9, p = 0.4). The capacitance (two-

way ANOVA, $F_{(2,37)}$ = 4.2, p = 0.02) and holding current (two-way ANOVA, $F_{(2,37)}$ = 4.9, p = 0.02) were

higher in responders than in non - responders (Table 2) and a greater proportion of responders were

localized to more medial regions of spinal slices (Fig. 7B), where putative commissural interneurons

reside [46].

We next set out to determine the type of interneurons that were hyperpolarized by quinpirole. Given that

many of the responding cells were located medially, we next targeted descending commissural

interneurons (dCINs; n = 10 cells across five animals; Table 2) since this population can be identified

based on anatomical connectivity [47,48], display intrinsic burst properties [49] and are rhythmically-

active during neurochemically-evoked fictive locomotion [50]. dCINs were retrogradely labelled with

tetramethylrhodamine-conjugated dextran amine (molecular weight (MW) 3000; Molecular Probes, Inc.)

inserted into the ventrolateral funiculus at the L4 segment (Fig. 7A). In contrast to our hypothesis, only

one dCIN responded with a sustained hyperpolarization and two were transiently hyperpolarized (Fig. 7B)

by quinpirole. Quinpirole did not alter any passive, spike, or repetitive firing properties of dCINs (n = 10;

Fig. 7D). These data suggest that the dCINs, although responsive in similar proportions to our global

interneuron survey, do not exclusively account for the 33% responding group and, therefore, are likely,

not responsible for the observed network effects. Interestingly, when all interneuron data were pooled (n

= 40 cells), irrespective of responsiveness as indicated by changes in resting membrane potential,

quinpirole had the greatest effect in cells that had a higher maximum steady-state firing rate at baseline

(Fig. 7DViii; r = −0.37, p = 0.026). There was no correlation between baseline FI slope and changes in FI

slope in response to quinpirole (r = −0.09, p = 0.6).

While dCINs can be identified anatomically, they are heterogeneous with respect to their neurotransmitter

phenotype [51] and as a result have varying contributions to network activities [46,50,52]. We therefore

next targeted V3 interneurons which are exclusively glutamatergic, contribute to the stabilization of

locomotor-like rhythmicity and can be identified genetically based on the expression of the Sim1

transcription factor [53]. Heterogeneity with respect to location, morphology and electrophysiological

properties has also been reported in V3 interneurons [54,55] which may be accounted for in part by a

recently described hierarchical microcircuit whereby a medial population projects to a lateral population

which provide glutamatergic excitation of ipsilateral motoneurons. Both populations also project

commissurally and receive recurrent glutamatergic inputs from intra and intersegmental ipsilateral

motoneurons [32]. Given that dopamine inhibits ventral root-evoked locomotor activity, which may be

mediated by this circuit [56], through $D_2$-receptor signaling [31], we hypothesized that V3 interneurons

may be a cellular locus for $D_2$-mediated inhibition of spinal network activity.

Consistent with our global interneuron survey, quinpirole produced a sustained hyperpolarization of the

resting membrane potential in a proportion of V3 interneurons (n=5 cells; 27%) that we recorded from

(total V3 interneurons n=23 cells, 7 animals) and transient hyperpolarization in 3 (13%) V3 interneurons.

The magnitude of the response in the 5 cells that responded with a sustained hyperpolarization was

variable and approached, but did not reach, significance (Supp Figure 1; paired t-test: $t_{(4)}$= 2.5, p = 0.06);

however, did reach significance when the cells that responded with a transient hyperpolarization were

included in the analysis (paired t-test: $t_{(6)}$= 2.7, p = 0.03). Consistent with our global interneuron survey

the change in resting membrane potential elicited by quinpirole was significantly greater in responding (n

= 5; dVm = -1.9 ± 1.9 mV) compared to non-responding (n = 15; dVm = -0.3 ± 1.0 mV) V3 interneurons

(unpaired t-test: $t_{(16)}$ = 3.8, p = 0.002).

**Discussion**

Dopamine is a monoamine neuromodulator that is important for the control of rhythmically active motor circuits across phyla (reviewed by [8]) but is probably best known in vertebrates for the control of dedicated circuits in the basal ganglia that control action selection (reviewed by [57] ). Work in small circuits of invertebrates has established that circuit connectomes define the constraints on which networks operate and that neuromodulators diversify outputs by altering intrinsic and synaptic properties of the neurons that compose the circuit (for reviews see [1–3]. In line with this, the distribution of receptors within circuits constrain the effect of neuromodulators on circuit output. For example, dopamine is exclusively inhibitory in spinal circuits of *Xenopus* tadpoles prior to free-swimming stages [58] due to expression of $D_2$ but not $D_1$ receptors. We show that dopamine has bidirectional concentration-dependent effects on spinal network output in neonatal mice where all dopamine receptor types are expressed which is consistent with what has been reported in tadpoles at free swimming-stages [12]. Our data highlights that neuromodulator concentration is also important because receptors have varying ligand affinities which underlie concentration-dependent actions of modulators. Although dopamine predominantly inhibits spinal output in neonatal mice, similar to pre-free-swimming tadpoles, it is primarily due to the concentration of spinal dopamine, not the distribution of receptors. Our previous work shows that neuromodulation of mammalian spinal networks is dependent on network excitability state [13] which is consistent with findings from invertebrates [14,15]. Our current work shows that receptor mechanisms and concentration-dependent control of spinal network output is also state-dependent. This is important because receptor expression, modulator concentration and network excitability are not fixed and fluctuate dynamically [59,60]. Therefore, these three factors need to be considered if we wish to understand how networks create diverse neuromodulator-dependent outputs (Figure 8A).

*A physiologically silent excitatory $D_1$ pathway?*

Previous work has shown predominantly excitatory $D_1$-receptor mediated effects of dopamine on fictive locomotion which is characteristic of the network operating in a high excitability state [13] - albeit in

these studies, higher concentrations of dopamine are necessary to elicit observable effects [20–25]. Here,

we show that endogenous levels of dopamine within the spinal cord of neonatal mice are low. Even

though HPLC suggests low concentrations of dopamine in the neonatal spinal cord, a critical question that

we addressed is what would happen when we manipulated endogenous dopamine? We accomplished this

by blocking dopamine reuptake and metabolism and found that the effects were not excitatory, but

inhibitory. Based on this, we suggest that although present, the excitatory $D_1$-mediated pathway is

'physiologically silent' during early postnatal stages given that endogenous levels are not sufficient to

activate this pathway. A similar phenomenon has been reported for glutamatergic synapses in developing

circuits of the hippocampus which express NMDA but not AMPA receptors. As a result, glutamatergic

synapses fall 'physiologically silent' as the release of glutamate does not produce sufficient

depolarization of the postsynaptic membrane to remove the magnesium block from the pore of the

NMDA channel [61,62].

Excitability state also influences the receptor mechanisms and therefore concentration-dependent control

of modulators on network output. In the spinal cord, $D_1$ pathways may be more important for the

regulation of spinal circuits operating in higher excitability states. For example, fictive locomotor rhythms

are drastically impacted when $D_1$ receptors are manipulated whereas manipulation of $D_2$ receptors elicit

only subtle changes in rhythm frequency, with rhythm robustness being maintained [23] (Figure 8Bii.). In

the adult animal; however, network excitability increases, and spinal dopamine levels are higher due to

increases in descending inputs. In line with this, optogenetic activation of the dopaminergic A11 leads to

an increase in motor activity [63]. Similarly, the excitatory $D_1$ system is more important for the control of

stepping movements whereas the inhibitory $D_2$ system plays less of a role [22,64]. Instead, the inhibitory

$D_2$ pathway may be more important in maintaining network quiescence during periods of immobility such

as has been inferred by the study of firing patterns in the zebrafish [9]. Our data demonstrates that in the

neonatal mouse, where spinal dopamine levels and network excitability are low, that inhibition of motor

output prevails (Figure 8Bii.). Thus, this points to the receptor-specific and therefore concentration-

dependent control of modulators on network output being strongly influenced by the state of the network

on which they are acting.

***Dedicated network components segregate excitatory and inhibitory control of spinal networks***

Dedicated circuits regulated by non-overlapping populations of neurons that express $D_1$ and $D_2$ receptors

compose the direct and indirect pathways of the basal ganglia in vertebrates and have also been reported

in the superior colliculus of rodents [65]. Our work suggests dedicated network elements within the spinal

cord that are regulated by $D_1$ and $D_2$ pathways (Figure 8Biii.). Specifically, we found that $D_1$ receptors

excite motoneurons through similar mechanisms that have been previously reported [44,45], but are not

affected by low concentrations of dopamine or $D_2$ agonists. This points to the possibility of a dedicated

$D_1$-dependent circuit that could underlie the generation of rhythmic activities elicited by high

concentrations of dopamine. Motoneurons compose key rhythm generating elements in invertebrate

circuits [66–68] and also participate in rhythm generation in vertebrates [69], including rodents

[32,40,56,70,71]. V3 interneurons are one subclass of genetically-defined spinal interneuron that are

important for the generation of rhythmic activities in mammalian spinal networks [53,55,72] and receive

recurrent excitatory collaterals from motoneurons in rodents [32]. Motoneurons in the rodent spinal cord

also form glutamatergic synaptic connections amongst each other [71] and activation of $D_1$ receptors

could serve to synchronize motor pools. Previous results demonstrating $D_1$- and not $D_2$-mediated

increases in AMPA conductances on motoneurons [45] support this possibility. We cannot rule out the

possibility that there is a degree of overlap between $D_1$ and $D_2$ controlled network elements within the

spinal cord. While we did not examine the $D_1$ control of ventral interneurons, we did find that $D_2$

receptors hyperpolarize a subset of V3 interneurons. This population is therefore a potential cellular locus

where cooperative excitatory $D_1$-$D_2$ interactions that we report here could occur. $D_1$ and $D_2$ receptors have

been reported in the brain to become co-activated or form heterodimeric complexes that augment

neuronal excitability through PLC-dependent increases in intracellular calcium [35–38]. Our

pharmacological and immunoprecipitation data indicate that this may also occur in the neonatal mouse

spinal cord. Although this pathway may be physiologically silent during early postnatal development,

increasing dopamine concentrations at later stages may activate this pathway.

The inhibitory $D_2$ pathway on the other hand does not appear to act through the modulation of

motoneuron intrinsic or synaptic properties [45] but instead hyperpolarizes a proportion of ventral

interneurons. Based on our responsiveness criteria, a majority of cells that responded to a $D_2$ agonist with

sustained hyperpolarization of the membrane potential were localized more medially in lumbar slices. We

tested the hypothesis that a subpopulation of interneurons was $D_2$ sensitive and tested both dCINs and

genetically identified V3 interneurons. Our hypothesis was not supported, and similar proportions of $D_2$

sensitive neurons were found; however, the data suggest that there is heterogeneity in the responsiveness

to neuromodulators within a class of genetically defined interneurons and that $D_2$ actions may not be

localized to a particular class of interneurons. One possibility is that neuromodulators elicit a robust effect

on network output through distributed control across multiple classes of genetically-defined interneurons

which would be consistent with findings that the locomotor rhythm generator is also distributed across

several classes of interneurons [73–75]. Our network data suggests that low concentrations of dopamine

inhibit network output by acting in parallel on $D_2$, $D_3$, $D_4$ and $α_2$ receptors. We may therefore have

underestimated the cellular targets that underlie dopamine's robust inhibitory effect on the network given

that in this series of experiments we looked at the activation of $D_2$ receptors alone.

***Developmental considerations for the endogenous dopaminergic system***

This robust inhibitory system may act as a brake on network activity during perinatal development when

chloride-mediated synaptic transmission still causes partial depolarization [76]. An inhibitory brake

would prevent runaway excitation of spinal network activity during this critical period of development. A

robust background inhibitory system mediated by the parallel inhibitory action of $D_2$, $D_3$, $D_4$, and α2-

adrenergic receptors activated by low levels of endogenous dopamine would counteract depolarizing

chloride-mediated transmission.

Signaling through [76] $D_2$-like receptors may also play a role in driving the maturation of spinal

networks. In larval zebrafish, $D_4$ receptors drive the maturation of spinal locomotor network organization

[77] and function leading to changes in locomotor behaviour [78]. Similar processes may also occur

perinatally in rodents, in that the preferential activation of the $D_2$ receptor system may favour intracellular

signaling that results in network reorganization. Serotonin receptors have been found to shape network

function and inhibitory synaptic transmission during early postnatal days of rodents [79,80]. Dopamine

could, therefore, act analogously via the $D_2$-system during perinatal development.

***Conclusions***

Here we present evidence for an inhibitory physiological role of dopamine in the regulation of developing

mammalian spinal networks. We also demonstrate an excitatory $D_1$-mediated pathway that acts through

excitation of motoneurons, and possibly recurrent excitatory collaterals to CPG neurons, however given

that endogenous levels of dopamine are low, propose that this pathway is physiologically silent. These

data advance our understanding of how neuromodulators regulate network output in light of dynamically

changing modulator concentrations and levels of network excitability.

**Methods**

*Ethical approval & animals*

Experiments were performed on male and female neonatal (P0–P4, n = 262) and adult (P50, n = 17)

C57BL/6 mice. A subset of experiments were performed on spinal cord slices obtained from neonatal

(P0-4, n=7) $Sim1^{Cre/+}$; $Rosa^{floxstop26TdTom}$ which were used to visualize and record from V3 interneurons. All

procedures performed were approved by the University of Calgary Health Sciences Animal Care

Committee University Committee on Laboratory Animals at Dalhousie University.

*Tissue preparation*

We anesthetized all animals by hypothermia. Pups were decapitated and eviscerated to expose the

vertebral column and rib cage. The isolated vertebrae and rib cage were transferred to a dish lined with a

silicone elastomer (Sylgard; DowDuPont, Midland, MI) and perfused with room-temperature (21 - 23ºC)

carbogenated (95% $O_2$ / 5% $CO_2$) aCSF (in mM, 4 KCl, 128 NaCl, 1 $MgSO_4$, 1.5 $CaCl_2$, 0.5 $Na_2HPO_4$, 21

$NaHCO_3$, 30 D-glucose; 310–315 mOsm.). We exposed the spinal cord with a ventral laminectomy and

isolated it by cutting the nerve roots that connected it to the vertebral column. The isolated spinal cord

was then transferred to a recording chamber, perfused with carbogenated aCSF, and placed ventral side

up. The bath temperature was gradually increased to 27°C [81]. We let the spinal cords stabilize for 1

hour before performing experiments.

*Spinal cord slice preparation*

Following isolation, spinal cords were transected above the tenth thoracic (T10) and below the first sacral

(S1) segments and transferred to a slicing chamber. Pre-warmed liquefied 20% gelatin was used to secure

cords to an agar (3%) block that was super-glued to the base of a cutting chamber and immersed in ice-

cold, carbogenated, high-sucrose slicing aCSF (in mM, 25 NaCl, 188 sucrose, 1.9 KCl, 10 Mg $SO_4$, 1.2

$Na_2HPO_4$, 26 $NaHCO_3$; 25 D-Glucose; 340 mOsm). Using a vibratome (Leica, Bussloch, Germany) we

cut 250-μm-thick lumbar slices, collected and transferred them to a recovery chamber containing regular

carbogenated aCSF (see Tissue Preparation) heated to 32ºC for one hour, then maintained them at room

temperature for at least 30 minutes before transferring them to a recording chamber.

*Labelling of descending commissural interneurons (dCINs)*

We retrogradely labelled dCINs by inserting tetramethylrhodamine-conjugated dextran amine crystals

(MW 3000; Molecular Probes, Inc.) into a cut in the ventrolateral funiculus at the L4 segment. Spinal

cords recovered for 4 hours to allow retrograde uptake of the fluorescent dye. Fluorescently-labelled cells

were visualized with epifluorescent illumination.

*Electrophysiological recordings*

Extracellular neurograms were recorded by drawing ventral roots of the second (L2) and fifth (L5) lumbar

segments into tight-fitting suction electrodes fashioned from polyethylene tubing (PE50). Signals were

amplified 1000× in total via 10× pre-amplification and 100× second-stage amplification (Cornerstone

EX4-400 Quad Differential Amplifier; Dagan Corporation, Minneapolis, MN). Amplified signals were

band-pass filtered (0.1–1000 Hz) and digitized at 2.5 kHz (Digidata 1440A/1550B; Molecular Devices,

Sunnyvale, CA). Data were acquired in Clampex 10.4/10.7 software (Molecular Devices) and saved on a

Dell computer for offline analysis. All experiments were performed on spinal cords naïve to drugs and

experimental treatment.

*Whole-cell patch-clamp recordings*

Spinal cord slices were gently transferred to a recording chamber perfused with room-temperature

carbogenated recording aCSF (in mM, 128 NaCl, 4 KCl, 1.5 $CaCl_2$, 1.0 $MgSO_4$, 0.5 $Na_2HPO_4$, 21

$NaHCO_3$, 30 D-glucose; approximately 310 mOsm) and stabilized in the recording dish with a stainless-

steel harp. We gradually heated aCSF to 27°C. Slices were visualized (Olympus BX51WI; Olympus

Corporation, Tokyo, Japan) under 5× magnification and putative motoneurons identified using a 40×

objective with infrared differential interference contrast (IR-DIC) illumination. We identified putative

motoneurons based on their location in the ventrolateral spinal cord and a soma diameter of greater than

20 μm. A cohort of motoneurons was passively filled with fluorescein dextran amine (FITC; MW 3000;

200μM; Molecular Probes, Inc.) added to the intracellular solution to visualize and localize the recorded

cells, for 20–60 minutes and motoneuron identity was verified post hoc by performing

immunohistochemistry for choline acetyltransferase (ChAT). To verify motoneuron identity in

subsequent experiments, basal biophysical properties of putative motoneurons were compared with the

FITC-filled ChAT-positive cells. We identified no differences in basal biophysical properties of ChAT-

positive identified cells, compared with cells in other experiments identified by position and size (Table

1). In one set of experiments, we recorded from ventral interneurons in lamina VII–X, descending

commissural interneurons (dCINs) and V3 interneurons of the lumbar spinal cord (Table 2). We measured

the distance of each interneuron from the central canal (LinLab2; Scientifica, Uckfield, UK) and

normalized position, relative to lateral and ventral borders, and plotted the normalized positions on a

template transverse spinal cord. Responsiveness was determined by the change in resting membrane

potential. All experiments were performed on one cell per slice to ensure that all cells were naïve to

treatment.

Recording electrodes were pulled from borosilicate capillary tubes (O.D. = 1.5mm, I.D. = 0.86 mm) using

a Flaming/Brown Model P-97 micropipette puller (Sutter Instrument, Novato, CA). Pipettes pulled to

record from motoneurons, and interneurons were within the range of 3–5 MΩ and 6–9 MΩ, respectively.

Pipettes were backfilled with intracellular solution (in mM, 130 K-gluconate, 0.1 EGTA, 10 HEPES, 7

NaCl, 0.3 $MgCl_2$, 2 ATP, 0.5 GTP, 5 phosphocreatine; 280 mOsm). Intracellular solutions were balanced

to a pH of 7.3 with 10 M KOH and osmolality was cross-checked to fall within the range of 275–290

mOsm. Data were acquired at 10 kHz using Clampex software (PClamp 10.4; Molecular Devices).

We examined intrinsic properties of motoneurons and ventral interneurons and terminated experiments if

access resistance was greater than 25 MΩ for motoneurons and 35 MΩ for interneurons, if cells had a

resting membrane potential greater than −50 mV, or if action potential amplitude was less than 60 mV at

baseline. Cells were excluded from analysis if access resistance deviated by more than 20% by the end of

the recording. We held all cells at a membrane potential of approximately −75 mV during experiments,

after correcting for a liquid junction potential of 14.3 mV. The liquid junction potential was calculated in

Clampfit (PClamp 10.4; Molecular Devices), using the ionic composition of our extracellular and

intracellular solutions. Protocols for the examination of intrinsic properties have been described elsewhere

[82].

*Pharmacology*

Dopamine hydrochloride (Sigma-Aldrich, Inc., St. Louis, MO) was bath applied in separate experiments

at 1 µM, 3 µM, 10 µM, 30 µM, 100 µM, and 300 µM to determine dose-dependent effects on motor

activity. The receptor-selective agonists we used included SKF 81297 for $D_1$-like receptors (10–50 µM;

Tocris, Minneapolis, MN); quinpirole for $D_2$-like receptors (10–50 µM; Tocris); and the $D_1/D_2$ receptor

co-agonist SKF 83959 (10–50 µM; Tocris). For dopamine receptor antagonists we used the $D_1$-like

antagonist SCH-23390 (10 µM; Tocris); the $D_2$-like antagonists sulpiride (20 µM) and L-741,626 (12

µM); the selective $D_3$ receptor antagonist SB 27701A (5 µM; Tocris); the selective $D_4$ receptor antagonist

L-745, 870 (5 µM; Tocris). We also used the $\alpha_2$ adrenergic receptor antagonist, yohimbine (2–4 µM;

Tocris). Endogenous dopamine levels were manipulated with the DAT inhibitor GBR-12909 (10 µM;

Hello Bio, Princeton, NJ) and the monoamine oxidase A & B inhibitor bifemelane (50 µM; Tocris).

*Immunoprecipitation for $D_1$ and $D_2$ receptors*

Spinal cords were dissected in ice-cold (4–8ºC) aCSF and homogenized in lysis buffer containing 50 mM

TrisHCl, 150 mM NaCl, 10 mM EDTA, 0.1% Triton-X, and 5% Glycerol. Lysis buffer contained

protease inhibitors (Sigma) and phosphatase inhibitors (GBiosciences). We homogenized three spinal

cords in 100 µL of buffer and incubated them on ice for 1 hour before centrifuging them at 10,000 rpm

for 30 minutes at 4ºC. Lysates were then extracted and stored at −20 ºC.

To reduce nonspecific binding, we first incubated lysates in anti-rabbit Ig agarose beads (Trueblot;

Rockland Inc., Limerick, PA) for 30 minutes, on ice and in the absence of primary antibody. We then

removed the supernatant and incubated the lysates on ice for 1 hour with rabbit antibody to $D_2$ receptors

(1 µg per 100 µL, Millipore). Anti-rabbit Ig IP beads were added and samples were incubated overnight at

4°C with gentle agitation. Immunoprecipitates were washed with lysis buffer, heated in loading buffer

(350 mM Tris, 30% glycerol, 1.6% SDS, 1.2% bromophenol blue, 6% β-mercaptoethanol) to 95°C for 10

min, electrophoresed on a precast SDS gel (4–12% Tris HCl; BioRad, Hercules, CA), and transferred

onto a nitrocellulose membrane. After blocking, the membranes were incubated with guinea pig antibody

to $D_1$ receptors (1:400; Alomone, Jerusalem, Israel) and rabbit antibody to $D_2$ receptors (1:500; Millipore,

Burlington, MA), washed, incubated for 2 hours at room temperature in fluorophore-conjugated

secondary antibodies (IRDye anti-guinea pig and Trueblot anti-rabbit IgG DyLight, 1:1000), and

visualized via antibody fluorescence at 680 or 800 nm (Odyssey CLx; LI-COR Biosciences, Lincoln,

NE).

***Post hoc verification of motoneurons***

Following the completion of experimental protocols, we post-fixed slices overnight in 4%

paraformaldehyde (PFA) at 4ºC, washed them the next morning in phosphate-buffered saline (PBS) for

three 20-minute intervals, and stored them at 4ºC. On the day we performed immunohistochemistry, we

first washed the slices for 30 minutes (3 × 10 minutes) in PBS with 0.5% Triton-X 100 (PBST) then

incubated them at room temperature (21 -23ºC) for 6 hours in blocking solution containing 10% donkey

serum in 0.5% PBST. Primary antibodies for choline acetyltransferase (ChAT) (1:500; goat anti-ChAT;

Millipore Cat No. AB144) and the fluorescence marker (1:1000; rabbit anti-FITC, ThermoFisher Cat No.

11090) were diluted in blocking solution and the slices incubated with them for at least 24 hours at room

temperature. Slices were then washed in PBST (0.5%) for 80 minutes (4 × 20 min) at room temperature

before secondary antibodies (donkey anti-rabbit Alexa Fluor 1:1000, Life Technologies Cat No. A-21206;

Donkey anti-Goat Alexa 568 1:1000; Cat No. A21447, Life Technologies, Carlsbad, CA) were applied

for 4–8 hours at room temperature, then washed for 80 minutes in PBST (0.5%). Slices were mounted and

coverslipped with fluorescent mounting medium (Vectashield; Vector Laboratories, Burlingame, CA);

coverslips were separated from slides by 500 μm spacers to prevent crushing.

***Imaging***

All sections processed via immunohistochemistry were imaged on a Nikon A1R MP$^+$ microscope (Nikon,

Tokyo, Japan) operating in confocal mode with a 16× water-immersion objective lens (numerical aperture

[NA] = 0.8, working distance [WD] = 3 mm). Image acquisition used a z-step of 1 μm and averaged two

frames with a resolution of 2048 × 2048. Pixel dwell time was 2.5 ms and exposure settings were

maintained for all sections. We used NIS-Elements software (Nikon) for image acquisition and ImageJ to

perform maximum intensity projections of z-stacks.

***High-performance liquid chromatography***

Monoamine content of neonatal and adult spinal cords was measured using high-performance liquid

chromatography (HPLC). We dissected spinal cords from neonatal (P3, n = 11) C57BL/6 mice in aCSF as

described above and extracted adult spinal cords (P60, n = 17) with a pressure ejection method. Tissue

was then flash-frozen with liquid nitrogen, stored at −80ºC, and analyzed for biogenic amines with a

modified version of the Parent et al. HPLC method [76]. Tissue was homogenized in ice-cold 0.1 M

perchloric acid. We centrifuged the homogenate and used 10 μl of supernatant in the assay, employing an

Atlantis dC18 column (Waters, Milford, MA) and an electrochemical detector.

*Data analysis*

We determined the relative inhibitory or excitatory effects of dopamine and dopamine receptor agonists

on spontaneous motor network activity using methods similar to those in our previous work [23]: we

calculated a response ratio from single ventral root neurograms between the root mean square of 5

minutes of basal spontaneous activity and 5 minutes of activity recorded 20 minutes after adding the drug.

We subtracted 1 from the response ratio so that positive values reflect excitation and negative values

reflect inhibition. The response ratio was used as a high throughput assay to detect global changes in

network activity. Neurogram data were analyzed with Spike2 software. Bursts of spontaneous activity

were analyzed using Clampfit (Molecular Devices) to determine how episode number and amplitude

contributed to changes in response ratio. Spectral analyses were conducted using Spinalcore software [83]

whenever we detected excitatory changes or observed rhythmicity.

Patterns of rhythmic motor activity recorded from single ventral roots were analyzed with autowavelet

spectral analysis. We created and analyzed frequency–power spectrograms by selecting regions of interest

around frequency ranges that coincided with those in the raw neurogram. The spectrograms revealed two

high power regions that reflect distinct rhythms evoked by dopamine at high concentrations: a slow 0.01–

0.04 Hz rhythm and a fast 0.8–1.2 Hz rhythm. Regions of interest were selected within these frequency

ranges and analyzed over the time course of each experiment. Frequency power within selected regions of

interest that corresponded to the fast and slow rhythms were used as a measure of rhythm robustness.

Data were segmented into 30 s bins and averaged over 5-minute intervals for statistical analysis. We used

tools available in Spinalcore for all analyses of rhythmic motor activity  [83], consistent with Sharples

and Whelan [13].

Motoneuron and interneuron intrinsic properties measured during whole-cell patch-clamp experiments

were analyzed as in our previous work [82,84], with the exception of repetitive firing analyses, which

examined the instantaneous firing rate for the first spike interval and steady-state firing separately.

*Experimental design and statistical analysis*

All experiments were repeated measures. We tested for differences in the magnitude of effects between

conditions with one-way ANOVAs, focusing on comparisons to time-matched vehicle controls. Two-way

ANOVAs compared baseline to multiple post-drug conditions. All effects surpassing a significance

threshold of $p < 0.05$ were further examined with post hoc analyses. We used Holm-Sidak post hoc tests

to compare all treatment conditions to the appropriate normalized time-matched vehicle control. Data that

violated assumptions of normality (Shapiro–Wilk test) or equal variance (Brown–Forsythe test) were

analyzed via nonparametric Mann–Whitney $U$ (if two groups) or Kruskal–Wallis (if more than two

groups) tests.

**Tables and Captions**

*Table 1: Baseline motoneuron intrinsic properties.*
Passive, spike, and repetitive firing properties of motoneurons with dopamine or dopamine agonists did not differ at baseline in this series of experiments. Motoneuron identity was verified in a subset of experiments where cells were filled with fluorescein and verified post hoc for expression of choline acetyltransferase (ChAT); 100% of cells were ChAT[+] indicating that they were indeed motoneurons. Data are means ± SD and analyzed using one-way ANOVAs for each property.

| Motoneuron basal properties (N= 65 MNs, 40 animals (P0-P4)) | | | | | | | | | |
|---|---|---|---|---|---|---|---|---|---|
| Property | Chat+ filled MNs (8,5)[1] | Vehicle (12,6)[2] | 1 µM DA (7,4)[3] | 5 µM DA (7,4)[4] | 10 µM DA (6,5)[5] | 100 µM DA (5,4)[6] | SKF 81297 (8,5)[7] | Quinpirole (12, 7)[8] | $F_{(7,56)}$, p |
| **Passive properties** | | | | | | | | | |
| Capacitance (pF) | 203±54 | 203±35 | 193±55 | 194.3±45 | 149±66 | 244±54 | 199±50 | 210±43 | 1.6, 0.15 |
| $V_m$ (mV) | -73±4.1 | -74±4.3 | -75±4.1 | -73±4.7 | -73±3.7 | -76±3.5 | -72±6.1 | -73±5.0 | 0.7, 0.7 |
| $I_{Holding\ -75mV}$ (pA) | -86±148 | -25±138 | 26±125 | -63±85 | 23±150 | -101±99 | -38±87 | -74±62 | 1.16, 0.3 |
| Rin (MΩ) | 50±15 | 48±24 | 59±44 | 70±83 | 115±84 | 40±9.2 | 71±28 | 49±14 | 2.0, 0.07 |
| Rheobase (pA) | 678±225 | 747±386 | 596±367 | 464±286 | 443±390 | 762±318 | 580±298 | 633±260 | 0.9, 0.5 |
| **Spike Properties** | | | | | | | | | |
| $AP_{TH}$ (mV) | -51±4.4 | -55±4.4 | -55±4.2 | -52±5.9 | -56±3.5 | -53±2.5 | -54±4.3 | -55±4.4 | 1.0, 0.4 |
| $AP_{AMP}$ (mV) | 66±5.0 | 71±5.9 | 75±6.9 | 76±13 | 76±4.2 | 69±5.7 | 71±9.6 | 72±6.1 | 1.6, 0.2 |
| $AP_{rise\ time}$ (ms) | 0.45±0.2 | 0.33±0.1 | 0.37±0.1 | 0.4±0.1 | 0.44±0.1 | 0.32±0.04 | 0.35±0.05 | 0.33±0.1 | 1.6, 0.2 |
| $AP_{half\ width}$ (ms) | 0.59±0.15 | 0.60±0.1 | 0.64±0.2 | 0.78±0.3 | 0.75±0.2 | 0.62±0.07 | 0.68±0.1 | 0.68±0.19 | 1.2, 0.3 |
| $AHP_{duration}$ (ms) | 73.7±29 | 70±18 | 88±34 | 76±13 | 108±38 | 74±16 | 69±23 | 76±26 | 1.5, 0.2 |
| $AHP_{AMP}$ (mV) | -3.9±1.1 | 3.6±2.2 | 3.1±1.4 | 3.2±1.4 | 3.4±2.1 | 4.7±1.2 | -4.2±1.9 | -4.3±1.4 | 0.8, 0.6 |
| **Repetitive Firing Properties** | | | | | | | | | |
| SS F/I Slope (Hz/pA) | 0.053±0.01 | 0.044±0.008 | 0.058±0.008 | 0.052±0.018 | 0.053±0.03 | 0.051±0.007 | 0.046±0.008 | 0.053±0.012 | 1.2, 0.3 |
| Repetitive firing TH (pA) | 605±446 | 498±254 | 410±271 | 354±200 | 338±327 | 515±253 | 367±186 | 494±173 | 0.8, 0.6 |
| Max SS firing rate (Hz) | 59±8.1 | 59±8.3 | 51±13 | 46±7.9 | 52±8.9 | 55±12 | 57±9.5 | 58±12 | 1.9, 0.09 |
| First spike latency (ms) | 146±86 | 116±57 | 124±77 | 91±34 | 113±79 | 136±55 | 132±67 | 142±50 | 0.5, 0.8 |

*Table 2: Baseline ventral interneuron intrinsic properties*
Passive, spike, and repetitive firing properties of ventral interneurons that responded to quinpirole with sustained hyperpolarization of membrane potential (responders) were compared with those that did not respond (non-responders) and also retrogradely labelled descending commissural interneurons (dCINs). Responders had higher capacitance and holding current than non - responders and dCINs. Data are means ± SD; F and p values are reported from one-way ANOVAs for each property. Red values highlight variables where significant main effects were detected with $p < 0.05$. Superscript numbers reflect significant differences between respective conditions from post hoc analysis.

| Ventral Interneurons (n=40 cells, 22 animals (P0-4)) | | | | |
|---|---|---|---|---|
| **Property** | **Non-responders[1] (n=20,8)** | **Responders[2] (n=10,9)** | **dCINs[3] (n=10,5)** | **(2,37) F, p** |
| **Passive Properties** | | | | |
| Capacitance (pF) | 47±15[2] | 73±40[1] | 46±17[2] | 4.2, 0.02 |
| $V_m$ (mV) | -66±4.3 | -66±5.2 | -66±6.0 | 0.02, 0.98 |
| $I_{Holding -75mV}$ (pA) | -10.8±16[2] | -57±52[1] | -26±18 | 4.9, 0.02 |
| Rin (MΩ) | 759±466 | 497±248 | 719±326 | 1.6, 0.1 |
| Rheobase (pA) | 84±103 | 59±13 | 47±13 | 0.9, 0.4 |
| **Spike Properties** | | | | |
| $AP_{TH}$ (mV) | -55±4.6 | -55±2.9 | -56±2.3 | 0.2, 0.8 |
| $AP_{AMP}$ (mV) | 71±9.3 | 71±8.2 | 72±7.3 | 0.05, 0.95 |
| $AP_{rise\ time}$ (ms) | 0.99±0.48[3] | 0.92±0.3[3] | 1.4±0.25[1,2] | 4.2, 0.023 |
| $AP_{half\ width}$ (ms) | 1.2±0.5 | 1.4±0.6[3] | 1.9±0.4[2] | 6.2, 0.005 |
| $AHP_{duration}$ (ms) | 173±96 | 193±76 | 172±81 | 0.2, 0.8 |
| $AHP_{AMP}$ (mV) | -4.9±2.7 | -4.8±2.1 | -4.8±2.7 | 0.02, 0.98 |
| **Repetitive Firing Properties** | | | | |
| SS F/I slope (Hz/pA) | 0.24±0.07 | 0.21±0.09 | 0.2±0.1 | 1.0, 0.4 |
| Repetitive firing TH (pA) | 26±14 | 31±12 | 28±13 | 0.5, 0.6 |
| Max firing rate (Hz) | 43±16.3 | 39±11.6 | 32±5.9 | 2.1, 0.13 |
| Fist spike latency (ms) | 143±81 | 146±95 | 117±87 | 0.45, 0.6 |

**Figures and Captions**

**Figure 1: Dopamine evokes bidirectional dose-dependent modulation of lumbar network activity**
A. Extracellular neurograms recorded from naïve single lumbar ventral root preparations after applying dopamine in various concentrations. Each row indicates a separate trial. Each recording began with spontaneous motor activity; the green section shows the effect of dopamine for each dose. The red-dashed box highlights expanded sections of data to the right of each neurogram. B. Mean normalized response ratios calculated from the root mean square of the raw neurogram during a 5-minute window, 20 minutes after dopamine application, compared to baseline activity. The negative response ratios represent inhibition and positive values represent excitation. The number in each green bar indicates the number of preparations in the average for each concentration. Error bars indicate standard deviation. Asterisks denote the significance level of Tukey post hoc tests between ratio concentrations (*$p < 0.05$, **$p < 0.01$, ***$p < 0.001$).

***Figure 2: Receptor mechanisms for dopaminergic inhibition of spontaneous network activity.***
A–C. Single ventral root (L5) extracellular neurograms of spontaneous activity in the presence of dopamine (A), a $D_1$ antagonist and dopamine (B), and with a cocktail of $D_2$ antagonists and dopamine (C). Spinal cords were perfused with receptor-preferring antagonists (red bars) 20 minutes prior to the application of low concentrations of dopamine (10 μM). D. Response ratio represents the root mean square of raw neurograms during a 5-minute window, 20 minutes after dopamine application, compared with prior to dopamine application. Negative response ratios indicate inhibition and positive values indicate excitation. The number in each bar indicates the number of preparations included in that condition's mean. Bars indicate the mean (± SD), number of episodes per minute (Dii), and amplitude (Diii) of spontaneous episodes recorded within the 5-minute epochs during which response ratios were calculated. Asterisks indicate the significance level of post hoc comparisons (*p < 0.05, **p < 0.01, ***p < 0.001).

***Figure 3: Receptor mechanisms for dopaminergic excitation of spinal network activity.***
High concentrations of dopamine excite spinal networks and produce episodic and continuous rhythmic patterns of activity. A. Single ventral root extracellular neurograms of spinal network activity from each condition: dopamine applied alone (Ai), in the presence of a $D_1$ antagonist (Aii), and with a cocktail of $D_2$ antagonists (Aiii). Horizontal bars indicate the timing of dopamine application (green bars) and the application of receptor-preferring antagonists (red bars). B. Spectrograms show autowavelet frequency power across time, evoked at high concentrations of dopamine (50 -100 µM). The colour bar indicates power magnitude, from high (warm) to low (cool colours). We selected regions of interest around regions that coincided with fast and slow rhythm frequency rhythms. Spinal cords were perfused with antagonists for 20 minutes prior to the application of dopamine. Ci. Mean response ratios (± SD), as in Fig. 2, for each condition. Spectral analysis of fast rhythm (Cii) and slow rhythm (Ciii) power following drug application for each experimental condition. Histograms present mean values ± SD, with asterisks denoting significance level of post hoc tests (*p < 0.05, **p < 0.01, ***p < 0.001) following one-way ANOVA (fast rhythm; Cii) or nonparametric one-way ANOVA (slow rhythm; Ciii).

**Figure 4: $D_1$ $D_2$ receptor co-activation contributes to the excitatory effect of dopamine.**
A. Single ventral root extracellular DC neurograms of spontaneous activity from conditions where a $D_1$ agonist (SKF 81297) was applied alone (red), co-applied with a $D_2$ agonist (SKF 81297 + Quinpirole; black), or $D_1$ and $D_2$ receptors were coactivated by a $D_1/D_2$ co-agonist (SKF 83959; blue). Bi-iii. For each agonist, (B) expands the region of spontaneous activity 20–25 minutes after agonist application, as outlined by the dashed line in (A). C. Ci depicts depolarization of DC potentials from DC neurograms, averaged across the number of preparations denoted by numbers in each condition's bar. Cii shows the response ratio for each condition (as in previous figures), which represents changes in the amount of spontaneous activity. Ciii. Co-immunoprecipitation of $D_1$ with $D_2$ receptors suggests that they may interact in the neonatal spinal cord. Histograms present mean values ± SD, with asterisks denoting significance level of post hoc comparisons (*$p < 0.05$, **$p < 0.01$, ***$p < 0.001$) following one-way ANOVA.

**Figure 5: Inhibitory actions of endogenous dopamine in the neonatal mouse spinal cord.**
A. Single ventral root (L5) neurogram of spontaneous activity after blocking the dopamine transporter (DAT) with the DAT antagonist GBR 12909 (purple) to increase the endogenous dopamine level. Dopamine reuptake was also blocked in the presence of monoamine oxidase A & B inhibitor bifemelane (green) to reduce dopamine metabolism. Bi illustrates the response ratio; negative and positive values indicate inhibition and excitation, respectively. The number below each bar represents the number of preparations for each experiment. Spontaneous episode occurrences per minute (Bii) and amplitude (Biii) were measured within the epochs where the response ratio was calculated. Histograms present mean values ± SD and asterisks indicate a significance level of post hoc analyses (*p < 0.05, **p < 0.01, ***p < 0.001) following one-way ANOVA for response ratios, and two-way ANOVA for burst occurrences and amplitude that compared between conditions. Biv. Endogenous levels of serotonin (5-HT), dopamine (DA) and norepinephrine (NE) were measured in neonatal (P3) and adult (P60) lumbar spinal cords with high-performance liquid chromatography.

***Figure 6: High concentrations of dopamine and D₁ agonists increase motoneuron excitability***
A. Whole-cell patch-clamp recordings were obtained from motoneurons visualized with infrared differential interference contrast (IR-DIC) in lumbar slices of neonatal (P0–P4) mice. A subset of cells were filled with fluorescein (FITC, green) and verified post hoc with immunohistochemistry for choline acetyltransferase (ChAT, red). B–D. High concentrations of dopamine (100 μM, green) and the D₁ agonist SKF 81297 (20 μM, purple) increased motoneuron excitability by modulating various passive, spike, and repetitive firing properties. All measures were compared to that of time-matched vehicle control (grey). Ci. A typical motoneuron trace following current injection. Di, Diii display representative frequency–current (FI) plots from a single cell before (black dots) and after (purple dots) application of the D₁ agonist. Box and whisker plots display interquartile range (boxes), median (horizontal black lines), max, and minimum value in data range (whiskers). Asterisks denote significance level of post hoc tests (*p < 0.05, **p < 0.01, ***p < 0.001) following one-way ANOVA.

**Figure 7: D₂ agonists hyperpolarize a proportion of ventral interneurons.**
Whole-cell recordings obtained from lamina VII–X interneurons. Ai. Descending commissural interneurons (dCINs) were retrogradely labelled with rhodamine-conjugated dextran amine crystals inserted into the ventrolateral funiculus at L4 level. Aii. Cells were visualized in transverse lumbar slices under infrared differential interference contrast (IR-DIC) or epifluorescent illumination. B. Ventral interneuron location was measured relative to the central canal and X-Y positions normalized to the distance of ventral and lateral borders of lumbar slices. C. Whole-cell current-clamp recordings depict responsiveness of interneuron resting membrane potential to the D₂ agonist (quinpirole, 20 µM) in responders, non - responders, and dCINs. A proportion of ventral interneurons responded to quinpirole with either a sustained (B, red and green dots) or transient (B, blue dots) hyperpolarization of the membrane potential. A reduction in input resistance and increase in spike rise time accompanied hyperpolarization in sustained responders (Dii, Div). Box and whisker plots display interquartile range (boxes), median (horizontal black lines), max, and minimum value in data range (whiskers). asterisks denote significance level from post hoc tests (*p < 0.05, **p < 0.01, ***p < 0.001) following two-way ANOVA. Dviii. Reductions in steady-state frequency-current (FI) slope of all cells following application of quinpirole was greatest in cells that had a higher maximum steady-state firing rate at baseline.

**Figure 8: Dopamine exerts state and receptor-dependent control of spinal motor networks.**
A. Neural network output is dependent on multiple factors. Circuit connectomes and intrinsic properties defining constraints of a network output. Similarly, the distribution of neuromodulator receptors defines the constraints on which neuromodulators can alter these properties. Varying ligand affinities of these receptors determine dose-dependent effect of neuromodulators and network state dictates the dose-dependent effects of neuromodulators. Bi. Previous work has demonstrated that tonic and phasic firing patterns of dopamine cells lead to the release of low and high concentrations of dopamine, respectively [10]. Bii. During a low excitability state (blue part of the circle), low concentrations of dopamine (denoted by the green gradient) acted on $D_2$, $D_3$, $D_4$ and $\alpha 2$-adrenergic receptors in parallel to inhibit (parallel stop lines) spinal motor output. As dopamine levels rise, activation of $D_1$ and co-activation of $D_2$ receptors increase (arrowheads) spinal network output. During a high excitability state (orange part of the circle), the actions of low dopamine are not apparent and at higher concentrations, activation of $D_1$ receptors boost spinal network output and activation of $D_2$ receptors slows rhythmic activity [13]. Biii. Higher concentrations of dopamine act through $D_1$ receptors to increase motoneuron excitability by reducing A-type and $SK_{Ca}$-dependent calcium conductance. $D_2$-mediated inhibition of spinal network output is triggered by reduced excitability in a proportion of ventral interneurons (red cells).

***Supplemental Figure 1*: *D₂ - receptor control of V3 interneurons.***
Whole-cell patch-clamp recordings obtained from lamina V3 interneurons. A. Visually guided patch showing pipette tip on a GFP labelled-V3 interneuron. B. V3 interneuron location was measured relative to the central canal and X-Y positions normalized to the distance of ventral and lateral borders of lumbar slices. Non-responders represented by black dots, responders by red dots, and transient responders by blue dots. C. A sample trace showing a hyperpolarization of the resting membrane potential following administration of 20 µM quinpirole in responders (top trace) and non-responders (bottom trace). Di. Box and whisker plots showing a change in resting membrane potential in responders and non-responders before (gray) and after (red) administration of quinpirole. Dii. Input resistance graphed in a similar manner as Di.

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