

Lhx3-Chx10 Reticulospinal Neurons in Locomotor Circuits

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Motor behaviors result from the interplay between the brain and the spinal cord. Reticulospinal neurons, situated between the supraspinal structures that initiate motor movements and the spinal cord that executes them, play key integrative roles in these behaviors. However, the molecular identities of mammalian reticular formation neurons that mediate motor behaviors have not yet been determined, thus limiting their study in health and disease. In the medullary reticular formation of the mouse, we identified neurons that express the transcription factors Lhx3 and/or Chx10, and demonstrate that these neurons form a significant component of glutamatergic reticulospinal pathways. Lhx3-positive medullary reticular formation neurons express Fos following a locomotor task in the adult, indicating that they are active during walking. Furthermore, they receive functional inputs from the mesencephalic locomotor region and have electrophysiological properties to support tonic repetitive firing, both of which are necessary for neurons that mediate the descending command for locomotion. Together, these results suggest that Lhx3/Chx10 medullary reticular formation neurons are involved in locomotion.

Introduction

Goal-directed movements result from interplay between the brain and the spinal cord. The brain selects and plans the motor programs, and transmits motor commands from the brainstem to the spinal cord, which in turn translates these commands into motor action. While progress has been made in understanding motor commands in the forebrain (Drew et al., 2004; Bretzner and Drew, 2005) and integration in the spinal cord (Brownstone and Bui, 2010), less is known about the brainstem neurons that play key integrative roles in producing movement.

One movement on which studies of neural circuits have been focused is locomotion. In the 1960s, Shik and Orlovsky identified the mesencephalic locomotor region (MLR) in the dorsal aspect of the mesopontine tegmentum as a key center that initiates locomotion (Shik et al., 1966). Tonic stimulation of the MLR in decerebrate cats initiates coordinated four-limbed locomotor activity. The MLR command signal is integrated in the medullary reticular formation (medRF; Garcia-Rill and Skinner, 1987b; Jordan, 1998; Deliagina et al., 2002; Zaporozhets et al., 2004; Dubuc et al., 2008; Mullins et al., 2011), likely within both gigantocellular

and magnocellular reticular nuclei (Mori et al., 1978; Shefchyk et al., 1984; Mori, 1987; Noga et al., 2003). The reticulospinal pathway that recruits spinal locomotor circuits is a fast-conducting (Orlovskii, 1970; Degtyarenko et al., 1998; Noga et al., 2003), glutamatergic (Douglas et al., 1993; Jordan, 1998; Häglund et al., 2010) pathway that works in concert with raphespinal serotonergic systems (Liu and Jordan, 2005). However, the identity and properties of the involved mammalian reticulospinal neurons remain largely unknown.

The advent of genetic techniques has led to the identification and characterization of specific populations of neurons involved in motor systems (Lee and Pfaff, 2001; Kiehn, 2006; Goulding, 2009; McLean and Fetcho, 2011). For example, it is possible to drive persistent expression of fluorescent proteins in genetically defined neuronal populations so as to target these neurons for anatomical, molecular, and physiological studies (Pierani et al., 2001; Lanuza et al., 2004; Wilson et al., 2005, 2010). Like in the spinal cord (Sharma et al., 1998; Al-Mosawie et al., 2007; Lundfald et al., 2007; Dougherty and Kiehn, 2010), the transcription factors Lhx3 and Chx10 are expressed in the medRF (Cepeda-Nieto et al., 2005; Hafner et al., 2008; Kinkhabwala et al., 2011; Koyama et al., 2011; Crone et al., 2012). In the zebrafish, hind-brain Chx10 cells are glutamatergic, rhythmically active, and provide excitation to the spinal cord during swimming (Kinkhabwala et al., 2011; Kimura et al., 2013). Therefore, in seeking neurons in the medRF that project to the spinal cord and may be involved in movement in limbed animals, we studied mouse Lhx3-expressing and Chx10-expressing neurons and their potential role in locomotion.

We show here that the transcription factors Lhx3 and Chx10 define populations of glutamatergic neurons in the medRF that project to the spinal cord, receive functional inputs from the MLR, and are activated in locomotion. These data support the

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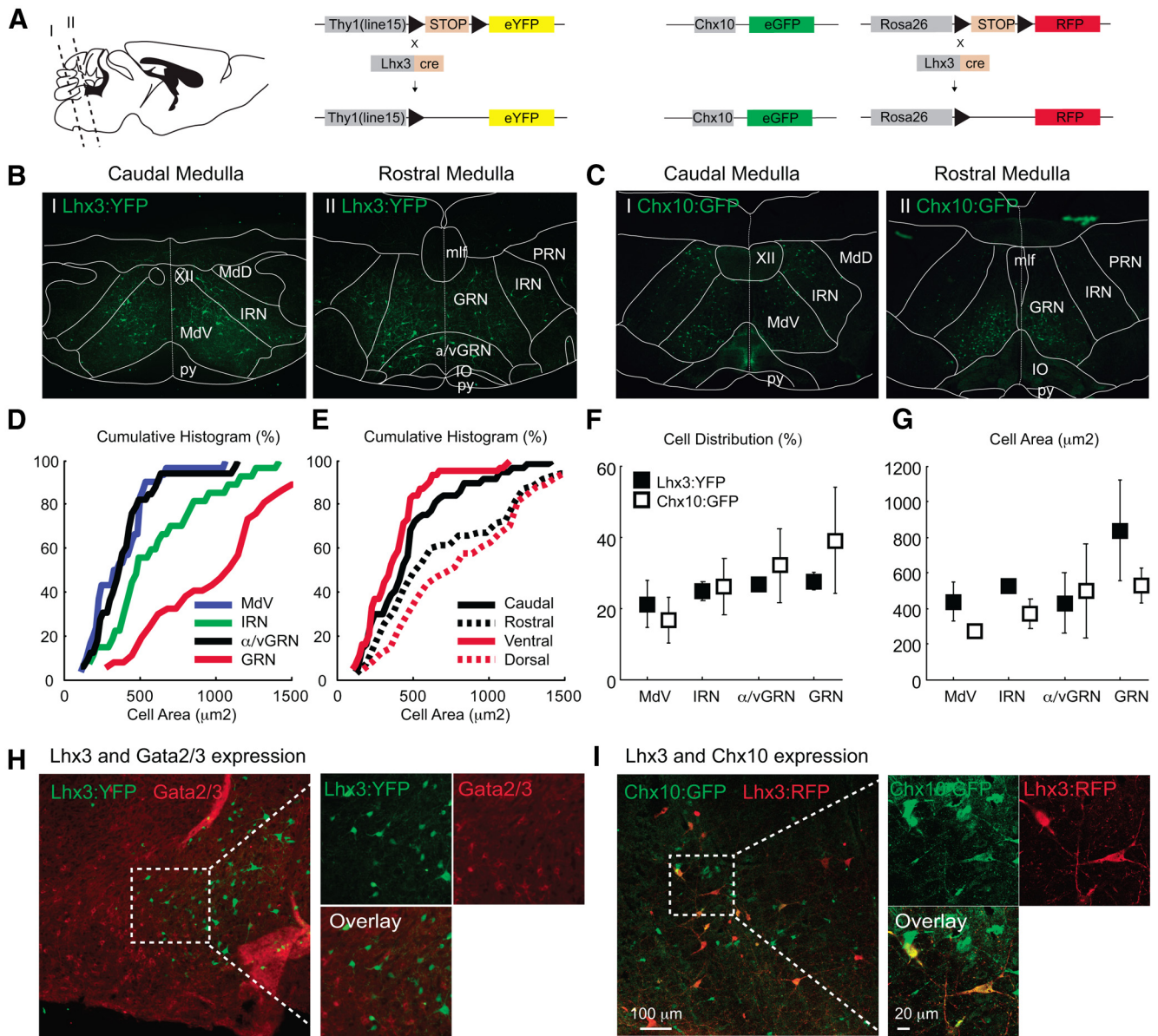


Figure 1. Anatomical characterization of *Lhx3* and *Chx10* medRF neurons. **A**, Genetic constructs and levels of transverse sections in the sagittal plane. **B**, **C**, Low-magnification images of adult *Lhx3*-YFP (**B**) and *Chx10*::GFP (**C**) brainstem coronal sections. **D**, Cumulative histograms for *Lhx3*-YFP medRF neuron area in four medRF nuclei. **E**, Cumulative histograms for *Lhx3*-YFP medRF neuron area across rostrocaudal and ventrodorsal axes. Note a similar pattern was observed for *Chx10*::GFP medRF neurons (data not shown). **F**, Cell distribution (mean and STD) of *Lhx3*-YFP (filled squares) and *Chx10*::GFP (open squares) medRF neurons across medRF nuclei. **G**, Areas (mean and STD) of *Lhx3*-YFP and *Chx10*::GFP medRF neurons across medRF nuclei. **H**, Absence of overlap between *Lhx3*-YFP (green) and *Gata2/3* (red) medRF neurons. **I**, Overlap of *Lhx3*-RFP (red) and *Chx10*::GFP (green) medRF neurons. py, Pyramide; mlf, medial longitudinal fasciculus; IO, inferior olive; MdD, dorsal aspect of the medullary reticular formation; XII, hypoglossal nucleus.

hypothesis that *Lhx3*/*Chx10* reticulospinal neurons are involved in mediating a descending motor command for locomotion.

Materials and Methods

All procedures were in accordance with protocols approved by the Dalhousie University Committee on Laboratory Animals and conformed to the standards of the Canadian Council for Animal Care. Figures reported and bars shown are means \pm SDs.

Mouse strains. The following mice of either sex were used in various combinations: *Lhx3*^{cre/+} (Sharma et al., 1998), *thy1-loxP-stop-loxP-YFP* reporter line 15 (*thy1-fs-YFP*, yellow fluorescent protein; Feng et al., 2000; Al-Mosawie et al., 2007), *Rosa26-loxP-stop-loxP-tdTomato* [*Rosa26-fs-tdTomato*, Jackson Labs; when crossed with *Lhx3*^{cre/+}, termed *Lhx3*-YFP or *Lhx3*-RFP (red fluorescent protein) mice respectively], BAC *Chx10*::eGFP

(Mutant Mouse Regional Resource Center, University of California, Davis; and BAC *GAD67::eGFP* (Gong et al., 2003; Jackson Labs).

The use of *Lhx3*^{cre/+} mice has the advantage that the neurons can be genetically manipulated, but suffers from incomplete recombination (Al-Mosawie et al., 2007). The *Chx10*::eGFP mice provided more complete labeling of this neuronal population. Therefore, both types were used in this study.

Immunohistochemistry, in situ hybridization, tracing, and imaging. Mice were anesthetized with ketamine and transcardially perfused with 0.9% saline followed by 4% paraformaldehyde in 0.1 M phosphate buffer, pH 7.4. Tissues were harvested, postfixed in 4% paraformaldehyde overnight, and subsequently cryoprotected in 30% sucrose. As previously described, immunohistochemistry was performed on cryostat cut sections mounted on slides (Bretzner et al., 2010) or microtome cut floating

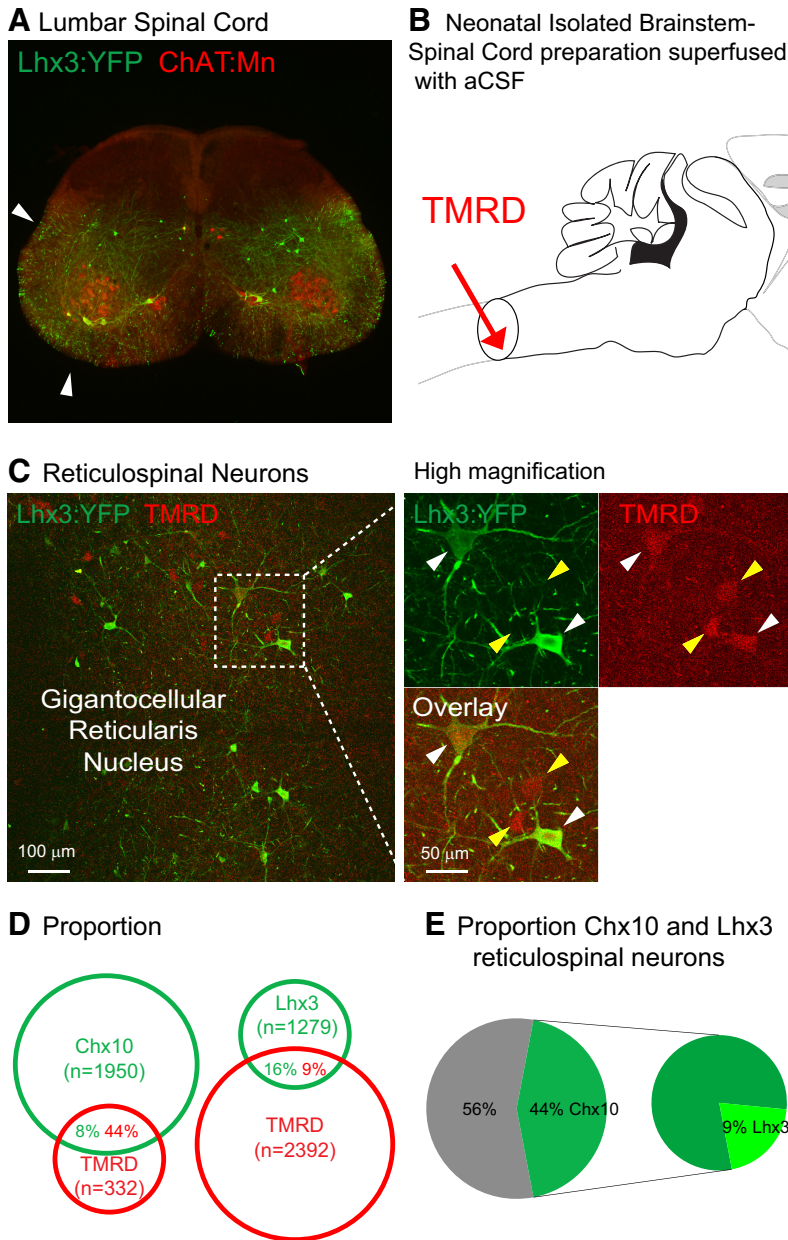


Figure 2. Lhx3 and Chx10 medRF neurons are reticulospinal neurons. **A**, Adult Lhx3-YFP upper lumbar spinal cord section with ChAT-positive motoneurons. Arrows indicate Lhx3-YFP axons in both ventrolateral and ventral funiculi. **B**, Schematic showing the application of retrograde tracer TMRD at cervical level in neonatal brainstem-spinal cord preparations. **C**, TMRD back-labeled Lhx3-YFP medRF neurons at low (left) and high magnifications (left, dashed window). White arrowheads indicate TMRD back-labeled Lhx3-YFP neurons. Yellow arrowheads indicate Lhx3-YFP-negative TMRD-positive reticulospinal neurons. **D**, Proportions of fluorescent protein-TMRD cells over fluorescent protein (green) and TMRD (red) cells ($n = 3$ Lhx3-YFP and $n = 3$ Chx10::GFP neonatal preparations). **E**, Percentages of Chx10 and Lhx3 neurons among TMRD-positive reticulospinal neurons.

sections (Wilson et al., 2010). Primary antibodies included mouse or sheep anti-GFP (1:500; Biogenesis), goat anti-5HT (1:500; Immunostar), goat anti-Chx10 (1:500; Millipore Bioscience Research Reagents), goat anti-Gata2/3 (1:500; Santa Cruz Biotechnology), goat anti-Fos (1:1000; Millipore Bioscience Research Reagents), mouse anti-GAD65 or anti-GAD67 (1:500), guinea-pig anti-vGluT1 (1:2500), guinea-pig anti-vGluT2 (1:2500), goat anti-choline acetyltransferase (ChAT, 1:100; all from Millipore Bioscience Research Reagents), and anti-Cav3.1 (1:2500; provided by Drs. Gerald Zamponi, University of Calgary, and Terry Snutch, University of British Columbia). Appropriate secondary antibodies were conjugated to Alexa 488, Alexa 555, Cy3, or Alexa 637 (1:500; Invitrogen), or rhodamine or Cy5 (1:100; Jackson ImmunoResearch).

Combined fluorescent *in situ* hybridization and immunohistochemistry was performed as previously described (Wilson et al., 2005).

Anterograde tracing. Biotinylated dextran amine (BDA, 10 kDa, 10% in sterile water; Sigma-Aldrich) was injected in the cuneiform nucleus [CnF; $n = 4$ mice; anteroposterior (AP), -5 mm; mediolateral (ML), 1.25 mm; depth, 1.9 mm] and the pedunculopontine nucleus (PPN; $n = 2$ mice; AP, -4.6 mm; ML, 1.25 mm; depth, 2.25 mm) of anesthetized adult Lhx3-YFP mice, as described previously (Bretzner et al., 2008, 2010). After 1 week, mice were perfused and tissues harvested and labeled.

Retrograde tracing. tetramethylrhodamine (TMRD; Sigma-Aldrich) was applied with insect pins to the cervical level of Lhx3-YFP ($n = 3$) and Chx10::GFP ($n = 3$) neonatal superfused brainstem-spinal cord preparations. After 6–8 h at room temperature, the tissues were prepared for subsequent immunohistochemistry.

Images were acquired on a Zeiss LSM510 Meta confocal microscope or Zeiss Axioplan II microscope. Analyses were performed using ImageJ for cell morphology (Fig. 1) and colocalization (Figs. 2, 3, 4). The number of observations was counted for bins of $50 \mu\text{m}^2$ for cumulative histograms (Fig. 1). All images presented are single optical sections.

Electrophysiology: patch clamp. Postnatal day 10–14 Lhx3-YFP mice were deeply anesthetized with ketamine/xylazine mix (100/10 mg/kg body weight; i.p.). The animal was decapitated, and the brainstems were harvested under cold ($<4^\circ\text{C}$) calcium-free sucrose-substituted artificial CSF (aCSF). Transverse brainstem slices ($250 \mu\text{m}$) were cut with a vibrating microtome (Vibratome 3000) and transferred to warm ($32\text{--}37^\circ\text{C}$) aCSF containing the following (in mM): 127 NaCl, 1.9 KCl, 1.3 MgCl_2 , 2.4 CaCl_2 , 2.4 NaH_2PO_4 , 26 NaHCO_3 , 10 D-glucose, and oxygenated with 95% O_2 –5% CO_2 for at least 30 min. Slices were transferred to the recording chamber and superfused with room temperature aCSF.

YFP neurons were identified under epifluorescence and visualized with infrared differential interference contrast microscopy using a Leica DMLFSA upright microscope. Whole-cell patch-clamp recordings were acquired with a MultiClamp 700A amplifier, a Digidata 1322A analog-to-digital converter (Molecular Devices), and pCLAMP software (version 9, Molecular Devices). The recording electrodes ($4\text{--}6 \text{ M}\Omega$) contained an internal solution made up of the following (in mM): 140 KMeSO_4 , 10 NaCl, 1 CaCl_2 , 10 HEPES, 1 EGTA, 3 Mg-ATP.

The pH was adjusted to 7.2 with KOH. The voltages presented were not corrected for junction potential, expected to be ~ 8 mV with these solutions.

Two-photon calcium imaging in the hemi-brainstem preparation. Postnatal day 1–5 Chx10::eGFP mice were deeply anesthetized with ketamine/xylazine mix (100/10 mg/kg body weight; i.p.) and decapitated. The brainstem was dissected under cold ($<4^\circ\text{C}$) aCSF, then hemisected through the midline to expose Chx10::GFP-expressing neurons in the medRF. The hemi-brainstem was then pinned, medial side up, in a customized recording chamber superfused with oxygenated aCSF. As previously described (Stosiek et al., 2003; Wilson et al., 2007), Chx10::GFP-expressing neurons were

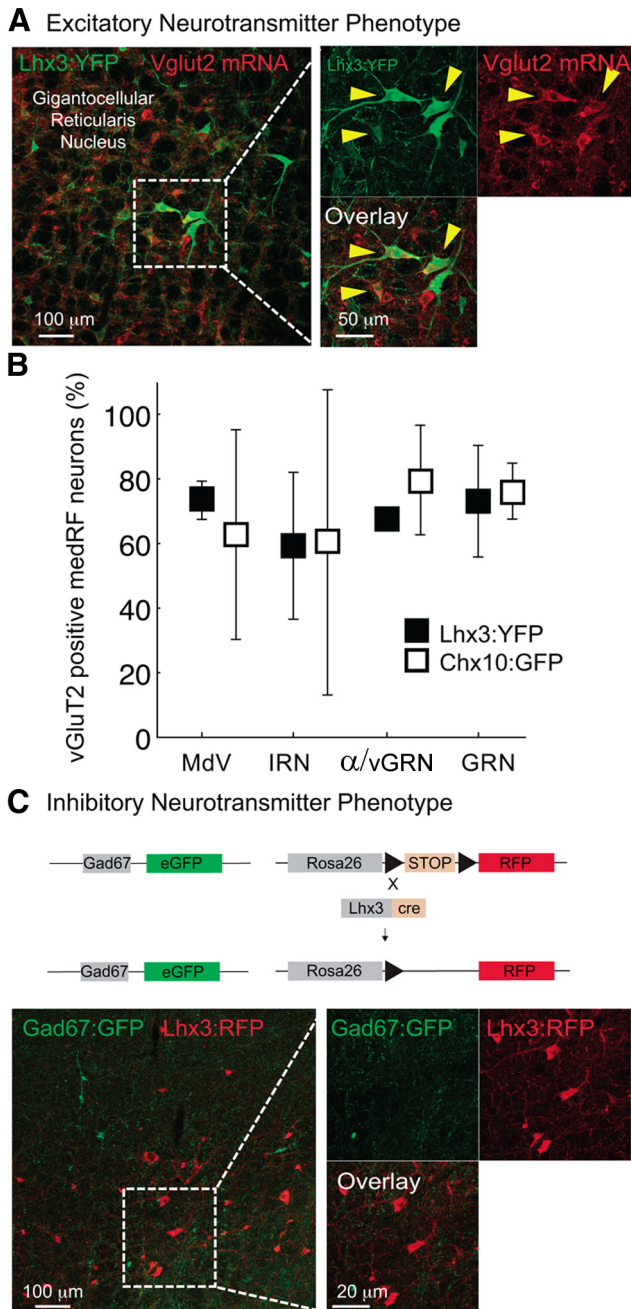


Figure 3. Lhx3 and Chx10 medRF neurons are glutamatergic. **A**, *In situ* hybridization for vGluT2 mRNA and immunohistochemistry for Lhx3-YFP medRF neurons. **B**, Mean and STD of proportions of vGluT2-expressing Lhx3-YFP (filled square) and Chx10:GFP (open square) medRF neurons across nuclei. **C**, Immunohistochemistry for Lhx3-RFP (red) and GAD67:GFP (green) medRF neurons showing lack of overlap.

loaded with Fluo-3 AM at 100 μ M (Invitrogen) using pressure injection. After 30 min recovery, brainstem neurons were visualized with a Zeiss Axioskop multiphoton microscope (Wilson et al., 2007).

GFP-expressing neurons could be readily identified, and time series of fluorescent images of calcium activity were collected with the following parameters: 1024 \times 1024 pixel images, digital zoom 4–8 times with 40 \times objective, 800 frames, 200–400 ms/frame. Bidirectional scanning was used to increase scan speed. Calcium transients were recorded as changes in mean pixel intensity in a defined region of interest over time and expressed as the percentage change in fluorescence divided by the baseline fluorescence ($\Delta F/F$).

Electrodes were placed in the region of the MLR identified by surface anatomy. This region was stimulated by delivering trains of stimuli [5–150 μ A (mean $117 \pm 7 \mu$ A), cathodal current dependent on threshold of response, 50 pulses at 10 Hz, pulse duration 0.2 ms, train duration 5 s] with monopolar tungsten microelectrodes (WPI), using a protocol similar to that described previously (Szokol et al., 2008). The stimulation train and a transistor–transistor logic (TTL) pulse initiated at the start of the frame scan were digitized and captured by the acquisition software to synchronize the onset and termination of stimulation to the calcium imaging traces offline.

The electrode was coated with the fluorescent dye 1,1'-dioctadecyl-3,3,3',3'-tetramethyl-indocarbocyanine perchlorate (DiI, solution saturated in absolute alcohol; Invitrogen) to localize electrode trajectories (Szokol et al., 2008). Sections containing DiI traces were imaged and used to identify the position of the effective stimulation sites. These sites were plotted on anatomical sections of a neonatal mouse brain adapted from (Paxinos and Franklin, 2004) and the Allen Mouse Brain Atlas (<http://mouse.brain-map.org/static/atlas>). Data obtained using stimulation sites outside of the cuneiform/PPN were discarded.

Fos expression. The activity-dependent immediate early gene *c-fos* was assessed in adult mice walking either on a wheel for 60 min or on a treadmill at 20 cm/s for 30 min. All mice were trained to walk on either device for 4 d, and assessed on the fifth day. Under control conditions, mice were kept in a quiet and illuminated environment for the same period. Mice were then anesthetized as above and perfused within 45–60 min of the test, and their brainstems harvested and processed for Fos immunoreactivity. The numbers of double-labeled neurons were quantified. Mice were paired—one walking with one control from the same litter at the same time of day—to remove variation based on time of day or litter, and the paired results presented normalized to the number of neurons that were Fos⁺ in the control animals.

All figures reported and bars shown are means \pm SDs.

Results

Lhx3⁺ neurons are Chx10⁺ and are located in locomotor regions of the medRF

We first sought to determine the location of Lhx3⁺ neurons in the medRF. In the caudal brainstem (Fig. 1A), YFP⁺ neurons are distributed across the medulla including the ventral part of the medullary reticular nucleus (MdV) and the intermediate reticular nucleus (IRN). In the rostral medulla, they are in the gigantocellular reticular nucleus (GRN) and in both ventral and α parts (sometimes referred to as “magnocellular”) of the gigantocellular nuclei (together termed α /vGRN; Fig. 1B). Lhx3⁺ neurons were typically large cells, with the largest neurons located in the most rostral and dorsal parts of the medulla in the GRN and IRN, and the smallest in the most ventral and caudal medulla in the MdV and α /vGRN (Figs. 1D–F, 2C). The location of these neurons in known brainstem motor regions (Noga et al., 2003; Szokol et al., 2008) together with the size suggesting they may be involved in fast-conducting pathways, indicates that they may be good candidates for neurons involved in locomotion.

In the spinal cord, Lhx3-expressing interneurons (V2 neurons) are known to differentiate into V2a neurons, which are Chx10⁺, V2b neurons, which express Gata2/3, and a subgroup of V2c neurons, which express Gata2/3 and Sox1 (Karunaratne et al., 2002; Li et al., 2005; Al-Mosawie et al., 2007; Lundfald et al., 2007; Panayi et al., 2010). We therefore asked whether a similar genetic lineage is seen in the brainstem. Initially, we labeled Gata2/3 and Chx10 protein in early postnatal Lhx3-YFP mice, and found Gata2/3⁺ cells located mainly in the ventral, midline, or dorsolateral medulla [Fig. 1H; similar to mouse (Hafler et al., 2008; Liu et al., 2010; Crone et al., 2012) and zebrafish (Kinkhabwala et al., 2011; Koyama et al., 2011)]. None of Gata2/3 cells coexpressed YFP, suggesting that differentiation pathways in the brainstem are different from those in the spinal cord.

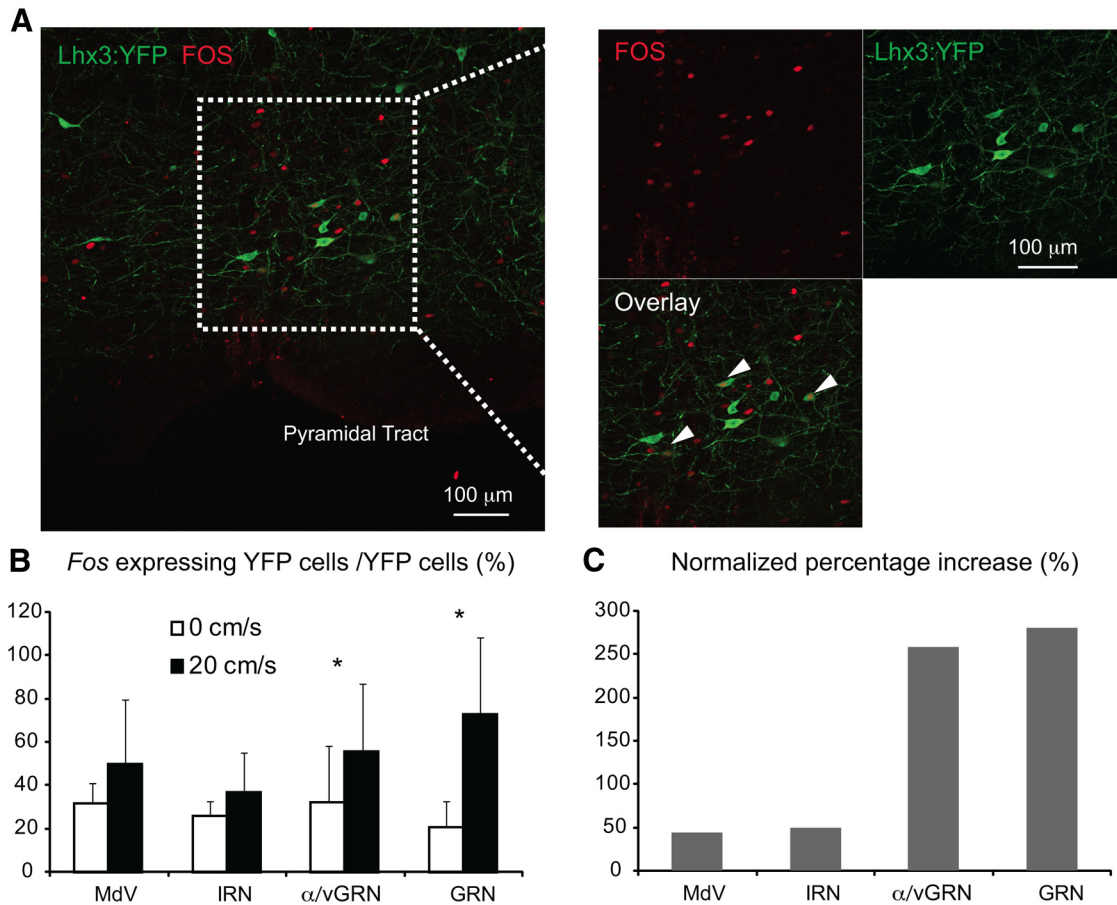


Figure 4. Genetically identified Lhx3 and Chx10 medRF neurons are active during *in vivo* locomotion. **A**, Fos expression in Lhx3 medRF neurons of a P21 Lhx3-YFP mouse following a locomotor task. **B**, Proportions (mean and STD) of Fos-expressing Lhx3 and Chx10 medRF neurons from three pairs of mice kept at rest ($n = 3$, open bar) or following locomotor task ($n = 3$, filled bar). **C**, Paired analysis showing the normalized percentage increase of Fos-expressing (Lhx3 and Chx10 together) genetically identified medRF neurons from the data illustrated in **B**.

In contrast, Chx10-immunoreactivity was seen in YFP+ neurons in the GRN, α/v GRN, IRN, and MdV (data not shown). Given this colocalization, we investigated the distribution of Chx10+ neurons in Chx10::eGFP mice and found the distribution and neuronal sizes to be similar to those of Lhx3+ neurons (Fig. 1C,F,G). Furthermore, in Chx10::eGFP;Lhx3-RFP triple-transgenic mice, almost all Lhx3-RFP+ neurons (excepting a few large dorsal neurons) across the four nuclei mentioned above also expressed GFP, indicating coexpression of Chx10 and Lhx3 ($n = 3$ mice; Fig. 1I). That is, Lhx3+ neurons in the medRF are Chx10+ and are found in known locomotor-related regions in the mouse (Häggglund et al., 2010). Given that Lhx3+ neurons are a subset of Chx10+ neurons, we will generically refer to these neurons as Chx10 medRF neurons.

Lhx3 and Chx10 medRF neurons are reticulospinal neurons

We next asked whether medRF Lhx3/Chx10 neurons are reticulospinal. We had noted Chx10::eGFP-labeled and Lhx3-YFP-labeled axons in the white matter of the spinal cord, especially in the ventral, ventrolateral, and ventromedial funiculi, where reticulospinal axons run (Fig. 2A; Al-Mosawie et al., 2007). We applied a retrograde tracer, TMRD, to the cervical spinal cords (C2–C3) of neonatal brainstem-spinal cord preparations (Fig. 2B, $n = 3$ postnatal day 10 Lhx3-YFP and $n = 3$ postnatal day 0 Chx10::GFP mice). Given that TMRD does not fill all axons (Lundfald et al., 2007), the number of

neurons labeled would underestimate the number of reticulospinal neurons and the data may be skewed toward the identification of reticulospinal neurons with larger axons. Approximately 1 of 10 TMRD-labeled reticulospinal neurons were Lhx3+, whereas almost one-half were Chx10+ (Fig. 2D,E). This difference is in keeping with the incomplete recombination seen with Lhx3-cre animals, and suggests that approximately one-half of reticulospinal neurons are Chx10+. The proportion of fluorescent protein-labeled neurons that were TMRD labeled was 16% for Lhx3-YFP neurons but only 8% for Chx10::eGFP neurons. This difference likely indicates that Chx10 neurons are also involved in non-reticulospinal pathways (Crone et al., 2012). Together these data suggest that, while these neurons may be involved in systems other than reticulospinal, Lhx3+ and Chx10+ neurons form a significant component of the reticulospinal system.

Lhx3 and Chx10 medRF neurons are glutamatergic

As the fast-descending system for initiating locomotion is glutamatergic (Douglas et al., 1993), we next sought to determine whether these neurons express the glutamate transporter vGluT2 (Fig. 3A). We found that ~80% of Lhx3-YFP neurons ($n = 2$ mice) and Chx10::eGFP neurons ($n = 4$ mice) in the GRN express vGluT2 (Fig. 3B), indicating that they are glutamatergic. We next crossed Lhx3-RFP mice with GAD67::eGFP mice to determine whether any Lhx3 neurons were GABAergic. None of Lhx3-RFP cells were positive for GAD67::eGFP (Fig. 3C; $n = 4$ mice, six sections per animal). Together with

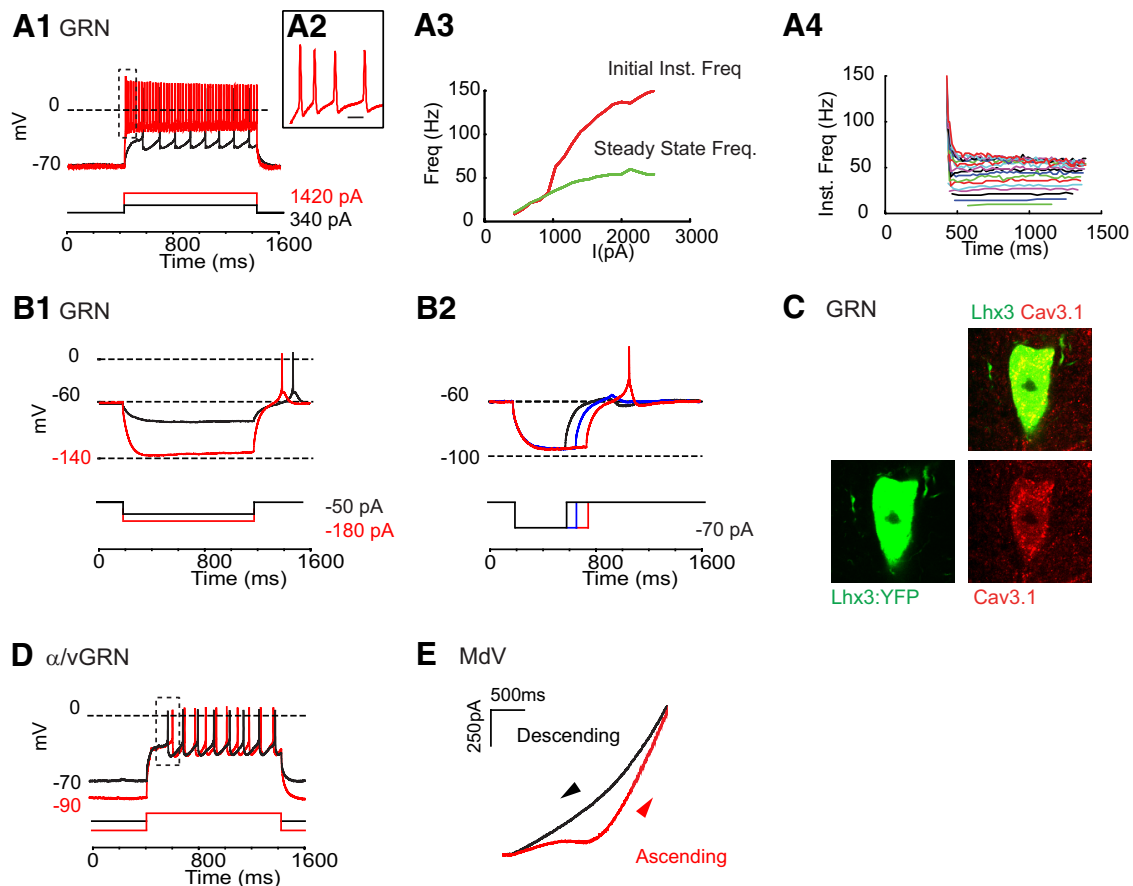


Figure 5. Electrophysiological properties of Lhx3 medRF neurons. **A**, Tonic repetitive firing of a Lhx3 GRN neuron at low (**A1**) and high (**A2**) time sweeps. Scale bar, 10 ms. **A3**, Instantaneous frequency versus current injection. **A4**, Instantaneous frequency versus time. Each trace represents a different amount of current injected. **B**, Postinhibitory rebound with spike of a GRN neuron depends on the amplitude (**B1**) and duration (**B2**) of membrane hyperpolarization. **C**, Immunolabeling for Cav3.1 onto a GRN neuron. **D**, First-spike latency delayed following membrane hyperpolarization of an α/v GRN neuron. **E**, Voltage clamp demonstrates region of negative slope conductance during voltage ramp (0.03 mV/ms for 3 s) in a MdV neuron.

the expected underestimation of mRNA expression when combining fluorescent *in situ* hybridization (for vGluT2) with fluorescent immunohistochemistry, we conclude that medRF Lhx3 and Chx10 neurons—which belong to the same population and are reticulospinal neurons—are glutamatergic.

Lhx3 and Chx10 neurons are activated during locomotion

We next asked whether Lhx3 medRF neurons are recruited during locomotion in adult mice (Fig. 4A). The expression of *c-fos*, an activity-dependent immediate early gene, has previously been used to determine whether neurons are activated in locomotion (Dai et al., 2005; Wilson et al., 2005; Dai and Jordan, 2010a,b; Wilson et al., 2010). The proportion of fluorescent protein-positive neurons in the GRN expressing Fos increased significantly following a locomotor task (Fig. 4B), with ~80% of fluorescent protein-expressing neurons in the GRN expressing Fos following locomotion. To account for variability between litters and experiments, we paired the data, comparing control and experimental animals from the same litter. In these paired experiments, the normalized percentage increase of Fos expression was ~250% in both the GRN and α/v GRN ($p < 0.05$; Fig. 4C) compared with only ~50% in the other regions. This indicates that a considerable proportion of Lhx3/Chx10 brainstem neurons in the α/v GRN and GRN are activated during locomotion in the adult mouse.

Table 1. Physiological properties of Lhx3-YFP medRF neurons (mean \pm SD)

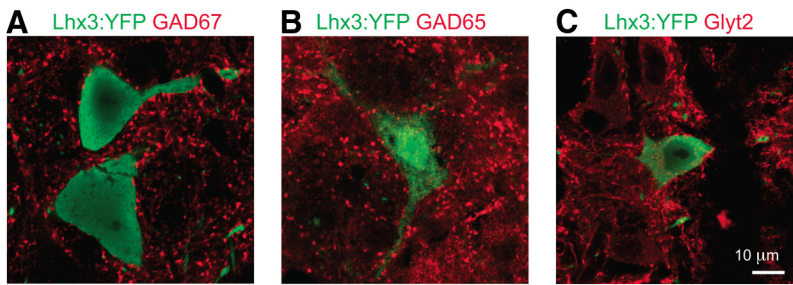
	All Lhx3 neurons (n = 72)	MdV (n = 12)	IRN (n = 8)	α/v GRN (n = 9)	GRN (n = 19)
Resting membrane potential (mV)	-50 ± 8	-51 ± 7	-49 ± 9	-50 ± 6	-53 ± 8
Capacitance (pF)	70 ± 50	68 ± 28	57 ± 23	70 ± 65	83 ± 64
Input resistance (M Ω)	83 ± 58	81 ± 29	59 ± 24	108 ± 81	96 ± 76
Spike threshold (mV)	-40 ± 5	-41 ± 4	-39 ± 5	-42 ± 3	-41 ± 6
Spike amplitude (mV)	71 ± 20	75 ± 24	68 ± 15	65 ± 18	77 ± 17
Max initial frequency (Hz)	52 ± 36	57 ± 27	47 ± 26	43 ± 22	73 ± 50
Max steady-state frequency (Hz)	24 ± 11	26 ± 11	23 ± 8	23 ± 5	32 ± 14
AHP duration (ms)	238 ± 115	282 ± 142	256 ± 112	293 ± 113	211 ± 102
f-I gain (Hz/100 pA)	9 ± 8	10 ± 10	8 ± 2	7 ± 3	11 ± 8

Chx10 medRF neurons have diverse electrophysiological properties

Because the physiological properties of neurons are critical to the function of the networks in which they are involved (Harris-Warrick, 1993; Marder and Bucher, 2007), we compared properties of YFP+ neurons of different nuclei in brainstem slices of postnatal day 10–14 Lhx3-YFP mice (Fig. 5, Table 1). Unfortunately, we were unable to successfully patch clamp healthy neurons in slices following the long incubation times needed for retrograde labeling, so neurons were identified based on their location and expression of fluorescent proteins. To ensure inclu-

Table 2. Proportion of Lhx3-YFP brainstem neurons

	All Lhx3 neurons	MdV	IRN	α/v GRN	GRN
Tonic firing	92% (66/72)	100% (12/12)	100% (8/8)	89% (8/9)	80% (15/19)
Doublet	22% (16/72)	23% (3/13)	43% (3/7)	10% (1/10)	25% (5/20)
Sag	60% (39/66)	64% (7/11)	75% (6/8)	67% (6/9)	42% (7/17)
PIR	43% (27/63)	36% (4/11)	37% (3/8)	33% (3/9)	59% (10/17)
Delayed first-spike latency (I_A)	50% (17/34)	45% (5/11)	72% (5/7)	67% (2/3)	28% (2/7)
Region of negative slope conductance	44% (24/55)	88% (7/8)	28% (2/7)	50% (4/8)	33% (5/15)

**Figure 6.** Inhibitory inputs to Lhx3-YFP MedRF neurons. **A–C**, GAD67 (**A**), GAD65 (**B**), and Glyt2 (**C**) boutons in apposition to Lhx3-YFP medRF neurons.

sion of healthy neurons only, neurons with peak-to-peak action potential amplitudes >60 mV were used in this analysis ($n = 72$ neurons). The resting potentials and input resistances of neurons across the four nuclei were similar, but as expected the larger cells of the GRN (Fig. 1D) tended to have higher whole-cell capacitances (Table 1).

Given that tonic descending commands can activate spinal locomotor circuits, we determined whether Chx10 medRF neurons could sustain repetitive firing. Injections of current led to sustained tonic repetitive firing in 66 of 72 (92%) neurons across the four nuclei (Fig. 5A, Table 2). The input–output relation, as assessed by frequency–current (f – I) gain, was similar across the four nuclei (9 ± 1 Hz/100 pA), although 1 of 9 neurons in the α/v GRN had a slope >10 Hz/100 pA (Fig. 5A3). Spike frequency adaptation was seen in neurons of all nuclei, particularly with the higher current injections (Fig. 5A4; 38 of 72 neurons had ratio of second to last instantaneous frequencies >1.5). There was no difference in afterhyperpolarization (AHP) duration (Table 1), although only neurons within the GRN (3 of 15) had AHP durations <110 ms and none of the neurons in the GRN (as opposed to each of the other nuclei) had AHP durations >400 ms. Together these data indicate that most neurons in medRF locomotor regions could repetitively fire in a steeply graded fashion, and could sustain this firing. Reticulospinal neurons would be a subset of those recorded and, given the ubiquity of these findings, would be expected to have these same properties.

We had also noticed that some neurons in each nucleus fired with doublets at the onset of the higher-magnitude step depolarizations (Fig. 5A2,A3; 16 of 72 neurons had ratio of first to second instantaneous frequencies >1.5). The maximum initial instantaneous frequency was similar across all four nuclei, but neurons in the α/v GRN and IRN never had doublets >100 Hz, and only 1 of 12 in MdV had a doublet >100 Hz. This contrasts with the GRN, in which 4 of 19 had doublets between 110 and 200 Hz. We next asked whether postinhibitory rebound (PIR) could account for the doublets. PIR ensures rapid firing of neurons following release of inhibition, and is prominent in some neurons involved in motor circuits (Perkel and Mulloney, 1974; Willows, 1980; Miller

and Selverston, 1982). As with the high-frequency initial doublets, PIR was found most commonly in GRN neurons and was dependent on the amplitude and duration of the preceding membrane hyperpolarization (10 of 17 neurons; Fig. 5B1,B2), which is consistent with it being produced by a Ca_v3 (T-type) current, and consistent with $Ca_v3.1$ expression (Fig. 5C). As GAD65-IR, GAD67-IR, and Glyt2-IR boutons are found in close apposition to these neurons (Fig. 6), it is possible that during circuit operation, inhibition could prime GRN neurons for rapid-onset firing.

Some central bursting neurons coexpress A-type and Ca_v3 -type currents (Molineux et al., 2005; Anderson et al., 2010), and together provide a mechanism by which PIR can be modulated (Harris-Warrick et al., 1995a,b). We therefore asked whether A-type currents, which lead to a delay to first spike following hyperpolarization, are present in Chx10 medRF neurons. A delay to first spike, dependent on magnitude of hyperpolarization, was most commonly seen in α/v GRN, MdV, and IRN neurons, and was conspicuously absent in GRN neurons, the latter group being more likely to express T-type currents (Fig. 5, compare D, B1). Thus, there was little evidence of

coexpression of A-type and T-type currents in YFP+ neurons. Finally, neurons involved in motor circuits may express persistent conductances (Na^+ and/or Ca^{2+}) to maintain firing (Miles et al., 2005) and/or amplify synaptic input (Hultborn et al., 2004). Slow ramp depolarizations in voltage-clamp experiments revealed that neurons, particularly in the MdV (88%), had regions of negative slope conductance (Fig. 5E; Schwandt and Crill, 1977). This contrasts with only 37% of neurons in the other nuclei (Table 2, Fisher's exact test $p = 0.06$) and suggests that MdV neurons in particular may have the capacity to fire in a sustained manner with minimal synaptic input, and/or to significantly amplify their inputs.

In summary, although some electrophysiological properties may differ in incidence between the nuclei, we have found no clear electrophysiological signatures of Lhx3/Chx10 neurons that correspond to the particular nuclei of the medRF. The electrophysiological properties of these neurons are heterogeneous across these four nuclei.

Lhx3 and Chx10 medRF neurons receive inputs from the MLR

If Chx10 medRF neurons mediate a descending command for locomotion, they should receive input from the MLR, a region known to initiate locomotion in various species (Shik et al., 1966; Orlovskii, 1970; Steeves and Jordan, 1984; Garcia-Rill et al., 1987; Mori, 1987; Takakusaki et al., 2003a). The MLR corresponds to the CnF, with the adjacent PPN also implicated in motor control (Garcia-Rill et al., 1985; 1990; Milner and Mogenson, 1988; Jordan, 1998; Takakusaki et al., 2003a, b). Two weeks following stereotactic injection of an anterograde tracer, BDA, into the CnF and PPN ($n = 6$) of adult Lhx3-YFP mice (Fig. 7A), BDA-labeled axon terminals were found in close proximity to Lhx3-YFP brainstem neurons, particularly in the rostral medRF in the α/v GRN (Fig. 7B,C), and some terminals in the GRN. These data support anatomically that neurons in both the α/v GRN and GRN receive MLR inputs.

To determine whether there is functional connectivity, we used multiphoton microscopy (Fig. 8A) for calcium imaging of Chx10

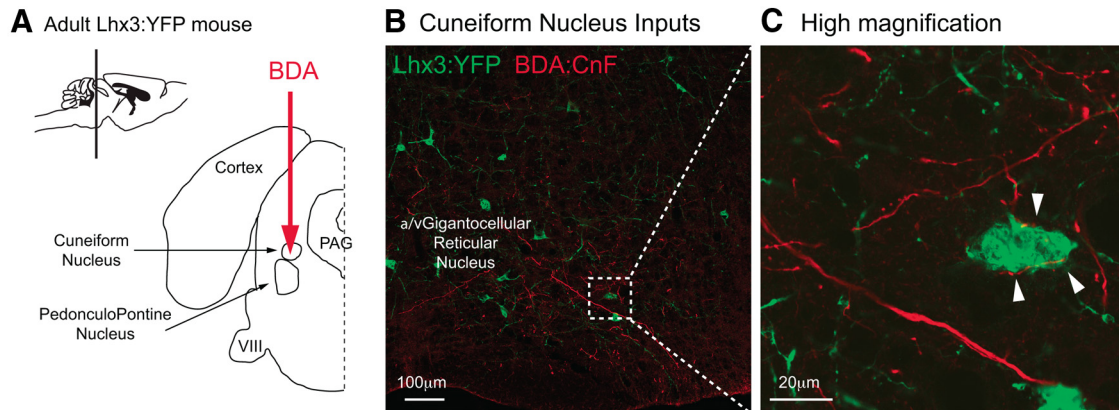


Figure 7. MLR projections to Lhx3-YFP medRF neurons. **A**, Schematic illustrating the injection site of an anterograde tracer, BDA, within the cuneiform nucleus (MLR) in adult Lhx3-YFP mice. **B**, BDA-filled axonal terminals in close proximity to Lhx3-YFP+ neurons located within the α/v GRN at low (**B**) and high (**C**, from the dashed window in **B**) magnification.

neurons while stimulating the region of the MLR (CnF and PPN). We detected calcium transients in 21% of Chx10::GFP neurons in the GRN and α/v GRN in response to MLR stimulation ($n = 30/144$ cells, $n = 13$ stimulating sites, 7 of which were identified by histology; Fig. 8B). The responses were not uniform: individual stimulation sites led to calcium transients in some Chx10::GFP neurons with no responses in others (Fig. 8D2). Stimulation in 5 of 7 identified sites (Fig. 8B, blue dots) led to increases in calcium transients in 23 of the 30 neurons (Fig. 8D1,D2). Stimulation in the other two sites (Fig. 8B, red dots) led to either decreases or poststimulation (i.e. after cessation of stimulation) increases in calcium transients in the other seven neurons (Fig. 8D3). These poststimulation increases in calcium transients could result from PIR (Fig. 5B). In general, stimulation sites in the dorsal area in the region of the CnF led to increases in calcium transients in Chx10::GFP neurons, while stimulation sites located in the region of the PPN evoked a mixture of decreases and increases in calcium transients in these neurons (Fig. 8D3). Taking the anatomical and physiological data together, we conclude that neurons in the region of the MLR project to and excite Chx10+ neurons in the medRF, thus supporting their potential role in locomotor circuits.

Discussion

While recent studies have furthered our understanding of spinal interneuronal networks that produce locomotor movements, little is known about the neurons that mediate the descending motor commands that initiate and modulate mammalian walking (Jordan et al., 2008). We have shown that Lhx3/Chx10-expressing neurons in the medRF are large glutamatergic neurons, a subset of which constitutes a subset of reticulospinal pathways. These neurons are activated during locomotion in adult mice, and receive functional synaptic inputs from the MLR, a supraspinal structure involved in initiation and maintenance of locomotion. Together, these data support a possible role for Lhx3/Chx10-expressing medRF neurons in mammalian brainstem locomotor networks.

Brainstem versus spinal cord Lhx3 and Chx10 neurons

Our data demonstrate a difference in differentiation between Lhx3-expressing neurons in the brainstem compared with the spinal cord. During spinal cord development, cells from the p2 progenitor domain express a combination of transcription factors, including Lhx3 (Zhou et al., 2000; Karunaratne et al., 2002; Li et al., 2005). These Lhx3+ V2 cells diversify into several distinct interneuronal populations: Chx10-expressing glutamatergic V2a and Gata2/3-expressing GABA/glycinergic V2b-c interneurons (Karunaratne et al., 2002; Li et al., 2005; Al-Mosawie et al., 2007; Lundfald et al., 2007; Batista et

al., 2008). In contrast, in the medulla, based on their size, location, and glutamatergic phenotype, YFP+ neurons in Lhx3-YFP animals seem to be a subset of GFP+ neurons in Chx10::eGFP mice. Furthermore, we found no Lhx3+/Gata3+ double-labeled cells in the medRF. Also, parallel to findings in chick (Cepeda-Nieto et al., 2005), all but a very few large Lhx3-expressing neurons in the dorsal aspect of the GRN expressed Chx10. The fact that the numbers of Lhx3-YFP neurons were lower than those of Chx10::eGFP neurons likely resulted from incomplete recombination using Lhx3^{cre/+} crossed with this thy1-fs-YFP reporter mouse (Al-Mosawie et al., 2007). This is not the first example in which neurons that developmentally express the same transcription factors in the brainstem and spinal cord differentiate into different subpopulations in each of these structures. For example, the majority of neurons expressing Dbx1 in the spinal cord are inhibitory interneurons (Pierani et al., 2001; Lanuza et al., 2004), whereas in the brainstem, Dbx1-expressing neurons are glutamatergic (Bouvier et al., 2010; Gray et al., 2010). In summary, our data support that, in contrast to the spinal cord, the majority of Lhx3+ neurons in the medRF differentiate into glutamatergic Chx10+ neurons.

Possible role of Chx10 medRF neurons in the descending control of locomotion

Reticulospinal neurons play a central role in locomotion, as demonstrated in *Xenopus* tadpoles (Roberts et al., 2008), fish (Brocard and Dubuc, 2003; Kinkhabwala et al., 2011; Koyama et al., 2011; Kimura et al., 2013), birds (Sholomenko et al., 1991), and mammals (Garcia-Rill and Skinner, 1987a; Zaporozhets et al., 2006; Häggglund et al., 2010). These neurons are activated by higher locomotor centers, such as the MLR (Noga et al., 2003), the cerebellar locomotor region (Homma et al., 1995; Mori et al., 1998), and subthalamic locomotor region/lateral hypothalamus (Sinnamon and Stopford, 1987), and are thus likely responsible for integrating these higher locomotor signals. In turn, they activate spinal locomotor networks (Steeves and Jordan, 1984; Shefchyk and Jordan, 1985; Garcia-Rill and Skinner, 1987a,b; Noga et al., 1988, 1991, 2003; Marlinsky and Voitenko, 1991). To determine whether a population of neurons in the reticular formation mediate the descending command for locomotion, they should meet the following criteria: (1) they should be reticulospinal; (2) they should have a glutamatergic transmitter phenotype; (3) they should be able to sustain repetitive firing; (4) they should be activated in locomotion; (5) they should receive inputs from higher locomotor centers; and at least one of: (6) their selective activation should produce sustained locomotion (sufficiency); and/or (7) their selective inhibition should inhibit locomotion (ne-

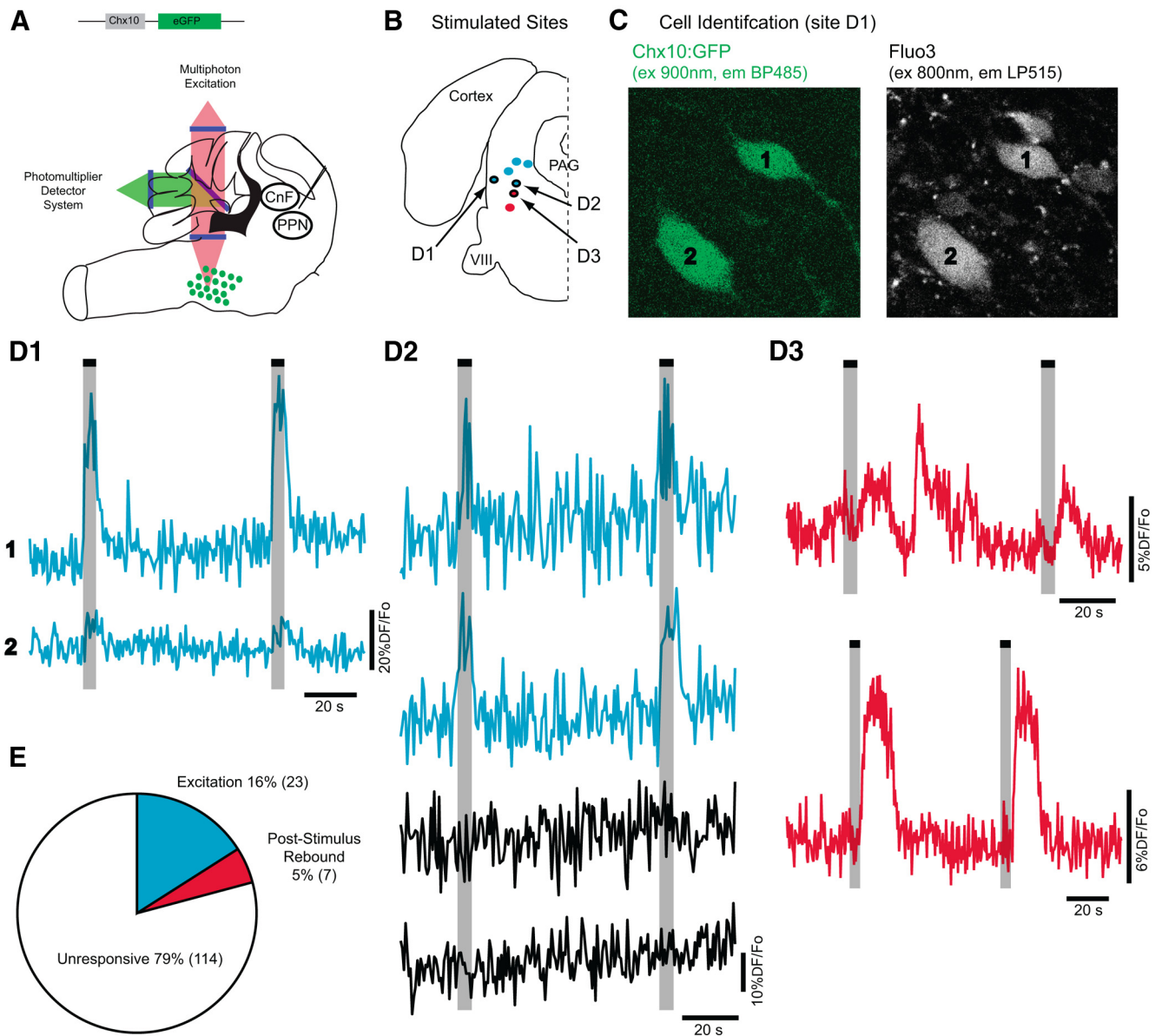


Figure 8. Functional connectivity of MLR to Chx10::GFP medRF neurons. **A**, Schematic demonstrating experimental paradigm. **B**, Stimulation sites confirmed by histology. Blue circles indicate sites that produced calcium increases, while red circles are sites that produced decreases or poststimulation increases in calcium signal. **C**, Identification of Chx10::GFP medRF neurons and calcium dye (Fluo3) by multiphoton microscopy. ex, Excitation wavelength; em, emission filter; BP, bandpass; LP, long pass. **D**, Calcium transients in medRF neurons in response to stimulation at the sites indicated with arrows and black circles in **B**. **D1**, Simultaneous calcium increases in two Chx10::GFP medRF neurons. **D2**, Another example of MLR-evoked calcium transients with prolonged calcium increases in two Chx10::GFP neurons (top row) but not in two neighboring GFP neurons (bottom row). **D3**, Example of calcium decrease and poststimulus rebound in two different Chx10::GFP neurons. **E**, Pie chart of responsiveness to MLR stimulation across 144 imaged Chx10::GFP medRF neurons. Poststimulus rebound responses were considered as inhibitory. A total of 21% Chx10::GFP cells were responsive (30 of 144 cells).

cessity). We have demonstrated that Lhx3-Chx10 neurons meet the first five of these criteria, and we thus suggest that they are excellent candidate neurons for mediating locomotor commands.

With respect to the first four criteria, we provide data demonstrating that Lhx3/Chx10 neurons are reticulospinal, similar to findings in chick and mouse embryos and larvae (Cepeda-Nieto et al., 2005; Kinkhabwala et al., 2011), and are glutamatergic, key requirements as previously reported in mammals (Douglas et al., 1993; Hägglund et al., 2010; Talpalar et al., 2011), and lower vertebrates (*Xenopus* tadpole, zebrafish, and lamprey; Brocard and Dubuc, 2003; Li et al., 2004; Kinkhabwala et al., 2011; Kimura et al., 2013). In addition, zebrafish Alx neurons (homologues of Chx10 neurons) were recently identified as hindbrain glutamatergic neurons, some of which are rhythmically active during swimming, project to the

spinal cord, and drive spinal locomotor circuits (Kinkhabwala et al., 2011; Kimura et al., 2013). We have demonstrated that mouse Chx10 brain stem neurons can sustain repetitive firing and, based on Fos expression, are activated in a locomotor task. While a necessary condition for neurons that would mediate locomotion, Fos expression does not demonstrate necessity for walking. For example, they may simply play an ancillary role in overground locomotion, such as in postural control, integration of locomotion with respiration, or integration of activity in ascending neurons. Thus the first four of the above seven criteria have been met by Lhx3-Chx10 medRF neurons.

We have also demonstrated that they receive functional input from higher locomotor centers. The projection of MLR neurons to Chx10 medRF neurons of the α /vGRN and GRN is in agreement with previous tracing and electrophysiological studies showing

monosynaptic connectivity from the MLR to reticulospinal neurons in various species, including cat, newt, lamprey, and salamander (Iwakiri et al., 1995; Bar-Gad et al., 1999; Brocard and Dubuc, 2003; Kagan and Shik, 2004). While our data do not define the anatomical correlates of the MLR, it is interesting that, as opposed to the few deeper stimulation sites, the more dorsal sites (in the region of the CnF nucleus) led to activation of Chx10 neurons. This is in agreement with previous studies in cat, which showed that stimulation of the CnF and dorsal PPN, through their effects on reticulospinal neurons, depolarized lumbar motoneurons, leading to increased muscle tone and locomotion, whereas stimulation of the ventral PPN led to tonic inhibition (Takakusaki et al., 1989, 2003b; Habaguchi et al., 2002). Excitation of Chx10 medRF neurons by MLR stimulation supports their potential role in mediating locomotor commands, and fulfills the fifth criterion above.

The last two criteria have proven to be much more difficult to demonstrate. Although a recent optogenetic study has demonstrated the critical role of Chx10 hindbrain neurons for swimming in the zebrafish (Kimura et al., 2013), photostimulation of glutamatergic hindbrain neurons in the mouse evoked only a few steps of locomotor-like activity (Hägglund et al., 2010). In addition, our attempts to selectively inactivate (using allatostatin receptor expression) or activate (using selective TRPV1 receptor or channelrhodopsin2 expression) brainstem Lhx3 neurons failed to consistently alter or drive locomotor activity (data not shown). This may be because Lhx3 is expressed in heterogeneous neurons. Even the reticulospinal subpopulation of Lhx3 neurons may be mixed, with some involved in activating movement, and others involved in inactivating movement (Takakusaki et al., 2003a). Thus we cannot definitively state whether these neurons, or subpopulations of these neurons, are either necessary or sufficient for locomotion. However, by fulfilling at least five of the seven criteria above, our data do support the idea that Lhx3/Chx10 medRF neurons in the GRN and α /vGRN are candidates for mediators of the descending control of mammalian locomotion.

Role of Chx10 medRF neurons in other motor functions

Chx10 medRF neurons are found in four discrete nuclei, each of which likely serves different functions. We have suggested that Chx10 neurons of the GRN and α /vGRN are involved in locomotor control. In contrast, we have shown that MdV neurons, which may be involved in licking behavior (Travers et al., 1997), have some distinct electrophysiological properties. It is possible that Lhx3/Chx10 medRF neurons could be involved in integrating supraspinal and spinal inputs to adapt and coordinate motor behaviors to environmental and intrinsic constraints (Prentice and Drew, 2001; Drew et al., 2004; Schepens and Drew, 2004, 2006). Recently, Chx10 brainstem neurons have been found to be important in the control of respiration: genetic deletion of Chx10 neurons led to irregular and low breathing frequency in isolated preparations and whole animals (Crone et al., 2012). Given that stimulation of the MLR, in addition to initiating locomotion, increases respiratory frequency (WalDROP and Iwamoto, 2006; Gariépy et al., 2012) and that locomotion modulates respiration through brainstem networks in the lamprey (Gravel et al., 2007), it is possible that Chx10 medRF neurons are involved in adjusting ventilation to motor output during locomotion. Such coupling of multiple motor circuits could adjust motor behavior to environmental constraints.

Possible role of Chx10 neurons in recovery of motor function

It has recently been shown that strategies that improve reticulospinal axon regeneration after spinal cord injury can lead to a worsened functional outcome (Lu et al., 2012). This, together

with the known roles of glutamatergic reticulospinal pathways in initiating locomotion (Hägglund et al., 2010) and their control over propriospinal (Zaporozhets et al., 2006) and spinal commissural neurons (Szokol et al., 2011), illustrates the importance of understanding the functions of identified reticulospinal neurons. Given their inputs from the MLR, it is possible that Chx10 reticulospinal pathways may contribute to reorganization after spinal cord injury. Moreover, given the convergence of diverse supraspinal, including cortical, inputs onto reticulospinal neurons (Drew et al., 2002; Jankowska and Edgley, 2006; Baker, 2011; Zaaïmi et al., 2012), they may also play a key role in rerouting descending motor commands after brain injury (Jankowska et al., 2005). Furthermore, recent trials of deep brain stimulation in the region of the dorsal mesopontine tegmentum (the region of the CnF and PPN) have shown some improvement in gait in people with Parkinson's disease (Stefani et al., 2007; Pereira et al., 2008). This improvement could be mediated via reticulospinal pathways, possibly Chx10 medRF neurons.

In the current study, we have identified reticulospinal neurons that may be involved in locomotor control. Such definition is necessary for the development of new strategies aimed at motor functional restoration in people with injuries and diseases affecting the CNS.

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