



# Guidance of longitudinally projecting axons in the developing central nervous system

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The directed and stereotypical growth of axons to their synaptic targets is a crucial phase of neural circuit formation. Many axons in the developing vertebrate and invertebrate central nervous systems (CNSs), including those that remain on their own (ipsilateral), and those that cross over to the opposite (commissural), side of the midline project over long distances along the anterior-posterior (A-P) body axis within precisely positioned longitudinally oriented tracts to facilitate the transmission of information between CNS regions. Despite the widespread distribution and functional importance of these longitudinal tracts, the mechanisms that regulate their formation and projection to poorly characterized synaptic targets remain largely unknown. Nevertheless, recent studies carried out in a variety of invertebrate and vertebrate model systems have begun to elucidate the molecular logic that controls longitudinal axon guidance.

**Keywords:** longitudinal axon guidance/targeting, Robo, CAMs, Sema, Wnt

## INTRODUCTION

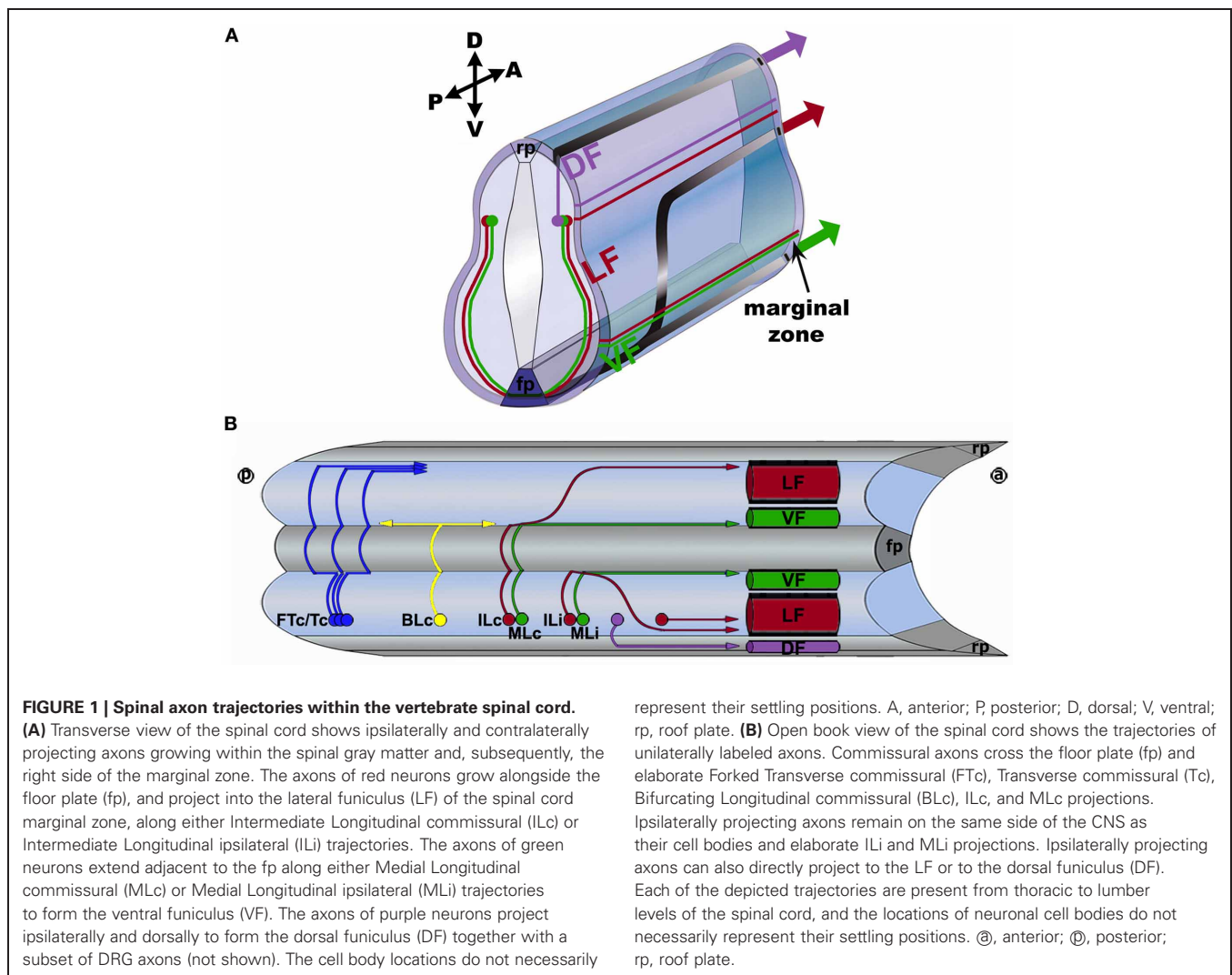
Longitudinally projecting axons connect different regions of the central nervous system (CNS) by extending over long distances along the anterior-posterior (A-P) body axis. Here, we review recent evidence supporting *in vivo* roles for long- and short-range guidance systems in regulating the pathfinding of longitudinally projecting ipsilateral and commissural axons. We first examine dye tracing and reporter gene expression data that reveal a previously unanticipated diversity and complexity of commissural and ipsilateral projections, which originate from neurons in the spinal cord and project longitudinally in the marginal zone; the outermost layer of the spinal cord proper that surrounds the gray matter. We then consider the reported roles of the morphogens, Sonic hedgehog (Shh) and Wnts, and potential interplay between these factors, in regulating the rostrally directed turn executed by post-crossing spinal commissural axons immediately after they cross the midline. Next, we address how Robo-Slit signaling, operating on its own or, potentially, in conjunction with particular cell adhesion molecules (CAMs), directs decussated commissural and ipsilateral axons into longitudinal tracts in the vertebrate CNS. We also evaluate the likely possibility that invertebrate model systems use a conserved set of, as well as unique, guidance cues and their receptors to ensure that commissural and ipsilateral axons are properly organized into stereotypically arranged longitudinal-oriented fascicles. In addition, we consider the mechanisms that appear to control the pathfinding of longitudinally projecting commissural and ipsilateral axons, which emanate from neurons located in various brain regions and descend into the spinal cord. Finally, we briefly discuss the

few studies aimed at identifying the synaptic targets of genetically distinct populations of neurons/axons in the vertebrate CNS.

## PATHFINDING OF LONGITUDINALLY PROJECTING AXONS IN VERTEBRATES: ASCENDING SPINAL PROJECTION AND DESCENDING MIDBRAIN NEURONS

### SPATIAL ORGANIZATION OF LONGITUDINAL AXON TRACTS WITHIN THE SPINAL CORD MARGINAL ZONE

By extending over long distances along the antero-posterior (A-P) axis of the CNS, ascending and descending spinal projection neurons transmit information to the brain and spinal cord, respectively. In vertebrates, projection neuron axons are contained within the spinal cord marginal zone (**Figure 1**). These axons assemble into longitudinal tracts or fascicles that are spatially organized in register with the positions of their brain targets (ascending axons) or brain origins (descending axons) in the spinal cord marginal zone (Burt, 1993; Brodal, 1998). Numerous anatomical and physiological studies have characterized the relative positioning of particular ascending and descending tracts within the spinal cord marginal zone (Burt, 1993; Brodal, 1998). For example, with specific regard to ascending projections, the spinothalamic tract is a major component of anterolateral system, which is housed within the ventral funiculus (VF) and ventrolateral funiculus (vLF) (Kerr, 1975; Giesler et al., 1981; Björkeland and Boivie, 1984; Willis, 2007). On the other hand, the spinocerebellar tract represents the predominant ascending pathway and is formed by axons located within both the dorsolateral funiculus (dLF) and the vLF (Xu and Grant, 1994, 2005; Willis, 2007). In addition, the dorsal funiculus (DF) contains the fasciculus gracilis



and the fasciculus cuneatus, which terminate in the medulla oblongata (Giesler et al., 1984).

#### DIVERSITY OF CONTRALATERAL COMMISSURAL AND IPSILATERAL PROJECTIONS IN THE VERTEBRATE SPINAL CORD

Although the spatial arrangement of longitudinally projecting tracts within the spinal cord marginal zone has long been appreciated, the trajectories that the component axons adopt in the spinal cord proper, and which presumably prefigure this organization, have only recently been characterized. Since most ascending and descending tracts are comprised of both ipsilateral and commissural projections, and given the bilateral symmetry of the spinal cord, “one-sided/unilateral” labeling strategies are required to visualize the individual axons or sets of axons contained within a particular tract and to clearly delineate its ipsilateral and commissural components. Utilizing the lipophilic axon-tracer, 1,1'-dioctadecyl-3,3,3',3' tetramethylindocarbocyanine perchlorate (DiI), unilateral labeling of open book spinal cord preparations derived from various age chick and mouse embryos was previously carried out to selectively characterize the projections

elaborated by post-crossing commissural axons. Importantly, this unilateral labeling strategy provided unobstructed views of post-crossing commissural axons and revealed a previously unappreciated complexity and diversity in their projections within the spinal cord marginal zone (Imondi and Kaprielian, 2001; Kadison and Kaprielian, 2004).

All spinal commissural axons initially project to their common intermediate target, the floor plate located at the ventral midline, along a simple linear pre-crossing trajectory (Bovolenta and Dodd, 1990; Imondi and Kaprielian, 2001) (Figure 1A). After crossing the floor plate, the contralateral segments of decussated commissural axons adopt at least five distinct trajectories (Figure 1B). The most commonly observed contralateral projections are elaborated by medial longitudinal commissural (MLc) axons, which extend in the longitudinal plane alongside the floor plate at the ventral midline and join/form the VF, and intermediate longitudinal commissural (ILc) axons, which initially grow alongside the floor plate and then project laterally, away from the ventral midline along an arcuate trajectory and ultimately turn longitudinally to form the LF (Bovolenta and Dodd, 1990; Imondi

and Kaprielian, 2001; Kadison and Kaprielian, 2004). After turning into the longitudinal plane most MLc and ILc axons project rostrally, however, a minor but significant subset of each population extends in the caudal direction (Bovolenta and Dodd, 1990; Kadison and Kaprielian, 2004). The other three, relatively minor projections are elaborated by: bifurcating longitudinal commissural (BLc) axons, which bifurcate into rostrally and caudally projecting branches either within the VF alongside the FP or at a significant lateral distance from the spinal cord, where they contribute to the LF, forked transverse commissural (FTc) axons, which bifurcate on the ipsilateral side first, cross the floor plate, project transverse to the floor plate, and appear to join the LF or DF, or transverse commissural (Tc) axons, which do not turn in the VF but, rather, project transverse to the floor plate, and likely join the LF and DF (Bovolenta and Dodd, 1990; Kadison and Kaprielian, 2004).

Over the past 10 years, the emergence of *in ovo* and *in utero* electroporation technologies, together with the development of a wide range of reporter constructs, has made it possible to visualize axon pathfinding in both the embryonic chick and mouse spinal cord (Krull, 2004; Saito, 2006). Most importantly, unilateral *in ovo* electroporation has been used to reproducibly label genetically distinct populations of spinal axons and to assess potential differences in the trajectories elaborated by specific spinal neuron populations. For example, dorsal spinal neurons, which represent a major class of projection neurons (Nunes and Sotelo, 1985; Burstein et al., 1990; Yeziarski and Mendez, 1991; Brodal, 1998), have been classified according to the particular transcription factor(s) they express (Helms and Johnson, 2003), and it has become possible to selectively visualize their pathfinding axons by electroporating reporter constructs harboring enhancer elements derived from the corresponding genes (Lumpkin et al., 2003; Nakada et al., 2004; Reeber et al., 2008; Avraham et al., 2009, 2010). These genetic labeling studies have revealed that the majority of d1 and d2 commissural axons adopt MLc and ILc trajectories and join the VF and LF, respectively, (Reeber et al., 2008; Avraham et al., 2009).

In addition to characterizing contralateral commissural projections, DiI tracing and unilateral *in ovo* electroporation of specific reporter constructs have also been used to identify three major projections elaborated by ipsilateral spinal neurons/axons. Specifically, major subsets of ipsilaterally projecting axons contribute to particular funiculi within the spinal cord marginal zone: (1) medial longitudinal ipsilateral (MLi) axons, which project along MLc axon-like trajectories, join the VF (Avraham et al., 2010), (2) intermediate longitudinal ipsilateral (ILi) axons, analogous to ILc axons, initially extend toward the ventral midline and then project away from the FP along an arcuate trajectory, or by directly projecting into lateral regions of the spinal cord, join the LF (Kadison and Kaprielian, 2004; Avraham et al., 2010), and (3) a subset of ipsilateral axons projects toward the marginal zone and turns into the longitudinal plane between the dorsal midline and the dorsal root, and join the DF (Avraham et al., 2010).

Whereas most spinal projection neurons extend ascending axons to the brain, some populations of spinal cord interneurons make local connections in segments located either above or below their cell bodies (Brodal, 1998; Kullander et al., 2003; Kiehn,

2006). Accordingly, DiI tracing and genetic labeling studies have identified a subset of caudally extending axons that contribute to the VF, LF and/or DF (Bovolenta and Dodd, 1990; Kadison and Kaprielian, 2004; Reeber et al., 2008; Avraham et al., 2009, 2010).

#### THE ROLE(S) OF MORPHOGENS IN DIRECTING THE ROSTRAL TURN EXECUTED BY DECUSSATED SPINAL COMMISSURAL AXONS

As described above, many commissural axons turn rostrally into the longitudinal plane after crossing the ventral midline. It has now become apparent that morphogens, once thought to exclusively control patterning events in the developing nervous system, regulate this key pathfinding decision faced by newly decussated commissural axons (Bovolenta, 2005; Charron and Tessier-Lavigne, 2005; Zou and Lyuksyutova, 2007; Sánchez-Camacho and Bovolenta, 2009). For example, Wnt4 (see **Table 1**) appears to have a critical role in controlling the rostral turn executed by MLc and ILc axons in the mouse spinal cord (Lyuksyutova et al., 2003). This is supported by the finding that floor plate expresses Wnt4 mRNA in a rostral (high) to caudal (low) gradient in mouse embryos, and that Wnt4-expressing cells attract decussated commissural axons *in vitro* (Lyuksyutova et al., 2003). In addition, mice lacking a Wnt receptor and the well-known planar cell polarity signaling molecule, *Frizzled3*, display a reduction in the number of DiI-labeled rostrally growing decussated axons (Lyuksyutova et al., 2003). Other planar cell polarity components, including *Drosophila* van Gogh ortholog Vangl2 and Flamingo ortholog Celsr3 (Tissir and Goffinet, 2010), determine the rostrocaudal polarity of axon growth by interacting with Frizzled3 (Shafer et al., 2011). Accordingly, it appears that Wnt4 and/or its receptor Frizzled3, as well as Vangl2 and Celsr3 have a major role in regulating the polarity of the stereotypical rostral turn executed by ascending post-crossing commissural axons.

In the chick spinal cord, the potent morphogen, Shh, which is selectively expressed at the ventral midline, also has an important role in regulating the rostral turning of post-crossing commissural axons (Bourikas et al., 2005). Contrasting the pattern of Wnt4 expression described in mouse embryos, Shh is expressed by floor plate cells in a caudal (high) to rostral (low) gradient in chick embryos. Accordingly, it has been proposed that Shh operates as a repellent for post-crossing commissural axons as they project along the A-P axis of the spinal cord. Supporting this notion, knock down of Shh expression through the use of long double-stranded RNA interference (dsRNAi) causes stalling of decussated DiI-labeled axons in the vicinity of the floor plate, and prevents them from turning in the rostral direction (Bourikas et al., 2005). Interestingly, although Wnt4 is not present within the floor plate of the chick spinal cord, a non-graded distribution of *Wnt5a* and *Wnt7a* mRNA has been detected at the ventral midline (Domanitskaya et al., 2010). In addition, a Wnt antagonist, *Secreted frizzled-related protein 1* (*Sfrp1*) mRNA is expressed within the floor plate, in an increasing rostrocaudal gradient, and ectopic expression of Shh induces *Sfrp1* expression. Together these intriguing observations raise the possibility that Shh guides post-crossing spinal commissural axons in the rostral direction, by inducing an attractive Wnt gradient in the chick spinal cord (Domanitskaya et al., 2010).

**Table 1 | Selected molecules associated with the pathfinding of longitudinally projecting axons.**

Name	Organism	Localization <sup>a</sup>	Suggested role in longitudinal guidance
Celsr3	Mouse	Spinal commissural axons; mdDA and hindbrain 5-HT axons	Rostral turn of spinal commissural axons (Shafer et al., 2011); rostrocaudal orientation of mdDA and hindbrain 5-HT axons (Fenstermaker et al., 2010)
EphA4	Mouse	Corticospinal axons	Prevents corticospinal axons from re-crossing in the spinal cord (Dottori et al., 1998; Kullander et al., 2001b; Yokoyama et al., 2001)
ephrinB3	Mouse	Ventral midline	Midline barrier for corticospinal axons in the spinal cord (Kullander et al., 2001a; Yokoyama et al., 2001)
FasII	<i>Drosophila</i>	Longitudinal axons	Fasciculation of FasII <sup>+</sup> longitudinal tracts (Lin et al., 1994)
lin-17 (Frizzled)	<i>C. elegans</i>	Posterior dendrite of PLM neurons	A-P orientation of PLM axon and dendrite (Hilliard and Bargmann, 2006)
mig-1 mom-5 (Frizzled)	<i>C. elegans</i>	?	Anterior orientation and growth of AVM and PVM longitudinal axons (Pan et al., 2006)
Frizzled3	Mouse	Spinal commissural axons; mdDA and hindbrain 5-HT axons	Rostral turn of spinal commissural axons (Lyuksyutova et al., 2003); rostrocaudal orientation of mdDA and hindbrain 5-HT axons (Fenstermaker et al., 2010)
L1	Mouse	Corticospinal axons at the pyramid	Pyramidal decussation; facilitating caudal growth below the pyramid (Cohen et al., 1997)
N-cadherin	<i>Drosophila</i>	Longitudinal and commissural axons	Fasciculation of FasII <sup>+</sup> and Apterous longitudinal tracts (Iwai et al., 1997)
NCAM	Mouse	Corticospinal axons at the pyramid	Pyramidal decussation; facilitating caudal growth below the pyramid (Rolf et al., 2002)
PSA-NCAM	Mouse	Corticospinal axons in the pyramid and in the DF	Facilitates collateral formation by corticospinal axons (Daston et al., 1996)
Npn2	Mouse	mdDA axons	Directs mdDA axons along specific routes and to the prefrontal cortex (Kolk et al., 2009; Yamauchi et al., 2009)
PlexinA	<i>Drosophila</i>	Longitudinal axons	Defasciculation of FasII <sup>+</sup> axons from intermediate to lateral fascicle (Winberg et al., 1998)
PlexinA3 PlexinA4	Mouse	(mRNA) corticospinal neurons; ( <i>PlexinA4</i> mRNA) along the corticospinal tracts in the hindbrain and inferior olive	Pyramidal decussation (Faulkner et al., 2008; Runker et al., 2008)
PlexinB	<i>Drosophila</i>	Intermediate and lateral region of neuropile; FasII <sup>+</sup> intermediate fascicle	Positioning and formation of FasII <sup>+</sup> intermediate fascicle (Wu et al., 2011)
Robo1 Robo2	Mouse Chick	Spinal commissural axons; descending midbrain axons; ascending mdDA axons; corticospinal and corticofugal tracts	Dorsoventral positioning of longitudinal tracts formed by spinal commissural axons (Reeber et al., 2008; Jaworski et al., 2010); fasciculation and organization of midbrain and mdDA axon tracts (Farmer et al., 2008; Dugan et al., 2011); restricting mdDA, midbrain, corticospinal and corticofugal axons to the ipsilateral side (Lopez-Bendito et al., 2007; Farmer et al., 2008; Dugan et al., 2011)
Robo1 Robo2 Robo3	<i>Drosophila</i>	Longitudinal axons in ventral nerve cord	Mediolateral positioning of longitudinal tracts (Rajagopalan et al., 2000; Simpson et al., 2000; Spitzweck et al., 2010)
Ryk	Mouse	Corticospinal axons	Facilitates the caudal growth of corticospinal tract (Liu et al., 2005)
Sema1A	<i>Drosophila</i>	Longitudinal and commissural axons	Defasciculation of FasII <sup>+</sup> axons from intermediate to lateral fascicle (Yu et al., 1998)
Sema2A	<i>Drosophila</i>	Midline and commissure in the ventral nerve cord	Repulsive boundary for positioning and growth of FasII <sup>+</sup> intermediate fascicle (Wu et al., 2011)
Sema2B	<i>Drosophila</i>	Medial and intermediate region of neuropile; FasII <sup>+</sup> intermediate fascicle	Fasciculation of FasII <sup>+</sup> intermediate fascicle (Wu et al., 2011)
Sema3F	Mouse	Midbrain-hindbrain border	Directs mdDA axons rostrally (Yamauchi et al., 2009); directs mdDA axons to the prefrontal cortex (Kolk et al., 2009)

(Continued)

Table 1 | Continued

Name	Organism	Localization <sup>a</sup>	Suggested role in longitudinal guidance
Sema6A	Mouse	(mRNA) along the corticospinal tracts in the hindbrain and inferior olive	Pyramidal decussation (Faulkner et al., 2008; Runker et al., 2008)
Sfrp1	Chick	(mRNA) ventral midline	Regulates rostral turn of spinal commissural axons by antagonizing Wnt5a and Wnt7a (Domanitskaya et al., 2010)
Shh	Chick	Ventral midline	Rostral turn of spinal commissural axons (Bourikas et al., 2005)
Slit1	Mouse	(mRNA) ventral midline; ( <i>Slit1</i> and <i>Slit2</i>	Positioning of longitudinal spinal axon tracts (Long et al., 2004);
Slit2		mRNA) forebrain-midbrain border and	fasciculation and organization of longitudinal midbrain and
Slit3		ventral forebrain structures	mdDA axon tracts (Farmer et al., 2008; Dugan et al., 2011); forming the ventral midline boundary in the brain (Bagri et al., 2002; Farmer et al., 2008; Dugan et al., 2011)
Unc5c	Mouse	(mRNA) cerebral cortical layer V and VI	Pyramidal decussation (Finger et al., 2002)
Vangl2	Mouse	mdDA and hindbrain 5-HT axons; (mRNA) spinal cord gray matter	Rostral turn of spinal commissural axons (Shafer et al., 2011); rostrocaudal orientation of mdDA and hindbrain 5-HT axons (Fenstermaker et al., 2010)
egl-20 (Wnt)	<i>C. elegans</i>	Tail	Anterior orientation and growth of AVM and PVM longitudinal axons via repulsion (Pan et al., 2006)
lin-44 (Wnt)	<i>C. elegans</i>	Posterior epidermal cells	A-P orientation of PLM axon and dendrite (Hilliard and Bargmann, 2006)
Wnt1 Wnt5a	Mouse	(mRNA) dorsal spinal gray matter; ( <i>Wnt5a</i> mRNA) ventral midline in the brain	Repels corticospinal tract <i>in vitro</i> (Liu et al., 2005); ( <i>Wnt5a</i> ) initial rostral bias of mdDA axon projection (Fenstermaker et al., 2010)
Wnt4	Mouse	(mRNA) ventral midline	Attracts spinal commissural axons rostrally <i>in vitro</i> (Lyuksyutova et al., 2003)
Wnt5a Wnt7a	Chick	(mRNA) ventral midline	Rostral turn of spinal commissural axons (Domanitskaya et al., 2010)

<sup>a</sup>Protein localization unless otherwise stated (i.e., mRNA).

Given that a subset of the ipsilaterally extending axons also project to the brain along MLI and ILI trajectories (see above), it would be interesting to determine whether ventral midline-associated Wnts and Shh regulate the rostral turn executed by these subsets of ascending axons. The findings of such experiments could conceivably reveal a previously unsuspected heterogeneity in the response of commissural and ipsilateral axons to morphogens that control the polarity of longitudinally projecting axons.

#### ORGANIZATION OF LONGITUDINAL AXON TRACTS: THE ROLE(S) OF ROBO-SLIT SIGNALING

After executing rostral or caudal turns into the longitudinal plane at various positions along the A-P axis of the CNS, the axons of both ascending and descending projection neurons are sorted into discrete longitudinal bundles or tracts located at specific distances from the midline (Burt, 1993; Brodal, 1998). In the vertebrate spinal cord, Robo receptors and their Slits ligands have well-established and critical roles in regulating the positioning of contralaterally ascending spinal commissural axon-containing longitudinal tracts (Dickson and Gilestro, 2006; Dickson and Zou, 2010). For example, Robo-Slit signaling normally drives spinal commissural axons along ILc trajectories and into the LF in the embryonic chick spinal cord (Reeber et al., 2008). By disabling Robo-Slit signaling via unilateral electroporation of spinal commissural neurons with cytoplasmic truncations (dominant-negative forms) of Robo1 or Robo2, a striking and

selective axon pathfinding phenotype was observed: post-crossing commissural axons failed to elaborate ILc projections and, instead, exclusively extended along MLc-like trajectories, forming a hyperfasciculated, inappropriately thick VF (Reeber et al., 2008). Essentially all of the spinal commissural axons transfected with Robo dominant-negative constructs, including those extended by d1 and d2 neurons, and both ascending and descending axons, exhibited this phenotype (Reeber et al., 2008). Similarly in the mouse spinal cord, the number of DiI-labeled ILc axons is reduced in the spinal cord of *Robo2* null and *Robo1*; *Robo2* double mutant mice (Jaworski et al., 2010). In addition, the expression of L1, which is a general marker for longitudinally projecting spinal axons (Imondi et al., 2000), in the spinal cord of mice lacking all three Slits, *Slits1–3*, revealed a reduction in the width of the LF (Long et al., 2004). Together, these observations support the view that Robo-Slit signaling has a major role in directing spinal commissural axons into the LF.

Robo-Slit signaling also appears to have a role in regulating the pathfinding of longitudinally projecting descending axons, which emanate from particular midbrain neurons. For example, analyses of various knockout mice have shown that Robo-Slit interactions normally prevent a variety of DiI-labeled descending axons from inappropriately crossing the midline in the brain (Bagri et al., 2002; Lopez-Bendito et al., 2007; Farmer et al., 2008). It is interesting to note in this regard that roles for Robos and Slits in barring longitudinally projecting spinal axons from the midline have not been clearly defined in the chick and mouse spinal cord.

For example, in chick embryos, spinal (commissural and ipsilateral) axons mis-expressing dominant-negative forms of Robo do not ectopically cross the ventral midline (Reeber et al., 2008). Similarly, the re-crossing of spinal commissural axons has not been observed in *Robo1*; *Robo2* double null mutant mice (Jaworski et al., 2010). In addition, whereas midline re-crossing events have been observed in mice lacking all three *Slits*, these apparently occur before the decussated axons turn into the longitudinal plane (Long et al., 2004). Therefore, the factors/mechanisms that prevent ascending spinal axons from invading the ventral midline have not been identified.

Consistent with additional roles for Robo-Slit signaling in the pathfinding of descending axons, the longitudinal tracts in the brain, which they normally assemble into, are disorganized or defasciculated in mice lacking Robos or Slits (Lopez-Bendito et al., 2007; Farmer et al., 2008). For example, in mouse embryos, Robo-Slit signaling is required for organizing ipsilateral descending axons extended by midbrain neurons within longitudinal fascicles, but not the dorsoventral positioning of these fascicles (Farmer et al., 2008). Although the ventral-most medial longitudinal fascicle invades the ventral midline in the midbrain and hindbrain of *Slit1*; *Slit2* and *Robo1*; *Robo2* double mutants, resulting in a ventral shift of this tract, the other two lateral tracts do not shift ventrally or fasciculate into one large bundle (Farmer et al., 2008). This contrasts with the ventral shift of the spinal commissural axon-containing LF observed in mice lacking Robos/Slits. Whereas both ascending spinal commissural axons and ipsilaterally descending midbrain axons appear to rely on Slits at the ventral midline, the roles of Robo-Slit signaling are clearly model system-specific. These differences might be due to distinct sources of Slits, different downstream signaling systems and/or the presence of non-canonical signaling molecules, which interact with Robo-Slit signaling system.

### PATHFINDING OF LONGITUDINAL AXONS IN VERTEBRATES: MIDBRAIN/FOREBRAIN DOPAMINERGIC TRACTS

In vertebrates, longitudinal axons emanating from spinal projection neurons pathfind through a rather homogenous spinal cord environment, which lacks conspicuous cellular specializations or choice points that could act as intermediate targets, before reaching the brain. In contrast, longitudinal axons originating from the neurons in higher brain areas encounter many distinct structures, which could act as guideposts and sources of key guidance cues, as they project to their appropriate synaptic targets. Therefore, the role of intermediate targets in regulating the pathfinding of longitudinal axons can often be more clearly delineated in the brain. One of the most prominent longitudinal tracts in the brain is the ascending mesodiencephalic dopaminergic (mdDA) pathway, which originates from dopaminergic neurons in the substantia nigra and ventral tegmental area in the midbrain and forebrain, and projects to the striatum as well as the cortex in the forebrain (Bjorklund and Dunnett, 2007).

### ESTABLISHING THE ROSTROCAUDAL POLARITY OF mdDA AXONS

Several independent studies have shown that canonical axon guidance molecules, in particular, Semaphorin-3F (Sema3F) regulates various aspects of mdDA axon pathfinding via its receptor,

Neuropilin-2 (Npn2), which is expressed on these axons (Kolk et al., 2009; Yamauchi et al., 2009). Initially, Sema3F, which is present at the midbrain-hindbrain boundary caudal to a sub-population of mdDA cell bodies, appears to direct ascending mdDA axons rostrally by operating as a repulsive guidance cue (Yamauchi et al., 2009). A subset of mdDA axons, labeled by an antibody against tyrosine hydroxylase (TH), inappropriately projects caudally in *Npn2* mutant mice, supporting the role of Sema3F-Npn2 in establishing the initial rostrocaudal polarity of this projection (Yamauchi et al., 2009). It is interesting to note in this regard that *Sema3F* expression at the midbrain-hindbrain boundary might be controlled by the morphogen, fibroblast growth factor 8 (Fgf8), since Fgf8-soaked beads can induce ectopic *Sema3F* mRNA expression and disrupt the rostral projection of mdDA axons, whereas reducing Fgf8 signaling using a Fgf receptor tyrosine kinase inhibitor eliminates *Sema3F* mRNA expression (Yamauchi et al., 2009).

Just as for ascending spinal commissural axons, Wnts and planar cell polarity signaling molecules appear to orient longitudinally projecting mdDA axons, as well as other monoaminergic neurons in the brain (i.e., serotonergic [5-HT] axons) (Fenstermaker et al., 2010). Here, in *Wnt5a* mutant, a small subset of ascending mdDA axons initially mis-projects caudally in E12.5 embryos although the phenotype is corrected by E17.5. In contrast, E17.5 mouse embryos lacking the planar cell polarity components, *Frizzled3*, *Vangl2*, or *Celsr3*, which are all expressed on ascending mdDA axons, exhibit more prominent and persistent aberrant caudal mis-projections (Fenstermaker et al., 2010). Hindbrain 5-HT axons also show aberrant rostrocaudal projections in these mutant mice (Fenstermaker et al., 2010). Whereas the rostrocaudal bias of mdDA and 5-HT axon projections appear to be established by *Frizzled3*, *Vangl2*, and *Celsr3*, the organization and/or orientation of these cell bodies are also disrupted in the corresponding mutants, thus, complicating phenotypic analyses. It will be important to examine the causal relations between the organization/orientation of mdDA cell bodies and axons, if these events are related.

### FASCICULATION OF mdDA AXONS

Robo-Slit signaling has a critical role in regulating fasciculation and rostral growth of ascending mdDA axons (Bagri et al., 2002; Dugan et al., 2011). *Slit1* and *Slit2* mRNA are expressed at the ventral midline and within the hypothalamus in the forebrain, as well as at the forebrain-midbrain border (Bagri et al., 2002; Dugan et al., 2011). Normally, TH-labeled mdDA axons, which express Robo1 and Robo2, project rostrally in close proximity to these *Slit1/Slit2*-expressing ventral structures (Dugan et al., 2011). However, in both *Slit1*; *Slit2* and *Robo1*; *Robo2* double mutants at E12.5, the initial projection of mdDA axons is defasciculated and disorganized (Dugan et al., 2011). Many mdDA axons appear to inappropriately project toward the ventral midline, which expresses high levels of *Slit1* and *Slit2*. One day later, at E13.5, although a subset of the defasciculated axons seems to recover and project in the appropriate direction within organized bundles, inappropriate invasion of the ventral midline is still evident and a part of the rostral track is hyperfasciculated, in both *Robo1* and *Slit* double mutants (Dugan et al., 2011). In addition, *Robo1*; *Robo2*

double mutants display dorsal mis-projections, which are not as prominent in *Slit1*; *Slit2* mutants. Thus, some roles of Robo1 and Robo2 in orienting mdDA axons into the rostral track appear to be independent of Slits (Dugan et al., 2011). Taken together, it appears that Robo-Slit signaling is required for the fasciculation and organization of ascending mdDA tracts, and prevents these axons from crossing the ventral midline in the forebrain.

Sema3F-Npn2 signaling also appears to control fasciculation of TH-positive mdDA longitudinal tracts as they project rostrally, since these axons are observed to defasciculate in *Sema3F* or *Npn2* mutant mice (Kolk et al., 2009). In addition, Npn2- or Sema3F-independent phenotypes have been reported in mice lacking *Sema3F* or *Npn2*, respectively. In *Sema3F* mutants, a subset of the defasciculated mdDA axons mis-projects ventrally toward the lateral hypothalamus, whereas mdDA axons in *Npn2* mutants follow an aberrant route to reach the prefrontal cortex (Kolk et al., 2009). Importantly, the targeting of these mdDA axons to the prefrontal cortex is disrupted in both *Sema3F* and *Npn2* mutant mice, but in disparate ways. At E18.5, the number of mdDA axons innervating the target area of the prefrontal cortex is significantly reduced in *Sema3F* mutants, whereas the number of these axons projecting to the same target area is increased in *Npn2* mutant mice (Kolk et al., 2009). Interestingly and likely reflecting an example of error correction, the majority of mdDA axons are reported to reach their appropriate target areas in adult *Sema3F* mutants (Kolk et al., 2009). Further studies focusing on elucidating the *in vivo* roles of Sema3F and Npn2 signaling, as well as considering the potential involvement of other Semas should help clarify these complicated phenotypes. Most importantly, understanding the underlying mechanisms that control the relevant pathfinding events should reveal the consequences of the targeting errors made by mdDA axons in mouse embryos lacking *Sema3F* or *Npn2*.

### PATHFINDING OF LONGITUDINAL AXONS IN VERTEBRATES: THE CORTICOSPINAL TRACT

One of the longest longitudinal axon-containing projections in the vertebrate CNS is the corticospinal tract, which connects the cerebral cortex to the spinal cord. Perhaps due to the relative ease of unilaterally labeling the component axons from the cerebral cortex, the complete trajectory followed by corticospinal tract axons has been carefully mapped. These axons are extended by pyramidal neurons located in layer V of the cerebral cortex, remain ipsilateral with respect to the midline (Stanfield, 1992; Brodal, 1998), and descend in the ventral region of the cerebral peduncle, through midbrain and hindbrain until they reach the caudal-most portion of the hindbrain, where most of these axons cross the midline dorsally to the contralateral side, forming the X-shaped pyramidal decussation. Corticospinal axons further project caudally within the marginal zone of the spinal cord, in a region containing the DF in rodents, and innervate neurons located within the spinal gray matter (Stanfield, 1992; Brodal, 1998).

### A ROLE FOR ROBO-SLIT SIGNALING IN REGULATING THE IPSILATERAL PROJECTION OF THE CORTICOSPINAL TRACT IN THE BRAIN

As described above, the corticospinal tract initially projects in a purely ipsilateral manner within the brain. Accordingly, the

mechanisms that maintain this laterality have a critical role in corticospinal tract formation. *Slit1* and *Slit2* mRNA are also expressed at the ventral midline of the forebrain, where these midline repellents create a barrier between the two sides of the rostral CNS (Bagri et al., 2002). Robo1 and Robo2 are expressed in particular populations of forebrain neurons/axons, including the major longitudinal projections that originate within the cortex, the corticospinal and corticofugal tracts (Lopez-Bendito et al., 2007), and Robo-Slit signaling prevents these tracts from crossing the midline in the forebrain (Bagri et al., 2002; Lopez-Bendito et al., 2007). Specifically, in *Robo1*; *Robo2* and *Slit1*; *Slit2* double mutants, the majority of the DiI-labeled corticospinal tract-associated axons aberrantly cross the midline at rostral levels before they would normally turn into the longitudinal plane (Bagri et al., 2002; Lopez-Bendito et al., 2007). Some of these misguided axons re-cross to the ipsilateral side of the brain, whereas others continue projecting into the contralateral side of the telencephalon (Lopez-Bendito et al., 2007). Thus, Robo-Slit signaling prevents corticospinal axons from inappropriately crossing the midline in the forebrain.

### DECUSSATION OF THE CORTICOSPINAL TRACT

The hindbrain pyramids represent the most conspicuous intermediate targets for longitudinally projecting corticospinal axons. Although several classes of molecules have been identified as candidates for regulating the decussation of corticospinal axons within this brain region, thus far, mainly contact-dependent, short-range guidance molecules have been directly implicated in this process.

Most notably, the immunoglobulin cell adhesion molecule (IgCAM), L1, is required for corticospinal axons to cross the ventral midline in the hindbrain (Cohen et al., 1997; Dahme et al., 1997). Although corticospinal axons appear to normally descend in the hindbrain of *L1* mutant mice, a subset of these axons fails to cross the midline in the lower hindbrain and inappropriately projects both ipsilaterally and contralaterally (Cohen et al., 1997). Some of mis-routed axons project dorsally to the ipsilateral DF or remain ventrally positioned but turn into the longitudinal plane and project caudally in the ventrolateral edge of the ipsilateral side of hindbrain (Cohen et al., 1997). Moreover, the number of caudally projecting axons in the spinal cord is severely reduced in adult mutants. Since the number of normally decussated corticospinal commissural axons is also reduced in the spinal cord of *L1* mutant mice, L1 might be required for the growth of these axons within the spinal cord. Although L1 is expressed on developing pyramidal fibers (Cohen et al., 1997), how L1 regulates the midline crossing of corticospinal axons remains to be determined. In this regard, it has been suggested that Sema3A secreted by the ventral spinal cord is capable of repelling cortical axons expressing L1 *in vitro*, and Neuropilin-1 (Npn1) and L1 form a receptor complex that likely transduces this Sema3A-mediated repulsive signal (Castellani et al., 2000). However, this is unlikely to be the mechanism underlying the formation of the pyramidal decussation, since corticospinal tracts form normally and on schedule in *Sema3A* mutant mice (Sibbe et al., 2007).

In addition to L1, another IgCAM, neural cell adhesion molecule (NCAM), has also been implicated in controlling the

pyramidal decussation of corticospinal axons (Rolf et al., 2002). The corticospinal axon pathfinding defects observed in *NCAM* mutants resemble the *L1* mutant phenotype in that they reveal the lack of decussation in the hindbrain and bilateral caudal projections in the spinal cord of early postnatal mice. Unlike *L1* mutant mice, however, only a small number of mis-routed axons are present on the ipsilateral side of the pyramid in *NCAM* adult mutants (Rolf et al., 2002). Whether these mis-projections were corrected or eliminated at later ages has not been investigated. Since corticospinal axons arrive later at the caudal hindbrain in *NCAM* mutant mice as compared to their wild type counterparts, the defects in corticospinal tracts appear to manifest themselves prior to the formation of the pyramidal decussation (Rolf et al., 2002).

The transmembrane *Sema6A* and its receptors *PlexinA3* and *PlexinA4* have also been implicated in regulating the decussation of corticospinal axons (Faulkner et al., 2008; Runker et al., 2008). In *Sema6A* single, *PlexinA4* single and *PlexinA3*; *PlexinA4* double mutant mice, corticospinal axons normally descend to the caudal hindbrain. Subsequently, however, some of these axons correctly decussate, but many others inappropriately projected to the ventrolateral edge of the ipsilateral hindbrain (Faulkner et al., 2008; Runker et al., 2008). Notably, however, the mis-guided axons continue projecting caudally (Faulkner et al., 2008; Runker et al., 2008). *PlexinA3* and *PlexinA4* mRNA is expressed by cortical neurons, and *Sema6A* and *PlexinA4* mRNA are found along the terrain through which corticospinal axons project in the caudal hindbrain and in the inferior olivary nuclei, which is located adjacent and lateral to the corticospinal tract in the pyramid (Faulkner et al., 2008; Runker et al., 2008). Collectively the loss of function phenotypes exhibited by the various *Sema* and *Plexin* mutant mice suggest that *Sema6A*-*PlexinA3/4* signaling-mediated repulsion normally drives corticospinal axons toward the midline where they undergo decussation (Faulkner et al., 2008; Runker et al., 2008).

A reduction in the size of the pyramidal decussation has been also reported in *Netrin1* receptor *Unc5c* (previously known as *Unc5h3*) mutant mice (Finger et al., 2002). The phenotypes observed in *Unc5c* mutants most closely resemble those displayed by *Sema6A* and *PlexinA3*; *PlexinA4* mouse mutants: the majority of corticospinal axons fail to cross the midline, and the misdirected axons continue projecting caudally in the lateral funiculus of the spinal cord. Moreover, these mis-routed corticospinal axons appear to fasciculate with each other in the lateral funiculus (Finger et al., 2002). Therefore, unlike in *L1* and *NCAM* mutants, the caudal projection of mis-routed corticospinal axons in *Sema6A*, *PlexinA3*; *A4*, and *Unc5c* mutants is intact and possibly regulated by other molecules found in both the DF and lateral funiculus. Since CAMs are known to promote axon outgrowth (Raper and Mason, 2010), *L1* and *NCAM* might, at least to some extent, facilitate the caudally directed growth of the corticospinal axons within the spinal cord. Alternatively, the differences between the phenotypes exhibited by the various mutant mice might simply reflect the different techniques and experimental approaches used to visualize corticospinal axons/tracts.

## CAUDAL GROWTH OF CORTICOSPINAL AXONS IN THE SPINAL CORD

After crossing the midline within the caudal hindbrain, rodent corticospinal axons project caudally in the DF of the spinal cord (Stanfield, 1992). Consistent with *Wnt4* being required for the polarity of ascending spinal commissural axons (Lyuksyutova et al., 2003), *Wnt1* and *Wnt5a* have been implicated as regulators of the caudal growth displayed by corticospinal axons within the spinal cord (Liu et al., 2005). Support for this model comes from the finding that *Wnt1* and *Wnt5a* mRNA is expressed in the dorsal spinal gray matter surrounding the DF in rostral (high) to caudal (low) gradients. In addition, the Wnt receptor, Receptor tyrosine kinase-related tyrosine kinase (Ryk)-like transmembrane receptor is expressed at high levels on corticospinal axons, and *Wnt1*/*Wnt5a* repel corticospinal axons expressing Ryk *in vitro* (Liu et al., 2005). Moreover, injection of a functional blocking anti-Ryk antibody into the cervical level spinal cord of neonatal mice retards the growth of descending corticospinal axons (Liu et al., 2005). With regard to these particular findings, it is important to note that the DF also contains the fasciculus gracilis and fasciculus cuneatus, which ascend from dorsal root ganglia to the lower hindbrain (Burt, 1993; Brodal, 1998). Thus, it would be interesting to determine whether *Wnt1*/*Wnt5a* facilitate the rostral growth of these dorsal-column-associated ascending axons, possibly through resident Wnt receptors that mediate attraction via Wnt gradients in the spinal cord. In addition, Ryk-mediated signaling through Wnts might be responsible for the caudal mis-projection of the corticospinal axons in mutant mice with pyramidal decussation defects, such as the *Sema6A* and *Unc5c* null mutants (Finger et al., 2002; Faulkner et al., 2008; Runker et al., 2008), mentioned above. Since *Wnt1*/*Wnt5a* mRNA expression is not present in the vicinity of the lateral funiculus (Liu et al., 2005), other Wnts found in this region could be presented to Ryk-bearing axons. In any case, assessing the role of Wnts *in vivo* might help to further understand the mechanisms that control the rostrocaudal polarity of corticospinal axons.

## PREVENTING CORTICOSPINAL AXONS FROM RE-CROSSING THE MIDLINE IN THE SPINAL CORD

As reflected by the somatotopic organization of corticospinal neurons in the sensorimotor cortex, these descending axons innervate neurons at specific levels of the spinal cord (Stanfield, 1992; Kuang and Kalil, 1994). Upon reaching the appropriate rostrocaudal segments of the spinal cord, corticospinal axons project into the spinal gray matter, where they form connections with their specific synaptic targets (Stanfield, 1992; Brodal, 1998). Since corticospinal axons transmit signals from one side of the cerebral cortex to neurons located on the opposite side of the spinal cord, once they cross, these descending longitudinal axons must terminate on the contralateral side of the CNS and never re-cross the midline. Within the spinal cord gray matter, repulsive interactions between the EphA4 receptor tyrosine kinase and the transmembrane ephrinB3 ligands facilitate this pattern of innervation by preventing decussated corticospinal axons from re-crossing back to the ipsilateral side (Dottori et al., 1998; Kullander et al., 2001a; Yokoyama et al., 2001). EphrinB3 is selectively expressed at the ventral midline of the spinal cord and this short-range repulsive



Eph ligand forms a boundary separating axons on either side of the spinal cord (Kullander et al., 2001a; Yokoyama et al., 2001). Corticospinal axons express one of the ephrinB3 receptors, EphA4, and thereby regulate ephrinB3-mediated repulsive signaling cell autonomously (Kullander et al., 2001b). In both *ephrinB3* and *EphA4* mutant mice, corticospinal axons normally cross at the caudal hindbrain and project caudally in the contralateral DF (Kullander et al., 2001a,b; Yokoyama et al., 2001). However, after these axons enter the spinal cord gray matter, some of them re-cross the midline and innervate targets on the ipsilateral side of the spinal cord (Kullander et al., 2001a,b; Yokoyama et al., 2001). Since the majority of re-crossing corticospinal axons enter the ipsilateral side of the CNS via the spinal cord gray matter rather than through the DF, in *ephrinB3* and *EphA4* mutant mice (Kullander et al., 2001a,b), the laterality of descending longitudinal projections in the dorsal funiculi appears to be regulated by some other yet to be described mechanism. It is important to note that the midline barrier formed by ephrinB3 is critical for preventing the aberrant crossing of not only corticospinal axons, but also other EphA4-expressing axons in the spinal cord, such as those extended by ventral spinal interneurons (Kullander et al., 2003).

## PATHFINDING OF LONGITUDINAL AXONS IN THE INVERTEBRATE CNS

### THE ROLE OF Wnt-FRIZZLED SIGNALING IN *C. elegans*

In a striking example of evolutionary conservation, morphogens also appear to control the guidance of longitudinally projecting axons along the A-P axis in invertebrates. Consistent with the likely role of Wnts in regulating the polarity of post-crossing spinal commissural axons in the embryonic mouse spinal cord (see above), Wnt-Frizzled signaling has a key role in regulating rostrocaudal pathfinding decisions made by longitudinally projecting axons in *C. elegans*. Specifically, Wnt (*lin-44*) via the Frizzled (*lin-17*) receptor regulates the A-P orientation of PLM mechanosensory axons and dendrites (Hilliard and Bargmann, 2006). PLM mechanosensory neurons are bipolar and Frizzled (*lin-17*) is highly expressed on posteriorly growing dendrites. This asymmetric expression of Frizzled (*lin-17*) receptors is induced by Wnt (*lin-44*) and thus, Wnt (*lin-44*) and Frizzled (*lin-17*) signaling regulate the A-P orientation of PLM neurons neither by repelling nor attracting these axons or dendrites. In addition to establishing the A-P orientation of axons, longitudinal projections of mechanosensory axons are guided by Wnt-Frizzled signaling in *C. elegans* (Pan et al., 2006). However, opposite to what has been observed in the mouse spinal cord, caudally expressed Wnt (*egl-20*) operates as a repellent via Frizzled (*mig-1* and *mom-5*) signaling to direct longitudinally projecting axons rostrally in *C. elegans* (Pan et al., 2006). Thus, although the role of Wnt-Frizzled signaling in regulating A-P polarity is conserved between mammals and worms, the mechanistic details are distinct across species.

### THE ROLES OF ROBO RECEPTORS IN POSITIONING LONGITUDINAL TRACTS IN THE *DROSOPHILA* VENTRAL NERVE CORD

Further consistent with evolutionarily conserved roles for guidance systems that control longitudinal axon pathfinding, the

involvement of Robos in regulating the lateral positioning of longitudinally projecting axons tracts was first demonstrated in the *Drosophila* ventral nerve cord (Rajagopalan et al., 2000; Simpson et al., 2000). The expression patterns of the three *Drosophila* Robo receptors, Robo1–3, on longitudinal axons define distinct medial-lateral zones in the neuropile of the ventral nerve cord; Robo1 is expressed on axons in the medial zone, the intermediate zone contains axons that express Robo1 and Robo3, and axons that express all three Robos are present in the lateral zone, and disrupting these collective expression patterns predictably perturbs the lateral positioning of longitudinal axons (Rajagopalan et al., 2000; Simpson et al., 2000). In addition to the pronounced midline re-crossing phenotype displayed by *Robo1*, and occasionally *Robo2*, mutants, the loss of function phenotype of each *Robo* mutant is similar to the consequences of disabling Robo-Slit signaling in the vertebrate spinal cord; a major medial shift in the positioning of many longitudinal axons (see above). In particular, anti-Fasciclin II [FasII, marker of three major longitudinal tracts in the *Drosophila* CNS (Grenningloh et al., 1991)] labeling of intermediate fascicle reveals that the loss of *Robo3* function results in an inappropriate fusion of these axons with the medial fascicle (Rajagopalan et al., 2000; Simpson et al., 2000). In addition, the knockdown of *Robo3* in a *Robo2* mutant background gives rise to one large medial axon bundle representing a fusion of all three FasII-positive fascicles, whereas *Robo1* and *Robo2* double mutants display a midline collapse phenotype, similar to that observed in *Slit* mutants (Kidd et al., 1999; Simpson et al., 2000). These observations strongly suggest that Robo2 and Robo3 have critical roles in the proper lateral positioning of longitudinal tracts, and that the primary functions of Robo1 and Robo2 are to bar these axons from re-entering the midline. Accordingly, the role of *Drosophila* Robo3 appears to be analogous to that of vertebrate Robo1 and Robo2, whereas vertebrate Robo3 (Rig-1) operates in a manner similar to *Drosophila* Commissureless, by facilitating the passage of commissural axons across the ventral midline (Seeger et al., 1993; Keleman et al., 2002; Sabatier et al., 2004; Chen et al., 2008).

Regarding the mechanism through which Robos regulate the lateral positioning of longitudinal axons, it has recently been established that structural differences between Robo receptors do not have a crucial role in this process (Spitzweck et al., 2010). Rather, the gene expression profile of each Robo controls longitudinal axon sorting. This was elegantly demonstrated by replacing the genomic loci of each *Robo* with full length *Robo* sequences through homologous recombination and assessing the effects of these manipulations on longitudinal axon pathfinding (Spitzweck et al., 2010). Together, the findings strongly suggest that the timing, location, and/or level of each Robo gene are essential for achieving proper sorting of longitudinal axons in *Drosophila* ventral nerve cord.

Thus far, the role of the Robo ligand Slit, which operates as a potent repulsive guidance cue for a variety of invertebrate and vertebrate axons (Mastick et al., 2010; Ypsilanti et al., 2010), in the lateral positioning of longitudinal tracts has not been clearly established within the *Drosophila* ventral nerve cord. Although multiple populations of axons collapse on the midline in *Slit* mutants (Battye et al., 1999; Kidd et al., 1999), and the overexpression of *Slit* in midline glia cells disrupts the pathfinding of

commissural axons (Battye et al., 1999), the role(s) of Slit in regulating the guidance of post-crossing commissural axons has not been explicitly addressed (Dickson and Gilestro, 2006). In contrast, and as discussed above, Slits appear to regulate the sorting of longitudinal axons in the embryonic mouse spinal cord.

### THE ROLE OF AXON FASCICULATION IN THE FORMATION OF LONGITUDINAL TRACTS WITHIN THE INVERTEBRATE CNS

The dynamic roles of axon fasciculation and defasciculation events on the pathfinding of longitudinally projecting axons were originally demonstrated in grasshopper embryos (Raper et al., 1983, 1984; Harrelson and Goodman, 1988; Raper and Mason, 2010). The observation that precise fasciculation patterns exist among populations of longitudinally projecting axons led to the identification of molecules that are selectively expressed on particular longitudinal tracts in *Drosophila*. One major class of fasciculation molecules is the IgCAM superfamily members, of which FasII is an example (Grenningloh et al., 1991). In the ventral nerve cord of *Drosophila FasII* mutants, intermediate and lateral fascicles are conspicuously defasciculated, and forced expression of *FasII* leads to partially fused longitudinal fascicles (Lin et al., 1994). These results support a role for FasII in mediating the fasciculation of longitudinally projecting axons in *Drosophila*. Notably, however, defasciculated or hyperfasciculated axons in genetically manipulated *FasII* embryos retain their ability to appropriately project along the A-P axis adjacent to the midline (Lin et al., 1994). Thus, it appears that FasII does not explicitly control the directed growth/pathfinding of longitudinal axons along the A-P axis of the *Drosophila* ventral nerve cord.

The classical cadherins represent another major class of CAMs, and N-cadherin, in particular, has been implicated in the fasciculation of FasII-positive axons and their assembly into longitudinal axon tracts (Iwai et al., 1997). N-cadherin is expressed broadly in the *Drosophila* CNS, with particularly high levels present on axons, and *N-cadherin* mutants display defects in the bundling and pathfinding of FasII-positive intermediate and lateral fascicles (Iwai et al., 1997). Reminiscent of the FasII mutant phenotype, the majority of improperly fasciculated axons appropriately project along the A-P axis in *Drosophila N-cadherin* mutants. Interestingly, the observed defects appear quite selective and specific given that N-cadherin is broadly and robustly expressed on axons within the *Drosophila* ventral nerve cord. To further dissect the role of N-cadherin in longitudinal tract formation, the pathfinding behavior of a genetically distinct population of neurons, defined by the expression of LIM homeodomain transcription factor, *Apterous*, was examined in N-cadherin null mutants. Axons extending from *Apterous*-positive neurons normally fasciculate tightly with each other and project ipsilaterally in the anterior direction along the most medial edge of the *Drosophila* ventral nerve cord (Lundgren et al., 1995). However, in *N-cadherin* mutants, these axons fail to fasciculate, and the dorsal-most axons project slightly lateral to their normal medial tract. Despite these defects *Apterous* axons continue to project rostrally in *N-cadherin* mutants (Iwai et al., 1997), indicating that like FasII, N-cadherin does not regulate this pathfinding decision. Notably, the perturbations in the pathfinding of N-cadherin-lacking *Apterous* axons resemble the defective

fasciculation of these axons observed in *Apterous* mutants (Lundgren et al., 1995), raising the possibility that *Apterous* regulates N-cadherin expression/function. Overall, these studies suggest that although N-cadherin might not be required for the directed growth of axons along the A-P axis, it does ensure the fidelity of longitudinal axon tract formation in *Drosophila*.

In addition to the CAMs, Semas, which are cell surface or secreted molecules, and their receptors, the Plexins have been implicated in the formation of longitudinal axon tracts within the *Drosophila* CNS. For example, transmembrane *Sema1A* and its receptor *PlexinA* regulate the defasciculation of FasII-labeled laterally positioned longitudinal axons from their neighboring intermediate fascicles (Winberg et al., 1998; Yu et al., 1998). Specifically, the lateral fascicle in *Sema1A* or *PlexinA* mutants is disrupted and fused with the intermediate fascicle, and *PlexinA* regulates this process via repulsive signaling. These mutant phenotypes likely reflect defects in defasciculation, rather than fasciculation, since the lateral fascicle-associated axons apparently fail to break away from the intermediate fascicle (Winberg et al., 1998). Whereas transmembrane *Sema1A*-*PlexinA* signaling selectively controls the assembly of laterally positioned FasII-positive axons into longitudinal tracts, secreted *Sema2A* and *Sema2B* appear to regulate the bundling of axons in the FasII-labeled intermediate fascicle, through the actions of *PlexinB* receptors (Wu et al., 2011). In a key recent study, it has been shown that in *Sema2A; Sema2B* double and *PlexinB* mutants, FasII-positive axons in intermediate fascicles defasciculate from each other and these tracts are severely disorganized (Wu et al., 2011). Here, *Sema2A* likely creates a repulsive boundary that constrains the positioning and growth of the intermediate fascicle. On the other hand, *Sema2B* facilitates the fasciculation of these axons, by acting as a short-range attractive cue (Wu et al., 2011). *PlexinB* is expressed at high levels in the region of the neuropile that contains intermediate and lateral tracts, and is also likely present on the first set of pioneer axons, which form the FasII-positive intermediate fascicle (Wu et al., 2011). Therefore, it appears that *PlexinB* controls the formation of the intermediate longitudinal fascicle by transducing *Sema2A*-dependent repulsive signals and *Sema2B*-mediated attraction (Wu et al., 2011).

In the *Drosophila* CNS, longitudinal axon tracts receive direct innervation from sensory axons (Boyan and Ball, 1993). Interestingly, *PlexinB* signaling activated by *Sema2A* and *Sema2B* also regulates the targeting of genetically distinct mechanosensory axons to the FasII-positive intermediate fascicle (Wu et al., 2011). Specifically, in *Sema2A; Sema2B* double or *PlexinB* mutant lines, particular mechanosensory axons fail to innervate axons located in the FasII-labeled intermediate fascicle. Since *PlexinB* appears to regulate the targeting of these mechanosensory axons in a cell autonomous manner, the failure of this sensory axon innervation is not the result of the defasciculation of the intermediate fascicle-associated axons, which has been observed in these mutant lines as mentioned above (Wu et al., 2011). This finding raises the possibility that targeting of other longitudinal fascicles to appropriate sensory axons is also coordinately regulated by Semas and their *Plexin* receptors.

Although these studies clearly show that the fasciculation of longitudinal tracts is tightly regulated by specific sets

and combinations of guidance molecules the consequences of the defective fasciculation events remain poorly understood. Ultimately, disrupted patterns of fasciculation might be expected to promote mis-targeting or defects in the pre-synaptic termination patterns on a given fascicle.

## TARGETING OF LONGITUDINAL AXONS IN THE VERTEBRATE CNS

In contrast to the progress made in identifying molecules and mechanisms that regulate the pathfinding of longitudinal axons in the vertebrate CNS, considerably less is known about what controls the targeting of these axons. Although it is generally assumed that mis-directed axons, which pathfind along aberrant trajectories, will ultimately fail to reach their appropriate targets, this may not always be the case (see examples above). Accordingly, identifying the synaptic targets of longitudinally projecting axons and elucidating the molecular mechanisms that control the targeting of these axons should clarify key aspects of neural circuit formation in wild type and mutant phenotypes. Part of the lack of progress in elucidating mechanisms that control axon targeting can most likely be attributed to the technical limitations associated with tracing axons over long distances. For example, since many axonal tracers have slow diffusion rates in fixed tissue (Vercelli et al., 2000), these reagents would have to be delivered *in vivo* or into appropriate *in vitro* preparations to label the full extent of longitudinal axon tracts, and these are not always feasible options. In addition, pan-axonal tracers, such as DiI, are incapable of reliably and reproducibly labeling specific populations of axons and, thus it has not been possible to visualize particular sets of axons over long distances and assess the consequences of molecular perturbations on their pathfinding and targeting. Analogous to approaches that are routinely employed in invertebrate systems, recently developed genetically labeling strategies have now made it possible to visualize, and study the targeting of, long-range longitudinal projections within the vertebrate CNS.

### ASCENDING LONGITUDINALLY PROJECTING SPINAL AXONS

Along with the technical limitations described above, the bilateral symmetry of the spinal cord and presence of both ipsilateral and commissural axons within the major longitudinal axon tracts of the marginal zone also confound attempts to visualize these long-range projections. The ascending spinal longitudinal tracts, in particular, are largely bilaterally symmetric with both sides of the LF and VE, but not the DF, containing contralaterally and ipsilaterally projecting axons (Kerr, 1975; Matsushita and Hosoya, 1979; Giesler et al., 1984; Burstein et al., 1990; Katter et al., 1991; Yeziarski and Mendez, 1991). In addition, some of the component axons, such as those that compose the spinocerebellar tract, re-cross the midline in the cerebellum (Matsushita and Hosoya, 1979; Matsushita and Ikeda, 1980; Brodal, 1998). Accordingly, unilateral labeling carried out in intact animals is required to obtain unobstructed views of ascending longitudinal axons, and to assess the consequences of molecular/genetic manipulations on their pathfinding and targeting.

Unilateral labeling of axons *in vivo* can be achieved via the use of, *in ovo* and *in utero* unilateral electroporation strategies, in

chick and mouse embryos, respectively, (Krull, 2004; Saito, 2006; Petros et al., 2009). In fact, *in ovo* electroporation of a pan-axonal reporter, has been used to show that both commissural and ipsilateral axons originating from a specific segment of the spinal cord (lumbosacral level 2) project rostrally to a common target, the cerebellum, in the E8–E13 chick brain (Arakawa et al., 2008).

To investigate the mechanisms that control the pathfinding and targeting of *particular classes* of spinal projection neurons, it is critical to first develop labeling methods that can be used to visualize genetically distinct populations of these neurons/axons. Dorsal spinal neurons represent a well-studied and major class of projection neurons (Nunes and Sotelo, 1985; Burstein et al., 1990; Yeziarski and Mendez, 1991; Brodal, 1998), and subtypes of these neurons have been defined on the basis of transcription factor expression profiles (Helms and Johnson, 2003). For example, d1 dorsal spinal neurons are derived from progenitors that express the basic helix-loop-helix (bHLH) transcription factor *Atoh1* (formerly *Math1*), and express the homeodomain (HD) transcription factors, *Lhx2*, *Lhx9*, *BarH1*, and *Brn3a* (Helms and Johnson, 2003). On the other hand, *Neurog1*- (bHLH family member) positive progenitors give rise to d2 neurons, which are defined by the expression of *Lhx1*, *Lhx5*, *FoxD3*, and *Brn3a* (Helms and Johnson, 2003). Based on these observations, a variety of transgenic reporter mice were generated to facilitate the mechanisms that control the development of d1 and d2 dorsal spinal neurons (Helms and Johnson, 1998; Nakada et al., 2004; Machold and Fishell, 2005; Saba et al., 2005; Wilson et al., 2008). However, the bilateral symmetry of the reporter gene expression patterns and the presence of endogenous labeling in the brain, has hampered the definitive identification of d1 and d2 targets (Helms and Johnson, 1998; Bermingham et al., 2001; Nakada et al., 2004; Saba et al., 2005). Nevertheless, DiI labeling in an *Atoh1-lacZ* knock-in (null for *Atoh1*) mouse line was carried out in an attempt to identify the brain targets of d1 neurons/axons (Bermingham et al., 2001). This study reported that the size of the DiI-labeled spinocerebellar tract within the hindbrain was reduced in *Atoh1<sup>lacZ/lacZ</sup>* mice compared to wild type and heterozygous littermates, raising the possibility that d1 axons target the cerebellum (Bermingham et al., 2001). More recently we have delivered *Atoh1* and *Neurog1* reporter constructs (Lumpkin et al., 2003; Nakada et al., 2004; Reeber et al., 2008) into the embryonic chick spinal cord via unilateral *in ovo* electroporation and achieved reproducible labeling of longitudinally projecting d1 and d2 axons. Moreover, we have found that both of these projection neurons target a variety of brain regions, including the cerebellum (N.S. and Z.K., unpublished observations). These observations support the feasibility of using genetic labeling strategies to visualize the long-range projections of genetically distinct spinal projection neurons and to carry out detailed analyses of spinal axon targeting in an *in vivo* setting.

### DESCENDING CORTICOSPINAL AXONS

Anatomical considerations suggest that the targeting of descending corticospinal axons is somatotopically regulated, since collaterals of the primary corticospinal axons exit the DF and innervate the spinal cord gray matter at specific positions along the rostro-caudal axis (O'Leary and Koester, 1993; Kuang and Kalil, 1994).

The PSA modification of NCAM (PSA-NCAM) appears to facilitate collateral formation by modifying the fasciculation state of corticospinal axons within the DF (Daston et al., 1996). Whereas NCAM is expressed all along pathfinding corticospinal axons, PSA-NCAM is selectively present in the DF, where collateral formation occurs. Removal of PSA through enzymatic degradation reduces the number of collaterals at particular levels of the spinal cord (Daston et al., 1996). Despite these observations, exactly how the formation of axon collaterals is somatotopically regulated at specific segments of the spinal cord remains poorly defined.

Although corticospinal tracts are composed of ipsilateral and contralateral projections located rostral and caudal to the pyramidal decussation, respectively, the component axons retain their unilaterality on either side of the spinal cord (Stanfield, 1992). Accordingly, unilateral labeling strategies, like those we have used to visualize ascending spinal projection neuron tracts (see above), should not necessarily be required to elucidate the mechanisms that control corticospinal axon targeting in the spinal cord. Rather, transgenic reporter mice that can be used to visualize corticospinal tract axons (Bareyre et al., 2005) should be sufficient. In this regard, the recent use of corticospinal axon-specific reporter mice has led to the identification of Clark's column neurons, which contribute to the spinocerebellar projection, as targets of corticospinal axons (Hantman and Jessell, 2010). This new and important observation will likely facilitate a detailed investigation into the molecular mechanisms that control the guidance and targeting of corticospinal axons.

## SUMMARY AND PERSPECTIVE

Just as longitudinally projecting axons must travel long distances in order to reach their synaptic targets and to transmit information between disparate regions of the CNS and from the environment to the brain, we have a long way to go before acquiring even a superficial understanding of the underlying

guidance and targeting mechanisms. Despite the recent progress made in elucidating mechanisms that control particular phases of longitudinal axon growth/targeting in specific model vertebrate and invertebrate systems, many fundamental "big picture" questions remain. For example: (1) Do morphogens have central roles in longitudinal axon guidance and/or targeting or are they only required to set the polarity of axons as they turn into the longitudinal plane? (2) Do the mechanisms that control longitudinal axon guidance also control the targeting of these axons? (3) Do short-range axon-axon interactions on their own facilitate the formation of precisely positioned longitudinal axon tracts or is the directed and stereotypical growth of longitudinal tracts controlled by a complex interplay between long-range and short-range/adhesive guidance systems? (4) Does the same molecular logic control the guidance and targeting of ipsilateral and commissural axons and/or the formation of ascending and descending axonal tracts? (. . . as discussed above the answer seems to be a resounding no) (5) Are the mechanisms that control the long-range pathfinding and targeting of longitudinally projecting axons conserved across species and organisms? As we have suggested above, obtaining the answers to these and other burning questions in the field will require that state-of-the-art genetic labeling strategies initially be used, in mouse and chick embryos as well as in *Drosophila* and *C. elegans*, to carefully map the projection patterns of longitudinal axons and to identify their synaptic targets. Once this has been achieved, a combination of complementary gain- and loss-of-function manipulations in the various model systems should, ultimately, make it possible to elucidate the molecular logic that controls longitudinal axon guidance and targeting. Importantly, these likely to be tedious, time-consuming and ground breaking studies will surely provide exciting new insights into neural circuit formation across species and the molecular genetic underpinnings of neurological dysfunction.

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