

Mechanisms of Axon Guidance in the Developing Nervous System

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The human brain assembles an incredible network of over a billion neurons. Understanding how these connections form during development in order for the brain to function properly is a fundamental question in biology. Much of this wiring takes place during embryonic development. Neurons are generated in the ventricular zone, migrate out, and begin to differentiate. However, neurons are often born in locations some distance from the target cells with which they will ultimately form connections. To form connections, neurons project long axons tipped with a specialized sensing device called a growth cone. The growing axons interact directly with molecules within the environment through which they grow. In order to find their targets, axonal growth cones use guidance molecules that can either attract or repel them. Understanding what these guidance cues are, where they are expressed, and how the growth cone is able to transduce their signal in a directionally specific manner is essential to understanding how the functional brain is constructed. In this chapter, we review what is known about the mechanisms involved in axonal guidance. We discuss how the growth cone is able to sense and respond to its environment and how it is guided by pioneering cells and axons. As examples, we discuss current models for the development of the spinal cord, the cerebral cortex, and the visual and olfactory systems. © 2005, Elsevier Inc.

I. Introduction

The establishment of correct neuronal connections is crucial for proper functioning of the nervous system. In the human brain, there are more than a billion neurons, of many different types, that assemble into highly complex neuronal networks with over 10^9 connections. To understand the brain it is essential to understand how this network of neuronal connections is achieved with such high precision as the nervous system develops. Santiago Ramón y Cajal studied the developing mammalian brain and proposed that each axon is attracted to its target cells by diffusible molecules secreted from it. Direct evidence supporting this guidance mechanism has only started to emerge during the past decade, with an increasing number of families of guidance molecules being identified. It is well established that developing axons are guided to their targets by a variety of environmental cues, including long-range diffusible and short-range surface-bound molecules that can either attract or repel the axon. The presence of these guidance cues in a temporal and spatial pattern enables the axon to navigate through the complex environment of the developing embryo to reach its correct target and to stop growing once it arrives. This chapter describes our current understanding of the mechanisms involved in the formation of neuronal connections.

II. The Growth Cone

In order to connect with targets, neurons extend processes called axons. At the tip of the extending axon is a specialized sensing device called a growth cone. In 1880, Santiago Ramón y Cajal observed and named the growth cone (“*cono de crecimiento*”) as the motile structure at the leading edge of extending axons. During development of the nervous system, growth cones navigate along specific pathways, recognize their targets, and then form synaptic connections by elaborating terminal arbors. En route to their target, growth cones show stereotyped behavior by following appropriate pathways and by interacting with intermediate targets. These characteristic behaviors of axons are regulated by a combination of various types of guidance cues, including attractive and repulsive cues. How growth cones respond to guidance cues is not fixed and changes dynamically during navigation. Growth cones derive their directional information from a variety of extracellular guidance cues. These cues can take the form of both secreted and membrane-bound attractive and repulsive factors. To date, a large number of these guidance cues and their receptors have been identified (Mueller, 1999; Tessier-Lavigne and Goodman, 1996). There is also an extensive collection of actin-associated proteins that regulate the structure and dynamics of the actin cytoskeleton that makes up the motile apparatus of the growth cone (Stossel, 1993).

The growth cone has two major domains: the central domain (C domain), rich in microtubules and membranous organelles, and the peripheral domain, rich in actin filaments (Fig. 1). The peripheral domain consists of lamellipodia surrounding a central domain and filopodia extending from the outer edge of the peripheral domain. Filopodial and lamellipodial projections are sent out by the growth cone to sense its environment; they contain dense and highly polymerized actin microfilaments. Both lamellipodia and filopodia are involved in growth cone motility and undergo continuous cycles of expansion and contraction (Bray and Chapman, 1985; Goldberg and Burmeister, 1986). The cytoskeleton of the growth cone is a combination of dynamic cytoplasmic filamentous structures, including microtubules and microfilaments (Forscher and Smith, 1988; Letourneau *et al.*, 1986; Luduena and Wessells, 1973; Yamada *et al.*, 1970, 1971).

The microtubules are involved in axon extension; they extend from the neurite shaft and splay into the central part of the growth cone. Microtubules are polar structures with one end termed the “plus” end and the other end termed the “minus” end. They are composed of heterodimers of α - and β -tubulin that self-assemble to form polymers. Microtubules form dense arrays within the neurite, while they are less stable within the growth cone (Ahmad *et al.*, 1993). As microtubules enter the growth cone from the neurite shaft, the bundled microtubules separate from each other and extend

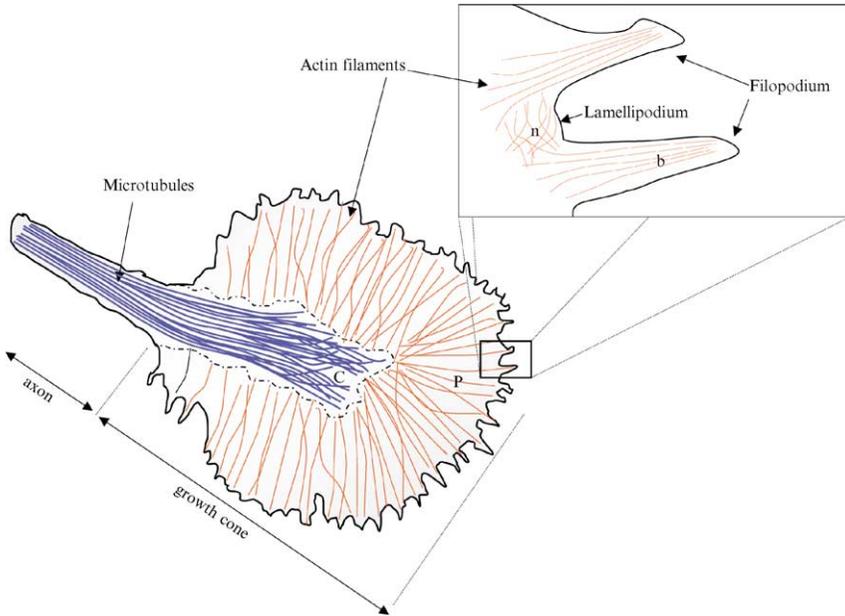


Figure 1 Schematic of the growth cone. The growing tip of the axon consists of a central domain C and a peripheral domain P with filopodium and lamellopodium. The C domain contains a dense network of microtubules, but some microtubules can also be found in the base of filopodia. Actin filaments are predominant in the P domain. Filopodium move like fingers exploring their environment. These movements are based on actin polymerization and depolymerization. Filopodial actin is organized into bundles (b) whereas actin filaments in lamellopodia and in the C region form an intricate network (n).

into the organelle-rich central domain of the growth cone in the form of individual microtubules (Tanaka and Kirschner, 1991). Growth cones also contain a large pool of soluble tubulin capable of assembling into microtubules (Letourneau and Ressler, 1984). The majority of growth cone microtubules are restricted to the C domain, but individual microtubules can also extend deep into the actin-rich peripheral domain (Gordon-Weeks, 1991; Tanaka and Kirschner, 1991).

The actin cytoskeleton is composed of two distinct microfilament subpopulations: one forms a meshwork of relatively short randomly oriented filaments in the lamellopodia, and the second is composed of parallel bundles of up to a dozen filaments that radiate from the leading edge of the lamellopodia and extend into the filopodium (Forscher and Smith, 1988; Letourneau and Ressler, 1984; Lewis and Bridgman, 1992; Luduena and Wessells, 1973; Yamada *et al.*, 1970, 1971). The actin contributes to the structure of the growth cone as well as the motility. It exits in both an

unassembled (often referred to as globular actin [G-actin] or filamentous [F-actin] form). The F-actin is polymerized from the monomeric G-actin subunits. Actin polymerization in growth cones occurs predominantly at the leading edge and is vital to growth cone motility and hence guidance.

A. Cytoskeletal Rearrangements and Growth Cone Turning

Recent studies indicate that the actin and microtubule cytoskeletons are a final common target of many signaling cascades influencing the developing neuron. Regulation of polymer dynamics and transport are crucial for proper growth cone motility. Three stages of axon outgrowth have been termed protrusion, engorgement, and consolidation (Dent and Gertler, 2003; Goldberg and Burmeister, 1986). Protrusion of the growth cone occurs by the rapid extension of filopodia and thin lamellar extensions while the plasma membrane advances forward. These extensions are primarily composed of bundled and mesh-like F-actin networks. In engorgement, the organelles and cytoplasm move forward and, finally, in consolidation, a new axon section is left behind (Bamburg, 2003; Goldberg and Burmeister, 1989). Within the cytoplasm are neurofilaments, which provide structure, and microtubules, which provide a mechanism for rapid vesicle transport (Rivas and Hatten, 1995; Schnapp *et al.*, 1986). Vesicle transport to the growth cone is vital to maintain and re-organize the membrane as it changes. Consolidation occurs when the majority of F-actin depolymerizes at the proximal part of the growth cone. These extension processes also occur during the formation of collateral branches from the main axon shaft (Dent *et al.*, 1999).

Axons are guided in new directions by the reorientation of their growth cones as well as by extension of collateral branches (O'Leary *et al.*, 1990). Axon branch formation occurs by controlled branch extension, retraction (including pruning), and stabilization. Similar to the primary growth cone, collaterals extending from the axon shaft are tipped by growth cones. In axon collateral initiation, extension and navigation require the controlled and coordinated assembly and disassembly of the neuronal cytoskeleton. At axon branch sites, the microtubules become unbundled and undergo local fragmentation within the axon shaft. This is accompanied by focal accumulation of F-actin. Similar to growth cone steering, axon branching is impaired by selective inhibition of microtubule or actin dynamics (Dent and Kalil, 2001; Rodriguez *et al.*, 2003). Cortical axon branching occurs *in vitro* through changes in growth cone morphologies and behaviors (Szebenyi *et al.*, 1998). Using a novel approach of visualizing simultaneous changes in both microtubules and actin filaments during different stages of axon branching in living cortical neurons, Dent and Kalil (2001) observed the

cytoskeletal reorganization underlying cortical axon branching. Branching from the growth cone and the axon shaft is always preceded by splaying apart of the looped or bundled microtubules, accompanied by localized accumulation of F-actin. Dynamic microtubules colocalize with F-actin in transition regions of growth cones and axon branch points, consistent with observations in fixed growth cones (Bridgman and Dailey, 1989; Challacombe *et al.*, 1996; Challacombe *et al.*, 1997; Rochlin *et al.*, 1999; Tanaka *et al.*, 1995; Williamson *et al.*, 1996), whereas F-actin is excluded from regions of stable microtubules. Recent evidence shows that several guidance molecules can also influence branching behavior (Dent *et al.*, 2004; Kornack and Giger, 2005). It has been shown that Netrin and Slit promote branching (Dent *et al.*, 2004; Wang *et al.*, 1999b) and Sema3A decreases branching (Dent *et al.*, 2004).

As the axon extends through the complex extracellular environment of the nervous system, its growth cone senses and responds to a variety of molecular guidance cues. For example, it can change from forward elongation to pauses or retraction, or it can change its direction. Growth cone turning is central to axonal navigation and is responsible for changing the direction of neurite elongation (Oakley and Tosney, 1991; Sabry *et al.*, 1991; Sretavan and Reichardt, 1993) (see Section VIII.C for pipette assay). A key event in growth cone turning may be the local realignment and advancement of microtubules (Lin *et al.*, 1994; Mitchison and Kirschner, 1988; Sabry *et al.*, 1991; Tanaka and Sabry, 1995) to establish the dominant side of a turning growth cone. When a growth cone contacts a positive guidance cue, filopodia become stabilized in the direction of neurite elongation, and microtubules are locally reoriented and advance toward the contact site (Bentley and O'Connor, 1994; Lin and Forscher, 1993; O'Connor and Bentley, 1993; Sabry *et al.*, 1991). In a similar manner, actin filaments and microtubules of growth cones may interact to accomplish turning away from a negative cue or stopping the extension of the axon altogether in a process called growth cone "collapse" (Fig. 2). Growth cone collapse involves the loss of lamellipodia and filopodia in response to a negative guidance cue.

In order to understand the mechanisms by which guidance cues direct axon growth it is important to determine how guidance cues affect F-actin dynamics and organization. Indeed, the rate of filopodial tip extension is determined by both the rate of F-actin polymerization at the filopodial tip and the retrograde displacement of polymerization filaments toward the base of the filopodium (Mallavarapu and Mitchison, 1999). The extension cycle of the lamellipodial edge is determined by both F-actin polymerization and retrograde filament transport (Lin and Forscher, 1995). Disruption of this cytoskeletal organization with drugs such as cytochalasin, nocodazole, and taxol results in alterations in growth cone morphology and motility (Forscher and Smith, 1988; Letourneau and Ressler, 1984; Letourneau *et al.*,

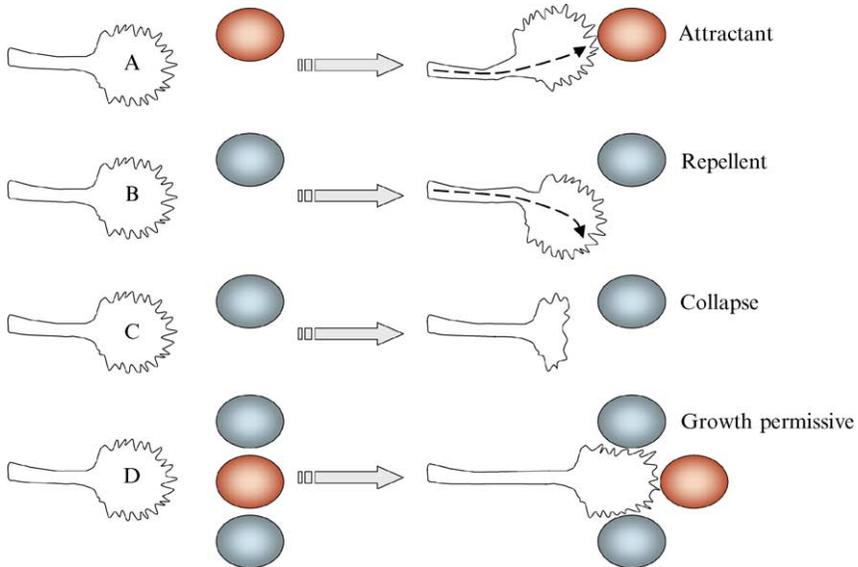


Figure 2 Growth cone behavior. The growth cone is growing toward an attractive molecule (A) or is turning away from a repulsive molecule (B). Some molecules can produce a total collapse of the growth cone (C). Some substrates are growth permissive; the growth cone grows into it with repellent cues on either side (D).

1986; Marsh and Letourneau, 1984; Neely and Gesemann, 1994; Yamada *et al.*, 1970, 1971). Disruption of either the microtubule system, by treatment with nocodazole, or the actin system, by treatment with cytochalasin, halted neuronal migration.

In vivo, growth cones are simultaneously exposed to a number of different guidance cues, both negative and positive. Repulsive guidance cues can either collapse the whole growth cone to arrest neurite outgrowth or cause asymmetric collapse leading to growth cone turning. How signals from repulsive cues are translated by growth cones into a morphological change through rearranging the cytoskeleton is unclear. Many studies examining axon repulsion have focused on the collapse response to semaphorin (Sema) 3A. Sema3A is a member of the class 3 semaphorin family that repels the axons of sensory and sympathetic neurons (Luo *et al.*, 1993). Sema3A-induced growth cone collapse is a process whereby lamellar protrusion and filopodial motility are paralyzed before the retraction of all growth cone specializations (Fan *et al.*, 1993). Fan *et al.* (1993) showed that growth cone collapse induced by Sema3A correlates with a 50% net loss of F-actin in the growth cone, suggesting that growth cone collapse could be mediated by the depolymerization of the F-actin. The morphological response to

Sema3A is similar to that caused by cytochalasin B, a plant alkaloid that inhibits actin polymerization, paralyzes normal growth cone motility, and inhibits axonal extension (Letourneau *et al.*, 1987). Collapsin response mediator proteins (CRMPs) and *rac1* have been implicated in the Sema3A signaling pathway. Treatment of neurons with either anti-CRMP-62 antibodies (Goshima *et al.*, 1995) or dominant-negative *rac1* (Jin and Strittmatter, 1997; Kuhn *et al.*, 1999) prevents growth cone collapse by Sema3A. Stimulation of endocytosis by Sema3A also correlates with growth cone collapse. Sema3A stimulates endocytosis by the focal and coordinated rearrangement of receptor and cytoskeletal elements (Fournier *et al.*, 2000). To examine the cytoskeletal events contributing to growth cone collapse, Zhou and Cohan (2001) used three factors able to induce the collapse of extending *Helisoma* growth cones: serotonin, myosin light chain kinase inhibitor, and phorbol ester. They found that all three factors induced the collapse of extending growth cones and caused actin bundle loss in poly-L-lysine-attached growth cones without loss of actin meshwork. In addition, actin bundle loss correlated with specific filamentous actin redistribution away from the leading edge, characteristic of repulsive factors. Using time-lapse studies of extending growth cones, they showed that actin bundle loss paralleled collapse. Taken together, these results suggest that F-actin reorganization through actin bundles is the cytoskeletal mechanism underlying growth cone collapse. Growth cone collapse, or at least the initiation of collapse, may be mainly a process involving F-actin reorganization resulting in decreased actin assembly at the leading edge rather than direct inhibition of actin polymerization (Zhou and Cohan, 2001). Growth cone collapse is mediated by several signaling pathways, including the Rho-GTPases (Liu and Strittmatter, 2001) ADF (actin depolymerizing factor)/cofilin (Carlier *et al.*, 1997), LIMK (LIM kinases) (Aizawa *et al.*, 2001), and GSK-3 β , FYN, and *cdk5* (Eickholt *et al.*, 2002; Sasaki *et al.*, 2002).

Positive guidance cues promote and polarize protrusive activity in the direction of growth cone migration. Gundersen and Barrett (1979) were the first to observe the turning of neuronal growth cones towards sources of nerve growth factor (NGF). Since then, additional members of the neurotrophin family have been shown to act as chemoattractants (Paves and Saarma, 1997; Song *et al.*, 1997, 1998). The Arp2/3 complex nucleates the formation of new actin branches and thus initiates the step that causes forward extension of actin meshwork and subsequent extension of the membrane leading edge (Mullins *et al.*, 1998).

Ena/VASP (enabled/vasodilator-stimulated phosphoprotein) proteins play important roles in axon outgrowth and guidance. Ena/VASP activity regulates the assembly and geometry of actin networks within fibroblast lamellipodia. Ena/VASP proteins are found in the leading edge of lamellipodia (Nakagawa *et al.*, 2001) and are concentrated in the F-actin-rich region. They function

by binding to the barbed ends of filaments and competing with capping proteins, allowing for longer filament extension (Bear *et al.*, 2002). A more recent study suggests that Ena/VASP proteins play a pivotal role in the formation and elongation of filopodia along the neurite shaft and at the growth cone (Lebrand *et al.*, 2004). Netrin-1-induced filopodia formation is dependent upon Ena/VASP function and is directly correlated with Ena/VASP phosphorylation at a regulatory protein kinase A (PKA) site. Ena/VASP function is required for filopodial formation from the growth cone in response to global PKA activation. Ena/VASP proteins likely control filopodial dynamics in neurons by remodeling the actin network in response to guidance cues (Lebrand *et al.*, 2004). Krause and colleagues (2004) identified Lamellipodin as a novel Ena/VASP-binding protein. Both proteins co-localize at the tips of lamellipodia and filopodia. Lamellipodin overexpression increases lamellipodial protrusion velocity, an effect observed when Ena/VASP proteins are overexpressed or artificially targeted to the plasma membrane. Conversely, knockdown of Lamellipodin expression impairs lamellipodia formation, reduces in velocity of residual lamellipodial protrusion, and decreases in F-actin content. Lamellipodin may act as a key convergence point linking polarized phospholipid signals and small GTPases with Ena/VASP proteins to regulate the actin cytoskeleton (Krause *et al.*, 2003, 2004). A recent study also showed that RIAM, an Ena/VASP and Profilin ligand, interacts with Rap1-GTP and mediates Rap1-induced adhesion. RIAM links Rap1 to integrin activation and plays a role in regulating actin dynamics (Lafuente *et al.*, 2004). More recently, a novel protein with high similarity to the *Caenorhabditis elegans* MIG-10 protein, called PREL1 (proline-rich Ena/VASP homology domain 1 [EVH1] ligand) has been identified. PREL1 directly binds to Ena/VASP proteins and co-localizes with them at lamellipodia tips and at focal adhesions in response to Ras activation. PREL1 provides a link from Ras signaling to the actin cytoskeleton via Ena/VASP proteins (Jenzora *et al.*, 2005). The role of Ena/VASP proteins in axon guidance will be discussed further in Section IV.G.1.

Another important component of the neuronal cytoskeleton is the microtubules. Challacombe *et al.* (1996) have shown that dynamic microtubule ends are rearranged in growth cone repulsion to avoid an inhibitory guidance cue. The importance of microtubule dynamics in axonal growth and guidance has been demonstrated by the pharmacological inhibition of dynamics without affecting microtubule assembly. Low concentrations of drugs such as nocodazole, vinblastine, and taxol not only reduce axonal elongation but also prevent growth cone turning (Challacombe *et al.*, 1997; Rochlin *et al.*, 1996; Williamson *et al.*, 1996).

Studies have shown that stathmin and SCG10 are potent microtubule destabilizing factors. While stathmin is expressed in a variety of cell types and shows a cytosolic distribution, SCG10 is neuron specific and membrane

associated. SCG10 accumulates in the central domain of the growth cone, a region that also contains highly dynamic microtubules. SCG10 appears to be an important factor for the dynamic assembly and disassembly of growth cone microtubules during axonal elongation. Phosphorylation negatively regulates the microtubule destabilizing activity of SCG10 and stathmin, suggesting that these proteins may link extracellular signals to the rearrangement of the neuronal cytoskeleton (Grenningloh *et al.*, 2004).

Zhou *et al.* (2002) report that local and specific disruption of actin bundles from the growth cone peripheral domain induced repulsive growth cone turning. Meanwhile, dynamic microtubules within the peripheral domain were oriented in areas where actin bundles remained and were lost from areas where actin bundles disappeared. This resulted in directional microtubule extension, leading to axon bending and growth cone turning. In addition, this local actin bundle loss coincided with localized growth cone collapse, as well as asymmetrical lamellipodial protrusion. Regional actin bundle reorganization can steer the growth cone by coordinating actin reorganization with microtubule dynamics. This suggests that actin bundles are potential targets of signaling pathways downstream of guidance cues, providing a mechanism for coupling changes in leading edge actin with microtubules at the central domain during turning (Zhou *et al.*, 2002).

III. The Role of Pioneering Axons and Glial Guidepost Cells in Axonal Guidance

A. Pioneering Axons

Pioneering axons are the first axons to extend along a given trajectory in the brain (Fig. 3). These axons are tipped with growth cones that respond to repellents and attractants within their environment. Here we discuss the role that pioneering axons play in guiding the formation of large axonal tracts within the brain. Once the pioneer axons have established a correct path of growth, later arriving axons, called follower axons, fasciculate with the pioneers to find their target. Often a series of pioneering axons growing over short distances may be used by the main bundle to find their targets. Ablation experiments in both vertebrates and invertebrates have established the importance of pioneering axons as a mechanism for wiring the nervous system.

1. Pioneering Axons in Invertebrates

Invertebrate organisms such as flies, worms, and grasshoppers have played a prominent role in elucidating the mechanisms that are involved in axonal guidance. The advantages of these organisms include their simple nervous

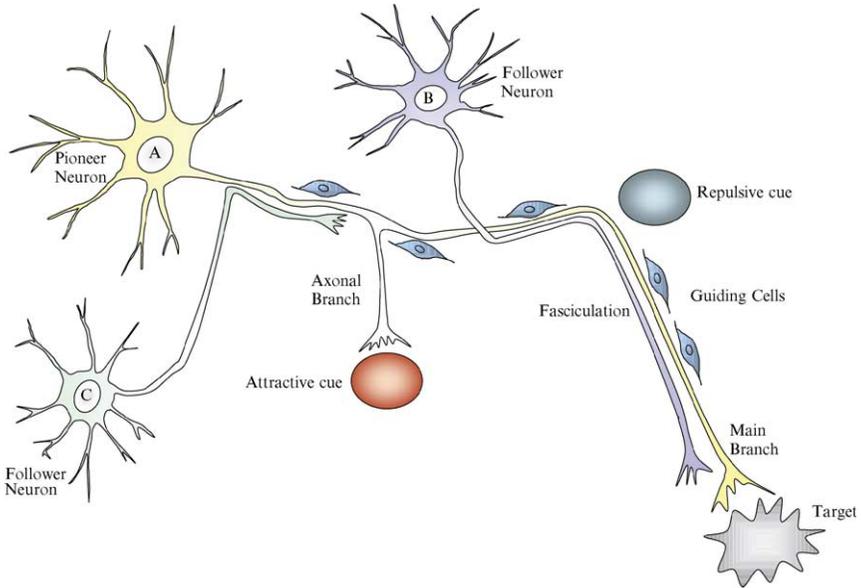


Figure 3 The pioneers. Axon of the pioneer neuron (A) grows along the pathway formed by the presence of guidepost cells as well attractive and repulsive molecules. An axonal branch can form and grow toward an attractive molecule. This axonal branch may become a stable axon, resulting in the retraction of the main branch. As pioneer neurons establish their route, follower neurons (B and C) extend their neurites. Axons of the follower neurons fasciculate with the pioneer to project to the appropriate target.

system architecture, for example, each hemisegment of the *Drosophila melanogaster* ventral nerve cord contains about 300 neurons, and the *C. elegans* nervous system has a total of 302 neurons. It is also possible to identify and follow individual neurons throughout development. Furthermore, the *Drosophila* and *C. elegans* systems allow the use of classical genetic analyses. Mutagenesis screens in both organisms have revealed many essential genes that are necessary for wiring the nervous system. In invertebrates as well as in vertebrates, axons must travel large distances to reach their final targets. Studies examining early neural development in crustacean embryos have shown the important role of pre-existing axons in guiding subsequent axon outgrowth (Whittington, 1993). Once pioneer neurons have established a route for axonal growth, axons that develop later frequently choose to selectively bundle or fasciculate with individual tracts to reach their target area.

Since axons are often pathfinding over long distances, they use guidepost cells such as pioneering axons or glial structures (also called intermediate targets) along the way to help them locate their final target. The first observation of guidepost cells came from an analysis of axon outgrowth in

the grasshopper limb bud (Bate, 1976). The T11 neuron navigates its way to the central nervous system (CNS) through key landmarks that are provided by neuronal somata. Ablation of these cells results in the failure of the neuron to efficiently complete its pathfinding. Guidepost cells in the limb bud of grasshopper embryos direct pioneer axons to the correct pathway (Bentley and Caudy, 1983; Keshishian and Bentley, 1983).

If particular pioneering axons are removed, the follower neurons stall and fail to extend (Bastiani *et al.*, 1986; du Lac *et al.*, 1986; Hidalgo and Brand, 1997; Raper *et al.*, 1984). Studies in grasshoppers (Bastiani *et al.*, 1984; Goodman *et al.*, 1984; Raper *et al.*, 1983a,b, 1984) demonstrate that follower axons can recognize molecules on the surface of the pioneering axon. The growth cones of two follower axons, the G and C cells, in the developing grasshopper neuropil follow four pioneering neurons (the A1, A2, P1, and P2 cells) to their targets. The G neuron extends anteriorly in the bundle and the C neuron extends posteriorly in the same bundle, but only after several other axons have joined the bundle. These results suggest that the growth cones of the G and C neurons can recognize and extend upon the four pioneering axons. This has been described as the labeled pathways hypothesis (Goodman *et al.*, 1984), which proposes that axon tracts have different molecular labels on their cell surface, labels that follower axons can specifically recognize to extend upon.

Some molecules are beginning to be identified for pioneering axon-mediated guidance. AcP neurons pioneer the anterior commissure of each grasshopper segment and extend their axons toward the midline. A cell surface glycoprotein, Lazarillo, is expressed during grasshopper embryogenesis on the surface of a subset of CNS neurons and by a group of neuroblasts, the precursors of neurons and glia (Sanchez *et al.*, 1995, 2000). When Lazarillo function was blocked, AcP neurons failed to grow into the midline; they either stopped growing or turned anteriorly. These results suggest that Lazarillo could play a role in the guidance of the AcP pioneering neurons (Sanchez *et al.*, 1995, 2000).

Goodman and Jacobs characterized neurons that pioneer the major CNS axon tracts in *Drosophila* (Jacobs and Goodman, 1989). In the embryonic *Drosophila* CNS, RP motor axons make stereotypic pathway choices involving distinct cellular contacts: (i) extension across the midline via contact with the axon and cell body of the homologous contralateral RP motoneuron, (ii) extension down the contralateral longitudinal connective through contact with connective axons and longitudinal glia, and (iii) growth into the intersegmental nerve (ISN) through contact with ISN axons and the segmental boundary glial cell (SBC). Removal of the longitudinal glia or the SBC did not adversely affect pathfinding. This suggests that the motor axons either utilized alternative axonal substrates or could still make filopodial contact with the cues in the next pathway. In contrast, RP motor axons did require contact

with the axon and soma of their contralateral RP homologue. Absence of this neuronal substrate frequently impeded RP axon outgrowth, suggesting that the next cues were beyond filopodial reach. Together, these direct ablations of putative guidepost cells in the CNS demonstrated a susceptibility by RP axons to the absence of specific cellular contacts (Whittington *et al.*, 2004).

In the nematode *C. elegans*, the gene *unc-6* is required to guide pioneer axons and mesoblasts in dorsal or ventral directions of the body wall (Hedgecock *et al.*, 1990). UNC-6 is required to direct the early circumferential extension of pioneer axons in *C. elegans*, as well as the circumferential migrations of mesoblast cells (which migrate along the basement membrane surfaces facing the central pseudocoelomic cavity). Nerves form as additional axons enter the tract and fasciculate with the pioneering axon and with each other. These followers enter the tract in a set order and at stereotyped positions. Key to the role that UNC-6 plays in this process is its dynamic expression pattern (Wadsworth, 2002; Wadsworth *et al.*, 1996). The *unc-129* gene, like the *unc-6* gene, is required to guide pioneer motor axons along the dorsoventral axis of *C. elegans*. UNC-129 mediates expression of dorsoventral polarity information required for axon guidance and guided cell migration (Colavita *et al.*, 1998).

In *C. elegans*, synaptic guidepost cells in the vulval epithelium initiate the formation of synapses between the HSNL (hermaphrodite-specific neurons left) and HSNR (hermaphrodite-specific neurons right) and either vulval muscle cells or ventral type C (VC) neurons (Shen and Bargmann, 2003). In the absence of the guidepost cells, synaptic vesicle markers in the HSNL fail to accumulate at the normal synaptic locations, and instead form ectopic aggregates in anterior locations. *Syg-1* encodes a transmembrane immunoglobulin superfamily protein acting in the presynaptic HSN axon. Analysis of the *syg-1* mutant shows that HSNL synapses have defects similar to those observed after guidepost cell ablation, suggesting a role for SYG-1 in guidepost signaling (Shen and Bargmann, 2003). Another protein, SYG-2, binds heterophilically to SYG-1 and is expressed by guidepost cells and vulval epithelial cells. *Syg-2* mutants lack synapses at the normal location and instead form synapses onto inappropriate target cells at ectopic locations (Shen *et al.*, 2004). Molecular interactions between SYG-1 and SYG-2 on neuron and guidepost cells, respectively, contribute to synaptic specificity but also serve to suppress the formation of inappropriate synapses.

2. Pioneering Axons in Vertebrates

Pioneering axons have also been described in zebrafish. The first axons to navigate the neuroepithelium of the zebrafish brain emerge from the ventrocaudal cluster (vcc) at approximately 16 hr postfertilization (hpf) (Chitnis and Kuwada, 1990; Ross *et al.*, 1992). These axons grow caudally to pioneer

the medial longitudinal fasciculus (MLF), which is part of the larger ventral longitudinal tract (VLT) (Chitnis and Kuwada, 1990; Ross *et al.*, 1992; Wilson *et al.*, 1990), the major longitudinal tract that connects the midbrain with the hindbrain. By 18 hpf, the first axons in the forebrain emerge from the ventrorostral cluster (vrc) and grow caudally to pioneer the tract of the postoptic commissure (TPOC) (Chitnis and Kuwada, 1990; Ross *et al.*, 1992; reviewed in Hjorth and Key, 2002). Nural and Mastick (2004) also studied the postoptic commissure (POC) and characterized a system of early neurons that establish the first two major longitudinal tracts in the embryonic mouse forebrain. Each of the early axon populations first grows independently, pioneering a short segment of new tract. However, each axon population soon merges with other axons to form one of only two shared longitudinal tracts, both descending: the TPOC, and, in parallel, the stria medullaris. Thus, the forebrain longitudinal tracts are pioneered by a relay of axons, with distinct axon populations pioneering successive segments of these pathways. They identified that the transcription factor Pax6 is critical for tract formation. In Pax6 mutants, both the TPOC and the stria medullaris failed to form due to pathfinding errors of the early pioneering axons. Their results show that Pax6 could regulate longitudinal tract formation by guiding a relay of pioneer longitudinal axons in the embryonic mouse forebrain (Nural and Mastick, 2004).

Fraser and colleagues (2003) studied the POC using *in vivo* microscopy of embryonic zebrafish expressing green fluorescent protein (GFP) in the vrcs of cells in the embryonic forebrain. Their data showed that the growth of the leader pioneering axons slows down at the midline, but not the follower axons. When the leading pioneer axon is ablated, the follower axons change their midline kinetics and behave as leaders. Similarly, once the leader axons have crossed the midline they change their midline kinetics when they encounter the leading axon from the contralateral side (Bak and Fraser, 2003). These data suggest a simple model in which the level of growth cone exposure to midline cues and the presence of other axons as a substrate shape the midline kinetics of commissural axons.

In the neocortex, subplate neurons have been shown to serve as pioneering axons for thalamocortical and corticothalamic axons (De Carlos and O'Leary, 1992; Ghosh and Shatz, 1992, 1993; Ghosh *et al.*, 1990; McConnell *et al.*, 1989). When subplate neurons are ablated both corticothalamic and thalamocortical targeting are disrupted demonstrating a direct requirement for pioneering subplate neurons in forming connections between the thalamus and cortex (Ghosh and Shatz, 1992; Ghosh *et al.*, 1990; McConnell *et al.*, 1994).

In the medial cortical projection the first axons to cross the rostral cortical midline (rostral to the hippocampal commissure) are derived from neurons in the cingulate cortex (Koester and O'Leary, 1994; Ozaki and Wahlsten,

1998; Rash and Richards, 2001). These axons begin to cross the midline at embryonic day (E) 17 in the rat and E15.5 in the mouse. Neurons in the cingulate cortex project to three different regions: across the midline into the contralateral cortex, into the fornix, and ventrally into the medial septum and the diagonal band of Broca. The cingulate axons cross the midline first, followed by the neocortical axons, which grow within the tract of the cingulate pioneering axons, possibly fasciculating with the cingulate axons (Rash and Richards, 2001).

The optic chiasm also contains a subset of early generated neurons, which have been shown to be involved in retinal axon guidance (Easter *et al.*, 1993; Sretavan *et al.*, 1994, 1995). Furthermore, Cajal-Retzius cells in the hippocampus are suggested to play a role in the guidance of entorhinohippocampal axons (Del Rio *et al.*, 1997; Soriano *et al.*, 1994). In fact, two groups of pioneer neurons, Cajal-Retzius cells and GABAergic neurons, form layer-specific scaffolds that overlap with distinct hippocampal afferents at embryonic and early postnatal stages. Before the dendrites of pyramidal neurons develop, these pioneer neurons act as synaptic targets for hippocampal afferents. These findings indicate that distinct pioneer neurons are involved in the guidance and targeting of different hippocampal afferents (Super *et al.*, 1998).

Recent evidence also suggests the presence of guidepost cells in the developing mouse olfactory system. Mitral cell axons, the major efferents of the olfactory bulb, caudally elongate in a very narrow part of the lateral telencephalon and make a stereotyped turn toward the amygdala (Brunjes and Frazier, 1986; Schwob and Price, 1984; Shipley *et al.*, 1995). The axons collectively form a fiber bundle called the lateral olfactory tract (LOT). Organotypic co-culture of olfactory bulb with various regions of the mouse telencephalon showed that mitral cell axons are guided by biochemical cues that are strictly localized to the telencephalon (Sugisaki *et al.*, 1996). Indeed, intrinsic cells in the telencephalon play a directional role in the guidance of mitral cell axons (Sugisaki *et al.*, 1996). Mitral cell axons selectively grew along the LOT cells *in vivo* and in co-culture. Ablation of LOT cells in organotypic cultures caused mitral cell axons to stall. These results suggest that LOT cells function as guidepost cells for mitral cell axons (Sato *et al.*, 1998).

B. Intermediate Targets and Glial Guidepost Cells

Often, the final target of an axon is a long distance away. In order to navigate toward their final target, axons use intermediate targets to guide them along the correct path of growth. Such intermediate targets can be pioneering axon populations such as those described above or glial cells or structures present along the pathway that secrete guidance factors (Fig. 3).

Once a growth cone encounters such an intermediate target, it slows and transforms its morphology, apparently looking for further molecular “directions.” Guidepost cells have been identified in *Drosophila* (Auld, 1999; Hidalgo, 2003) in which glia secrete guidance cues and express cellular cues on their surface that guide axonal outgrowth. Glia act as intermediate targets in growth cone guidance. Because guidepost cells were found in invertebrates, similar guiding mechanisms have been postulated in the mammalian CNS. In vertebrates, primitive glial cells are involved in guiding pioneering growth cones in the developing spinal cord (Kuwada, 1986; Singer *et al.*, 1979), the ventral roots (Nordlander *et al.*, 1981), the optic nerve (Silver and Sapiro, 1981; Silver and Sidman, 1980), the auditory system (Carney and Silver, 1983) and the developing corpus callosum (Silver *et al.*, 1982, 1993). During embryonic development, glia cells are required for the formation of the CNS (Fitch and Silver, 1997). They also define boundaries between different brain areas or between functional subdomains within the same area (Cooper and Steindler, 1986; Garcia-Abreu *et al.*, 1995; Mastick and Easter, 1996; Silver, 1994; Silver *et al.*, 1993; Yoshida and Colman, 2000). These glial boundaries serve to prevent axons from straying from their correct path of growth (Fitch and Silver, 1997). At the *Drosophila* midline, glia function as guidepost cells for commissural and ipsilaterally projecting axons to determine which axons cross the midline and which do not (Hidalgo and Booth, 2000; Jacobs and Goodman, 1989; Kidd *et al.*, 1999). To better understand the origin of commissures, a mutant screen was carried out for flies with defective commissure formation. Commissureless (Comm) is one of the mutations isolated. In Comm mutants, commissural growth cones initially orient toward the midline, but none actually cross it. Rather, any short medially oriented processes are retracted, and the axons remain exclusively on their own side, producing the commissureless axon guidance phenotype (Seeger *et al.*, 1993). Robo was isolated in the same screen, and leads to the opposite misrouting; some growth cones that normally extend only on their own side project across the midline in Robo mutants. The phenotype of these two genes suggests that they encode components of attractive and repulsive signaling systems at the midline. Comm is able to downregulate levels of the Robo protein on the cell surface, which is necessary for axons to cross the midline. Comm is expressed and required in both commissural neurons and midline cells for correct midline crossing (Georgiou and Tear, 2002). It is suggested that the presence of Comm in the commissural neurons may encourage midline crossing. Comm protein accumulates at the axon surface within the commissural region, using a mechanism that is likely to involve Comm expressed by midline glia (Couch and Condron, 2002; Georgiou and Tear, 2002; Keleman *et al.*, 2002; Keleman *et al.*, 2005; Myat *et al.*, 2002).

Three pairs of midline glia, as well as the medial precursor 1 (MP1) and ventral unpaired median (VUM) neurons, are present at the midline of the *Drosophila* ventral nerve cord (Klambt *et al.*, 1991). Analysis of mutants defective in midline cell development reveal essential roles for these cells in the formation of the *Drosophila* ventral nerve cord. In the single-minded (*sim*) mutant, midline cells fail to differentiate and ultimately die. Consequently, commissures do not form and the longitudinal connectives collapse into a single fused tract at the midline (Klambt *et al.*, 1991; Thomas *et al.*, 1988).

Glial cells may also function to sort ipsilaterally from contralaterally projecting axons. At the optic chiasm, retinal ganglion cell (RGC) axons contact glial cells known as the glial palisade. These glia may contribute to retinal axonal divergence at the chiasm (Erskine *et al.*, 2000; Marcus *et al.*, 1995). Glial populations are also associated with the formation of commissures such as the anterior commissure (Cummings *et al.*, 1997; Pires-Neto *et al.*, 1998), the corpus callosum (Shu and Richards, 2001; Silver *et al.*, 1993) (see Section V.B), and decussating axons in the hindbrain, brain stem, and the corticospinal tract (Joosten *et al.*, 1989; Mori *et al.*, 1990; Van Hartesveldt *et al.*, 1986). Glial cells in the floor plate of the spinal cord guide commissural axons of the dorsal spinal cord (Altman and Bayer, 1984; Tessier-Lavigne *et al.*, 1988a).

IV. Molecules Involved in Axonal Guidance

Neuronal growth cones are guided to their targets by both short- and long-range cues. Both attractive and repulsive cues are equally important for the guidance of growth cones to their appropriate targets. Most axon guidance molecules identified were discovered first in *Drosophila* or *C. elegans*. In some cases, guidance factors were simultaneously directly purified from vertebrate systems through massive purification efforts. There are four major families of classic axon guidance molecules: Netrins, Slits, semaphorins, and ephrins and their receptors. However, a number of other molecules have been shown to guide axons, including morphogens, steroids, and extracellular matrix and adhesion molecules. Here we briefly review these molecules and some of the intracellular signaling mechanisms they use.

A. Netrins and DCC

Netrin-1, whose name stems from the Sanskrit term “one who guides,” was originally purified in vertebrates as a floorplate-derived chemoattractant by an exhaustive purification from over 10,000 chick brains that can promote commissural axon outgrowth (Kennedy *et al.*, 1994; Serafini *et al.*, 1994). Netrins are secreted proteins that act on neural cells through transmembrane

receptors of the Neogenin A2b receptor, DCC (deleted in colorectal carcinomas), and UNC5H families (Mehlen and Mazelin, 2003) (Fig. 4). Netrins make up a small family of secreted, laminin-related molecules with multi-functional roles in axon guidance, acting as context-dependent chemoattractants or chemorepellents. Several members of the Netrin family have been identified in a variety of species: the UNC-6 gene product in nematodes;

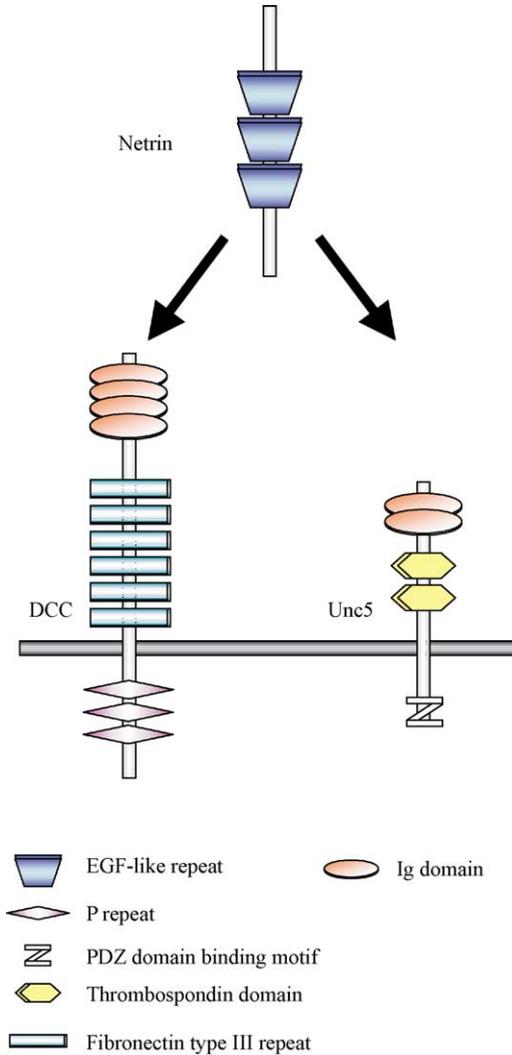


Figure 4 Netrins and DCC.

netrin-A and netrin-B in *Drosophila*; netrin-1 and netrin-2 in chicks; netrin-1, netrin-3, netrin-G1, netrin-G2, and netrin-4/ β -netrin in mice; and the NTN2L (netrin-2-like) in humans (Koch *et al.*, 2000; Nakashiba *et al.*, 2002; Serafini *et al.*, 1994, 1996; Wang *et al.*, 1999a). The Netrins encode ~60–80 kDa secreted proteins that share homologous domains with laminin (Banyai and Patthy, 1999; Serafini *et al.*, 1994). The *C. elegans* netrin (UNC6) was the first member of this family identified by examining mutants with uncoordinated (unc) phenotypes (Hedgecock *et al.*, 1990; Ishii *et al.*, 1992). Netrins regulate the development of commissural axons in both the spinal cord and brain, including the corpus callosum, hippocampal commissure, and the optic chiasm (Serafini *et al.*, 1996). Netrins are not only involved in axon guidance but also play central roles in the migration of neurons, glial oligodendrocyte precursors, and mesodermal cells during embryogenesis (Alcantara *et al.*, 2000; Bloch-Gallego *et al.*, 1999; Hamasaki *et al.*, 2001; Lim and Wadsworth, 2002; Spassky *et al.*, 2002; Su *et al.*, 2000; Sugimoto *et al.*, 2001; Tsai *et al.*, 2003; Yee *et al.*, 1999). More recently, it has been shown that the Netrin-1 receptor DCC is phosphorylated by Fyn and that phosphorylation is required for DCC function. This suggests that Fyn is essential to initiate the responses of axons to Netrin-1 (Meriane *et al.*, 2004).

B. Semaphorins, Neuropilins, and Plexins

Semaphorins (Semas) were originally identified in invertebrates (Kolodkin *et al.*, 1992). *Sema3A*, the prototype vertebrate member of the semaphorin family, was initially purified from chick brain extracts on the basis of its collapse-inducing activity on cultured dorsal root ganglion (DRG) growth cones (Luo *et al.*, 1993). The semaphorin family contains both secreted and membrane-bound members, divided into eight classes (Semaphorin Nomenclature Committee, 1999). The first two classes represent invertebrate semaphorins, classes 3–7 represent vertebrate semaphorins, and the eighth class comprises viral semaphorins. Of the vertebrate classes (Fig. 5), class 3 contains secreted semaphorins and classes 4–7 contain transmembrane or membrane-anchored semaphorins. All semaphorins share a conserved, 500-amino-acid motif, termed the sema domain (Kolodkin *et al.*, 1993; Luo *et al.*, 1993). The first semaphorin receptor identified was neuropilin (Chen *et al.*, 1997; Giger *et al.*, 1998; He and Tessier-Lavigne, 1997; Kolodkin *et al.*, 1997). The two main protein families now known to be involved in mediating semaphorin responses are the neuropilins (two members, Npn1 and Npn2; Fig. 5) and plexins (nine members), Plexin A1–4, Plexin B1–3, Plexin C1, and Plexin D1 (de Wit and Verhaagen, 2003; Fujisawa and Kitsukawa, 1998; Pasterkamp and Kolodkin, 2003; Raper, 2000; Tamagnone *et al.*, 1999; Winberg *et al.*, 1998). Invertebrate semaphorins, membrane-associated

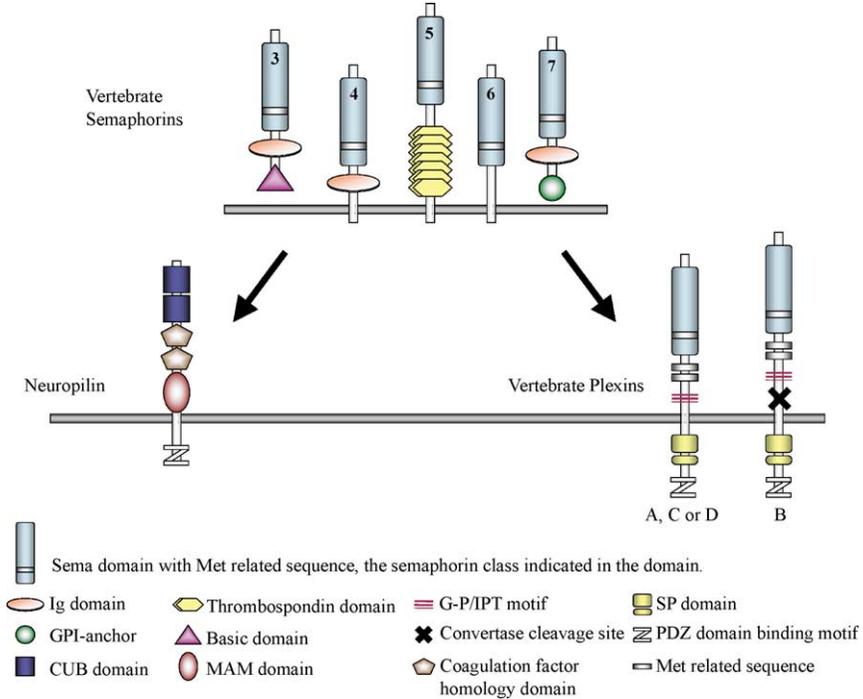


Figure 5 Semaphorins, Neuropilins, and Plexins.

semaphorins in vertebrates, and viral semaphorins have been shown to interact directly with plexins (Comeau *et al.*, 1998; Tamagnone *et al.*, 1999; Winberg *et al.*, 1998). Vertebrate class 3 secreted semaphorins, however, utilize neuropilin proteins as ligand-binding obligate co-receptors, which assemble a semaphorin/neuropilin/plexin signaling complex (Chen *et al.*, 1997; He and Tessier-Lavigne, 1997; Takahashi *et al.*, 1999; Tamagnone *et al.*, 1999).

All four members of the plexin-A subfamily can function as a signal-transducing component in class 3 semaphorin receptor complexes. Plexin A1, Plexin A2, and Plexin A4, when expressed in a complex with Npn-1 or Npn-2, induce a contraction of the cytoskeleton of COS cells in response to class 3 semaphorins (Suto *et al.*, 2003; Takahashi and Strittmatter, 2001). Cultured sympathetic axons derived from Plexin A3 knockout mice, however, are no longer repelled by Sema3F and partially lose their responsiveness to Sema3A (Cheng *et al.*, 2001). The response to class 3 semaphorins is determined by the complement of neuropilins on the cell surface, and

additional specificity is achieved by the combination of plexin-As expressed in the neuron.

Npn1 can bind another ligand, vascular endothelial growth factor (VEGF) (Soker *et al.*, 1998), and several proteins other than neuropilins and plexins can participate in semaphorin receptor complexes. These include the cell adhesion molecule L1, which transduces a chemorepulsive response to *Sema3A* in cortical neurons, together with Npn1 (Castellani *et al.*, 2000, 2002). To study the role of Npn1-*Sema* signaling independent of VEGF/Npn1 signaling, Gu and colleagues specifically mutated the *Sema*-binding domain of Npn1, while leaving the VEGF-binding domain intact (Gu *et al.*, 2003). These mice have axonal guidance defects in the formation of the entorhino-hippocampal pathway, cranial and spinal nerves, sensory projections to the inner ear, and the corpus callosum (Gu *et al.*, 2003).

Recently, integrins have been shown to function as receptors for *Sema7A*, which induces olfactory axon growth without the need for plexins (Pasterkamp *et al.*, 2003). Finally, recent data suggest that the attractive/repulsive signaling of *Sema5A* can be modulated by interactions with the extracellular matrix (Kantor *et al.*, 2004; see further below).

C. Slits and Robos

Slit proteins make up a family of multifunctional guidance cues with putative roles in regulating neuronal migration (Wu *et al.*, 1999), axonal and dendritic branching (Ozdinler and Erzurumlu, 2002; Wang *et al.*, 1999b; Whitford *et al.*, 2002), and axon guidance (Brose *et al.*, 1999; Kidd *et al.*, 1999). These large glycoproteins are conserved across species with three family members (Slit1, Slit2, and Slit3) identified in the developing and adult mammalian nervous systems (Brose *et al.*, 1999; Marillat *et al.*, 2002). They contain several protein motifs: leucine-rich repeats, EFG repeats, and a laminin G domain (Rothberg *et al.*, 1990) (Fig. 6). The three vertebrate Slits have overlapping, but distinct, patterns of expression throughout development and in adulthood (Holmes *et al.*, 1998; Itoh *et al.*, 1998; Marillat *et al.*, 2002; Piper *et al.*, 2000; Yuan *et al.*, 1999). Knockout studies demonstrate that Slits 1 and 2 play critical roles in the formation of several mammalian fiber tracts, including corticofugal, thalamocortical, callosal, optic, and the lateral olfactory tract (Bagri *et al.*, 2002; Keleman *et al.*, 2002; Nguyen-Ba-Charvet *et al.*, 2002; Plump *et al.*, 2002). The Robo family of transmembrane proteins are the receptors for Slits (Brose *et al.*, 1999; Fricke *et al.*, 2001; Kidd *et al.*, 1998; Zallen *et al.*, 1998). Three mammalian homologues of *Drosophila* Robo have been identified (Robo1, Robo2, and Rig-1). Robos are members of the immunoglobulin (Ig) superfamily; their ectodomain contains five Ig-like repeats followed by three fibronectin type III

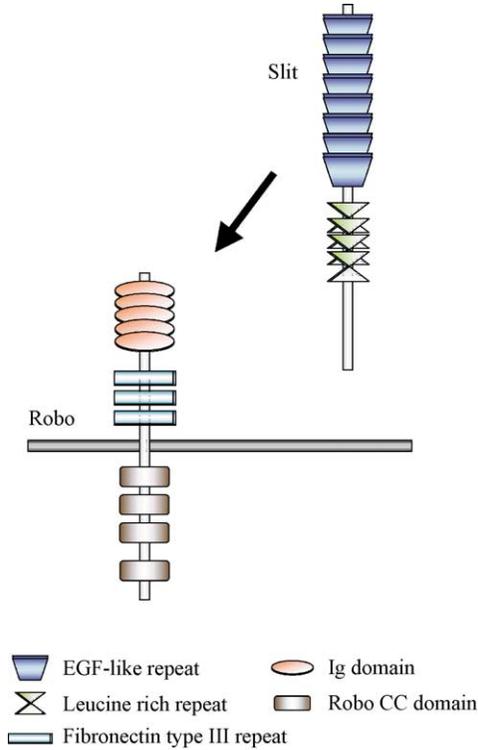


Figure 6 Slits and Robos.

repeats (Kidd *et al.*, 1998). In *Drosophila*, ipsilaterally projecting axons and decussated commissural axons expressing Robo are prevented from inappropriate crossing/re-crossing of the midline via interaction with Slit (Kidd *et al.*, 1999). Genetic mutations in either Robo or Slit lead to aberrant crossing and re-crossing, or failure of these axons to leave the midline (Kidd *et al.*, 1999). In *Drosophila*, commissural axons acquire a postcrossing sensitivity to Slit resulting from increased surface expression of Robo. This occurs via a mechanism that includes inactivation of Comm, an intracellular sorting protein that normally targets Robo for endosomal degradation (Keleman *et al.*, 2002). In Comm mutants, commissural growth cones initially orient toward the midline, but none actually cross. Rather, any short medially oriented processes are retracted, and the axons remain exclusively on their own side, producing the commissureless phenotype (Seeger *et al.*, 1993). Downregulation of Robo is necessary for axons to cross the midline. Comm is expressed and required in both commissural neurons and midline

cells for correct midline crossing (Georgiou and Tear, 2002). It is suggested that the presence of Comm in commissural neurons may encourage midline crossing. Comm protein accumulates at the axon surface within the commissural region, using a mechanism that is likely to involve Comm in the midline glia. However, Comm activity does not extend beyond the midline, allowing Robo levels to increase at the growth cone surface and initiate sensitivity to the midline inhibitor Slit that encourages axon growth away from the midline and prevents re-crossing (Couch and Condrón, 2002; Georgiou and Tear, 2002). Thus, Comm controls axon guidance at the midline by regulating surface levels of Robo. Two different models have been proposed to explain how Comm regulates Robo. The first model proposes that Comm controls the sorting of Robo at the trans-Golgi network (Keleman *et al.*, 2002). The second model proposes that Comm controls Robo by acting at the plasma membrane. In this model, Comm does not block the delivery of Robo to the growth cone but instead rapidly removes it by endocytosis (Myat *et al.*, 2002). In a genetic rescue assay for Comm, Dickson and colleagues showed that midline crossing does not require the presence of Comm in midline cells (as proposed in the second model). They also showed by monitoring the trafficking of Robo that Comm prevents the delivery of Robo to the growth cone (as predicted in the first model) (Keleman *et al.*, 2005).

Slit proteins are also alternatively spliced in both mouse and human, implying that multiple Slit protein isoforms may exist (Little *et al.*, 2002). Recent studies of the Slit1 protein show an alternatively spliced mRNA product for *slit1* found specifically in the vertebrate nervous system (Tanno *et al.*, 2004). This variant was designated *slit1 α* . *Slit1 α* is specifically expressed in rat brain, but not heart or kidney, suggesting that *slit1 α* plays a role in nervous system development. *Slit1 α* expression was found in the cerebral cortex and the hippocampus. *In vivo*, *slit1 α* acts as a chemorepellent of olfactory bulbs axons (Tanno *et al.*, 2004).

D. Ephrins and Eph

Ephrins and their tyrosine kinase receptors, the Eph molecules, are divided into two classes: ephrin-As, which are anchored to the membrane via glycosyl-phosphatidylinositol (GPI) linkage, and ephrin-Bs, which are transmembrane proteins (Fig. 7). The ephrin-A subclass contains ephrins A1 to A5 and the ephrin-B subclass has three members, ephrins B1 to B3 (Kullander and Klein, 2002). Eph receptors are divided into an A subclass that contains eight members (EphA1–EphA8), and a B subclass that contains five members (EphB1–EphB4, EphB6) (Cutforth and Harrison, 2002; Huot, 2004; Wilkinson, 2001). A-type receptors bind to most or all A-type ligands, and

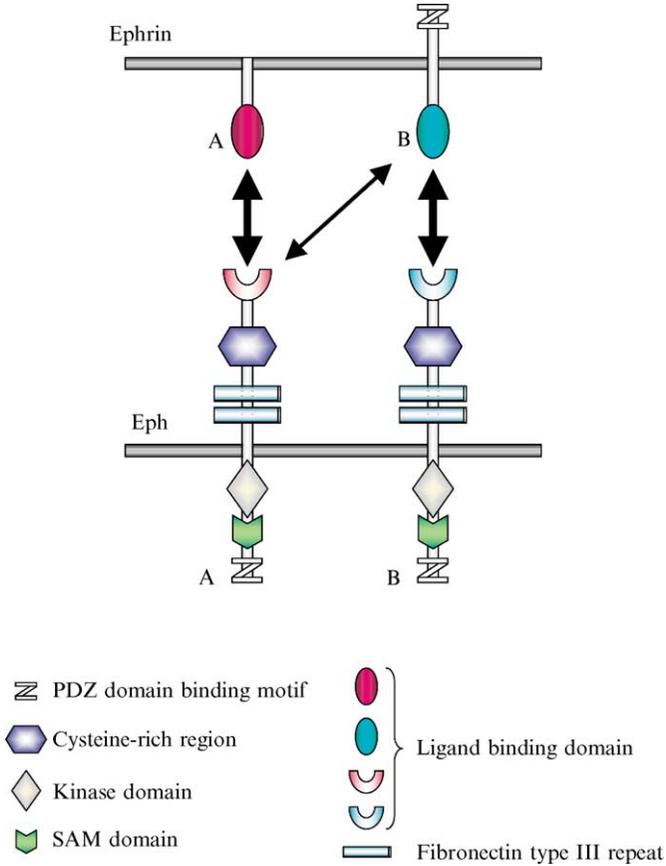


Figure 7 Ephrins and Eph.

B-type receptors bind to most or all B-type ligands. The primary exception is the EphA4 receptor, which has been shown to interact with members of both class A and class B ephrins. The extracellular domain of the Eph receptors contains the ligand-binding domain, a cysteine-rich region, and two fibronectin type III repeats. The cytoplasmic domain of the Eph receptors can be divided into four functional units: the juxtamembrane domain that contains two conserved tyrosine residues, a classical protein tyrosine kinase domain, a sterile- α -motif, and a PDZ domain binding motif (Huot, 2004; Kullander and Klein, 2002). Since both the ephrin ligands and the Eph receptors are membrane bound, the interactions between ephrins and the Eph receptors require intracellular contact. An Eph receptor can also act as a ligand in the same way that an ephrin ligand can act as a receptor (Mellitzer *et al.*, 1999).

Ligand binding induces forward signaling, but ephrins can also signal via the receptor, which is called reverse signaling (Bruckner *et al.*, 1997; Holland *et al.*, 1996).

The ephrin–Eph receptor system regulates many cellular functions that depend upon cytoskeletal remodeling, such as axon guidance and synaptic plasticity. The influence of Eph–ephrin activation on cell behavior differs depending on the cell type. They are generally associated with repulsion of neighboring cells or of cellular processes, such as the neuronal growth cone. However, in some cases, Eph–ephrin activation leads to increased adhesion or attraction. Both classes have been implicated as regulators of axon guidance. Eph/ephrin molecules have also been implicated in guiding commissural axons in the spinal cord and brain (Henkemeyer *et al.*, 1996; Hu *et al.*, 2003; Imondi and Kaprielian, 2001; Imondi *et al.*, 2000; Orioli *et al.*, 1996; Palmer and Klein, 2003; Yokoyama *et al.*, 2001). Ephrin–Eph complexes also regulate axon guidance in the visual system. In particular, ephrin-A ligands and Eph A receptors mediate repulsion that is typically involved in regulating the mapping of retinal axons along the anterior-posterior axis (reviewed in Huot, 2004; see also Section VIII.B for the stripe assay). Topographic mapping of the anterior-posterior tectal/superior collicular axis is dependent upon expression patterns of EphA/ephrin-A (Brown *et al.*, 2000; Feldheim *et al.*, 2000), while EphB/ephrin-B signaling is important for mapping along the dorsoventral axis (Hindges *et al.*, 2002; Mann *et al.*, 2002b).

E. Morphogens

Morphogens are known for their effects on cell fate determination during development. Local concentration gradients convey positional information used during organization of the major body axes, limb development, and patterning of the nervous system. Recent studies suggest that in addition to these roles, morphogens, including members of the Wnt family, bone morphogenetic protein (BMP) family, and Sonic Hedgehog (Shh), might also function in axon guidance. For instance, a knockout of the Wnt receptor, Frizzled-3, results in severe defects in several major fiber tracts in the vertebrate forebrain (Wang *et al.*, 2002c), and Wnt-3 slows axon outgrowth and may mediate terminal branching of vertebrate muscle afferents (Krylova *et al.*, 2002). Wnt proteins are a large family of diffusible factors (7 members in *Drosophila* and 19 in humans) that play several important roles in both embryonic development and adult function. Wnt proteins have well-established roles in early cell fate decisions and embryonic patterning (Wodarz and Nusse, 1998), but they have also been implicated in synaptic remodeling and terminal arborization within the developing CNS (Hall *et al.*, 2000;

Krylova *et al.*, 2002; Packard *et al.*, 2002) and in planar cell polarity (Bhanot *et al.*, 1996; Vinson and Adler, 1987). In the *Drosophila* ventral nerve cord, Wnt-5 binding to the receptor tyrosine kinase Derailed is required for targeting axons to the appropriate midline commissure (Yoshikawa *et al.*, 2003). Genetic and biochemical data indicate that Wnt5 binds Derailed and prevents Derailed-expressing axons from entering the posterior commissure (Garrity, 2003). The dorsal spinal cord commissural neurons form several ascending somatosensory pathways. During embryonic development, they project axons to the ventral midline. At the floor plate, commissural axons cross the midline, enter the contralateral side of the spinal cord, and make a sharp anterior turn toward the brain (Bovolenta and Dodd, 1990). *In situ* hybridizations of Wnts in developing mouse embryos revealed that Wnt4, Wnt7b, and Wnt5a are expressed in areas where postcrossing axons turn anteriorly. Wnt4 was found specifically enriched in the floor plate and the ventricular zone, exhibiting a decreasing anterior-to-posterior gradient along the entire length of the floor plate (Lyuksyutova *et al.*, 2003). A similar Wnt4b gradient in the floor plate was found in zebrafish embryos at similar stages (Liu *et al.*, 2000). A directed source of Wnt4 protein attracted postcrossing commissural axons (Lyuksyutova *et al.*, 2003). Commissural axons in mice lacking the Wnt receptor Frizzled3 displayed anterior-posterior guidance defects after midline crossing. Thus, Wnt-Frizzled signaling guides commissural axons along the anterior-posterior axis of the spinal cord (Lyuksyutova *et al.*, 2003).

A role for BMPs in axonal guidance in vertebrates has emerged from studies of commissural axon trajectories in the developing spinal cord. Many commissural neurons differentiate adjacent to the roof plate at the dorsal midline of the spinal cord and extend axons ventrally (Dodd *et al.*, 1988; Holley, 1982). Signals derived from the floor plate contribute to the ventral trajectory of commissural axons (Colamarino and Tessier-Lavigne, 1995). The initiation of ventral growth of commissural axons may be mediated by a chemorepellent signal emanating from the roof plate. *In vitro* studies have shown that the roof plate is the source of a diffusible repellent activity that orients commissural axons in explants and that this repellent activity can be blocked by antagonists of BMPs (Augsburger *et al.*, 1999). At the time that commissural axon extension is initiated, *Bmp6*, *Bmp7*, and *Gdf7* are expressed in the rodent roof plate (Augsburger *et al.*, 1999; Lee *et al.*, 1998). BMP7 can mimic the repellent activity of the roof plate on commissural axons in explants *in vitro*, whereas BMP6 has only a low level of repellent activity, and GDF7 is inactive. Moreover, BMP7 elicits commissural growth cone collapse, illustrating the direct nature of its action. Roof plate tissue isolated from *Bmp7* mutant mice exhibits a marked reduction in roof plate-repellent activity *in vitro* (Augsburger *et al.*, 1999). Together, these findings suggest that BMPs can act as axon guidance signals that contribute to the

chemorepellent activity of the roof plate. A more recent study analyzing roof plate-repellent activity in mice lacking *Bmp7*, *Bmp6*, and *Gdf7* alone and in pair-wise combinations show that both *Gdf7* and *Bmp7* but not *Bmp6* are required for the ability of the roof plate to orient commissural axons. GDF7 and BMP7 heterodimerize, and the heterodimer is a more potent repellent than the BMP7 homodimer for commissural axons. These results suggest that the GDF7:BMP7 heterodimer functions as a roof plate-derived repellent that establishes the initial ventral trajectory of commissural axons (Butler and Dodd, 2003).

Other evidence for morphogen involvement in axon guidance includes defects in retinal ganglion cell projections in mice deficient in the BMP receptor, BMPR-IB (Liu *et al.*, 2003). Shh can inhibit the outgrowth of neurons from retinal explants *in vitro* (Trousse *et al.*, 2001). Shh is a secreted protein that interacts with two transmembrane proteins, Patched (*ptc*) and Smoothed (*smo*). *Ptc* binds to *shh*, whereas *smo* is involved in signal transduction. In the absence of *shh*, *ptc* inhibits *smo*. A recent study by Charron and colleagues (2003) shows that Shh from midline structures collaborates with netrin-1 to guide commissural axons. This new role for the morphogen Shh raises the possibility that principles similar to those used to establish positional information in embryonic patterning are also employed during axon navigation (reviewed in Salinas, 2003).

F. Steroids

Steroid hormones may also induce directed neurite outgrowth. Estrogen elicits a significant enhancement of neurite outgrowth and differentiation within organotypic explant cultures of hypothalamus, preoptic area, and cerebral cortex (Toran-Allerand, 1976, 1980, 1984). Forebrain neurons coexpress nerve growth factor (NGF) receptors and estrogen receptor mRNA (Miranda *et al.*, 1993; Toran-Allerand *et al.*, 1992), and NGF significantly increases nuclear estrogen binding in cortical but not basal forebrain explants (Miranda *et al.*, 1996). Steroid/neurotrophin interactions may stimulate the synthesis of proteins required for neuronal differentiation, survival, and maintenance of function (Toran-Allerand, 1996). Estrogen and neurotrophin may regulate the same broad array of cytoskeletal and growth-associated genes involved in neurite growth and differentiation (Singh *et al.*, 1999).

The estrogen receptor, a nuclear transcription factor, is widely expressed in the developing forebrain (Toran-Allerand, 2004). Two mammalian estrogen receptors have been described, ER- α (White *et al.*, 1987), mediating most of the transcriptional action of estrogen in the brain, and ER- β (Kuiper *et al.*, 1996; Tremblay *et al.*, 1997). Mice lacking ER- β exhibit a reduction in the expression of genes involved in neuronal migration and axonal guidance,

such as semaphorin G, syndecan 3, and reelin. Therefore, it appears that ER- β influences migration of neurons during development and neuronal survival throughout life (Wang *et al.*, 2002a, 2003).

The principal nucleus of the bed nuclei of the stria terminalis (BSTp) pathway to the anteroventral periventricular nucleus of the preoptic region (AVPV) develops in a sexually dimorphic pattern, suggesting a directed mechanism of axonal guidance (Hutton *et al.*, 1998). *In vitro*, addition of testosterone to BSTp to AVPV co-cultures induces neurite extension (Ibanez *et al.*, 2001). Testosterone induces a target-derived, diffusible chemotropic activity that results in a sexually dimorphic pattern of connectivity (reviewed in Simerly, 2002). The AVPV projects to the gonadotropin-releasing hormone (GnRH) neurons and the tuberoinfundibular dopaminergic (TIDA) neurons. Expression of ER- α by GnRH neurons (Skyner *et al.*, 1999) could suggest that estrogen may also direct the development of projections from the anteroventral periventricular nucleus to the GnRH neurons (Simerly, 2002).

The role of estrogen in branching has also been reported in invertebrates. Ecdysteroids, the insect steroids that trigger metamorphosis, control both regression and outgrowth *in vivo* and stimulate neuritic growth in cultured pupal leg motor neurons. Ecdysteroid enhances neuritic branching by altering growth cone structure and function, suggesting that hormonal modulation of cytoskeletal interactions contributes significantly to neuritic remodeling during metamorphosis (Matheson and Levine, 1999).

G. Intracellular Signaling Mechanisms

1. Ena/VASP

Ena/VASP proteins are a conserved family of actin regulatory proteins made up of Ena/VASP homology domain 1 (EVH1) and EVH2 domains and a proline-rich central region (Fig. 8). Mammalian Ena/VASP members are Mena, n-Mena, Ena/VASP like (EVL), and VASP. The members are 60–70% identical to each other. The Ena/VASP family is involved in Abl and/or cyclic nucleotide-dependent protein kinase signaling pathways. They have been implicated in actin-based processes such as fibroblast migration, axon guidance, and T-cell polarization and are important for the actin-based motility of the intracellular pathogen *Listeria monocytogenes* (Chakraborty *et al.*, 1995; Gerstel *et al.*, 1996; Gertler *et al.*, 1996; Pistor *et al.*, 1995; Smith *et al.*, 1996). Vertebrate Ena/VASP proteins are substrates for PKA/PKG serine/threonine kinases. Phosphorylation by these kinases appears to modulate Ena/VASP function within cells, although the mechanism underlying this regulation remains to be determined. Ena/VASP are also crucial factors in regulating actin dynamics and associated processes such as cell–cell adhesion.

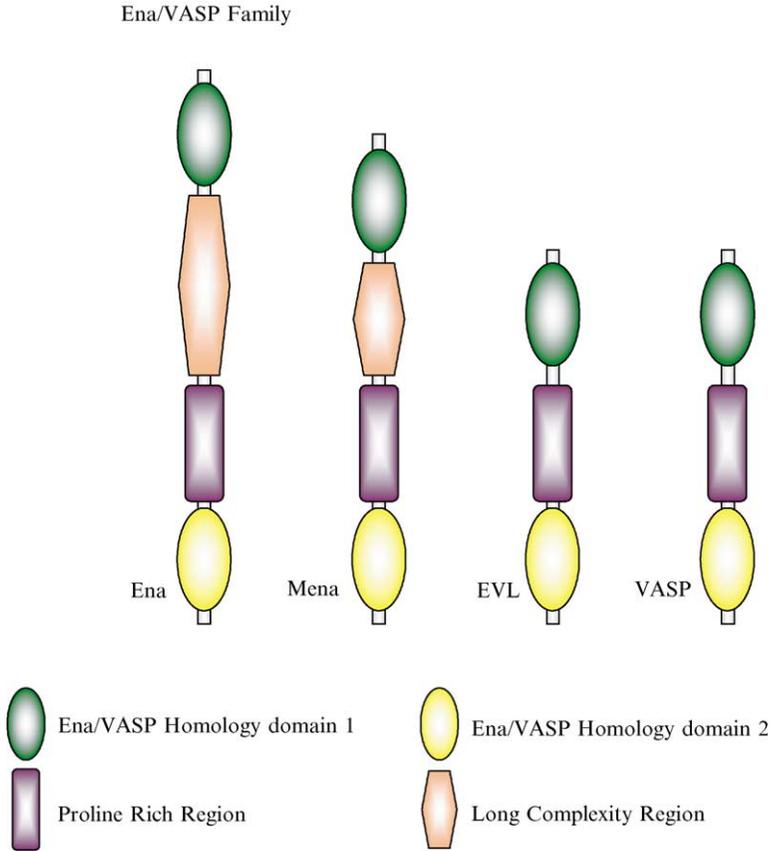


Figure 8 Ena/VASP.

Evidence also suggests that these have inhibitory functions in integrin regulation, cell motility, and axon guidance (Krause *et al.*, 2003).

Mena and Mena/VASP mouse mutants have defects in several major axonal tracts (Lanier *et al.*, 1999), including the corpus callosum, the hippocampal commissure, the anterior commissure, and the pontocerebellar pathway. Ena/VASP proteins are highly expressed in the developing cortical plate in cells bordering reelin-expressing Cajal-Retzius cells and in the intermediate zone. Inhibition of Ena/VASP function through retroviral injections *in utero* leads to the aberrant migration of early-born pyramidal neurons in the superficial layers of both the embryonic and the postnatal cortex in a cell-autonomous fashion. The results demonstrate that Ena/VASP proteins play a key role in regulating neuronal migration and layering within the developing mouse neocortex (Goh *et al.*, 2002).

2. Rho GTPases

Rac, Rho, and Cdc42 are small GTPases of the Rho family. They have been shown to regulate actin organization in non-neuronal cells (Hall, 1998), as well as cytoskeletal dynamics in neuronal growth cones (Luo *et al.*, 1997). A model has been proposed (Hall, 1998; Mueller, 1999) in which attractive guidance cues activate Rac and Cdc42 to promote growth cone advance, whereas repulsive guidance cues activate Rho to inhibit growth and induce retraction (Dickson, 2001). For example, ephrin-A5 activates Rho and inhibits Rac in cultures of retinal ganglion cells. Both Rho and its downstream effector Rho kinase are required for growth cone collapse (Wahl *et al.*, 2000). Rac proteins mediate axon guidance, outgrowth, and branching as well as suppress the formation of ectopic axon growth (Lundquist, 2003). Calcium-dependent regulation of Rho GTPases triggers turning of nerve growth cones (Jin *et al.*, 2005). The regulators of the Rho GTPases, GTPase-activating proteins and guanine exchange factors, play important roles in axon guidance. Cross GTPase-activating proteins (CrGAPs) are involved in Robo-mediated repulsive axon guidance. Too much or too little CrGAP activity leads to defects in Robo-mediated repulsion at the midline. CrGAP directly interacts with Robo both biochemically and genetically and acts as a GTPase-activating protein specifically for Rac to regulate midline crossing (Hu *et al.*, 2005).

H. Extracellular Matrix Molecules

The extracellular matrix (ECM) is an important source of extrinsic cues that influence the response of growth cones to guidance cues (Condic *et al.*, 1999; Diefenbach *et al.*, 2000; Hopker *et al.*, 1999; Nguyen-Ba-Charvet *et al.*, 2001). They can act to promote or inhibit neurite outgrowth and modulate the response of axons to particular guidance cues. Laminin, tenascin, collagen, fibronectin, and a number of proteoglycans have been implicated in modulating axonal outgrowth. For example, laminin can promote, while tenascin can inhibit, neurite extension. Receptors for ECM molecules include integrins as well as Ig family members. For example, Ig CAM F3 can function as a receptor for a type of tenascin. The laminin family and its receptors are one of the best-studied examples of ECM molecules with regard to neuronal development. The laminins are heterotrimers, in which different subunits combine to form at least 10 different isoforms with growth-promoting or -inhibiting effects depending on the cell type. The axonal receptors for the laminins are the integrins. Integrins are heterodimers whose subunit composition determines their laminin binding specificity. The integrins link the ECM signals to the cytoskeleton and various

signal transduction pathways. The exact role of laminins and other ECM molecules in neuronal development is not clear, although most evidence suggests a role in axonal guidance. Integrin signaling regulates cytoskeletal dynamics, adhesion, and migration events, through associated proteins such as talin, vinculin, integrin-linked kinase (ILK), focal adhesion kinase (FAK), paxillin, p130Cas, Abl kinase, and many other signaling or cytoskeletal proteins (Hynes, 2002).

In the developing nervous system, proteoglycans predominantly carry either chondroitin sulfate or heparan sulfate glycosaminoglycans (GAGs) (Bovolenta and Feraud-Espinosa, 2000). Heparan sulfate proteoglycans (HSPGs) are a group of extracellular and cell surface proteins essential for proper axonal pathfinding during nervous system development (Bulow and Hobert, 2004; Walz *et al.*, 1997; Wang and Denburg, 1992), and it is increasingly evident that the major mechanism by which HSPGs influence axon pathfinding is by regulating the function of axon guidance cues. HSPGs affect several axon guidance cues, including fibroblast growth factor (FGF), heparin-binding growth associated molecule (HB-GAM), Slits, and Anosmin/Kallman syndrome gene (KAL-1) (Bulow and Hobert, 2004; Hu, 2001; Inatani *et al.*, 2003; Irie *et al.*, 2002; Johnson *et al.*, 2004; Kinnunen *et al.*, 1998; Steigemann *et al.*, 2004; Walz *et al.*, 1997). Chondroitin sulfate proteoglycans (CSPGs) are a heterogeneous set of proteins bearing GAGs of the chondroitin sulfate class (Lander, 1998). The CSPGs are also ECM molecules involved in the regulation of axon growth as demonstrated by *in vitro* studies on CSPGs such as NG2 (Dou and Levine, 1994), neurocan, and phosphacan (Margolis *et al.*, 1996). They influence the behavior of neuronal growth cones during development and, importantly, following CNS injury (Bovolenta and Feraud-Espinosa, 2000; Morgenstern *et al.*, 2002). CSPGs are known to modulate the response of growth cones to other matrix components such as laminin (Condic *et al.*, 1999). This raises the possibility that CSPGs are components of the developmental environment capable of regulating how growth cones respond to surrounding guidance cues. The biological activity of CSPGs may also be determined by distinct proteins that bind to glycosaminoglycans and interact with receptors on the surface of neuronal growth cones (Anderson *et al.*, 1998; Brittis and Silver, 1994; Emerling and Lander, 1996; Golding *et al.*, 1999). Although CSPGs are known to interact with growth factors, adhesion molecules, and other matrix components, the specific binding proteins capable of mediating the effects of CSPGs on neuronal growth cones remain to be identified (Bovolenta and Feraud-Espinosa, 2000; Morgenstern *et al.*, 2002).

A recent study showed that the thrombospondin repeats of *Sema5A* physically interact with the glycosaminoglycan portion of both CSPGs and HSPGs. CSPGs function as bound localized extrinsic cues that convert *Sema5A* from an attractive to an inhibitory guidance cue. Therefore,

glycosaminoglycans provide a molecular mechanism for CSPG-mediated inhibition of axonal extension. Further, axonal HSPGs are required for Semaphorin 5A-mediated attraction, suggesting that HSPGs are components of functional Semaphorin 5A receptors. Therefore, the nature of a growth cone's response to Semaphorin 5A depends on the types of sulfated proteoglycans present in the developmental environment (Kantor *et al.*, 2004).

I. Adhesion Molecules

Several cell adhesion molecules (CAMs) of the Ig superfamily have also been implicated in regulating axon guidance at the midline, including mammalian L1CAM, NrCAM, and TAG-1 (mammalian ortholog of chick Axonin-1). The superfamily includes several subfamilies that are found in a number of tissues during development and in the adult (reviewed in Edelman and Crossin, 1991). Homologues of most of these have also been found in invertebrate animals. Each CAM has an extracellular region containing six Ig domains, as well as two (NrCAM), four (L1-CAM), or five (TAG-1) fibronectin type III extracellular domains (Walsh and Doherty, 1997) (Fig. 9). The cytoplasmic domain is highly conserved among individual members of the L1 subfamily and between invertebrate and vertebrate species (Hortsch, 1996). L1 is linked to the cytoskeleton through two regions in the cytoplasmic domain. An ankyrin binding site is located in the C terminus of the cytoplasmic domain (Davis and Bennett, 1994). CAMs,

Adhesion Molecules

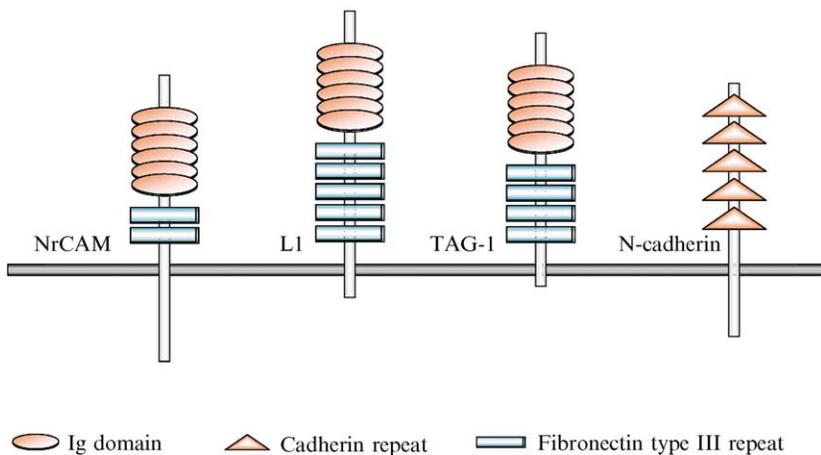


Figure 9 Adhesion molecules.

including N-CAM, L1, and N-cadherin, promote neurite outgrowth (Doherty *et al.*, 1989; Doherty *et al.*, 1990).

L1 is one of the most intensely studied adhesion molecules expressed in the developing central and peripheral nervous system (Kamiguchi *et al.*, 1998). L1 is important in neuronal migration, axon growth, guidance, fasciculation, and synaptic plasticity. L1 is also expressed in non-neuronal cells such as the immune system, kidney, pigment cells, and a variety of cancers. Commissural axons extending toward the ventral midline of the rodent spinal cord express TAG-1, but not L1, while axonal segments on the contralateral side of the floor plate express L1, but not TAG-1 (Dodd *et al.*, 1988; Imondi *et al.*, 2000; Tran and Phelps, 2000). Consistent with an altered-responsiveness mechanism, it has been postulated that the switch in expression from TAG-1 to L1, presumably triggered by contact with the floor plate, delays the rostral turn exhibited by commissural axons until after they cross the floor plate (Dodd *et al.*, 1988). NrCAM, another commissural axon-associated IgCAM, is expressed at low levels on commissural axons both as they extend toward the floor plate on the ipsilateral side of the midline and as they project in the longitudinal direction on the contralateral side of the floor plate in the chick and mouse spinal cord (Matise *et al.*, 1999; Stoeckli and Landmesser, 1995).

Knockouts of L1CAM (Cohen *et al.*, 1998; Dahme *et al.*, 1997; Fransen *et al.*, 1998) display a reduced corticospinal tract, an abnormal pyramidal decussation, a decreased axonal association with non-myelinating Schwann cells, ventricular dilatation, and hypoplasia of the cerebellar vermis. Demyanenko and colleagues (1999) reported abnormal morphogenesis of cortical dendrites, showing that pyramidal neurons in layer V exhibited undulating apical dendrites that did not reach layer I. They also found that L1 mutants had a smaller hippocampus with fewer pyramidal and granule cells (Demyanenko *et al.*, 1999) and an altered distribution of dopaminergic neurons in the brain of L1 null mice (Demyanenko *et al.*, 2001). There is also a reduced size of the corpus callosum because of the failure of many callosal axons to cross the midline, as well as the formation of other commissural tracts in the brain. L1 has been shown to interact with neuropilin via the first Ig domain (Castellani *et al.*, 2002) and to participate in Sema3a signaling (Castellani *et al.*, 2000) to mediate axonal repulsion.

V. Axon Guidance Mechanisms at the Midline of the Nervous System

A. Commissural Axons in Spinal Cord

The spinal cord has proven to be a useful system for identifying molecules that guide axons to their appropriate targets. During spinal cord development, commissural neurons, which differentiate in the dorsal neural tube,

send axons that project toward and subsequently across the floor plate, forming axon commissures (Colamarino and Tessier-Lavigne, 1995). These commissural axons project toward the midline in part because they are attracted by Netrin-1 (Kennedy *et al.*, 1994; Placzek *et al.*, 1990; Serafini *et al.*, 1996, 1994; Tessier-Lavigne *et al.*, 1988b). Once on the contralateral side, axons are no longer attracted but are repelled by Slit expressed by ventral midline cells (Brose and Tessier-Lavigne, 2000). This change in the response of axons is due to the silencing of Netrin-1 attraction by signaling through DCC interacting with the Slit receptor, Robo (Stein and Tessier-Lavigne, 2001). In *Netrin-1* or *DCC* mutant mice, many commissural axon trajectories fail to invade the ventral spinal cord and are misguided (Fazeli *et al.*, 1997; Serafini *et al.*, 1996). However, some of them do reach the midline, indicating that other guidance cues cooperate with Netrin-1 to guide these axons (Serafini *et al.*, 1996). Sonic hedgehog (Shh), a morphogen secreted by the floor plate, functions as a gradient signal for the generation of distinct classes of ventral neurons along the dorsoventral axis (Ingham and McMahon, 2001; Jessell, 2000; Marti and Bovolenta, 2002). Shh is an axonal chemoattractant that provides the Netrin-1-independent chemoattractant activity of the floor plate. Shh collaborates with Netrin-1 in commissural axon attraction *in vitro* and is required for normal guidance of these axons *in vivo* (Charron *et al.*, 2003). Several studies show that other members of the morphogen family, Wnt and BMP, are also involved in the spinal cord development. A directed source of Wnt4 protein attracts postcrossing commissural axons and Frizzled3 mutant mice display anterior-posterior guidance defects after midline crossing. This indicates that Wnt-Frizzled signaling guides commissural axons along the anterior-posterior axis of the spinal cord (Lyuksyutova *et al.*, 2003). Bmp7, Bmp6, and growth differentiation factor 7 (Gdf7) are expressed by the roof plate and are potential dorsal repellent cues for commissural axons (Augsburger *et al.*, 1999).

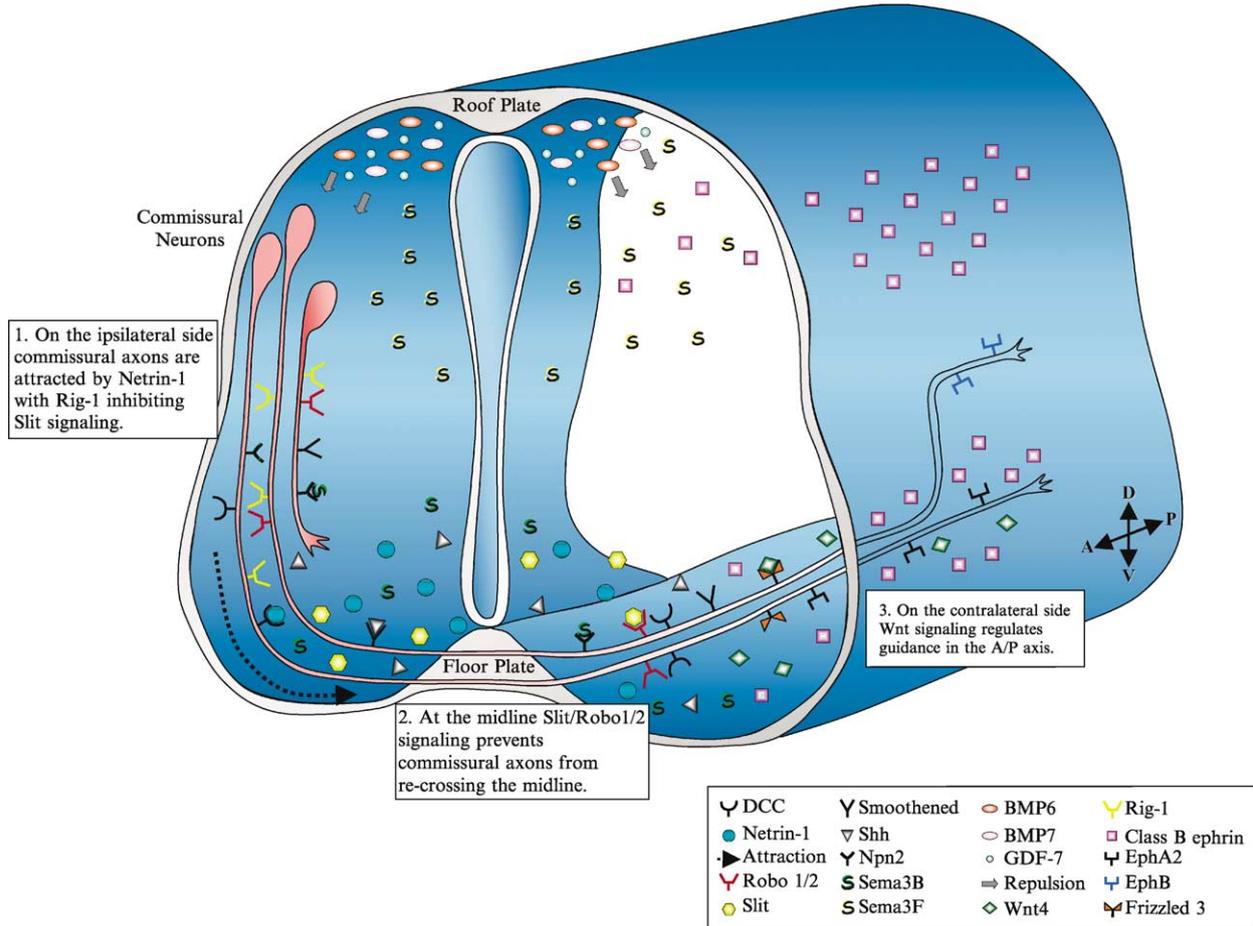
The *Slit* genes are also expressed in the floor plate at the ventral midline of the spinal cord, and *Robo1* and *2* are expressed in regions that include commissural neuron cell bodies (Brose *et al.*, 1999; Itoh *et al.*, 1998; Kidd *et al.*, 1998; Li *et al.*, 1999). *In vitro* commissural axons are repelled by Slit2 only after they have crossed the floor plate (Zou *et al.*, 2000). In *Slit1/Slit2* double mutant mice, although the formation of several major forebrain tracts (corticofugal, callosal, and the thalamocortical tracts) and the optic chiasm are defective (Bagri *et al.*, 2002; Plump *et al.*, 2002), no obvious commissural axon guidance defects were observed in the spinal cord (Plump *et al.*, 2002). Another member of the Slit family, Slit3, is also expressed by floor plate cells (Brose *et al.*, 1999), and analysis of triple Slit mutant revealed that many commissural axons stalled at the floor plate and failed to cross (Long *et al.*, 2004). This indicates that Slits contribute to the repulsion of axons away from the midline. Rig-1 is a divergent member of

the Robo family (Yuan *et al.*, 1999) that is highly expressed before midline crossing and is downregulated after crossing (Sabatier *et al.*, 2004). Rig-1 prevents commissural axons from sensing Slit in the floor plate through their cognate receptor Robo1 as they grow toward the floor plate, allowing them to enter and cross to the contralateral side. At the midline, the downregulation of Rig-1 protein expression helps the axons to sense the floor plate as a repulsive environment, thus preventing them from re-crossing the midline (Sabatier *et al.*, 2004). Commissural axons also express another receptor, Neuropilin-2, mediating the repulsive effects of Sema3B, found in the floor plate, and Sema3F, expressed widely in the spinal cord, except in the floor plate (Zou *et al.*, 2000; Fig. 10). Analysis of homozygous *neuropilin-2* mutant mice shows disorganized axons at the midline while crossing (Zou *et al.*, 2000), suggesting a role for neuropilin-2 in commissural axon pathfinding.

Once the axons have crossed the midline, they execute a rostral turn at the contralateral floor plate margin and extend for a short distance within the ventral funiculus, a longitudinal fiber tract that forms in close apposition to the floor plate (Bovolenta and Dodd, 1990). EphA2 and EphB expression is upregulated on contralateral commissural axons (Brittis *et al.*, 2002; Imondi *et al.*, 2000), and class B ephrins are expressed in the floor plate as well as in the dorsal part of the spinal cord. A subset of decussated commissural axons takes a more dorsal trajectory before turning at the border of class B ephrin expression, indicating a role in excluding these axons from the dorsal spinal cord (Imondi and Kaprielian, 2001; Fig. 10).

B. Guidance of Cortical Axons at the Midline

Contralateral cerebral cortical projections through the corpus callosum integrate sensory and motor information between the two brain hemispheres. In split-brain animals and in people whose corpus callosum has been severed, interhemispheric transfer of sensory and motor information is deficient (Gazzaniga, 1995). In these individuals, visual and tactile information presented to one hemisphere is not available for analysis by the other hemisphere. In addition, perceptual interactions between the two hemispheres are absent in these individuals. These observations, pioneered by Roger Sperry in the 1960s (Sperry, 1968, 1982), defined the critical roles of contralateral cortical projections in human consciousness and behavior. Data in humans and in mice suggest the possibility that different mechanisms may regulate the development of the corpus callosum across its rostro-caudal extent (Richards *et al.*, 2004). The complex developmental processes required for formation of the corpus callosum may provide some insight into why such a large number of human congenital syndromes are associated with agenesis of this structure (Richards *et al.*, 2004). Anatomical studies



have demonstrated that the majority of contralaterally projecting (callosal) neurons are located in layers 2/3 and 5 (Innocenti, 1986; Wise and Jones, 1976). In rodents callosal axons project to corresponding, homotopic areas in the contralateral cortex. During development, callosal axons grow ventrally to the intermediate zone (the future cortical white matter) and then turn medially, cross the midline, and re-enter the appropriate contralateral cortical area to form synapses with their targets (Fig. 11). The development of the corpus callosum depends on guidance by midline glial populations, and their expression of specific molecules, and fasciculation pioneering axons derived from neurons in the cingulate cortex (as described in Section III.A.2). Four midline populations at the corticoseptal boundary have been described: the glial wedge, the indusium griseum glia (Shu and Richards, 2001; Shu *et al.*, 2003c), the midline zipper glia (Silver *et al.*, 1993), and the subcallosal (glial) sling (Silver *et al.*, 1982) (see Section III.B). The sling is a glial fibrillary acidic protein (GFAP)-negative population of cells (in rodents) with neuronal properties (Shu *et al.*, 2003b) that migrates from the lateral sub-ventricular zone to underlie the developing corpus callosum (Silver *et al.*, 1982). Both ablation and rescue experiments (Silver and Ogawa, 1983; Silver *et al.*, 1982) have shown that the glial sling is required for the development of the corpus callosum. The indusium griseum glia and the midline zipper glia have many phenotypic and molecular characteristics in common, indicating that they may represent a common population of glia that becomes spatially distinct by the formation of the corpus callosum (Shu *et al.*, 2003c). The glial wedge is part of the radial glial scaffold (Shu *et al.*, 2003c) and, together with the indusium griseum glia, expresses Slit2. In the brain, unlike the spinal cord, Slit2 mediates both precrossing and postcrossing axonal guidance (Shu *et al.*, 2003d). Robo1 and Robo2 mRNAs are expressed in the neocortex during callosal axon targeting (Shu and Richards, 2001), and Robo proteins are expressed on callosal axons (Sundaresan *et al.*, 2004). In the Slit2 mutant, the corpus callosum fails to form. Instead, axons grow into large ectopic bundles of fibers on either side of the midline that resemble Probst bundles (Bagri *et al.*, 2002). Taken together, these data

Figure 10 Molecules involved in the guidance of commissural axons in the spinal cord. Commissural neurons send their axons toward and across the midline. Bmp6 and 7 and GDF-7 expressed by the roof plate act as dorsal repellents for commissural axons. Axons express DCC, Robo1/2, Smoothed, and Npn2. The axons are attracted by Netrin-1 and Shh but initially are not responsive to Slit because Rig-1 inhibits Slit/Robo1/2 signaling. Sema/Npn2 signaling is required to avoid inappropriate targeting. Once the axons cross the midline, they are then repelled by Slit, expressed by the floor plate, and lose their attraction to Netrin-1, allowing them to leave the floor plate and preventing them from re-crossing the midline. Wtn4/Frizzled3 signaling regulates guidance in the anterior-posterior axis. A subset of commissural axons expressing EphB takes a more dorsal trajectory but grows between regions of Class 3 ephrin-B expression both dorsally and ventrally.

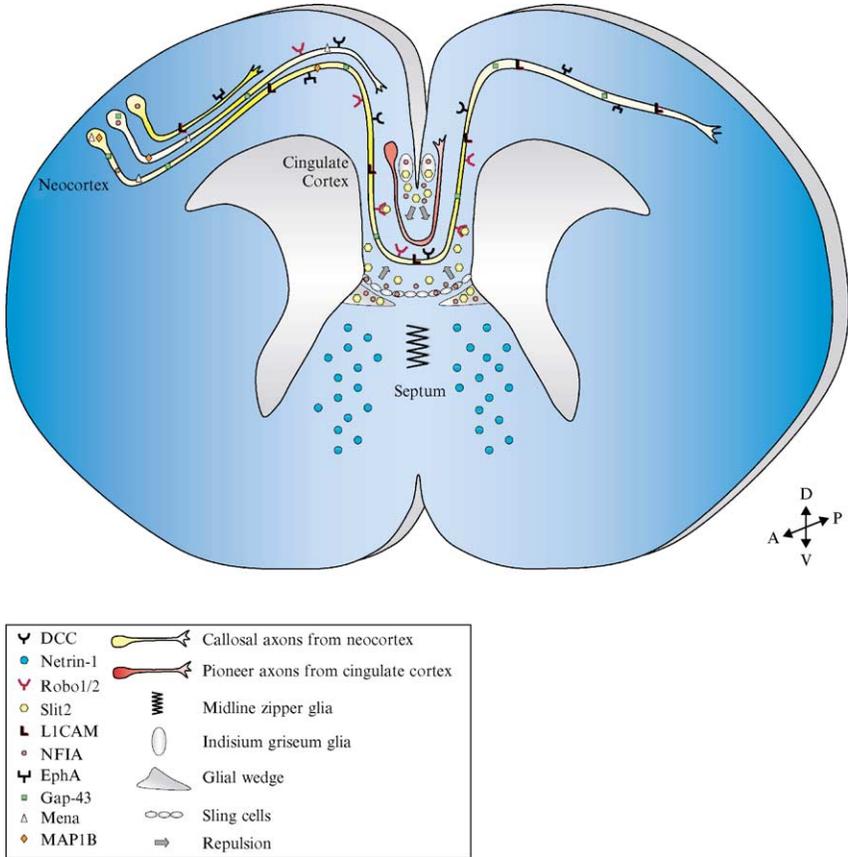


Figure 11 Molecules involved in the formation of the cortical commissural axons. Axons from the cingulate cortex pioneer a path across the midline. They are followed by neocortical callosal axons that probably fasciculate with them to cross the midline. Slit2, expressed by the glial wedge and the indisium griseum glia, provides a surround repulsion mechanism that keeps callosal axons within the tract, causing them to turn and cross the midline and preventing them from entering the septum. After axons have crossed the midline, Slit2 then repels the postcrossing axons away from the midline area. Callosal axons express several different molecules, including DCC, Robo1/2, EphA, NFIA, Gap-43, L1, Mena, and MAP1B. Netrin-1 is expressed within the septum under the corpus callosum; thus, unlike commissural axons in the spinal cord, callosal axons do not grow through the region of Slit and Netrin expression. NFIA is present in the sling cells, glial wedge, and the indisium griseum. Mouse mutants for DCC, Netrin-1, EphA, NFIA, Gap-43, L1, Mena, and MAP1B have defects in callosal formation (see Section V.B for more details) but how these genes regulate callosal axon pathfinding has not yet been elucidated.

suggest an important role for Slit/Robo signaling in corpus callosum formation.

In the mammalian brain, Netrin-1 mutants and DCC knockout mice do not develop a corpus callosum or a hippocampal commissure and have a greatly reduced or absent anterior commissure (Fazeli *et al.*, 1997; Serafini *et al.*, 1996). In the developing forebrain, Netrin-1 has been shown to attract laterally directed cortical axons *in vitro* (Metin *et al.*, 1997; Richards *et al.*, 1997), indicating that these molecules are important for additional axonal guidance systems in the forebrain other than commissural axon guidance. Netrin-1 acts *in vitro* as an attractant and growth promoter for dorsal thalamic axons and is required for the proper development of the thalamo-cortical axon (TCA) projection *in vivo* (Braisted *et al.*, 2000). DCC protein is expressed predominantly in large fiber tracts such as the lateral olfactory tract, the internal capsule, the corpus callosum, the anterior commissure, the fimbria/fornix, the fasciculus retroflexus, and the stria medularis (Shu *et al.*, 2000), as well as in dorsal thalamus (Braisted *et al.*, 2000). DCC knockout mice have defects in multiple commissures, including the corpus callosum, the hippocampal commissure, and the anterior commissure (Fazeli *et al.*, 1997). The basilar pons is absent in both Netrin-1 and DCC knockout mice (Fazeli *et al.*, 1997; Yee *et al.*, 1999).

The nuclear factor I (Nfi) family of transcription factors regulates both adenoviral DNA replication and viral and cellular gene expression, including the control of olfactory-specific genes (Baumeister *et al.*, 1999; Behrens *et al.*, 2000; Gronostajski *et al.*, 1985; Hennighausen *et al.*, 1985; Leegwater *et al.*, 1985; Nagata *et al.*, 1982, 1983; Nowock *et al.*, 1985). The Nfi family is made up of four members, Nfia, Nfib, Nfic, and Nfix. The Nfia mutant exhibits both agenesis of the corpus callosum and a reduction in GFAP expression (das Neves *et al.*, 1999). Nfia is expressed in midline glial structures, and the development of these structures is severely impaired in Nfia mutant mice. These data indicate that Nfia regulates both commissural development and the development of midline glia (Shu *et al.*, 2003a). Nfib mutant mice also display agenesis of the corpus callosum and abnormalities in midline glial development, as well as enlargement of the lateral ventricles (Steele-Perkins *et al.*, 2005).

In knockouts of the L1 gene (Cohen *et al.*, 1998; Dahme *et al.*, 1997; Fransen *et al.*, 1998) the corpus callosum failed to form properly due to the failure of many callosal axons to cross the midline (Demyanenko *et al.*, 1999, 2001). These findings suggest a variety of biological roles for L1 that are critical in brain development in different brain regions. Nr-CAM, a member of the L1 subfamily of cell adhesion molecules, is not expressed on callosal axons until postnatal day zero (P0), suggesting that Nr-CAM may be involved in the later stages of axonal growth or tract maintenance (Lustig *et al.*, 2001).

Mice lacking p35, an activator of cdk5 in the CNS, exhibit defects in a variety of CNS structures, most prominently characterized by a disruption in the laminar structure of the neocortex (Chae *et al.*, 1997). In these mutant mice, the corpus callosum appears bundled at the midline, but dispersed lateral to the midline. After crossing the midline, cortical axons defasciculate prematurely from the corpus callosum and take similarly oblique paths through the cortex. These results suggest that defective axonal fasciculation and guidance may be primary responses to the loss of p35 in the cortex (Kwon *et al.*, 1999).

In EphA5 mutant mice, callosal axons failed to grow into the corpus callosum, indicating that the EphA receptors and their ligands, the A-ephrins, play critical roles in the development of callosal axon projection to their contralateral targets (Hu *et al.*, 2003). EphA4 is expressed in the developing corpus callosum, with an interesting differential expression of EphA4 within different parts of the corpus callosum. In rostral regions, the entire corpus callosum was EphA4 positive whereas more caudally (around the hippocampal commissure) EphA4 was restricted to the most dorsal part of the corpus callosum (Greferath *et al.*, 2002). EphB2 (Nuk) and EphB3 (Sek4) mutant mice have been described (Henkemeyer *et al.*, 1996; Orioli *et al.*, 1996). EphB2 and EphB3 are members of the Eph-related family of receptor protein-tyrosine kinases. These receptors interact with a set of cell surface ligands that have recently been implicated in axon guidance and fasciculation. Whereas mice deficient in EphB2 exhibit defects in pathfinding of anterior commissure axons, EphB3 mutants have defects in corpus callosum formation. The phenotype in both axon tracts is markedly more severe in EphB2/EphB3 double mutants, indicating that the two receptors act in a partially redundant fashion (Orioli *et al.*, 1996).

MAP1B, a microtubule-associated protein, is expressed in axons, dendrites, and growth cones throughout the CNS during development. MAP1B is implicated in the crosstalk between microtubules and actin filaments. Homozygous MAP1B mutant mice display agenesis of the corpus callosum (Meixner *et al.*, 2000). A recent study showed that MAP1B phosphorylation is controlled by Netrin-1 (Del Rio *et al.*, 2004). Map1B mutant mice have severe abnormalities, similar to those described in netrin-1-deficient mice, in axonal tracts and in the pontine nuclei. These data indicate MAP1B may be a downstream effector in the Netrin-1-signaling pathway (Del Rio *et al.*, 2004).

A number of other genes are associated with agenesis of the corpus callosum in mice (reviewed in Richards *et al.*, 2004). The most common phenotype observed when cortical axons fail to reach the midline was that the axons do not stop growing but instead form swirled ipsilateral bundles of axons, called Probst bundles. Probst bundles form in mutants such as *Vax-1* (Bertuzzi *et al.*, 1999), *Gap-43* (Shen *et al.*, 2002), or heparan sulfate (Inatani

et al., 2003), indicating that these genes may regulate callosal axon guidance at the midline. However, in *GAP-43* mutant mice, callosal axons respond normally to Slit-2, although glial abnormalities may contribute to the phenotype (Shen *et al.*, 2002).

C. Guidance of Retinal Ganglion Cell Axons at the Optic Chiasm

In animals with binocular vision (such as mammals), retinal ganglion cell (RGC) axons originating from the nasal retina cross the midline to project into the contralateral optic tract, while a population of RGC axons from the temporal retina do not cross, but project away from the midline into the ipsilateral optic tract (Fig. 12). In mouse, ipsilaterally projecting RGCs are found in the ventro-temporal crescent of the retina, whereas contralaterally projecting RGCs are found throughout the retina (Guillery *et al.*, 1995; Mason and Sretavan, 1997; Sretavan, 1993).

After retinal ganglion cell axons exit each eye at the optic nerve head, forming the optic nerve, they traverse the ventral diencephalon toward the midline. Axons from the two eyes cross over each other to form the chiasm (X shape). The proportion of uncrossed to crossed retinal fibers varies across species. The ipsilateral projection represents about 40% of all RGCs in humans (Kandel *et al.*, 2000), less than 15% in ferrets (Cucchiari, 1991; Thompson and Morgan, 1993), and about 3–5% in mice (Rice *et al.*, 1995). Adult birds and fish do not have an ipsilateral projection (O’Leary *et al.*, 1983) and thus lack binocular vision.

Neuronal cells within the chiasm are postulated to provide guidance cues for the earliest axons (Marcus and Mason, 1995; Sretavan *et al.*, 1994). When these neurons are destroyed by complement-mediated cytolysis, the growth of all axons entering the chiasm is halted (Sretavan *et al.*, 1995). Later axons, however, derive guidance cues from a midline palisade of radial glia. The cells in this region provide a generalized negative signal, as the growth cones of both crossed and uncrossed axons pause when entering this region *in vivo* (Godement, 1994) and *in vitro* (Mason and Wang, 1997; Wang *et al.*, 1996). After the optic chiasm, axons then continue through the optic tract to their targets: the superior colliculus in mammals or the optic tectum in fish, frogs, or birds. The main visual nuclei that receive retinal input are the lateral geniculate nucleus (LGN), the superior colliculus (SC), and the pretectal nuclei (Wassle, 1982). They also project to the superchiasmatic nucleus and the accessory optic system (Zhang and Hoffmann, 1993). Retinal axons express DCC (as well as the repulsive receptor UNC-5), but Netrin is absent from the chiasm region (Anderson and Holt, 2002; Deiner and Sretavan, 1999; Shewan *et al.*, 2002). Instead, Netrin is expressed at the optic nerve head, and in Netrin or DCC mutants RGC axons fail to exit this

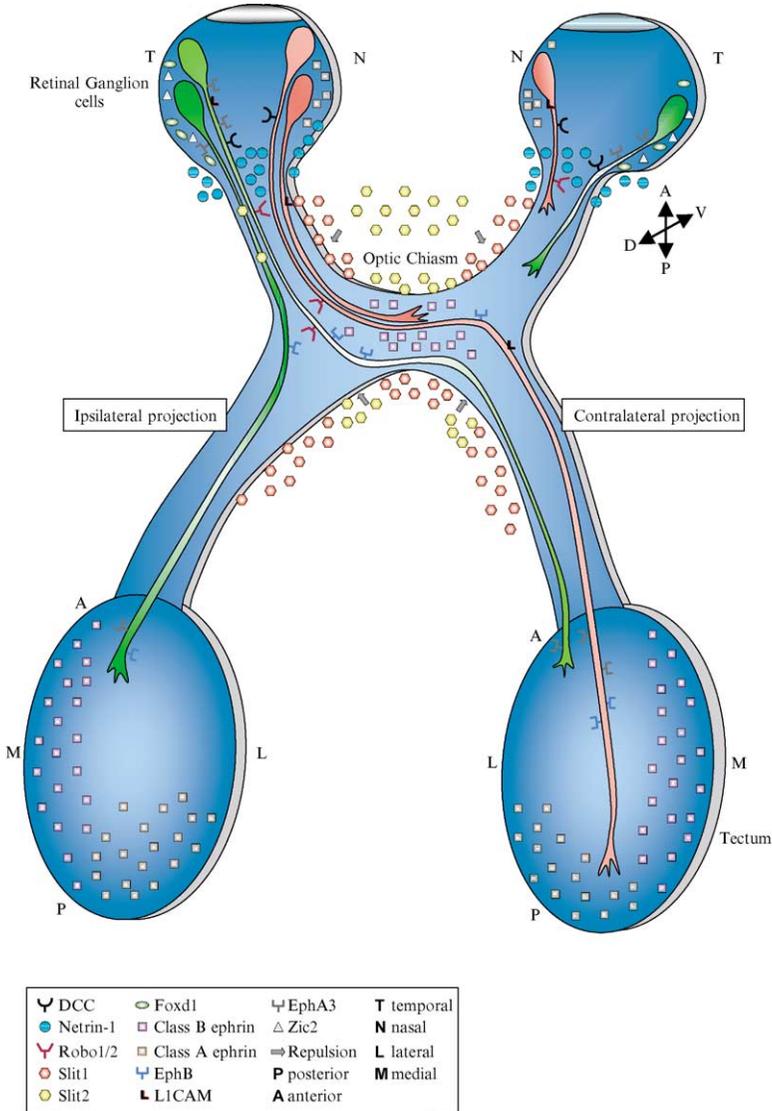


Figure 12 Molecules involved in the guidance of RGC axons to the tectum. The majority of RGC axons projects into the contralateral optic tract, while a small population of temporal RGC axons does not cross and projects ipsilaterally in mice. A diverse group of molecules acts to guide the RGC axons toward their final target in the tectum. Netrin-1 is expressed in the optic nerve head and Slits are expressed in the optic chiasm region, channeling the axons through the tectum and determining the position of the chiasm. RGC axons express DCC, Robo1/2, L1, and EphB and, as they enter the tectum, they express EphA3. Class B ephrins are expressed in the medial part of the tectum and mediate dorsoventral targeting, whereas Class A ephrins are expressed in the posterior tectum and mediate anterior-posterior topographic targeting.

region (Deiner *et al.*, 1997). In the retina, *robo2* is expressed by RGCs before any axons have reached the ventral midline of the diencephalon and continues to be strongly expressed during later stages of development. In contrast, *robo1* is not detected until after a number of axons have started to cross the midline and then only in a subset of cells. This suggests that Robo2 is likely to be the principle receptor in RGC axon guidance (Erskine *et al.*, 2000). Two slit genes are expressed around the chiasm: Slit1 and Slit2 in mouse, Slit2 and Slit3 in zebrafish. Furthermore, Slit1 and Slit2 both repel retinal axons in culture (Erskine *et al.*, 2000; Niclou *et al.*, 2000; Ringstedt *et al.*, 2000). As shown by experiments in zebrafish and mouse, Robo/Slit signaling controls the formation of the optic chiasm (Fricke *et al.*, 2001; Hutson and Chien, 2002; Plump *et al.*, 2002). Zebrafish mutant for the *astray/robo2* receptor show multiple guidance errors near the chiasm, including ipsilateral, anterior, and retinorectal projections. Slit1 or Slit2 single knockout mice have little or no retinal projection phenotype, presumably because of the partial overlap in the expression of the two genes. However, Slit1 and Slit2 double knockouts show a strong phenotype, most notably the expansion of the chiasm more anteriorly. Developmental and time-lapse analyses in zebrafish embryos show that, even in wild-type animals, RGC growth cones occasionally leave their pathway and misproject, but these errors are quickly corrected (Hutson and Chien, 2002). In *astray* mutants, many more errors occur and persist. Thus, Slit/Robo signaling has two functions: preventing errors in the first place and correcting them if they do occur (Rasband *et al.*, 2003).

Optic nerve fibers grow out from the eye in fasciculated bundles, with contact between them mediated in part by membrane glycoproteins. The first of these glycoproteins to be identified was neural cell adhesion molecule (NCAM) (Jessell, 1988). NCAM plays a crucial role in axonal fasciculation and substrate adhesion. The localization of the cell adhesion molecules L1, NCAM, and myelin-associated glycoprotein (MAG) was studied in the developing and adult mouse optic nerve and retina. At all stages of development, NCAM is expressed by fasciculating axons, growth cones, and their contact sites with glial cells, and contacts between glial cells. MAG is first associated with the endoplasmic reticulum and Golgi apparatus in oligodendrocytes and then moves at cell surface. L1 is expressed by optic axons that are grouped into fascicular bundles and by their growth cones, but not by the growth cones that contact glial cells (Bartsch *et al.*, 1989). L1 was differentially expressed on unmyelinated axons, but absent on myelinated axons. This finding supports the notion that L1 may be involved in the stabilization of axonal fascicles but not of axon-myelin contacts (Bartsch *et al.*, 1989). Recent reports have shown that the absence of L1 in the optic pathway in L1 knockout mice has no obvious effect on the development pattern of axon divergence in the chiasm (Cohen *et al.*, 1998; Demyanenko and Maness, 2003). L1 and the polysialic acid-associated form of NCAM

(PSA-NCAM) are dynamically expressed in a regionally specific pattern in the retinotectal pathway (Chung *et al.*, 2004). At the chiasm, the level of L1 expression is high, whereas that of PSA-NCAM is low. However, within the tract, intense expression of both molecules is found predominantly on axons from the dorsal but not ventral retina. These changes are observed when axons arrive at the junction of the chiasm and the optic tract, indicating a site-specific switch in expression of cell adhesion molecules on the optic axons. Moreover a population of PSA-NCAM-rich cells also projects axons to the TPOC. These results suggest that CAM expression on the optic axons may control formation of the partial retinotopic axon order within the optic tract (Chung *et al.*, 2004).

CSPGs provide an unfavorable environment for axon growth and have been implicated in the changing patterns of fiber order in the developing retinotectal pathway (Brittis and Silver, 1995). In mouse, CSPGs are expressed by an early population of neurons in the ventral diencephalon, and enzymatic removal of the chondroitin moieties affects retinal ganglion cell guidance (Chung *et al.*, 2000a,b). CSPGs are expressed in the retina as well (Chung *et al.*, 2000a) and are also important for maintaining fiber order as axons approach the chiasm (Leung *et al.*, 2003).

Heparan sulfates are also involved in the formation of the optic chiasm (Inatani *et al.*, 2003). Analysis of mutant mice for EXT1, an enzyme indispensable for HS synthesis, revealed that retinal axons projected ectopically into the contralateral optic nerve, similar to Slit1/Slit2 double-knockout mice (Plump *et al.*, 2002).

B-class ephrins are required for the sorting of axons at the optic chiasm. In *Xenopus laevis*, ephrin-B2 is present at the chiasm coincident with the formation of the uncrossed component at metamorphosis, and premature misexpression of ephrin-B2 in the ventral diencephalon induced an ectopic ipsilateral projection (Nakamura *et al.*, 2000). EphB receptors are expressed in the retina (Birgbauer *et al.*, 2000; Braisted *et al.*, 1997; Hindges *et al.*, 2002; Mann *et al.*, 2002b; Williams *et al.*, 2003). A receptor for ephrin-B2, EphB1, is found exclusively in regions of retina that give rise to the ipsilateral projection. EphB1 null mice exhibit a dramatic reduction in the ipsilateral projection, suggesting that this receptor contributes to the formation of the ipsilateral retinal projection, most likely through its repulsive interaction with ephrin-B2 (Williams *et al.*, 2003).

Several regulatory genes expressed in the developing retina have been reported to play a role in retinal axon guidance at the optic chiasm. Mice lacking *Vax1*, *Vax2*, *Pax2*, or *Brn3b* exhibit different defects in retinal axon pathfinding at the chiasm (Barbieri *et al.*, 2002; Bertuzzi *et al.*, 1999; Mui *et al.*, 2002; Torres *et al.*, 1996; Wang *et al.*, 2002b). It has also been shown that mutual regulation of *Pax-2* and *Shh* are important for the formation of the chiasm region (Alvarez-Bolado *et al.*, 1997). Overexpression of *Pax-2* affects

axon navigation through the chiasm and in the ascending postchiasmatic pathway, including the optic tract up to the tectum (Thanos *et al.*, 2004).

The zinc finger transcription factor *Zic2* is expressed in retinal ganglion cells (Fig. 12). Loss- and gain-of-function analyses indicate that *Zic2* is necessary and sufficient to regulate RGC axon repulsion by cues at the optic chiasm and to determine the ipsilateral projection (Herrera *et al.*, 2003).

The winged helix transcription factor *Foxd1* (previously known as BF-2, brain factor 2) is expressed in the ventrotemporal retina, as well as in the ventral diencephalon during the formation of the optic chiasm. Both retinal development and chiasm morphogenesis are disrupted in embryos lacking *Foxd1*. In the *Foxd1*-deficient retina, proteins designating the ipsilateral projection, such as *Zic2* and *EphB1*, are missing. In addition, in the *Foxd1*-deficient ventral diencephalon, *Foxg1* expression invades the *Foxd1* domain, *Zic2* and *Islet1* expression are minimized, and *Slit2* expression prematurely expands, changes that could contribute to axon projection errors. *Foxd1* plays a dual role in the establishment of the binocular visual pathways: first, in specification of the ventrotemporal retina, acting upstream of proteins directing the ipsilateral pathway, and second, in the patterning of the developing ventral diencephalon where the optic chiasm forms (Herrera *et al.*, 2004).

After forming the optic chiasm, RGC axons extend laterally and dorsally to establish the optic tract along the lateral wall of the diencephalon to reach visual targets in the thalamus and midbrain.

VI. Axon Guidance in the Retinotectal System

The retinotectal projection is one of the best-studied model systems for the examining of the mechanisms regulating precise topographic connectivity in the embryonic CNS. In 1963, Roger Sperry showed that after optic nerve section, retinal ganglion cell axons regenerate to their normal topographic positions in the tectum regardless of whether the entire retina is present (Attardi and Sperry, 1963). He proposed the chemoaffinity hypothesis: that each point in the target area has a unique molecular address determined by a specific distribution of cell surface molecules (Sperry, 1963). The retinotectal projection from the temporal retina is connected to the anterior tectum and the retinotectal projection from the nasal retina is connected to the posterior tectum (Fig. 12). Dorsal retina is connected to the dorsal tectum and ventral retina is connected to the ventral tectum (Holt and Harris, 1993; Mey and Thanos, 1992; Thanos and Mey, 2001; van Horck *et al.*, 2004). Time-lapse analyses of the retinotectal projection have demonstrated that retinal ganglion cell axons grow rapidly within the optic tract but move slowly after reaching the tectum, suggesting the presence of target-derived cues that suppress axonal growth (Harris *et al.*, 1987).

A combination of *in vitro* and genetic studies has revealed that the mapping of RGC axons along the antero-posterior axis of the optic tectum is controlled in large part by the matched gradients of EphA receptor tyrosine kinases in the retina and the GPI-linked ephrin ligands in the tectum (McLaughlin *et al.*, 2003a). Activation of EphA by ligand binding leads to axon repulsion (Drescher *et al.*, 1997; Flanagan and Vanderhaeghen, 1998) and inhibition of axon branching (Yates *et al.*, 2001). Ephrin-A2 and ephrin-A5 are expressed in overlapping gradients in the tectum, while the corresponding receptors are expressed in complementary gradient across the retina (Cheng *et al.*, 1995). Axons from temporal retinal ganglion cells, where EphA expression is high, are inhibited from innervating the caudal tectum, where ephrin-A levels peak (Ruthazer and Cline, 2004). Ephrin-As control the temporal-nasal mapping of the retina in the optic tectum/superior colliculus by regulating the topographically specific interstitial branching of retinal axons along the anterior-posterior tectal axis. This branching is mediated by relative levels of EphA receptor repellent signaling (McLaughlin *et al.*, 2003a). In ephrin-A2/A5 double knockout mice the retinotectal map is severely disrupted (Feldheim *et al.*, 2000), indicating complementary functions of these two molecules in retinotectal mapping.

Members of the EphB family of receptor tyrosine kinases and their transmembrane ligands ephrin-Bs have been implicated in dorsoventral patterning of the vertebrate retinotectal projection (Flanagan and Vanderhaeghen, 1998). EphB/ephrin-B interactions mediate axon attraction (Hindges *et al.*, 2002; Mann *et al.*, 2002a). It has been shown in mice that EphB in ventral retina contributes to axon targeting to the medial part of the superior colliculus, where ephrin-B levels are high (Hindges *et al.*, 2002). In *Xenopus*, high levels of ephrin-B in dorsal RGCs guide axons to the EphB-rich ventral tectum (Mann *et al.*, 2002a). EphB/ephrin-B interactions are known to result in bidirectional signaling, characterized by signaling into cells expressing EphB receptors (i.e., forward signaling) and into cells expressing ephrin-B (i.e., reverse signaling) (Holland *et al.*, 1996). Both reverse and forward ephrin-B/EphB signaling are involved in regulating dorso-ventral topography (Hindges *et al.*, 2002; Mann *et al.*, 2002a; Pittman and Chien, 2002). EphB receptor forward signaling and ephrin-B reverse signaling mediate axon attraction to control dorsal-ventral retinal mapping along the lateral-medial tectal axis (McLaughlin *et al.*, 2003b).

VII. Axon Guidance in the Olfactory System

The sense of smell is a primal sense for humans as well as animals. For both humans and animals, it is an important means with which we sense our environment. Compared to other sensory systems, the olfactory system is

unique in that new olfactory receptor neurons are continuously produced throughout life (Graziadei and Monti Graziadei, 1978). This process is essential for the replacement of mature olfactory receptor neurons that have a limited life span. Axons derived from newly differentiating olfactory receptor neurons in the olfactory epithelium project to glomeruli in the main olfactory bulb, where they synapse on the dendrites of mitral and tufted cells, whose neurons then send axons to the primary olfactory cortex (Shiple and Ennis, 1996). Therefore, all new olfactory axons exiting the olfactory epithelium, in embryos and adults, must be guided to their specific glomerular targets in the olfactory bulb. The precise olfactory axon targeting depends on the combination of guidance cues they encounter along their pathway.

A. Guidance Cues in the Main Olfactory System

The main olfactory system is formed by the olfactory epithelium, the olfactory nerve, and the olfactory bulbs. Olfactory sensory neurons located in the epithelium of the principal nasal cavity project their axons to the main olfactory bulb (MOB), where they synapse with second-order neurons (mitral and tufted cells) within specialized compartments of neuropil called glomeruli (reviewed in Dulac, 2000; Halpern, 1987). In rodents, axons extending from the olfactory epithelium grow initially through mesenchyme rich in NCAM (Croucher and Tickle, 1989; Key and Akeson, 1990; Miragall *et al.*, 1989), retinoic acid, FGF-8 (LaMantia *et al.*, 2000, 1993), L1, laminin, and HSPG and CSPG (Gong and Shipley, 1996; Raabe *et al.*, 1997; Treloar *et al.*, 1996). Several additional candidate guidance factors have been identified in the developing and mature olfactory system, but functional evidence for their involvement in the precise connectivity of olfactory sensory neurons is limited. Directional outgrowth toward distinct regions in the developing bulb involves a variety of signaling molecules, including semaphorins and neuropilins (Kobayashi *et al.*, 1997; Pasterkamp *et al.*, 1998, 1999; Renzi *et al.*, 2000; Schwarting *et al.*, 2000; Walz *et al.*, 2002; Williams-Hogarth *et al.*, 2000), adhesion molecules (Kafitz and Greer, 1998; Puche and Key, 1996; Yoshihara *et al.*, 1997), and Eph/ephrins (Maisonpierre *et al.*, 1993; St John and Key, 2001; St John *et al.*, 2000, 2002; Zhang *et al.*, 1996). Olfactory sensory neurons expressing a given odorant receptor project with precision to specific glomeruli in the olfactory bulb, generating a topographic map. This implies a tight linkage between the choice of a specific odorant receptor and the choice of a glomerular target. Mombaerts *et al.* (1996) developed a genetic approach to visualize these projections and examine the relationship between receptor expression and axon targeting. They observed that neurons expressing P2, an olfactory receptor gene, project with precision to two of

the 1800 glomeruli within the olfactory bulb. These data provide direct support for a model in which a topographic map of the receptor topographically fixed glomeruli in the mouse olfactory bulb. Substitution of the olfactory receptor gene leads to the conclusion that the olfactory receptor may be one determinant in the guidance process (Mombaerts *et al.*, 1996; reviewed in Mombaerts, 1996). Neurons expressing different odorant receptors express different levels of ephrin-A protein on their axons. Moreover, alterations in the level of ephrin-A alter the glomerular map. Deletion of the ephrin-A5 and ephrin-A3 genes posteriorizes the glomerular locations for neurons expressing either the P2 or SR1 receptor, whereas overexpression of ephrin-A5 in P2 neurons results in an anterior shift in their glomeruli. Thus the ephrin-As are differentially expressed in distinct subpopulations of neurons and are likely to participate, along with the odorant receptors, in governing the targeting of like axons to precise locations in the olfactory bulb (Cutforth *et al.*, 2003).

Expression of dominant-negative neuropilin-1 protein causes chick olfactory neurons to enter the telencephalon prematurely (Renzi *et al.*, 2000). Mutations in galectin-1, Sema3A, neuropilin-2, p75NTR, CHL1, or N-CAM-180 (Montag-Sallaz *et al.*, 2002; Puche *et al.*, 1996; Schwarting *et al.*, 2000; Tisay *et al.*, 2000; Treloar *et al.*, 1997; Walz *et al.*, 2002) result in subtle perturbations in the overall pattern of olfactory axon projections; however, the role of these molecules in the precision of glomerular targeting has not been examined.

Netrin-1 expression has also been observed within the olfactory epithelium, and its receptor DCC was detected in cells under the neuroepithelium (Livesey and Hunt, 1997). DCC has also been found in the embryonic rodent olfactory bulb, on the granule cells, and on pioneer axons from output neurons (Gad *et al.*, 1997; Shu *et al.*, 2000). A detailed analysis of the spatio-temporal expression patterns of both Netrin-1 and DCC proteins in the developing rat olfactory system showed that the association of Netrin-1 expression near DCC-expressing olfactory axons is restricted to the initial period of olfactory nerve pathfinding, which suggests that Netrin-1 may play a role in the directed outgrowth of the nascent olfactory axons toward the telencephalon (Astic *et al.*, 2002). Another member of the netrin family, Netrin-4, is localized within the lateral olfactory tract and may be involved in promoting neurite elongation (Koch *et al.*, 2000).

Diffusible factors secreted from the bulb that attract axons of olfactory sensory neurons have been proposed in the past (Ressler *et al.*, 1994; Vassar *et al.*, 1994), and putative guidance molecules in olfactory tissue have been identified (Kafitz and Greer, 1997, 1998; Key and Akeson, 1990; Puche and Key, 1996; Schwarting *et al.*, 2000; St John *et al.*, 2000; Tisay and Key, 1999; Yoshihara *et al.*, 1997), but their direct involvement in axon outgrowth and pathfinding is still unknown. In addition, *in vitro* experiments suggest that a

diffusible signal of unknown identity from the olfactory bulb may also direct receptor axons toward the bulb (Goetze *et al.*, 2002).

As the axons grow toward and into the rostral telencephalon, they encounter a variety of ECM molecules, some of which are associated with both axons and glial cells. These include NCAM, L1, laminin, OCAM (olfactory cell adhesion molecule), chondroitin sulfate proteoglycan (Gong and Shipley, 1996; Miragall and Dermietzel, 1992; Treloar *et al.*, 1996; Yoshihara *et al.*, 1997), and galectin-1 (Crandall *et al.*, 2000).

B. Guidance Cues in the Accessory Olfactory System

In many mammalian species, a second olfactory system, the vomeronasal system, is thought to be specialized in the perception of stimuli related to social and reproductive behaviors (Halpern, 1987; Keverne, 1999). The sensory neurons reside within an oblong-shaped structure, the vomeronasal organ (VNO), situated at the rostral end of the nasal cavity. The vomeronasal receptor neurons are bipolar neurons with a dendrite reaching the surface of the epithelium and an axon projecting through the vomeronasal nerves to the accessory olfactory bulb (AOB). The AOB is an oval structure situated on the dorso-posterior surface of the main olfactory bulb. The incoming vomeronasal nerve axons make synaptic contact with dendrites of AOB neurons in specialized structures called glomeruli (Farbman, 1992). Sensory neurons located in the apical portion of the vomeronasal organ innervate glomeruli restricted to the anterior region of the AOB. In contrast, neurons of the basal region of the VNO project their axons to glomeruli located in the posterior half of the AOB (Jia and Halpern, 1996). The molecular mechanisms that orchestrate the segregation of vomeronasal projections to the anterior and posterior halves of the AOB are just beginning to be understood. Two independent families of vomeronasal receptor genes have been characterized, the V1Rs and V2Rs, which encode seven-transmembrane domain proteins thought to represent the mammalian pheromone receptors (Dulac and Axel, 1995; Herrada and Dulac, 1997; Matsunami and Buck, 1997; Ryba and Tirindelli, 1997). The V1R and the V2R receptor families are expressed by two spatially segregated populations of VNO sensory neurons, such that neurons lining the apical half of the VNO neuroepithelium co-express V1Rs and the G-protein α -subunit, $G_{\alpha i2}$, whereas neurons of the basal half of the VNO are both V2R and G_{α} positive (Dulac, 2000).

One member of the ephrin family, ephrin-A5, is required for the targeting of V1R-expressing vomeronasal axons to the anterior AOB, presumably through interactions with the Eph receptor EphA6 (Knoll *et al.*, 2001). The targeting of apical vomeronasal axons to the anterior AOB is also

dependent on repulsive forces from the posterior AOB that signal, at least in part, through neuropilin-2 (Npn-2) (Cloutier *et al.*, 2002). Npn-2 is required for the targeting of apical vomeronasal axons to the anterior AOB as well as fasciculation of the vomeronasal nerve and also for separation of the main and accessory olfactory projections during development (Cloutier *et al.*, 2002; Walz *et al.*, 2002).

Secreted semaphorins and slits also participate in establishing the glomerular map in the accessory olfactory system. It is likely that a combination of attractive and repulsive cues in the AOB, and possibly in the vomeronasal sensory neurons themselves, allows for the precise targeting of these sensory afferent projections to either posterior or anterior regions of the AOB. The Slit and Robo families of axon guidance molecules function as cues for basal vomeronasal axons (Knoll *et al.*, 2003). Vomeronasal neurons are repelled by Slit proteins *in vitro*, and Robo2 is expressed in vomeronasal neurons located in the basal region of the VNO (Knoll *et al.*, 2003; Marillat *et al.*, 2002).

Sema3F-Npn-2 signaling is essential for fasciculation of the vomeronasal nerve as it courses past the MOB but is largely dispensable for the targeting of apical vomeronasal axons to the anterior AOB. Moreover, Sema3F is required for the segregation of sensory neuron projections within the main and accessory olfactory systems and for accurate laminar targeting of main olfactory sensory neuron axons (Cloutier *et al.*, 2004). Slit-1 is not required for fasciculation of the vomeronasal nerve but is critical for the targeting of basal vomeronasal sensory neuron axons to the posterior AOB (Cloutier *et al.*, 2004). These results show that two families of secreted repellents play complementary roles in the development of primary sensory neuron projections in the accessory olfactory system.

VIII. Analysis of Axon Growth and Guidance

Several techniques have been used to study axonal guidance. The collagen gel assay, the stripe assay, the pipette assay, and the pump assay are very powerful techniques that have been used to study the effects (attraction, repulsion, or collapse) of different molecules on given axonal populations. In order to study the formation of axon tracts *in vivo*, researchers have used both tract tracing techniques using fluorescent dyes such as DiI or *in vivo* imaging by diffusion tensor magnetic resonance imaging (DTI-MRI). This nondestructive technique can be used to survey the development of multiple axonal tracts in three dimensions in the embryonic, postnatal, and adult brains.

A. The Collagen Gel Assay

The three-dimensional collagen gel assay is one of the most widely used techniques for analyzing axon growth from explants. Collagen is a physiologically relevant biological matrix that allows neurites to grow and soluble factors to diffuse. Generally, two explants or one explant and a source of a putative guidance factor are placed side-by-side in the gel. One explant contains the cells being studied (for example, the dorsal spinal cord containing commissural neurons). As these axons grow into the collagen, they encounter guidance factors released from a second explant (such as the floor plate). The number of axons emanating from the side toward the second, or target, explant can be compared with the number of axons projecting from the side away from the target. The number of axon turning within the gel can also be analyzed. The collagen gel assay was first used by [Lumsden and Davies \(1983\)](#), using co-cultured embryonic mouse sensory neurons, but has since been used to identify and characterize a number of different axonal guidance molecules.

B. The Stripe Assay

In order to elucidate the mechanisms involved in axonal guidance by membrane-bound cues, especially with regard to the topographic targeting of retinal ganglion cell axons within the tectum, Friedrich Bonhoeffer and colleagues developed the “stripe assay” ([Walter *et al.*, 1987a,b](#)). In this assay, retinal explants are grown on membrane stripes made from alternating rostral and caudal parts of the tectum. Membrane fragments of two different sources are arranged as a carpet of very narrow alternating strips. Axons growing on such striped carpets are simultaneously confronted with the two substrates at the stripe borders. If there is a preference of axons for one or the other substrate, axons become re-oriented and grow within the lanes of the preferred substrate. Such preferential growth can be due to an increased affinity for attractive factors on the preferred stripes or avoidance of repulsive factors on the alternate stripes, and thus adequate controls must be used to differentiate between these two possibilities. Tectal cell membranes are an excellent substrate for the growth of retinal axons. Results using tissues from developing chicks or rodents show that temporal RGC axons avoid stripes made up of caudal tectal tissue, while nasal RGC axons grow equally well on either rostral or caudal tectal tissue stripes ([Godement and Bonhoeffer, 1989](#); [Roskies and O’Leary, 1994](#); [Walter *et al.*, 1987a,b](#)), or show a preference for nasal stripes providing specific pre-treatments ([von Boxberg *et al.*, 1993](#)).

These experiments suggest the existence of biochemical labels for the specification of axon–target interactions and provide the conceptual basis for the work of Friedrich Bonhoeffer and others that ultimately led to the identification of specific molecular guidance factors for retinotectal topography.

C. The Pipette Assay

Originally pioneered by Mu-Ming Poo and colleagues, the “*Xenopus* growth cone turning assay” has recently become a popular assay for assessing growth cone turning responses to gradients of guidance factors. The turning assay uses cultured *Xenopus* spinal neurons to examine the cytoplasmic events associated with neurite growth and the response of the growth cone toward extracellular guidance cues. This method uses repetitive pulse application to create reproducible chemical gradients. Microscopic gradients are generated by repetitive pulsatile ejection of picoliters of chemical solution near the growth cone. The gradient generated is a stable gradient over a distance of tens of microns within several minutes. Growth cones of isolated *Xenopus* spinal neurons in culture exhibit chemotactic turning responses when exposed to a gradient of Netrin-1 (Ming *et al.*, 2002). The gradient is produced by repetitive pulsatile ejection of picoliters of solution containing Netrin-1 from a micropipette, positioned at a 45° angle with respect to the direction of initial neurite extension and a distance of 100 μm away from the center of the growth cone. By applying defined extracellular gradients of guidance molecules, early responses of the growth cone to specific guidance cues can be examined as well as the involvement of various cytoplasmic signaling pathways in regulating the turning decision of the growth cone.

Using this assay, Poo and colleagues had previously demonstrated that growth cones from young *Xenopus* retinal explants extend toward a gradient of Netrin-1 protein and away from a gradient of Sema3A (de la Torre *et al.*, 1997; Song *et al.*, 1998). In addition to inducing transient collapse and branching, Sema3A can act as a directional guidance cue in the growth cone turning assay for retinal growth cones (Campbell *et al.*, 2001).

D. The Pump Assay

Geoffrey Goodhill and colleagues designed a technique for generating precise and reproducible gradients of diffusible molecules in collagen gels (Narasimhan, 2004; Rosoff *et al.*, 2004). The assay establishes molecular gradients by printing drops of solution onto the surface of a thin collagen gel. Based on the mechanics of diffusion, the shape and steepness of the

gradient can be controlled, and the actual concentration of molecules produced by this method can be measured quantitatively with fluorescence imaging. Using this technology, they showed that growth cones are capable of detecting a concentration difference as small as about one molecule across their spatial extent. Furthermore, this sensitivity exists across only a relatively small range of ligand concentrations, indicating that adaptation in these growth cones is limited. Their gradient generation method allows for the production of large numbers of identical gradients that require only limited quantities of chemotropic molecules. Moreover, the gradients are stable for at least a day or two after they have been generated. This technique could also be used to generate gradients of multiple factors with different shapes and arbitrary spatial relationships. This powerful new technology can be applied to quantitative studies of other biological processes controlled by molecular gradients, such as cell migration, as well axonal regeneration following injury.

E. The Organotypic Slice Assay

In mammals, surgical manipulation is difficult before birth. Thus, many researchers have turned to using slice preparations to study axonal growth and guidance. Slices maintained in stationary culture with the interface method are ideally suited for manipulations that demonstrate the guidance properties of a prospective target tissue (Gahwiler *et al.*, 1997). For example, to better understand the projection of the commissural axons in the spinal cord, an “open book” organotypic slice assay was used, in which sections of spinal cord are split at the midline, flattened, and cultured in collagen to analyze the trajectories of the commissural axons. Advances in live optical imaging, including confocal and two-photon microscopy using GFP-based markers, allow the direct observation of axonal growth within the slice. Thus, using this assay, live growing axons can be studied *in situ* with maximum resolution and precision.

F. MRI/DTI

Magnetic resonance imaging (MRI) uses a strong electromagnetic field to align hydrogen atoms in the body in parallel with the magnetic field, in either the same or the opposite direction of the field. Hydrogen atoms become excited and resonate with the exciting wave. As the hydrogen atoms return to their original energy state, energy is released in the form of radio waves, which are detected by the MRI machine. Diffusion tensor imaging (DTI) measures the alignment of water molecules in the brain. Water molecules

preferentially align within ordered structures in the brain such as axonal tract. The vector orientation of their alignment can be observed and color-coded to give a visual image of the position and direction of axon tracts within the brain. DTI can be combined with conventional MRI to capture T1- and T2-weighted images such that a three-dimensional map of the brain can be generated (Zhang *et al.*, 2003).

Recent advances in magnetic resonance research have opened up new opportunities for gathering functional information about the brain. The development of DTI has offered the possibility to go beyond anatomical imaging to study tissue structure at a microscopic level *in vivo*. Currently, DTI provides an invaluable tool for the study and diagnosis of white-matter diseases. An important application of DTI is called “fiber tracking,” which follows the vector trajectories from given coordinates with the brain to “trace” axonal pathways. Combined with fMRI, information about white-matter tracts reveals important information about neurocognitive networks and may improve our understanding of brain function. Future advances could involve the use of magnetically labeled molecules to analyze axons expressing specific molecules *in vivo*, possibly in living brains.

IX. Conclusions

The establishment of correct neuronal connections is crucial for proper functioning of the vertebrate nervous system. Such precise wiring of neuronal connections is also required in the adult brain for functional recovery after brain injury and disease. To reach their proper targets, axons rely upon the expression of highly conserved families of attractive and repulsive guidance molecules, including the Netrins, Slits, semaphorins, and ephrins. These guidance systems are used to generate an astonishingly varied set of neuronal circuits. It is difficult to understand how such complexity in brain wiring is derived from such a relatively small set of molecules. The regulation of guidance receptors and ligands allows a single guidance system to generate a variety of different responses, but even these cannot account for the huge number of different connections made in the nervous system. What is clear is that the same molecules are used over and over in establishing different projections within the brain and spinal cord. Furthermore, although they may function in slightly different ways, the molecules themselves are conserved from flies and worms to humans. Recent evidence suggests that additional layers of complexity may exist where molecules and receptors from different molecular families interact in signaling to create more complicated responses. We do not know exactly how growth cones are able to respond to molecular gradients within the environment, nor exactly how these signals are relayed intracellularly. Furthermore, little is known about

how axons are able to respond to multiple gradients simultaneously. Because neuronal regeneration is, to a certain extent, a recapitulation of development, understanding the mechanisms regulating axon guidance will not only help delineate the pathoetiology of various neurological disorders, due to erroneous axon pathfinding, but also shed light on potential ways of clinically repairing the injured nervous system.

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