

Chapter 1

Semaphorin Receptors and Their Signaling

Manabu Negishi and Izumi Oinuma

Abstract Semaphorins, which are a large family of secreted and membrane-bound molecules, were initially identified as neuronal axon-guidance signaling molecules but are now known as important key regulators for cell adhesion and motility in a wide range of organ systems, such as angiogenesis and immune response. The semaphorin receptors, neuropilins and plexins, are expressed in a variety of cell types, including neurons, endothelial cells, and cancer cells. Plexins are primarily receptors responsible for intracellular semaphorin signalings. Plexins possess an intrinsic GAP (glyceraldehyde-3-phosphate) activity for R-Ras subfamily GTPases, and this GAP activity is one of the crucial signals of semaphorins. In addition, plexins associate with a variety of signaling molecules, such as Rho GEFs and Rho GAP, and these associated molecules determine the characters of semaphorin signals. On the other hand, their signalings are critically modulated by their associated co-receptor molecules, including tyrosine kinase receptors. Semaphorins provide attractive and repulsive responses in a variety of cells, but associated co-receptors of plexins frequently hold the key to conversion between attraction and repulsion. In this chapter, we focus attention on the molecular signaling systems of plexins.

Keywords Semaphorin • Plexin • GAP • Small GTPase

1.1 Introduction

The semaphorins were initially identified as axon-guidance factors that induce axonal growth cone collapses. However, it is now clear that semaphorins and their receptors are widely expressed in a variety of cells, including neurons, endothelial cells, and various tumor cells, and they are implicated in a variety of physiological functions, including control of nervous and endothelial development and immune systems. Semaphorins display their functions through direct binding to either

M. Negishi (✉) • I. Oinuma
Laboratory of Molecular Neurobiology, Graduate School of Biostudies, Kyoto University,
Sakyo-ku, Kyoto 606-8501, Japan
e-mail: mnegishi@pharm.kyoto-u.ac.jp

their specific receptors, plexins, or their co-receptors, neuropilins, which form complexes with plexins. Plexins function as crucial receptors of semaphorins, and their intracellular domains are responsible for initiating cellular signal transduction inducing cellular functions of semaphorins. Initially, Plexin-B1, the receptor for *Sema4D*, has been shown to activate RhoA through binding to RGS-RhoGEF, but the plexin family has been identified to display GAP activity for R-Ras and Rap small GTPases through intrinsic GAP domains within the intracellular tails of plexins (Negishi et al. 2005; Zhou et al. 2008). In this chapter we focus attention on the molecular signaling systems of plexins.

1.2 Semaphorins and Their Receptors, Plexins and Neuropilins

Semaphorins are a large family of secreted or membrane-bound proteins. The semaphorin family contains 21 vertebrate genes and 8 additional genes found in invertebrates, and they are now classified into 8 subclasses on the basis of amino acid sequence similarity and distinctive structural features (Neufeld and Kessler 2008). Classes 1 and 2 are invertebrate semaphorins, classes 3 through 7 are vertebrate semaphorins, and class V are viral-encoded semaphorins, found in the genome of non-neurotrophic DNA viruses. Among these, classes 1, 4, 5, and 6 are transmembrane molecules, and class 7 is a membrane-associated form through a glycosylphosphatidylinositol-anchor motif. In the case of *Sema4D*, membrane-bound *Sema4D* is cleaved proteolytically, generating a biologically active soluble *Sema4D* fragment in some conditions (Zhu et al. 2007). On the other hand, classes 2, 3, and V are secreted proteins. Semaphorins are characterized by the sema domains and PSI (plexins, semaphorins, and integrins) domains, located in amino-terminal regions. In contrast, the carboxyl-terminal regions are more variable, and some of them have immunoglobulin-like loops. Class 3 and 4 semaphorins are homodimerized through cysteine disulfide bonds, and dimerization appears to be critical for their biological function. Semaphorins of classes 4 to 7 as well as *Sema3E* bind directly to specific plexins and produce plexin-mediated signal transductions. Other class 3 semaphorins do not bind to plexins but instead bind to neuropilins.

The function of semaphorins is mediated by their receptors, the plexins, and mammalian plexins are classified into four subfamilies: Plexin-A1-4, Plexin-B1-3, Plexin-C1, and Plexin-D1 (Tamagnone et al. 1999). Plexins are single-pass transmembrane receptors, and they contain sema domains, PSI domains, and G-P (glycine-proline)-rich motifs in the extracellular regions. On the other hand, in the cytoplasmic regions, plexins directly encode GAP domains for Ras family small GTPases split by the docking site for several Rho family small GTPases such as Rnd, RhoD, and Rac1. In addition, B-type plexins contain a C-terminal PDZ-binding motif for RGS-RhoGEFs. The neuropilins, neuropilin-1 and neuropilin-2, are another type of receptor and were initially identified as direct binding receptors

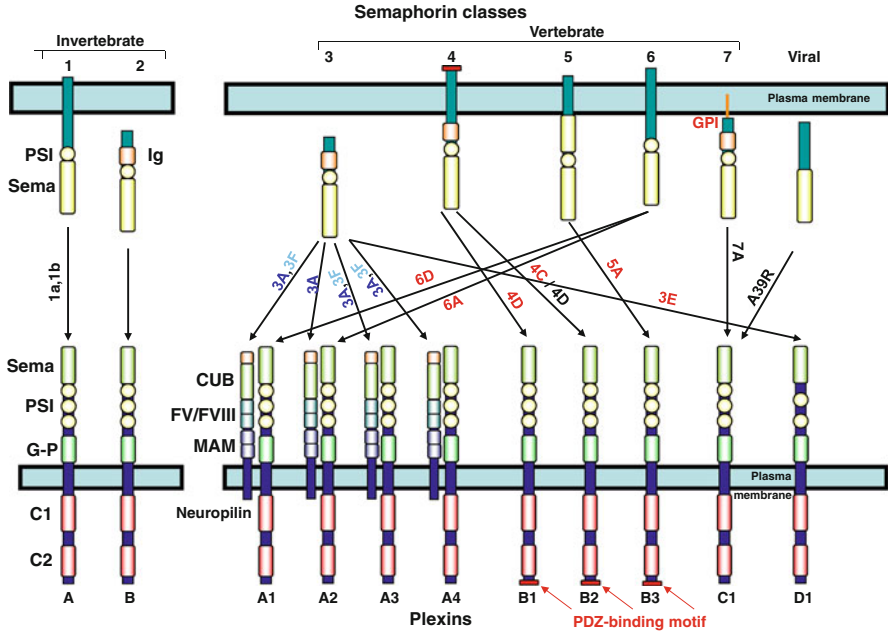


Fig. 1.1 The structures of semaphorins and their receptors. The structural features of the various classes of semaphorins and their neuropilin and plexin receptors are depicted. Semaphorins exist as the secreted, transmembrane, or GPI-anchored form shown *above*. The two neuropilins are single-pass transmembrane receptors. The nine vertebrate plexins are subdivided into four type A and three type B plexins and plexins C1 and D1. Plexins are single-pass transmembrane receptors distinguished by the presence of a split cytoplasmic GAP domain. Domains: *PSI* plexin, semaphoring, and integrin, *Sema* semaphorin, *Ig* immunoglobulin-like, *G-P* glycine-proline-rich, *CUB* complement binding, *FV/FVIII* FV/FVIII coagulation factor-like, *MAM* Meprin, A5, Mu

for class 3 semaphorins. The B-type plexins also contain a conserved cleavage site in their extracellular domains for furin-like pro-protein convertases that is not found in other plexins. The two neuropilins are single-pass transmembrane receptors, and they have two CUB (complement-binding) domains, two FV/FVIII coagulation factor-like domains, and a MAM domain in their extracellular regions. However, neuropilins have very short cytoplasmic tails, and they alone have no ability to transduce the signals of Sema3. Neuropilins form complexes with Plexin-As, and associated Plexin-As are signaling moieties. Classification of semaphorins and their receptors are depicted in Fig. 1.1.

1.3 Invertebrate Plexins

Invertebrate semaphorins have been characterized in *Drosophila melanogaster*. *D. melanogaster* semaphorins consist of Sema-1a, Sema-1b, and Sema-2a. Sema-1a and Sema-1b are transmembrane semaphorins and bind to PlexA, whereas Sema-

Sema-2a is a secreted semaphorin and binds to PlexB, inducing axon repulsion, dendritic targeting, and synapse formation (Fig. 1.2). PlexA forms a complex with Off-track, a transmembrane-type putative receptor tyrosine kinase, and this association has been thought to be required for axon repulsion (Winberg et al. 2001). In addition, PlexA probably associates with Gyc76C, a transmembrane-type guanylate cyclase, and Gyc76C-mediated cyclic GMP production has been thought to function in PlexA signaling (Ayoob et al. 2004). Downstream signaling systems of PlexA have been studied and several associated molecules have been identified. PlexA associates with Nervy, a member of the A-kinase anchoring proteins, and this association links to A-kinase, suggesting the involvement of cyclic AMP-dependent phosphorylation in the functions of Sema-1a/PlexA (Terman and Kolodkin 2004). Another important association molecule of PlexA is MICAL (molecule interacting with CasL). MICAL is a flavoprotein monooxygenase, and this enzyme activity has been shown to be required for Sema-1a-mediated axon repulsion (Terman et al. 2002). Recently, it has been shown that MICAL oxidizes actin, inducing the severing of actin

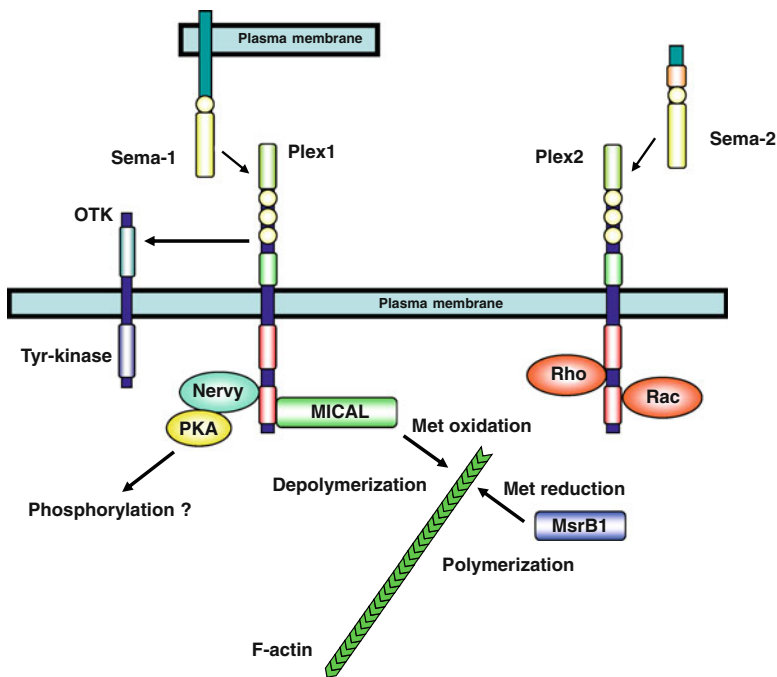


Fig. 1.2 Model for signal transduction of invertebrate semaphorin receptors. Two *Drosophila melanogaster* plexins have been described, PlexA and PlexB. Sema-1 directly binds to PlexA. A novel protein, MICAL, directly associates with the C2 domain of PlexA, inducing repulsive response through actin cytoskeleton reorganization. MICAL and MsrB1 are an oxidase and a reductase, respectively, and promote depolymerization and polymerization of actin filament, respectively, through regulation of actin oxidation. PlexB is a functional Sema-2a receptor and interacts directly with both Rac and Rho

filaments, and decreasing polymerization (Hung et al. 2011), and this redox activity of MICAL links semaphorins to F-actin disassembly (Hung et al. 2010), proposing a striking novel mechanism for semaphorin-mediated cytoskeletal reorganization. Furthermore, methionine-R-sulfoxide reductase B1 (MsrB1) has been shown to reduce the methionine residues of actin oxidized by MICAL and support actin assembly, indicating that MICAL and MsrB1 regulate actin assembly via oxidation and reduction of the methionine residues of actin (Lee et al. 2013). In contrast to PlexA, the signaling systems of PlexB are not yet understood. It has been reported that PlexB associates with the Rho family small GTPases, Rac and Rho, but this signaling system is not known (Hu et al. 2001).

1.4 Plexin-As

The *Sema3* subfamily does not directly bind to Plexin-As but does bind to neuropilins, and the *Sema3*-bound Plexin-A/neuropilin complex triggers downstream signaling. Among the *Sema3* subfamily, *Sema3A* and *Sema3F* selectively associate with the neuropilin-1/Plexin-A4 complex and the neuropilin-2/Plexin-A3 complex, respectively.

Semaphorins induce guidance signaling through regulation of cytoskeletal reorganization. One of the key regulators of cytoskeletal reorganization is Rho family small GTPases (BurrIDGE and Wennerberg 2004). The Rho family GTPases serve as molecular switches by cycling between an inactive GDP-bound state and an active GTP-bound state, and, once activated, they can interact with their specific effectors, leading to a variety of biological functions. Activation of the Rho family GTPases requires GDP-GTP exchange catalyzed by various guanine nucleotide exchange factors (GEFs) in response to a variety of extracellular stimuli, and their activation is turned off by GTPase-activating proteins (GAPs), which promote the intrinsic GTPase activities of the G proteins. Extracellular stimuli regulate the activities of Rho family GTPases through control of either GEFs or GAPs. *Sema3A*-induced growth cone collapse has been known to require Rac1 activity in dorsal root ganglion (DRG) neurons and spinal motoneurons (Jin and Strittmatter 1997). Plexin-A1 has been shown to associate with FARP2 (FERM, Rho GEF, and pleckstrin domain protein 2), a Rac GEF, and *Sema3A* binding to the receptor triggers the dissociation of FARP2 from Plexin-A1 and induces Rac1 activation (Toyofuku et al. 2005) (Fig. 1.3). Concerning the role of Rac1 activity in *Sema3A* signaling, it was reported that Rac1 is required for *Sema3A*-mediated endocytosis of the growth cone plasma membranes and reorganization of F-actin in DRG neurons (Jurney et al. 2002). Endocytosis of plasma membranes is supposed to be an important step for growth cone collapse. *Sema3A*-induced Rac1 activation may drive endocytosis of the plasma membranes instead of membrane protrusions, resulting in growth cone collapse. In addition, Rac1 activation transduces signaling through PAK (p21-activated kinase)-LIM kinase-cofilin to reorganization of actin filaments. Another role of Rac1 in the signaling of Plexin-A is the regulation of

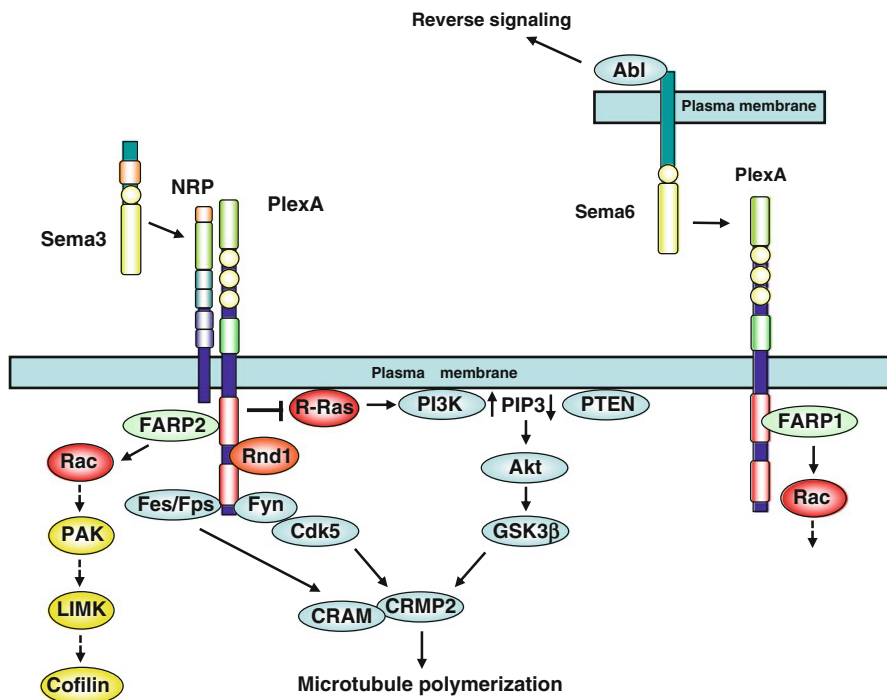


Fig. 1.3 Model for Plexin-A signal transduction. Sema3A and Sema3F bind to neuropilin-1 and -2, respectively, which associate with Plexin-A, and Plexin-A transduces repulsive signals. Several tyrosine and serine/threonine protein kinases, including Fes/Fps, Fyn, Cdk5, and GSK3 β , are activated by Sema3A, and eventual phosphorylation of CRMP2 is involved in Sema3A-induced repulsive response. Sema3A stimulates R-Ras GAP activity of Plexin-A, suppressing R-Ras activity, whereas it activates Rac through a Rac GEF, FARP2. Sema6A directly binds to Plexin-A2 and -A4, whereas Sema6C binds to Plexin-A1 to elicit axon repulsion

Plexin-A GAP activity. Plexins have a GAP domain in the intracellular tail, and the GAP domain is split by the Rho family GTPase-binding domain (Negishi et al. 2005; Zhou et al. 2008). Rac1 activated by FARP2 facilitates the binding of Rnd1, a member of Rho family GTPases, to the inserted region between the split GAP domains, inducing GAP activity for R-Ras (Toyofuku et al. 2005). Dissociated FERP2 also inhibits type 1 phosphatidylinositol phosphate kinase (PIP1 γ 661), resulting in suppression of integrin activation.

Tyrosine phosphorylation is involved in Sema3A/Plexin-A signaling, and Fes/Fps tyrosine kinase has been implicated in Sema3A-induced growth cone collapse (Mitsui et al. 2002). Fes/Fps is a non-receptor-type tyrosine kinase and directly binds to the cytoplasmic tail of Plexin-A1. In the unstimulated condition, neuropilin associates with Plexin-A1 and blocks the binding of Fes/Fps to Plexin-A1. Sema3A binding to neuropilin permits Fes/Fps to associate with and phosphorylate Plexin-A1. This tyrosine phosphorylation stimulates a repulsive

response in the receptor. Collapsin response mediator protein 2 (CRMP2) was originally identified as an intracellular signaling molecule of *Sema3A* (Goshima et al. 1995). CRAM was identified as another member of CRMPs and was shown to associate with CRMP2, forming a CRMP2–CRAM complex (Inatomi et al. 2000). Fes/Fps tyrosine-phosphorylates the CRMP2/CRAM complex as well as Plexin-A1. In addition to Fes/Fps, another kinase signaling is involved in *Sema3A*-mediated repulsive response. Fyn, a member of the Src family of tyrosine kinases, associates with and phosphorylates Plexin-A2 in response to *Sema3A* (Sasaki et al. 2002). Furthermore, serine/threonine kinase Cdk5 associates with Plexin-A2 and is activated through *Sema3A*-mediated Fyn; activated Cdk5 initially phosphorylates CRMP2. Then, another Ser/Thr kinase, GSK3 β , subsequently phosphorylates the Cdk5-phosphorylated CRMP2, and this dually phosphorylated CRMP2 is essential for the *Sema3A*-induced growth cone collapse (Brown et al. 2004). CRMP2 binds to tubulin heterodimers, promoting microtubule assembly (Fukata et al. 2002). It is thus plausible that Cdk5 and GSK3 β participate in *Sema3A* signaling through regulation of microtubule dynamics by phosphorylation of CRMP2.

In addition to *Sema3s*, *Sema6s* are ligands for Plexin-As. In contrast to *Sema3s*, secreted semaphorins, membrane-associated *Sema6s* directly bind to Plexin-As and do not require neuropilins (Fig. 1.3). *Sema6A* binds to Plexin-A2, controlling cerebellar granule cell migration (Kerjan et al. 2005). *Sema6A* binds to both Plexin-A2 and Plexin-A4, controlling lamina-restricted projection of hippocampal mossy fibers (Suto et al. 2007). *Sema6A* has been shown to associate with Plexin-A4 and promote the dendritic growth of spinal motor neurons through FARP1, a member of the Rho GEFs (Zhuang et al. 2009). In optic chiasm retinal ganglion cells, *Sema6D* has been shown to regulate axon guidance through Plexin-A1, switching the responses between repulsion and attraction dependent on the combination with Nr-CAM (Kuwajima et al. 2012). Interestingly, it has been shown that *cis* interaction between *Sema6A* and Plexin-A4 modulates the repulsive response to *Sema6A* (Topper et al. 2010). Further, reverse signaling of *Sema6D* has been reported, meaning that *Sema6D* is a Plexin-A1 ligand and also a receptor for Plexin-A1 (Toyofuku et al. 2004). *Sema6D*/Plexin-A1 regulates ventricular expansion, whereas *Sema6D* reverse signaling induces the expansion of the outer myocardial layer and trabeculation through the recruitment of Abl to the C-terminal domain of *Sema6D*. Thus, *Sema6s*/Plexin-As functions are diverse and complicated, and their downstream signaling pathways are not well characterized.

In addition to the aforementioned molecular systems, several other signaling molecules have been reported to associate with Plexin-As, including RanBPM (Togashi et al. 2006), DAPI2 (Takegahara et al. 2006), and p75 neurotrophin receptor (Zvi et al. 2007). Furthermore, it has been reported that *Sema3A* induces intra-axonal translation of RhoA mRNA, and this local translation of RhoA is necessary and sufficient for *Sema3A*-mediated axonal growth cone collapse, indicating regulation of the neuronal cytoskeleton by local RhoA translation (Wu et al. 2005). These signaling molecules may contribute to many functions of the *Sema3* subfamily.

1.5 Plexin-Bs

Plexin-Bs are responsible for a wide range of Sema4 subfamily functions, including axon guidance, dendrite remodeling, and angiogenesis, and Sema4s directly bind to Plexin-Bs, transducing signals. Plexin-B1 is the receptor for Sema4D, and it has been well characterized in its signaling pathways. Rho family small GTPases are also thought to be required for Plexin-B1 signaling. Active Rac1 was first reported to directly bind to the CRIB-like domain located between two GAP domains in the cytoplasmic tail of Plexin-B1 (Driessens et al. 2001). Plexin-B suppresses Rac activity by sequestering active Rac from its downstream effector PAK, inducing repulsive response in both *Drosophila* and mammalian cells (Hu et al. 2001; Vikis et al. 2002). Inhibition of Rac-mediated PAK activation by Plexin-B may suppress Rac-induced membrane protrusions, supporting the repulsive response. Furthermore, active Rac was shown to stimulate the localization of Plexin-B1 to the cell surface, enhancing Sema4D binding to the receptor (Vikis et al. 2002). Thus, signaling of Rac and Plexin-B1 appears to be bidirectional; Rac modulates Plexin-B1 activity, and Plexin-B1 modulates Rac function.

Among plexin families, the Plexin-B subfamily has a PSD-95/Dlg/ZO-1 (PDZ) domain-binding motif at the C-terminus (Fig. 1.4). Plexin-B1 directly interacts with two Rho GEFs containing PDZ domains, PDZ-RhoGEF and leukemia-associated Rho GEF (LARG), through the C-terminal PDZ binding motif (Swiercz et al. 2002; Aurandt et al. 2002; Perrot et al. 2002). They have PDZ domains at the N-terminal region and are members of the RGS-RhoGEF family, which has G12-family binding domains and specifically activates RhoA through G12-family binding. Sema4D binding to Plexin-B1 regulates the activity of PDZ-RhoGEF/LARG, leading to activation of RhoA, and this RhoA activation is involved in the Plexin-B1-induced repulsive response. It was shown that Sema4D/Plexin-B1-induced repulsive response is inhibited by an inhibitor of Rho-kinase, a well-studied Rho effector (Swiercz et al. 2002). Rho-kinase is known to phosphorylate the myosin-binding subunit of myosin phosphatase and myosin light chain, causing contraction of actomyosin (Fukata et al. 2001). The activation of myosin by Rho-kinase and the resultant contraction of actomyosin trigger cellular contraction and neurite retraction. In angiogenic systems, Sema4D/Plexin-B1 promotes endothelial cell motility through RhoA/Rho-kinase activation, and this action is mediated by integrin-dependent Pyk2 activation and the resultant stimulation of PI3kinase, Akt, and ERK through RhoA/Rho-kinase activation (Basile et al. 2007). Plexin-B1 constitutively interacts with PDZ-RhoGEF through its C-terminal PDZ domain-binding motif. How does Sema4D trigger the activation of associated PDZ-RhoGEF? It is known that ErbB-2, a tyrosine kinase receptor, associates with Plexin-B1 and tyrosine-phosphorylates Plexin-B1 in response to Sema4D (Swiercz et al. 2004). It has been shown that this ErbB-2-mediated tyrosine phosphorylation of Plexin-B1 promotes the interaction of phosphorylated Plexin-B1 with phospholipase C γ through its SH2 domain and that phospholipase C γ activates PDZ-RhoGEF via its SH3 domain

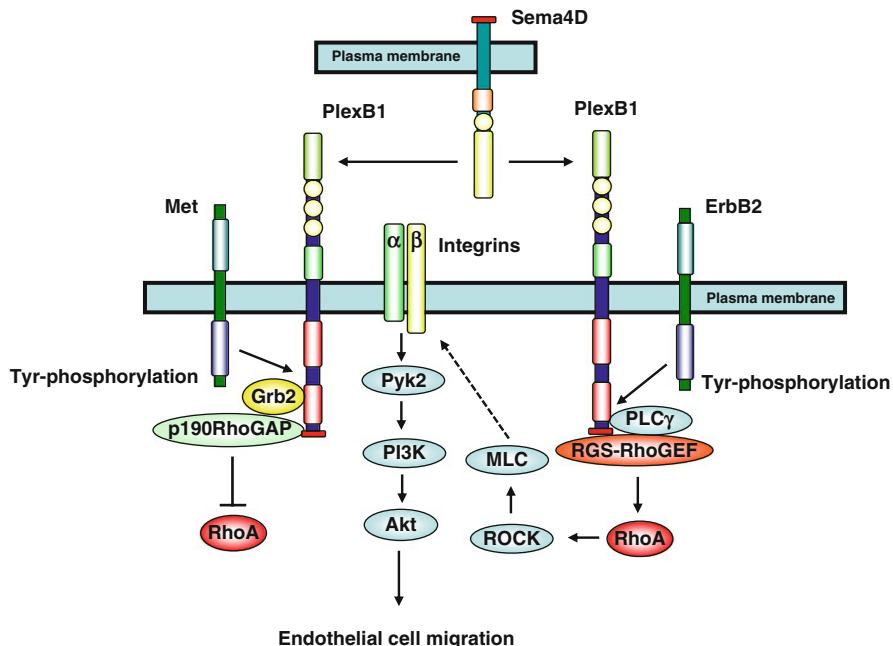


Fig. 1.4 Model for signal transduction of regulation of Rho activity by Plexin-B1. The C-terminal tail of Plexin-B1 has a PDZ-binding motif and stably associates with Rho-specific GEF, PDZ-RhoGEF, and LARG through this motif. Tyrosine phosphorylation of Plexin-B1 by ErbB2 in response to Sema4D induces phospholipase γ association with Plexin-B1, inducing RGS-RhoGEF and resultant RhoA activation. On the other hand, tyrosine phosphorylation of Plexin-B1 by Met triggers the association of Grb2/p190RhoGAP with Plexin-B1, activating p190RhoGAP and inhibition of RhoA activity

(Swiercz et al. 2009). This phospholipase γ -mediated activation of PDZ-RhoGEF is independent of its lipase activity.

Sema4D/Plexin-B1 stimulation frequently evokes different and sometimes opposing cellular responses depending on the cellular context; Sema4D/Plexin-B1 inhibits integrin-mediated cell attachment and cell migration in NIH-3T3 cells, whereas Sema4D/Plexin-B1 stimulates migration of SK-BR3 cells and endothelial cells through RhoA activation (Zhou et al. 2008). In contrast to RhoA activation by Plexin-B1 through PDZ-RhoGEF, Plexin-B1 has been shown to associate with p190 RhoGAP, leading to inhibition of cell migration (Barberis et al. 2005). Furthermore, it has been shown that interaction of Plexin-B1 with Met, another tyrosine kinase receptor, promotes the association of Plexin-B1 with p190 RhoGAP and subsequent RhoA inactivation, leading to Sema4D-mediated inhibition of cell migration (Swiercz et al. 2008). The activation of Plexin-B1 by Sema4D induces tyrosine phosphorylation of Plexin-B1 by Met, creating a docking site for the SH2 domain of Grb2, and associated Grb2 recruits p190 RhoGAP into the Plexin-B1/Grb2 complex through its SH3 domain, inducing RhoA inactivation (Sun et al. 2012).

Therefore, Plexin-B1 associates with two types of tyrosine kinase receptors, ErbB-2 and Met, and reciprocally regulates RhoA activity; the Plexin-B1/ErbB-2 complex activates RhoA through PDZ-RhoGEF, whereas Plexin-B1/Met complex inactivates RhoA through p190 RhoGAP, explaining how Sema4D exerts differential activities through governing different signaling pathways.

Plexin-B1 activates RhoA through association of PDZ-RhoGEF with the C-terminal PDZ-binding motif of Plexin-B1. However, among the entire plexin family, the C-terminal PDZ-binding motif is found only in the Plexin-B subfamily, and it is not found in invertebrate Plexin-B. Thus, it is unlikely that PDZ-RhoGEF-mediated RhoA activation is a common signaling pathway for the plexin family. The cytoplasmic tails of plexins have GAP domains. Plexin-B1 was first identified to display GAP activity for R-Ras (Oinuma et al. 2004a, b), and Plexin-A1, -D1, and -C1 have been shown to have R-Ras GAP activity in succession (Uesugi et al. 2009) (Fig. 1.5). Plexin-B1 has two subdomains, C1 and C2, showing GAP activity within the C-terminal tail, and these domains contain primary and secondary arginine motifs, respectively, which are critical motifs for GAP activity for small GTPases. Rnd1, a constitutively active atypical Rho family GTPase, binds to the region between C1 and C2, and this Rnd1 binding is essential for the expression of R-Ras GAP activity. Both Rnd1 binding to Plexin-B1 and Sema4D stimulation are indispensable for this activity (Oinuma et al. 2004a, b). In the absence of Rnd1, C1 and C2 domains interact intramolecularly, rendering the receptor inactive for R-Ras GAP activity. Rnd1 binding to the region between C1 and C2 domains disrupts the interaction of C1 and C2 domains, indicating that Rnd1 relieves the closed conformation of the C-terminal tail of Plexin-B1. This Rnd1-bound open conformation acquires an ability to associate with GTP-bound R-Ras. The Rnd1-Plexin-B1 complex can hold the GTP-bound R-Ras but not promote GTPase activity. Sema4D is homodimerized through cysteine disulfide bonds. Sema4D ligand binding to Plexin-B1 induces clustering of the Rnd1-bound monomeric receptor, and this clustering triggers the hydrolysis of GTP on R-Ras. Therefore, GAP activation of Plexin-B1 consists of two steps, interaction with GTP-bound R-Ras and promotion of GTP hydrolysis by R-Ras; the former is stimulated by Rnd1, and the latter is a process induced by the clustering by Sema4D. In addition to Plexin-B1, interaction of C1 and C2 domains has also been reported in Plexin-A1 and Plexin-D1, suggesting that this activation mechanism is a common system for the plexin family (Uesugi et al. 2009). The expression of R-Ras GAP activity of Plexin-B1 in response to Sema4D is required for the Sema4D/Plexin-B1-mediated axonal growth cone collapse. Sema4D/Plexin-B1 inhibits integrin-mediated cell migration in a variety of cells through R-Ras GAP activity (Oinuma et al. 2006). R-Ras has been shown to play a key role in cell adhesion and its activation is known to promote cell adhesion and neurite outgrowth through integrin activation (Kinbara et al. 2003). Therefore, downregulation of R-Ras activity by Sema4D/Plexin-B1 suppresses integrin activation and thereby reduces cell adhesiveness, leading to growth cone collapse and inhibition of cell migration.

Downstream signaling pathways of R-Ras GAP activity of Plexin-B1 were characterized. PI3-kinase is one of the prominent downstream effectors of R-Ras. It

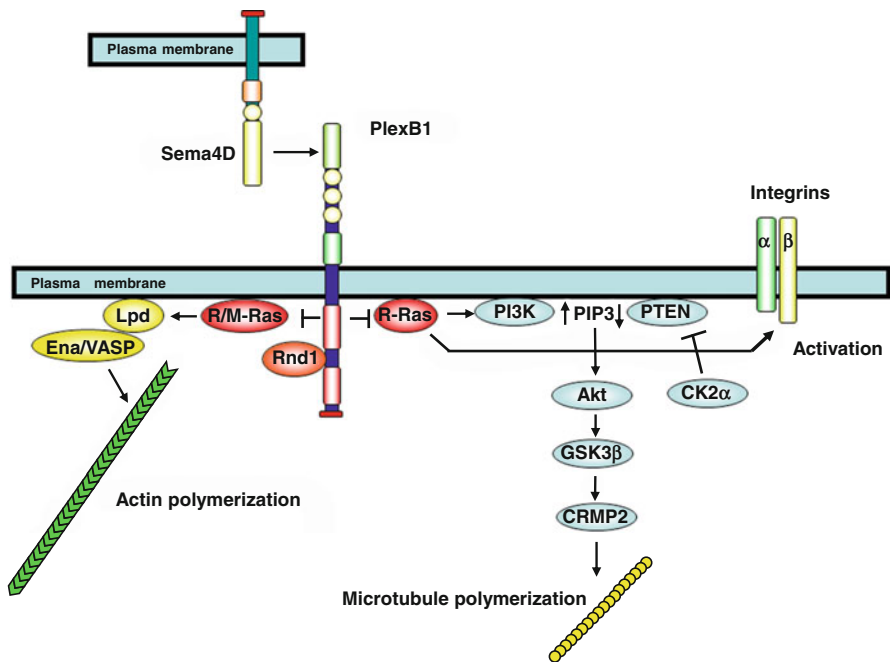


Fig. 1.5 Model for signal transduction of R-Ras GAP activity of Plexin-B1. The C1 and C2 domains of the cytoplasmic tail of Plexin-B1 encode R-Ras GAP. The C1 and C2 domains interact with each other, and Rnd1 binding to the region between C1 and C2 domains disrupts this interaction, allowing the receptor to associate with GTP-bound R-Ras. Sema4D-induced clustering of the Plexin-B1/Rnd1 complex promotes the hydrolysis of GTP by R-Ras. Downregulation of R-Ras activity inhibits integrin activity, reducing cell adhesion. Sema4D/Plexin-B1 suppresses the PI3-kinase pathway through both inhibition of R-Ras activity and PTEN activation, leading to CRMP2 phosphorylation and resultant microtubule depolymerization. R-Ras/M-Ras GAP activity suppresses R-Ras/M-Ras-mediated lamellipodin activation, leading to inhibition of actin filament polymerization

has been shown that Sema4D stimulation of Plexin-B1 inactivates PI3-kinase and Akt and activation of GSK3β through R-Ras GAP activity, and that active GSK3β phosphorylates and inactivates CRMP2, inducing axonal growth cone collapse (Ito et al. 2006). On the other hand, the phosphatidylinositol-3-phosphate level is critically regulated by PI3-kinase and PTEN (phosphatase and tensin homologue deleted chromosome 10). Phosphorylation of PTEN is known to suppress its phosphatase activity. It has been shown that Sema4D stimulation of Plexin-B1 induces dephosphorylation and activation of PTEN through R-Ras GAP activity and that this PTEN activation is also involved in Sema4D/Plexin-B1-mediated axonal growth cone collapse (Oinuma et al. 2010). Furthermore, Sema4D-induced PTEN activation via R-Ras GAP activity is mediated by the inhibition of casein kinase 2α. Thus, R-Ras GAP activity of Plexin-B1 dually regulates the phosphatidylinositol-3-phosphate level, inhibition of PI3-kinase activity, and stimulation of PTEN activity,

leading to Akt activity suppression, GSK3 β activation, and CRMP2 phosphorylation and then induces repulsive response, probably through suppression of microtubule polymerization mediated by CRMP2.

The R-Ras subfamily consists of three G proteins: R-Ras, M-Ras, and TC21. Among these, Plexin-B1 displays GAP activity for R-Ras and M-Ras but not for TC21 (Saito et al. 2009). During neuronal development, R-Ras is expressed in the stage of axon specification and elongation, whereas M-Ras expression is upregulated during dendritic development, and R-Ras and M-Ras localize at the axon and dendrite, respectively (Saito et al. 2009). *Sema4D/Plexin-B1* induces axonal growth cone collapse through inhibition of R-Ras activity, whereas *Sema4D/Plexin-B1* induces reduction of dendrite growth through inhibition of M-Ras activity (Saito et al. 2009). Thus, Plexin-B1 is a dual functional GAP for R-Ras and M-Ras, remodeling axon and dendrite morphology, respectively. Concerning downstream signaling of M-Ras, lamellipodin was identified to be a novel effector of M-Ras (Tasaka et al. 2012). Lamellipodin is a ligand for Ena/VASP, which is an actin polymerization-promoting factor (Legg and Machesky 2004). Furthermore, *Sema4D/Plexin-B1* has been shown to induce inhibition of lamellipodin action and the resultant disappearance of F-actin from distal dendrites through M-Ras GAP activity, reducing dendrite outgrowth (Tasaka et al. 2012). Thus, the M-Ras GAP activity of Plexin-B1 induces repulsive response through suppression of actin polymerization mediated by lamellipodin. Recently, plexins, including Plexin-B1, have been reported to display GAP activity for Rap1, and Rnd1 binding to plexins does not contribute to the activation of Rap GAP activity (Wang et al. 2012). However, it is not well characterized how both GAP activities, Rap GAP and R-Ras GAP, coordinately contribute to functions of plexins.

In contrast to Plexin-B1, signaling systems of the other Plexin-B subfamily receptors, Plexin-B2 and -B3, are not well characterized. *Sema5A* is a ligand for Plexin-B3, and *Sema5A* induces cell collapse of NIH3T3 fibroblasts through Plexin-B3, whereas *Sema5A* elicits attractive responses for epithelial and endothelial cells through the Plexin-B3/Met receptor complex (Artigiani et al. 2004). Thus, *Sema5A/Plexin-B3* appears to provide reciprocal responses in cell contexts and Met may convert the *Sema5A* signaling.

1.6 Plexin-C1 and -D1

Plexin-D1 is a receptor for *Sema3E*. In contrast to the other *Sema3s*, *Sema3E* directly binds to Plexin-D1. Plexin-D1 is expressed in vascular endothelium, and *Sema3E* acts as a repulsive cue for the endothelial cells (Gu et al. 2005). Plexin-C1, also named VESPR for its viral origin, serves as a receptor for the virally encoded *SemaVA*, and Plexin-C1 stimulation induces inhibition of integrin-mediated adhesion and the chemokine-induced migration of dendritic cells (Walzer et al. 2005). Plexin-D1 and -C1 also display GAP activity for R-Ras and M-Ras, but differ from Plexin-A and -B subfamilies in the requirement for Rnd GTPases of R-

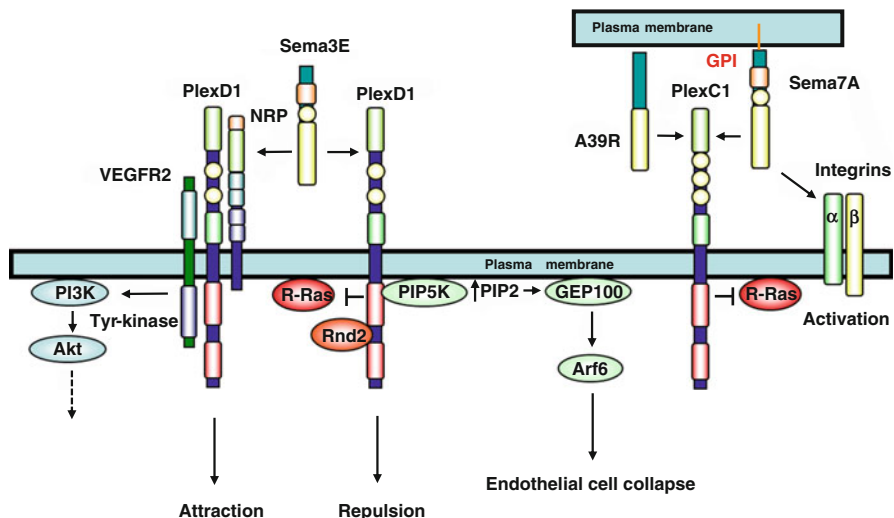


Fig. 1.6 Models for signal transduction of Plexin-C1 and D1. Sema3E directly binds to Plexin-D1 to either induce repulsive response through R-Ras GAP activity or attract response through the Plexin-D1/neuropilin/VEGF receptor complex. Plexin-C1 is a receptor for viral SemaVA or Sema7A. Plexin-C1 encodes R-Ras GAP, inhibiting R-Ras activity. Sema7A also binds to and activates integrins

Ras GAP activity (Fig. 1.6). Although Plexin-A and -B subfamilies require Rnd1 for displaying R-Ras GAP activity among Rnd subfamily GTPases, Plexin-C1 exhibits R-Ras GAP activity without the Rnd subfamily, whereas Plexin-D1 requires Rnd2 for displaying R-Ras GAP activity (Uesugi et al. 2009). Therefore, R-Ras GAP activity is a common signaling of plexin subfamilies, but the regulation of R-Ras GAP activity of plexins by Rnd proteins differs among plexin subfamilies.

Concerning antiangiogenic signaling through Plexin-D1, Sema3E/Plexin-D1 has been reported to initiate a two-pronged mechanism, R-Ras inactivation and Arf6 stimulation, which affect the status of activation of integrins and their intracellular trafficking, respectively (Sakurai et al. 2010). Furthermore, Sema3E stimulation has been shown to recruit phosphatidylinositol-4-phosphate-5-kinase to Plexin-D1, and its product, phosphatidylinositol-4,5-bis-phosphate, binds to GEP100/Brag2, a GEF for Arf6, stimulating Arf6 activity and inducing endothelial cell collapse (Sakurai et al. 2011). Sema3E/Plexin-D1 acts as the critical regulator for angiogenesis through cooperative linkage of two small GTPase signaling pathways. In contrast, Sema3E/Plexin-D1 provides reciprocal responses; Sema3E acts as a repellent for corticofugal and striatonigral neurons, expressing Plexin-D1 but not neuropilin-1, and Sema3E acts as an attractant for subiculomammillary neurons, coexpressing Plexin-D1 and neuropilin-1 (Chauvet et al. 2007). Furthermore, the latter neurons express VEGFR2 (vascular endothelial growth factor receptor type 2), and VEGFR2 associates with the Plexin-D1/neuropilin-1 receptor complex (Bellon et al. 2010). Sema3E triggers VEGFR2-dependent activation of the PI3-kinase/Akt signaling

pathway, leading to axonal growth; this activation is independent of VEGF, a ligand for VEGFR2. Thus, gating of Sema3E/Plexin-D1 signaling by neuropilin-1 switches axonal repulsion to attraction, controlling the formation of select forebrain projections.

Sema7A was reported to associate with Plexin-C1, influencing immune cell functions, such as cytokine production. However, it was clearly shown that Sema7A promotes axon outgrowth through integrin activation and initiates T-cell-mediated inflammatory responses through the integrin receptor, indicating that integrin is a receptor for Sema7A (Pasterkamp et al. 2003; Suzuki et al. 2007). On the other hand, Sema7A has been shown to suppress melanoma progression through activation of Plexin-C1-mediated R-Ras GAP activity (Scott et al. 2009). Further work is needed for characterization of the endogenous ligands of Plexin-C1.

1.7 Concluding Remarks

Here we have summarized great advances in our understanding of the molecular mechanisms for intracellular signalings of semaphorin receptors. The main receptors for semaphorins are plexins, which directly encode R-Ras GAP and exert guidance signaling through R-Ras GAP activity and the resultant regulation of integrin activity, and this GAP activity is probably a common feature of semaphorin signaling. In addition, plexins differentially regulate Rho family GTPase activities through associating with either Rho GEF or Rho GAP. Identifying various molecules involved in semaphorin signal transduction pathways provides a good understanding of the diverse molecular mechanisms for the intricate guidance functions of semaphorins. In addition to semaphorins, many guidance molecules, such as netrins, ephrins, and slits, have been identified, and their signaling systems are extensively studied. Navigation of many cells is frequently determined by the integration of various guidance cues, and the combined actions of guidance molecules influence the outcome of guidance (Dudanova and Klein 2013). To unravel the complicated signaling systems, studies of the signaling by each guidance molecule and their combined actions are required. It seems certain that semaphorins are important guidance molecules in a wide range of cellular functions. Future research will undoubtedly reveal the entire picture of the signaling cascades of semaphorins and plexins for their diverse functions.

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