

Review



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The Role of the Slit/Robo Signaling Pathway

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Abstract

The Slit family is a family of secreted proteins that play important roles in various physiologic and pathologic activities via interacting with Robo receptors. Slit/Robo signaling was first identified in the nervous system, where it functions in neuronal axon guidance; nevertheless, an increasing number of studies have shown that Slit/Robo signaling even regulates other activities, such as angiogenesis, inflammatory cell chemotaxis, tumor cell migration and metastasis. Although the precise role of the ligand-receptor in organisms has been obscure and the conclusions drawn are sometimes paradoxical, tremendous advances in understanding the Slit/Robo signaling pathway have been made. As such, our review summarizes the characteristics of the Slit/Robo signaling pathway and its role in various cell types.

Key words: Slit, Robo, neuro, angiogenesis, chemotaxis, cancer, motility

Introduction

Slits are highly conserved secreted glycoproteins that regulate many physiologic processes, such as neuronal axon guidance, cell proliferation, cell migration, and vascularization, via binding to Robo receptors [1]. Slit/Robo signaling was originally recognized by humans as attracting and repelling neuron axons to across the midline [2]. Since then, the Slit/Robo signaling pathway has also been found to play an important role in the development of organs, such as diaphragm, kidney, heart and mammary gland, in addition to the nervous system [3-6]. More recently, accumulating studies reported that Slit/Robo signaling was altered in various cell types and proved that it acts as a vital regulator in keeping physical or pathological function. Herein, we summarize advances in Slit/Robo signaling in various functional events.

Characteristics of Slit and Robo

Structural characteristics of Slit protein

Slit proteins are a class of single peptides with approximately 1500 amino acids. Invertebrates have only one Slit, while vertebrates have Slit1, Slit2, and Slit3 [7]. The Slit1 gene is located on human chromosome 10q24.1, the Slit2 gene is located on human chromosome 4p15.31, and the Slit3 gene is located on human chromosome 5q34-35.1 [8]. The Slit protein consists of five regions: one N-terminal signal peptide, four leucine-rich domains (LRR, D1-D4) in tandem with disulfide bonds, six epidermal growth factor-like (EGF-like) domains, an Agrin-Perlecan-Laminin-Slit/Laminin-G-like domain, one (invertebrates) or three (vertebrates) epidermal growth factor (EGF) domains, and a C-terminal cysteine-rich knot [9, 10] (Figure 1A). Protein structural studies showed that Slit protein plays a regulatory role by binding to the first Ig of Robo1 at the second LRR domain [11] (Figure 2B), whereas two Slit2 proteins can unexpectedly bind to each other at the fourth LRR domain to form homodimers [12]. The Slit protein is cleaved by proteolytic enzymes between the fifth and sixth EGF-like domains to generate the long N-terminal Slit segment (SlitN) and the short C-terminal Slit segment (SlitC) (Figure 2A).

The SlitN fragment combines with Robos to mediate various life activities, while the SlitC

fragment cannot bind to Robo [13]. SlitC has long been considered as a fragment without a regulatory function until recent studies reported that the SlitC fragment was found to be involved in the regulation of the protein kinase A (PKA) pathway in adipocyte thermoregulation and glucose metabolism [14] and mediated neuronal axon guidance by binding to the plexin receptor [13, 15] (Figure 2A). More recently, a report argued that a Slit2 variant that lacks exon 15 in the D2 domain plays a different role in lung cancer [16] (Figure 3). Overall, the function of different Slit protein fragments after translation or different types must be determined further.



Figure 1. Structure of the Slit/Robo protein family and their interaction. (A). Structure of the human Slit protein. Slits consist of five regions as follows: one N-terminal signal peptide, four leucine-rich domains (LRR, D1-D4) in tandem with disulfide bonds, six epidermal growth factor-like (EGF-like) domains, an Agrin-Perlecan-Laminin-Slit (ALPS)/Laminin-G-like domain, three epidermal growth factor-likedomains, and a C-terminal cysteine-rich knot. Slits are proteolytically cleaved between EGF-like domains. (B). Structure of the human Robo protein. The extracellular domains of the Robo1-3 proteins have the same structures, including 5 immunoglobulin domains, 3 fibronectin domains and one transmembrane domain. The Robo4 extracellular domain has only 2 immunoglobulin domains, 2 fiber connexin domains and one transmembrane region; the Robo1 and Robo2 intracellular region contains four conserved proline-rich domains, referred to as CC0-CC3. The Robo3 intracellular domain contains CC0, CC2 and CC3, and Robo4 only contains CC0 and CC2.



Figure 2. Slit/Robo protein proteolytic processing. (A). Slit protein proteolytic processing. Full-length Slit ligands are cleaved between the fifth and sixth EGF-like domains to create an N-terminal fragment (Slit N) and a C-terminal fragment (Slit C), both of which combine with different receptors and serve strikingly different functions. (B). Robo protein proteolytic processing. Slit protein in the extracellular matrix binds to the first lg of Robo I at the second LRR structure and creates tension in the Robo juxtamembrane domain, allowing metalloprotease ADAM10 Kuzbanian to cleave the Robo ectodomain. The remaining segment may be hydrolyzed further by γ-secretase and enter the nucleus to initiate downstream molecules.



Figure 3. The function of two kinds of Slit2 in lung cancer. (A). Slit2-WT (presence of exon 15) possesses only invasion inhibitory activity. (B). Empty Slit2 is a control. (C). Slit2- Δ E15 (absence of exon 15) inhibits both the growth and invasion of CL1–5 lung cancer cells.

Structural characteristics of Robo protein

Robo proteins are a class of transmembrane receptor proteins with 1000 to 1600 amino acids and have highly conserved intracellular domains with no autocatalytic or intrinsic enzymatic activity. Nematodes have one Robo receptor (Sax-3, sensory axon guidance receptor 3) [17], Drosophila, chickens, and Xenopus have three Robo receptors called Robo1, Robo2 and Robo3 [18, 19], and zebrafish and mammals have four Robo receptors (Robo1-4) [20]. The Robo1 and Robo2 genes located in the human chromosome at 3p12.3, and Robo3 and Robo4 genes located in the human chromosome 11q24.2. The extracellular domains of the Robo1-3 proteins have the same structure, which includes 5 immunoglobulin domains, fibronectin domains and 3 one transmembrane domain. The extracellular Ig4 (D4) of Robo2 is the dimerization domain to facilitate the recruitment and activation of enzymatic effectors to instigate intracellular signaling [21]. The Robo4 extracellular domain consists of only 2 immunoglobulin domains and 2 fiber connexin domains and one transmembrane region. The Robo1 and Robo2 intracellular region contains four conserved proline-rich domains, referred to as CC0-CC3; the Robo3 intracellular domain contains CC0, CC2 and CC3, whereas Robo4 only contains CC0 and CC2 [7, 20] (Figure 1B).

Robo proteins with no autocatalytic and intrinsic enzymatic activity in the intracellular region mediate downstream signaling through the recruitment of different adaptors or proteins. Notably, Robos can be selectively cleaved into different subtypes after transcription, and the consequence of its extracellular domain loss after translation and the interaction between different Robo receptors increase the diversity and complexity of their function [22-24] (Figure 2B).

Divergent members of the Robo family

Robo4 was long regarded as a specific receptor for vascular endothelia cells until very recently, when a study indicated that Robo4 can also exist in the newborn cerebral cortex to regulate the radial migration of newborn neurons [25]. Although Robo4 can be coimmunoprecipitated together with Slit2 in earlier studie [20], latter it was proved that it cannot bind to Slits directly but binds to the complex of Slit2 and Robo1 using biostructural techniques [11, 26]. Nevertheless, another studie showed that Robo4 as a ligand binds to UNC5B (a netrin receptor, also called Protein unc-5 homolog B) and inhibits VEGF-induced angiogenesis and vascular permeability, not as a receptor binds to Slit2 [27]. Meanwhile, some researchers proposed that Robo4 transduces signals through the development of a coreceptor with other molecules, such as Robo1, heparan sulfate proteoglycans (HSPGs) [11, 28]. Like the controversial results of Robo4, current evidences suggested that the role of Robo3 is also uncertain. For example, Zelina et al. contended that Robo3 cannot bind to Slits because of its specific difference in the Ig1 region in axon guidance but binds to Netrin-1/ Deleted in colorectal carcinoma (DCC) [29]. Another study observed that antagonized the Slit2-Robo1/2-induced Robo3 repulsion effect by binding to to neural EGFL Like 2 (NELL2), not by binding to Slit or Netrin-1, allowing the commissural axons to pass through the midline [30]. Therefore, Robo4 and Robo3 are two unique receptors which were endowed with much uncertainty, how and whether it binds to Slit ligand in regulating cell activities is yet to be determined.

Slit/Robo signaling pathway

Slits interact with Robos and subsequently play an important role in muscle cell formation, cell migration, stem cell growth, angiogenesis, organ development, and tumor formation by recruiting different adaptor molecules or proteins to cause a cascade of signaling pathways, but little is known about how exactly Slit binding to Robo is transmitted across the cell membrane. Many analysises showed that the ectodomain (ECD) is mainly responsible for Robo1-Robo1interactions [31, 32], and further studies indicate that this was largely mediated by the Ig domains of Robo1/2 in vitro [21]. Aleksandrova N recently indicated that the Robo1 ECD folds back on itself in a looping configuration, thereby forming a larger tetrameric structural arrangement consisting of "dimer-of-dimers" in a putative inactive а conformation [33]. The result was consistent with a mechanistic model in which a Slit2-N-induced conformational change of Robo1/2 is required for receptor activation [21, 31, 32].

However, increasing evidences suggested that Slit binds to other molecules, receptors or coreceptors in addition to Robo. Heparan sulfate proteoglycan (HSP) binds to both Slit and Robo to form a ternary complex to stabilize their interaction at the membrane [34]. Slits can also bind to extracellular matrix (ECM) molecules such as Plexin A [15], IV collagen [35], Netrin-1 [19], and dystroglycan [36]. Additionally, another study demonstrated that Slit combines with Dscam1, subsequently enhances the binding between Down syndrome cell adhesion molecule 1 (Dscam1) and Receptor Protein tyrosine phosphatase 69D (RPtp69D), and promotes Dscam1 dephosphorylation, which ultimately promotes axon formation (Figure 2A) [37]. In terms of the function of combining with other ECM, Barak et al. investigated the Robo1 membrane-proximal domain and hypothesized that the combination of Slit and extracellular matrix molecules increases intermolecular tensions and allows Slit to bind tightly to the juxtamembrane region of Robo1 [38].

A similar phenomenon was also shown in numerous reports for Robo, which binds to other molecules apart from Slit, such as fibronectin leucine rich transmembrane protein 3 (FLRT3), NELL2, Unc5B [27, 30, 39]. The ability of Robo to bind to other molecules will be elucidated in following parts of our review and so will not be discussed further here. Overall, the variable combinations of Slit and Robo depending on different environments may explain the complexity of the Slit/Robo signaling pathway and may provide a rational explanation for the paradoxical observations in different studies.

Downstream targets of Slit/Robo

The signaling molecules downstream of Robos are mainly cytoplasmic kinases, and regulatory molecules associate with actin polymerization and microtubule cytoskeleton reorganization (Figure 4), which all contribute to impact the cell mobility, including kinases comprising Hakai, Myo9b, and GTPases containing the Rho-family (Rac, Cdc42 and RhoA), and some key regulators of cytoskeleton, like Abl, Ena.

Ab1 is actual involved in the Robo signaling pathway. It was recently demonstrated that Abl tyrosine kinase (Abl) can inhibit Robo signaling by phosphorylation of the Robo CC1 domain after binding to Robo [40] and promotes Robo signaling by binding to Capulet protein or Cables protein to affect the activity of β -catenin and N-cadherin, eventually alter cell adhesion [41, 42]. P-cadherin can also be impacted by Robo3 to regulate cell-cell adhesion [43]. GTPases are small GTP-binding proteins that regulate cell polarity and cell motility by modulating the cytoskeleton. The activity of GTPases is regulated by different functional proteins, such as Dock/Nck (Nck in mammals), s/rGAPs (Slit/RoboGTPase activating proteins) and GEFs (guanine nucleotide exchange factors), all which can be recruited by Robos in published papers [44-46]. The enabled (Ena) protein, a family of proline-rich proteins that positively regulate cellular actin formation and cell motility, which plays an important role in the axonal repulsion of filopodia assembly and extension mediated by Slit [47, 48].

There are de novo targets published recently that Slit/Robo can bind and recruit to promote cancer cell migration and weaken cancer cell adhesion, such as a ubiquitin kinase Hakai, which leads to the ubiquitination of E-cadherin, and Myo9b (or myosin IXB), which inhibits the growth and migration of lung cancer cells by inhibiting RhoA protein [49, 50]. Altogether, it is well-estimated that these melocules like Abl, Ena, GTPases and other targets of Slit/Robo regulate cell motility and migration. Hence, Slit/Robo may be a potential therapeutic target to impede disease development, like tumor.

Regulatory molecules of Slit/Robo

Many studies demonstrated that Slits/Robos are regulated at either the genetic or protein level in a variety of ways, such as by DNA, RNA and protein modification (Figure 5), which plays an important role in physiologic and pathologic processes.

The promoter region of the Slit and Robo genes are frequently downregulated via hypermethylation in tumors, including non-small cell lung cancer, breast cancer, glioma, hepatocellular carcinoma, colorectal cancer, and leukemia [51-57]. In addition, the expression of Slit2 can also be downregulated by the epigenetic modification enzyme EZH2 (enhancer of zeste 2 polycomb repressive complex 2 subunit) in prostate and brain cancer [58, 59].

The result first reported in our laboratory that Robo1 and Robo2 are negatively regulated by miR-218 [60-62], was eventually verified by many other investigators [53, 56, 63, 64]. More recently, it was shown that Robo3 is negatively regulated by miR-383 and Robo1 is suppressed by miR92 [65, 66]. Robo2 protein may be a downstream target of the JAK/STAT pathway [67]. It has reported that Robo3 can antagonize Robo1/2 signaling [30], while Robo1/2 and Robo3 have synergistic effects in another study [68], so it can speculate that molecules unrecognized in between can regulate Robo. Moreover, Ubiquitin-specific protease 33 (USP33), a deubiquinating enzyme, stalblizes Robo1 by deubiquitinating it in lung cancer and colorectal cancer [55, 69]. Like the expression of Slit downregulated by hypermethylation, there are also reports showed that Robo can be hypermethylated in a few of cancer diseases [70, 71]. In summary, these datas suggested that Slits and Robos can be regulated in many patterns to influence cell activities. Given that Slit/Robo plays an important role in tumor or dysplastic disease development, it is rational to speculate that upstream molecules may also be a cluster of promising targets to restrain these diseases' progression.



Figure 4. Downstream targets of Slit/Robo. Slit/Robo signaling participates in various activities through different kinases to influence cell motility. These molecules include Ab1, GTPases, Ena, Hakai, and Myo9b.



Roles of Slit/Robo in cell motility

protein modification.

Slit/Robo was first reported in the nervous system and acted as an extracellular signpost to guide neuronal axon path finding, branching and control neuronal migration. Cells that are positive for Slit protein attract or repulse cells positive for Robo protein in one direction, which leads to cell migration. However, the function of Slit/Robo has been found to extend far beyond neurogenesis, including neurocyte development, angiogenesis, leukocytic chemotaxis, and cancer metastasis. The following section provides an overview of how this ligand-receptor functions in a variety of cells, which can be summed up as the roles of Slit/Robo in cell motility.

The role of Slit/Robo in neurocyte navigation

Nervous system development requires that the axons of neurocytes navigate to their correct location. Pathfinding is directed by molecular cues sensed by receptors in the axonal growth cone. Slits normally transmit signals through Robo family receptors. All three Slits are expressed by the floor plate, and commissural neurons express Robo1, Robo2, and Robo3. Slits bind Robo1 and Robo2 with high affinity to mediate neuro axon migration [7, 66, 72-74]. Interestingly, several developments suggested that untraditional receptors or coreceptors bind with ligands to regulate neuron axguidance (Figure 6).

Recently, one study established that Plexin A1, a well-known semaphoring receptor, is also a receptor for Slits during commissural axon guidance and reported for the first time that the C-terminal fragment generated by Slit processing has crucial bioactivity in axon branching [15]. Additionally, in the formation of thalamocortical connections, FLRT3 has been identified as a coreceptor for Robo1 and can modulate axonal responsiveness to Netrin [39]. Although Robo3 mentioned above does not bind to Slit proteins due to its specific difference in the Ig1 region [29], it was even reported that Robo3 can regulate midline crossing by antagonizing Robo1/2-mediated repulsion from midline-expressed Slits and potentiating DCC-mediated midline attraction to Netrin-1, without binding to Slits [29, 75]. Another unhomogeneous ligand was also reported to bind to Robo3 to control midline crossing, like NELL2 [30]. Robo4, which was initially considered to be solely expressed in endothelial cells, was not identified as a regulator in the neuronal system until a study argued that Robo4 is expressed in the developing brain and regulates the radial migration of newborn neurons in the neocortex [25]. Taken together, it is undoubted to conclude that Slit/Robo signaling de facto takes part in the growth of neuro axons through either its cognate or nonhomologus counterparts in different contexts. Notably, more determining researches should be conducted to make it as a possible potential target in neurological disorder.

The role of Slit/Robo in angiogenesis

Normally, Slit2 and Slit3 are secreted by vascular muscle cells, endothelial cells smooth and perivascular cells. Robo4, Robo1 and Robo2 are expressed in all kinds of endothelial cells [1]. Slit/Robo signaling regulates angiogenesis by altering endothelial cell motility and polarity. An increasing

number of studies highlighted the role of Slit/Robo signaling during angiogenesis and the formation of blood vessels from existing vessels, but the conclusions drawn were not completely consistent with each other (Figure 7).

For a long time, the role of Slit/Robo signaling in angiogenesis was deemed to be to promote angiogenesis by binding to Robo1 or Robo1/Robo4 heterodimers [26, 76-79]. Investigations carried out in our laboratory also proved that Slit2/Robo1 triggers angiogenesis in gastric cancer [61]. Nevertheless, the opposite effect of inhibiting angiogenesis was later reported for Slit2/Robo4 signaling [80-83]. One research viewed that the angiogenic effect of Slit2 on endothelial cells is related to the ratio of Robo1 and Robo4 in endothelial cells [84]. These researchers found that Slit2 promoted the migration of human umbilical vein endothelial cells (HUVECs) by binding to Robo1 only after Robo4 was knocked down [84]. However, another explanation for why Slit2 plays two



Figure 6. Roles of Slit/Robo in neurocyte navigation. Robo receptors can interact with other receptors, such as FLRT3, DCC, and NELL2, and form homodimers and/or heterodimers to elicit their functions in different contexts.



Figure 7. Roles of Slit/Robo in angiogenesis. Slits/Robos can alternatively bind with Robo1 or Robo4 or cooperate with VEGF to balance vascular vessel development.

implausible contrary roles in angiogenesis, proposed by Dunaway, C.M. is that Slit2 alone promotes tumor angiogenesis but suppresses tumor angiogenesis when coexisting with ephrinA1 [85]. In addition to the role of Slit2 in angiogenesis, Slit3 had been shown to induce endothelial cell migration through Robo4, which activates Rac1 and RhoA [86]. These envidences indicated that Slit/Robo signaling can either promotes angiogenesis or inhibites angiogenesis via different receptors.

Other angiogenetic factor like VEGF/VEGFR can be impacted by Slit/Robo signaling in angiogenesis. For example, Slit2/Robo4 can inhibit VEGF-induced endothelial permeability and maintain vascular stability by recruiting GAPs to inhibit the activity of Arf6 and Rac [87]. Another study further viewed that Robo4 binds UNC5B and inhibits VEGF/VEGFR signaling, thus stabilizing blood vessels and inhibiting angiogenesis [27].

Interestingly, Slits expressed by different cell sources have distinct functions in angiogenesis. Marlow R et al found that vascularization of the breast gland is not affected by loss of Slit expression in the epithelial compartment [81]. Instead, they identified a stromal source of SLIT, mural cells encircling blood vessels, and showed that loss of Slit in the stroma leads to elevated blood vessel density and complexity [81]. Eventually, they clarified that Slit2/Slit3 expressed by mural cells encircling blood vessels in the stroma, rather than those expressed by the epithelium, inhibits VEGF-induced angiogenesis via Robo4 but not Robo1 [81].

Together with these findings, it is difficult to determine exactly the regulating role of Slit/Robo in angiogenesis. In view of studies had shown that Slit/Robo signaling inhibits VEGF-induced vascular formation and so another method can be utilized for VEGF targeting drug resistance in many carcinomas' treatment, such as bevacizumab and sorafenib.

The role of Slit/Robo in leukocyte chemotaxis

Slit was initially identified as an inhibitor of leukocyte chemotaxis by Wu et al [88]. They used transwell migration assays to domenstrate that lymphocytes and leukocytes in response to various chemoattractants is inhibited by Slit2 [88]. Tole S et al. further domenstrated that Slit2 selectively impaired neutrophil migration toward other chemoattractants, namely, C5a and IL-8, and inhibited neutrophil chemotaxis by preventing chemoattractant induced actin barbed end formation and cell polarization [89]. In the model of renal ischemia-reperfusion injury, Slit2 blocked the capture and firm adhesion of human neutrophils to the inflamed vascular endothelial barrier by suppressing the inducible activation of Cdc42 and Rac2, which are critical mediators for cell migration [90]. Another observation in a mouse model that Slit2 significantly reduces the recruitment of neutrophils to the site of inflammation is in accordance other with inflammation models. including glomerulonephritis-associated kidney injury, global cerebral ischemia, and skin sensitization to allergens [91-93]. Interestingly, one study from a different perspective found that Slit/Robo4 strengthens the vascular barrier and diminishes deleterious aspects of the host's response to the pathogen-induced cytokine reaction [94]. Collectively, Slit/Robo these evidences showed inhibits chemotaxis of leukocytes toward chemoattractants, and it may have a therapeutic role as an inhibitor of inflammatory cell infiltration.

On the contrary, a recent paper suggested that Slit2 can also cause chemorepulsion of human neutrophils other than chemoattraction [95]. In their report, the authors showed that Slit2 comprises ~140-kDa N-terminal Slit2 fragment (Slit2-N), which serves as a chemoattractant for human neutrophils, and ~110-kDa N-terminal Slit2 fragment (Slit2-S), which serves as a chemorepellent for human neutrophis [95]. They further found that the effects of both Slit2 fragments were blocked by Abs to the Slit2 receptor Roundabout homolog 1 or the Slit2 coreceptor Syndecan-4, but involved in different intracellular signaling pathways [95]. These evidences remind of us that numerous works are needed to determining the clear role of Slit/Robo in leukocyte regulation.

The role of Slit/Robo in cancer metastasis

Like the role of Slit/Robo signaling in axon guidance, in which the proteins have dual attraction and repulsion effects, the role of Slit/Robo in cancer cell migration varies in certain contexts. A variety of studies showed that the expression of Slit is downregulated or not detected in most tumors, including breast cancer [96], gastric cancer [53], lung cancer [97], liver cancer [98], esophageal cancer [54] and others [99], and is largely related to promoter hypermethylation, indicating the inhibitory effect of Slit/Robo in these cancers, which is consistent with our studies in gastric cancer [60-62]. In contrast, overexpression of Slit and Robo appear in some tumors, such as melanoma [76], gastric cancer [100], pancreatic cancer tissues and cell lines [65], and hepatocellular carcinoma [101], which demonstrates that Slit/Robo signaling has a facilitating effect in certain cancers. The process of migration and metastasis of tumor cells was driven by stimulatory molecules such as Slit/Robo within the environment, which involves the process of reducing adhesion between cells, reassembling the cytoskeleton, inducing chemotaxis, enhancing angiogenesis, interrupting lymphogenesis and innervation.

Emerging evidences suggested that Slit/Robo signaling alters tumor cell-cell adhesion by regulating the connection between E-cadherin, a signaling target that is important for the maintenance and stability of cells, tissues and organs, and β -catenin, a signaling target that translocates into the nucleus to activate Lef/Tcf after activating [42, 49, 65, 97, 102, 103]. Of note, P-cadherin was also involved in regulating cell-cell adhesion by combining with Robo3 in oral squamous cell carcinoma [43].

In addition to reducing cell-cell adhesion, the assembly of cytoskeletal actin and the dissolution of the extracellular matrix can be impacted by Slit/Robo signaling to regulate cancer cell metastasis. Parray et al. used wound healing migration assay to demonstrate that prostate cancer cell migration was inhibited by Slit2 through inactivating Rac protein in the experimental model of Robo1 mutations in the C2 and C3 loci [70]. Another study suggested that Slit2 inhibits esophageal cancer cell metastasis by inhibiting Cdc42, FAK and Paxillin protein activities [54]. We and Kong et al. showed that in lung cancer, Slit2/Robo1 inhibited lung cancer cell migration via inhibiting the Myob/RhoA signaling pathway [50]. Meanwhile, Yuan, M et al. showed high expression of slit2 and down-regulation of Robo1 markedly enhanced migration of hepatic cancer cell [104]. The authors demonstrated that Robo1 overexpression upregulated matrix metalloproteinase (MMP)-2, MMP-9, and membrane-type1 MMP (MT1-MMP) expression, thereby promoting tumor metastasis [104]. In malignant melanoma, MMP2 and MMP14 was showed to be downregulated by Slit3/Robo2 and thereby inhibiting cancer cell migration [105]. They demonstrated that by binding to Robo3, the Slit3 suppressed the expression of AP1 and then downregulated its targeted genes expression [105].

To survive under hypoxic hazardous, conditions, cancer cells shift their metabolism pattern, known as Warburg effect (or hypoxic adaption) to meet demands of sustainable growth and metastasis. By knocking down Slit2 expression, Shi, R. L et al. demonstrated that Warburg effect was utilized by Slit2 to inhibit the metastasis of thyroid cancer cells [106]. More recently, Jeon MJ et al. observed a similar role of Slit in thyroid cancer cell migration that Slit3 also involved in inhibiting thyroid cancer cell migration but by regulating beta-catenin and Rho GTPase activity [107].

Additional mechanisms were also reported in papers. CXCL12/CXCR4- induced breast cancer cell chemotaxis, chemoinvasion and the adhesion, the foundmental components that promote metastasis was inhibited by slit2 [108, 109]. The authors revealed Slit2 inhibits CXCL12-induced that tyrosine phosphorylation of focal adhesion components such as RAFTK/Pyk2 at residues 580 and 881, focal adhesion kinase at residue 576, and paxillin [108]. Another study demonstrated that in breast cancer patients, glioblastoma cells with high Slit2 promoted the migration of siRobo1/breast cancer cells compared with the Slit2 absent group [110]. Thereby, the authors hypothesized that high level of Slit2 in brain serving as a chemoattractant to attract breast cancer cells with lower level of Robo1 [110]. Taken together, these evidences suggest that Slit can impact other chemokines or serve as chemokine to regulate cancer cell migration other than traditional pathway.

By using immune competent Robo4 knockout mouse model, Robo4 was showed that it played a very important role for suppressing breast cancer growth and metastasis by regulating tumor angiogenesis, endothelial leakage and tight junction zonula occludents protein protein (ZO-1) downregulation [82]. Due to the lymphatic system plays critical roles in the maintenance of fluid homeostasis, immune response, and tumor al. metastasis, Yu, J. et demonstrated that Slit2N/Robo4 modulates lymphatic dysfunction characterized by VEGF-C/VEGFR3 activation and this ligand-receptor pair may be a potential drug target to inhibit cancer metastasis [111]. Meanwhile, another study suggested that Slit2 Inhibits Neural Invasion and Metastasis in Pancreatic Cancer [112]. The authors demonstrated that SLIT2 mRNA expression was reduced in Pancreatic ductal adenocarcinoma (PDAC) compared with nontransformed pancreatic tissues and cell lines, suggesting a reduction in Slit2/Robo pathway activity in PDAC [112]. In according with the interpretation, restoring the SLIT2 expression in SLIT2- deficient PDAC cells inhibited their bidirectional chemoattraction with neural cells, and impaired unidirectional PDAC cell navigation along outgrowing neurites in models of neural invasion [112]. Collectively, these studies revealed that Slit/Robo can also impact cancer metastasis indirectly by interrupting angiogenesis, lymphogenesis and innervation.

Conclusions and perspectives

From the findings mentioned above, despite the high diversity of functional processes mediated by Slit/Robo signaling in different cells, it is reasonable to speculate that the regulation in various activities by Slit/Robo are correlated with altering cell motility. The reason that complex phenotypes related to Slit/Robo signaling can be attributed to the innumerous ligands or receptors involved in this pathway in different environments, and diverse outcomes could be caused by various molecules participating in this signaling pathway.

Encouragingly, several reports revealed beneficial outcomes by using Robo mAb therapy to treat cancer and vascular diseases. For instance, Fujiwara et al. uncovered a significant therapeutic effect by intravenous injection of Robo1 mAb in mouse hepatoma and non-small cell lung cancer tumor models [113, 114]. Mounting studies have demonstrated the potential anti-angiogenic role of Slit2 in corneal and retinal neovascularization, endometriosis and renal ischemia-reperfusion injury [77, 79, 90, 115]. Therefore, the Slit/Robo signaling pathway might be a promising target in therapy for tumor or intractable diseases, like VEGF/VEGFR or EGF/EGFR. Further investigations should focus on context- and cell-dependent downstream signaling pathways that underlie cellular responses in different environments. Additionally, more research is needed to interpret the potential therapeutic effect of the Slit/Robo signaling pathway in animal and human genetic models because gene modification is the most vital experiment in studying genetic function.

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Competing Interests

The authors have declared that no competing interest exists.

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