

Deubiquitinating enzyme USP33/VDU1 is required for Slit signaling in inhibiting breast cancer cell migration

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Slit regulates migration of not only neurons, but also nonneuronal cells, such as leukocytes and cancer cells. Slit effect on cancer cell migration has not been well-characterized. In this study, we used several different assays to examine Slit effect on breast cancer cell migration in vitro. We show that ubiquitin-specific protease 33 (USP33)/VDU1, originally identified as a von Hippel–Lindau tumor suppressor (VHL) protein-interacting deubiquitinating enzyme, binds to the Robo1 receptor, and that USP33 is required for Slit responsiveness in breast cancer cells. Slit induces redistribution of Robo1 from intracellular compartments to the plasma membrane in a USP33-dependent manner. Slit impairs directional migration of breast cancer cells without affecting their migration speed. This inhibitory effect is Robo-mediated and USP33-dependent. These data uncover a previously unknown function of USP33 and reveal a new player in Slit-Robo signaling in cancer cell migration.

cell migration and motility | metastasis | Slit-Robo signaling

Cell migration is a fundamental process critical for not only embryonic development but also homeostasis in adult animals. A number of molecular cues guide axons and migrating neurons (1–4). Recent studies suggest that molecular mechanisms modulating migration of cells in different tissues/organs are conserved. For example, guidance cues, receptors, and the intracellular signaling pathways for neuronal migration are also used for cells outside of the nervous system, ranging from immune cells, myoblasts, and endothelial cells to tumor cells (4–7). The *Slit* gene was first identified in *Drosophila*, and subsequent studies indicated that secreted proteins of the Slit family and their receptors of the Roundabout (Robo) family play important roles in neuronal guidance (8–13). The *Slit* genes are frequently inactivated in cancer (14–19).

Accumulating evidence supports that chemokines and their receptors play important roles in tumorigenesis and cancer metastasis, including chemotactic invasion and migration of cancer cells (20, 21). In some cases, cancer cells show increased expression of chemokine receptors not expressed in normal nontumor cells, providing a plausible explanation for distant metastasis to organs that secrete corresponding chemokine ligands (22). Studies from our group and other groups have shown that Slit suppresses chemokine-directed chemotaxis of leukocytes and breast cancer cells (23–25) and inhibits medulloblastoma cell invasion (26). These observations suggest a potential therapeutic strategy in controlling aberrant cell migration during cancer metastasis. Cellular and molecular mechanisms underlying Slit signaling in cancer cells remain to be elucidated.

To dissect the Slit-Robo signaling pathway, we carried out yeast two-hybrid screens by using the intracellular domain of Robo1 as bait (27). From such screening, we have identified ubiquitin-specific protease 33 (USP33)/von Hippel–Lindau tumor suppressor protein (pVHL)-interacting deubiquitinating enzyme 1 (VDU1) as a protein interacting with Robo1 intracellular domain. USP33 was originally discovered as a deubiquitinating enzyme (DUB) binding to pVHL-containing E3 ligase complex targeted for ubiquitin (Ub)-mediated degradation (28).

The substrates identified so far for USP33 and/or its homolog USP20/VDU2 are type 2 iodothyronine deiodinase (D2) and hypoxia-inducible factor-1 α (HIF-1 α) (29–31). The interaction between Robo1 and USP33 suggests that Robo1 is a previously unknown substrate for USP33. Our experiments demonstrate that in breast cancer cells, Slit inhibits chemokine-directed chemotaxis by affecting the directionality of cell migration, and that Slit responsiveness requires USP33. Our study reveals a functional link between USP33 and Robo1 and uncovers a previously uncharacterized role for USP33 in cancer cell migration.

Results

Slit Inhibits Breast Cancer Cell Migration Induced by SDF1 in a Robo-Dependent Manner. Our previous studies indicate that Slit inhibits the chemokine SDF1-induced leukocyte chemotaxis (23, 25). SDF1 activation has been implicated in breast cancer metastasis (22). To investigate the involvement of Slit in breast cancer metastasis, we examined expression of its receptor Robo1 and CXCR4 in tissue samples from primary breast cancer patients and two metastatic breast cancer cell lines: MDA-MB-231 and MDA-MB-435 (abbreviated as MDA231 and MDA435, respectively). Consistent with previous reports (22, 24), CXCR4 and Robo1 were expressed in cancer samples (Fig. 1A, lanes 1–6) and both of the cell lines (Fig. 1A, lanes 7 and 8). Robo1 expression was also confirmed by Western blotting using a specific anti-Robo1 antibody (Fig. 1B).

To examine the effect of Slit in cancer cell migration, we set up a transwell migration assay by using either MDA231 or MDA435 cells. Both types of breast cancer cells exhibited SDF1-induced chemotaxis in a similar manner (Fig. 1C). We therefore used MDA231 cells for subsequent experiments. SDF1 induced the migration of MDA231 cells, and this SDF1-induced chemotaxis was inhibited by Slit in a dose-dependent manner, although Slit did not significantly affect the baseline cell migration (Fig. 1D), consistent with a previous study (24). To test the involvement of Robo in Slit signaling in cancer cells, RoboN, a soluble extracellular domain of Robo blocking Slit-Robo signaling (12, 13, 23), was used together with Slit. Addition of RoboN effectively blocked the inhibitory effect of Slit on SDF1-induced chemotaxis (Fig. 1D), indicating that Slit signaling in MDA231 cells is mediated by Robo.

Slit Affects the Directionality of Cancer Cell Migration. Slit treatment inhibits SDF1-induced chemotaxis of cancer cells in the transwell assay (Fig. 1C and D). However, this assay does not allow direct visualization or detailed characterization of cell migratory be-

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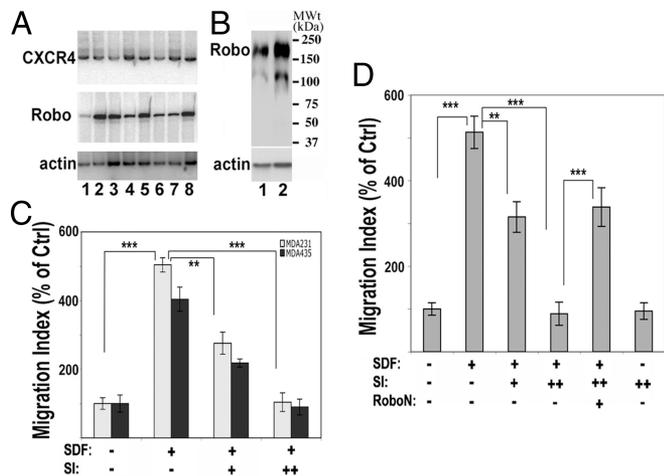


Fig. 1. Slit inhibits SDF1-induced chemotaxis of breast cancer cells. (A) Expression of CXCR4 and Robo1. Primary breast cancer samples (lanes 1–6) as well as cell lines (MDA231 and MDA435, lanes 7 and 8, respectively) were examined by RT-PCR with primers specific for CXCR4 or Robo1. Actin was used as a control. (B) Western blotting was used to detect Robo1 protein in MDA231 and MDA435 cells (lanes 1 and 2, respectively). A band of ≈ 210 kDa (the full-length Robo1 protein) and a lower band (a possible proteolytic product) were detected in these cells. (C and D) Slit2 inhibits cell migration in a Robo-dependent manner. In C, MDA231 and MDA435 cells were tested in a transwell migration assay in the presence of the vehicle control, SDF1 (10 nM), Slit2 (+, 30 pM; ++, 100 pM), or combinations. The cell migration index was measured as the ratio of cells that migrated into the lower well in the test group to those in the control group with only vehicle treatment. In D, similar transwell experiments were carried out by using MDA231 cells in the presence or absence of RoboN (10 nM). In C and D, data are presented as the mean \pm SEM, representing the results from three independent experiments, and the differences between the marked groups are statistically significant (**, $P < 0.01$; ***, $P < 0.001$ by Mann–Whitney test).

haviors. We then established a wound-healing assay by using MDA231 cells grown to confluence on fibronectin-coated coverslips and tested Slit effect on wound-induced cancer cell migration. Immediately after wound formation by scratching the cell monolayer using a fine tip, cells were cultured in the presence of the control, SDF1, Slit, or combinations, and then subjected to time-lapse imaging to monitor cell migration. Wound scratching induced cell migration in the direction toward the center of the wound area. Directional cell migration was measured as the forward distance that individual cells migrated from the wound edge line at the zero time point. A uniform application of SDF1 did not show an additive effect on wound-induced directional cell migration. The treatment with Slit alone significantly reduced the wound-induced cell migration (Fig. 2A and C; see also [Movie S1](#) and [Movie S2](#)). Similar to that observed in the transwell assay (Fig. 1C), addition of RoboN blocked the inhibitory effect of Slit on wound-induced directional migration (Fig. 2A and C), indicating that Slit effect is Robo-dependent. Slit treatment did not significantly affect the cell migration speed (Fig. S1).

In a migrating cell, microtubules (MTs) are polarized, with the MT-organizing center (MTOC) positioned between the nucleus and the leading edge of cell migration (32, 33). We examined the effect of Slit treatment on MTOC orientation in the wound-healing assay by pericentrin immunostaining, as described previously (33, 34). At different time points, cells were fixed and stained by using an anti-pericentrin antibody to visualize MTOCs. In this experiment, the percentage of mock-treated cells exhibiting MTOCs redirected toward the wound edge increased during the first 3 h, reaching a plateau of $\approx 60\%$ (Fig. 2B and D). The percentage of cells with the polarized MTOCs in the Slit-treated group similarly increased during the first 3 h

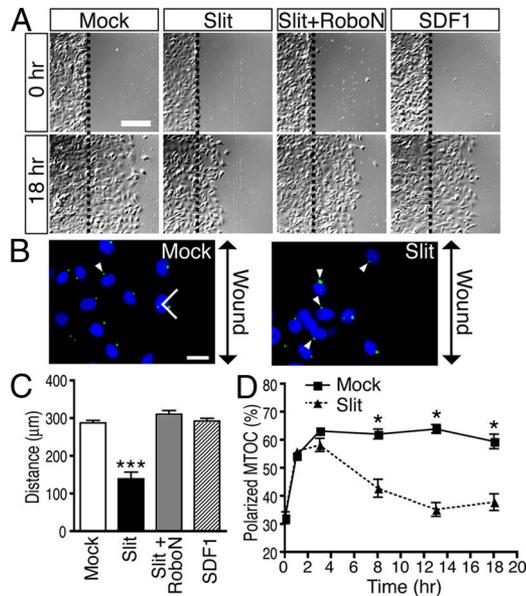


Fig. 2. Slit impairs wound-induced migration of cancer cells. (A) Time-lapse imaging in a wound-healing assay of MDA231 cells was performed in the presence of mock control, Slit2 (25 pM), RoboN, or SDF1 (15 nM). Wound scratches were made so that the cells at the wound edge were migrating toward the right side in each panel. Representative images taken at 0 and 18 hr after wound scratching are shown. The black dotted lines mark the wound edge at 0 hr. (Scale bar: 200 μm .) (B) Cells were fixed after 13 hr of incubation, and MTOC was visualized by immunostaining with an anti-pericentrin antibody followed by Alexa488-conjugated anti-mouse secondary antibody (green). Cell nuclei were stained with Hoechst 33342 (blue). Cells in the first and second rows at the wound edge that had their MTOC in the forward-facing quadrant toward the wound edge were counted as having polarized MTOC. Arrowheads in each panel mark the cells with nonpolarized MTOC. (Scale bar: 20 μm .) (C) Quantification of the forward migration distance of individual cells during the 18-hr imaging period. Dividing cells were excluded from the analysis. Data are presented as the mean \pm SEM ($n = 30$ from three independent experiments; 10 cells per experiment). ***, $P < 0.0001$ by Mann–Whitney test. (D) Time course of the ratio of cells with polarized MTOC among the total number of cells examined in the first and the second rows of the wound edge quantified from four independent experiments. The difference between the mock control and Slit-treated groups is significant at 8, 13, and 18 hr (*, $P < 0.05$, Mann–Whitney test).

but reduced to less than 40% afterward, with the maximal inhibitory effect ≈ 13 h after Slit treatment (Fig. 2B and D).

Although the wound-healing assay allowed direct visualization of cell migration, it is difficult to monitor individual migrating cells and measure their trajectories. To examine the migrating behavior of isolated individual cells, we established a Dunn chamber chemotaxis time-lapse microscopy assay (35) using MDA231 cells (Fig. 3A). We first tested whether Slit acted as a repellent for cancer cells, as was reported for neurons (13, 36). When placed under a Slit concentration gradient, MDA231 cells did not show repulsive response or change in migration speed (Slit, $0.40 \pm 0.03 \mu\text{m}/\text{min}$ vs. control, $0.42 \pm 0.03 \mu\text{m}/\text{min}$; $n = 30$; not statistically significant). Thus, Slit treatment alone did not affect the migration direction or speed of MDA231 cells.

We next tested MDA231 cells in SDF1 gradients in the presence of mock or Slit protein preparation (Fig. 3A). Fig. 3B shows the distribution of cells in the absence or presence of Slit. It is important to note that if cells display completely random migration, 33% of cells should distribute in each 120° arc. The cells treated with the mock preparation migrated toward the SDF1 gradient with relatively straight paths, such that $77\% \pm 1\%$ of the cells were in the 120° arc facing the SDF1 source at the top of the y axis, compared with randomly distributed cells in the

(GFP-USP33), or control vector plasmids were cotransfected in combinations into human embryonic kidney (HEK) 293 cells. After immunoprecipitation using an anti-HA antibody, GFP-USP33 was detected only in the immunoprecipitates of the cells expressing both Robo1 and USP33 proteins, not of the control cells (Fig. 4A, lanes 1–3). Slit treatment did not affect Robo1–USP33 interaction (Fig. 4A, compare lanes 4 and 5 to lane 3). To test whether the endogenously expressed Robo1 and USP33 interacted with each other, we used MDA231 cell lysates for immunoprecipitation by using the control IgG, anti-Robo1, or anti-USP33 antibodies (Fig. 4B, lanes 2–4). Anti-USP33 antibody did not cross-react with Robo protein, but it coimmunoprecipitated Robo1 (Fig. 4B, lane 4), whereas control IgG did not precipitate Robo1 (lane 2), indicating that Robo1 interacts with USP33 in MDA231 cells. Detailed characterization of the domains involved in Robo1–USP33 interaction will be described elsewhere.

USP33 Is Required for Slit Signaling in Breast Cancer Cells. To test the role of USP33 in Slit signaling in cancer cells, we transfected MDA231 cells with siRNAs targeted against either *luciferase* (used as a control siRNA; ref. 40) or *USP33* (siUSP33). Two different siRNAs specifically against *USP33*, siUSP33 nos. 1 and 2, were efficient for knocking down USP33 expression (Figs. 4C and 6C and Fig. S2C). Data obtained by using siUSP33 no. 2 are presented in Fig. S2. In the subsequent experiments, siUSP33 no. 1 was used. Immunoblotting experiments confirmed that siUSP33 efficiently and specifically inhibited USP33 expression in both HEK293 cells and MDA231 cells (Fig. 4C and Fig. S2C), reducing USP33 expression to $20.7\% \pm 0.6\%$ of the level in control siRNA-transfected MDA231 cells. In the Dunn chamber chemotaxis assays, $67\% \pm 1\%$ or $49\% \pm 2\%$ of control siRNA-transfected cells treated with mock or Slit preparation, respectively, ended up within the 120° arc facing the SDF1 source, similar to the parental cells (Fig. 5A). By contrast, $78\% \pm 4\%$ or $74\% \pm 3\%$ of USP33 knockdown cells that were treated with mock or Slit preparation, respectively, distributed within the arc facing the SDF1 source, indicating that siUSP33, but not control siRNA, eliminated Slit effect in inhibiting SDF1-induced chemotaxis (Fig. 5A). Moreover, tracking the centroids of the migrating cells showed significantly more straight trajectories of the USP33 knockdown cells treated with Slit compared with the control siRNA-treated cells exposed to Slit (FMI: siUSP33, 0.52 ± 0.05 vs. control, 0.17 ± 0.04 , $P < 0.001$; Fig. 5B and D). Therefore, siUSP33, but not control siRNA, specifically abolished Slit activity in inhibiting SDF1-induced chemotaxis. The siUSP33-transfected cells showed higher FMI than the control siRNA-treated cells in the absence of Slit (SDF1 plus mock) (siUSP33, 0.56 ± 0.03 vs. control siRNA, 0.40 ± 0.06 ; $P < 0.05$; Fig. 5B and D). Knockdown of USP33 did not significantly affect the migration speed of these cancer cells (Fig. 5C). In addition, wound-healing experiments with MDA231 cells transfected with two independent siRNAs directed against *USP33* revealed that the inhibitory effect of Slit on wound-induced directional migration depends on USP33 (Fig. S2). Therefore, USP33 is required for Slit signaling in the cancer cells.

Slit Stimulates the Redistribution of Robo to the Plasma Membrane in a USP33-Dependent Manner. To understand the role of USP33 in Slit signaling, we examined whether USP33 affected the Robo protein level. USP33 overexpression or knockdown did not affect the total Robo1 protein level as determined by immunoblotting (Fig. 4C and Fig. S2C), suggesting that USP33 may not regulate Robo1 protein stability. We then examined whether USP33 affected Slit responsiveness by altering the subcellular distribution of Robo1. In nontreated MDA231 cells, a significant portion of Robo1 immunostaining signal was in the perinuclear compartment, and only a low level of Robo1 signal was detected at the plasma membrane. Slit treatment stimulated redistribution

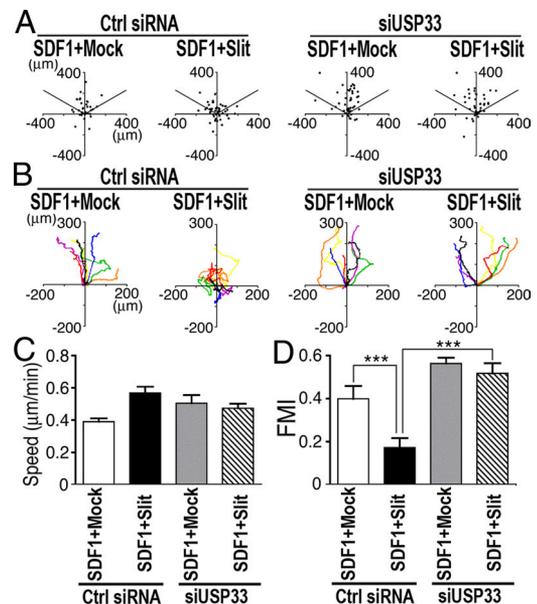


Fig. 5. Knocking down USP33 eliminates Slit activity in suppressing SDF1-induced chemotaxis. Dunn chamber chemotaxis assay was performed as described for Fig. 3, except that MDA231 cells were transfected with control or USP33-specific siRNAs. (A) Distribution of mock- or Slit2-treated cells under SDF1 gradients after transfection with the corresponding siRNA. (B) Representative trajectories of seven migrating cells expressing control siRNA or siUSP33, in the presence of the control or Slit2 under SDF1 gradients. (C and D) Quantitative analyses are shown in C (migration speed) and D (FMI). Data are presented as the mean \pm SEM ($n = 30$ from three independent experiments; 10 cells per experiment). ***, $P < 0.001$ by Mann–Whitney test.

of Robo1 immunostaining signals from the perinuclear region to the plasma membrane. Robo1 redistribution was not affected by the control siRNA (Fig. 6A Top). In the control siRNA-transfected group, Slit treatment induced a ≈ 3 -fold increase in the Robo1 signal at the plasma membrane, whereas in the siUSP33-treated cells, there was no significant change in Robo1 immunostaining signal at the plasma membrane upon Slit stimulation (Fig. 6A and B). Thus, siUSP33, but not control siRNA, abolished Slit-induced Robo1 redistribution process, with the majority of Robo1 signals remaining in the perinuclear region (Fig. 6A Bottom).

By extracellular biotinylation, we examined the surface levels of Robo1 in control and USP33-depleted HEK cells. After Slit stimulation, cell surface proteins were immediately biotinylated at 4°C and were collected by using avidin-immobilized agarose beads. As shown in Fig. 6C, lanes 1 and 2, in control cells, the surface level of Robo1 increased by ≈ 6 -fold 5 min after Slit stimulation. In contrast, Slit-induced up-regulation of Robo1 surface presentation was dramatically inhibited by siUSP33 (Fig. 6C, lanes 3 and 4). Together, these results reveal an essential role for USP33 in Slit-induced Robo redistribution from the intracellular compartments to the plasma membrane.

Discussion

Slit Signaling in Breast Cancer Cells. In this study, we used several assays to characterize Slit effect on cancer cell migration in vitro. Consistent with a previous study using a transwell assay (24), Slit inhibited SDF1-induced cancer cell migration. The transwell assay allowed examination of Slit effect on a large number of cells at the same time. The wound-healing assay enabled us to examine Slit effect in cancer cells in monolayer cultures. The Dunn chamber time-lapse microscopy allowed us to directly visualize migration of isolated cells and investigate Slit effects on chemotactic behaviors of the cells, with quantitative measure-

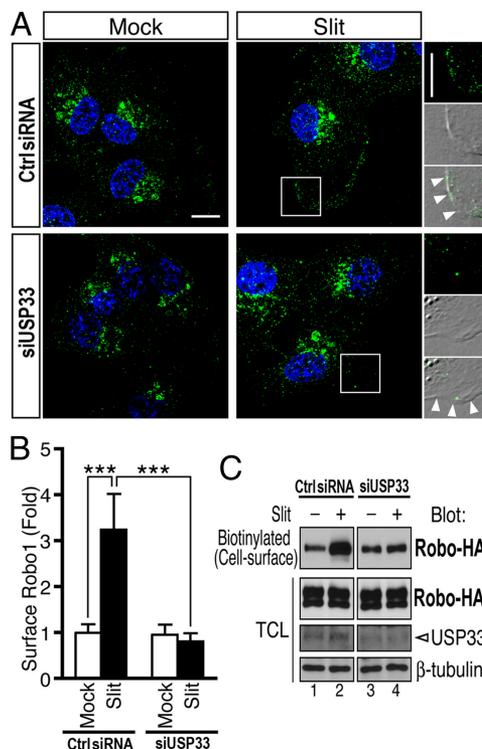


Fig. 6. USP33 is required for Slit-induced Robo1 relocation to the plasma membrane. (A) Robo1 immunostaining of MDA231 cells transfected with either control or USP33 siRNA at 10 min after Slit stimulation (25 pM), with Robo1 signals in green and nuclear staining in blue (Hoechst 33342). Images represent two-dimensional images of three-dimensional reconstruction. (Right) High-power views of the boxed areas, merged with differential interference contrast (DIC) images, showing Robo1 signals distributed at the plasma membrane (arrowheads). (Scale bars: 10 μ m.) (B) Quantification of surface levels of Robo1 in MDA231 cells. The ratio of Robo1 signal intensity at the plasma membrane to its total signal intensity in the entire cell was normalized to that in mock-stimulated control cells. To quantify cell surface-localized versus intracellularly localized Robo1, the plasma membrane area was manually marked by tracing the cell circumference. Twenty cells per group were analyzed. Data are presented as the mean \pm SEM. ***, $P < 0.001$ by Mann-Whitney test. (C) Cell surface biotinylation shows that Slit stimulation significantly increases cell surface expression of Robo1. Robo1-HA-expressing HEK cells transfected with control siRNA or siUSP33 were stimulated with Slit for 5 min. Extracellular domains were probed by cell surface biotinylation. Biotinylated proteins were then harvested by avidin-agarose pulldown and examined by Western blotting using anti-HA.

ments of migration speed and directionality. Slit clearly impairs directional migration induced by SDF1 or by wounding without affecting the migration speed. The loss of directionality in cell migration provides an explanation for a Slit inhibitory effect on cancer cells in the transwell assays (ref. 24 and the present study). It is interesting to note that a recent study suggested that N-terminal fragment of Slit may act as a chemoattractant for breast cancer cells in transwell assays (41). By using the full-length Slit protein, we did not observe such activity in the transwell assay.

Microarray studies implicate high levels of Robo1 expression in breast and liver cancers (42, 43) and suggest that the role of Slit-Robo signaling in cancer cells is complex and multifold, similar to netrin-deleted in colorectal cancer (DCC) signaling in cancer (7). Our data suggest that Slit plays a role in blocking chemokine-induced metastasis toward tissues expressing high levels of chemokines, such as SDF1. Consistent with this is the observation that Slit is frequently inactivated by methylation in cancer cells (15–19). These data support the role of Slit in tumor suppression. Recent evidence further indicates that Slit exerts the tumor-suppressive effect by regulating β -catenin (44).

The Role of USP33 in Slit Signaling in Breast Cancer Cells. The critical role of Ub-mediated modification in regulating protein stability/function has been well-established (45). Endocytic trafficking of receptors is also regulated by ubiquitination (46). DUBs have important regulatory roles in receptor trafficking and degradation of oncoproteins or tumor suppressor proteins (47–49). For example, USP7/HAUSP interacts with the tumor suppressor p53 and influences its activity (50, 51). Mutations in a tumor suppressor gene encoding the deubiquitinating enzyme CYLD have been identified in familial cylindromatosis. CYLD interacts with proteins regulating NF- κ B and TNF receptors (52–54). DUBs often act to protect target proteins from degradation. For example, AMSH inhibits down-regulation of EGF receptor signaling via deubiquitination in endosomes (55), although it is not clear whether EGF responses are completely abolished by AMSH depletion. In our study, USP33 interacts with Robo1 and is required for Slit-induced Robo1 redistribution to the plasma membrane without affecting the total Robo1 protein level. In contrast to the previously reported role of DUBs in preventing down-regulation of receptors and other substrate proteins by the Ub-mediated pathway (30, 31, 55), our data reveal a previously unknown activity of USP33 in modulating subcellular distribution of Robo.

Our understanding of functional relationships between the Slit-Robo and other signal transduction pathways in cancer cells remains limited. Together with previous studies (23, 24), our study supports a role for the Slit-Robo pathway in regulating cancer cell responses to chemokines. USP33 is a downstream target for ubiquitination by pVHL E3 ligase and is degraded via the Ub-proteasome system (28). It will be important to test in future studies whether the Slit-Robo pathway cross-communicates with those involving pVHL and its substrates, such as HIFs, and to address what role Slit signaling plays in tumorigenesis and tumor angiogenesis.

Materials and Methods

Cell Migration Assays. Transwell assay was carried out as described previously (23), except that 8- μ m pore polycarbonate filter membranes coated with fibronectin (5 μ g/mL) were used to separate the upper and lower wells. In wound-healing assay, MDA231 cells were seeded on fibronectin-coated coverslips and grown to confluence. A wound scratch was made on the cell monolayer by using a micropipette tip. After removing the dislodged cells by extensive washing with culture media, cells were treated with mock control or Slit. Cell migration was monitored by time-lapse microscopy, and the behaviors of individual, randomly selected cells were tracked by using ImageJ software (National Institutes of Health, Bethesda, MD). These data were processed with Microsoft Excel software and were subjected to statistical analysis using Prism 4.0b software (GraphPad). Dunn chamber chemotaxis assay was performed with modification of published protocols (35, 56).

MTOC Orientation Assay. Wound-healing assay was carried out as described above. After incubation at 37 $^{\circ}$ C for different periods of time, the cells were fixed in methanol (-20° C) for 5 min. MTOC was visualized by immunostaining using anti-pericentrin and Alexa488-conjugated anti-mouse secondary antibody. The cells in the first and second rows at the wound edge were scored for MTOC orientation (at least 150 cells in each group).

Immunoprecipitation Assay and Immunostaining. For a detailed description of experimental procedures, see *SI Materials and Methods*.

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