

Published in final edited form as:

Dev Cell. 2012 January 17; 22(1): 116–130. doi:10.1016/j.devcel.2011.10.030.

Phosphoinositide Signaling Regulates the Exocyst Complex and Polarized Integrin Trafficking in Directionally Migrating Cells

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Summary

Polarized delivery of signaling and adhesion molecules to the leading edge is required for directional migration of cells. Here, we describe a role for the PIP₂ synthesizing enzyme, PIPKIγ₂, in regulation of exocyst complex control of cell polarity and polarized integrin trafficking during migration. Loss of PIPKIγ₂ impaired directional migration, formation of cell polarity, and integrin trafficking to the leading edge. Upon initiation of directional migration PIPKIγ₂ via PIP₂ generation controls the integration of the exocyst complex into an integrin-containing trafficking compartment(s) that requires the talin-binding ability of PIPKIγ₂, and talin for integrin recruitment to the leading edge. A PIP₂ requirement is further emphasized by inhibition of PIPKIγ₂-regulated directional migration by an Exo70 mutant deficient in PIP₂ binding. These results reveal how phosphoinositide generation orchestrates polarized trafficking of integrin in coordination with talin that links integrins to the actin cytoskeleton, processes that are required for directional migration.

Keywords

Cell Migration; Phosphatidylinositol-4; 5-biphosphate; Cell Polarity; Integrin; Exocyst

Introduction

Cell migration is critical for many biological processes including embryogenesis, inflammation, and the metastasis of cancer cells. At the onset of migration, cells undergo a spatial reorganization of the cytoskeleton and membrane proteins to establish polarity (Insall and Machesky, 2009; Ling et al., 2006; Ridley et al., 2003; Rorth, 2009; Vicente-Manzanares et al., 2009). Coordinated cell migration hinges on the ability of cells to traffic signaling molecules and proteins toward the leading edge (Caswell and Norman, 2008; Fletcher and Rappoport, 2010; Ulrich and Heisenberg, 2009), a process that requires the tight regulation of cytoskeletal and vesicle trafficking machineries. The trafficking of newly

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synthesized or recycled integrin molecules to and from the plasma membrane is required for directional cell migration (Caswell and Norman, 2008; Caswell and Norman, 2006; Ulrich and Heisenberg, 2009). A prevailing theory is that migrating cells assemble adhesion sites at the leading edge and disassemble at the trailing edge resulting in a continual endo- and exocytosis of integrins (Bretscher, 1984, 1989; Ridley et al., 2003). Impairment of the endo-exocytic trafficking of integrins profoundly affects the polarity and directionality of cell migration (Caswell et al., 2009; Kuo et al., 2006; Nishimura and Kaibuchi, 2007).

Phosphatidylinositol-4,5-bisphosphate (PIP₂) is a lipid messenger that modulates many diverse biological processes including regulation of actin cytoskeletal dynamics, cell migration, cell-cell contact formation, endocytosis and exocytosis (Heck et al., 2007; Ling et al., 2006; van den Bout and Divecha, 2009). PIP₂ is a lipid messenger that is spatially and temporally generated, making it an ideal messenger for polarized signaling (Anderson et al., 1999; Heck et al., 2007; Ling et al., 2006). Type I PIPKs (α , β and γ isoforms) represent the predominant class of PIP₂ generating enzymes in mammalian cells (Anderson et al., 1999). The spatiotemporal generation of PIP₂ by the coordinated activity and/or recruitment of PIPKs and phosphatases is a central hypothesis in PIP₂ signaling (Anderson et al., 1999; Heck et al., 2007; Ling et al., 2006). PIPK γ has roles in vesicle trafficking both at the plasma membrane and in endosomal structures (Baird et al., 2006; Ling et al., 2007; Schill and Anderson, 2009a). In addition, PIP₂ generation is required for vesicle exocytosis (Hay et al., 1995; Martin, 1998) and endocytosis (Jost et al., 1998).

PIP₂ is generated at many cellular compartments, although its cellular content does not vary significantly suggesting that PIP₂ signals differently than other messengers (Anderson et al., 1999). The signaling specificity of PIP₂ is defined by the interaction of the PIP kinases with PIP₂ effectors or compartments containing PIP₂ effectors (Anderson et al., 1999; Heck et al., 2007; Ling et al., 2006; Schill and Anderson, 2009a). Multiple PIPK γ isoforms exist in mammals that have different C-terminal extensions and these sequences specifically interact with PIP₂ effectors and the PIP₂ generated regulates these effectors (Heck et al., 2007; Schill and Anderson, 2009b). Previously, PIPK γ 2 has been demonstrated to interact with adaptor molecules AP2 and AP1B, regulating the endocytosis and basolateral trafficking of E-cadherin molecules in polarized epithelial cells (Baird et al., 2006; Ling et al., 2007; Schill and Anderson, 2009a, b; Thieman et al., 2009). PIPK γ 2 is also specifically recruited to focal adhesions by an association with talin (Ling et al., 2002) and this requires the same sequence in the PIPK γ 2 C-terminus that interacts with the AP complexes (Baird et al., 2006; Ling et al., 2007; Thieman et al., 2009). PIPK γ 2 is also specifically required for chemotaxis towards growth factors and the interaction of PIPK γ 2 with talin appears to be required for chemotaxis (Sun et al., 2007).

The exocyst protein complex has a pivotal function in polarized trafficking of membrane proteins during cell migration (He and Guo, 2009). The exocyst complex consists of eight different subunits (Sec3, Sec5, Sec6, Sec8, Sec10, Sec15, Exo70 and Exo84) that mediate tethering of post-Golgi and endocytic recycling endosomes to the plasma membrane (He and Guo, 2009; Yeaman et al., 2001) and is associated with all stages of endosomal trafficking (Oztan et al., 2007). The exocyst complex is important for the polarized trafficking of LDL receptor, E-cadherin, integrin and Glut4-containing vesicles (Grindstaff et al., 1998; Inoue et al., 2003; Spiczka and Yeaman, 2008) and serves as an effector of the small GTPases Rab11 and Arf6 (Oztan et al., 2007). The small GTPases Rab11 and Arf6 also regulate integrin trafficking (Powelka et al., 2004). Two subunits, Sec3 and Exo70, directly interact with PIP₂ via conserved basic residues in their C-terminus suggesting that PIP₂ generation could be an important mechanism in regulating the exocyst complex in vesicle trafficking (Liu et al., 2007).

Here, we report that PIPKI γ 2 regulates the exocyst complex trafficking of β 1-integrin to the leading edge in directionally migrating cells. In this pathway, PIPKI γ 2 interacts with the exocyst complex and β 1-integrin upon initiation of directional cell migration and regulates β 1-integrin trafficking to focal adhesion complexes at the leading edge membrane. This requires an interaction between PIPKI γ 2 and talin.

Results

PIPKI γ 2 is Required for Directional Cell Migration

To define the mechanistic role of PIPKI γ 2 in cell migration, we specifically knocked down endogenous PIPKI γ 2 expression using a lentiviral vector-mediated delivery system. The expression levels of PIPKI γ 2 were reduced >90% using this approach (Fig. 1A). PIPKI γ 2 knockdown cells were morphologically indistinguishable from control cells and showed no obvious effect on cell proliferation (Fig. S1A, B). We quantified the impact of PIPKI γ 2 knockdown on cell migration using both wound-healing and haptotactic migration assays using a modified Boyden chamber. PIPKI γ 2 knockdown significantly impaired cell migration in MDA-MB-231 cells (Fig. 1B, C) and HeLa cells (Fig. S1D, E, F). Haptotactic cell migration was performed to assess the role of PIPKI γ 2 during integrin-dependent cell migration towards extracellular matrix (ECM) proteins. These data show that PIPKI γ 2 knockdown impaired cell migration toward fibronectin (FN) and collagen I (Col.I) (Fig. 1D), suggesting that PIPKI γ 2 knockdown could regulate integrin dynamics. The re-expression of PIPKI γ 2 but not a kinase dead mutant, rescued integrin-dependent cell migration (Fig. 1E). However, PIPKI γ 2 knockdown did not show any obvious defect on directionality nor velocity in non-directionally migrating cells (Fig. S1C and Movie S1, S2).

Knockdown of PIPKI γ 2, Exocyst Complex Components or Rab11 Impairs Polarized Recruitment of β 1-integrin and Cell Migration

During migration the reorganization of the actin cytoskeleton, microtubules and the Golgi apparatus induces polarity in the direction of migration, resulting in polarized membrane trafficking toward the leading edge (Caswell and Norman, 2008; Caswell et al., 2009; Ulrich and Heisenberg, 2009). In directionally migrating cells, PIPKI γ 2 is recruited to the leading edge (Fig. S1G). Knockdown of PIPKI γ 2 resulted in impaired actin assembly at the leading edge and impaired microtubule orientation (Fig. 1F). These cells also lost Golgi orientation in the direction of migration (Fig. 1F, G) indicating that PIPKI γ 2 is required for cell polarization during migration.

Polarization of cells during migration is regulated by vesicular trafficking, cytoskeletal dynamics, small G-proteins, and cell adhesion receptors (Caswell and Norman, 2008; Etienne-Manneville, 2008; Ridley et al., 2003). The endosomal recycling of integrin molecules controlled by Rab11 plays an integral role in polarity (Caswell and Norman, 2008; Powelka et al., 2004). Similarly, the exocyst complex has been implicated in polarized vesicle trafficking and integrin recruitment to focal adhesions (Spiczka and Yeaman, 2008). As PIPKI γ 2 modulates both focal adhesion dynamics and membrane trafficking (Bairstow et al., 2006; Ling et al., 2007; Ling et al., 2006; Sun et al., 2007) we used siRNA-mediated knockdown of PIPKI γ 2, Rab11, or Exo70 (Fig. 2A) to compare the role of each of these molecules in establishing polarity and cell migration. Individual knockdown of PIPKI γ 2, Rab11 or Exo70 similarly impaired cell orientation towards the direction of migration (Fig. 2B and 2C) and haptotactic cell migration towards FN, a β 1-integrin-dependent process (Fig. 2D).

The polarized trafficking of integrins is required for formation of nascent focal adhesion complexes and the stabilization of the leading edge in migrating cells (Caswell and Norman,

2008; Caswell and Norman, 2006; Choma et al., 2004). As shown in Fig. 2E, the knockdown of PIPKI γ 2, exocyst components or Rab11 disrupted focal adhesion complex assembly as evidenced by a loss of FAK at the migrating cell front. Further, the loss of PIPKI γ 2, Exo70 or Rab11 all impaired the polarized recruitment/trafficking of β 1-integrin to the leading edge (Fig. 2F, G). The phenotypes resulting from the loss of PIPKI γ 2 were specific, as knockdown of PIPKI γ 5 isoform had no impact on these processes (Fig. S2A, B, C, D and E). PIPKI γ 2 knockdown cells lost accumulation of β 1-integrin at membrane ruffles/protrusions and a loss of colocalization with cortactin (Fig. S3A, B). These data indicate a role for PIPKI γ 2 in the targeting of β 1-integrin to the leading edge in migrating cells.

The adhesion of cells to ECM protein mimic some events that take place in cell migration such as activation of integrins (Ginsberg et al., 2005). PIPKI γ 2 knockdown cells were morphologically indistinguishable from control cells and showed no obvious impairment in adhering or spreading when plated on FN (10 μ g/ml) or Col.I (20 μ g/ml) coated plates for 30 minutes (Fig. S3D, E). There were subtle changes in cell adhesion at lower FN concentration with shorter incubation time (Fig. S3F, G). These data are also consistent with a selective role for PIPKI γ 2 in polarized β 1-integrin trafficking and cell migration.

Cell Migration Stimulates a PIPKI γ 2 Association with β 1-integrin

In non-migrating confluent MDA-MB-231 or HeLa cells, PIPKI γ 2 and β 1-integrin are localized at the cell periphery in addition to intracellular compartments (Fig. 3A). Upon initiation of directional migration, PIPKI γ 2 and β 1-integrin re-localized to the leading edge and perinuclear vesicle-like compartments (Fig. 3A). Consistent with previous data, PIPKI γ 2 colocalizes with talin at focal adhesion complexes of migrating cell fronts and also the recycling endosome (Ling et al., 2007; Ling et al., 2002), but did not localize to the Golgi, early endosomes or lysosomes (not shown). PIPKI γ 2 also showed partial colocalization with Rab4 and Rab11-containing compartments (Fig. 3B), GTPases with established roles in endosomal recycling of integrins (Caswell and Norman, 2006; Powelka et al., 2004). At the onset of migration, a large increase in PIPKI γ 2 association with β 1-integrin and talin was observed in both MDA-MB-231 and HeLa cells (Fig. 3C). This is consistent with the colocalization of PIPKI γ 2 with these molecules.

PIPKI γ 2 directly interacts with talin, and talin directly associates with the cytoplasmic domain of β 1-integrin, therefore, a role for talin in mediating complex assembly between PIPKI γ 2 and β 1-integrin in migrating cells was examined. PIP₂ modulates the talin interaction with β 1-integrin (Martel et al., 2001) and consistent with this, PIPKI γ 2 knockdown severely impaired the association of talin with β 1-integrin in migrating cells (Fig. 3D). PIPKI γ 2 and β 1-integrin associate with talin's FERM domain (de Pereda et al., 2005; Ling et al., 2003; Wegener et al., 2007) and the interaction of PIPKI γ 2 with talin is required for chemotaxis (Sun et al., 2007). Talin forms a homo-dimer that would interact with both PIPKI γ 2 and β 1-integrin *in vivo* and talin also contains a second β 1-integrin binding site in the rod domain (Critchley and Gingras, 2008). GST-pull down approach was used to demonstrate that talin can bind both PIPKI γ 2 and β 1-integrin. For this GST-fused to the cytoplasmic domain of β 1 or α 5-integrin was purified and incubated with cell lysates prepared from cells expressing PIPKI γ 1 or PIPKI γ 2 or PIPKI γ 2Y649F mutant defective in talin binding (Ling et al., 2003). The GST- β 1 cytoplasmic domain pulled down both talin and PIPKI γ 2 but not PIPKI γ 1 (lacking the C-terminal talin binding region) or PIPKI γ 2Y649F indicating the requirement of talin in mediating PIPKI γ 2 association with β 1-integrin (Fig. 3E). Similarly, direct binding assays using GST- β 1 or α 5-integrin with purified His-tagged PIPKI γ 2 indicated no binding (Fig. 3F). These data demonstrate that PIPKI γ 2 forms a complex with talin and the PIPKI γ 2-talin interaction enhanced the

binding of $\beta 1$ -integrin to talin. Knock down of PIPKI $\gamma 2$ results in loss of $\beta 1$ -integrin targeting to the leading edge (Fig. 2F, G) indicating a defect in trafficking.

PIPKI $\gamma 2$ Knockdown Impairs $\beta 1$ -integrin Exocytosis

To define the role of PIPKI $\gamma 2$ in integrin trafficking, we examined the recycling of $\beta 1$ -integrin in control and PIPKI $\gamma 2$ knockdown cells (Powelka et al., 2004). When $\beta 1$ -integrin was surfaced labelled and then internalized, there was enhanced accumulation of $\beta 1$ -integrin in the perinuclear region of PIPKI $\gamma 2$ knockdown cells (Fig. 4A, B and C). The isolation of the $\beta 1$ -integrin-antibody complex following endocytosis at 37°C for 10 minutes did not show a difference in the endocytosis of $\beta 1$ -integrin in PIPKI $\gamma 2$ knockdown cells (Fig. 4D). This demonstrated that internalization of $\beta 1$ -integrin was not impaired in PIPKI $\gamma 2$ knockdown cells, suggesting that PIPKI $\gamma 2$ regulates exocytosis.

To define if exocytosis was impacted by PIPKI $\gamma 2$ loss, we quantified the trafficking of perinuclear $\beta 1$ -integrin to the plasma membrane upon stimulation of serum starved cells with 10% FBS. PIPKI $\gamma 2$ knockdown cells resulted in diminished plasma membrane trafficking of $\gamma 1$ -integrins (Fig. 4E, F and G) indicating a role for PIPKI $\gamma 2$ in integrin exocytosis. These data were also confirmed biochemically by demonstrating more internal $\beta 1$ -integrin remaining in PIPKI $\gamma 2$ knockdown cells after FBS stimulation (Fig. 4H). In addition, we measured the $\beta 1$ -integrin recycling using a cell surface biotinylation approach. Quantification of $\beta 1$ -integrin recycling indicated that the exocytosis of $\beta 1$ -integrin was diminished in PIPKI $\gamma 2$ knockdown cells but was rescued by re-expression of PIPKI $\gamma 2$ (Fig. 4I, J). Yet, there was no detectable change in the total surface content of $\beta 1$ or $\alpha 5$ integrin in either confluent or migrating cells upon knockdown of PIPKI $\gamma 2$ (Fig. S3C), supporting a role for PIPKI $\gamma 2$ in polarized trafficking of integrin. We focused on $\beta 1$ -integrin trafficking as it represents the predominant integrin in epithelial cells and interacts with the most abundant ECM proteins, FN and collagen (Caswell and Norman, 2006; Caswell et al., 2007). The loss of $\beta 1$ -integrin impaired microtubule orientation, nascent focal adhesion complex formation at migrating cell fronts and haptotactic cell migration towards FN (Fig. S2F, G).

PIPKI $\gamma 2$ Directly Associates with the Exocyst Complex

The data indicates a role for PIPKI $\gamma 2$ in the polarized trafficking of integrins and the involvement of PIP $_2$ -regulated proteins in $\beta 1$ -integrin trafficking during cell migration. The exocyst is a conserved octomeric protein complex involved in polarized vesicle trafficking and is required for directional cell migration (Hertzog and Chavrier, 2011; Zuo et al., 2006). Components of the exocyst complex also serve as effectors of Rab11 and Arf6 GTPases, which regulate integrin trafficking and cell migration (Caswell and Norman, 2006). In addition, the docking of the exocyst complex to membrane is regulated by PIP $_2$ through interactions with Exo70 and Sec3 (He et al., 2007; Liu et al., 2007). As PIP kinases often associate with PIP $_2$ effectors (Anderson et al., 1999; Heck et al., 2007), an interaction of PIPKI $\gamma 2$ with the exocyst complex was explored. The exocyst components were co-immunoprecipitated with PIPKI $\gamma 2$ (Fig. 5A). Cell migration induced the association between PIPKI $\gamma 2$, exocyst complex and $\beta 1$ -integrin (Fig. 5B). This migration induced association was also observed between endogenous PIPKI $\gamma 2$, the exocyst complex and $\beta 1$ -integrin (Fig. 5C). PIPKI $\gamma 2$ expression specifically promoted the complex formation between $\beta 1$ -integrin and the exocyst complex in migrating cells whereas expression of PIPKI $\gamma 1$, PIPKI $\gamma 2$ KD (kinase dead mutant) or PIPKI $\gamma 2$ Y649F (mutant deficient in talin binding) poorly enhanced these associations (Fig. 5D). Consistent with this co-immunoprecipitation of the exocyst complex with $\beta 1$ -integrin was also reduced in PIPKI $\gamma 2$ knockdown cells (Fig. S6B). The immunoprecipitation of PIPKI γ further confirmed these associations and showed that PIPKI $\gamma 2$ Y649F lost interactions with both talin and $\beta 1$ -

integrin (Fig. 5E). This indicates a requirement for PIP₂ generation and talin binding ability of PIPKI γ in regulating the complex formation. Yet, PIPKI γ 1 and PIPKI γ 2Y649F were equally efficient in their interaction with exocyst complex in migrating cells (Fig. 5E). PIPKI γ 2KD had a reduced association with exocyst components but not talin (Fig. 5E) supporting a requirement for both talin interaction and PIP₂ production. These data indicate that PIPKI γ 2 and PIP₂ generation specifically provide the platform for association of exocyst complex with β 1-integrin in migrating cells and this requires PIPKI γ 2 interaction with talin.

To investigate direct interactions between PIPKI γ 2 and the exocyst complex, components of the exocyst complex were purified as GST-fusion proteins. GST-pull down assays demonstrated Sec6 and Exo70 as direct binding partners of PIPKI γ 2 (Fig. 5F), although Exo70 interacted more strongly than Sec6. The interaction between Sec6 or Exo70 with PIPKI γ 2 *in vitro* was not specific for PIPKI γ 2 as all splice variants interacted (not shown). Co-expression and co-immunoprecipitation studies in HEK293 cells indicated that all the isoforms of PIPKI γ interact with Sec6 and Exo70 (S4A, B). This indicates that PIPKI γ interacts with Sec6 and Exo70 through regions conserved in all PIPKI γ isoforms (Heck et al., 2007). Expression of Flag-tagged Sec6 or Exo70 in HeLa cells co-immunoprecipitated endogenous PIPKI γ 2 along with other exocyst complex components (Fig. 5G). In cells, Sec6 and Exo70 mediate the PIPKI γ 2-association with the exocyst complex as knockdown of either Sec6 or Exo70 abrogated co-immunoprecipitation of the exocyst complex with PIPKI γ 2 (Fig. 5H).

The Exocyst Complex is Required for PIPKI γ 2-Regulated Cell Migration

PIPKI γ 2 forms a complex with β 1-integrin and exocyst components, but PIPKI γ 2KD is poorly incorporated into this complex indicating that PIP₂ generation is required (Fig. 5D, E). Consistent with this, the expression of PIPKI γ 2, but not PIPKI γ 2KD, promoted the haptotactic cell migration of HeLa cells towards FN (Fig. 6A). Similar results were obtained using a HeLa Tet-off cell line expressing PIPKI γ 2 or PIPKI γ 2KD (not shown). The expression of PIPKI γ isoforms is variable between cell lines and tissues (Schill and Anderson, 2009b). In breast cancers, increased PIPKI γ expression correlates with disease progression (Schrapf et al., 2011; Sun et al., 2010), indicating that changes in PIPKI γ content is an *in vivo* mechanism to modulate cellular function.

To define the specificity of PIPKI γ 2 in promoting cell migration, each PIPKI γ isoform was expressed in HeLa cells. Enhanced cell migration toward FN was promoted exclusively in cells expressing PIPKI γ 2 (Fig. 6B) and only the PIPKI γ 2 isoform associated with talin and β 1-integrin (Fig. S4E, F). PIPKI γ 2 expression specifically promoted complex formation between β 1-integrin and the exocyst complex in migrating cells (Fig. 5D, E). To define the functional link between the exocyst complex and PIPKI γ 2 in cell migration, we used siRNAs to knockdown exocyst components (Sec5 or Sec8) in cells ectopically expressing PIPKI γ 2. As shown in Figure 6C, knockdown of exocyst components reduced PIPKI γ 2-stimulated cell migration towards FN. Furthermore, knockdown of Sec5 or Sec8 blocked PIPKI γ 2-enhanced β 1-integrin recruitment to the migrating cell front (Fig. 6D). The knockdown of any exocyst complex component impaired the PIPKI γ 2-regulated cell migration and polarized β 1-integrin targeting (Fig. 6C, D and not shown). Overexpression of individual exocyst complex components did not rescue the cell migration defect in PIPKI γ 2 knockdown cells (not shown). These data indicate that PIPKI γ 2 and PIP₂ generation regulates the assembly of exocyst complex required for driving polarized recruitment/trafficking of integrin molecules required for directional migration. To explore this, the polarized recruitment of Exo70, Sec6 and Sec8 to the migrating cell fronts was examined. As shown in Fig. 6E and F, PIPKI γ 2 knockdown impaired the polarized recruitment of exocyst complex components to the leading edge membrane.

PIPKI γ 2 and PIP₂ generation regulates cell migration, and the exocyst complex is required for driving polarized recruitment/trafficking of β 1-integrin. The exocyst complex binds to PIP₂ and is regulated by this interaction (He et al., 2007; Liu et al., 2007). The role of PIP₂ binding was explored using the Exo70 mutant, Exo70-1, deficient in PIP₂-binding (He et al., 2007). Expression of Exo70 modestly enhanced directional migration but co-expression with PIPKI γ 2 synergistically increased migration (Fig. 6G). However, expression of Exo70-1 did not enhance migration and co-expression of Exo70-1 with PIPKI γ 2 blocked PIPKI γ 2-induced cell migration (Fig. 6G). This demonstrates that PIP₂ binding to Exo70 regulates PIPKI γ 2-induced directional cell migration.

PIPKI γ 2 directly associates with Exo70, is a link to the exocyst complex, and these interactions are enhanced during cell migration (Fig. 5B, C). The PIP₂ generation by PIPKI γ 2 is required for its association with the exocyst complex in migrating cells as PIPKI γ 2KD weakly associated with the exocyst complex (Fig. 5E). When co-expressed PIPKI γ 2 and Exo70 tightly co-localize in cells, but PIPKI γ 2KD did not co-localize with Exo70, which was diffusely localized (Fig. S6C). PIP₂ binding is required for Exo70 localization to the membrane and the Exo70-1 mutant loses this localization (Liu et al., 2007). Similarly, the Exo70-1 mutant poorly interacted with PIPKI γ 2 compared to Exo70 (Fig. 6I). Yet, Exo70-1 retained the ability to interact with other exocyst components (Fig. S4G). These data indicate that the PIPKI γ 2 interaction with Exo70 and the exocyst complex is regulated by PIPKI γ 2 generation of PIP₂ and PIP₂ binding to Exo70.

Integrated Role of PIPKI γ 2, Exocyst Complex and Talin in Integrin Trafficking

The exocyst regulates polarized membrane trafficking (He and Guo, 2009) and cell migration (Hertzog and Chavrier, 2011; Zuo et al., 2006). Expressed GFP-Exo70 targets to the plasma membrane and co-localized with β 1-integrin at the plasma membrane (Fig. 6H). GFP-Exo70-1 poorly targeted to plasma membrane and did not co-localize with β 1-integrin at plasma membrane (Fig. 6H). This is consistent with previous reports demonstrating Exo-70 binding to PIP₂ is required for the trafficking of membrane proteins (Liu et al., 2007). GFP-Exo70 colocalizes with α 5-integrin (a β 1-integrin partner) both at the plasma membrane and intracellular compartments (Fig. S6A), but, in PIPKI γ 2 knockdown cells, GFP-Exo70 poorly colocalized with α 5-integrin, specifically in the intracellular compartment (Fig. S6A).

The role of PIPKI γ 2 and the exocyst complex to drive polarized trafficking of integrin molecules to focal adhesion complexes was further supported by *in vivo* colocalization. PIPKI γ 2 and components of the exocyst complex (Exo70, Sec6 and Sec8) colocalized at focal adhesion complexes in HeLa cells adhering and spreading to FN (Fig. 7A). In PIPKI γ 2 expressing cells, Exo70 and α 5 integrin colocalized with PIPKI γ 2 at plasma membrane and vesicular intracellular compartments (Fig. 7B, upper panel) whereas Exo70 poorly co-localized with α 5-integrin in cells not ectopically expressing PIPKI γ 2 (Fig. 7B, middle panel). As a control Exo70 and PIPKI γ 2 were highly colocalized but not with GFP (Fig. 7B, lower panel). These data indicate that PIPKI γ 2 expression promotes Exo70 localization/association with α 5-integrin. In consistent to this, the knockdown of PIPKI γ 2 resulted in diminished exocyst association with β 1-integrin (Fig. S6B).

PIPKI γ 2 directly and specifically interacts with talin (Di Paolo et al., 2002; Ling et al., 2002; Ling et al., 2003) (Fig. S4E) and this interaction mediates PIPKI γ 2 association with β 1-integrin in migrating cells (Fig. 3E and Fig. 5D, E). This interaction is also required for growth factor stimulated chemotaxis (Sun et al., 2007). Talin physically links integrins at focal adhesions to the actin cytoskeleton, a process controlled by PIP₂ (Gilmore and Burridge, 1996; Ling et al., 2006; Martel et al., 2001). Talin is required for focal adhesion targeting of PIPKI γ 2 and talin knockdown cells were defective in focal adhesion formations

(Fig. S5A). Also, targeting of the exocyst complex and β 1-integrin to the migrating cell front and polarization of cells was impaired in talin knockdown cells (Fig. S5B, C, D). Exo70 is targeted to talin containing focal adhesions in PIPKI γ 2 expressing cells (Fig. 7C, top panel) compared to non-expressing cells (bottom panel). Consistent with this observation, PIPKI γ 2 expression specifically promoted the association of talin with the exocyst complex and β 1-integrin (Fig. 7D). These linkages are further supported by the migration dependent integration of PIPKI γ 2 into a complex with talin, α 5 and β 1-integrin but not PIPKI γ 2KD indicating that these are PIP₂ regulated processes (Fig. 7E).

Discussion

The precisely controlled trafficking of integrin molecules to and from plasma membrane is a fundamental process of migrating cells (Caswell and Norman, 2008; Caswell et al., 2009; Muller et al., 2009). The mechanisms for trafficking of integrins toward the membrane front in directionally migrating cells are emerging with roles for Rab4, Rab11, ARF6, Rab25, and spatial growth factor receptor signaling (Caswell and Norman, 2008; Caswell and Norman, 2006; Caswell et al., 2009). The exocyst complex plays a role in polarized secretion and also cell migration (He and Guo, 2009; Zuo et al., 2006). Upon initiation of migration, components of the exocyst complex are redistributed from cell-cell contact sites to focal adhesions (Spiczka and Yeaman, 2008). Here, we describe a role of PIPKI γ 2 in the polarized trafficking of integrin molecules in directionally migrating cells via its association with and regulation of the exocyst complex.

Generation of PIP₂ in a spatiotemporal manner controls vesicle trafficking at the plasma membrane (Di Paolo et al., 2004; Schill and Anderson, 2009a). A role for PIP₂ in trafficking to the plasma membrane or between intracellular compartments is also emerging, as PIP₂ is synthesized on intracellular membrane compartments (Vicinanza et al., 2008) and PIP₂ generation modulates E-cadherin sorting to the basolateral membrane from the recycling endosome (Ling et al., 2007). Exocyst complex components also bind PIP₂ and may regulate trafficking (Liu et al., 2007). This suggests that the exocyst complex coordinates with PIP₂ synthesizing enzymes to modulate integrin trafficking during cell migration.

PIPKI γ 2 directly interacts with exocyst components Sec6 and Exo70. The association of PIPKI γ 2 with both Sec6 and Exo70 may be functionally significant as Sec6 is associated with vesicle containing cargo, whereas Exo70 may mediate plasma membrane docking via PIP₂ interactions (Yu and Hughson, 2010). The interaction with both Sec6 and Exo70 is consistent with the localization of PIPKI γ 2 in cytosolic compartments and at the plasma membrane/focal adhesions. This indicates that PIPKI γ 2 regulates the exocyst complex in multiple compartments positioning PIPKI γ 2 to modulate polarized trafficking of molecules required for cell migration.

The exocyst complex, PIPKI γ 2, β 1-integrin and talin are all individually required for cell migration (Ling et al., 2006; Sun et al., 2007; Zuo et al., 2006). We show that these components integrate together to orchestrate directional migration. Based on these results, upon migration PIPKI γ 2 integrates the exocyst complex with β 1-integrin. The ability of PIPKI γ 2 to interact with talin through its unique C-terminal domain enables the targeting of the exocyst/ β 1-integrin complex to the leading edge where integrin delivery/activation is required for nascent focal adhesion complex formation (Fig. 7F).

The interaction between PIPKI γ 2 and talin targets the exocytosis of β 1-integrin to talin-enriched focal adhesion complexes at leading edge plasma membrane (Fig. 7F) suggesting that talin serves as a tethering factor to guide β 1-integrin trafficking (Yu and Hughson, 2010). In this context, PIPKI γ 2 acts both as a signaling scaffold that links the exocyst

complex and $\beta 1$ -integrin vesicle to talin-based adhesive complexes and generates PIP_2 that regulates vesicle docking with the plasma membrane through PIP_2 regulation of Exo70. This would place $\beta 1$ -integrin, talin, $PIPKI\gamma 2$ and PIP_2 generation in spatial proximity, where PIP_2 enhances the interaction of talin with $\beta 1$ -integrin (Martel et al., 2001). Talin mediated the $PIPKI\gamma 2$ interaction with $\beta 1$ -integrin and PIP_2 generation enhanced the interaction of the exocyst complex with $\beta 1$ -integrin. The intrinsic ability of talin to integrate into focal adhesion complexes in concert with $PIPKI\gamma 2$ regulation of exocyst function facilitates the polarized delivery of $\beta 1$ -integrin to the leading edge of migrating cells. This would lead to the formation of adhesive complexes at the leading edge an event critical for cell migration.

Vinculin links adhesive complexes to actin and its incorporation into the talin/integrin complexes required $PIPKI\gamma 2$ kinase activity (PIP_2 synthesis). The interaction of vinculin with talin has been reported to be both PIP_2 dependent and independent (Chandrasekar et al., 2005; Gilmore and Burridge, 1996). Talin interaction with vinculin is also enhanced by talin stretching (del Rio et al., 2009). As PIP_2 regulates the interaction of talin with integrin this interaction would serve as an anchor for talin such that stretching that would expose vinculin binding sites; indicating that PIP_2 regulates multiple talin interactions.

In vivo migration and invasion occur in three-dimensional matrices. This requires cells to form highly polarized membrane projections to migrate or invade through the matrix. Compared to two-dimensional migration, the role of $PIPKI\gamma 2$, the exocyst, integrin trafficking, and talin in 3-D migration are likely to be accentuated as membrane structures are more polarized and the polarized trafficking of molecules to the leading edge is essential. The increased expression of $PIPKI\gamma 2$ enhanced cell migration and formation of the exocyst/integrin complexes. Increased $PIPKI\gamma$ expression also correlates with disease progression in breast cancer patients (Schramp et al., 2011; Sun et al., 2010). This implicates a role for $PIPKI\gamma 2$ in the metastasis of breast cancers a process requiring cell migration and invasion.

Experimental Procedures

Cell Migration and Wound Healing Assays

Cell migration assays were performed using modified Boyden chambers. The underside of polycarbonate membrane (8- μ m pore size, Neuroprobe) were coated overnight in 10 μ g/ml FN or Col.I, air dried and placed in the chamber-filled with DMEM containing 0.1% BSA with coated surface facing down. Cells after overnight serum starvation were suspended in DMEM containing 0.1% BSA and then, introduced into the upper compartment and incubated for 8-12 hours at 37°C in the incubator. Membranes were fixed after removing the non-migrated cells from upper side with cotton swab, stained with crystal violet stain. Migrated cells were counted from at least 10 randomly selected areas at x200 microscopic fields (HPF). Each experiment was reproduced at least in triplicate for each cell type and matrix. For wound healing assay, MDA-MB-231 or HeLa cells grown to confluence on FN-coated culture dishes were wounded using 200 μ l pipette tips and incubated for 12-16 hours before taking several randomly selected fields to measure wound width.

Immunoprecipitation and GST-pulldown Assay

Cells were lysed in lysis buffer (50 mM Tris-HCl [pH 7.4], 150 mM NaCl, 1% NP40, 1 mM EDTA, 10 mM NaF, 5 mM Na_3VO_4 and proteases inhibitors). Clear supernatant were incubated with indicated antibodies for overnight followed by isolation of immunocomplexes using protein G sepharose 4B beads (Amersham). The beads were washed three times with lysis buffer before eluting immunocomplexes with 2 \times sample buffer and then subjected to immunoblotting with indicated antibodies.

GST fusion proteins of exocyst complex components were purified from *E. coli*. GST-fusion proteins (5 µg each) immobilized on glutathione-agarose beads were incubated with His-tagged PIPKI γ 2 (purified from BL21) in binding buffer (20 mM HEPES [pH 7.4], 150 mM NaCl, 1 mM EDTA, 0.5% Triton X-100 and protease inhibitors) at 4°C for 1 hour. After washing the beads 3 times with binding buffer, bound PIPKI γ 2 were analyzed by immunoblotting using anti-His antibody.

For Pull down assay using cell lysates, GST-fusion proteins (5 µg each) of the cytoplasmic domain of β 1- or α 5-integrin immobilized on glutathione-agarose beads were incubated with cell lysates prepared from MDA-MB-231 cells stably expressing PIPKI γ 1, PIPKI γ 2 or PIPKI γ 2Y649F. After incubation at 4°C for 2-3 hours, beads were washed 2-3 times with lysis buffer before eluting the bound proteins for immunoblotting.

Immunofluorescence Microscopy

To examine the polarized recruitment of PIPKI γ 2, β 1-integrin, exocyst complex or talin, the MDA-MB-231 or HeLa cells were cultured into FN-coated coverslips until confluency. The scratch-wound were created using 200 µl pipette tips and detached cells removed. The remaining cells were allowed to migrate towards the denudated area by incubating the cells at 37°C for 4-5 hours before fixing the cells with 3.7% PFA. The cells were permeabilized with 0.1% Triton-X before blocking with 3% BSA followed by overnight incubation with indicated antibodies at 4°C.

For examining the recruitment of PIPKI γ 2 and exocyst complex at focal adhesion sites, cells were seeded onto FN-coated coverslips in serum-free DMEM medium and incubated for 2-3 hours at 37°C in the incubator before processing the cells for immunofluorescence study as described above. Images were acquired using metamorph in fluorescence microscope (Nikon Eclipse TE2000-U).

Integrin Trafficking Assay

Control or PIPKI γ 2 knockdown MDA-MB-231 cells grown on FN-coated coverslips were labeled with anti- β 1 (MAB2000, Millipore) or anti- α 5 integrin (610633, BD Bioscience) at concentration of 10 µg/ml in serum-free DMEM medium/0.1% BSA by incubating the cells at 4°C for one hour. After removing the unbound antibody with cold DMEM medium, internalization of antibody-integrin complex were initiated by incubating the cells at 37°C in DMEM/10% FBS. At different time points, cell surface antibodies were removed by acid wash (0.5% glacial acetic acid and 0.5 M NaCl, pH 3.0) followed by fixation with 3.7% PFA and cell permeabilization with 0.1% Triton-X. Cells were incubated with Alexa555-labeled goat anti-mouse antibody (Molecular Probe) to visualize internalized antibody-integrin complex. Fluorescence intensity of internalized integrin was measured using metamorph. For biochemical assay of endocytosis, cells were lysed after removing the cell surface antibodies by acid wash. The internalized β 1-integrin antibody complex was isolated from clear supernatants using protein G Sepharose beads followed by SDS-PAGE and immunoblotting of β 1-integrin in the isolated complex.

For integrin exocytosis assay, cell surface β 1-integrins were labeled as described above. Integrin-antibody complex were allowed to internalize and accumulate at perinuclear region by incubating the cells at 37°C for 1-2 hours in DMEM/0.1% BSA medium. The β 1-integrin antibody remaining on cell surface were removed by acid wash. Then, internalized integrin β 1-antibody complex were induced to recycle to plasma membrane by stimulating cells with DMEM/10% FBS for 3-6 minutes. Cells were processed for staining as above without cell-permeabilization (except for examining the internalized β 1-integrin before cell stimulation).

For biochemical assay, recycled β 1-integrin antibody on cell surface was removed by acid wash before cell lysis.

For cell surface biotinylation approach to assess integrin recycling, cells were serum starved overnight and incubated with biotin (0.2 mg/ml) (Sulf-NHS-SS-Biotin, Thermo Scientific) to label surface protein at 4°C for 30 minutes. Then, biotinylated cell surface proteins were allowed to undergo endocytosis by incubating the cells at 37°C for 1-2 hours in DMEM medium. After incubation, biotin remaining on cell surface was removed by MeSNa wash. The recycling of internalized integrins to cell surface were stimulated by incubating the cells in DMEM containing 10% FBS at 37°C. Biotin from recycled integrins to cell surface was removed by second wash with MeSNa. Biotinylated cell surface proteins remaining inside the cells were isolated using Streptavidin affinity gel (Sigma) followed by immunoblotting to examine β 1-integrin and transferrin receptor (TRFR). The % of β 1-integrin recycled was calculated as % of the difference in the amount of β 1-integrin in zero time and selected time point divided by total internalized integrin.

Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

Acknowledgments

This work was supported by National Institute of Health (NIH) grants (CA104708, GM057549, and P30-CA-014520) to RAA and American Heart Association Postdoctoral Fellowship (10POST4290052) to N Thapa. We thank Dr. Shu-Chan Hsu at Rutgers University for generously sharing the antibodies to exocyst complex components and Dr. Wei Guo at University of Pennsylvania for providing the plasmids for GFP-Exo70 and GFP-Exo70-1.

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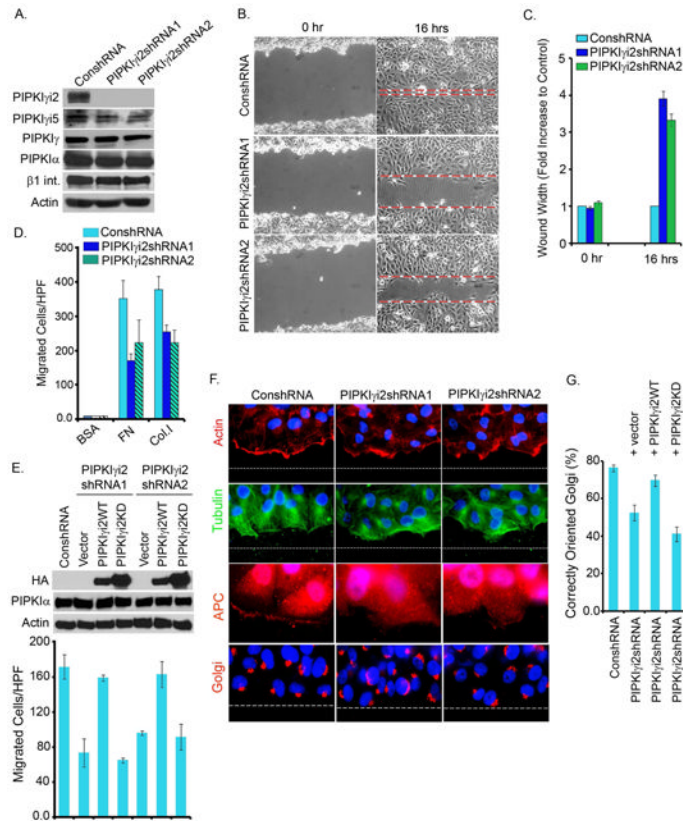


Figure 1. PIPKI γ 2 is Required for Directional Cell Migration

(A) shRNA/lentiviral system was used to knockdown endogenous PIPKI γ 2. Isolated cell lines (designated as PIPKI γ 2shRNA1 and PIPKI γ 2shRNA2) were examined by immunoblotting for knockdown of PIPKI γ 2.

(B) Control or PIPKI γ 2 knockdown cells grown to confluency were wounded and wound width measured at zero and 16 hours post-wounding (representative images at zero and 16 hours post-wounding).

(C) The results are expressed as average fold increase in wound width compared with control cells at zero and 16 hours post-wounding (mean \pm SD from three independent experiments).

(D) For haptotactic cell migration towards FN or Col.I, the modified Boyden chamber was used. Results expressed as the total number of cells migrated/HPF (mean \pm SD from three independent experiments).

(E) Rescue of cell migration defect in PIPKI γ 2 knockdown cells. Lentiviral expression system was used to express PIPKI γ 2 or its kinase dead mutant into the PIPKI γ 2 knockdown cells. Haptotactic cell migration was examined as described above (results are mean \pm SD from three independent experiments)

(F) Confluent cell cultures were wounded and processed 6-hours post-wounding to examine orientation of actin (red), microtubules (green), APC (red) and Golgi (red) towards the direction of cell migration.

(G) Quantitative data for Golgi orientation (mean \pm SD of three independent experiments). See also Figure S1, Movie S1 and S2

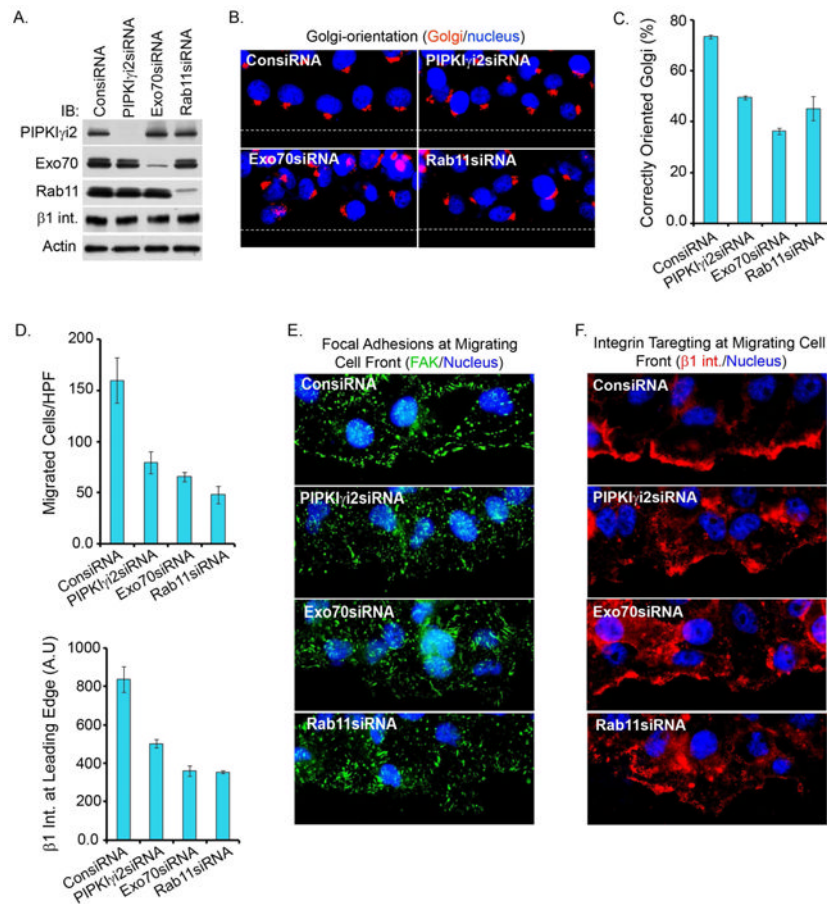


Figure 2. Knockdown of PIPKI γ 2, Exocyst Complex components or Rab11 Impairs Polarized Recruitment of β -integrins and Cell Migration

(A) Knockdown of PIPKI γ 2, Exo70 or Rab11 in MDA-MB-231 cells. Cells were transfected with specific siRNAs and knockdown of indicated proteins was examined by immunoblotting.

(B) Confluent cell culture (48-72 hours post-transfection with siRNA), were wounded and processed 6 hours post-wounding to examine Golgi (red) orientation towards the direction of cell migration.

(C) Quantitative data of Golgi orientation (mean \pm SD of three independent experiments).

(D) The modified Boyden chamber assay as described above was used to examine the migration of siRNA-treated cells towards FN. Results were expressed as the total number of cells migrated/HPF (mean \pm SD from three independent experiments).

(E, F) Confluent cell culture (48-72 hours post-transfection with siRNA) were fixed 5-6 hours post-wounding and immunostained for FAK (green) and β 1-integrin (red) to examine the focal adhesion complex formation and β 1-integrin recruitment at the leading edge.

(G) Quantitative data of β 1 integrin recruitment at the migrating cell fronts. The average fluorescence intensity (AU) of β 1 integrin at migrating cell front was measured using the metamorph (mean \pm SD of three independent experiments). See also Figure S2

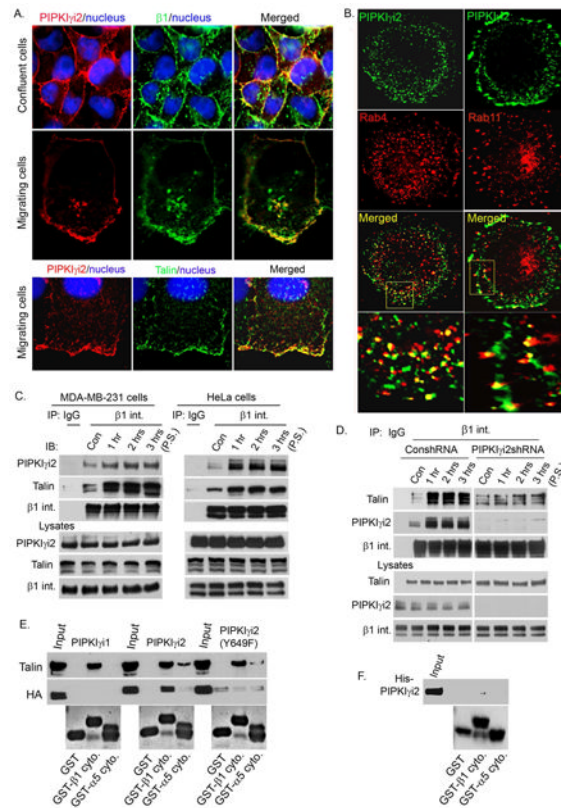


Figure 3. Cell Migration Promotes PIPKI β 2 Reorganization and Association with β 1-integrin Complexes

(A) PIPKI β 2 colocalizes with β 1-integrins and talin. MDA-MB-231 cells expressing moderate level of HA-tagged PIPKI β 2 were wounded and processed for immunofluorescence (4-5 hours post-scratching). PIPKI β 2 (red) and β 1-integrin or talin (green) are recruited to and colocalize at migrating cell fronts and at intracellular compartments.

(B) PIPKI β 2 colocalizes with Rab4 and Rab11. MDA-MB-231 cells expressing PIPKI β 2 were seeded on FN-coated coverslips and cultured for 2-3 hours prior to cell fixation and immunostaining for PIPKI β 2 (green) and Rab4 or Rab11 (red).

(C) Cell migration enhances a PIPKI β 2 association with β 1-integrin and talin. Confluent MDA-MB-231 or HeLa cells were wounded extensively so that about 50% of cells were detached from culture plates. Cells were harvested at different time points and β 1-integrin was immunoprecipitated followed by immunoblotting to examine the co-immunoprecipitation of PIPKI β 2 and talin. (P.S., post-scratch).

(D) PIPKI β 2 is required for talin association with β 1-integrin in migrating Cells. Confluent cultures of control or PIPKI β 2 knockdown cells (HeLa) were wounded as described above before immunoprecipitation of β 1-integrin to examine the co-immunoprecipitation of talin and PIPKI β 2 by immunoblotting.

(E) Cytoplasmic domain of β 1-integrin pulled down both talin and PIPKI β 2. GST-fusion protein of cytoplasmic domain of β 1- or α 5-integrin was incubated with cell lysates prepared from MDA-MB-231 cells expressing PIPKI β 1, PIPKI β 2 or PIPKI β 2Y649F. Pull down of talin and PIPKI β 2 were examined by immunoblotting.

(F) *In vitro* binding study. GST-fusion protein of cytoplasmic domain of β 1- or α 5-integrin was incubated with His-tagged PIPKI β 2 purified from bacteria and PIPKI β 2 binding examined by immunoblotting using anti-His antibody. See also Figure S3

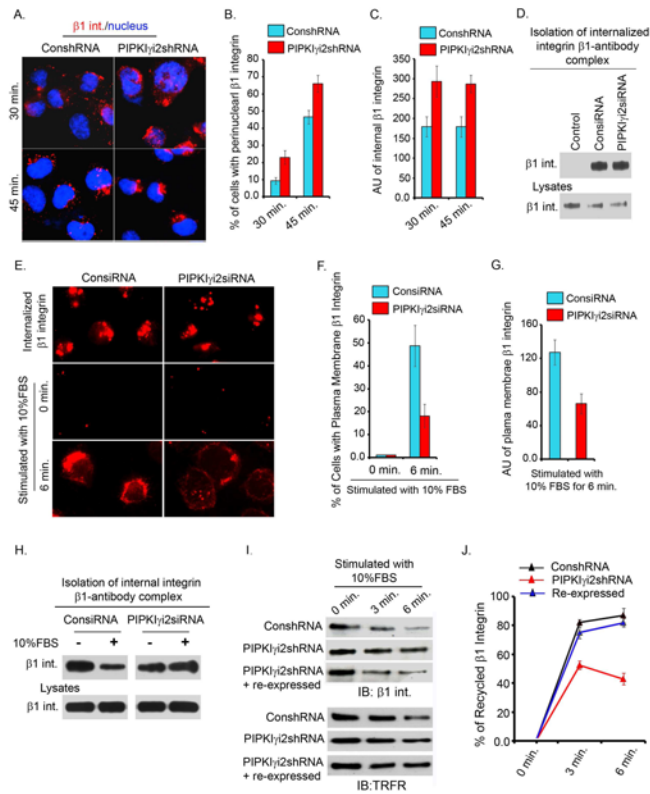


Figure 4. PIPKI γ 2 Knockdown Impairs β 1-integrin Exocytosis

(A) For β 1-integrin endocytosis, cell surface β 1-integrin were labeled with anti- β 1 antibody at 4°C. Cells were incubated at 37°C to induce internalization. Shown is the β 1-integrin internalized after 30 and 45 minutes incubation.

(B) Cells with distinct perinuclear accumulation of β 1-integrin-antibody complex were counted and expressed as % of total cells. A total of 150-200 cells were counted for each condition (results are mean \pm SD of three independent experiments).

(C) Average fluorescence intensity (AU) of internalized β 1-integrin in knockdown and control cells was measured (around 150 cells included for each condition; results are mean \pm SD of three independent experiments).

(D) For biochemical assay of β 1-integrin endocytosis, cell surface β 1-integrin were labeled with anti- β 1 antibody at 4°C followed by incubation of cells at 37°C for 10 minutes to induce internalization. The content of internalized β 1-integrins in control or PIPKI γ 2 knockdown cells was examined by immunoblotting.

(E) For examining β 1-integrin accumulation at perinuclear regions, cells were permeabilized (top panel) before immunostaining as described in “Experimental Procedures”. Cells were processed for immunostaining without cell permeabilization to examine the β 1-integrin (red) trafficking to the plasma membrane before (middle panels) or after (bottom panels) cell stimulation with FBS.

(F) The number of cells with distinct plasma-membrane localization of β 1-integrin in control vs. PIPKI γ 2 knockdown cells was quantified (around 150 cells counted each time; values are mean \pm SD of three independent experiments).

(G) Average fluorescence intensity (AU) of plasma-membrane localization of β 1-integrin in control vs. PIPKI γ 2 knockdown cells (around 150 cells counted each time; values are mean \pm SD of three independent experiments).

(H) The content of internal β 1-integrin after FBS stimulation. Representative image of three independent experiments showing that PIPKI γ 2 knockdown slowed β 1-integrin trafficking to the plasma membrane.

(I) Integrin recycling was examined by cell surface biotinylation assay as described in “Experimental Procedures”. Biotinylated cell surface proteins remaining inside the cells were isolated using streptavidine affinity gel followed by examination of β 1-integrin and transferrin receptor (TRFR) by immunoblotting.

(J) Quantitative data of β 1-integrin recycling. The % of β 1-integrin recycled was calculated as described in “Experimental Procedures” (values are mean \pm SD from three independent experiments).

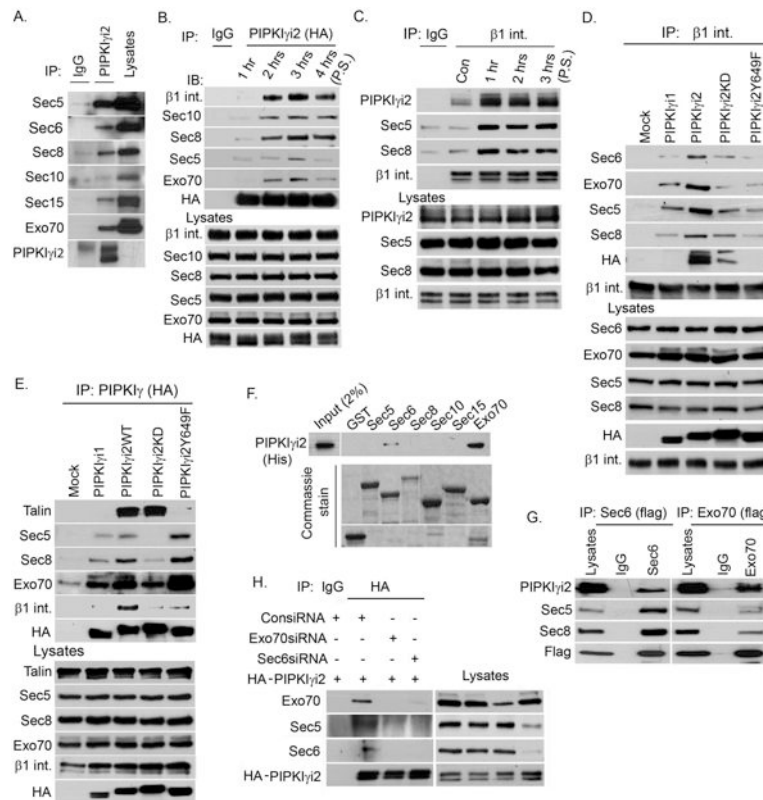


Figure 5. PIPKI γ 2 Directly Associates with the Exocyst Complex

(A) Endogenous PIPKI γ 2 was immunoprecipitated from MDA-MB-231 cells and co-immunoprecipitation of exocyst complex examined by immunoblotting.

(B) MDA-MB-231 cells expressing moderate level of HA-tagged PIPKI γ 2 grown to confluence were harvested at different time points after scratch-wounding. PIPKI γ 2 was immunoprecipitated using anti-HA antibody and co-immunoprecipitation of β 1-integrin and exocyst complex examined by immunoblotting.

(C) HeLa cells grown to confluence were harvested at different time points after scratch-wounding. β 1-integrin was immunoprecipitated and co-immunoprecipitation of PIPKI γ 2 and exocyst components examined by immunoblotting.

(D) MDA-MB-231 cells expressing PIPKI γ 1, PIPKI γ 2, PIPKI γ 2KD or PIPKI γ 2Y649F were harvested 2-3 hours post-wounding. β 1-integrin was immunoprecipitated and co-immunoprecipitation of exocyst complex proteins examined by immunoblotting.

(E) MDA-MB-231 cells expressing PIPKI γ 1, PIPKI γ 2, PIPKI γ 2KD or PIPKI γ 2Y649F were harvested 2-3 hours post-wounding. PIPKI γ 2 and other mutants were immunoprecipitated using anti-HA antibody and co-immunoprecipitation of exocyst complex examined by immunoblotting.

(F) Sec6 and Exo70 directly interact with PIPKI γ 2. GST-fusion protein of exocyst complex components were incubated with His-PIPKI γ 2 purified from bacteria. PIPKI γ 2 binding was examined by immunoblotting using anti-His antibody.

(G) Exo70 and Sec6 co-immunoprecipitate endogenous PIPKI γ 2. HeLa cells were transiently transfected with Flag-tagged Exo70 or Sec6 and immunoprecipitated using anti-Flag antibody. Co-immunoprecipitation of PIPKI γ 2 and other components of exocyst complex were examined by immunoblotting.

(H) Knockdown of Exo70 or Sec6 impairs PIPKI γ 2 association with exocyst complex. HeLa cells were transfected with siRNA for Exo70 or Sec6. 24 hours after the siRNA transfection, cells were transfected with HA-tagged PIPKI γ 2. Next day, cells were

harvested to immunoprecipitate PIPKIγ2 and co-immunoprecipitation of exocyst complex examined by immunoblotting. See also Figure S4

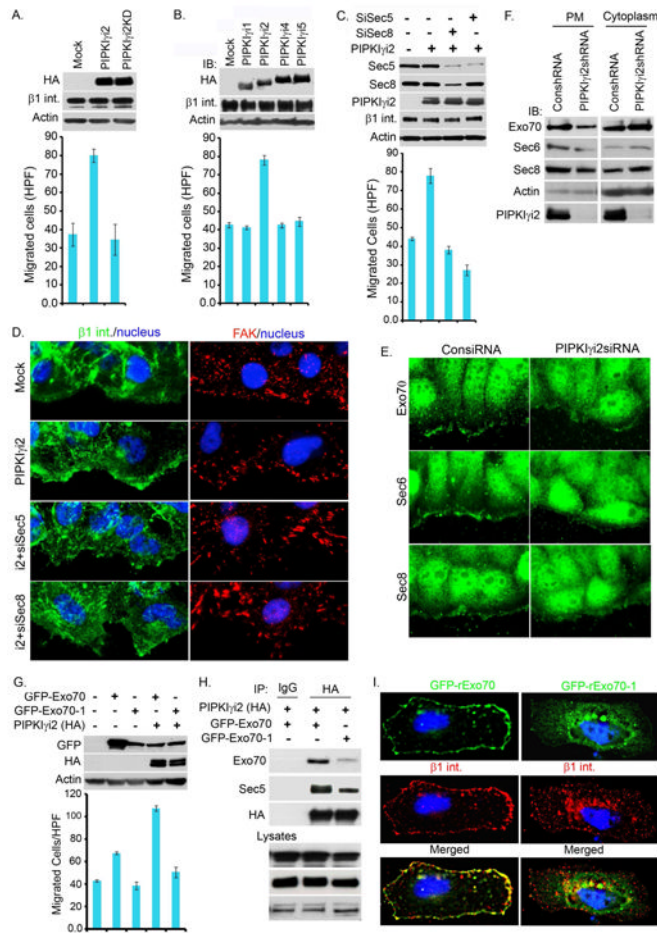


Figure 6. The Exocyst Complex is Required for PIPKI γ 2-regulated Cell Migration

(A) PIPKI γ 2 expression promotes cell migration. HeLa cells transiently transfected with PIPKI γ 2 or PIPKI γ 2KD were monitored for changes in haptotactic cell migration using a modified Boyden chamber assay. The results are expressed as migrated cells/HPF (mean \pm SD of three experiments). Immunoblots were used to examine PIPKI γ expression using anti-HA antibody.

(B) Cell migration assays were performed as above in HeLa cells transiently transfected with PIPKI γ isoforms and results expressed as migrated cells/HPF (mean \pm SD of three experiments). Immunoblots were used to examine PIPKI γ expression using anti-HA antibody.

(C) Cell migration assays were performed in HeLa cells treated with siRNA to knockdown exocyst components (Sec5 or Sec8) followed by PIPKI γ 2 overexpression as described above. Results expressed as migrated cells/HPF (mean \pm SD of three experiments). Knockdown of Sec5 or Sec8 and expression of PIPKI γ 2 were monitored by immunoblotting.

(D) Exocyst complex is required for polarized recruitment of β 1-integrin. HeLa cells stably expressing PIPKI γ 2 were treated with siRNA to knockdown Sec5 or Sec8. 48-72 hours post-transfection, cells were scratch-wounded and immunostained for β 1-integrin (green) and FAK (red) to examine the recruitment of β 1-integrin and focal adhesion formation at migrating cell fronts.

(E) Confluent culture of cells (48-72 hours post-transfection with siRNA) were processed 2-3 hours post-wounding. The polarized recruitment of endogenous exocyst complex

(Exo70, Sec6 and Sec8) (green) to migrating cell fronts were examined using their specific antibodies.

(F) Crude plasma membrane was isolated from control or PIPKI γ 2 knockdown cells followed by examination of exocyst complex components in plasma membrane and cytosol.

(G) HeLa cells were transfected with GFP-Exo70 or GFP-Exo70-1 or cotransfected with PIPKI γ 2. Haptotactic cell migration towards FN was examined as described above. The results expressed as migrated cells/HPF (mean \pm SD of three experiments).

(H) HeLa cells were transfected with either GFP-Exo70 or GFP-Exo70-1. GFP-Exo70 colocalized with β 1-integrin (red) at plasma membrane whereas GFP-Exo70-1 was found either diffusely distributed into the cytoplasm or accumulated around perinuclear regions. β 1-integrin was poorly recruited to plasma membrane and accumulated around perinuclear regions in GFP-Exo70-1 expressing cells.

(I) Exo70-1 poorly associates with PIPKI γ 2 and impairs the PIPKI γ 2 association with the exocyst complex. HeLa cells were cotransfected with PIPKI γ 2 and Exo70 or Exo70-1. Cells were harvested 24 hours post-transfection to immunoprecipitate PIPKI γ 2 and co-immunoprecipitation of exocyst complex examined by immunoblotting. See also Figure S5

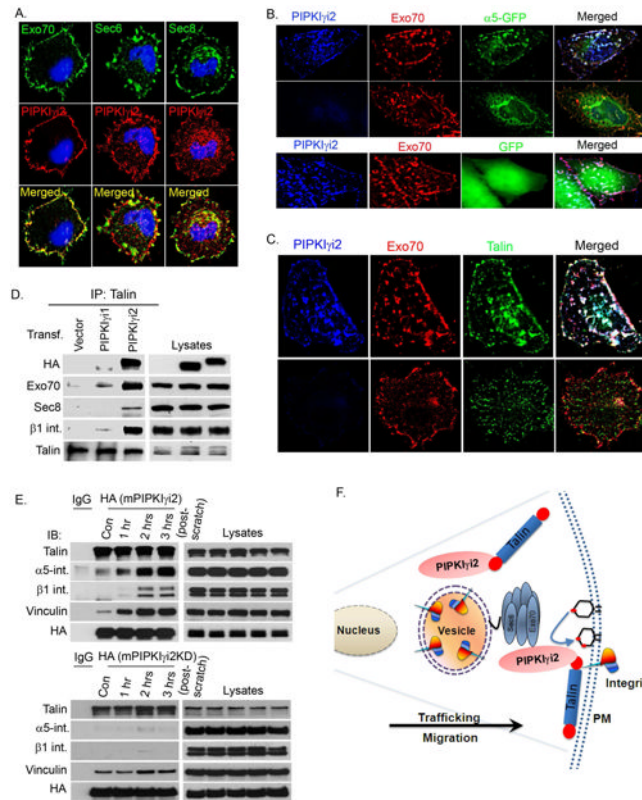


Figure 7. PIPKIγ2 Integrates Exocyst Complex and Talin with Integrin, and is Required for Integrin Trafficking

(A) HeLa cells cotransfected with Flag-tagged exocyst complex components (Sec6 or Sec8 or Exo70) and HA-tagged PIPKIγ2 were allowed to adhere on FN-coated coverslips for 1-2 hours before immunostaining with anti-Flag (green) and anti-HA (red) antibodies.

(B) PIPKIγ2, Exo70 and α5-GFP integrin colocalize at cell membrane and intracellular compartments. HeLa cells (bottom) or HeLa cells stably expressing HA-tagged PIPKIγ2 (top) were co-transfected with Flag-tagged Exo70 and α5 integrin-GFP or GFP. Cells were fixed and immunostained using anti-HA (blue) or anti-Flag (red) antibodies.

(C) Exo70 colocalizes with talin at focal adhesions and intracellular sites. HeLa cells transiently transfected with Flag-tagged Exo70 alone or cotransfected with HA-tagged PIPKIγ2 were allowed to adhere on FN-coated coverslips for 1-2 hours before immunostaining with anti-HA (blue), anti-Flag (red) and anti-talin (green) antibodies.

(D) PIPKIγ2 integrates talin, β1-integrin and exocyst complex in the same complex. HeLa cells were transfected with PIPKIγ1 or PIPKIγ2. Talin were immunoprecipitated 24 hours post-transfection followed by immunoblotting for exocyst complex and β1-integrin.

(E) PIPKIγ2 were immunoprecipitated from HeLa cells stably expressing PIPKIγ2 or PIPKIγ2KD at different time points following wounding to induce migration. Immunocomplexes were examined for presence of integrins and/or talin by immunoblotting.

(F) Model depicting the role of PIPKIγ2 in integrin trafficking in directionally migrating cells. Cell migration induces the integration of PIPKIγ2, talin, β1-integrin into the complex either in plasma membrane or in intracellular recycling compartments. Further, PIP₂ generation by PIPKIγ2 into the complex facilitates the assembly of the exocyst complex. Thus, coordinated activity of PIPKIγ2 and the exocyst complex in concert with talin promotes the polarized recruitment and trafficking of integrin molecules to migrating cell fronts. Loss of PIPKIγ2 or the exocyst complex or talin compromises the polarized

recruitment/trafficking of integrin impairing cell polarization and directional cell migration.
See also Figure S6