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The hematopoietic stem cell polarization and migration

A dynamic link between RhoA signaling pathway, microtubule network and ganglioside-based membrane microdomains

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Abbreviations: ERM, ezrin/radixin/moesin; HSPCs, hematopoietic stem and progenitor cells; MSCs, multipotent mesenchymal stromal cells; PSGL-1, P-selectin glycoprotein ligand-1; ROCK, Rho-associated coiled-coil protein kinase; RNAi, RNA interference; SDF-1α, stromal cell-derived factor-1α

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The polarization and migration of L eukaryotic cells are fundamental processes for the development and maintenance of a tissue. These aspects gain especial interest when it comes to stem and progenitor cells in the way that their manipulation might open new avenues in regenerative therapy. In recent years, novel biological facets of migrating hematopoietic stem cells were revealed by several groups, including ours. Among these features, the polarization of their membranous (proteins and lipids) and cytoplasmic constituents, which leads to the formation of a specialized sub-cellular structure located at the rear pole—the uropod—has gained increasing interest. In a new study we have demonstrated that such phenomena involve a coordinated mechanism between Rho GTPase signaling and the microtubule network. Specifically, our results based on the use of synthetic inhibitors and RNA interference suggest that the activity of RhoA and its effector ROCK I is indispensable for cell polarization and the active reorganization of microtubules that are required for migration.

Understanding the cellular and molecular trafficking mechanisms that regulate the migration of hematopoietic stem and progenitor cells (HSPCs) throughout the development of an organism and later on its homeostasis are important not only from a biological standpoint, but also with regard to therapeutic purposes. For instance, bone marrow transplantation is one recognized procedure for treating

hematological diseases. However, the accurate mechanism underlying the migration and engraftment of HSPCs into the bonemarrow niche is not fully characterized. In order to gain novel insights we have developed an ex vivo co-culture system consisting of human HSPCs from healthy donors growing on primary human multipotent mesenchymal stromal cells (MSCs) as feeder cell layer (for cell isolation and culture conditions see literatures). 1-3 Such cellular system reproduces numerous characteristics found within bone marrow cavities4 including adhesive interactions⁵ and the essential chemotactic axis⁶ based on the G-protein-coupled receptor CXCR4, which is expressed by HSPCs and its chemokine ligand CXCL12 (alias stromal cell-derived factor- 1α ; SDF- 1α) secreted by MSCs.1,7

Under these conditions, HSPCs display various morphologically identifiable types of plasma membrane protrusions.1 Interestingly, in migrating HSPCs a noteworthy protrusion called uropod is formed at the rear pole (Fig. 1A).1,8 Like in leukocytes (e.g., T cells), the uropod might play a role in intercellular adhesion, communication and motility.^{9,10} Numerous proteins with adhesive properties are found therein including P-selectin glycoprotein ligand-1 (PSGL-1; Fig. 1A).11,12 The presence of the stem cell marker CD133 (prominin-1),¹³⁻¹⁵ a 5-span transmembrane glycoprotein that binds plasma membrane cholesterol and associates with a specific membrane microdomain (lipid raft),16 was instructive with regard to its membrane organization. Indeed, a membrane microdomain

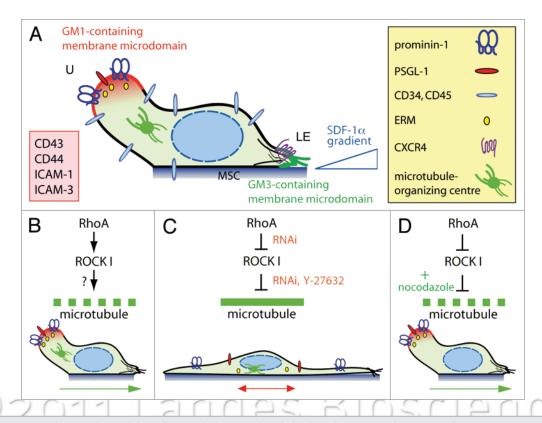


Figure 1. RhoA/ROCK I pathway and remodeling of microtubule network underlie the polarization and migration of HSPCs. (A) A migrating HSPC growing on MSC displays a polarized morphology with the formation of a uropod (U) at the rear pole and a leading edge (LE) at the front. Both types of plasma membrane protrusions contain a specific ganglioside-based membrane microdomain—the uropod being enriched in GM_1 (red) whereas the leading edge in GM_3 (green). In addition to prominin-1, a plethora of cell adhesion molecules (inset) including PSGL-1 are concentrated in the uropod whereas the chemokine receptor CXCR4 is found at the leading edge consistent with its sensory role towards an SDF-1α gradient.⁶ ERM proteins seem to be actively involved in the subcellular localization of some uropod-associated membrane proteins.¹⁸ Other molecules such as CD34 and CD45 are evenly distributed. The microtubule-organizing centre is found between the nucleus and the uropod. (B) The activity ($_1$) of RhoA and its downstream effector ROCK I contributes to the formation of the uropod, and hence polarization and migration of HSPCs. The downstream target(s) remain to be identified (?), but it might engage a protein involved in microtubule destabilization (dashed green line). (C) Inhibition ($^\perp$) of ROCKs using Y-27632 or the specific knockdown of ROCK I or its upstream regulator RhoA by means of RNAi results in an elongated morphology where the uropod is lost. Both membrane (prominin-1, PSGL-1) and cytoplasmic (ezrin) proteins are redistributed. These cells display an impairment of migration caused by microtubule stability (solid green line). (D) In RhoA/ROCKI-deficient HSPCs, the addition (+) of nocodazole restores their proper polarization and migration highlighting the implication of unidentified microtubule-destabilizing proteins. Green and red arrows indicate the direction of migration.

enriched in ganglioside GM, is concentrated in the uropod, 10 and its spatiotemporal regulation might engage molecules such as flotillins.17 From the cytoplasmic side, certain proteins of the ezrin/ radixin/moesin (ERM) family might link membrane proteins (e.g., PSGL-1) via their juxta-membrane domains to the underlying actin cytoskeleton.¹⁸ The microtubule-organizing centre is found at the base of the uropod.12 At the front pole, a migrating HSPC exhibits a lamellipodium, which concentrates CXCR4 at its tips in agreement with a chemotactic role.11 As reported for T cells, a distinct membrane microdomain based on ganglioside GM₂ instead of GM₁ is found therein (Fig. 1A).11,19 From both morphological and phenotypical angles, the migrating HSPC develops thus a highly polarized structure that underlies coordinated but opposite actions at both cell sides. While retracting the uropod, the cell extends its lamellipodium at the leading edge. As a result, a net cell movement can be achieved by continuous attachment to and de-adhesion from the substratum at the front and rear pole, respectively (Fig. 1B and green arrow).

From a mechanistical and/or biochemical perspective, our recent study has focused on the implication of Rho GTPase signaling pathway in these orchestrated processes (Fig. 1B). ¹² It is known that Rho GTPases such RhoA, Rac and Cdc42 are key players in cell polarity and migration by modulating cytoskeletal dynamics. ²⁰ As the most important downstream

effectors of RhoA, Rho-associated coiledcoil protein kinases (ROCK) are implicated in various cellular functions such as actin organization and transformation. Using Rho kinase inhibitor Y-27632 and RNA interference (RNAi) directed against either RhoA or ROCK I we demonstrated that both proteins are indispensable for the polarization of HSPCs, and hence their migration. For instance, the use of the synthetic drug resulted in the complete loss of the uropod and the formation of two to three long and thin plasma membrane protrusions (Fig. 1C). Narrowed lamellipodia were formed at the tip of those protrusions rather than close to the cell body, as in untreated cells (Fig. 1C).12 Such a drastic morphological alteration was followed at the molecular level by a redistribution of plasma membrane (prominin-1 and PSGL-1) and cytoskeleton (ezrin, an ERM protein) constituents of the uropod. The asymmetric distribution of microtubule-organizing centre was also lost (Fig. 1C). 12 As a functional consequence, Y-27632-treated cells displayed a net impairment of migration as evaluated by time-lapse video microscopy and Transwell-filter assay.¹² Specifically, Y-27632-treated cells showed a defect in retracting the long plasma membrane protrusion located at the rear pole and frequently changed their directional movement by 180°, suggesting a perturbation in the front-rear orientation mediated by CXCR4/SDF-1α axis (Fig. 1C and red double-headed arrow). All the characteristics described above for Y-27632-treated cells were reproduced upon the use of RNAi-mediated knockdown of RhoA and ROCK I, thus confirming the direct implication of Rho GTPase signaling pathway in the polarization and migration of HSPCs (Fig. 1C). These outcomes appeared highly specific since the knockdown of ROCK II did not provide such radical effects.

Surprisingly, our study also revealed that the defect in cell polarization including the formation of the uropod could be fully rescued by the nocodazole-mediated depolymerization of the microtubule network (Fig. 1D). Not only was the polarized morphology of RhoA/ROCK I-deficient HSPCs restored, but also their locomotion. The actin depolymerization triggered by latrunculin B did not produce such reversible effects.¹² The precise way that RhoA/ROCK I signaling contributes to the microtubule instability at the uropod cortex is currently unknown. Nevertheless our experiments with nocodazole/RNAi seem to exclude a feedback loop involving microtubule-associated guanine nucleotide exchange factor (GEF)-H1, which activates RhoA21 upon its release from microtubules after the disruption of the latter. However such phenomenon might occur in a natural context. The potential direct or indirect targets of ROCK remain to be identified, but they might engage enzymes that mediate tubulin detyrosination and acetylation, e.g., histone deacetylase 6, the activity of which is modulated by RhoA/ROCK.^{22,23} Active crosstalk between players at the leading edge and the uropod as well as a dynamic balance of the actomyosin and microtubule systems also need to be considered.24 The lack of the front-rear orientation of RhoA/ROCK I-deficient HSPCs and altered lamellipodia are consistent with it.12 Microtubuledestabilizing protein stathmin/OP18 might participate in these biochemical reactions via Rac/Cdc42,25 and the regulation of Rac by ROCK via the filamin A-binding RhoGTPase-activating protein reveals the complexity of the system.²⁶ Similarly, members of the ERM protein family such as ezrin or moesin might be involved as well in the integrity of the uropod, and it might be more than a coincidence that these adaptor proteins, which are also potential substrates of ROCK, are playing an active role in membrane microdomain dynamics.²⁷

Lastly, it is noteworthy that comparable data showing the implication of RhoA and microtubule network in the migration of T cells were recently reported independently^{28,29} indicating that our current observations might extend to cells of hematopoietic origin in general. Further studies based on a quantitative proteomic approach should lead to an exhaustive list of ROCK I substrates involved in these processes, which might represent potential therapeutic targets in the development of new strategies to improve the efficiency of bone marrow transplantation.

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