A novel caveolin-1/p85 α /Rab5/Tiam1/Rac1 signaling axis in tumor cell migration and invasion

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Keywords: Caveolin-1, cell migration, Rab5, invasion, p85α, Rac1

Abbreviations: CAV1, Caveolin-1; GAP, GTPase activating protein; GEF, guanine exchange factor; PI3K, phospahtidylinositol-3-kinase.

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Submitted: 07/07/2014

Accepted: 07/14/2014

http://dx.doi.org/10.4161/19420889.2014.972850

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The small GTPase Rab5 has been fre-I quently studied in the context of intracellular trafficking, but evidence obtained more recently has implicated Rab5 as a critical regulator of cell adhesion, migration and invasion in both normal and tumor cells. These recent findings showing that Rab5 promotes Rac1 activation and focal adhesion dynamics have highlighted the question as to what the upstream regulators of Rab5 activity might be and how these are connected to cell migration. The efforts to shed light on this issue identified in metastatic cancer cells a novel Caveolin-1/p85\alpha/Rab5/Tiam1/Rac1 signaling axis relevant to cancer cell migration and invasion. In this addendum, we highlight aspects concerning Rab5 regulation in this context.

Rab5 in Cell Migration

Rab proteins are small GTPases that control intracellular trafficking, by acting as molecular switches that cycle between an inactive GDP-bound and an active GTP-bound state. More than 60 members of this family have been identified and classified according to sequence, function and subcellular localization.^{1,2} Among these proteins, Rab5 is of particular interest, because it coordinates early endosome fusion, dynamics and transport, as well as vesicle formation.1 Moreover, recent observations extend the functional repertoire of Rab5 beyond its cannonical role in membrane trafficking (reviewed in³). Specifically, in the context of cell migration, Rab5 was shown to promote

Rac1-GTP loading, remodeling of the actin cytoskeleton and formation of lamellipodia.4-7 In addition, Rab5 associates in a complex with integrin β 1, increasing integrin recycling and sustaining cell migration.^{7,8} Consistent with these observations, our more recent findings indicate that Rab5 is required for focal adhesion disassembly, tumor cell migration and invasion,9 and, therefore, that Rab5 is a critical player in cell migration.^{10,11} Thus, to advance further in understanding the non-canonical functions of Rab5, the identification of putative Rab5 regulators became essential. As a small GTPase, Rab5 function (i.e. GTP-loading) is tightly controlled by guanine exchange factors (GEFs) and GTPase activating proteins (GAPs). In this context, the regulatory subunit of Class I phospahtidylinositol-3-kinase (PI3K), p85α, became a particularly interesting candidate, because it is implicated not only in cell migration via the regulation of PI3K, but also as a Rab5-GAP.^{12,13}

The Caveolin-1/p85α/Rab5/ Tiam1/Rac1 Axis

Rab5 activation is followed by the recruitment of Tiam1, a Rac1-GEF, to early endosomes, leading to Rac1 activation, localized actin remodeling and cell migration.⁵ However, the identity of regulators and upstream elements implicated in this signaling axis remained poorly understood. In this respect, we recently identified Caveolin-1 (CAV1), a membrane-bound protein, as a critical regulator of Rab5 activity in metastatic cancer

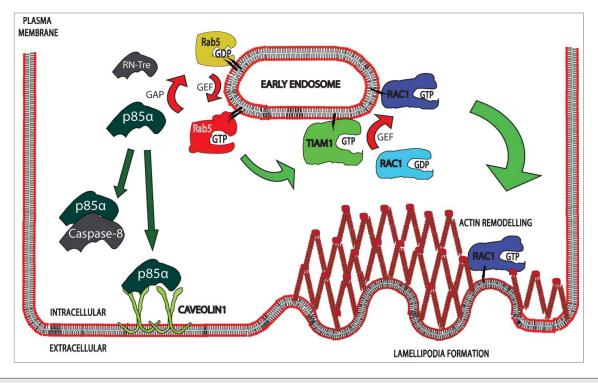


Figure 1. The figure depicts key signaling proteins implicated in CAV1 and Rab5 dependent cell migration. CAV1 promotes Rab5-GTP loading by sequestering $p85\alpha$, which is a Rab5-GAP, leading to Rab5-dependent recruitment of Tiam1, a Rac1-GEF, to early endosomes. This cascade of events is followed by the activation of Rac1, leading to local remodeling of the actin cytoskeleton, cell migration and invasion. This mechanism is reminiscent of that previously described for Caspase-8, which is included in the scheme to the left.

cells.14 CAV1 was previously known to stimulate Rac1 activity and focal adhesion turnover in metastatic melanoma and breast cancer cells,15,16 but the precise connection between CAV1 and Rac1 remained unknown. Our recent work¹⁴ showed that CAV1 promotes Rab5 activation, leading to Rac1 activation, migration and invasion in 3 different metastatic lines, B16-F10 mouse melanoma, HT-29 (US) human colon adenocarcinoma and MDA-MB-231 breast cancer cells. Mechanistically, CAV1 increased the amount of Tiam1 at Rab5-positive early endosomes and, accordingly, Tiam1 was shown to be necessary to link the CAV1-Rab5 axis to enhanced Rac1 activation, cell migration and invasion.

The search for molecules implicated in CAV1-dependent activation of Rab5 led to the identification of $p85\alpha$, which was shown to associate in a complex with CAV1. Sequestration of $p85\alpha$ by CAV1 precluded Rab5-GTP hydrolysis, leading to increased Rab5 activation, as well as migration and invasion in metastatic

cancer cells.¹⁴ Intriguingly, this sequestration mechanism is reminiscent of that previously described for Caspase-8-dependent activation of Rab5 in neuroblastoma cells.¹⁷ In that case, Caspase-8 phosphorylation on Y380 created an SH2-binding site that recruited p85 α , preventing Rab5 inactivation.¹⁷⁻¹⁹ Whether this mechanism is valid for other proteins that connect to Rab5 and favoring migration as well as invasion, remains to be elucidated.

Concluding Remarks

Recent studies support the notion that Rab5 is a key regulator not only of intracellular trafficking, but also of additional non-canonical functions, such as cell migration and invasion. Because enhanced migratory and invasive capacity represent hallmarks characteristic of malignant tumor cells, the identification of mechanisms leading to deregulation of Rab5 function is currently of great interest. The

recently described Caveolin-1/p85α/ Rab5/Tiam1/Rac1 signaling axis sheds light on how this may occur, especially since the pathway was found to be constitutively activated (Fig. 1) in the metastatic cancer cell lines analyzed. However, more experiments are required to demonstrate the functional importance in animal models in vivo and also in clinically relevant settings. Indeed, the future for this novel signaling axis looks bright given that it brings together a number of proteins that are commonly deregulated in cancer and other human pathologies.

Disclosure of Potential Conflicts of Interest

No potential conflicts of interest were disclosed

Funding

This work was supported by Fondecyt 1140907 and 11100287 (VT); CONI-CYT-FONDAP 15130011, Anillo ACT 1111, Fondecyt 1130250 (AFGQ); Conicyt PhD Student Fellowship (JD, PS).

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