

## Rho GEFs in endothelial junctions

### Effector selectivity and signaling integration determine junctional response

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**R**ho GTPases are cytoskeleton-regulating proteins that mediate the formation of intercellular junctions. Their localized activation by Rho GEFs (guanine-nucleotide exchange factors) and the selective activation of downstream effectors have emerged as areas of active research in the cell adhesion field. We reported recently that the Rho-specific GEFs Syx (Synectin-binding RhoA exchange factor) and TEM4 (Tumor Endothelial Marker 4) are both essential for endothelial junction maturation and barrier function. Syx is recruited to cell contacts via its C-terminal PDZ binding motif and its interaction with Muppl and the Crumbs polarity complex, while the junctional localization of TEM4 requires its N-terminal domain and interaction with the cadherin-catenin complex. Our findings support multiple roles for RhoA in junction formation and maintenance. They also suggest that selective coupling of RhoA activation to Dial1 and/or ROCK signaling is critical for determining endothelial junction integrity.

Cell junctions act as gatekeepers to maintain the adhesion and barrier functions of epithelial and endothelial monolayers. The proper regulation of cell contacts is important for physiological events such as vascular development, tissue regeneration, and organ morphogenesis. Under circumstances where cell-cell adhesion is impaired, pathological conditions such as infection, inflammation, edema, ischemia, and cancer can arise, leading to severe and potentially life-threatening consequences. The adherens junctions (AJs) and tight junctions (TJs) are two

specialized adhesive structures that regulate the barrier function of epithelial and endothelial monolayers. The main adhesion-related proteins at AJs include cadherins, nectins, and catenins, whereas TJs are defined and regulated by the claudins, occludins, junctional adhesion molecules (JAMs), and zonula occludens (ZO) proteins.<sup>1</sup> Although these AJ and TJ-associated proteins provide some level of adhesiveness to maintain nascent cell-cell contacts, stable cell-cell cohesion requires adhesion receptor clustering and re-organization of the underlying actin cytoskeleton, processes regulated by Rho GTPases.

The Rho GTPases form a subfamily of the Ras superfamily and exist in an active, GTP-bound state and an inactive, GDP-bound state. It is well documented that Rho GTPases regulate cell adhesion by manipulating the dynamics and reorganization of the actin cytoskeleton.<sup>2</sup> Rho proteins are themselves controlled by three classes of molecules: the guanine-nucleotide exchange factors (GEFs), which activate Rho proteins at specific sub-cellular locations; the GTPase activating proteins (GAPs), which inactivate Rho GTPases; and the guanine nucleotide-dissociation inhibitors (GDIs), which interact with and sequester inactive Rho proteins in the cytosol. As the molecular “on” switches, the GEFs have become interests of study in the field of cell-cell adhesion.

#### The Rho GEF Syx in intercellular junctions

We recently published that the Rho-specific synectin-binding exchange

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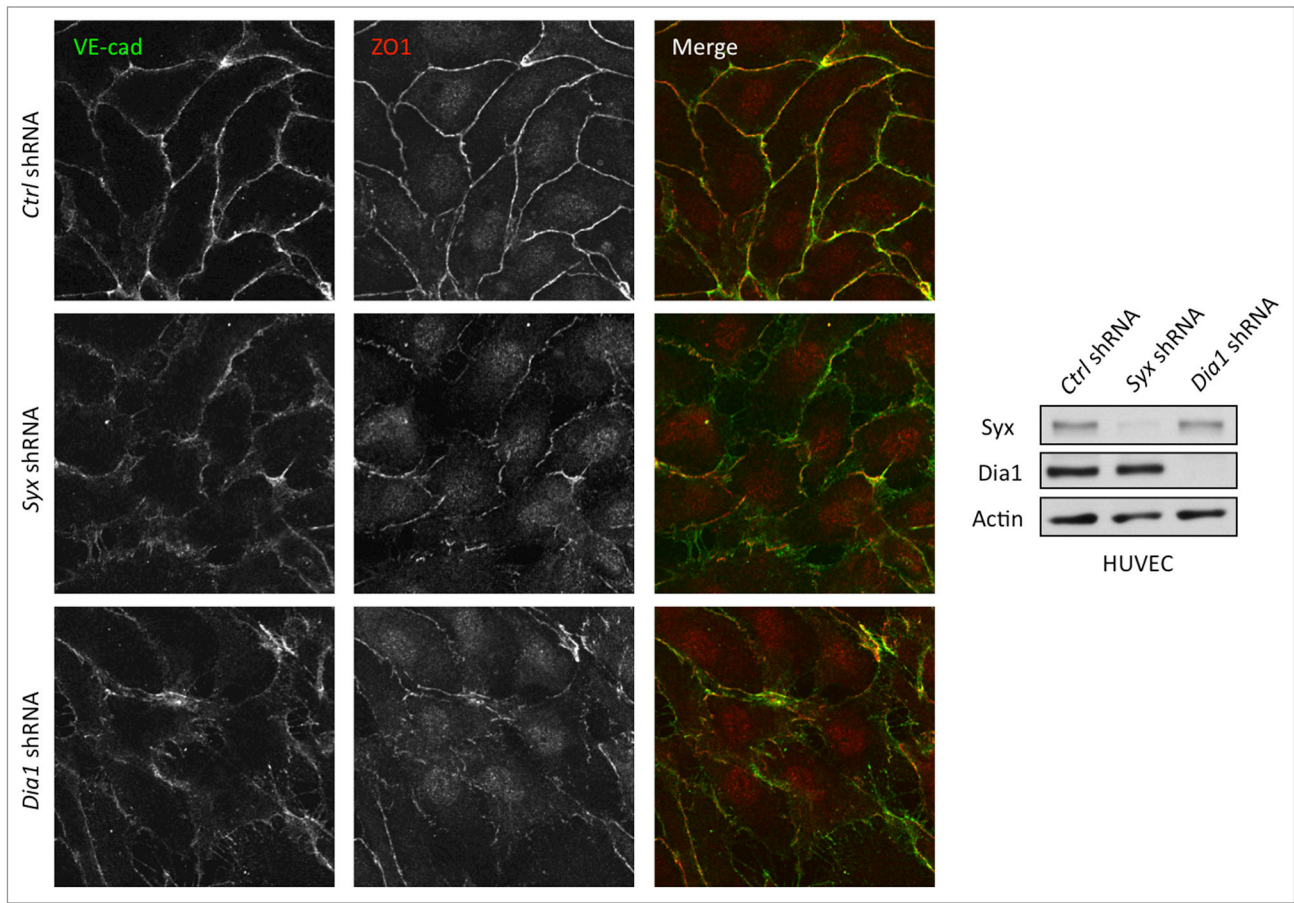
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**Figure 1.** Effect of silencing endogenous Syx or Dia1 on VE-cadherin and ZO1 localization in HUVECs. Western blot analysis shows the silencing efficacy of the *syx* or *dia1* shRNA constructs in cells.

factor (Syx; also known as PLEKHG5 or TECH) is essential for the maintenance of intercellular junctions and the barrier function of endothelial cells (ECs).<sup>3</sup> Previous studies had established that Syx selectively activates RhoA,<sup>4-6</sup> is involved in EC migration,<sup>7</sup> and regulates angiogenesis in both the zebrafish and mouse.<sup>8</sup> Syx binds to and co-localizes with multi-PDZ domain protein 1 (Mupp1) and protein associated with Lin7 (Pals1),<sup>3,9,10</sup> members of the Crumbs polarity complex whose function in ECs is largely unclear. shRNA-mediated downregulation of Syx in human umbilical vein endothelial cells (HUVECs) produced a pronounced reduction in trans-endothelial impedance, a measure of barrier integrity and function, which was coupled to disrupted junctional localization of cortical actin, VE-cadherin and ZO1.<sup>3</sup> Consistent with these observations, when compared with *syx*<sup>+/+</sup> mice, *syx* knockout mice exhibited increased capillary leakage, as evidenced

by a higher rate of microsphere extravasation from tracheal venules and increased Evan blue dye leakage from vessels in the skin. Collectively, the data strongly suggest that Syx plays a key role in determining junction stability and vessel permeability both in vitro and in vivo.

Our data also indicated that the opposing effects of Vascular Endothelial Growth Factor (VEGF) and Angiopoietin-1 (Ang1) exert on cell junctions are mediated, at least in part, by junctional Syx. VEGF caused translocation of Syx from cell junctions, resulting in junction disassembly, whereas Ang1 maintained Syx at the junctions, promoting junction stabilization.<sup>3</sup> In agreement with the observation that Syx is required for Ang1-mediated stabilization of EC junctions, Ang1 is known to regulate vascular leakiness, vascularization, inflammation, as well as tumor cell intra- and extravasation.<sup>11-14</sup> On the other hand, the VEGF-induced translocation of Syx from EC

junctions was caused by protein kinase D1 (PKD1)-mediated phosphorylation of Syx at Ser806, which reduced Syx association to its junctional anchor, Mupp1. Removal of junctional Syx resulted in the activation of Src, which can destabilize both AJ<sup>15,16</sup> and TJ complexes,<sup>17,18</sup> inducing junction disassembly.

RhoA and Src signaling are thought to be involved in a number of conditions that depend on endothelial junction disassembly, including angiogenesis, leukocyte transendothelial migration, viral hemorrhagic fever, sepsis, and acute respiratory distress syndrome (ARDS).<sup>16,19-21</sup> The ability of Ang1 to antagonize VEGF-induced junction disassembly by regulating Syx's subcellular localization, potentially implicates Syx in these pathological conditions. Interestingly, we found that in addition to endothelial cells, Syx regulates junction stability and monolayer integrity of MDCK cells (unpublished observations), a prototypical epithelial cell model.

Disruption of monolayer integrity results in inflammation.<sup>22</sup> Consistent with this premise, *Syx* resides in the highly unstable 1p36.3 region, which is commonly altered in developmental and inflammatory diseases,<sup>4,23</sup> as well as in cancer.<sup>24</sup> Therefore, it is possible that *Syx* regulates inflammatory responses and/or acts as a tumor suppressor by promoting cell junction integrity. Alternatively, *Syx* may affect tumor cell invasion and metastasis by regulating tumor cell migration or intra/extravasation, as is the case with *Ang1*. *Syx* is already known to affect endothelial cell migration,<sup>7,10</sup> and the *Syx*-associated proteins *Pals1* and *Pals1*-associated TJ protein (*PATJ*) are required for the directional migration of epithelial cells.<sup>25</sup> Importantly, of 303 *Syx* polymorphisms identified to date, 13 are missense mutations that could affect *Syx* activity (NCBI and Applied Biosystems SNP databases). It is intriguing to speculate that some of these SNPs may be markers of compromised vascular integrity and elevated susceptibility of individual patients to pathological conditions such as viral hemorrhagic fever, sepsis, ARDS, or tumor metastasis.

Importantly, we identified *Diaphanous 1* (*Dia1*) as a *Syx* effector that modulates junction stability, which was opposed by *ROCK* (Rho kinase) activity in HUVECs. *Dia* and *ROCK*, two known effectors of Rho signaling, are involved in regulating cytoskeleton dynamics. They are important candidates proposed to impact cell adhesion and junction integrity, through mechanisms that are still largely unclear. The remaining of this article will review the findings of *Dia* and *ROCK* in cell junction remodeling, and discuss the possibility that balanced signaling of these two effectors downstream of specific Rho GEFs is required for proper cell junction formation.

### **Dia and ROCK: friends or foes?**

*Dia1* and *ROCK* are crucial regulators of the actin cytoskeleton: the former drives *de novo* nucleation and elongation of actin filaments,<sup>26</sup> the latter stabilizes existing actin filaments and facilitates actomyosin contraction,<sup>27</sup> and the two work together to modulate the actin cytoskeleton and regulate cellular behavior.<sup>28</sup>

Our work demonstrates the importance of *Syx*-mediated *Dia1* signaling in promoting EC adhesion.<sup>3</sup> In agreement, downregulation of *Dia1* results in junctional defects in HUVECs (Fig. 1). In contrast, activated *Dia1* preserves junction stability by sequestering *Src*, a known mediator of junction disassembly, to prevent  $\beta$ -arrestin-mediated *VE-cadherin* endocytosis.<sup>16,29</sup> In epithelial cells, expression of *Dia1* reinforces the adhesion zone, whereas *Dia1* knockdown results in decreased junctional *E-cadherin*.<sup>30</sup> *Formin-1*, a *Dia1* related protein, is required at the intercellular junctions to direct actin cable formation and to support cell-cell contact in mouse keratinocytes *in vivo*.<sup>31</sup> Furthermore, *Dia1* knockout mice exhibit severe defects in neuroepithelial junction formation and polarization, supporting its indispensable role in regulating cell-cell adhesion and monolayer integrity.<sup>32</sup> The observation that *Syx* is coupled to a polarity complex and acts through *Dia1* to stabilize cell-cell adhesion suggests that it acts as a critical checkpoint for maintaining normal cell monolayer physiology.

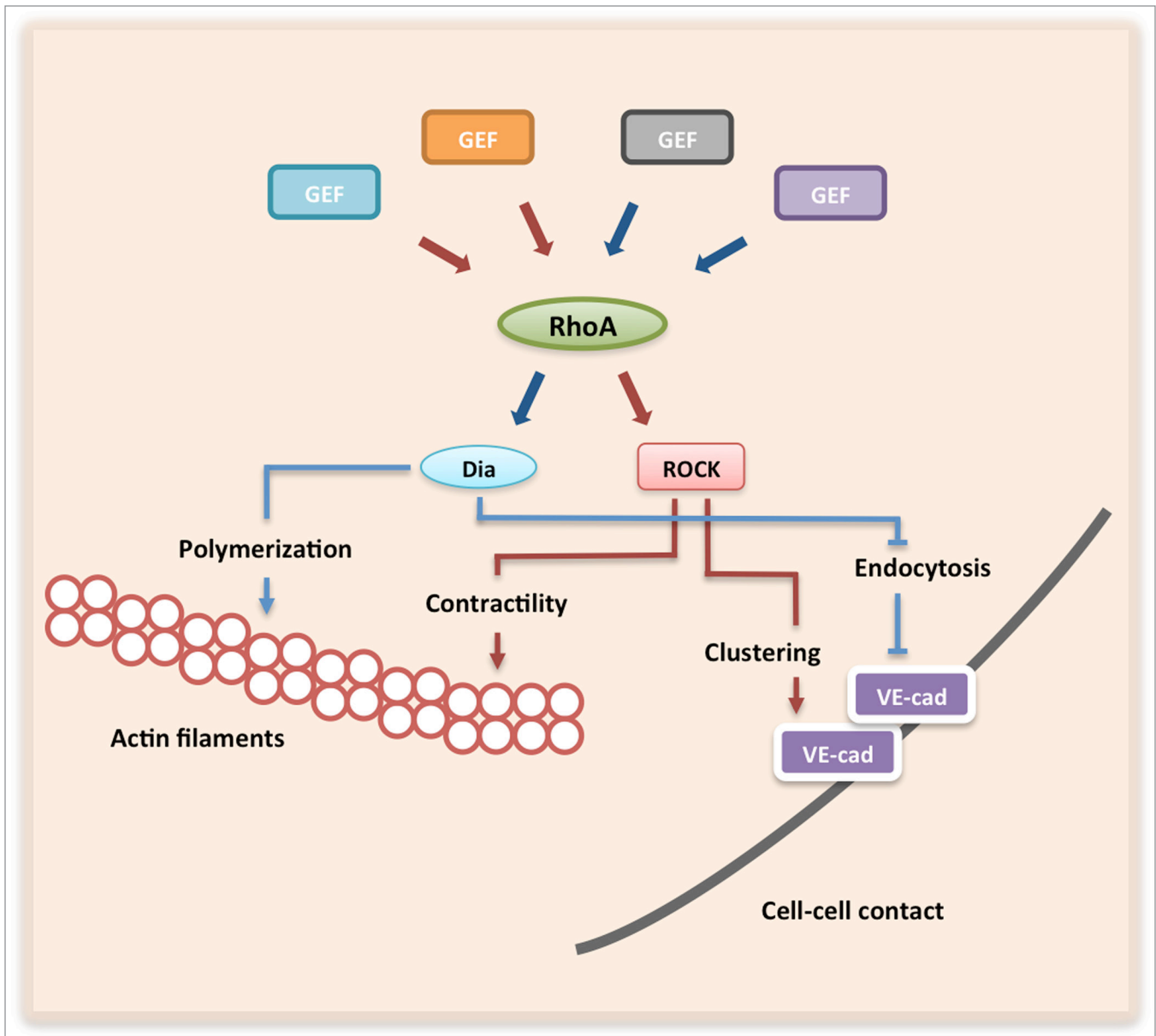
*ROCK*, on the other hand, appears to play a negative role in affecting EC junction stability, especially in the absence of *Syx*.<sup>3</sup> Indeed, *ROCK* can disrupt AJ and TJ integrity by increasing actomyosin contractility.<sup>33-36</sup> Inhibition of *ROCK* activity is directly correlated with an increase in junction stability and monolayer barrier function, suggesting that *ROCK* plays a role in the induction of monolayer leakiness.<sup>37,38</sup> Consistent with this, *ROCK* is required for *VEGF*, histamine, thrombin, and bradykinin-induced permeability, indicating that it is one of the key regulators of endothelial junction disassembly.<sup>39-41</sup> Interestingly, evidence has also emerged to support a role for *ROCK* in promoting junction formation and stabilization. Depletion of endogenous *ROCK* was shown to affect endothelial barrier function and junctional *VE-cadherin* in ECs,<sup>42</sup> as well as cell-cell adhesion in epithelial cells.<sup>43</sup> Inhibition of *ROCK* activity compromised the organization of the actin cytoskeleton and the proper distribution of AJ components, thus preventing normal junction formation.<sup>44,45</sup> These effects likely involve *ROCK*-mediated Myosin

II activation at areas of cell-cell contact, which is thought to promote cadherin clustering and stabilization at the AJs.<sup>46</sup> It is important to note that the two isoforms of *ROCK*, *ROCK1* and *2*, have both overlapping and unique functions in endothelial cells. Both *ROCK* isoforms regulate stress fiber formation. However, *ROCK2* is a more potent driver of endothelial cell migration and *in vitro* angiogenesis,<sup>47,48</sup> perhaps attributed to its selective ability to induce myosin phosphatase and cofilin phosphorylation.<sup>49</sup> In contrast, *ROCK1* is enriched at areas of endothelial cell-cell contact and physically associates with the cadherin complex via p120 catenin, suggesting that it is more relevant in regulating cell-cell adhesion.<sup>43,49</sup> Differences in experimental design, cellular background, and isoform expression could possibly explain the contradicting findings of *ROCK* in junction regulation; however, the phenotypes observed may represent the outcome of highly regulated signaling, where the function of both *Dia1* and *ROCK* and their crosstalk should be taken into consideration.

### **Working in synergy: selective and balanced signaling by *Dia* and *ROCK***

The following propositions, which take into consideration recent findings, may clarify how *Dia1* and *ROCK* can work together to modulate cell-cell cohesion.

First, the recruitment to specific subcellular locations and the strength of *Dia1* and *ROCK* activities are critical determinants for cellular outcome. *Dia1* is proposed to have a housekeeping function in stabilizing cell junctions.<sup>30</sup> Even though the mechanism of *Dia1* recruitment to cell-cell contacts is unclear, its basal activity is likely maintained by upstream *RhoA* signaling at the junctions. *ROCK* binds several junctional proteins such as p120, *Shroom 3*, and p114*RhoGEF*,<sup>43,50,51</sup> suggesting that its recruitment to the cell membrane is highly regulated and that its activation is dependent on its immediate associated protein complex. In a quiescent cell monolayer, *ROCK* and *Dia1* may function together<sup>28</sup> to stabilize the junctions through: a) the clustering of adhesion receptors and the suppression of



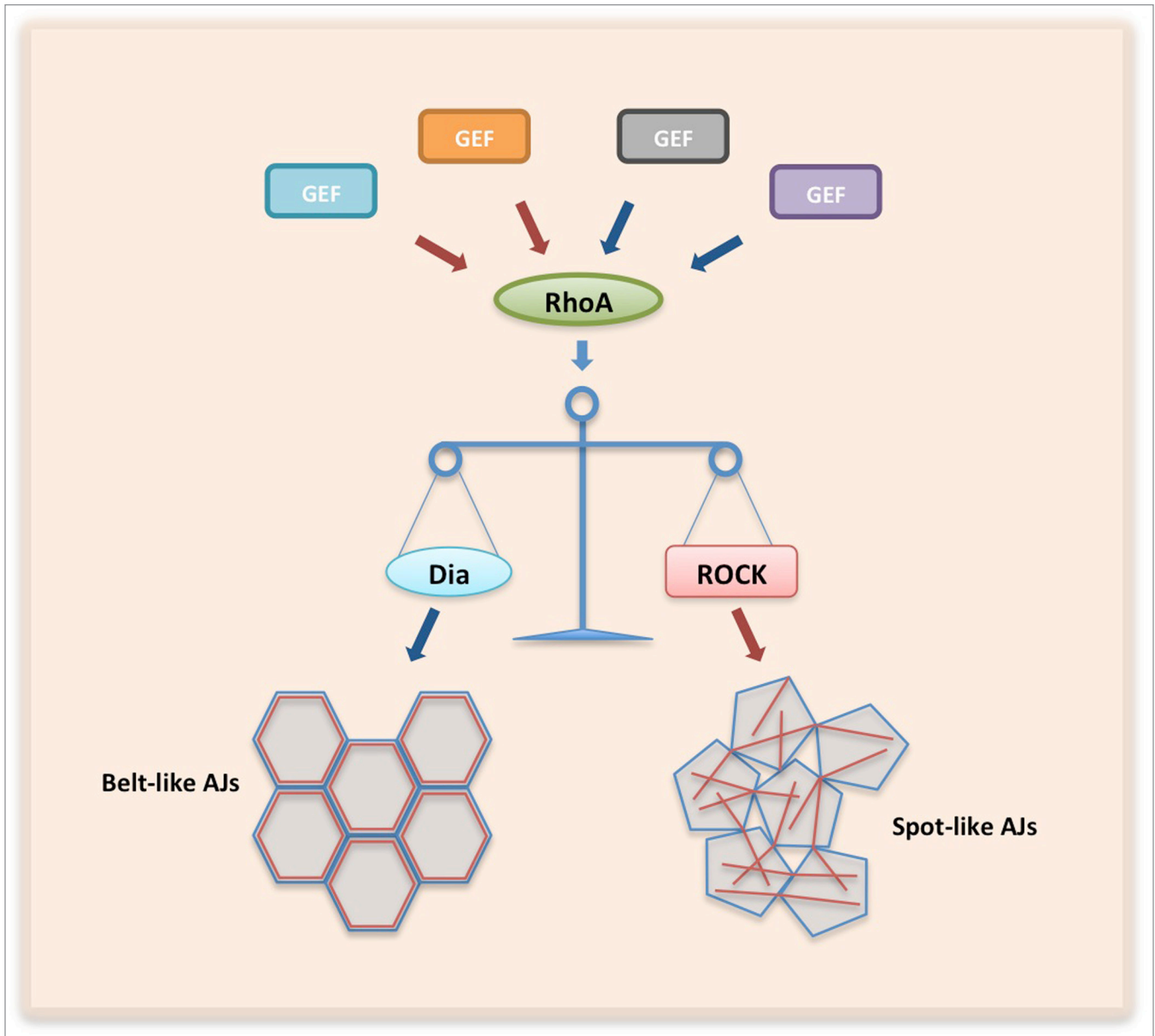
**Figure 2.** Effects of Dia1 and ROCK signaling on endothelial junctions and their underlying actin cytoskeleton. Junctionally localized RhoGEFs selectively activate Dia1 (blue arrows) or ROCK (red arrows) via RhoA. Dia1 promotes actin polymerization while ROCK induces actomyosin-induced contractility. Furthermore, Dia1 and ROCK promote junction integrity independently by inhibiting VE-cadherin endocytosis and promoting VE-cadherin clustering, respectively, at sites of cell-cell contact.

cadherin endocytosis, and b) by promoting the formation and alignment of actomyosin bundles parallel to the cell border at the mature AJs of the Zonula Adherens (ZA) (Fig. 2). In the context of a sealed, patent monolayer, the parallel orientation of the actomyosin bundles could minimize contractile forces directly impacting the cell-cell junctions, allowing adhesive forces to prevail. We postulate that when junction integrity is altered due to depletion of essential junctional proteins (e.g.,

in cancer), suppression of Dia1, or presence of extracellular signaling inducing further ROCK activation, the balanced act between Dia1 and ROCK is compromised, causing either unopposed or increased actomyosin contractility that induces junction disassembly and promotes cell migration<sup>52</sup> (Fig. 3). While not tested directly, it is possible that these conditions induce the transition of the belt-like ZA to the recently reported spot-like (or punctate) AJ morphology (pAJs),<sup>53</sup> and the change

from parallel circumferential actomyosin bundles to actin fibers laterally attached to the pAJs (Fig. 3). Our data are consistent with such a transition, which could lead to increased contractile forces exerted at the pAJs, leading to ROCK-mediated junction disassembly.<sup>3</sup> Indeed, we showed that ROCK inhibition partially reverses junctional defects in Syx downregulated cells,<sup>3</sup> supporting the hypothesis that the combined effect of Dia and ROCK determines EC junction stability.





**Figure 3.** Proposed model of balanced Dia1 and ROCK activities in cell-cell cohesion. Under normal physiological conditions, the two Rho effectors work in synergy to align and maintain a circumferential actomyosin bundle, and to promote the formation of mature belt-like endothelial AJs. However, loss of Dia1 signaling, and/or increased signaling via ROCK promotes the transition of belt-like AJs to punctate AJs. This transition is accompanied by a shift from the parallel, circumferential actomyosin bundles into unorganized, radially attached actin stress fibers that exert contractile forces able to disrupt junction stability.

Additionally, the direct crosstalk between Dia and ROCK may be an integral mechanism for modulating cell-cell cohesion. Formin homology domain protein 1 (FHOD1), an endothelial enriched protein closely related to Dia1, is activated via phosphorylation by ROCK.<sup>54</sup> FHOD1 also physically associates with ROCK in a Src-dependent manner and facilitates cytoskeleton rearrangement.<sup>55</sup> Furthermore, phosphorylation by ROCK

increases the ability of mDia2, another Dia family member, to polymerize actin filaments.<sup>56</sup> Interestingly, Dia1 can function upstream of Rho/ROCK signaling in mediating neutrophil chemotaxis and regulating cancer cell morphology.<sup>57,58</sup> Although additional studies are needed to elucidate the specific roles of these two Rho effectors in the context of the intercellular junctions, it is likely that junctional Dia and ROCK corroborate with

each other to promote the formation and stabilization of intercellular junctions, while Dia1 or ROCK misregulation leading to unopposed contractility at the pAJs induces junction disassembly.

An important question for further study is how junctional Rho activation is coupled to either Dia or ROCK signaling. A potential mechanistic explanation is the existence of different junctional RhoGEF-Rho modules that determine

selective effector signaling. For example, p114RhoGEF-associated ROCK II activation<sup>51</sup> may cooperate with Syx/Dial in mediating junction stability. The close proximity of the two Rho GEF complexes at the junctions could facilitate the synergistic effect of Dia and ROCK, thus positively influencing intercellular junctions. A similar condition may exist between the Rho GEFs TEM4 and Syx. As with Syx, TEM4 is essential for EC junction integrity and barrier function. Unlike Syx, which utilizes a C-terminal PDZ binding motif to interact with a polarity complex and localize to the junctions, TEM4 utilizes its N-terminal domain to associate with the cadherin-catenin complex at the AJs.<sup>59</sup> While the downstream effector of TEM4-mediated RhoA activation is currently unknown, the data indicate that at least two Rho-specific exchange factors, TEM4 and Syx, are essential for EC junction integrity and barrier function. The inability of these Rho GEFs to complement each other's function at the junctions strongly suggests that RhoA activation is highly regulated and is critical at multiple steps in junction formation and maintenance.

In conclusion, the integration of fine-tuned Rho GTPase activation by junctional Rho GEFs coupled with selective effector signaling represents a highly versatile mechanism for the precise spatio-temporal regulation of cell-cell adhesion. Despite a recent surge of studies showing how selective Rho GEFs influence junction integrity and plasticity,<sup>3,51,59,60</sup> uncovering additional Rho regulators (GEFs, GAPs, or GDIs) that affect the formation and maintenance of intercellular junctions is essential for understanding the function of Rho GTPases and their effectors at intercellular junctions and their misregulation in human disease, including inflammation and cancer.

#### Disclosure of Potential Conflicts of Interest

No potential conflicts of interest were disclosed.

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