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ZAK negatively regulates RhoGDIβ-induced RacI-mediated hypertrophic growth and cell migration

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Abstract

RhoGDI β , a Rho GDP dissociation inhibitor, induced hypertrophic growth and cell migration in a cultured cardiomyoblast cell line, H9c2. We demonstrated that RhoGDI β plays a previously undefined role in regulating RacI expression through transcription to induce hypertrophic growth and cell migration and that these functions are blocked by the expression of a dominant-negative form of RacI. We also demonstrated that knockdown of RhoGDI β expression by RNA interference blocked RhoGDI β -induced RacI expression and cell migration. We demonstrated that the co-expression of ZAK and RhoGDI β in cells resulted in an inhibition in the activity of ZAK to induce ANF expression. Knockdown of ZAK expression in ZAK-RhoGDI β -expressing cells by ZAK-specific RNA interference restored the activities of RhoGDI β .

Background

The mitogen-activated protein kinase (MAPK) signaling pathway consists of the sequentially acting upstream kinases MAPK kinase kinase (MAP3K) and MAPK kinase (MAP2K), and the downstream MAPKs, p38MAPK, extracellular signal-regulated kinase (ERK1/2), and c-jun N-terminal kinase (JNK). The mixed lineage kinases are a family of serine/threonine kinases, all of which are classified as MAP3Ks. The seven mixed lineage kinases cloned over the past several years can be classified into three subfamilies based on domain organization and sequence similarity: the MLKs (MLK1–4), the dual leucine zipperbearing kinases (DLK and LZK), and the zipper sterile- α motif (SAM) kinases (ZAK α and ZAK β) [1,2]. ZAK can activate the JNK pathway and the nuclear factor κ B (NF κ B) pathway [3], and it induces JNK activation through a dual phosphorylation kinase, JNKK2/MKK7 [4]. Overexpression of wild-type ZAK induced apoptosis in a hepatoma cell line [3], and a recent report indicated that ZAK expression in a rat cardiac cell line, H9c2, induced hypertrophic growth and re-expression of atrial natriuretic factor (ANF) [5]. ZAK also mediates TGF- β - induced cardiac hypertrophic growth via a novel TGF- β signaling pathway [6]. In our previous study [5], we showed that the leucine zipper of ZAK mediates homodimerization and promotes autophosphorylation and JNK activation.

We identified RhoGDI_β(Rho GDP dissociation inhibitor beta) as a ZAK effector. RhoGDIB, also known as Ly-GDI or D4-GDI, belongs to a family of Rho GDP dissociation inhibitors, and it is thought to regulate the activity and localization of Rho family proteins [7-10]. The RhoGT-Pase family includes Rho, Rac, and Cdc42, which differentially regulate the actin cytoskeleton [11-13] and function as molecular switches in cellular signal transduction by alternating between an inactive GDP-bound form that is maintained in cytosolic complexes with GDIs and a GTPbound form that usually associates with the plasma membrane and interacts with downstream target proteins therein [14,15]. RhoGTPases regulate the reorganization of the actin cytoskeleton and the integrity of the integrinassociated adhesion complexes [16]. Rho facilitates stress fiber and focal adhesion assembly, Rac regulates the formation of lamellipodia and membrane ruffles at the leading edge of migrating cells, and Cdc42 triggers filopodia at the cell periphery [17]. RhoGDIs regulate RhoGTPase activity by inhibiting GDP dissociation to keep RhoGT-Pases in an inactive state.

A recent study indicated that stimulation of T lymphocytes and myelomonocytic cells with phorbol esters leads to RhoGDI β phosphorylation on serine/threonine residues [18], raising the question of whether RhoGDI β is involved in a signal transduction pathway in these cells. Thus, RhoGDI β may play numerous roles in the regulation of biological activities; however, many details of the regulatory roles of RhoGDI β remain to be elucidated.

Materials and methods Northern blot analysis

Trizol reagents (Life Technologies) were used to isolate total RNA from H9c2 cells transiently transfected with the recombinant RhoGDI β plasmids or from cells stably expressing RhoGDI β . Total RNA (20 µg) was separated on a formaldehyde agarose gel, transferred to a nylon filter, and then hybridized with a probe corresponding to the full-length Rac1 cDNA labeled using the NEBlot random labeling kit (New England BioLabs) in the presence of [α -³²P]dCTP. The blot was washed with SSC/SDS solutions (Sodium Chloride, Sodium Citrate/SDS) before autoradiography. Ethidium bromide staining was used to check the integrity of all samples.

Wound healing assay

H9c2 cells seeded on 10-cm plates were cultured to confluency. They were then scratched with a 200 μl pipette tip

and further incubated in DMEM supplemented with 10% FBS. Images were taken at 24, 48, and 72 h with a Zeiss Axiovert 200 microscope. The Image-Pro image analysis system was used to measure the lesion area. The data were expressed as the percentage of recovery (WC%) using the equation: WC% = [1 - (wounded area at T_t/wounded area at T₀)] × 100%, where T_t is the number of hours post-injury and T₀ is the time of injury.

Membrane and cytosol fractionation

H9c2 cells were cultured with 1 μ g/ml doxycycline for 48 h and then treated with lysis buffer (20 mM Tris-HCl, pH 7.5, 100 mM NaCl, 5 mM EDTA, 2 mM PMSF, 1× protease inhibitor) at 4°C for 30 min. The samples were centrifuged at 500 × g at 4°C for 10 min, and the pellets were dissolved in lysis buffer plus 0.1% (w/v) Triton X-100 for the membrane fractions. The supernatants were re-centrifuged at 15,000 rpm at 4°C for 20 min, and the supernatants were saved as cytosolic fractions.

Results

Expression of RhoGDI β induces hypertrophic growth via modulation of Rac1 expression in cardiac cells

The RhoGTPases act as molecular switches by cycling between the inactive GDP-bound form located in the cytoplasm and an active membrane-associated GTPbound form. The activities of Rho family proteins are regulated by various proteins, such as guanine nucleotide exchange factors (GEFs), GTPase activating proteins (GAPs), and GDIs. The functions and binding of RhoG-DIα to RhoA, Rac1, and Cdc42 are well studied; however, the functions and targets for RhoGDIβ remain unclear, as it binds poorly to RhoA, Rac1, and Cdc42. We therefore sought to determine whether RhoGDIß stimulates the expression or activities of these GTPases in cardiac cells by western blotting. The total RhoA and Cdc42 levels remained the same; however, cells overexpressing RhoGDIβ had increased levels of Rac1 (Figure 1A). Moreover, the expression of RhoGDIa in H9c2 cells was not affected by the overespression RhoGDI β (Figure 1A). The expression levels of Rac1 may be causally linked to RhoGDIβ expression or merely an epiphenomenon of the selection of a stable clone. If the former is the case, then Rac1 might be a functionally important downstream target of RhoGDIB. Northern blot analysis of total RNA indicated that cardiac cells either stably or transiently overexpressing RhoGDIB had increased Rac1 mRNA levels (Figure 1B). We therefore concluded that RhoGDIß plays a role in the transcriptional regulation of Rac1 and that the increased level of Rac1 mRNA was not a secondary effect of the selection of a stable RhoGDIB-expressing clone. Rac1 association with membranes reflects its biological activity [19]. To further address the question of whether induction of Rac1 may also influence Rac1 activity, a detergent-insoluble membrane fraction was pre-

RhoGDIß



Figure I

Effects of RhoGDI β **on Rac1 expression and subcellular localization of RhoGTPases**. (A) RhoGDI β , RhoGDI β , Rho, CDC42, and Rac1 were detected by western blotting of cell lysates from H9c2 cells stably expressing RhoGDI β . (B) Total RNA was isolated from H9c2 cells stably or transiently expressing RhoGDI β . Rac1 transcripts were analyzed by northern blotting. The lower EtBr panels of (B) represent the 28S and 18S loading controls. (C) Membrane (Mem.) and cytosolic (Cyto.) fractions from H9c2 control (C) and RhoGDI β -expressing cells were analyzed by immunoblotting for Rac1, Cdc42, and RhoGDI β . (D) GTP loading in Rac1 was determined utilized PAK PBD binding assay in H9c2 cells stably expressing RhoGDI β .

pared from RhoGDIβ-overexpressing cells, and the levels of membrane-associated Rac1 were determined by western blotting. Both the levels of membrane-associated Rac1 and cytosolic Rac1 increased in RhoGDIβ-overexpressing cells (Figure 1C). Furthermore, the membrane-associated and cytosolic forms of Cdc42 remained unchanged in RhoGDIβ-expressing cells when compared to parental cells. We further detect the amount of GTP-bound Rac1 in H9c2 RhoGDIβ-overexpressing cells. We found that RhoGDIβ increased GTP loading in Rac1 (Figure 1D). Therefore, overexpression of RhoGDI β in H9c2 cells increases the level of membrane-associated Rac1 and GTP loading in RAC1 by upregulating Rac1 transcripts. However, the increase in membrane-associated Rac1 in RhoGDI β -expressing cells may be a secondary effect of increased expression of Rac1, as we were not able to detect any physical interaction between RhoGDI β and Rac1 by co-immunoprecipitation.

We next examined whether Rac1 mediates RhoGDIBinduced hypertrophic growth. We found that H9c2 cells stably expressing a dominant-negative form of Rac1 (Rac1N17) and RhoGDIß significantly reduced the twofold increase in cell size (Figure 2A) and actin organization induced by RhoGDIB. Moreover, overexpression of wild-type or a constitutively active (V12) Rac1 in H9c2 cells was sufficient to induce hypertrophic growth (Figure 2A) and actin organization. These findings indicate that the RhoGDIβ-induced hypertrophic growth in H9c2 cells is mediated through increased expression of Rac1, and probably through increased levels of membrane-associated Rac1.

H9c2 cell migration promoted by RhoGDI β is Rac1 dependent

Rac1 is able to regulate cell migration [20]. To test whether the effect of RhoGDIß on cell migration is Rac1 dependent, confluent monolayers of cells stably expressing RhoGDIB, RhoGDIB and Rac1, or RhoGDIB and Rac1N17 were scrape-wounded with a sterile plastic pipette, and the migration of cells into the wound was monitored. The RhoGDIβ-expressing cells closed the



48h

Figure 2

RhoGDIβ-induced hypertrophic growth and cell migration is RacI-dependent, but suppression of cell cycle progression is not. (A) Control H9c2 and H9c2 RhoGDI β -expressing cells stably transfected with Rac1, Rac1N17, or RacIV12 were grown in 10% fetal bovine serum with doxycycline for three days, and the cell size was determined. (B) Woundhealing assay. The cell lines were seeded on plates, as previously described. After reaching confluency, the cell layer was wounded with a 200 μ l pipette tip and incubated for 48 h with medium and doxycycline. (C) SiRNA knockdown of RhoGDI β inhibited RhoGDIβ-induced Rac1 expression levels in H9c2 cells. U6-Ci is the scramble control siRNA, and both U6-353i and U6-568i are two specific RhoGDI β siRNAs.

wound area faster than control cells (Figure 2B). RhoGDIβ-expressing cells that also expressed a dominantnegative form of Rac1N17 migrated slower than RhoGDIβ-expressing or RhoGDIβ- and Rac1-expressing cells (Figure 2B), suggesting that Rac1 plays a key role in mediating cell migration in RhoGDIβ-expressing cells.

RhoGDI β -induced cell migration does not correlate with cell proliferation

To determine whether Rac1 might play a role in the RhoGDIβ-mediated cell cycle arrest, we examined the growth rate of cells expressing both RhoGDIβ and Rac1N17 and found that it was substantially slower than the growth rate of RhoGDIβ-expressing cells or control cells (Additional file 1). Therefore, the RhoGDIβ-regulated cell arrest was not mediated through increased levels of membrane-bound active Rac1.

Since the cyclin-dependent kinase inhibitors $p21^{Waf1/Cip1}$ and $p27^{Kip1}$ were expressed in RhoGDI β -expressing cells at higher levels than they were in control cells, we examined $p21^{Waf1/Cip1}$ and $p27^{Kip1}$ levels in RhoGDI β - and Rac1N17expressing cells to study the effects of this dominant-negative form of Rac1 on the expression of p21 and p27. RacN17 was unable to block the expression of p21^{Waf1/Cip1} and p27^{Kip1} induced by RhoGDI β in H9c2 cardiac cells (Additional file 2), suggesting that the RhoGDI β -induced cell cycle arrest was not mediated through Rac1.

Knockdown of overexpressed RhoGDI β by siRNA reduces H9c2 cell migration

To this point, our results indicated that RhoGDIß stimulates cell migration through the induction of Rac1 expression due to a proportional increase in the activity of Rac1 in H9c2 cells. To confirm the role of RhoGDIß in cell migration, two RhoGDIB knockdown cell lines were used to assess whether RhoGDIB directly stimulates Rac1 expression to induce cell migration. The specific knockdown of RhoGDIB using siRNA in H9c2 cells was confirmed by immunoblotting (Fig. 2C). We found that targeted disruption of RhoGDIB by siRNA effectively blocked expression of Rac1 (Fig. 2C); therefore, RhoGDIβ depletion is associated with Rac1 downregulation. We used migration assays to confirm the role of RhoGDIß in cell migration. Cells were seeded in an upper chamber of a Transwell on a porous filter, and the migration through the filter pores of H9c2 cells expressing RhoGDIB and cells expressing both RhoGDIß and RhoGDIß- specific siRNA was compared. RhoGDIβ-expressing cells showed increased migration compared to parental cells, whereas migration was inhibited in the siRNA RhoGDIB knockdown cells relative to the RhoGDIβ-expressing cells (Additional file 3). These results suggest that RhoGDI^β may play a critical role in the regulation of Rac1 expression and H9c2 cell migration.

RhoGDI β -induced wound healing is negatively regulated by ZAK

Our experiments demonstrated that RhoGDIB increased the rate of wound closure. However, our data also suggested that the physical association of ZAK with RhoGDIß and phosphorylation of RhoGDIß by ZAK could abolish RhoGDIß function. To identify the regulatory role of ZAK on cell migration responses in RhoGDIβ-expressing cells, a wound-healing assay was performed using cells expressing RhoGDIB, ZAK and RhoGDIB, or ZAKdn and RhoGDIB. After wounding, control, ZAK-expressing, and ZAKdn-expressing cells closed the gap at a similar rate (Figure 3A). As demonstrated above, RhoGDIβ-expressing cells were highly migratory, but ZAK- and RhoGDIBexpressing cells were substantially slower to close the wound than ZAKdn- and RhoGDIB-expressing cells. At 36 h post-wounding, control, ZAK-expressing, ZAKdnexpressing, and ZAK- and RhoGDIβ-expressing cells closed the gap by about 20%, 24%, 25%, and 22%, respectively (Figure 3B). RhoGDIβ-expressing cells and ZAKdn- and RhoGDIβ-expressing cells closed 50% and 39% of the gap. These data suggested a negative regulatory role for ZAK in RhoGDIβ cell migratory function.

To elucidate the role of RhoGDIß in controlling the localization of Rac1, we performed confocal microscopy. The majority of Rac1 was present in the perinuclear region with some present at the plasma membrane of the control, ZAK-expressing, and ZAKdn-expressing cells (Additional file 4). More Rac1 was present at both the plasma membrane and the perinuclear region in RhoGDIβexpressing cells. However, co-expression of ZAK, but not ZAKdn, with RhoGDIβ in H9c2 cardiac cells decreased the amount of Rac1 at the plasma membrane (Additional file 4). These data also suggested that ZAK might decrease the total amount of cellular Rac1, probably due to ZAK binding and phosphorylation of RhoGDIB. To determine whether ZAK negatively regulates RhoGDIB through expression of Rac1, especially membrane-associated Rac1, we performed western blotting. Consistent with the above experiment, membrane-associated Rac1 increased in RhoGDI\beta-expressing cells, whereas, in ZAK- and RhoGDIβ-expressing cells, the levels of membrane-associated Rac1 decreased. However, co-expression of ZAKdn and RhoGDIB had no such effect (Figure 4A). In this regard, it was of interest whether RhoGDIB increases the amount of Rac1 at the plasma membrane as a consequence of increasing the total cellular levels of Rac1 or whether RhoGDIß facilitates the translocation of Rac1 to the plasma membrane. We attempted to co-immunoprecipitate RhoGDIß and Rac1to investigate this possibility; however, the antibody used was insufficient for this purpose (data not shown). This suggests that it is unlikely that RhoGDIB facilitates translocation of Rac1 to the plasma membrane.



Figure 3

ZAK reverses the effects of RhoGDI β on the induction of cell migration in H9c2 cells. (A) Wound healing assay. H9c2 cells and H9c2 cells ectopically expressing ZAK, dominant-negative ZAK (ZAKdn), RhoGDI β , ZAK and RhoGDI β , or ZAKdn and RhoGDI β were seeded onto plates. After reaching confluency, the cell layer was wounded with a 200 μ l pipette tip and incubated for 72 h with medium and doxycycline. (B) The cell migration capacity at 36 h was estimated by measuring the percentage wound closure (WC%). Values are means (SEM from three independent experiments).



Figure 4

ZAK specifically downregulates the activities of RhoGDI β as a consequence of decreasing the amount of membrane-associated Rac1. (A) ZAK decreases the levels of membrane-associated Rac1 induced by RhoGDI β . Control H9c2 and transfected H9c2 cells were collected and fractionated into membrane (Mem.) and cytosolic (Cyto.) fractions by centrifugation. (B) SiRNA knockdown of ZAK restores the effects of RhoGDI β in ZAK- and RhoGDI β -expressing cells (GRi1 and GRi2) upon Rac1 membrane association. (C) RhoGDI β inhibits ZAK functions upon the induction of ANF mRNA expression. (D) The effect of ZAK-RhoGDI β and RhoGDI β on total amount of Rac1 expression.

To test the importance of ZAK in regulating RhoGDIβmediated membrane-associated Rac1 and hypertrophic growth, we reduced the levels of ZAK using siRNA. We were able to reduce the levels of ZAK by expressing two different human ZAK siRNAs: ZAKGRi1 (ZAK-RhoGDIβ U6-460i) and ZAKGRi2 (ZAK-RhoGDIβ U6-1712i) (data not shown). The reduced levels of ZAK in these two individual clones were able to restore the levels of membraneassociated Rac1 to levels similar to those of RhoGDIβexpressing cells, whereas the introduction of a scrambled ZAK siRNA (GRCi) into ZAK- and RhoGDIβ-expressing cells had no effect (Figure 4B). These results confirmed the importance of ZAK as a negative regulator of the effect of RhoGDIβ on the expression of Rac1 and membrane-associated Rac1. We found that ZAK was able to regulate ANF expression. We then studied the effects of RhoGDI β on the regulation of ANF expression by ZAK by examining the levels of ANF mRNA in ZAK- and RhoGDI β -expressing cells when compared with ZAK-expressing cells. The levels of ANF mRNA induced by ZAK were decreased in cells expressing both ZAK and RhoGDI β (Fig. 4C). We also found that ZAK-RhoGDI β cells has less total amount of Rac1 than RhoGDI β cells. (Figure 4D). This result suggested that RhoGDI β negatively regulates the functions of ZAK. Moreover, the data presented here (and consistent with our previous studies) indicate that both ZAK and RhoGDI β may be hypertrophic growth inducers; however, ZAK physically interacts with RhoGDI β and phosphor-

ylates RhoGDI β , thus inhibiting the ability of RhoGDI β to induce Rac1 expression. The levels of Rac1 induced by RhoGDI β are associated with the closure rate of wound healing (Figure 2B) and hypertrophic growth (Figure 2A), but they are not associated with cell cycle inhibition (Additional file 1). Thus, RhoGDI β appears to play a role in signaling pathways regulating Rac1 expression that govern wound healing and hypertrophic growth in cardiac cells.

Discussion

Upon introduction of RhoGDI β into rat cardiac H9c2 cells, the cells exhibited hypertrophic growth, had a slower cell cycle, and migrated to a greater extent. We previously demonstrated that RhoGDI β is phosphorylated by ZAK *in vitro*. It is striking that the co-expression of ZAK and RhoGDI β in H9c2 cardiac cells results in the inhibition of the biological functions of RhoGDI β , indicating that not only does RhoGDI β possibly physically interact with ZAK, but it may also be negatively regulated by ZAK, and this regulation might occur via phosphorylation. These phenomena regulated by ZAK were correlated with Rac 1 expression and especially with the levels of membrane-associated Rac1 in H9c2 cells.

In H9c2 cells transiently and stably expressing RhoGDIβ, the levels of Rac1 transcripts increased compared with control cells. In this study, we described these surprising findings and, to our knowledge, the first demonstration that expression of RhoGDIB induces Rac1 transcripts and increases the levels of membrane-associated Rac1. The results from western blotting and confocal microscopy experiments indicate that RhoGDIß regulates Rac1 expression, which leads to increased levels of membrane-associated Rac1. We propose either that the increased levels of membrane-associated Rac1 in RhoGDIß cells are merely a consequence of RhoGDIβ-induced expression of Rac1 or that RhoGDIβ regulates membrane translocation of Rac1. We were unable to detect an association between RhoGDIβ and Rac1 using co-immunoprecipitation; therefore, it is unlikely that RhoGDIβ and Rac1 directly interact. However, we still have not ruled out the possibility that RhoGDIß regulates Rac1 translocation. However, the signaling pathway between RhoGDIß and Rac1 has not yet been elucidated, and there is currently no evidence that RhoGDIß can directly bind to any gene promoter. RhoGDIß can be translocated into the nucleus upon certain stimuli [21], leaving the possibility that RhoGDIß can regulate gene expression directly. It is also possible that RhoGDIβ regulates Rac1 expression via signaling pathway effector proteins. Studies have also demonstrated that RhoGDIβ is cleaved at its N-terminus during apoptosis in a caspase-dependent manner and that the cleaved RhoGDI β is retained in the nuclear compartment [22]. This suggests that RhoGDI^β could function in the nucleus.

We previously found that RhoGDI^β was able to associate with a mixed lineage kinase, ZAK, resulting in the phosphorylation of RhoGDI^β. To further study the role of ZAK in regulating the activities of RhoGDIB, we used a bi-directional tet-on inducible system to express both ZAK and RhoGDIß in H9c2 cardiac cells. Our results demonstrate that the levels of membrane-associated Rac1 and the hypertrophic growth phenotype were inhibited by coexpression of ZAK and RhoGDIB; however, we did not observe the inhibitory effect with a dominant-negative form of ZAK. Clearly, the kinase activities of ZAK are necessary for the negative regulation of RhoGDIB functions, including cell cycle arrest, hypertrophic growth, alterations in the amount of membrane-associated Rac1, and cell migration. Among all the biological functions that are regulated by RhoGDI^β, the phenomena of hypertrophic growth and cell migration are Rac1-dependent, whereas the regulation of the cell cycle arrest is Rac1-independent, as shown by the results of expression of a dominant-negative Rac1 (Rac1N17) in RhoGDIβ-expressing cells. It should be pointed out that the cell migration phenotype induced in RhoGDIβ-expressing cells seems to result primarily from the rate of migration rather than cell division. Therefore, RhoGDIB may stimulate cell migration in a manner dissociated from its effects on cell cycle progression.

Competing interests

The authors declare that they have no competing interests.

Authors' contributions

CYH was responsible for the pull-down assay. LCY was responsible for the SiRNA knockdown. KYL was responsible for the cell migration assay. ICC, JIY, MYC and WWL were responsible for Western blot analysis. PHL was responsible for cell staining. JJY wrote the manuscript.

Additional material

Additional file 1

Figure S1. The growth rate of H9c2 cells expressing RhoGDIβ and Rac1N17 grown in 10% fetal bovine serum with doxycycline. Click here for file [http://www.biomedcentral.com/content/supplementary/1423-0127-16-56-S1.tiff]

Additional file 2

Figure S2. Expression levels of cyclin-dependent kinase inhibitors p21 and p27 in H9c2 RhoGDI β -expressing cells and H9c2 cells expressing both RhoGDI β and Rac1N17. Click here for file [http://www.biomedcentral.com/content/supplementary/1423-

[http://www.biomedcentral.com/content/supplementary/1423-0127-16-56-S2.tiff]

Additional file 3

Figure S3. SiRNA knockdown of RhoGDIβ inhibited cell migration of H9c2 RhoGDIβ-expressing cells. Click here for file [http://www.biomedcentral.com/content/supplementary/1423-0127-16-56-S3.tiff]

Additional file 4

Figure S4. Confocal microscopy of H9c2 and H9c2 cells expressing the indicated proteins.

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