Gα12/13 regulate epiboly by inhibiting E-cadherin activity and modulating the actin cytoskeleton

Fang Lin, Songhai Chen, Diane S. Sepich, Jennifer Ray Panizzi, Sherry G. Clendenon, James A. Marrs, Heidi E. Hamm, and Lilianna Solnica-Krezel

1 Department of Pharmacology, Vanderbilt University Medical Center, Nashville, TN 37232
2 Department of Pharmacology and 3 Department of Anatomy and Cell Biology, Carver College of Medicine, University of Iowa, Iowa City, IA 52242
4 Department of Biological Sciences, Vanderbilt University, Nashville, TN 37235
5 Department of Medicine, Indiana University University Medical Center, Indianapolis, IN 46202

Introduction

During vertebrate gastrulation, an embryo of simple and symmetrical morphology is reshaped to reveal its fundamental body plan. This process is accomplished by cooperation of four morphogenetic movements—epiboly, internalization, and convergence and extension (C&E)—that are largely conserved among vertebrates (Arendt and Nubler-Jung, 1999; Leptin, 2005; Solnica-Krezel, 2005). Epiboly starts at the late blastula stage as the yolk cell pushes into the blastoderm, which thins and expands vegetally until it encloses the entire yolk cell (Warga and Kimmel, 1990; Solnica-Krezel, 2006; Rohde and Heisenberg, 2007). At this stage, the embryo is comprised of four cell layers: the enveloping layer (EVL), deep cells, the yolk syncytial layer (YSL), and the yolk cell. The EVL is a superficial epithelial layer that covers a mass of deep cells, which give rise to embryonic tissues. The YSL is a shallow and superficial cytoplasmic layer within the yolk cell (Solnica-Krezel and Driever, 1994; Rohde and Heisenberg, 2007). Proper epiboly involves coordinated movements of all of these layers, and the underlying cellular and molecular mechanisms remain to be fully defined (Solnica-Krezel, 2006; Rohde and Heisenberg, 2007).

Recent studies indicate that E-cadherin–mediated cell–cell adhesion plays a critical role in zebrafish epiboly. In both E-cadherin mutant embryos and embryos injected with E-cadherin morpholino oligonucleotides (MOs) to block its translation, the epibolic movement of the deep cells is delayed or arrested at midgastrulation, although the YSL and EVL expand vegetally in a relatively normal fashion (Babb and Marrs, 2004; Kane et al., 2005; McFarland et al., 2005; Shimizu et al., 2005). This epibolic delay has been attributed to impaired radial intercalation resulting from decreased adhesion among the deep cells and between the deep cells and the EVL (Kane et al., 2005; Montero et al., 2005; Shimizu et al., 2005). An additional cell–cell adhesion

Correspondence to Fang Lin: fang.lin@uiowa.edu; Heidi E. Hamm: heidi.hamm@vanderbilt.edu; or Lilianna Solnica-Krezel: lilianna.solnica-krezel@vanderbilt.edu

Abbreviations used in this paper: C&E, convergence and extension; cad1, cadherin 1; CT, carboxy terminal; CyT, cytoplasmatic terminus; dcm, deep cell margin; df, dorsal forerunner; EVL, enveloping layer; GEF, guanine nucleotide exchange factor; GPFR, G protein-coupled receptor; hab, half-baked; LWR, length-to-width ratio; MO, morpholino oligonucleotide; nif, no tail; WT, wild type; YCL, yolk cytoplasmic layer; YSL, yolk syncytial layer; YSN, yolk syncytial nuclei.

© 2009 Lin et al. This article is distributed under the terms of an Attribution–Noncommercial–Share Alike License for the first six months after the publication date (see http://www.jcb.org/misc/terms.shtml). After six months it is available under a Creative Commons License Attribution–Noncommercial–Share Alike 3.0 Unported license, as described at http://creativecommons.org/licenses/by-nc-sa/3.0/.

Supplemental Material can be found at: /content/suppl/2009/03/21/jcb.200805148.DC1.html
defect was observed in E-cadherin–deficient embryos, with cells bulging and detaching from the embryonic surface (Babb and Marrs, 2004; Kane et al., 2005; McFarland et al., 2005; Shimizu et al., 2005).

E-cadherin is a plasma membrane glycoprotein that is indirectly linked to the actin cytoskeleton through β-catenin (Barth et al., 1997). The involvement of E-cadherin in morphogenesis and differentiation during the early development has been also demonstrated in many species including mouse, chick, and frog (Halbleib and Nelson, 2006). In addition, E-cadherin is essential for cell migration and polarity, as well as neuronal synapse function. E-cadherin expression is regulated at various levels including gene expression, protein stability, and intracellular protein distribution (Halbleib and Nelson, 2006). Downregulation of E-cadherin is regarded as the hallmark of the epithelial–mesenchymal transition, and is often observed in invasive tumor cells (Behrens, 1999).

In comparison to our fairly detailed knowledge about the regulation of E-cadherin expression, we know very little about regulation of its activity. However, recent studies in cell culture indicate that heterotrimeric G proteins of the Go12 family (Go12 and Go13) can modulate E-cadherin function: Go12/13 can bind E-cadherin at its cytoplasmic domain to block the β-catenin–binding site, resulting in inhibition of cell–cell adhesion (Kaplan et al., 2001; Meigs et al., 2001; Meigs et al., 2002). Nevertheless, the significance of the Go12/13 and E-cadherin interaction during morphogenesis remains to be tested.

During epiboly, the yolk cell may serve as a towing motor to drive the movements of epiboly. Nuclei of the YSL move vegetally even after removal of the blastoderm (Trinkaus, 1951), which indicates that the YSL can undergo epiboly autonomously. Because the EVL and the YSL are tightly attached (Betchaku and Trinkaus, 1986), EVL epiboly is believed to depend on the YSL expansion. In addition, endocytosis in the YSL near the blastoderm margin results in removal of the yolk cytoplasmic membrane and could play a role in epiboly by drawing the blastoderm to the vegetal pole (Trinkaus, 1993; Solnica-Krezel and Driever, 1994).

The cytoskeleton plays many important roles during epiboly. Extensive microtubule networks in the yolk cell may facilitate the epibolic movements, as microtubule disruption completely inhibits the movement of yolk syncytial nuclei (YSN) and impairs the epibolic movements of the deep cells and the EVL (Strahle and Jesuthasan, 1993; Solnica-Krezel and Driever, 1994). A decrease in the amount of polymerized microtubules in the yolk cell also leads to epiboly delay (Hsu et al., 2006). Actin microfilaments throughout the embryo contribute to epiboly as well (Zalik et al., 1999; Cheng et al., 2004; Koppen et al., 2006). Three distinct actin structures are elaborated during late epiboly stages: two rings at the margin of the deep cells and the EVL, and a punctate band of actin accumulation in the external YSL adjacent to the EVL margin (Cheng et al., 2004). It has been proposed that the actin rings act as a “purse string” to pull the EVL vegetally, thereby advancing the epiboly process (Cheng et al., 2004), whereas the punctate band of contractile elements including actin and myosin 2 in the YSL contributes to the shortening of the actin bands and the EVL margin, moving the EVL toward the vegetal pole (Koppen et al., 2006). Disruption of these actin structures, as a consequence of either cytochalasin B treatment (Cheng et al., 2004), interference with myosin2 (Koppen et al., 2006), or the homeobox transcription factor Mtx2 (Wilkins et al., 2008), results in a delay or failure in epiboly. Similarly, an abnormal cytoskeleton contributes to the epibolic delay in Pou5f1-deficient embryos (Lachnit et al., 2008).

We previously demonstrated that Go12/13 are required for C&E gastrulation movements in zebrafish (Lin et al., 2005). Here, we find that Go12/13 regulate epiboly in zebrafish, and provide evidence that Go12/13 interact with E-cadherin to negatively modulate E-cadherin–mediated cell–cell adhesion. Moreover, we show that Go12/13 also regulate epiboly by promoting actin microfilament assembly through a Rho guanine nucleotide exchange factor (GEF)-dependent signaling pathway. Our results therefore identify a novel Go12/13-dependent mechanism for modulating epiboly during vertebrate gastrulation.

Results

Disruption of Go12/13 function results in epiboly delay

We have previously identified one Go12 and two Go13 (Go13a and Go13b isoforms) in zebrafish and demonstrated that proper Go12/13 signaling is essential for C&E movements, as well as for epiboly during zebrafish gastrulation (Lin et al., 2005). To define further the mechanisms whereby Go12 and Go13 regulate epiboly, we used gain- and loss-of-function approaches. To enhance Go13 function, we injected embryos with a synthetic RNA encoding Go13a. To inhibit Go12/13 function, we injected embryos with a mixture of antisense MOs (3MO) that interfere with translation of the three Go12/13 transcripts (gna13a, gna13b, and gna12; 4 ng each). Alternatively, we injected a synthetic RNA encoding the carboxy-terminal (CT) peptides of Go12/13, which have been shown to disrupt the coupling of Go12/13 to their cognate receptors (Akhter et al., 1998; Gilchrist et al., 1999; Arai et al., 2003; Lin et al., 2005). In zebrafish, epiboly initiates at the sphere stage and is complete when the blastoderm encloses the yolk cell (Warga and Kimmel, 1990). Embryos with either an excess or deficiency of Go12/13 expression initiated epiboly and progressed through early stages at rates comparable to those of their uninjected siblings. Furthermore, they underwent internalization normally and formed embryonic shields of normal morphology (unpublished data). However, when the blastoderm covered 70% of the yolk cell in control embryos (70% E), embryos with altered Go12/13 activity lagged in epibolic movements behind their siblings by 10–20%. By 80% E, only a very small fraction of uninjected embryos (1.1 ± 1.9%; 136 embryos) showed epiboly defects, yet a majority of embryos overexpressing Go13a (98.5 ± 2.6%, 258 embryos), injected with the Go13-CT RNA (86.5 ± 6.5%, 106 embryos) or injected with 3MO (84.4 ± 5.7%, 143 embryos), exhibited epiboly defects (Fig. 1 M).

The YSL consists of an internal YSL and an external YSL that is populated with YSN (Solnica-Krezel and Driever, 1994). The EVL is tightly linked to the YSL margin. Therefore, as
Figure 1. Gα12/13 regulate epibolic movements of the deep cells. (A and B) Nomarski images of control WT embryos (A) and embryos overexpressing Gα13a [B] at 80% epiboly (A′ and B′ are schematic drawings of A and B), showing the dcm and YSL nuclei (YSLn; green arrows and dots), which move together in control embryos (A and A′) but are separated in embryos overexpressing Gα13a (B and B′). (C and D) Nomarski images of yolk cell region at high magnification in a control WT embryo (C) and an embryo overexpressing Gα13a (D), showing distortions in the YCL (white arrowheads). (A–D) Lateral view, with dorsal shown toward the right and vegetal toward the bottom. (E–H) Nomarski images of control WT embryos (E), embryos overexpressing either full-length Gα13a (F), or the CT fragment of Gα13a (Gα13a-CT; G), and embryos injected with 3MOs against gna13a, gna13b, and gna12 (4 ng each; H) at 95% epiboly. (E′–H′) Schematic drawings of E–H. Vegetal view is shown. df, df cells (red arrowheads). Note: in F–H versus E, the vegetal opening is much larger, and dfs are separated from the dcm; in F and H, the dfs are split. (I–L) Expression of the ntl mRNA at 90% epiboly. Images show ntl expression domains at dcm and df. Dorsal view, with the vegetal pole (VP; blue lines) toward the bottom. Yellow lines with double arrows, distance from dcm to VP. Bars, 100 μm. (M) The percentage of embryos with epibolic defects. Data are compiled from two to three different experiments. Error bars represent mean ± SEM.

* p < 0.01, vs WT
† p < 0.01, vs Gα13a
# p > 0.5, vs WT
seen in Fig. 1 (A and A’), during the course of normal epiboly, the YSN and the deep cell margin (dcm) stay together (the EVL is invisible, as it is not in the focal plane; Trinkaus, 1984; Solnica-Krezel and Driever, 1994). However, in embryos over-expressing Gα13Δa, a sizable gap was formed between the YSN and the dcm (Fig. 1, B and B’), which indicates that epibolic movement of the deep cells lags behind the movement of the YSL. Thus, the distance between the dcm and the vegetal pole is significantly greater in embryos overexpressing Gα13Δa than in the uninjected control (Fig. 1, A and B). In addition, we noted that in contrast to the uniform and smooth appearance of the yolk cytoplasmic layer (YCL; a thin anuclear cytoplasmic layer covering the yolk mass) in control embryos (Fig. 1 C), this structure was frequently distorted in Gα13Δa-expressing embryos, exhibiting an uneven thickness (Fig. 1 D; arrowheads). As epiboly progressed to 95% E, the dcm of wild-type (WT) embryos moved closer to the vegetal pole (Fig. 1, E and E’), but those of embryos overexpressing Gα13Δa, the dominant-negative Gα13-CT peptide, or injected with 3MO had a much larger vegetal opening (Fig. 1, E–H’). Moreover, the dorsal forerunners (dfs), a small dorsal cell population that normally moves toward the vegetal pole as a single cluster in close association with the dcm (Fig. 1 E; Cooper and D’Amico, 1996), were well separated from the dcm and far ahead of the remaining deep cells in embryos with reduced or excess Gα12/13 function. Interestingly, in these embryos, the df cells split and formed several smaller clusters (Fig. 1, F–H). These observations in live embryos were confirmed by analyzing the expression of the no tail (ntl) gene, which marks the mesodermal precursors at the dcm and the df cells (Schulte-Merker et al., 1994). As seen in Fig. 1 (I–L), the distance between the dcm and the vegetal pole was significantly greater in embryos with either reduced or excess Gα12/13 function than that in control embryos, and dfs were separated from the dcm and divided into several smaller clusters. The observed delay in epiboly and abnormal behavior of the dfs resemble aspects of the phenotypes that have been described for the half-baked (hab) mutants, which harbor mutations in the cadherin1 (cdh1; E-cadherin) gene (Kane et al., 1996; Kane and Warga, 2004), and in embryos injected with an MO that targets cdh1 (Babb and Marrs, 2004). This observation suggested a possible link between Gα12/13 function and E-cadherin activity.

In embryos overexpressing Gα13Δa, but not those injected with Gα13-CT RNA or 3MO (not depicted), cells frequently dissociated from the embryonic surface (Fig. 2, A–C’), and gaps formed between the paraxial and axial mesoderm during segmentation (Fig. 2 E). These phenotypic changes have also been observed in hab mutant embryos and have been attributed to defects in cell–cell adhesion (Kane et al., 2005; McFarland et al., 2005). Together, these observations suggest that Gα12/13 signaling may negatively regulate E-cadherin–mediated cell–cell adhesion during zebrafish gastrulation.

**Gα12/13 do not influence E-cadherin expression or intracellular distribution**

To test the hypothesis that Gα12/13 regulate epiboly by modulating the function of E-cadherin, we first determined if Gα12/13 affect the expression of E-cadherin in embryos with reduced or excess signaling. We performed Western blot analyses using an anti-E-cadherin antibody with protein extracts prepared either from gastrulae injected with gna13a RNA or 3MO, or from uninjected control siblings. As shown in Fig. 3 A, two prominent bands of E-cadherin, which may correspond to two glycosylation forms of E-cadherin, were detected, as described previously (Babb and Marrs, 2004). There was no clear difference in the expression level of E-cadherin protein between the control embryos and embryos with excess or reduced Gα12/13 signaling (Fig. 3 A). We then performed whole-mount immunostaining to determine if Gα12/13 regulate the cellular distribution of E-cadherin. It has been shown that E-cadherin is expressed at a higher level at the anterior region of the hypoblast during gastrulation (Babb and Marrs, 2004). To identify this region, we used embryos obtained from transgenic TG[gtsc-GFP] fish, in which GFP is expressed in the dorsal midline (Doitsidou et al., 2002; Inbal et al., 2006). As shown in Fig. 3 B, in control embryos at 70% E, E-cadherin was expressed in all blastomeres, predominantly on the cell membranes, but also in a punctate pattern in the cytosol, as described previously (Babb and Marrs, 2004; Montero et al., 2005). Our analyses revealed that neither Gα13Δa overexpression nor Gα12/13 down-regulation (3MO-mediated) affected the expression level or the cellular distribution of E-cadherin (Fig. 3 B).
**Gα_{12/13} regulate epiboly by inhibiting E-cadherin activity**

Next, we aimed to determine whether Gα_{12/13} modulate E-cadherin function in vivo by testing their genetic interactions. We took advantage of a zebrafish mutant, hab^{vu44}, harboring a premature stop codon at amino acid residue L553 within the EC4 domain of the extracellular portion of the cdh1 gene (Kane et al., 2005). hab^{vu44} embryos display an epiboly delay/arrest after midgastrulation, probably due to a moderating effect of the maternal contribution of E-cadherin, which has been shown to cooperate with the zygotically expressed E-cadherin to regulate epiboly (Shimizu et al., 2005). We injected embryos derived from crosses among hab^{vu44} heterozygous fish with either a small dose of synthetic RNA encoding Gα_{13a} (10 pg) or a single MO against Gα_{13a} (4 ng) to elevate or reduce the function of Gα_{13}, or Gα_{12}, respectively. Such treatments alone had no effect on the epiboly in WT embryos (unpublished data). We then assessed whether this manipulation of Gα_{12/13} function can modulate the phenotypic changes caused by E-cadherin deficiency by analyzing the ntl expression profile. We reasoned that if Gα_{12/13} negatively regulate the E-cadherin activity, then excess Gα_{12/13} function exacerbates it, and decreased Gα_{12/13} signaling should suppress the phenotypic changes caused by E-cadherin deficiency. Among the uninjected progeny from hab^{vu44} parents, 63 ± 11% embryos showed a normal pattern of ntl expression (Fig. 4 A); 16 ± 9% exhibited mild defects in epiboly, in which their df cells were divided into smaller clusters in spite of being tightly associated with the margin (type I defect; Fig. 4 B); and 20 ± 3.3% showed a strong epiboly delay in the deep cells and obvious separation of the df cells from the dcm (type II defect; Fig. 4 C). This phenotypic distribution is consistent with a partial penetrance of both the dominant df defect and the recessive epiboly phenotype of hab^{vu44} mutation (Kane et al., 2005). A reduction in the expression of either Gα_{12} or Gα_{13} in the progeny of hab^{vu44} heterozygotes partially suppressed the mutant epibolic defects, as indicated by a significant increase in the proportion of embryos showing normal ntl expression in the blastoderm margin and df cells, and a decrease in the percentage of embryos with severe epibolic defects (type II; Fig. 4 E). Conversely, a slight increase in Gα_{13} activity exacerbated these defects (Fig. 4, D–E). These results support the notion that Gα_{12/13} regulate epiboly through E-cadherin by acting as negative regulators of E-cadherin activity.

![Figure 3. Altered Gα_{12/13} expression does not change the levels and distribution of E-cadherin and β-catenin.](image-url)
**Gα_{12/13} interact with E-cadherin and inhibit cell adhesion**

To better understand the mechanisms by which Gα_{12/13} regulate E-cadherin activity, we set out to test these two proteins for physical interactions in vivo. Because previous studies in cultured cells had shown that mammalian Gα_{12} or Gα_{13} can bind the cytoplasmic domain of E-cadherin (Kaplan et al., 2001; Meigs et al., 2001), we performed the following procedures. First, we cotransfected HEK 293 cells with zebrafish Gα_{13a} and a GST-tagged construct encoding the E-cadherin cytoplasmic terminus (E-cad–CyT) or GST only, and performed a GST pull-down assay. As shown in Fig. 5 A, Gα_{13a} was pulled down by GST–E-cad–CyT but not by GST alone, which suggests a specific association between zebrafish Gα_{13a} and the E-cad–CyT. In addition, we demonstrated that β-catenin can compete with Gα_{13a} for the binding to E-cadherin in a dose-dependent manner (Fig. 5 B), confirming that the β-catenin– and Gα_{13a}-binding sites on E-cadherin are close to one another (Kaplan et al., 2001). However, we did not observe any obvious change in the expression level and intracellular distribution of β-catenin in embryos with altered Gα_{12/13} expression (Figs. 3 C and S1).

E-cadherin is known to regulate cell–cell adhesion in zebrafish (Meigs et al., 2002; Montero et al., 2005) and many other animals (Halbleib and Nelson, 2006). Moreover, the binding of Gα_{13} to E-cadherin interferes with its cell adhesive function in mammalian cultured cells (Meigs et al., 2002). To determine if Gα_{12/13} can influence cell adhesion in zebrafish, we performed a cell tracing experiment in embryos (Warga and Kane, 2003). In this assay, zygotes were first injected with gna13a RNA to enhance Gα_{13} function. At the 256-cell stage, a single cell at the animal pole of an uninjected or gna13a-RNA–injected blastula was then injected with fluorescein dextran, then the distribution of the progeny of the labeled cells at several time points up to 50% E was analyzed. During embryonic development, blastomeres at the animal pole become separated from each other by intercalating radially from deeper layers to the more superficial layers without significant directional migration (Warga and Kimmel, 1990). This phenomenon is thought to be mediated by E-cadherin–dependent cell–cell adhesion interactions, because in hab mutants (E-cadherin deficient), cells intercalate from the deeper to the more superficial layers but fail to maintain this position and often fall back into the deeper layer (Warga and Kane, 2003; Kane et al., 2005). We found that progeny of the labeled cells gradually dispersed over time in control embryos (Fig. 5 C) and in embryos overexpressing Gα_{13a} (Fig. 5 D). To quantify the scattering, we marked the outside edge of the regions containing the labeled cells, and calculated the areas. We then determined a scattering factor by comparing the areas at different time points to the initial area for each embryo. 1 h after injection, the scattering factor for the Gα_{13a}-expressing embryos was similar to that of control embryos. However, by the second and third hour, the ratio in embryos overexpressing Gα_{13a} was significantly greater than that in control embryos (Fig. 5, C–E). These results indicate that overexpression of Gα_{13a} enhanced dispersion in the blastoderm during epiboly, which suggests that Gα_{13a}-expressing cells have a reduced tendency to adhere to one another. These findings provide further support for the notion that signaling via Gα_{13} negatively regulates E-cadherin activity.

**Gα_{12/13} regulate actin cytoskeleton assembly during epiboly via a RhoGEF/Rho-dependent pathway**

Although embryos with enhanced or decreased Gα_{12/13} showed similar epibolic defects in deep cells as E-cadherin mutant embryos, we noted that Gα_{13a}-overexpressing embryos exhibited additional defects such as a distorted YCL (Fig. 1 D), which suggests that Gα_{13a} signaling may contribute to the regulation of epiboly via additional mechanisms that are independent of
E-cadherin. Such distortions of the YCL have also been observed in embryos with cytoskeleton abnormalities; e.g., in embryos treated with taxol to stabilize microtubules (Solnica-Krezel and Driever, 1994) or in Pou5fl mutants (Lachnit et al., 2008). We have previously shown that, like their mammalian counterparts, zebrafish Gα13 can promote actin stress fiber formation in cultured cells (Lin et al., 2005). Based on these observations, we tested whether Gα13 can also regulate cytoskeletal function in zebrafish gastrulae.

To assess the organization of actin cytoskeleton during gastrulation, we visualized actin by whole-mount immunostaining with phalloidin. As shown in Fig. 6 (A–D), the confocal images revealed the periphery of the superficial EVL cells and the deep cells beneath, as well as two actin rings at the margins of the deep cells and the EVL (Fig. 6, A–D, red and green arrowheads, respectively), as reported previously (Cheng et al., 2004). In WT embryos, the actin rings adjacent to the deep cells and the EVL are closely associated (Fig. 6 A), which indicates that EVL and the deep cells move together toward the vegetal pole during epiboly. Consistent with previous papers on studies performed in hab^vu44/vu44 mutant embryos, the deep cells exhibited impaired epiboly and lagged behind the EVL margin (Fig. 6 D); whereas the EVL underwent epiboly at a relatively normal rate, as revealed by the observation that the distance between the EVL margin and the vegetal pole (Fig. 6, yellow lines with arrows) in the mutant was comparable to that in WT embryos (Fig. 6, A and D; Kane et al., 2005; Koppen et al., 2006). As expected, embryos with reduced or excess Gα13 function displayed similar epibolic defects of the deep cells (separation from EVL margin), although the defects were more minor than those in hab^vu44/vu44 mutant embryos (Fig. 6, B–D). However, embryos with altered Gα13 function exhibited an epibolic delay of the EVL, as the distance between the EVL margin and vegetal pole (Fig. 6, yellow lines with arrows) was significantly increased relative to that in the age-matched uninjected WT embryos (Fig. 6, A–C).

![Diagram](https://via.placeholder.com/150)

**Figure 5.** Gα13 interacts with E-cadherin and inhibits cell adhesion. (A) Gα13 interacts with the cytoplasmic domain of E-cadherin. The GST pull-down assay was performed on cell extracts from HEK 293 cells cotransfected with Gα13 and either GST or a GST-tagged cytoplasmic domain of E-cadherin (GST–E-cad–CyT). The precipitates were immunoblotted with anti-Gα13 and anti-GST antibodies. The level of Gα13 expression in the lysates is shown at the bottom of the panel. (B) β-catenin competes with E-cadherin for binding to Gα13 in a dose-dependent manner. HEK 293 cells were transfected with Gα13 and GST–E-cad–CyT with or without β-catenin at various doses, and the GST pull-down assay was performed. The expression levels of β-catenin and Gα13 in the lysates are shown. (C and D) Overexpression of Gα13 enhances cell scattering in the blastoderm. Shown are representative images of labeled cells in the blastoderm of control WT embryos and embryos overexpressing Gα13 scattering over time. The area of cell scattering is indicated by the yellow broken lines, which mark the cells at the outer edge. Bars, 100 μm. (E) Quantitative data from four separate experiments (eight embryos in each group), showing the ratio of the area of cell scattering relative to the starting point, at different time points. Error bars represent mean ± SEM. * P < 0.05 versus control.
During epiboly, the constriction of the marginal EVL cells leads to dramatic cell-shape changes in the EVL cells, and to the elongation of the EVL cells along the animal–vegetal axis. Failure of such cell-shape changes has been implicated in epibolic defects (Koppen et al., 2006). To further evaluate the morphology of the EVL cells in embryos with altered Gα12/13 function, we took confocal images of phalloidin-stained embryos at higher magnification, and analyzed cell shape (length-to-width ratio [LWR]) and orientation (the angle of the long axis of the EVL cells relative to a line parallel to the EVL margin) of the EVL cells near the margin. As shown in Fig. 6 (E–H), there was no significant difference in the intensity of F-actin staining in the EVL cells between the uninjected WT embryos and embryos injected with 3MO or the Gα13a RNA. However, both the shape and orientation of the EVL cells in embryos with reduced or excess Gα12/13 function were significantly altered with respect to those in the control embryos (Fig. 6 E–G). In the uninjected control embryos, the EVL cells were elongated, with a mean LWR of 1.73 ± 0.3 (220 cells, 6 embryos), and were orientated at an angle of 67 ± 20°. Of 220 cells counted, 73% aligned their cell bodies at an angle in the range of 60–120° with respect to the EVL margin; this indicates that most of these cells elongate vegetally along a line roughly perpendicular to the EVL margin, which is consistent with the direction of the epibolic movement. Similarly, the EVL cells in habvu44/vu44 mutant embryos were elongated and aligned properly (LWR = 1.79 ± 0.54; angle = 64 ± 16°; 171 cells, 3 embryos; P > 0.05 vs. WT). In contrast, in embryos with reduced or excess Gα12/13 function, the EVL cells were significant rounder (smaller LWR; 3MO: LWR = 1.46 ± 0.32, 379 cells, 8 embryos; Gα13a: LWR = 1.55 ± 0.44, 342 cells, 8 embryos; P < 0.001 vs. control; Fig. 6, E–H). In addition, these EVL cells were more disorganized and failed to align their cell bodies along the direction of epibolic movement, with orientations of 52 ± 26° or 54 ± 25° (P < 0.001 vs. control) in Gα12/13-depleted or Gα13a-overexpressing embryos, respectively. Moreover, only 41–49% of the cells from these embryos exhibited an angle within the range of 60–120°, which suggests that most EVL cells in these embryos were oriented in random directions (Fig. 6, E–G and I–K). Interestingly, the punctate actin accumulation adjacent to the EVL margin was markedly reduced in embryos overexpressing Gα13a (Fig. 6 G; compare the yellow arrows Fig. 6, E and G). Moreover, these embryos exhibited abnormal formation of actin bundles in the YCL, although these are rarely observed in the yolk cells of WT embryos. This is possibly due to the aggregation or contraction of F-actin, which was absent in some areas of the cortical cytoplasmic layer (Fig. 6, C and G).

Work in mammalian systems has established that Gα12/13 regulate actin cytoskeleton dynamics to modulate cell shape.
and migration via a RhoGEF/Rho-dependent signaling pathway (Buhl et al., 1995; Gohla et al., 1998; Hart et al., 1998; Kozasa et al., 1998). We have shown previously that, like zebrafish Gα12/13, one of the zebrafish RhoGEFs, PDZRhoGEF (Arhgef11), can induce stress fiber formation in HEK 293 cells (Lin et al., 2005; Panizzi et al., 2007), which suggests that zebrafish Gα12/13 also function through RhoGEF to regulate actin organization. Furthermore, we showed that, when coexpressed in HEK 293 cells, Gα13a specifically coprecipitated with myc-tagged full-length Arhgef11, and this interaction was not observed when an Arhgef11 mutant lacking the RGS domain, known to be required for target binding, was coexpressed. This indicates that Gα12/13 physically interact with PDZRhoGEF via the RGS domain (Fig. 7 J).

To determine if zebrafish Gα12/13 modulate epiboly via a RhoGEF-dependent signaling pathway, we first examined the effect of Arhgef11 overexpression on epiboly. The overexpression of Arhgef11 resulted in similar epiboly defects and distortions in the YCL, similar to those observed for Gα12/13 overexpression (Fig. 7 D and not depicted). Actin staining revealed that embryos overexpressing Arhgef11 also exhibited delayed epiboly of the deep cells and the EVL as well as the formation of thick actin bundles in the YCL (Fig. 7 H). Similar defects were observed in embryos overexpressing a constitutively activated zebrafish RhoA (data not shown). To test whether RhoGEF acts downstream of Gα12/13 in the regulation of the actin cytoskeleton, we coexpressed Gα13a together with a dominant-negative form of Arhgef11 lacking the DH and PH

Figure 7. Gα<sub>13a</sub> promotes actin assembly via a PDZ RhoGEF-dependent pathway. (A–D) Nomarski images of live WT embryos (A), embryos overexpressing Gα<sub>13a</sub> alone (B), embryos overexpressing Gα<sub>13a</sub> and a dominant-negative mutant zebrafish Arhgef11, ΔDHPH (C), or embryos overexpressing Arhgef11 (D) at 80% epiboly. Bar, 250 μm. (E–H) Confocal z-projection images show phalloidin staining of F-actin. Red and green arrowheads indicate the dcm and the EVL, respectively; pink asterisks show the actin bundles in the yolk. Note the gap between dcm and the EVL, respectively; pink asterisks show the actin bundles in the yolk. Note the gap between dcm and the EVL, respectively; pink asterisks show the actin bundles in the yolk.

- Pellet
- IP: anti-Myc
- Lysate

Embryos with actin bundles (%)

<table>
<thead>
<tr>
<th>Gα&lt;sub&gt;13a&lt;/sub&gt;</th>
<th>Gα&lt;sub&gt;13a&lt;/sub&gt;+ΔDHPH</th>
</tr>
</thead>
<tbody>
<tr>
<td>+</td>
<td>+</td>
</tr>
<tr>
<td>+</td>
<td>+</td>
</tr>
<tr>
<td>−</td>
<td>−</td>
</tr>
<tr>
<td>−</td>
<td>−</td>
</tr>
</tbody>
</table>

*P < 0.05 versus Gα<sub>13a</sub>. (J) Gα<sub>13a</sub> interacts with zebrafish Arhgef11. Coimmunoprecipitation was performed on cell extracts from HEK 293 cells transfected with Gα<sub>13a</sub> or Arhgef11 alone, or with both Gα<sub>13a</sub> and myc-tagged Arhgef11 forms (WT, dominant-negative mutants lacking the RGS domain [ΔRGS], or lacking the DH and PH domains [ΔDHPH]). Immunoblotting was performed with the indicated antibodies. Error bars represent mean ± SEM.
domains (∆DHPH), which are needed for interacting with downstream proteins (Panizzi et al., 2007). We found that Arhgef11 ∆DHPH bound to Go13 (Fig. 7 J) and suppressed both the formation of actin bundles in the YCL and the epible defects associated with Go13a overexpression (Figs. 1 M and 7). Although actin bundles were found in 86 ± 5% of the embryos overexpressing Go13a, only 33 ± 3% of the embryos coexpressing Go12/13 and Arhgef11 ∆DHPH showed this phenotype (Fig. 7 I). However, we observed that coexpression of Arhgef11 ∆DHPH did not fully rescue the epible delay in the deep cells (Fig. 7 G), which suggests that cytoskeletal assembly regulated by Arhgef11 only partially accounts for the function of Go12/13 in epible. Collectively, these results indicate that Go12/13 can regulate epible through a PDZ RhoGEF/RhoA-dependent signaling pathway to modulate the function of the actin cytoskeleton.

**Discussion**

In this paper, we demonstrate that Go12/13 signaling can regulate different aspects of epible movements by two distinct mechanisms: inhibiting E-cadherin activity and modulating actin cytoskeleton organization.

Excess or reduced Go12/13 signaling during gastrulation resulted in delayed epible of the deep cells and in the splitting of the df cell cluster (Fig. 1). Moreover, excess Go12/13 activity led to the detachment of cells from embryonic tissues, which suggests that cell adhesion is defective under these circumstances (Fig. 2). All of these phenotypic characteristics resemble those observed in hab (cdh1) mutant embryos (Kane et al., 1996; Kane and Warga, 2004), which suggests a possible link between Go12/13 signaling and E-cadherin. Indeed, although altered Go12/13 expression did not change the expression level and cellular distribution of E-cadherin (Fig. 3), our in vivo genetic experiments demonstrated that Go12/13 can inhibit the function of E-cadherin. In particular, we found that a reduction in the expression of either Go12 or Go13 function by MO injection partially suppressed, whereas an increase in Go13 activity exacerbated the epible defects in hab mutant mutants (Fig. 4). Interestingly, decreased Go12/13 function reduced the fraction of embryos with a weak epible defect, as well as the fraction with a strong epible delay (Fig. 4). This suggests that reduced Go12/13 function may suppress not only the epible defects in heterozygous embryos, but also those in homozygous mutants. We speculate that such an effect might be caused by the reduced inhibition of the maternal E-cadherin protein by Go12/13 in homozygous mutants (Babb and Marrs, 2004; Kane et al., 2005).

Our biochemical studies support the notion that Go12/13 can interact with E-cadherin. We showed that zebrafish Go13a was pulled down with the CT fragment of zebrafish E-cadherin (E-cad–CyT) in HEK cells (Fig. 4), which is consistent with the physical interaction between mammalian Go12/13 and E-cadherin shown previously (Kaplan et al., 2001; Meigs et al., 2001). The E-cad–CyT has been shown to act as a dominant-negative protein (Sadot et al., 1998). Accordingly, we observed that embryos expressing this fragment exhibited cleavage defects, and the detachment of blastodermal cells during early development and coinjection of Go13a-RNA exacerbated these phenotypes (unpublished data). The region of E-cadherin that binds Go12/13 was found to be located near the binding site for β-catenin (Kaplan et al., 2001). This is supported by our result showing that β-catenin can compete with Go13 for E-cadherin binding in HEK cells (Fig. 5 B). Furthermore, it has been hypothesized that the binding of Go12/13 to E-cadherin interferes with the ability of E-cadherin to form a complex with β-catenin. In fact, Go12/13 overexpression can cause β-catenin to dissociate from E-cadherin and to translocate from the membrane to the cytosol in cultured cells (Meigs et al., 2001). We speculate that the competition of Go12/13 with β-catenin for binding to E-cadherin may be one of the underlying mechanisms in embryos. However, we did not observe any overt change in expression level or intracellular distribution of β-catenin in embryos overexpressing Go13a (Fig. 3). The reason for this discrepancy between studies in cell culture and our studies in zebrafish embryos is unclear. However, one possibility is that the levels of Go12/13 we used were sufficient to alter E-cadherin activity but not to produce detectable changes in β-catenin distribution.

Studies from hab mutant embryos indicate that E-cadherin regulates epible in part by impinging on cell–cell adhesion (Warga and Kimmel, 1990; Montero et al., 2005; Shimizu et al., 2005). Considering these findings in the light of our biochemical and genetic data showing that Go12/13 functionally interact with E-cadherin in zebrafish, we propose that Go12/13 modulate epible movement by inhibiting E-cadherin–mediated cell–cell adhesion. Accordingly, cells in embryos overexpressing Go13a during early epible scattered across a larger area (Fig. 5). In addition, we also demonstrated that a larger scattering area in embryos overexpressing Go13a is not due to an increase in cell number (Fig. S2). However, we cannot rule out the possibility that other functions of Go12/13 could contribute to reduced cohesion or abnormal cell movements.

In addition to the impaired epible of the deep cells, altered Go12/13 signaling resulted in epible defects of the EVL (Fig. 6). This is in contrast to hab−/− and maternal-zygotic cdh1−/− mutant embryos, in which the EVL appears to undergo normal epible in spite of the fact that the deep cells exhibit severe epible defects (Fig. 6 D; Shimizu et al., 2005). These results suggest that Go12/13 may impinge on pathways other than the E-cadherin pathway to regulate epible in EVL cells. Recent evidence indicates that proper organization of the F-actin–based cytoskeleton plays critical roles in the normal epible of zebrafish embryos (Zalik et al., 1999; Cheng et al., 2004; Koppen et al., 2006). The actin contractile elements in the YSL are necessary for facilitating the proper EVL cell shape changes during late gastrulation (Koppen et al., 2006). Notably, in E-cadherin–deficient embryos, actin organization appeared to be normal, and EVL cells were elongated and oriented properly (Fig. 6, D and H; Shimizu et al., 2005), which indicates that E-cadherin does not play a significant role in actin organization and EVL epible in zebrafish.

In mammalian cultured cells, Go12/13 are known to be involved in the regulation of actin polymerization and the maintenance of proper cell morphology, which suggests that Go12/13 may affect EVL epible by regulating actin cytoskeleton organization.
and/or function. Although there is no significant change in the organization of filamentous actin of EVL cells in embryos with altered Gα12/13 signaling, these cells displayed defects in cell shape and orientation (Fig. 6), which may contribute to the epibolic defects of the EVL. In addition, in embryos with excess Gα12/13 signaling, the punctate F-actin ring adjacent to the EVL was significantly reduced (Fig. 6 G), and abnormal thick actin bundles, separated by F-actin–free regions, were frequently found in the yolk (Fig. 6 C and G). We speculate that the thick actin bundles may cause abnormal “contractile” forces that disrupt the YCL; alternatively, these forces may create resistance to vegetal pulling of the EVL. Altogether, the changes in actin architecture resulting from altered Gα12/13 activities could prevent the EVL from undergoing active cell rearrangement and shape changes, ultimately affecting normal EVL epiboly. In E-cadherin mutants, in contrast, the abnormal actin fibers are not observed, and thus they are unlikely to be caused by a decrease in E-cadherin function. Interestingly, although the deep cells in embryos with Gα12/13 mutants, in contrast, the abnormal actin fibers are not observed, and thus they are unlikely to be caused by a decrease in E-cadherin function. Interestingly, although the deep cells in embryos with altered Gα12/13 signaling displayed severe defects in epiboly, they did not exhibit corresponding changes in F-actin organization and cell shape (unpublished data). This further underscores the notion that epiboly of the deep cells might involve distinct mechanisms. However, we cannot exclude the possibility that in the deep cells, Gα12/13 also influence epiboly by modulating the actin cytoskeleton. Indeed, in differentiated leukocyte-HEK60, Gα12/13 were shown to influence the actomyosin network during retraction of the trailing edge (Xu et al., 2003). It will be interesting in the future to investigate how cell migration contributes to epiboly in zebrafish. Furthermore, in Xenopus laevis, it has been shown that two G protein–coupled receptors (GPCRs; the phospholipid lysophosphatidic acid [LPA] receptor and Xfl p) that couple to Gα12/13 in some cell types (Ishii et al., 2004) can regulate expression of a calcium-dependent EP-cadherin and modulate the assembly of cortical actin (Lloyd et al., 2005; Tao et al., 2005, 2007). Therefore, it will be important in the future to investigate if LPA functions in a similar manner in zebrafish.

In addition, we found that embryos overexpressing Gα13 exhibited microtubule organization defects similar to those observed for F-actin, showing thick bundles of microtubules surrounded by areas devoid of microtubules (Fig. S3). However, embryos with reduced Gα13 do not show significant defects in microtubule organization (unpublished data), which suggests that at their normal expression levels, Gα12/13 do not play an essential role in microtubule stabilization in zebrafish.

Gα12/13 are known to regulate cytoskeletal function via Gα12/13 signaling pathway to regulate actin organization during EVL epiboly. First, coexpression of a dominant-negative zebrafish PDZ RhoGEF, Arhgef11, with Gα13 significantly reduced the formation of actin bundles, and suppressed the epiboly defects in the EVL (Figs. 1 M and 7). Conversely, overexpression of Arhgef11 or a constitutively active RhoA resulted in similarly abnormal actin organization in the yolk, and impaired epiboly (Fig. 7, D and H; and unpublished data). Notably, embryos expressing Arhgef11 did not exhibit the detachment of cells from the embryo surface (unpublished data), and Arhgef11 LOF did not result in obvious epiboly and C&E defects but gave rise to defects associated with ciliated epithelia (Panizzi et al., 2007). We speculate that Arhgef11 and Gα12/13 may act in both overlapping and different signaling pathways during gastrulation. These results further support the idea that Gα12/13 regulate actin organization and cell adhesion via distinct mechanisms.

In summary, our studies establish Gα12/13 as novel regulators of epiboly in zebrafish. Our data indicate that Gα12/13 may regulate different aspects of epibolic movements by two distinct pathways. In the deep cells, Gα12/13 bind the intracellular domain of E-cadherin and inhibit its activity to modulate epiboly; in the EVL and the yolk cell, Gα12/13 promote actin cytoskeleton assembly through RhoGEF/RhodA in order to regulate epibolic movement of the EVL. It has been shown that Gα12/13 can transmit signals from different GPCRs and suggested that different GPCRs may activate distinct signaling pathways through Gα12/13 (Riobo and Manning, 2005). It will be interesting to determine if zebrafish epiboly involves different extracellular signals acting through distinct GPCRs via Gα12/13 to specify distinct cell behaviors in different cell types. Gα12/13 are oncogenes with transforming potential and growth-promoting activity (Chan et al., 1993; Voyno-Yasenetskaya et al., 1994; Radhika and Dhanasekaran, 2001). Furthermore, the down-regulation of E-cadherin is associated with tumor metastasis and cancer progression (Behrens, 1999). Thus, our findings on the in vivo role of Gα12/13 in epiboly may have significant implications for the mechanisms whereby Gα12/13 function during tumorigenesis and metastasis, as well as during other morphogenetic processes in multicellular systems.

Materials and methods

Zebrafish strain and maintenance

WT, transgenic Tg[gsc:GFP] (Doitsidou et al., 2002), and habredd mutant strains of zebrafish were maintained as described previously (Solnica-Krezel et al., 1994). Embryos were obtained by natural mating and staged according to morphology as described previously (Kimmel et al., 1995). Krezel et al., 1994). Embryos were obtained by natural mating and staged according to morphology as described previously (Kimmel et al., 1995). Whole-mount in situ hybridization using an antisense RNA probe was performed as described previously (Thisse and Thisse, 1998), except that BM Purple (Roche) was used for the chromogenic reaction.

Capped sense mRNAs were synthesized using the SP6 mMessage machine (Applied Biosystems). The injection of synthetic mRNAs encoding Gα12/13 (60 pg), Gα13 (800 pg), myc-tagged PDZ RhoGEF [Arhgef11, 2pg], the dominant-negative mutant Arhgef11 [3DHPH, 600 pg], constitutively activated RhodA [10 pg], and antisense MOs targeting zebrafish gna12a, gna13a, and gna12b transcripts [4 ng each] has been described previously (Liu et al., 2005). Whole-mount in situ hybridization using an antisense mRNA probe was performed as described previously (Thiese and Thiese, 1998), except that BM Purple (Roche) was used for the chromogenic reaction.

Western blotting

Embryos at 80% epiboly stage were manually deyolked and homogenized in lysis buffer (Chen et al., 2004) to prepare embryo extracts. Equal amounts of protein were used for Western blot analysis. The following primary antibodies were used: anti-E-cadherin antibody (1:10,000; Babb and Marrs, 2004), anti-Gα12/13 antibody (1:5,000; Santa Cruz Biotechnology, Inc.), and anti-β-catenin antibody (1:250; Sigma-Aldrich).

GST pull-down and coimmunoprecipitation assays

HEK 293 cells were transiently cotransfected with cDNAs encoding Gα12/13 and GST or the GST-tagged CT fragment of E-cadherin, or zebrafish Gα12/13.
and a myc-tagged full-length Arhgef11, or myc-tagged Arhgef11 mutants lacking the RGS domain (ARGS) or DH and PH domains (ADPH; Panuzzi et al., 2007). After serum starvation overnight, cells were washed twice with serum-free medium and lysed in PBS containing 1% Igepal, 0.2% deoxycholate, and protease inhibitors. For the GST pull-down assay, protein extracts were incubated with glutathione-Sepharose beads (GE Health-care). The presence of Gβ13a in the lysates and precipitate was detected by using unpaired Student’s t tests, unequal variance.

Whole-mount immunostaining

Embryos were fixed at appropriate stages in 4% PFA/PBS/4% sucrose at 4°C overnight. For F-actin staining, Alexa Fluor 546 phalloidin (1:100; Invitrogen) was used as described previously (Koppens et al., 2006). In addition, the following primary antibodies were used: anti-E-cadherin (1:11,000; Babb and Moris, 2004), anti-α-tubulin (DM1A; 1:300; EMD), and anti-α-catenin (1:250; Sigma-Aldrich). Embryos were then mounted in 75% glycerol in PBS for analysis by microscopy.

Quantification of cell shape and alignment

Confocal images of E12.5 embryos were collected using a 20x/0.8 NA objective lens (Carl Zeiss, Inc.). LVRs [a ratio of the longest to shortest axis of the cell] and the angle of the long axis of the cells relative to a line parallel to the EVL margin were determined using ImageJ software (Schneider et al., 2012). The acquired images were exported and edited using Photoshop (Adobe), and then compiled in Illustrator software (Adobe).

Statistical analysis

Data are presented as the mean ± SEM. Statistical analyses were performed using unpaired Student’s t tests with 2 tails, unequal variance.

Online supplemental material

Fig. S1 shows that the distribution of β-catenin and α-catenin is not changed in cells expressing Gα13a. Fig. S2 shows that Gα13a overexpression does not promote cell proliferation. Fig. S3 shows the microtubules in WT control embryos and embryos overexpressing Gα13a revealed by anti-α-tubulin staining. Online supplemental material is available at http://www.jcb.org/cgi/content/full/jcb.200805148/DC1.

References


