

## RESEARCH ARTICLE

# Vangl2-dependent regulation of membrane protrusions and directed migration requires a fibronectin extracellular matrix

Anna M. Love, Dianna J. Prince and Jason R. Jessen\*

## ABSTRACT

During zebrafish gastrulation the planar cell polarity (PCP) protein Vang-like 2 (Vangl2) regulates the polarization of cells that are engaged in directed migration. However, it is unclear whether Vangl2 influences membrane-protrusive activities in migrating gastrula cells and whether these processes require the fibronectin extracellular matrix. Here, we report that Vangl2 modulates the formation and polarization of actin-rich filopodia-like and large lamellipodia-like protrusions in ectodermal cells. By contrast, disrupted Glypican4/PCP signaling affects protrusion polarity but not protrusion number or directed migration. Analysis of fluorescent fusion protein expression suggests that there is widespread Vangl2 symmetry in migrating cells, but there is enrichment at membrane domains that are developing large protrusions compared with non-protrusive domains. We show that the fibronectin extracellular matrix is essential for cell-surface Vangl2 expression, membrane-protrusive activity and directed migration. Manipulation of fibronectin protein levels rescues protrusion and directed migration phenotypes in *vangl2* mutant embryos, but it is not sufficient to restore either PCP, or convergence and extension movements. Together, our findings identify distinct roles for Vangl2 and Glypican4/PCP signaling during membrane protrusion formation and demonstrate that cell-matrix interactions underlie Vangl2-dependent regulation of protrusive activities in migrating gastrula cells.

**KEY WORDS:** Vangl2, Fibronectin, Protrusion, Polarity, Migration, Gastrulation, Zebrafish

## INTRODUCTION

Establishment of planar cell polarity (PCP) is a complex process that involves many proteins functioning cell autonomously and/or non-autonomously. First described in reference to the orientation of insect hairs and bristles (Lawrence, 1966; Nubler-Jung et al., 1987), PCP and its associated proteins are linked to a variety of morphogenetic processes, including gastrulation movements (Goodrich and Strutt, 2011; Gray et al., 2011; Jessen and Solnica-Krezel, 2005). The six core PCP genes identified in fly include *vang*, *prickle*, *dishevelled*, *frizzled*, *flamingo* and *diego* (Strutt and Strutt, 2007; Wong and Adler, 1993; Wu and Mlodzik, 2009). Genetic evidence that a core PCP protein is required for vertebrate gastrulation came with the identification of *vang-like 2* (*vangl2*), which encodes a four-pass transmembrane protein, as the defective gene in *trilobite* mutant zebrafish (*Danio rerio*) embryos (Jessen

et al., 2002; Solnica-Krezel et al., 1996). Loss of the Vangl2 protein produces a strong convergence and extension gastrulation phenotype, characterized by shortened and broadened embryonic body axes. Underlying normal convergence and extension movements are several cell behaviors, among them directed migration and mediolateral intercalation. Here, PCP is defined as the elongation and mediolateral alignment (MLA) of ectodermal and mesodermal cells perpendicular to the dorsal embryonic axis. Loss of Vangl2 causes lateral cells to meander, moving dorsally along indirect trajectories (Jessen et al., 2002). How Vangl2 regulates directed migration during zebrafish gastrulation is not understood.

A growing body of data indicates that a non-canonical Wnt pathway controls PCP during gastrulation. With the aid of the glycosylphosphatidylinositol-anchored Glypican4 co-receptor, Wnt11 and Wnt5b bind Frizzled receptors to effect changes in the subcellular distribution of Dishevelled (Heisenberg et al., 2000; Kilian et al., 2003; Ohkawara et al., 2003; Topczewski et al., 2001). Downstream, Rho family small GTPases and their effector proteins (e.g. Rho kinase) are thought to stimulate changes in actin arrangement and cell polarization (Habas et al., 2003, 2001; Marlow et al., 2002). Vangl2 and its cytoplasmic binding partner, Prickle1a, somehow influence the Glypican4/PCP pathway (Carreira-Barbosa et al., 2003; Dohn et al., 2013; Jessen et al., 2002; Veeman et al., 2003), perhaps by regulating Dishevelled signaling (Seo et al., 2017). Zebrafish with null mutations in *vangl2* and *glypican4* (*knypek*) or *vangl2* and *wnt11* (*silberblick*) have convergence and extension phenotypes stronger than the individual mutants (Heisenberg and Nüsslein-Volhard, 1997; Marlow et al., 1998). These findings indicate that Vangl2 and Glypican4 have distinct functions during gastrulation movements.

A hallmark of PCP in fly epithelium is asymmetrical expression of certain core proteins (Strutt, 2002). In the wing, Van Gogh and Prickle localize to the proximal side of the cell, whereas Frizzled, Dishevelled and Diego are enriched distally in the area of actin-rich trichome formation. Asymmetrical protein-containing membrane domains on adjacent cells interact in order to regulate the global alignment of trichome polarity in relation to the wing axes. Unlike fly proteins, asymmetrical expression of vertebrate core PCP proteins has proven challenging to demonstrate. In polarized non-migrating neuroectoderm or axial mesoderm, fluorescent fusion proteins for Vangl2 and Prickle are enriched at anterior cell boundaries, whereas Dishevelled preferentially localizes to posterior membranes (Ciruna et al., 2006; Roszko et al., 2015; Yin et al., 2008). It is unknown whether PCP proteins are asymmetrically expressed in lateral ectodermal or mesodermal cells that are undergoing directed migration. Data from non-migrating cells suggest an anterior/posterior axis of PCP. However, migrating gastrula cells exhibit dynamic changes in cell shape, orientation and position relative to neighboring cells (Jessen et al., 2002; Roszko et al., 2015; Yin et al., 2009) and a plane of polarization may be difficult to discern. Whatever the subcellular

Department of Biology, Middle Tennessee State University, Murfreesboro, TN 37132, USA.

\*Author for correspondence (jason.jessen@mtsu.edu)

 J.R.J., 0000-0002-8209-3620

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distribution of PCP proteins, understanding their function in migrating gastrula cells remains a key issue.

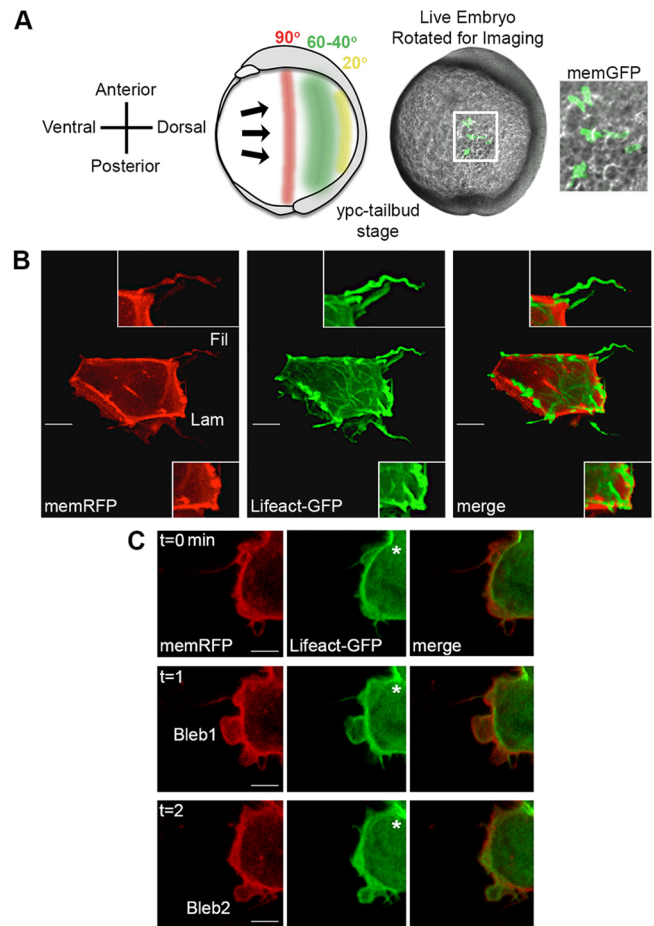
For migrating cells, adhesion to the extracellular matrix (ECM) is usually required to generate the traction needed for membrane protrusion formation and cell body translocation (Friedl and Gilmour, 2009). During migration, *vangl2* mutant cells lack directionality, which indicates a possible relationship between Vangl2, the ECM and membrane-protrusive activity. Experiments using the frog gastrula demonstrate that integrin  $\alpha 5\beta 1$  and fibronectin interactions suppress inappropriate membrane-protrusive activity (Davidson et al., 2006). Moreover, overexpression of frog Vangl2, Prickle or Frizzled7 disrupts fibronectin fibril assembly and organization, in a manner correlating with the severity of the PCP phenotype (Goto et al., 2005). Similar to frog gastrulation movements, zebrafish convergence and extension occur in the context of a fibronectin-containing ECM network (Boucaut and Darribere, 1983; Latimer and Jessen, 2010; Winklbauer and Keller, 1996). During gastrulation, a layer of fibronectin forms between the ectoderm and superficial mesoderm and another layer between the yolk and mesoderm, with individual fibrils protruding between cells (Latimer and Jessen, 2010). Therefore, by late gastrulation, ectodermal and mesodermal cell migration is associated with a fibrillar ECM. Notably, whereas loss of either Vangl2 or Prickle1a results in reduced fibronectin, *glypican4* mutant embryos exhibit increased fibronectin assembly (Dohn et al., 2013; Williams et al., 2012). These different effects on ECM structure further support the notion that Vangl2/Prickle1a and Glypican4/PCP signaling have distinct effects on cell behaviors. In addition, these data suggest that fibronectin may be necessary for certain aspects of Vangl2 function.

The major goal of this study was to determine how Vangl2 and fibronectin regulate membrane protrusion dynamics in migrating zebrafish gastrula cells. We used time-lapse imaging combined with mosaic expression of fluorescent fusion proteins to visualize protrusions in live embryos. We have shown that Vangl2 regulates distinct aspects of protrusion formation compared with Glypican4. We found GFP-VANGL2 expression to be generally symmetrical in migrating gastrula cells, but enriched in forming membrane protrusions compared with non-protrusive domains. Our work implicates fibronectin in the regulation of protrusion formation and polarization, and Vangl2 cell-surface expression. Finally, we have shown that increasing fibrillar fibronectin in *vangl2* mutant embryos rescues the protrusion phenotype, but not PCP. These results uncover a previously unrecognized interaction between Vangl2, fibronectin and membrane-protrusive activity, which are required for the dorsal convergence of gastrula cells.

## RESULTS

### Vangl2 and Glypican4 differentially regulate membrane protrusion formation and directed migration

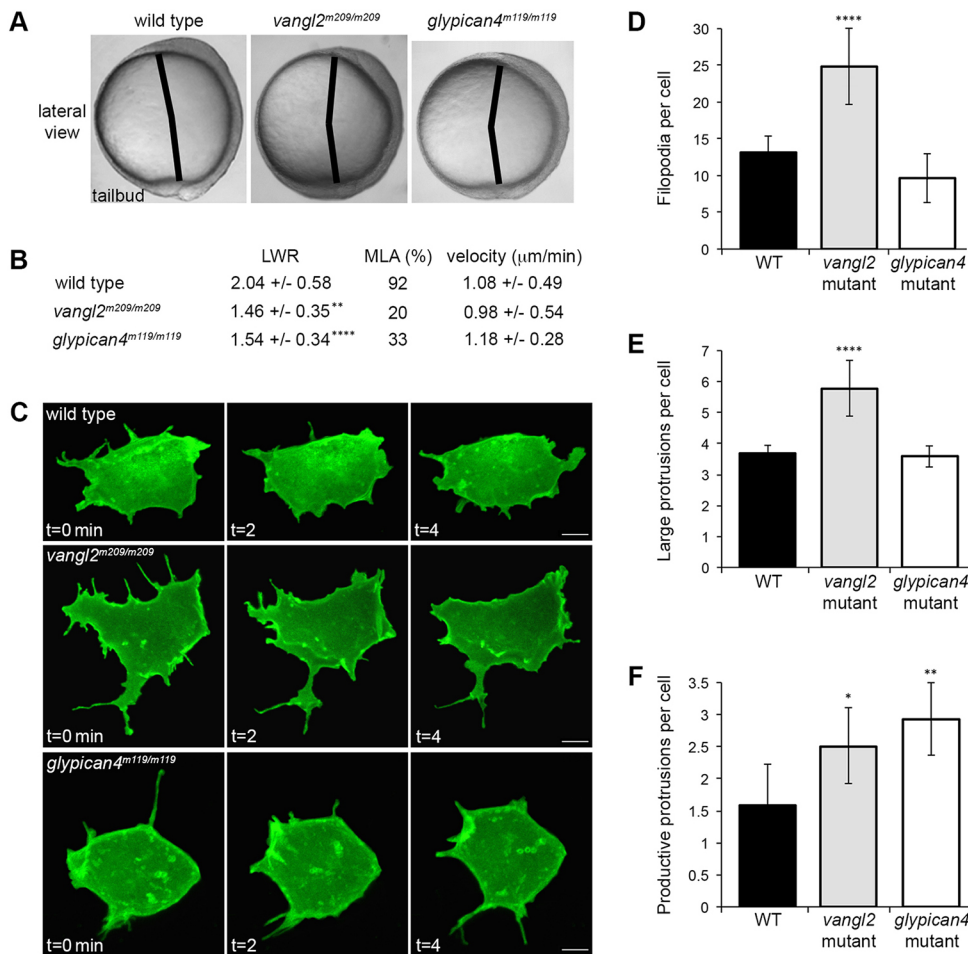
Multiple cell behaviors contribute to the processes of convergence and extension during zebrafish gastrulation, including the directed migration of lateral cells toward the dorsal body axis (Jessen and Solnica-Krezel, 2005). Whereas wild-type gastrula cells are elongated and mediolaterally aligned, this type of PCP is disrupted in *vangl2* and *glypican4* homozygous mutant embryos (Jessen et al., 2002; Topczewski et al., 2001). To better understand the mechanism whereby Vangl2 regulates dorsal convergence, we analyzed and compared membrane protrusion formation in *vangl2*<sup>m209/m209</sup> and *glypican4*<sup>m119/m119</sup> mutants. Time-lapse confocal microscopy was used to image late gastrula lateral ectodermal cells 40–60 degrees from the notochord (Fig. 1A). We achieved mosaic labeling by injecting single blastomeres from eight-cell-stage embryos with



**Fig. 1. Time-lapse imaging of membrane protrusions.** (A) Schematic (left) and live wild-type zebrafish embryo (right) highlighting the 40°–60° lateral region that was analyzed. Lateral ectodermal cells with mosaic memGFP expression from an embryo used for confocal imaging at 60° from dorsal (white boxed area, enlarged far right). ypc, yolk-plug closure. (B) Wild-type ectodermal cell expressing memRFP and Lifeact-GFP with filopodia (Fil) and large lamellipodia-like (Lam) membrane protrusions. Insets show enlarged images of protrusion classifications. (C) Wild-type ectodermal cell undergoing division. Asterisks indicate forming cleavage furrow. Two blebs are observed during the three time points. Scale bars: 5  $\mu$ m.

synthetic mRNA that encoded fluorescent fusion proteins. Lifeact-GFP and membrane-targeted RFP (memRFP) were used to assess the types of membrane protrusions that are formed by lateral ectodermal cells. Three distinct protrusions were identified: small spike-like actin-rich protrusions that resemble filopodia; large sheet-like actin-rich protrusions that resemble lamellipodia/filolamellipodia (henceforth large protrusions); and spherical bleb-like protrusions that are initially devoid of actin (Blaser et al., 2006) (Fig. 1B,C). Filopodia-like protrusions (henceforth filopodia) predominate, followed by large protrusions, and therefore were the focus of our study. Bleb-like protrusions are much less abundant and are often associated with dividing cells (Fig. 1C).

For the analysis of membrane-protrusive activity, single blastomeres were injected with only memGFP, to minimize mRNA cytotoxicity. Homozygous *vangl2* and *glypican4* mutant embryos are identified by their strong convergence and extension phenotypes at tailbud stage (Fig. 2A) (Solnica-Krezel et al., 1996). First, we confirmed that mutant lateral ectodermal cells exhibited defective PCP, as indicated by a reduced length-width ratio (LWR) and MLA (Fig. 2B). We next quantified the number of protrusions



**Fig. 2. Membrane protrusion formation in *vangl2* and *glypican4* mutant embryos.** (A) Live embryo images at late yolk-plug closure/tailbud stage. Black lines denote the polster-tailbud angle. (B) PCP and migration velocity quantitation in the ectoderm. LWR and MLA values were obtained from: wild type (WT),  $n=50$  cells, 13 embryos; *vangl2*<sup>m209/m209</sup>,  $n=10$  cells, 8 embryos; *glypican4*<sup>m119/m119</sup>,  $n=49$  cells, 5 embryos. PCP data for *vangl2* mutant ectoderm was previously published (Jessen et al., 2002), which explains the lower  $n$ . (C) Time-lapse confocal images of ectodermal cells expressing memGFP over three time points. (D-F) Quantitation of the total numbers of filopodia (D), large protrusions (E) and productive protrusions (F) formed by WT ( $n=12$  cells, 8 embryos), *vangl2*<sup>m209/m209</sup> ( $n=10$  cells, 7 embryos) and *glypican4*<sup>m119/m119</sup> ( $n=12$  cells, 5 embryos). Data are mean  $\pm$  s.d. \* $P<0.05$ , \*\* $P<0.01$ , \*\*\*\* $P<0.0001$ ;  $P$  values are versus WT; one-way ANOVA significance test followed by Tukey HSD post-hoc tests. Scale bars: 5  $\mu\text{m}$ .

that were formed by ectodermal cells. Wild-type embryos averaged 13.16 filopodia and 3.68 large protrusions per cell (Fig. 2C-E). In *vangl2* embryos, these averages increased to 24.84 filopodia and 5.77 large protrusions per cell (Fig. 2C-E). By contrast, *glypican4* mutant ectodermal cells showed no difference in the number of filopodia (9.58) or large protrusions (3.59) compared with wild type (Fig. 2C-E). From the total number of large protrusions, we determined the proportion that were productive, meaning protrusion formation in one direction is followed by cell body translocation in the same direction. We found the average number of productive protrusions formed by cells in *vangl2* and *glypican4* embryos to be increased (Fig. 2F; wild type=1.58, *vangl2*=2.50, *glypican4*=2.92).

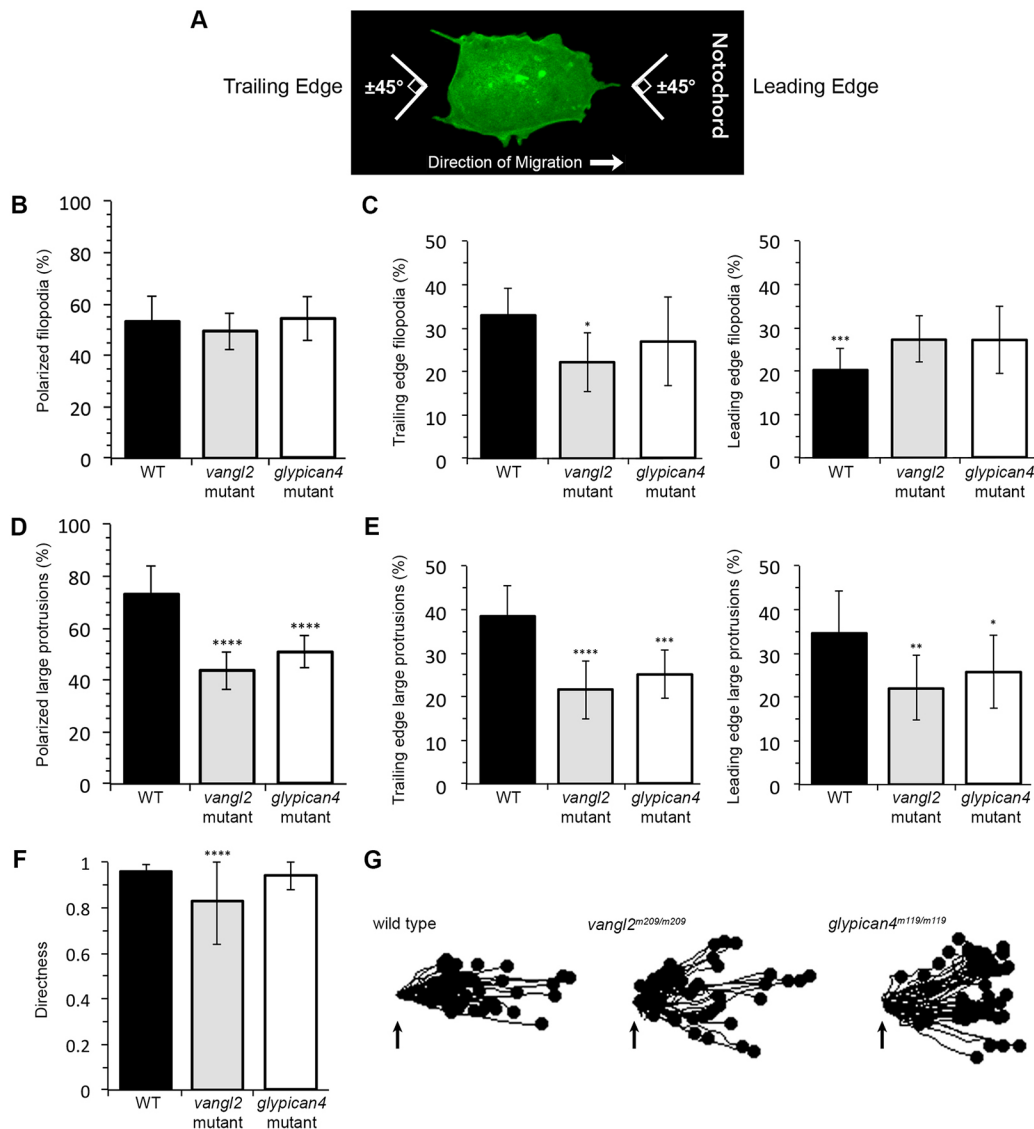
Given the difference in protrusion number between *vangl2* and *glypican4* mutant embryos, we examined the polarity of filopodia and large protrusions. Similar to MLA, protrusion polarity was determined in relation to the notochord, running anterior-posterior along the dorsal body axis. Protrusions were considered polarized if their base was oriented  $\pm 45^\circ$  from the path of dorsal migration and included protrusions at both the leading and trailing edges (Fig. 3A). We did not detect a polarized distribution of filopodia in ectodermal cells (Fig. 3B; wild type=53.23%, *vangl2*=49.52%, *glypican4*=54.27% polarized protrusions). However, there was a significant difference between the numbers of leading versus trailing edge wild-type filopodia (Fig. 3C; 32.97% and 20.26%, respectively). We also found that a decreased number of *vangl2* mutant filopodia localized to the trailing edge compared with in the wild type (Fig. 3C; *vangl2*=21.69%). Unlike filopodia, 73.10% of wild-type large protrusions are oriented along the path of dorsal migration (Fig. 3D).

By contrast, a reduced number of polarized large protrusions are observed in ectodermal cells from both *vangl2* and *glypican4* mutant embryos (Fig. 3D; *vangl2*=43.76%, *glypican4*=50.97%). We detected no difference in the numbers of leading versus trailing edge large protrusions in wild-type or PCP mutant embryos (Fig. 3E). Rose diagrams indicate that *vangl2* and *glypican4* mutant ectodermal cells generate large protrusions at multiple positions around their circumference (Fig. S1). The reduced number of polarized large protrusions in *vangl2* and *glypican4* mutant ectoderm suggests that these cells have indirect migration trajectories. Migration directness values were calculated by dividing the Euclidean distance by the accumulated or total migration distance (Table S1); a value of 1 indicates a straight migration path. Whereas *vangl2* mutant ectodermal cells have reduced directness (0.83) compared with wild-type cells (0.96), loss of Glypican4 function has a negligible effect (0.94) on directness (Fig. 3F). However, trajectory plots indicate that both mutations disrupt overall net dorsal convergence (Fig. 3G). These data imply that Vangl2 and Glypican4 are both necessary for PCP and polarized membrane-protrusive activity, but these proteins have distinct effects on protrusion formation and directed cell movement.

### Vangl2 overexpression affects membrane-protrusive activity, similar to the loss of Vangl2

Another hallmark of core PCP protein function is that excess activity results in cell polarity defects (Yang and Mlodzik, 2015). Previously published work has not reported Vangl2 overexpression relative to Vangl2 loss-of-function cellular phenotypes. Therefore,





**Fig. 3. Membrane protrusion polarity and directed migration in *vangl2* and *glypican4* mutant embryos.** (A) Wild-type (WT) ectodermal cell image illustrating polarized protrusion angles. (B, C) Quantitation of the total percentages of polarized filopodia (B) and filopodia localized at the leading versus trailing edge (C) in WT ( $n=11$  cells, 8 embryos), *vangl2*<sup>m209/m209</sup> ( $n=10$  cells, 7 embryos) and *glypican4*<sup>m119/m119</sup> ( $n=11$  cells, 5 embryos). (D, E) Quantitation of polarized large protrusions (D) and large protrusions localized at the leading versus trailing edge (E) in WT ( $n=12$  cells, 8 embryos), *vangl2*<sup>m209/m209</sup> ( $n=10$  cells, 7 embryos) and *glypican4*<sup>m119/m119</sup> ( $n=12$  cells, 5 embryos). (F) Directed migration values in WT ( $n=50$  cells, 11 embryos), *vangl2*<sup>m209/m209</sup> ( $n=51$  cells, 13 embryos) and *glypican4*<sup>m119/m119</sup> ( $n=50$  cells, 8 embryos). (G) Schematic of the migration paths of individual ectodermal cells. Origins (arrows) standardized for comparison. Dorsal is to the right. Data are mean $\pm$ s.d. \* $P<0.05$ , \*\* $P<0.01$ , \*\*\* $P<0.001$ , \*\*\*\* $P<0.0001$ ;  $P$  values are versus WT, except WT leading edge  $P$  value which is versus WT trailing edge; one-way ANOVA significance test followed by Tukey HSD post-hoc tests.

we performed *Vangl2* gain-of-function studies to substantiate its role in membrane protrusion formation and polarity. We verified with immunofluorescence (Fig. S2) and western blot (Fig. S3) that *vangl2* mRNA injection at the single-cell stage increases protein expression. We used live imaging and whole-mount *in situ* hybridization to visualize the severity of the convergence and extension phenotype at the end of gastrulation. Compared with controls, *vangl2*-injected wild-type embryos are broadened mediolaterally and have shorter anterior-posterior axes (Fig. S3). Next, using mosaic memGFP expression and time-lapse imaging, we analyzed PCP and membrane protrusions in *vangl2*-injected embryos. Again, *vangl2* mRNA injection was done at the single-cell stage to promote ubiquitous protein expression. The LWRs and MLA of *vangl2* overexpressing lateral ectodermal cells are reduced compared with wild type (Fig. S3). Increased *Vangl2* causes lateral

ectodermal cells to average twice as many filopodia (27.63) and large protrusions (6.44) as control cells (Fig. S3). *Vangl2* overexpression also disrupts filopodium and large protrusion polarity (Fig. S1; Fig. S3; 40.74% and 49.64% polarized protrusions, respectively). Lastly, increasing *Vangl2* protein levels has a strong effect on directed ectodermal cell migration (Fig. S3; wild type=0.96, *vangl2* mRNA=0.69). These results indicate that a specific level of cell-surface *Vangl2* expression is essential for PCP, membrane-protrusive activity, and directed migration.

#### ***Vangl2* expression is enriched in forming membrane protrusions**

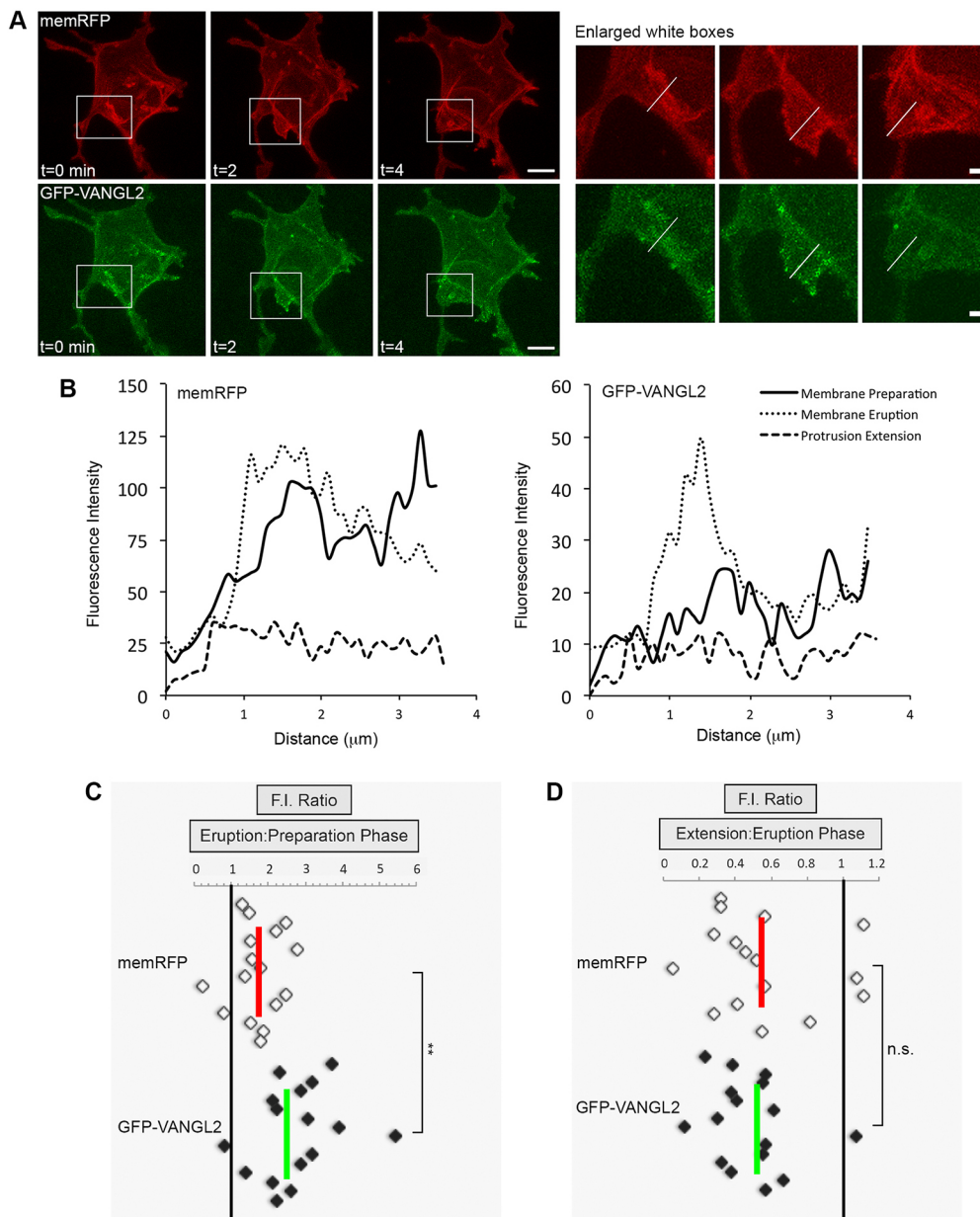
Previous work using polarized zebrafish notochord cells has demonstrated asymmetrical anterior-membrane-biased *Vangl2* protein expression at post-gastrulation stages (Roszko et al.,



2015). Axial cells are unlike the lateral gastrula cells that are engaged in directed migration, in terms of dynamic changes in membrane-protrusive activity and position relative to neighboring cells. Therefore we examined the distribution of a human GFP-VANGL2 fusion protein (Roszko et al., 2015) in anterior, posterior, leading edge and trailing edge plasma membrane domains of wild-type lateral ectodermal cells. Here, anterior/posterior refers to the cell axis perpendicular to the path of dorsal migration, whereas the leading edge/trailing edge axis is parallel to the migration path (Fig. S4). Single blastomeres from eight-cell-stage embryos were injected with synthetic mRNA that encoded GFP-VANGL2 to generate mosaic expression, and cells were imaged at the tailbud-1-somite stage. We injected GFP-VANGL2 at low doses so as not to perturb cell polarity (Roszko et al., 2015). Blastomeres were co-injected with memRFP to control for plasma membrane variations and changes in fluorescence intensity due to membrane thinning or thickening. Consistent with published data (Roszko et al., 2015), GFP-VANGL2 localizes in a punctate pattern at the plasma

membrane and within putative vesicular compartments (Fig. S4, Fig. 4A). By analyzing the fluorescence intensity ratios at multiple positions along the anterior to posterior membrane domains, we concluded that GFP-VANGL2 expression is not broadly asymmetrical in polarized lateral ectodermal cells (Fig. S4). Similar results were obtained for the leading edge and trailing edge membrane domains (Fig. S4).

The increased number of membrane protrusions in *vangl2* mutant cells suggests that Vangl2 may normally function to inhibit protrusive activity, or limit protrusion formation or lifespan at specific membrane domains. Therefore, we quantified GFP-VANGL2 protein expression during the formation of large protrusions. Large protrusions were chosen because they are associated with migratory behaviors, such as cell body translocation. We quantified fluorescence intensity at three subjectively named time points: before protrusion formation (preparation phase); during early protrusion development (eruption phase); and after protrusion elongation or remodeling (extension phase). Blastomeres were



**Fig. 4. Increased Vangl2 expression in forming membrane protrusions.**

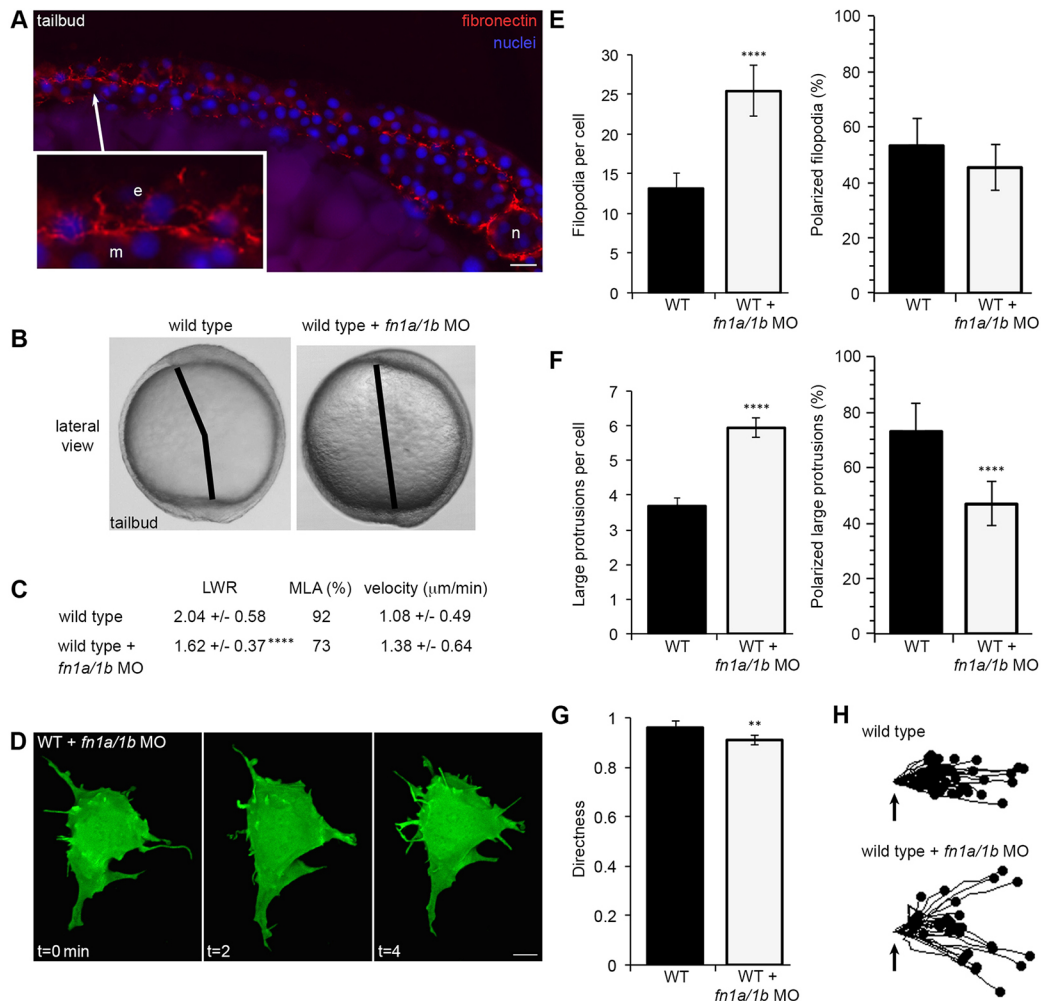
(A) Time-lapse confocal images of wild-type lateral ectodermal cells expressing memRFP and GFP-VANGL2 (three time points shown). White boxes highlight a forming large protrusion (enlarged images on the right; white lines approximate the regions analyzed). (B) Plot profiles of fluorescence intensity (F.I.) across the plasma membrane of an individual large protrusion expressing memRFP (left) and GFP-VANGL2 (right) before protrusion formation (preparation), after formation (eruption) and during extension. (C, D) memRFP and GFP-VANGL2 F.I. ratios for individual large protrusions [eruption phase/preparation phase (C),  $n=16$  protrusions, 6 embryos; extension phase/eruption phase (D),  $n=16$  protrusions, 6 embryos]. Vertical red and green lines indicate the averages. \*\* $P<0.01$ ; two-tailed paired  $t$ -test. n.s., not significant. Scale bars: 5  $\mu\text{m}$  in A, left; 1  $\mu\text{m}$  in A, right.

injected with a combination of GFP-VANGL2 and memRFP, and time-lapse imaging was performed near the tailbud stage (Fig. 4A). We report a significant increase in GFP-VANGL2 protein localization within erupting membrane protrusions (Fig. 4B,C), that becomes reduced as protrusions extend or remodel (Fig. 4B,D). GFP-VANGL2 expression within erupting protrusions is enriched compared with non-protrusive membrane domains (Fig. S5). However, our studies found that there was no difference in GFP-VANGL2 localization between polarized and non-polarized protrusions (Fig. S5). In conclusion, although GFP-VANGL2 does not exhibit widespread asymmetrical membrane expression in migrating gastrula cells, our data demonstrate protein enrichment in membrane domains that are associated with protrusive activity.

### Fibronectin is required for membrane protrusion formation and polarity

Our published data report that *vangl2* mutant embryos and *prickle1a* morphants have reduced fibronectin (Dohn et al., 2013; Williams

et al., 2012). Experiments using the frog gastrula indicate that integrin-fibronectin interactions are essential for polarized membrane protrusion formation (Davidson et al., 2006). Therefore we hypothesized that loss of fibronectin disrupts protrusion formation and, perhaps, PCP in lateral ectodermal cells. Zebrafish express two fibronectin genes (*fn1a* and *fn1b*) during gastrulation (Sun et al., 2005; Thisse et al., 2001; Trinh and Stainier, 2004). Correlating with the establishment of PCP, fibronectin protein expression and fibril assembly increase near the end of gastrulation (Fig. 5A) (Latimer and Jessen, 2010; Roszko et al., 2015). We used a combination of *fn1a* and *fn1b* antisense morpholino oligonucleotides to knockdown protein expression, which produced a mild convergence and extension phenotype in wild-type embryos (Fig. 5B) (Jülich et al., 2005; Latimer and Jessen, 2010). These morpholinos reduce fibronectin expression during gastrulation, a phenotype that is rescued by injection of *fn1a/1b* mRNA (Fig. S6) (Latimer and Jessen, 2010). Fibronectin knockdown alters LWRs and MLA compared with wild type, but does not reduce PCP to the extent that



**Fig. 5. Fibronectin regulates membrane protrusion dynamics.** (A) Cryosection through the dorsolateral region of a wild-type (WT) tailbud-stage embryo showing fibronectin and nuclei. Inset shows an enlarged image illustrating the relationship between ECM, ectodermal (e) and mesodermal (m) cells. n, notochord. (B) Live embryo images at late yolk-plug closure/tailbud stage in WT and *fn1a/1b* morpholino (MO)-injected WT embryos. Black lines denote the polster-tailbud angle. (C) PCP and migration velocity quantitation of ectodermal cells. LWR and MLA values were obtained from: WT,  $n=50$  cells, 13 embryos; *fn1a/1b* MO-injected WT,  $n=48$  cells, 10 embryos. (D) Representative ectodermal cell expressing memGFP over three time points from time-lapse data. (E,F) Quantitation of the average total number (left) and the total percentage of polarized (right) filopodia (E) and large protrusions (F) in WT ( $n=12$  cells, 8 embryos) and *fn1a/1b* morphant ( $n=10$  cells, 7 embryos) embryos. (G) Directed migration values in WT ( $n=50$  cells, 11 embryos) and *fn1a/1b* morphants ( $n=32$  cells, 10 embryos). (H) Schematic of the migration paths of individual ectodermal cells. Origins (arrows) standardized for comparison. Dorsal is to the right. Data are mean  $\pm$  s.d. \*\* $P < 0.01$ , \*\*\*\* $P < 0.0001$ ;  $P$  values are versus WT; two-tailed unpaired  $t$ -test. Scale bars: 20  $\mu\text{m}$  in A; 5  $\mu\text{m}$  in D.

was observed during Vangl2 loss-of-function (Fig. 5C). Reduced fibronectin expression also causes ectodermal cells to have excess numbers of filopodia and large protrusions (Fig. 5D-F; 25.42 and 5.94, respectively), similar to those found in *vangl2* mutant cells. Unlike the filopodia of *fn1a/1b* morphant ectodermal cells, the percentage of polarized large protrusions is markedly lower (Fig. S1; Fig. 5E,F; 46.88% polarized protrusions). Further, loss of fibronectin disrupts the directed migration of ectodermal cells (Fig. 5G,H; wild type=0.96, *fn1a/1b* MO=0.91). These data provide evidence that fibronectin supports proper membrane-protrusive activity, but that the ECM plays a lesser role during the establishment of PCP and convergence and extension movements.

### Fibronectin regulates cell-surface Vangl2 expression

Vangl2 protein expression at the plasma membrane increases at mid-gastrulation, immediately before the onset of Vangl2-dependent PCP and the appearance of the *vangl2* mutant convergence and extension phenotype (Roszko et al., 2015; Sepich et al., 2000). At mid-gastrulation, loss of Vangl2 increases the formation of large membrane protrusions but does not affect protrusion polarity (Fig. S7). Vangl2 membrane localization, and therefore cell-surface function, is regulated by Prickle1a and an unknown extracellular signal or signals (Dohn et al., 2013; Roszko et al., 2015). As the late gastrula undergoes Vangl2-dependent changes in cell behavior, fibrillar fibronectin increases (Latimer and Jessen, 2010). Therefore, we reasoned that fibronectin may control Vangl2 membrane accumulation. To determine the role of fibronectin in Vangl2 translocation to the cell surface, wild-type embryos were injected with *fn1a/fn1b* morpholinos and fixed for Vangl2 immunolabeling at two stages: early gastrulation (60% epiboly) and the end of gastrulation (tailbud). Vangl2 is primarily cytoplasmic during early gastrulation (Fig. S8) and translocates to the plasma membrane at mid-gastrulation stages (Roszko et al., 2015). Here, we used a polyclonal antibody that recognizes an epitope near the Vangl2 N-terminus (Fig. S2). Vangl2 expression at 60% epiboly is similar between morphant and control embryos (Fig. S8). However, by the tailbud stage, Vangl2 does not localize effectively to the plasma membranes of ectodermal (Fig. 6A,B) or mesodermal cells (Fig. 6C,D) in *fn1a/1b* morphants. Though the immunofluorescence images suggest that reduced cell-surface Vangl2 expression could be attributed to protein loss, western blot analyses indicate that morphant embryos have near-normal levels of Vangl2 protein (Fig. S9). Together, these data are evidence that fibronectin is necessary for Vangl2 translocation to the plasma membrane and suggest that the *fn1a/1b* morphant phenotype may partially be because of the loss of Vangl2 at the cell surface.

### Fibronectin rescues membrane-protrusive activity but not PCP in *vangl2* mutant embryos

The ability of fibronectin to promote cell-surface Vangl2 expression indicates a complex relationship between ECM, PCP and the formation of polarized protrusions. This is supported by our *in vitro* data that implicates human VANGL2 in the regulation of integrin-mediated cell adhesion to multiple ECM proteins (Jessen and Jessen, 2017). Loss of Vangl2 decreases fibronectin protein and fibril assembly (Williams et al., 2012) and morpholino knockdown of fibronectin in *vangl2* mutant embryos does not enhance the protrusion phenotypes (Fig. S10). Similarly, *vangl2* mRNA overexpression in *fn1a/1b* morphant embryos does not enhance the individual protrusion phenotypes (Fig. S11). To further elucidate the relationship between fibronectin and Vangl2, we tested whether an increase in fibronectin suppresses the protrusion

phenotypes in *vangl2* mutants. We injected single-cell-stage embryos with an equal mixture of synthetic mRNA that encoded *fn1a* and *fn1b*. Both *fn1a/1b* mRNA-injected *vangl2* mutant and wild-type embryos showed increased fibronectin, but *vangl2* mutants, in particular, had increased fibrillogenesis (Fig. 7A). In addition to *fn1a/1b* mRNA, embryos were injected with memGFP at the eight-cell-stage to achieve mosaic labeling and cells were imaged using time-lapse microscopy. Ectodermal cells from *fn1a/1b*-injected *vangl2* mutants have near wild-type numbers of filopodia and large membrane protrusions (Fig. 7B-D; 15.69 and 4.13, respectively). Despite the rescue of protrusion number, injected *vangl2* mutant embryos retain the convergence and extension phenotype that is seen in control embryos (Fig. S12). Therefore, we further examined membrane protrusion polarity and the PCP of ectodermal cell bodies. Total filopodia polarity is unaffected in *vangl2* mutant embryos, and remains unchanged in *fn1a/1b*-injected mutants (Fig. 7C). Defective polarization of large protrusions in *vangl2* mutants was partially rescued by increasing fibronectin expression (Fig. S1; Fig. 7D; 54.37% polarized protrusions). By contrast, neither the elongation nor orientation of *vangl2* mutant ectodermal cells was improved (Fig. 7E). Nonetheless, increasing fibronectin in *vangl2* mutant embryos rescued the directed migration phenotype (Fig. 7F,G). These data confirm that PCP is required for convergence and extension movements and illustrate that ECM supports proper membrane-protrusive activity and directed cell migration. Fibronectin however, is not sufficient for establishment of PCP.

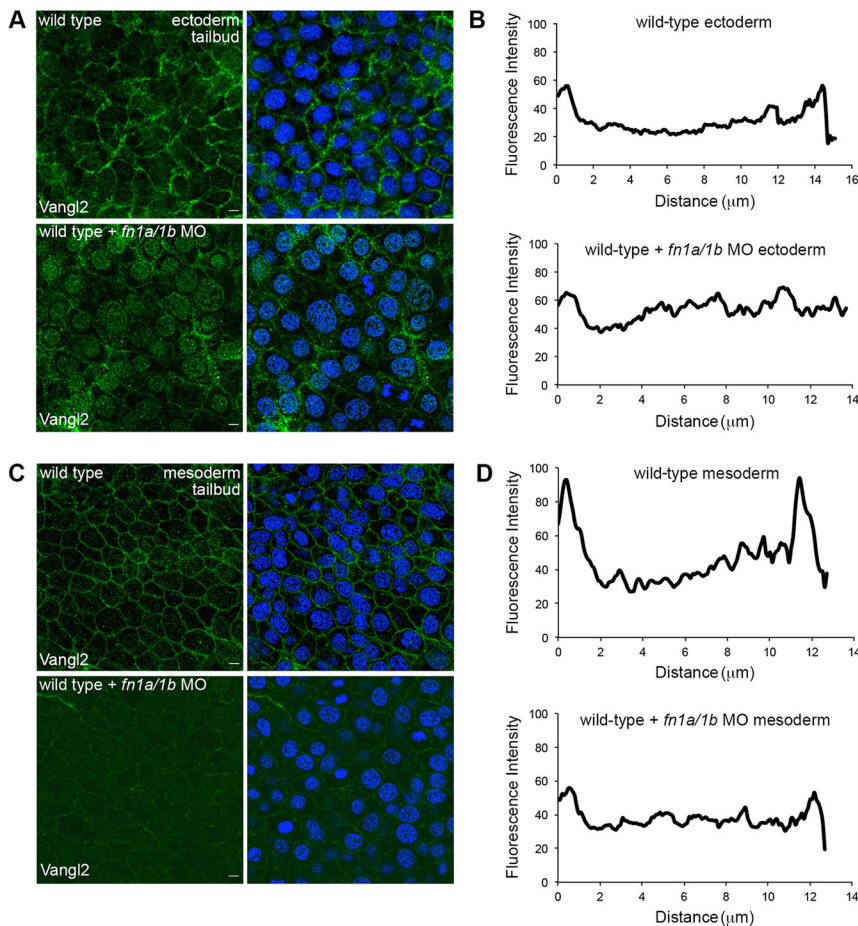
### DISCUSSION

PCP proteins are linked to several morphogenetic processes, including the gastrulation movements of convergence and extension (Gray et al., 2011). At ~80% epiboly lateral gastrula cells engage in directed migration behaviors, which continue through the tailbud stage (Sepich et al., 2005). As they migrate, round cells elongate and align mediolaterally (Roszko et al., 2015). This event is accompanied by a straightening of migration trajectories toward the dorsal axis, a process that fails to occur in *vangl2* mutant embryos (Jessen et al., 2002). In the fly wing, core PCP proteins regulate the production and asymmetrical positioning of a single actin-rich trichome in each epithelial cell (Strutt and Strutt, 2007; Taylor et al., 1998). Thus regulation of actin rearrangement is thought to underlie the establishment of PCP in vertebrate tissues. However, it has been unclear whether PCP proteins regulate actin-rich membrane protrusions in migrating lateral gastrula cells. Here, we have systematically analyzed formation and polarization of protrusions in cells with disrupted Vangl2 function and compared these phenotypes with those obtained after manipulation of Glypican4/PCP signaling. We defined cellular domains of Vangl2 protein expression and report that Vangl2 enrichment at the cell surface requires fibronectin. We have established a role for the ECM during protrusion formation and polarization, and link fibronectin with Vangl2 function.

### Regulation of membrane-protrusive activity by PCP proteins

During early zebrafish gastrulation, before the onset of PCP-dependent cell movements, dorsal prechordal plate progenitor cells use a combination of blebs and actin-containing filopodia and large lamellipodia-like protrusions (Diz-Muñoz et al., 2010). Here, large protrusions are associated with directed migration and blebs are associated with cell tumbling and meandering (Diz-Muñoz et al., 2016). Using time-lapse imaging and mosaic memGFP expression, we examined for the first time the dynamic membrane-protrusive





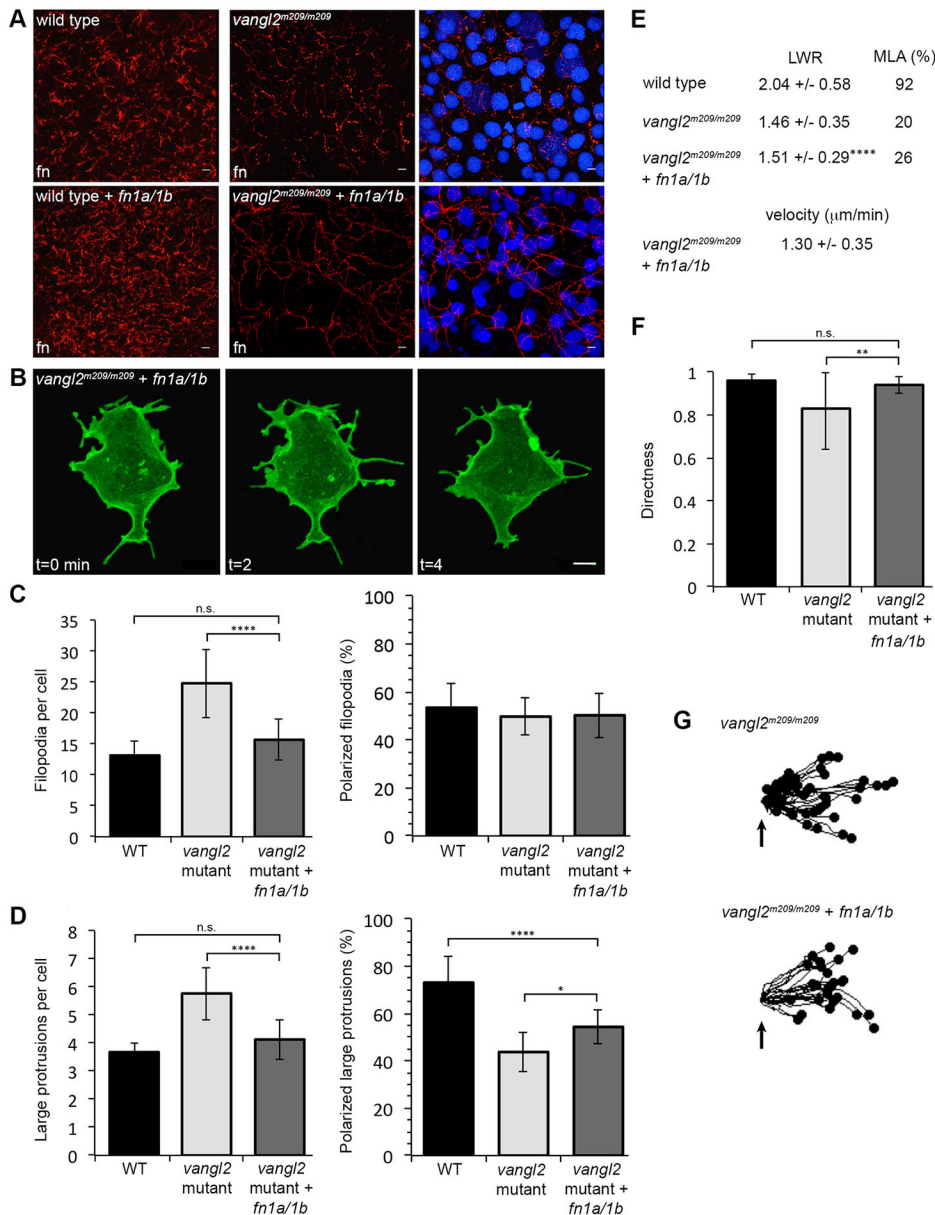
**Fig. 6. Fibronectin is required for cell-surface Vangl2 expression.** (A,C) Zebrafish Vangl2 protein expression in wild type and *fn1a/1b* morpholino (MO)-injected tailbud-stage embryos without (left) and with (right) DAPI nuclear labeling. Immunofluorescence labeling of Vangl2 in the ectoderm (A) and mesoderm (C). (B,D) Representative plot profiles showing average fluorescence intensities across single ectodermal cells in wild type ( $n=30$  cells, 3 embryos) and *fn1a/1b* morphant embryos ( $n=30$  cells, 3 embryos) in ectoderm (B) and mesoderm (D). Scale bars: 5  $\mu\text{m}$ .

activity that occurs as polarized lateral ectodermal cells undergo directed migration. Our analysis identified filopodia and large protrusions as the predominant protrusions produced by ectodermal cells. Blebs are less abundant, except in dividing cells, in which they form at cell poles to stabilize cell shape (Sedzinski et al., 2011).

Unlike *glypican4* mutant ectodermal cells, *vangl2* mutant cells have an increased number of filopodia and large protrusions. Glypican4 likely acts as a Wnt co-receptor to activate downstream proteins and, subsequently, actin regulators (Roszko et al., 2009). By contrast, Vangl2 interacts with Prickle1a to potentially contain or limit Glypican4/PCP signaling within specific cellular domains. Although the *vangl2* and *glypican4* mutant convergence and extension phenotypes are similar, our data assert that these proteins have distinct effects on the ECM. In *glypican4* mutant embryos, fibrillar fibronectin is enhanced because of increased cell-surface cadherin expression and cell adhesion (Dohn et al., 2013). Conversely, *vangl2* mutant and *prickle1a* morphant embryos exhibit decreased fibronectin expression and assembly, possibly because of increased matrix metalloproteinase activity and/or decreased cadherin expression (Dohn et al., 2013; Williams et al., 2012). We propose that fibronectin is necessary for proper formation and polarization of membrane protrusions (see further discussion below). Consequently, although it is increased in *vangl2* mutant embryos, protrusion number is not higher in *glypican4* mutants, as the fibronectin ECM remains intact. The ability of fibronectin to function as a suppressor of excess protrusion formation is supported by previous work in frog (Davidson et al., 2006).

Notably, *vangl2*- and *glypican4*-deficient ectodermal cells have excess non-polarized large protrusions, suggesting a relationship

between protrusion polarity and the establishment of PCP. Previous work reports that *vangl2* mutant mesodermal cell MLA is slightly biased along the anterior/posterior axis (Roszko et al., 2015). Although rose diagrams suggest a slight anterior/posterior bias in the polarization of large protrusions from *vangl2* and *glypican4* mutant ectodermal cells, in general, protrusions were detected around the entire cell circumference. These data support the notion that loss of protrusion polarity may represent an expansion of membrane domains that are capable of forming protrusions. A bias in wild-type filopodia polarity was not detected, indicating that these thin protrusions may play a sensory role. After surveying the extracellular environment for specific signals, we hypothesize that filopodia promote the formation of polarized large lamellipodia-like protrusions (Faix and Rottner, 2006). When wild-type lateral gastrula cells migrate dorsally, they generate more cell-cell contacts as they align and pack. One assumption is that mediolaterally oriented protrusions aid cell packing and thus directed movement. As a result, individual cell behaviors would affect morphogenesis at the tissue level (Davidson et al., 2010). In addition, manipulation of Vangl2 function interferes with ectodermal cell directed movement. Despite having a protrusion polarity phenotype similar to that found in *vangl2* mutant embryos, *glypican4* mutant cells migrate along relatively straight trajectories, indicating that polarized-protrusive activity may not be sufficient for directed migration. Unlike *vangl2* mutant embryos however, *glypican4* mutants have normal numbers of membrane protrusions, and both increased cadherin-mediated cell adhesion and increased fibronectin fibrillogenesis (Dohn et al., 2013). We hypothesize that these features contribute to more efficient directed migration of ectodermal cells in *glypican4* mutant



**Fig. 7. Fibronectin expression rescues the *vangl2* membrane protrusion phenotype but not PCP.** (A) Fibronectin expression in tailbud-stage wild type (WT) and *vangl2<sup>m209/m209</sup>* mutant control embryos (top) or WT and mutant embryos injected with *fn1a/1b* synthetic mRNA (bottom). Nuclei labeled with DAPI (right). fn, fibronectin. (B) Representative ectodermal cell expressing memGFP over three time points from time-lapse data. (C,D) Quantitation of the average total number of protrusions and the total percentage of polarized protrusions in WT ( $n=12$  cells, 8 embryos), *vangl2<sup>m209/m209</sup>* mutant control embryos ( $n=10$  cells, 7 embryos) and *vangl2<sup>m209/m209</sup>* mutants injected with *fn1a/1b* mRNA ( $n=10$  cells, 7 embryos). (E) PCP quantitation in the ectoderm. LWR and MLA values obtained from: *vangl2<sup>m209/m209</sup>* PCP and migration velocity quantitation in the ectoderm. LWR and MLA values were obtained from: WT,  $n=50$  cells, 13 embryos; *vangl2<sup>m209/m209</sup>*,  $n=10$  cells, 8 embryos; *fn1a/1b*-injected *vangl2<sup>m209/m209</sup>* mutants,  $n=50$  cells, 8 embryos. (F) Directed migration values in WT ( $n=50$  cells, 11 embryos), *vangl2<sup>m209/m209</sup>* ( $n=51$  cells, 13 embryos) and *fn1a/1b*-injected *vangl2<sup>m209/m209</sup>* mutants ( $n=26$  cells, 8 embryos). (G) Schematic of the migration paths of individual ectodermal cells. Origins (arrows) standardized for comparison. Dorsal is to the right. Data are mean  $\pm$  s.d. \* $P<0.05$ , \*\* $P<0.01$ , \*\*\*\* $P<0.0001$ ;  $P$  values are versus WT, except where indicated on the graphs; one-way ANOVA significance test followed by Tukey HSD post-hoc tests. n.s., not significant. Scale bars: 5  $\mu\text{m}$ .

embryos compared with *vangl2* mutants. The mechanism behind Glypican4/PCP signaling and its effects on cell packing and migration trajectories need to be further explored, as directed mesodermal cell migration is also Vangl2 dependent (Jessen et al., 2002).

#### Vangl2 expression in migrating ectodermal cells

A feature of fly PCP is that certain core proteins are asymmetrically expressed in the wing epithelium (Strutt and Strutt, 2009). Fluorescent fusion proteins and cell transplantation methods have allowed the identification of an anterior/posterior axis of asymmetrical PCP protein expression in non-migratory polarized axial cells. Here, Vangl2 and Prickle are enriched at anterior membranes, whereas Dishevelled localizes posteriorly (Ciruna et al., 2006; Roszko et al., 2015; Yin et al., 2008). It is important to note that axial cells do not exhibit Vangl2 or Prickle asymmetry in gastrula-stage embryos (Roszko et al., 2015; Yin et al., 2008). Based on our examination of lateral ectodermal cells that are engaged in directed migration, GFP-VANGL2 does not localize

asymmetrically along either the anterior/posterior or leading edge/trailing edge cell axes. These cells undergo rapid changes in morphology and cohesiveness as they migrate and, because Vangl2 expression is dynamic in migrating cells, an asymmetry may remain undetected. It is possible that subtle differences in Vangl2 protein stability exist at certain regions of the membrane (Chien et al., 2015). Our data do confirm that GFP-VANGL2 is enriched at the plasma membrane domains that are engaged in protrusive activity, compared with non-protrusive domains. Specifically, GFP-VANGL2 expression exhibits a transient increase within newly developed large protrusions before protrusion elongation or remodeling. These data are reminiscent of zebrafish hindbrain motor neurons, in which GFP-Vangl2 expression localizes to filopodia tips before protrusion retraction (Davey et al., 2016). Our study did not find a difference in GFP-VANGL2 expression between polarized and non-polarized protrusions. Wild-type ectodermal cells form protrusions around their circumference, but more polarized protrusions are formed along the path of migration. By contrast, *vangl2* mutant cells have both an increased number of



protrusions and an increased number of non-polarized protrusions. We hypothesize that, rather than acting as a general inhibitor of membrane protrusion formation, Vangl2 acts to destabilize, restrict or limit protrusive activity. Further work is needed to determine how Vangl2 expression and function are regulated at distinct membrane domains and how this translates into polarized cell behaviors. For example, Prickle1a may influence subtle differences in Vangl2 expression or stability at polarized versus non-polarized protrusions. Notably, *vangl2* mRNA overexpression does not inhibit protrusion formation, which suggests that a balanced level of core PCP protein expression is necessary for proper protrusive activity and directed migration. Perhaps Vangl2-dependent communication of signals within and between cells regulates localization of other PCP proteins to membrane protrusions (Strutt and Strutt, 2009).

At early gastrulation, Vangl2 protein localizes within presumptive cytoplasmic vesicles but enriches at the plasma membrane by mid-gastrulation. Vangl2 cell-surface expression at mid-gastrulation precedes the Vangl2-dependent onset of PCP and protrusion polarization. Our previous work demonstrates Prickle1a can control Vangl2 expression (Dohn et al., 2013), suggesting that *prickle1a* morphant phenotypes may be partially because of disrupted Vangl2 function. Interestingly, disruption of other polarity proteins (Glypican4, Frizzled7, Dishevelled, and Scribble1) or tissue patterning regulators does not significantly affect cell-surface Vangl2 expression (Roszko et al., 2015). Because of a temporal correlation between the requirement for Vangl2 and fibronectin fibrillogenesis at late gastrulation (Latimer and Jessen, 2010; Sepich et al., 2000), we hypothesized that the ECM influences plasma membrane levels of Vangl2. We report that loss of fibronectin inhibits translocation to the membrane, but whether fibronectin impacts Vangl2 expression directly or indirectly is unclear. Human VANGL2 interacts with integrin  $\alpha v$  and promotes cell adhesion to ECM substrates, including fibronectin (Jessen and Jessen, 2017). We hypothesize that integrin-fibronectin interactions may affect Vangl2 protein stability and function at the plasma membrane. Other signaling pathways or post-translational modifications may also contribute to Vangl2 intracellular trafficking and expression (Gao et al., 2011; Merte et al., 2009).

### Interaction of fibronectin and Vangl2 during protrusion formation and polarization

Fibronectin has long been recognized for its roles in vertebrate gastrulation. Among these, fibronectin affects cell polarity and protrusive activity, as well as providing a substrate or tracks for migration (Darribère and Schwarzbauer, 2000; Davidson et al., 2006; Winklbaauer and Keller, 1996). However, the requirement for fibronectin during zebrafish gastrulation and its relationship with PCP protein function are still emerging. Lateral ectodermal cells interact with a network of fibronectin and laminin, which forms a boundary at the ectoderm/mesoderm interface (Latimer and Jessen, 2010). Laminin mutant embryos do not have obvious gastrulation cell movement phenotypes (Parsons et al., 2002). Assembly of fibronectin fibrils relies on actin mechanotension, which is generated by integrin-mediated cell-matrix interactions and cadherin-mediated adhesion (Schwarzbauer and DeSimone, 2011). Here, we show that *fn1a/1b* morphant embryos have defects in PCP and membrane protrusion number and polarity. Although the protrusion phenotypes are similar to those of *vangl2* mutant embryos, loss of fibronectin has a lesser effect on MLA and directed cell migration. We hypothesize that the ECM is not essential for all cell behaviors underlying dorsal convergence and, further, reduced fibronectin in *vangl2* mutant gastrulae is not

sufficient to explain the severity of the *vangl2* phenotype. Supporting this notion, neither zygotic *fn1a/matter* mutant embryos (Trinh and Stainier, 2004) nor *fn1a/1b* morphants (Latimer and Jessen, 2010) exhibit strong convergence and extension defects. Moreover, the *vangl2* cell elongation, MLA and gastrulation phenotypes cannot be rescued by increasing fibronectin. Loss of Vangl2 increases matrix metalloproteinase activity and decreases cell-surface cadherin expression (Cantrell and Jessen, 2010; Dohn et al., 2013; Jessen and Jessen, 2017; Williams et al., 2012), either of which compromises the ability to assemble a normal ECM. Although fibronectin expression cannot be restored to wild-type levels in *vangl2* mutant embryos, *fn1a/1b* mRNA rescues protrusion number, directed migration and, to a lesser extent, protrusion polarity. It is unclear how a uniform fibronectin ECM promotes polarized protrusive activity, given that GFP-VANGL2 localizes to both polarized and non-polarized protrusions. It will be important to understand in more detail the dynamics of Vangl2 expression in polarized gastrula cells and to determine the localization patterns of other core PCP proteins. Our results indicate that fibronectin is required to suppress excess protrusion formation. Moreover, we argue that fibronectin regulates protrusion number and polarity directly, but by inhibiting cell-surface Vangl2 expression it influences PCP indirectly. The ability of Vangl2 to affect fibronectin proteolysis and cadherin expression (Dohn et al., 2013; Williams et al., 2012) provides possible mechanisms to regulate ECM assembly and, subsequently, membrane-protrusive activity. Future studies will further address the complex relationships between Vangl2 function, Glypican4/PCP signaling and the fibronectin ECM.

## MATERIALS AND METHODS

### Zebrafish lines and husbandry

Adult zebrafish (*Danio rerio*) 1-2 years of age were maintained following standard procedures (Solnica-Krezel et al., 1994). Embryos were collected after natural spawning, grown at 28.5°C in egg water (purified water with 60 mg/l Instant Ocean) and staged according to morphology (Kimmel et al., 1995). Embryo ages were between 6 and 10.5 h postfertilization. Strains used in this study included: wild type (AB\*, TL and WIK), *vangl2/trilobite<sup>m209</sup>* (Jessen et al., 2002; Solnica-Krezel et al., 1996), *vangl2/trilobite<sup>vu7</sup>* (Jessen et al., 2002) and *glypican4/knypek<sup>m119</sup>* (Solnica-Krezel et al., 1996; Topczewski et al., 2001). The *vangl2<sup>m209</sup>* and *glypican4<sup>m119</sup>* alleles behave like null mutations, whereas the *vangl2<sup>vu7</sup>* allele is a chromosomal deletion lacking the *vangl2* gene.

The Middle Tennessee State University Institutional Animal Care and Use Committee approved the animal research described in this paper. All procedures were conducted following approved guidelines. The Office of Laboratory Animal Welfare assurance number is A4701-01.

### Morpholinos, synthetic mRNA and embryo microinjection

Antisense morpholino oligonucleotides were obtained from Gene Tools. The following morpholinos, which have been previously published, were used: *fn1a* (5'-TTTTTTCACAGGTGCGATTGAACAC-3'), *fn1b* (5'-TACTGACTCAGGGGTCATTTTACC-3' and 5'-GCTTCTGGCTTTGAC-TGTATTTCCGG-3') and *vangl2* (5'-AGTCCACCTTACTCTGAGAGAAT-3') (Dohn et al., 2013; Jülich et al., 2005; Latimer and Jessen, 2010; Trinh and Stainier, 2004; Veeman et al., 2003; Williams et al., 2012). Each morpholino was injected at a dose of 3-10 ng/embryo. The *fn1a/1b* morpholino combination reduces fibronectin protein expression (Fig. S3) and produces the reported anterior somite defect (Jülich et al., 2005; Koshida et al., 2005; Latimer and Jessen, 2010). The *vangl2* morpholino produces convergence and extension phenotypes (Dohn et al., 2013; Williams et al., 2012) and late gastrulation ectodermal cell protrusion phenotypes similar to the *vangl2<sup>m209/m209</sup>* mutant (24.60 filopodia and 5.77 large protrusions on average per cell). None of the morpholinos used in this study stimulates obvious p53-dependent cell death.



DNA constructs encoding either Vangl2, the F-actin marker Lifeact-GFP (Riedl et al., 2008), memGFP, memRFP, GFP-VANGL2, Fibronectin1a or Fibronectin1b were cloned in the pCS2 vector and linearized by *NorI* restriction endonuclease. The fibronectin clones are not targeted by the *fn1a/1b* morpholinos. Synthetic mRNA was made using the mMACHINE™ SP6 Transcription Kit as directed by the manufacturer (Ambion) and purified by Sephadex G-50 columns (Millipore-Sigma). Microinjection into single-cell-stage or eight-cell-stage embryos was performed following standard methods (Gilmour et al., 2002). Synthetic mRNA was injected at the following doses: *vangl2* (100-300 pg); *Lifeact-GFP* (100 pg), *memGFP* and *memRFP* (100 pg); *GFP-VANGL2* (50 pg); *fn1a* and *fn1b* (200 pg).

### Whole-mount *in situ* hybridization, immunofluorescence and antibodies

Staged embryos were fixed overnight in 4% paraformaldehyde in PBS and washed with PBS that contained 0.1% Tween-20 before high resolution *in situ* hybridizations were performed as described (Thisse and Thisse, 2008). Antisense RNA probes were generated by *in vitro* transcription and purified following standard methods (Westerfield, 2000). For immunofluorescence, embryos were fixed in either 4% paraformaldehyde/PBS/sucrose fix buffer [1:1 mixture of 8% paraformaldehyde/PBS and 2× sucrose buffer; 8.0 g sucrose, 0.15 ml 0.2 M CaCl<sub>2</sub>, 90 ml 0.2 M Na<sub>2</sub>HPO<sub>4</sub> buffer (pH 7.3)] overnight or in Prefer Fixative (Anatech) for 2 h, and then manually dechorionated and permeabilized for 30 min at room temperature (5% normal donkey serum, 0.5% Triton X-100, PBS). Embryos were blocked for 2 h at room temperature [5% bovine serum albumin, 1% dimethyl sulfoxide (DMSO), 0.1% Triton X-100, PBS] with gentle rocking. Vangl2 or fibronectin primary antibodies (see below) were diluted in block solution and applied before incubating embryos overnight at 4°C with gentle rocking. Embryos were washed for 6×15 min in PBST with DMSO (0.1% Triton X-100, 1% DMSO, PBS). Embryos were treated with secondary antibodies (711-545-152 and 711-585-152, Jackson ImmunoResearch) diluted 1:500 in 50% block solution and 50% PBST (0.1% Triton X-100, PBS), and incubated for 2 h at room temperature under UV protection with gentle rocking. For nuclei labeling, embryos were washed with PBS before 30 min incubation in 2-(4-amidinophenyl)-1H-indole-6-carboxamide (DAPI). Embryos were washed 3× with PBS and stored at 4°C. Sections were obtained by embedding fibronectin immunolabeled/DAPI stained embryos in 1.2% agarose in a 5% sucrose solution followed by cryosectioning at a thickness of 10 μm. Fibronectin was detected using a rabbit polyclonal antibody generated against an epitope within the human protein (1:100, F3648, Millipore-Sigma). An affinity purified rabbit polyclonal antibody capable of recognizing an epitope near the N-terminus (NH<sub>2</sub>-RSKSRDSSSRGDKSC-COOH) of zebrafish Vangl2 was generated (1:100, ProSci). This antibody will not bind Vangl1 because of lack of epitope sequence homology, and Vangl1 is not expressed during gastrulation (Jessen and Solnica-Krezel, 2004). We validated the antibody by immunolabeling embryos overexpressing *vangl2* mRNA and embryos with a *vangl2* chromosomal deletion (Fig. S2).

### Embryo protein extraction, western blot and antibodies

For zebrafish embryo whole-cell lysates, pools of ~20-50 embryos were manually dechorionated, devolged, lysed in RIPA [50 mM Tris (pH 7.4), 150 mM NaCl, 1% NP-40, 0.5% deoxycholate, 0.1% SDS, mammalian protease inhibitor cocktail] and clarified by centrifugation. Whole-embryo lysates were boiled for 10 min following addition of Laemmli sample buffer (1× final concentration). Protein extracts were separated by 10% SDS-PAGE and transferred to a PVDF membrane using a Trans-Blot Turbo Transfer System following the manufacturer's instructions (Bio-Rad). Non-specific binding to membranes was blocked with 5% non-fat milk in TBS-Tween [50 mM Tris (pH 7.4), 150 mM NaCl, 0.1% Tween-20] and membranes were incubated overnight with rabbit polyclonal Vangl2 antibody (1:1000) in block solution at 4°C with gentle rocking. Membranes were incubated with a peroxidase-conjugated secondary antibody (1:5000, 711-035-152, Jackson ImmunoResearch), developed using Clarity ECL substrate (Bio-Rad) and imaged using a GelDoc-It Imaging System (UVP). During each experiment, blots were stripped at

room temperature for 15 min using 25 mM glycine and 1% SDS (pH 2.0) and re-probed in a 1:1000 dilution of rabbit β-actin antibody (#4967, Cell Signaling Technology) or a 1:1000 dilution of mouse GAPDH antibody (AM4300, ThermoFisher). GAPDH was also detected using a peroxidase-conjugated secondary antibody (715-035-150, Jackson ImmunoResearch). Densitometry was performed on western blots with UVP Visionworks software.

### Microscopic imaging

Staged and dechorionated live or fixed embryos were mounted as previously described (Roszko et al., 2015) and imaged with a Zeiss LSM700 confocal microscope with a 63× oil-immersion objective (N.A. 1.4). Time-lapse images were collected for 20 min at 1-2 min intervals with a 0.39 or 0.50 μm z-axis step size. Embryo orientation was documented for each time-lapse series by taking low-magnification (10×) reference images under transmitted light. Yolk-plug closure/tailbud stage live embryo images were obtained with an Olympus SZX16 stereomicroscope equipped with a Q-Color5 CCD camera. Sectioned embryo images were acquired using an Olympus IX83 inverted microscope equipped with a 20× objective and a Hamamatsu Flash 4.0 CMOS camera.

### Quantitation of cell shape, alignment, velocity and directness

Cell LWRs and MLA analyses were performed using Fiji software (fiji.sc) (Schindelin et al., 2012) as described (Jessen et al., 2002). To obtain cell migration velocity and directionality data, we tracked cell movement during our 20 min time-lapse image series (10 frames) using Fiji's Manual Tracking tool. This tool outputs data points that are distributed as scaled latitude and longitude values, which were exported to Ibidi's Chemotaxis and Migration Tool standalone software (ibidi.com) to generate plot diagrams and directness and velocity statistics.

### Analysis of membrane protrusions

We quantified and annotated membrane protrusions using Fiji software. Protrusion number and polarity were quantified as the number formed during a 20 min time interval (10 image frames). A spike-like structure distinguished actin-rich filopodia from other types of protrusions. Large actin-rich protrusions were subjectively classified by their irregular shapes and base widths >2.5 μm and included lamellipodia-like protrusions and filolamellipodia. Characterized by their spherical appearance, blebs were rare in lateral ectodermal cells. Productive protrusions were quantified as the number of large protrusions formed that were associated with cell body translocation during a 20 min time interval (10 image frames). Protrusions were classified as polarized if they had a base to tip orientation within ±45° of the path of migration. Rose diagrams were generated using the angles of individual ectodermal cell large protrusions.

### GFP-VANGL2 and Vangl2 localization

Time-lapse confocal image series were collected at the tailbud-1-somite stage as described above. Before analysis, we subtracted the image background and created sum slice projections. For the GFP-VANGL2 localization studies, we used Fiji's Plot Profile tool to quantify fluorescence intensities along the length of a line and generate an output of distance in μm (*x*-value) and grey value (*y*-value). To test for asymmetrical membrane expression bias, we drew a line through each cell from either anterior to posterior or leading edge to trailing edge membranes before producing a plot profile. During our membrane domain fluorescence intensity analysis, we avoided membrane-protrusive domains. We selected peak values at multiple positions for each membrane domain to generate anterior/posterior and leading edge/trailing edge fluorescence intensity ratios for memRFP and GFP-VANGL2. To assess GFP-VANGL2 expression dynamics in large membrane protrusions, we used Fiji's Plot Profile tool to gather fluorescence intensity data. Drawing a line inward from the plasma membrane at three time points, the Plot Profile tool produced plot profile data similar to that of our Vangl2 asymmetry study. Because membrane protrusion cycles of gastrula cells are highly dynamic in space and time, we examined GFP-VANGL2 expression at subjective time points associated with membrane protrusion preparation, eruption and extension. We selected the eruption plot apex, and the fluorescence intensity values for the corresponding

distance points ( $x$ -values) were collected for the preparation and extension phases. These data points produced eruption/preparation and extension/eruption fluorescence intensity ratios for both memRFP and GFP-VANGL2. We used the eruption phase of large protrusion formation for comparison of GFP-VANGL2 expression between protrusive and non-protrusive domains and between polarized and non-polarized protrusions. To analyze endogenous zebrafish Vangl2 expression, we collected confocal  $z$ -stack images of epiblast cells at 60% epiboly and ectodermal and mesodermal cells at tailbud stage. We drew a line across the center of each cell from anterior to posterior membranes, and using Fiji's Plot Profile tool we generated a grey value plot profile, representative of fluorescence intensities along the length of the drawn line. As we used Vangl2 antibody labeling and not mosaic expression, we could not discern adjacent cell membranes. For consistency, we used congruent techniques when performing plot profile analyses on control and experimental cells. The fluorescence intensities for each distance point along the drawn line were averaged to generate a mean grey value plot profile.

### Statistics

Data were exported to Microsoft Excel for graphing and StatCrunch software used for statistical analysis. The statistical tests performed and significance values obtained are indicated in each figure legend. The data presented are normally distributed.

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### Competing interests

The authors declare no competing or financial interests.

### Author contributions

Conceptualization: J.R.J.; Methodology: A.M.L., J.R.J.; Validation: A.M.L.; Formal analysis: A.M.L., J.R.J.; Investigation: A.M.L., D.J.P.; Resources: J.R.J.; Writing - original draft: A.M.L., J.R.J.; Writing - review & editing: A.M.L., D.J.P., J.R.J.; Visualization: A.M.L., J.R.J.; Supervision: J.R.J.; Project administration: J.R.J.; Funding acquisition: J.R.J.

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### Supplementary information

Supplementary information available online at <http://dev.biologists.org/lookup/doi/10.1242/dev.165472.supplemental>

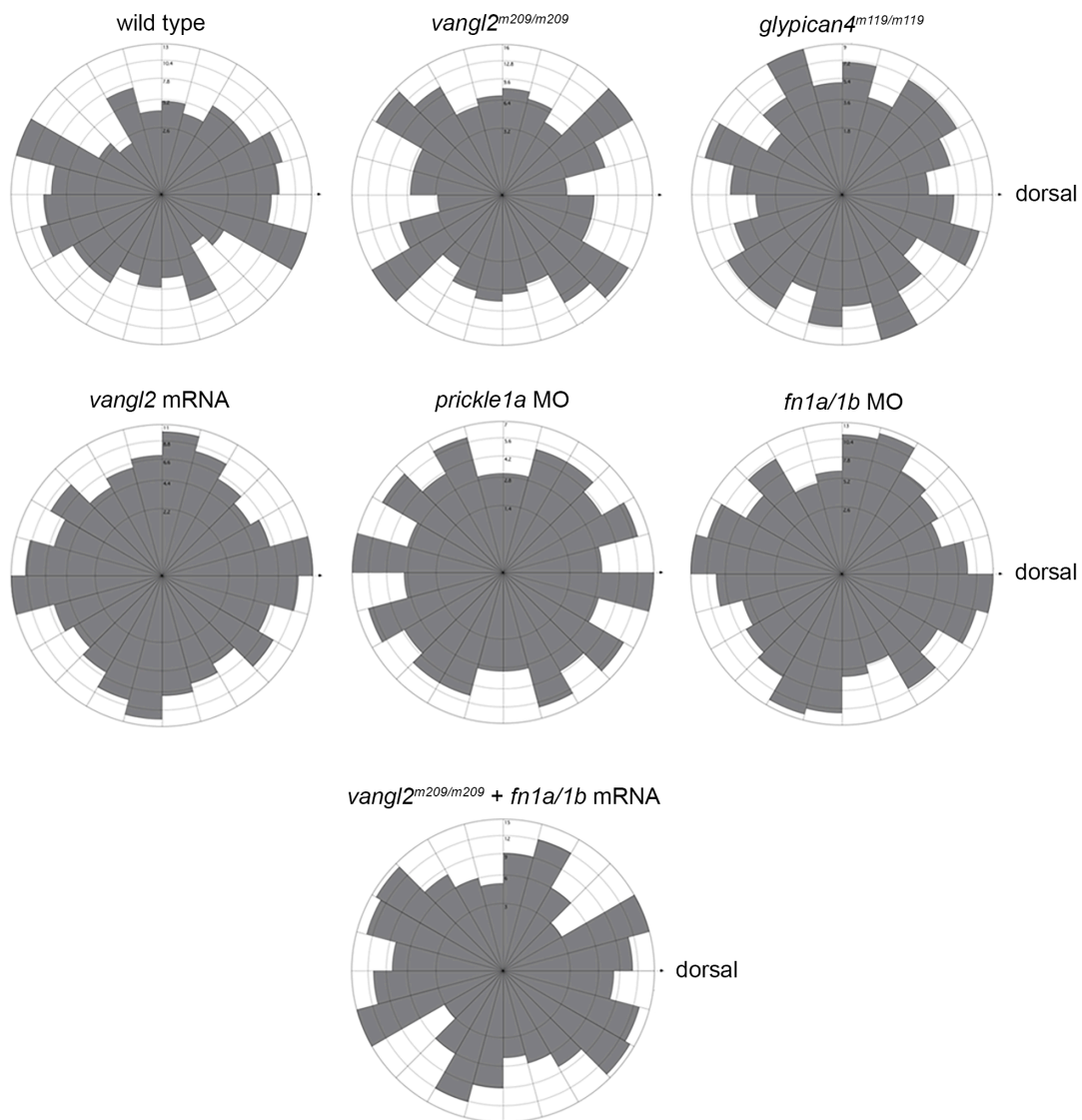
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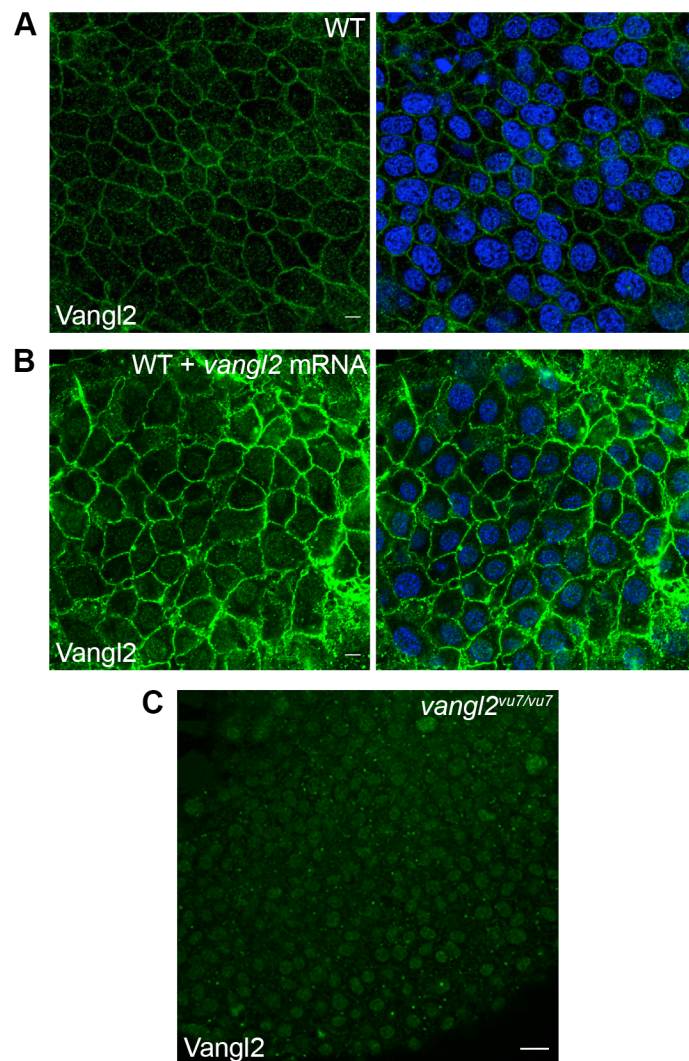


**Fig. S1. Rose diagrams showing large membrane protrusion directionality around the cell circumference.** Diagrams depicting the circular frequency and distribution of ectodermal cell large protrusions are shown for each experimental condition: wild type, *vangl2*<sup>m209/m209</sup> mutant embryos, *glypican4*<sup>m119/m119</sup> mutant embryos, *vangl2* mRNA injected wild-type embryos, *prickle1a* morpholino (MO) injected wild-type embryos, *fn1a/1b* MO injected embryos, and *fn1a/1b* mRNA injected *vangl2*<sup>m209/m209</sup> mutant embryos. The horizontal plane represents 0° and a mediolateral protrusion orientation while the vertical plane represents 90° and an anterior-posterior protrusion orientation. The position of the dorsal embryonic axis is indicated.



**Fig. S2. Zebrafish Vangl2 polyclonal antibody validation and Vangl2 overexpression.**

(A) Immunofluorescence labeling of endogenous Vangl2 protein expression in wild-type (WT) tailbud stage embryo (right panel, nuclei labeled with DAPI). (B) Vangl2 expression in a wild-type embryo injected with 200 pg synthetic *vangl2* mRNA (right panel, nuclei labeled with DAPI). (C) Endogenous Vangl2 expression in a tailbud stage *vangl2<sup>vu7/vu7</sup>* mutant embryo. Only background fluorescence from the secondary antibody is visible. Scale bars: 5  $\mu$ m in panels A and B; 20  $\mu$ m panel C.



**Fig. S3. Vangl2 overexpression disrupts PCP and membrane protrusion dynamics.** (A) Western blot of Vangl2 protein expression in *vangl2* mRNA-injected wild-type (WT) embryos. Raw unmodified densitometry values are shown. (B) Upper panels, live images of late yolk-plug closure/tailbud stage embryos. Black lines denote the polster-tailbud angle. Lower panels, whole-mount *in situ* hybridizations performed using antisense RNA probes labeling the neural/non-neural ectoderm boundary (*dlx3b*), midline (*shha*), and prechordal mesendoderm (*ctslb*). (C) PCP and migration velocity quantitation in the ectoderm. LWR and MLA values were obtained from: wild type n values are as in Fig. 2; *vangl2*-injected wild type n=50 cells, 9 embryos. (D) Representative ectodermal cell expressing memGFP over three time points from time-lapse data. (E,F) Quantitation of the average total number of protrusions and the total percentage of polarized protrusions in wild type (n values are as in Figs 2 and 3) and *vangl2*-injected wild-type embryos (n=10 cells, 7 embryos). (G) Directed migration values (wild type n values are as in Fig. 3; *vangl2*-injected n=43 cells, 9 embryos). (H) Schematic representations of the migration paths of individual ectodermal cells. Origins (arrows) standardized for comparison. Dorsal is to the right. Average values are shown  $\pm$  standard deviation.  $**P<0.01$ ,  $****P<0.0001$ ; *P* values are versus wild type; two-tailed unpaired *t*-test. Scale bar, 5  $\mu$ m.



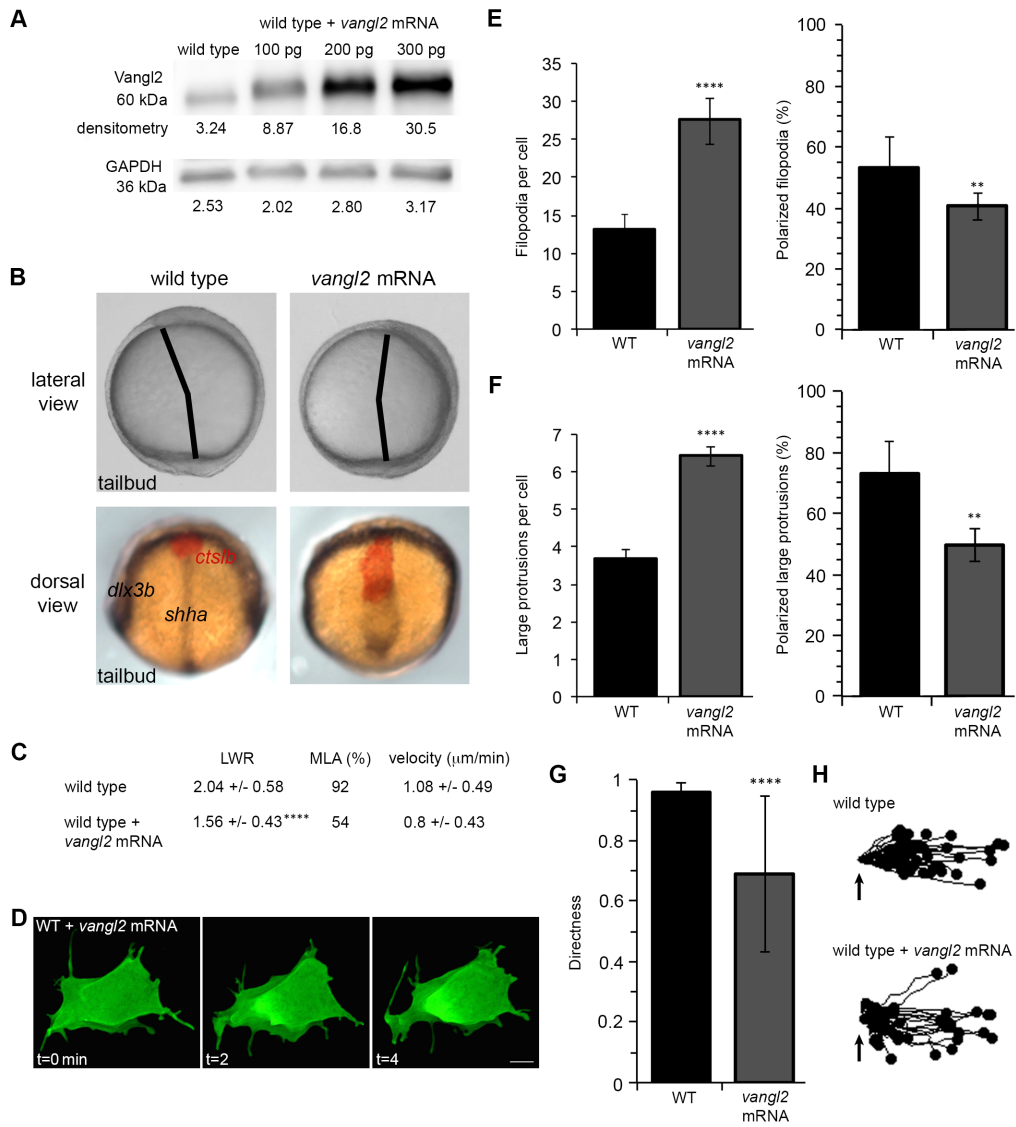
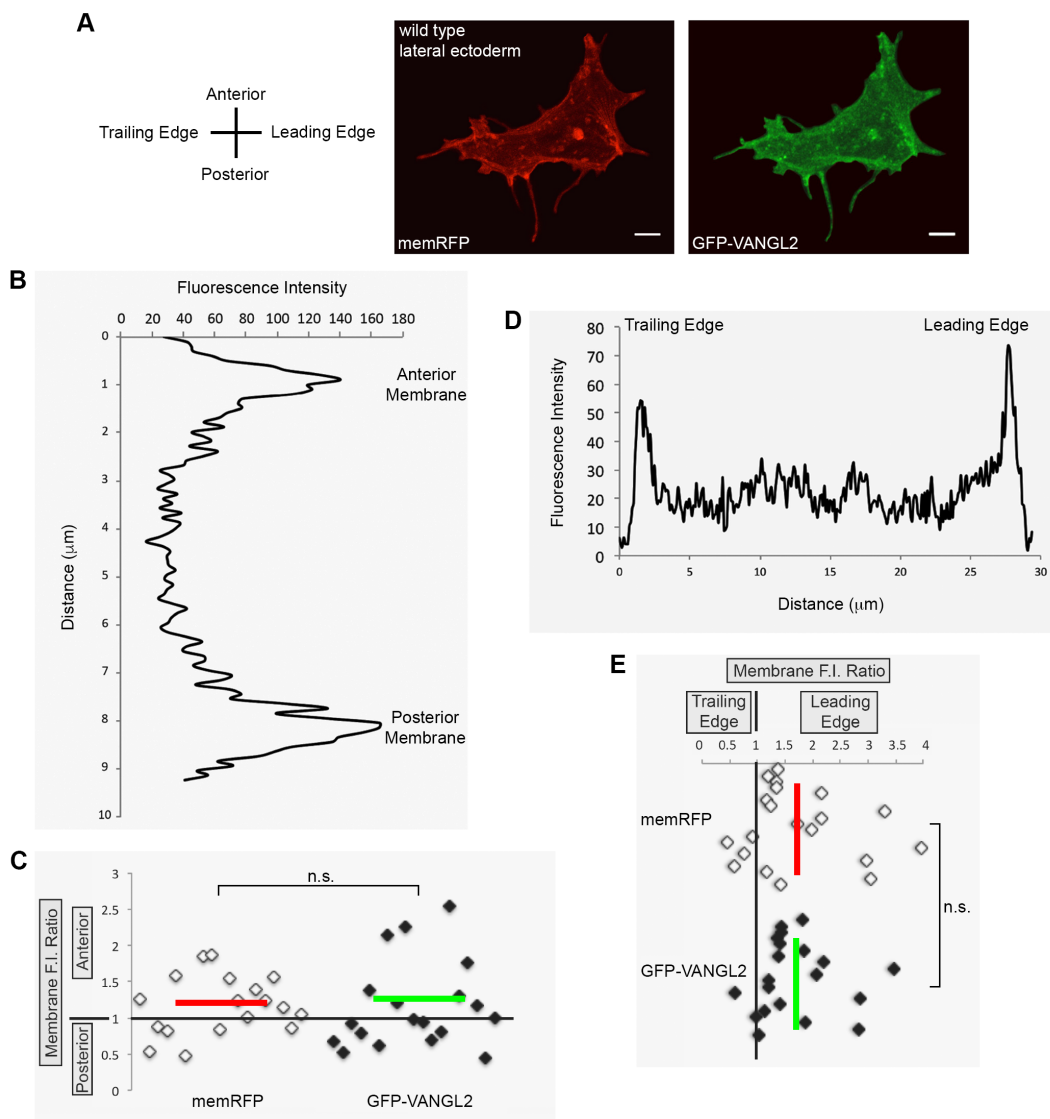
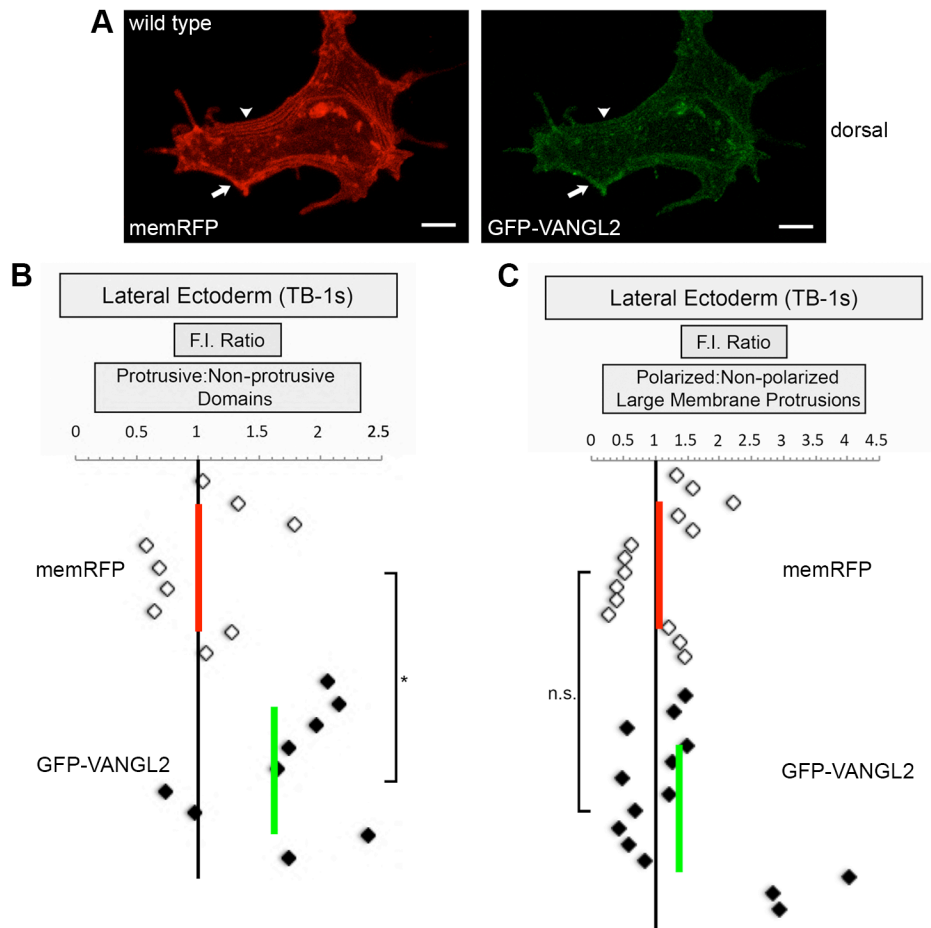


Figure S3

**Fig. S4. Vangl2 expression in polarized migrating gastrula cells.** (A) Tailbud-1-somite stage wild-type lateral ectodermal cells injected with memRFP and GFP-VANGL2 synthetic mRNA. (B,D) Plot profiles of fluorescence intensity (F.I.) across the anterior/posterior (B) and leading edge/trailing edge (D) axes of single ectodermal cells. (C,E) memRFP and GFP-VANGL2 anterior/posterior (C) and leading edge/trailing edge (E) membrane F.I. ratios for individual lateral ectodermal cells (anterior/posterior n=18 cells, 15 embryos; leading edge/trailing edge n=20 cells, 15 embryos). Horizontal and vertical red and green lines indicate averages. n.s. not significant; two-tailed paired *t*-test. Scale bars, 5  $\mu$ m.

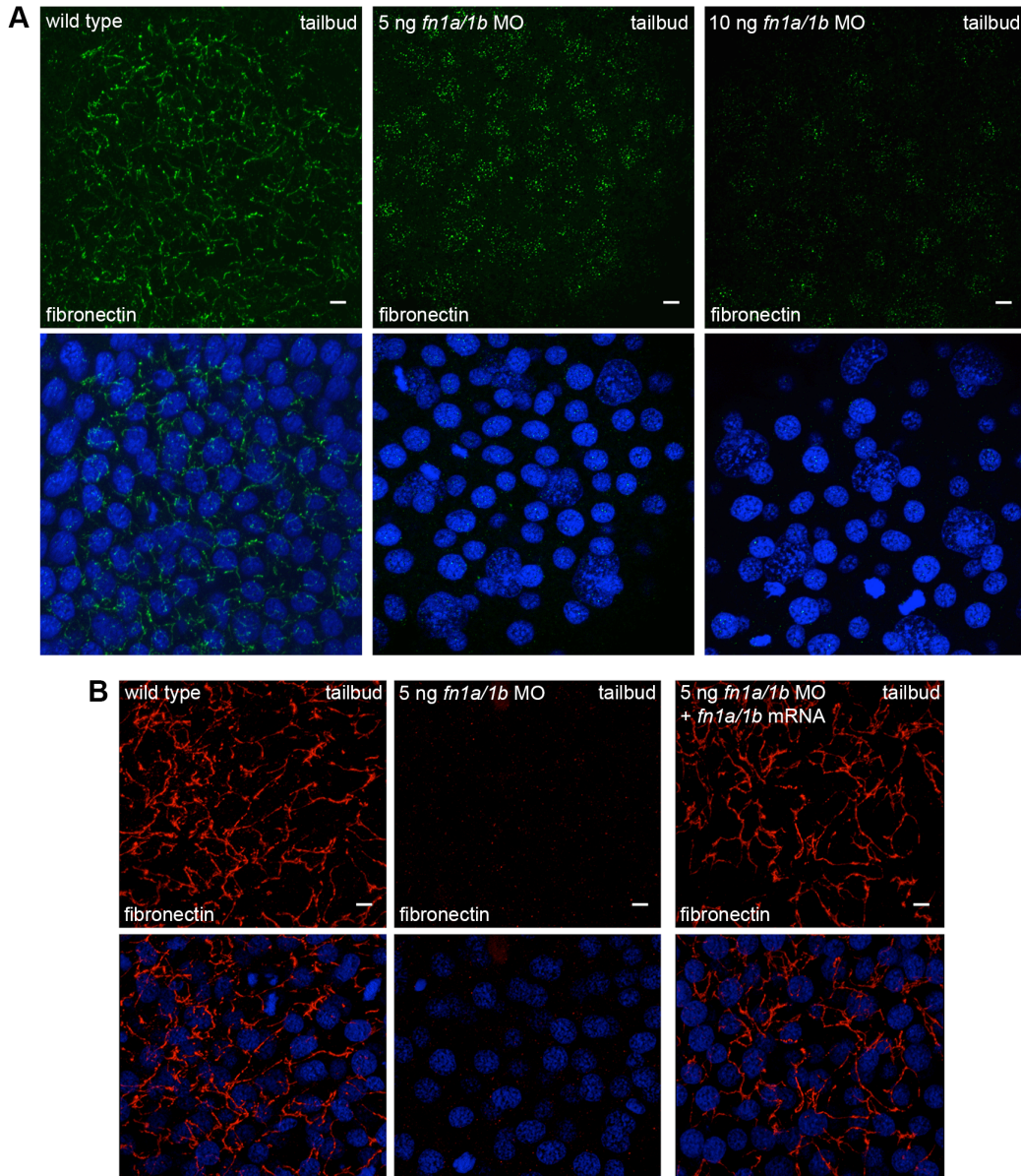


**Fig. S5. GFP-VANGL2 localization in protrusive and non-protrusive membrane domains.** (A) Lateral ectodermal cell labeled with memRFP and GFP-VANGL2. Arrows denote a forming non-polarized large protrusion. Arrowheads show a non-protrusive membrane domain. memRFP and GFP-VANGL2 fluorescence intensity (F.I.) ratios for (B) protrusive/non-protrusive membrane domains (n=9 protrusions, 6 embryos) and (C) polarized/non-polarized large protrusions (n=14 protrusions, 5 embryos). Vertical red and green lines indicate the averages. \* $P < 0.05$ , n.s. not significant; two-tailed paired *t*-test. Scale bars, 5  $\mu\text{m}$ .

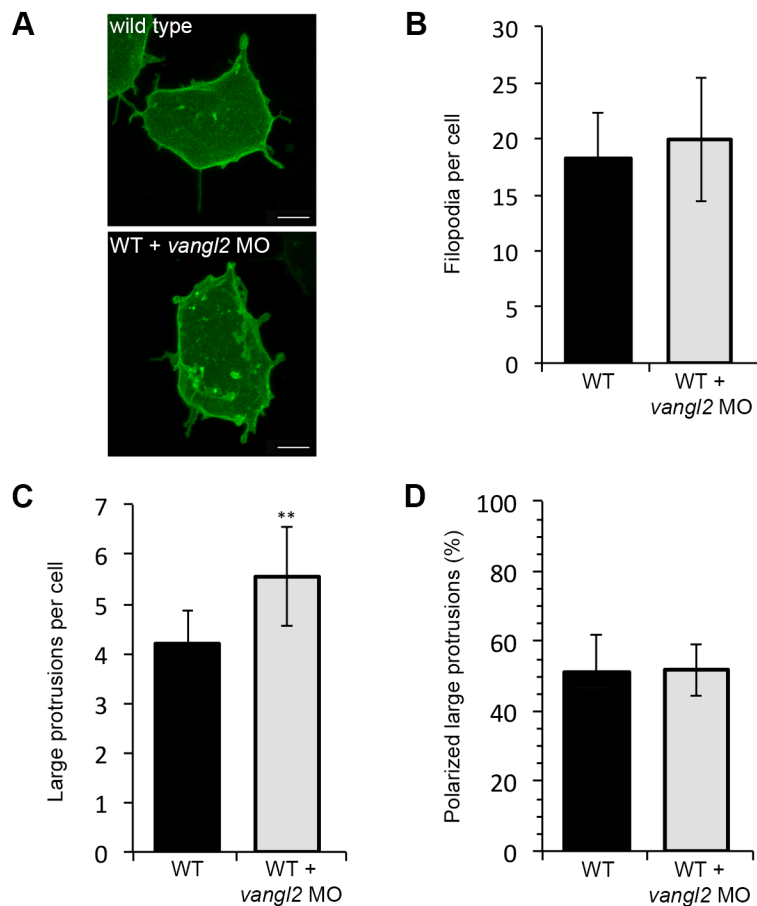




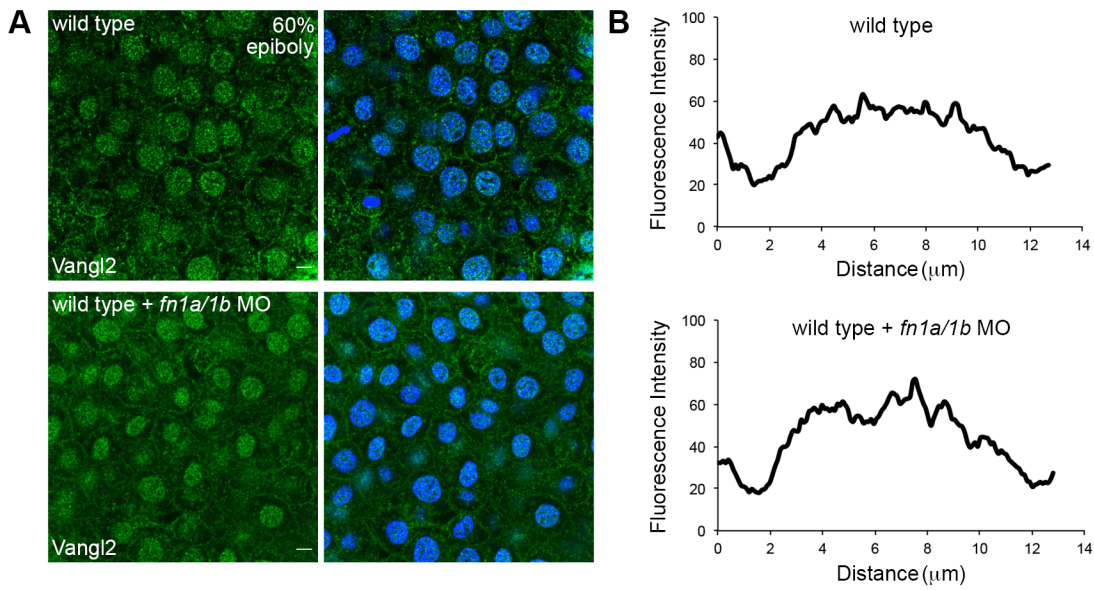
**Fig. S6. Morpholino knockdown disrupts fibronectin expression and fibrillogenesis.** (A) Immunofluorescence labeling of fibronectin expression in tailbud stage wild-type embryo and wild-type embryos injected with either 5 ng or 10 ng of each *fibronectin* morpholino (MO). (B) Injection of *fn1a/1b* synthetic mRNA rescues the *fn1a/1b* morpholino ECM phenotype. Bottom images in (A) and (B) show DAPI-labeled nuclei. Scale bars, 5  $\mu$ m.



**Fig. S7. Wild type and *vangl2* morphant membrane protrusive activity at mid-gastrulation.** (A) Time-lapse confocal images of 80% epiboly epiblast cells expressing memGFP. WT, wild type. (B,C) Quantitation of the total numbers of membrane protrusions formed by wild type (n=9 cells, 7 embryos) and wild-type embryos injected with *vangl2* morpholino (MO) (n=10 cells, 6 embryos). (D) Quantitation of the total percentage of polarized large protrusions. Average values are shown  $\pm$  standard deviation.  $**P < 0.01$ ;  $P$  values are versus wild type; two-tailed unpaired  $t$ -test. Scale bars, 5  $\mu$ m.

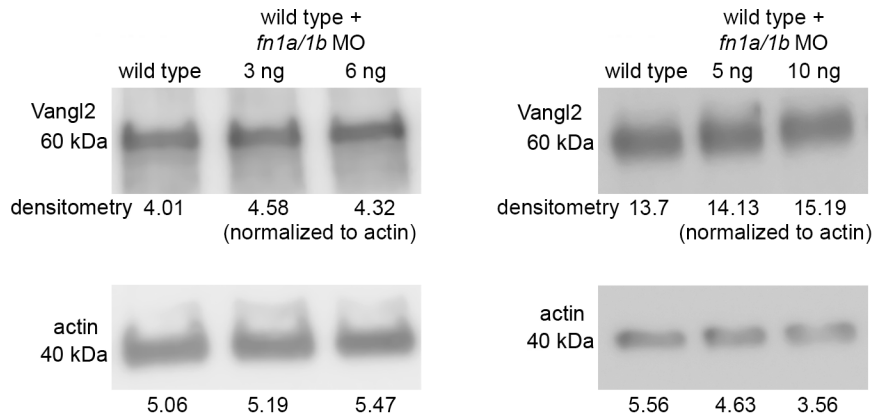


**Fig. S8. Fibronectin is not required for Vangl2 expression at 60% epiboly.** (A) Immunofluorescence labeling of Vangl2 protein expression in wild type and *fn1a/1b* morpholino (MO)-injected wild-type embryos without (left panels) and with (right panels) DAPI nuclear labeling. (B) Representative plot profiles showing the average fluorescence intensities across single epiblast cells in wild type (n=30 cells, 3 embryos) and *fn1a/1b* morphant embryos (n=30 cells, 3 embryos). Scale bars, 5  $\mu\text{m}$ .

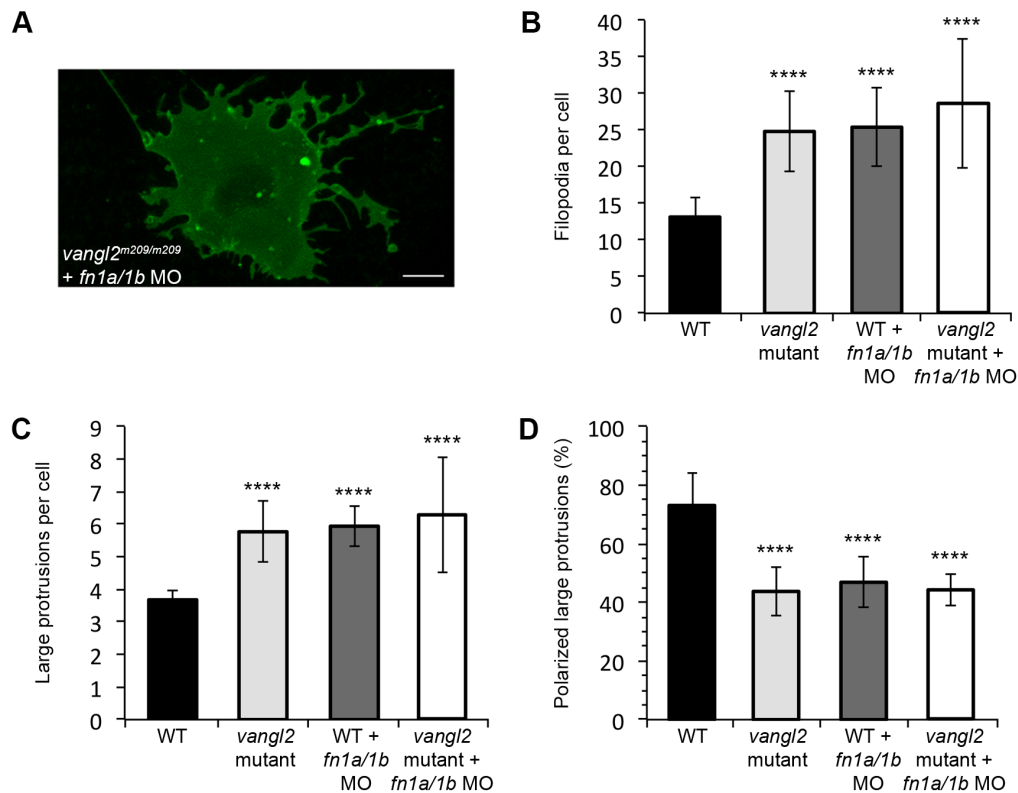




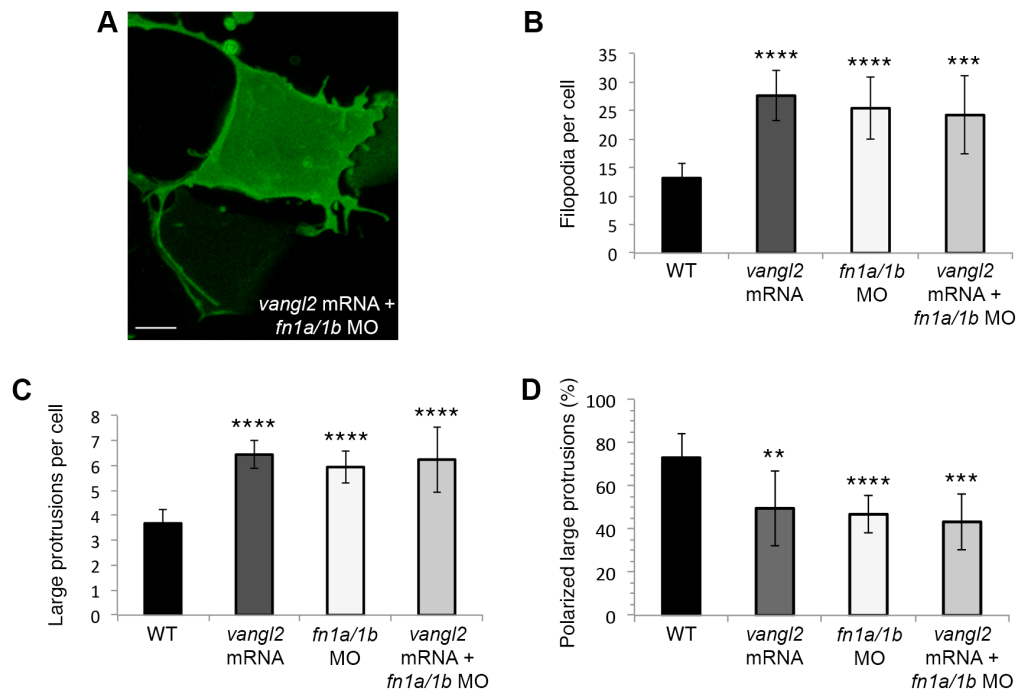
**Fig. S9. Western blot analysis of Vangl2 expression in *fibronectin* morphants.** Blots of total protein extracts from wild-type control embryos and wild-type embryos injected with 3, 6, 5, and 10 ng of each *fibronectin* morpholino (MO). Vangl2 labeled with a rabbit polyclonal antibody and actin loading control. Densitometry numbers for Vangl2 are normalized to actin.



**Fig. S10. Knockdown of fibronectin in *vangl2* mutant embryos.** (A) Time-lapse confocal image of late gastrula stage cell expressing memGFP. *vangl2* mutant embryo injected with *fn1a/1b* morpholinos (MO). (B,C) Quantitation of the total numbers of membrane protrusions formed by *vangl2* mutant embryos injected with *fn1a/1b* MO (n=5 cells, 3 embryos). The data from wild type (WT), *vangl2* mutant embryos, and wild-type embryos injected with *fn1a/1b* MO are shown for comparison. (D) Quantitation of the total percentage of polarized large protrusions. Average values are shown  $\pm$  standard deviation. \*\*\*\* $P < 0.0001$ ;  $P$  values are versus wild type; one-way ANOVA significance test followed by Tukey HSD post-hoc tests. Scale bar, 5  $\mu$ m.

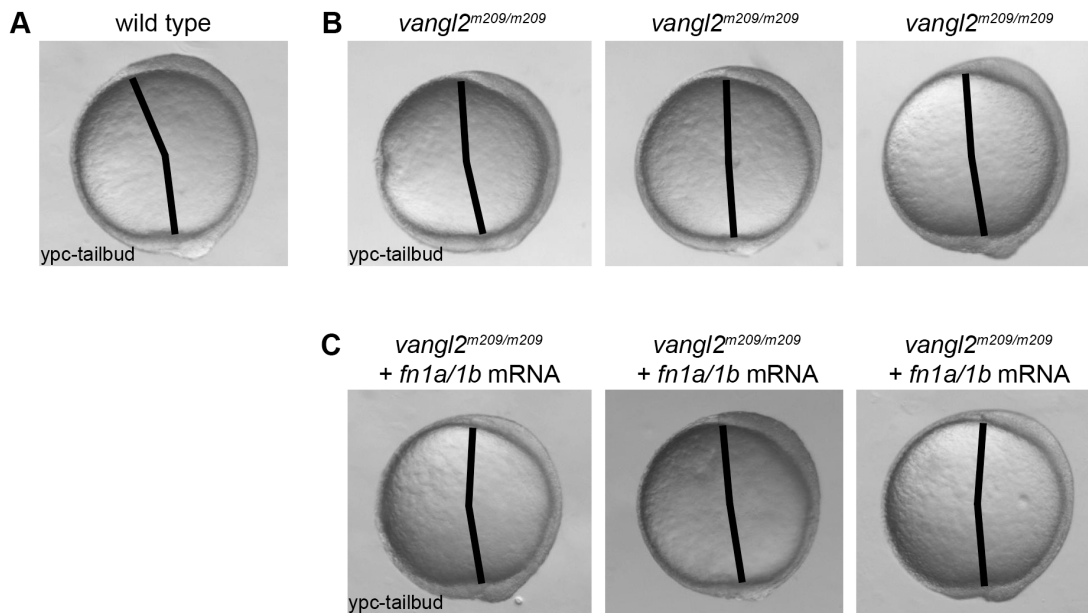


**Fig. S11. *vangl2* mRNA overexpression in *fn1a/1b* morphant wild-type embryos.** (A) Time-lapse confocal image of late gastrula cell expressing memGFP. *fn1a/1b* morpholino (MO) injected wild-type embryo co-injected with synthetic *vangl2* mRNA. (B,C) Quantitation of the total numbers of membrane protrusions formed by wild-type embryos injected with *fn1a/1b* MO and *vangl2* mRNA (n=5 cells, 4 embryos). The data from wild type (WT), *vangl2* mRNA injected wild-type embryos, and *fn1a/1b* MO injected wild-type embryos are shown for comparison. (D) Quantitation of the total percentage of polarized large protrusions. Average values are shown  $\pm$  standard deviation. \*\* $P < 0.01$ , \*\*\* $P < 0.001$ , \*\*\*\* $P < 0.0001$ ;  $P$  values are versus wild type; one-way ANOVA significance test followed by Tukey HSD post-hoc tests. Scale bar, 5  $\mu\text{m}$ .





**Fig. S12. Injection of *fn1a/1b* mRNA fails to rescue the *vangl2* mutant embryo convergence and extension phenotype.** (A) Lateral view of a wild-type embryo at the yolk-plug closure (YPC)-tailbud stage. Black lines denote the polster-tailbud angle. (B) Lateral views of three *vangl2*<sup>m209/m209</sup> embryos highlighting subtle variations in the mutant convergence and extension phenotype. (C) Lateral views of three *vangl2*<sup>m209/m209</sup> embryos injected with *fn1a/1b* mRNA (n=50 mRNA injected homozygous mutant embryos analyzed).



**Table S1.** Accumulated and Euclidean ectodermal cell translocation distances.

	Accumulated distance ( $\mu\text{m}$ )	Euclidean distance ( $\mu\text{m}$ )
wild type	19.49 +/- 8.82	18.81 +/- 8.78
<i>vangl2</i> <sup>m209/m209</sup>	17.68 +/- 9.70	15.76 +/- 10.54
<i>glypican4</i> <sup>m119/m119</sup>	21.22 +/- 5.05	20.02 +/- 5.18
wild type + <i>vangl2</i> mRNA	14.36 +/- 7.75	11.45 +/- 9.06
wild type + <i>prickle1a</i> morpholino	17.74 +/- 6.18	15.68 +/- 8.04
wild type + <i>fn1a/1b</i> morpholino	24.78 +/- 11.49	22.88 +/- 11.26
<i>vangl2</i> <sup>m209/m209</sup> + <i>fn1a/1b</i> mRNA	23.45 +/- 6.34	22.03 +/- 6.15