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## Planar cell polarity breaks the bilateral symmetry by controlling ciliary positioning

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### Abstract

Defining the three body axes is a central event of vertebrate morphogenesis. Establishment of left-right (L-R) asymmetry in development follows the determination of dorsal-ventral (D-V) and anterior-posterior (A-P) body axes<sup>1,2</sup>, though the molecular mechanism underlying precise L-R symmetry breaking in reference to the other two axes is still poorly understood. Here by removing both *Vangl1* and *Vangl2*, the two mouse homologues of a *Drosophila* core planar cell polarity (PCP) gene *Van Gogh* (*Vang*), we have uncovered a previously unappreciated function of PCP in initial breaking of lateral symmetry. The leftward nodal flow across the posterior notochord (PNC, also referred to as “the node”) has been identified as the earliest event in the de novo formation of L-R asymmetry<sup>3,4</sup>. We found that PCP is essential in interpreting the A-P patterning information and linking it to L-R asymmetry. In the absence of *Vangl1* and *Vangl2*, cilia are positioned randomly around the center of the PNC cells and nodal flow is turbulent, which results in disrupted L-R asymmetry. Importantly, PCP in mouse, unlike what has been implicated in other vertebrate species, is not required for ciliogenesis, cilium motility, Sonic hedgehog (Shh) signaling or apical docking of basal bodies in ciliated tracheal epithelial cells. Our data suggest that PCP acts earlier than the unidirectional nodal flow during bilateral symmetry breaking in vertebrates and provide insight into the functional mechanism of PCP in organizing the vertebrate tissues in development.

Identifying the mechanism underlying the initial bilateral symmetry breaking has been an overriding goal in understanding how nature is able to produce organisms with consistent and distinct left and right anatomies. In mice, bilateral symmetry is broken by a leftward flow across a pit-like, teardrop-shaped PNC generated by posteriorly localized motile cilia<sup>3–6</sup>. However, an earlier asymmetric event that positions the cilia to the posterior sides of PNC

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### Author contributions

Hai Song generated most of the data in the manuscript. He designed and performed the experiments, interpreted and presented the data in figures. Jianxin Hu designed and performed some of the experiments, carried out statistical analysis, interpreted and put his data together in figures. Wen Chen analyzed and maintained the *Vangl1* mutant mice. Gene Elliott generated the *Vangl1* and *Vangl2* mutant mice by performing blastocyst injection and teratoid aggregation. Philipp Andre and Bo Gao provided some critical technique support and help in writing the manuscript. Yingzi Yang designed and supervised the entire project, interpreted all data and wrote the manuscript.

cells in response to the A-P patterning signals remains to be identified. PCP, originally refers to asymmetry of epithelial cells along an axis orthogonal to the apical-basal axis, is an evolutionarily conserved mechanism by which positional information is transmitted and maintained among and within cells<sup>7</sup>. A group of core PCP proteins including the transmembrane protein Vang have been identified in *Drosophila*<sup>8</sup>. While mutants of core PCP genes have thus far failed to show L-R asymmetry defects, it is likely due to functional redundancy among members of a core PCP component. To overcome this and uncover important regulatory functions of PCP, we inactivated both *Vangl1* and *Vangl2*, the two mouse homologues of the *Drosophila* Vang.

*Vangl1* and *Vangl2* were broadly expressed during L-R patterning (Supplementary Fig. 1). Importantly, prior to appearance of ciliated PNC cells, *Vangl1* and *Vangl2* protein levels were too low to be detected. However, after E7.75, both *Vangl1* and *Vangl2* protein levels were selectively and progressively upregulated in the PNC cells and asymmetrical localization of *Vangl1* and *Vangl2* protein was observed along the anterior-posterior (A-P) axis (Supplementary Fig. 2a–c), indicating that the PNC cells are indeed polarized and PCP may be required by the PNC cells to regulate early embryonic development. The *Vangl1* and *Vangl2* localization patterns are also consistent with the role of PCP in regulating convergent extension (CE) of neurulating vertebrate embryos<sup>9</sup>. In the sensory epithelia of the cochlea, *Vangl1* and *Vangl2* demonstrated similar asymmetrical localization pattern in both hair cells and supporting cells (Supplementary Fig. 2d, e). These data suggest that *Vangl1* and *Vangl2* play redundant roles in PCP, which should be severely disrupted by removing both *Vangl* genes.

A bona fide null allele of *Vangl1* (*Vangl1<sup>gt</sup>*) was generated using a gene trap ES cell line (Supplementary Fig. 3). We also generated a floxed *Vangl2* allele, *Vangl2<sup>f</sup>*, from which the null *Vangl2* allele was derived (Supplementary Fig. 3). *Vangl1* and *Vangl2* regulated PCP in a gene dose-dependent manner (Fig. 1 and Supplementary Fig. 4). In the cochlea of the *Vangl2<sup>-/-</sup>* mutant, asymmetrical *Vangl1* localization was still detected (Fig. 1a). The *Vangl2<sup>-/-</sup>; Vangl1<sup>gt/gt</sup>* embryos showed the most severe developmental defects compared to embryos with other genotypes, particularly when sensory hair cell polarity and CE were examined to assess PCP. Thus, *Vangl1* and *Vangl2* function of regulating PCP was removed in the *Vangl2<sup>-/-</sup>; Vangl1<sup>gt/gt</sup>* embryos.

The *Vangl2<sup>-/-</sup>; Vangl1<sup>gt/gt</sup>* embryos exhibited multiple laterality defects, which were not detected in embryos with other genotypes (Fig. 2a). The *Vangl2<sup>+/-</sup>; Vangl1<sup>gt/+</sup>* mice were fertile, morphologically normal and served as controls in our study. The *Vangl2<sup>+/-</sup>; Vangl1<sup>gt/+</sup>* and *Vangl2<sup>-/-</sup>; Vangl1<sup>gt/+</sup>* embryos always turned to the right side and their hearts correctly looped to the right (n>50), but in all of the *Vangl2<sup>-/-</sup>; Vangl1<sup>gt/gt</sup>* embryos assayed (n>10), both embryonic turning and heart looping failed. In addition, the normal mouse lung has five lobes, four on the right and one on the left. In the *Vangl2<sup>-/-</sup>; Vangl1<sup>gt/gt</sup>* embryos, however, mild left lung isomerism was observed with three lobes on the right side (4/9). Since PCP in visceral organs may also control later stage morphogenesis, to further assess the early L-R asymmetry defects in the *Vangl2<sup>-/-</sup>; Vangl1<sup>gt/gt</sup>* embryos, we examined the components in the Nodal signal cascade that are asymmetrically expressed and required for L-R patterning<sup>10</sup>. In control embryos, *Nodal* was first expressed symmetrically around

the PNC (somite 0–1). Then between somite 2–7 stages, *Nodal* expression on the left side of the PNC became progressively stronger and its expression in the left lateral plate mesoderm (LPM) was also detected<sup>11,12</sup>. In the *Vangl2*<sup>-/-</sup>; *Vangl1*<sup>gt/gt</sup> embryos, however, *Nodal* expression around the PNC was reduced (6/11) (Fig. 2b, c). Strikingly, more robust *Nodal* expression on the right side of the PNC was also observed in the *Vangl2*<sup>-/-</sup>; *Vangl1*<sup>gt/gt</sup> embryos (5/11) (Fig. 2b, c). In the LPM of *Vangl2*<sup>-/-</sup>; *Vangl1*<sup>gt/gt</sup> embryos, reversed (1/8) or bilateral (3/8) *Nodal* expression sites were observed (Fig. 2c). Controlled by *Nodal* signaling, expression of *Lefty1*, *Lefty2* and *Pitx2* was only observed on the left side of the control embryos<sup>10,13,14,15</sup>. In the double mutant embryos, *Lefty1* and 2 expression sites (1/2) were reversed (Fig. 2d) and bilateral *Pitx2* expression (2/7) was found (Fig. 2e). These results demonstrate that PCP is required upstream of *Nodal* to pattern the L-R axis. The incomplete penetrance of the laterality defects in the *Vangl1/2* mutant embryo is possibly due to other input(s) to the PCP pathway<sup>16</sup>.

Since PCP has been implicated in ciliogenesis<sup>17</sup>, we examined the morphology and movement of cilia in PNC cells. Cilia are long, thin microtubule-based projections from basal bodies of most mammalian cells<sup>18</sup>. The PNC cell has a single motile cilium<sup>3</sup>, which is essential for establishing leftward fluid flow across the ventral surface of the PNC. When examined by scanning electron microscopy and immunofluorescent labeling, cilia appeared normal in the PNC of the *Vangl2*<sup>-/-</sup>; *Vangl1*<sup>gt/gt</sup> embryos (n=4 in each case) at E8.0 (Fig. 3a, b). Thus, the L-R asymmetry defects in the *Vangl2*<sup>-/-</sup>; *Vangl1*<sup>gt/gt</sup> embryos are not secondary to defects in ciliogenesis. The leftward nodal flow is enabled by the chirality of the cilium, as it rotates in a clockwise manner<sup>3</sup>, and the posterior ciliary localization in the PNC cell, which results in a posterior tilt of the cilium and its rotation axis<sup>5,6</sup>. Since it has been postulated that the posterior positioning of the basal body might be controlled by PCP<sup>6</sup>, we examined basal body localization in the PNC cells of the *Vangl* mutant embryos (Fig. 3c, d). The basal bodies were mostly localized posteriorly in both *Vangl2*<sup>+/-</sup>; *Vangl1*<sup>gt/+</sup> and *Vangl2*<sup>-/-</sup>; *Vangl1*<sup>gt/+</sup> embryos. However, in the *Vangl2*<sup>-/-</sup>; *Vangl1*<sup>gt/gt</sup> embryos, the PNC cell density was slightly decreased (Supplementary Fig. 5a) and the number of posteriorly localized basal bodies was significantly reduced. Most basal bodies remained around the center and anteriorly localized basal bodies were increased, resulting in an even distribution of basal bodies on the anterior and posterior sides of the PNC cells in the *Vangl2*<sup>-/-</sup>; *Vangl1*<sup>gt/gt</sup> embryos (Fig. 3d and supplementary Fig. 5b, c). Such loss of basal body posterior localization in the *Vangl* mutant is not a result of developmental delay (Supplemental Fig. 5c). Thus, posterior positioning of basal bodies in the PNC cells is indeed controlled by PCP, in the absence of which, basal bodies are mostly localized around the center. Cilia in the PNC emerge from the center of each cell at late streak (LS) stage (~E 7.25). By early head fold (EHF) stage (E7.75), most cilia are located in the posterior side of the PNC cells<sup>6,19</sup>. As positioning of the kinocilium in sensory hair cells is also an event downstream of PCP<sup>20</sup>, ciliary positioning may be a common readout of PCP.

To test whether altered ciliary positioning in the PNC cells affect function, we examined nodal flow (Fig. 4a and Supplementary movie 1, 2). In control embryos, fast and smooth leftward bead movement across the PNC was observed. In the *Vangl2*<sup>-/-</sup>; *Vangl1*<sup>gt/gt</sup> embryos, bead movement was still biased leftward, though beads were often trapped in

swirling vortices and requires more time to cross the PNC. While the nodal flow of control embryos generated straight and parallel bead paths, in the mutant, bead paths contained “knots”, abrupt turns and were often crossed with each other (2/2). When observed directly, the cilia in the PNC of the *Vangl2*<sup>-/-</sup>; *Vangl1*<sup>gt/gt</sup> embryo were beating at a speed and direction similar to that of the control (Fig. 4b and Supplementary movie 3, 4). The abnormal nodal flow in the *Vangl2*<sup>-/-</sup>; *Vangl1*<sup>gt/gt</sup> embryos also disrupted the left-biased perinodal rise of intracellular calcium in control embryos<sup>21</sup> (Fig. 4c, d). Incidence of bilateral rise or reduction of intracellular calcium was increased in the *Vangl2*<sup>-/-</sup>; *Vangl1*<sup>gt/gt</sup> embryos. The ectopic calcium signal on the right side was weaker and the domain was narrower and shorter. Taken together, unidirectionality of nodal flow was significantly compromised in the *Vangl2*<sup>-/-</sup>; *Vangl1*<sup>gt/gt</sup> embryos. Thus, we have identified PCP as the first step in breaking L-R symmetry by transmitting the A-P positional information to the PNC cells to enable posterior ciliary positioning (Fig. 4e).

Our data show that PCP is not required for ciliogenesis in mouse (Fig. 1b and supplementary Fig. 6c–f). In addition, Vangl1 and Vangl2 were not colocalized with cilia in any of the cells we had examined including the PNC cells, sensory hair cells, tracheal epithelial cells and floor plate cells (Supplementary Figs. 2a, d and 6a, b). In the *Vangl2*<sup>-/-</sup>; *Vangl1*<sup>gt/gt</sup> embryos at E18.5, there were no defects in ciliogenesis or docking of the basal bodies along the apical-basal axis of the multiciliated tracheal epithelial cells (Supplementary Fig. 6c, d). Normal Hedgehog signaling in the *Vangl2*<sup>-/-</sup>; *Vangl1*<sup>gt/gt</sup> embryos and cells provides further evidence that primary cilia are functionally normal (Supplementary Figs. 6g, h and 7).

To promote a robust break in symmetry, vertebrates have evolved mechanisms of early asymmetry initiation followed by asymmetry amplification and stabilization<sup>22</sup>. As PCP is a major conserved molecular mechanism by which directional cellular and tissue behaviors are controlled through reorganizing cytoskeleton, it could lead to both cilia-dependent and cilia-independent regulation of L-R asymmetry by controlling asymmetrical ciliary localization in ciliated PNC or cell movement around the node in chick which lacks ciliated PNC<sup>5,6,23</sup>. Involvement of PCP in regulating L-R asymmetry has been implicated before since mutations in *Inversin* (*Inv*), *Bicaudal C* and *Seahorse* lead to L-R asymmetry defects upstream of *Nodal* expression and these genes can interact with a core PCP gene *Dishevelled* (*Dvl*)<sup>24–26</sup>. *Dvl* has recently been found to regulate ciliary localization in the PNC and implicated to regulate L-R patterning<sup>27</sup>. However, it is unlikely that *Inv* acts mainly through PCP since *Inv*, but not Vangl1/2, is localized to PNC cilia and the phenotypes of the *Inv* and *Vangl1/2* mutants are different in many aspects. As abnormal L-R patterning is also associated with multiple congenital diseases<sup>28</sup>, the availability of *Vangl1* and *Vangl2* conditional double mutants has opened a door to investigate the roles of PCP in many fundamental aspects of vertebrate development and diseases.

Note added in proof: While this work was in press, two independent studies<sup>29,30</sup> were published, both also found that planar cell polarity mediated by *Vangl2* is required for orienting and positioning motile cilia.

## Methods Summary

### Scanning electron microscopic analysis

Embryos were dissected at E7.75 and fixed in 2% glutaraldehyde, 4% paraformaldehyde, 0.1M sodium cacodylate, pH 7.4 at 4 °C overnight. Standard procedures were followed to prepare samples. Samples were examined by a Hitachi S-570 scanning electron microscope.

### Analysis of calcium signaling

Embryos were dissected at E7.75–E8.0 and analyzed according to the established procedure<sup>21</sup>.

### Video Microscopy

Embryos were dissected at E7.75–E8.0 in DMEM with 20% FBS. The node was placed facing down in a chamber with a cover slip on the bottom. Fluospheres (Invitrogen) were added into the medium to visualize the direction of nodal flow. Wide-field images were collected using a Personal DeltaVision system (Applied Precision Inc, Issaquah, WA, USA) mounted on an inverted Olympus IX71 microscope with an oil immersion PlanApo N 60×/1.42 objective lens. Images were acquired using a CoolSNAP ES2 camera with 2×2 binning and 128 pixels ×128 pixels imaging field.

## Methods

### Generation of and genotyping mouse lines

Generation of the floxed *Vangl2* allele is described in Supplementary Fig. 1e. *Vangl2* was generated by deleting exon 4 and 5 using the *Sox2-Cre* line<sup>31</sup>. *Vangl1<sup>gt</sup>* allele was generated using a Baygenomic gene trap ES cell line (XL802) (Supplementary Fig. 1a). Genotyping was done by PCR using genomic DNA prepared from yolk sac or tail. *Vangl2P1* (5'-CCATCGATGCACATGTGGTATC-3') and *Vangl2P2* (5'-GACACTGTCTCCATGTCCTTG-3') were used to genotype the *Vangl2<sup>f</sup>* allele and *Vangl2<sup>+</sup>* allele. *Vangl2P1* and *Vangl2P3* (5'-GCTATGACCAGCTACCTGAAGTC-3') were used to genotype the *Vangl2* allele. *Vangl1P1* (5'-CCCTGGCTTTCTTGTGTC-3') and *Vangl1P3* (5'-GGGCTGGCTCTTGAGTCAT-3') were used to genotype the *Vangl1<sup>+</sup>* allele. *Vangl1P1* and *Vangl1P2* (5'-CCTGGGGTTCGTGTCCTACA-3') were used to genotype the *Vangl1<sup>gt</sup>* allele.

### Whole mount in situ hybridization and LacZ staining

Whole mount in situ hybridization and LacZ staining were performed as described previously<sup>32</sup>. Embryos were cryosectioned at 20 µm.

### Immunofluorescence and antibodies

Embryos were fixed in 4% paraformaldehyde for 30 min at 4 °C, washed in PBS and incubated overnight in 30% sucrose/PBS. Embryos were embedded in OCT compound (Tissue-Tek) and sectioned at 14 µm. Sections were permeabilized with 0.5% Triton X-100 in PBS and blocked in 3% bovine serum albumin, 0.1% Triton X-100 in PBS for 1 hour.

Sections were incubated with primary antibodies at 4°C overnight. Secondary antibodies were applied at room temperature for 1 hour. Primary antibodies used in immunofluorescence include mouse anti-Pax6, Pax7, Nkx2.2, Lim1/2, Nkx6.1, Hnf3 $\beta$  and Islet-1 (Developmental Studies Hybridoma Bank, 1:50), rabbit anti-Olig2 (Abcam, 1:100), mouse anti-acetylated-tubulin (Sigma, 1:5,000), goat anti- $\gamma$ -tubulin (Santa Cruz, 1:200), goat anti-Vangl2 (Santa Cruz, 1:100), rabbit anti-Vangl2 (gift of Dr. Mathew Kelley's lab, 1:50) and goat anti-E-Cadherin (R&D, 1:500). Vangl1 antibodies were generated against Vangl1 (1-110 aa) polypeptides fused to glutathione S-transferase (GST) in rabbit. Anti-Vangl1 antibodies were purified using the immunizing peptide coupled to Affigel (Bio-Rad). Alexa Fluor 488 and 568 of donkey anti-mouse, rabbit and goat secondary antibodies were from Invitrogen. Confocal images were acquired using a Zeiss LSM 510 NLO microscope. Excitation wavelengths of 488, 561 and 740 nm were used for detection of secondary antibodies conjugated by Alexa Fluor 488, 568 and DAPI respectively.

### **Whole-mount Immunostaining of cochlea or early E6 – E8.5 embryos**

Dissected cochlea or embryos were fixed in 4% PFA for 30 min at 4 °C. Cochlea or embryos were permeabilized with 0.5% Triton X-100 in PBS for 5 min and blocked in PBS with 3% BSA, 0.1% Triton X-100 at room temperature for 1 hour, followed by overnight incubation with primary antibodies at 4°C. Cochlea or embryos were washed in PBS with 0.1% Triton X-100 at room temperature for 1 h, followed by incubation with fluorescent secondary antibodies and Alexa Fluor-594-conjugated phalloidin (Invitrogen 1:200) at room temperature for 1 hour. The actin-based stereocilium bundles were shown by phalloidin staining whereas the microtubule-based kinocilia or PNC cilia were shown by staining using the mouse anti-acetylated-tubulin antibody (Sigma, 1:5,000). After washed with 0.1% Triton X-100 in PBS at room temperature for 1 hr, cochlea or the PNC region were flat mounted. Confocal images were acquired using a Zeiss LSM510 confocal microscope.

### **Immunoblot Analysis**

Embryonic tissues or cells were lysed in RIPA buffer. Cell lysates were analyzed by Immunoblot. Primary antibodies used include goat anti-Vangl2 (1:2000), rabbit anti-Vangl1(1:1000), rabbit anti-Gli3 (gift from Dr. Susan Mackem, 1:5000).

### **Analysis of calcium signaling**

Embryos were dissected at E7.75–E8.0 and incubated with the fluorescent calcium indicator Fluo-4 (Invitrogen) at a concentration of 10  $\mu$ M in DMEM with 20% FBS for 20 minutes at 37 °C. Embryos were then washed for 3 times in DMEM with 20% FBS and placed node facing up in a silicon chamber under a cover slip. Embryos were incubated at 37 °C for 20 minutes before confocal imaging. Confocal images were acquired using a Zeiss LSM510 confocal microscope. Excitation wavelength of 488 was used to detect Fluo-4. Embryos were genotyped after imaging. Data were analyzed with Image-Pro 6.3 software. High intensity was shown in red and low intensity in blue. Intensity scales of the calcium signal were plotted from left to right side of the node.



## Video Microscopy

Embryos were dissected at E7.75–E8.0 in DMEM with 20% FBS. The node was placed facing down in a chamber with a cover slip bottom. For the nodal flow assay, Fluospheres (Invitrogen) were added into the medium to visualize the direction of nodal flow. Embryos were genotyped after imaging. Wide-field images were collected using a Personal DeltaVision system (Applied Precision Inc, Issaquah, WA, USA) mounted on an inverted Olympus IX71 microscope with an oil immersion PlanApo N 60×/1.42 objective lens. All images were acquired using a CoolSNAP ES2 camera with 2×2 binning and 128 pixels × 128 pixels imaging field. Videos were created using Applied Precision's SoftWoRx software package version 4.0.0. Individual bead movement was traced and shown as a line with different color. The mean cilia beating frequency was calculated from at least 10 nodal cilia of embryos with indicated genotypes at E8.0. Data are means ±SD. The p value is calculated according to the student's t-test.

## Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

## Acknowledgement

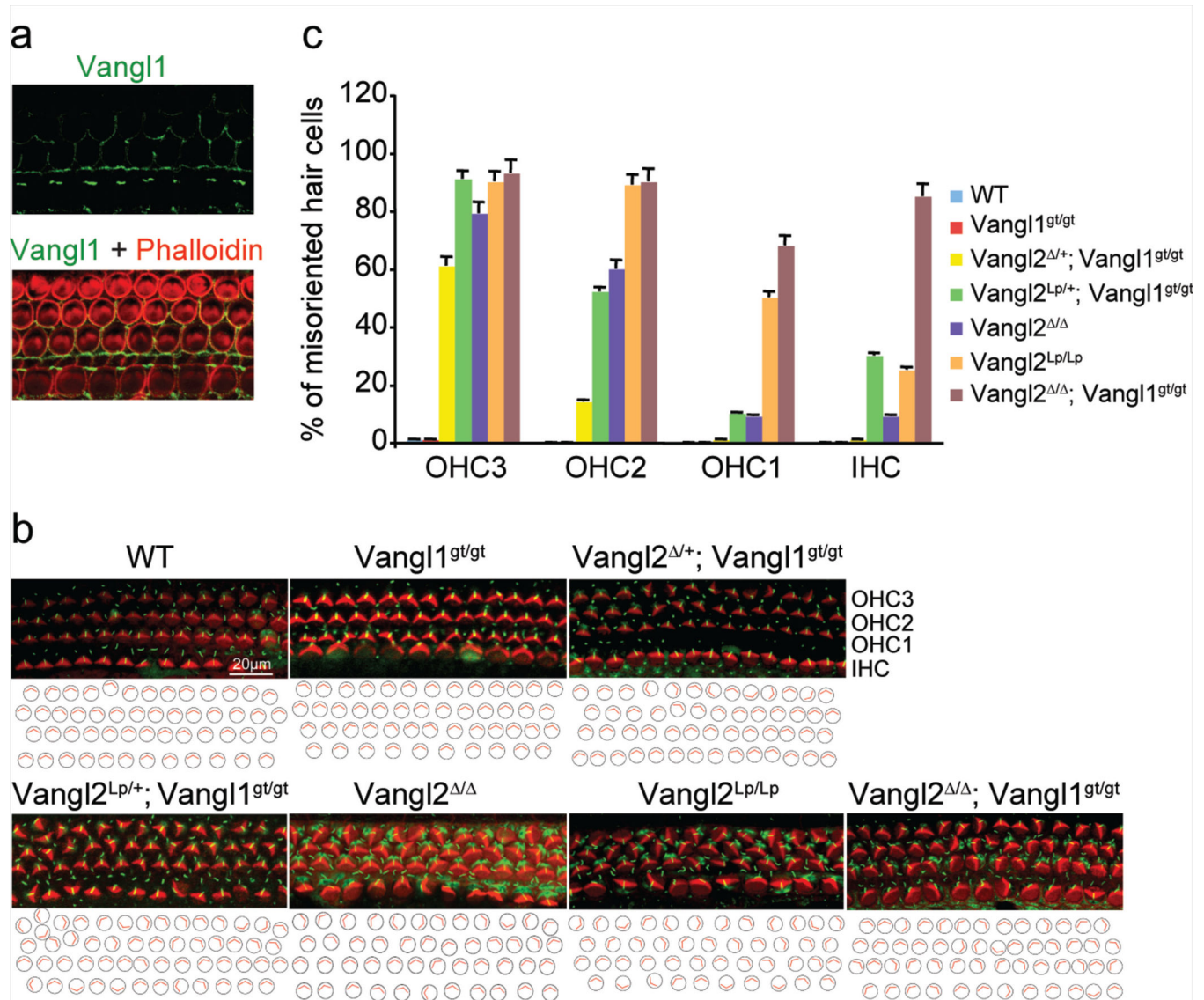
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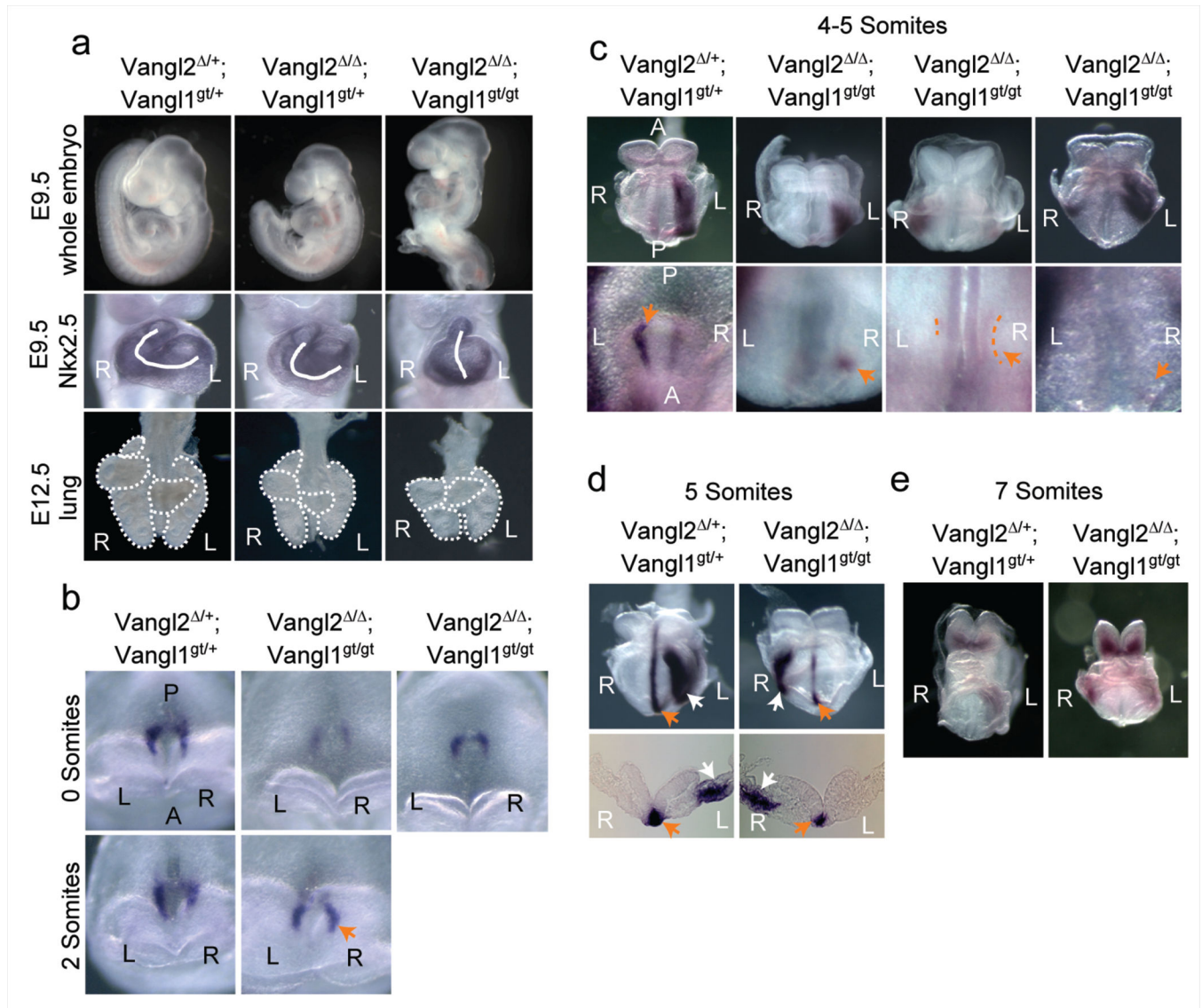
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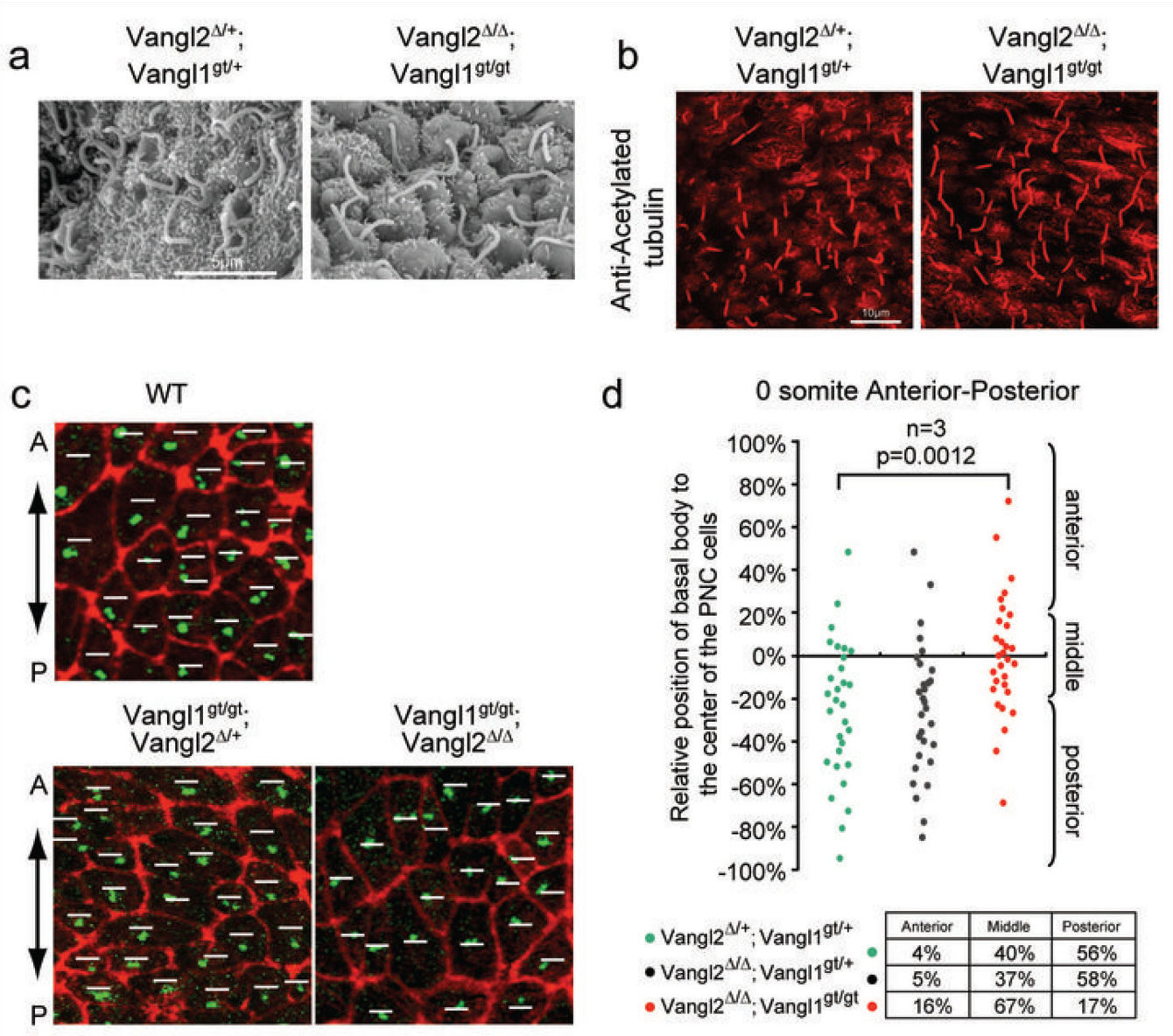
**Figure 1. The *Vangl2*<sup>Δ/+</sup>; *Vangl1*<sup>gt/gt</sup> embryos exhibit the most severe polarity defects**  
**a**, Asymmetrical localization of Vangl1 protein (green) in the sensory hair cells of the *Vangl2*<sup>Δ/+</sup> cochlea at E18.5. **b**, The sensory hair cell polarity at E18.5 is shown by the orientation of actin-based stereocilium bundles (red) and the microtubule-based kinocilia (green). **c**, Quantification of misoriented hair cells in embryos with different genotypes. Data are means ±SD from 3 samples. The orientation defects in the cochlea ranged from few misoriented cells (*Vangl2*<sup>Δ/+</sup>; *Vangl1*<sup>gt/gt</sup>) to several affected cells (*Vangl2*<sup>Δ/+</sup>, *Vangl2*<sup>Lp/+</sup>; *Vangl1*<sup>gt/gt</sup> and *Vangl2*<sup>Lp/Lp</sup>) to almost complete loss of polarity (*Vangl2*<sup>Δ/Δ</sup>; *Vangl1*<sup>gt/gt</sup>).



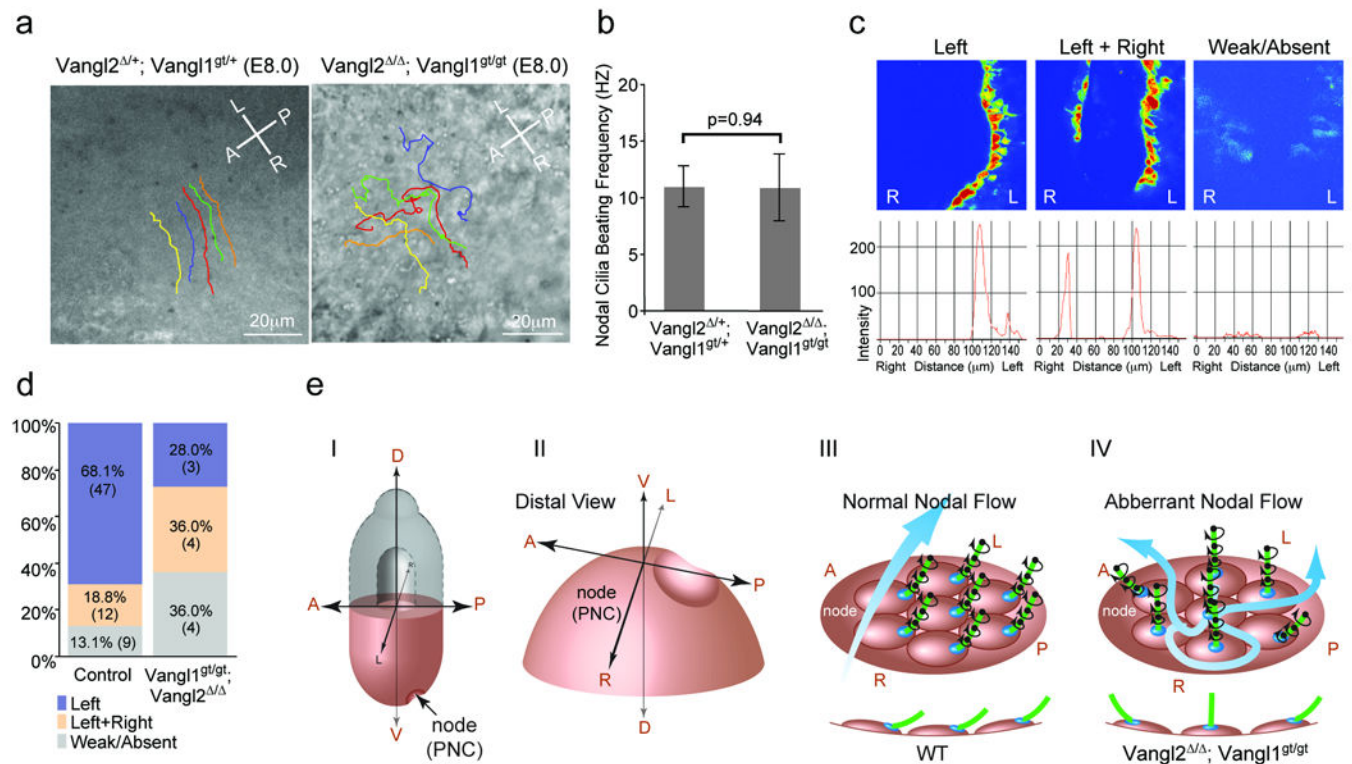
**Figure 2. Laterality defects in the *Vangl2*<sup>-/-</sup>; *Vangl1*<sup>gt/gt</sup> embryos**

**a**, Failure of embryonic turning and heart (marked by *Nkx2.5* expression) looping in the *Vangl2*<sup>-/-</sup>; *Vangl1*<sup>gt/gt</sup> embryos. The *Vangl2*<sup>-/-</sup>; *Vangl1*<sup>gt/gt</sup> lung has three lobes (outlined) instead of four on the right side. R: right; L: left. **b**, Perinodal *Nodal* expression is enhanced on the right side of the *Vangl2*<sup>-/-</sup>; *Vangl1*<sup>gt/gt</sup> embryo (arrow). **c**, Ventral views of *Nodal* expression in LPM (upper panel) and around the node (lower panel, arrows). Weak perinodal *Nodal* expression was outlined. **d**, *Lefty1* expression (orange arrow) on the left side of prospective floor plate and *Lefty2* expression (white arrow) in the LPM were shown in ventral views (upper panel) and transverse sections (lower panel). **e**, Ventral views of *Pitx2* expression.





**Figure 3. Randomized ciliary positioning in the *Vangl2*<sup>Δ/+</sup>; *Vangl1*<sup>gt/gt</sup> embryos**  
**a**, Scanning electron microscopy of the PNC showing normal cilia at E8.0. **b**, Normal cilia (acetylated tubulin, red) in the PNC of the *Vangl2*<sup>Δ/+</sup>; *Vangl1*<sup>gt/gt</sup> embryo at E8.0. **c**, Basal 16 bodies (γ-tubulin, green) and cell boundaries (F-actin, red) of PNC cells at somite 0 stage. White lines indicate the middle position along the A-P axis of each PNC cell. A: anterior; P: posterior. **d**, Relative positions of basal bodies to the midline (along the A-P axis) of the PNC cells were plotted. Posteriorly biased basal body positioning was lost in the *Vangl2*<sup>Δ/+</sup>; *Vangl1*<sup>gt/gt</sup> embryos.



**Figure 4. *Vangl1* and *Vangl2* are required for normal leftward nodal flow**

**a**, Nodal flow shown by tracing latex bead movement across the PNC with lines in different colors (Supplementary movie 1, 2). A: anterior; P: posterior; L: left; R: right. **b**, Ciliary beating was recorded by live imaging (Supplementary movie 3, 4) and the mean beating frequency was calculated from at least 10 nodal cilia of 3 embryos with indicated genotypes. Data are means  $\pm$ SD. **c**, Intracellular calcium release around the node, intensity scales are shown in the lower panels. **d**, Distribution of embryos (percentage and number) according to calcium patterns. **e**, Schematic diagram showing the role of *Vangl1* and *Vangl2* in breaking the L-R symmetry. I. A mouse embryo at E8.0. II. Enlarged distal view of (I). III. Normal nodal flow generated by posteriorly positioned cilia. A side view of ciliated cells are shown in the lower panel. IV. In the *Vangl2*<sup>Δ/+</sup>; *Vangl1*<sup>gt/gt</sup> embryo, randomly positioned cilia around the center of the PNC cells resulted in turbulent nodal flow.