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New insights into regulation and function of planar polarity in the inner ear

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Abstract

Acquisition of cell polarity generates signaling and cytoskeletal asymmetry and thus underpins polarized cell behaviors during tissue morphogenesis. In epithelial tissues, both apical basal polarity and planar polarity, which refers to cell polarization along an axis orthogonal to the apical-basal axis, are essential for epithelial morphogenesis and function. A prime example of epithelial planar polarity can be found in the auditory sensory epithelium (or organ of Corti, OC). Sensory hair cells, the sound receptors, acquire a planar polarized apical cytoskeleton which is uniformely oriented along an axis orthogonal to the longitudinal axis of the cochlear duct. Both cell-intrinsic and tissue-level planar polarity are necessary for proper perception of sound. Here we review recent insights into the novel roles and mechanisms of planar polarity signaling gained from genetic analysis in mice, focusing mainly on the OC but also with some discussions on the vestibular sensory epithelia.

Keywords

Planar Cell Polarity; stereocilia; kinocilium; hair bundle; hair cell; cochlea; deafness

A. Intercellular PCP signaling in cochlear morphogenesis

Located in the cochlea, the OC consists of one row of inner hair cells (IHCs) and three rows of outer hair cells (OHCs) interdigitated by intervening supporting cells (SCs), forming a highly stereotyped cellular mosaic that extend along the longitudinal axis of the cochlea. Each hair cell (HC) has an actin-based stereociliary hair bundle on its apical surface that converts sound vibrations into electrical signals [81, 93]. HCs manifest several salient

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features of planar polarity (Figure 1). First, on a cell-intrinsic level, the hair bundle adopts a polarized structure with rows of stereocilia arranged in a V-shaped staircase pattern and is thus directionally sensitive to mechanical stimuli [81, 93]. On a tissue level, asymmetric hair bundles are uniformly aligned along the medial-lateral (or neural-abneural) axis of the OC, pointing toward the lateral edge of the cochlea duct, and thus respond only to deflections along the medial-lateral axis. The tissue-wide alignment of HC orientation is referred to as planar cell polarity (PCP).

A1. Regulation of cochlear extension and hair cell orientation

The OC develops from a pool of progenitor cells in the prosensory domain expressing Sox2 and p27kip1 (reviewed in [76]). Following cell cycle exit at around embryonic day (E) 14, prosensory cells undergo myosin II-dependent cellular rearrangements resulting in thinning and elongation of the OC [20, 112], and HC differentiation begins around E15 and proceeds in a base-to-apex gradient along the cochlear duct. The first physical evidence of planar polarity at the HC apex is the centrifugal migration of the HC primary cilium, the kinocilium, and its associated basal body towards the lateral pole of the cell [18, 74, 104] (Figure 2A). This is followed by the growth of neighboring microvilli into stereocilia, and nascent V-shaped hair bundles form by E17, with the kinocilium tethered to adjacent stereocilia at the vertex. During the same time period, neighboring HCs adopt a similar orientation to align their kinocilium and nascent hair bundle along the medial-lateral axis. This manifestation of PCP is likely influenced by "tug of war" interactions between HCs and SCs as a result of active cellular movements in the OC. This notion is supported by the identification of three major intercellular signaling pathways that act in concert in both HCs and SCs to coordinate HC orientation and control cellular patterning in the OC.

A1.1 The core PCP pathway—Ground-breaking discoveries in 2003 [12, 77] followed by numerous studies have shown that the evolutionarily conserved core PCP pathway regulates OC patterning by coordinating HC orientation in inner ear sensory epithelia. Mammalian core PCP proteins comprise orthologs of Drosophila Frizzled (Fzd3 and Fzd6) [109], Van Gogh (Vangl1-2) [11, 97, 98], Flamingo (Celsr1-3) [12, 22], Dishevelled (Dvl1-3) [24, 108], Prickle (Pk1–2) [16] and Diego (Ankrd6) [48]. Core PCP mutants have severe neural tube closure defect and often die at birth. In inner ear sensory epithelia, the uniform orientation of hair bundles is disrupted, although the asymmetry of the apical cytoskeleton including the polarized structure of the hair bundle appears unaffected. Consistent with a role in cellular rearrangements during OC extension, the cochlear ducts of core PCP mutants are shorter, with HCs in the apex organized into supernumerary rows (reviewed in [34, 67]). Similar to other systems, core PCP proteins form two complexes asymmetrically localized along the medial-lateral axis in both HCs and SCs to propagate tissue polarity information across the entire OC. Another conserved feature of intercellular PCP signaling is the cell non-autonomous function of transmembrane core PCP proteins. Mosaic analysis in the Drosophila wing epithelium showed that Van Gogh and Frizzled mutant clones can repolarize neighboring wild-type cells, a phenomenon coined "domineering non-autonomy" (reviewed in [34]). A recent study showed that Vangl1/2 deletion in the lateral extrastriolar region of the utricle, one of the vestibular sensory organs, can likewise cause misorientation in neighboring striolar wild-type HCs [98].

Besides conserved functional properties, it is worth noting that there are also intriguing differences that hint at functional divergence of core PCP signaling in the OC. For example, Drosophila Van Gogh and Frizzled proteins are enriched on opposite sides of cell junctions where they form intercellular bridges stabilized by Flamingo on both sides. Intracellularly, Frizzled is thought to regulate Rho family GTPases through membrane recruitment of the cytoplasmic scaffold protein Dishevelled. In the OC, Dvl2/3 proteins are asymmetrically localized along the lateral junctions of OHCs [24, 108]. However, Fz3 and Fz6 are enriched on the medial junctions of HCs [109], where they have been proposed to interact with Vangl2 that is asymmetrically enriched at the lateral junction of the neighboring SC [33]. Dvl2 localization on lateral membranes of HCs has been independently demonstrated by electroporation of a Dvl2-EGFP plasmid in cochlear explants [94], while Fzd6 localization on the medial membranes of HCs was demonstrated using genetic mosaics [109]. Phenotypically, Fz3/Fz6 double mutants showed reversal of hair bundle orientation in IHCs without significantly affecting OHCs, whereas Vangl1/2 double mutants and Dvl compound mutants affect both IHCs and OHCs. [24, 97, 109]. These discrepancies in protein localizations and mutant phenotypes suggest that additional Fzd receptors might be involved in PCP in the OC. Fzd2/7 are likely candidates as they have redundant functions in convergent extension during gastrulation, genetically interact with core PCP genes [114] and are expressed in the cochlea, although their protein localizations are unknown [30].

A1.2 A vertebrate specific, Ptk7-mediated PCP pathway—Protein tyrosine kinase 7 (Ptk7) was identified as a vertebrate-specific regulator of PCP signaling [66]. Similar to core PCP mutants, Ptk7 deficiency causes a wide spectrum of developmental defects, including OHC3 misorientation [66, 84, 113]. Ptk7 encodes a conserved receptor-like pseudo-kinase that lacks endogenous kinase activity. In the developing OC, Ptk7 acts in parallel to the core PCP pathway and is required in SCs for hair bundle orientation [59]. Mechanistically, Ptk7 signals in part through Src family kinases (SFKs) to regulate actomyosin contractility and intercellular tension [1, 59]. Unlike core PCP proteins, Ptk7 is uniformly localized at cell junctions, and thus may act as a permissive factor or partner with other PCP regulators. The ligands for Ptk7 in the OC remain unknown.

A1.3 Nectin- and Notch-mediated cell adhesion signaling—Nectins are Ca²⁺independent IG-like cell-adhesion molecules that interact in trans to induce intercellular adhesion. The cellular mosaic in the OC is mediated by heterotypic interactions between Nectin-1 and -3, which are differentially expressed in HCs and SCs, respectively [105]. Moreover, in *Nectin-3* deficient mice, some HCs are aberrantly in contact with one another, accompanied by hair bundle morphology and orientation defects [28]. The PCP pathways do not overtly affect heterotypic HC-SC adhesion, suggesting that Nectin-mediated cell adhesion signaling is independent of the PCP pathways.

Notch signaling also mediates HC-SC interactions, and some Notch ligand and receptor mutants display patterning defects in line with their role in cell fate regulation, including abnormal contact between HCs. Perhaps as a consequence, these mutants also display hair bundle orientation and shape defects [52, 58, 119].

It is worth noting that defective Nectin and Notch signaling disrupts both hair bundle orientation and shape, possibly indicating a role in both cell-intrinsic and tissue-level planar polarity. How Nectin and Notch signaling regulates HC planar polarity at a molecular level remains to be determined, but likely involves cross-talk with neighboring cells and the recruitement of intrinsic polarity modules in HCs (see section B2).

A1.4 HC reorientation during normal and abnormal development.—

Presumably because the cuticular plate that anchors the hair bundle is initially loosely organized [85], the initial orientation of nascent hair bundles is imperfect and undergoes continuous refinement of up to 30 degrees during embryonic and postnatal development. This process might involve Wnt signaling based on in vitro evidence [13]. In principle, core PCP proteins might participate in normal refinement, as asymmetric core PCP protein localization is maintained after birth [11, 95]. However, OHC3 hair bundles in a viable conditional knockout (cKO) for *Vangl2* mediated by *Pax2-Cre* were able to correct their initial misorientation and reorient by as much as ~180 degrees at postnatal stages, except in the apex region of the cochlea [11]. As asymmetric core PCP complexes were disrupted in the *Vangl2* cKO cochlea, the core PCP pathway is dispensable or acts redundantly for the correction process. Consequently, it is likely that a still undefined molecular mechanism is involved in postnatal HC reorientation.

A2. Additional roles of PCP signaling in the cochlea

A2.1 Roles in SC cytoarchitecture and type II SGN afferent guidance.—SCs in the OC not only propagate apical PCP information to orient neighboring HCs, but also have a planar polarized cytoarchitecture themselves. There are several types of SCs in the OC, including the inner phalangeal cells, the inner and outer pillar cells and the Deiters' cells. During cochlear extension, the OC thins from a pseudo-stratified epithelium to a two-layer stratified epithelium with the nuclei of SCs below those of HCs (reviewed in [76]). On the luminal surface, HCs are separated from one another by phalangeal processes projected from the cell body of SCs. Interestingly, the phalangeal processes of outer pillar cells and Deiters' cells show a slant toward the cochlear apex, contacting OHCs located several cell bodies away along the longitudinal axis [47] (Figure 3). In *Vang12* cKO mutants, the apically-directed slant of the Deiters' phalangeal processes was lost, accompanied by disorganized HC-SC contacts [11]. Core PCP signaling is therefore involved in the longitudinal mechanical coupling between Deiters' cells and OHCs, a property essential for OHC electromotility and cochlear amplification [80, 115]. Accordingly, *Vang12* cKO mutants had hearing deficits as shown by elevated ABR thresholds [11].

Besides regulating their cytoarchitecture, PCP signaling in SCs is increasingly recognized to polarize other processes during cochlear development. Early studies in cochlear explants had shown that cochlear extension is unidirectional from the base to the apex and requires myosin II activity [108, 112]. Recent time-lapse imaging experiments showed that SCs, but not HCs, extend dynamic protusions along the basement membrane and migrate toward the cochlear apex, thus providing a cellular basis for the unidirectional extension [20]. Interestingly, Vangl2 is asymmetrically localized at the SC-SC baso-lateral contacts along the longitudinal axis [32], parallel to the trajectory of SC movements and orthogonal to its

medial-lateral polarization at apical HC-SC junctions, raising the possibility that SC directional migration is regulated by core PCP signaling. While this remains to be determined, a recent study uncovered a novel, non-autonomous role of PCP signaling in SCs to guide type II spiral ganglion neurons (SGN) afferents [32].

Each SGN extends a peripheral process toward the OC and a central process that branches into the cochlear nuclei. The most abundant type I SGNs form ribbon-like synaptic contacts exclusively at the single row of IHCs. In the remaining 5–10% type II SGNs, the peripheral process crosses the tunnel of Corti into the Deiters' cell region and turns towards the cochlear base, extending several hundred micrometers to innervate several OHCs [2, 10]. Unlike type I afferents, type II do not encode sound information but are thought to detect acoustic trauma [64]. When Vangl2 was deleted in the cochlear epithelium using Emx2-Cre, type II SGN afferent turning appeared randomized, and about 50% of fibers turned erroneously towards the cochlear apex. By contrast, deletion of Vangl2 in SGNs did not cause turning defects, establishing its non-autonomous role in SCs. Similar but milder SGN afferent turning defects were recorded in Fzd3 and Celsr1 null mutants. Vangl2 is asymmetrically enriched on the SC basolateral surface facing towards the cochlear apex, where it is proposed to recruit and stabilize type II growth cones turning towards the base. It remains unclear whether complementary core PCP complexes mediate growth coneepithelial substrate interactions, as reported for anterior guidance of spinal commissural axons and migration of hindbrain facial branchiomotor neurons (reviewed in [14]).

A2.2. PCP signaling in Kolliker's organ orients the apical extracellular matrix

-Outside the OC, PCP signaling in a subset of cells in Kolliker's organ was recently shown to orient the overlying tectorial membrane. The tectorial membrane is a specialized apical extracellular matrix structure secreted by epithelial cells in the ventral cochlear duct, and extending over the OC to contact the hair bundles of OHCs (reviewed in [29]). Collagen fibrils in the tectorial membrane are oriented radially along the medial-lateral axis with a slant towards the cochlear apex, a spatial organization participating in auditory function. Interestingly, Pk2 is asymmetrically localized at apical junctions along the longitudinal axis in Kolliker's organ, orthogonal to its enrichment at apical SC-HC junctions, but parallel to Vangl2 enrichment at basolateral SC-SC junctions. Intriguingly, Pk2 polarized enrichment is only observed in a band of cells whose radial junctions are aligned with the same apicallydirectly slant as the overlying collagen fibrils. Moreover, the apically-directed slant of collagen fibrils, as well as asymmetric Pk2 localization, were abolished in Vangl2 and Ptk7 cKO mutants, indicating a requirement for PCP signaling in the planar orientation of collagen fibrils within the tectorial membrane [35]. In summary, core PCP proteins are widely expressed in the developing cochlear epithelium, and their planar polarized distribution serves as a "compass" to orient a range of polarized cellular processes. At junctions between non-sensory cells, core PCP proteins are polarized along the longitudinal axis, orthogonal to the medial-lateral (radial) axis of PCP at HC-SC junctions. In every case, the observed PCP features correspond to the local vector of tissue polarity.

B. Establishment of hair cell-intrinsic planar polarity

Even the most severe PCP mutants [77, 97, 109] retain a normal planar polarization of the apical cytoskeleton in individual HCs at birth. This includes a peripheral basal body/ kinocilium situated at the vertex of the V-shaped hair bundle edge, a heart-shaped cuticular plate stabilizing the hair bundle, and apical microtubules (MTs) confined laterally by the cuticular plate (Figure 1B). Therefore, intercellular PCP signaling is not required for planar polarization of individual HCs. In the last decade, several signaling modules have been identified and likely act in concert to control planar asymmetry of the cytoskeleton in single HCs.

B1. Influence of the basal body and kinocilium on hair bundle morphogenesis

Hair bundle development is both temporally and spatially coupled to the lateral placement of the kinocilium and its associated basal body. The kinocilium recedes around postnatal day (P) 9 when hair bundle maturation is close to complete, and is absent in mature auditory HCs.

B1.1. A microtubule-based mechanism for basal body/kinocilium placement

-The lateral migration of the basal body is required to establish the asymmetric shape of the hair bundle, as illustrated by the circular bundles observed in a small fraction of Ift88 mutant HCs where the basal body failed to migrate [49]. While the mechanisms that trigger the early centrifugal migration of the kinocilium are still under debate (see sections B3, F2), accumulating evidence indicates that the final lateral basal body/kinocilium placement is controlled by a MT-based machinery (Figure 1B). The basal body anchors both the kinocilium and the apical MTs, and its placement was proposed to be mediated by interactions between the dynamic plus ends of apical MTs and the HC cortex at the lateral junction. First, the plus end tracking protein EB1 is enriched near the lateral HC junction [26]. Second, proper basal body placement requires both plus-end and minus-end directed MT motors. In addition to its role in intraflagellar transport, the kinesin-II subunits Kif3a mediates coupling between the hair bundle and the basal body as well as cortical Rac1 GTPase signaling via MT-mediated intracellular transport [96]. Deletion of Lis1, an activator of cytoplasmic dynein, disrupted MT organization and cortical attachment, leading to basal body placement and hair bundle defects [95]. These results indicate that dynein-mediated cargo transport and/or pulling forces on apical MTs play a critical function in basal body/ kinocilium placement. In contrast, early basal body migration was unaffected by Kif3a or Lis1 inactivation. Lastly, several planar polarized signaling modules appear to stabilize MTcell cortex interactions to anchor the basal body at the lateral pole, as described in Section B2.

B1.2 A role for the kinocilium and Usher syndrome type I (Ush1) proteins-

As a signaling organelle physically attached to the hair bundle (Figure 1B, 2A) the kinocilium may play both a signaling and mechanical role during hair bundle morphogenesis. Its importance was first demonstrated by inactivating genes required for intraflagellar transport, such as Ift88 and Kif3a, thereby preventing ciliogenesis [49, 96]. The mechanical role of the kinocilium was then demonstrated by specifically ablating the

CD2 isoform of Pcdh15, a component of the kinociliary links that tether the kinocilium to neighboring stereocilia [111]. In the absence of the kinocilium or kinociliary links, hair bundles lost their chevron shape and had a flattened morphology. On rare occasions, circular hair bundles either with a central or peripheral basal body were also observed [49, 111]. However, most HCs still formed a planar polarized hair bundle in the absence of a kinocilium, even when the remaining basal body was spatially uncoupled from the forming hair bundle [49, 96]. This challenges the popular but unproven assumption that following its migration, the kinocilium somehow prompts neighboring microvilli to grow into stereocilia. Interestingly, core PCP protein localization was largely normal in *Ift88, Kif3a* and *Pcdh15-CD2* mutants, suggesting that core PCP signaling operates independently of the kinocilium to position stereocilia. Of note, trafficking of Pcdh15-CD2 to the kinocilium was recently shown to be mediated by DAB2/clathrin/Ift-B transport particles and regulated by ciliary Fgfr1 signaling, providing one of the first examples of signaling functions of the kinocilium in HC planar polarity [41].

Ciliopathies are developmental disorders generally associated with defects in primary cilia. Accordingly, a number of mouse models for human ciliopathies exhibit a variable spectrum of hair bundle and kinocilium defects, including *Mkks/Bbs6*, *Bbs4* and *Alms1* mutants [26, 42, 89]. Importantly however, some ciliopathy proteins were found to have non-ciliary functions that also influence HC planar polarity. For example, unique among Ift and Bbs proteins, Bbs8 and Ift20 were shown to interact with the core PCP protein Vangl2 and to regulate its polarized localization, presumably through MT-based trafficking [73]. In addition, many ciliopathy gene products localize to or around the basal body, and may therefore regulate basal body/centrosome-associated functions such as proteasomal degradation and endosomal trafficking [31, 39].

Besides Pcdh15, other Usher syndrome 1 (Ush1) proteins, including Cdh23, the scaffold proteins harmonin and SANS, myosin VIIa, and the calcium and integrin-binding protein CIB2, are also important for hair bundle structure and kinocilium positioning. In addition to kinociliary links, heterophilic Pcdh15 and Cdh23 adhesion complexes form various interstereocilia links that maintain hair bundle cohesion during HC differentiation [60, 111]. Therefore, it is tempting to speculate that kinocilium and hair bundle positioning are regulated in a reciprocal manner; as a cohesive unit, the hair bundle may also regulate the final position of the basal body/kinocilium by pulling on kinociliary links. In Ush1 mutants, this reciprocal regulation is disrupted, leading to planar polarity defects.

B2. Several signaling modules act in concert during HC planar polarization.

While ciliary and Ush1 proteins influence the structural polarity of the hair bundle, they are not themselves planar polarized in HCs. By contrast, several signaling modules with a planar polarized distribution have been identified and likely play a more instructive and broader role in establishing HC cytoskeleton planar polarity, as discussed below.

B2.1 The Par3-Rac1-Pak module—The small GTPase Rac1 and its effector Pak were among the first HC-intrinsic planar polarity regulators identified [36]. Cortical Rac-Pak signaling activity is asymmetrically enriched along the lateral HC junction, where it was

proposed to preferentially stabilize interactions between the apical MTs and the HC junction, thereby controlling kinocilium placement and hair bundle shape [36, 96].

How is the polarized domain of Rac signaling established? A recent study revealed a critical function of the PDZ-domain scaffold protein Par3, a conserved regulator of cell polarity [57, 69]. In epithelial cells, Par3 helps define the apical domain by forming a tripartite complex with Par6 and the kinase aPKC. The complex occupies apical cell junctions, while aPKC-Par6 but not Par3 are also enriched at the apical membrane [37, 38, 71, 78]. In HCs, Par6 and aPKC are planar polarized on the medial side while Par3 is polarized on the lateral side at the onset of hair bundle formation [26, 57, 102]. Par3 was shown to have both cell-autonomous and non-autonomous functions and to stimulate Rac1 signaling in both HCs and SCs through interactions with the guanine nucleotide exchange factors (GEFs) Tiam1 and Trio. Strikingly, expression of constitutively active Rac1 rescued most, if not all, planar polarity defects in *Par3* cKO, arguing that the key function of Par3 in HC planar polarity is to regulate Rac1 signaling [57]. As a scaffold protein, Par3 can interact with multiple planar polarity regulators and thereby likely participates in signal integration (see Section C).

B2.2. The cdc42-aPKC-Par6 module—Cdc42 is another member of the Rho family GTPases with conserved functions in cell polarity and cytoskeletal regulation. Inactivation of Cdc42 starting at E13.5 in the OC resulted in defects in OHC patterning, MT organization and hair bundle orientation, indicating a requirement of Cdc42 for HC planar polarity. In *Cdc42* mutants, the asymmetric localization of aPKC, a known effector of Cdc42, was misoriented in register with the hair bundle, while asymmetric localization of core PCP proteins was unaffected [54]. Although their function in HCs has not been reported, Par6 and aPKC have a similar medial localization in HCs [26] and thus likely form a Cdc42/ aPKC/Par6 module.

B2.3. The Insc-Gpsm2-Gai module establishes a molecular blueprint for hair cell planar polarity—Gpsm2 (also known as LGN) stands for G protein signaling modulator 2, and Gai is the inhibitory alpha subunit of the heterotrimeric G protein complex. Along with partners, GPSM2-Gai serves as an evolutionary conserved cortical complex polarizing pulling forces that orient the mitotic spindle during cell division [79]. Mechanistically, Gai binds Gpsm2 and is myristoylated to anchor the complex at the plasma membrane, while Gpsm2 recruits other partners including Numa (Fig. 4A). Numa in turn recruit effectors including the dynein motor to pull on astral MTs. The net result is control over the spindle axis, daughter cell placement and thus tissue architecture. Biochemically, Gpsm2 is thought to act as a G protein dissociation inhibitor (GDI), sequestering Gai in a GDP-bound form that is usually considered to be an inactive signaling state.

Gpsm2-Gai is rapidly planar polarized at the lateral aspect of post-mitotic HCs [6, 26, 102] (Figure. 3). This behavior roughly coincides in time with the early lateral migration of the basal body/kinocilium. However these processes appear to be independent, as the position of the kinocilium and the Gpsm2-Gai crescent are initially only loosely correlated. Gpsm2-Gai and the Gpsm2 binding partner Insc occupy the "bare zone", an expanding lateral region of apical membrane devoid of microvilli that appears after the migration of the basal body [102] (Figure 2). There, Insc-Gpsm2-Gai antagonizes aPKC that initially spans the

entire apical membrane, and appear to exclude microvilli previously covering the entire apical surface. Consistent with the notion that segregation of polarity proteins at the apical membrane acts as a blueprint to pattern the cytoskeleton at large, stereocilia, the kinocilium, the cuticular plate and apical MTs are abnormally distributed in *Gpsm2* mutants [26, 102].

To circumvent functional redundancy among Gai proteins, Pertussis toxin (PTX) was used to globally downregulate Gai activity in HCs. Surprisingly, treating cochlear explants with PTX [26] or expressing PTX catalytic subunit (PTXa) in HCs using transgenic approaches [102, 103] produced cytoskeleton placement defects as in *Gpsm2* mutants, but caused also a unique and severe PCP phenotype (Figure 1C). Specifically, PTXa-expressing OHCs showed graded orientation reversal by row where OHC1 were cleanly reversed, OHC2 were variably oriented medially and OHC3 were variably oriented laterally. These results suggest that Gai is involved independently from Gpsm2 in the Emx2-directed switch of HC orientation recently uncovered in vestibular organs [46] (see section D1).

It is unclear how the polarized localization of Gpsm2-Gai is initially established. Interestingly, a recent study identified SorCS2, a member of the Vps10 family of sorting receptors involved in protein trafficking, as essential for Gpsm2-Gai localization at the bare zone. Supporting the blueprint model, the medial aPKC domain was expanded in *SorCS2* mutants, coinciding with severe hair bundle defects [27]. New studies are needed to identify other components of the trafficking machinery underlying cell-intrinsic planar polarity.

B3. Common signaling modules polarize diverse cellular processes

Regulation of HC intrinsic planar polarity by several components of the mitotic spindle orientation machinery has suggested mechanistic parallels between these processes (Figure 4A). Based on their shared lateral bias at the HC apex, it was proposed that Insc-Gpsm2-Gai and Par3, similar to their role as cortical landmarks in mitotic progenitors, might recruit partners including Lis1/dynein to exert pulling forces on centriolar MTs. By analogy with mitotic spindle orientation, this mechanism could trigger the early lateral migration of the basal body/kinocilium to break radial symmetry at the HC apex, and/or regulate their placement at later stages of HC differentiation [26, 95]. These compelling hypotheses are discussed in Future Directions (see section F2, F3).

Another polarized cellular process that shares striking parallels with HC planar polarity is directional migration/chemotaxis (Figure 4C). Persistent directional migration towards a source of chemoattractant requires a protrusive leading edge and a retracting trailing edge that define the front-to-rear polarity axis. During directed migration, MTs are oriented and show persistent growth towards the leading edge. MTs and the dynein motor were shown to regulate direction persistence by maintaining front-rear polarity [21, 117]. In addition to a structural role, MT trafficking can deliver a number of GEFs to activate Rac and Cdc42 at the leading edge. Reciprocally, the Par3 complex links Cdc42 and Rac signaling to MT regulation (reviewed in [25]). The Insc-Gpsm2-Gai module is also enriched at the leading edge and can recruit Par3-aPKC via Insc during neutrophil chemotaxis [50](Figure 4C). Importantly, in that case, the Gpsm2-Gai complex regulates actin dynamics, underscoring that Gpsm2-Gai has MT-independent functions, as also evident during stereocilia elongation (see section E1). While "canonical" G protein receptor-coupled (GPCR) signaling is well-

known to detect chemoattractants and promote cell motility (reviewed in [51]), noncanonical signaling involving Gpsm2-Gai was proposed to stabilize the leading edge and thus ensure sustained directionality in neutrophils (Figure 4C). It is interesting to speculate that core PCP signaling in HCs may play an analogous role as chemokine gradients in migrating cells to orient cell-intrinsic polarity modules in the tissue.

C. Integration of intercellular and cell-intrinsic polarity signaling

In core PCP mutants, intrinsic polarity modules including planar polarized Rac activity and Insc-Gpsm2-Gai domains are intact in individual HCs, but misoriented at the tissue level [26, 36, 102]. Reciprocally, core PCP protein distribution is also unaffected in *Gpsm2* or Gai mutants [26, 102]. Furthermore, Gpsm2-Gai is only planar polarized in HCs, while core PCP proteins are polarized in both HCs and SCs to propagate polarity information in the OC. Finally, Gpsm2-Gai are planar polarized at the apical membrane above tight junctions [94], whereas core PCP proteins are planar polarized at cell-cell contacts below tight junctions [26], two distinct yet abutting subcellular domains (Figure 5). Taken together, intercellular (PCP) and cell-intrinsic planar polarity appear to function independently, but must be coordinated during development to ensure that asymmetric hair bundles are all oriented laterally. A strategy to functionally bridge the two systems may employ proteins that can physically bind both junctional core PCP proteins and cell-intrinsic players at the apical membrane. Two candidates fulfilling some of these requirements have been reported to date.

Daple (Dishevelled-associating protein with a high frequency of leucines), also known as Ccdc88c, is a regulator of Wnt signaling that can bind Dvl with its C-terminal PDZ-binding domain [83] and bind and regulate Gai, acting as a non-receptor GEF in vitro [3]. Interestingly, Daple mutant HCs show severe apical cytoskeleton defects including a grossly mishappen hair bundle and randomized kinocilium position [94]. This complex phenotype can be interpreted as a mixture of PCP and planar asymmetry defects. Early kinocilium migration defects similar to those of *Vangl2* mutants [77] suggest disrupted PCP signaling. By contrast, complementary but aberrant Gpsm2-Gai and aPKC domains at the apex suggest defects in the apical blueprint. At the molecular level, Daple is planar polarized laterally near the HC tight junction, sitting at the interface between Gai at the bare zone and Dvl2 at the adherens junction [94] (Figure 5). Although the exact relationship between Dvl2, Daple and Gai remains to be solved, they show co-dependence for their normal enrichment and are all spontaneously planar polarized and spatially coordinated in ectopic HC-like cells induced in Kolliker organ [94]. Together, this suggests that Daple could bridge intercellular and cell-intrinsic planar polarity modules by binding to Dvl2 and Gai.

Par3 is another candidate signal integrator of planar polarity. Par3 is enriched at the lateral HC junction similar to Daple and Dvl2, and was previously found to be a binding partner of Nectins [57, 94, 101]. Par3 is also enriched at low levels at the bare zone along with Insc-Gpsm2-Gai [102], and can bind Insc in the context of spindle orientation [79]. However, Par3 is not required for asymmetric Gpsm2-Gai localization in HCs. Instead, Par3 regulates several GEFs that activate Rac signaling in both HCs and SCs, thereby likely influencing both intercellular and cell-intrinsic planar polarity [57]. Interestingly, Par3 was found to bind

Daple in a yeast assay and their early distribution is very similar [94], suggesting potential crosstalk between the Dvl2-Daple-Gai-Gpsm2 and Par3-Rac-Pak modules (Figure 4). Moreover, Par3 may dynamically interact with aPKC-Par6 to establish a crosstalk between the lateral Par3-Rac-Pak and medial Cdc42-aPKC-Par6 modules.

In addition to physical interactions between PCP and cell intrinsic polarity proteins, signal integration may also occur through mechanical cues, as both PCP and front-rear polarity can be reoriented by tissue and cell mechanics [56, 68]. In support of this notion, the Par3-Rac1 [23] signaling axis has been shown to reinforce asymmetric core PCP protein localization in the OC, likely indirectly through cytoskeletal remodeling [36, 57].

D. Transcriptional regulation of hair cell planar polarity

D1. Transcription factors known to be required for hair cell planar polarity

As a component of the HC differentiation program, planar polarization is probably under the control of transcription factors that drive OC development. Although transcriptomic analysis of FACS-purified HCs and other cochlear cell types have been performed [8, 23, 91], presently very little is known about specific regulation of planar polarity genes by key TFs that control OC differentiation. Through genetic analyses, two TFs, namely Six1 and Emx2, have been shown to influence HC planar polarity. The homeodomain TF Six1 and its cofactor Eya1 play essential roles of in prosensory cell differentiation. Deletion of Six1 in OC progenitor cells disrupted the OC cellular mosaic and HC planar polarity [118], although it is unclear whether Six1 directly regulates planar polarity genes, or whether polarity defects are an indirect effect of impaired OC differentiation.

Of particular interest, the homeobox transcription factor Emx2 has emerged as a global regulator of HC orientation in the vestibular system. HCs in the crista ampullae detect angular head acceleration and adopt a uniform orientation matching the spatial trajectory of the associated semicircular canal. HCs in the macular organs (the utricle and saccule in mammals) detect linear head acceleration and gravity, and adopt a radiating range of orientations where HCs are aligned locally. PCP in the maculae also includes an abrupt transition that separates each organ in two distinct HC populations with opposite orientations. HCs across this virtual line of polarity reversal (LPR) thus show opposite responses to the same head movement (reviewed in [15]).

Interestingly, core PCP proteins are not reversed in their asymmetric enrichment across the LPR, demonstrating that the PCP vector of tissue polarity is not sufficient to define HC orientation [16]. Expression of the transcription factor Emx2 was recently shown to be limited to one orientation domain, and to be necessary and sufficient to reverse HC orientation [40, 46]. Remarkably, Emx2 restriction to one orientation domain is conserved in chicken macular organs and in the zebrafish lateral line, where HCs grouped in neuromasts respond to bi-directional water movement. In zebrafish, as in the mouse, Emx2 is necessary and sufficient to reverse HC orientation.

Presumably, HCs respond to Emx2 by reversing how cell-intrinsic planar polarity modules associate to invariant organ-level core PCP patterning at cell junctions. Effectors in the

Emx2 transcriptional program remain to be identified, but Gai activity appears to be required. PTXa expression can indeed reverse HC orientation in the lateral utricle, and thus phenocopy the loss of Emx2 [46]. As PTXa expression can prevent HC reversal induced by ectopic Emx2 expression, Gai clearly functions downstream of Emx2. Interestingly, the auditory epithelium is Emx2-positive but cristae are Emx2-negative [46], and PTXa was previously shown to provoke the graded orientation reversal of OHCs in the cochlea (see section B2.3) [26, 102]. These results suggest that Emx2-based patterning might be an evolutionary novelty used to reverse a "ground" state of HC orientation present in ancestral vestibular organs. Emx2 is required for OHC differentiation however [40], precluding direct testing of this hypothesis in the cochlea.

D2. Approaches to identify TF target genes crucial for cochlear morphogenesis

Along with many other TFs, Six1 and Emx2 form an intricate and dynamic gene regulatory network for hair cell specification, differentiation and survival. Only a handful of direct target genes are known, in part because popular techniques for genomic mapping of TF binding sites such as chromatin immunoprecipitation (ChIP)-seq and ATAC-seq have not been widely applied to cochlear tissues due to the small number of HCs in the cochlea [44, 118]. To overcome this limitation, investigators have used a multi-pronged approach by identifying differentially expressed gene in loss- and gain-of function TF mutants, and cross-referencing with available ChIP-seq and TF-binding site datasets in other tissues and cell lines [8, 23]. With the rapid advance of next-generation sequencing technologies, genomewide mapping of TF targets in space and time during cochlear morphogenesis will be achievable in the near future, which will shed light on the mechanisms underlying global patterns of planar polarity in inner ear sensory epithelia.

E. Novel roles of cell-intrinsic planar polarity factors in hair cells

E1. Role of Gpsm2-Gai in stereocilia elongation and hair bundle maturation

While Insc-Gpsm2-Gai proteins are first planar polarized at the bare zone (Figure 2; see section B2.3), Gpsm2-Gai, but not Insc, is then also enriched in the hair bundle during embryogenesis already [99, 103]. At postnatal stages, Gpsm2-Gai is enriched at the distal tip of stereocilia in the first, tallest row. Enrichment in the bundle is first observed in central stereocilia adjacent to the kinocilium, and then spread to more peripheral ones. Broad and premature Gpsm2-Gai enrichment at stereocilia tips was observed in *Insc* and *Gpsm2* mutants, with a concomitant loss of enrichment at the bare zone. This imbalance in protein distribution suggests that protein anchoring at the bare zone prevents premature trafficking to stereocilia tips, establishing a relationship between these subcellular compartments [103]. Based also on the close segregation of Gpsm2-Gai at the bare zone with the base of row 1 stereocilia, it was proposed that intrinsic planar polarity information is used to establish the characteristic asymmetric architecture of vertebrate hair bundles in the apicobasal axis, with stereocilia rows of graded heights.

This idea is attractive because in absence of Gpsm2 or Gai function, postnatal stereocilia elongation is compromised, a phenotype most striking in IHC stereocilia [72, 103]. IHC hair bundles are largely flat and maintain supernumerary rows, very reminiscent of defects

reported previously for *Myo15a*, *Whrn* and *Eps8* mutants [70, 75, 86, 116] (Figure 6). Myo15a, Whrn and Eps8 localize to stereocilia tips and were proposed to promote actin incorporation to stimulate postnatal stereocilium elongation [5, 17, 70, 90]. In fact, genetic interaction studies demonstrated that Gpsm2-Gai and Myo15a-Whrn function in the same elongation/maturation pathway [99]. Stereocilia stunting is a likely cellular etiology for profound congenital hearing loss reported when Gpsm2-Gai function or trafficking is defective in humans or mouse models [19, 27, 72, 103, 106].

Teasing apart protein localization in normal and mutant contexts revealed that Myo15a-Eps8 forms an early complex at stereocilia tips, with a later Whrn-Gpsm2-Gai module added in first row stereocilia only [99]. Myo15a and Whrn are required for Gpsm2-Gai localization at tips, and Whrn can bind both Gpsm2 and Gai [72, 99]. This suggests that Whrn acts as an adapter to include Gpsm2-Gai to Myo15a cargo [72]. In contrast, Myo15a-Eps8 were still trafficked to tips in absence of Gpsm2-Gai function, but protein amounts never built up to high levels observed in normal row 1 [99]. In addition, Myo15a-Eps8 lost its characteristic graded enrichment mirroring stereocilia height across rows. As protein markers for shorter rows were found in the first row when Gpsm2-Gai function was disrupted, and reciprocally, markers for the first row (Gpsm2-Gai themselves) were found in all rows in PTXa-expressing HCs, mutant stereocilia appear to maintain a generic molecular and morphological identity. The Whrn-Gpsm2-Gai module thus appears to be added to an ubiquitous Myo15a-Eps8 stereocilia tip complex in order to specify the unique tallest identify of row 1 stereocilia, and to promote differential row identity in the planar-asymmetric hair bundle [99].

E2. Role of Gpsm2-Gai and Emx2 in hair cell synaptic properties

Each IHC forms ribbon synapses with multiple SGN type I afferent fibers to transmit auditory signals to the brain. Although they innervate the same sound receptor, these afferents have different firing properties and contact heterogeneous active zones on the IHC body, which helps explain the astounding range of sound pressure processed by the inner ear. Heterogeneity is spatial in nature and potentially includes planar polarized features. For example, ribbon synapses are larger on the medial than the lateral side of the cell [61, 63]. This coincides with graded electrophysiological synaptic properties, and could potentially instruct well-established graded afferent behaviors at the post-synaptic level [62]. Interestingly, Gpsm2 or Goi3 inactivation and PTXa expression collapsed the medio-lateral gradient of ribbon synapse size while maintaining overall ribbon size and number [43] (Figure 6). In addition, PTXa provoked enlarged pre- and post-synaptic clusters, increased Ca2+ influx, and the collapse of the spatial gradient of maximal synaptic Ca2+ influx. These results suggest that the Gpsm2-Gai complex polarizes not only the cytoskeleton at the HC apex, but also the spatial organization and the properties of ribbon synapses at the basal end of the IHC. An independent study showed that Gai2 and Gai3 are redundantly required for the normal maturation of the ribbon synapses in IHCs [4]. More work is needed to determine how Gpsm2-Gai influences the assembly and function of heterogeneous active zones. It would also be interesting to test a potential role of PCP proteins.

A link between cell-intrinsic planar polarity and synapse development was also found in HCs in the zebrafish neuromasts. Single neuromast afferents neurons exhibit directional selectivity, contacting only HCs that share the same orientation. Remarkably, recent work demonstrated that a single neuron either contacts Emx2-positive or Emx2-negative HCs [45, 65]. *Emx2* inactivation or Emx2 ectopic expression in HCs led to an increase or decrease in the number of HCs contacted, depending on the afferent selectivity. As Emx2-based selectivity was retained in a core PCP mutant with randomized hair bundle orientation, afferent synaptic choices strictly reflect Emx2 expression in HCs, and not HC orientation. It remains unknown however how navigating fibers discriminate between Emx2-positive or Emx2-negative HCs to establish synaptic contacts. In conclusion, as Gpsm2-Gai, Emx2 influences both the polarization of the apical cytoskeleton and basal synaptic properties in HCs.

F. Future Directions

Despite significant advances in the last decade, our understanding of the mechanisms of planar polarity and their functions in the inner ear is still fragmentary. Below, we discuss several incompletely understood areas that warrant future investigations.

F1. Unsolved questions in intercellular PCP signaling

Despite conservation of the principle features of core PCP signaling, a number of lingering questions remain about intercellular PCP signaling in the inner ear. First, the puzzling opposite localization of Fzd3/6 and Dvl2/3 in HCs suggest that additional Frizzled proteins may regulate Dvl at the lateral junction, and/or that Fzd3/6 may partner with Dvl1 or function independently of Dvl. Second, it remains unknown how the vector of tissue polarity changes in different parts of the cochlear epithelium. While tissue mechanics likely plays a role [53], junctional rearrangement and actomyosin dynamics have not been analyzed outside the OC. It also remains unclear how the uniformly distributed Ptk7 protein can generate signaling and cytoskeletal asymmetry. To solve these questions, it would be important to catalog the precise localization of all core PCP proteins in the auditory and vestibular epithelia. This is a challenging task for several reasons. 1) The OC alone includes 2 HC and at least 6 SC types, and therefore the heterogeneity of bi-cellular junctions could influence core PCP protein distributions differently. 2) The cell(s) of origin for junctional enrichment has proved difficult to determine, requiring high-resolution microscopy or mosaic analyses. 3) Difference of core PCP protein organization in a single cell type could exist between the auditory and vestibular epithelia, or even among vestibular organs. Although a comprehensive catalog is work-intensive and might appear a "low impact" descriptive goal, it would greatly facilitate future studies required to fully understand intercellular PCP. An important future direction is to probe dynamic protein localization and its links to junctional actomyosin contractility in space and time, as has been initiated in other systems using quantitative live imaging [7]. Finally, mapping the protein interactomes of known PCP genes in the inner ear may identify novel PCP regulators and/or modes of signaling integration.

F2. Is Gai function required for the early peripheral migration of the basal body/ kinocilium?

Gai was proposed to be the driving force for the centrifugal migration of the basal body (see B3), but to date limited evidence supports this model. Treating E16-E18 cochlear explant cultures with PTX resulted in a centrally positioned basal body and consequently a circular hair bundle in a small fraction of HCs [26]. However PTXa expression in vivo, even with an early Cre driver (Foxg1-Cre), did not recapitulate this phenotype [46, 102, 103]. Furthermore, HCs were apparently able to break radial symmetry in conditional double Gai2/3 mutants using Foxg1-Cre [4], and when the Gai regulator Gpsm2 was missing [26, 102]. Another caveat is the subcellular localization of Gpsm2-Gai, which occupies the lateral apical membrane and only abutt the lateral junction where a planar pulling force would be expected to originate. In summary, it remains unclear whether the MT-pulling role of Gpsm2-Gai during mitotic spindle orientation is conserved in HCs. As Gpsm2-Gai was closely associated to actin dynamics [72, 99, 103], notably in stereocilia (see section E1) and during chemotaxis (Figure 4), Gai activity might not be required for the early basal body migration, and alternative mechanisms yet to be identified. Alternatively, it is also possible that we still lack adequate tools to fully block Gai function in HCs. Definitive support for a mitotic spindle-like role in HCs awaits an experimental model where the basal body remains central in HCs in vivo.

F3. Role of dynein-mediated pulling force in subsequent kinocilium positioning?

While there is no evidence for dynein mediating the early migration of the basal body, dynein is required for its subsequent positioning at the lateral apex. Dynein's precise mechanism of action is not fully understood, but likely involves both cargo transport and force generation. Specifically, both dynein and Lis1 are concentrated around the basal body where many ciliary and polarity proteins are localized, suggesting a possible cargo transport role. In addition, dynein is also localized near the cell junctions, consistent with an involvement in generating pulling force [26, 95]. In the latter case, how dynein is tethered at the HC cortex remains a key unresolved question. In mitotic cells, dynein is recruited through binding to NuMA that is in complex with Gpsm2-Gai (Figure 4A). Of note, NuMA is a nuclear protein and is only relocalized to the spindle poles and the cell cortex upon nuclear envelope breakdown [87]. In post-mitotic HCs, NuMA is found sequestered in the nucleus as expected, and is therefore unlikely to function in HC planar polarization [6]. What protein(s), then, could tether dynein at the cell cortex? Par3 and Daple again emerge as possible candidates, as both have been shown to interact with dynein in other systems [88, 92]. Moreover, both Par3 and Daple can bind MTs [9, 100], and they colocalize at lateral junctions from early stages of HC differentiation [94], an appropriate location to anchor planar pulling forces.

F4. Additional roles of the kinocilium during hair bundle morphogenesis?

While kinocilium tethering to central stereocilia in the first row is important to impart a chevron shape to the early hair bundle, the potential role of the kinocilium during postnatal hair bundle maturation remains relatively unexplored. In light of recent results, it would be interesting to ask whether the kinocilium is somehow implicated in the restricted trafficking

of Gpsm2-Gai to first row stereocilia, or the fact that these proteins are first enriched in central stereocilia in the kinocilium vicinity before being enriched in more peripheral stereocilia. Interestingly, cross-talk between the kinocilium and the Gai domain seems essential for hair bundle integrity, as hair bundle fragmentation was only observed in *Daple* mutants when the aberrantly medial kinocilium was outside of the Gai domain [94]. Alternatively, it also remains possible that the kinocilium only plays a limited early role to shape the hair bundle. A comprehensive understanding of the kinocilium contribution would greatly benefit from in vivo models that disrupt kinocilium function without confounding non-ciliary defects.

F5. Molecular mechanism(s) of Gai function during apical HC morphogenesis?

In the context of mitotic spindle orientation, Gai is thought to be sequestered in its GDP form by Gpsm2. By analogy, Gai activity in HCs at the bare zone and at stereocilia tips is likely to be non-canonical, perhaps serving as a cortical anchor point rather than a bona fide signaling relay for a GPCR. However, the citical role of the non-receptor GEF Daple suggests that guanine nucleotide exchange could be important for Gai function during HC planar polarization. Daple might compete with Gpsm2 for Gai binding at the apical membrane (bare zone)/apical junction interface, for example. In addition, Gai appears to function in a Gpsm2-independent pathway to regulate HC orientation reversal downstream of Emx2. While PTX is best known to disrupt GPCR signaling by preventing heterotrimeric Gai-G $\beta\gamma$ complex dissociation, PTX appears to effectively inhibit Gpsm2-Gai function as well, as illustrated by stunted stereocilia, for example. Nevertheless, it remains possible that some aspects of Gai function are insensitive to PTX. Further studies need to clarify 1) what possibly independent purposes Gai serves during HC differentiation, 2) what partners are regulating Gai activity in each case, and 3) to what extent each role involves classic signaling or a more structural function.

F6. Solving and leveraging planar polarity mechanisms for hair cell regeneration

Studying developmental mechanisms underlying auditory HC development has both fundamental and translational significance. In particular, recent insights indicate that planar polarity signaling regulates many aspects of auditory HC and SC development critical for mechanotransduction (section E1) and synaptic transmission (section E2). Thus, reactivation of planar polarity signaling pathways is absolutely necessary for regeneration of fully functional auditory HCs. Indeed, stem cell derived and experimentally regenerated HC-like cells often form vestibular-like or misshapen hair bundles with limited potential to function properly as sound receptors [55, 82, 107]. While restoration of intercellular PCP has been observed during HC regeneration of non-mammalian species [110], it will be important to probe and manipulate both PCP and cell-intrinsic signaling in mammalian models of HC regeneration.

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Highlights

- Intercellular and cell-intrinsic signaling coordinately regulates hair cell planar polarity.
- Planar polarity of supporting and other non-sensory cells is also crucial for hearing.
- Hair bundle stereocilia elongation is coupled to hair cell planar polarity.
- Epithelial planar polarity influences hair cell-spiral ganglion neuron connectivity.



Figure 1. Two levels of planar polarity in the organ of Corti.

A) Diagram depicting tissue-level, intercellular planar cell polarity (PCP), manifested by the uniform orientation of HCs along the medial-lateral axis of the cochlear duct. PCP is regulated by intercellular communication through asymmetric core PCP protein complexes at HC-SC junctions. B) By contrast, cell-intrinsic planar polarity refers to cytoskeleton asymmetry along the planar axis in individual HCs. Different structures displaying intracellular planar polarity are indicated. This behavior involves protein modules acting cell-intrinsically. C) Scanning electron microscopy (SEM) images showing PCP defects in

representative mouse mutants. Yellow arrowheads indicate OHC orientation. **D**) SEM images showing cell-intrinsic planar polarity defects in representative mouse mutants. The kinocilium is highlighted in orange. Note OHCs in *Rac1* mutants with off-center kinocilium relative to an abnormally flat hair bundle (arrows). PTXa mutants show IHCs where the kinocilium is disconnected from a split hair bundle. *Ptk7* mutants strictly exhibit orientation (PCP) defects, whereas *Rac1* and PTXa mutants exhibit both orientation and cell-intrinsic planar polarity defects. PTXa mutants have Cre-induced expression of the catalytic subunit of Pertussis toxin (PTXa) in HCs. *Ptk7* and *Rac1* SEM images are modified from [36, 66].



Figure 2. A molecular blueprint for planar polarization of the apical cytoskeleton. A) SEM images of individual OHCs representative of different stages of apical

A) SEM images of individual OHCs representative of different stages of apical differentiation. The kinocilium is highlighted in pink and the approximate OHC junction indicated in red. **B**) Diagram depicting changes at the HC apex from the onset of differentiation (E15.5, left) to around birth (P0, right). Initially, the aPKC kinase is uniformly enriched at the apical membrane, which is covered with microvilli, and the kinocilium occupies a central position. The first morphological evidence of planar asymmetry is the approximately lateral position of the kinocilium, which occurs at about the time the Insc-Gpsm2-Gai complex becomes planar polarized at the lateral aspect of the cell. The Insc-Gpsm2-Gai complex expands in surface area and labels the bare zone, the lateral region of apical membrane devoid of stereocilia or microvilli (asterisks). Insc-Gpsm2-Gai prevents aPKC enrichment at the bare zone, establishing a molecular blueprint at the apical membrane that helps position and coordinate the hair bundle and the kinocilium. The expansion of the bare zone coincides with a relocalization of the kinocilium, from its post-migration position juxtaposed to the lateral junction to a more central position at the vertex of the chevron-shaped hair bundle around birth.



Figure 3. Deiters' cell phalangeal processes are planar polarized along the longitudinal axis of the cochlea.

A) Diagram depicting one Deiters' cell (green) that extends a slanted phalangeal process towards the luminal surface further towards the cochlear apex compared to its cell body position. B) A maximum-intensity projection of confocal Z stacks of a P18 cochlea whole-mount stained for F-actin (magenta) and alpha-tubulin (green). Note the planar polarized phalangeal processes of the three rows of Deiters' cells (arrowheads). Arrows indicate the direction of the cochlear apex.



hair cell apical view

Figure 4. Shared protein modules regulate HC planar polarity and other polarized processes.

A) During oriented cell division, the Gpsm2-Gai complex, and in some cases the Par3-Par6aPKC complex, are polarized at the cell cortex and recruit effectors to pull on astral microtubules and position the mitotic spindle. Gai is myristoylated and anchored at the membrane. Gpsm2-Gai can interact with the Par3 complex through binding to the Insc adapter, or recruit the minus-end microtubule motor dynein through the Numa adapter. Lis1 is an activator for dynein required for spindle orientation. This evolutionary conserved process controls daughter cell positioning and tissue architecture across multiple tissues. B) During HC planar polarization, dynein-mediated pulling forces on apical microtubules may play a role in the lateral placement of the basal body. Junctional proteins including Dvl2, Daple and Par3, may serve as cortical dynein tethers to anchor pulling forces, and/or stabilize microtubule cortical attachment by activating Rac-Pak signaling. The Insc-Gpsm2-Gai complex provides a blueprint on the HC apical membrane, and may also plays a role in basal body placement. C) During chemotaxis, for example in neutrophils, motility requires G protein coupled receptors (GPCRs) to sense extracellular cues in the environment and activate heterotrimeric G proteins. GBy dissociated from activated, GTP-bound Gai in turn activates downstream effectors to regulate actin assembly and pseudopod extension at the leading edge. In addition to canonical GPCR signaling, it has been proposed that Gai-GDP bound to Gpsm2 might more specifically regulate sustained directionality by signaling through Insc and Par3. In other cell types, sustained directionality was shown to require microtubules and Lis1/dynein to regulate transport and polarized activation of signaling

molecules such as Rac1 GTPase. It is interesting to speculate that control over Gai guanine nucleotide exchange involving GEF proteins like GPCRs or Daple might ultimately interconnect different polarity modules that act in concert in HCs.



Figure 5. Distinct subcellular localizations for different polarity modules at the HC apex.

Proteins identified as planar polarized at the apical HC surface have been shown to occupy distinct subcellular domains. Core PCP proteins are enriched at adherens junctions and form antagonistic lateral (e.g. Dvl2, blue band) or medial (pink band) protein complexes. In contrast, Par3 and Daple proteins largely coincide with lateral tight junctions (orange), slightly more apical than core PCP proteins. Finally, Insc-Gpsm2-Gai are enriched above tight junctions at the bare zone, the region of apical membrane devoid of stereocilia or microvilli (green). Close vicinity between Gpsm2-Gai, Daple-Par3 and Dvl2 at the lateral aspect of HCs suggests that cross-talk between these protein modules and compartments may integrate intercellular PCP and cell-intrinsic cytoskeleton polarization.



Figure 6. Novel roles for Gpsm2-Gai and planar polarity in hair cells.

A) At the IHC apical membrane, Gpsm2-Gai is required for stereocilia elongation and graded heights across stereocilia rows. Gpsm2 and Gai proteins (green, arrows) are localized at the tip and near the base of stereocilia in the first row, being planar polarized at the lateral bare zone. This dual distribution suggests that planar polarity might play a role in the restriction of Gpsm2-Gai trafficking to first row stereocilia, establishing their tallest identity and regulating hair bundle staircase-like architecture. At the basolateral IHC membrane, both Gpsm2 and Gai are required for the medial-lateral gradient of ribbon synapse (orange) size and synaptic activity. The IHC ribbon synapse releases neurotransmitters onto spiral ganglion neuron (SGN) afferent terminals (blue). **B**) In absence of Gpsm2-Gai function, stereocilia fail to elongate at postnatal stages and the hair bundle retains an immature morphology with an excess number of rows, and the gradient of ribbon synapse size is abolished. It remains unclear how Gpsm2-Gai proteins influence synapse size and activity.