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Mutant mice reveal the molecular and cellular basis for specific sensory connections to inner ear epithelia and primary nuclei of the brain

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Abstract

We review the *in vivo* evidence for afferent fiber guidance to the inner ear sensory epithelia and the central nuclei of termination. Specifically, we highlight our current molecular understanding for the role of hair cells and sensory epithelia in guiding afferents, how disruption of certain signals can alter fiber pathways, even in the presence of normal hair cells, and what role neurotrophins play in fiber guidance of sensory neurons to hair cells. The data suggest that the neurotrophin *BDNF* is the most important molecule known for inner ear afferent fiber guidance to hair cells *in vivo*. This suggestion is based on experiments on *Ntf3* transgenic mice expressing *BDNF* under *Ntf3* promoter that show deviations of fiber growth in the ear to areas that express *BDNF* but have no hair cells. However, fiber growth can occur in the absence of *BDNF* as demonstrated by double mutants for *BDNF* and *Bax*. We directly tested the significance of hair cells or sensory epithelia for fiber guidance in mutants that lose hair cells (*Pou4f3*) or do not form a posterior crista (*Fgf10*). While these data emphasize the role played by *BDNF*, normally released from hair cells, there is some limited capacity for directed growth even in the absence of hair cells, *BDNF*, or sensory epithelia. This directed growth may rely on semaphorins or other matrix proteins because targeted ablation of the *sema3* docking site on the sema receptor *Npn1* results in targeting errors of fibers even in the presence of hair cells and *BDNF*. Overall, our data support the notion that targeting of the afferent processes in the ear is molecularly distinct from targeting processes in the central nuclei. This conclusion is derived from data that show no recognizable central projection deviation, even if fibers are massively rerouted in the periphery, as in *Ntf3*^{tgBDNF} mice in which vestibular fibers project to the cochlea.

1. Introduction

Neurosensory hearing loss is, next to conductive hearing loss, one of the more frequently encountered ailments of the elderly and may afflict as many as 1 in six over the age of 70. Neurosensory hearing loss consists of two distinct but interrelated processes, hair cell loss and loss of sensory neurons. Sensory neuron loss may be a consequence of hair cell loss or may happen through direct neuronal loss in certain neuropathies. Loss of neurons limits the usefulness of cochlear and vestibular implants, currently the only remedy to minimize the devastating personal and social effects neurosensory losses have on affected individuals. The molecular basis of sensory neuron maintenance in the absence of hair cells as well the molecular basis for directed growth of neuronal processes to regenerated hair cells or stimulating electrodes of implants is, next to the regeneration of hair cells, the second most challenging process in inner ear neurosensory development and regeneration. Studying

neurosensory development can provide insights into the molecular biology of directed nerve fiber growth in sensory regeneration and thus may contribute to our understanding and clinical application of such information.

Sensory neurons of the ear form through the action of the bHLH gene Neurogenin 1 (*Neurog1*; formerly *Ngn1*) in the wall of the developing otocyst. In the absence of *Neurog1*, no sensory neurons in either the vestibular or cochlear part of the ear ever form (Ma et al., 2000, 1998). Downstream of *Neurog1* is another bHLH gene, *Neurod1* (formerly *NeuroD* or *Beta*). This gene mediates certain aspects of pathfinding and migration as well as survival of sensory neurons (Kim et al., 2001; Liu et al., 2000). How *Neurog1* and/or *Neurod1* activate other downstream genes relevant for connecting developing sensory neurons to specific hair cells of a given sensory epithelium is not fully understood (Fritzsche, 2003a). Numerous data show that hair cells will be innervated by sensory fibers no matter how unusual their position is. Innervation of sensory epithelia will be in a rather specific and highly conserved pattern (Fritzsche et al., 2001) that can be used to identify sensory epithelia across phyla no matter what shape and form (Fritzsche, 2003b). In vitro data indicate that this is so because an unknown factor is released, apparently from hair cells (Bianchi and Cohan, 1991, 1993; Hemond and Morest, 1991). It has been suggested that this substance is not one of the two neurotrophins known to be important for inner ear innervation survival (Fritzsche et al., 2004), BDNF and Ntf3 (formerly NT-3). Specifically, BDNF seems not to play the role of a major attractant of nerve fibers (Bianchi and Cohan, 1993), despite the fact that it is almost exclusively expressed in hair cells of the developing ear (Farinas et al., 2001; Pirvola et al., 1992). However, it cannot be ruled out that other factors generated by supporting cells contribute to the apparent attraction generated by hair cells. This is so because hair cells and supporting cells are clonally related and always occur together in a normal ear (Fekete et al., 1998). Also, supporting cells express the neurotrophin Ntf3 (Farinas et al., 2001; Pirvola et al., 1992).

Beyond the role attributed to hair cells, it is clear that other factors that are not associated with hair cells may play additional roles in overall targeting of fibers across the developing otocyst to specific sensory epithelia (Fritzsche, 2003b; Rubel and Fritzsche, 2002). For example, tissue culture experiments showed that spiral ganglion fibers can extend toward the organ of Corti even if the hair cells have been removed (Sobkowicz, 1992). The initial contact formation will ultimately be refined through a process of pruning and fiber growth (Echteler and Nofsinger, 2000). However, none of these processes have been analyzed in the background of specific neurotrophin mutations to provide information independent of neurotrophin mediated nerve attraction.

Equally important and even less well understood is how the various central projections are determined. It is unknown what molecular mechanisms make cochlear fibers connect selectively to the cochlear nuclei (Rubel and Fritzsche, 2002) and vestibular fibers connect selectively to specific subsets of the vestibular nuclear complex or the cerebellum (Maklad and Fritzsche, 2003a). So far, we know only that the central projection is specifically targeted from earliest fiber growth onward (Maklad and Fritzsche, 2003b), but neonatal refinement of the overall properly targeted projections through activity mediated processes likely plays a role as well (Leake et al., 2002; Rubel and Fritzsche, 2002). Obviously, such specificity of both central and peripheral connections requires that both the peripheral dendrite and the central axon are properly targeted through either similar or complementary mechanisms co-expressed in the same sensory neuron.

Recent years have seen dramatic progress in the understanding of the molecular and cellular basis of connection formation in several sensory systems using loss-of-function (knockout) and transgenic approaches. For example, such analyses have clarified that in the olfactory

system there is a close correlation between the type of olfactory receptor formation and the specific projection of such receptor neurons to given glomeruli, their targets in the olfactory bulb (Mombaerts et al., 1996; Zou et al., 2004). While details of the guidance are still unclear, it appears that differential expression of ephrin family members are involved in proper targeting (Cutforth et al., 2003). Ephrin family members also play a major role in the formation of the retino-topical map as revealed by numerous in vitro studies (McLaughlin et al., 2003) and these suggestions were recently confirmed in loss-of-function experiments in vivo (Feldheim et al., 2004). Recent work on the central organization of taste bud projections showed the importance for fiber and target interaction in loss-of-function mutants in this nucleus (Qian et al., 2001). Together these data on nonotic sensory systems highlight the strength of the genetic approach as a tool to molecularly dissect, in vivo, the known or suspected cell and molecular interactions. While progress has been made, it is still a long way before the digital information of the genome can be used in a systems approach to understand the network perturbations caused by mutations and this information can then be used to guide therapeutic intervention (Hood et al., 2004). Such a detailed understanding of the function of individual genes in the context of a developmental network of gene interactions is, in our opinion, an essential prerequisite for any attempt to manipulate fiber growth to generate, for example, more profound connections with cochlear implants.

In this review, we will therefore highlight data based on genetic approaches in vivo. However, where appropriate, we will also outline the information gathered by other approaches. We will show that most of the peripheral targeting mechanisms currently known seem to have little effect on the central targeting specificity, indicating that different networks are involved on the dendritic and axonal end of a sensory neuron's processes.

Overall, we will address the following points:

1. What is the role of hair cells in guiding afferents?
2. Can fibers project correctly even in the absence of sensory epithelia?
3. Can the disruption of certain signals alter fiber pathways even in the presence of normal hair cells?
4. What is the role of neurotrophins in fiber guidance of sensory neurons?

1.1. The role of hair cells in guiding afferents to and within sensory epithelia

Assessing the role of hair cells in guiding afferents requires elimination of hair cells early in development to investigate whether other mechanisms for guidance exist which do not require hair cells. The effects of loss of some or all hair cells has been studied in the past in tissue culture or using mutants such as the Bronx-Waltzer mutation (Sobkowicz, 1992). These data suggest that at least some pathfinding properties are not mediated by hair cells and also showed the importance of hair cells. In recent years, several additional targeted mutations have become available that allow us to continue this investigation of the relative role played by non-hair cell mediated guidance mechanisms. Abrogation of hair cell formation on fiber extension to the cochlea can now be studied in embryos with two gene mutations, *Atoh1* (formerly *Math1*) and *Pou4f3* (formerly *Brn3c* or *Brn3.1*). Loss of hair cells can also be achieved later in neonatal development using various mutations such as the *Barhl1* homeobox gene (Li et al., 2002) or the *Gfi1* zinc finger factor (Hertzano et al., 2004; Wallis et al., 2003) and might present excellent models to study the effect of the absence of hair cells on maintenance of central and peripheral nerve fibers.

Null mutations for *Pou4f3* show initial formation of hair cells that fail to differentiate fully and disappear progressively in late neonates and early postnatal mice (Erkman et al., 1996; Hertzano et al., 2004; Vahava et al., 1998; Xiang et al., 1997, 1998). It needs to be stressed

that these cochlear hair cells *never* fully develop their apical specialization (Hertzano et al., 2004) and that most hair cells can not be recognized any more around postnatal day 8 (Xiang et al., 1998). Analysis of the peripheral projection pattern shows almost normal innervation of all cochlear and vestibular organs until approximately 8 days of age (Xiang et al., 2003). In fact, in the apex of the cochlea there is retention of sensory neurons and their innervation to the undifferentiated organ of Corti in the absence of hair cells for at least 6 months (Fig. 1). These data refute the importance of differentiated hair cells for long term maintenance of afferents. However, it cannot be ruled out that even undifferentiated hair cells and supporting cells that form before they degenerate have some attraction for afferents. Indeed, close examination shows that the neurotrophins BDNF and Ntf3 are still expressed in the sensory epithelia but at an apparently reduced level (Xiang et al., 2003). No hair cells could be detected in the cochlea apex of these mutants during the time fibers are extending to the outer hair cells in normal mice and no fibers grow beyond the topographical level equivalent to the inner hair cells in these mutants. These data suggest that growth of fibers to the cochlea is independent of differentiated hair cells, that retention of afferents is independent of hair cells but that growth of fibers to outer hair cells depends on the presence of hair cells. Further analysis is needed to show quantitatively the level of neurotrophin expression using real time PCR (Stankovic and Corfas, 2003) as well as the cellular localization and correlate that directly with the survival of fibers and their specific innervation pattern. Absence of hair cells seems to have little effect on the formation of crudely topographically restricted projection to the apex and the base of the cochlea which seem to develop rather normally in neonates (Fig. 1). Overall, these data are consistent with an analysis showing such topographically restricted projections in cats prior to the onset of hearing (Leake et al., 2002). Whether the topographical refinement of the central projection will be affected in older *Pou4f3* null mice has not been investigated yet. Likewise, whether these findings can be extended to the specific vestibular projection to the cerebellum (Maklad and Fritzsche, 2003a) remains to be demonstrated.

We have recently begun to investigate the pattern of innervation in *Atoh1* null mice (Bermingham et al., 1999). These mutants never develop recognizable hair cells likely because of a failure to initiate postmitotic differentiation (Chen et al., 2002). Our preliminary data suggest that at least some primordial hair cells form that express a low level of the neurotrophin BDNF and are apparently specifically innervated (Fritzsche et al., submitted). These data indicate that complete elimination, even of hair cell primordia, is needed to exclude the possible attraction of fibers to these primordia. An additional complication arises through the hair cell supporting cell interaction via the delta-notch system (Eddison et al., 2000). The importance of the delta-notch system mediated upregulation of bHLH genes in the CNS development has recently been established through mutational analysis (Hatakeyama et al., 2004). Since hair cell and supporting cell formation is clonally linked (Fekete et al., 1998) and hair cells are involved in the regulation of supporting cell maturation through the delta-notch system (Zine et al., 2001), elimination of hair cell precursors might result in abrogation of entire sensory epithelia. It is, therefore, possible that elimination of the entire formation of sensory patches may be required to investigate the role of other guiding mechanisms to bring fibers to specific sensory epithelia. Alternatively, suspected molecules for guidance such as neurotrophins can be eliminated and the remaining pattern of innervation analyzed in the presence of hair cells but in the absence of such presumptive guidance molecules.

1.2. Fibers project correctly even in the absence of sensory epithelia

Fibroblast growth factors have been shown to be essential for ear formation (Pirvola et al., 2002; Wright and Mansour, 2003a) and may also play a role in patterning of innervation (Brumwell et al., 2000; Wright and Mansour, 2003b). We recently showed that FGF10 is

essential for semicircular canal formation but is also necessary for the formation of the posterior canal crista (Pauley et al., 2003). Interestingly, while no hair cell or sensory primordia of the posterior crista seem to form, there is, although reduced, an initial formation of posterior crista-specific sensory neurons. Those neurons project toward the absent posterior crista (Fig. 2) but disappear, likely because of failure to receive neurotrophic support in the absence of any target, within two days (Pauley et al., 2003). These data suggest that initial pathfinding properties may not be related to sensory epithelia and their attraction for fibers. In addition, other factors may play a role such as the topology of sensory neuron formation in the otocyst wall, allowing neurons that delaminate from specific areas to project crudely correctly at least to the areas they delaminated from (Carney and Silver, 1983; Fritzscht et al., 2002). Alternatively, sensory neurons may leave their dendrites behind as trailing processes thus requiring only limited near target guidance of fibers toward and within sensory epithelia (Fritzscht, 2003a). These preliminary assessments require further confirmation by the demonstration that no molecules known to be associated with sensory epithelia formation are ever expressed in the area of future posterior crista in *Fgf10* null mutant mice. It also needs to be shown that posterior crista sensory neurons derive from distinctly different areas than the sensory epithelia, a possibility recently suggested (Raft et al., 2004).

1.3. Disruption of certain signals results in axonal targeting defects despite otherwise normal hair cells

The data on *Fgf10* null mutants clearly suggest that some pathfinding information derives from outside the sensory epithelia. In other sensory systems, pathfinding decisions may be intrinsically specified by their receptors, as is the case for odorant receptors in the olfactory system (Zou et al., 2004). Alternatively, projections are guided and refined by graded distributions of specific guidance cues and their receptors, such as ephrin ligands and their cognate receptors in the visual system (O'Leary and Wilkinson, 1999) or semaphorins and their receptors in both the peripheral and central nervous system (Pasterkamp and Verhaagen, 2001). Ephrin receptors and ligands are distributed in the developing ear (Bianchi and Liu, 1999) but no alterations in projection patterns have been found in the only mutant analyzed thus far, the EphB2 null mutant mouse (Cowan et al., 2000), possibly because of redundancy of ephrin ligand signaling through several ephrin receptors. Conditional mutations that eliminate several ephrin ligands and/or receptors simultaneously will be needed to address this complex problem. Such approaches are now possible using the growing availability of ear specific *Cre* transgenic mice combined with increasingly available floxed ephrin ligands and receptors.

Comparable to the ephrin ligands and their receptors, intriguing patterns of expression of some members of the semaphorin family of guidance cues and their receptors, neuropilin 1 and neuropilin 2, have been described in the ear (Miyazaki et al., 1999; Murakami et al., 2001). Null mutations of some of the ligands (*Sema3a*) and both neuropilin receptors exist (Cloutier et al., 2002). The *Npn-1* null mice exhibit an early embryonic lethal phenotype, prior to ear innervation. Thus, studying the effect of the semaphorin receptor *Npn-1* during ear innervation development required the targeted elimination of a single semaphorin docking site in *Npn-1* or the conditional elimination of semaphorin ligands and/or receptors only in the ear with either approach being likely to generate viable mutations. The former was recently done by altering the endogenous *Sema3a* docking site of the *Npn-1* receptor such that the receptor could not bind to semaphorin ligands, and the pattern of inner ear innervation was studied (Gu et al., 2003). These *Npn-1*^{Sema} mice exhibited profound reorganization of fibers into unusual trajectories around the ear. For example, projections to the posterior crista were found to pass along the anterior side of the otocyst as an extension of fibers targeted for the utricle (Fig. 3). Other fibers were found to overshoot sensory

epithelia and terminate instead in the skin above the ear (Fig. 3). A somewhat similar, transient phenotype was reported for some *Pou4f1* (*Brn3a*) null mutant mice (Huang et al., 2001), but it is unknown whether this Pou domain factor regulates *Npn-1* or plexin expression in sensory neurons.

We have also analyzed the central termination of such fibers in the skin in the *Npn-1*^{Sema}-knock-in mice by applying lipophilic dyes to the fibers in the skin at embryonic day 15 mutants. The findings show that these fibers form very few collaterals to vestibular sensory organs, derive from vestibular ganglia and terminate centrally in vestibular nuclei (Fig. 3). Most importantly, these data suggest that *Sema3a* mediated signaling via *Npn-1* provides a stop signal at or near the sensory epithelia. The absence of this stop signal, even in the presence of otherwise normal hair cells, leads to an overshooting of fibers which extend outside of the ear and into the skin. A more detailed analysis of mice lacking the various semaphorin ligands, and the plexin and neuropilin receptors needs to be performed to fully understand the effects thus far described in the *Npn-1*^{Sema}- knock-in mice (Gu et al., 2003). Floxed alleles for *Npn-1* and *Npn-2*, when combined with ear or hair cell specific *Cre* expressing mice such as the recently available *Pax2-Cre* (Ohshima and Groves, 2004) or a hair cell specific *Cre*, such as the *Prestin-Cre* (Tian et al., 2004), provided a proper time of *Cre* upregulation is achieved, would allow for organ specific or hair cell specific dissection of the function of each receptor, alone or in combination. Together such future directions could help to clarify the apparently significant role of this large family of ligands and receptors in the formation of the ear innervation pattern.

1.4. The role of neurotrophins in fiber guidance of sensory neurons

Only two neurotrophins and their cognate high affinity receptors are necessary for the maintenance of all inner ear innervation, *BDNF* with *Ntrk2* and *Ntf3* (formerly *NT-3*) with *Ntrk3* (Fritzsche et al., 2004; for review). No sensory neurons survive in the absence of both neurotrophins (Ernfors et al., 1995; Liebl et al., 1997) or both neurotrophin receptors (Fritzsche et al., 1995; Silos-Santiago et al., 1997). In the ear, in single receptor or ligand null mutants, the neurotrophin ligands show complex alterations in their spatiotemporal pattern of expression (Farinas et al., 2001) that appears to be directly related to the spatially restricted loss of sensory neurons (Fritzsche et al., 1997, 2004). For example, loss of *BDNF* results in absence of all crista innervation except for an occasional fiber (Fig. 4). In contrast, gravistatic organs and the cochlea show only reduced innervation and a limited loss of spiral ganglion neurons (Bianchi et al., 1996; Fritzsche et al., 1997). *Ntf3* null mutants show a loss of 85% of spiral ganglion cells with complete loss of all basal turn sensory neurons (Farinas et al., 1994; Fritzsche et al., 1997).

Recently, we directly tested the prediction that hair cells can have some attraction for sensory fibers independent of *BDNF* by combining *BDNF* null mutation with the *Bax* mutation. *Bax* null mice do not show neuronal cell death even in the absence of *BDNF*. Our analysis shows a more profound innervation of the anterior and horizontal crista, but not of the posterior crista (Hellard et al., 2004). Close examination shows that many, but not all fibers, are targeted to the cristae and seem to innervate hair cells (Fig. 4). However, several fibers were found outside the cristae and those inside the cristae tended to be near the base where the *Ntf3* expression was found (Figs. 4 and 6). Consistent with the data on the transgenic mice, these data suggest that hair cells do exert some very limited attraction on nerve fibers that is not mediated by *BDNF*.

From this analysis, one might predict a functional equivalence of either neurotrophin mediated by the apparently redundant expression of both neurotrophin receptors in all inner ear sensory neurons. (Farinas et al., 2001) Indeed, this co-expression of both receptors allowed direct testing of this prediction by studying the remaining sensory neurons to the

cochlea of transgenic mice expressing *BDNF* under *Ntf3* promoter control (Coppola et al., 2001) or of *Ntf3* under *BDNF* promoter control (Agerman et al., 2003). In *Ntf3^{tgBDNF}* (*BDNF* under *Ntf3* promoter control) mutants, *BDNF* is expressed instead of *Ntf3* such that there is no *Ntf3* expression, and *BDNF* is expressed in both its normal location and in the *Ntf3* location. The opposite is true for *Ntf3^{tgBDNF}* mutants. Both studies agree that, at least in the cochlea, there is functional equivalence of these neurotrophins for sensory neuron survival.

Using these transgenic mice, we showed that premature expression of *BDNF* under *Ntf3* promoter control leads to massive rerouting of vestibular fibers to the basal turn of the cochlea (Fig. 5). *Ntf3* is only expressed late in development in the crista of the semicircular canals and is around the sensory cristae rather than in supporting cells of the cristae (Fig. 4). Importantly, in the *Ntf3^{tgBDNF}* mutants, fibers to the posterior crista stall on their way to the crista and turn around to innervate the basal turn of the cochlea (Fig. 5). This effect depends directly on the lack of early *BDNF* expression in the posterior crista and is most profound in the combined *Ntf3^{tgBDNF}/BDNF* null mutants described below. Most interesting are the innervation defects in anterior and horizontal crista. Simple misexpression of *BDNF* under *Ntf3* promoter control results in fibers innervating areas outside the sensory epithelia, suggesting that *BDNF* acts as short range attraction for fibers even in the presence of hair cells (Figs. 5 and 6). More recently, we combined the *Ntf3^{tgBDNF}* mice with the *BDNF* null mutation, to test the effect of misexpression of *BDNF* on vestibular fiber pathfinding (Tessarollo et al., 2004). In these mice, there is no *Ntf3* expression and *BDNF* is only expressed in the *Ntf3* pattern and not in its normal pattern. Eliminating *BDNF* expression under its own promoter control, and therefore in hair cells, showed profound rescue of fibers to the cristae organs (Fig. 5). These fibers target hair cells inside the cristae organs as well as the *BDNF* expressing epithelial cells outside the cristae organs (Figs. 5 and 6) suggesting that factors other than *BDNF* can cause a very short range attraction to hair cells, provided neurons survive in *BDNF* null mutants and have fibers near the crista epithelia. This suggestion is supported by the complete rerouting of posterior cristae fibers into the basal turn of the cochlea in *Ntf3^{tgBDNF}* mice combined with *BDNF* null mutation, presumably because of the limited and delayed upregulation of *BDNF* around the posterior crista epithelium that can not compete with the more extensive and earlier expression of *BDNF* under *Ntf3* promoter control in the basal turn of the cochlea. Moreover, the presence of fibers outside sensory epithelia in areas that express *BDNF* in the transgenic misexpressors suggest that *BDNF* can override the attraction of hair cells, even if hair cells *do* express *BDNF*.

We also analyzed the central projection of the vestibular fibers that enter the sensory epithelia but do not, at least in the vast majority, innervate cochlear hair cells. Different colored lipophilic tracers injected into the posterior crista and the basal turn of the cochlear directly show that many of these fibers terminate not in the auditory nuclei but rather in the nearby vestibular nuclei (Tessarollo et al., 2004). These data suggest that the central projection of vestibular fibers is not dependent on the target but rather reflects properties that are distinct for the central projection independent of their peripheral connection. Moreover, these data fully agree with our data on the *Npn-1^{Sema}*-knock-in mice and together argue that guidance of central fiber projection is molecularly distinct from peripheral dendrite to hair cell targeting mechanisms. This implies that each end of the neurite growth process has to develop its own, unique, yet closely coupled molecular targeting machinery to accomplish proper navigation independent of, and yet crucially dependent upon, each other.

2. Uncited reference

(Fritzsche et al., 1997).

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Fig. 1.

The Pou domain factor *Pou4f3* (*Brn3c*) is required for hair cell differentiation and maintenance of hair cells past birth. Only few basal turn hair cells can be identified in neonatal animals using Myosin VII immunocytochemistry (a,b). Despite this absence of differentiated hair cells, afferent fibers are targeted to both vestibular and cochlear sensory epithelia (c,d). Such fibers project in a crudely topographical fashion to the cochlear nuclei as revealed by the injection of different fluorescent tracers into the apex and base of the cochlea, respectively (e,f). These data establish that an overall fairly normal peripheral and a crude central projection can develop in the absence of functional hair cells. AC, anterior crista; HC, horizontal crista; HCs, hair cells; IHCs, inner hair cells; OHCs, outer hair cells; PC, posterior crista; VCN, ventral cochlear nucleus. Modified after (Xiang et al., 2003).

Fig. 2.

Fgf10 null mutant mice do not develop semicircular canals and have no posterior canal crista (a,b). Such animals do not show a projection of afferent or efferent fibers to the posterior region of the ear either in late embryonic stages. However, at early embryonic stages posterior crista sensory neurons apparently form and project toward the area of the posterior crista almost like in wildtype or heterozygotic littermates (c,d). Within two days such projections are lost, presumably because of lack of neurotrophic support. These data support the idea that some pathfinding properties of vestibular afferents reside outside sensory epithelia. AC, anterior cristae; HC, horizontal cristae; PCS, posterior crista sensory neurons; PC, posterior crista; S, sacculle. Modified after (Pauley et al., 2003).

Fig. 3.

Mice with a targeted replacement of the *Sema3a* docking site at the neuropilin1 receptor (*Npn-1*^{Sema-} knock-in mice) show distinct defects in the inner ear innervation. Specifically, in these mutants, fibers may not stop near hair cells or sensory epithelia but continue to grow until they reach the skin above the ear (a,b). Injections of a lipophilic tracer into the skin above the ear will label the trigeminal system but also fibers that show side-branches to the vestibular endorgans as they pass through the ear (c). In addition, these fibers can be traced to vestibular ganglion cells (d) and can be shown to project centrally like vestibular fibers into the vestibular nuclei rather than like trigeminal fibers. These data suggest that *Sema3a* is at least one of the stop signals at or near sensory epithelia that directs fibers to hair cells. The data also show that at least during embryonic development specification of central projections of vestibular neurons does not depend on being connected to hair cells. AC, anterior crista; HC, horizontal crista; MesV, nucleus mesencephalicus V; MVN, medial vestibular nucleus; PC, posterior crista; S, saccule; SVN, superior vestibular nucleus; U, utricle. Modified after (Gu et al., 2003).

Fig. 4.

The neurotrophins *BDNF* (a) and *Ntf3* (b) are both expressed in late embryonic canal cristae. However, only hair cells are positive for *BDNF* (a) while *Ntf3* is expressed in the stroma of the crista as well as in adjacent dark cells near the crista (b). In wildtype mice afferent fibers are targeted to cristae as well as the utricle and innervate densely hair cells (c). In mutants, in which *BDNF* has been eliminated, hardly any fiber projects to canal cristae (e). However, breeding *BDNF* null mice into a *Bax* null background results in survival of some neurons and growth of some neurites in the absence of *BDNF* (d,f). These fibers extend toward cristae (d) but innervate predominantly areas of *Ntf3* expression with only an occasional fiber extending to hair cells (f). These data suggest that *BDNF* is not only a major survival factor but also helps direct growth of fibers to hair cells. Nevertheless, the residual ability of fibers to grow toward hair cells suggests that some additional attractive substances may be released from hair cells. AC, anterior crista; HC, horizontal crista; U, utricle. Modified after (Hellard et al., 2004).

Fig. 5.

The effect of misexpression of *BDNF* under the *Ntf3* promoter control on vestibular fiber pathfinding is shown without (a) or with a combined elimination of *BDNF* (b–f). Replacing of *Ntf3* with *BDNF* results in rerouting of saccular and posterior crista afferents even in the presence of normal *BDNF* expression (a). However, combining the transgenic expression of *BDNF* under *Ntf3* promoter control with absence of *BDNF* results in more profound rerouting of fibers to the basal turn of the cochlea (b) as well as rescue of fibers to the anterior and horizontal crista (d,e). Despite the fact that no neurotrophin is expressed in hair cells in these animals, fibers preferentially innervate hair cells (e) but also project outside the sensory epithelia to areas of expression of *Ntf3* (c–e). These data suggest that attractors other than *BDNF* must exist in hair cells but also suggest that such attractors can be overridden by *BDNF* which by itself can attract fibers outside the sensory epithelia. Central projections from the cochlea (f) show fibers not only to the cochlear nuclei but also to vestibular nuclei. These data suggest that the central projection is regulated molecularly distinct from the peripheral projection. AC, anterior crista; DCN, dorsal cochlear nucleus; HC, horizontal crista; MVN, medial vestibular nucleus; PC, posterior crista; U, utricle. Modified after (Tessarollo et al., 2004).

Fig. 6.

This schematic illustrates (a) the normal areas of expression of *BDNF* and *Ntf3* in the utricle, hair cells of the anterior canal crista, and cells surrounding the sensory area of the crista, and the normal pattern of innervation of the crista. (b–f) The variations in expression pattern of neurotrophins and resultant patterns of crista innervation are shown. *BDNF* is required for neuron growth and survival (b) whereas *Ntf3* is not (c). When *Bax* is also knocked out in the *BDNF* null mutant, fibers survive and project to areas of *Ntf3* expression (d). When *Ntf3* is replaced with *BDNF*, fibers innervate the sensory cells as usual, but also project to non-sensory, *BDNF*-expressing areas, demonstrating a positive attraction of fibers to *BDNF* even when expressed in an abnormal location (e). Combining the transgenic mouse in which *Ntf3* is replaced with *BDNF* with *BDNF* null mutant shows that fibers can innervate hair cells that do not express *BDNF*, suggesting that other hair cell or supporting cell factors are involved in fiber attraction, but are also attracted to *BDNF* expressing non-sensory patches (f). White areas are areas in which neither *BDNF* or *Ntf3* are expressed. Red represents *Ntf3* expression. Blue represents *BDNF* expression. Purple represents normal overlap of *BDNF* and *Ntf3* expression in hair cells and supporting cells, respectively. Green represents *BDNF* expression in place of *Ntf3* expression and teal represents an overlap of normal *BDNF* expression in hair cells with *BDNF* expressed under the *Ntf3* promoter in supporting cells. AC, anteroventral crista.