



Npr2 null mutants show initial overshooting followed by reduction of spiral ganglion axon projections combined with near-normal cochleotopic projection

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Abstract

Npr2 (natriuretic peptide receptor 2) affects bifurcation of neural crest or placode-derived afferents upon entering the brain stem/spinal cord, leading to a lack of either rostral or caudal branches. Previous work has shown that early embryonic growth of cochlear and vestibular afferents is equally affected in this mutant but later work on postnatal Npr2 point mutations suggested some additional effects on the topology of afferent projections and mild functional defects. Using multicolor lipophilic dye tracing, we show that absence of Npr2 has little to no effect on the initial patterning of inner ear afferents with respect to their dorsoventral cochleotopic-specific projections. However, in contrast to control animals, we found a variable degree of embryonic extension of auditory afferents beyond the boundaries of the anterior cochlear nucleus into the cerebellum that emanates only from apical spiral ganglion neurons. Such expansion has previously only been reported for Hox gene mutants and implies an unclear interaction of Hox codes with Npr2-mediated afferent projection patterning to define boundaries. Some vestibular ganglion neurons expand their projections to reach the cochlear apex and the cochlear nuclei, comparable to previous findings in Neurod1 mutant mice. Before birth, such expansions are reduced or lost leading to truncated projections to the anteroventral cochlear nucleus and expansion of low-frequency fibers of the apex to the posteroventral cochlear nucleus.

Keywords Natriuretic peptide receptor 2 · Npr2 null mutants · Cochleotopic projection · Axonal branching · C-type natriuretic peptide

Introduction

Previous work has identified the receptor guanylyl cyclase natriuretic peptide receptor 2 (Npr2, also designated guanylyl cyclase-B), its ligand C-type natriuretic peptide (CNP) and the cGMP-dependent kinase I (cGKI) as leading factors that regulate bifurcation of central axons of dorsal root ganglion (DRG) neurons along the spinal cord (Schmidt et al. 2002, 2007, 2009). More recent work has expanded these initial findings to neural crest and placode-derived fibers associated

with the V, VII, IX and X cranial nerves as well as to the placode-derived inner ear ganglia (Ter-Avetisyan et al. 2014) and most recently, the midbrain sensory neurons innervating muscles of mastication, the mesencephalic trigeminal (MesV) neurons (Ter-Avetisyan et al. 2018). Without CNP, Npr2, cGKI, DRG, or cranial sensory ganglion (CSG), axons do not bifurcate when entering the spinal cord or the hindbrain, respectively. Instead, they either turn in an ascending or descending direction. Collateral formation from these stem axons is not impaired (Dumoulin et al. 2018; Schmidt et al. 2007; Tröster et al. 2018). Expression of receptor or ligand suggests a progressive alteration over time from a restricted initial expression in sensory neurons (Npr2, cGKI) and rhombomeres (CNP) that have cranial nerve entry points (Ter-Avetisyan et al. 2018).

Detailed growth of fibers either rostral or caudal in the hindbrain has previously been reported for the vestibular system and sparse labeling suggests immediate bifurcation of inner ear afferents comparable to other sensory neurons (Ter-Avetisyan et al. 2014). However, judging from mass labeling

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of all vestibular afferents, many appear to behave differently from other sensory systems by developing for a given ganglion neuron either a rostral or caudal directed axon that only later is supplemented by a secondary branch in the opposite direction (Maklad et al. 2010). This development appears to help sort sensory organ-specific polarity information flow for the gravistatic vestibular organs to the hindbrain (Fritzsche and López-Schier 2014). In contrast, other sensory organs of the ear such as canal cristae and the organ of Corti of the cochlea have hair cells of only one polarity and organize their afferents according to different principles reflecting angular acceleration in three dimensions (Chagnaud et al. 2017) or project the tonotopic organization of the organ of Corti as regular cochleotopic projection onto the cochlear nuclei (Fritzsche et al. 2015; Jahan et al. 2010b; Muniak et al. 2016) even if such nuclei do not form properly (Elliott et al. 2017). Additional work using the *Npr2*-cn mouse mutant that lacks activity due to a point mutation in the guanylyl cyclase domain (Tsuji and Kunieda 2005) has implicated *Npr2* in some aspects of cochleotopic sorting of spiral ganglion (auditory) afferents leading to a “blurred tonotopic organization” (Goodrich 2016; Lu et al. 2014) and confirmed branching defects of *Npr2* mutants (Lu et al. 2011; Schmidt et al. 2009; Schmidt et al. 2007). Surprisingly, this reported “blurred projection of cochleotopic afferents” had little effect on multiple physiological properties in the *Npr2*-cn mutant (Lu et al. 2014) implying that in the auditory system, the mapping of the primary organization axes does not profoundly influence physiological properties of higher order neurons. Likewise, physiological analysis of *Npr2* null mutant mice shows only very limited effect with a slightly altered auditory brain stem response (ABR) and distortion product otoacoustic emission (DPOAE) (Wolter et al. 2018). This is in sharp contrast to a recent report showing that a massively disorganized cochleotopic projection in mutants with a conditional deletion of *Neurod1* causes loss of tuning in higher order neurons (Macova et al. 2019).

In contrast to previous work (Lu et al. 2014), we here use the well-established *Npr2* null mouse model (Ter-Avetisyan et al. 2014) to investigate in more detail the development of the cochleotopic central projections of the ear with an emphasis on the development over a larger range of stages compared to previous reports (Lu et al. 2014) to better elucidate how lack of bifurcation branching influences tonotopic and polarity-specific projections. Notably, a blurred cochleotopic projection was only reported for 2 out of 4 mutants at E16.5 implying incomplete penetrance of the point mutation. Moreover, we also like to better understand how *Npr2* and other guanylyl cyclase signaling intersects with the Wnt/planar cell polarity pathway recently implicated in a massive reorganization of some otherwise cochleotopically organized spiral ganglion neuron projections (Fritzsche et al. 2019; Yang et al. 2017). Our data reveal near-normal cochleotopic and vestibular projections in *Npr2*-deficient mutants suggesting the limited

influence of lack of bifurcation on cochleotopic or vestibulotopic projections. Surprisingly, we find that *Npr2* alters anterior cochlear nucleus afferent projections by partially removing blocks that prevent auditory afferents to grow beyond rhombomere 2 to enter the cerebellum, like vestibular fibers. Later stages show an unexpected reduction of afferents to the anteroventral cochlear nucleus and a bias toward the posteroventral and dorsal cochlear nucleus as well as an unusual cochleo-vestibular anastomosis interconnecting vestibular organs with the apex of the cochlea and the most ventral cochlear nucleus through unusual bifurcations in the ear.

Material and methods

Mice We used the previously described (Ter-Avetisyan et al. 2014) *Npr2*-lacZ (B6.129P2-*Npr2*^{tm1.1(NLS-lacZ)Fgr}) mouse line to generate homozygous *Npr2*-lacZ/lacZ mice, here referred to as *Npr2* null mice. Timed pregnancies were initiated by placing males and females together and time of detection of a vaginal plug was designated as embryonic day (E) 0.5. Embryos were collected at E13.5, E16.5 and E18.5. Toes of embryos were clipped and used for genotyping as previously described (Ter-Avetisyan et al. 2014). Embryos were immersion fixed in 4% paraformaldehyde (PFA) in 0.1 M phosphate buffer with 0.3 M sucrose added (Fritzsche et al. 2016). Fixed heads were transferred to 0.4% PFA in the same buffer and shipped from Germany to the USA for analysis. All procedures were performed in accordance with the relevant ethical regulations and had been approved by the local animal use and care authority (LaGeSo, Berlin: Nos. T0370/97 and G0370/13).

Expression analysis Heterozygous *Npr2*-lacZ [B6.129P2-*Npr2*tm1.1(nslacZ)/Fgr] (Ter-Avetisyan et al. 2014) and heterozygous *CNP*-lacZ [B6;129P2-*Nppc*^{tm1.1(nslacZ)/Fgr}] (Schmidt et al. 2009) mice were used to monitor the expression of *Npr2* and *CNP*, respectively, in embryonic brains and sensory neurons. Both transgenic reporter lines encode a β -galactosidase expression cassette containing a nuclear localization signal.

For whole mount β -galactosidase (β -gal) staining, embryos were fixed in Zamboni’s fixative (Stefanini et al. 1967): PBS containing 2% paraformaldehyde and 15% picric acid, pH 7.3 at room temperature for 15–30 min according to the developmental stage of the specimen. After 3 incubations for 10 min each in β -gal wash solution (2 mM MgCl₂, 0.02% Nonidet-P40, 0.01% sodium deoxycholate in PBS), tissues were transferred to β -gal wash solution containing 0.5 mg/mL 5-bromo-4-chloro-3-indolyl- β -D galactopyranoside (X-gal; Life Technologies, Germany), 5 mM potassium ferrocyanide and 5 mM potassium ferricyanide. Tissues were incubated at 37 °C under light agitation until the development of a blue color. The staining reaction was then terminated by

rinsing the embryos 3 times for 10 min each in ice-cold PBS containing 2 mM MgCl₂. Subsequently, the tissues were postfixed in PBS containing 4% paraformaldehyde and shipped to the University of Iowa for further analysis.

Following X-gal staining, whole heads were dissected and either the whole mounted brain (CNP-Z/+) or the ganglia (Npr2-Z/+) were imaged using a compound Nikon E800 microscope (Fig. 1) to validate expression changes in particular in rhombomere 4, the entry point of the VIIIth cranial nerve (Maklad et al. 2010) and the sensory neurons delaminating from the developing ear (Elliott et al. 2017) as previously described (Matei et al. 2006).

Dye tracing Heads of Npr2 mutants and control animals were transferred to 4% PFA in 0.1 M phosphate buffer with 0.3 M sucrose added and stored at 4 °C until tracing was initiated after shipment to the University of Iowa. For tracing, heads were split to allow insertion of dyes into the CNS and the ear (Fritzscht et al. 2016; Maklad et al. 2010). We previously established that long wavelength lipophilic dyes allow for better signal-to-noise ratio due to lack of background fluorescence that is particularly high in the 488-nm excitation range and also established that the use of dye-soaked wavers provides a more reliable way of dye application compared to crystals or blind injections (Duncan et al. 2011, 2015; Tonniges et al. 2010). Specifically, we used the NeuroVue dyes Maroon (NVM; 635/650–700 excitation/emission), Red (NVR; 543/565–615) and Jade (NVJ; 488/500–540), which we previously developed in collaboration with Molecular Targeting Technologies (Duncan et al. 2011, 2015; Fritzscht et al. 2016; Tonniges et al. 2010) that are sold through Polyscience (<http://www.polysciences.com/default/catalog-products/life-sciences/molecular-biology/neuronal-tracing-neurovue-sup-r-sup-dyes>). NeuroVue wavers were cut into appropriately sized and shaped pieces and inserted into either the cochlea, vestibular organs, or different parts of the brain as previously described (Elliott et al. 2017; Fritzscht et al. 2016; Macova et al. 2019). We always inserted NeuroVue Maroon (NVM) into the apex and false colored it green and NeuroVue Red (NVR) into the base and false colored it red. NeuroVue Jade (NVJ) was used in addition to reveal in some stages the vestibular afferents through waver application into the anterior and horizontal canal as well as the utricle. Insertion of appropriately sized slivers of the dye-soaked waver was for the base transverse to the basal turn hook region through the round window. For the apex, we used a lateral approach that inserted the dye filter near the apical tip into the modiolus. For the vestibular insertion, we removed the otic wall near the cristae of the anterior and horizontal canal using the pigment epithelium surrounding these organs as a guide. Care was taken to apply dyes as identical as possible to control and mutant animals to reveal differences in a direct comparison of littermates with identical diffusion times. The

analysis was mostly done in whole-mounted ears and brains to generate overviews of most profound qualitative differences. Diffusion of dyes was initiated at 60 °C for 2–7 days in 4% PFA with sucrose added as previously described (Fritzscht et al. 2016).

Brain stem analysis After diffusion, the cochlear and vestibular nerves were cut close to the cochlear nuclei and the half brain stem was trimmed to better show cochlear and vestibular nuclei and the cerebellum in younger stages. Diffusion timing was optimized for a given age of the embryos. We previously had shown that lipophilic dye diffusion leads to unreliable results in older stages due to massive dye uptake in myelin (Duncan et al. 2011) and low osmolality of the carrier leads to swelling of neurons, membrane disruption and unspecific dye leaking. We therefore terminated our investigation at P7, the oldest stage we previously showed leads to reliable results showing single-labeled fibers using lipophilic dye tracing for the above outlined reasons (Maklad and Fritzscht 2003a, b). For each preparation, the microdissected ear and the cochlear/vestibular nuclei were mounted on a glass slide in glycerol with spacers added to avoid compression. Preparations were imaged within an hour after mounting to avoid excessive washout effects of lipids and, by logical extension, lipophilic dyes with them. Preparations were viewed on either a Leica SP5 or SP8. Stacks of images were taken covering the entire area of cochlear nuclei or the microdissected inner ear. Using the intensity settings of the Leica system, laser power and gain were set to stay within the linear dynamic range of the system. Stacks were taken at 8–12 μm (lower power lenses such as × 10; 0.5 NA) or at 6 μm (higher power lenses such as × 20; 0.75 NA or 40 × 1.35 NA). The resolution was set at 1026 × 1026 pixels. Using Leica software, the image stacks were rendered into a single 2D image. Since our previous work had demonstrated that NVM is the fastest diffusing NeuroVue dye (Fritzscht et al. 2005; Jensen-Smith et al. 2007), we always used this dye to label the apex. Unfortunately, the 635-nm excitation also stimulates hemoglobin and thus shows all remaining red blood cells in particular in younger stages. Only preparations, in which the labeling in the ear showed that only intended parts were labeled, were used for the central projection analysis.

Ear analysis In several cases, we inserted dye into the brain stem and cerebellum to label afferents to the vestibular part of the ear as previously described (Maklad and Fritzscht 2003a, b; Maklad et al. 2010). As before, we verified the intended insertion and accepted only preparations confirming to the intent. After appropriate diffusion of 2–5 days at 60 °C, the nerves connecting the ear with the brain were cut with micro-scissors; the ears were microdissected and mounted with the brain stem in glycerol using spacers and viewed as described above.

An initial assessment of projections was conducted in a blinded fashion to ensure that the data generated by the tracing could identify the mutants without knowing which of the traced projections belonged to a mutant. Only after it became obvious that only *Npr2* mutant embryos had fiber projections passing beyond the rostral boundaries of the cochlear nucleus to reach the cerebellum (Figs. 2, 3) were the genotypes

disclosed prior to the tracing experiment. For each data point, we analyzed at least 3 *Npr2* null mutants and control wild-type littermates for a total of six half head preparations per each stage and genotype analyzed.

Statistics We used the Wilcoxon-Mann-Whitney *U* test (Mann and Whitney 1947) to establish at one stage

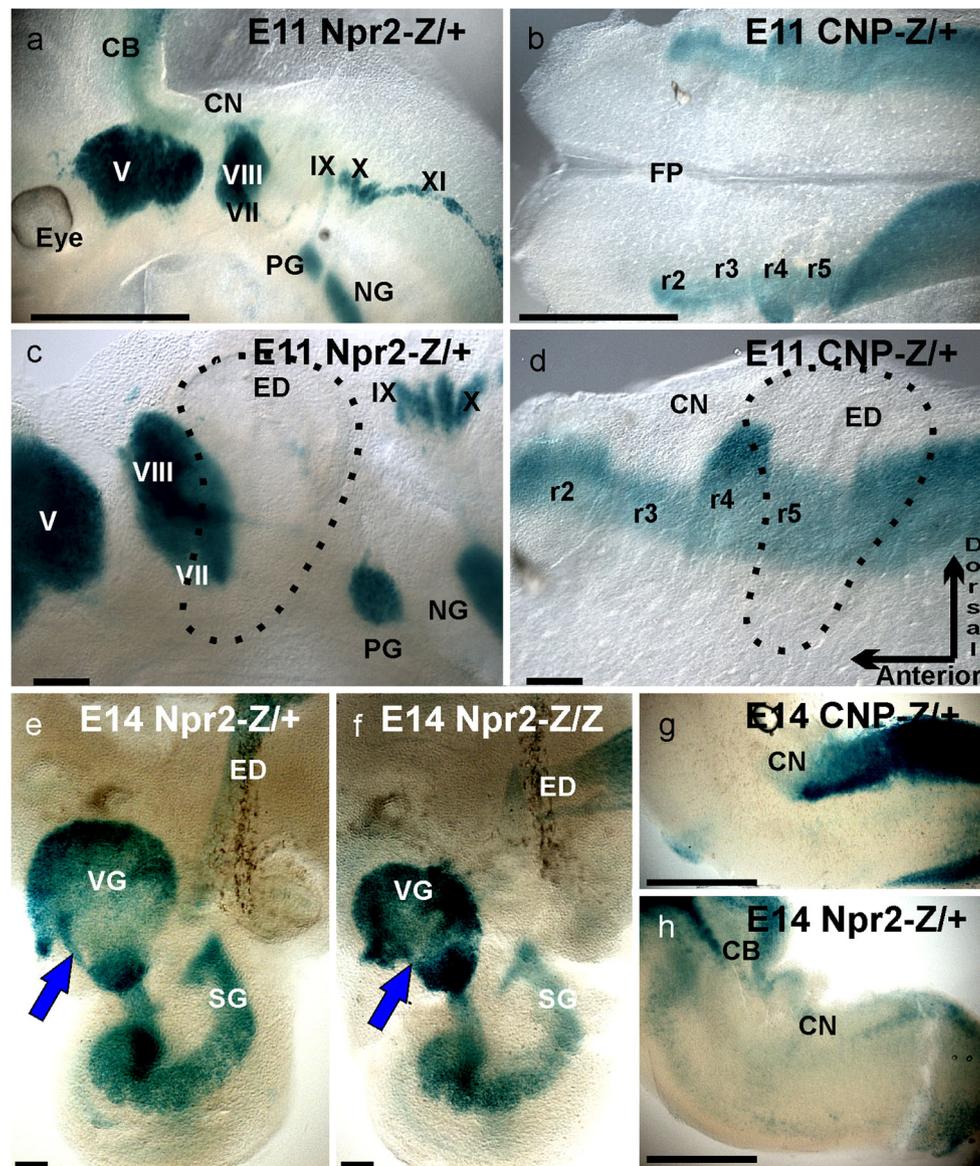
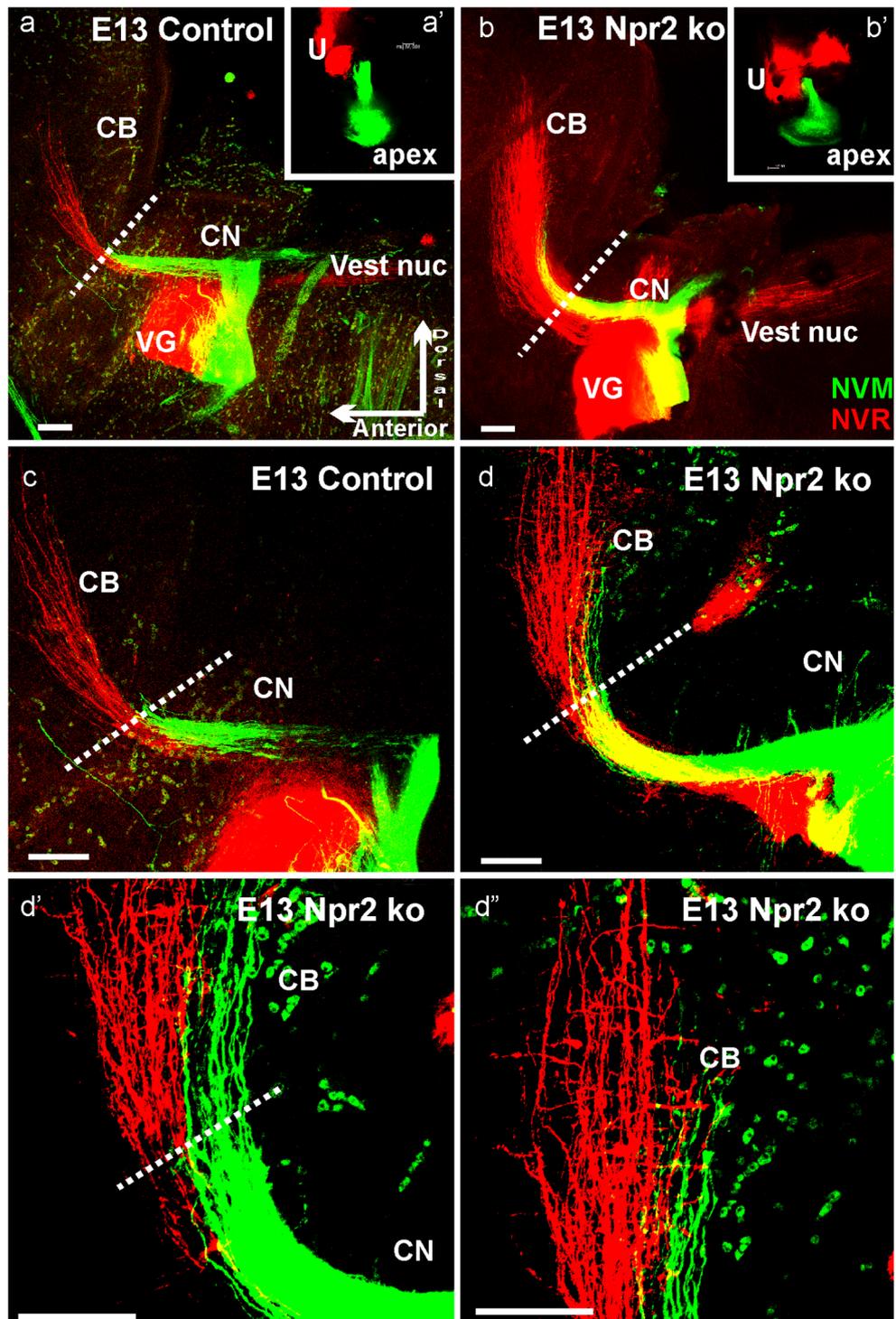


Fig. 1 X-gal staining of tissue from *Npr2*- or *CNP*-lacZ reporter mice to monitor the expression of the receptor *Npr2* and its ligand *CNP*. Whole-mounted hemisected heads (**a**), peripheral ganglia (**c**) and whole mounted hindbrains (**b**, **d**) show the β -galactosidase reaction product indicating expression of *Npr2* in all cranial ganglia (**a**, **c**) and expression of *CNP* in nearly all rhombomeres (**b**, **d**). Note neurons delaminating from the otocyst (indicated by dotted line in **c,d**) already express *Npr2* and the ligand *CNP* is especially highly expressed dorsally in rhombomere 4(r4) where the VIIIth cranial nerve enters. Later stages show continued expression of

Npr2 in inner ear sensory neurons (**e**, **f**), hindbrain (**h**) and of *CNP* in the hindbrain (**g**). Comparing *Npr2* mutant lacZ expression (**f**) with mice heterozygous for *Npr2* deletion shows only a minor size reduction of vestibular ganglion neurons (**e**, **f**). Note the *Npr2*-positive population of cells near the anterior margin of the vestibular ganglion (blue arrowhead). *CNP* expression remains high in the caudal hindbrain but is not detected in the cochlear nuclei (**g**). Bar indicates 1 mm in **a**, **b**, **g**, **h** and 100 μ m in **c**, **d**, **e**, **f**

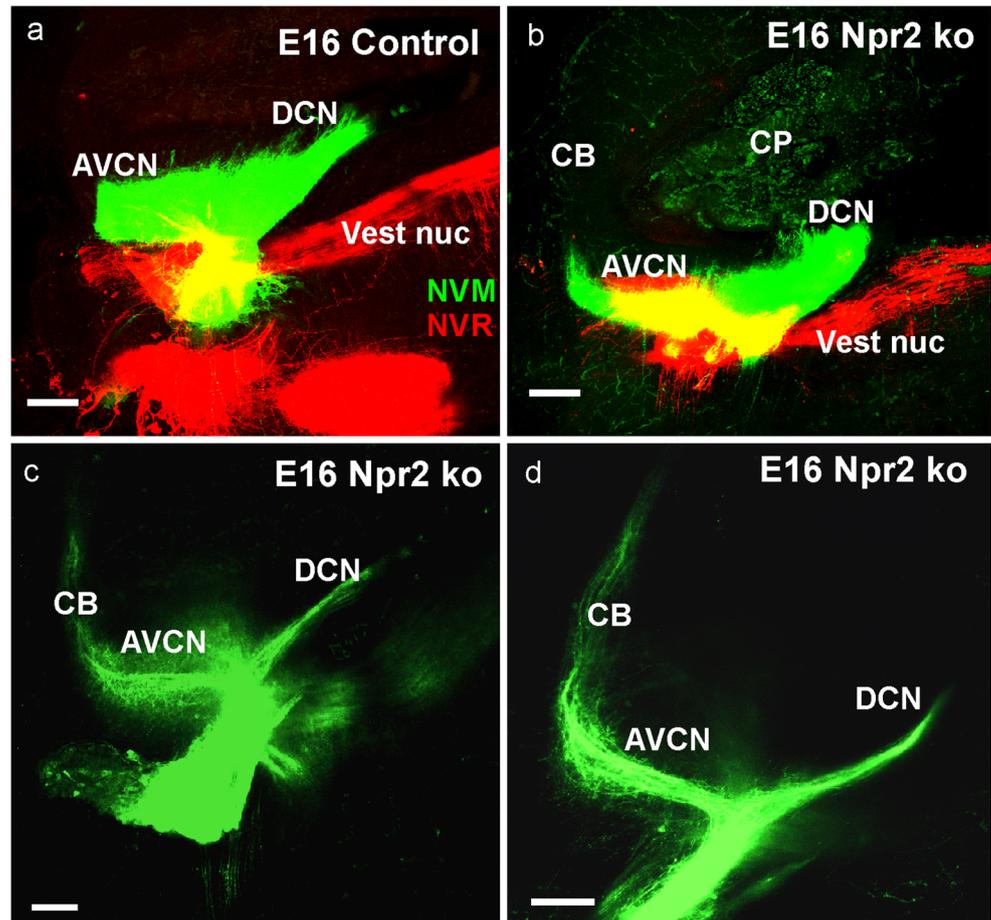
Fig. 2 Cerebellar afferent projections studied in 3 *Npr2* mutants. E13.5 *Npr2* mutants (**b**, **d**) showed more vestibular (NVR, red) and cochlear (NVM, green) afferents compared to littermate control animals (**a**, **c**) after comparable applications of dyes to the cochlea and vestibular organs (**a'**, **b'**). Higher power shows that many but not all, cochlear afferents go past the r1/2 border (indicated by dotted lines) in *Npr2* mutants (**d**, **d'**, **d''**) but not in control animals (**a**, **c**). Cochlear afferents project at this stage almost as far medial as vestibular afferents (**d''**). CB, cerebellum; CN, cochlear nucleus. Red blood cells (green dots) are highly excited by the 635 nm used for NVM. Orientation shown in (**a**) is identical for all images with dorsal up and anterior to the left. Bar indicates 100 μ m



(E18.5) the anterior extent of cochlear afferents labeled from base and apex in *Npr2* mutant and control animals. This measurement started at the anterior edge of the entering basal turn bundle to provide a uniform landmark for rostral fiber extension of both apical and basal turn spiral ganglion neuron afferents. A line was drawn along this rostral bundle

and another line at 90° was extended to the most rostral area that was showing above the background label. This likely overrates the mutant extension as it tended to be more gradual compared to the sharper end of the horizontal fibers in the control animals. We accepted a one-tailed difference as significant at $p > 0.05$.

Fig. 3 All 4 *Npr2* mutants show different cochlear projections to the cochlear nucleus complex. At E16.5, labeling of nearly all cochlear afferents by large insertions (a) resulted in fiber distributions to all parts of the cochlear nuclei. At this stage, the control shows more massive projections to the AVCN compared to DCN (a) whereas projections to AVCN and DCN in *Npr2* ko were more profound in DCN (b). Small applications to the cochlea revealed that the origin of cochlear afferents to the cerebellum is predominantly from the apex (c, d). Bar indicates 100 μ m



Results

Expression of *Npr2* and *CNP* as revealed by X-gal staining

Previous work has demonstrated the expression of *Npr2* and *CNP* in developing whole heads, demonstrating initial expression of *CNP* selectively in the even rhombomeres r2, r4 and r6 as early as E8.5 followed by expansion at E9.5 into the odd rhombomeres ahead of afferent growth from the entry points at even rhombomeres into odd rhombomeres (Ter-Avetisyan et al. 2018). Here, we expand this initial finding to E11.5 and E14.5 (Fig. 1). At E11.5, the first afferents of spiral ganglion neurons enter the hindbrain to form a distinct cochlear projection as early as E12.5 (Fritzsche et al. 2015) and this selective projection of spiral ganglion neurons to cochlear nuclei has been verified using *Gata3-taulacZ* (Fritzsche et al. 2006; Karis et al. 2001). At this stage, all inner ear-derived ganglia are strongly *Npr2* positive (Fig. 1a, c). *CNP* is particularly strongly expressed in the most dorsal part of r4 that is the entry point of vestibular and spiral afferents (Fig. 1b, d). Later stages show the *CNP* expression to be limited to the caudal part of the hindbrain with no expression in the cochlear nuclei (Fig. 1g). *Npr2* expression

remains strong in all sensory neurons of the ear and the endolymphatic duct (Fig. 1e). Comparison of mice heterozygous or homozygous for the *LacZ* reading frame replacement showed overall similar expression but somewhat reduced vestibular ganglia size (Fig. 1e, f). A particularly intensely stained population of seemingly vestibular neurons interconnected the vestibular and cochlear nerve foramina (arrowhead, Fig. 1e, f). In the brain, *Npr2* expression was detected in the cerebellum and the caudal hindbrain but not in the cochlear nuclei (Fig. 1h). Overall, these data demonstrate that there is a somewhat complementary distribution of *Npr2* and its ligand *CNP* suggesting an instructive function of *CNP/Npr2* signaling for *Npr2*-expressing neurons.

Dye tracing data

E13.5

Inner ear vestibular afferents reach the brain around E10.5 (Fritzsche 2003) and the cochlear spiral ganglion neurons reach the cochlear nucleus by E12.5 (Fritzsche et al. 2006, 2015; Karis et al. 2001; Yang et al. 2011). Afferent development follows with a 1–2-day delay in the cell cycle exit of vestibular and

spiral ganglion neurons (Matei et al. 2005; Ruben 1966), in line with progressive ventral to dorsal projection development of other cranial afferents over time (Fritzscht and Elliott 2017). At E13.5, vestibular and spiral afferents reach the entire rostro-caudal targets in the cochlear nucleus and vestibular nuclei, including the cerebellum for vestibular fibers (Fig. 2). Control animals show a sharp stop at the r1/2 boundary (Fig. 2) with no fiber extending beyond this rostral boundary of the cochlear nucleus complex (Farago et al. 2006; Fritzscht et al. 2006; Maricich et al. 2009). In contrast, all *Npr2* mutant animals had fibers labeled from the cochlea apex extending beyond this boundary to reach nearly as far to the midline of the cerebellum as vestibular afferents (Fig. 2d–d’). There was also a notable projection difference in vestibular fibers reaching far more medial across the cerebellum in *Npr2* mutants compared to control animals (Fig. 2). Combining these data suggests that in the absence of *Npr2* embryonic afferent growth is expanded toward r1 (Glover et al. 2018) with both vestibular and cochlear afferents projecting more extensively to the cerebellum. Whether this enhanced anterior expansion is a consequence of reduced branching allowing remaining fibers to extend further or indicates an impairment of axonal guidance remains unclear and requires future analysis using sparingly labeled individual fibers in later stages (Ter-Avetisyan et al. 2014). It should be noted that in *Npr2* and *CNP* mutants, some *TrkA*-positive neurites prematurely entered the spinal cord (Schmidt et al. 2002, 2009) suggesting a loss of responsiveness to repulsive guidance cues such as semaphorins known to play such a function in the ear (Gu et al. 2003). Previous work in a point mutation of *Npr2* using limited sparing spiral ganglion label analysis at E12.5 (Lu et al. 2011) and a more extensive analysis at E16.5 (Lu et al. 2014) did not report such fiber extensions. This could be due to a failure of labeling these fibers or due to the differences between the *Npr2*-cn mutant (Tsuiji and Kunieda 2005) they used, which is known to cause premature axonal ingrowth of some DRG neurons into the spinal cord compared to the *Npr2* null mutant (Ter-Avetisyan et al. 2014) used here.

E16.5

We next investigated whether the cerebellar projections of cochlear spiral ganglion afferents were transient or persisted by investigating the afferent organization in later stage animals. At E16.5, we found the similar expansion of cochlear afferents to the cerebellum in *Npr2* mutants only (Fig. 3b–d). In addition, we noted an overall asymmetry of cochlear afferent distribution to the different parts of the cochlear nuclei. Afferent projections were overall more profound in the large anteroventral cochlear nucleus in all control animals (Fig. 3a) whereas the proportion of afferent distribution was equal to or even inverse with more afferents projecting to the posterior part of the cochlear nucleus complex with occasionally enlarged afferents to the dorsal cochlear nucleus (Fig. 3b). In

an attempt to localize the origin of the cochleo-cerebellar fibers, we inserted very small amounts of dye into the extreme apex or base and obtained consistent labeling of cochlear afferents extending to the cerebellum only from the apex (Fig. 3c, d). These data suggest that perhaps more spiral ganglion afferents turn caudal, reducing the projection to the anteroventral cochlear nucleus. Those cochlear afferents projecting anteriorly from the apex expand beyond the r1/2 boundary into the cerebellum.

We next injected different colored dyes into the cerebellum and brain stem, respectively, to label afferents to the ear that project to these different parts of the hindbrain as previously described (Maklad et al. 2010). Such injections consistently labeled a differential afferent distribution to vestibular organs, indicating that different hair cell polarities in the utricle and saccule (not shown) project differentially to the brain. Overall, this segregation was maintained in the *Npr2* mutants with more fibers projecting to the cerebellum (Fig. 4a–d). These data suggest that there is no specific vestibular ganglion population targeted by the loss of *Npr2* but either population reaching the distinct polarities of the gravistatic sensory organs (Jiang et al. 2017) project nearly normally to their rostral and caudal targets as previously described (Maklad and Fritzscht 2003b; Maklad et al. 2010).

Consistent with cochlear spiral ganglion afferents reaching the cerebellum in particular from the apex (Fig. 3c, d), we also found afferents labeled from the cerebellum to pass through the ventral part of the cochlear nucleus complex but also labeled cochlear nucleus afferents from injections into the r6 area that spares cochlear afferents entirely in control animals (Fig. 4a). As suggested by the cochlear tracing (Fig. 3c, d), we could only label fibers in the ventral part of the cochlear nucleus complex after cerebellar injections. These data suggest that at least some spiral ganglion afferents develop collaterals that project past the rostral and caudal boundaries of the cochlear nucleus and some caudally projecting spiral ganglion afferents have at this stage-developed collaterals that expand to the anteroventral cochlear nucleus as also reported for other *Npr2* point mutants and control animals (Goodrich 2016; Lu et al. 2014; Muniak et al. 2016).

We next investigated the origin of these unusual cochlear nucleus projections by flat mounting the cochlea after lateral cerebellar as well as r6 dye insertions. While such injections do not label any cochlear afferents in control animals (Fig. 4b’), we found many basal turn spiral ganglion neurons to be labeled after r6 dye insertions in the *Npr2* null mutant (Fig. 4b, b’’) whereas more apical spiral ganglion neurons were labeled after cerebellar injections (Fig. 4e). Importantly, spiral ganglion neurons projecting either rostrally or caudally past the cochlear nucleus complex were never double labeled, suggesting that each of these afferents is likely forming a single unbranched central fiber consistent with our analysis on initial afferent projections (Ter-Avetisyan et al. 2014).

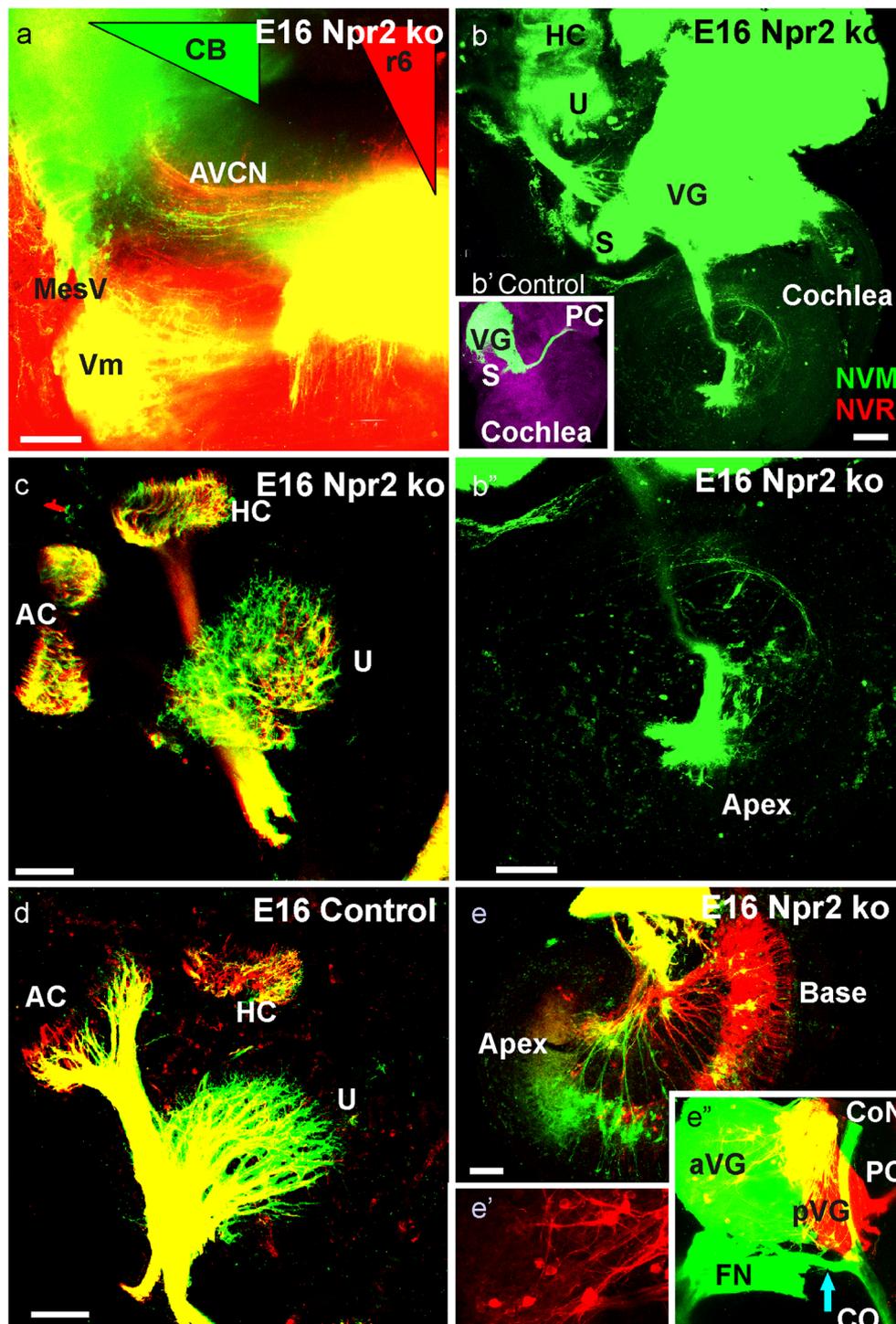


Fig. 4 All 4 *Npr2* mutants have altered cerebellar projections. These E16.5 *Npr2* mutant and control data show the injection of different colored dyes in the cerebellum (NVM, green triangle in **a**) and rhombomere 6 (NVR, red triangle in **a**). Cochlear afferents (labeled green in AVCN) became retrogradely labeled from the cerebellum and r6 cochlear afferents (labeled red) from the r6 injection only in mutants. These dye insertions labeled vestibular fibers to the vestibular organs in *Npr2* mutants (**c**) and control animals (**d**). While the overall segregation was comparable, fibers from the cerebellum occupied a larger area in the utricle (compare **c**, **d**). Only *Npr2* mutants had spiral ganglion cells labeled from r6 or cerebellar injections (**b**, **b'**) whereas no such fibers

to the cochlea are found in control animals (**b'**). Cerebellar labeling resulted in prominent apical labeling (**e**) that was exclusive from the apex in more medial cerebellar injections (**b**, **e**). In contrast, r6 injections labeled only basal turn spiral ganglion neurons in some animals (**e**). Injections into vestibular endorgans labeled the posterior vestibular ganglion (pVG) from the posterior canal crista (PC) and anterior vestibular ganglion (aVG) from the utricle. Only mutants showed massive labeling of a fiber bundle expanding from the aVG to the modiolus to bifurcate toward the cochlear nuclei and the cochlea (**e'**). Note the distributed and highly branched vestibular neurons in the aVG (**e''**). Bar indicates 100 μ m

We next aimed to investigate whether the most medial cerebellar projections are even more restricted in their origin or derive from widespread distributed spiral ganglion neurons. Such near midline cerebellar dye injections revealed an unusual projection of only a limited number of a very apical population of spiral ganglion neurons that projected to the apex of the cochlea (Fig. 4b, b''). This filling of only apical spiral ganglion neurons is consistent with data after point applications to the apex showing more profound anterior cochlear nucleus projections that always extended far into the cerebellum (Fig. 3c, d). Given that we found some spiral ganglion neuron labeling after cerebellar applications (Fig. 4e), we next investigated if perhaps vestibular neurons project to the cochlea as recently reported in *Neurod1* (Macova et al. 2019) and *Foxg1* mutant mice (Pauley et al. 2006). Our data show both an unusual distribution of vestibular ganglion neurons across the vestibular ganglion complex and a bundle of fibers passing ventral to the vestibular ganglion to bifurcate upon reaching the auditory afferents to extend toward the cochlear nucleus and the cochlea (Fig. 4e, e'). Such additional branching was also obvious in vestibular ganglion neurons labeled from the posterior canal crista application in the anterior vestibular ganglion (Fig. 4e, e', e'').

In summary, tracing data show that mostly apical spiral ganglion neurons project to the cerebellum and have a bifurcation to reach also anterior vestibular sensory epithelia.

E18.5

At this stage, the cochlear nucleus could be fully analyzed in terms of cochleotopic projections using differently colored dye injections into the base and apex as previously described (Fritzsch et al. 2016; Macova et al. 2019; Xiang et al. 2003; Yang et al. 2017). These tracings revealed identical cochleotopic trajectories of basal and apical fiber bundles within the cochlear nerve as previously described (Macova et al. 2019; Xiang et al. 2003) but also revealed profound differences between the anterior projections to the AVCN and the posterior projections to the PVCN and DCN. Whereas the projections to the PVCN and, to a lesser degree, the DCN were enlarged relative to that of control mice (Fig. 5a, b), the anterior extent of the projection to AVCN in *Npr2* null mice was reduced (Table 1). Instead of extending for 333 μm (± 25) into the dorsal part of the AVCN, only very few fibers passed anteriorly for only 63 μm (± 7) (Fig. 5a'', b''). Furthermore, the base and apex showed differences with apical fibers projecting more profoundly to the DCN with expansions across the basal fibers beyond the limited expansion found in control animals at this stage (Fig. 5a', b'). Interestingly, the apical insertions resulted in many more fibers to the AVCN compared to basal insertions but fibers extended only about half as far anteriorly with a sharp reduction in longitudinal extent in more dorsally located fibers

(Table 1). Quantification of rostral fibers showed a clear and significant difference between the control and mutant for both basal and apical projections ($N=6$ for control and *Npr2* null mice, $p > 0.05$) indicating that anterior growth of spiral ganglion axons was reduced in *Npr2* null mice at this stage. In contrast, projections of the posteroventral cochlear nucleus were more profound in all preparations (Fig. 5). We could not trace fibers at this stage reaching the cerebellum and injections into the cerebellum failed to label any spiral ganglion neurons suggesting that all of these fibers had either withdrawn, or the parental neurons had died.

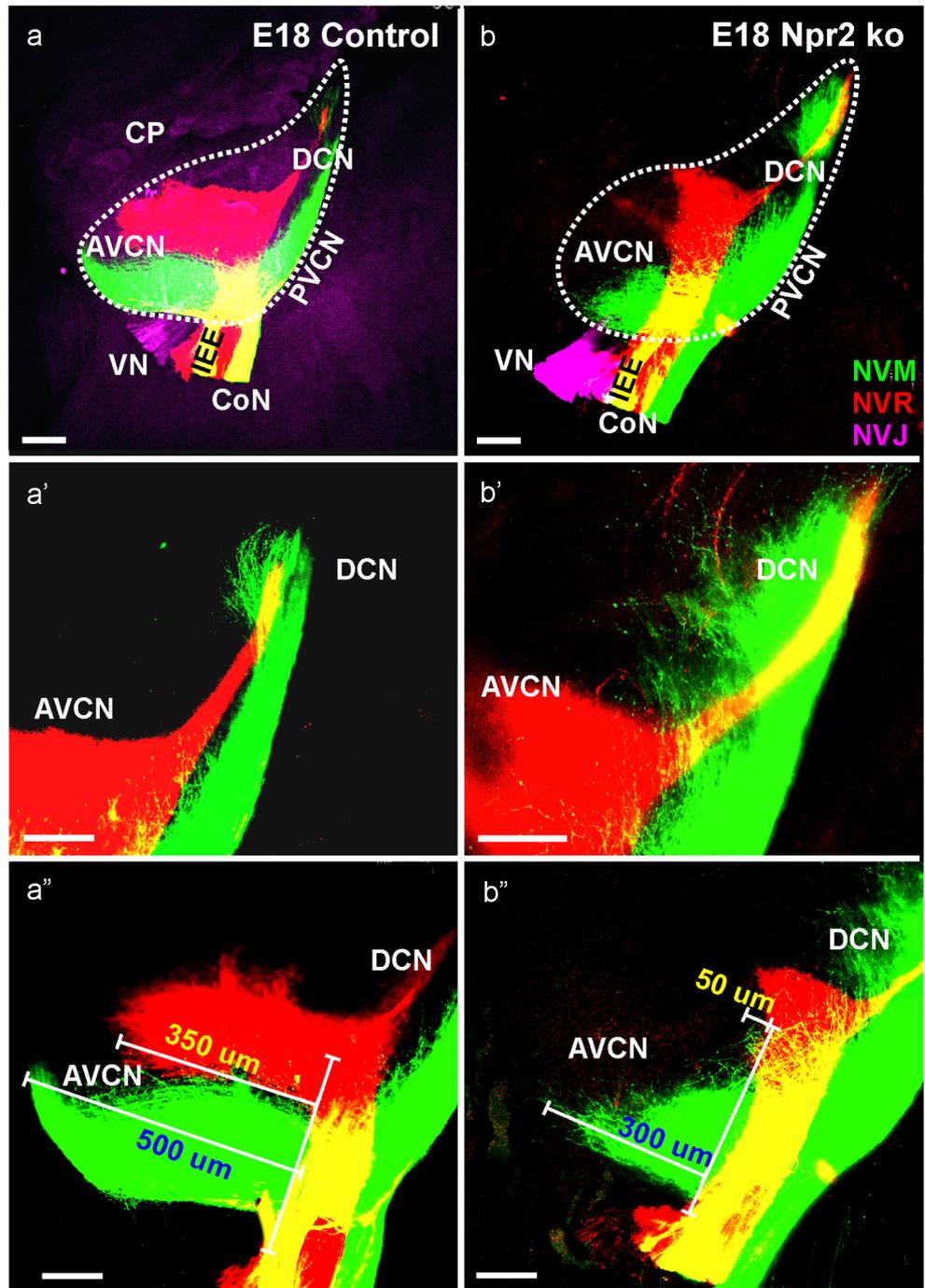
We next wanted to verify if the unusual fibers interconnecting vestibular endorgans with the cochlea remained and injected dye into the cochlear apex, base and vestibular organs (Fig. 6a, c; green, red, lilac, respectively). In most animals, we found comparable small projections emanating from cochlear labeling toward the vestibular ganglion (Fig. 6a, c). However, in three of three mutant mice, we labeled more excessive fibers and cells in this cochlear vestibular anastomosis (Fig. 6e; CVA). Preparing the vestibular organs in one of these animals revealed several fibers emanating from a ganglion formed along this anastomosis to reach the three anterior vestibular endorgans, the utricle, horizontal and the anterior canal (Fig. 6f). These data suggest that many but not all of the unusual vestibulo-cochlear connections are eliminated in older embryos. We also investigated the collaterals reaching from the PVCN/DCN to the AVCN by injecting dye into the DCN and cerebellum (Fig. 6b, d). This approach labeled expectedly some fibers in the restiform body adjacent to the cochlear nucleus in both control and mutant mice but no cochlear nucleus afferents, suggesting that the expansion to the cerebellum is reduced or eliminated at this stage. Most importantly, such an approach labels multiple parallel fibers covering all of AVCN in control animals and shows only limited expansion to the more posterior half of the AVCN in mutants (compare Fig. 6b, d) consistent with our labeling from cochlear injections (Fig. 5).

We also investigated the insertion sites in the cochlea not only for the precision of injections as intended but also for unusual interconnections of spiral ganglion neurons between the base and the apex as recently reported in a mutant with a near complete loss of cochleotopic projections (Macova et al. 2019). Our data showed virtually identical labeling of a gradual blending of basal- and apical-labeled neurons and fibers between control and *Npr2* mutant mice (Fig. 6g, h) giving no indication of either differential migration or multiple connections of a given spiral ganglion neuron within the cochlea.

P2

After birth, the differences in control (Fig. 7a, b) and mutant animals became more obvious. Neither small (Fig. 7a) nor very large (Fig. 7b) injections in control animals showed any

Fig. 5 Cochleotopic projections reveal profound differences at E18.5 in all six Npr2 mutants studied. Triple dye injections into the vestibular organs (NVJ, lilac), apex of the cochlea (NVM, green) and base of the cochlea (NVR, red) show profound afferent distribution differences between control (a, a', a'') and Npr2 null mutants (b, b', b'') but show no effect on overall cochleotopic dorsoventral patterning. Most notable were reduced basal projections to the AVCN (a, b, a'', b'') and expanded projections of the apex to the PVCN and DCN (a', b'). Measuring the extension of afferents showed a significant reduction in the length of apical or basal cochlear projections in Npr2 ko mice to the AVCN (Table 1). NVJ is excited at 488 nm and thus results in background fluorescence that was used to outline the cochlear nuclei (a) and also labels the choroid plexus (CP in a). Bar indicates 100 μm



deviation from the cochleotopic projection pattern as previously reported in neonates of a mutant with a disorganized cochleotopic projection (Macova et al. 2019). While Npr2 mutants also showed well-organized fiber stratification after large

(Fig. 7c, d) or small (Fig. 7e) dye applications, there were noticeable differences in the labeling of afferents and their rostro-caudal extent. All mutants showed a profound projection to the PVCN from the apex that was nearly absent in the basal turn

Table 1 Rostral extent of spiral ganglion afferents from apical and basal turn bundle (length in μm)

	Control	Mean + SD	Npr2 ko	Mean + SD
Apex	468, 512, 488, 532, 490, 500	498 + 23	285, 300, 412, 375, 423, 328	354 + 30
Base	325, 346, 385, 350, 304, 290	333 + 25	75, 55, 50, 65, 45, 85	63 + 7

projections. Conversely, the projection to the AVCN was reduced in a number of fibers and their anterior extent (Fig. 7c–e). In fact, in some cases, we obtained, despite massive PVCN labeling from the apex, hardly any fibers to the AVCN (Fig. 7e). Importantly, fibers did not show the parallel fiber bundles of cochleotopic projections (Muniak et al. 2016) but showed a more random distribution as in *Neurod1* mutant mice (Macova et al. 2019) and as was previously reported based on single fiber tracings (Lu et al. 2014). While afferents in control animals showed multiple parallel fibers with short collaterals (Fig. 7a), fibers in the *Npr2* null mutant were erratic and random with many branches suggesting them to be collaterals, comparable to those expanding to the ventral part of the DCN (Fig. 7c–e) instead of primary afferents. In three out of six cases with anterior vestibular injections, we obtained labeling not only of vestibular afferents entering with the vestibular nerve but also of some afferents entering with the cochlear nerve (Fig. 7f), consistent with the vestibulo-cochlear anastomosis shown in younger animals (Figs. 4 and 6).

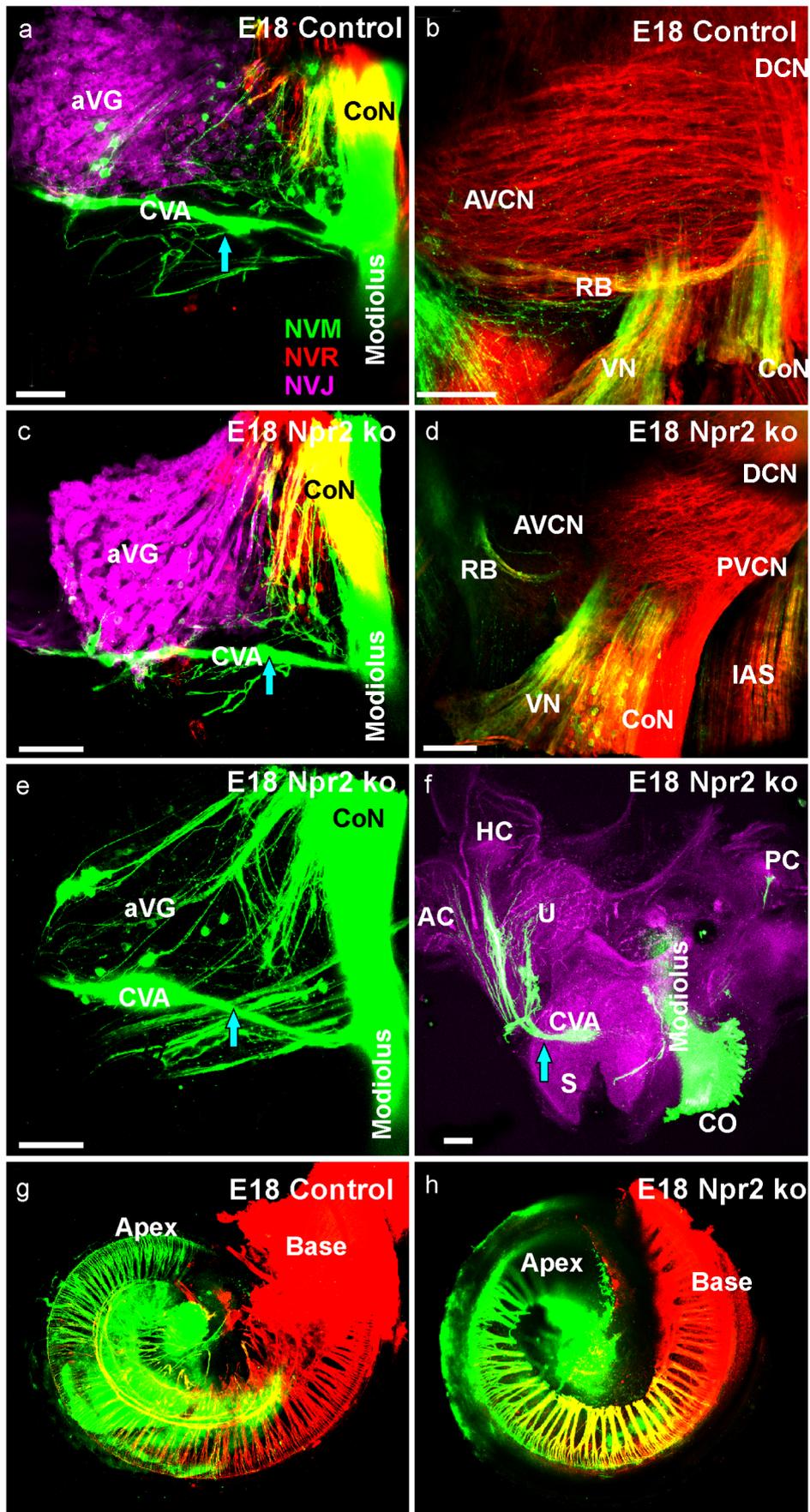
P7

Some *Npr2* mutants can survive for some time (Wolter et al. 2018) and we harvested two mutants at P7 to investigate late stages in cochleotopic projections. The data collected on 4 half brains show features mostly consistent with the earlier data as far as apical projections are concerned. Clearly, expansion of anterior directed fibers is truncated to about 2/3 the length of control animals (Fig. 8). Both, control and *Npr2* mutants, show comparable expansion to the DCN and PVCN. In particular, the projection of the apex to the most ventral part of the DCN showed a comparable expansion at this stage (Fig. 8a', b') with fibers reaching the area of the octopus cells (Malmierca 2015; Osen 1969), which were occasionally transcellularly labeled by the lipophilic dye. The more profound projection of the apex to the PVCN as compared to the AVCN found in earlier stages (Figs. 5 and 7) remained, suggesting that caudal branches of spiral ganglion neurons are more prominent in *Npr2* mutant mice. Beyond the obvious reduction of the apical projection to the AVCN was a blurring of the dorsal boundary of the projection area. In control animals, there is space left between the basal (red) and apical (green fibers; Fig. 8a, a"). In contrast, in the mutants, there is an expansion of branches from primary apical fibers to fill in the gap to basal fibers with the most profound expansion in the posterior part of the AVCN (Fig. 8b, b"). These data imply a differential overlap of cochleotopic projection by branches mostly from apical fibers in the AVCN that could affect in a subtle way tonotopic aspects of hearing.

Discussion

Our data extend previous descriptions of the *Npr2* phenotype of the inner ear and in particular cochlear afferent projections (Lu et al. 2011; Lu et al. 2014; Ter-Avetisyan et al. 2014) that focused on erroneous single fiber branching with little data provided on the development of the cochleotopic projection (4 *Npr2* point mutants each at E16.5 and P14) in addition to physiological work (Lu et al. 2014). Auditory physiology was recently extended to the genetically targeted *Npr2* null mutants we analyzed here (Wolter et al. 2018). Our data show for the first time an unusual cochlear afferent expansion past the rostral and caudal boundaries of the cochlear nuclei in early embryos followed by a base to apex progressing reduction of AVCN projections and expansion of PVCN projection (Figs. 2, 3, 4, 9). Our data show that these expansions form early and suggest that extension of the single unbranched afferent in *Npr2* null mutants may drive the extensive longitudinal growth that normally may be more truncated due to the initial branching, assuming comparable growth potential in control and *Npr2* null spiral ganglion neurons. Alternatively, cochlear afferents of *Npr2* mutants might have lost the ability to read a r1/2 stop signal that normally restricts anterior expansion of cochlear afferents unless the Hox code of rostral rhombomeres has been manipulated (Oury et al. 2006). However, given that we also obtained a caudal expansion of a different set of spiral ganglion neuron afferents, we suggest that the expansions reflect mostly additional longitudinal growth possibly due to the absence of primary branching.

The differential origin of caudally projecting versus extensively rostrally projecting spiral ganglion neurons is a novel finding. Obviously, only the most apical spiral ganglion neurons project the most like vestibular afferents suggesting that these fibers behave without *Npr2* as a mix of spiral ganglion neurons (cochlear nucleus projection) and vestibular neurons (projection to the cerebellum). We here provide for the first time evidence for an unusual connection between the auditory afferents and vestibular endorgans, the cochleo-vestibular anastomosis (Figs. 1, 4, 6, 7). Past work on *Neurod1* mutant mice has shown that a spiro-vestibular ganglion forms in these mutants due to migration of spiral ganglion neurons to mix with vestibular neurons (Jahan et al. 2010a; Macova et al. 2019). These mixed ganglia also have mixed peripheral fibers in the cochlea and a mixed central projection of vestibular and spiral ganglion neurons. In contrast, our data on *Npr2* null mutants do not show such mixing of spiral and vestibular ganglia and a limited projection of some vestibular fibers to the cochlear nuclei (Fig. 7). Importantly, beyond this limited overlap, our data show no blurring of basal and apical projections within the cochlear nerve as recently shown for the *Npr2*-cn mouse mutant (Lu et al. 2014) but basal and apical fibers remained always discrete within the nerve (Figs. 3, 5, 6, 7, 8). In addition, we never found basal turn afferents to be



◀ **Fig. 6** Triple labeling of cochlear apex (NVM, green, **a, c, e, f, g, h**), cochlear base (NVR, red, **a, c, e, f, g, h**) and utricle (NVJ, lilac **a, c**) at E18.5 show a cochlea-vestibular anastomosis of fibers. This anastomosis extends from the apex to the anterior vestibular nerve foramen and is in three cases of *Npr2* mutants more profound (**c, e**) compared to littermate control (**a**). In all three mutants, fibers can be traced to utricle, anterior and horizontal canal crista (**f**) not found in control littermates. Dye injection into the cerebellum (green in **b, d**) and DCN (red, **b, d**) labels most fibers in the entire AVCN in control (**b**) but not *Npr2* mutant mice (**d**). Afferents in the restiform body (RB) are equally labeled and so are vestibular afferents exiting the brain stem. Dye injections into the cochlea reveal a comparable distribution of labeled neuronal profiles in control (**g**) and *Npr2* mutant (**h**). Lilac in (**f**) is the background fluorescence. Bar indicates 100 μm in all images

most ventro-caudal relative to apical afferents in the PVCN as reported (Lu et al. 2014). Whether these differences between our findings and previous work are due to technical issues (overlapping injections in base and apex, co-labeling of vestibular fibers after basal turn injection) or relate to the different genotype (*Npr2* deletion versus *Npr2* point mutation) remains to be resolved. It should be mentioned that the previous work published no evidence that the injection was accomplished as intended by showing the injected cochlea as we provide here (Fig. 6). Interestingly, the added schematics show segregated fibers in the modiolus and cochlear nerve, identical to our findings.

Our data go beyond previous findings on other mutants by demonstrating a unique connection between the two ear foramina for the anterior vestibular nerve and the posterior vestibular/cochlear nerve (Figs. 4 and 6) and also it appears that many spiro-vestibular afferents in *Neurod1* mutants project along that pathway (Macova et al. 2019). It remains unclear if *Neurod1* regulates transcription of *Npr2* and whether the absence of *Npr2* alters interactions with neural crest-derived Schwann cells known to affect peripheral afferent distribution (Mao et al. 2014). Clearly, data on cGKI null mice are needed to fully reveal the signal/receptor deletion effects in the cochleotopic projection development.

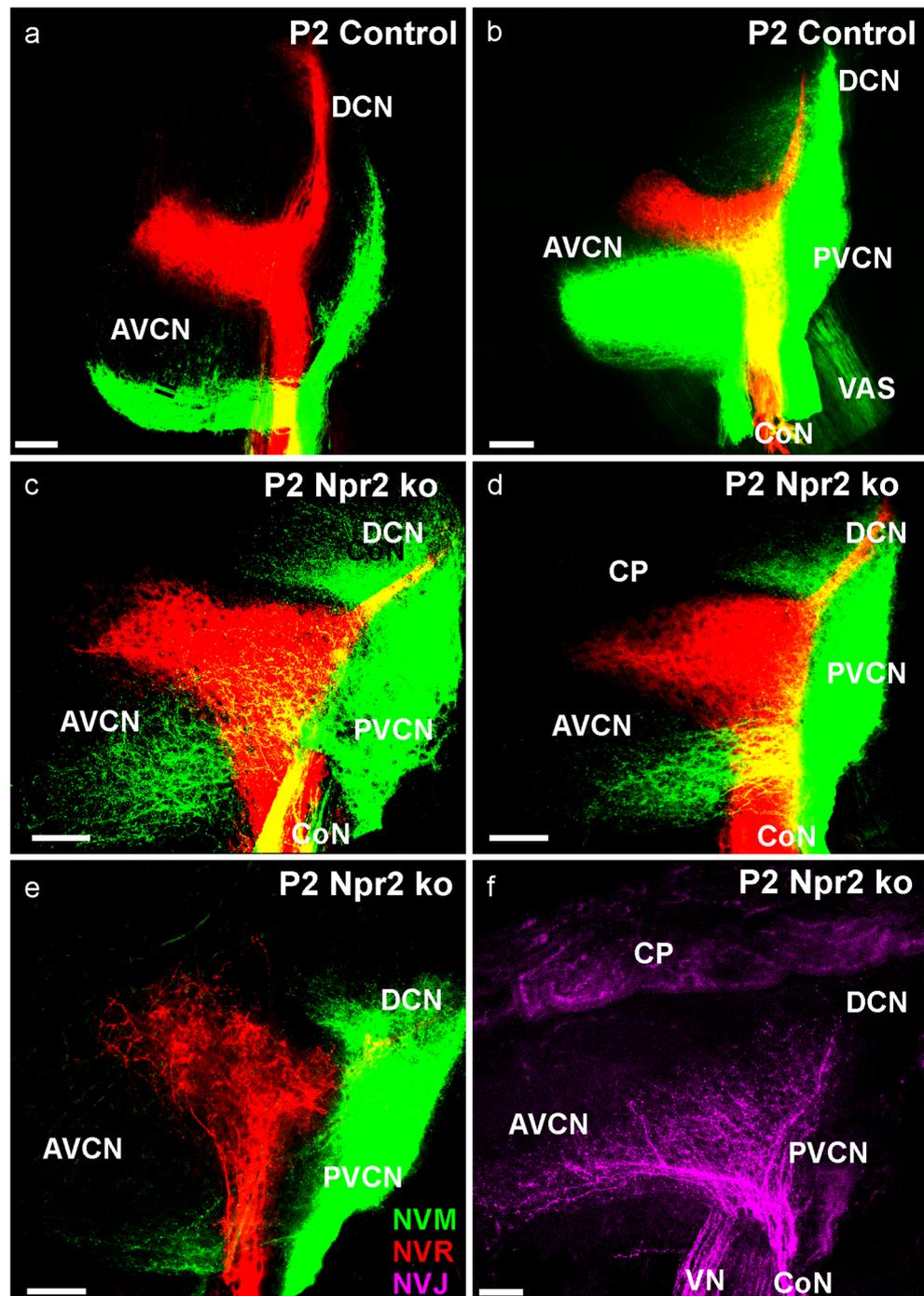
While early embryonic stages show overshooting projections, later stages show a mix of reduction and expansion of different longitudinal spiral ganglion populations to different parts of cochlear nuclei. Basal fibers show reduced growth to the AVCN combined with normal DCN projection whereas apical fibers show reduced AVCN combined with expanded PVCN and premature DCN projections. These differences in the behavior of apical and basal fibers have previously been noted for the DCN projection in another *Npr2* mutant line (Lu et al. 2014) and are particularly obvious in *Prickle 1* mutants (Yang et al. 2017). In the latter mutants, the basal projections are nearly normal whereas apical afferents develop collaterals that expand across the entire cochlear nucleus complex (Yang et al. 2017) indicating radically different effects of the same mutation to different spiral ganglion neurons coding different

frequencies in line with the observed expansion of apical fibers in the caudal part of the AVCN we show here (Figs. 7 and 8). This expansion of apical afferents is particularly obvious in the consistent posterior projection expansion but variably reduced in the anterior projection (Fig. 6; Table 1). Somewhat similar effects were previously reported as intensity differences that were claimed to be more profound for the apex compared to the base (Lu et al. 2014). Since no data were provided in this work for the stage we analyzed here we do not know how the dynamics of fiber expansion and retraction compares to our mutant (Fig. 9).

Cochleotopic projection is essential for the normal function of frequency-specific hearing (Muniak et al. 2016) and complete loss of tonotopic organization results in blurred tuning that cannot be corrected by activity (Macova et al. 2019). Previous work has indicated that some blurring of this frequency-specific projection happens in certain areas of the cochlear nucleus complex in mice harboring the *Npr2*-cn point mutation (Lu et al. 2014). Our data do not confirm such a blurring in *Npr2* knockout mutant primary afferents (Fig. 9) but show a much more dynamic effect of early fiber overshooting followed by later retraction of some branches to some parts of the cochlear nucleus combined with the expansion of other branches. These data imply that *Npr2* does not only exert a profound initial effect on ingrowing afferents (Schmidt et al. 2007; Ter-Avetisyan et al. 2014) but may play a second role in differential modeling of fibers to different parts of their targets. The secondary expansion of apical afferents seemingly has little effect on tonotopic performance (Lu et al. 2014) compared to the massive tuning distortion resulting from disorganization afferents of *Neurod1* conditional deletion mutants (Macova et al. 2019). Further work is needed to reconcile the minor physiological differences of the two *Npr2* mutants (Lu et al. 2014; Wolter et al. 2018) with the more obvious tonotopic differences we found here compared to the *Npr2* point mutant.

Finally, while *Npr2* regulates bifurcation of afferents upon entering the brain stem or spinal cord (Ter-Avetisyan et al. 2014; Tröster et al. 2018), we here describe unusual peripheral branching at the ear that may represent an exaggeration of a transient natural branching, possibly regulated by *Neurod1* (Macova et al. 2019). Whether other developing sensory processes exhibit not only the truncation of central branches but an exaggeration of peripheral branches remains to be seen. It is also unclear what other factor(s) may compensate for *Npr2* in some inner ear neurons (Booth et al. 2018) and why there is a graded effect of anterior versus posterior branch expansion in base versus apical spiral ganglion neurons (Figs. 7 and 9). Obviously, the overshooting and retraction in anterior projecting branches combined with more profound posterior projections correlates with the differential distribution of the *Npr2* ligand CNP that is more profound in caudal rhombomeres at later stages but initially expands to r2

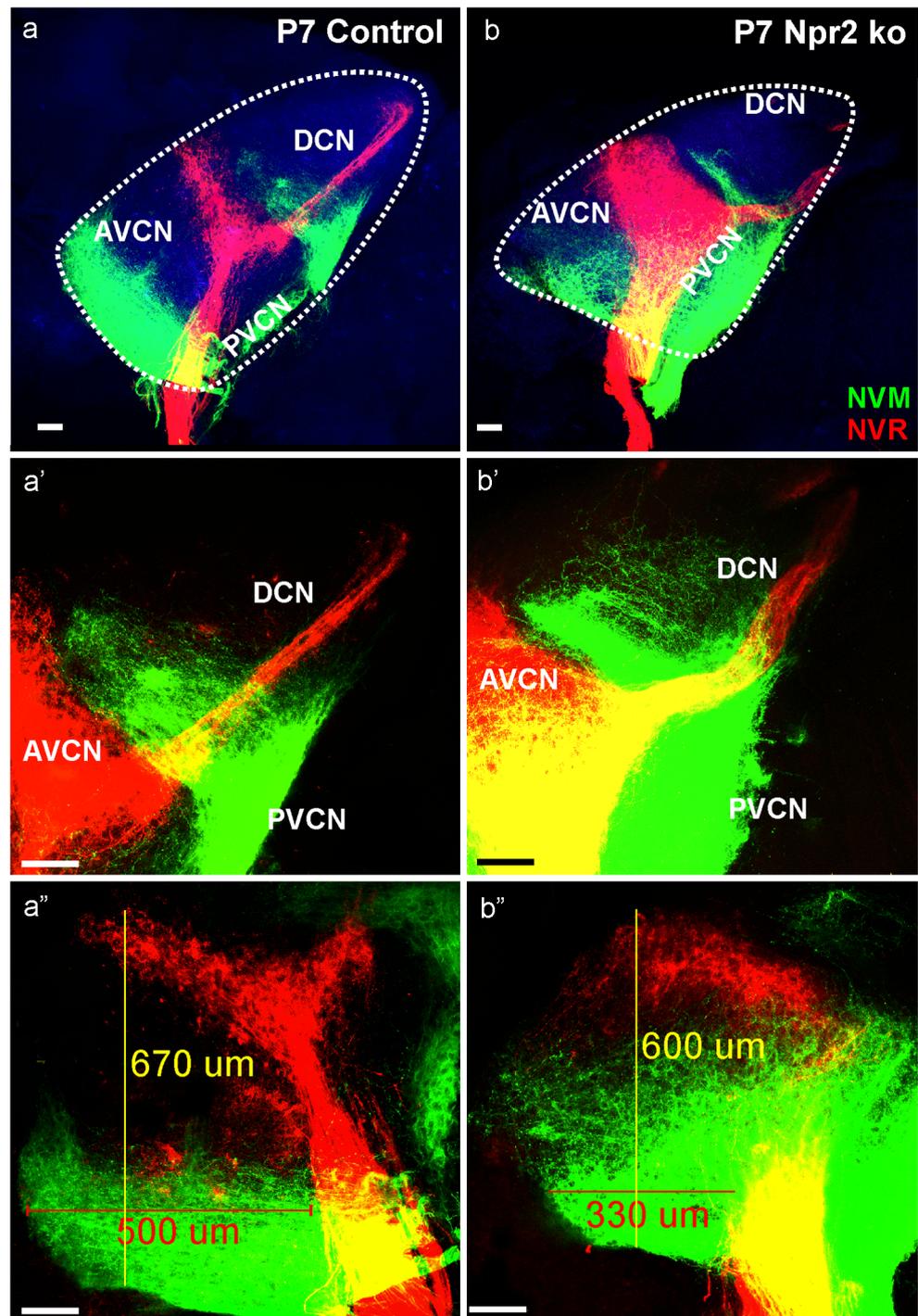
Fig. 7 Small (a) or large (b) dye applications to the base (NVR, red) and apex (NVM, green) show cochleotopic projections expanding the entire length of the cochlear nucleus complex in control animals. All 3 mutants shown here display a more restricted fiber expansion of apparent collaterals to the AVCN (c, d, e) but a more profound projection of apical cochlear afferents to the PVCN. Only in 3 of 6 mutants could we trace vestibular fibers from the utricle not only through the vestibular nerve to the vestibular nucleus but also through the cochlear nerve to the most ventral aspect of the cochlear nucleus complex. Note the branches off the stem fibers in vestibular fibers entering into the ventral part of the AVCN. A single optical plane reveals a highly disorganized fiber projection that is not organized in parallel lines (e). Bar indicates 100 μm



(Fig. 1b, d, g). Investigating inner ear projections in CNP mutants could reveal causality of this observation if *Npr3* receptor is expressed in inner ear sensory neurons that can read that gradient. Clearly, more data on quantitative expression variation on possible co-factors in spiral ganglia are needed to mechanistically explain this continued variation using quantifiable in situ hybridization approaches (Kersigo et al. 2018) combined with fiber tracing.

Comparative implications Previous mutant analysis has shown that the cochlear apex and base respond differentially to the loss of *Lmx1a* (Nichols et al. 2008), *N-Myc* (Kopecky et al. 2011) and *Neurod1* (Jahan et al. 2010b) implying that the apex is part of an ancestral vestibular sensory organ, the lagena of tetrapods (Schultz et al. 2017), incorporated into the expanding eutherian organ of Corti (Fritsch et al. 2013). The

Fig. 8 Detailed analysis of two P7 Npr2 mutant (**b, b', b''**) and control littermate (**a, a', a''**) shows overall cochleotopic matching projections with apical fibers (NVM, green) projecting ventrally and basal fibers (NVR, red) projecting dorsally in the anteroventral cochlear nucleus (AVCN). However, while fibers in control animals remain restricted to narrow bands, in particular, cochlear afferents from the apex expand to reach almost the basal turn projection in the mutant (compare **a'', b''**). Projection of the basal turn reaches unbranched to the dorsal cochlear nucleus (DCN, **b'**) whereas basal turn neurons show a profound collateral expansion along the ventral border of the DCN with the PVCN, which is comparable in Npr2 mutant (**b, b'**) and control littermates (**a, a'**). In addition to the AVCN dorsal expansion, the anterior expansion of the basal turn is truncated to about 2/3 of the rostral extent (in the presented case from 500 μm to 330 μm). Note that branches emerge from the apical fibers to nearly fill the gap between apical and basal tonotopic tracts (**a'', b''**). Blue indicates the 488-background fluorescence that was used to outline with a dotted line the cochlear nuclei complex. Bar indicates 100 μm



data presented here show that only the most apical spiral ganglion neurons can behave nearly identical to vestibular afferents by projecting to the cerebellum (Figs. 3 and 4) and form an anastomosis with fibers that apparently branch to reach the apex of the cochlea and project to the most ventral aspect of the cochlear nuclei. We assume that these mixed spiro-vestibular

neurons are the ones that form the projection to the cerebellum indicating that loss of Npr2, possibly regulated by Neurod1 (Pataskar et al. 2016), may unmask an ancient vestibular-auditory connection. It should be noted that no such anastomosis nor cochlear nucleus projections were described in chicken lagena projections (Kaiser and Manley 1996; Mahmoud et al.

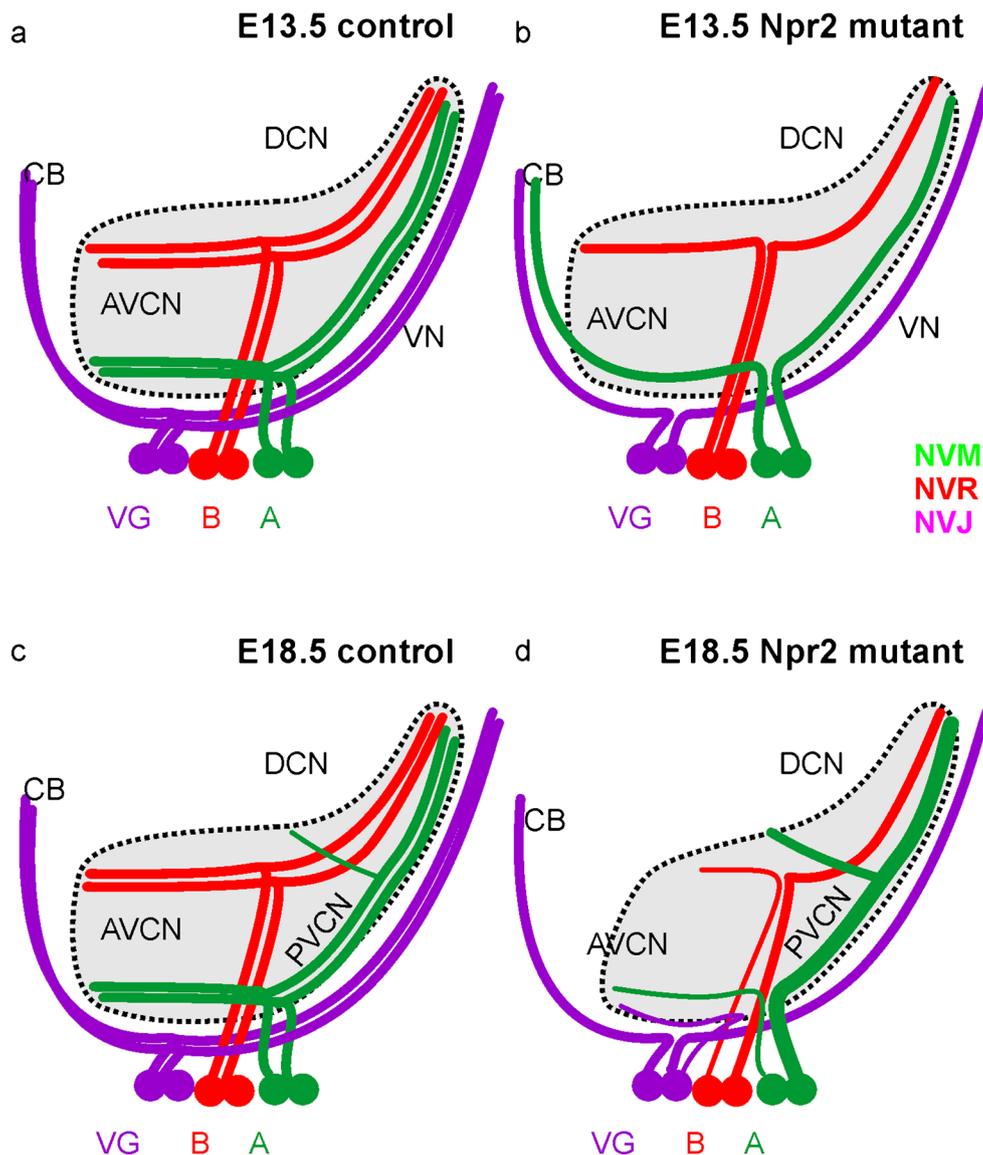


Fig. 9 This summary diagram shows the differences in afferent projections of spiral and vestibular ganglion neurons in control and *Npr2* mutant mice at two different stages (E13.5, **a, b**; E18.5, **c, d**). For each projection, we depict only two representative neurons for the vestibular ganglion (VG), the base (B) and the apex (A) of the spiral ganglion. Color coding is consistent with the color rendering of the dye tracing. All neurons project in parallel and cochleotopic to both anterior and posterior aspects of the vestibular and cochlear nuclei in control animals with a small, late-forming branch of only the apical fibers to the octopus region. In mutants, there is an initial overshooting of

projections from apical spiral ganglion neurons to the cerebellum whereas in later mutants, the projection of the apex is expanded to the PVCN and DCN (thick green fiber) and reduced in rostro-caudal extent and projection size (thin fibers) to the anteroventral cochlear nucleus. Note also the expansion of vestibular fibers to the most ventral aspect of the cochlear nucleus complex only in mutants. Thus, while the overall cochleotopic projection is retained in *Npr2* mutants, in particular, the apical afferents show unusual expansions and contractions to different parts of the cochlear nuclei

2013) but more refined connectational analysis in developing chicken and in particular, monotremes as the only mammal that retain the lagena (Schultz et al. 2017) with the above information in mind, are clearly warranted.

In summary (Fig. 9), our data on *Npr2* knockout mutants are in line with physiological data (Lu et al. 2014; Wolter et al. 2018) and reveal no major tonotopic disorganization as previously claimed based on limited

cochleotopic tracing experiments (Lu et al. 2014). However, we reveal surprising dynamics of cochlear afferent branching to various subnuclei, confirming and extending previous indications of reduced anteroventral projection and expanded posteroventral projection (Lu et al. 2014). In addition, we find unusual connections of only the apical spiral ganglion neurons with the cerebellum as well as limited projection of vestibular neurons to the cochlear nuclei.

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Compliance with ethical statements

Conflict of interest The authors declare that they have no conflict of interest.

Ethical approval All applicable international, national and/or institutional guidelines for the care and use of animals were followed.

Abbreviations AC, anterior canal crista; AVCN, anteroventral cochlear nucleus; CB, cerebellum; CN, cochlear nucleus; CO, cochlea; CoN, cochlear nerve; CP, choroid plexus; CVA, cochleo-vestibular anastomosis; DCN, dorsal cochlear nucleus; ED, endolymphatic duct; FN, facial nerve; FP, floor plate; HC, horizontal canal crista; MesV, mesencephalic trigeminal projection; NG, nodose ganglion; NVJ, NeuroVue Jade; NVM, NeuroVue Maroon; NVR, NeuroVue Red; PC, posterior canal crista; PG, petrosal ganglion; PVCN, posteroventral cochlear nucleus; r, rhombomere; RB, restiform body; S, saccule; SG, spiral ganglion; U, utricle; VAS, ventral acoustic stria; (a, p) VG, (anterior, posterior) vestibular ganglion; VN, vestibular nerve; V, trigeminal ganglion; Vm, trigeminal motor neurons; VIII, octaval ganglion; VII, facial (geniculate) ganglion; IX, proximal glossopharyngeal ganglion; X, proximal vagal ganglion; XI, transient accessory ganglia

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