

# Role of Bone Morphogenetic Proteins on Cochlear Hair Cell Formation: Analyses of *Noggin* and *Bmp2* Mutant Mice

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The mammalian organ of Corti of the inner ear is a highly sophisticated sensory end organ responsible for detecting sound. *Noggin* is a secreted glycoprotein, which antagonizes bone morphogenetic proteins 2 and 4 (*Bmp2* and *Bmp4*). The lack of this antagonist causes increased rows of inner and outer hair cells in the organ of Corti. In mice, *Bmp2* is expressed transiently in nascent cochlear hair cells. To investigate whether *Noggin* normally modulates the levels of *Bmp2* for hair cell formation, we deleted *Bmp2* in the cochlear hair cells using two *cre* strains, *Foxg1<sup>cre/+</sup>* and *Gfi1<sup>cre/+</sup>*. *Bmp2* conditional knockout cochleae generated using these two *cre* strains show normal hair cells. Furthermore, *Gfi1<sup>cre/+</sup>;Bmp2<sup>lox/-</sup>* mice are viable and have largely normal hearing. The combined results of *Noggin* and *Bmp2* mutants suggest that *Noggin* is likely to regulate other *Bmps* in the cochlea such as *Bmp4*. *Developmental Dynamics* 239:505–513, 2010. Published 2010 Wiley-Liss, Inc.<sup>†</sup>

**Key words:** *Noggin*; *Bmp2*; *Foxg1-cre*; *Gfi1-cre*; hair cell; cochlea; mouse; hearing

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## INTRODUCTION

The organ of Corti of the mammalian cochlear duct consists of one row of inner hair cells (IHCs), three rows of outer hair cells (OHCs), and several specialized types of supporting cells such as Deiters' and pillar cells. Cells within the prosensory domain that give rise to the organ of Corti, start to exit from the cell cycle and express p27Kip1 at embryonic day 12.5 (E12.5; Chen and Segil, 1999). Proper differentiation of the various cell types within the organ of Corti requires diverse genetic pathways and interac-

tions (Kelley, 2007; Doetzlhofer et al., 2009).

Bone morphogenetic proteins (*Bmps*) are diffusible molecules, which play important roles in the formation of multiple organs during embryogenesis such as limb, heart, kidney, brain, and eye (Dudley et al., 1995; Furuta et al., 1997; Delot et al., 2003; Bandyopadhyay et al., 2006). *Bmps* mediate cellular changes by binding to specific *Bmp* receptors, which lead to phosphorylation of cytosolic Smad proteins. Phosphorylated Smads, in turn, form specific complexes and translocate to the nu-

cleus to activate or repress transcription of *Bmp* target genes (von Bubnoff and Cho, 2001). Extracellularly, *Bmp* signaling pathways can be negatively regulated by antagonists such as *Noggin* and *Chordin* (Balemans and Van Hul, 2002). Both *Noggin* and *Chordin* are thought to interfere with *Bmp* signaling by directly binding *Bmps* and preventing their interactions with *Bmp* receptors (Holley et al., 1996; Piccolo et al., 1996). In fact, *Noggin* knockout (*Nog<sup>-/-</sup>*) mouse embryos are often used as an animal model for studying the role of *Bmp* functions in a gain-of-function context (Brunet

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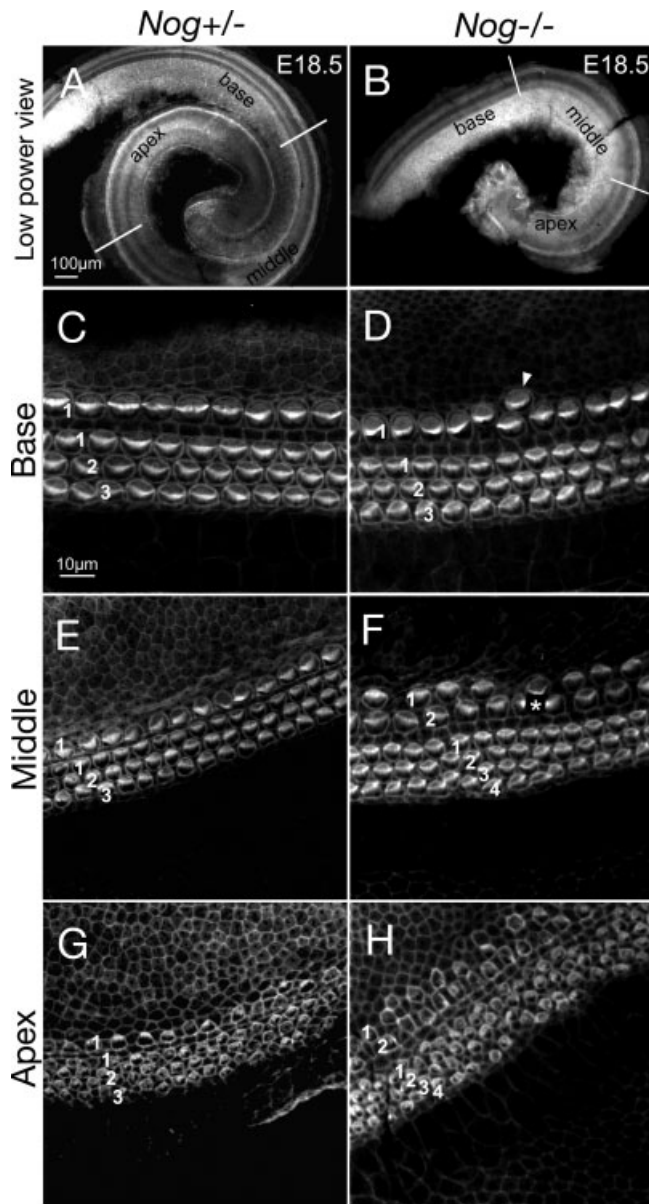
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**Fig. 1.** Increased inner and outer hair cells in *Nog*<sup>-/-</sup> cochlea. A–H: Phalloidin-labeled *Nog*<sup>+/-</sup> (A,C,E,G) and *Nog*<sup>-/-</sup> (B,D,F,H) cochleae at embryonic day (E) 18.5. **A,B:** *Nog*<sup>+/-</sup> cochlea shows one and three-fourth turns (A), whereas *Nog*<sup>-/-</sup> cochlea shows only one turn (B). Representative images from the base, middle, and apical regions are shown for *Nog*<sup>+/-</sup> (C,E,G) and *Nog*<sup>-/-</sup> cochleae (D,F,H). **C,E,G:** A *Nog*<sup>+/-</sup> cochlea showing the typical one row of inner hair cells (IHC; 1) and three rows of outer hair cells (OHC; 1,2,3), which are less organized at the apex. **D,F,H:** *Nog*<sup>-/-</sup> cochlea showing occasional extra IHC at the base (D) and two rows of IHC (1,2) and four rows of OHC (1,2,3,4) in the middle and apical turns (F,H). Asterisk in F indicates damaged hair cells. Scale bar in A applies to B. Scale bar in C applies to D–H.

et al., 1998; Choi et al., 2007; Davis and Camper, 2007).

Using *Nog*<sup>-/-</sup> mouse embryos, we showed that proper levels of Bmps are required for the formation of the stapes in the middle ear and cochlear duct in the inner ear (Bok et al., 2007; Hwang and Wu, 2008). *Nog* expression is not detected in the otic cup and otocyst stages, and the early defects of the

mis-shapen cochlear duct in *Nog*<sup>-/-</sup> embryos is attributed to a misalignment of the hindbrain and inner ear primordia (Bok et al., 2007). Nevertheless, *Nog* expression is detected later in the developing cochlear duct starting at mid-gestation and its expression domain is associated with other Bmps (Bok et al., 2007). Here, we show that

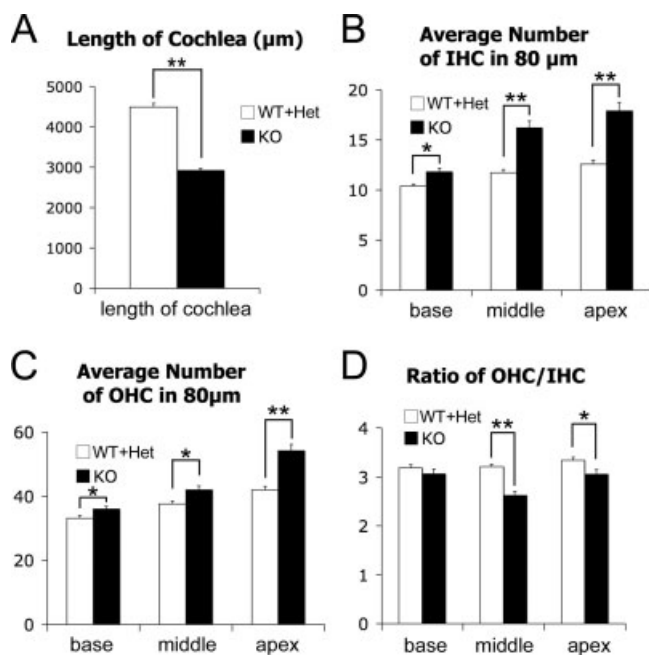
the *Nog*<sup>-/-</sup> cochlear duct is disrupted with an increase in the rows of IHC and OHC. While this hair cell phenotype in *Nog*<sup>-/-</sup> cochleae could be due to mis-regulation of multiple Bmps, we focused our attention on *Bmp2*, which we show to be transiently expressed in nascent cochlear hair cells. To address whether *Bmp2* could be regulating hair cell numbers in the organ of Corti, we conditionally deleted *Bmp2* expression in the mouse cochlear duct using *Foxg1*<sup>cre/+</sup> and *Growth Factor Independent-1-cre* (*Gfi1*<sup>cre/+</sup>) strains. Our results indicate that *Bmp2* is not required for proper hair cell formation and is unlikely to affect normal hearing in the mouse inner ear.

## RESULTS

### Increased Hair Cells in the Cochlea of *Nog*<sup>-/-</sup> Embryos

At E18.5, the length of *Nog*<sup>-/-</sup> cochleae was significantly shorter than cochleae from heterozygotes (Fig. 1A,B). While *Nog*<sup>+/-</sup> cochleae showed an organized single row of IHCs and three rows of OHCs along the entire duct (Fig. 1C,E,G), *Nog*<sup>-/-</sup> cochleae showed occasional extra IHCs at the base (Fig. 1D, arrowhead), and two rows of IHCs and four rows of OHCs in the middle and apical turns (Fig. 1F,H). There was no clear difference in hair bundle orientation between mutants and heterozygotes.

Cochlear lengths and hair cell numbers in each turn were quantified and compared (see the Experimental Procedures section). No obvious difference between wild-type and heterozygotes (Supp. Fig. S1, which is available online) was observed and they were grouped together as controls for quantification. *Nog*<sup>-/-</sup> cochleae showed a 35% reduction in length, compared with controls (Fig. 2A; knockout mice [KO], 2932.8 ± 41.6 μm, n = 10; wild-type and heterozygotes [WT+Het], 4493.9 ± 94.7 μm, n = 7; *P* < 0.001). In the mutant cochlea, there was a significant increase in inner hair cell numbers in a given region, which was more pronounced in middle and apical turns (Fig. 2B, single asterisk; *P* < 0.05, double asterisks; *P* < 0.001). The OHC numbers were found to increase significantly throughout each turn of the cochlea as well (Fig. 2C). Despite the



**Fig. 2.** Quantification of the length and hair cell number in *Nog*<sup>-/-</sup> cochleae. **A:** Reduced cochlear lengths in *Nog*<sup>-/-</sup> cochleae (WT+Het, n = 7; KO, n = 10). **B:** Comparisons of the average IHC numbers from the base, middle, and apex of the cochlea between controls and mutants. The average IHC numbers are significantly increased throughout the *Nog*<sup>-/-</sup> cochlea (WT+Het, n = 6; KO, n = 12). **C:** Comparison of the average OHC numbers in each turn. Statistically significant increase of the average OHC number is also observed throughout the *Nog*<sup>-/-</sup> cochlea (WT+Het, n = 6; KO, n = 12). **D:** Comparison of the ratio of the average outer hair cell versus the average inner hair cell number. The ratio is significantly decreased in middle and apical turns, suggesting that IHCs are preferentially affected in middle and apical turns of *Nog*<sup>-/-</sup> cochlea (WT+Het n = 6, KO n = 12). Each error bar represents standard error of the mean. \**P* < 0.05, \*\**P* < 0.001. OHC, outer hair cell; IHC, inner hair cell; KO, knockout.

increase in hair cell counts in individual turns of the *Nog*<sup>-/-</sup> organ of Corti, the total estimated hair cell number based on the entire length of cochlear duct compared with wild-type and heterozygotes showed a 11% and 24% decrease in IHC and OHC, respectively (KO, IHC  $572.7 \pm 16.8$ ; OHC  $1587.2 \pm 52.0$ ; n = 7; WT+Het, IHC  $642.6 \pm 16.8$ , OHC  $2085.7 \pm 47.3$ ; n = 6). Although these results indicate that the length of the mutant cochlear duct affected the total hair cell number, a shortened cochlear duct could not easily account for the lower ratios of outer versus inner hair cell number observed in the middle and apical turns (Fig. 2D). Therefore, it is likely that both longitudinal and radial patterning of the organ of Corti in *Nog*<sup>-/-</sup> cochleae are affected.

### Normal Expression Patterns of Bmps in the Developing Cochlea

In other tissues where Noggin has been shown to modulate the levels of

Bmps, its expression domain is intimately associated with those of Bmps (Brunet et al., 1998; Warren et al., 2003; Zhou et al., 2006). Similar relationships were observed in the developing cochlear duct (Morsli et al., 1998; Bok et al., 2007). At E16.5, *Bmp4* expression is juxtaposed to the abneural side of the organ of Corti, a region presumed to give rise to Hensen's and Claudius' cells (Fig. 3A; Morsli et al., 1998; Bok et al., 2007). The expression of *Nog* overlaps partially with the *Bmp4* expression domain (Fig. 3G; Bok et al., 2007). *Bmp7* expression in the cochlear duct appears broad, but its strongest expression domain also colocalized in the *Bmp4*-positive region (Fig. 3C). In contrast, *Bmp2* expression does not overlap with that of *Bmp4* but coincides within the sensory region of the cochlea (Figs. 3E, 4A'; bracket), which is positive for *Lunatic fringe* (*Lfng*; Fig. 4A). Comparing adjacent sections probed for *Myosin XVa* (*Myo15a*) transcripts, which are local-

ized in sensory hair cells, *Bmp2* transcripts seem to be detected only in OHC and not IHC (Fig. 4A', A''). *Bmp2* expression in the hair cells appears transient, starting at E15.5 and is only detectable in the apical cochlear region by postnatal day 1 (data not shown). Nevertheless, the timing of this expression is at a critical developmental period for organ of Corti patterning (Shim et al., 2005; Puligilla et al., 2007).

### Targeted Deletion of *Bmp2* Using *Foxg1*<sup>cre/+</sup> Mice

Noggin could be functioning to regulate the levels of any one of the aforementioned Bmps in the cochlea, even though there was no obvious change in the expression patterns of *Bmp2*, *Bmp4*, and *Bmp7* in *Nog*<sup>-/-</sup> cochleae (Fig. 3B,D,F). Because Bmps are specifically expressed in cochlear and not vestibular hair cells in both chicken and mouse (Oh et al., 1996; Morsli et al., 1998), we focused our attention on the role of *Bmp2* in cochlear hair cell formation. We generated *Bmp2* conditional knockout embryos using *Foxg1*<sup>cre/+</sup> mice (Hebert and McConnell, 2000). In these mutant cochleae (*Foxg1*<sup>cre/+</sup>*Bmp2*<sup>lox/-</sup>), *Bmp2* transcripts were not detectable, indicating that *Bmp2* was successfully deleted (Fig. 4B'). Yet, the organ of Corti as well as the sensory hair cells within appear normal, based on the expression patterns of *Lfng* and *Myo15a*, respectively (Fig. 4B,B'). Phalloidin labeling of the whole-mount cochlea showed no obvious abnormality in *Foxg1*<sup>cre/+</sup>*Bmp2*<sup>lox/-</sup> mutants compared with controls (Fig. 4C,D), indicating that *Bmp2* is not required for hair cell formation. Because *Foxg1*<sup>cre/+</sup>*Bmp2*<sup>lox/-</sup> mice die at birth, we used a different *Bmp2* conditional mutant to evaluate the hair cell function in the absence of *Bmp2*.

### Targeted Deletion of *Bmp2* Expressed in Outer Hair Cells Using *Gfi1*<sup>cre/+</sup> Mice

To evaluate the role of *Bmp2* in hair cell function, we generated a more specific targeted deletion of *Bmp2* using *Gfi1*<sup>cre/+</sup> mice, in which *cre* is driven by the promoter of *Gfi1*. *Gfi1* is expressed in hair cells in developing



mouse inner ear starting from E14.5 (Wallis et al., 2003). *Gfi1* is thought to function downstream of *Math1* in sensory hair cells and contribute to the maintenance of *Math1* function (Wallis et al., 2003; Hertzano et al., 2004). Our gene expression study showed that *Gfi1* is strongly expressed in cochlear hair cells at E15.5 (data not shown). In *Gfi1<sup>cre/+</sup>;Bmp2<sup>lox/-</sup>* cochleae, *Bmp2* expression was not detectable in sensory hair cells (Fig. 5B; bracket), which were *Myo15a*-positive (Fig. 5B'; bracket). Whole-mount cochlear preparations also showed no change in hair cell formation and patterning, confirming the results of *Foxg1<sup>cre/+</sup>;Bmp2<sup>lox/-</sup>* embryos that *Bmp2* is not required for hair cell formation (Fig. 5D).

### Hearing and Presbycusis of *Gfi1<sup>cre/+</sup>;Bmp2<sup>lox/-</sup>* Mice

Auditory brainstem response (ABR) measurements at click, 8 kHz, and 16 kHz were conducted on heterozygous and *Gfi1<sup>cre/+</sup>;Bmp2<sup>lox/-</sup>* mice at 3–6 months of age and they showed comparable hearing thresholds (Fig. 6A). Although the observed 6.5 dB threshold shift at 16 kHz is statistically significant, it is not clear if this minimal threshold shift noticeably compromises hearing because a 20-dB threshold shift is considered only a mild hearing loss (Zheng et al., 1999).

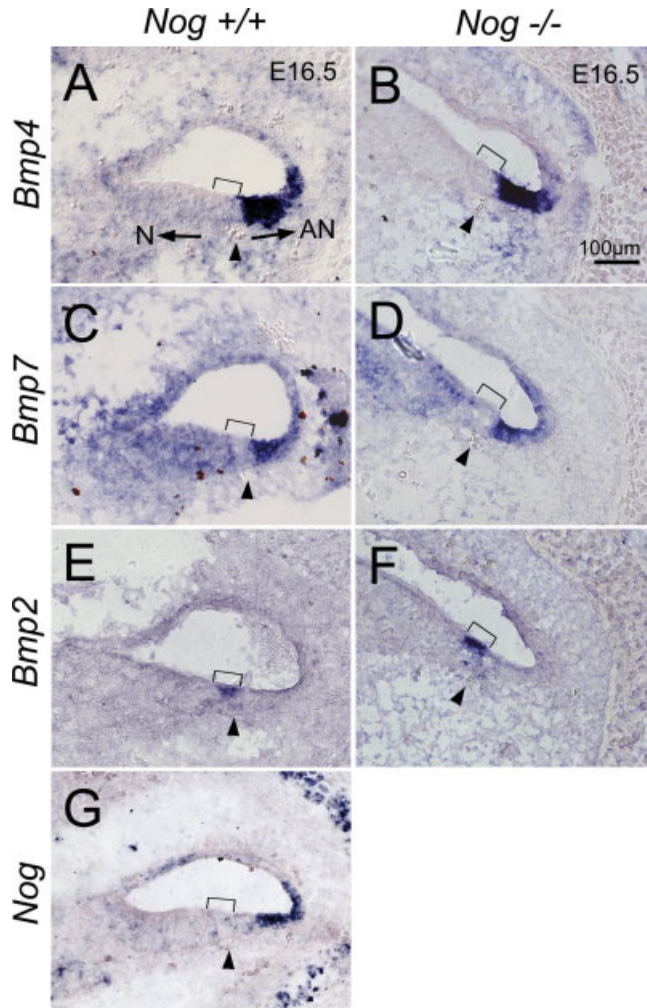


Fig. 3.

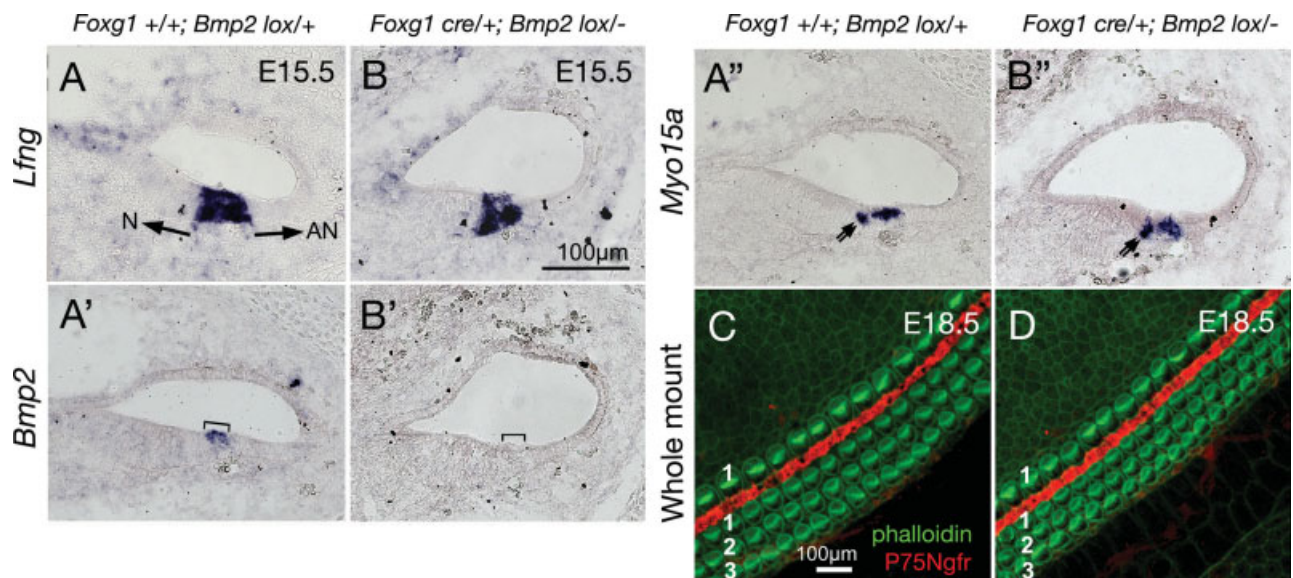


Fig. 4.

To investigate the effect of lack of *Bmp2* on hearing loss due to aging (presbycusis), ABR was measured at 9–10 months of age. At this age, heterozygous littermates started to show significant age-related hearing loss (Fig. 6B;  $P < 0.05$ ), yet the hearing thresholds of *Gfi1<sup>cre/+</sup>;Bmp2<sup>lox/-</sup>* mice remained comparable to heterozygous littermates (Fig. 6B). These results suggest that *Bmp2* is not required for proper hair cell function.

## DISCUSSION

### Role of *Noggin* in the Growth of the Cochlear Duct and Hair Bundle Orientation

The anlagen of the cochlear duct in mice emerges from the ventral otocyst around E10.75 (Morsli et al., 1998). As the duct elongates, it starts to coil forming half a turn by E13.5 and reaches its final shape of one and three-quarter turns by E16.5 (Morsli

et al., 1998). The elongation of the cochlear duct and patterning of the organ of Corti are thought to be mediated by the process of convergent extension whereby the cochlear duct lengthens along its longitudinal axis and at the same time narrows along the radial axis (Keller, 2002; McKenzie et al., 2004; Wang et al., 2005; Yamamoto et al., 2009). Thus, the prosensory domain begins as an epithelium of several cells thick, which thins to a layer of two cells (Wang et al., 2005). Genes involved in establishing planar cell polarity such as *Vangl2*, *Scrb1*, *Dvl*, and *Wnt5* are important in dictating cochlear duct elongation and hair cell alignment (Montcouquiol and Kelley, 2003; Wang et al., 2005; Qian et al., 2007; Simons and Mlodzik, 2008). Lack of these gene products results in a similar phenotype of shortened cochlear duct with increased hair cell numbers in the apical region, as well as disrupted hair cell arrangement along the entire duct (Montcouquiol et al., 2003).

Cochlear ducts from *Nog<sup>-/-</sup>* mutants are shortened and misshapen. However, the planar cell polarity pathway does not appear to be affected in these mutants based on the relatively normal hair cell alignment in the organ of Corti, compared with *Nog<sup>+/-</sup>* cochlea (Fig. 1). In fact, the shortened cochlear duct in these mutants is thought to be indirect, due to misalignment of the rhombomeres and otocysts at an earlier stage of development (Bok et al., 2007). This misalignment presumably affects signaling molecules from the hindbrain, which confer the body plan to the inner ear and guide the proper outgrowth and patterning of the cochlear duct. Examples of other genes that are required for proper cochlear duct outgrowth but appear to be independent of the planar cell polarity pathway include *Gli3* and *Lmx1a* (Driver et al., 2008; Koo et al., 2009).

### *Noggin* Regulates Hair Cell Numbers in the Organ of Corti

The observed increase in the rows of sensory hair cells in *Nog<sup>-/-</sup>* cochleae could be due to defects in the longitu-

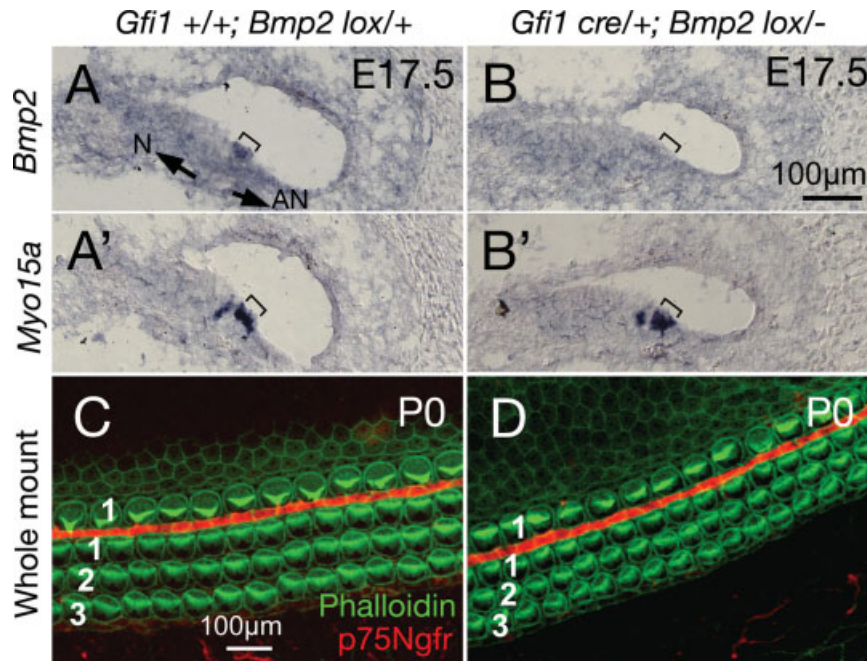
dinal growth of the cochlear duct. Nevertheless, the preferential increase in IHC number in the middle and apical turns of the cochlea cannot be explained easily by the lengthening defects of the cochlear duct. It is likely that *Noggin* is involved in the radial patterning of the organ of Corti as well (Fig. 7). Targeted deletion of *Noggin* in the cochlear duct will address this issue. Those experiments are currently under way. The notion of *Noggin* being required for radial patterning of the organ of Corti is partially based on the expression patterns of *Nog* and *Bmps* in the developing cochlea. The relationships of these genes' expression domains are conserved between chicken and mouse (Chang et al., 1999; Bok et al., 2007). *Bmp4*, *Bmp7*, and *Nog* are expressed in overlapping domains in the abneural side of the developing sensory region in both species (Oh et al., 1996; Chang et al., 1999; Bok et al., 2007). The intimate relationships among *Bmps* and *Nog* expression domains in the cochlea are similar to reports in other tissues where *Noggin* functions to regulate the levels of *Bmps* (Brunet et al., 1998; Warren et al., 2003; Zhou et al., 2006).

Furthermore, several in vitro studies have implicated a role of *Bmps* in hair cell formation, which could be extrapolated to a role in vivo. For example, *Bmp4* has been shown to induce hair cell formation in mouse cochlear explants as well as cultures of dissociated otic epithelial cells from chicken (Li et al., 2005; Puligilla et al., 2007). However, other studies using explants of chicken otocysts appear to arrive at an opposite conclusion showing that hair cell formation is induced by *Noggin* but inhibited by *Bmp4* (Pujades et al., 2006). These various in vitro results are not necessarily in conflict with each other. In the chicken otocyst explants studies, the anterior and posterior cristae were the main sensory organs being analyzed, rather than an auditory sensory organ. *Bmp4* has multiple functions in the sensory cristae promoting both sensory and nonsensory fates (Chang et al., 2008). It is possible that, at the developmental period of the otocyst cultures, *Bmp4* is required to co-ordinate the nonsensory fate, which could account for the observed loss of

**Fig. 3.** Expression patterns of *Bmps* and *Nog* in wild-type and *Nog<sup>-/-</sup>* cochleae at embryonic day (E) 16.5. **A:** *Bmp4* is strongly expressed in the abneural region of wild-type cochlea. **C:** *Bmp7* expression is broad but its strongest expression domain overlaps with the *Bmp4*-positive region. **E:** *Bmp2* is expressed in the sensory region of the cochlea, not overlapping with the *Bmp4*-positive region. **G:** The expression domain of *Nog* partially overlaps with *Bmp4* expression. **B,D,F:** There is no noticeable change in expression patterns of *Bmp4*, *Bmp7*, and *Bmp2* in *Nog<sup>-/-</sup>* cochleae, respectively. Arrowhead indicates the spiral blood vessel beneath the prosensory domain (bracket). Arrows in A point toward neural (N) and abneural (AN) sides of the organ of Corti. Scale bar in B applies to A–G.

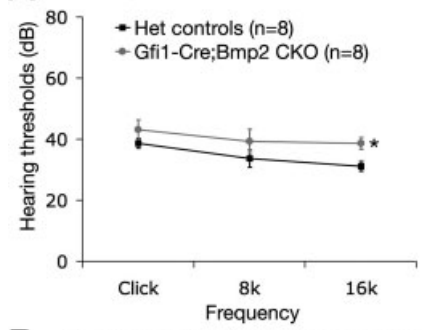
**Fig. 4.** Targeted deletion of *Bmp2* using *Foxg1<sup>cre/+</sup>* mice. **A–A'':** Serial sections of a *Foxg1<sup>+/+</sup>;Bmp2<sup>lox/+</sup>* cochlea at embryonic day (E) 15.5 showing *Lfn3* expression in the organ of Corti (A), within which *Bmp2* transcripts are detected in outer hair cell (OHC; A', bracket), and *Myo15a* transcripts are detected in both OHC and inner hair cell (IHC; A''). **B–B'':** Serial sections of a *Foxg1<sup>cre/+</sup>;Bmp2<sup>lox/-</sup>* cochlea at E15.5. *Lfn3* and *Myo15a* expression patterns in the organ of Corti are similar to *Foxg1<sup>+/+</sup>;Bmp2<sup>lox/+</sup>* cochlea, whereas *Bmp2* expression is not detected (B', bracket). Double arrows points to *Myo15a*-positive IHC in (A'') and (B''). Arrows in (A) point toward neural (N) and abneural (AN) sides of the organ of Corti. **C,D:** Phalloidin-labeled *Foxg1<sup>+/+</sup>;Bmp2<sup>lox/+</sup>* (C) and *Foxg1<sup>cre/+</sup>;Bmp2<sup>lox/-</sup>* (D) cochleae at E18.5. Scale bar in B applies to A–A'', B'–B'', and C applies to D.



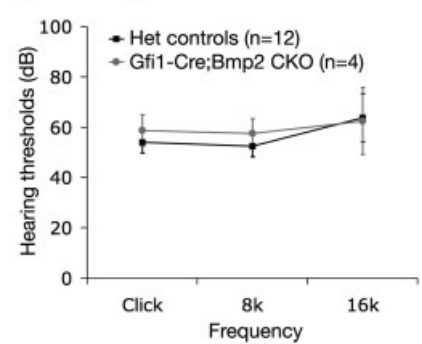


**Fig. 5.** Targeted deletion of *Bmp2* using *Gfi1<sup>cre/+</sup>* mice. **A–A'**: Serial sections of a *Gfi1<sup>+/+</sup>;Bmp2<sup>lox/+</sup>* cochlea at embryonic day (E) 17.5, showing *Bmp2* expression overlapping with *Myo15a* in outer hair cell (OHC; bracket). **B–B'**: Serial sections of a *Gfi1<sup>cre/+</sup>;Bmp2<sup>lox/-</sup>* cochlea at E17.5 showing the absence of *Bmp2* expression in OHC (compare bracketed regions). **C,D**: Whole-mounts of *Gfi1<sup>+/+</sup>;Bmp2<sup>lox/+</sup>* (C) and *Gfi1<sup>cre/+</sup>;Bmp2<sup>lox/-</sup>* (D) cochleae at postnatal day (P) 0, showing the three rows of OHC and one row of inner hair cell (IHC; phalloidin labeling in green), separated by pillar cells (p75Ngfr staining in red). No difference in hair cell formation and organization are detected in *Gfi1<sup>cre/+</sup>;Bmp2<sup>lox/-</sup>* cochlea. Scale bar in B applies to A–A', B', and C applies to D. Arrows in A point toward neural (N) and abneural (AN) sides of the organ of Corti.

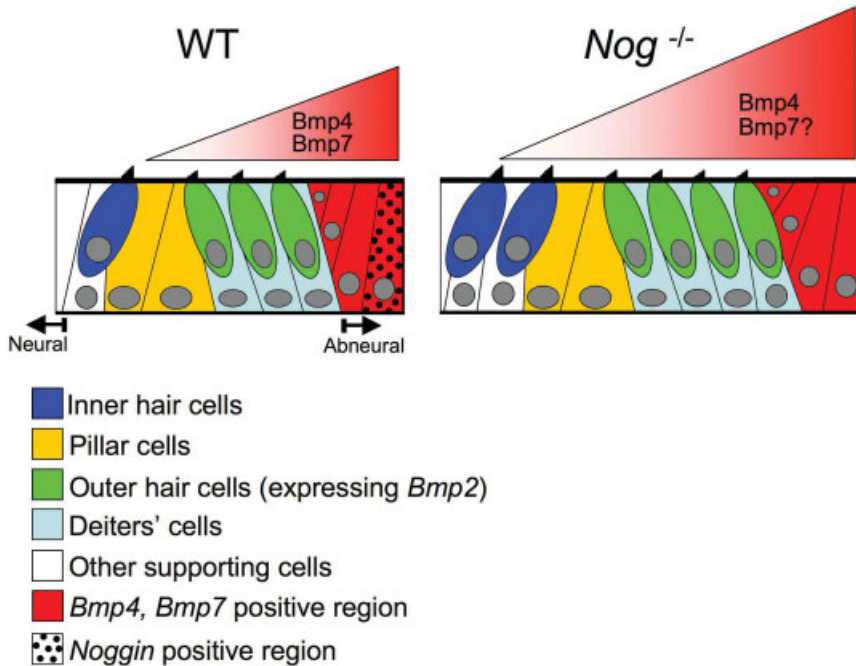
### A Hearing Thresholds at 3–6 months



### B Hearing Thresholds at 9–10 months



**Fig. 6.** Hearing and presbycusis of *Gfi1<sup>cre/+</sup>;Bmp2<sup>lox/-</sup>* mice. The hearing thresholds at click, 8 kHz, 16 kHz are compared between heterozygotes (*Gfi1<sup>cre/+</sup>;Bmp2<sup>lox/+</sup>* or *Gfi1<sup>+/+</sup>;Bmp2<sup>lox/-</sup>*) and *Gfi1<sup>cre/+</sup>;Bmp2<sup>lox/-</sup>* mice at 3–6 months and 9–10 months of age. Asterisk indicates a 6.5 dB threshold shift in *Gfi1<sup>cre/+</sup>;Bmp2<sup>lox/-</sup>* mice compared with heterozygotes at 16 kHz ( $P < 0.05$ ). Both heterozygotes and conditional mutant mice start to show age-related hearing loss at 9–10 month ( $P < 0.05$ ) but there is no significant difference in threshold shift between heterozygotes and *Gfi1<sup>cre/+</sup>;Bmp2<sup>lox/-</sup>* mice at the ages measured ( $P > 0.05$ ).



**Fig. 7.** Model of the role of bone morphogenetic proteins (Bmps) in organ of Corti formation. In wild-type cochlea, *Bmp4* and *Bmp7* secreted from the abneural region regulate hair cell formation. The levels of Bmps are modulated by the antagonist, *Noggin*. In the absence of *Noggin*, the net level of *Bmp4* and may be the level of *Bmp7* as well across the prosensory domain are increased leading to extra hair cell formation in the organ of Corti.

sensory hair cells with exogenous *Bmp4*. While the nature of the increased hair cells in the dissociated cell culture studies is not clear (Li et al., 2005), the cochlear explant results are direct indications that Bmps promote cochlear hair cell formation (Puligilla et al., 2007). The expression patterns of Bmps and *Noggin* in the cochlea as well as the increased rows of auditory hair cells of *Nog<sup>-/-</sup>* mutants is consistent with the hypothesis that Bmps and *Noggin* are involved in radial patterning of the organ of Corti. The difference between the two gain-of-Bmp function models, cochlear explants, and *Nog<sup>-/-</sup>* mutants, is that increased IHC and OHC rows are observed in the *Nog<sup>-/-</sup>* cochlea, whereas only OHC number is observed to increase in cochlear

explants treated with human *Bmp4* (Puligilla et al., 2007). The lack of changes in IHC number in the explants could be related to the late timing of *Bmp4* treatments.

A recent preliminary report showed that the lack of *Bmp receptor Ia* in the mouse cochlea caused a duplicated organ of Corti (T. Ohyama and A.K. Groves; personal communication). Further reduction of Bmp signaling in compound *Bmp receptor Ia* and *Ib* mutants resulted in loss of the prosensory domain. Thus, these authors suggested that organ of Corti development is sensitive to a gradient of Bmp levels across the prosensory region. Such levels of Bmps could be regulated by *Noggin*. Nevertheless, the various resulting phenotypes from the Bmp receptor knockouts or *Nog* mutants cannot be easily reconciled. It is possible that Bmps have multiple roles during the course of the organ of Corti formation. The profound sensory phenotypes obtained from knocking out the two Bmp receptors suggest an early role of Bmps in sensory organ specification. In contrast, the increase in rows of hair cells in *Nog*<sup>-/-</sup> cochlea or *Bmp4* treated cochlear explants could be the consequence of a later, modulatory role of Bmps in promoting hair cell formation. This idea is consistent with the fact that *Nog* expression is not detected in the cochlea until mid-gestation (Bok et al., 2007).

### Role of *Bmp2* in Hair Cell Formation

It is not clear though which Bmps are being regulated by *Noggin* within the cochlear duct. The published data thus far do not exclude the possibility of *Bmp2* playing a role in regulating hair cell formation in the cochlea. *Bmp2* and *Bmp4* are thought to have similar affinities in binding to *Noggin* and Bmp receptors (Rosen et al., 1996; Zimmerman et al., 1996). Thus, the increased OHC number in explant cultures using *Bmp4* could be mimicking either one or both sources of Bmps in the cochlea. Even though expression of *Bmp2* in the sensory hair cells is transient, it is at a developmental period when hair cell fates are amenable to change (Shim et al., 2005; Puligilla et al., 2007). Further-

more, expression of *Bmps* in auditory hair cells is conserved between chicken and mice, even though it is *Bmp4* rather than *Bmp2* that is expressed in auditory hair cells of chicken (Oh et al., 1996). Here, using two different *cre* expressing mouse strains to delete *Bmp2* expression in auditory hair cells, we concluded that *Bmp2* is not required for proper hair cell formation and is likely not needed for normal hearing in mice. Based on these results, *Bmp4* expression in the abneural region seems to be the most likely candidate to be regulated by *Noggin*. Moreover, while there is no report of inner ear defects in *Bmp7*<sup>-/-</sup> embryos (Karsenty et al., 1996) and *Noggin* does not bind *Bmp7* as effectively as *Bmp2* and *Bmp4* (Zimmerman et al., 1996), a role of *Bmp7* in cochlear hair cell formation cannot be ruled out.

### Interactions Between Fgfs and Bmps in the Organ of Corti

Multiple fibroblast growth factors are expressed in the developing organ of Corti such as *Fgf8*, *Fgf10* and *Fgf20* (Jacques et al., 2007; Pauley et al., 2003; Hayashi et al., 2008). Conditional deletion of *Fgfr1* in the inner ear resulted in patches of sensory islands along the cochlear duct (Pirvola et al., 2002). *Fgf20* expressed in the prosensory domain was proposed to be the main ligand activating *Fgfr1* in the cochlea (Hayashi et al., 2008). In addition, loss of *Fgfr3*, which is also expressed in the prosensory domain, results in the absence of pillar cells and increased OHC (Puligilla et al., 2007; Hayashi et al., 2007). *Fgf8* emanating from the IHC is thought to be the main ligand activating *Fgfr3* (Jacques et al., 2007).

The interactions between Fgfs and Bmps in organ of Corti formation was implicated when *Noggin* treatment rescued the increase in OHC number in *Fgfr3*<sup>-/-</sup> cochlear explants (Puligilla et al., 2007). These results and the demonstration that *Bmp4* induces OHC formation in cochlear explants led to the hypothesis that Fgfs interact with *Bmp4* in the abneural region of the cochlear duct in specifying the number of OHC (Puligilla et al.,

2007). While these results do not preclude the possibility that *Bmp2* in the sensory hair cells could be interacting with Fgf signaling, our results here indicate that *Bmp2*'s function in the organ of Corti is dispensable and further implicate the importance of other Bmps expressed in the abneural region.

Given the antagonistic relationships between Fgf and Bmp signaling in other tissues (Pera et al., 2003; Cushing et al., 2008; Huang et al., 2009), the effects of a *Bmp4* gradient, if it exists, could be further fine-tuned by Fgfs expressed in the prosensory region such as *Fgf10* and *Fgf20* or *Fgf8* in the IHC (Jacques et al., 2007; Pauley et al., 2003; Hayashi et al., 2008). If this hypothesis is correct, one would predict a change in hair cell fate or number when either *Bmp4* or Fgfs signaling is affected. On the contrary, only OHC and not IHC appear to be affected when Fgf signaling is perturbed in both *Fgfr3*<sup>-/-</sup> and *Sprouty2*<sup>-/-</sup> mutants (Shim et al., 2005; Hayashi et al., 2007; Puligilla et al., 2007). Functional redundancy among the Fgfs and their receptors could account for the lack of a more profound IHC phenotype in some of these mutants (Hayashi et al., 2007; Puligilla et al., 2007). Future studies will focus on the requirements of Fgfs and Bmps in establishing inner and outer hair cell fates.

## EXPERIMENTAL PROCEDURES

### Mice and Genotyping

The *Nog*<sup>+/*tm1Amc*</sup> (*Nog*<sup>+/-</sup>) mice (McMahon et al., 1998) on a congenic C57BL/6J background were obtained from Dr. Richard Harland (Univ. of California at Berkeley) and maintained on a C57BL/6J;FVB hybrid background. Genotyping for *Nog*<sup>-/-</sup> embryos was determined by PCR as previously described (McMahon et al., 1998). The *Bmp2*<sup>*tm1.1Mis/tm1.1Mis*</sup> (*Bmp2*<sup>*lox/lox*</sup>) and *Bmp2*<sup>+/*tm1Brd*</sup> (*Bmp2*<sup>+/-</sup>) mice, generated as previously described (Zhang and Bradley, 1996; Singh et al., 2008) were maintained on a 129SvEv; C57BL6/J mixed background and genotyped according to previous description (Singh et al., 2008). The generation of



*Gfi1*<sup>cre/+</sup> knock-in mouse, in which *cre* is driven by the promoter of *Gfi1*, will be described elsewhere (Lin Gan, Univ. of Rochester). Mice with conditional deletion of *Bmp2* were generated by mating *Foxg1*<sup>cre/+</sup>;*Bmp2*<sup>+/-</sup> or *Gfi1*<sup>cre/+</sup>;*Bmp2*<sup>+/-</sup> with *Bmp2*<sup>lox/lox</sup> mice.

### In Situ Hybridization, Immunohistochemistry, and Hair Cell Number Quantification

For in situ hybridization, RNA probes for *Bmp4*, *Lfn3* (Morsli et al., 1998), *Bmp2* (McMahon et al., 1998), *Bmp7* (Bellusci et al., 1996), *Nog* (Brunet et al., 1998), *Gfi1* (Yang et al., 2003), and *Myo15a* (Anderson et al., 2000) were prepared as previously described. Immunohistochemistry for whole-mount cochlear duct was done as previously described (Koo et al., 2009). To quantify the hair cell number, each cochlea was divided into the base, middle and apical turns of equal length demarcated in Figure 1A,B. In each turn, hair cell numbers were counted in three evenly spaced 80- $\mu$ m segments when available, and the mean hair cell number was calculated and used to compute the average hair cell numbers among controls and mutants. Statistical analyses on hair cell numbers and hearing thresholds between mutants and controls were conducted using Student's *t*-test.

### Hearing Measurement

Auditory brainstem response (ABR) was recorded using Smart EP (Intelligent Hearing System, FL) as previously described (Hwang and Wu, 2008). Briefly, mice were anesthetized with an intra-peritoneal injection of the mixture of Ketamine (56 mg/kg body weight, Ketalar, Generamedix, USA) and Dexmedetomidine (0.375 mg/kg body weight, Dexdomitor, Pfizer Animal Health, USA) and kept on a heating pad in a soundproof chamber. Auditory stimuli used were click, 8 kHz, and 16 kHz tone pips and hearing of right ears was measured. After the procedure, mice were recovered by intra-peritoneal injection of Atipamezole (5 mg/kg body weight, Antisedan, Pfizer Animal Health, USA).

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