

LMO4 Functions As a Negative Regulator of Sensory Organ Formation in the Mammalian Cochlea

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In mammals, formation of the auditory sensory organ (the organ of Corti) is restricted to a specialized area of the cochlea. However, the molecular mechanisms limiting sensory formation to this discrete region in the ventral cochlear duct are not well understood, nor is it known whether other regions of the cochlea have the competence to form the organ of Corti. Here we identify LMO4, a LIM-domain-only nuclear protein, as a negative regulator of sensory organ formation in the cochlea. Inactivation of *Lmo4* in mice leads to an ectopic organ of Corti (eOC) located in the lateral cochlea. The eOC retains the features of the native organ, including inner and outer hair cells, supporting cells, and other nonsensory specialized cell types. However, the eOC shows an orientation opposite to the native organ, such that the eOC appears as a mirror-image duplication to the native organ of Corti. These data demonstrate a novel sensory competent region in the lateral cochlear duct that is regulated by LMO4 and may be amenable to therapeutic manipulation.

Key words: inner ear development; LIM-domain; LMO4; organ of Corti; cochlea; transcriptional regulator

Introduction

Unlike the sensory regions in fish, amphibians, and birds, loss of mammalian cochlear hair cells and supporting cells causes permanent hearing deficits due to the inability of the sensory cells to spontaneously regenerate. The mammalian organ of Corti (OC) has a unique structure, in which the hair cells and supporting cells are arrayed in a tightly organized pattern composed of rows of hair cells and intervening supporting cells. Although considerable inroads have been made in understanding the molecular mechanisms that induce sensory or hair cell formation in the ear (Birmingham et al., 1999; Zheng and Gao, 2000; Mizutani et al., 2013), the factors that regulate OC formation in the cochlea are largely unknown.

Previously, we reported that the LMO family of nuclear proteins LMO1, LMO3, and LMO4 are expressed in the developing inner ear in temporally and spatially distinct regions (Deng et al.,

2006, 2010). To understand the role of LMO4 in the developing inner ear, we have inactivated it using the lacZ reporter knock-in and conditional knock-out approaches (Deng et al., 2010). Both knock-outs demonstrate a similar phenotype, in which the vestibular region is severely malformed (Deng et al., 2010). The morphogenetic defects are due to the loss of early expression of LMO4 in the lateral portion of the otocyst, leading to early patterning defects (Deng et al., 2010). Nevertheless, cochleae are formed in the *Lmo4*-null mice, although shorter than the wild-type, allowing an examination of the development of the OC in the absence of LMO4 function.

Here we identify a novel sensory competent region in the lateral cochlear duct that is negatively regulated by *Lmo4*. We found that *Lmo4* is continuously expressed in the developing mouse cochlea and inactivation of *Lmo4* in mice leads to the formation of an ectopic organ of Corti (eOC) located in the lateral cochlea. Interestingly, the eOC retains the features of the native OC. Structurally, the eOC shows a high degree of similarity with the native OC. More importantly, the eOC contains all cell types that the native OC has, including inner and outer hair cells, supporting cells, and other nonsensory specialized cell types. Our study presents the first case that an eOC can be regenerated by manipulating the *Lmo4* gene, which may have important implications for the treatment of hearing loss.

Materials and Methods

Mice. *Lmo4*^{lacZ} knock-in and the *Lmo4*^{loxP} conditional knock-out mice were previously reported (Deng et al., 2010). Noon of the day on which a vaginal plug was observed was designated as embryonic day (E)0.5. All animal procedures used in this study were approved by The University

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Committee of Animal Resources at University of Rochester. All of the experiments were performed in mice of either sex ($n \geq 3$ for each genotype).

X-gal staining. To determine the expression pattern of *Lmo4-lacZ* gene, we used X-Gal staining to detect the activity of β -galactosidase (Gan et al., 1999). Briefly, embryos of either sex were fixed in 4% paraformaldehyde (PFA) in PBS (pH 7.2) at 4°C for 1 h. Whole-mount embryos or 20- μ m-thick cryosections were stained overnight at room temperature with 0.1% X-Gal, 5 mM potassium ferricyanide, 5 mM potassium ferrocyanide, 2 mM $MgCl_2$ in PBS.

Immunohistochemistry and scanning electron microscopy. Scanning electron microscopy (SEM) experiments were performed as previously described (Deng et al., 2010; Pan et al., 2013). For immunohistochemistry of cryosections, the heads of the embryos of either sex from different stages were fixed in 4% PFA in PBS for 1 h. After three washes of PBS, the tissues were dehydrated in 20% sucrose in PBS for overnight, embedded in OCT compound (Tissue-Tek), and cryosectioned at a thickness of 14–20 μ m. Immunolabeling of the sections was then conducted as previously described (Deng et al., 2010; Luo et al., 2013). For whole mount immunohistochemistry, the inner ears of P5 pups were isolated in PBS and then treated with collagenase (1 mg/ml; Worthington) and neutral protease (1 mg/ml; Worthington) for 8 min in cold HBSS (Invitrogen). The HBSS solution was replaced by DMEM/F12 (contained 5% FBS) and incubated for 30 min. The cochlea was then dissected under the dissecting microscope, fixed in 4% PFA in PBS for 1 h at 4°C, and then washed with PBS three times. For immunostaining, the cochlea was permeabilized and blocked in PBS plus 0.1% Triton X-100, 0.03% saponin, 10% horse serum. After incubation for 30 min at room temperature, primary antibody (diluted in PBS plus 0.1% Triton X-100, 0.03% saponin, 3% horse serum, and 3% BSA) was added and incubated at 4°C overnight. The following day, the cochlea was washed in PBS for ~8 h (PBS changed every 2 h) and incubated with fluorescently labeled secondary antibody at 4°C overnight.

The primary antibodies and concentrations used in this study were as follows: rabbit anti-MYO6 (Proteus Biosciences; 1:500), goat anti-SOX2 (Santa Cruz Biotechnology; 1:500), rabbit anti-SOX2 (Millipore; 1:500), rabbit anti-S100A1 (Dako; 1:50), mouse anti-p27kip1 (BD Pharmingen; 1:200), goat anti-JAG1 (Santa Cruz Biotechnology; 1:200), rabbit anti-P75 (Millipore; 1:100), rabbit anti-pSMAD1/5/8 (Cell Signaling Technology; 1:75), rabbit anti-CASP3 (R&D Systems; 1:500), and AlexaFluor 488-conjugated phalloidin (Invitrogen; 1:100). AlexaFluor-conjugated secondary antibodies were obtained from Invitrogen and were used at a concentration of 1:1000. Images were captured with a Zeiss 510 META confocal microscope.

In situ hybridization. *In situ* hybridization was performed as previously described (Yang et al., 2003). Briefly, embryos were dissected in cold PBS and fixed in 4% paraformaldehyde in PBS for overnight. After cryopreserved in 20% sucrose for overnight, the embryos were embedded in OCT medium (Tissue-Tek) and frozen with dry ice quickly. Tissues were sectioned at 20 μ m and mRNA expression was detected with digoxigenin-labeled riboprobes specific for *Bmp4* (a gift from Dr James F. Martin, Baylor College of Medicine, Houston, TX), *Fgf8*, *Hes5*, *Jag2* (a gift from Dr Rulang Jiang, Cincinnati Children Hospital, Cincinnati, OH), *Lfng* and *Ped* (a gift from Dr Doris K. Wu, NIH, Bethesda, MD), *Lmo3*, *Lmo4* (a gift from Dr Terence H. Rabbitts, University of Oxford, Oxford, United Kingdom), *Atoh1*, *Ped*, and *Sox2*.

Results

We first investigated the detailed spatiotemporal expression pattern of *Lmo4* in the developing mouse cochlea using a *lacZ* reporter gene knock-in at *Lmo4* locus (*Lmo4-lacZ*) (Deng et al., 2010). Before the sensory formation at E12.5–E13.5, *Lmo4-lacZ* was broadly expressed in the ventral portion of the cochlea (Fig. 1A,B), including the *Sox2*-expressing presumptive prosensory domain (Fig. 1F,G). From E14.5, the expression of *Lmo4* is downregulated in the developing OC (Fig. 1C–E), which is marked by *Atoh1* expression (Fig. 1H–J). *Lmo4-lacZ* expression was confined in two specific regions in the cochlea: the greater

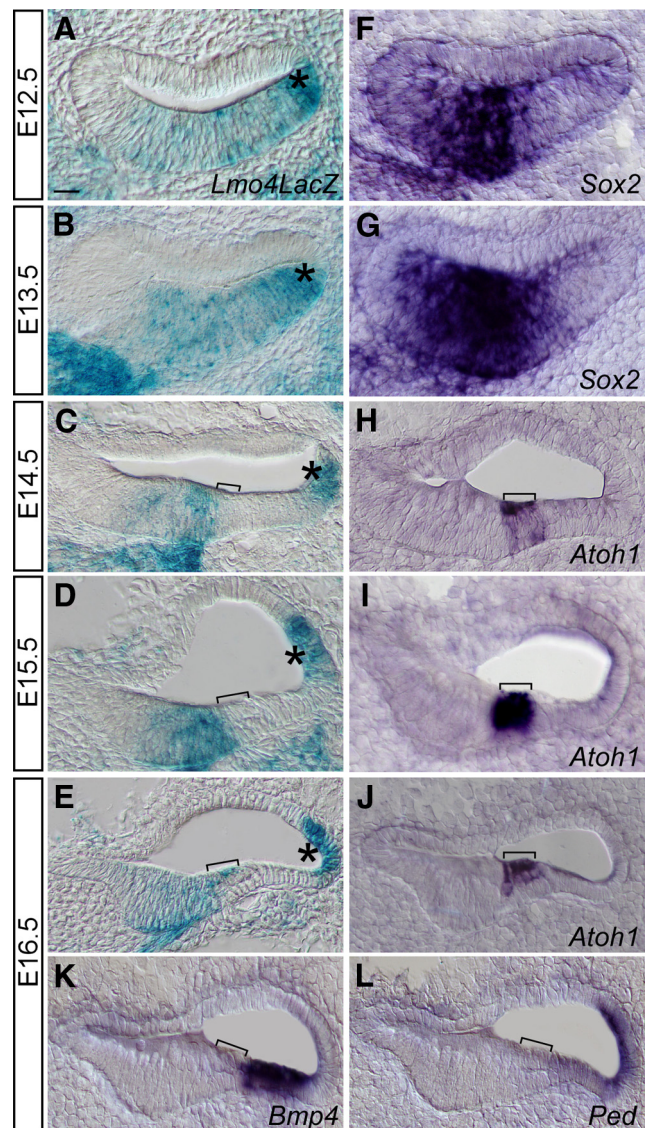


Figure 1. *Lmo4* is continuously expressed in the developing mouse cochlea, with a region-specific expression pattern. **A–E**, X-Gal staining of *Lmo4-lacZ* knock-in reporter activities in *Lmo4^{lacZ}* cochlea sections reveals the spatiotemporal expression pattern of *Lmo4* in the developing mouse cochlea. **F–L**, *In situ* hybridization with region-specific markers confirms the spatiotemporal expression pattern of *Lmo4* in the developing mouse cochlear duct. Brackets indicate the OC. Note *Lmo4* is continuously expressed in the ES at high level (asterisks in **A–E**). Scale bar, 50 μ m.

epithelium ridge (GER) adjacent to the OC and the external sulcus (ES) lateral to the lesser epithelium ridge (LER; Fig. 1C). The specific expression of *Lmo4* in these two regions was maintained after OC formation until at least E16.5 (Fig. 1D,E). The expression of *Lmo4* in the ES region was further confirmed by comparing with the expression of *Bmp4*, a marker for the LER region and *Pendrin* (*Ped*), a marker of the ES region (Yoshino et al., 2004, 2006). *Lmo4* is coexpressed with *Ped* in the ES region but not with *Bmp4* at E16.5 (Fig. 1E,K,L). Later at E16.5, the expression of *Lmo4* is detected in hair cells marked by *Atoh1* expression (Fig. 1E,J). Together, these results indicate that *Lmo4* is continuously expressed in the developing mouse cochlea, with a region-specific expression pattern.

We then characterized the function of LMO4 in the development of mouse cochlea by examining both a traditional knock-out and a conditional knock-out (*Foxg1-Cre; Lmo4^{loxP/loxP}*).

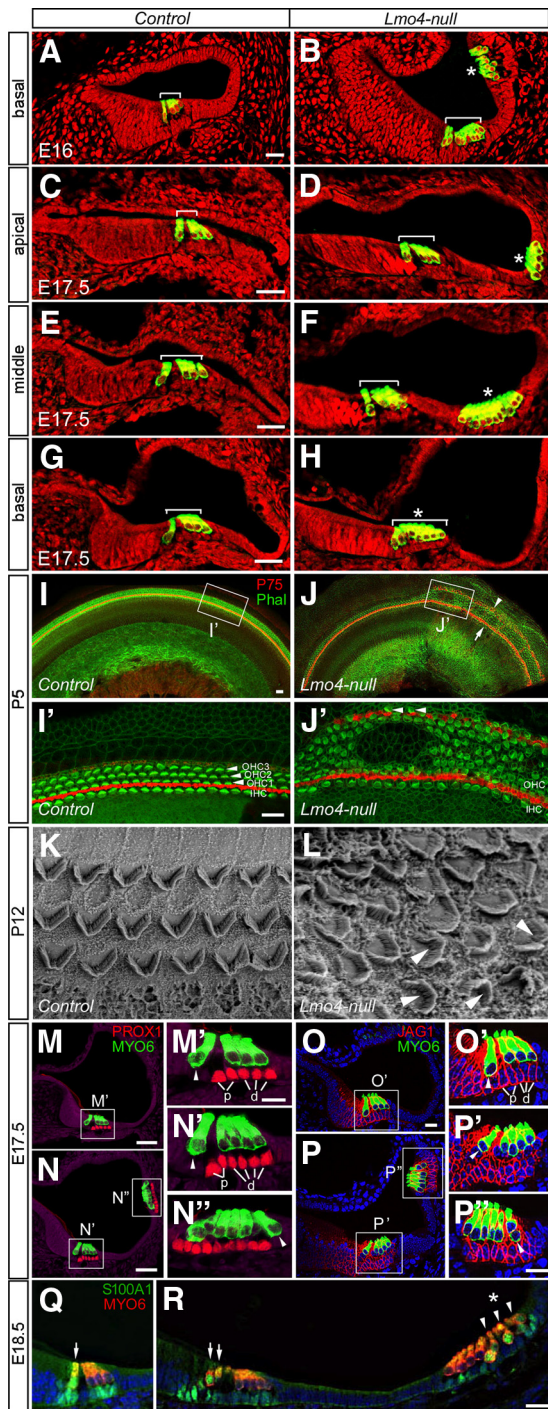


Figure 2. Formation of an ectopic organ of Corti in the *Lmo4*-null cochlea. **A–H**, Ectopic hair cells are observed in *Lmo4*-null cochlea. Cross-sections through the indicated turns of the cochlea duct were immunolabeled with anti-MYO6 (green) and nuclear counterstained with propidium iodide (red). Brackets indicate the native OC, which contains one row of inner hair cells and three rows of outer hair cells. Asterisks indicate the ectopic hair cells. **I–J'**, Ectopic hair cells and pillar cells are formed in the *Lmo4*-null cochlea. Whole-mount immunocytochemistry using anti-P75 (red) and FITC-conjugated phalloidin (green) reveals the formation of eOC in the *Lmo4*-null cochlea. **I'** and **J'** are higher-magnification images of **I** and **J**, respectively. Arrow and arrowhead in **J** indicates pillar cells in the native OC and eOC. Arrowheads in **I'** indicate outer hair cells in control cochlea. Arrowheads in **J'** indicate ectopic inner hair cells in the eOC. **K, L**, SEM image shows that planar cell polarity of hair cells is affected in *Lmo4*-null cochlea. Arrowheads show that the orientation of hair cell stereocilia in the region of ectopic hair cell formation is disrupted in *Lmo4*-null cochlea. **M–P'**, The *Lmo4*-null cochlea contains an ectopic and reverse orientated OC. Compared with the control (**M, O**), immunolabeling of the *Lmo4*-null cochlea (**N, P**) with anti-MYO6 (green) and anti-PROX1 or anti-JAG1 (red) at E17.5 shows the formation of

Compared with the wild-type control (Fig. 2A), ectopic hair cells were detected at the basal turns of the *Lmo4*-null cochlea at E16 (Fig. 2B, asterisk). By E17.5, this region of hair cells appeared in the middle and apical regions of the lateral cochlea (Fig. 2C–H). Subsequently, the two regions of hair cells merge together to form an expanded domain of hair cells (Fig. 2H), perhaps due to apoptosis of intervening cells.

By postnatal day (P)5, expanded regions of hair cells were present in the middle and apical regions of the cochlea, which sometimes include two domains of hair cells, separated by a small island of nonsensory cells (Fig. 2I, J, I', J'). Compared with one row of P75-positive pillar cells seen in the control cochlea, an ectopic row of pillar cells (arrowhead) was observed in the *Lmo4*-null cochlea in addition to the normal row of the pillar cells (Fig. 2J, arrow) seen in the middle turn of the cochlea duct. Interestingly, the ectopic hair cells appeared disorganized within the plane. We therefore asked whether the planar cell polarity (PCP) of hair cells was affected in *Lmo4*-null cochlea using SEM. In wild-type controls, hair cells were orderly arranged and all the stereocilia of the outer hair cells point in the same direction (Fig. 2K). In *Lmo4*-null mice, the orientation of hair cells in the region of ectopic hair cell formation was disrupted (Fig. 2L). Compared with normal PCP observed in the wild-type controls, the PCP of hair cells was disrupted in *Lmo4*-null mice and the stereocilia of outer hair cells pointed randomly and irregularly in the region of ectopic hair cell formation (Fig. 2L), which presumably results from the high hair cell density in the shortened *Lmo4*-null cochlea.

Structurally, the ectopic region of hair cells resembles the native OC (Fig. 2A–H, brackets). We then characterized this ectopic region of hair cells using markers for hair cells, supporting cells, and other cochlear regions. We found the ectopic region expressed well characterized hair cell markers, including MYO6, phalloidin, *Atoh1*, and *Jag2* (Figs. 2B, D, F, H, M–P'; 3A–B'). Analysis of S100A1 expression, which marks inner hair cells as well as supporting cells (Dieter's cells and inner phalangeal cells; Woods et al., 2004; Kiernan et al., 2006), identified inner hair cells on the lateral side of the ectopic region (Fig. 2Q, R). Moreover, *Fgf8* expression, which marks inner hair cells (Pirvola et al., 2002; Shim et al., 2005), revealed the *Fgf8*-expressing inner hair cells in the dorsolateral region of the cochlea, confirming that the ectopic region also contains inner hair cells (Fig. 3C, C'). By comparing the location of *Fgf8* expression with *Jag2* expression, it appeared the ectopic *Fgf8*-expressing inner hair cell was located on the lateral rather than medial side of the ectopic sensory region.

We next sought to determine whether supporting cells were formed in this ectopic region. Using a number of supporting cell markers, we demonstrated that supporting cells are present in the ectopic region and they express the appropriate markers that are normally expressed in the OC region, including P75, PROX1, JAG1, *Hes5*, and *Lfng* (Figs. 2I–J', M–P'; 3D–E'). Unlike JAG1, *Hes5*, and *Lfng*, which are expressed in all OC supporting cells, P75 is uniquely expressed in pillar cells (Mueller et al., 2002; Doetzlhofer et al., 2009), which are specialized supporting cells that separate the inner and outer hair cells. We examined P75

an eOC completed with hair cells and supporting cells. Arrowheads indicate the inner hair cells. **Q, R**, Cross-sections through the control (**Q**) and *Lmo4*-null (**R**) cochlea at E18.5 stained for MYO6 (red), S100A1 (green), and DAPI (blue) confirms that the ectopic region is oriented as a mirror image to the native organ. Arrowheads indicate the inner hair cells in the eOC and arrows show inner hair cells in the native OC. Scale bars: **A–H, I', J', M, N**, 50 μ m; **I, J**, 200 μ m; **M', N', O', P', P', Q, R**, 20 μ m.

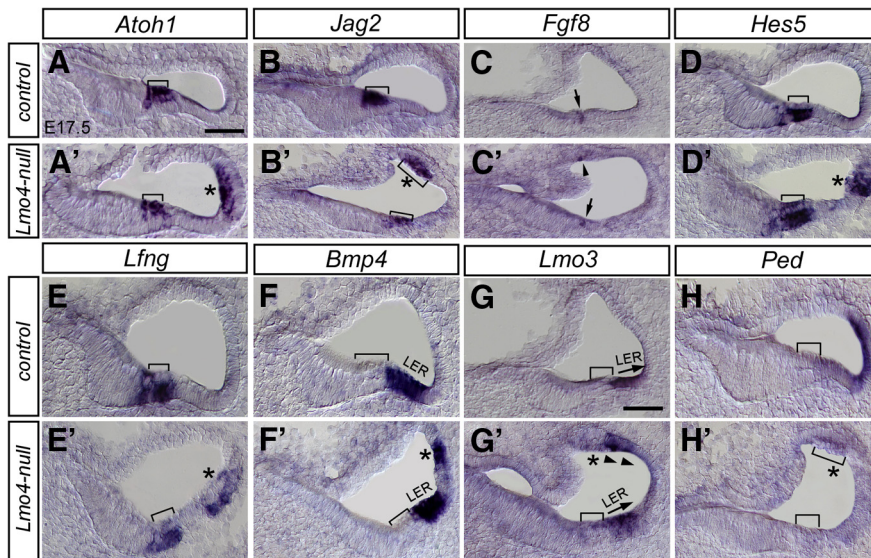


Figure 3. The eOC expresses the appropriate hair cell and supporting cell markers. *In situ* hybridization of E17.5 cochlea sections at E17.5 with indicated cellular markers. **A–C'**, The eOC in the *Lmo4*-null cochlea expresses hair cell markers (*Atoh1* and *Jag2*) and inner hair cell marker (*Fgf8*). Brackets show the normal OC and asterisks indicate the eOC. Arrows in **C** and **C'** show the inner hair cell. Arrowhead in **C'** shows the inner hair cell in the eOC. **D–E'**, Supporting cell markers, *Hes5* and *Lfng*, are expressed in the eOC. **F–G'**, *Bmp4* and *Lmo3* expression identifies a second LER region adjacent to the eOC in the *Lmo4*-null cochlea, suggesting that not only is the OC duplicated, but also the surrounding LER region. Arrows in **G** and **G'** indicate opposite orientation of the LER in the native OC and eOC. **H, H'**, The expression of *Ped* was lost in the *Lmo4*-null cochlea, indicating an ES to OC conversion in *Lmo4*-null cochlea. Scale bar, 50 μ m.

expression in whole mount cochleae using phalloidin to mark the hair cells at P5. We found that the *Lmo4*-null mutants had a duplicated domain of P75 expression in the middle and apical turns of the cochlea (Fig. 2*I–J'*). The extra domain was located on the nonmodiolar (lateral) side of the cochlea, with most of the hair cells lying between the normal and ectopic domain (Fig. 2*J'*), indicating that the ectopic sensory region contains pillar cells. Together, our results demonstrate the formation of an eOC completed with hair cells and supporting cells in the *Lmo4*-null cochlea and that the eOC is generated in an opposite orientation to the native OC.

We also examined selective regional markers outside the OC, such as *Bmp4* and *Lmo3*, both markers for the LER region that lies immediately distal to the OC on the nonmodiolar side. Both of these genes show a duplicated region of expression in the lateral cochlea (Fig. 3*F–G'*), suggesting that not only is the OC duplicated, but also the surrounding regions. The ectopic LER region lies on the modiolar side to the ectopic sensory region, consistent with its opposite orientation to the native organ. Furthermore, the expression of *Ped* was lost in the *Lmo4*-null cochlea (Fig. 3 compare *H* and *H'*), indicating an ES to OC conversion in *Lmo4*-null cochlea. Thus, loss of *Lmo4* not only leads to the formation of an eOC as a mirror-image duplication of the native OC, but also the duplication of its immediate surrounding areas in the ES. Considering that, *Lmo4* is specifically expressed in the GER and ES regions from E14.5, these data also suggest that *Lmo4* normally represses sensory formation in the ES region of the cochlea and that loss of *Lmo4* leads to eOC formation in the ES.

We then investigated whether the eOC developed in a similar fashion to the native organ, including whether the established prosensory markers, such as SOX2, P27KIP1, and JAG1, were expressed before eOC formation. When examined at E14.5, the developmental time period when all of these markers can be de-

tected in the nascent OC, none of these markers demonstrated an ectopic lateral domain. In the control cochlea at E15.5, JAG1 is specifically expressed in the GER, and SOX2 and P27KIP1 are expressed in sensory domain (Fig. 4*A–C*). However, in the *Lmo4*-null cochlea, both SOX2 and JAG1 showed an ectopic lateral domain, whereas the P27KIP1 domain was expanded laterally. These ectopic or expanded domains were consistent with the location of the eOC (Fig. 4*A'–C'*). These results demonstrate that the eOC develops similarly to the native organ, albeit in a later and accelerated time frame, because the prosensory markers appear 1 d after those in the native organ, but the eOC appears to differentiate at approximately the same time as the native organ (Fig. 1).

Previous study has shown that BMP signaling plays a pivotal role in the development of OC (Ohyama et al., 2010). To further explore the potential mechanism underlying the formation of the eOC, we examined the phospho-Smad (pSmad) expression in *Lmo4*-null cochlea. In wild-type controls, pSmad (pSMAD1/5/8) was specifically expressed in the organ of Corti (marked by SOX2 staining; Fig. 4*D–F*). Interestingly, in addition to its expression in the native organ of Corti, we found that pSMAD1/5/8 was also expressed in the ectopic organ of Corti (Fig. 4*G–I*). These results indicate that BMP signaling pathway is activated in the eOC, presumably by the ectopic *Bmp4* expression on the modiolar side of the eOC.

Discussion

The mammalian OC is a highly specialized sensory organ that forms in a discrete ventral region of the cochlea. In this study, for the first time, we have demonstrated the existence of a region in the lateral portion of the mammalian cochlea that is competent to make an OC in the absence of LMO4 function. Previous studies have shown that manipulation of a number of genes and pathways, including *Notch*, *Wnt*, *Atoh1*, *Six1*, *Eya1*, and *Sox2* (Zheng and Gao, 2000; Kawamoto et al., 2003; Shou et al., 2003; Izumikawa et al., 2005; Cafaro et al., 2007; Hartman et al., 2010; Pan et al., 2010; Jeon et al., 2011; Lin et al., 2011; Ahmed et al., 2012; Chai et al., 2012; Kelly et al., 2012; Yang et al., 2012; Mizutari et al., 2013), can induce the formation of hair cells outside the normal region of the OC. However, in none of these cases has an eOC been induced, complete with its diversity of hair cell types, as well as specialized surrounding cell types. Currently, the specification and patterning of the OC is not well understood, although a number of signaling pathways have been implicated, including FGF, Notch, BMP, Wnt, and SHH (Kelly and Chen, 2009). Our results indicate that a second OC-competent region exists in the lateral region of the Corti, and its formation is normally repressed by LMO4. Moreover, we show that loss of *Lmo4* leads to the activation of prosensory markers, such as SOX2, P27KIP1, and JAG1 in the ES region before the formation of the eOC (Fig. 4*A–C'*), suggesting that LMO4 could function upstream of these prosensory proteins and suppress the formation of the prosensory domain.

The reverse orientation of the eOC in relation to the endogenous OC suggests that LMO4 may interact with signaling that originates from a region that lies between the two organs, such as the LER. For example, if a signal arises from the LER that induces OC formation as well as dictating the orientation of the OC, that signal would necessarily lead to two organs on either side of that signal that are mirror images of each other. In comparison, a signal arising from a medial source, such as the GER, to induce and pattern the OC and a lateral competent region would lead to two organs that have similar orientations. A possible candidate for an LER signal is BMP4 (Morsli et al., 1998), as a previous study has suggested that BMP levels are important for OC induction (Ohyama et al., 2010). Here, we show that there is phosphorylated SMAD1/5/8 in the eOC region (Fig. 4G–I), suggesting that LMO4 might normally act to suppress BMP signaling on the lateral side and to specify it into structures such as spiral prominence. In the absence of LMO4, cells in the lateral wall show signs of ectopic responsiveness to BMP signaling, allowing the entire lateral wall to be repatterned as prosensory domain using the same signaling pathways that are normally confined to the native prosensory domain. Intriguingly, we found that the formation of the ectopic prosensory region induces an ectopic region of *Bmp4* expression in the lateral region of cochlea (Fig. 3F'). Thus, an alternative possibility is that during normal development, LMO4 acts to suppress the ectopic domain of *Bmp4* expression in the lateral region of cochlea and that the phosphorylation of SMAD1/5/8 in the eOC (Fig. 4G–I) in *Lmo4*-null mice is secondary to the ectopic production of BMP4. A third possibility is that specification of prosensory domain could precede the specification of the LER region, which in turn influences the differentiation of prosensory domain into the OC.

Although their exact function is unclear, LMO proteins contain two tandem zinc-finger LIM domains for protein-protein interactions but lack the DNA-binding domain typical of LIM-homeodomain (LIM-HD) proteins (Kenny et al., 1998). It has been speculated that LMO proteins could function antagonistically toward LIM-HD proteins by competing for binding to their essential cofactor LDB (LIM-domain binding) proteins. Alternatively, they could function to promote the formation of multimeric transcriptional regulatory complexes by bridging factors, such as basic helix-loop-helix and GATA proteins (Joshi et al., 2009). In hematopoiesis, a large DNA-binding complex containing GATA1, LMO2, and LDB1, has been isolated from erythroid cells (Wadman et al., 1997). Furthermore, it has been shown that the interaction of LMO2 and LDB1 within such a complex acts as a negative regulator of erythroid differentiation (Visvader et al., 1997). Interestingly, in the developing cochlea, GATA3 is expressed in the prosensory domain, LER, and the presumptive ES region and is required for the specification

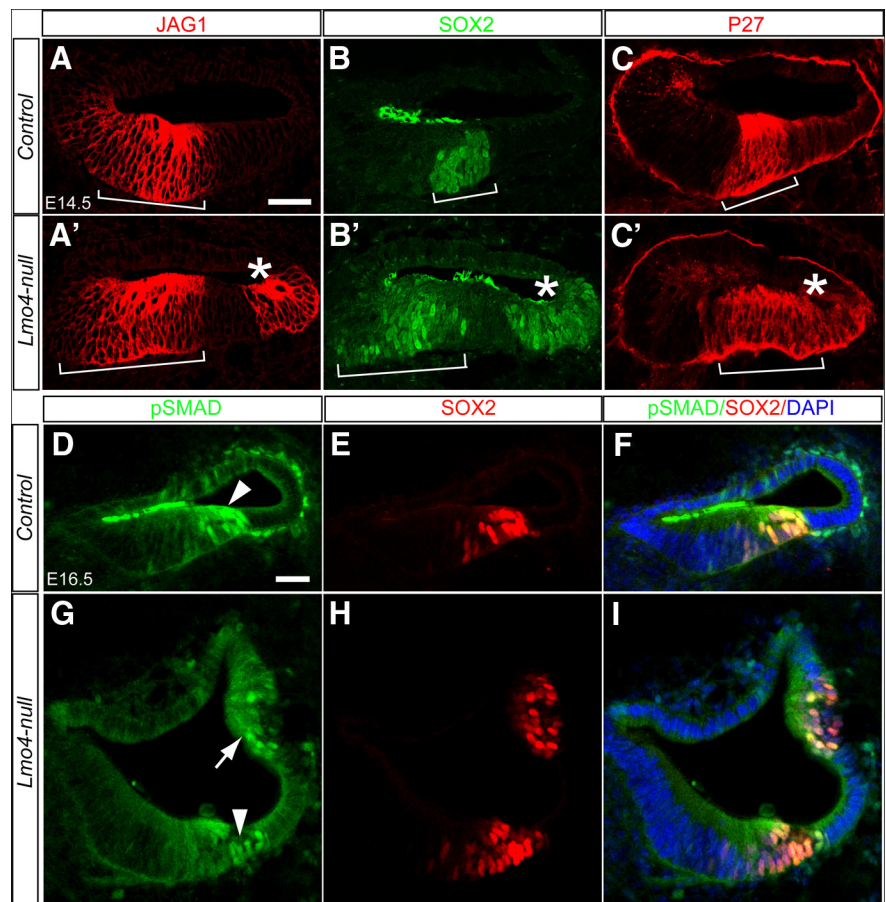


Figure 4. The establishment of ectopic prosensory domain is preceded by the activation of prosensory genes in the *Lmo4*-null cochlea. **A–C'**, Immunostaining reveals regions of ectopic expression of prosensory markers, JAG1 (**A, A'**), SOX2 (**B, B'**), and P27KIP1 (**C, C'**) in the *Lmo4*-null cochlea at E14.5. Brackets show the native prosensory region and asterisks indicate the ectopic prosensory region. **D–I**, Colabeling of *Lmo4*-null cochlea sections at E17.5 with antibodies against SOX2 and pSMAD1/5/8 shows the activation of BMP signaling pathway in the ectopic prosensory region in addition to the native one. Arrowheads indicate the pSMAD1/5/8 staining in the native OC and arrow shows the activation of pSMAD1/5/8 in the eOC in *Lmo4*-null cochlea. Scale bar, 50 μ m.

of prosensory domain. Targeted deletion of *Gata3* in the inner ear significantly impairs the formation of prosensory domain (Luo et al., 2013). Thus, it is conceivable that the expression of GATA3 in the ventral cochlear epithelium (including the ES region) makes it competent to generate prosensory domain but LMO4 inhibits GATA3's role in the ES. More studies are required to resolve these different possibilities and are likely to shed light on the nature, timing, and specificity of the signaling that is required for OC formation in the mammalian cochlea.

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