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Migration of Oligodendrocyte Progenitor Cells Is Controlled by Transforming Growth Factor β Family Proteins during Corticogenesis

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During embryonic development oligodendrocyte precursor cells (OPCs) are generated first in the ventral forebrain and migrate dorsally to occupy the cortex. The molecular cues that guide this migratory route are currently completely unknown. Here, we show that bone morphogenetic protein-4 (Bmp4), Bmp7, and Tgf β 1 produced by the meninges and pericytes repelled ventral OPCs into the cortex at mouse embryonic stages. Ectopic activation of Bmp or Tgf β 1 signaling before the entrance of OPCs into the cortex hindered OPC migration into the cortical areas. OPCs without Smad4 signaling molecules also failed to migrate into the cortex efficiently and formed heterotopia in ventral areas. OPC migration into the cortex was also dramatically reduced by conditional inhibition of Tgf β 1 or Bmp expression from mesenchymal cells. The data suggest that mesenchymal Tgf β family proteins promote migration of ventral OPCs into the cortex during corticogenesis.

Introduction

Long-range cellular migration in the embryonic forebrain is critical for orderly telencephalic development. The germinative zones that generate both neurons and glia are distant from the final positions of these cells, thus understanding regulation of migration is an important question. Forebrain oligodendrocyte precursor cells (OPCs) are born successively in three germinative zones and are distributed widely in the brain where they mature after birth (Rowitch, 2004; Kessaris et al., 2006). The distinct germinative zones also have characteristic timing such that OPCs are produced ventrally first and migrate dorsally to the developing cortex, while the cortex does not produce OPCs from the cortical germinative zone until around birth in mice (Kessaris et al., 2006; Langseth et al., 2010). This raises the question of what cues regulate migration of OPCs generated ventrally to the dorsal forebrain and ultimately whether these same cues might regulate migration of OPCs in the adult brain and in response to injury.

There has been a fairly limited study thus far of factors that regulate directional OPC migration. The migration pattern of OPCs during embryonic development suggests that OPCs might migrate using axonally produced factors that guide OPCs by ax-

onophilic attraction (Prestoz et al., 2004; Tsai et al., 2006; Bribián et al., 2008; Klämbt, 2009; Piaton et al., 2011), but these factors have not been definitely identified (although PDGF-A may be one of these; Yeh et al., 1991; Nakahira et al., 2006). We wondered whether an additional contributor to OPC dorsal migration might be signaling by repellents produced in the ventral forebrain that drive migration of OPCs dorsally.

Bone morphogenetic proteins (Bmps) play critical roles during dorsalization of the neural tube such as the choroid plexus and skull bone development (Hogan, 1996; Zhou et al., 2006), and the expression areas of Bmps expand into the craniofacial mesenchyme forming an ectodermal signaling center that directs craniofacial development starting at E12.5 (Furuta et al., 1997; Foppiano et al., 2007). There is also extensive literature showing that Bmps regulate the development of cells in the oligodendrocyte lineage mostly focusing on two aspects of oligodendrocyte development. Bmps have been shown *in vitro* and *in vivo* to drive the fate of unspecified neural precursor cells away from the oligodendrocyte lineage and toward the production of astrocytes (Gross et al., 1996; Gomes et al., 2003). There is also evidence that Bmps act to retard maturation of OPCs into myelinating oligodendrocytes (See et al., 2004). Much of the focus of this work has presumed that Bmps are mostly produced dorsally in the neural tube during embryonic development (Furuta et al., 1997).

Our recent work shows that the meninges surrounding the forebrain are a major source of secreted Bmps during cortical development (Choe et al., 2013), so we wondered whether the meninges and mesenchymal cells surrounding the ventral forebrain might influence oligodendrocyte lineage cells at the embryonic stages when they are first produced and migrate to the cortex.

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The authors declare no competing financial interests.

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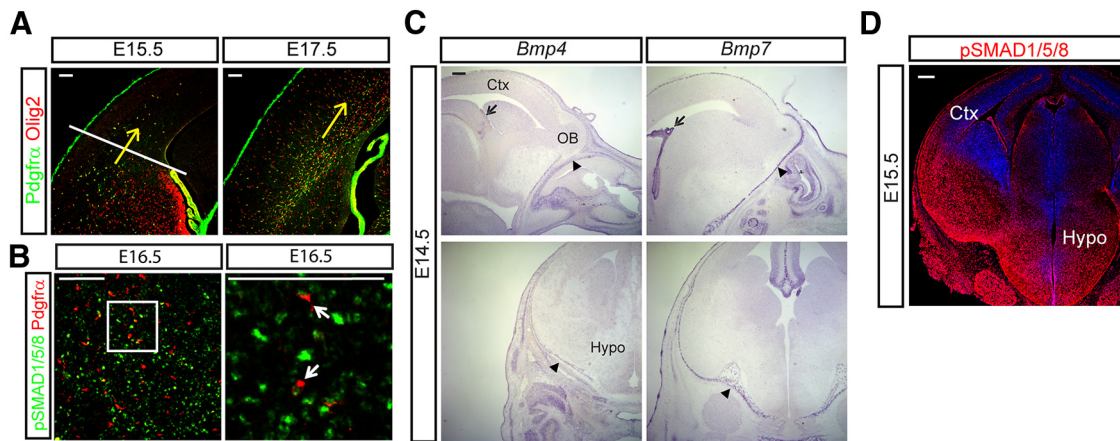


Figure 1. Migration of OPCs into the cortex away from the mesenchymal BMP signaling. **A**, OPC migration into the cortex at E15.5 is shown with double staining for Olig2 and Pdgfr α . An arrow indicates the direction of OPC migration. OPCs localized below the line before entering into the cortex at E15.5. At E17.5, migrating OPCs spread in the cortex. **B**, At E16.5, Pdgfr α -labeled OPCs were stained for pSMAD1/5/8 from the ventral forebrain. A high-magnification image of the box is shown on the right to show Pdgfr α + OPCs are activated by Bmp signaling (arrows). **C**, *In situ* hybridization was conducted to show the expression of Bmp4 and Bmp7 in the head mesenchyme, the meninges, and the choroid plexus at E14.5. Sagittal sections (top) and coronal sections (bottom) are presented. Arrow, choroid plexus; arrowhead, meninges. **D**, Expression of pSMAD1/5/8 at E15.5 embryonic heads. Representative images from at least four independent experiments are presented. Ctx, cortex; Hypo, hypothalamus; OB, olfactory bulb. Scale bars: **A**, **B**, 100 μ m; **C**, **D**, 200 μ m.

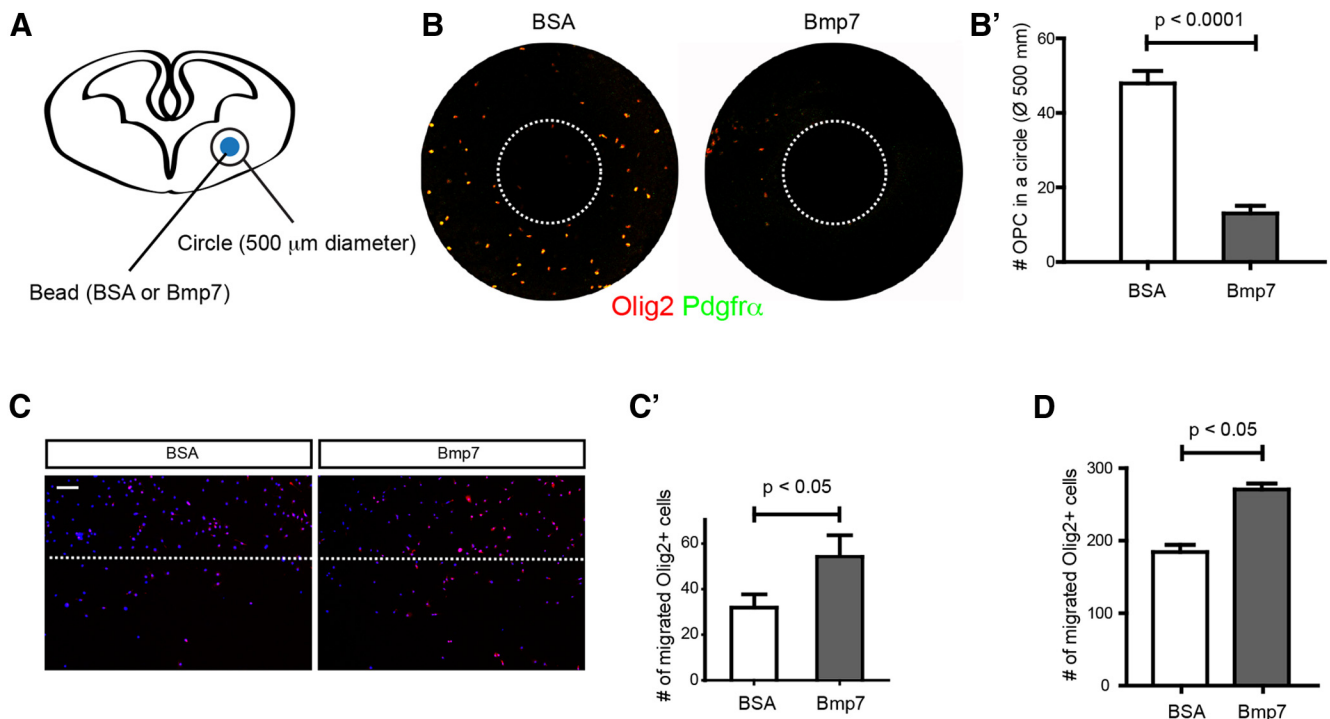


Figure 2. Stimulation of OPC migration by Bmp7 treatment *in vitro*. **A**, A schematic drawing of the bead assay. **B**, Representative images of OPCs stained using vibratome slices (see Materials and Methods). Dotted circles represent the location of a bead. **B'**, Number of OPCs was counted from two independent experiments ($n = 6$). **C**, **C'**, *In vitro*-enriched OPCs were used for the wound-healing assay. Bmp7 (10 ng/ml) was treated for 24 h and cells labeled for Olig2 were counted ($n = 4$). **D**, *In vitro*-enriched OPCs were used for the Boyden chamber assay using culture inserts with 8 μ m pores. Bmp7 (10 ng/ml) was treated for 24 h and Olig2 + cells penetrated to the opposite side were counted ($n = 4$). BSA (0.1%) was used for vehicle treatment. Scale bars: 100 μ m. Error bars depict SEM.

Materials and Methods

Animals. Mice used in this study were previously described [Pdgfr β -Cre (Foo et al., 2006), Sox10-Cre (Matsuoka et al., 2005), Wnt1-Cre (Danielian et al., 1998), Foxc1flx (Hayashi and Kume, 2008), Smad4flx (Bardesy et al., 2006), Tgf β 1flx (Shull et al., 1992), Bmp4flx (Chang et al., 2008), and Bmp7flx (Zouvelou et al., 2009)]. Experimental mice were obtained by crossing male mice carrying an allele of a Cre recombinase and a heterozygous allele of floxed gene to female mice, which carry homozygous floxed genes. The day of vaginal plug was considered to be E0.5. Mouse colonies were housed at the University of California, San

Francisco, in accordance with UCSF Institutional Animal Care and Use Committee (IACUC) guidelines.

***In vitro* OPC migration assay.** For bead experiments, E14.5 CD1 embryos were dissected in 1 \times Neurobasal-A with 10 mM HEPES and 250 μ m coronal sections were prepared using a vibratome (Leica). Vibratome slices were grown on Millicell culture inserts (Millipore) in serum-free culture media containing Neurobasal-A, 1 \times GlutaMAX, and 1 \times B27 (all from Invitrogen). Beads (Bio-Rad; Affi-Gel blue gel, 150–300 μ m) were soaked in Bmp7 (10 ng/ml, 10 μ l; ProSpec) or 0.1% BSA overnight and washed three times in the culture medium before loading

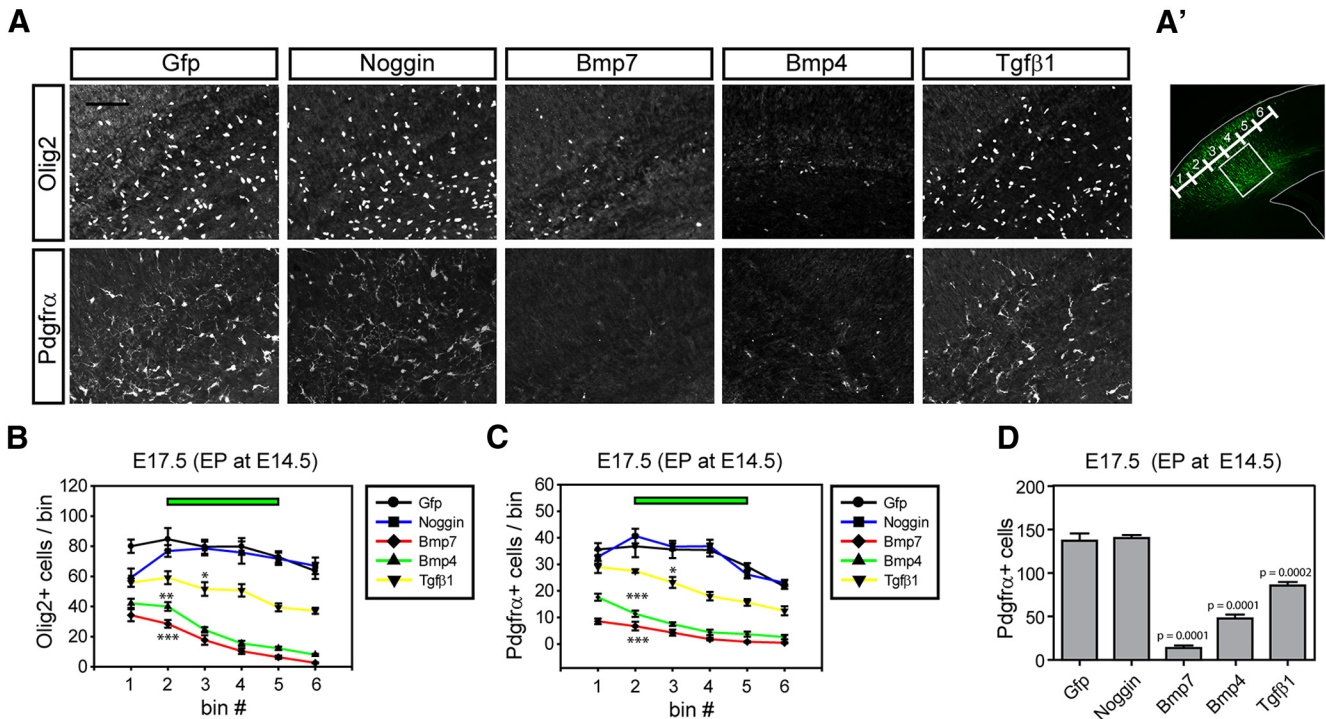


Figure 3. Inhibition of OPC entrance into the cortex by ectopic cortical Bmp or Tgf β signaling. **A**, CD1 mouse embryos were electroporated with Gfp, Noggin, Bmp7, Bmp4, and Tgf β 1 expression vectors at E14.5 and analyzed at E17.5. A representative figure shows the cortex stained for Olig2 and Pdgfr α in the electroporated area ($n = 3$). Images were obtained from the boxed area (**A'**). **B**, **C**, Plots were made from three independent electroporation experiments to show the effect of electroporated gene products on the migration of OPCs labeled with Olig2 (**B**) or Pdgfr α (**C**; $n = 3$). The number of cells was plotted by bins divided by ventral (1) to dorsal (6) electroporated areas as noted in **A'**. **D**, A plot shows total number of Pdgfr α + OPCs in the electroporated area ($n = 3$). Scale bars: 100 μ m. Error bars depict SEM; *** $p < 0.001$, ** $p < 0.005$, * $p < 0.05$.

onto the specific location of slices. Beads were diluted in the culture media at a concentration of 1 bead per 10 μ l and 5 μ l of bead-containing media was loaded onto the slice with a pipette, and slices with a single bead were selected for the experiments. After 24 h, slices with a stabilized bead were briefly fixed in 4% PFA and used for floating staining of Olig2 and Pdgfr α . Olig2 and Pdgfr α + cells around a bead were counted in a circle with a diameter of 500 μ m. To examine the effect of Bmp7 on the motility of Olig2 or Pdgfr α + cells, the wound-healing assay and Boyden migration assay using chambers with 8 μ m pores were conducted. For these assays, E13.5 embryonic forebrains were used to collect and expand Olig2 and Pdgfr α + cells in serum-free culture media containing DMEM/F12 (50:50; Invitrogen), 1 \times GlutaMAX, 1 \times B27, EGF (10 ng/ml; Sigma), FGF (20 ng/ml; Sigma), and PDGF-AA (2 ng/ml; ProSpec). Cells were plated on poly-L-lysine-coated cover glasses for 1 d before making wounds with pipette tips. For Boyden cell migration assay, cells were plated on the poly-L-lysine-coated culture inserts 4 h before treatment. Cells were treated with Bmp7 (10 ng/ml) or 0.1% BSA for 24 h and shortly fixed and stained for Pdgfr α or Olig2 to count cells.

In utero electroporation. The surgery was conducted using timed pregnant CD1 mice according to IACUC-approved protocols at UCSF. Briefly, pregnant mice were anesthetized with Nembutal. The uterine horns were exposed and embryos were injected with 0.5 μ l of 1 mg/ml DNA in TE into the lateral ventricle. Electroporation was conducted at 35 V and 50 and 950 ms with five pulses. All DNA constructs were cloned into the pCIG2-IRES-Gfp vector as described previously (Choe et al., 2012). All experiments were repeated to get four electroporated embryos from more than three independent electroporations. For *in utero* LDN-193189 injection, all procedures were conducted as *in utero* electroporation without electric pulses.

Immunostaining and in situ hybridization. Embryos were collected at noon of embryonic days. Collected brains were fixed in 4% PFA overnight and cryo protected in 20–30% sucrose/PBS for an additional day. OCT-embedded tissues were processed in a cryostat at 12 μ m sections for immunostaining and 20 μ m sections for *in situ* hybridization. All exper-

iments were done by comparing control and mutant sections stained on the same slides to minimize variation between slides. Primary antibodies used for immunostaining are chicken anti-GFP (Aves Labs; 1:1000), rabbit anti-Pdgfr β (Cell Signaling Technology; 1:200), rabbit anti-Ki67 (LabVision; 1:200), rabbit anti-Olig2 (Millipore; 1:300), rabbit anti-Blp (Millipore; 1:300), rat anti-Pdgfr α (Millipore; 1:300), rabbit anti-pSMAD1/5/8 (Cell Signaling Technology; 1:200), rabbit anti-Prox1 (Bagri et al., 2002), and rat anti-Ctip2 (Abcam; 1:1000). Templates for RNA probes used for *in situ* hybridization were designed according to the Allen Developing Mouse Brain Atlas. *In situ* hybridization was conducted as described in a recent publication (Choe et al., 2013). To double stain Pdgfr β proteins on the sections, which were first stained with a Tgf β 1 mRNA probe by *in situ* hybridization, signals for the antigen were amplified using Tyramide Signal Amplification plus kits (PerkinElmer). Images were acquired at the Nikon Imaging Center at UCSF using an upright Nikon C1 spectral confocal microscope equipped with 405, 488, and 561 nm lasers with 10 \times , 20 \times , and 63 \times objectives.

Statistics. Values are presented as the SEM in the graphs. For the statistical analysis of samples, we used Student's *t* test or one-way ANOVA with Tukey's multiple-comparison test using a SigmaPlot program (Systat Software) and a GraphPad program (GraphPad Software) unless otherwise stated in the figure legend.

Results

Extensive activation of Bmp signaling in the embryonic ventral forebrain

During mid-corticalogenesis, facial mesenchymal and meningeal cells produce Bmps, which regulate craniofacial morphogenesis (Nomura and Li, 1998; Foppiano et al., 2007) and cortical development (Choe et al., 2012, 2013). OPCs born in the ventral germinal areas migrate into the cortex at E14.5 and occupy the cortex by E17.5 when the cortical OPCs form a migratory stream in the ventricular zone and SVZs (Fig. 1A, arrows show direction

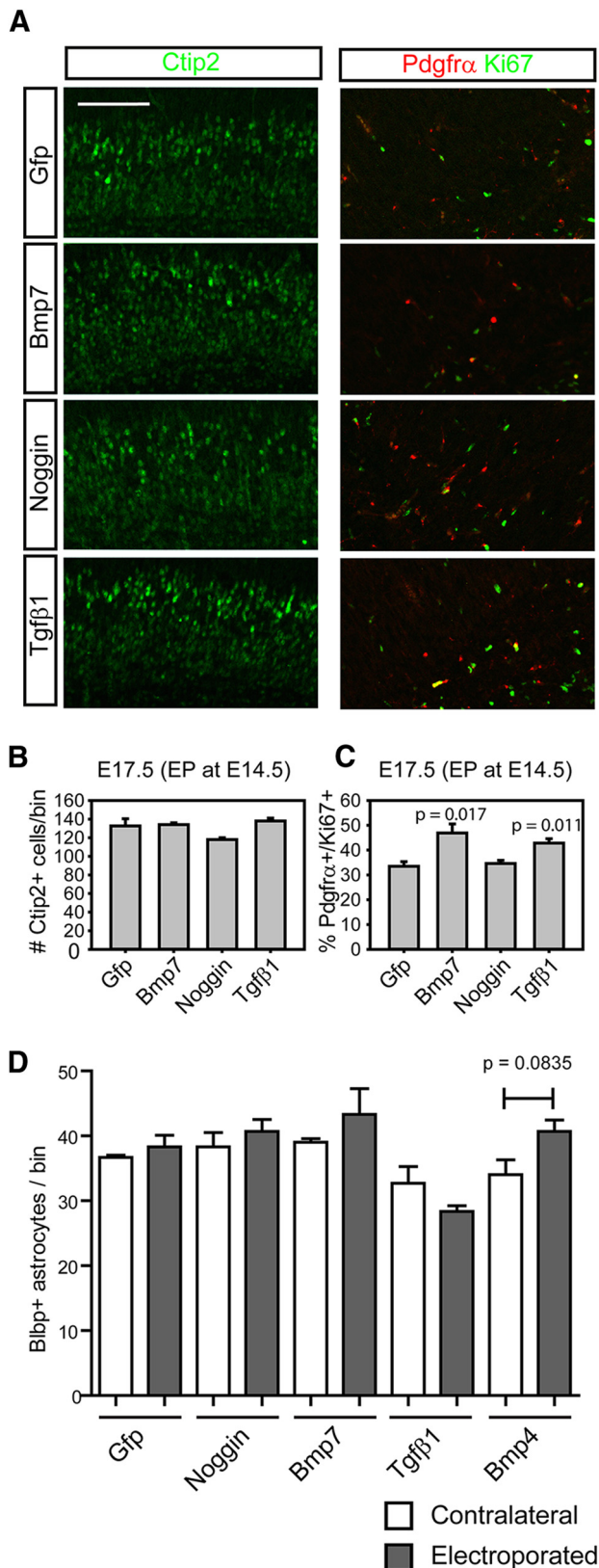


Figure 4. The effects of Bmp or Tgfβ activation on the cortical neurogenesis and OPC proliferation. **A**, Using the electroporated cortices from Figure 3, we stained CtIp2 to see the neurogenesis that occurs at the time of the electroporation and double stained Pdgfrα and Ki67 to see the OPCs in the active cell cycle. **B**, **C**, Plots were made from data collected from **A** to show the effects of the electroporated gene products on neurogenesis (**B**) and the proliferation of OPCs (**C**; $n = 3$). **D**, A plot was made from counting Blbp+ cells from the sections used in Figure 3 ($n = 3$). Scale bars: 100 μm. Error bars depict SEM.

of OPC migration). If ventral Bmp signaling is involved in the development of OPCs or their migration dorsally into the cortex, the ventral OPCs should be exposed to Bmp activation before entering the cortex. Staining for phospho(p)SMAD1/5/8, an indicator of active Bmp signaling, and Pdgfrα, an OPC marker, at E16.5 when ventral OPCs were motile showed that OPCs in the ventral forebrain had evidence of Bmp signaling (Fig. 1B). *In situ* hybridization for *Bmp4* and *Bmp7* showed that the facial mesenchymal cells, the meninges, the choroid plexus, and hippocampal neuroepithelial cells were the main source of Bmps at E14.5 (Fig. 1C). Interestingly, despite a lack of expression of the relevant ligands, Bmp4 or Bmp7, within the ventral forebrain (Fig. 1C), there was very strong pSMAD1/5/8 staining widely seen in the ventral forebrain (Fig. 1D). These data suggest that Bmp4 and Bmp7 expressed by the meninges and facial mesenchyme are likely to be driving significant BMP signaling in the ventral embryonic forebrain.

Bmp7 regulates OPC migration *in vitro*

Our results showing that the ventral forebrain has prominent Bmp signaling at embryonic stages suggest that these signals may have specific effects on the behavior of OPCs. Previous studies showed that Bmp signaling in the germinative zones increases astrocyte production (Gross et al., 1996; Gomes et al., 2003), but the area of high Bmp signaling is excluded from the germinative zones of the ventral forebrain. The other published effect of Bmp signaling on the oligodendrocyte lineage is the effect on differentiation, which tends to keep OPCs in an undifferentiated state blocking myelin production (See et al., 2004) and the distribution of pSMAD1/5/8 staining seems possibly consistent with this effect, but we also wondered whether it is possible that polarized production of Bmp7 by the meninges might have an instructive effect on OPC migration. To test this we soaked beads in Bmp7 (10 ng/ml, 10 μl) and placed them on a 250 μm slice from the E14.5 ventral forebrain and then examined the effect on local OPC migration (Fig. 2A). We found that Bmp7-soaked beads dramatically reduced the number of OPCs in the area surrounding the bead after 24 h, suggesting that Bmp7 drives migration of ventral OPCs as a repellent (Fig. 2B,B'; $p < 0.0001$, $n = 6$). To examine whether Bmp7 induces cellular motility of OPCs, wound-healing (Fig. 2C,C') and Boyden chamber assays (Fig. 2D) were conducted using *in vitro*-expanded OPCs. In both assays, OPCs labeled using Olig2 showed increased cell motility after Bmp7 treatment ($p < 0.05$, $n = 4$). These *in vitro* assays are consistent with the data showing that Bmps can act as repellents driving OPC migration.

Tgfβ family ligands act as repellents for OPC migration *in vivo*

Ventrally produced OPCs migrate into the cortex during embryogenesis and very few OPCs found in the cortex are locally produced before P1 (Kessar et al., 2006; Langseth et al., 2010). Previous studies showed that conditional mutation of the type I Bmp receptor, Bmpr1a, in the ventral oligogenic niche results in a twofold decrease in the number of oligodendrocytes in the forebrain at birth (Samanta et al., 2007). These previous findings suggest that disrupted Bmp signaling might affect oligodendrocyte lineage specification or also migration of OPCs in the forebrain. Bmp signaling controls astroglial specification but analysis of OPC migration was not previously examined in the spinal cord explicitly (Gross et al., 1996; Mekki-Dauriac et al., 2002; Gomes et al., 2003; Miller et al., 2004). We hypothesized that mesenchymal/meningeal Bmp ligands are involved in OPC migration to the cortex, which might be a feature of OPC migration in the fore-

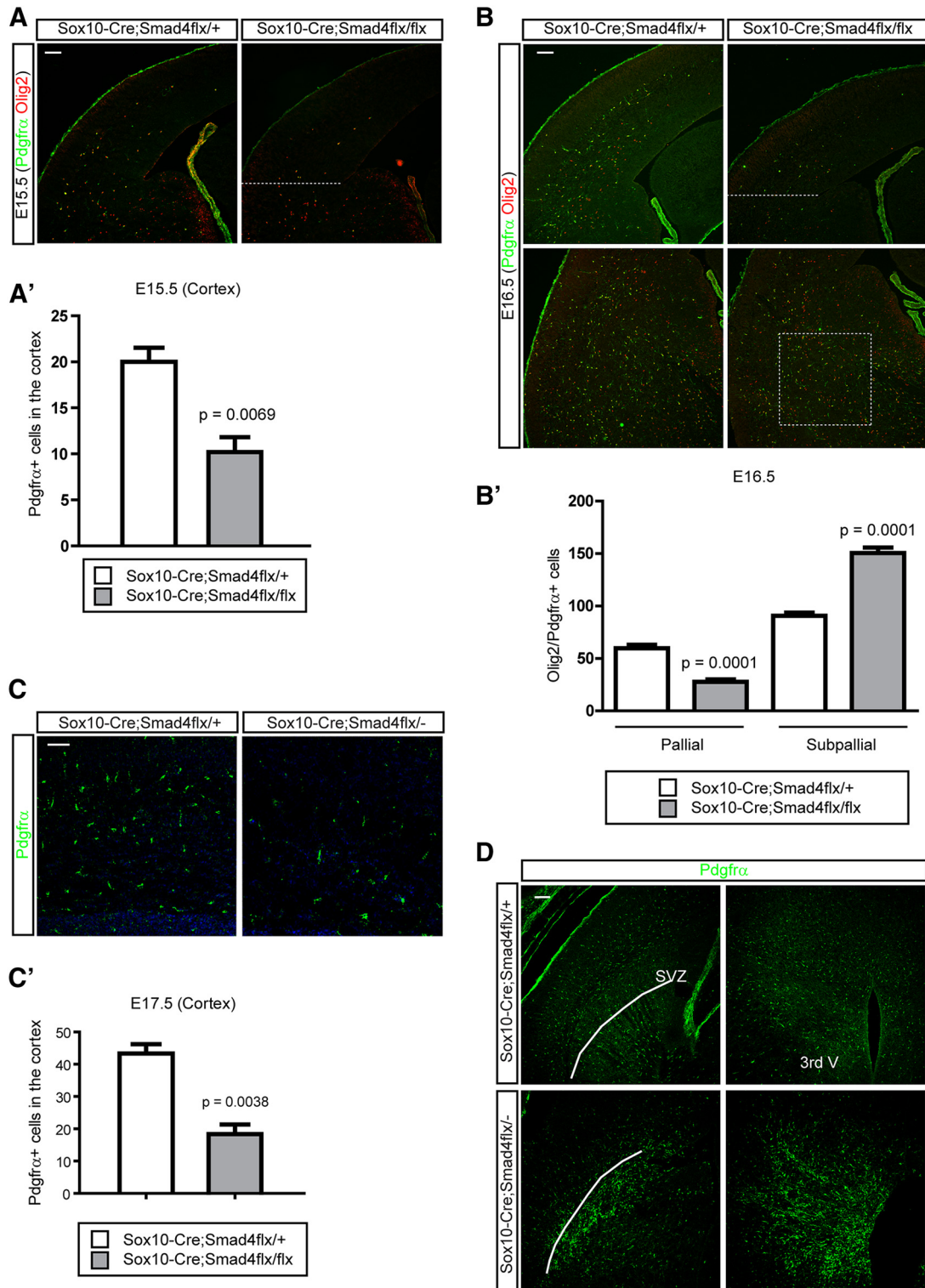


Figure 5. The effects of genetic loss of Smad4 gene expression on OPC migration. **A, A'**, To remove the Smad4 gene expression in the OPC lineage, we used Sox10-Cre to conditionally inhibit Smad4. At E15.5 when OPCs migrate into the cortex we counted OPCs labeled with Pdgfrα and Olig2 in the cortex above the dotted line. A graph depicts the quantitative results from four litters (**A'**; $n = 4$). **B, B'**, A day later at E16.5, we counted OPCs in the cortex (pallial) and subcortical boxed regions (subpallial). A graph depicts the quantitative results from four litters ($n = 4$; **B'**). **C, C'**, A day before the mutant embryos die, we stained the cortex with Pdgfrα to show the OPCs at E17.5. A plot was made from data collected from three litters at E17.5 ($n = 4$). **D**, E17.5 forebrains were stained for Pdgfrα to show the heterotopic OPCs in the two main germinal areas, SVZ, and third ventricle (3rd V) in the mutant embryos. Scale bars: 100 μm. Error bars depict SEM.

brain. To address this, we first decided to examine the effects of ectopic expression of relevant Bmp ligands by *in utero* electroporation to target the cortex laterally, in the route taken by OPCs as they migrate. We electroporated Bmp7, Bmp4, Noggin, and

Tgfβ1 expression vectors into the E14.5 cortex before the initiation of ventral OPC migration into the cortex and examined the migration pattern of OPCs at E17.5. OPCs in the contralateral cortex or control Gfp-electroporated cortex did not show

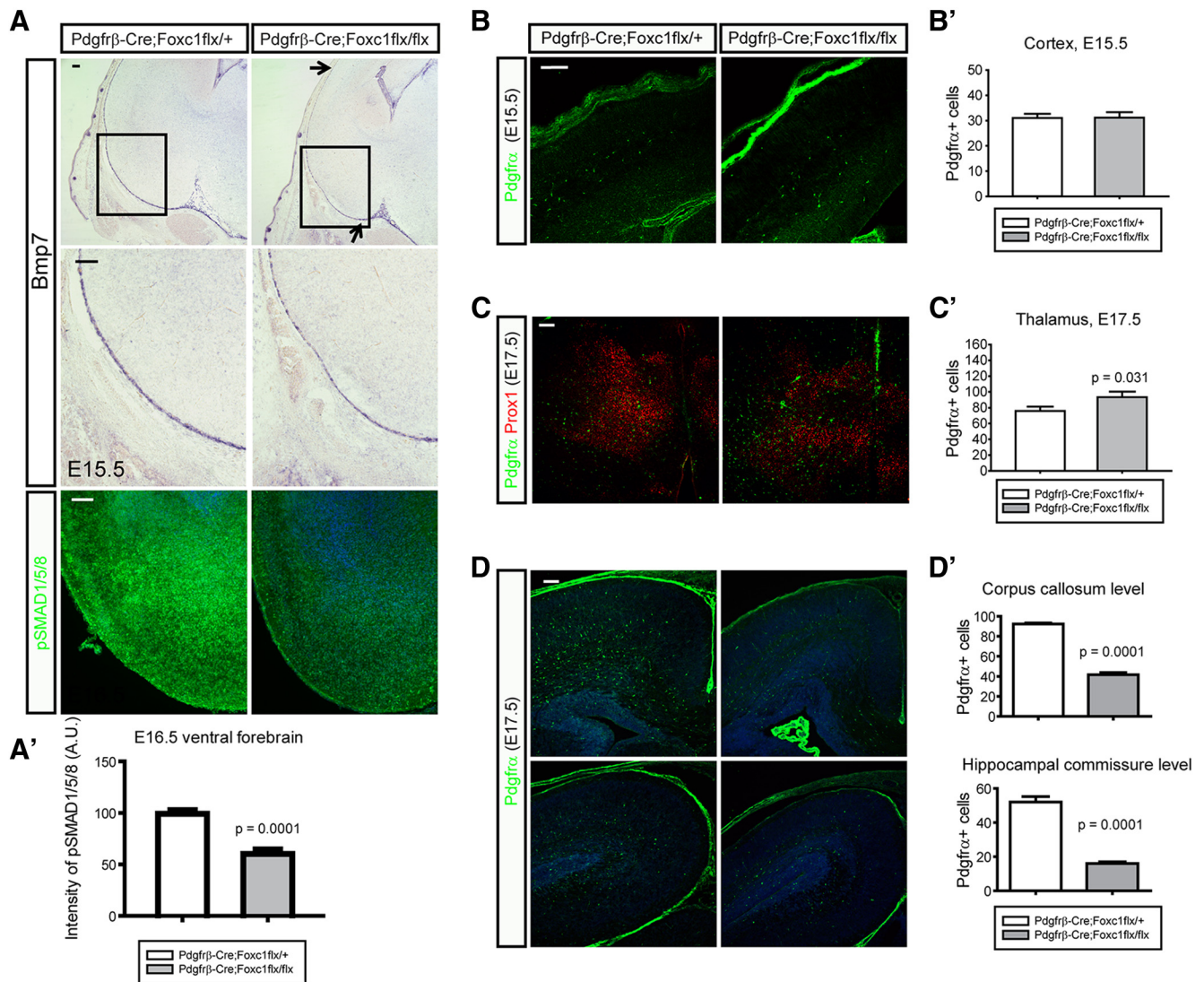


Figure 6. Meninges-mediated migration of OPCs into the cortex. **A**, To obtain the reduced meningeal expression of Bmps, a Pdgfr β -Cre mouse was bred to Foxc1 conditional mutant to get Pdgfr β -Cre;Foxc1flx/flx mutants. *In situ* hybridization of Bmp7 shows reduced Bmp7 expression in the meninges of ventral forebrain at E15.5 (arrows). Higher magnification images of boxed areas are presented in the middle. Staining of pSMAD1/5/8 shows the reduced activation of Bmp signaling in the mutant ventral forebrain at E16.5 (bottom). **A'**, A graph shows the intensity of pSMAD1/5/8 in the ventral forebrain ($n = 4$). A measure function of ImageJ software was used. **B, B'**, Cortical OPCs were stained for Pdgfra and counted at E15.5. **C, C'**, Ventral OPCs were stained for Pdgfra and counted at E17.5. Prox1 was used to stain the thalamic neurons to match the sections. $p = 0.031$ (Mann–Whitney rank sum test; $n = 4$). **D, D'**, Cortical OPCs were counted in the anterior (top) to the posterior (bottom) cortex ($n = 4$). Error bars depict SEM. Scale bars: 100 μ m.

changes of OPC migration patterns; however, most of the OPCs (Olig2+/Pdgfra+) in the Bmp7 or Bmp4-electroporated cortex stalled before they confronted Bmp expression in the electroporated area and resulted in a large decrease in OPCs within the electroporated regions (Fig. 3A–D; $n = 3$). OPCs in the Noggin-electroporated cortex showed a similar migration pattern to the Gfp-electroporated cortex while OPCs in the Tgf β 1-electroporated cortex showed a moderate decrease in the numbers of migrating OPCs. These results support the idea that the mesenchymal Tgf β family molecules such as Bmp7, Bmp4, and Tgf β 1 may repel OPCs from ventral sources helping to drive them to migrate toward the cortex.

To be sure that our manipulations did not lead to changes in neuronal production in the E14.5 cortex, which might somehow lead to secondary changes in OPC distribution, we stained Ctip2 to label the deep-layer neurons born at the time of electroporation. We could not see any significant changes of Ctip2-expressing cells by ectopic Bmp7 expression (Fig. 4A, B; $n = 3$).

We also costained Ki67 in the Pdgfra-expressing cells to check whether the Bmp expression inhibited the proliferation of OPCs in the cortex. Interestingly we found that 10% more OPCs expressed Ki67 in Bmp7 or Tgf β 1-electroporated area, which rules out the possibility that Bmp7 or Tgf β 1 inhibited OPC proliferation in the electroporated area leading to the changes in local OPC numbers (Fig. 4A, C; $n = 3$). Bmp signaling is also reported to promote astrocyte differentiation in the gliogenic niche (Gross et al., 1996; Zhu et al., 1999); such an effect could decrease the number of OPCs if astrocyte production is increased. We stained the sections with an astrocyte marker, Blbp, and found the Blbp staining intensity was increased by Bmps but not by Tgf β 1 and that the number of Blbp+ cells was not significantly changed compared with nonelectroporated contralateral cortex (Fig. 4D; $n = 3$). This result suggests that Bmps reduced OPC migration not diverting OPCs into astrocytes.

This combination of results indicates that presentation of ectopic sources of Bmp ligands (either by beads or electroporation) leads to

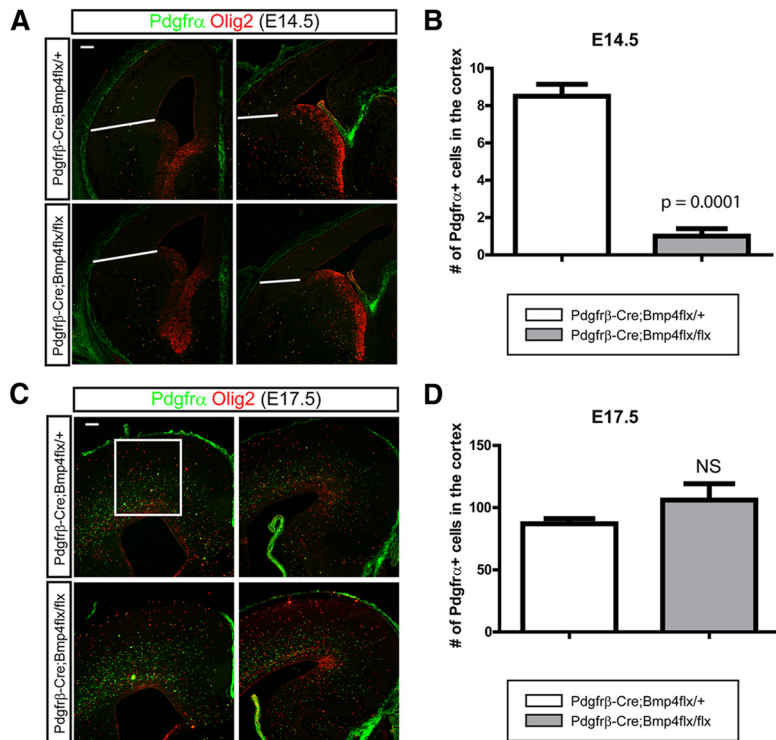


Figure 7. The effects of inactivation of Bmp4 from the mesenchymal tissues on the migration of OPCs into the cortex. To inhibit the expression of Bmp4 from the mesenchymal tissues, we used Pdgfrβ-Cre to ablate the Bmp4 allele in the mesenchymal tissues including meninges. **A**, Immunostaining of OPCs for Olig2 and Pdgfra at E14.5. **B**, OPCs in the cortex were counted from the area above the line ($n = 3$). **C**, Immunostaining of OPCs for Olig2 and Pdgfra at E17.5. **D**, OPCs in the cortex were counted from the area in the box ($n = 3$). Scale bars: 100 μm. Error bars depict SEM.

altered migration behavior of OPCs. This effect is most consistent with a chemorepellent effect of Bmps supplied by the ventral forebrain acting to direct migration of OPCs toward the developing cortex.

Cell-autonomous inhibition of Tgfβ signaling in the OPCs inhibits OPC migration

We next wanted to test whether inhibition of Tgfβ ligand family signaling in OPCs would affect OPC migration in a cell-autonomous manner. We used Sox10-Cre (Matsuoka et al., 2005) to target OPCs to delete Smad4, a coregulator of the Smad1/5/8 and Smad2/3 signaling pathways. We counted OPCs in the cortex just as OPCs enter the cortex at E15.5 and again a day later at E16.5 in Sox10-Cre;Smad4flx/flx compared with littermate controls (Fig. 5A,A',B,B'). Fewer OPCs were found in the cortex at both E15.5 and E16.5 and more OPCs remained in the subpallial region in the mutant at E16.5 (Fig. 5B,B'; $p = 0.0001$, $n = 4$). At E17.5, mutant cortices also contained significantly fewer Pdgfra + cells (Fig. 5C,C'; $p = 0.0038$, $n = 4$). These results are most consistent with an effect on OPC migration leading to increased numbers of OPCs remaining in the ventral forebrain and failing to reach the cortex. Interestingly, we found that mutant OPCs in the Sox10-Cre;Smad4flx/flx embryos formed heterotopias near their sites of origin in the anterior entopeduncular region and the pallial–subpallial border, probably because of their failure to appropriately migrate and spread the remaining heterotopias in the germinative areas (Fig. 5D). These genetic loss-of-function data are consistent with our other data indicating that the Tgfβ and Bmp signaling pathways are repellent for OPCs supporting migration into the cortex and perhaps also involved in the spreading of OPCs in the cortex.

Meningeal Bmps regulate entrance and scattering of OPCs in the cortex

In recent studies we showed that the meninges are a major source of Bmp ligands during forebrain development and that these ligands regulate both neuronal axon guidance and the formation of the dentate gyrus (Choe et al., 2012, 2013). This led us to examine mutant mice with defects in Bmp production in the meninges to determine whether this leads to effects on OPC migration in the cortex. We used Pdgfrβ-Cre to remove Foxc1 expression during mid-corticogenesis resulting in reduced Bmp expression in the cortical meninges after E14.5 (Choe et al., 2012). For exam-

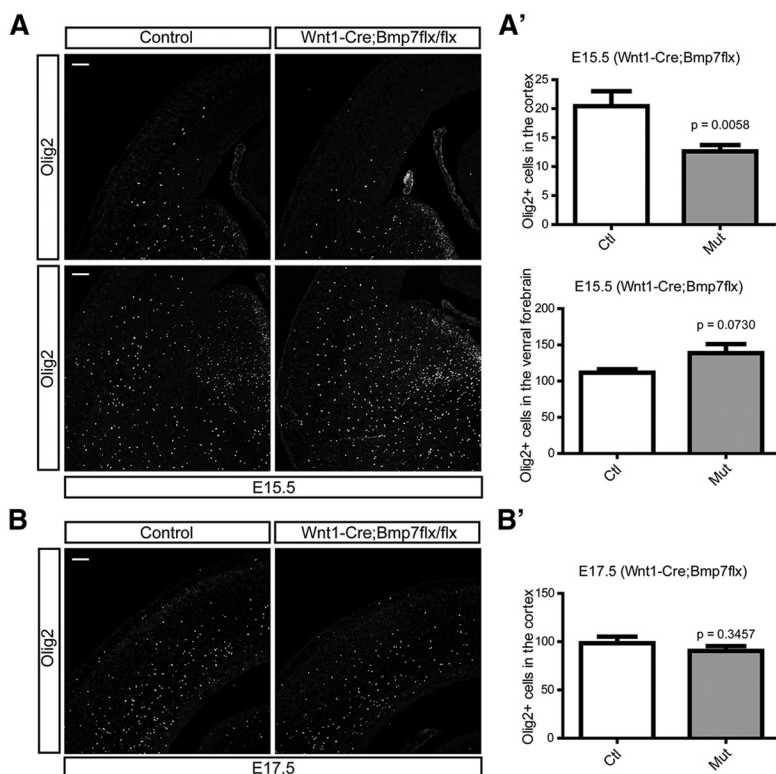


Figure 8. The effects of meningeal Bmp7 on the migration of OPCs into the cortex. To inhibit the expression of Bmp7 from the meninges, we used Wnt1-Cre to ablate the Bmp7 conditional allele in the neural crest cell lineages. **A,A'**, Immunostaining for Olig2 shows cortical (top) and ventral OPCs (bottom) at E15.5 (**A**) and graphs show number of Olig2 + OPCs ($n = 3$; **A'**). **B,B'**, OPCs in the cortex were stained using the E17.5 embryonic cortex ($n = 3$) and the number of cortical OPCs was plotted in a graph (**B'**). Scale bars: 100 μm. Error bars depict SEM.

ple, Bmp7 expression in the mutant meninges covering the ventral forebrain was modestly decreased (similar to our previous observations in the midline meninges; Choe et al., 2012; Fig. 6A). Consistent with this reduction in Bmp7 expression, pSMAD1/5/8 signaling was also reduced in the ventral forebrain at E16.5 (Fig. 6A'; $p = 0.0001$, $n = 4$). Next, we counted cortical and ventral OPCs in the heterozygote control and mutant embryos and found that the OPCs in the cortex were similar at E15.5 when meningeal Bmp7 expression starts to decrease (Fig. 6B,B') but that the number of the ventral OPCs at E17.5 was slightly higher in the mutant (Fig. 6C,C'; $p = 0.031$, $n = 4$). At this embryonic stage, the cortical OPCs at the anterior (corpus callosum level) and the posterior (hippocampal commissure level) regions were both significantly reduced (Fig. 6D,D'; $p = 0.0001$, $n = 4$). The results support the idea that mesenchymal tissues are the main source of Bmps regulating OPC migration during corticogenesis.

The results with Foxc1 mutant mice indicate that there is a likely role for Bmp ligands from the meninges in regulating OPC migration; however, since the meninges express a variety of ligands we wished to more specifically address the role of Bmp ligands. To do this we conditionally ablated Bmp4 expression, enriched in mesenchymal and meningeal tissues during early corticogenesis, using the mesenchymal tissue- and pericyte-specific Pdgfr β -Cre (Foo et al., 2006). At E14.5, we found a significant decrease of OPCs in the cortex accompanying the loss of Bmp4 expression (Fig. 7A,B; $p = 0.0001$, $n = 3$) but we found that the migration of OPCs into the cortex had recovered by E17.5 (Fig. 7C,D). We felt it likely that this represents compensation by the other Tgf β family proteins such as Bmp7 also expressed by mesenchymal and meningeal cells (Fig. 1C). To address this possibility, we used Wnt1-Cre to inactivate a Bmp7 conditional allele to inhibit Bmp7 expression from neural crest-derived meningeal tissues (Choe et al., 2013). At E15.5, similar to Bmp4 conditional mutants, cortical OPCs were significantly reduced ($p = 0.0058$, $n = 3$) while the ventral OPCs were slightly increased (Fig. 8A,A'; $p = 0.0730$, $n = 3$). However, as we saw in Bmp4 mutants, the cortical OPC number also recovered by E17.5 (Fig. 8B,B'). This implies functional redundancy in the roles of Bmp4 and Bmp7 in regulating OPC migration.

Inhibition of type I Bmp receptors hinders OPC migration into the cortex

Because of the apparent redundancy between Bmp ligands and the fact that Smad4 is involved in both Bmp and Tgf β signaling, we wished to devise another means to selectively examine the role

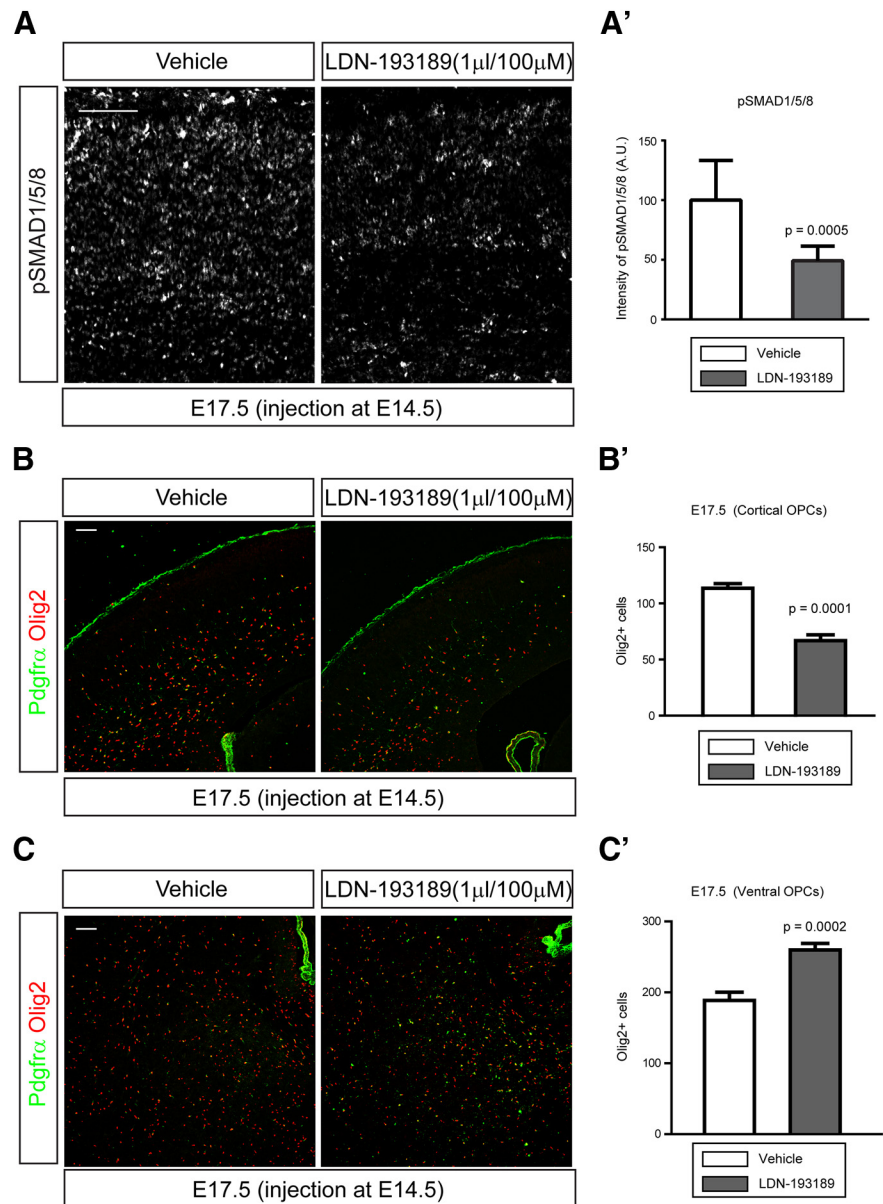


Figure 9. Delayed migration of OPCs into the cortex by inhibition of type I Bmp receptors. To inhibit type I Bmp receptors, 1 μ M of LDN-193189 (100 μ M) was injected into the lateral ventricle of E14.5 embryos and cortices were analyzed at E17.5. One percent DMSO was injected as a vehicle control. **A**, Representative images of cortices stained for pSMAD1/5/8. **A'**, A graph shows the intensities of pSMAD1/5/8 ($n = 4$). A measure function of ImageJ software was used. **B**, Representative images of cortices stained for both Pdgfra and Olig2 at E17.5. **B'**, A graph shows the number of OPCs in the cortex ($n = 4$). **C**, Representative images of ventral OPCs stained for both Pdgfra and Olig2 at E17.5. **C'**, A graph shows the number of OPCs in the ventral forebrain ($n = 4$). Scale bars: 100 μ m. Error bars depict SEM.

of Bmp signaling in OPC migration. To do this we used LDN-193189, a cell-permeable, small molecule inhibitor of type I Bmp receptors (Cuny et al., 2008). *In utero* injection of LDN-193189 (1 μ M/100 μ M in 1% DMSO) into the lateral ventricle of E14.5 embryos led to reduced staining for pSMAD1/5/8 in the E17.5 forebrain indicating a general effect on Bmp signaling (Fig. 9A,A'; $n = 4$). We examined the distribution of OPCs in the E17.5 cortex by staining with Pdgfra and Olig2 antibodies and compared the number of OPCs between inhibitor- and vehicle-injected embryos. The total number of OPCs was significantly reduced in the LDN-193189-injected cortex (Fig. 9B,B'; $p = 0.0001$, $n = 4$) while more OPCs remained in the ventral forebrain (Fig. 9C,C'; $p = 0.0002$, $n = 4$) compared with controls. This strongly suggests that activation of

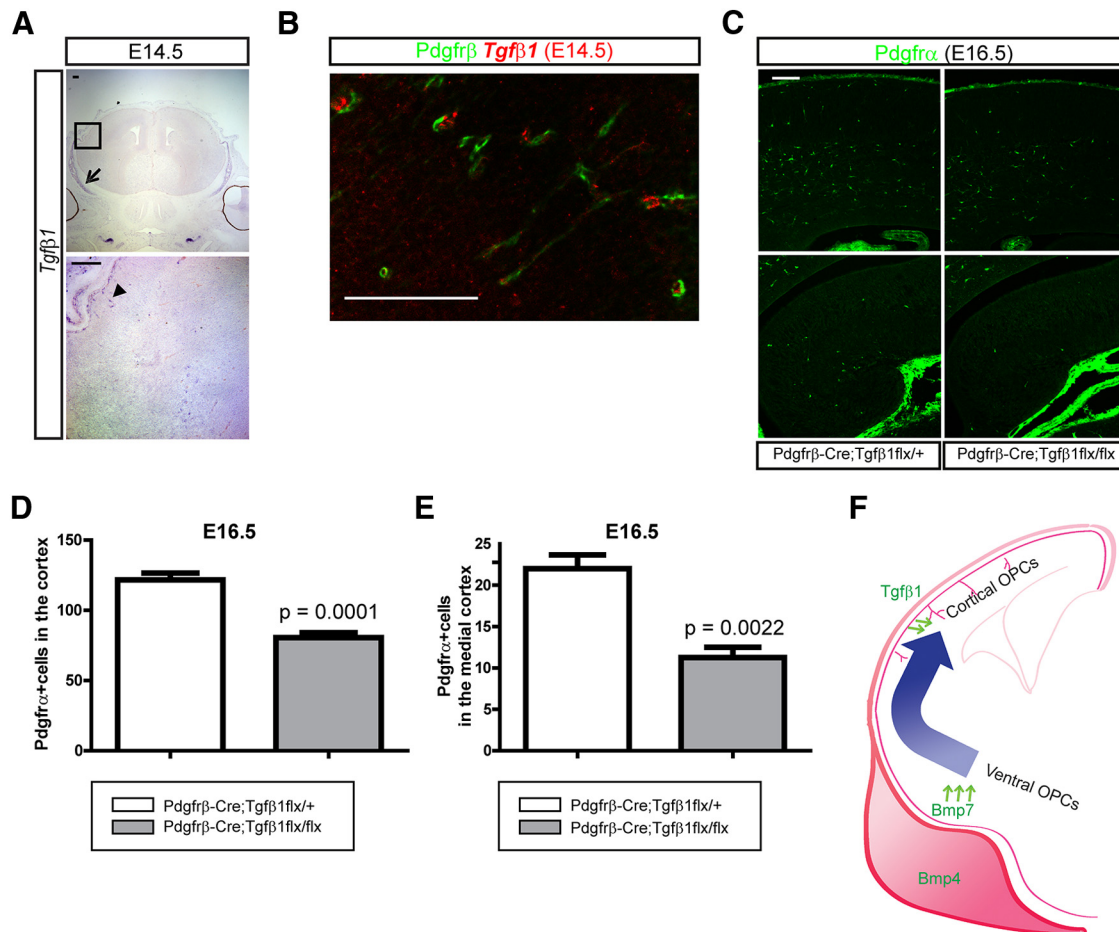


Figure 10. Tgf β 1-mediated migration of OPCs into the cortex. **A**, Expression of Tgf β 1 in the E14.5 embryonic heads was examined by *in situ* hybridization. An arrow indicates calvarial mesenchymal cells and an arrowhead indicates blood vessels. A high-magnification image of the boxed area of the upper image is shown in the bottom. **B**, Double *in situ* hybridization and immunostaining of Tgf β 1 mRNA and Pdgfr β (pericytes) in the E14.5 cortex. **C**, Immunostaining of the Pdgfr α using E16.5 embryos from Pdgfr β -Cre;Tgf β 1flx/+ and Pdgfr β -Cre;Tgf β 1flx/flx embryos shows reduced migrating OPCs in the cortex (top; **D**) and in the mediadorsal cortex and the hippocampus (bottom; **E**) of mutant embryos. **D, E**, Pdgfr α + OPCs was counted from the cortex (**D**) and the mediadorsal cortex (**E**) as shown in **C** ($n = 3$). **F**, A schematic drawing illustrates migration of ventral OPCs into the cortex by the repelling activities of Tgf β family proteins from the mesenchymal tissues such as calvarial mesenchyme (Bmp4), meninges (Bmp4 and Bmp7), and the pericytes (Tgf β 1). Scale bars: 100 μ m. Error bars depict SEM.

type I Bmp receptors is important for OPC migration into the cortex.

Mesenchymal and vascular Tgf β 1-mediated OPC migration in the cortex

In our ectopic expression studies we showed that Tgf β 1 also had a mild repelling activity for cortical OPC migration (Fig. 3). There are many instances where Bmp and Tgf β signaling pathways exhibit cross talk during development (Spinella-Jaegle et al., 2001; Keller et al., 2011) and, as we showed, deletion of Smad4, which signals in both pathways, has a significant effect on OPC distribution. To examine whether the repelling effect of Tgf β 1 on OPC migration could also be observed from the direct loss of Tgf β 1 expression, we wished to generate conditional mutants lacking Tgf β 1 expression. First, we checked expression of Tgf β 1 in the mesenchymal tissues at E14.5. Tgf β 1 was expressed in the calvarial mesenchymal cells and blood vessels at E14.5 (Fig. 10A), which raises the possibility that Tgf β 1 could play a role to fine-tune the localization of OPCs within the cortex. We performed double staining of Tgf β 1 mRNA and Pdgfr β (which marks pericytes in the cortex) using E14.5 embryonic cortex. Tgf β 1 was mostly expressed by pericytes, which face neuronal and glial cells in the cortical vasculature and a few unidentified cells (Fig. 10B).

To ablate Tgf β 1 expression in the calvarial mesenchymes and cortical pericytes, we crossed Pdgfr β -Cre mice to Tgf β 1 conditional mutant mice. Staining of OPCs with Pdgfr α in the E16.5 embryonic cortex showed that less OPCs were found in the neocortex (Fig. 10C,D; $p = 0.0001$, $n = 3$) and in the dorsomedial cortex and the hippocampus (Fig. 10C,E; $p = 0.0022$, $n = 3$). This demonstrates that the mesenchymal and vascular Tgf β 1 molecules also directly regulate migration of OPCs in the cortex and indicate a complex set of overlapping roles for Tgf β superfamily ligands in regulation of OPC positioning and migration.

Discussion

In this study we showed that Bmps from the meninges and vascular Tgf β 1 coordinate OPC migration into the cortex during mouse embryonic life. Previous studies showed that Bmp signaling regulates migration of neural crest cells and osteoblast cells, thus playing a critical role in craniofacial morphogenesis (Lind et al., 1996; Dudas et al., 2004; Goldstein et al., 2005; Sotobori et al., 2006). Our data show that the same cues regulate migration of OPCs into the cortical areas. It is interesting to consider whether this role of Bmps in developmental distribution of OPCs is relevant to adult situations where OPCs are also required to migrate. Support for this possibility comes from studies where the Bmp

inhibitor Noggin enhanced remyelination in chemically induced demyelination (Sabo et al., 2011) although some of this was probably due to effects on the differentiation of OPCs (See et al., 2007; Wang et al., 2011). It is also reported that the demyelinating lesions have increased expression of Bmps in neurons, astrocytes, and OPCs (Fuller et al., 2007; Ara et al., 2008; Dummula et al., 2011), suggesting that changes in Bmp ligands are perhaps physiologically relevant in demyelination and might be responsible in part for failure of OPCs to fully occupy demyelinated lesions in the forebrain. In this respect, developing cocktails of Tgf β and Bmp inhibitors that block signaling in the vicinity of demyelinating injuries might provide therapeutic opportunities for multiple sclerosis.

One of the more fascinating features of OPCs is their even distribution in the CNS. A recent study clearly suggests that this is in part due to mutual repulsion of OPCs in the cortex (Hughes et al., 2013). Our study also provides a potential mechanism helping to control OPC distribution, i.e., expression by OPCs of Bmp ligands. In fact, OPCs in the spinal cord were known to express Bmp4 (Kondo and Raff, 2004) and our unpublished work shows that OPCs in the postnatal corpus callosum express Bmp4, which we have shown is a repellent for OPCs. Thus OPCs would be “tiled” in the forebrain by mutual repulsion via the Bmp signaling pathway. We intend to revisit this idea using genetic approaches allowing inhibition of the Bmp signaling pathways in individual postnatal OPCs and examining their distribution in the cortex. It is also possible that there are other OPC-attracting signaling pathways that would counterbalance the repelling activities of Tgf β family molecules further ensuring the even tiling of OPCs. Embryonic OPCs migrate on the thalamocortical axon fibers (Nakahira et al., 2006), a putative source of PDGF-A (Yeh et al., 1991), and the elongating thalamocortical axons could act as an axophilic path that confines OPCs in the migratory stream by counteracting repelling activities of Tgf β family molecules. In the end, we suspect that the even distribution of OPCs in the adult brain will be the result of interactive tuning by both repulsive and attractive signals.

Previous studies showed that the Bmp and Tgf β signaling pathways are also involved in the proliferation, differentiation, and specification of OPCs (McKinnon et al., 1993; Gross et al., 1996; Gomes et al., 2003; See et al., 2004, 2007; Zhang et al., 2010; Sabo et al., 2011). Bmp signaling is a potent stimulator of astrocyte fates in multipotential neural precursor cells and Bmp inhibitors are involved in biasing these precursors toward neuronal or OPC lineages (Gross et al., 1996; Lim et al., 2000). Of particular interest, increased Chordin, an inhibitor of Bmp signaling, in lysoclethrin-induced demyelination of the corpus callosum leads to recruitment of more neuronal progenitor-derived oligodendrocytes to the corpus callosum (Jablonska et al., 2010). This evidence suggests that Bmp signaling regulates fate determination of neuronal progenitors into oligodendrocyte lineage cells in normal contexts as well as in pathologic situations. Interestingly, there is also extensive evidence that Bmp signaling modulates the differentiation of OPCs into myelinating oligodendrocytes, with Bmps acting as inhibitors of maturation and promoting persistence of cells in an immature state (See et al., 2004; Sabo et al., 2011). These included several studies examining the *in vitro* effects of Bmps on OPCs as well as others examining Bmp receptor mutant mice, all showing that Bmps are important brakes on the differentiation of OPCs into oligodendrocytes (See et al., 2007; Wang et al., 2011). This led to examination of Bmp inhibitors as potential regulators of repair of demyelination in toxic or inflammatory models and showed that Noggin, a Bmp inhibitor, accel-

erates repair and remyelination (Dizon et al., 2011; Sabo et al., 2011). Thus Bmps inhibit specification of cells in the oligodendrocyte lineage while at the same time maintaining their undifferentiated state once produced. In our work we show that OPCs have potent migratory responses to Bmp signaling, particularly when the Bmps are presented as they are expressed during embryonic development, from limited anatomic sources such as the meninges. It is interesting to consider the possibility that the block in OPC maturation and migratory responsiveness to Bmps are linked mechanistically. It seems likely that the transcriptional program responsible for maintaining OPCs in an undifferentiated state may also support the expression of factors involved in the long-range migration and homing of OPCs in development and perhaps after injury.

In this study we have shown that Bmps act as regulators of OPC migration and distribution in the developing forebrain. It is interesting to consider the potential roles of Bmps within areas of injured white matter in demyelinating or ischemic injury as a key regulator of the repair process in these situations. The previously published roles of Bmp ligands as inhibitors of OPC maturation have made examination of these as therapeutic nodes for disease important and our study implies that these measures may also have potent effects on OPC migration in disease states.

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