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# Bone Morphogenic Protein Signaling Is a Major Determinant of Dentate Development

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To understand life-long neurogenesis in the dentate gyrus (DG), characterizing dentate neural stem cells and the signals controlling their development are crucial. In the present study, we show that bone morphogenic protein (Bmp) signaling is a critical regulator of embryonic dentate development, required for initiating neurogenesis in embryonic DG progenitors and required for the establishment of dentate neural stem cells postnatally. We tested the hypothesis that Bmp signaling regulates dentate development in part by controlling the expression of *Lef1*, a Wnt responsive transcription factor expressed in dentate stem cells and absolutely required for dentate granule cell production. Bmp activation through the *Acvr1* receptor induced *Lef1* expression and neurogenesis in the embryonic DG. Ectopic expression of *Bmp7* in the embryonic midline increased DG neurogenesis and inhibition of local Bmp signaling decreased embryonic DG neurogenesis. Mice with selective loss of Bmp expression due to defective meningeal development or with selective conditional deletion of meningeal *Bmp7* also have dentate developmental defects. Conditional deletion of Activin receptor type I (*Acvr1*) or *Smad4* (a downstream target nuclear effector of Bmp signaling) in DG neural stem cells resulted in defects in the postnatal subgranular zone and reduced neurogenesis. These results suggest that *Acvr1*-mediated meningeal Bmp signaling regulates *Lef1* expression in the dentate, regulating embryonic DG neurogenesis, DG neural stem cell niche formation, and maintenance.

## Introduction

The mammalian forebrain has two regions with active ongoing neurogenesis into adulthood: the subgranular zone (SGZ) in the dentate gyrus (DG), and the subventricular zone lining the lateral ventricles. In both sites the stem cell niche is established and maintained by a panoply of signals including Wnt, Sonic hedgehog (Shh), and bone morphogenic proteins (Bmps), which cooperate to maintain the neurogenic capacity of the niche (Galceran et al., 2000; Chenn and Walsh, 2002; Machold et al., 2003; Zhou et al., 2004; Lie et al., 2005; Machon et al., 2007; Favaro et al., 2009; Caronia et al., 2010; Mira et al., 2010; Munji et al., 2011). While many studies address the roles of individual morphogenic signaling pathways, these neural stem cell niches, illuminating the interplay of these signaling pathways is critical to understanding the physiologic and pathophysiologic regulation of new neuron production.

Previous studies showed that Wnt signaling is pivotal in the development of the embryonic DG and in postnatal DG stem cell niche signaling. Mice with mutations in critical components of the Wnt signaling pathway show defective DG development and

loss of DG neural stem cells (Galceran et al., 2000; Lee et al., 2000; Zhou et al., 2004; Li and Pleasure, 2005). In particular, *Lef1*, a Wnt-activated transcription factor selectively expressed in the developing dentate, is required for dentate granule neuron production (Galceran et al., 2000). Activation of the Wnt signaling pathway also directs the restricted expression of *Prox1* in DG granule neurons and regulates DG neurogenesis in the adult (Lie et al., 2005; Machon et al., 2007; Karalay et al., 2011). The role of Bmp signaling in dentate development is far less established, although there have been a few relevant publications. Conditional deletion of *Smad4*, a common transcriptional regulator for transforming growth factor  $\beta$  (*Tgfb*) signaling pathways, in the adult DG radial neural stem cells reduced dentate neurogenesis (Colak et al., 2008; Caronia et al., 2010) and compromised quiescence of adult dentate neural stem cells (Mira et al., 2010). In addition, a recent study showed that post-translational control of *Noggin* (*Nog*) expression is probably involved in DG neurogenesis as well (Guo et al., 2011). However, the role of Bmp signaling in establishing the dentate has been far less clear. The one study that explicitly examines this found only modest effects of loss of function for both *Bmpr1a* and *1b* (Caronia et al., 2010). Thus, how Bmp signaling is integrated with other niche signals in regulating DG neurogenesis remains largely unexplored.

In the present study, we show that *Lef1* is a landmark signaling molecule expressed by the DG neuronal stem cells throughout development of the DG and that activation of the Bmp signaling pathway through Activin receptor type I (*Acvr1*, also known as *Alk2*) regulates the expression of *Lef1* in the DG stem cells at embryonic and postnatal stages. This study provides novel in-

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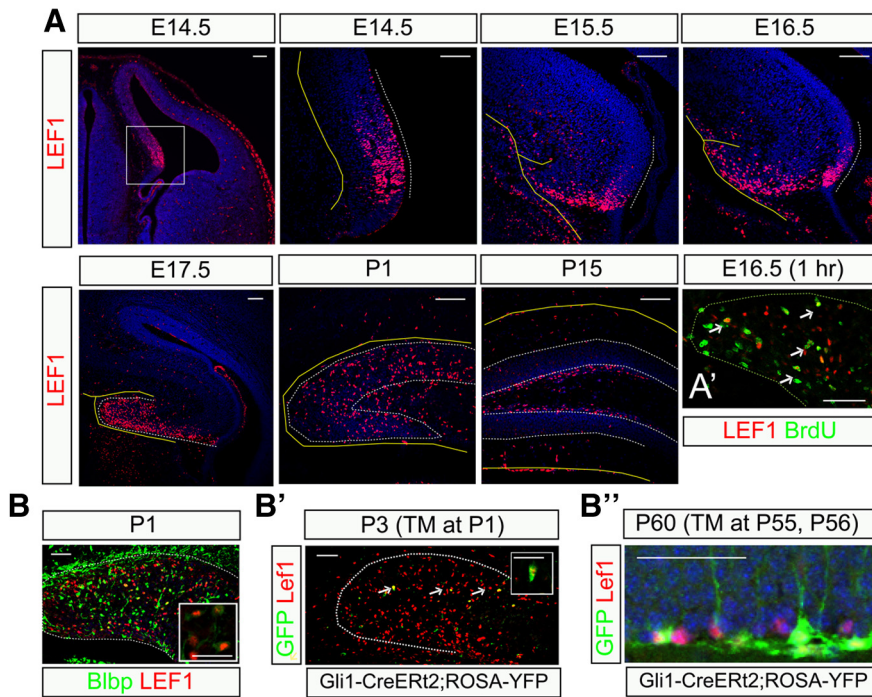
Author contributions: Y.C. and S.J.P. designed research; Y.C. performed research; A.K. and D.G. contributed unpublished reagents/analytic tools; Y.C. and S.J.P. analyzed data; Y.C. and S.J.P. wrote the paper.

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**Figure 1.** Expression of Lef1 during DG development. **A**, Lef1 expression was seen in the DG neural epithelium at E14.5 and in migrating cells toward the meninges (yellow lines) from E15.5 to E17.5. Lef1+ cells were observed in the granule cell layer and the hilus of the P1 DG and the P15 SGZ. Dashed lines in the top depict the dentate neural epithelial ventricular zone and dashed lines in the bottom outline the DG. **A'**, Pregnant CD1 mice were injected with BrdU (50 mg/kg) at 1 h before collection at E16.5. The DG was stained with Lef1 and BrdU to show the expression of Lef1 in the dividing cells. The dashed line outlines the DG and yellow lines mark the meninges covering the area of the DG. **B**, The DG of P1 pups was stained with Lef1 and Blbp with higher power images in the inset. The dashed line outlines the DG. **B'**, Shh-responsive DG neural stem cells were marked in Gli1-CreERT2 mice. TM was injected at P1 and Cre recombinant cells visualized with ROSA-YFP reporter mice at P3. Cells costained with GFP and Lef1 are indicated by arrows (inset). The dashed line outlines the DG. **B''**, TM was given at P55, P56 of Gli1-CreERT2; ROSA-YFP animals and stained for GFP and Lef1 at P60. Scale bars: 100  $\mu$ m; insets of **B** and **B'**, 50  $\mu$ m.

sight into how multiple signaling pathways regulating DG neurogenesis might cooperate.

## Materials and Methods

**Animals.** Mice used in this study were previously described (Gli1-CreERT2, Ahn and Joyner, 2004; hGFAP-Cre, Zhuo et al., 2001; Smad4<sup>lox</sup>, Bardeesy et al., 2006; Acvr1<sup>lox</sup>, Kaartinen and Nagy, 2001; Tgfr2<sup>lox</sup>, Chytil et al., 2002; Pdgfr $\beta$ -Cre, Foo et al., 2006; Foxc1<sup>lox</sup>, Hayashi and Kume, 2008; Wnt1-Cre, Danielian et al., 1998; Bmp7<sup>lox</sup>, Zouvelou et al., 2009) and ROSA-YFP Cre reporter mice were obtained from The Jackson Laboratory. 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 carrying homozygous floxed alleles. All mice were maintained in a mixed background and experimental mice were compared with littermate controls. The day of vaginal plug was considered to be embryonic day 0.5 (E0.5). Mouse colonies were housed at the University of California, San Francisco (UCSF), in accordance with UCSF Institutional Animal Care and Use Committee (IACUC) guidelines.

**In utero electroporation.** Timed pregnant CD1 mice were purchased from Charles River and the surgery was conducted according to IACUC approved protocols at UCSF. Briefly, the CD1 pregnant females were anesthetized with Nembutal. The uterine horns were exposed and embryos were injected with 1  $\mu$ l (1 mg/ml) DNA in TE into the lateral ventricle. Electroporation was conducted at 33 V, 50 ms, 950 ms with five pulses. All DNA constructs were cloned into the pCIG2-IRES-EGFP vector from Dr. Franck Polleux by basic DNA cloning techniques. All template full-length cDNA was purchased from Open Biosystems. All experiments were repeated to get six electroporated embryos from more than four different electroporated mice. A constant active variant of

Acvr1 (CA-Acvr1) and a kinase dead (KD) variant of Acvr1 (KD-Acvr1) depict Q207D (Macias-Silva et al., 1998) and K235R (Visser et al., 2001) point mutants respectively.

**Tamoxifen and bromodeoxyuridine injection.** Tamoxifen (TM; Sigma) stock was prepared by dissolving the powder in corn oil (Sigma) at 20 mg/ml. Neonates were single dosed subcutaneously with 2 mg of TM and adult mice were dosed daily for 2 d with intraperitoneal injection of 4 mg of TM/40 g per animal. Timed pregnant mice were subcutaneously injected with bromodeoxyuridine (BrdU) (Roche) dissolved in saline (10 mg/ml) at the dose of 50 mg/kg animal.

**Immunostaining and in situ hybridization.** Embryos were collected at noon of embryonic days. Collected brains were fixed in 4% paraformaldehyde (PFA) in PBS overnight and cryoprotected in 20% sucrose/PBS for an additional day. OCT-embedded tissues were processed in a cryostat at 12  $\mu$ m sections for immunostaining and 20  $\mu$ m sections for *in situ* hybridization. Primary antibodies used for the immunostaining are chicken anti-GFP (Aves Labs; 1:1000), rabbit anti-Ki67 (LabVision; 1:200), rat anti-PECAM (Abcam; 1:1000), mouse anti-Reelin (Millipore; 1:1000), rabbit anti-pSMAD1/5/8 (Cell Signaling Technology; 1:100), rabbit anti-BLBP (Millipore Bioscience Research Reagents; 1:500), rabbit anti-GFAP (DAKO; 1:1000), rabbit anti-Tbr2 (Abcam; 1:200), rabbit anti-Prox1 (Bagri et al., 2002), and rabbit anti-Lef1 (Cell Signaling Technology; 1:200). Templates for RNA probes used for *in situ* hybridization were designed according to the Allen Developing Mouse Brain Atlas. All experiments were done by comparing control and mutant sections stained on the same

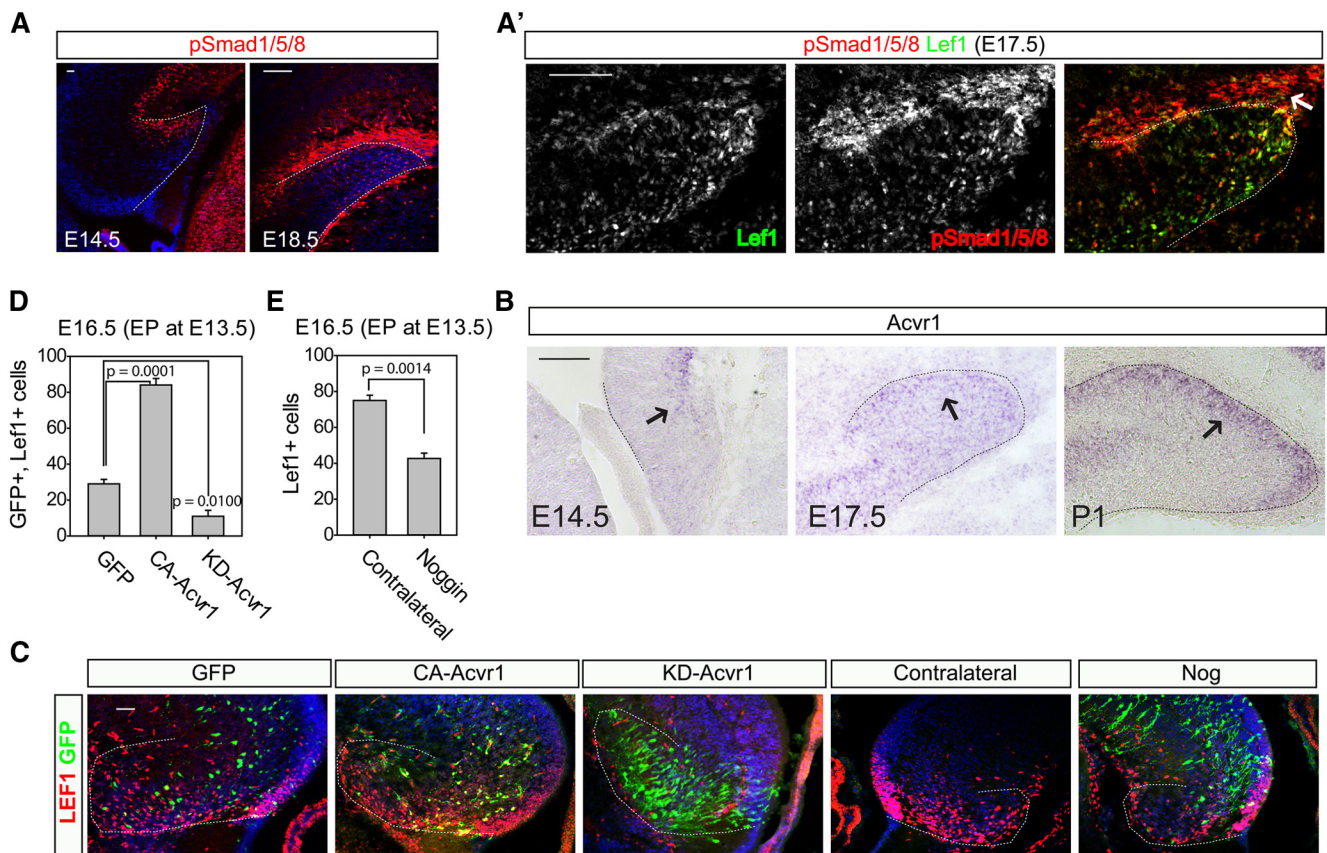
slides to minimize variation in an experiment. For *in situ* hybridization, slides with 20  $\mu$ m sections were warmed to room temperature and treated with proteinase K (50  $\mu$ g/ml) for 1 min, and fixed with 4% PFA for 10 min. Acetylation was performed using 0.25% acetic anhydride in 0.1 M triethanolamine, pH 8.0, for 10 min, followed by three PBS washes. Slides were incubated with a hybridization buffer (50% formamide, 5 $\times$  SSC, 0.3 mg/ml yeast tRNA, 100 mg/ml heparin, 1 $\times$  Denhardt's, 0.1% Tween 20, 0.1% 3-[(3-cholamidopropyl) dimethylammonio]-1-propane-sulfonate, 5 mM EDTA) for 10 min at 65°C, followed by overnight incubation with digoxigenin (DIG)-labeled probes (500 ng/ml) in a hybridization buffer. Five high-stringency washes were performed with 0.2 $\times$  SSC at 65°C. Slides were then incubated with alkaline phosphatase (AP)-conjugated anti-DIG antibodies (Roche) followed by the detection of signals with nitroblue tetrazolium/5-bromo-4-chloro-indolyl phosphate (Roche). Images were acquired at the Nikon Imaging Center at UCSF using an upright Nikon C1 spectral confocal microscope.

**Statistics.** For the pairwise analysis of samples we used Student's *t* test of the SigmaPlot program (Systat Software). For multiple group comparison, we used one-way ANOVA with Tukey's multiple-comparison test using the GraphPad Prism5 program (Graphpad Software). Error bars indicate  $\pm$  SEM.

## Results

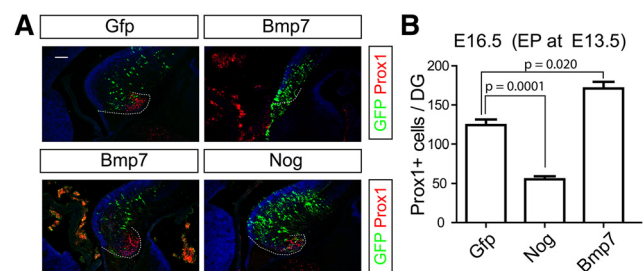
### Expression of Lef1 in DG stem cells across development

Lef1 mutant mice have a prominent dentate phenotype that goes along with the restricted expression of Lef1 mRNA within the developing forebrain to the dentate neuroepithelium and forming DG (Galceran et al., 2000; Zhou et al., 2004). We first set out to check the expression pattern of a Lef1 protein during the de-



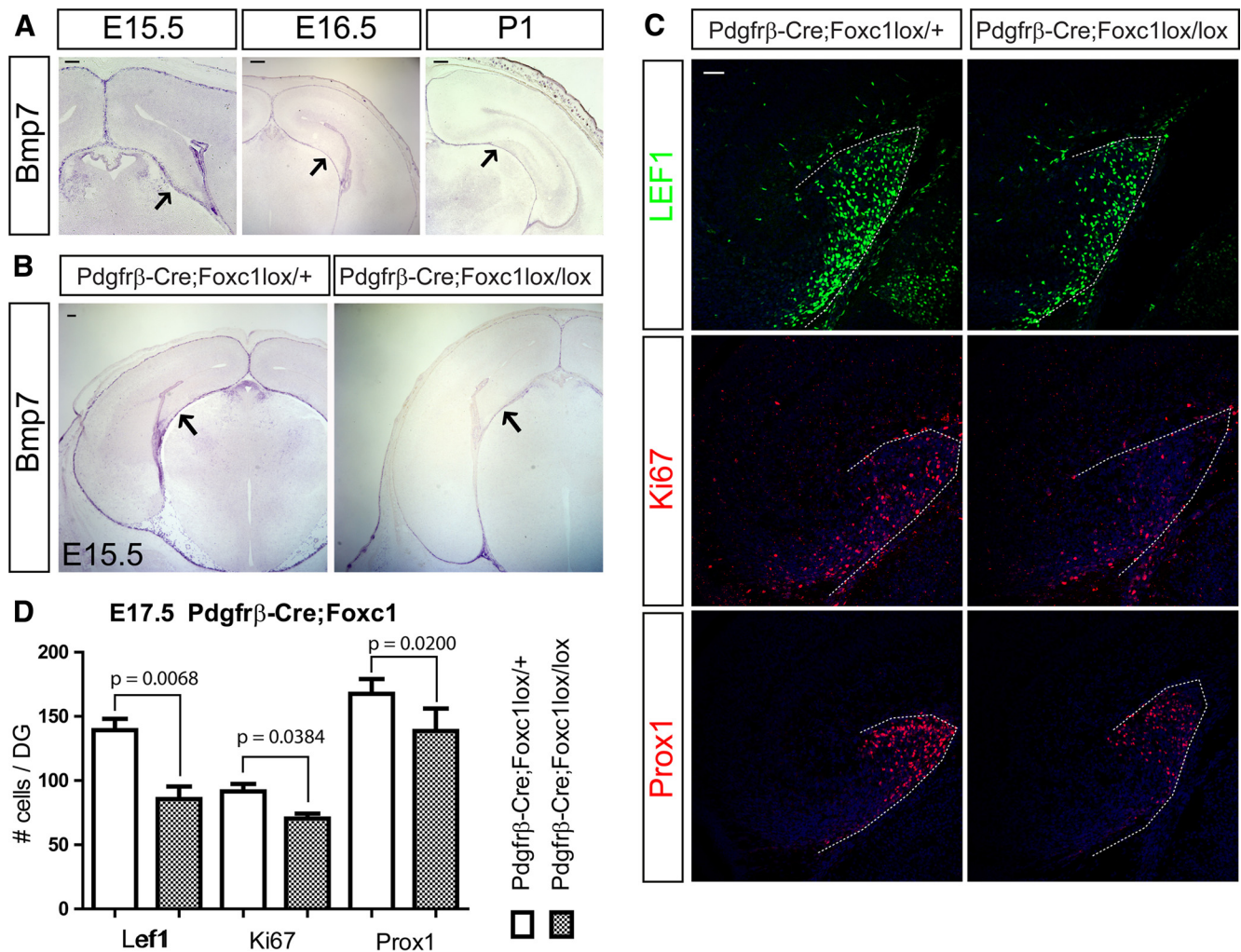
**Figure 2.** Bmp signaling in perinatal DG neural stem cells activates Lef1 expression. **A**, Expression of phospho-Smad1/5/8 in the DG at E14.5 and 18.5. Dashed lines outline the DG. **A'**, Expression of phospho-Smad1/5/8 in the Lef1+ DG stem cells and the hippocampal fissure (arrow) at E17.5. Dashed lines outline the DG. **B**, Expression of *Acvr1* by *in situ* hybridization in the DG at E14.5, E17.5, and P1. *Acvr1* is expressed in the DG migratory stream, the developing DG (arrow), and the CA3 region. **C**, Induction of Lef1 expression by activation of the Bmp signaling pathway through *Acvr1* expression was examined by *in utero* electroporation. CA and KD variants of *Acvr1* expression constructs were used to examine *Acvr1*-mediated Lef1 expression in the embryonic DG. GFP-electroporated DG was used as a control. Nog was used to show the effect of inhibiting the Bmp signaling pathway on Lef1 expression in the DG. The nonelectroporated contralateral DG was compared as a control group for Nog electroporation. Dashed lines outline the embryonic DG. **D, E**, The number of Lef1+ cells is presented from four independent experiments. Electroporation was conducted at E13.5 when the DG neural epithelium is exposed to the lateral ventricle and Lef1 expression was examined at E16.5 in the DG. Error bars indicate SEM. Scale bars, 100  $\mu$ m.

development of DG stem cells using a recently available commercial antibody that is useful for immunohistologic analysis. Lef1 expression was detected in the dorsal midline at E14.5 and the expression of Lef1 in later developmental stages depicted the migration pattern of the presumptive DG stem cells (Altman and Bayer, 1990a,b; Li and Pleasure, 2005) (Fig. 1A). Lef1 expression was also seen in non-neural meningeal tissues and endothelial cells but was not seen in Cajal–Retzius cells within the dentate (data not shown). At P1, Lef1 was expressed in the forming dentate in the transient proliferative zone including the hilus and DG blades where there is abundant production of granule neurons. Expression of Lef1 at 2 weeks after birth correlated with the establishment of the SGZ stem niche and is essentially limited to this zone (Fig. 1A). To examine whether the cells expressing Lef1 were proliferative, we traced cells in S-phase with a single BrdU injection 1 h before analysis at E16.5. Cells colabeled for Lef1 and BrdU were seen in the DG migratory stream at E16.5 (Fig. 1A'). At P1, Lef1+ cells were numerous and largely restricted to the dentate, some showing colabeling with Blbp (a marker of the glia-like dentate stem cells beginning to appear at this age) (Fig. 1B). Although many of the Lef1+ cells were colabeled with Blbp and BrdU, we did not find examples of colabeling with Tbr2 (transit-amplifying cells) or Prox1 (DG granule neurons) (Fig. 1B; data not shown). Previous studies showed that around birth the DG stem cells also become Shh-responsive (Ahn and Joyner,



**Figure 3.** Bmp7 regulates embryonic DG neurogenesis. **A**, Constructs as indicated were electroporated into the medial cortex at E13.5 and Prox1+ DG granule neurons were stained at E16.5. Dashed lines outline the embryonic DG. **B**, The neurogenic effect of the signaling molecules was compared by counting Prox1-stained cells in the E16.5 DG from four independent experiments. The *p* values from the Tukey's *post hoc* comparisons are presented after ANOVA. Error bars indicate SEM. Scale bars, 100  $\mu$ m.

2005). We used Gli1-CreERT2 mice for short-term lineage tracing to label DG stem cells under the influence of the Shh signaling pathway by using ROSA-YFP reporter mice and introducing TM (2 mg/pup) at P1 to Gli1-CreERT2 mice. This allowed us to identify Shh-responsive DG stem cells at P3 and we found that GFP+ cells were costained with Lef1 (Fig. 1B'). Similarly, adult Shh-responsive DG stem cells expressed Lef1 and were predominantly radially oriented SGZ-based cells (Fig. 1B').



**Figure 4.** Decreased DG neurogenesis in mice with reduced meningeal Bmp expression. **A**, Expression of Bmp7 at the level of the hippocampus was revealed by *in situ* hybridization using E15.5, E17.5 embryos and P1 pups. **B**, Pdgfrβ-Cre-mediated inhibition of meninges-specific Foxc1 expression (Pdgfrβ-Cre; Foxc1<sup>lox/lox</sup>) resulted in the decrease of Bmp7 expression in the meninges at E15.5 (arrows). **C**, Lef1, Ki67 (a marker for proliferating cells), and Prox1 immunostaining was examined in the E17.5 embryonic DG with DAPI counterstaining (blue). Dashed lines outline the embryonic DG. **D**, We counted the number of cells stained with Lef1, Ki67, and Prox1 in the DG. Four different litters were used for comparison. Error bars indicate SEM. Scale bars, 100 μm.

Thus, our data show that Lef1 is expressed in the cells in the dentate neuroepithelium and also in the dentate stem cells migrating to establish the ectopic proliferative zone where granule neurons are produced locally. Lef1 also continues to be expressed in radially oriented dentate progenitors through later postnatal ages as well (Choe and Pleasure, 2012). Thus, determining how Lef1 is expressed in such a restricted spatial domain in the developing medial pallium and dentate is of some interest since it should reveal important insights into how the dentate forms.

#### Acvr1, a Bmp receptor, signaling regulates Lef1 expression in the medial cortex

Previous studies showed that in other non-neural tissues Bmp signaling can induce Lef1 expression providing a potential mechanism whereby Bmp signaling can render receiving cells responsive to specific Wnt signaling (Kratochwil et al., 1996). Since Bmp ligands are expressed both by the developing meninges and the cortical hem (Choe et al., 2012; Segklia et al., 2012) it seemed quite possible that Bmps might play a role in inducing the expression of Lef1 specifically in the hem-adjacent neuroepithelium and then later in the dentate, which develops in close proximity to the meninges. To determine whether signaling is active in the

early dentate, we stained for phospho-Smad1/5/8, which allows visualization of cells where Bmp signaling is active. Staining for phospho-Smad1/5/8 was detected in dentate cells close to the meninges at E14.5 and in the hilar germinal area at E18.5 as well as in many cells in the hippocampal fissure (Fig. 2A). To examine whether phospho-Smad1/5/8 is found in Lef1+ DG stem cells, we doublestained for phospho-Smad1/5/8 and Lef1 at E17.5. Lef1+ DG stem cells with evidence of Bmp signaling were readily seen in the hilus (Fig. 2A'). Among type I Bmp receptors, Bmpr1a and Bmpr1b were examined in previous studies but Acvr1 expression was not included in previous studies of DG development (Caronia et al., 2010). Previous studies of double mutant mice for Bmpr1a and 1b showed that these mice have largely intact dentate development with fairly normal formation of the dentate SGZ, but it seemed clear that there should be some more potent effect of Bmp signaling in the dentate. Thus we wondered whether Acvr1 might be part of the missing picture of Bmp signaling in the dentate. Acvr1 is a serine/threonine transmembrane receptor related to Bmpr1a and 1b that phosphorylates Smad1 and 5 relatively specifically after activation by Bmp7 (Macias-Silva et al., 1998), thus we examined expression of Acvr1 and found that Acvr1 is expressed at high levels in the hippocampus with highest

levels of expression in the DG (arrow) and pyramidal regions of the developing hippocampus (Fig. 2B).

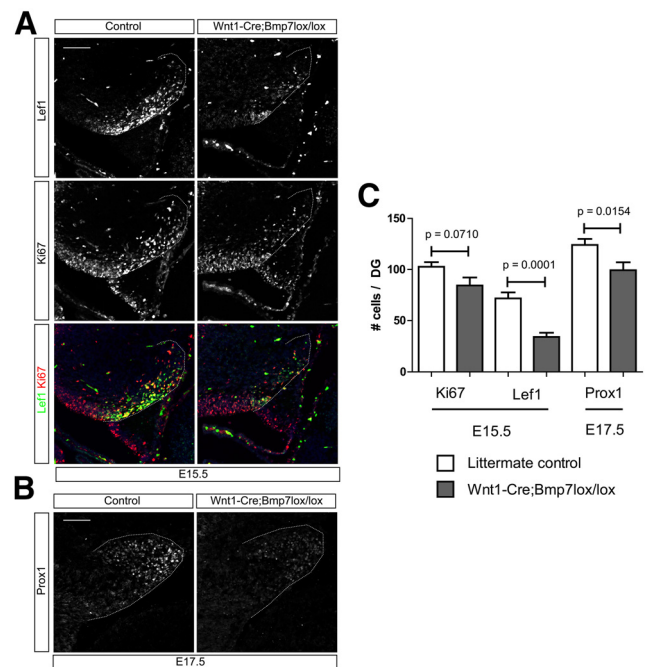
To examine the potential effects of *Acvr1* signaling in the developing dentate, we tested whether *Acvr1* directly regulates *Lef1* expression. We electroporated a constitutively active (CA) variant of *Acvr1* (Macias-Silva et al., 1998) into the developing DG and found that CA-*Acvr1* cell autonomously induced expression of *Lef1* in the DG ( $n = 4$ , Tukey's *post hoc* test,  $p = 0.001$ ), while a KD variant of *Acvr1* (Visser et al., 2001) blocked expression in GFP-expressing cells ( $n = 4$ , Tukey's *post hoc* test,  $p = 0.0100$ ). Importantly *Nog* dramatically reduced the expression of *Lef1* in the entire expressing DG, thus indicating that endogenous Bmp signaling regulates *Lef1* expression (Fig. 2C–E). These results indicate that activating Bmp signaling through *Acvr1* robustly induces *Lef1* expression in the embryonic DG and that Bmp signaling is an important regulator of *Lef1* expression. Thus, we believe that Bmp signaling in the dentate is a key regulator of dentate development by regulating expression of *Lef1*, a component of Wnt signaling required for dentate formation.

### Bmp signaling regulates dentate granule neurogenesis in the embryonic DG

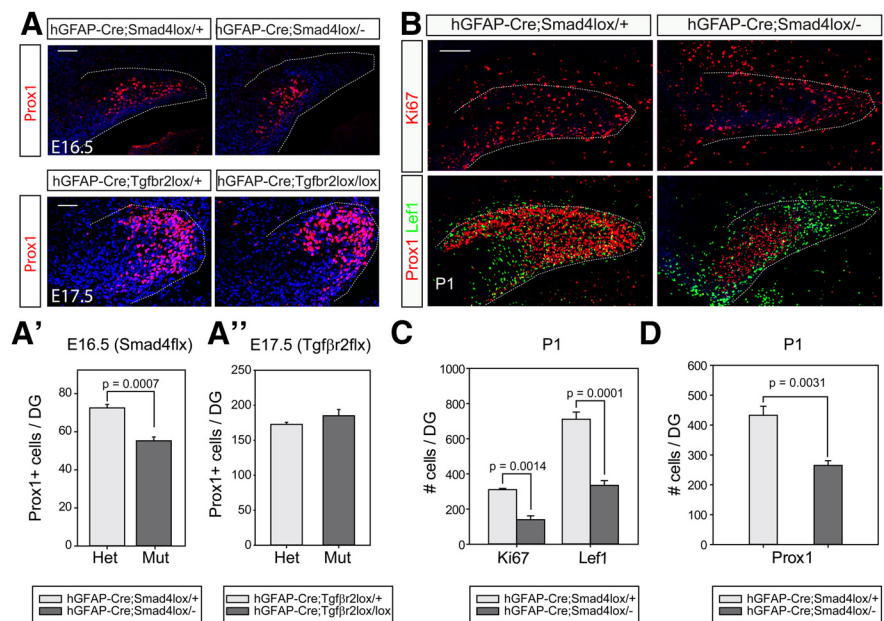
Activation of the Bmp receptor *Acvr1* regulates the expression of *Lef1* in the developing dentate, but does Bmp signaling also regulate the production of dentate granule neurons? To address this we electroporated Bmp7 into the medial cortical wall at E13.5 and examined expression of *Prox1*, a marker for postmitotic granule cells at E16.5. We compared these effects to the effects of *Nog*, to inhibit endogenous Bmp signaling. Interestingly we found that high-level overexpression of Bmp7 induced loss of the DG, perhaps through the death of the DG neural epithelium and agenesis of the choroid plexus (see the cell debris in the ventricle) similar to that seen previously with high dosage Bmps in cortical cultures (Mabie et al., 1999), so we also examined animals with the low-level expression of Bmp7 and compared the number of *Prox1*+ cells with the DG electroporated with GFP or *Nog*. As shown in Fig 3, A and B, lower level mis-expression of Bmp7 increased *Prox1*+ DG granule neurons and expression of *Nog* partially inhibited DG neurogenesis ( $n = 4$ ,  $p = 0.001$  (*Nog*),  $p = 0.020$  (*Bmp7*)). Thus, Bmp signaling does indeed regulate the production of *Prox1*+ dentate granule neurons along with the expression of the required Wnt signaling component *Lef1*.

### Bmp7-expressing meninges are important in regulating the late embryonic DG stem cell niche

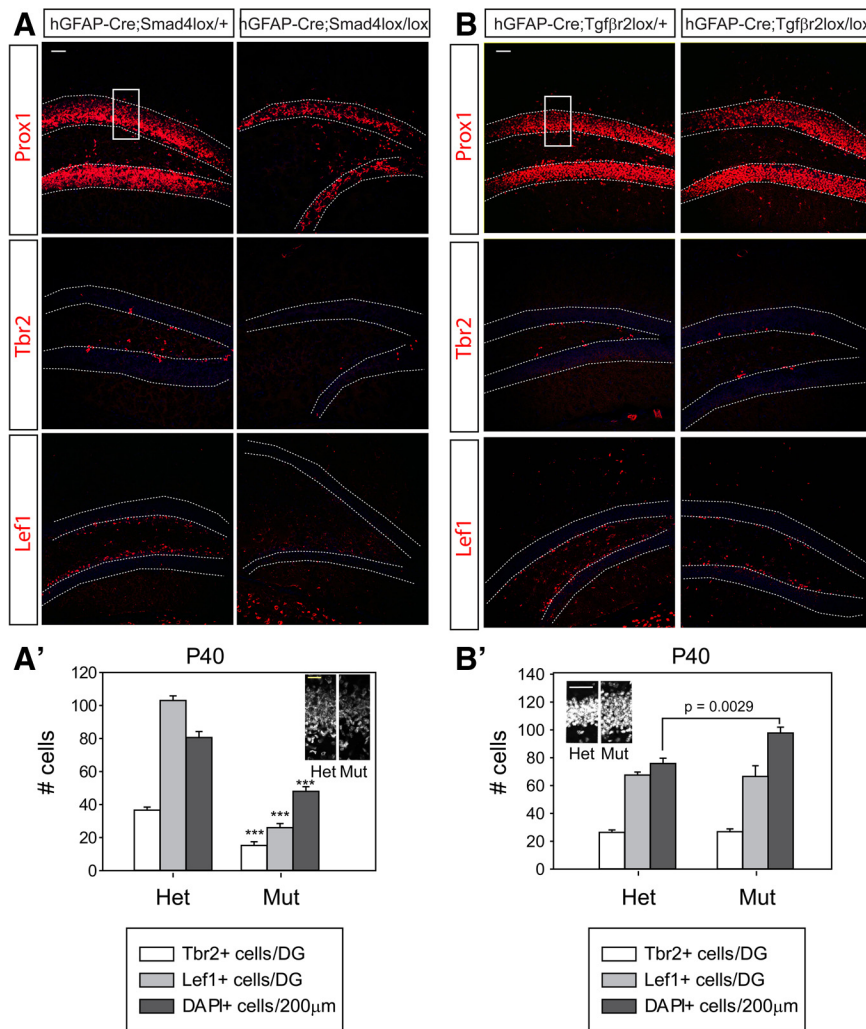
Wnt expression persists throughout embryonic life in the cortical hem and its derivative, the fimbria, and activation of Wnt signaling induces DG neurogenesis and expression of *Prox1* (Grove et al., 1998; Karalay et al., 2011). Wnts are expressed by the fimbria next to the DG, but what is the source of the Bmp ligands that might regulate DG development? We examined expression of Bmp ligands and



**Figure 5.** Loss of meningeal Bmp7 leads to inhibition of dentate neurogenesis. **A**, *Lef1* and *Ki67* immunostaining was conducted using the DG of *Wnt1-Cre; Bmp7lox/lox* mutants and littermate controls at E15.5. Dashed lines depict the DG. **B**, E17.5 embryos were used to stain dentate granule neurons using *Prox1*. Dashed lines depict the DG. **C**, Quantification of immunostained cells reveals significant decreases of *Lef1*+ dentate stem cells and *Prox1*+ dentate granule neurons at E15.5 and E17.5, respectively ( $n = 3$ ). Error bars indicate SEM. Scale bars, 100  $\mu\text{m}$ .



**Figure 6.** Failure of DG neurogenesis by cell-autonomous inhibition of Bmp signaling in DG neural stem cells. **A**, Embryonic DG neurogenesis is decreased by loss of *Smad4* expression in hGFAP-Cre lineage cells. The DG of embryos at E16.5 (*Smad4*) and E17.5 (*Tgfr2*) was stained with *Prox1*. **A'**, The number of *Prox1*+ cells in the DG of hGFAP-Cre; *Smad4*<sup>lox/+</sup> (Het) and hGFAP-Cre; *Smad4*<sup>lox/-</sup> (Mut) embryos is presented ( $n = 4$ ,  $p = 0.0007$ ). **A''**, The number of *Prox1*+ cells in the DG of hGFAP-Cre; *Tgfr2*<sup>lox/+</sup> (Het) and hGFAP-Cre; *Tgfr2*<sup>lox/lox</sup> (Mut) embryos is presented ( $n = 3$ ). **B**, hGFAP-Cre-mediated inhibition of *Smad4* expression at P1 shows decreased neurogenesis along with decreased expression of *Lef1* and *Ki67*. Dashed lines outline the DG. **C, D**, The number of *Lef1*, *Ki67* (**C**)-positive, and *Prox1* (**D**)-positive cells was counted from P1 pups of four different litters ( $n = 4$ ). Error bars indicate SEM. Scale bars, 100  $\mu\text{m}$ .



**Figure 7.** Compromised postnatal neurogenesis by developmental inhibition of Bmp signaling. **A**, P40 mice showed malformed SGZ neural stem cell niches and granule cell layers when the Bmp signaling pathway is inhibited in the DG stem cells from development onward. Effects of Smad4 inhibition in the hGFAP-Cre-expressing DG neural stem cells persisted to P40 as revealed with staining for Prox1, Tbr2, and Lef1. **A'**, The number of granule neurons (DAPI+ cells excluding the SGZ) as well as transit-amplifying cells (Tbr2+ cells) were counted in control (Het, hGFAP-Smad4<sup>lox/+</sup>) and mutant animals (Mut, hGFAP-Smad4<sup>lox/lox</sup>). Inset, DAPI-stained image of the boxed area of **A** ( $n = 3$ ,  $***p = 0.0001$ ). Number of granule cells are presented as number of cells in a 200  $\mu\text{m}$  box. **B**, Inhibition of Tgfb2 signaling pathway led to the slightly enlarged DG at P40 as shown with staining for Prox1, Tbr2, and Lef1. **B'**, The number of Tbr2+ cells, Lef1+ cells and granule cells (DAPI+) was counted from control (Het, hGFAP-Cre; Tgfb2<sup>lox/+</sup>) and mutant animals (Mut, hGFAP-Cre; Tgfb2<sup>lox/lox</sup>). Inset, DAPI-stained images of the boxed area of **B** ( $n = 3$ ,  $p = 0.0029$ ). Dashed lines outline the DG granule cell layer. Error bars indicate SEM. Scale bars: **A, B**, 100  $\mu\text{m}$ ; **A', B'**, 50  $\mu\text{m}$ .

found that meningeal tissue is a major source of Bmp expression at the time of late embryonic DG neurogenesis. Considering that the migration route of DG progenitors is dependent on meningeally produced Cxcl12 and these cells spend time in a subpial transient stem cell niche (Li et al., 2009), it is reasonable to hypothesize that Bmps from the meninges regulate dentate progenitor development and might drive Lef1 expression in DG stem cells. Staining of Bmp7 at the level of the DG from E15.5 to P1 showed strong expression in the meninges (Fig. 4A, arrows). We previously found that major defects in meningeal development from early in development result in major defects in the medial forebrain, making it hard to test dentate development specifically (Zarbali et al., 2007). To assess later conditional loss of Bmps, we used the *Pdgfr $\beta$ -Cre* to conditionally knock out *Foxc1*, a meningeal transcription factor, to decrease the expression of Bmp7 in the meninges covering the DG (Zarbali et al., 2007; Siegenthaler

et al., 2009; Choe et al., 2012). *Pdgfr $\beta$ -Cre; Foxc1<sup>lox/lox</sup>* mutant mice showed decreased expression of Bmp7 in the meninges surrounding the DG (Fig. 4B, arrows); this persisted until E18.5 when the mutant embryos died (data not shown). We counted the number of Lef1, Ki67 (cells actively in the cell cycle), and Prox1+ DG granule neurons in the DG at E17.5. *Pdgfr $\beta$ -Cre; Foxc1<sup>lox/lox</sup>* mutant embryos showed significant loss of Lef1, Ki67, and Prox1 expression ( $n = 4$ ,  $p = 0.0068$  (Lef1),  $p = 0.0384$  (Ki67),  $p = 0.2000$  (Prox1)) (Fig. 4C,D), supporting the idea that the meninges provide factors critical for the development of the embryonic DG and production of early DG neurons. These data are consistent with an important role for meningeal Bmps in this process. We wished to exclude a role for Cxcl12 in this phenotype since this chemokine is expressed by the meninges and also has some role in DG precursor behavior, so we examined the expression of Cxcl12 in *Pdgfr $\beta$ -Cre; Foxc1<sup>lox/lox</sup>* mutant embryos and found no significant change in expression (data not shown).

#### Neural crest-derived meningeal Bmp7 mediates embryonic DG neurogenesis

Our data show that meningeal morphogens regulate dentate neurogenesis by regulating Lef1 expression in DG stem cells. To be certain that it is Bmp7 that is one of the main regulators of dentate Lef1 expression from the meninges, we used *Wnt1-Cre* to delete Bmp7 in neural crest-derived meningeal cells (Danielian et al., 1998; Zouvelou et al., 2009). At E15.5, Lef1+ cells are significantly ( $p = 0.0001$ ) reduced in the mutant (*Wnt1-Cre; Bmp7<sup>lox/lox</sup>*) and resulted in decreased production of DG granule neurons at E17.5 (Fig. 5A–C). This result supports our idea that meningeally produced Bmp7 is a component of the DG niche signal that controlling Lef1 expression in DG stem cells during development.

#### Inhibition of embryonic Bmp signaling in DG stem cells impairs DG stem cell niche integrity

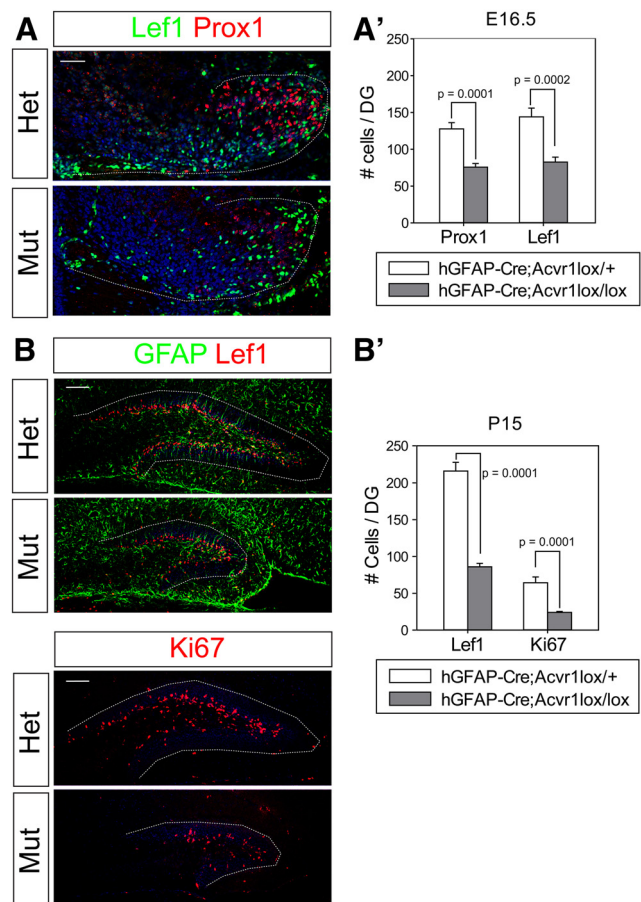
*Bmpr1a* is known to regulate adult neural stem cell survival and we provided evidence that *Acvr1* is involved in DG neurogenesis by regulating Lef1 expression in DG stem cells. Previous studies have also demonstrated a clear but modest role for *Bmpr1a/1b* signaling in the developing dentate (Caronia et al., 2010). Given our new data on *Acvr1*, we wanted to further address the role of the Bmp signaling pathway on DG stem cell niche formation by using hGFAP-Cre mice to target the developing dentate to ablate the expression of Smad4, a common transcription factor for both Bmp and Tgfb signaling. Tgfb2 is the only Tgfb receptor that binds all Tgfb isoforms and is required for activating the Smad4-

dependent canonical Tgf $\beta$  signaling pathway (Wrana et al., 1992; Derynck and Zhang, 2003). To rule out the involvement of Tgf $\beta$  signaling pathway in dentate development, we compared Smad4 mutants to Tgf $\beta$ 2 conditional mice using the same Cre driver to selectively block the canonical Tgf $\beta$  signaling pathway. Analysis of the hGFAP-Cre; Smad4<sup>lox/-</sup> mutants at late embryonic development showed that by this stage, when the dentate is just becoming established, that there was a modest but significant loss of Prox1+ granule neurons ( $n = 4$ ,  $p = 0.0007$ ) while the hGFAP-Cre; Tgf $\beta$ 2<sup>lox/lox</sup> mutants had no apparent phenotype (Fig. 6A,A',A''). Several days later, at P1, proliferating Ki67+ cells as well as Lef1+ cells were dramatically reduced in hGFAP-Cre; Smad4<sup>lox/-</sup> mutants ( $n = 4$ ,  $p = 0.0014$  (Ki67),  $p = 0.0001$  (Lef1)) (Fig. 6B,C). These reduced numbers of progenitor cells led to reduced neurogenesis as measured by staining for Prox1 at P1 ( $n = 4$ ,  $p = 0.0031$ ) (Fig. 6B,D).

We examined animals at older ages to see the end results of the developmental defects in Bmp signaling using markers such as Lef1, Prox1, and Tbr2, a marker of the transit-amplifying cells in the SGZ (Hodge et al., 2008). There were dramatic defects in the number of granule neurons (Fig. 7A). There were also major losses of transit-amplifying cells (Tbr2+) and dentate stem cells (Lef1+) ( $n = 3$ ,  $p = 0.0001$ ) (Fig. 7A,A'). In contrast, the Tgf $\beta$ 2 conditional mutants showed slight thickening of the DG granule layer without significant difference of Tbr2 or Lef1+ cells as was expected from the previously reported inhibitory function of Tgf $\beta$ 1 on adult neurogenesis ( $n = 3$ ,  $p = 0.0029$  (DAPI)) (Fig. 7B,B') (Buckwalter et al., 2006). These results suggest that late embryonic loss of Smad4 has major consequences for dentate development and the establishment of the dentate stem cell niche due in part to the loss of Bmp signaling during dentate formation.

### Acvr1 mediates DG neurogenesis and SGZ stem cell niche formation

The dramatic effect of overall blockade of the Bmp signaling pathway on DG dentate development and ongoing neurogenesis is quite distinct from the published effects of loss of function of Bmpr1a/b (Caronia et al., 2010). This led us to wonder whether a key part of the signaling equation may be Acvr1. We wanted to examine whether loss of Acvr1 would have dramatic effects on DG progenitors and resulting DG neurogenesis as well as the maintenance of Lef1+ DG stem cells *in vivo*. We crossed hGFAP-Cre with Acvr1 conditional mice and examined the Lef1 and Prox1 expression at E16.5. As shown in Figure 8, A and A', loss of Acvr1 in the hGFAP-Cre-expressing DG progenitors reduced both Lef1 and Prox1 expression [ $n = 4$ ,  $p = 0.0001$  (Prox1),  $p = 0.0002$  (Lef1)]; this strongly supports our *in utero* electroporation and conditional loss of the meningeal Bmp expression. We also examined whether loss of Acvr1 at embryonic stages leads to disruption of dentate development and effects on the adult DG stem cell niche by examining Ki67, Lef1, Tbr2, and Prox1 at P15, P30, and P40. The numbers of cells expressing markers of proliferating cells (Lef1 and Ki67) were reduced in the mutant at P15 ( $n = 3$ ,  $p = 0.0001$ ) (Fig. 8B,B'). At both P30 and P40, we also observed decreases in Lef1 expression in the hGFAP-Cre; Acvr1<sup>lox/lox</sup> mutants ( $n = 4$ ,  $p = 0.0001$ ). Tbr2 expressing transit amplifying cells were also reduced at P30 ( $n = 4$ ,  $p = 0.0053$ ) (Fig. 9A,A'). The number of Ki67+ cells was low in the P30 hGFAP-Cre; Acvr1<sup>lox/lox</sup> mutant ( $n = 4$ ,  $p = 0.0515$ ); however, similar at P40 ( $n = 4$ ,  $p = 0.8331$ ), which might be due to increased gliogenesis associated with gliosis (Fig. 9A,A',B,B'). To support this idea, we examined the expression of the dentate granule marker Prox1 and found a dramatic loss of Prox1+ cells in the hGFAP-Cre;



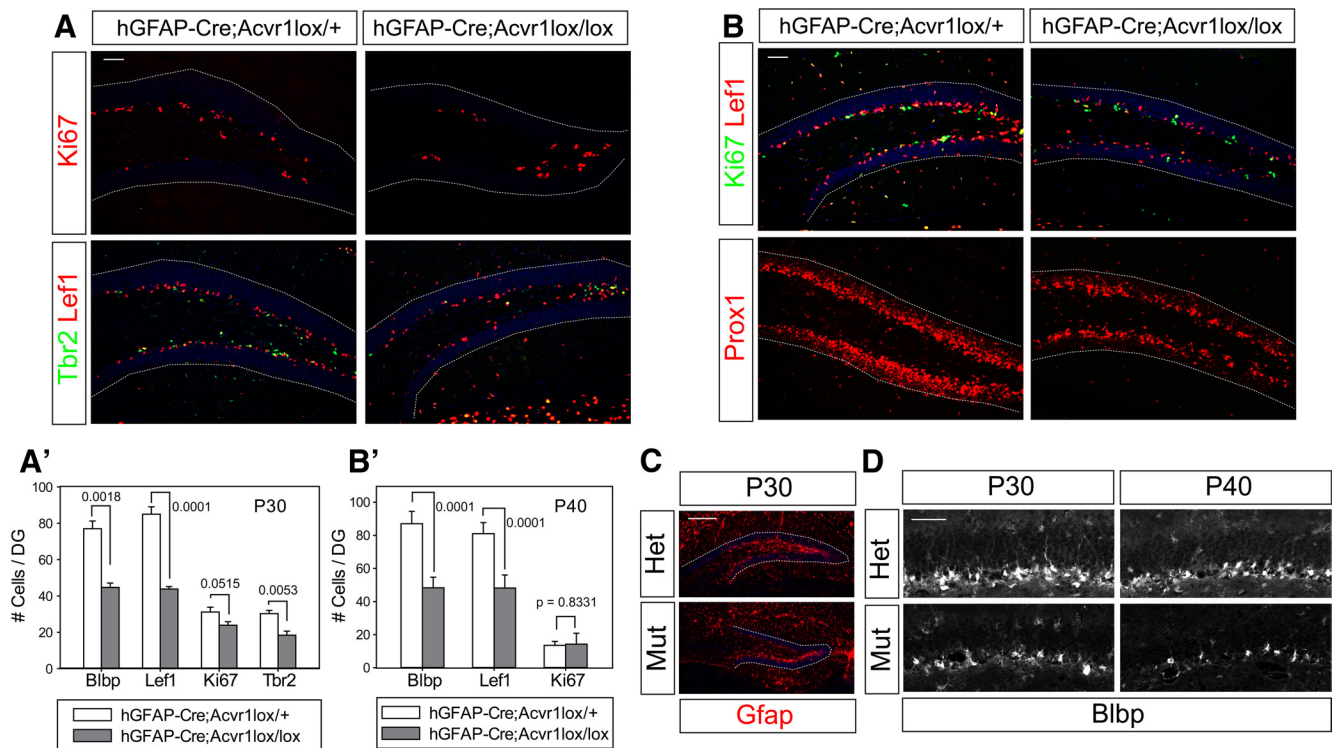
**Figure 8.** Loss of Acvr1 from DG neural stem cells leads to defects in dentate development. **A**, Lef1 and Prox1 immunostaining of the E16.5 embryonic DG from hGFAP-Cre; Acvr1<sup>lox/+</sup> (Het) and hGFAP-Cre; Acvr1<sup>lox/lox</sup> (Mut). Dashed lines outline the DG. **A'**, The number of Lef1 or Prox1+ cells was counted from four control and mutant embryos obtained from three litters ( $n = 4$ ). **B**, Lef1, GFAP, and Ki67 immunostaining of the DG obtained from hGFAP-Cre; Acvr1<sup>lox/+</sup> and hGFAP-Cre; Acvr1<sup>lox/lox</sup> mice at P15. Dashed lines outline the DG. **B'**, Quantification of the immunostained cells from **B** ( $n = 3$ ). Error bars indicate SEM. Scale bars: 100  $\mu$ m.

Acvr1<sup>lox/lox</sup> mutant at P40 (Fig. 9B). As expected by the decreased number of DG progenitors, the size of the DG was also reduced in the mutants (Figs. 8B,9C). Since the Lef1-expressing cells in the adult DG are mostly radial glial cells in the adult DG (Fig. 1B'), we also examined the number of Blbp- and GFAP-expressing radial glial DG neural stem cells. At both P30 and P40, the number of radial glial neural stem cells (we counted Blbp+ cells as these are more easily identified) was significantly reduced in the mutant SGZ and GFAP+ scaffolding cells were somewhat disorganized in hGFAP-Cre; Acvr1<sup>lox/lox</sup> mutant ( $n = 4$ ,  $p = 0.0018$  (P30),  $p = 0.0001$  (P40) for Blbp) (Fig. 9A',B',C,D).

### Discussion

In this study we show that Bmp signaling strongly influences the development of the DG by regulating the formation of the dentate stem cell niche. Specifically we provide evidence that Bmp ligands from the meninges act through Acvr1 to regulate the expression of Lef1 in the developing DG. These new findings provide an important framework for understanding the events that control the ability of the dentate to maintain long-term neurogenic capacity. In addition, our identification of a major role for the previously little studied Acvr1 receptor, a member of the Bmpr1 family, and identification of the meninges as a source of





**Figure 9.** Loss of *Acvr1* leads to persistent defects in neurogenesis in the dentate stem cell niche. **A**, Ki67, Tbr2, and Lef1 immunostaining was conducted using the DG of hGFAP-Cre; *Acvr1*<sup>lox/+</sup> (Het) and hGFAP-Cre; *Acvr1*<sup>lox/lox</sup> (Mut) at P30. Dashed lines depict the DG. **A'**, Quantification of immunostained cells using four control and mutant animals from three litters ( $n = 4$ ). **B**, P40 animals were used to stain Ki67, Lef1, and Prox1. **B'**, Quantification of Blbp, Lef1, and Ki67+ cells at P40 using four control and mutant animals from three litters ( $n = 4$ ). **C, D**, Immunostaining of GFAP using P30 (**C**) and staining of Blbp using P30 and P40 (**D**) animals ( $n = 4$ ). Error bars indicate SEM. Scale bars: 100  $\mu\text{m}$ .

Bmp7 are important developments revealing for the first time the central role of Bmp signaling in development of the dentate stem cell niche.

### Bmp signaling controls Wnt responsiveness in the developing dentate

A number of studies over the last decade have made it very clear that Wnt signaling is particularly important in the regulation of dentate granule cell production during development and in postnatal life (Galceran et al., 2000; Zhou et al., 2004; Li and Pleasure, 2005; Lie et al., 2005). The function of Lef1, a Wnt-responsive transcription factor, is absolutely required for granule cell production (Galceran et al., 2000). Recent studies have also shown that Prox1, a transcription factor required for dentate granule cell fate, is directly regulated by Wnt signaling (Machon et al., 2007; Karalay et al., 2011). In this study we show that Lef1 is quite selectively expressed in the dentate progenitor lineage at a variety of stages. Since Lef1 plays a truly central role during dentate neurogenesis, factors that regulate the selective expression of Lef1 in the dentate lineage should be important regulators of dentate development and neurogenesis. One of the key regulators of Lef1 expression is Wnt signaling itself (Hovanes et al., 2001; Filali et al., 2002); this makes sense since the dentate forms in proximity to the cortical hem, the strongest source of Wnt ligands in the developing cortex (Grove et al., 1998).

A number of studies indicate that Wnt signaling alone is insufficient to generate the appropriate milieu for dentate development. A mixture of ligands from the cortical hem is apparently necessary and sufficient to induce ectopic dentate gyri in the cortex (Mangale et al., 2008). Also, previous studies showed that Bmps produced by the cortical hem are required at early times to

allow formation of the hippocampus as a whole (Monuki et al., 2001). This work suggested to us a potential role for Bmp ligands as an additional regulator of dentate development at later stages, when the dentate is established. Since Bmp ligands can induce Lef1 expression in non-neural tissues we formed the hypothesis that Bmps from the hem and later from the meninges regulate the expression of Lef1. Our results showing that activating and blocking the Bmp signaling pathway selectively in the dentate regulates the number of Lef1-expressing cells are clearly consistent with this hypothesis. We believe this provides an important new advance in our understanding of the selective mechanisms governing dentate development and neurogenesis.

### The distinct role of *Acvr1* in dentate development

The developmental ablation of type I Bmp receptors, *Bmpr1a* and *1b*, results in defective DG formation partially through the inhibition of Wnt signaling molecules from the cortical hem and not solely through cell-autonomous inhibition of Bmp signal transduction in the DG (Caronia et al., 2010). The specification of the choroid plexus, the most dorsomedial derivative, is also dependent on the *Bmpr1a* signaling pathway (Hébert et al., 2002), which imposes difficulties in interpreting results from inhibiting Bmp receptor signaling in a nontissue-specific manner. *Acvr1*, however, is not expressed in the cortical hem, unlike *Bmpr1a* (Caronia et al., 2010). This implied to us that this additional member of the *Bmpr1* family potentially has a specific role inducing Lef1 expression. This idea is amply supported by our results showing cell-autonomous regulation of Lef1 expression by *Acvr1* in electroporation experiments and in conditional mutant mice. It is possible that there is some compensation by other *Bmpr1* family members that tends to

limit the severity of the *Acvr1* phenotype, however, it is abundantly clear that this receptor has a key role in dentate development and probably ongoing neurogenesis.

### Roles of the meninges in dentate development

What is the source of Bmp ligands regulating dentate development? Almost certainly at early stages Bmps are made by the hem and are also available from the CSF (made by the choroid plexus) and are important for patterning the medial cortical wall and future hippocampus (Furuta et al., 1997; Segklia et al., 2012). It is likely that these sources are also responsible in part, along with Wnt signaling, for the distribution of *Lef1* in the medial wall before the dentate is formed. Once dentate development commences, *Lef1*-expressing cells are seen relocating to the developing dentate field, away from the ventricular lumen, where they expand dramatically and then ultimately settle in the SGZ to become dentate stem cells. We showed quite clearly that ectopic *Bmp7* and Bmp inhibitors expressed in the forming dentate regulate the number of *Lef1* + cells as well as the production of *Prox1* + granule neurons. This suggests that a local source of *Bmp7* might be important in the developing dentate, and indeed, we found that mice with mutations causing loss of meningeal *Bmp7* expression have reduced numbers of *Lef1* + cells and impaired dentate neurogenesis. Thus, *Bmp7* produced by the meninges is likely to be a key regulator of dentate development, and given the continued postnatal phenotype seen in the *Acvr1* and *Smad4* mutant mice, it is possible that production of *Bmp7* by the meninges has a continuing role in postnatal neurogenesis. This is quite consistent with previous studies showing roles of Bmp signaling in adult neurogenesis, although these studies never considered the relevant physiologic sources of Bmp ligands in the adult dentate (Bonaguidi et al., 2008; Caronia et al., 2010; Mira et al., 2010).

In summary, we provide clear evidence that Bmp signaling, via the lesser-studied *Acvr1* receptor, is a key player in the development of the DG and the establishment of the dentate stem cell niche. Our results show that this is likely to be related to the control of Wnt responsiveness in the dentate niche as controlled by the expression of the key Wnt responsive transcription factor *Lef1*. This work helps to establish the development of the dentate as a process critically regulated by the intersection of a number of fundamental developmental signaling pathways.

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