

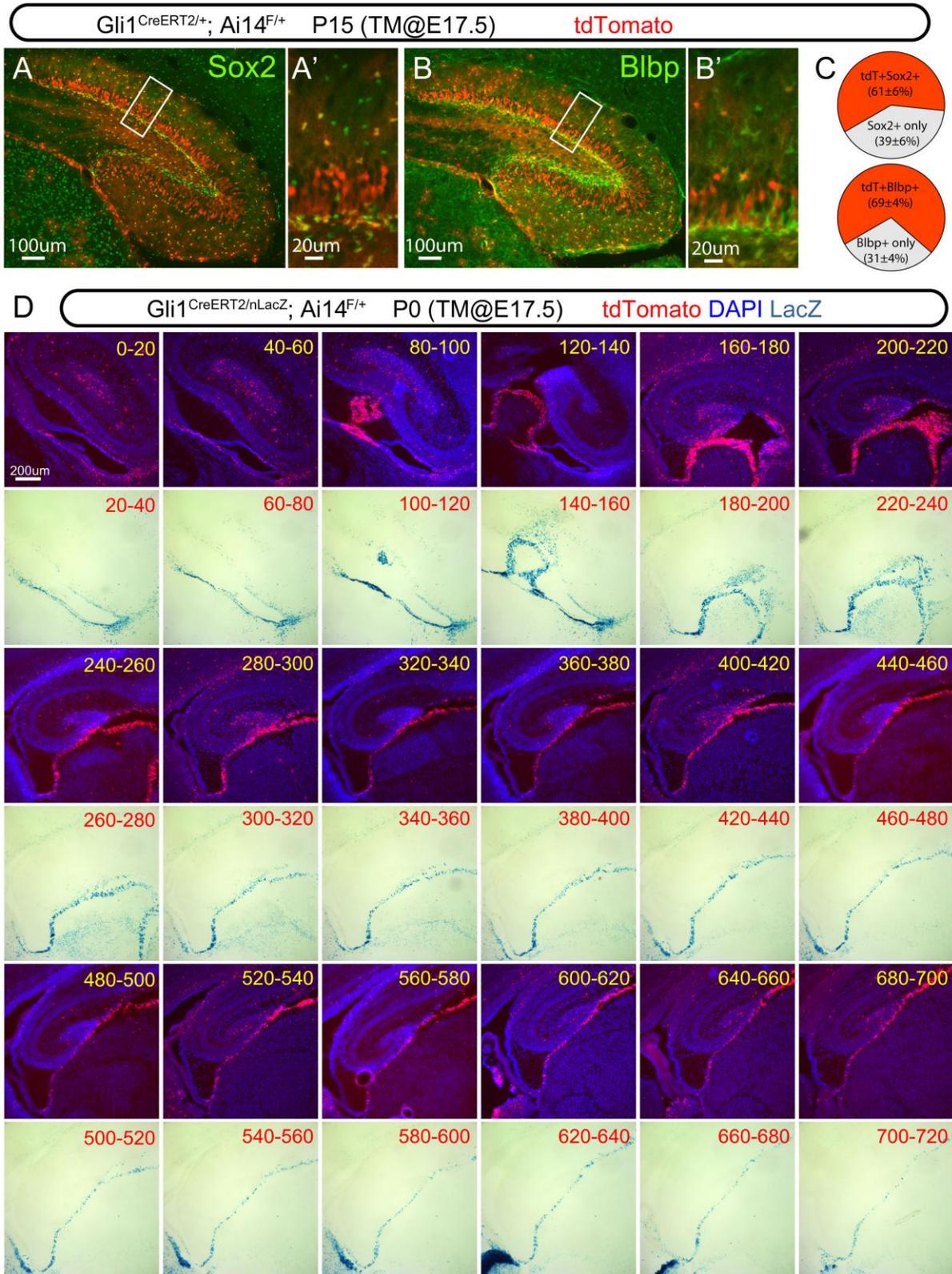
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**Supplemental Information**

**The Ventral Hippocampus Is the Embryonic Origin  
for Adult Neural Stem Cells in the Dentate Gyrus**

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Figure S1 (related to Figure 2)

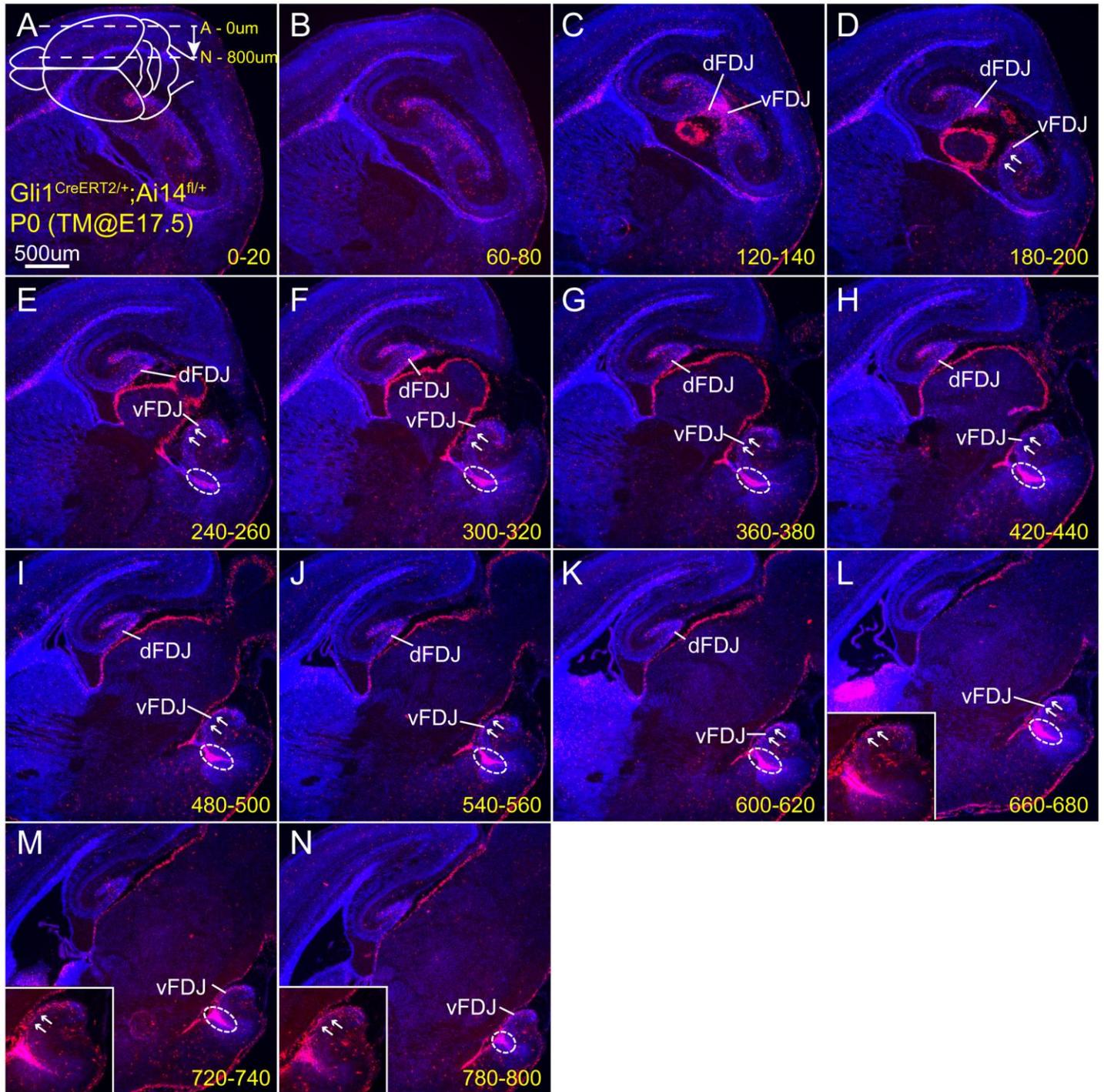


**Figure S1 (related to Figure 2):**

**(A-C)** The labeling efficiency of Gli1-CreERT2 in the SGZ was assessed by P15 with tamoxifen injection (3mg/40g animal) at E17.5 after crossing to the cre reporter Ai14. tdT+ cells in the SGZ were shown to express the radial glia markers Sox2 (**A**) and Blbp (**B**). Boxed areas in (**A**) and (**B**) were shown at the higher magnification in (**A'**) and (**B'**), respectively. 61±6% of the total Sox2+ cells in the SGZ along the septotemporal axis were tdT+ (upper panel in **C**), and 69±4% of the Blbp+ cells with radial processes in the SGZ were also tdT+ (lower panel in **C**).

**(D)** Alternate sections were used for analyzing the spatial distribution of Gli1-nLacZ+ and tdT+ cells respectively for the Gli1<sup>CreERT2/nLacZ</sup>;Ai14<sup>F/+</sup> mice. The tdT+ cells at P0 represented the descendants of Hh-responding cells labeled by tamoxifen induction at E17.5. The number in each panel represents the thickness distance from the first panel. tdT+ cells clearly occupied the septal levels (480µm away from the first panel) where the Gli1-nLacZ expression was no longer detectable.

Figure S2 (related to Figure 3)

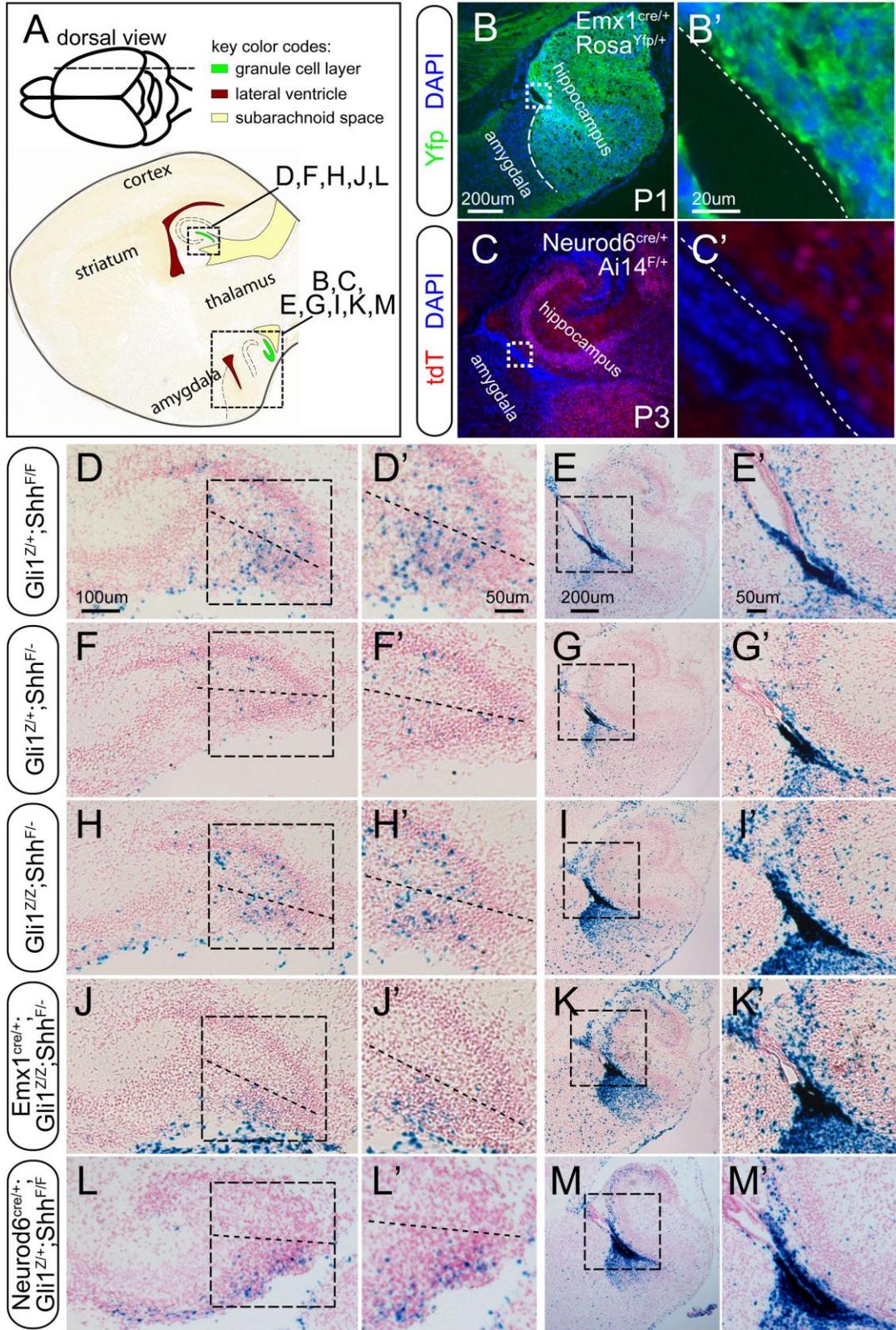


**Figure S2 (related to Figure 3)**

**(A-N)** The distribution of tdT+ cells in the  $Gli1^{CreERT2/+};Ai14^{F/+}$  animals were shown every 60 $\mu$ m by birth after tamoxifen (TM) induction at E17.5. The thickness distance as micron from the first section was shown at the bottom-right corner. Throughout the whole hippocampus **(A-N)**, the VZ of the temporal hippocampus was the most heavily labeled by tdT (white ovals in **E-N**). The tdT+ cell stream was noticeable in the vFDJ region (arrows in **D-N**). The tdT+ cells were continuous from the vFDJ into the dFDJ at the transitional level **(C)** and tapered off in the dorsal hippocampus toward the septal pole.

dFDJ, dorsal fimbriodentate junction; vFDJ, ventral fimbriodentate junction.

Figure S3 (related to Figure 4)



**Figure S3 (related to Figure 4)**

**(A)** The schemas were illustrated for the sagittal sections from the P1 animals shown in **B-M**.

**(B)** In the  $Emx1^{Cre/+};Rosa^{Yfp/+}$  animal at P1,  $Emx1$ -cre showed the recombination activity in the ventral hippocampus as the expression of Yfp (**B**), which also covered its VZ (along the dotted line in **B'**). Boxed area in **B** was shown at the higher magnification in **B'**.

**(C)** In the  $Neurod6^{Cre/+};Ai14^{F/+}$  animal at P3,  $Neurod6$ -cre showed the recombination activity in the ventral hippocampus as the expression of tdTomato (**C**), which didn't cover its VZ (along the dotted line in **C'**). Boxed area in **C** was shown at the higher magnification in **C'**.

**(D-E)** In the  $Gli1^{Z/+};Shh^{F/F}$  animals,  $Gli1$ -nLacZ+ cells were localized at the entrance of the hilus in the dorsal DG (**D** and **D'**). The dorsal dentate can be divided into upper and lower portions by a line connecting the tip of the CA3 field and the apex of the dentate pole. Some of  $Gli1$ -nLacZ+ cells could be detectable in the upper portion (**D'**).  $Gli1$ -nLacZ+ cells were clearly identifiable to form a stream out of the VZ of ventral hippocampus into the forming DG (**E** and **E'**). Boxed areas in **D** and **E** were shown at the higher magnification in **D'** and **E'**.

**(F-G)** In the  $Gli1^{Z/+};Shh^{F/-}$  animals, one copy of the  $Shh$  flox allele was replaced with a  $Shh$  null allele. The number of  $Gli1$ -nLacZ+ cells in the dorsal dentate was dramatically reduced in both portions (**F** and **F'**). The stream of  $Gli1$ -nLacZ+ cells leaving the VZ of ventral hippocampus was slightly diminished and there was a decrease in the expression of nLacZ in the VZ (**G** and **G'**). Boxed areas in **F** and **G** were shown at the higher magnification in **F'** and **G'**.

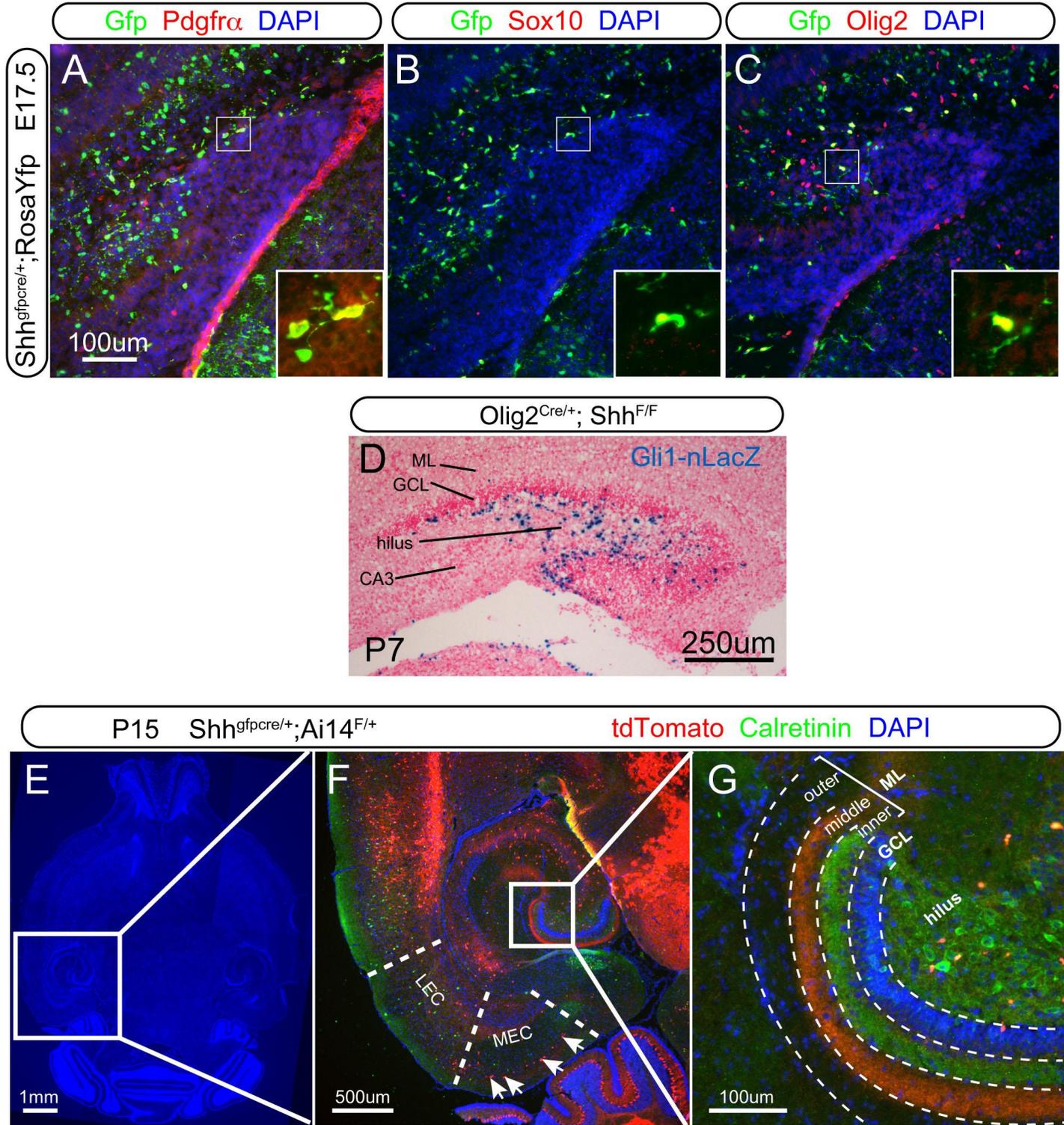
**(H-I)** In the  $Gli1^{ZZ};Shh^{F/-}$  animals, an extra copy of  $Gli1$ -nLacZ was present in relation to the animals in **F-G**. The distribution of  $Gli1$ -nLacZ+ cells was mostly restored in both dorsal DG (**H** and **H'**) and ventral hippocampus (**I** and **I'**) close to the level seen in the  $Gli1^{Z/+};Shh^{F/F}$  animals. Boxed areas in **H** and **I** were shown at the higher magnification in **H'** and **I'**.

**(J-K)** In the  $Emx1^{cre/+};Gli1^{ZZ};Shh^{F/-}$  animals, the remaining  $Shh$  flox allele was floxed out in the  $Emx1$  domain. Most of the  $Gli1$ -nLacZ+ cells were abolished in the upper portion of the dorsal dentate whereas residual  $Gli1$ -nLacZ+ cells were still present at the entrance of the hilus in the lower portion (**j** and **j'**), which appeared to be

continuous with the Gli1-nLacZ+ cell stream from the VZ of the ventral hippocampus (**K** and **K'**). Boxed areas in **J** and **K** were shown at the higher magnification in **J'** and **K'**.

(**L-M**) In the Neurod6<sup>cre/+</sup>;Gli1<sup>Z/+</sup>;Shh<sup>F/F</sup> animals, two copies of the Shh flox alleles were removed from the pallial neuronal lineage. The distribution of Gli1-nLacZ+ cells (**L**, **L'**, **M** and **M'**) was quite similar to the Emx1<sup>cre/+</sup>;Gli1<sup>Z/Z</sup>;Shh<sup>F/-</sup> animals. Boxed areas in **L** and **M** were shown at the higher magnification in **L'** and **M'**.

Figure S4 (related to Figure 6)



**Figure S4 (related to Figure 6)**

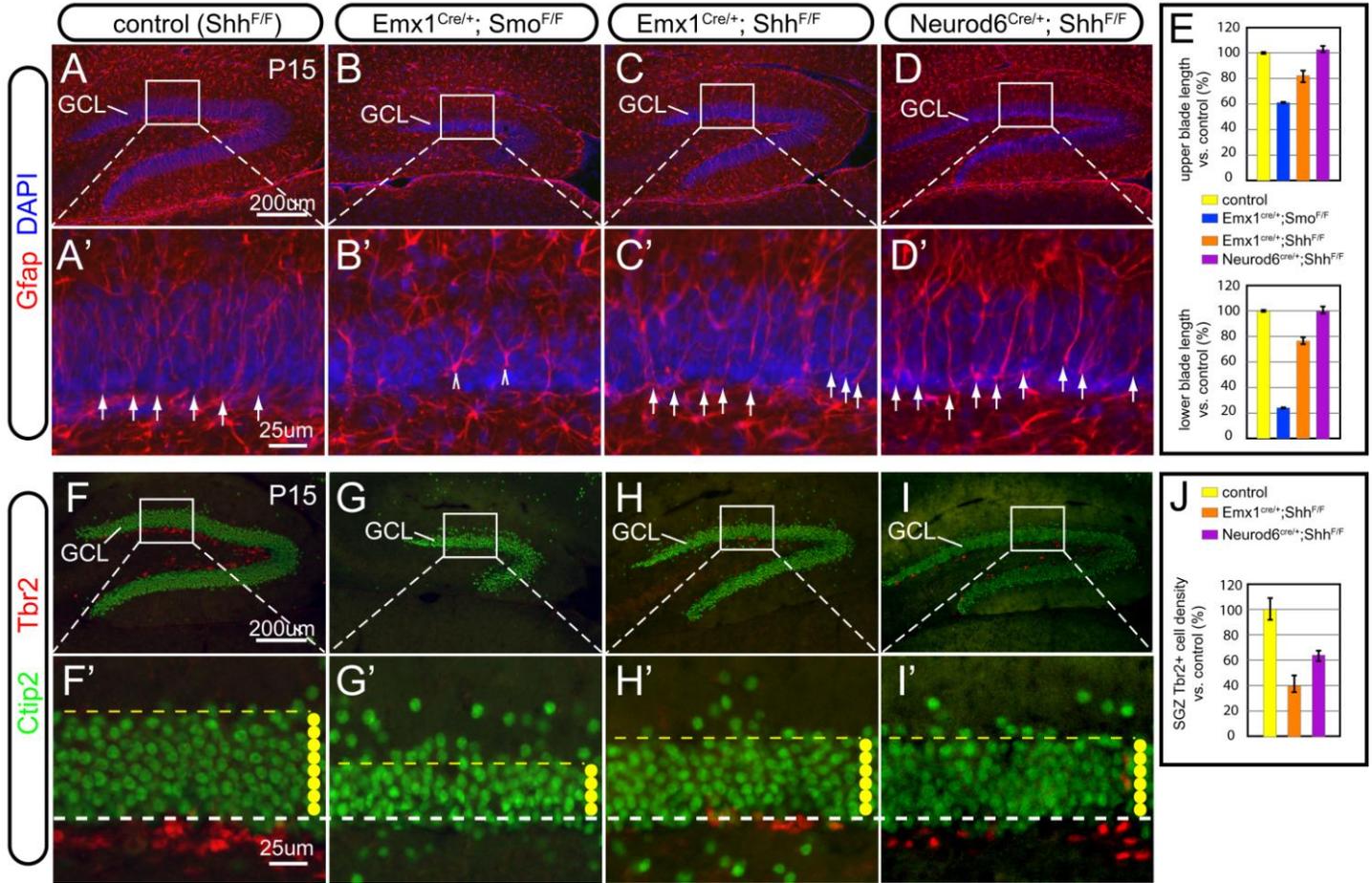
**(A-C)** The Shh lineage expressed markers for oligodendrocyte progenitor cells (OPCs) in the dentate gyrus at E17.5. By crossing Shh-gfpcre with Rosa-Yfp, the Shh lineage marked as Yfp+ cells (enhanced with Gfp antibody staining) in the dentate gyrus at E17.5 was colabeled with oligodendrocyte progenitor cell (OPC) markers -- Pdgfr $\alpha$  **(A)**, Sox10 **(B)** and Olig2 **(C)**.

**(D)** In the Olig2<sup>cre/+</sup>;Shh<sup>F/F</sup> animals, Shh was removed from the oligodendrocyte lineage. However, the distribution of the Gli1-nLacZ+ cells in the dorsal DG was unaffected by P7.

**(E-G)** By crossing Shh-Gfpcre with the cre reporter Ai14, a horizontal section at P15 **(E)** showed the Shh lineage also marked the cells in the MEC (arrows in **F**), which made projection to the middle molecular layer in the DG **(G)**.

LEC, lateral entorhinal cortex; MEC, medial entorhinal cortex.

Figure S5 (related to Figure 7)



**Figure S5 (related to Figure 7)**

(A-D) and (A'-D') Gfap staining at P15 showed the radial glial scaffolding in the control (A, A'), Emx1-Smo cKO (B, B'), Emx1-Shh cKO (C, C'), Neurod6-Shh cKO (D, D'). Compared to the control (arrows in A'), the radial glia spanning the GCL poorly developed in the Emx1-Smo cKO (arrowheads in B'), whereas the radial glial fibers were rather intact in the Emx1-Shh cKO (arrows in C') and Neurod6-Shh cKO (arrows in D').

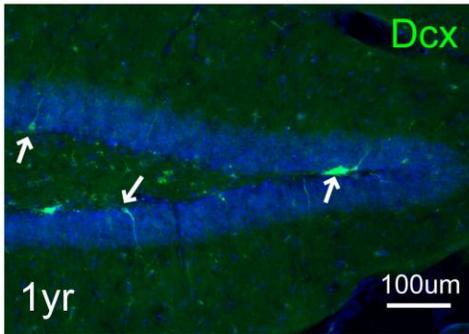
(E) Quantification was made for the length of the upper and lower blades in the control, Emx1-Smo, Emx1-Shh and Neurod6-Shh cKOs. In relation to the control, the length of the upper and lower blades were  $61\pm 1\%$  (n=5,  $p<0.001$ ) and  $24\pm 1\%$  (n=5,  $p<0.001$ ) in the Emx1-Smo cKO, whereas they were  $81\pm 5\%$  (n=5,  $p<0.01$ ) and  $76\pm 3\%$  (n=5,  $p<0.01$ ) in the Emx1-Shh cKO, and  $102\pm 4\%$  (n=5,  $p=0.25$ ) and  $98\pm 3\%$  (n=5,  $p=0.31$ ) in the Neurod6-Shh cKO.

(F-I) and (F'-I') Ctip2+ cells in the GCL and Tbr2+ neurogenic precursors in the SGZ were shown for the control (F, F'), Emx1-Smo cKO (G, G'), Emx1-Shh cKO (H, H') and Neurod6-Shh cKO (I, I'). Emx1-Smo cKO displayed complete deficiency for the Tbr2+ cells in the SGZ and the thinnest GCL (~4 cells thick, yellow dots in G'). Emx1-Shh cKO also had diminished number of Tbr2+ cells in the SGZ and thinner GCL (~6 cells thick, yellow dots in H' and I'), compared to the control (~8 cells thick, yellow dots in F').

(J) Quantification of Tbr2+ cell densities in the SGZ was made for the control, Emx1-Shh cKO and Neurod6-Shh cKO. As compared to the control ( $100\pm 9\%$ ), it was  $40\pm 7\%$  (n=5,  $p<0.001$ ) in the Emx1-Shh cKO and  $62\pm 4\%$  (n=5,  $p<0.001$ ) in the Neurod6-Shh cKO.

Data were shown as mean $\pm$ SEM and p values for the indicated sample sizes were returned by Student's *t*-test for two-tailed distribution with unequal variance.

**Figure S6 (related to Figure 8)**



**Figure S6 (related to Figure 8)**

Only very few Dcx+ immature neurons (arrows) were detectable in the DG in one year old animals