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Supplemental Information

Hippocampal TERT Regulates Spatial Memory Formation through Mod-

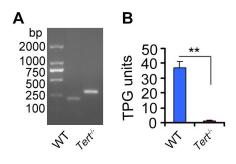
ulation of Neural Development

Qi-Gang Zhou, Meng-Ying Liu, Han-Woong Lee, Fuyuki Ishikawa, Sushil Devkota, Xin-Ru Shen, Xin Jin, Hai-Yin Wu, Zhigang Liu, Xiao Liu, Xun Jin, Hai-Hui Zhou, Eun Jeoung Ro, Jing Zhang, Yu Zhang, Yu-Hui Lin, Hoonkyo Suh, and Dong-Ya Zhu

1	Supplementary Information
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4	Hippocampal TERT regulates spatial memory formation
5	through modulation of neural development
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11 Supplemental Figures

- 14 Figure S1. Validation of the Tert -/- mice. Related to Figure 1



- 18 Figure S2. Latency to the platform in visible platform training. Related to Figure 1, 2, 6 and 7

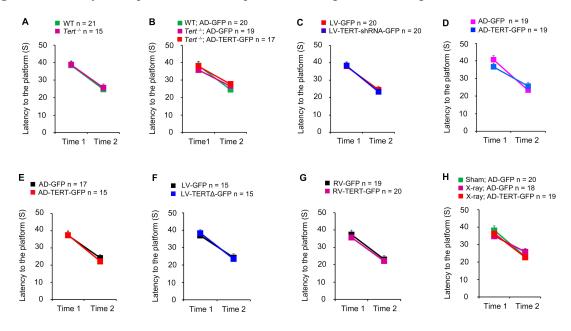
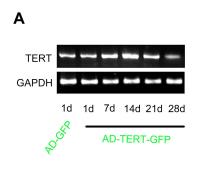
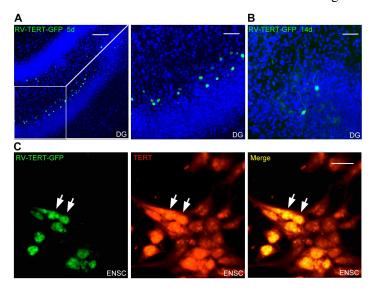


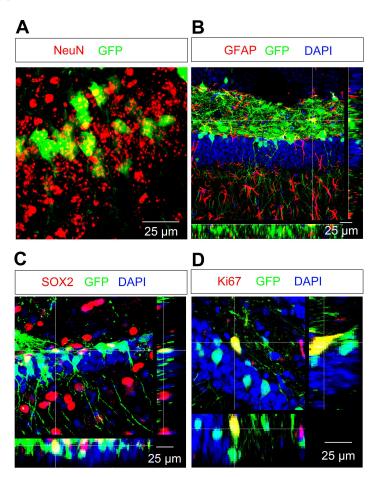
Figure S3. TERT expression dynamic induced by AD-TERT-GFP. Related to Figure 2



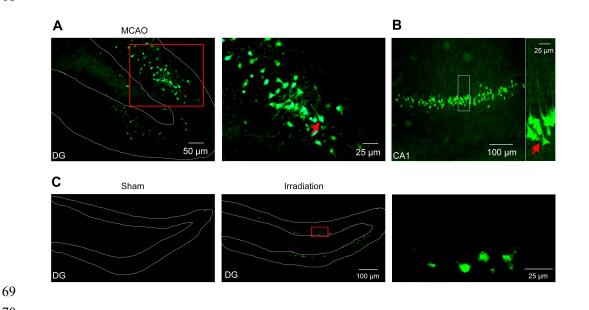
- **Figure S4.** Validation of RV-TERT-GFP *in vivo* and *in vitro*. Related to Figure 4



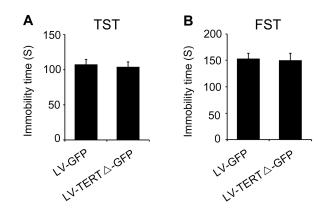
49 Figure S5. Cell types of LV-TERT-GFP infected cells in the DG. Related to Figure 7



- Figure S6. X ray-irradiation causes cell death selectively in the sub-granular layer of the DG. Related to
- Figure 7



- Figure S7. Expression of shortened TERT with non-activity in the DG did not cause depression-related
- behavior. Related to Figure 6



Experiments	Groups	n	Swim speed (mm/s) Mean ± SEM	<i>P</i> value
Figure 1A	WT	21	507.4667 ± 14.9994	0.2199
Figure TA	Tert ^{-/-}	15	479.3620 ± 16.3210	0.2199
	WT;LV-GFP	20	445.1270 ± 15.7291	
D ' 1D	<i>Tert-</i> '-;LV-GFP	19	451.6412 ± 18.0740	0.2430
Figure 1E	<i>Tert^{-/-};</i> LV-TERT-GFP	17	413.9472 ± 15.7541	
	LV-GFP	20	431.5740 ± 12.4375	
Figure 1 H	LV-TERT-shRNA-GFP	20	420.2310 ± 16.9280	0.5923
	AD-GFP	19	452.5030 ± 22.5279	0 7259
Figure 2A	AD-TERT-GFP	19	442.1970 ± 20.2784	0.7358
E. 0D	AD-GFP	17	457.2770 ± 9.8820	0.2354
Figure 2 B	AD-TERT-GFP	15	441.4620 ± 15.3795	
	LV-GFP	15	439.3920 ± 10.4737	
Figure 6E	LV-TERTΔ-GFP	15	447.8219 ± 12.3251	0.1592
	RV-GFP	19	467.9337 ± 15.2172	0.3116
Figure 7A	RV-TERT-GFP	20	489.0720 ± 13.5414	
	Sham + AD-GFP	20	458.0568 ± 12.9293	
Figure 7 D	X ray + AD-GFP	18	477.3935 ± 15.0006	0.8916
	X ray + AD-TERT-GFP	19	484.0584 ± 16.8535	

Table S1. The swimming speeds in probe test of mice in each experiment. Related to Figure 1, 2, 6and 7

79 Movie S1. Related to Figure 1

This movie showed the performance of a mouse in WT group in MWM test at day 5 during 5-day
training. The WT mouse found the platform in a short time.

Movie S2. Related to Figure 1

This movie showed the performance of a mouse in *Tert^{-/-}* group in MWM test at day 5 during 5-day training. The *Tert^{-/-}* mouse did not found the platform.

91 Supplemental Figure Legends

- Figure S1. Validation of the *Tert* --- mice (related to Figure 1). (A) RT-PCR showing the genotype of the *Tert* --- mice. (B) Deletion of telomerase activity in the embryonic stem cells from *Tert* --- mice (n = 3
- 95 independent experiments, P = 0.0012, Student's t-test.). Error bars, s.e.m.
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97 Figure S2. Latency to the platform in visible platform training (related to Figure 1, 2, 6 and 7). (A)

98 Related to Figure 1A. (B) Related to Figure 1E. (C) Related to Figure 1H. (D) Related to Figure 2A. (E)

99 Related to Figure 2B. (F) Related to Figure 6E. (G) Related to Figure 7A. (H) Related to Figure 7D.

100 One-way ANOVA in (**B** and **G**) and Student's *t*-test in the others. Error bars, s.e.m.

101

102 Figure S3. TERT expression dynamic induced by AD-TERT-GFP (related to Figure 2). (A)

103 RT-PCR showing the expression dynamic of TERT mRNA after injection of AD-TERT-GFP into the

- 104 DG. n = 3 independent experiments.
- 105

Figure S4. Validation of RV-TERT-GFP in vivo and in vitro (related to Figure 4). (A) 106 Representative image of the DG infected by RV-TERT-GFP 5 days after RV infection. On the right: a 107 108 high-magnification image from the box area, showing the location of RV-TERT-GFP in the 109 sub-granular layer. (B) Representative image of the DG infected by RV-TERT-GFP 14 days after RV 110 infection. Note that the major GFP fluorescence was observed in the cell body. (C) Representative 111 image of the cultured NSCs infected by RV-TERT-GFP (left) with immunostaining with anti-TERT 112 primary antibody (middle). The merge of RV-TERT-GFP signal with TERT immunostaining signal (yellow) was shown (right). Note: the signal of TERT in the NSCs infected by RV-TERT-GFP (arrow 113 indicated) was significantly stronger than others, indicating an enhancement of TERT expression in 114 115 NSCs by RV-TERT-GFP 24 hours after infection in NSCs culture. Scale bars: A (left), 100 µm; A 116 (right) and **B**, 50 µm, **C**, 25 µm.

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Figure S5. Cell types of LV-TERT-GFP infected cells in the DG (related to Figure 7). (A) Representative photo showing that NeuN⁺ neurons located at the granular layer were infected by LV-TERT-GFP. (**B**) Representative photo showing that GFAP⁺ astrocytes in the hillus were infected by LV-TERT-GFP. (**C**) Representative photo showing that SOX2⁺ cells located in the sub-granular layer were infected by LV-TERT-GFP. (**D**) Representative photo showing that Ki67⁺ cells located in the sub-granular layer were infected by LV-TERT-GFP.

124

125 Figure S6. X ray-irradiation causes cell death selectively in the sub-granular layer of the DG

126 (related to Figure 7). (A) Representative photos showing FJ-positive cells located in the DG 7 days

127 after MCAO. A magnification of the box area was shown on the right. Red arrow indicated that

FJ-positive cells located in the granular layer with neuronal morphology. (B) Representative photos showing FJ-positive cells located in the CA1 7 days after MCAO. A magnification of the box area was shown on the right. Red arrow indicated that FJ-positive cells located in the granular layer with neuronal morphology. (C) Representative photos showing FJ-positive cells in the DG of mice with/without exposure to irradiation. A magnification of the box area was shown on the right. Red arrow indicated that FJ-positive cells located in the granular layer with the morphology of NPCs. n = 3independent experiments.

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136 Figure S7. Expression of shortened TERT with non-activity in the DG did not cause 137 depression-related behavior (related to Figure 6). (A) Bar graph showing the immobility time of 138 mice in TST 28 days after injection of LV-GFP or LV-TERT Δ -GFP into the DGs. n = 15. P = 0.7148. 139 Student's *t*-test. (B) Bar graph showing the immobility time of mice in FST 28 days after injection of 140 LV-GFP or LV-TERT Δ -GFP into the DGs. n = 15. P = 0.8810. Student's *t*-test.

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- 155 Supplemental Experimental Procedures
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157 **Genotyping**

Genotyping Primers of *Tert^{-/-}* mice: (1) gcg tgg agt atc ctc ctg cat ctc ta, (2) ctg tct cct aaa gga ctt gtg gac tt, (3) agg att ggg aag aca ata gca ggc at. The primer stock should be diluted to make 100 picomoles/ μl. Then mix 10 μl of each primer 1, 2, and 3 with 70 μl 3-distilled water to make final volume of 100 μl. This 100 μl is the working stock. 1 μl of the working stock can be used for 1 PCR reaction. PCR cycles: 32 cycles. Denaturation: 94 °C, 30 sec. Annealing: 55 °C, 30 sec. Extension: 72 °C, 1 min. Final extension: 72 °C, 10 min.

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165 Virus construction

166 RV-TERT-GFP The sequence of TERT-GFP in the plasmid pDC315-TERT-GFP was cutted by using 167 restriction enzymes and inserted into the plasimd pCAG-GFP (For RV-GFP) by T4 DNA ligase to 168 replace the sequence of GFP in pCAG-GFP. GFP was fused to the C-terminal of TERT. The plasmid 169 was named as pCAG-TERT-GFP, which was used to identify by DNA sequencing. Using 100µl 170 Lipofectamine 2000, 293T cells were co-transfected with 22.5 µg of pCAG-TERT-GFP, 15 µg of 171 pCMV-GP, and 7.5 µg of pCMV-VSVG to generate the recombinant retrovirus, RV-TERT-GFP. After 48 h, supernatant was harvested from 293T cells, filtered at 0.45 μ m, and pelleted by 172 173 ultracentrifugation at 18000 × g for 2 h at 4 °C. After resuspension by PBS, serially diluted retrovirus 174 was used to transduce 293T cells; 4 days later, labeled 293T cells were counted to calculate the viral 175 titer (~ 2×107 transducing units/ml). As a control, we also generated a retroviral vector that expresses 176 GFP (RV-GFP) or RFP (RV-RFP) alone.

177 AD-TERT-GFP Briefly, the coding sequence of mouse TERT was amplified by RT-PCR. The 178 sequences follows: forward, primer were as 179 5'-GTAGAACGCAGATCGAAT-TCATGACCCGCGCTCCTCG-3'; reverse, 180 5 '-CCCTTGCTCACCATG-AATTCGTCCAAAATGGTCTGAAAGTC-3'. The PCR fragments and 181 the pDC315-GFP plasmid were digested with EcoR I and ligated with T4 DNA ligase to produce 182 pDC315-mTERT-IRES-GFP. The plasmid was used to trans-form competent DH5a Escherichia coli 183 bacterial strains for identification. Using 100µl of Lipofectamine 2000 mixed with 50 µl of DMEM, 184 HEK293 cells were cotransfected with 5 µg of the pDC315-GFP plasmid with a cDNA encoding 185 mTERT and 5 μ g of the pBHG lox Δ E1,3 Cre plasmid as a helper plasmid to generate the recombinant 186 adenovirus, AD-TERT-GFP. After 8 d, supernatant was harvested from HEK293 cells. After 3 times 187 the virus amplification, the supernatant was filtered at 0.45µm and purified using the Adeno-X Virus 188 Purification kit. After resuspension, serially diluted adeno-virus was used to transduce HEK293 cells. 189 Seven days later, labeled HEK293 cells were counted to calculate the viral titer (~ $2.5 \times 10 \ 10 \ pfu/ml$). 190 As a control, we also generated a adenoviral vector that expresses GFP alone (AD-GFP).

191 LV-TERT-GFP and LV-TERTA-GFP Briefly, the coding sequence of mouse TERT was amplified by 192 RT-PCR. The follows: forward. primer sequences were as 193 5'-GTAGAACGCAGATCGAAT-TCATGACCCGCGCTCCTCG-3'; reverse, 194 5 '-CCCTTGCTCACCATG-AATTCGTCCAAAATGGTCTGAAAGTC-3'. The PCR fragments was 195 cutted by restriction enzymes and inserted into the plasimd pCMV-GFP (For LV-GFP) by T4 DNA 196 ligase to replace the sequence of GFP in pCMV-GFP. GFP was under control of another promotor 197 pUbi. Mice mTERTA gene was amplified by our previous consructed plasmid LV-TERT-GFP carring the whole gene encoding TERT by deletion mutant PCR. The primer sequences were as follows: 198 199 forward. 5'-GAGGATCCCCGGGTACCGGTCGCCACCATGACCCGCGCTCCTCGTTGCC-3'; 200 reverse, 5 '-TCCTTGTAGTCCATACCGTCCAAAATGGTCTGAAAGTCTGTGCTTAG -3'. The 201 PCR fragments and the pGV287-GFP plasmid were digested with Age 1 and BamH 1 and ligated with 202 T4 DNA ligase to produce LV-TERTΔ-GFP. Using 100µl Lipofectamine 2000, 293T cells were 203 co-transfected with 20 µg of pCMV-TERT-GFP, 10 µg of VSVG, 7.5 µg of RSV-REV and 3.5 µg 204 pMDL g/p RRE to generate the recombinant lentiovirus, LV-TERT-GFP. After 48 h, supernatant was 205 harvested from 293T cells, filtered at 0.45 μ m, and pelleted by ultracentrifugation at 18000 \times g for 2 h at 4 °C. After resuspension by PBS, serially diluted retrovirus was used to transduce 293T cells; 4 206 207 days later, labeled 293T cells were counted to calculate the viral titer (~ 2×109 transducing units/ml). 208 As a control, we also generated a retroviral vector that expresses GFP alone (LV-GFP).

209 LV-TERT-shRNA-GFP Mouse TERT shRNA (m) Lentiviral Particles, we named it 210 LV-TERT-shRNA-GFP, and its control shRNA Lentiviral Particles (LV) were purchased (Santa Cruz, 211 CA, USA). LV-TERT-shRNA-GFP is a pool of concentrated, transduction-ready viral pa rticles 212 containing 4 target-specific constructs that encode 19- 25 nt (plus hairpin) shRNA designed to knock 213 down gene expression. Each vial contains 200 μ l frozen stock containing 1.0 × 106 infectious units of 214 virus (IFU) in Dulbecco's Modified Eagle's Medium with 25 mM HEPES pH 7.3.

215 RV-SYN-GTR, RV-SYN-GT, and EnvA- Δ R-mCh The plasmids of RV-SYN-GTR and RV-SYN-GT 216 were constructed based on the plasmids from Callaway's lab purchased from addgene (Osakada and 217 Callaway, 2013). RV-SYN-GTR and RV-SYN-GT were produced by transient transfection of the 218 RV-SYN-GTR or RV-SYN-GT vector (7.5 µg), CMVGagPol (5 µg) and CMV-VSVG (2.5 µg) in 60% 219 confluent 293T cells grown in 10-cm plates for 5 hours. Virus-containing supernatant was harvested 220 48 hours later. The final titers were estimated to be ~ 2 \times 107 pfu/ml or ~ 1 \times 107 pfu/ml as 221 determined by infection of serially diluted virus into 293T cells. The production of EnvA-ΔR-mCh 222 was carried out as described previously (Osakada and Callaway, 2013; Osakada et al., 2011). Briefly, 223 glycoprotein–gene-deleted rabies virus vector (ΔR -mCherry) was generated in which a mCherry 224 reporter gene was replaced with the cDNA encoding the rabies virus glycoprotein. The helper cell line, 225 BHK-EnvARGCD, was infected with ΔR -mCherry, to produce rabies virus pseudotyped with 226 envelope protein EnvA. After 2 h and 4 h, the cells were washed with PBS three times and the media

reapplied. Supernatants containing Δ R-mCherry rabies virus pseudotyped with EnvA were harvested 5 days later, filtered and concentrated by ultracentrifugation. EnvA- Δ R-mCh titre was estimated to be ~1.2×107 pfu/ml and diluted for use to ~4×106 pfu/ml.

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231 Western blot

Western bolt analysis of samples from cultured hippocampal neurons, NSC, and hippocampal tissues of animals was performed as described previously (Zhou et al., 2011b). The primary antibodies: Rabbit Anti-Flag (Bioss, bs-0287R), Mouse Anti-GAPDH (Kangchen Bio-Tech, kc-5G4), and Rabbit Anti-Synapsin 1 (Millipore, AB1543). Appropriate horseradish peroxidase-linked secondary antibodies were used for detection by enhanced chemiluminescence (Pierce).

237

238 MCAO

239 Focal cerebral ischemia was induced by MCAO, as described previously (Zhou et al., 2010). In brief, 240 under chloral hydrate anesthesia (350 mg/kg, i.p.), a 8/0 surgical nylon monofilament with rounded tip 241 was introduced into the left internal carotid artery through the external carotid stump, advanced 16–17 242 mm past the carotid bifurcation until a slight resistance was felt. At this point, the intraluminal 243 filament blocked the origin of the middle cerebral artery and occluded all sources of blood flow from 244 the internal carotid artery, anterior cerebral artery, and posterior cerebral artery. Throughout the 245 procedure, body temperature was maintained at 37 ± 0.5 °C. The filament was left in place for 90 min 246 and then withdrawn. In the sham-operated animals, the occluding filament was inserted 7 mm above 247 the carotid bifurcation.

248

249 Fluoro-Jade staining

Neuronal degeneration and death were determined by Fluoro-Jade (FJ; Histo-Chem) staining (Zhou et al., 2011a). In brief, sections were washed and mounted on glass slides and dried overnight. The slides were immersed for 3 min in absolute ethanol solution, for 1 min in 70% ethanol solution, and for 1 min in distilled water. Then, the slides were transferred into a solution containing 0.01% Fluoro-Jade and 0.1% acetic acid for 30 mintues on a shaker. After three 10 mintues washes, the slides were finally coverslipped for analysis.

256

257 Immunocytochemistry

Mice were anesthetized with with 0.07 ml of a mixture of ketamine (90.9 mg/ml) and xylazine (9.1 mg/ml) and perfused transcardially with saline followed by about 50 ml of 4% paraformaldehyde (PFA). Brains were removed and postfixed overnight in 4% PFA. Serial sections (40µm) were made on an oscillating tissue slicer in a bath of physiological saline. Every sixth section throughout the hippocampus was processed for 5-bromo-2'-deoxyuridine (BrdU) (Rat, Bio-rad, OBT0030) immunohistochemistry as described previously. All the sections throughout the brain were collected for tracing analyis. All the cells were counted in each section by another experimenter blinded to the
study code. Other primary antibodies: Rabbit Anti-DCX (Santa Cruz Biotechnology, sc-28939),
Rabbit Anti-GFAP (Abcam, ab7260), Mouse Anti-GFP (Santa Cruz Biotechnology, sc-101525),
Rabbit Anti-RFP (Abcam, ab62341), Rabbit Anti-Ki67 (Abcam, ab66155), Rabbit Anti-SOX2
(Abcam, ab97959), Rabbit Anti-NeuN (Cell Signaling, #24307), and Rabbit Anti-Synapsin 1
(Millipore, AB1543).

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271 **RT-PCR**

272 Total mRNA was extracted from the hippocampus using Trizol reagent according to the 273 manufacturer's instructions (Sigma). The primers used for RT-PCR: TERT: forward, 5' -ATGGCGTTCCTGAGTATG - 3', reverse, 5' -AGCCAGAGGCCTTTAGT - 3'; GAPDH: forward, 274 5' - CAAGGTCATCCATGACAACTTTG - 3', reverse, 5' - GTCCACCACCCTGTTGCTGTAG -275 276 3'. PCR conditions were 30 cycles of denaturation at 94°C for 45 s, annealing at 55°C for 45 s, and 277 extension at 72°C for 45 sec. PCR products were separated by electrophoresis through 1.5% agarose 278 gel containing 0.5% lg/mL ethidium bromide and imaged using a BioDoc-IT imaging system 279 (Bio-Rad); band intensities were determined using GS-710 calibrated imaging Densitometer 280 (Bio-Rad). The mRNA for GAPDH was detected as a standard.

281

Telomerase activity assay

283 Telomerase activity was detected using TRA-PEZE XL telomerase detection kit (Millipore, Billerica, 284 MA) as described previously (Zhou et al., 2011a). Following the manufacturer's instructions, 285 telomeric repeat amplification protocol reactions (TRAP) were performed using the TRAPEZE XL 286 telomerase detection kits (Millipore) for analysis of the telomerase activity. The fluorescence energy 287 transfer primers were used to generate fluorescently labeled TRAP products, quantitatively measured 288 with a fluorescence plate reader (SpectraMax M2e) or visualized after terminal deoxynucleotidyl 289 transferase-mediated dUTP nickend labeling on a 10% nondenaturing gel and SYBR Green I 290 (Invitrogen) staining.

291

292 Analysis of spine density and morphology of neurons

Images of GFP-labeled neurons or DCX⁺ neurons were acquired at 1 μ m intervals with the Leica confocal system with a plane apochromatic 40 × oil lens and a digital zoom of 1.5. The image files were used for reconstruction of the morphology with the Imaris v7.2.3 program. The length of each dendritic segment was determined by tracing the center of the dendritic shaft, and the number of spines was counted manually from the two-dimensional projections. The linear spine density was calculated by dividing the total number of spines by the length of the dendritic segment. For classification of mushroom spines, major and minor axes of each spine head were identified. When the 300 width size of the spine > $0.6 \mu m$, the spines regonized as mushroom spines. Confocal imaging and 301 data quantification were done by the same person, who was blinded to the experimental conditions.

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303 Stereotaxic surgery

Mice were anesthetized with 0.07 ml of a mixture of ketamine (90.9 mg/ml) and xylazine (9.1 mg/ml) and placed in a stereotaxic apparatus. Stereotaxic surgery was performed to deliver viruses or drugs into the hippocampal DG (1 μ l, coordinates: 2.3 mm posterior to bregma, 1.35 mm lateral to the midline, and 2.3 mm below dura), The mice were recoverd on hot pad (37 °C) until waken up and then retrun back to homecages (Zhou et al., 2011).

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310 Hippocampal X-irradiation

As described previously (Zhou et al., 2011), X-irradiation was performed. Briefly, mice were anesthetized with 0.07 ml of a mixture of ketamine (90.9 mg/ml) and xylazine (9.1 mg/ml). Then mice were placed in a stereotaxic apparatus and the hippocampal zone were exposed to cranial irradiation using therapeutical x-ray equipment operated at 300 kVp and 20 mA. Mice were protected with a lead shield that covered the entire body, with the exception of a 3.2×11 mm treatment field above the hippocampus. The corrected dose rate was ≈ 3 Gy/min at a source-toskin distance of 100 cm. The procedure lasted 1 min and 42 s, delivering a total of 5 Gy each time.

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319 Cell Cultures

320 Primary hippocampal neurons were cultured from embryonic hippocampus, on embryonic day 18 321 (E18), of WT or Tert^{-/-} mice in neurobasal medium (Gibco) containing 2% B27 supplement as reported (Luo et al., 2010). NSCs differentiation was also performed from embryonic hippocampus on 322 embryonic day 18 (E18), of WT or Tert^{-/-} mice. Monolayer-cultured NSCs were allowed to 323 differentiate in growth factor-free DMEM/F12 (1:1) medium containing 2% B27 and 0.5% fetal 324 325 bovine serum. Four days later, the cultures were fixed and stained with DCX and GFAP antibodies to 326 mark neurons and astrocytes, respectively. The percentages of neurons and astrocytes were calculated 327 in 10 high-power fields systematically across the coverslip.

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329 Supplemental References

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