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AhR Deletion Promotes Aberrant Morphogenesis and Synaptic Activity of Adult-Generated Granule Neurons and Impairs Hippocampus-Dependent Memory

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39

40 **ABSTRACT**

Newborn granule cells are continuously produced in the subgranular zone of 41 42 dentate gyrus throughout life. Once these cells mature, they integrate into preexisting circuits modulating hippocampus-dependent memory. Subsequently, 43 mechanisms controlling generation and maturation of newborn cells are 44 essential for proper hippocampal function. Therefore, we have studied the role 45 46 of Aryl Hydrocarbon Receptor (AhR), a ligand-activated bHLH-PAS transcription factor, in hippocampus-dependent memory and granule neuronal morphology 47 and function using genetic loss-of-function approaches based on constitutive 48 and inducible-nestin AhR^{-/-} mice. The results presented here show that the 49 impaired hippocampus-dependent memory in AhR absence is not due to its 50 effects on neurogenesis but to aberrant dendritic arborisation and to an 51 52 increased spine density albeit with a lower number of mature mushrooms spines in newborn granule cells, a finding that is associated to an immature 53 54 electrophysiological phenotype. Together, our data strongly suggest that AhR 55 plays a pivotal role in the regulation of hippocampal function, by controlling hippocampal granule neuron morphology and synaptic maturation. 56

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58 SIGNIFICANCE STATEMENT

Hippocampus-dependent memory depends on the generation and maturation of dentate gyrus (DG) newborn granule cells. Aryl Hydrocarbon Receptor (AhR) is a ligand-activated bHLH-PAS transcription factor recently implicated in dendrite branching in the CNS. Since its role in the modulation of dendrite branching and plasticity of adult hippocampal newborn granule neurons and subsequent impact on hippocampus-dependent memory remains unknown, we have undertaken its study using genetic loss-of-function approaches in adult mice. Our study provides evidence indicating that AhR is a regulator of dendrite arborisation and proper synaptic maturation of adult hippocampal newborn neurons and showing its critical role for learning and memory function. These findings point out AhR as a new potent druggable target for the treatment of several cognitive disorders.

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72 INTRODUCTION

73 In the adult mammal brain, the hippocampus is one of the main regions 74 implicated in cognitive function. Throughout life, hippocampal newborn neurons 75 migrate into the granule cell layer to become new dentate granule cells where 76 they integrate synaptically into the pre-existing circuits providing potential 77 substrates for new learning and memories (Altman and Das, 1965; 78 Kempermann et al., 1997; Squire and Zola-Morgan, 1991; van Praag et al., 79 1999). The correct morphogenesis of these newborn neurons is a critical feature for the adequate function of the hippocampus. Although the aberrant 80 81 integration of adult newborn granule neurons is able to disrupt cognitive function and is linked to several neurological disorders (Winkle et al., 2016; Zhou et al., 82 2013), the molecular factors that drive the morphogenesis and maturation of 83 84 these cells are still quite unknown.

The aryl hydrocarbon receptor (AhR), a ligand-activated transcription 85 86 factor that belongs to the basic Helix-loop-Helix Per-Arnt-Sim (bHLH-PAS) 87 superfamily, has been traditionally studied in association with toxic effects of the environmental pollutants and xenobiotic compounds metabolism and its role in 88 89 the immune system (Fernandez-Salguero et al., 1995; Fernandez-Salguero et 90 al., 1996; Mandal, 2005; Mulero-Navarro and Fernandez-Salguero, 2016; Murray et al., 2014). However, recent studies across different species suggest 91 92 other important biological roles of AhR in other systems, as the CNS. In 93 invertebrates, AhR homologs are implicated in dendrite branching in neurons: in 94 Drosophila, the loss-of-function of the AhR homolog (spineless) promotes more complex dendritic arborisation in sensory neurons (Crews and Brenman, 2006; 95 96 Kim et al., 2006); in C. elegans, ahr-1 mutant (AhR homolog) neurons also turn

97 into a highly-branched architecture (Smith et al., 2013). In mammals, AhR is 98 expressed in the adult brain (Kimura and Tohyama, 2017) and its constitutive activation drastically reduces dendritic arborisation and aberrant neuronal 99 positioning in cortical pyramidal neurons and olfactory bulb interneurons 100 (Kimura et al., 2016; Kimura et al., 2017; Kimura and Tohyama, 2017). AhR 101 mRNA is also expressed in the dentate gyrus (DG) granule cells of the adult 102 103 hippocampus (Kimura and Tohyama, 2017). In this area, even though it has been reported that AhR might modulate hippocampal neurogenesis (Latchney 104 et al., 2013), its role in the modulation of dendrite branching and plasticity in 105 106 adult hippocampal newborn granule neurons remains unknown. Therefore, we have studied the role of AhR in hippocampus-dependent memory and granule 107 neuronal morphology and function using genetic loss-of-function approaches in 108 109 adult mice. Our data demonstrate that the transcription factor AhR plays a crucial role in hippocampus-dependent function, by controlling dendritic 110 111 arborisation and dendritic spine growth in granule neurons.

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113 METHODS

114 Animals and tamoxifen treatment

Experiments were performed in male WT and AhR^{-/-} knockout mice (C57BL/6) at 4, 8 and 14 weeks of age, obtained from Taconic. Both WT and AhR^{-/-} mice were generated by crossing heterozygous AhR^{+/-} mice. AhR^{f/f} mice were acquired from The Jackson Laboratory and were maintained through homozygous breeding pairs. AhR icKO mice (tamoxifen-inducible AhR conditional knockout mice) were generated by crossing AhR^{f/f} mice (Walisser et al., 2005) with nestin-Cre^{ERT2} mice (Imayoshi et al., 2008) and then maintained 122 through homozygous breeding pairs on a C57BL/6 background. In these transgenic mice (nestin-Cre^{ERT2}/AhR^{f/f}), tamoxifen treatment suppresses the 123 expression of AhR in the neuroprogenitor cells present at the hippocampal 124 subgranular zone (SGZ). The tamoxifen protocol used in this study was as 125 described before (Cancino et al., 2013). Briefly, both AhR^{f/f} and AhR-icKO mice 126 were administered tamoxifen intraperitoneally in two different rounds. The first 127 128 round was performed at p30 and the second at p60, each round consisting of a daily injection of tamoxifen (180mg/Kg) in sunflower oil for 5 consecutive days. 129 Behavioural and histological analyses were performed 3 weeks after the last 130 131 tamoxifen administration. Mice had access to rodent chow and water ad libitum in a 12 h light/dark cycle room. This study was approved by the Animal Welfare 132 Committee of the Authors' Institution. 133

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135 BrdU treatment

For quantification of the proportion of proliferating SGZ neural precursors, a total of 4 injections of the cell proliferation marker BrdU (5-bromo-2⁻deoxyuridine; 100 mg/kg; Sigma-Aldrich) were administered intraperitoneally every 2h to 4, 8 and 14-weeks-old control and AhR^{-/-} mice. Twenty-four hours after the last administration, mice were sacrificed.

For the quantification of the integrated adult newborn neurons (BrdU⁺/calbindin⁺ cells), 8-weeks-old WT and AhR^{-/-} mice were injected daily with BrdU (100 mg/kg) intraperitoneally for 5 consecutive days, and mice were sacrificed 28 days after the last administration.

145

146 Histology

147 For histology and immunohistochemistry studies, mice were perfused transcardially with 0.1M PBS followed by 4% paraformaldehyde (PFA) in 0.1M 148 PBS (pH 7.4). Brains were post-fixed in PFA and transferred to 30% sucrose. 149 For SVZ (from bregma +1.70 mm to bregma 0.02 mm) and dentate gyrus (DG; 150 from bregma -1.46 mm to bregma -2.03 mm), coronal sections (30µm) were cut 151 152 using a microtome (Leica SM2000R) and stored in cryoprotective solution. Unless indicated otherwise, brain samples from AhR^{-/-} knockout and AhR icKO 153 mice after tamoxifen treatment were analysed at 2 and 3 months of age, 154 155 respectively.

Immunohistochemistry: Immunofluorescence was performed on free-156 floating sections. Briefly, sections were first permeabilized and blocked in 157 158 0.25% TritonX100 in PBS with 10% normal serum for 1 hour and then, incubated overnight at 4°C with the following primary antibodies in 0.25% Triton 159 160 X100 in PBS with 5% normal serum: goat anti-calbindin (neuronal marker; 161 1:500, SantaCruz), goat anti-DCX (doublecortin; neuroblast marker) (1:250, SantaCruz), rabbit anti-Ki67 (nuclear protein specifically expressed in cells 162 163 undergoing active proliferation; 1:500, Abcam), chicken anti-GFAP (glial 164 fibrillary acidic protein; astrocyte marker; 1:750, Thermo Scientific), mouse antinestin-PE (neural stem cell marker; 1:50, BD Biosciences), rabbit anti-AhR 165 166 (1:200, Enzo Life Sciences) and chicken anti-GFP (1:700, Thermo Scientific). 167 For BrdU staining, free-floating sections were pre-treated with 2 N HCl for 30 minutes at 37°C and, after blocking in 0.25% Triton X100 in PBS with 10% 168 normal serum for 1 hour, incubated overnight at 4°C with rat monoclonal anti-169 170 BrdU (1:200, Abcam) in 0.25% Triton X100 in PBS with 5% normal serum. The 171 secondary antibodies used were donkey Alexa-488 anti-goat (1:500, Invitrogen), donkey Cy3 anti-mouse (1:500, Vector Laboratories), goat anti-rat 172 biotinylated (1:250, Vector Laboratories), streptavidin Alexa-488 conjugate 173 (1:500, Thermo Scientific), goat Alexa-647 anti-chicken (1:500, Thermo 174 Scientific), donkey Cy3 anti-rabbit (1:500, Thermo Scientific) and donkey Alexa-175 488 anti-chicken (1:500, Thermo Scientific) in 0.25% Triton X100 in PBS with 176 177 5% normal serum. Controls performed in parallel without primary antibodies showed very low levels of nonspecific staining. 178

Image processing and quantitative analysis of immunostained sections: 179 180 Image acquisition was performed with a laser-scanning confocal imaging system (Zeiss LSM710) and image analysis was accomplished with the 181 ZEN2009 software (Zeiss). Image quantification was performed with ImageJ 182 183 Software (NIH) and Volocity Software (Improvision). Ki67⁺, DCX⁺, nestin⁺ and BrdU⁺ cells were counted in confocal z-stack images. In all cases, guantification 184 was performed using non-stereological methods. Specifically, every 5th section 185 186 (30 µm, separated 150 µm apart) was selected for a total of 5 representative matched sections per hippocampus (from bregma -1.46mm to bregma -2.06 187 188 mm) and SVZ (from bregma +1.70 mm to bregma 0.02 mm). Cells were 189 counted manually in frames of 212.55µm x 212.55µm (1024x1024) and data were expressed as the number of cells per 1000 µm². Since assessment of 5 190 191 sections may not reflect changes in hippocampal size and extent, quantification 192 data should not be considered in terms of absolute numbers. Although stereological assessment would be more accurate, the fact that cell numbers in 193 AhR^{-/-} mice ranged from being in excess to no differences with age suggests 194 195 that the methods we have employed are sensitive to major changes that occur with development. Co-localisation of calbindin⁺/BrdU⁺, nestin⁺/BrdU⁺,
nestin⁺/GFAP⁺, nestin⁺/GFAP⁺/BrdU⁺ was confirmed by orthogonal projection of
z-stack files.

In p60 AhR^{-/-} and p90 AhR-icKO mice and their respective controls, apical dendrite length of DCX⁺ cells was assessed in 5 serial sections (30 μ m) from dorsal hippocampus (from bregma -1.46mm to bregma -2.06 mm). Apical dendrite (considered the segment between the soma and the first dendrite ramification) was manually traced in confocal z-stack images taken at 63X and then dendritic length was measured by ImageJ. A total of 20-50 cells per animal from each group were quantified.

DCX⁺ dendritic staining was performed in confocal z-stack images taken 206 at 40X. Briefly, rectangular ROIs were generated around neuroblast dendritic 207 208 arborisation and somas. Total dendritic arborisation (distribution pattern of neuroblast dendrites along granular and molecular layers of the DG) was also 209 210 subdivided in 2 ROIs for differentiating proximal (GL) and distal (ML) DCX⁺ 211 dendritic staining. Integrated density was quantified in each compartment after 212 background subtraction. Total dendritic arborisation/soma, GL/soma and 213 ML/soma ratios were calculated from these values. This method was internally 214 normalized for immunostaining variability, since immunofluorescence values were always acquired in pairs of dendrites and adjacent somas. 215

216

217 Ex vivo flow cytometry from SGZ and SVZ-derived NPCs

To quantify the proportion of proliferating SGZ or SVZ neural precursors, mice were injected four times with 100 mg/Kg BrdU intraperitoneally every 2 h and sacrificed 24 h later. Brains were rapidly removed and SVZ and SGZ-

derived NPCs of WT and AhR^{-/-} mice were dissected, placed in ice-cold PBS 221 and dissociated into a single cell suspension. Cell suspensions were filtered on 222 40 um nylon mesh strainers and centrifuged at 300g for 10min at room 223 temperature. Next, cells were fixed, permeabilised and stained with anti-BrdU-224 APC and anti-Nestin-PE according to manufacturer's instructions (BD 225 Cytofix/Cytoperm Kit, BrdU Flow Kits BD Biosciences). Finally, cells were 226 227 washed and resuspended in 300 µl FACS Flow (BD Pharmingen); isotype controls (Miltenyi) were run in parallel. Whole suspensions were examined in a 228 FACSCalibur flow cytometer using CellQuest software (BD Pharmingen) and 229 230 data were analysed using FlowJo software (Tree Star Inc),

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232 Golgi-Cox staining

The fresh brain from 8-weeks-old AhR^{-/-} and WT mice without perfusion 233 were used for Golgi-Cox staining with FD Rapid GolgiStain Kit (FD Neuro 234 Technologies, Columbia, MD, USA) according to the user manual. Briefly, the 235 236 brain was first placed in impregnation solution for 2 weeks followed by 2 days in 237 30% sucrose. Then, brains were cut into 100-µm coronal sections using a 238 vibratome (Leica VT1000s) and stained. Neuronal reconstruction from each 239 animal were randomly drawn at 40X magnification for the different analysis by using the Neurolucida neuron tracing system (Microbrightfield, Colchester, VT). 240 241 Determination of total dendritic length of branches and Sholl analysis were 242 performed by using the Neuroexplorer software (Microbrightfield, Colchester, 243 VT). Sholl analysis was carried out by counting the number of dendrites that crossed a series of concentric circles at 10 µm intervals from the cell soma. To 244 245 calculate spine density of Golgi-stained neurons in the DG, a random 10-µm long dendrite segment in the molecular layer was measured (100-200 µm from the soma) and total number of spines was traced. The spine density was determined by dividing the total number of spines by the 10-µm length of the dendritic segment.

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Retrovirus-mediated labelling and morphological analysis of GFP⁺ newborn neurons

New neurons were labelled using a murine Moloney leukemia virus-253 254 based retroviral vector (CAG-GFP, a gift from Fred Gage, Salk Institute, La 255 Jolla, CA) (Zhao et al., 2006). Concentrated viral solutions were prepared by transfection of retroviral vectors into Gryphon Eco cells, followed by 256 ultracentrifugation of viral supernatant and concentrated virus solution by ultra-257 speed centrifugation (average 3 x 10⁷ iu/ml). Mice (8-weeks-old) were 258 anesthetized with isoflurane and placed in a stereotaxic frame. Two-µl retrovirus 259 260 was infused at a rate of 0.2 µl/min into the DG (-2 mm AP, -1.4 mm ML relative 261 to bregma, and 2.4 mm DV from skull) with a 5-µl 32-gauge Hamilton syringe. After the infusion, we allowed 5 extra min to avoid the retrovirus flow during 262 263 syringe releasing. Twenty-eight days after the viral infusion, mice from all 264 groups were sacrificed for GFP morphological experiments.

For morphometric analysis, 50-µm sections were used. A total of 10 sections of two series from each animal were used for immunohistochemical detection of GFP-labelled neurons. Eight-ten randomly selected neurons from each group were reconstructed. Confocal 40X stacks of images were obtained and z-projections were analysed to determine total dendritic length and to perform Sholl analysis. All cells were traced using *NeuronJ* plugin for ImageJ software. Sholl analysis was performed in order to determine dendritic
complexity using the plugin Sholl Analysis for ImageJ (Ferreira et al., 2014).

Primary apical dendrite length was manually traced from the soma until
the first dendrite ramification point in randomly selected neurons, and measured
by ImageJ.

For spine analysis, images of GFP-labelled dendritic processes at the 276 277 outer molecular layer were acquired at 0.5-µm intervals with a Zeiss LSM710 confocal microscope with a plan apochromatic 63X oil lens [numerical aperture 278 (NA), 1.4] and a digital zoom of 3X. The lsm images files were subjected to two 279 280 iterations of deconvolution with the AutoDeblur program (AutoQuant, Troy, NY). The length of each dendritic segment was measured, and total number of 281 spines in proximal (50-100 µm from the soma) and distal (100-200 µm from the 282 283 soma) dendritic segments were manually counted. The spine density was determined by dividing the total number of spines by the length of the dendritic 284 285 segment.

Mushroom spines in proximal (50-100 μ m from the soma) and distal (100-200 μ m from the soma) dendritic segments were identified when the estimated surface area (= π X D_{major} X D_{minor} /4) was \geq 0.4 μ m² (Zhao et al., 2014; Zhao et al., 2006). For the quantification, 8-12 dendritic segments from each animal per group were used. Confocal imaging and data quantification were performed blinded to the experimental conditions.

292

293 Electrophysiology experiments

AhR^{-/-} and WT mice (30- to 45-days-old) were anaesthetized with isoflurane (1.5-2% in a mixture of 80% synthetic air/20% oxygen) and

296	decapitated. The brain was quickly removed and placed in ice-cold artificial CSF
297	(ACSF) containing (in mM): NaCl 124, KCl 2.69, KH_2PO4 1.25, $MgSO_4$ 2,
298	NaHCO ₃ 26, CaCl ₂ 2, ascorbic acid 0.4 and glucose 10, continuously bubbled
299	with carbogen (95% O_2 and 5% CO_2 ; pH 7.3). Sagital hippocampal slices (325-
300	μ m thick) were obtained using a Leica VT 1200S vibratome and incubated (≥ 1
301	h) in a holding chamber at room temperature (21-24°C) in ACSF. Slices were
302	transferred to an immersion recording chamber and superfused at 1mL/min with
303	gassed ACSF including 50 μ M picrotoxin to block GABA _A receptors.
304	Experiments were performed at 25°C by using a temperature controller (Warner
305	Instruments). Granule cells from dentate gyrus were visualized under a 40x
306	water immersion objective and a Nomarski condenser combined with infrared
307	microscopy using differential interface contrast (DIC) in an Eclipse FN1 Nikon
308	microscope. Whole-cell electrophysiological recordings from granule cells were
309	performed using patch pipettes (3-4 $\mbox{M}\Omega$ resistance) pulled from thick-walled
310	borosilicate glass (1.5 mm outer diameter and 1.1 mm inner diameter) on a P-
311	97 puller (Sutter-Instrument) and filled with the internal solution containing (in
312	mM): K-Gluconate 135, KCl 10, HEPES 10, MgCl ₂ 1, ATP-Na ₂ 2 (pH = 7.3
313	adjusted with KOH; osmolality 280-290 mOsm/L). After formation of a whole-cell
314	configuration (-70 mV holding potential), current- or voltage-clamp protocols
315	were applied. For the analysis of the firing pattern and current-voltage
316	relationship, in current-clamp mode, a series of increasing currents (30 pA step,
317	500-ms duration with a 3-s interval) were injected. Evoked EPSCs were
318	recorded from dentate gyrus granule cells (voltage-clamp conditions, -70 mV
319	holding potential) by stimulation of glutamatergic afferents from perforant path
320	using a bipolar theta capillary (2-5 μm tip) filled with ACSF and placed in the

molecular layer of dentate gyrus. Stimuli were delivered at 0.33 Hz. Paired 321 Pulse Ratio (PPR) was obtained (2nd EPSC/1st EPSC) by delivering paired 322 pulses (20, 50, 75, 100, 150 and 200-ms interstimulus interval). AMPA/NMDA 323 ratio was obtained (2nd EPSC at +40 mV/1st EPSC at -60 mV holding potential) 324 by delivering paired pulses at 50-ms interstimulus interval. Series and input 325 resistances were monitored throughout the experiment using a -5 mV pulse. 326 327 Recordings were considered stable when the series and input resistances, 328 resting membrane potential and stimulus artefact duration were not changed > 20%. Cells that did not meet these criteria were discarded. The same procedure 329 was carried out for electrophysiology recordings in 50-days-old AhR^{f/f} and AhR-330 icKO mice (four weeks after 5 tamoxifen injections starting at 21-days-old). 331 Recordings were obtained by a PC-ONE amplifier and signals were fed to a 332 333 Pentium-based PC through a DigiData1322A interface board. The pCLAMP 10.2 software was used for stimulus generation, data display, acquisition, 334 335 storage and analysis.

336

337 Behavioural testing

338 Contextual fear conditioning. For mice, contextual fear conditioning 339 occurred in test chambers (31 cm x 24 cm x 21 cm) with shock-grid floors. The front, top and back of the chamber were clear acrylic and the sides were 340 modular aluminium. During training, AhR^{-/-} or AhR-icKO and their respective 341 342 controls were placed in the chamber and, after 2 min of habituation, they 343 received a mild or a weak conditioning protocol based on 3-foot shocks (0.6 mA, 2 s duration, 1 min apart) or 1-foot shock (0.48 mA, 2 s duration), 344 345 respectively. After conditioning, mice were removed from the chamber 1 min after the last shock. Behaviour was recorded by overhead cameras. Freezing
(i.e. absence of movement except for breathing) was measured using
automated scoring system for mice.

Barnes maze. A white circular platform (100-cm diameter, 70 cm above 349 350 the floor) contained 20 holes equally spaced around its perimeter. Under one of the holes, there was an escape box (17 x 13 x 7 cm) filled with paper bedding. 351 352 The location of this escape hole was always in the same place for all mice. For 353 avoiding navigation based on olfactory or proximal cues within the maze, the platform was rotated before each trial, and the spatial location of the escape 354 355 hole remained in a fixed location with respect to the distal room cues. During habituation, mice were allowed 5 min to freely explore the maze, with no escape 356 box present. During training, mice were given 3 trials per day for 6 days. On 357 358 each trial, the mouse was released in the centre of the maze and allowed for 5 min to enter the escape box, where it remained for 30 s. If a mouse failed to find 359 360 the escape box, it was guided by the experimenter. During probe test, the 361 escape box was removed from the maze, and the mouse was allowed to search for 5 min. Time spent around each hole was recorded. Search paths were 362 363 recorded by an overhead video camera and tracked using automated 364 Ethowatcher software.

Novel object recognition (NOR) and novel object location (NOL). Mice were placed in a rectangular arena (30 cm x 20 cm x 30 cm) with clear sidewalls containing two objects (A and B) for 8-min training session and returned to their home cages. The memory tests were performed as described (Nakashiba et al., 2008). Briefly, during the 8-min training phase of object recognition, two identical objects were placed in the arena. For the NOR test, 371 the animal's memory of one of the original objects was assessed by comparing 372 the amount of time spent exploring a novel object compared with that spent 373 exploring the familiar one during 8 min. For the NOL test, after the training 374 period, the animal's memory of one of the original objects was assessed by 375 comparing the amount of time spent exploring the new located object with that 376 spent exploring the original located one during 8 min. In both tests, the time 377 spent exploring each object was expressed as a percentage of exploration time 378 [% Exploration time = $(t_{novel}-t_{familiar})/(t_{novel}+t_{familiar})]$. Behaviour was recorded by 379 overhead cameras and videos analyses were performed by a blinded 380 experimenter.

Y-maze. The Y-maze was made of three solid white arms of equal size 381 (35-cm long and 5-cm wide with 10-cm high walls) joined in a Y-configuration. 382 383 The maze was cleaned with 70% ethanol between animals to eliminate traces of odour. For working memory, during habituation phase mice were placed 384 385 where the arms joined and allowed to freely explore the three arms for 6 min. 386 Arm entry was defined as having forelimbs inside an arm. The number of entries was recorded in order to calculate the alternation percentage (defined as 387 388 a set of consecutive arm entries), which was calculated by dividing the number 389 of triads by the number of possible alternations multiplied by 100. For spatial memory, during training phase mice were allowed to explore two of the three 390 391 arms for a total of 6 min while the third arm was blocked. Six hours later mice 392 were placed back in the maze for 6 min with all arms open, and the number of 393 entries in each arm was recorded. Behaviour was recorded by overhead cameras and videos analyses were performed by a blinded experimenter. 394

In order to avoid a possible interference due to the manipulation, training and testing on the histological studies, we used different mouse cohorts for each set of experiments. Specifically, two separate mouse cohorts were used for both contextual fear conditioning and Barnes maze. For the rest of the behavioural testing, we used two different groups of mice. One group was first trained in NOR and one week later in NOL and, in the other one, mice were first tested on Y-maze and one week later in NOL.

402

403 Statistical analysis

404 Results are expressed as mean ± SEM. Sample sizes for each experiment are indicated in the figure legends. In most experiments, sample 405 406 size was estimated based on previous extensive experience with similar 407 approaches. For specific experiments, we performed power analysis to adequate sample size with data from pilot studies (usually data from 3 408 409 animals/group) by calculating power analysis (http://www.biomath.info) with a 410 significance level of 5% and \geq 80% of power. Cohen's d was used to calculate effect size, with observed strong size effects (d values) ≥ 0.8 (Cohen, 1988). 411 412 Statistical significance was determined by use of a non-parametric, 2-tailed 413 Mann-Whitney t test; or a nonparametric, 2-way ANOVA followed by Bonferroni post hoc testing. Values of p<0.05 were considered statistically significant. All 414 415 statistical analyses were performed with Prism version 5.0 (GraphPad Software, 416 Inc).

417

418 **RESULTS**

419 Hippocampus-dependent memory is impaired in AhR^{-/-} mice

420 In order to test whether AhR plays a role in hippocampus-dependent memory. 8-weeks-old AhR^{-/-} mice and their WT littermates were subjected to a 421 battery of hippocampus-dependent tests. First, mice were trained in contextual 422 fear conditioning (CFC), an associative learning which involves the 423 hippocampus (Anagnostaras et al., 2001). No differences were found between 424 WT and AhR^{-/-} mice in response to the foot shock (data not shown), indicating 425 similar levels of nociception in both genotypes. During the retrieval, AhR^{-/-} mice 426 displayed a reduced freezing response in both 1h- and 24h-tests (p<0.05; 427 Figure. 1A), and a decreased percentage of activity suppression (88.94 ± 3.52 428 vs. 61.50 ± 9.89% activity suppression in AhR^{+/+} vs. AhR^{-/-} mice, respectively, 429 p<0.05) compared with the control group, indicating a reduced fear memory in 430 AhR^{-/-} mice, in agreement with previous evidence (Latchney et al., 2013). To 431 432 confirm our results we checked other types of hippocampus-dependent memory such as the novel object recognition (NOR) task that relies on mouse natural 433 exploratory behaviour (Ennaceur and Delacour, 1988). During the training 434 435 session (two similar objects), no preference was detected for one object over 436 the other in both genotypes (data not shown); however, during the test session, whereas WT mice presented a preferential exploration toward the new object, 437 AhR^{-/-} mice exhibited impaired NOR performance with a lack of net preference 438 for any of the objects (significant interaction between genotype and novel/old 439 object recognition two-way ANOVA; F_(1, 18)=7.46; p=0.0137; Fig. 1B). In 440 addition, we checked whether spatial memory was also altered in AhR^{-/-} mice. 441 442 First, mice were subjected to a novel object location task (NOL) (Antunes and Biala, 2012), in which WT mice spent more time investigating the new location 443 whereas, on the contrary, AhR^{-/-} mice explored similarly both locations 444

445	(significant interaction between genotype and novel/familiar location exploration
446	two-way ANOVA; $F_{(1, 42)}$ =7.93; p=0.0074; Fig. 1C). The impairment in spatial
447	memory shown by AhR ^{-/-} mice was corroborated by using the Y-maze and the
448	Barnes maze. In the first one, no differences were observed in spatial working
449	memory calculated either as percentage of spontaneous alternation (SAP) or as
450	arm entries (% SAP: 63.14 \pm 2.40 vs. 55.21 \pm 2.81 in AhR+/+ vs. AhR-/- mice,
451	respectively, p>0.05; arm entries: 26.43 \pm 2.86 vs. 24.40 \pm 5.39 in AhR^{+/+} vs.
452	AhR ^{-/-} mice, respectively p>0.05; Fig. 1D) but, relative to WT, AhR ^{-/-} mice did not
453	show any preference toward the novel arm (two-way ANOVA; $F_{(2, 27)}$ =3.84;
454	p=0.0342; Fig. 1D). In the Barnes maze (Fig. 1E), AhR KO mice and their WT
455	littermates spent similar time to find the escape box during all training sessions.
456	However, during the 24h-probe test, whereas WT animals spent most part of
457	the time in the target hole, $AhR^{-/-}$ ones did not (two-way ANOVA; $F_{(1.8, 19)}=260$;
458	p=0.0204; Fig. 1G). These results indicate impairment of the spatial memory
459	and support that AhR is required for a proper function of hippocampus-
460	dependent memory.

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462 Hippocampus-dependent memory deficits in AhR^{-/-} mice are independent 463 of the levels of hippocampal neurogenesis

AhR-dependent impairment of hippocampal function in AhR^{-/-} mice could be due to a decrease in the levels of hippocampal neurogenesis, consistent with previous studies of AhR in this setting performed at 3 months (P90) (Latchney et al., 2013). Interestingly, when animals were pulsed at different ages (p30, p60 and p100) with 4 consecutive BrdU injections (see Methods for details) at 2h-intervals and euthanized 24h after the last injection, higher levels of

proliferation at p30 and p60 but not at p100 were detected in AhR^{-/-} mice vs 470 their WT littermates (Fig. 2A). The results at p30 and p100 were confirmed 471 when we quantified the number of cells in cell cycle (using Ki67, marker of cells 472 in active proliferation) in the SGZ of WT and AhR^{-/-} mice (Fig. 2B-C). Since 473 memory studies had been performed at p60, deficits in proliferation are 474 therefore unlikely to account for hippocampal deficits in AhR KO mice, though 475 476 they may contribute to those reported previously at 3 months (P90; Latchney et al., 2013). Further confirming this finding, at p60. AhR^{-/-} mice displayed around 477 60% increase in the number of proliferative nestin⁺ precursors (nestin⁺/BrdU⁺) in 478 479 the DG when compared with WT ones, determined by flow cytometry 24h after BrdU administration (p<0.05; Fig. 2D-F). Furthermore, these data were 480 corroborated by analysing both type-1 (nestin⁺/GFAP⁺) and type-2 481 482 (nestin⁺/GFAP⁻) neural progenitor cells, as well as their respective proliferation rates at p30 and p100 (Fig. 2G-L). At p30, AhR^{-/-} mice presented significantly 483 484 higher levels of type-1 and -2 NPCs and its proliferative population than the control group but, even though at p100 the population of type-1 cells was similar 485 (p>0.05; Fig. 2G), type-2 and both type-1 and type-2 proliferative AhR^{-/-} NPCs 486 487 populations were dramatically decreased (p<0.05, Fig. 2G-J) compared to WT 488 populations.

To check whether the increase in SGZ proliferation induced by AhR absence was accompanied with a parallel increase in the number of adult newborn neurons, we first studied immunostaining of DCX, a marker of neuroblasts which is transiently expressed in new neurons. Supporting our previous data, p60 AhR^{-/-} mice displayed higher numbers of DCX⁺ cells, indicating immature neurons (p<0.05, Fig. 2M). In addition, we administered 495 BrdU during 5 days and, 28d later, we analysed the number of newborn mature neurons as BrdU⁺ cells expressing the mature neuronal marker calbindin. 496 Quantifications demonstrated that AhR^{-/-} mice present a significant increase in 497 the number of total BrdU⁺ cells and of double $BrdU^+/calbindin^+$ cells (p<0.05; 498 Fig. 2O), denoting that AhR absence leads to an increase in the number of 499 newly generated neurons. Furthermore, a comparable percentage of 500 BrdU⁺/calbindin⁺ cells was estimated for both groups (56% AhR^{-/-} vs. 55% WT), 501 suggesting that AhR deletion does not modify differentiation rate. 502

Similarly, neurogenesis enhancement attributable to the absence of AhR, 503 504 determined by quantification of $BrdU^+/nestin^+$ cells (4862 ± 561 vs. 8072 ± 851 cells in AhR^{+/+} vs. AhR^{-/-} mice, respectively, p<0.05, n=6), Ki67⁺ cells (276 \pm 29 505 vs. 430 \pm 23 cells in AhR^{+/+} vs. AhR^{-/-} mice, respectively, p<0.05, n=5) and 506 DCX⁺ volume (3707 ± 364 vs 6291 ± 545 μ m³ in AhR^{+/+} vs. AhR^{-/-} p60 mice, 507 respectively, p<0.05, n=5), was also observed in the other adult neurogenic 508 509 niche, the subventricular zone (SVZ). Thus, a common shared mechanism 510 underlying the actions of AhR in both neurogenic niches can be suggested.

Therefore, despite previously suggested role of AhR (Latchney et al., 2013), our data indicate that, at p30 and at p60, the absence of AhR enhances or does not affect hippocampal neurogenesis, thus discarding that hippocampus-dependent deficits observed in AhR^{-/-} mice at p60 are due to decreased levels of adult neurogenesis.

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517 Absence of AhR is associated to aberrant morphology of hippocampal 518 granule cells

519 A high rate of adult hippocampal neurogenesis has been previously demonstrated to correlate positively with increased learning and memory ability 520 (Kempermann et al., 1997; van Praag et al., 2002). Given that AhR^{-/-} mice 521 522 exhibit impaired memory when hippocampal neurogenesis is still enhanced and considering the high AhR mRNA expression in the DG granular layer (Kimura 523 and Tohyama, 2017) and the reported role of AhR orthologues in the control of 524 525 neuronal growth and dendritic arborisation in invertebrates (Crews and Brenman, 2006; Kim et al., 2006; Qin and Powell-Coffman, 2004; Smith et al., 526 2013), we hypothesised that defects observed when AhR is deleted could be 527 528 due to variations in the morphology of granule neurons and its subsequent functional activity. Therefore, we examined granule neuronal morphology in 529 hippocampi of WT and AhR^{-/-} p60 mice. First, dendritic branching of DG granule 530 531 neurons was quantified by Sholl analysis of Golgi-Cox stained sections (Fig. 3A-C). Relative to WT, AhR^{-/-} mice displayed a higher number of dendritic branches 532 533 close to the soma (40-70 µm from soma; p<0.05; Fig. 3A), increased total 534 dendritic length (p<0.05; Fig. 3B), and a larger number of dendritic spines 535 (p<0.05; Fig. 3C), supporting our hypothesis that the absence of AhR is 536 associated to an altered dendritic structure of granule neurons.

Interestingly, similar changes were found in immature neurons (DCX⁺ neuroblasts) in WT and AhR^{-/-} p60 mice, indicating that AhR effects on granule neuronal morphology seem to be also apparent in adult-born granule neurons. Interestingly, quantification of DCX⁺ labelling distribution between total (GL+ML), granular (GL) and molecular layers (ML) revealed significant differences in the arborisation pattern of both genotypes: compared with WT subjects, AhR^{-/-} mice showed increased DCX⁺ staining in GL while no

544 differences were found in the ML (p<0.05; Fig. 3D). In addition, immature AhR 545 KO cells also presented a significant reduction in the apical dendrite length (p<0.05; Fig. 3E). To confirm that the absence of AhR affects the morphology of 546 adult newborn neurons, high titres of CAG-GFP retrovirus were delivered into 547 the hilar region of WT and AhR^{-/-} KO p60 mice to selectively target proliferating 548 neuronal progenitors in vivo (Zhao et al., 2006). Twenty-eight days post-549 550 infection, Sholl analysis of GFP newborn neurons in the DG showed an increased dendritic branching in the proximal segment close to the soma of 551 AhR^{-/-} neurons compared to those of WT (50-150 μ m from the soma; p<0.05; 552 553 Fig. 3I). While total dendritic length was significantly greater in AhR^{-/-}/GFP⁺ neurons (p<0.05; Fig. 3J) indicating a higher pattern of branching, the apical 554 dendrite length was shorter than in control/GFP⁺ cells (p<0.05; Fig. 3K). Taking 555 556 all these results together, it can be concluded that the absence of AhR promotes retraction of the apical dendrite and altered branching of the dendritic 557 558 tree. The aberrant morphology of granular neurons was additionally supported by the finding of a higher dendritic spine density in AhR^{-/-}/GFP⁺ neurons 559 compared to those of WT mice in both, proximal and distal segments of 560 561 dendritic branches in the molecular layer (p<0.05; Fig. 3L). Despite this, the 562 number of mushroom spines in both segments, typically more abundant in mature neurons (Zhao et al., 2006), was significantly lower in AhR^{-/-}/GFP⁺ cells 563 than in AhR^{+/+}/GFP⁺ neurons along the dendrites in the molecular layer (p<0.05; 564 565 Fig. 3M), thus suggesting a more immature phenotype in the absence of AhR. 566

567 The lack of AhR alters granule cells intrinsic excitability, synapses 568 maturation and the correct function of the hippocampus

569 Morphological changes resulting from AhR ablation can lead to alterations of the hippocampal physiological properties. This possibility was 570 investigated by performing electrophysiological analysis of granule cells in acute 571 slices from AhR^{-/-} and control mice (Fig. 4A). Previously, the absence of 572 significant differences between groups in resting membrane potential and 573 membrane resistance was verified (Fig. 4B-C). Then, the ability of AhR^{-/-} 574 575 granule cells to fire repetitive action potentials, a hallmark of neuronal maturation (Deng et al., 2010), was assessed. Under the whole-cell current-576 clamp, AhR^{-/-} granule cells showed an increased firing rate in response to a 577 578 depolarizing current injection (40-190 pA; p<0.05; Fig. 4D-E), consistent with immature neuronal excitability (Dieni et al., 2016). It is known that the ratio of 579 active/silent synapses is changing over neuron maturation as a result of 580 581 changes in the content of glutamate AMPA receptors (AMPARs) (Carlisle and 582 Kennedy, 2005; Chater and Goda, 2014; Paoletti et al., 2013; Tada and Sheng, 583 2006). In order to assess functional AMPAR content, we quantified the 584 AMPAR/NMDAR ratio by comparing AMPA excitatory postsynaptic currents (EPSCs) at -70mV and glutamate NMDA receptors (NMDARs) EPSCs at 585 +40mV in WT and AhR^{-/-} granule cells. We found a significantly lower 586 AMPAR/NMDAR ratio in AhR^{-/-} granule cells, indicating a lower proportion of 587 AMPAR to NMDAR on their granule cell dendrites (p<0.05; Fig. 4F-G), thus 588 suggesting that AhR^{-/-} granule cells have a more immature phenotype than 589 590 those from WT. Finally, we studied the pair-pulse ratio (PPR) as a 591 hippocampus-dependent function measurement for integration of inputs coming from the entorhinal cortex to the hippocampal granule cells. The PPR 592 (EPSC2/EPSC1) was significantly higher in AhR^{-/-} KO compared with control 593

mice (p<0.05; Fig. 4H), indicating that the initial probability of release is lower in $AhR^{-/-}$ KO mice.

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597 Absence of AhR in neural progenitor cells is enough to promote 598 hippocampal deficits

In order to check the specific role of AhR in adult-born granule neurons, 599 600 we used a transgenic strategy to conditionally delete AhR from neural progenitor cells in adult mice (AhR-icKO mice; Fig. 5A). For such purpose we 601 crossed mice expressing tamoxifen (TAM)-inducible Cre-recombinase driven by 602 a progenitor specific promoter (nestin-Cre^{ERT2} mice) (Imayoshi et al., 2008) with 603 mice in which AhR is floxed (AhR^{f/f}) (Walisser et al., 2005). In double mutant 604 offspring from this cross (AhR-icKO), TAM treatment (Fig. 5A, bottom) induced 605 606 the excision of the exon 2 flanked sequences and the deletion of AhR in neural progenitor cells and their progeny resulted in a marked decrease of AhR 607 608 expression in the DG determined by immunofluorescence (Fig. 5B). 609 Examination of 3-month-old animals after tamoxifen administration revealed a 610 higher number of DCX⁺ cells in the SGZ of the AhR-icKO mice than in that of control mice (p<0.05; Fig. 5C) although not significant differences were found 611 for Ki67⁺ cells (29.28 ± 2.93 vs. 33.47 ± 1.86 in AhR^{+/+} vs. AhR^{-/-} mice, 612 respectively, p>0.05). Consistent with these data, AhR-icKO mice treated with 613 TAM presented an increased DCX staining in total (GL+ML) and GL but not in 614 the ML (p<0.05; Fig. 5D), similar to AhR^{-/-} mice. In order to study whether 615 616 morphological alterations present in newborn neurons were affecting cognitive ability, memory tests were carried out. First, we analysed episodic memory in 617 618 CFC test using a weak fear conditioning protocol based on one-foot shock

619 stimuli in order to detect subtle differences which are not masked by generalisation (Lonsdorf et al., 2017). Remarkably, in agreement with our 620 results in AhR^{-/-} mice, AhR-icKO mice displayed comparable cognitive 621 hippocampal deficits in CFC, showing a reduced freezing response compared 622 to AhR^{f/f} (p<0.05; Fig. 5E). Similarly, in the Y-maze test, only AhR^{f/f} mice 623 showed a preference towards the closed arm (significant interaction between 624 genotype and closed arm two-way ANOVA; F_(2, 57)=5,91; p=0.0047; Fig. 5F). 625 Finally, NOL test revealed that the specific ablation of AhR in adult 626 neuroprogenitor cells in AhR-icKO mice pattern separation skills is altered, 627 showing a worst discrimination efficiency than AhR^{f/f} mice, not demonstrating 628 any kind of preference for the new location (Fig. 5G). 629

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Acute conditional deletion of AhR in neuroprogenitor cells is enough to alter granule cell intrinsic excitability, synapses maturation and the correct function of the hippocampus

634 To examine whether the functional properties of the aberrant newborn neurons might underlie hippocampus-dependent memory impairments 635 636 observed in our AhR-icKO, we acutely ablated AhR in adult neuroprogenitor cells by treating p21 AhR^{f/f} and AhR-icKO mice with tamoxifen, and 637 electrophysiological studies were performed 4 weeks after the last tamoxifen 638 639 administration (Fig. 6A). We did not see any differences between groups in 640 resting potential and membrane resistance (Fig. 6B-C). When we analysed the intrinsic excitability of AhR^{f/f} and AhR-icKO granule neurons measured as the 641 ability to spike action potentials by stimulating them with increased electrical 642 643 depolarization currents, we found that acute ablation of AhR in adult

644 neuroprogenitor cells led to a significant increase in the intrinsic excitability of 645 AhR-icKO granule neurons (p<0.05; Fig. 6D-E), consistently with our previous 646 results in constitutive AhR KO mice. Given the importance of AMPA receptors in maturation and functionality of the synapses, AMPAR and NMDAR-mediated 647 currents were measured in order to explore the ability to integrate and respond 648 to the inputs coming from the entorhinal cortex into the hippocampus by the 649 aranule neurons. Confirming our results in AhR-/-, AhR-icKO granule neurons 650 displayed a reduction in AMPA/NMDA ratio indicating that the specific absence 651 of AhR in newborn granule neurons impairs a correct dendritic spine maturation 652 653 (p<0.05; Fig. 6F-G). Finally, we studied the PPR in these mice to explore the connectivity between EC and DG. According with our previous results in AhR 654 KO mice, AhR-icKO granule neurons showed significant higher PPR 655 (EPSC2/EPSC1) compared with AhR^{f/f} cells (Fig. 6H), meaning an initial lower 656 neurotransmitter probability of release when AhR is acutely ablate. 657

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659 DISCUSSION

We have investigated the role of the transcription factor AhR in hippocampal function and granule neuronal morphology in adult mice brain. Our data demonstrate that AhR absence is associated to a severe impairment of hippocampal-dependent memory, concomitantly to an increased dendrite arborisation pattern and a decreased dendritic spine maturity of hippocampal granule neurons, as well as to aberrant electrophysiological properties and functions of these cells.

667 AhR (Aryl hydrocarbon Receptor) is a basic helix-loop-helix-PER-ARNT-668 SIM (bHLH-PAS) transcription factor that mediates the toxic and carcinogenic

669 effects of xenobiotics. Interestingly, AhR is widely expressed in the CNS and, in 670 this context, its physiological and pathological roles are just beginning to be 671 unravelled. Specifically, AhR mRNA is enriched in the dentate gyrus granule 672 cells of the adult hippocampus (Kimura and Tohyama, 2017), a crucial structure for high cognition tasks, such as episodic, spatial learning and memory, which 673 are often disrupted in neurological disorders. AhR is the only PAS member 674 675 known to be activated by endogenous or exogenous ligands (Boitano et al., 2010; Cuartero et al., 2014; Mandal, 2005; Mimura and Fujii-Kuriyama, 2003; 676 Mimura et al., 1997; Nguyen and Bradfield, 2008). The potential therapeutic 677 678 implications derived from this fact prompted us to study its role in hippocampal function. Here, we demonstrate that the absence of AhR affects different types 679 of hippocampal-dependent tasks such as episodic (NOR and CFC) and spatial 680 681 memories (Y-maze, NOL and Barnes maze).

Hippocampal newborn neurons have been implicated in the acquisition 682 683 and recall of hippocampus-dependent memories (Aimone et al., 2014; Anacker 684 and Hen, 2017) and, therefore, adult hippocampal neurogenesis is an essential process for cognitive function (Kempermann et al., 1997; van Praag et al., 1999; 685 686 van Praag et al., 2002; Zhao et al., 2006). The reduced NPC proliferation in the 687 SGZ in parallel to deficits in fear conditioning memory tests which has been described in 12-weeks old AhR-deficient mice (Latchney et al., 2013) could 688 689 therefore explain the AhR-dependent impairment of episodic and spatial 690 hippocampus-dependent memory mice that we observe in 2-month old (p60) AhR^{-/-}. However, on the contrary, we have found that 2-month old (p60) AhR^{-/-} 691 mice display enhanced SGZ neurogenesis, as shown by a higher number of 692 693 newborn NPCs, neuroblasts and fully integrated mature neurons, thus discarding that hippocampus-dependent memory deficits are due to decreased adult neurogenesis. Indeed, we have found that at both p30 and p60, the absence of AhR increases SGZ NPCs proliferation. However, at p100, the process is reversed, with a decrease in NPCs proliferation, similarly to the reported results at 12-weeks old (Latchney et al., 2013). It is plausible that niche depletion induced by the enhanced NPCs proliferation at earlier times accounts for the decreased proliferation at later times in AhR^{-/-} mice. Several studies have reported an increased proliferative response, so-called compensatory proliferation, that takes place in order to counteract an excess of newborn cell death (rev. in Yamaguchi and Miura, 2015). Although the parallel increase in the number of BrdU⁺/calbindin⁺ cells suggests that this mechanism is not taking place in AhR^{-/-} mice, additional studies assessing precursor apoptosis and survival would be required to define the outcomes of excess proliferation in the absence of AhR.

In any case, as discussed above, decreased proliferation is not responsible for hippocampal memory deficits in 2-month old mice, supporting that other mechanisms are involved.

711 AhR is a highly-conserved protein from invertebrates to mammals. 712 Although in mammals AhR has been traditionally known to participate in the xenobiotic metabolism of toxic compounds like dioxins (Barouki et al., 2007; 713 714 Fernandez-Salguero et al., 1996; Fernandez-Salguero et al., 1997; Fujii-715 Kuriyama and Kawajiri, 2010; Fujii-Kuriyama and Mimura, 2005), invertebrate 716 AhR orthologues do not have a toxic response to dioxin, neither have dioxin binding capacity, what suggests another ancestral role for this receptor that 717 718 could have remained throughout the evolution. In fact, previous works in C.

719 elegans (Huang et al., 2004; Qin and Powell-Coffman, 2004; Smith et al., 2013) 720 and Drosophila (Crews and Brenman, 2006; Kim et al., 2006) identified AhR orthologues as regulators of dendrite branching in different types of neurons. 721 More recently, in mammals, AhR activation has been reported to disrupt 722 migration and dendritic growth of olfactory interneurons and hippocampal CA1 723 neurons, respectively, in the mouse brain (Kimura et al., 2016; Kimura et al., 724 725 2017). However, the specific role of AhR in dendrite morphology and functional activity of DG granule neurons in the adult murine brain is not known. Our 726 studies using Golgi-Cox staining or doublecortin immunofluorescence show 727 728 altered dendritic structure of granule neurons in the absence of AhR. GFP labelling confirmed that AhR absence affects the morphology of adult newborn 729 neurons, an effect that is likely contributing to the memory deficits exhibited by 730 these mice. AhR^{-/-} newborn neurons show an altered morphology characterised 731 732 by a shorter apical dendrite and a profuse dendritic branching close to the 733 soma. Since ectopic migration was not observed, the shortening of the apical dendrite in AhR^{-/-} cannot be ascribed to an ectopic pattern of migration in the 734 GL. All these changes could affect the synaptic partners of these neurons, with 735 detrimental consequences for hippocampal-dependent behaviour, and suggest 736 that modifications of AhR function might underlie some pathological situations 737 inducing aberrant granule neuronal morphology. 738

Regarding the morphology of dendritic spines, whereas filopodia and stubby spines are often associated with immature neurons, thin and mushroom spines are more abundant in mature neurons (Carlisle and Kennedy, 2005; Spruston and Johnston, 2008; Zhao et al., 2006). Of note, GFP labelling revealed that AhR^{-/-} granule neurons displayed a higher spine density but a much lower abundance of mature mushroom spines. Thus, reduced dendritic
mushroom spine density would further alter excitatory inputs and the number of
synaptic inputs and integration (Spruston and Johnston, 2008).

747 Consistently, AhR loss-of-function was associated to an increased intrinsic excitability which could suggest a more immature phenotype (Dieni et 748 al., 2016; Lopez-Rojas and Kreutz, 2016). Previous studies have reported that 749 750 neuronal membrane resistance as well action potential firing rate decrease along neuronal maturation (Mongiat et al., 2009; Dieni et al., 2016), resulting in 751 an adequate integration of the inputs coming from the entorhinal cortex. These 752 753 changes are critical factors that contribute to learning and memory in the hippocampus. Our results suggest that AhR could be necessary for this 754 electrophysiological change. We also found a lower AMPAR/NMDAR ratio in 755 granule neurons from AhR^{-/-} mice, probably indicating a selective depression in 756 757 AMPAR synaptic responses. During physiological granule cell maturation, there 758 is an increase in the number of mature dendritic spines due to a progressive 759 incorporation of AMPARs into the synaptic sites (Bassani et al., 2013). As 760 commented above, AhR deficiency was linked to a higher proportion of 761 immature spines albeit an increased spine density in mature granule cells. Our 762 results support the idea that AhR is necessary for this maturation, very likely by modulating the levels of AMPAR and NMDAR in the dendritic spines. In 763 addition, the PPR (EPSC2/EPSC1) was significantly higher in AhR^{-/-} 764 765 hippocampal slices compared with those from wild-type mice, strongly 766 suggesting that the initial probability of neurotransmitter release in the terminals coming from the entorhinal cortex to the dentate gyrus is reduced in AhR^{-/-} mice 767 768 (Fioravante and Regehr, 2011). This could suggest an immature phenotype of granule cell dendritic spines that affects synapses in a retrograde fashion, byaltering neurotransmitter release from perforant pathway terminals.

771 In contrast with our results, increased rates of neurogenesis and/or high 772 excitability on immature newborn granule cells are two features that have been associated to the role of these cells in learning and memory (Lopez-Rojas and 773 Kreutz, 2016). However, several pieces of evidence in the literature also 774 775 support that increased neurogenesis may not always result in improved 776 function. For instance, manipulations that increase neurogenesis may have positive effects on anterograde memories but not in retrograde memories 777 778 (Akers et al., 2014; Frankland et al., 2013). In agreement with our results, pathological situations that impair hippocampal function such as epilepsy (Cho 779 et al., 2015; Zhao et al., 2008) or stroke (Niv et al., 2012; Woitke et al., 2017) 780 781 trigger an augmented hippocampal neurogenic response which is accompanied by aberrant features of newborn neurons. Of note, these aberrant neurons show 782 a retraction in apical dendrite and an increased aberrant dendrite branching, a 783 784 phenotype which is considered an immature feature (Woitke et al., 2017) and clearly resembles the one that we have observed in AhR^{-/-} mice. This altered 785 786 dendritic morphology may, on its turn, underlie the "immature" 787 electrophysiological features described. As commented above, an interesting possibility suggested by our study is that AhR controls dendritic spine formation 788 789 and maturation, very likely by regulation of the expression and/or the post-790 translational trafficking of the AMPA receptor, a hypothesis that remains to be 791 studied. A sustained immature status due to the lack of proper dendritic maturation could trigger, as compensatory mechanisms, an 792 intrinsic 793 hyperexcitability and a subsequent lower initial probability of release, that will translate into an increased PPR. Although the underlying mechanisms are likely to differ depending on each pathophysiological circumstance, it has been described that animal paradigms of Alzheimer's disease (AD) show aberrant increases in network excitability in the dentate gyrus hippocampus that may contribute to the neurological deficits shown by these models (Frazzini et al., 2016; Hazra et al., 2013; Palop et al., 2007).

800 To further confirm the role of AhR in hippocampus-dependent memory 801 through the modulation of neurogenesis and granule neurons maturation and 802 activity, we analysed those mechanisms in our specific neuroprogenitors AhR-803 ablated mice. Of note, we evaluated SGZ NPCs proliferation using animals with specific AhR deletion in nestin-expressing cells. Our data showing higher 804 numbers of immature neurons in the SGZ of AhR icKO mice, and altered 805 806 dendritic arborisation and electrophysiological properties, similar to those of AhR-/- mice confirm our previous data and allow us to discard indirect effects 807 arising from the total lack of AhR in AhR^{-/-} mice. Besides, similar cognitive 808 809 impairment was observed after the specific deletion of AhR in neuroprogenitor 810 cells. First, by using a weak protocol in the CFC test, our data showed that the 811 absence of AhR in newborn neurons is enough to disturb episodic memory. 812 Furthermore, similar memory impairment was observed in AhR-icKO mice when they were tested on the Y-maze, NOR and NOL tests, supporting that alteration 813 814 of the AhR pathway in newborn granule cells, or even changes in the availability 815 of endogenous or exogenous AhR activators, not only may affect their 816 morphological properties, but also could have detrimental repercussions in the function of the hippocampus. 817

818 As several other members of the bHLH superfamily and consistent with 819 its dynamic pattern expression in the embryonic and postnatal mouse brain at different time points (Kimura & Tohyama, 2017), brain AhR is likely to play both 820 time- and region-specific functions. In agreement with this, it has been shown 821 822 that AhR activity regulates cerebellar granule neuron number and 823 differentiation, possibly by coordinating granule neuron precursor 824 developmental transition (Dever et al., 2016). Although with different features, 825 our longitudinal study illustrates how hippocampal neurogenesis varies along time from p30 to p100, strongly supporting a role of AhR in the control of 826 827 proliferation and pool maintenance of adult-hippocampal neural stem cells in an 828 age-dependent way.

In addition, several studies report the involvement of AhR in the migration 829 830 and dendritogenesis of olfactory interneurons and of cortical piramidal neurons during development (Kimura et al., 2016, 2017), as well as in neuronal 831 differentiation in adult mice (Dever et al., 2016). In line with these results, in the 832 833 present study our data demonstrate the role of AhR as a key modulator of newborn hippocampal granule neuron morphology, including dendritic spine 834 835 growth and maturation, which are essential for hippocampal function, thus 836 supporting AhR as a main player in neuronal maturation/differentiation. Hence, our study expands the biological implications of AhR receptor in such a relevant 837 838 function showing that AhR plays time-specific functions in the regulation of 839 neurogenesis and granule neuron maturation in hippocampus. Interestingly, 840 several studies have demonstrated that dioxin exposure interferes with developmental neurogenesis (Latchney et al., 2013; Williamson et al., 2005) so 841

abnormal modulation of AhR by ligands such as environmental pollutants
throughout life might lead to hippocampus-related dysfunctions.

Summing up, deletion of AhR induces a severe impairment of 844 845 hippocampal-dependent memory, concomitantly to a high dendrite arborisation pattern in newborn hippocampal neurons, increased density of dendritic spines 846 but a reduction in mature mushroom spines, which drive a phenotype showing 847 848 aberrant electrophysiological properties and function of these cells. Our study provides evidence indicating that AhR is a regulator of dendrite arborisation and 849 functional activity of adult hippocampal newborn neurons and showing its critical 850 851 role for learning and memory function. An additional novel line of research opened by our study is the possible implication of AhR, as a new potent 852 druggable target, in disorders in which cognitive deficits are accompanied by 853 854 hippocampal morphological alterations, such as epilepsy, schizophrenia or stroke. 855

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1044 FIGURE LEGENDS

Figure 1. Ablation of AhR impairs hippocampus-dependent memory. (A) 1045 Percentage of freezing in the CFC task for AhR^{+/+} and AhR^{-/-} mice 1h (left) and 1046 24h (right) after the foot shocks (*p<0.05 vs. AhR^{+/+}; n=8 AhR^{+/+} and 7 AhR^{-/-} 1047 animals/group). (B) Percentage of exploration time between familiar and new 1048 object in the NOR test for AhR^{+/+} and AhR^{-/-} mice 6h after training. Two-way 1049 1050 ANOVA demonstrated a significant interaction between the object and genotype $[F_{(1, 18)}=7,46; p<0.05]$ (*p<0.05 vs. AhR^{+/+}; n=6 AhR^{+/+} and 5 AhR^{-/-} 1051 animals/group). (C) Percentage of exploration time between old and new object 1052 location in the NOL test for AhR^{+/+} and AhR^{-/-} mice 6h after training. Two-way 1053 ANOVA demonstrated a significant interaction between the object and genotype 1054 [F_(1, 42)=7.93; p<0.05]; (*p<0.05 vs. AhR^{+/+}; n=12 AhR^{+/+} and 11 AhR^{-/-} 1055 1056 animals/group). (D) Percentage of time spent in each arm in the Y-maze test for AhR^{+/+} and AhR^{-/-} mice 6h after training. Two-way ANOVA showed a significant 1057 interaction between the arm/genotype [F_(2, 27)=3.84; p<0.05] (n=6 1058 animals/group). (E) Time spent to find the escape box during the Barnes maze 1059 training sessions in both AhR+/+ and AhR-/- mice. Two-way ANOVA 1060 demonstrated a significant effect during training sessions [F_(5, 65)=5.04; p<0.05] 1061 (n=7-8 animals/group). (F) Density plots for grouped data showing where the 1062 AhR^{+/+} and AhR^{-/-} mice concentrated their searches during retention test day. 1063 1064 (G) Percentage of time spent around each hole (s) in the Barnes maze platform for AhR^{+/+} and AhR^{-/-} mice during the retention test day. Two-way ANOVA 1065 demonstrated a significant interaction between the holes and genotype [F(19, 1066 ₂₆₀₎=1.83; p<0.05] (*p<0.05 vs. AhR^{+/+}; n=7-8 animals/group). Data are mean ± 1067 SEM. Data were compared by using non-parametric 2-tailed Mann-Whitney test 1068

1069 (A), 2-way ANOVA (B-G) or a nonparametric 2-way ANOVA followed by1070 Bonferroni post hoc testing (B-E).

1071

Figure 2. The absence of AhR exacerbates adult hippocampal 1072 neurogenesis. (A) BrdU⁺ cells in the DG of AhR^{+/+} and AhR^{-/-} mice measured 1073 at p30, p60 and p100 24h after the last BrdU injection. Two-way ANOVA 1074 demonstrated a significant interaction between the age and genotype $[F_{(2)}]$ 1075 24)=9.51; p<0.05] (*p<0.05 vs. AhR^{+/+}; n=5 animals/group), (B-C) Ki67⁺ cells in 1076 the DG of $AhR^{+/+}$ and $AhR^{-/-}$ mice measured at p30, p60 and p100. Two-way 1077 ANOVA demonstrated a significant interaction between the age and genotype 1078 $[F_{(2.28)}=8.56; p<0.05]$ (*p<0.05 vs. AhR^{+/+}; n=5-6 animals/group). Representative 1079 images of Ki67⁺ cells in AhR^{+/+} and AhR^{-/-} mice at different time points are 1080 1081 shown in (C). (D-F) Quantification of the number of nestin/BrdU⁺ cells by flow cytometry 24h after BrdU administration. Data are expressed as the percentage 1082 of control group (*p<0.05 vs. AhR^{+/+}; n=6 animals/group) (D). Representative 1083 dot plots of double-stained cells for nestin and BrdU in the DG of AhR^{+/+} (E) and 1084 AhR^{-/-} (F) mice. (G-L) Characterization of type-1 (nestin⁺/GFAP⁺) (G) and type-2 1085 progenitors (nestin⁺/GFAP⁻) (I) and their proliferative capacity (BrdU⁺) (H and J) 1086 in the DG of AhR^{+/+} and AhR^{-/-} mice determined at p30 and p100. 1087 Representative images of nestin (red), GFAP (green), BrdU (grey) of DG of 1088 AhR^{+/+} and AhR^{-/-} mice at p30 (K) and p100 (L) (*p<0.05 vs AhR^{+/+}; #p<0.05 vs 1089 AhR^{-/-}; n=4-6 animals/group). (M-N) Quantification of the number of DCX⁺ cells 1090 in the DG of WT and AhR^{-/-} mice (*p<0.05 vs. AhR^{+/+}; n=7-8 animals/group). 1091 Representative images of DCX⁺ cells in AhR^{+/+} and AhR^{-/-} mice are shown in 1092 (N). (O-P) Quantification of the number of BrdU⁺ cells (left) and newborn 1093

1094 integrated neurons (BrdU⁺/Calbindin⁺) (right) determined 28 days after BrdU administration (*p<0.05 vs. AhR+/+; n=7-9 animals/group). Representative co-1095 localisation images for BrdU and calbindin in AhR^{+/+} and AhR^{-/-} mice are shown 1096 in (P). Insets display high magnifications images. Numbers of cells are 1097 expressed per 1000 μ m². Data are mean \pm SEM. Scale bar is 50 μ m in C and K-1098 L and J. 70um in N. 30um in P. Data were compared by using non-parametric 1099 2-tailed Mann-Whitney test in D, M and O or a non-parametric 2-way ANOVA 1100 followed by Bonferroni post hoc testing (A, B, G-J). 1101

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Figure 3. The absence of AhR alters granule cells morphology and 1103 dendritic spine density and maturation. (A) Sholl analysis of granule 1104 dendritic branching of Golgi-Cox stained DG from AhR^{+/+} and AhR^{-/-} p60 mice. 1105 1106 Two-way ANOVA demonstrated а significant interaction between distance/genotype [F (24, 125)=2.45; p<0.05]; (*p<0.05 vs. AhR^{+/+}) (n=23 WT and 1107 37 AhR KO neurons from 3-4 animals/group). Representative reconstructions of 1108 Golgi-Cox stained neurons are shown for AhR^{+/+} (left) and AhR^{-/-} mice (right). 1109 (B) Total dendritic length of Golgi-Cox-stained granular cells (*p<0.05 vs. AhR). 1110 (C) Quantification of spine density in Golgi-Cox stained neurons (*p<0.05 vs. 1111 AhR^{+/+}: n=20-30 dendrite segments from 3-4 animals/group). (D) Densitometric 1112 analysis of dendrite DCX⁺ labelling distribution in the DG of WT and AhR^{-/-} mice 1113 Data is displayed as the DCX⁺ integrated density found in total 1114 at p60. (GL+ML), GL or ML normalized by values got from soma. (E) Quantification of 1115 1116 apical neuroblast length. Representative images of DCX⁺ labelling distribution in AhR^{+/+} and AhR^{-/-} mice are shown in (F). (*p<0.05 vs. AhR^{+/+}; n=5-6 1117 animals/group). (G-N) GFP-retroviral infection of newborn neurons. Schematic 1118

protocol followed for CAG-GFP retrovirus infusion in AhR^{+/+} and AhR^{-/-} p60 mice 1119 (G). Representative GFP newborn AhR^{+/+} (left) and AhR^{-/-} neurons (right) are 1120 shown in (H). Two-way ANOVA of Sholl analysis in GFP-labelled dentate 1121 granule cells demonstrates a significant interaction between the distance and 1122 genotype [F_(4,30)=3.02; p<0.05]; (*p<0.05 vs. AhR^{+/+}; n=19 WT and 28 AhR KO 1123 neurons from 4 animals/group; I). Quantification of total dendritic length (J) and 1124 1125 apical dendritic length (K). Density of dendritic (L) and mushroom spines (M) in proximal and distal dendritic segments in the molecular laver of AhR^{+/+}/GFP⁺ 1126 and AhR^{-/-}/GFP⁺ granule cells 4 weeks post-infection (*p<0.05 vs. AhR^{+/+}/GFP⁺; 1127 n=26 and 22 segments from 4 animals/group). Representative images of GFP-1128 labelled spines in both proximal and distal segments of AhR^{+/+}/GFP⁺ and AhR^{-/-} 1129 /GFP⁺ granule cells (N). Insets display high magnifications images. Data are 1130 mean ± SEM. Data were compared by using non-parametric Mann-Whitney 1131 tests in B-C, D-E and J-M or a non-parametric 2-way ANOVA followed by 1132 Bonferroni post hoc testing (A and I). Scale bar is 70µm in F and H, and 2µm in 1133 О. 1134

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Figure 4. AhR deletion alters the synaptic properties of dentate gyrus 1136 1137 granule cells. (A) Schematic representation of a hippocampal slice showing 1138 stimulating and recording electrode sites. (B-C) The resting membrane potential (B) and the current-voltage relationship (C) were not significantly different 1139 between DG granule cells from AhR^{+/+} and AhR^{-/-} mice (n=4 animals/group). (D-1140 1141 E) DG granule cells firing rate is significantly increased by AhR deletion (*p<0.05 vs. AhR^{+/+}). Representative sample traces (D) and averaged values 1142 (E) in response to increasing depolarising currents. (F-G) AMPA/NMDA ratio is 1143

decreased by AhR deletion (*p<0.05 vs. AhR^{+/+}). Representative traces (F) and 1144 averaged values (G) of NMDA- and AMPA-mediated EPSCs recorded at +40 1145 and -60 mV, respectively. (H) Averaged values showing a significant increase in 1146 the PPR at interstimulus intervals of 75, 100, 150 and 200 ms in cells lacking 1147 AhR (*p<0.05 vs. AhR^{+/+}. EPSC sample traces represent the mean of 20 1148 consecutive EPSCs at 0.33Hz. Data are mean \pm SEM (14 neurons from 8 slices 1149 from n=4 AhR^{+/+} mice and 12 neurons from 8 slices from n=4 AhR^{-/-} mice). Data 1150 were compared by using non-parametric 2-tailed Mann-Whitney tests. 1151

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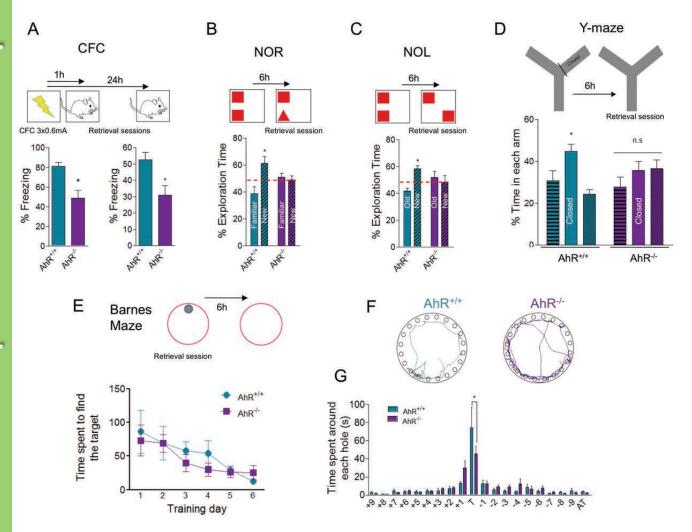
Figure 5. Acute ablation of AhR in adult neural precursors impairs 1153 hippocampus-dependent memory by promoting aberrant immature 1154 neurons. (A) Schematic diagram of the strategy for conditional deletion of AhR 1155 nestin-Cre^{ERT2+/}AhR^{f/f} mice (AhR-icKO). AhR^{f/f} and NPCs in nestinin 1156 Cre^{ERT2+/}AhR^{f/f} mice were administered two rounds of tamoxifen (TAM; at p30 1157 and p60) at a dose of 180mg/Kg. (B) AhR immunostaining in AhR^{1/f} (left) and 1158 AhR-icKO (right) 3 weeks after the second round of TAM 1159 injection. (C) Quantification of the number of DCX⁺ cells per 1000 μ m². in the 1160 DG of AhR^{f/f} and AhR-icKO mice 3 weeks after the last TAM injection (*p<0.05 1161 vs. AhR^{f/f;} n=8-8 animals/group) (left). Representative images of DCX⁺ staining 1162 in AhR^{f/f} and AhR-icKO mice (right). (D) Densitometric analysis of dendrite 1163 DCX⁺ labelling distribution in the DG of AhR^{f/f} and AhR-icKO mice 3 weeks after 1164 the last TAM injection. Data is displayed as the DCX⁺ integrated density found 1165 in total (GL+ML), GL or ML normalized by values got from soma (*p<0.05 vs. 1166 AhR^{f/f}; n=6 AhR^{f/f} and 5 AhR-icKO animals/group) (left). Representative images 1167 of the arborisation of immature newborn cells in AhR^{t/f} and AhR-icKO mice 1168

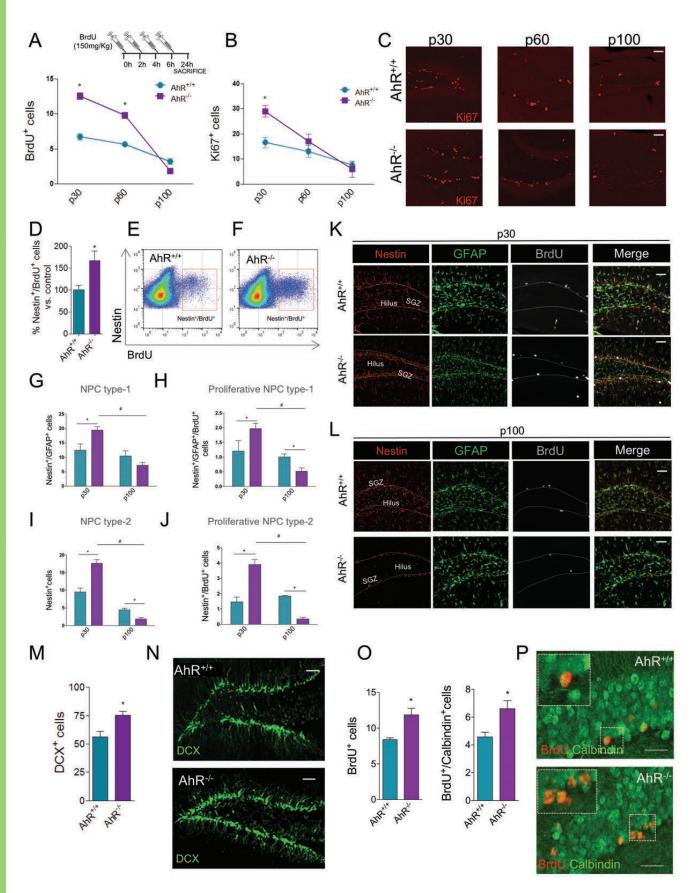
1169 (right). (E) Protocol followed for a weak contextual fear conditioning paradigm (0.4mA x1) in AhR^{f/f} and AhR-icKO treated with tamoxifen. Retrieval was 1170 performed 24h after training. (*p<0.05 vs AhRf/f; n=10-11 animals/group). (F) 1171 Percentage of time spent in each arm in the Y-maze test for AhR^{f/f} and AhR-1172 icKO mice 3 weeks after the last TAM injection 6h after training. Two-way 1173 ANOVA demonstrated a significant interaction between the arm and genotype 1174 [F_(2, 57)=5.91; p<0.05] (n=10-11 animals/group). (G) Percentage of exploration 1175 time between old and new object location in the NOL test for AhR^{f/f} and AhR-1176 icKO mice 3 weeks after the last TAM injection 6h after training (n=10-11 1177 animals/group). Data are mean \pm SEM. Scale bar is 50µm in B, C and D. Data 1178 were compared by using non-parametric 2-tailed Mann-Whitney test (C-E), or a 1179 non-parametric 2-way ANOVA followed by Bonferroni post hoc testing (F-G). 1180

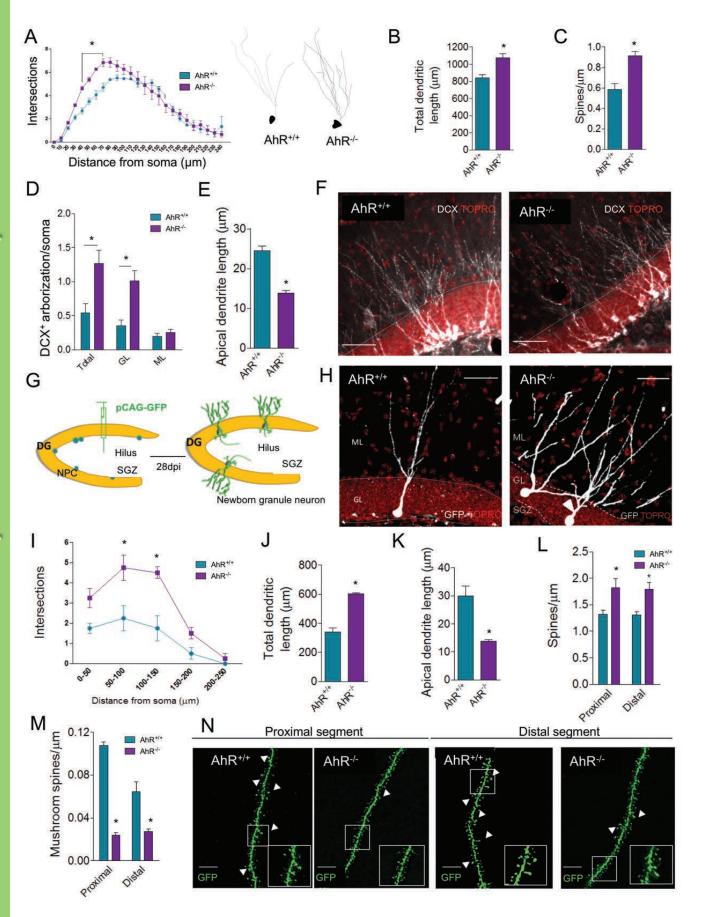
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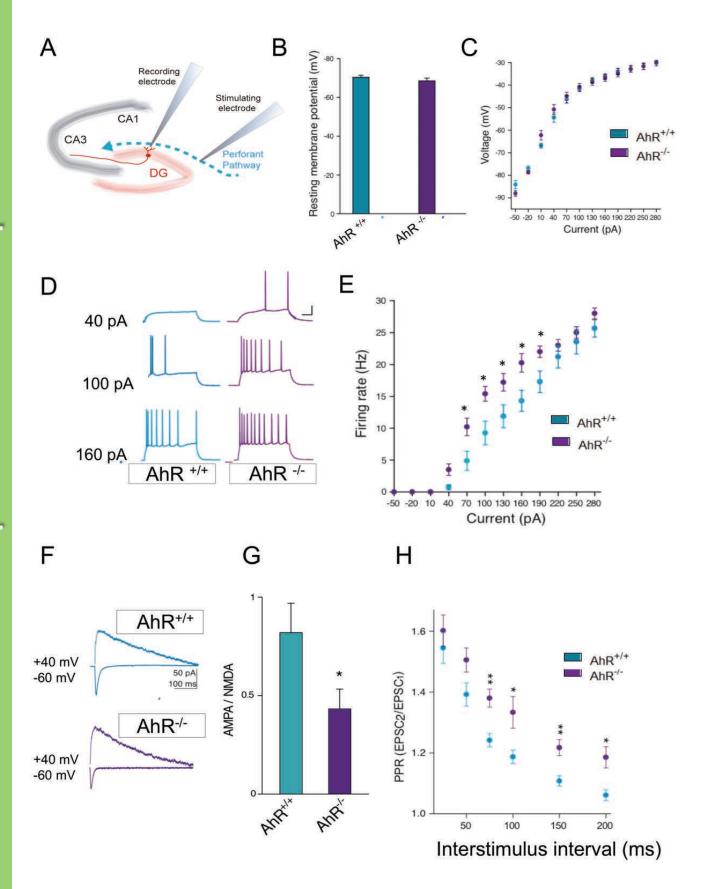
Figure 6. Acute ablation of AhR in adult neural precursors alters the 1182 synaptic properties of dentate gyrus granule cells. (A) Experimental 1183 protocol for tamoxifen administration and electrophysiological recordings. (B-C) 1184 The resting membrane potential (B) and the current-voltage relationship (C) did 1185 not show significant differences between DG granule cells from AhR^{f/f} and AhR-1186 icKO mice (n=4-6 animals/group). (D-E) DG granule cells firing rate is 1187 significantly increased by specific AhR ablation (*p<0.05 vs. AhR^{f/f}). 1188 Representative sample traces (D) and averaged values (E) in response to 1189 increasing depolarising currents. (F-G) AMPA/NMDA ratio is decreased by AhR 1190 deletion (*p<0.05 vs. AhR^{f/f}). Representative traces (F) and averaged values (G) 1191 of NMDA- and AMPA-mediated EPSCs recorded at +40 and -60 mV, 1192 respectively. (H) Averaged values showing a significant increase in the PPR at 1193

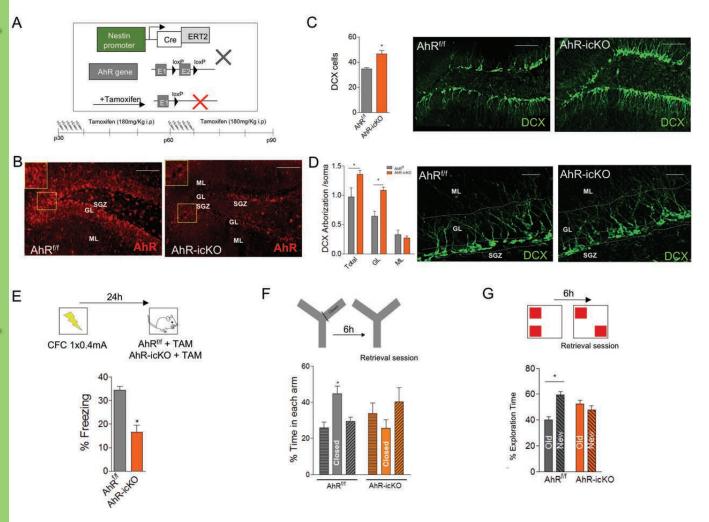
1194	interstimulus 75, 100, 150 and 200-ms intervals in cells lacking AhR (*p<0.05 $$
1195	vs. AhR ^{f/f}). Data are mean \pm SEM (25 neurons from 8 slices from n=6 AhR ^{f/f}
1196	mice and 24 neurons from 8 slices from n=4 AhR-icKO mice). Data were
1197	compared by using non-parametric 2-tailed Mann-Whitney tests.



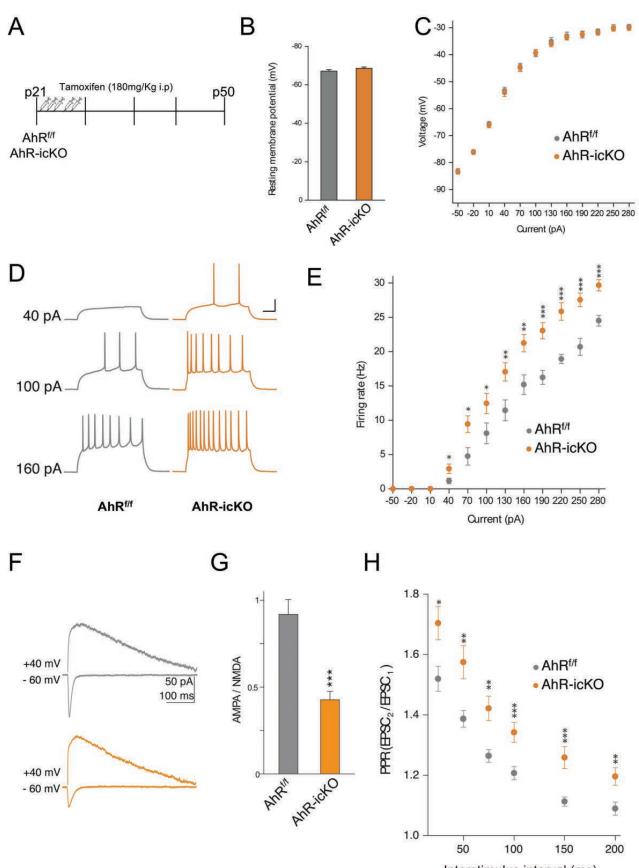












Interstimulus interval (ms)