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# Continuous exposure to amorphous formula of curcumin from the developmental stage facilitates anti-anxiety-like behavior and fear-extinction learning in rats

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## ABSTRACT

An amorphous formula of curcumin (CUR) has shown to enable an improved bioavailability after ingestion. The aim of this study was to investigate the hypothesis that exogenously administered CUR has an advantage in ameliorating post-traumatic stress disorder at low doses. To this end, Long-Evans rats were dietary exposed to CUR at 0.1% or 0.5% from gestational day 6 to postnatal day (PND) 74 or 77. Offspring exposed to 0.1% CUR revealed fa-

**Abbreviations:** AGIQ,  $\alpha$ -glycosyl isoquercitrin; ARC, activity-regulated cytoskeleton-associated protein; BW, body weight; CALB2, calbindin-D-29 K; COX2, cyclooxygenase-2; C<sub>T</sub>, threshold cycle; CUR, amorphous formula of curcumin; DAB, 3,3'-diaminobenzidine; DCX, doublecortin; EGR1, early growth response protein 1; FOS, Fos proto-oncogene, AP-1 transcription factor subunit; GABA,  $\gamma$ -aminobutyric acid; *Gapdh*, glyceraldehyde-3-phosphate dehydrogenase; GCL, granule cell layer; GD, gestational day; GFAP, glial fibrillary acidic protein; HPLC, high performance liquid chromatography; *Hprt1*, hypoxanthine phosphoribosyltransferase 1; IBA1, ionized calcium-binding adaptor molecule 1; IEG, immediate-early gene; IL, interleukin; LTP, long-term potentiation; mPFC, medial prefrontal cortex; NeuN, neuronal nuclei; NF- $\kappa$ B, nuclear factor  $\kappa$ B; NMDA, N-methyl-D-aspartate; NOS, nitric oxide synthase; OFT, open field test; OLT, object location test; ORT, object recognition test; PBS, phosphate-buffered saline; PCNA, proliferating cell nuclear antigen; p-ERK1/2, phosphorylated extracellular signal-regulated kinase 1/2 (phosphorylated p44/p42 MAP kinase); PFA, paraformaldehyde; PND, postnatal day; PTSD, post-traumatic stress disorder; PVALB, parvalbumin; PVP, polyvinylpyrrolidone; ROS, reactive oxygen species; RT, reverse transcription; SGZ, subgranular zone; SOX2, SRY-box transcription factor 2; SST, somatostatin; TBR2, T-box brain protein 2; TNF- $\alpha$ , tumor necrosis factor alpha; TUBB3, tubulin, beta 3 class III; TUNEL, terminal deoxynucleotidyl transferase-mediated dUTP-biotin nick end-labeling.

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cilitation of anti-anxiety-like behavior in the open field test and fear-extinction learning tested during PND 62 to 74, increases in hippocampal granule cells expressing immediate-early gene proteins and a decrease in prelimbic cortical neurons expressing phosphorylated extracellular signal-regulated kinase 1/2 after the last trial of the fear-extinction learning test on PND 74. The constitutive gene expression levels of *Gria1*, *Gria2*, *Grin2d*, *Slc17a6*, and *Slc17a7* were altered in the hippocampal dentate gyrus and amygdala on PND 77. These results suggest alterations in synaptic plasticity to strengthen neural circuits in promoting the behavioral effects by 0.1%-CUR. In contrast, 0.5% CUR revealed a lack of any of the changes in behavioral tests that were observed at 0.1%; however, this dose upregulated oxidative stress and neuroinflammation-related genes in the hippocampal dentate gyrus, and increased neural stem cells and proliferation activity of the subgranular zone in the dentate gyrus. These results suggest a possible preventive use of CUR at low doses in mitigating some stress disorders; however, excessively absorbed doses may prevent behavioral changes by inducing neuroinflammation that affects hippocampal neurogenesis involving neural stem cells.

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## 1. Introduction

Curcumin, a natural polyphenol contained in the rhizomes of *Curcuma longa* Linne, has pleiotropic properties such as anti-inflammatory, antioxidant, antiproliferative, and antiangiogenic activities [1]. In the brain, curcumin administration has improved hippocampal neurogenesis in chronically stressed rats [2], as well as antidepressant effects in rat depression models [3]. However, curcumin exhibits poor bioavailability because of poor absorption, rapid metabolization, and rapid systemic elimination [1]. Therefore, many methods have been proposed for improving the solubility of curcumin [4], such as amorphization, nanonization, and chemical conjugation. San-Ei Gen F.F.I., Inc. (Osaka, Japan) has recently developed an amorphous formula of curcumin (CUR) that has enabled an improved bioavailability after oral intake [5].

Classical fear conditioning is thought to contribute to anxiety disorders [6], and great efforts are being made to understand the neural circuits that underlie the acquisition, development, and disappearance of emotional memory. Three brain structures, including the medial prefrontal cortex (mPFC), hippocampus, and amygdala, form important brain circuits involved in fear conditioning and extinction [6]. The mPFC, which receives inputs from the hippocampus, projects directly onto the amygdala and midbrain, regulates the development of anxiety-like behavior and avoidance behavior against aversive stimuli [7]. There is also a prevailing notion that the prelimbic cortex, a dorsal region of the mPFC, controls the development of fear, and that the infralimbic cortex, a ventral region of the mPFC, suppresses fear [6]. The amygdala is the central brain region of fear response, and suppressive input from the mPFC is received by the basolateral amygdala and suppresses fear response by suppressing the neurons of the central nucleus of the amygdala [8].

The hippocampal dentate gyrus, known as a typical brain substructure that conducts adult neurogenesis and is responsible for memory formation and antidepressant activity, includes all processes of neuronal development of the granule

cell lineage, such as self-renewal of neural stem cells, proliferation and differentiation of neural progenitor cells in the subgranular zone (SGZ), and maturation of postmitotic granule cells in the SGZ and granule cell layer (GCL) [9]. In the hilar region of the hippocampal dentate gyrus,  $\gamma$ -aminobutyric acid-ergic (GABAergic) interneurons connect with adult-born dentate granule cells and play a functional role in adult neurogenesis [10,11].

Synaptic plasticity is the ability of neurons to bring about changes in the connections between neuronal networks in response to use or disuse in their activity. Because memories are postulated to be represented by vastly interconnected neural circuits in the brain regions including the hippocampus [12], synaptic plasticity is an important neurochemical foundation of learning and memory [13]. Excess production of reactive oxygen species (ROS) in the hippocampus leads to detrimental effects that are involved in impairments in synaptic plasticity and memory function [14], and administration of antioxidants is effective for ameliorating cognitive deficits by protecting against oxidative damage [15].

Recently, some polyphenolic antioxidants have been shown to exert an ameliorating effect on post-traumatic stress disorder (PTSD), a trauma and stressor-related disorder, in animal models [16,17], and as a result, more attention has been given to these antioxidants. Therefore, it could be hypothesized that CUR has an advantage in ameliorating PTSD and also in strengthening other behavioral functions in normal animals at low doses. In the present study, we tested the exposure effect of CUR at lower doses than those reported with nonamorphous curcumin formula on behavioral functions of rat offspring with exposure from the gestational stage until the adult stage by setting 2 doses to observe dose responses. By means of immunohistochemistry, we evaluated the effect of CUR on synaptic plasticity changes in the GCL of the hippocampal dentate gyrus and mPFC. We also examined gene expression changes related to synaptic plasticity and chemical mediator signaling by means of real-time reverse transcription (RT)-PCR. Changes in hippocampal neurogenesis were also examined.

## 2. Methods and materials

### 2.1. Chemicals and animals

Curcumin mixed with polyvinylpyrrolidone (PVP) and sucrose fatty acid ester (amorphous formula of curcumin; curcumin content: 19.9%) and PVP and sucrose fatty acid ester (carrier solvent) were provided by San-Ei Gen F.F.I. Inc. Forty-one mated female Iar: Long-Evans rats were purchased from Japan SLC, Inc. (Hamamatsu, Japan) at gestational day (GD) 3, where GD 0 is the day the appearance of vaginal plug. Rats were individually housed with their offspring in polycarbonate cages with paper bedding until postnatal day (PND) 21, where PND 0 was defined as the day of delivery. Animals were kept in an air-conditioned animal room (temperature:  $23 \pm 2$  °C, relative humidity:  $55 \pm 15\%$ ) with a 12-h light/dark cycle. Mated female rats were provided ad libitum a commercial rodent powder diet (CRF-1; Oriental Yeast Co., Ltd., Tokyo, Japan) until exposure to CUR began on GD 6 and tap water during the experiment.

While the detailed ingredient composition of the CRF-1 is not available (commercial formulation), this diet was mainly formulated from corn, wheat (bran), defatted soybeans, defatted rice bran, alfalfa, fish flour, skim milk powder, soybean oil, and brewer's yeast. The macronutrient composition of the CRF-1 is 8.2% moisture, 21.9% crude protein, 5.4% crude fat, 6.3% crude ash, 2.9% crude fiber, 55.3% nitrogen-free extract, and total energy of 3570 kcal/kg. Vitamin content/kg was as follows: vitamin A, 32,450 IU; vitamin D3, 6430 IU; vitamin E, 203 mg; vitamin K3, 1.6 mg; vitamin B1, 47.4 mg; vitamin B2, 33.1 mg; vitamin C, 120 mg; vitamin B6, 12.7 mg; vitamin B12, 172 µg; inositol, 4530 mg; biotin, 271 µg; pantothenic acid, 50.7 mg; niacin, 156.2 mg; choline, 2.6 g; folic acid, 2.7 mg. Mineral content/kg is as follows: calcium, 12.2 g; phosphorus, 8.1 g; magnesium, 2.3 g; sodium, 2.6 g; potassium, 8.6 g; iron, 138 mg; aluminum, 21 mg; copper, 9.5 mg; zinc, 61.5 mg; cobalt, 3.3 mg; manganese, 72.7 mg. Amino acid content/kg is as follows: isoleucine, 8.2 g; leucine, 16.1 g; lysine, 11.2 g; methionine, 4.4 g; cysteine, 3.4 g; phenylalanine, 9.6 g; tyrosine, 6.4 g; threonine, 8.3 g; tryptophan, 2.7 g; valine, 10.2 g; arginine, 13.0 g; histidine, 5.6 g; alanine, 11.3 g; aspartic acid, 19.2 g; glutamic acid, 37.5 g; glycine, 11.0 g; proline, 12.6 g; serine, 10.3 g.

From PND 21 onward, offspring were reared and provided the CRF-1 diet with or without CUR and tap water ad libitum throughout the experimental period. Dietary concentration of curcumin was measured using the diets kept for 17 days after preparation of 0.1% and 0.5% in diet by high performance liquid chromatography (HPLC; analyzed by San-Ei Gen F.F.I., Inc.), and the same concentration of curcumin was confirmed in diet after the preservation.

### 2.2. Experimental design

Mated female rats were randomly divided into three groups of vehicle controls (14 animals), 0.1% CUR (13 animals) and 0.5% CUR (14 animals) (Fig. 1). These animals were treated from GD 6 to day 21 post-delivery with PVP and sucrose fatty acid ester at 1.95 w/w% (vehicle controls), curcumin at 0.1 w/w% with 0.39 w/w% vehicle (0.1% CUR), or curcumin at 0.5 w/w%

**Table 1 – Composition of the experimental diets fed to rats**

	g/kg		
	Ctrl	0.1% CUR <sup>a</sup>	0.5% CUR
Basal diet	980.5	995.0	974.8
Curcumin powder	0	1.1	5.7
Polyvinylpyrrolidone	13.2	2.6	13.2
Sucrose fatty acid ester	6.3	1.3	6.3

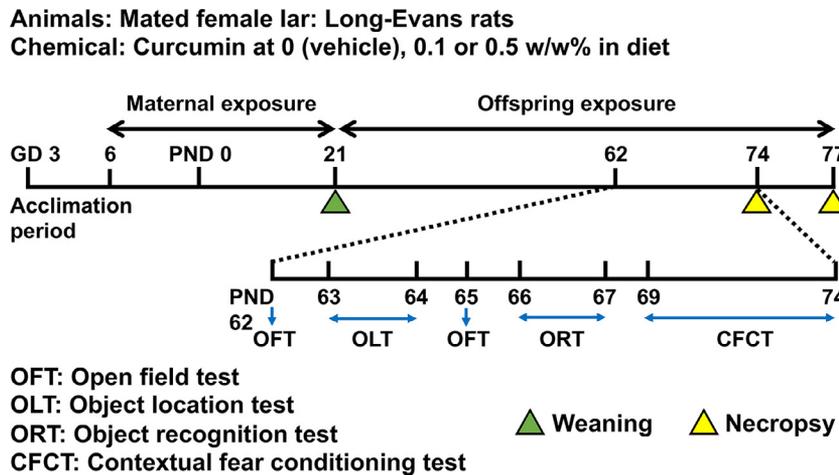
Abbreviations: Ctrl, vehicle controls; CUR, amorphous formula of curcumin.

<sup>a</sup> Curcumin-containing diets were prepared by adding the amorphous formula of curcumin, a mixture of curcumin powder with polyvinylpyrrolidone and sucrose fatty acid ester as a carrier solvent (curcumin content: 19.9%), to the basal diet. Basal diet was mixed with vehicle at 1.95 w/w% for vehicle controls, curcumin at 0.1 w/w% with 0.39 w/w% vehicle for 0.1% CUR, or curcumin at 0.5 w/w% with 1.95 w/w% vehicle for 0.5% CUR.

with 1.95 w/w% vehicle (0.5% CUR) in the basal diet (Table 1). Considering 5.7-fold higher absorption rate than nonamorphous fine-particulate formula of curcumin calculated by AUC<sub>0 to 24 hours</sub> after single oral administration of 100 mg/kg body weight (BW) [18], the doses of CUR were determined based on the study results of nonamorphous formula of curcumin with the dietary dose at 1.5% that impaired fear memory consolidation and reconsolidation by treatment for 5 days in a PTSD-induction model using rats [19]. In the present study, the rationale for setting the dose levels of CUR was to examine continuous exposure effect of CUR from fetal stages with the doses lower than nonamorphous curcumin formula, and therefore, we set 2 doses, 1 lower than and the other higher than the absorbed value of 1.5% dietary dose of aforementioned nonamorphous formula of curcumin. Concentrations of curcumin in plasma removed from male offspring cardiac blood on PND 21 (N = 8 each from the vehicle controls and 0.5% CUR) and in milk from female offspring stomach on PND 14 (N = 8 each from the vehicle controls and 0.5% CUR) were measured using HPLC by Japan Food Research Laboratories (Tokyo, Japan). For measuring curcumin concentrations in milk, one dam each from the vehicle controls and 0.5% CUR was used.

Dams were subjected to measurement of BW and food and water consumption twice a week from GD 6 to day 21 post-delivery. On PND 4, litters were randomly culled to preserve 7 or 8 male and 0 or 1 female pups per dam. If dams had fewer than 7 male pups, more female pups were included to maintain a total of 8 pups per litter. The offspring were weighed twice a week until PND 21. Dams were euthanized by exsanguination from the abdominal aorta under CO<sub>2</sub>/O<sub>2</sub> anesthesia on day 21 post-delivery. In this study, male offspring were selected for behavioral tests and immunohistochemical and gene expression analyses because animal behaviors and hippocampal neurogenesis are influenced by circulating levels of steroid hormones during the estrous cycle [20,21].

On PND 21, 28 to 35 male offspring (1–2 pups per dam) and all female offspring of all groups were euthanized by exsanguination from the abdominal aorta under CO<sub>2</sub>/O<sub>2</sub>-induced anesthesia and subjected to necropsy for brain tissue sampling for other purposes of analysis. The remaining male



**Fig. 1 – Experimental design of continuous exposure to amorphous formula of curcumin from developmental stage in rats.**

offspring were fed a diet containing CUR or vehicle at the same concentration to dams until PND 77 (4 or 5 animals/cage during PND 21–PND 40; 2 or 3 animals/cage from PND 41 onwards); BW and food and water consumption were measured once weekly. Behavioral tests were performed using 10 to 26 male offspring per group (1–2 pups per dam) during the period from PND 62 to PND 74.

On PND 74, animals that were subjected to behavioral tests were euthanized for brain sampling. Ten male offspring per group (1 pup per dam) were subjected to perfusion fixation for brain immunohistochemistry through the left cardiac ventricle with ice-cold 4% (w/v) paraformaldehyde (PFA) in 0.1 M phosphate buffer (pH 7.4) at a flow rate of 35 mL/min under CO<sub>2</sub>/O<sub>2</sub>-induced anesthesia. For transcript-level expression analysis, brain samples from 6 male offspring per group (1 male offspring per dam) were prepared. Animals were sacrificed at 90 min after the 3rd trial of fear-extinction learning.

On PND 77, animals that were not subjected to behavioral tests were euthanized for brain sampling. Ten male offspring per group (1 male offspring per dam) were subjected to perfusion fixation for brain immunohistochemistry in the same way with PND 74 samples. For transcript-level expression analysis of constitutive expression changes, brain samples from 6 male offspring per group (1 male offspring per dam) were prepared.

All dams and offspring were checked each day to assess their general appearance (abnormal gait and behaviors). All procedures in this study were conducted in accordance with the National Institutes of Health guide for the care and use of laboratory animals (NIH Publications No. 8023, revised 1978) and according to the protocol approved by the Animal Care and Use Committee of The Tokyo University of Agriculture and Technology (Approved No.: 30–63). All efforts were made to minimize animal suffering.

### 2.3. Behavioral tests

In each behavioral test, male animals were transported from the animal room to the behavioral test room 1 hour to 2 hours before starting the tests. After the end of each behavioral test,

animals were promptly returned to the home cage and transferred to the animal room. Apparatuses were cleaned with 70% ethanol solution before and after each test. All experiments were conducted during the period from 8:00 am to 7:00 p.m., and the order of animal selection for tests among groups was counter-balanced across test time to avoid any bias in the trial times of each group.

#### 2.3.1. Open field test (OFT)

The OFT was performed on PND 62 and PND 65 to assess locomotor activity and anxiety-like behaviors and to habituate the animals to the arena as an acclimation phase in the object location test (OLT) and object recognition test (ORT) on the following days. Two pups from one dam were assigned to OFT. The arena comprised a square stainless-steel tray with a matte black polyvinyl plastic surface and stainless-steel walls with matte black polyvinyl plastic surface surrounding the tray (900 mm width × 900 mm depth × 500 mm height; O'Hara & Co., Ltd., Tokyo, Japan). The illumination was set at 20 Lux at the middle of the arena. Each animal was placed at the corner of the arena with the head facing the wall and allowed to explore the arena freely for 10 min. Time spent in the central region (540 mm × 540 mm) or wall side and the total distance moved were recorded by a CCD camera (WAT-902B; Watec Co., Ltd., Tsuruoka, Japan) mounted above the arena and evaluated by an automatic video-tracking system (TimeOFCR1 software; O'Hara & Co. Ltd.).

#### 2.3.2. Object location test and object recognition test

The OLT was performed on PND 63 and PND 64 to assess spatial object memory, and the ORT was performed on PND 66 and PND 67 to assess nonspatial object memory. All experiments in both tests were conducted in the same arena that was used in the OFT. One different pup from one dam was assigned to each of OLT and ORT. Both of the OLT and ORT comprised three steps: acclimation, sample phase, and test phase. As aforementioned, OFT was conducted as acclimation of animals in both of the OLT and ORT. After 24 hours, each animal was placed in the middle of the wall along the inside of the field with the head facing the wall and allowed to explore for 5

min with 2 identical sample objects (sample phase in the OLT and ORT). Objects were placed equidistant to this location to the right and left sides behind the animal in the arena. Test phase of the OLT and ORT was conducted 24 hours after the sample phase, and the animals were placed in the same position as the sample phase. In the OLT, each animal was allowed to explore for 3 min with one object on the same location and the other object moved to the opposite side of the arena. In this experiment, objects were identical to the sample phase, and the position of one of the two objects was displaced 35 cm away from the original position. In the ORT, each animal was allowed to explore for 3 min with one sample object and one novel object on the same location as sample phase. The illumination was set at 20 Lux on the middle of the arena. The sample object was a gray cone of polyvinyl plastic with a rough surface and a stainless-steel tip. The novel object was a sphere with a patchy pattern of black and white consisting of smooth polyvinyl plastic. These objects were heavy enough that could not be moved by the animals. In both of the sample and test phases, the total distance traveled and exploration time toward each object were recorded by a CCD camera (WAT-902B; Watec Co., Ltd.) mounted above the arena and evaluated by automatic video-tracking system (TimeSSI software; O'Hara & Co., Ltd.). When a rat's nose approached to within 3 cm of an object, the video-tracking system automatically counted it as "exploration" and the cumulative exploration time was recorded.

A discrimination index for the OLT or ORT was determined using the following formula: discrimination index = exploration time with displaced object or novel object / (exploration time with familiar object + exploration time with displaced object or novel object).

### 2.3.3. Contextual fear conditioning test

The contextual fear conditioning test was performed during PND 69 and PND 74 using all the animals subjected to OFT, OLT, and ORT. Conditioning and testing took place in a rodent observation cage (370 mm width × 300 mm depth, 250 mm height; CL-3001, O'Hara & Co., Ltd.) that was placed in a sound-attenuating chamber (CL-4211; O'Hara & Co., Ltd.). The sidewalls and door of the observation cage were constructed of Plexiglas. The floor comprised 21 steel rods through which a scrambled footshock from a shock generator (SGA-2020; O'Hara & Co., Ltd.) could be delivered. During experiments, the chamber was ventilated, kept at a background white noise level of 50 dB, and illuminated at 200 Lux by white light emitting diode bulbs. This test comprised four steps: habituation, contextual fear conditioning, fear acquisition, and fear extinction.

**Habituation (Day 1):** To familiarize with test surroundings, animals were placed into the observation cage for 5 min without footshock.

**Contextual fear conditioning (Day 2):** Twenty-four hours after the habituation, each animal was moved to the observation cage, and after 88, 148, and 238 sec, they received 2 sec footshocks (0.3 mA intensity, a total of 3 footshocks). Animals were removed from the observation cage 60 sec after the final footshock and returned to the home cage. Thus, it took 5 min for a test.

**Fear acquisition (Day 3):** Twenty-four hours after the conditioning, each animal was placed back into the same context as the conditioning context for 5 min without footshock.

**Fear extinction (Day 4, 5, 6):** After 2, 3, and 4 days of the conditioning, each animal was placed back into the same context as the conditioning context for 5 min without footshock.

The animals' behavior was video recorded by the CCD camera (WAT-902B; Watec Co., Ltd.) mounted above the observation cage and analyzed using an automatic video-tracking system (TimeFZ2 software; O'Hara & Co., Ltd.). Body freezing time was automatically measured, and the freezing time was defined as the percent of time the rat spent in freezing behavior for  $\geq 5$  sec during each of the experiment time of 5 min.

## 2.4. Immunohistochemistry and apoptotic cell detection

For immunohistochemistry analysis, perfusion-fixed brains of male offspring were additionally fixed with the same PFA buffer solution overnight. Coronal cerebral slices were prepared at +3.0 mm and -3.5 mm from the bregma of the brains at PND 74 and PND 77 ( $N = 9-10$ /group) using a brain-matrix cast (Muromachi Kikai Co., Ltd., Tokyo, Japan). Due to a mistake in slicing at +3.0 mm, one sample each of the vehicle controls and 0.1% CUR was excluded from counting of immunoreactive cells. Brain slices were further immersion-fixed with the same PFA buffer solution overnight at 4°C and were routinely processed for paraffin embedding and sliced into 3- $\mu$ m-thick sections.

Brain sections were subjected to immunohistochemical analysis using primary antibodies against the following (Table 2): Glial fibrillary acidic protein (GFAP), which is expressed in type-1 neural stem cells (radial glial cells) in the SGZ and astrocytes [9]; SRY-box transcription factor 2 (SOX2), which is expressed in type-1 neural stem cells and type-2a progenitor cells in the SGZ [9]; T-box brain protein 2 (TBR2), expressed in type-2b progenitor cells in the SGZ [9]; doublecortin (DCX), which is expressed in type-2b and type-3 progenitor cells and immature granule cells in the SGZ and GCL [9]; tubulin, beta 3 class III (TUBB3, also known as Tuj-1), which is expressed mainly in postmitotic immature granule cells in the SGZ and GCL [22]; neuronal nuclei (NeuN), which is expressed in postmitotic neurons of both immature and mature granule cells in the SGZ and GCL [9]; and parvalbumin (PVALB), calbindin-D-29 K (CALB2), and somatostatin (SST), which are expressed in GABAergic interneurons [10]; Ionized calcium-binding adaptor molecule 1 (IBA1), a microglia-specific molecule in the brain [23]; proliferating cell nuclear antigen (PCNA), a cell proliferation marker; Fos proto-oncogene, AP-1 transcription factor subunit (FOS), activity-regulated cytoskeleton-associated protein (ARC), cyclooxygenase-2 (COX2), early growth response protein 1 (EGR1, also known as Zif268), which are members of the immediate-early gene (IEG) proteins involved in synaptic plasticity [13]; phosphorylated extracellular signal-regulated kinase 1/2 (p-ERK1/2; phosphorylated p44/p42 MAP kinase), a member of the mitogen activated protein kinase family that is activated by phosphorylation to promote transcriptional programs leading to the induction of *Arc*, *Fos*, and *Egr1* [24,25].

**Table 2 – Antibodies used in this study**

Antigen	Abbreviated name	Host species	Clonality	Clone name	Dilution	Antigen retrieval	Manufacturer (city, state, country)
Activity-regulated cytoskeleton-associated protein	ARC	Rabbit	Polyclonal IgG	n.a.	1:2000	Microwaving <sup>a</sup>	Synaptic Systems GmbH (Goettingen, Germany)
Calbindin-D-29K	CALB2	Rabbit	Monoclonal IgG	EP1798	1:600	Microwaving <sup>a</sup>	Abcam plc (Cambridge, UK)
Cyclooxygenase-2	COX2	Mouse	Monoclonal IgG <sub>1</sub>	33/Cox-2	1:200	Microwaving <sup>a</sup>	BD Biosciences, Inc. (San Jose, CA, USA)
Doublecortin	DCX	Rabbit	Polyclonal IgG	n.a.	1:1000	None	Abcam plc
Early growth response protein 1	EGR1	Rabbit	Monoclonal IgG	15F7	1:200	Microwaving <sup>a</sup>	Cell Signaling Technology, Inc. (Danvers, MA, USA)
Fos proto-oncogene, AP-1 transcription factor subunit	FOS	Rabbit	Polyclonal IgG	n.a.	1:500	None	Santa Cruz Biotechnology, Inc. (Dallas, TX, USA)
Glial fibrillary acidic protein	GFAP	Mouse	Monoclonal IgG <sub>1</sub>	GA5	1:200	None	Merck Millipore (Burlington, MA, USA)
Ionized calcium-binding adaptor molecule 1	IBA1	Rabbit	Polyclonal IgG	n.a.	1:300	None	Fujifilm Wako Pure Chemical Corporation (Osaka, Japan)
Neuronal nuclei	NeuN	Mouse	Monoclonal IgG <sub>1</sub>	A60	1:100	None	Merck Millipore
Parvalbumin	PVALB	Mouse	Monoclonal IgG <sub>1</sub>	PARV-19	1:1000	Microwaving <sup>a</sup>	Merck Millipore
phosphorylated extracellular signal-regulated kinase 1/2 (phosphorylated p44/p42 MAP kinase)	p-ERK1/2	Rabbit	Monoclonal IgG	137F5	1:100	Microwaving <sup>a</sup>	Cell Signaling Technology, Inc.
Proliferating cell nuclear antigen	PCNA	Mouse	Monoclonal IgG <sub>2a</sub>	PC10	1:200	None	Agilent Technologies (Santa Clara, CA, USA)
SRY-box transcription factor 2	SOX2	Mouse	Monoclonal IgG <sub>1</sub>	9–9–3	1:4000	None	Abcam plc
Somatostatin	SST	Rabbit	Polyclonal IgG	n.a.	1:400	Microwaving <sup>a</sup>	Abcam plc
T box brain 2	TBR2	Rabbit	Polyclonal IgG	n.a.	1:500	Autoclaving <sup>b</sup>	Abcam plc
Tubulin, beta 3 class III	TUBB3	Mouse	Monoclonal IgG <sub>1</sub>	TU-20	1:500	Microwaving <sup>a</sup>	Abcam plc

Abbreviation: n.a, not applicable.

<sup>a</sup> 90°C for 10 min in 10 mM citrate buffer (pH 6.0).

<sup>b</sup> 121°C for 10 min in 10 mM citrate buffer (pH 6.0).

Immunodetection was conducted using a Vectastain Elite ABC kit (Vector Laboratories Inc., Burlingame, CA, USA). In detail, paraffin-embedded brain sections were deparaffinized in xylene, rehydrated in a graded ethanol series and immersed in a tap water. The antigen retrieval was applied for some antibodies and the retrieval conditions are listed in Table 2. Endogenous peroxidase was blocked by exposure to 0.3% (v/v) H<sub>2</sub>O<sub>2</sub> solution in absolute methanol for 30 min and sections were rinsed in phosphate-buffered saline (PBS; pH 7.4). Block-

ing was conducted with 1.5% normal serum in PBS for 30 min. Horse serum was used in case of mouse monoclonal primary antibodies and goat serum was used for rabbit monoclonal and polyclonal primary antibodies. Incubation with respective primary antibodies was performed at 4 °C overnight. All antibodies were diluted with PBS containing 0.5% casein. This step was followed by incubation with: (a) Biotinylated horse antimouse secondary antibody or biotinylated goat antirabbit secondary antibody in PBS containing 1.5% normal serum for

30 min; (b) peroxidase-conjugated avidin in PBS for 30 min. After each step, the sections were rinsed in PBS and all steps except for incubation with primary antibody were performed at room temperature. All incubations were performed in a humidity chamber. Peroxidase activity was visualized by reaction with freshly prepared 0.03% 3,3'-diaminobenzidine (DAB) plus 0.01% hydrogen peroxide in Tris-buffered saline (pH 7.6). Hematoxylin counterstaining was then performed, and coverslips were mounted on immunostained sections for microscopic examination. One section per animal was subjected to immunohistochemistry of each molecule.

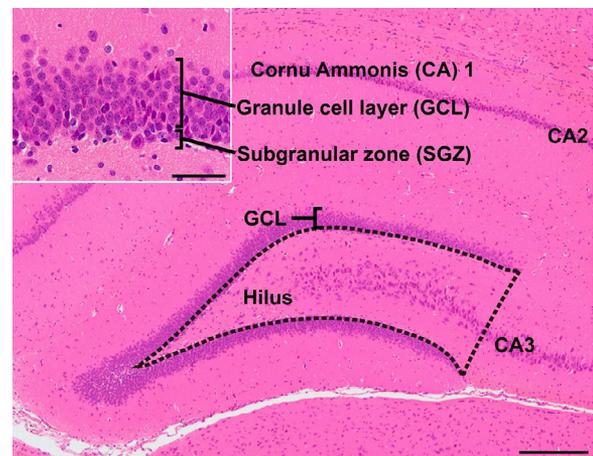
To evaluate apoptosis in the SGZ of the dentate gyrus in the offspring, terminal deoxynucleotidyl transferase-mediated dUTP-biotin nick end-labeling (TUNEL) assay was performed using the ApopTag Peroxidase *In Situ* Apoptosis Detection Kit (Merck Millipore, Burlington, MA, USA) according to the manufacturer's instructions, with DAB/H<sub>2</sub>O<sub>2</sub> as the chromogen. One section per animal was subjected to TUNEL assay.

## 2.5. Evaluation of immunoreactive cells and apoptotic cells

Immunoreactive cells (GFAP<sup>+</sup>, SOX2<sup>+</sup>, TBR2<sup>+</sup>, DCX<sup>+</sup>, TUBB3<sup>+</sup>, NeuN<sup>+</sup>, FOS<sup>+</sup>, ARC<sup>+</sup>, COX2<sup>+</sup>, EGR1<sup>+</sup>, p-ERK1/2<sup>+</sup>, and PCNA<sup>+</sup> cells) or TUNEL<sup>+</sup> apoptotic cells in the SGZ and/or GCL were bilaterally counted and normalized for the length of the SGZ (Fig. 2). Immunoreactive cells (PVALB<sup>+</sup>, CALB2<sup>+</sup>, SST<sup>+</sup>, GFAP<sup>+</sup> and IBA1<sup>+</sup> cells) distributed within the hilus of the hippocampal dentate gyrus were bilaterally counted and normalized per area unit of the hilus area (Fig. 2). Immunoreactive neurons located inside of the Cornu Ammonis region 3, consisting of large pyramidal neurons that can be morphologically distinguished from relatively small interneurons, were excluded from counting immunoreactive cells in the hilus of the dentate gyrus. Immunoreactive cells (FOS<sup>+</sup>, ARC<sup>+</sup>, COX2<sup>+</sup>, and p-ERK1/2<sup>+</sup> cells) in the infralimbic cortex and prelimbic cortex of the mPFC were bilaterally counted and normalized per area unit of each cortical area (Fig. 3). The position and shape of each area were determined with reference to a cresyl violet staining image adjacent to the immunostained sections and the atlas of the rat brain [26].

Brain samples of animals that were subjected to behavioral analysis and euthanized on PND 74 were examined for immunohistochemistry of FOS, ARC, COX2, EGR1, and p-ERK1/2. Because expression peak of IEG proteins in response to acute stimuli occurs within approximately 90 to 120 min [27], animals were euthanized for perfusion fixation 90 min after the last trial of the contextual fear conditioning test. Brain samples of animals that were not subjected to behavioral analysis and euthanized on PND 77 were examined for immunohistochemistry of GFAP, SOX2, TBR2, DCX, TUBB3, NeuN, PCNA, PVALB, CALB2, SST and IBA1, as well as FOS and ARC, and for TUNEL assay.

The number of each immunoreactive cellular population (except for NeuN<sup>+</sup> cells in the GCL) or TUNEL<sup>+</sup> apoptotic cells was manually counted while blinded to the treatment conditions under microscopic observation using a BX53 microscope (Olympus Corporation, Tokyo, Japan). In case of GFAP<sup>+</sup> cells in the SGZ, only those having apparent vertically extending radial processes were counted as type-1 neural stem cells. In

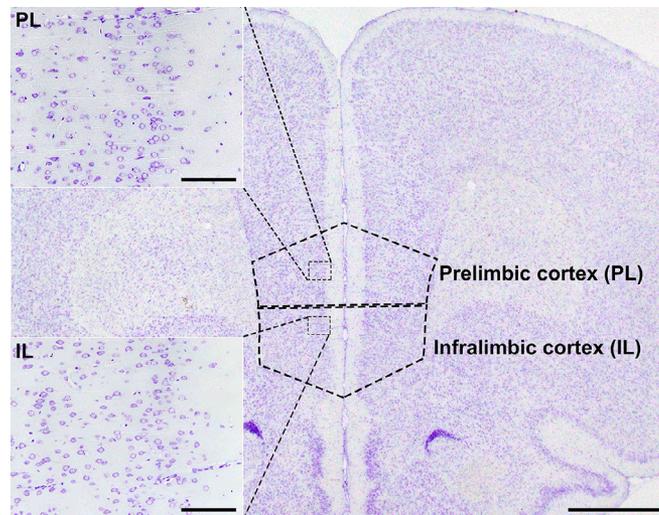


**Fig. 2 – Overview of the hippocampal formation of a male rat stained with hematoxylin and eosin. (Inset) Higher magnification of the granule cell layer (GCL) and subgranular zone (SGZ). The distribution of immunoreactive cells for GFAP, SOX2, TBR2, DCX, TUBB3, NeuN, ARC, FOS, COX2, EGR1, p-ERK1/2, and proliferating cells or apoptotic cells were measured in the SGZ and/or GCL. The number of cells in the hilus of the dentate gyrus (as enclosed by the dotted line) displaying immunoreactivity for PVALB, SST, CALB2, GFAP or IBA1 was counted and normalized for the unit area. Only small-sized cells with positive immunoreactivity for these antigens were counted, and larger Cornu Ammonis (CA) 3 neurons were excluded from counting. Magnification  $\times 40$ ; bar 250  $\mu\text{m}$ . (Inset) Higher magnification of the GCL. Magnification  $\times 400$ ; bar 50  $\mu\text{m}$ .**

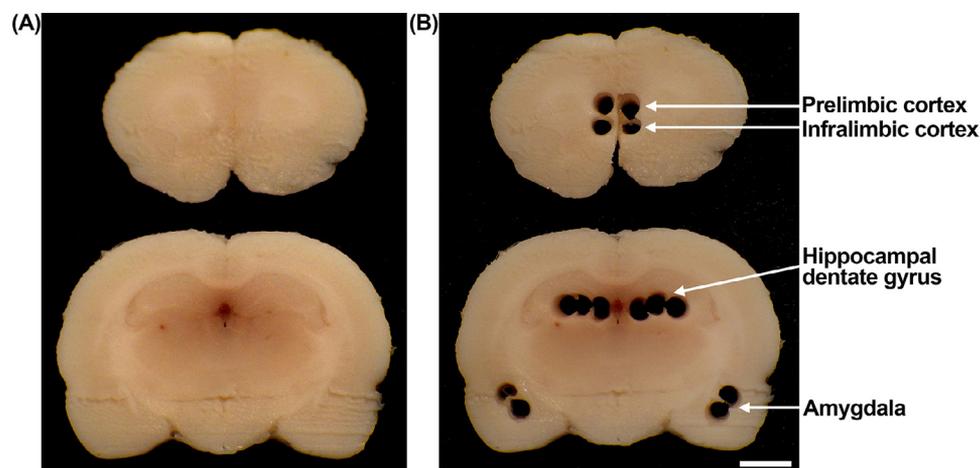
case of NeuN<sup>+</sup> cells in the GCL, the number of immunoreactive cells for counting was high, and therefore, an image analysis-assisted automatic cell counting method was applied. More specifically, digital photomicrographs at  $\times 200$ -fold magnification were taken using a DP72 Digital Camera System (Olympus Corporation), attached to a BX53 microscope, and positive cell counting was performed applying the WinROOF image analysis software package (version 5.7; Mitani Corporation, Fukui, Japan). The length of the SGZ and the hilar area were measured in microscopic images at  $\times 40$ -fold magnification by applying the cellSens Standard (version 1.9; Olympus Corporation).

## 2.6. Transcript-level expression analysis

Transcript levels in the hippocampal dentate gyrus, amygdala, infralimbic cortex and prelimbic cortex were examined using real-time RT-PCR in male offspring. Brain tissues were dissected according to the whole-brain fixation method using methacarn solution as previously reported [28]. In brief, 2-mm coronal cerebral slices were prepared at  $-3.0$  mm from the bregma using a brain-matrix cast to collect tissue samples from the hippocampal dentate gyrus and amygdala using a punch-biopsy device with a pore-size diameter of 1 mm in diameter (BP-10F; Kai Industries Co., Ltd., Gifu, Japan) (Fig. 4). For sampling the infralimbic cortex and prelimbic cortex tissues, 2-mm-thick coronal cerebral slices were prepared



**Fig. 3** – Overview of the medial prefrontal cortex formation of a male rat stained with cresyl violet. (Inset) Higher magnification of the prelimbic cortex (PL) and infralimbic cortex (IL). The distribution of immunoreactive cells for ARC, FOS, COX2, and p-ERK1/2 were measured in the PL and IL. Magnification  $\times 12.5$ ; bar 1 mm. (Inset) Higher magnification of the PL or IL. Magnification  $\times 200$ ; bar 100  $\mu\text{m}$ .



**Fig. 4** – Overview of the dissected brain regions for transcript-level expression analysis of rat offspring. Sampling was made from methacarn-fixed brain tissue slices using punch-biopsy devices bilaterally. For sampling of the tissues of the infralimbic cortex and prelimbic cortex, a 2 mm-thick slice anterior to the coronal cerebral plane at +2.0 mm from the bregma was prepared. For sampling of the tissues of the hippocampal dentate gyrus and amygdala, a 2 mm-thick slice posterior to the  $-2.0$  mm coronal plane from the bregma was prepared. (A) The coronal cut surface views at +3.0 mm and  $-3.0$  mm from the bregma before dissection. (B) The identical surface view to the panel (A) after dissection of the prelimbic cortex, infralimbic cortex, hippocampal dentate gyrus and amygdala. Bar 2 mm.

at +2.0 mm from the bregma. Total RNA was extracted from brain tissue samples from all groups ( $n = 6$  per group) using RNeasy Mini kit (Qiagen, Hilden, Germany). First-strand cDNA was synthesized using SuperScript III Reverse Transcriptase (Thermo Fisher Scientific, Waltham, MA, USA) in a 20- $\mu\text{L}$  total reaction mixture with 1  $\mu\text{g}$  of total RNA. Analysis of the transcript levels for the gene targets shown in Table 3 was performed using PCR primers designed with Primer Express software (Version 3.0; Thermo Fisher Scientific). Real-time PCR with Power SYBR Green PCR Master Mix (Thermo Fisher Scientific) was conducted using a StepOnePlus Real-time PCR

System (Thermo Fisher Scientific). The relative differences in gene expression between the vehicle controls and each exposure group were calculated using threshold cycle ( $C_T$ ) values that were first normalized to that of glyceraldehyde-3-phosphate dehydrogenase (*Gapdh*) or hypoxanthine phosphoribosyltransferase 1 (*Hprt1*), which served as endogenous controls in the same sample, and then relative to a control  $C_T$  value using the  $2^{-\Delta\Delta C_T}$  method [29].

Transcript levels of all genes were measured in the hippocampal dentate gyrus using samples of PND 77 animals that were not subjected to behavioral tests, except for *Il1a*, *Il1b*, *Il4*,

**Table 3 – Sequence of primers used for real-time RT-PCR**

Gene	Accession no.	Forward primer (5' → 3')	Reverse primer (5' → 3')
<b>Glutamate receptors</b>			
<i>Gria1</i>	NM_031608	GTGAGCGTCG TCCTTCTCT	TCTTCGCTGT GCCATTGTA
<i>Gria2</i>	NM_001083811.1	CATCACACCT AGTTCCCAA CA	CTTTGAGGTC AGGTCCGATC T
<i>Gria3</i>	NM_032990	GAAACATAAA GGACGTCCAG GAA	TTCTGCCTT CTGTCCATTT CT
<i>Grin2a</i>	NM_012573	GGCTGTCAGC ACTGAATCCA	GGTTTAGAGA ATCCTGGCGT AGAG
<i>Grin2b</i>	NM_012574	TGTGCAAGAC ACAAGATTA AAACCA	GGAGGATAAA GGAACGGAAG AAA
<i>Grin2c</i>	NM_012575.3	TGGGTGATGA TGTTCGTGAT G	GTGAGGTTCT GGTTGTAGCT GACA
<i>Grin2d</i>	NM_022797	TGTGCTCAC ACCCAAGGA	GGTCACTGCC ACAAGGATG T
<b>Glutamate transporters</b>			
<i>Slc17a6</i>	NM_053427.1	GGCAGACCCT GAGGAAACAA	TCCCCTGTTT CTTCATCCA
<i>Slc17a7</i>	NM_053859.2	TTGTGGCTAC CTCACCCTA A	CGAAGATGAC ACAGCCATAG TGA
<b>Synaptic plasticity-related genes</b>			
<i>Fos</i>	NM_022197	CAACGAGCCC TCCTCTGACT	TGCCTTCTCT GACTGCTCAC A
<i>Jun</i>	NM_021835	CTAACCCCGC GTGAAGTGA	GCATCGTCGT AGAAGGTCGT TT
<i>Arc</i>	NM_019361.1	AGTGTCTGAA AGGCAATGAA AAGTAG	CCTTCGGCCA TCTCTGATTC
<i>Ptgs2</i>	NM_017232.3	GCTTTCTCCA ACCTCTCCTA CTACA	GGGAGTTGGG CAGTCATCAG
<i>Egr1</i>	NM_012551.2	CGAGCGAACA ACCCTACGA	CGTTATTTCAG AGCGATGTCA GAA
<b>Oxidative stress-related genes</b>			
<i>Nfe2l2</i>	NM_031789	TGCCCTGGA AGTGTCAAA	GGCTGTACTG TATCCCAGA AGA
<i>Keap1</i>	NM_057152	CAGAACAAGC CATGCCTTCT T	TCTGGTCTTC CACAAGTCC TT
<i>Nos1</i>	NM_052799.1	TCCAATGTTC ACAAAAACG AGTCT	TCCGGTGGAC TTAGGGCTTT
<i>Nos2</i>	NM_012611.3	GGATTTTCCC AGGCAACCA	ACAATCCACA ACTCGCTCCA A
<i>Nos3</i>	NM_021838.2	CACCAGGAAG AAGACTTTTA AGGAA	CACACGCTTC GCCATCAC
<b>Chemical mediators and related molecules</b>			
<i>Nfkb1</i>	NM_001276711.1	CACTAAATCC AACACAGGCA TCAC	GGCACAATCT CTAGGCTCGT TT
<i>Il1a</i>	NM_017019.1	GAGGCCATAG CCCATGATTT AG	TGGAAGCTGT GAGGTGCTGA T
<i>Il1b</i>	NM_031512	TGACAGACCC CAAAAGATTA AGG	CTCATCTGGA CAGCCCAAGT C
<i>Il4</i>	NM_201270	CAAATTTTAC TTCCCACGTG ATGT	CACCCGAGAAC CCCAGACTTG
<i>Il6</i>	NM_012589	GAGGATACCA CTCACAACAG ACC	AAGTGCATCA TCGTTGTTC A TA
<i>Tnf</i>	NM_012675.3	GCCCGAGGCA ACACATCT	CCAGTCCAC ATCTCCGATC A
<b>Interneuron marker genes</b>			
<i>Pvalb</i>	NM_022499	TGCCCACAAA AAGTTCTTCC A	TCTTCACATC ATCCGCACTC TT
<i>Calb2</i>	NM_053988	AGCTCCAGGA GTACACCCAA AC	CCCAATTTGC CGTCTCCAT
<i>Sst</i>	NM_012659.1	ACCCCGGGAA CGCAA	CAGGATGTGA ATGTCTTCCA GAAG
<i>Reln</i>	NM_080394	GCCAGCTTTC GACTACCCTA TTAAC	CGTAGTGGCA CAGAAGCTAT CG
<b>Normalization control genes</b>			
<i>Gapdh</i>	NM_017008	GGCCGAGGGC CCACTA	TGTTGAAGTC ACAGGAGACA ACCT
<i>Hprt1</i>	NM_012583	GCCGACCGGT TCTGTCAT	TCATAACCTG GTTCATCATC ACTAATC

Abbreviations: *Arc*, activity-regulated cytoskeleton-associated protein; *Calb2*, calbindin 2 (also known as calbindin-D-29K); *Egr1*, early growth response protein 1; *Fos*, Fos proto-oncogene, AP-1 transcription factor subunit; *Gapdh*, glyceraldehyde-3-phosphate dehydrogenase; *Gria1*, glutamate ionotropic receptor AMPA type subunit 1; *Gria2*, glutamate ionotropic receptor AMPA type subunit 2; *Gria3*, glutamate ionotropic receptor AMPA type subunit 3; *Grin2a*, glutamate ionotropic receptor NMDA type subunit 2A; *Grin2b*, glutamate ionotropic receptor NMDA type subunit 2B; *Grin2c*, glutamate ionotropic receptor NMDA type subunit 2C; *Grin2d*, glutamate ionotropic receptor NMDA type subunit 2D; *Hprt1*, hypoxanthine phosphoribosyltransferase 1; *Il1a*, interleukin 1 alpha; *Il1b*, interleukin 1 beta; *Il4*, interleukin 4; *Il6*, interleukin 6; *Jun*, Jun proto-oncogene, AP-1 transcription factor subunit; *Keap1*, Kelch-like ECH-associated protein 1; *Nfe2l2*, nuclear factor, erythroid 2-like 2; *Nfkb1*, nuclear factor kappa B subunit 1; *Nos1*, nitric oxide synthase 1; *Nos2*, nitric oxide synthase 2; *Nos3*, nitric oxide synthase 3; *Ptgs2*, prostaglandin-endoperoxide synthase 2; *Pvalb*, parvalbumin; *Reln*, reelin; RT-PCR, reverse transcription-polymerase chain reaction; *Slc17a6*, solute carrier family 17 member 6; *Slc17a7*, solute carrier family 17 member 7; *Sst*, somatostatin; *Tnf*, tumor necrosis factor.

and *Il6* examined using samples of PND 74 animals that were subjected to behavioral tests. In the amygdala, prelimbic cortex and infralimbic cortex, transcript levels of genes encoding glutamate receptors and transporters were measured using samples of PND 77 animals.

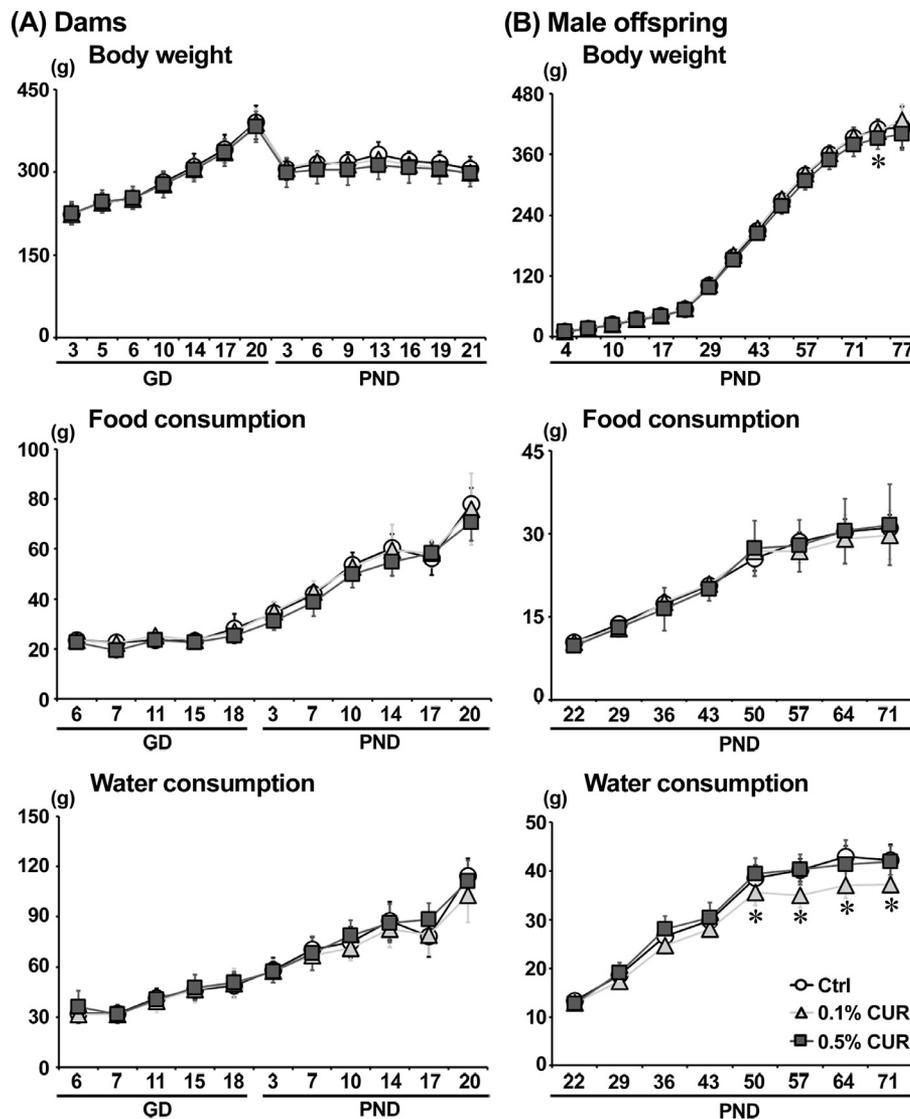
## 2.7. Statistical analyses

Maternal data, such as BWs, food and water consumption were analyzed using the individual animal as the experimental unit. Offspring data, such as body and brain weights at necropsy, the number of immunoreactive cells for each antigen, number of apoptotic cells, and the transcript-level ex-

pression data, were analyzed using the litter as the experimental unit.

Because some animals were resistant to fear-extinction learning, the Smirnov-Grubbs test was performed to detect the outliers in each group. Identified outliers were excluded as abnormal animals from the fear-extinction test analysis. The Smirnov-Grubbs test was performed using Excel Statistics 2013 software package version 2.02 (Social Survey Research Information Co. Ltd., Tokyo, Japan).

Data were analyzed using Levene's test for homogeneity of variance. If the variance was homogenous, numerical data were evaluated using Dunnett's test for comparisons between the vehicle controls and 0.1% CUR or 0.5% CUR. For heteroge-



**Fig. 5 – Body weight, food and water consumption of dams and offspring during experiment. (A) Dams. (B) Male offspring. Values are expressed as means  $\pm$  SD. \* $P < .05$ , compared with the vehicle controls by Dunnett's test or Aspin-Welch's t test with Bonferroni correction.**

neous data, Aspin-Welch's t-test with Bonferroni correction was used. These analyses were performed using IBM SPSS Statistics ver. 25 (International Business Machines Corporation, Armonk, NY, USA), and  $P < 0.05$  was considered statistically significant.

### 3. Results

#### 3.1. Maternal parameters

In the vehicle controls and 0.1% CUR, one each of nonpregnant animal was excluded from the experiment. One pregnant animal each of the vehicle controls and 0.5% CUR was used for measurement of curcumin concentrations in the gastric milk content of offspring. Therefore, the effective numbers of dams were 12, 12, and 13 for the vehicle controls, 0.1% CUR, and 0.5%

CUR, respectively. Dams in each exposure group showed no significant changes in BW and in food and water consumption during gestation and lactation periods compared with the vehicle controls (Fig. 5A).

Based on the mean food consumption values, dams in the 0.1% CUR and 0.5% CUR received 85.8 and 398.9 mg/kg BW/day curcumin, respectively, during the gestation period. Dams consumed 172.3 and 825.8 mg/kg BW/day curcumin during the lactation period, respectively.

#### 3.2. Curcumin concentration in the plasma and gastric content of offspring

Curcumin concentrations in the plasma content of male offspring at PND 21 were below the detection limit (0.1  $\mu\text{g/mL}$ ) in the vehicle controls and 2.78  $\mu\text{g/mL}$  in the 0.5% CUR.

Curcumin concentrations in the gastric milk content of female offspring at PND 14 were below the detection limit (1.0 µg/g) in the vehicle controls and 25.6 µg/g in the 0.5% CUR.

### 3.3. In life parameter data and necropsy data in offspring

No changes in the gait and behaviors were observed in offspring of any exposure group before necropsy on PND 21 and after weaning, but yellow dyeing of the fur and soft stools were observed in the 0.5% CUR animals. BW of male offspring of the 0.5% CUR significantly decreased on PND 71 compared to the vehicle controls ( $P < .05$  by Dunnett's test or Aspin–Welch's t-test with Bonferroni correction; Fig. 5B). There were no significant differences in food consumption in any exposure group. Water consumption was significantly decreased from PND 50 to PND 71 only in the 0.1% CUR compared with the vehicle controls ( $P < .05$  by Dunnett's test or Aspin–Welch's t-test with Bonferroni correction).

Based on the mean food consumption values after weaning, offspring in the 0.1% CUR and 0.5% CUR received 106.4 and 556.0 mg/kg BW/day curcumin, respectively.

### 3.4. Behavioral testing scores in male offspring

#### 3.4.1. OFT

Animals in the 0.1% CUR showed significantly higher center region time compared with the vehicle controls ( $P < .01$  by Dunnett's test or Aspin–Welch's t-test with Bonferroni correction), while the 0.5% CUR did not differ from the vehicle controls (Fig. 6A, Table 4). There were no significant differences in activity scores (total distance and total movement duration) between the vehicle controls and each exposure group.

#### 3.4.2. OLT and ORT

In both tests, the discrimination index in the test phase after a 24 hours interval showed no significantly different memory performance in any exposure group, as the discrimination index was about 0.5 or less (Table 4).

#### 3.4.3. Contextual fear conditioning test

Because there was a certain number of poor learners on contextual fear conditioning, an outlier test was performed and 5 outliers out of 24 animals in the 0.1% CUR and 3 outliers out of 26 animals in the 0.5% CUR were identified, respectively (Table 4).

There were no significant differences in the freezing time in the fear-acquisition test at 24 hours after fear conditioning between the vehicle controls and each exposure group (Fig. 6B, Table 4). Regarding the freezing time in the fear-extinction test, the 0.1% CUR showed a significantly lower rate at the 2nd and 3rd trials compared with the vehicle controls ( $P < .05$  by Dunnett's test or Aspin–Welch's t-test with Bonferroni correction), while the 0.5% CUR did not differ from that of the vehicle controls.

### 3.5. Immunohistochemical results in male offspring

#### 3.5.1. Synaptic plasticity-related molecule<sup>+</sup> cells after the last trial of fear-extinction test

In the hippocampal GCL, the 0.1% CUR showed a significantly higher number of FOS<sup>+</sup> and ARC<sup>+</sup> cells compared with the ve-

hicle controls ( $P < .05$  by Dunnett's test or Aspin–Welch's t-test with Bonferroni correction), while the 0.5% CUR did not differ significantly from the vehicle controls (Fig. 7A, Table 5). There were no significant differences in the number of COX2<sup>+</sup>, EGR1<sup>+</sup>, and p-ERK1/2<sup>+</sup> cells in the GCL between the vehicle controls and each exposure group.

In the prelimbic cortex of the mPFC, the 0.1% CUR showed a significantly lower number of p-ERK1/2<sup>+</sup> cells compared with the vehicle controls ( $P < .05$  by Dunnett's test or Aspin–Welch's t-test with Bonferroni correction), while the number of p-ERK1/2<sup>+</sup> cells in the 0.5% CUR did not differ significantly from the vehicle controls (Fig. 7A, Table 5). There were no significant differences in the number of FOS<sup>+</sup>, ARC<sup>+</sup>, and COX2<sup>+</sup> cells between the vehicle controls and each exposure group.

In the infralimbic cortex of the mPFC, there were no significant differences in the number of FOS<sup>+</sup>, ARC<sup>+</sup>, COX2<sup>+</sup>, and p-ERK1/2<sup>+</sup> cells between the vehicle controls and each exposure group (Table 5).

#### 3.5.2. Neurogenesis-related cell populations and glial cells in the hippocampal dentate gyrus of animals not used for behavioral studies

Regarding granule cell lineage subpopulations, the 0.5% CUR showed a significantly higher number of GFAP<sup>+</sup> cells in the SGZ compared with the vehicle controls ( $P < .05$  by Dunnett's test or Aspin–Welch's t-test with Bonferroni correction), while the 0.1% CUR did not differ from the vehicle controls (Fig. 7B, Table 6). There were no significant differences in the number of SOX2<sup>+</sup> and TBR2<sup>+</sup> cells in the SGZ, and DCX<sup>+</sup>, TUBB3<sup>+</sup>, and NeuN<sup>+</sup> cells in the SGZ and GCL between the vehicle controls and each exposure group. Regarding the PCNA<sup>+</sup> proliferating cells in the SGZ, the 0.5% CUR showed a significantly higher number compared with the vehicle controls ( $P < .05$  by Dunnett's test or Aspin–Welch's t-test with Bonferroni correction), while PCNA<sup>+</sup> cells in the 0.1% CUR did not differ statistically from the vehicle controls. There were no significant differences in the number of TUNEL<sup>+</sup> apoptotic cells in the SGZ between the vehicle controls and each exposure group. Regarding synaptic plasticity-related molecule-immunoreactive cells in the GCL, there were no significant differences in the number of FOS<sup>+</sup> and ARC<sup>+</sup> cells between the vehicle controls and each exposure group. There were no significant differences in the densities of PVALB<sup>+</sup>, CALB2<sup>+</sup>, and SST<sup>+</sup> interneurons, GFAP<sup>+</sup> astrocytes, and IBA1<sup>+</sup> microglia in the hilus of the dentate gyrus between the vehicle controls and each exposure group.

### 3.6. Transcript-level expression in male offspring

#### 3.6.1. Hippocampal dentate gyrus

In the 0.1% CUR, the transcript levels of *Grin2d*, *Slc17a6*, *Fos*, and *Calb2* were significantly increased ( $P < .01$  with *Grin2d*, *Slc17a6*, and *Fos*;  $P < .01$  or  $0.05$  with *Calb2* by Dunnett's test or Aspin–Welch's t-test with Bonferroni correction), while the transcript levels of *Gria1*, *Gria2*, *Slc17a7*, *Arc*, *Ptgs2*, and *Sst* were significantly decreased ( $P < .01$  or  $.05$  with *Arc* and *Sst*;  $P < .05$  with *Gria1*, *Gria2*, *Slc17a7*, and *Ptgs2* by Dunnett's test or Aspin–Welch's t-test with Bonferroni correction) after normalization with *Gapdh* and *Hprt1* compared with the levels in the vehicle controls (Table 7).

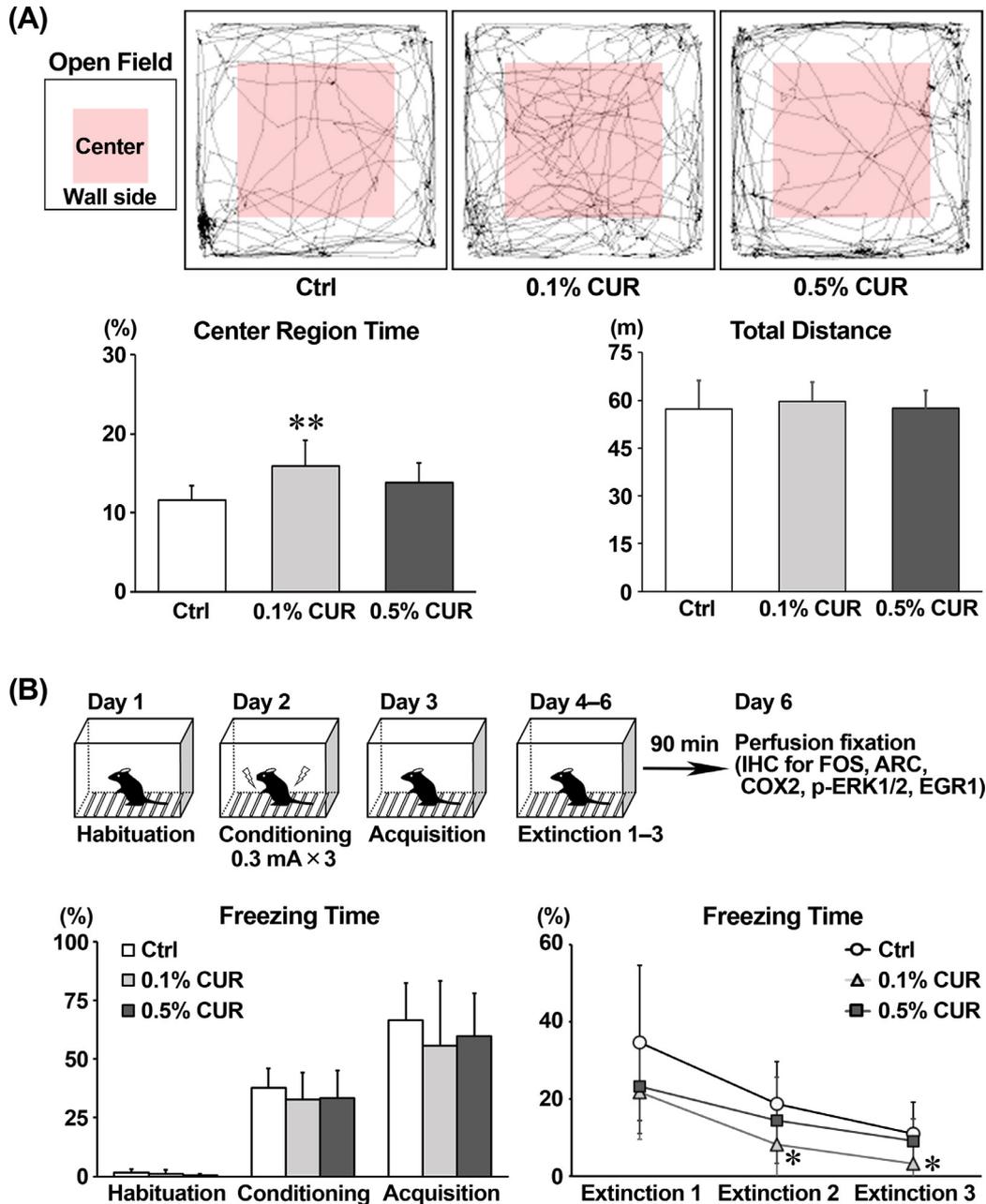


Fig. 6 – Behavioral tests. (A) Open field test. Upper panel shows representative examples of animal track in each group. Graphs show the percent of time in the center region and total distance traveled.  $N = 24$  to  $26$  in each group (2 animals per dam). (B) Contextual fear conditioning test. Upper panel shows experimental design. Graphs show the rate of the freezing time in the habituation, fear conditioning, fear acquisition and fear extinction.  $N = 24$  to  $26$  in each group (2 animals per dam; 5 and 3 animals that were detected as fear-extinction outliers by the Smirnov-Grubbs test were excluded from data analysis in the 0.1% CUR and 0.5% CUR, respectively). Values are expressed as means + SD or means ± SD. \* $P < .05$ , \*\* $P < .01$ , compared with the vehicle controls by Dunnett’s test or Aspin-Welch’s t test with Bonferroni correction.

In the 0.5% CUR, the transcript levels of *Gria3*, *Grin2b*, *Grin2c*, *Fos*, *Keap1*, *Nos2*, *Nos3*, *Nfkb1*, *Il4*, *Il6*, and *Tnf* were significantly increased ( $P < .01$  with *Grin2c*, *Keap1*, *Nos3*, *Nfkb1*, and *Il4*;  $P < .01$  or  $.05$  with *Grin2b*, *Nos2*, and *Tnf*;  $P < .05$  with *Gria3*, *Fos*, and *Il6* by Dunnett’s test or Aspin-Welch’s t-test with Bonferroni correction) after normalization with *Gapdh*

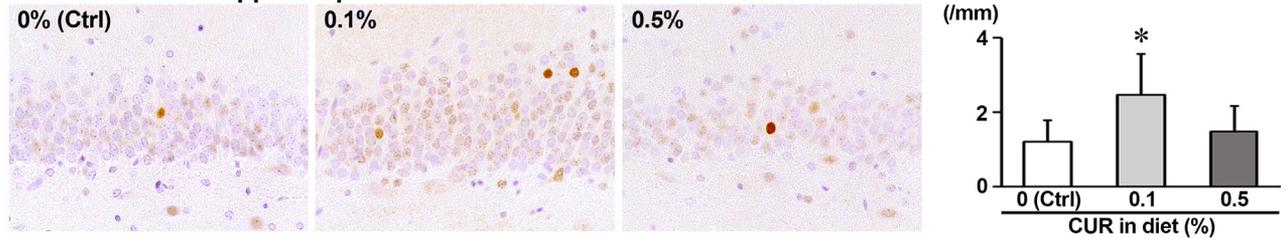
and/or *Hprt1* compared with the levels in the vehicle controls (Table 7).

### 3.6.2. Amygdala

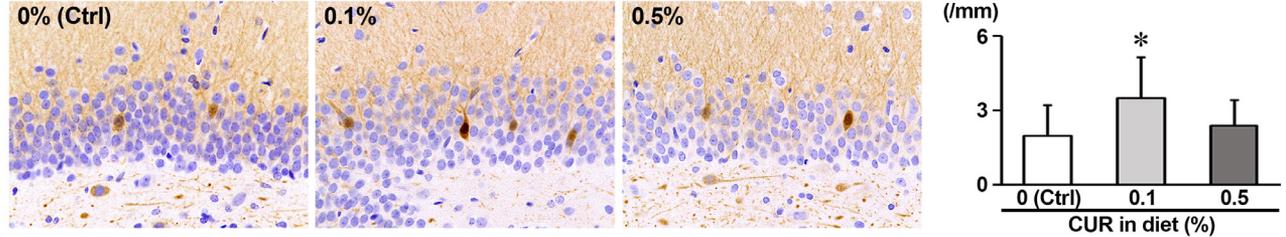
In the 0.1% CUR, the transcript levels of *Gria1*, *Gria2*, *Grin2d*, and *Slc17a6* were significantly increased ( $P < .01$  or  $.05$  with

**(A) Synaptic plasticity-related molecule<sup>+</sup> cells after the last trial of fear extinction test**

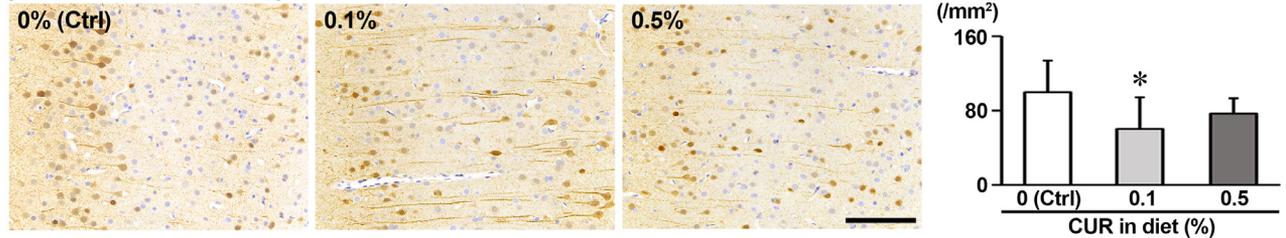
**FOS<sup>+</sup> cells in the hippocampal GCL**



**ARC<sup>+</sup> cells in the hippocampal GCL**

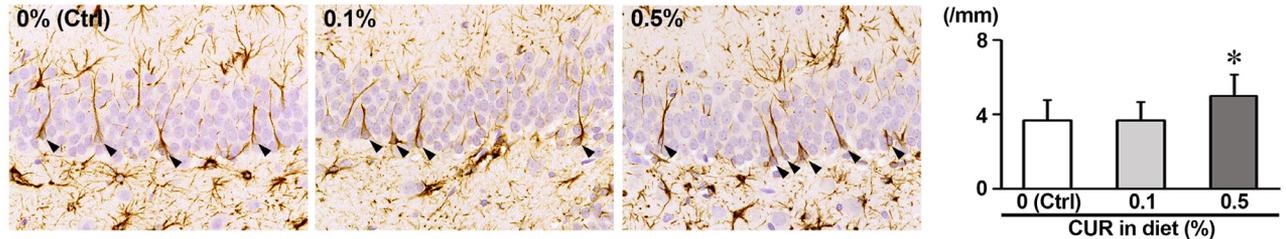


**p-ERK1/2<sup>+</sup> cells in the prelimbic cortex**

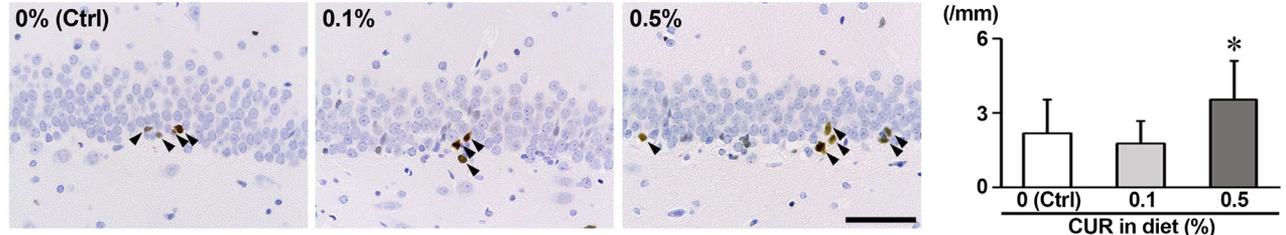


**(B) Neurogenesis-related cell populations in the hippocampal dentate gyrus**

**GFAP<sup>+</sup> cells in the SGZ**



**PCNA<sup>+</sup> cells in the SGZ**



**Fig. 7 – Representative immunohistochemistry results. (A) The number of immunoreactive cells for synaptic plasticity-related molecules in the hippocampal GCL and the prelimbic cortex after the last trial of fear-extinction test. Representative images from the vehicle controls (left), 0.1% CUR group (middle) and 0.5% CUR group (right) on PND 74. Magnification  $\times 400$ ; bar =  $50 \mu\text{m}$ . The graphs show the number of immunoreactive cells/unit length (mm) of the SGZ of the bilateral sides or the number of immunoreactive cells/unit area ( $\text{mm}^2$ ) in the prelimbic cortex of the bilateral sides. Values are expressed as means + SD.  $N = 9$  to  $10$  in each group (1 animal per dam). (B) The number of neurogenesis-related cell populations in the hippocampal dentate gyrus of animals that were not subjected to behavioral tests. Representative images from the vehicle controls (left), 0.1% CUR group (middle) and 0.5% CUR group (right) on PND 77. Arrowheads indicate immunoreactive cells. Magnification  $\times 400$ ; bar  $50 \mu\text{m}$ . The graphs show the number of immunoreactive cells/unit length (mm) of the SGZ of the bilateral sides. Values are expressed as means + SD.  $N = 10$  in each group (1 animal per dam). \* $P < .05$ , compared with the vehicle controls by Dunnett's test or Aspin-Welch's t test with Bonferroni correction.**

**Table 4 – Changes in the behavioral tests in male offspring**

	Ctrl	0.1% CUR	0.5% CUR
<b>Open field test</b>			
No. of offspring examined (2 pups/dam)	24	24	26
Total distance (cm)	5737.2 ± 887.1 <sup>a</sup>	5972.9 ± 606.6	5744.5 ± 564.3
Total movement duration (s)	288.7 ± 33.9	304.6 ± 27.4	291.0 ± 22.5
Wall side time (s)	530.8 ± 11.6	504.2 ± 19.0**	517.3 ± 14.8
Center region time (s)	69.2 ± 11.6	95.8 ± 19.0**	82.8 ± 14.8
<b>Object location test</b>			
No. of offspring examined (1 pup/dam)	12	12	13
Total distance (cm)			
Sample phase	3599.8 ± 529.0	3688.7 ± 460.4	3578.6 ± 306.1
Test phase	2205.5 ± 315.4	2446.1 ± 315.9	2302.6 ± 314.3
Discrimination index			
Sample phase	0.47 ± 0.15	0.49 ± 0.08	0.51 ± 0.14
Test phase	0.52 ± 0.12	0.48 ± 0.17	0.43 ± 0.15
<b>Object recognition test</b>			
No. of offspring examined (1 pup/dam)	12	12	13
Total distance (cm)			
Sample phase	3569.0 ± 390.1	3799.0 ± 400.6	3577.6 ± 388.0
Test phase	2216.6 ± 303.3	2431.4 ± 321.6	2386.8 ± 149.6
Discrimination index			
Sample phase	0.48 ± 0.11	0.42 ± 0.13	0.44 ± 0.07
Test phase	0.54 ± 0.11	0.49 ± 0.10	0.50 ± 0.11
<b>Contextual fear conditioning test</b>			
No. of offspring examined (1-2 pups/dam)	24	24 (19) <sup>b</sup>	26 (23) <sup>c</sup>
Freezing time (%)			
Day 1	Habituation	1.7 ± 1.5	1.2 ± 1.6
Day 2	Fear conditioning	37.7 ± 8.3	32.8 ± 11.4
Day 3	Fear acquisition	66.4 ± 15.9	55.6 ± 27.6
Day 4	Fear extinction 1	34.5 ± 20.1	21.7 ± 14.5
Day 5	Fear extinction 2	18.6 ± 11.1	8.1 ± 3.6*
Day 6	Fear extinction 3	10.9 ± 8.2	3.3 ± 1.5*

Abbreviations: Ctrl, vehicle controls; CUR, amorphous formula of curcumin.

<sup>a</sup> Means ± SD.

<sup>b</sup> Five animals that were detected as fear-extinction outliers by the Smirnov-Grubbs test were excluded from data analysis. The number in parenthesis is the effective number of animals.

<sup>c</sup> Three animals that were detected as fear-extinction outliers by the Smirnov-Grubbs test were excluded from data analysis. The number in parenthesis is the effective number of animals.\* $P < .05$ , \*\* $P < .01$ , compared with the vehicle controls by Dunnett's test or Aspin-Welch's t-test with Bonferroni correction.

*Grin2d*;  $P < .05$  with *Gria1*, *Gria2*, and *Slc17a6* by Dunnett's test or Aspin-Welch's t-test with Bonferroni correction) after normalization with *Gapdh* and/or *Hprt1* compared with the levels in the vehicle controls (Table 8).

In the 0.5% CUR, the transcript level of *Grin2d* was significantly increased ( $P < .05$  by Dunnett's test or Aspin-Welch's t-test with Bonferroni correction) after normalization with *Gapdh* and *Hprt1* compared with the levels in the vehicle controls (Table 8).

### 3.6.3. mPFC

In the 0.1% CUR, the transcript levels of *Gria1*, *Grin2c*, and *Slc17a6* were significantly increased ( $P < .01$  with *Gria1* and *Grin2c*;  $P < .05$  with *Slc17a6* by Dunnett's test or Aspin-Welch's t-test with Bonferroni correction) in the prelimbic cortex, while the transcript level of *Grin2a* was significantly decreased ( $P < .05$  by Dunnett's test or Aspin-Welch's t-test with Bonferroni correction) in this region after normalization with *Gapdh* or *Hprt1* compared with the levels in the vehicle controls (Table 8). In the infralimbic cortex, the transcript level of *Slc17a6* was significantly increased ( $P < .01$  by Dunnett's test

or Aspin-Welch's t-test with Bonferroni correction), while the transcript levels of *Grin2a*, *Grin2b*, *Grin2d*, and *Slc17a7* were significantly decreased ( $P < .05$  with *Grin2a*, *Grin2b*, *Grin2d*, and *Slc17a7* by Dunnett's test or Aspin-Welch's t-test with Bonferroni correction) after normalization with *Gapdh* and/or *Hprt1* compared with the levels in the vehicle controls. In the 0.5% CUR, the transcript level of *Gria1* was significantly increased ( $P < .05$  by Dunnett's test or Aspin-Welch's t-test with Bonferroni correction) in the prelimbic cortex after normalization with *Gapdh* and *Hprt1* compared with the levels in the vehicle controls (Table 8). In the infralimbic cortex, there were no genes encoding glutamate receptors and transporters to change the transcript level compared with the levels in the vehicle controls.

## 4. Discussion

We recently found that continued exposure to  $\alpha$ -glycosyl isoquercitrin (AGIQ), a polyphenolic flavonol glycoside derived by enzymatic glycosylation of rutin, from developmental

**Table 5 – Number of immunoreactive cells for synaptic plasticity-related molecules after the last trial of fear extinction test in male offspring**

	Ctrl	0.1% CUR	0.5% CUR
<b>Hippocampal GCL (No./mm SGZ length)</b>			
No. of animals examined	10	10	10
FOS	1.21 ± 0.57 <sup>a</sup>	2.47 ± 1.10*	1.50 ± 0.68
ARC	1.98 ± 1.23	3.50 ± 1.64*	2.40 ± 1.04
COX2	6.26 ± 2.03	7.43 ± 3.27	5.76 ± 2.80
EGR1	9.19 ± 3.09	9.05 ± 4.19	8.56 ± 3.76
p-ERK1/2	3.47 ± 1.27	3.84 ± 1.95	3.44 ± 1.41
<b>Prelimbic cortex (No./mm<sup>2</sup> measured area)</b>			
No. of animals examined	10	9	10
FOS	13.49 ± 6.89	17.29 ± 8.66	17.26 ± 8.26
ARC	75.40 ± 38.75	90.73 ± 32.13	74.60 ± 25.80
COX2	10.69 ± 6.79	14.03 ± 12.04	15.81 ± 13.03
p-ERK1/2	100.68 ± 33.16	60.33 ± 34.10*	77.20 ± 16.46
<b>Infralimbic cortex (No./mm<sup>2</sup> measured area)</b>			
No. of animals examined	9	9	10
FOS	9.35 ± 3.01	18.01 ± 8.59	15.93 ± 10.03
ARC	42.98 ± 18.06	58.21 ± 38.89	33.37 ± 17.73
COX2	7.96 ± 4.71	11.24 ± 6.47	14.54 ± 11.15
p-ERK1/2	138.31 ± 51.04	119.17 ± 41.42	136.23 ± 17.14

Abbreviations: ARC, activity-regulated cytoskeleton-associated protein; COX2, cyclooxygenase-2; Ctrl, vehicle control; CUR, amorphous formula of curcumin; FOS, Fos proto-oncogene, AP-1 transcription factor subunit; GCL, granule cell layer; EGR1, early growth response protein 1; p-ERK1/2, phosphorylated extracellular signal-regulated kinase 1/2 (phosphorylated p44/p42 MAP kinase); SGZ, subgranular zone.

<sup>a</sup> Means ± SD.

\* P < .05, compared with the vehicle controls by Dunnett's test or Aspin–Welch's t test with Bonferroni correction.

**Table 6 – Number of immunoreactive cells in the SGZ/GCL and hilar region of the hippocampal dentate gyrus of male offspring not subjected to behavioral tests**

	Ctrl	0.1% CUR	0.5% CUR
No. of animals examined	10	10	10
<b>Granule cell lineage subpopulations (No./mm SGZ length)</b>			
GFAP	3.70 ± 1.08 <sup>a</sup>	3.71 ± 0.97	5.02 ± 1.13*
SOX2	14.17 ± 3.43	15.30 ± 4.51	13.73 ± 4.91
TBR2	2.56 ± 1.76	2.75 ± 1.98	3.34 ± 1.90
DCX	8.91 ± 3.65	11.94 ± 4.13	9.80 ± 3.37
TUBB3	2.28 ± 1.67	3.41 ± 2.71	4.48 ± 3.00
NeuN	521.04 ± 42.24	552.84 ± 57.37	505.89 ± 52.88
<b>Cell proliferation and apoptosis (No./mm SGZ length)</b>			
PCNA	2.20 ± 1.34	1.77 ± 0.91	3.56 ± 1.54*
TUNEL	0.18 ± 0.20	0.07 ± 0.11	0.16 ± 0.22
<b>Synaptic plasticity-related molecules (No./mm SGZ length)</b>			
FOS	1.03 ± 0.40	1.02 ± 0.41	1.11 ± 0.64
ARC	2.66 ± 1.09	3.44 ± 1.54	2.17 ± 1.04
<b>Interneuron subpopulations (No./mm<sup>2</sup> hilar region)</b>			
PVALB	17.41 ± 4.43	19.58 ± 5.16	23.83 ± 10.71
CALB2	1.37 ± 1.38	0.82 ± 1.11	1.43 ± 1.15
SST	33.94 ± 15.32	36.04 ± 12.10	24.10 ± 10.83
<b>Astrocytes and microglia (No./mm<sup>2</sup> hilar region)</b>			
GFAP	191.03 ± 23.55	179.85 ± 23.72	194.65 ± 29.39
IBA1	100.94 ± 20.51	88.95 ± 11.04	115.23 ± 24.36

Abbreviations: ARC, activity-regulated cytoskeleton-associated protein; CALB2, calbindin-D-29 K; Ctrl, vehicle control; CUR, amorphous formula of curcumin; DCX, doublecortin; FOS, Fos proto-oncogene, AP-1 transcription factor subunit; GCL, granule cell layer; GFAP, glial fibrillary acidic protein; IBA1, ionized calcium-binding adaptor molecule 1; NeuN, neuronal nuclei; PCNA, proliferating cell nuclear antigen; PVALB, parvalbumin; SGZ, subgranular zone; SOX2, SRY-box transcription factor 2; SST, somatostatin; TBR2, T-box brain protein 2; TUBB3, tubulin, beta 3 class III; TUNEL, terminal deoxynucleotidyl transferase-mediated dUTP-biotin nick end-labeling.

<sup>a</sup> Means ± SD.

\* P < .05, compared with the vehicle controls by Dunnett's test or Aspin–Welch's t test with Bonferroni correction.

**Table 7 – Transcript level expression changes in the hippocampal dentate gyrus of male offspring not subjected to behavioral tests**

Normalization control No. of animals examined	Ctrl		0.1% CUR		0.5% CUR	
	<i>Gapdh</i> 6	<i>Hprt1</i> 6	<i>Gapdh</i> 6	<i>Hprt1</i> 6	<i>Gapdh</i> 6	<i>Hprt1</i> 6
<b>Glutamate receptors</b>						
<i>Gria1</i>	1.08 ± 0.45 <sup>a</sup>	1.07 ± 0.41	0.40 ± 0.16*	0.40 ± 0.18*	1.10 ± 0.65	1.14 ± 0.60
<i>Gria2</i>	1.06 ± 0.41	1.06 ± 0.36	0.45 ± 0.15*	0.45 ± 0.17*	1.11 ± 0.62	1.15 ± 0.51
<i>Gria3</i>	1.03 ± 0.27	1.03 ± 0.25	0.85 ± 0.15	0.86 ± 0.22	1.26 ± 0.37	1.36 ± 0.18*
<i>Grin2a</i>	1.06 ± 0.37	1.05 ± 0.33	0.72 ± 0.17	0.72 ± 0.21	1.37 ± 0.61	1.45 ± 0.50
<i>Grin2b</i>	1.04 ± 0.33	1.05 ± 0.31	1.09 ± 0.14	1.11 ± 0.27	1.45 ± 0.19*	1.61 ± 0.18**
<i>Grin2c</i>	1.03 ± 0.26	1.02 ± 0.20	1.42 ± 0.16	1.43 ± 0.27	2.02 ± 0.56**	2.35 ± 1.16**
<i>Grin2d</i>	1.03 ± 0.28	1.04 ± 0.30	2.17 ± 0.43**	2.17 ± 0.44**	1.41 ± 0.57	1.62 ± 0.87
<b>Glutamate transporters</b>						
<i>Slc17a6</i>	1.16 ± 0.58	1.18 ± 0.65	4.09 ± 1.33**	4.04 ± 1.18**	1.52 ± 0.75	1.80 ± 1.17
<i>Slc17a7</i>	1.03 ± 0.25	1.02 ± 0.24	0.58 ± 0.25*	0.58 ± 0.24*	0.82 ± 0.38	0.87 ± 0.36
<b>Synaptic plasticity-related genes</b>						
<i>Fos</i>	1.06 ± 0.36	1.08 ± 0.46	2.76 ± 0.84**	2.73 ± 0.75**	2.11 ± 0.67*	2.46 ± 1.15*
<i>Jun</i>	1.03 ± 0.28	1.01 ± 0.19	0.84 ± 0.18	0.85 ± 0.26	0.90 ± 0.26	0.98 ± 0.20
<i>Arc</i>	1.05 ± 0.37	1.04 ± 0.29	0.50 ± 0.19*	0.50 ± 0.17**	1.06 ± 0.40	1.13 ± 0.31
<i>Ptgs2</i>	1.08 ± 0.44	1.07 ± 0.40	0.38 ± 0.25*	0.39 ± 0.25*	0.87 ± 0.51	0.90 ± 0.45
<i>Egr1</i>	1.02 ± 0.21	1.04 ± 0.31	1.08 ± 0.23	1.08 ± 0.23	1.25 ± 0.23	1.37 ± 0.16
<b>Oxidative stress-related genes</b>						
<i>Nfe2l2</i>	1.05 ± 0.39	1.03 ± 0.27	1.00 ± 0.14	1.03 ± 0.29	0.97 ± 0.05	1.09 ± 0.22
<i>Keap1</i>	1.01 ± 0.18	1.02 ± 0.22	1.26 ± 0.36	1.31 ± 0.40	2.00 ± 0.36**	1.25 ± 0.20
<i>Nos1</i>	1.18 ± 0.71	1.16 ± 0.68	0.65 ± 0.21	0.66 ± 0.27	1.32 ± 0.66	1.39 ± 0.56
<i>Nos2</i>	1.31 ± 0.97	1.48 ± 1.28	4.47 ± 2.63	4.59 ± 2.88	37.80 ± 15.52**	43.93 ± 26.96*
<i>Nos3</i>	1.05 ± 0.33	1.03 ± 0.26	1.41 ± 0.30	1.44 ± 0.49	2.50 ± 0.47**	2.85 ± 0.99**
<b>Chemical mediators and related molecules</b>						
<i>Nfkb1</i>	1.06 ± 0.42	1.02 ± 0.21	1.22 ± 0.57	1.22 ± 0.42	2.00 ± 0.41**	1.26 ± 0.29
<i>Il1a</i>	1.13 ± 0.63 <sup>b</sup>	1.05 ± 0.34	1.11 ± 0.66	1.03 ± 0.76	1.49 ± 0.83	0.99 ± 0.49
<i>Il1b</i>	1.12 ± 0.62 <sup>b</sup>	1.05 ± 0.36	0.84 ± 0.59	0.74 ± 0.59	1.10 ± 0.21	0.75 ± 0.12
<i>Il4</i>	1.10 ± 0.53 <sup>b</sup>	1.03 ± 0.26	1.10 ± 0.43	0.98 ± 0.33	2.64 ± 0.52**	1.82 ± 0.41**
<i>Il6</i>	1.47 ± 1.28 <sup>b</sup>	1.41 ± 1.04	2.28 ± 1.29	1.99 ± 0.96	3.38 ± 1.46*	2.32 ± 1.08
<i>Tnf</i>	1.32 ± 0.95	1.22 ± 0.64	1.28 ± 1.36	1.14 ± 0.83	14.81 ± 6.47**	9.39 ± 4.60*
<b>Interneuron marker genes</b>						
<i>Pvalb</i>	1.40 ± 1.57	1.23 ± 1.01	0.93 ± 0.37	0.98 ± 0.43	1.25 ± 0.71	0.79 ± 0.51
<i>Calb2</i>	1.07 ± 0.39	1.07 ± 0.44	2.22 ± 0.65*	2.22 ± 0.56**	1.21 ± 0.92	1.28 ± 0.76
<i>Sst</i>	1.06 ± 0.36	1.02 ± 0.23	0.40 ± 0.19*	0.41 ± 0.20**	0.69 ± 0.48	0.69 ± 0.39
<i>Reln</i>	1.08 ± 0.44	1.03 ± 0.26	0.89 ± 0.23	0.91 ± 0.30	0.86 ± 0.39	0.91 ± 0.23

Abbreviations: *Arc*, activity-regulated cytoskeleton-associated protein; *Calb2*, calbindin 2 (also known as calbindin-D-29K); *Ctrl*, vehicle control; *CUR*, amorphous formula of curcumin; *Egr1*, early growth response protein 1; *Fos*, Fos proto-oncogene, AP-1 transcription factor subunit; *Gapdh*, glyceraldehyde-3-phosphate dehydrogenase; *Gria1*, glutamate ionotropic receptor AMPA type subunit 1; *Gria2*, glutamate ionotropic receptor AMPA type subunit 2; *Gria3*, glutamate ionotropic receptor AMPA type subunit 3; *Grin2a*, glutamate ionotropic receptor NMDA type subunit 2A; *Grin2b*, glutamate ionotropic receptor NMDA type subunit 2B; *Grin2c*, glutamate ionotropic receptor NMDA type subunit 2C; *Grin2d*, glutamate ionotropic receptor NMDA type subunit 2D; *Hprt1*, hypoxanthine phosphoribosyltransferase 1; *Il1a*, interleukin 1 alpha; *Il1b*, interleukin 1 beta; *Il4*, interleukin 4; *Il6*, interleukin 6; *Jun*, Jun proto-oncogene, AP-1 transcription factor subunit; *Keap1*, Kelch-like ECH-associated protein 1; *Nfe2l2*, nuclear factor, erythroid 2-like 2; *Nfkb1*, nuclear factor kappa B subunit 1; *Nos1*, nitric oxide synthase 1; *Nos2*, nitric oxide synthase 2; *Nos3*, nitric oxide synthase 3; *Ptgs2*, prostaglandin-endoperoxide synthase 2; *Pvalb*, parvalbumin; *Reln*, reelin; *Slc17a6*, solute carrier family 17 member 6; *Slc17a7*, solute carrier family 17 member 7; *Sst*, somatostatin; *Tnf*, tumor necrosis factor.

<sup>a</sup> Means ± SD.

<sup>b</sup> Values were obtained from animals that were subjected to behavioral tests. \**P* < .05, \*\**P* < .01, compared with the vehicle controls by Dunnett's test or Aspin-Welch's *t* test with Bonferroni correction.

stage facilitated fear-extinction learning and upregulated the constitutive expression level of *Fos* in the hippocampal dentate gyrus and *Grin2d* in the amygdala [28]. In the present study, we tested CUR that has enabled an improved bioavailability after oral intake and found that exposure of rats to 0.1% CUR led to an increase in the time spent in the center region of the open field test device without changing motor activity in an OFT, suggesting an antianxiety-like effect,

and facilitation of fear-extinction learning in the contextual fear conditioning test. However, there were no changes in the OLT or the ORT that evaluated spatial and nonspatial memory. Regarding the changes in the constitutive gene expression level, 0.1% CUR upregulated *Fos* in the hippocampal dentate gyrus and glutamate receptors, represented by *Grin2d* in the dentate gyrus and amygdala. While we did not perform OFT in our previous AGIQ study, these results

**Table 8 – Transcript level expression changes of glutamate receptors and transporters in the amygdala, prelimbic cortex and infralimbic cortex of male offspring not subjected to behavioral tests**

Normalization control No. of animals examined	Ctrl		0.1% CUR		0.5% CUR	
	<i>Gapdh</i> 6	<i>Hprt1</i> 6	<i>Gapdh</i> 6	<i>Hprt1</i> 6	<i>Gapdh</i> 6	<i>Hprt1</i> 6
<b>Amygdala</b>						
<i>Gria1</i>	1.01 ± 0.13 <sup>a</sup>	1.01 ± 0.12	1.29 ± 0.15*	1.19 ± 0.22	1.20 ± 0.26	1.14 ± 0.28
<i>Gria2</i>	1.01 ± 0.13	1.00 ± 0.10	1.18 ± 0.11*	1.08 ± 0.16	1.15 ± 0.08	1.09 ± 0.11
<i>Gria3</i>	1.03 ± 0.25	1.01 ± 0.18	1.14 ± 0.12	1.05 ± 0.18	1.15 ± 0.11	1.08 ± 0.11
<i>Grin2a</i>	1.01 ± 0.18	1.01 ± 0.13	1.08 ± 0.16	1.00 ± 0.23	1.16 ± 0.16	1.10 ± 0.20
<i>Grin2b</i>	1.01 ± 0.16	1.01 ± 0.12	1.10 ± 0.10	1.02 ± 0.17	1.15 ± 0.07	1.09 ± 0.10
<i>Grin2c</i>	1.01 ± 0.17	1.01 ± 0.16	1.02 ± 0.08	0.94 ± 0.13	1.01 ± 0.11	0.96 ± 0.13
<i>Grin2d</i>	1.04 ± 0.30	1.04 ± 0.33	2.22 ± 0.47**	2.02 ± 0.38*	2.06 ± 0.86*	1.95 ± 0.83*
<i>Slc17a6</i>	1.02 ± 0.23	1.02 ± 0.23	1.40 ± 0.19*	1.28 ± 0.22	1.29 ± 0.31	1.23 ± 0.34
<i>Slc17a7</i>	1.04 ± 0.28	1.02 ± 0.22	1.06 ± 0.24	0.99 ± 0.31	1.30 ± 0.27	1.23 ± 0.26
<b>Prelimbic cortex</b>						
<i>Gria1</i>	1.02 ± 0.24	1.00 ± 0.06	1.00 ± 0.30	1.21 ± 0.12**	1.45 ± 0.20*	1.18 ± 0.12*
<i>Gria2</i>	1.03 ± 0.24	1.00 ± 0.07	0.81 ± 0.16	1.01 ± 0.13	1.26 ± 0.23	1.02 ± 0.15
<i>Gria3</i>	1.03 ± 0.25	1.00 ± 0.09	0.79 ± 0.18	0.99 ± 0.14	1.14 ± 0.23	0.94 ± 0.24
<i>Grin2a</i>	1.03 ± 0.25	1.01 ± 0.15	0.62 ± 0.12*	0.78 ± 0.17	1.01 ± 0.44	0.83 ± 0.37
<i>Grin2b</i>	1.02 ± 0.22	1.00 ± 0.07	0.82 ± 0.18	1.02 ± 0.12	1.26 ± 0.30	1.03 ± 0.26
<i>Grin2c</i>	1.05 ± 0.36	1.02 ± 0.21	1.22 ± 0.34	1.50 ± 0.11**	1.38 ± 0.27	1.12 ± 0.13
<i>Grin2d</i>	1.05 ± 0.35	1.01 ± 0.17	0.69 ± 0.21	0.84 ± 0.04	1.44 ± 0.32	1.17 ± 0.26
<i>Slc17a6</i>	1.03 ± 0.28	1.01 ± 0.17	1.42 ± 0.62	1.66 ± 0.33*	1.59 ± 0.63	1.29 ± 0.46
<i>Slc17a7</i>	1.02 ± 0.21	1.00 ± 0.07	0.78 ± 0.22	0.98 ± 0.21	1.12 ± 0.35	0.94 ± 0.34
<b>Infralimbic cortex</b>						
<i>Gria1</i>	1.02 ± 0.19	1.00 ± 0.11	1.19 ± 0.27	1.29 ± 0.25	1.39 ± 0.35	1.08 ± 0.28
<i>Gria2</i>	1.02 ± 0.19	1.00 ± 0.09	1.07 ± 0.23	1.17 ± 0.20	1.21 ± 0.24	0.95 ± 0.22
<i>Gria3</i>	1.02 ± 0.21	1.01 ± 0.15	0.98 ± 0.17	1.06 ± 0.11	1.10 ± 0.08	0.88 ± 0.23
<i>Grin2a</i>	1.02 ± 0.24	1.01 ± 0.13	0.74 ± 0.16*	0.80 ± 0.05*	0.94 ± 0.18	0.76 ± 0.23
<i>Grin2b</i>	1.02 ± 0.23	1.01 ± 0.16	0.85 ± 0.15*	0.93 ± 0.08	1.01 ± 0.12	0.80 ± 0.19
<i>Grin2c</i>	1.06 ± 0.37	1.03 ± 0.26	1.18 ± 0.30	1.29 ± 0.33	1.45 ± 0.49	1.14 ± 0.43
<i>Grin2d</i>	1.03 ± 0.27	1.02 ± 0.19	0.71 ± 0.15*	0.77 ± 0.08	1.32 ± 0.21	1.05 ± 0.27
<i>Slc17a6</i>	1.05 ± 0.37	1.03 ± 0.27	1.45 ± 0.31	1.57 ± 0.18**	1.65 ± 0.74	1.29 ± 0.60
<i>Slc17a7</i>	1.01 ± 0.16	1.00 ± 0.10	0.70 ± 0.24	0.76 ± 0.19*	0.88 ± 0.26	0.72 ± 0.29

Abbreviations: Ctrl, vehicle control; CUR, amorphous formula of curcumin; *Gapdh*, glyceraldehyde-3-phosphate dehydrogenase; *Gria1*, glutamate ionotropic receptor AMPA type subunit 1; *Gria2*, glutamate ionotropic receptor AMPA type subunit 2; *Gria3*, glutamate ionotropic receptor AMPA type subunit 3; *Grin2a*, glutamate ionotropic receptor NMDA type subunit 2A; *Grin2b*, glutamate ionotropic receptor NMDA type subunit 2B; *Grin2c*, glutamate ionotropic receptor NMDA type subunit 2C; *Grin2d*, glutamate ionotropic receptor NMDA type subunit 2D; *Hprt1*, hypoxanthine phosphoribosyltransferase 1; *Slc17a6*, solute carrier family 17 member 6; *Slc17a7*, solute carrier family 17 member 7.

<sup>a</sup> Means ± SD.\*P < .05, \*\*P < .01, compared with the vehicle controls by Dunnett's test or Aspin-Welch's t test with Bonferroni correction.

suggest that low doses of CUR may cause similar effects to AGIQ in normal animals by exposure from developmental stage.

In the present study, CUR did not show any effect on fear acquisition. In the fear-extinction test, 0.1% CUR revealed decrease in freezing time during the second and third trials of fear extinction, as compared with the vehicle controls. These results suggest that continued CUR exposure for long term from developmental stage does not alter the formation of fear memory but promotes learning of fear extinction at low dietary concentrations. Continued exposure to environmental stressors of mice results in long-term behavioral abnormality that is implicated in stress-related disorders [30]. The present study results may reinforce curcumin as an effective means of extinction-based exposure psychotherapy for prevention of stress-related disorders represented by PTSD. Interestingly, a recent study has shown that dietary treatment with 1.5% curcumin for 5 days is capable of impairing fear-memory consolidation and reconsolidation processes in rats [19]. This result suggests that curcumin may also be useful as an adjunct in the

treatment of stress disorders even with brief oral treatment. Further studies may be warranted to examine the dose-effect relationship on behaviors at lower doses of CUR than those examined in the present study.

In the present study, we found a significant increase in the number of granule cells immunoreactive for FOS and ARC in the hippocampal dentate gyrus by 0.1% CUR among synaptic plasticity-related molecules 90 min after conducting the last trial of fear-extinction learning. In contrast, an increase in FOS<sup>+</sup> or ARC<sup>+</sup> cells was not confirmed at this dose in animals that had not been subjected to behavioral test, whereas constitutive expression of Fos was upregulated in the dentate gyrus. These results suggest an increase in synaptic plasticity mediated by FOS and ARC in association with the facilitation of fear-extinction learning. In the dentate gyrus, the expression of IEG proteins in granule cells suggested that these cells are integrated into the neural network [31]. Increased Fos expression in hippocampal neurons is regulated by learning-related neuronal activity [32], and mutant mice deficient in FOS in the brain show impaired spatial and contextual long-

term memory [33]. ARC also plays a critical role in the maintenance of synaptic change and in consolidation of long-term memory [13]. Moreover, 0.1% CUR exposure decreased p-ERK1/2<sup>+</sup> cells in the prelimbic cortex after the last trial of fear-extinction learning in the present study. In the mPFC, both of the prelimbic cortex and infralimbic cortex project similarly to the amygdala, and prelimbic cortex microstimulation increases conditioned fear development and inhibits extinction of fear memory, in contrast to the opposite function of the infralimbic cortex [34]. While we did not observe the alteration in the number of immunoreactive cells of any synaptic plasticity-related molecule in the infralimbic cortex, the decrease of p-ERK1/2<sup>+</sup> cells in the prelimbic cortex suggests decrease of synaptic plasticity to cause suppression of amygdala-mediated fear response that may be related to facilitation of fear-extinction learning. Thus, changes in the number of synaptic plasticity-related molecule-expressing cells in the hippocampal dentate gyrus and mPFC due to continued exposure to CUR at low doses may cause alterations in synaptic plasticity in the neural circuits involved in suppressing anxiety-like behavior and promoting fear-extinction learning.

In the present study, we examined changes in constitutive expression levels of genes encoding glutamate receptors and transporters in animals that have not been subjected to behavioral tests, and found that the glutamate receptor subunit genes were mostly expressed differently among the brain regions by 0.1% CUR exposure. Among them, *Grin2d* was strongly upregulated in the hippocampal dentate gyrus and amygdala. *Grin2d* encodes an N-methyl-D-aspartate (NMDA) receptor subunit and functions in enhancing the synaptic plasticity associated with long-term memory [35]. In the rat, the synaptic plasticity associated with learning and storage of both conditioned fear and fear extinction occurs within the basolateral amygdala [36]. The mechanism of memory storage in this region is thought to be long-term potentiation (LTP) by NMDA receptor activations [36]. Therefore, constitutive upregulation of *Grin2d* in the amygdala by 0.1% CUR may be involved in enhancing elimination of fear memory in the present study. In the hippocampus, the role of GRIN2D in the formation or resolution of anxiety or fear memory has not been reported until now; however, NMDA receptor hypofunction in the dentate gyrus of adult *Fmr1* knockout mice impaired extinction of contextual fear [37]. In the dentate gyrus, we also observed upregulation of *Calb2* and downregulation of *Sst* by 0.1% CUR in the present study, while the number of CALB2<sup>+</sup> or SST<sup>+</sup> GABAergic interneurons were unchanged in the hilar region. It has been reported that CALB2 expression positively contributes to the control of LTP in learning and memory in the mouse dentate gyrus by indirectly regulating the activity of GABAergic interneurons [38]. Regarding SST, while the precise role on cognitive and emotional functions is still poorly understood, one study reported that SST and the related peptide cortistatin in the dentate gyrus reduced the likelihood of generating LTP, suggestive of their role in increasing the threshold of input required for acquisition of new memories [10]. However, we did not find any influence of CUR on fear-acquisition memory. From these perspectives, altered expression of NMDA glutamate receptors and probably some interneuron proteins may be involved in suppressing anxiety or promoting fear-extinction learning.

Curcumin can attenuate oxidative stress by increasing the activity of antioxidant enzymes and activating the Nrf2-Keap1 pathway; however, the attenuating effect of oxidative stress can be abrogated and results in ROS production at high doses in cultured cells [39]. In the present study, 0.5% CUR upregulated *Keap1* transcripts in the dentate gyrus. At this dose, transcript upregulation of *Nos2* and *Nos3* was also observed in this brain region. While induction of nitric oxide synthase (NOS)-3 is sometimes neuroprotective, NOS2 induction in the brain causes detrimental oxidative damage [40]. These results suggest that 0.5% CUR causes induction of oxidative stress responses at least in the hippocampal dentate gyrus. Considering a highly bioavailable characteristics of CUR, 0.5% CUR dose may result in excess level of absorption to cause toxic effects.

Nuclear factor  $\kappa$ B (NF- $\kappa$ B) is an important upstream modulator of inflammatory cytokines, and inhibition of NF- $\kappa$ B activation suppresses the expression of inflammatory cytokines such as interleukin (IL)-1 $\beta$ , IL-6, and tumor necrosis factor (TNF)- $\alpha$  [41]. Early neonatal exposure to TNF- $\alpha$  increases levels of anxiety and depression-related behaviors [42]. Therefore, *Nfkb1* transcript upregulation by 0.5% CUR in the hippocampal dentate gyrus and also in the mPFC (data not shown for the latter) in the present study might be the signature of increased production of inflammatory cytokines that cause neuroinflammation, as evidenced by transcript upregulation of *Il4*, *Il6*, and *Tnf* in the hippocampal dentate gyrus, and may inhibit the antianxiety-like effect and extinction of fear memory at this dose.

Curcumin at a low concentration promotes neurogenesis in mice and increases proliferation of cultured neural progenitor cells of mice [43]. In the present study, 0.5% CUR increased the number of GFAP<sup>+</sup> cells in the SGZ. In the hippocampus, the GFAP<sup>+</sup> SGZ cells represent type-1 neural stem cells [11]. We also observed an increase in the number of PCNA<sup>+</sup> cells in the SGZ in this group. Because the number of no other subpopulations of granule cell lineage changed, these results suggest that 0.5% CUR causes neural stem cell proliferation. Brain inflammation can cause neural stem/progenitor cells to increase in hippocampal neurogenesis, and activation of microglia plays a role in promotion of neurogenesis [44]. Microglial TNF- $\alpha$  is involved in nerve damage and the neurotrophic effect in hippocampal neurogenesis [23], suggesting that upregulation of *Tnf* by 0.5% CUR may contribute to the proliferation of neural stem cells.

PVP, a polymer commonly used as a stabilizer and suspending agent, was used to increase the solubility of curcumin in the present study. This chemical property may contribute to better bioavailability of curcumin and its therapeutic effect, which is consistent with our results. In the present study, 0.1% CUR showed less water consumption from PND 50 until the end of the experiment compared with the vehicle controls. The increased thirst is considered to be due to the high volume of PVP and sucrose fatty acid ester that was used as the vehicle in the vehicle controls and 0.5% CUR. There was a report that rats bred for 2 years with a diet containing 10% PVP did not show toxic effects or gross lesions [45]. Therefore, we believe that dietary concentrations of PVP in the present study are not toxic to either dams or offspring in any way that affects test results.

In conclusion, CUR at 0.1% facilitated anti-anxiety-like behavior and fear-extinction learning. The induction of synaptic plasticity-related proteins may be linked to the altered behavioral attitudes, but these changes were lacking at the 0.5% CUR concentration, probably due to induction of oxidative stress responses and neuroinflammation that affects hippocampal neurogenesis involving neural stem cells. Thus, CUR may have a better preventive effect on some stress disorders by continuous administration at low doses. In contrast, excessively absorbed doses can negate its positive effects. Further research is needed to explore the detailed mechanisms by which curcumin prevents stress disorder. The prospect of using CUR as a new effective drug for stress disorders seems promising.

### CRedit authorship contribution statement

**Junta Nakahara:** methodology, validation, formal analysis, investigation, data curation, writing - original draft, visualization. **Yasunori Masubuchi:** investigation, writing - review and editing. **Kazumi Takashima:** investigation, writing - review & editing. **Yasunori Takahashi:** investigation, writing - review and editing. **Ryo Ichikawa:** investigation, writing - review and editing. **Tomohiro Nakao:** resources, investigation, writing - review and editing. **Mihoko Koyanagi:** resources, investigation, writing - review and editing. **Robert R Maronpot:** writing - review and editing. **Toshinori Yoshida:** writing - review and editing. **Shim-mo Hayashi:** resources, writing - review and editing, project administration. **Makoto Shibutani:** conceptualization, writing - review and editing, visualization, supervision, funding acquisition.

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