

# Ginsenoside Rb1 Induces a Pro-Neurogenic Microglial Phenotype via PPAR $\gamma$ Activation in Male Mice Exposed to Chronic Mild Stress

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## Research Article

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# Abstract

## Background

Anti-inflammatory approaches are emerging as a new strategy for treatment of depressive disorders. Ginsenoside Rb1 (GRb1), a major component of *Panax ginseng*, can inhibit inflammatory cascade and alleviate depressive behaviors. Microglia can promote or inhibit adult hippocampal neurogenesis according to their functional phenotypes. Here, we examined whether GRb1 may exert antidepressant effects by promoting a pro-neurogenic phenotype of microglia and thereby increasing neurogenesis.

## Methods

The antidepressant effects of GRb1 or the licensed antidepressant imipramine (IMI) were assessed in chronic mild stress (CMS)-exposed male mice. The depressive-like behaviors of mice were evaluated by sucrose preference test, forced swimming test (FST), and tail suspension test (TST). The microglial phenotypes were identified by molecular markers and morphological properties, analyzed by RT-qPCR, western blotting and immunofluorescence staining. Effect of GRb1-treated microglia on adult hippocampal neurogenesis *in vivo* and *in vitro* were detected using immunofluorescence staining.

## Results

Behavioral assessment indicated that GRb1 or IMI treatment alleviated depressive-like behaviors in CMS-exposed mice. Immunofluorescence examination

demonstrated that GRb1 induced a pro-neurogenic phenotype of microglia via activating PPAR $\gamma$  *in vivo* and *in vitro*, which were reversed by PPAR $\gamma$  inhibitor GW9662. In addition, GRb1-treated microglia increased the proliferation and differentiation of neural precursor cells.

## Conclusions

These findings demonstrated that GRb1 alleviated depressive-like behaviors of CMS-exposed male mice mainly through PPAR $\gamma$ -mediated microglial activation and improvement of adult hippocampus neurogenesis.

## 1. Introduction

Major depressive disorder (MDD), a major public health burden, remains underdiagnosed and undertreated [1]. As a disease involving neuroplasticity dysfunction, MDD is related to chronic neuroinflammation impairment [2]. The imbalance of inflammatory process was observed in rodent models of stress-induced depression, with the increase of pro-inflammatory cytokines interleukin (IL)-1 $\beta$ , IL-6, and IL-18 and the decrease of anti-inflammatory cytokines IL-10, transforming growth factor (TGF)- $\beta$ , IL-4 [2–4].

Microglia play a key role in immune surveillance of the central nervous system (CNS). Microglia can be activated to show the classical M1 phenotype, associated with up-regulation of pro-inflammatory mediators or the alternative M2 phenotype, associated with up-regulation of anti-inflammatory cytokines [5]. Depending on their phenotypes, activated microglia exert different effects on the differentiation of neural precursor cells (NPCs) *in vivo* and *in vitro*. M1 microglia impair survival and proliferation of NPCs [6], whereas M2 microglia increase the generation of new neurons [7].

There is a close relationship between pathogenesis of depression and the impairment of adult hippocampal neurogenesis. In depression patients, hippocampus atrophy has been reported [8, 9]. In animal models of depression, the depressive-like behaviors are associated with the impairment of adult hippocampal neurogenesis [10]. Neuronal plasticity within the hippocampus is thought to play a significant role in neurobiological responses to stress. Therefore, modulating the switch of microglial phenotypes from neurotoxicity to pro-neurogenesis should be a strategy for treating depression. It has been reported that anti-inflammatory agents such as minocycline inhibited the microglial M1 phenotype [11]. The activation of M2 microglia by anti-inflammatory herbal medicine salvianolic acid B, enhanced neurogenesis and ameliorated depressive-like behaviors [4].

Traditional Chinese medicines are widely used in China and other countries for treatment of mental disorders [12]. Ginseng has been widely applied to treat various conditions in China and other countries over 5000 years, with beneficial effects on immune functions and stress resilience [13, 14]. It was shown high safety of ginseng in clinical prescription [15]. Clinical observations show that ginseng plays significant antidepressant effect [16]. Ginsenoside Rb1 (GRb1), one of main components of Ginseng, has various neuropharmacological effects, including modulating monoamine neurotransmitters, reconstructing neuronal plasticity, regulating the function of hypothalamic-pituitary-adrenal axis, and anti-inflammatory activities [17, 18]. GRb1 exerts significant antidepressant effects in chronic mild stress (CMS)-exposed rodents [19, 20]. However, few studies have focused on the relationship between the antidepressant effects of GRb1 and microglia-mediated neuroinflammatory processes. Already known that GRb1 regulates activation of microglia [21], protecting neurons from inflammatory, oxidative injury and promoting neurogenesis [22].

The peroxisome proliferator-activated receptor gamma (PPAR $\gamma$ ), a ligand-activated transcription factor, triggers the expression of anti-inflammatory cytokines that attenuate neurodegeneration [23]. The shifts of M1 microglia to a M2 phenotype with PPAR $\gamma$  agonists such as pioglitazone, down-regulate pro-inflammatory mediators, and up-regulate pro-neurogenic factors in stress-exposed animals [24]. Conversely, PPAR- $\gamma$  antagonist, GW9662, inhibits the polarization of microglia to M2 phenotype [25]. Here, we tested the hypothesis that GRb1 alleviated depressive-like behaviors mainly through PPAR $\gamma$ -mediated microglial activation and enhancing adult hippocampus neurogenesis in CMS-exposed mice.

## 2. Material And Methods

### 2.1 Animals

Adult male C57BL/6J mice 8 weeks old (weighing 18–22 g) were obtained from the Laboratory Animal Center of the Sichuan Academy of Medical Sciences (Chengdu, China). The mice were housed individually under controlled conditions (temperature  $23 \pm 1.5^{\circ}\text{C}$ , humidity  $65 \pm 5\%$ , specific pathogen free, single cage) on a 12 h light/dark cycle (7 PM to 7 AM). Mice were acclimated in 27.5 x 15.5 x 18.5 cm plastic cages with sterile cotton wood sawdust. All experimental procedures were approved by the Ethics Committee of the University of Electronic Science and Technology of China and carried out in strict accordance with the National Institutes of Health Guide for the Care and Use of Laboratory Animals (8th edition, revised 2010).

## 2.2 Chronic mild stress

Male mice were subjected to CMS as described previously [26]. Briefly, mice were subjected to 1 or 2 kinds of random stressors per day. The stressors included cage tilting ( $45^{\circ}$ , 24 h), reversal of the light-dark cycle (24 h), food or water deprivation (12 h), empty or wet cage (12 h), lights-off (3 h), restraint (2 h), cage shaking (1 h), tail clamping (15 min) and ice water stimulation (5 min).

## 2.3 Drug administration

GRb1 (purity  $\geq 97\%$ , C54H92O23, Cat# P0088) was purchased from Pureone biotechnology company of shanghai. Lipopolysaccharide (LPS, E. Coli, 0127: B8) were purchased from Sigma-Aldrich Chemical Co. (St Louis, MO, USA) and extremely soluble in water. The experiment administered the following treatments once per day (at 16:00 h): saline, GRb1 (20 mg/kg/d, given as a 2 mg/ml solution in 0.9% saline (intra-gastric administration; Herbpurify, Chengdu, China), or imipramine hydrochloride (IMI, 20 mg/kg/d, intraperitoneally (i.p.); Sigma-Aldrich, Darmstadt, Germany). A subset of stressed animals was pretreated with PPAR $\gamma$  inhibitor GW9662 (1 mg/kg/day in 1% DMSO, i.p., 28 d; Med Chem Express, Monmouth Junction, NJ, USA), then treated 1 h later with GRb1 (control mice were administrated with 1% DMSO solution in 0.9% saline). The doses of GRb1 and IMI were chosen based on previous studies [27, 28].

## 2.4 Behavioral tests

### 2.4.1 Locomotor activity test

Locomotor activity of mice was evaluated using a mouse autonomic activity tester (Techman Software-zz6, Chengdu, China) in a quiet environment. In each trial, each mouse was placed in a chamber to acclimatization for 5 min. The numbers of movements and standing were recorded during 10 min test period. After each trial, the apparatus was cleaned using 75% ethyl alcohol.

### 2.4.2 Sucrose preference test and body weight measurement

The SPT was performed as described previously [4]. Before the test, mice were habituated to consume 1 % sucrose solution for 24 h. In the test, mice were deprived of food and water for 12 h, then provided with two containers of 1% sucrose and the same amount of water for 2 h. Sugar preference (%) was

calculated according to the following formula: Sugar preference (%) = sugar consumption (g) / [sugar consumption (g) + water consumption (g)] × 100%. The SPT and body weight was performed weekly.

#### 2.4.3 Tail suspension test

Each mouse was placed on the end of a rod suspended 30 cm above a tabletop and placed in an individual compartment. The mice were continuously monitored for 6 min using a digital camera. The duration of immobility during 6 min was analyzed.

#### 2.4.4 Forced swim test

The FST was carried out as described previously [29]. Briefly, the glass cylinders (height 21 cm, diameter 12 cm, volume 1000 ml) were filled with tap water (25 ± 2°C). Each mouse was monitored for 6 min using a computer-assisted video camera system (FST-100 Forced Swimming Analysis System, Techman Soft, Chengdu, China). Duration of immobility during the last 4 min was analyzed. FST was performed on week 8 after all other behavioral assays.

### 2.5 Cell cultures

#### 2.5.1 Primary microglial culture

Primary cultures of microglia from neonatal C57BL/6J mice brain (P0–P3 neonates) were prepared as described previously [30]. Briefly, the entire brain region was dissociated into a single-cell suspension using 0.25% pancreatin (Gibco, California, USA). Neonates were decapitated under sterile conditions, cut off the scalp and skull, removed the brain tissue. And brain tissue was placed in a dish containing cold D-Hank's solution, pH 7.2, without calcium or magnesium (Gibco, California, USA, Cat# C14175500BT). Tissues were enzymatically dissociated into a single-cell suspension using 0.25% pancreatin (Gibco, California, USA), then the mixed glial cells were cultured for 1 week in DMEM/F12 (Gibco, California, USA, Cat# C11330500BT) containing 10% fetal bovine serum (Gibco, California, USA) at 37 °C and 5% CO<sub>2</sub>. The isolated microglia were activated with phosphate-buffered saline (PBS, Servicebio, Cat# G4202) or LPS (100 µg/ml; Sigma-Aldrich, Darmstadt, Germany) in the presence or absence of (10, 20, 40) µg/ml GRb1. Some cultures were pretreated for 1 h with 10 µM GW9662, then treated with GRb1 for 24 h, and finally with LPS for 24 h. The collected microglia were transferred to a 6-well plate (2 × 10<sup>5</sup> cells/cm<sup>2</sup>) for subsequent analysis.

#### 2.5.2 NPCs culture

NPCs from C57BL/6J mice (P0–P3, n = 30) were isolated and cultured. Disinfection of infant mice was following the procedure described above in method 2.5.1, and subsequently the hippocampus tissue was dissociated from sagittal brains in ice-cold DMEM/F12, then digested using 0.25% pancreatin (Gibco, California, USA, Cat# 25200056). The digested cells were divided into two parts. One part of the cells (5 × 10<sup>4</sup> cells/cm<sup>2</sup>) was incubated in proliferation medium [high-glucose DMEM/F12, 20 ng/ml epidermal growth factor (Peprotech, Cat# 500-P174G), 20 ng/ml fibroblast growth factor (Peprotech, Cat# 450-33),

40 ng/ml N2 (Gibco, Cat# 17502-048), and 80 ng/ml B27 supplement (Gibco, Cat# 17504-04)]. The second ( $5 \times 10^4$  cells/cm<sup>2</sup>) was incubated in differentiation medium (high-glucose DMEM/F12, 40 ng/ml N2, 80 ng/ml B27 supplement, and 10% fetal bovine serum).

### 2.5.3 Conditioned microglial medium for NPCs culture

Primary microglia were first treated for 24 h with 20 µg/ml GRb1, then with PBS or 100 µg/ml LPS. Some cultures were pretreated for 1 h with 10 µM GW9662, then treated for 24 h with GRb1 and finally treated for 24 h with LPS. These media from microglia were harvested and used as conditioned medium to treat primary NPC cultures that had previously been cultured under the conditions described in the preceding section. Therefore, there were five NPC culture conditions: PBS-conditioned microglial medium (CM), GRb1-conditioned microglial medium (GRb1-CM), LPS-conditioned microglial medium (LPS-CM), LPS + GRb1-conditioned microglial medium (LPS+GRb1-CM), and LPS + GRb1 + GW9662-conditioned microglial medium (LPS+GRb1+GW-CM). NPC proliferation was analyzed after 24 h in the conditioned medium, while NPC differentiation was analyzed after 10 days in the conditioned medium. Areas with the highest cell density were imaged with 20 × on a Zeiss confocal microscope (LSM 800, Germany). Data were analyzed by an investigator blinded to animal treatment.

### 2.6 RNA extraction and reverse transcription-quantitative PCR (RT-qPCR)

Five mice from each group were randomly selected and sacrificed by decapitation. In accordance with the previous method [31], mice were anesthetized with pentobarbital diluted in 0.9% saline (50 mg/kg, i.p.; R&D Systems, Minneapolis, USA, Cat# 4579/50). Brain tissues were collected after intracardial perfusion with 120 ml cold 0.9% saline solution using a peristaltic pump at 11.5 ml/min. Both hippocampus and cortex were dissected out on ice. Each hemisphere was collected separately for subsequent analyses. Brain tissues were homogenized by the GeneUP Total RNA Mini Kit (Biotechrabbit, Cat# BR0702303). Total RNA (10 ng) was converted to cDNA by reverse transcription with TaKaRa reagent (Takara, Cat# 6210A). The cDNA was stored at -20°C. The following primer pairs were used: β-actin, 5'-CCG TGA AAA GAT GAC CCA GAT C-3' and 5'-CAC AGC CTG GAT GGC TAC GT-3'; IL-1β, 5'-CCA GCA GGT TAT CAT CAT CAT CC-3' and 5'-CTC GCA GCA GCA CAT CAA C-3'; tumor necrosis factor (TNF)-α, 5'-TAC TGA ACT TCG GGG TGA TTG GTC C-3' and 5'-CAG CCT TGT CCC TTG AAG AGA ACC-3'; TGF-β, 5'-GAC CGC AAC AAC GCC ATC TA-3' and 5'-GGC GTA TCA GTG GGG GTC AG-3'; Arginase (Arg)-1, 5'-AGA CAG CAG AGG AGG TGA AGA G-3' and 5'-CGA AGC AAG CCA AGG TTA AAG C-3'; PPARγ, 5'-CCC TGG CAA AGC ATT TGT AT-3' and 5'-CAC CTC TTT GCT CTG CTC CT-3'.

The reaction mixture for RT-qPCR consisted of 1 µl template cDNA, 0.3 µl primer and 5 µl SsoFast EvaGreen Supermix (Bio-Rad, California, USA) in a total volume of 10 µl. Data were reported as fold increase in mRNA levels relative to β-actin.

### 2.7 Western blotting

Protein extraction, tissue processing, and western blot analysis were based on described method [32]. Protein extraction kit were as follows: Total Protein Extraction kit (Sangon Biotech, Cat# 786-225). Nucleoprotein Extraction Kit (Sangon Biotech, no. c500009). Protein concentration of extracts BCA kit (Beyotime Institute of Biotechnology). Extracts were assayed for total protein using a BCA kit (Beyotime Institute of Biotechnology, Shanghai, China) according to the manufacturer's guidelines. Equal amounts of protein (2 µg/µl, 30 µg) were separated by 12% SDS/PAGE gel electrophoresis, then transferred to nitrocellulose membranes (Servicebio, Cat# G6014-15\*15CM). The membranes were covered with 5% skim milk (Servicebio, Cat# G5002) and incubated for 2 h. Then membrane with target protein were incubated overnight at 4°C with rabbit antibodies against PPAR $\gamma$  (1:2,000; IgG, Abcam, 52 KDa, Cat# ab59256, RRID: AB\_944767), rabbit antibodies against activated PPAR $\gamma$  (p-PPAR $\gamma$ ) (1:500; Thermo Fisher Scientific, 54 KDa, Cat# PA5-36763, RRID: AB\_2553712) or GAPDH (1:1,000; 37 KDa, Cat# JM-3777-100, RRID: AB\_843142). Lamin B (1:1,000; Wanleibio, 67 KDa, Cat# WL01775). The secondary antibodies goat anti-rabbit IgG (1:10,000; Thermo Fisher Scientific, Cat# 31460, RRID: AB\_228341) were incubated for 2 h. Primary and secondary antibodies were all diluted with 5% skimmed milk. Densitometry of protein bands was performed using Image J (version 1.45J; National Institutes of Health, Bethesda, MD, USA).

## 2.8 Immunofluorescence

To label proliferating cells in the brain, bromodeoxyuridine (BrdU, 50 mg/kg, i.p.; Sigma-Aldrich, Darmstadt, Germany, Cat# B5002) was received 12 h apart. To examine progenitor differentiation in SGZ, animals were injected with a double dose of BrdU and sacrificed 8 weeks after injection. (Fig. 1A). In cell experiment, BrdU was into the medium for 2 h before the test. Mice were anesthetized with pentobarbital sodium (50 mg/kg, i.p.; R&D Systems, Minneapolis, USA, Cat# 4579/50) and perfused transcardially with 120 ml of 0.9% saline using a peristaltic pump, followed by 4% paraformaldehyde with 120 ml. Dissected brains were fixed in 4% paraformaldehyde for 48 h, then dehydrated in 10%, 20%, 30% sucrose solution at 4°C for 24 h each. For immunostaining against DCX, Iba1 and PPAR $\gamma$ , p-PPAR $\gamma$ . Nuclei were stained with 4',6-diamidin-2-phenylindol (DAPI, Roche, USA, Cat# 10236276001).

The primary antibodies included rat anti-BrdU (1:500; Abcam, Cambridge, UK, Cat# ab6326, RRID: AB\_305426), goat anti-DCX (1:400; goat polyclonal antibody, Santa Cruz, Cat# sc-271390, RRID: AB\_10610966), The secondary antibodies (1:1,000; Jackson ImmunoResearch, Pennsylvania, USA) included AffiniPure Donkey Anti-Goat IgG, (Cat# 705-585-003, RRID: AB\_2340432), AffiniPure Donkey Anti-Rabbit IgG (Cat# 711-545-152, RRID: AB\_2313584), AffiniPure Donkey Anti-Rabbit IgG (Cat# 711-515-152, RRID: AB\_2340621). AffiniPure Donkey Anti-Rat IgG, (Cat# 705-585-003, RRID: AB\_2340432).

## 2.9 Morphometry of microglia

Images of hippocampal or cerebral cortical slices (1024 × 1024 pixels) were obtained using a Zeiss confocal microscope (LSM 800, Germany) and the Plan-Apochromat × 63/1.40 NA oil-immersion DIC M27 objective (Zeiss). The mouse brains were sliced into 35 µm thickness slices and were chosen to sample every sixth section based on unbiased counting methods [33]. The brain slice per well was selected sequentially in a 1:6 ratio for a mouse, with each brain slice including the intact hippocampus

and cortex. Microglial density was determined by dividing the number of cells by the total area ( $\text{mm}^3$ ). Microglial morphology was quantified using skeleton analysis method [34]. Images of each slice ( $1024 \times 1024$  pixels) were obtained using a Zeiss confocal microscope (LSM 800, Germany). The resulting images were skeletonized using Image J software.

## 2.10 The quantification of neurogenesis regions

The number of BrdU<sup>+</sup> cells and DCX<sup>+</sup> cells in the dentate gyrus (DG) was quantified on the basis of the Cavalieri principle with every sixth section of the hippocampus [35] at 20 $\times$  objective. In each section, the number of cells was divided by per  $\text{mm}^2$  DG in that section [36]. To estimate the total volume of the subregions in the hippocampus, the hippocampal slices were chosen according to the Cavalieri's method, and every sixth section was stained with DAPI antibody at a 40 $\times$  magnification [37]. We performed the volume measurements of DG and granular cell layer (GCL) in the hippocampal area of 12 sections per animal using the Image J software (version 1.45 J; National Institutes of Health, Bethesda, MD, USA). The volume of subregions was obtained by multiplying the sum of the section areas per animal by 35  $\mu\text{m}$ .

## 2.11 Statistical analyses

Experiments were performed at least three samples and three times independently. Samples and animals were randomly allocated to experimental groups. The data and statistical analysis comply with the documentation requirements [38]. All experiments were performed in a blinded manner since the measurements could be influenced by personal bias. All data were expressed as mean  $\pm$  SEM. Statistical analyses were performed using SPSS for Windows® (version 17; Chicago, USA). In sucrose preference and body weight from week 4 to 8 group, Ctrl and CMS+GRb1 groups at 4<sup>th</sup> week and 8<sup>th</sup> week were assessed by three-way repeated measures (ANOVA). CMS and CMS+IMI groups at 4<sup>th</sup> week and 8<sup>th</sup> week were analyzed by two-way repeated measures (ANOVA). CMS+GRb1 and CMS+GRb1+GW groups at 4<sup>th</sup> week and 8<sup>th</sup> week were performed using two-way repeated measures (ANOVA). A post hoc test was applied only when the F value reached significance and there was no significance in homogeneity of variance. Differences among three or more groups were assessed using one-way or two-way analysis of variance as appropriate, followed by Bonferroni test as post-hoc. The level of confidence was set at 95% ( $P < 0.05$ ), and were done only on datasets of  $n \geq 5$ . To exclude the misinterpretation of results by the difference in the area of region of interest, the number of proliferating cells in mice, or the total number of cells in each image, normalization of the data was carried out for some analyses in the study.

# 3. Results

## 3.1 GRb1 alleviates depressive-like behaviors in male mice exposed to CMS

Previous pharmacokinetics and pharmacodynamics experiments have showed that GRb1 plays a role in a variety of CNS disorders [39]. In the present study, we adopted CMS paradigm, one of the most extensively validated and realistic models of depression [40]. Following exposure to CMS for 8 weeks,

C57BL/6J mice were treated GRb1 at 4th week. The procedure shown in Fig. 1A. We first assessed the locomotor ability of mice. In spontaneous activity, there was no statistically difference in activity level or standing time among groups, indicating that the drug treatment had no effect on motor ability (Fig. 1B). The weight gain was inhibited by CMS exposure at 4th and 8th week. The CMS-induced reduction of weight gain was attenuated by GRb1 and IMI at week 8 (Fig. 1C). The FST and TST were performed to test the desperate behaviors of mice. Chronic stress caused mice to remain immobile for longer periods, whereas the GRb1-treated and IMI-treated mice displayed decreased immobility time in both TST and FST, similar to that of Ctrl mice (Fig. 1D–1E). The sucrose preference test is commonly used to evaluate anhedonia in animals. The CMS groups showed reduction in sucrose consumption between week 4 and 8. Both GRb1 and IMI-treated mice displayed increase the sucrose consumption compared with CMS group (Fig. 1F). The obtained results indicated that GRb1 was able to recover depressive-like behaviors in CMS-exposed mice.

### 3.2 GRb1 promotes microglial M2-polarization in the hippocampus and cortex.

Depression as a microglial disease [41], we next explored whether GRb1 could activated microglial polarization. The microglial morphological property using immunofluorescence labeling. Iba1 as microglia-specific marker (Fig. 2A–2B). Results showed that CMS significantly increased the density of Iba1<sup>+</sup> cells in hippocampus and cortex, while treatment with GRb1 significantly prevented the increase in cell density of CMS mice (Fig. 2C). When microglia are activated, they switch from ramified to amoeboid morphology. Correspondingly, hippocampal microglia in CMS animals had larger and bigger soma, fewer and shorter processes than Ctrl group. GRb1 treatment of CMS-exposed mice reduced the cell area (Fig. 2D), increased the number and the total length of microglia processes (Fig. 2E–2F). Whereas cortical microglia had shorter processes in CMS animals than Ctrl animals, and GRb1 treatment increased the cell area, the number of processes in cortex (Fig. 2D–2F). Polarized microglia also are distinguished by their expression profiles of signature surface markers. Stress up-regulated the pro-inflammatory (M1 microglia) markers TNF- $\alpha$  and IL-1 $\beta$  in hippocampus and cortex, while down-regulated M2 microglial markers TGF- $\beta$  and Arg-1. GRb1 inhibited these changes (Fig. 2G–2J). These results indicated that GRb1 activated M2-like polarized microglia.

### 3.3 GRb1 rescues neurogenesis impairment in hippocampus

Adult hippocampal neurogenesis as a potential candidate mechanism for the etiology of depression [42]. We then examined whether GRb1 was involved in neurogenesis. Proliferative cells were labeled with BrdU, and newborn neurons were labeled with DCX (Fig. 3A). Chronic stress decreased the mRNA level of DCX, which was increase by GRb1 (Fig. 3B). The immunofluorescence results indicated that CMS also decreased the number of BrdU<sup>+</sup> cells in DG, which was restored by GRb1 (Fig. 3C). Here, at 8 weeks post-BrdU injection, chronic stress significantly reduced the number of DCX<sup>+</sup> cells and the “differentiation ratio” in DG, which was significantly increased in GRb1 group (Fig. 3D–3E). In addition, in order to determine whether the effects of GRb1 administration on hippocampus volume, we performed analysis of volume in DG and GCL. The results revealed that GRb1 reversed the reduction the volume of DG and GCL in CMS

group (Fig. 3F–3H). The CMS group also showed a tendency to cause thinning of GCL, which GRb1 reversed (Fig. 3I). In contrast, no alterations were found in administering GRb1 in the absence of CMS in neurogenesis of hippocampus.

### 3.4 GRb1 activates PPAR $\gamma$ expression in CMS-induced depression model

Activation of the nuclear receptor PPAR $\gamma$  prevents microglial activation, and PPAR $\gamma$  antagonist aggravates microglia-regulated neuroinflammation [43]. We therefore hypothesized that GRb1 treatment could mediate PPAR $\gamma$  activation. We performed GW9662 as PPAR $\gamma$  effective inhibitors. CMS down-regulated PPAR $\gamma$  mRNA in hippocampus, which GRb1 up-regulated PPAR $\gamma$  mRNA in hippocampus (Fig. 4A). GRb1 had slight downward trend of PPAR $\gamma$  mRNA levels in cortex of CMS group. (Fig. 4B). We further verified the PPAR $\gamma$  expression or p-PPAR $\gamma$  expression at the protein level (Fig. 4C–4D). Western blotting showed that in hippocampus, the CMS significantly reduced levels of PPAR $\gamma$  and p-PPAR $\gamma$  in total protein and nuclear protein. The levels of p-PPAR $\gamma$  in nuclei were also decreased in CMS-treated mice, GRb1 inhibited CMS-induced reduction of PPAR $\gamma$  and p-PPAR $\gamma$ ; In cortex, Reduction of PPAR $\gamma$  protein in CMS group compared with control group (Fig. 4E–4J). Whether this function was reversed by PPAR $\gamma$  inhibitor GW9662. Indeed, the above PPAR $\gamma$  expression were significantly reversed by PPAR $\gamma$  inhibitor GW9662 in hippocampus. Administering GRb1 alone had no effect on PPAR $\gamma$  expression at mRNA or protein level.

### 3.5 PPAR $\gamma$ activation mediates effects of GRb1-induced microglia

We first assessed whether the effect of antidepressant of GRb1-induced microglia was mediated by PPAR $\gamma$  activation. Mice were treated with GW9662 treatment, blocking the PPAR $\gamma$  pathway, before GRb1 administration on CMS condition. We found that the GW9662 treatment notably blocked the effects of antidepressant of GRb1 on the decrease of body weight, the increase of the immobility time in FST, the reduction of sucrose consumption, without affecting autonomic activity. The GW9662 administration also caused similar trends in both weight change and sucrose preference as those in the CMS group at week 4 and week 8 (Fig. 1B–1F). We then examined the interception of GW9662 administration on GRb1-induced microglial activation. Indeed, the GW9662 administration weakened the effects of GRb1 on Iba1<sup>+</sup> cell number (Fig. 2C) and microglial morphology in the hippocampus (Fig. 2E–2F). In cortex, the GW9662 treatment blocked only the effects of GRb1 on Iba1<sup>+</sup> cell number in CMS-exposed mice (Fig. 2C). We also found that GW9662 treatment dramatically reversed GRb1-induced down-regulation of M1 markers and up-regulation of M2 markers in hippocampus (Fig. 2G–2H). Similar results were observed in cortex, except that GW9662 treatment did not affect GRb1-induced down-regulation of TNF- $\alpha$  (Fig. 2I–2J). We then examined PPAR $\gamma$  inhibitor in neurogenesis in CMS mice. The GW9662 treatment dramatically decreases in the level of DCX mRNA in compared with GRb1 + CMS group (Fig. 3B), the GW9662 treatment reduced the neurogenesis (Fig. 3C–3I). The results provided evidence that GRb1 activated M2 microglia and promoted neurogenesis dependent on PPAR $\gamma$  pathway.

### 3.6 PPAR $\gamma$ activation mediates effects of GRb1-induced microglia *in vitro*

Primary cultures of microglia were stimulated by LPS in the presence or absence of GRb1, with or without GW9662. To maximize drug effectiveness *in vitro*, 20 µg/ml of GRb1 was chosen to be used in the following experiments to determine the regulatory effect of GRb1 on the activation of microglia and neurogenesis (Fig. S1). Cultures were analyzed by immunofluorescence to assess microglial morphology and PPAR $\gamma$  expression (Fig. 5A). GRb1 increased the PPAR $\gamma$  expression in LPS group, which GW9662 reversed the increase (Fig. 5B). GRb1 up-regulated the PPAR $\gamma$  expression in LPS treatment. LPS-treated primary microglia exhibited round soma and developed longer processes. GRb1 pretreatment inhibited LPS-induced microglial activation, decreasing the Iba1<sup>+</sup> cell area and length of processes, and this change was significantly suppressed by GW9662 (Fig. 5C–5D). To correlate these effects with neurogenesis, we treated NPCs with some stimuli (Fig. 5E). Proliferative cells were labeled with BrdU, and newborn neurons were labeled with DCX (Fig. 5F). GRb1 increased the ratio of BrdU<sup>+</sup> to DAPI<sup>+</sup> NPCs, which GW9662 did significantly reduced (Fig. 5G). The GRb1 treatment increased the ratio of DCX<sup>+</sup> to DAPI<sup>+</sup> cells (Fig. 5I). The GRb1 alone showed no significant effects on the ratio of BrdU<sup>+</sup> to DAPI<sup>+</sup> and DCX<sup>+</sup> to DAPI<sup>+</sup> cells NPCs.

Next we asked whether the effects of GRb1 on neurogenesis were mediated by microglia. Primary cultures of NPCs were incubated in conditioned medium from microglia. Conditioned medium from microglia stimulated with LPS reduced the ratios of BrdU<sup>+</sup> to DAPI<sup>+</sup> cells and DCX<sup>+</sup> to DAPI<sup>+</sup> cells, and GRb1-CM increase the ratios, whereas the effect was inhibited by pretreated with GW9662 (Fig. 5H–5J). These results indicated that GRb1-mediated M2 microglia stimulated neurogenesis dependent on PPAR $\gamma$  activation *in vitro*.

## 4. Discussion

GRb1 shows antidepressant effects in rodent models of stress-induced depression. Our present study discloses that GRb1 alleviates depressive-like behaviors by regulating inflammatory cytokines in hippocampus and cortex of CMS-exposed mice. Moreover, the results revealed that GRb1 treatment shifted microglia to M2 phenotype and stimulating neurogenesis in hippocampus. The data from *in vivo* and *in vitro* both indicated that GRb1-induced shift in microglial phenotype was dependent on PPAR $\gamma$  pathway. These findings provided first evidence that GRb1 induced a pro-neurogenic microglial phenotype via PPAR $\gamma$  activation in hippocampus of CMS-exposed mice.

In order to prove the antidepressant effect of GRb1, we conducted *in vivo* studies using male mice exposed to CMS. CMS model is widely used to investigate the underlying mechanism animal lines of depression [45]. The most obvious feature of CMS is anhedonia, that is a decreased preference for sugar water. Another feature is an extinction-like inhibitory learning behavior, that is longer immobility time in both FST and TST. GRb1 and IMI administration can reverse the effects of anhedonia and desperate behaviors. Since no alterations in the spontaneous locomotor activity were found, the results indicated that antidepressant-like activity of GRb1 exerted antidepressant effects in CMS depression model, not affect the locomotor ability. Sex as an important biological variable from basic and preclinical research

[46]. It has been previously reported that GRb1 ameliorated the depressive-like behaviors in ovariectomized or menopausal female mice. [47, 48], indicating that the potential function of the sex hormone effects of GRb1 on depression would be achieved further investigation in the future studies.

Accumulating evidence supports an association between depression and inflammatory processes [49]. As a primary source of inflammatory molecules in the CNS, microglia show a spectrum of polarization phenotypes responsible for balance of pro- and anti-inflammatory mediators [50]. Chronic stress activates hippocampal microglia, microglia tended to show an M1 phenotype, causing soma enlargement, shortening and thickening of processes. Activated microglia were also characterized by increasing expression of TNF- $\alpha$  and IL-1 $\beta$  and decreasing expression of the M2 markers TGF- $\beta$  and Arg-1. GRb1 treatment *in vivo* significantly reduced the area of hippocampal Iba1<sup>+</sup> cells and increased the number and total length of microglial processes. Meanwhile, GRb1 decreased the expression of M1 markers and increased the expression of M2 markers. These results suggested that GRb1 shifted the phenotype of microglia from M1 to M2 polarization in CMS-exposed mice. Since there were interactions between various cell types in brain, the direct effect of GRb1 on microglial morphology was further tested on primary microglia. The result showed that GRb1 directly prevented LPS-induced effects on microglial activation, consistent with the effects of GRb1 *in vivo*.

Impairment of hippocampal neurogenesis has been associated with inflammation-mediated depressive-like symptoms. Neurogenesis occurs predominantly in the SGZ of hippocampus and subependymal ventricular zone [51]. Our results showed that GRb1 significantly increased the number of BrdU<sup>+</sup> and DCX<sup>+</sup> cells, as well as the differentiation ratio in SGZ. Several studies performed using magnetic resonance imaging showed similar total brain volume but smaller hippocampal volume in patients with depression than in controls [52]. Post mortem analysis of depressed patients has shown a reduction in hippocampal volume [53]. Consistent with these results, our data showed that stress decreased the volume of DG and GCL, while long-term treatment with GRb1 significant amelioration of neuronal structural changes. Previous studies have reported that persistent production of pro-inflammatory cytokines is detrimental to neurogenesis [54]. Based on our observation that a shift of microglial phenotypes from M1 to M2, pretreatment of GRb1 reversed chronic stress-induced the impairment in hippocampal neurogenesis. we speculated that GRb1-induced M2 microglia played a phenotype-associated neurogenic role.

We next analyzed the role of microglia on neurogenesis *in vitro* using conditioned media. LPS-CM reduced the proliferation and differentiation of NPCs, while GRb1 + LPS-CM promoted the proliferation and differentiation of NPCs. GRb1 appeared to play a subtle neurogenic effect on immune-stimulated NPCs, promoting their proliferation but not differentiation. Consistent with our results, M1 microglia promoted astrocytogenesis, while M2 microglia supported neurogenesis *in vitro* [55]. Liu et al. [56] reported that GRb1 increased survival of newborn neurons and didn't influence NPCs proliferation in non-stressed rat. Our different findings of GRb1 on NPCs proliferation may come from different animal models, which remain to be confirmed. These *in vivo* and *in vitro* results implied that GRb1 enhanced

neurogenesis via switching microglial phenotype from M1 to M2, contributing to neurogenesis in CMS-exposed mice.

Mounting evidence has been presented demonstrating that GRb1 has anti-inflammatory and neuroprotective effects. Ying et al. [57] suggested that GRb1 exerted antidepressant effects via multiple cellular and molecular pathways. PPAR $\gamma$  is a ligand-activated transcription factor that regulates the expression of inflammatory cytokines and microglial phenotypes [58]. We thus reasoned that GRb1-activated microglia may be related to PPAR $\gamma$  pathway. As expected, our study showed that GRb1 up-regulated PPAR $\gamma$  expression and enhanced PPAR $\gamma$  activation in the hippocampus of CMS-treated mice. GRb1 also altered phenotype of microglia transition from M1 to M2, increasing neurogenesis in hippocampus. As an approach to corroborate these findings, we used PPAR $\gamma$  antagonist GW9662 to intercept microglial activation. The results indicated that the GW9662 treatment strongly prevent GRb1-induced microglial activation *in vivo* and *in vitro*. Furthermore, the culture NPCs using conditioned medium from microglia showed that GRb1 enhanced the proliferation and differentiation of NPCs via PPAR $\gamma$ -mediated microglia *in vitro*. The results of this experiment showed that inhibition of PPAR $\gamma$  activation produced a significant dysregulation in neurogenesis mediated by microglia. Our experiments highlighted the functional role of microglia as components of a neurogenic niche in the brain, and implicated the role of PPAR $\gamma$  activation in NPCs proliferation and differentiation.

## 5. Conclusion

Our study provides strong *in vivo* and *in vitro* evidence that GRb1 exerts antidepressant effects by activating PPAR $\gamma$  to shift microglia towards an anti-inflammatory, pro-neurogenic phenotype. Our findings may shed light on the potential contribution of GRb1-treated microglia to promote neurogenesis as a therapeutic strategy against MDD.

## Abbreviations

Arg, Arginase; BrdU, Bromodeoxyuridine; CMS, chronic mild stress; CNS, central nervous system; DAPI, 4',6-diamidin-2-phenylindol; DG, dentate gyrus; FST, force swimming test; GCL, granular cell layer; GRb1, Ginsenoside Rb1; IL, interleukin; IMI, imipramine; LAT, locomotor activity test; LPS, lipopolysaccharide; MDD, Major depressive disorder; NPC, neural precursor cell; OFT, open field test; PPAR $\gamma$ , peroxisome proliferator-activated receptor gamma; RT-qPCR, reverse transcription-quantitative PCR; SGZ, subgranular zone; SPT, sucrose preference test; TGF, transforming growth factor; TNF, tumor necrosis factor; TST, tail suspension test.

## Declarations

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### **Authors' contributions**

Author's contributions: L.Z. designed the study, performed experiments and wrote the paper. M.T., Q. Z. and X.X. designed and performed statistical analysis of animal behavior experiments. N.H. and H.H. performed experiments and analyzed quantitative PCR and western blot data. G.L. and S.H. performed experiments and analyzed immunofluorescence data. C.P., Y.X. and Z.Y. designed the study and revised the paper.

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### **Availability of data and materials**

All data generated and materials supporting the conclusion of the study are included within the article and its supplementary information files.

### **Competing interests**

All authors declare that there are no conflicts of interest.

### **Consent for publication**

Not applicable

### **Ethics approval and consent to participate**

All animals care and experimental procedures were approved by the Ethics Committee of the University of Electronic Science and Technology of China and carried out in strict accordance with the National Institutes of Health Guide for the Care and Use of Laboratory Animals.

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## Tables

**Table 1** The F value and P value in multiple comparisons of figure 1

<i>figure</i>	group	F or T	P	N
<i>Figure 1B-1</i>	CMS vs. Ctrl		0.0970	
	GRb1 vs. Ctrl	4.612	0.1135	12
	CMS+GRb1 vs. CMS		0.9942	
	CMS+IMI vs. CMS		0.2171	
	CMS+IMI vs. CMS+GRb1	1.903	0.0702	
	CMS+GRb1+GW vs. CMS+GRb1		0.0968	12
		1.734		
<i>Figure 1B-2</i>	CMS vs. Ctrl		0.2054	
	GRb1 vs. Ctrl	1.411	0.6570	12
	CMS+ GRb1 vs. CMS		0.6361	
	CMS+IMI vs. CMS		0.9489	
	CMS+ IMI vs. CMS+GRb1	0.076	0.9399	
	CMS+GRb1+GW vs. CMS+GRb1		0.7401	12
		0.336		
<i>Figure 1C-1</i>	Treatment vs Basal (Ctrl)		0.0020	
	Treatment vs Basal (GRb1)		0.0010	
	Treatment vs Basal (CMS)		0.0750	
	Treatment vs Basal (CMS+GRb1)		< 0.0001	12
		14.211		
	Treatment vs Basal (CMS+IMI)		< 0.0001	
	Treatment vs Basal (CMS+GRb1+GW)		0.0960	
	Ctrl vs CMS (Basal)		< 0.0001	
	CMS+GRb1 vs CMS (treatment)		< 0.0001	
	CMS+IMI vs CMS (treatment)	4.540	0.0631	12
CMS+GRb1+GW vs. CMS+GRb1 (treatment)		< 0.0001		
CMS vs. Ctrl		0.0343		

<i>Figure 1C-2</i>	GRb1 vs. Ctrl	4.636	0.9032	12
	CMS+ GRb1 vs. CMS		0.0035	
	CMS+IMI vs. CMS		0.1519	
	CMS+ IMI vs. CMS+GRb1	0.472	0.6416	
	CMS+GRb1+GW vs. CMS+GRb1		0.0003	12
		4.255		
<i>Figure 1D</i>	CMS vs. Ctrl		0.0002	
	GRb1 vs. Ctrl	43.820	>0.9999	12
	CMS+ GRb1 vs. CMS		< 0.0001	
	CMS+IMI vs. CMS		0.0006	
	CMS+ IMI vs. CMS+GRb1	1.097	0.2846	
	CMS+GRb1+GW vs. CMS+GRb1		0.0001	12
		4.721		
<i>Figure 1E</i>	CMS vs. Ctrl		< 0.0001	
	GRb1 vs. Ctrl	6.458	0.6054	12
	CMS+GRb1 vs. CMS		< 0.0001	
	CMS+IMI vs. CMS		0.0001	
	CMS+ IMI vs. CMS+GRb1	0.513	0.6127	
	CMS+GRb1+GW vs. CMS+GRb1		0.0003	12
		4.228		

	Treatment vs Basal (Ctrl)		0.0720	
	Treatment vs Basal (GRb1)		0.1241	
<i>Figure 1F-1</i>	Treatment vs Basal (CMS)		0.4190	
	Treatment vs Basal (CMS+GRb1)		< 0.0001	12
		7.804		
	Treatment vs Basal (CMS+IMI)		< 0.0001	
	Treatment vs Basal (CMS+GRb1+GW)		0.4515	
	Ctrl vs CMS (Basal)		< 0.0001	
	CMS+GRb1 vs CMS (treatment)		< 0.0001	
	CMS+IMI vs CMS (treatment)	18.730	< 0.0001	12
	CMS+GRb1+GW vs. CMS+GRb1 (treatment)		< 0.0001	
	CMS vs. Ctrl		< 0.0001	
	GRb1 vs. Ctrl	33.920	0.1233	12
<i>Figure 1F-2</i>	CMS+ GRb1 vs. CMS		< 0.0001	
	CMS+IMI vs. CMS		< 0.0001	
	CMS+ IMI vs. CMS+GRb1	0.809	0.4287	
	CMS+GRb1+GW vs. CMS+GRb1		0.0057	12
		3.135		

**Table 2** The F value and P value in multiple comparisons of figure 2

<i>figure</i>	group	F or T	P	N
<i>Figure 2C-h</i>	CMS vs. Ctrl		< 0.0001	
	GRb1 vs. Ctrl	33.920	0.1233	6
	CMS+GRb1 vs. CMS		< 0.0001	
	CMS+GRb1+GW vs. CMS+GRb1	8.648	< 0.0001	6
<i>Figure 2C-c</i>	CMS vs. Ctrl		0.0002	
	GRb1 vs. Ctrl	7.923	0.2627	6
	CMS+ GRb1 vs. CMS		0.0004	
	CMS+GRb1+GW vs. CMS+GRb1	2.444	0.0193	6
<i>Figure 2D-h</i>	CMS vs. Ctrl		0.0207	
	GRb1 vs. Ctrl	2.347	0.6339	5
	CMS+ GRb1 vs. CMS		0.0155	
	CMS+GRb1+GW vs. CMS+GRb1	3.191	0.0057	6
<i>Figure 2D-c</i>	CMS vs. Ctrl		0.2343	
	GRb1 vs. Ctrl	8.559	0.2388	6
	CMS+GRb1 vs. CMS		0.0441	
	CMS+GRb1+GW vs. CMS+GRb1	2.107	0.0682	
<i>Figure 2E-h</i>	CMS vs. Ctrl		0.0006	
	GRb1 vs. Ctrl	5.422	0.1039	6
	CMS+GRb1 vs. CMS		0.0001	
	CMS+GRb1+GW vs. CMS+GRb1	6.878	< 0.0001	
<i>Figure 2E-c</i>	CMS vs. Ctrl		0.0695	
	GRb1 vs. Ctrl	0.380	0.0732	6
	CMS+GRb1 vs. CMS		0.0056	

CMS+GRb1+GW vs. CMS+GRb1

0.6611

0.445

<i>Figure 2F-h</i>	CMS vs. Ctrl		0.0142	
	GRb1 vs. Ctrl	1.530	0.3609	6
	CMS+GRb1 vs. CMS		0.0017	
	CMS+GRb1+GW vs. CMS+GRb1	2.139	0.0482	
<i>Figure 2F-c</i>	CMS vs. Ctrl		0.0151	
	GRb1 vs. Ctrl	1.862	0.4787	6
	CMS+GRb1 vs. CMS		0.9884	
	CMS+GRb1+GW vs. CMS+GRb1	0.933	0.3783	
<i>Figure 2G-1</i>	CMS vs. Ctrl		<0.0001	
	GRb1 vs. Ctrl	1.443	0.7468	5
	CMS+GRb1 vs. CMS		0.0057	
	CMS+GRb1+GW vs. CMS+GRb1	4.834	0.0013	5
<i>Figure 2G-2</i>	CMS vs. Ctrl		0.0329	
	GRb1 vs. Ctrl	3.340	0.8778	5
	CMS+GRb1 vs. CMS		0.0015	
	CMS+GRb1+GW vs. CMS+GRb1	3.610	0.0069	5
<i>Figure 2H-1</i>	CMS vs. Ctrl		0.0109	
	GRb1 vs. Ctrl	9.361	0.8942	5
	CMS+GRb1 vs. CMS		0.0004	
	CMS+GRb1+GW vs. CMS+GRb1	4.258	0.0028	5
<i>Figure 2H-2</i>	CMS vs. Ctrl		0.0159	
	GRb1 vs. Ctrl	6.216	0.6888	5
	CMS+GRb1 vs. CMS		0.0011	
	CMS+GRb1+GW vs. CMS+GRb1		0.0037	

		4.040		5
<i>Figure 2I-1</i>	CMS vs. Ctrl		0.0003	
	GRb1 vs. Ctrl	6.799	0.9950	5
	CMS+GRb1 vs. CMS		0.0048	
	CMS+GRb1+GW vs. CMS+GRb1		0.1812	
		1.464		5
<i>Figure 2I-2</i>	CMS vs. Ctrl		0.0005	
	GRb1 vs. Ctrl	30.290	0.2315	5
	CMS+GRb1 vs. CMS		< 0.0001	
	CMS+GRb1+GW vs. CMS+GRb1		0.0294	
		2.646		5
<i>Figure 2J-1</i>	CMS vs. Ctrl		0.0073	
	GRb1 vs. Ctrl	11.820	0.9568	5
	CMS+GRb1 vs. CMS		0.0002	
	CMS +GRb1+GW vs. CMS+GRb1		< 0.0001	
		7.331		5
<i>Figure 2J-2</i>	CMS vs. Ctrl		0.0282	
	GRb1 vs. Ctrl	16.090	0.3674	5
	CMS+GRb1 vs. CMS		0.0010	
	CMS+GRb1+GW vs. CMS+GRb1		0.0123	
		3.215		5

h, hippocampus; c, cortex

**Table 3** The F value and P value in multiple comparisons of figure 3

<i>figure</i>	group	F or T	P	N
<i>Figure 3B</i>	CMS vs. Ctrl		0.0008	
	GRb1 vs. Ctrl	27.880	0.2877	5
	CMS+GRb1 vs. CMS		< 0.0001	
	CMS+GRb1+GW vs. CMS+GRb1		< 0.0001	
		7.291		
<i>Figure 3C</i>	CMS vs. Ctrl		< 0.0001	
	GRb1 vs. Ctrl	47.020	0.7149	6
	CMS+GRb1 vs. CMS		< 0.0001	
	CMS+GRb1+GW vs. CMS+GRb1		< 0.0001	
		8.922		
<i>Figure 3D</i>	CMS vs. Ctrl		< 0.0001	
	GRb1 vs. Ctrl	30.200	0.9995	6
	CMS+GRb1 vs. CMS		< 0.0001	
	CMS+GRb1+GW vs. CMS+GRb1		0.0003	
		6.145		
<i>Figure 3E</i>	CMS vs. Ctrl		< 0.0001	
	GRb1 vs. Ctrl	4.820	0.9971	6
	CMS+GRb1 vs. CMS		0.0094	
	CMS+GRb1+GW vs. CMS+GRb1		0.0064	
		3.659		
<i>Figure 3G</i>	CMS vs. Ctrl		0.0399	
	GRb1 vs. Ctrl	10.990	0.4946	6
	CMS+GRb1 vs. CMS		0.0021	
	CMS+GRb1+GW vs. CMS+GRb1		0.0212	
		2.858		
<i>Figure 3H</i>	CMS vs. Ctrl		0.5954	
	GRb1 vs. Ctrl	1.460	0.8912	6
	CMS+GRb1 vs. CMS		0.0934	

	CMS+GRb1+GW vs. CMS+GRb1	0.1439	
		1.528	
	CMS vs. Ctrl	0.4818	
	GRb1 vs. Ctrl	3.167	6
<i>Figure 3I</i>	CMS+GRb1 vs. CMS	0.0378	
	CMS+GRb1+GW vs. CMS+GRb1	0.0062	
		3.678	

**Table 4** The F value and P value in multiple comparisons of figure 4

<i>figure</i>	group	F or T	P	N
<i>Figure 4A</i>	CMS vs. Ctrl		0.0174	
	GRb1 vs. Ctrl	2.407	0.6867	5
	CMS+GRb1 vs. CMS		< 0.0001	
	CMS+GRb1+GW vs. CMS+GRb1	5.368	0.0007	5
<i>Figure 4B</i>	CMS vs. Ctrl		0.9993	
	GRb1 vs. Ctrl	3.246	0.3731	5
	CMS+GRb1 vs. CMS		0.0422	
	CMS+GRb1+GW vs. CMS+GRb1	1.179	0.2723	5
<i>Figure 4E-1</i>	CMS vs. Ctrl		0.0119	
	GRb1 vs. Ctrl	9.619	0.9997	5
	CMS+GRb1 vs. CMS		0.0010	
	CMS+GRb1+GW vs. CMS+GRb1	4.306	0.0026	5
<i>Figure 4E-2</i>	CMS vs. Ctrl		0.0003	
	GRb1 vs. Ctrl	2.197	0.0913	5
	CMS+GRb1 vs. CMS		0.0147	
	CMS+GRb1+GW vs. CMS+GRb1	4.670	0.0009	5
<i>Figure 4F-1</i>	CMS vs. Ctrl		0.0134	
	GRb1 vs. Ctrl	2.242	0.9577	5
	CMS+GRb1 vs. CMS		0.0590	
	CMS+GRb1+GW vs. CMS+GRb1	1.005	0.3443	5
<i>Figure 4F-2</i>	CMS vs. Ctrl		0.1676	
	GRb1 vs. Ctrl	7.645	0.9676	5
	CMS+GRb1 vs. CMS		0.0538	

	CMS+GRb1+GW vs. CMS+GRb1	0.0067	
		3.692	5
<i>Figure 4I</i>	CMS vs. Ctrl	<0.0001	
	GRb1 vs. Ctrl	33.790	5
	CMS+GRb1 vs. CMS	<0.0001	
	CMS+GRb1+GW vs. CMS+GRb1	0.0026	
		4.299	5
<i>Figure 4J</i>	CMS vs. Ctrl	0.1916	
	GRb1 vs. Ctrl	5.065	5
	CMS+GRb1 vs. CMS	0.3793	
	CMS+GRb1+GW vs. CMS+GRb1	0.4853	
		0.732	5

**Table 5** The F value and P value in multiple comparisons of figure 5

<i>figure</i>	group	F or T	P	N
<i>Figure 5B</i>	LPS vs. Ctrl		0.0175	
	GRb1 vs. Ctrl	0.101	0.0331	5
	LPS+GRb1 vs. LPS		0.0131	
	LPS+GRb1+GW vs. LPS+GRb1	2.352	0.0318	
<i>Figure 5C</i>	LPS vs. Ctrl		< 0.0001	
	GRb1 vs. Ctrl	16.850	0.9873	5
	LPS+GRb1 vs. LPS		< 0.0001	
	LPS+GRb1+GW vs. LPS+GRb1	2.953	0.0112	
<i>Figure 5D</i>	LPS vs. Ctrl		< 0.0001	
	GRb1 vs. Ctrl	25.380	0.1650	5
	LPS+GRb1 vs. LPS		< 0.0001	
	LPS+GRb1+GW vs. LPS+GRb1	1.615	0.1150	
<i>Figure 5G</i>	LPS vs. Ctrl		< 0.0001	
	GRb1 vs. Ctrl	20.280	0.2200	5
	LPS+GRb1 vs. LPS		< 0.0001	
	LPS+GRb1+GW vs. LPS+GRb1	1.514	0.1523	
<i>Figure 5H</i>	LPS vs. Ctrl		< 0.0001	
	GRb1 vs. Ctrl	26.160	0.7954	5
	LPS+GRb1 vs. LPS		< 0.0001	
	LPS+GRb1+GW vs. LPS+GRb1	5.004	0.0002	
<i>Figure 5I</i>	LPS vs. Ctrl		0.7890	
	GRb1 vs. Ctrl	1.794	0.0089	5
	LPS+GRb1 vs. LPS		0.2532	

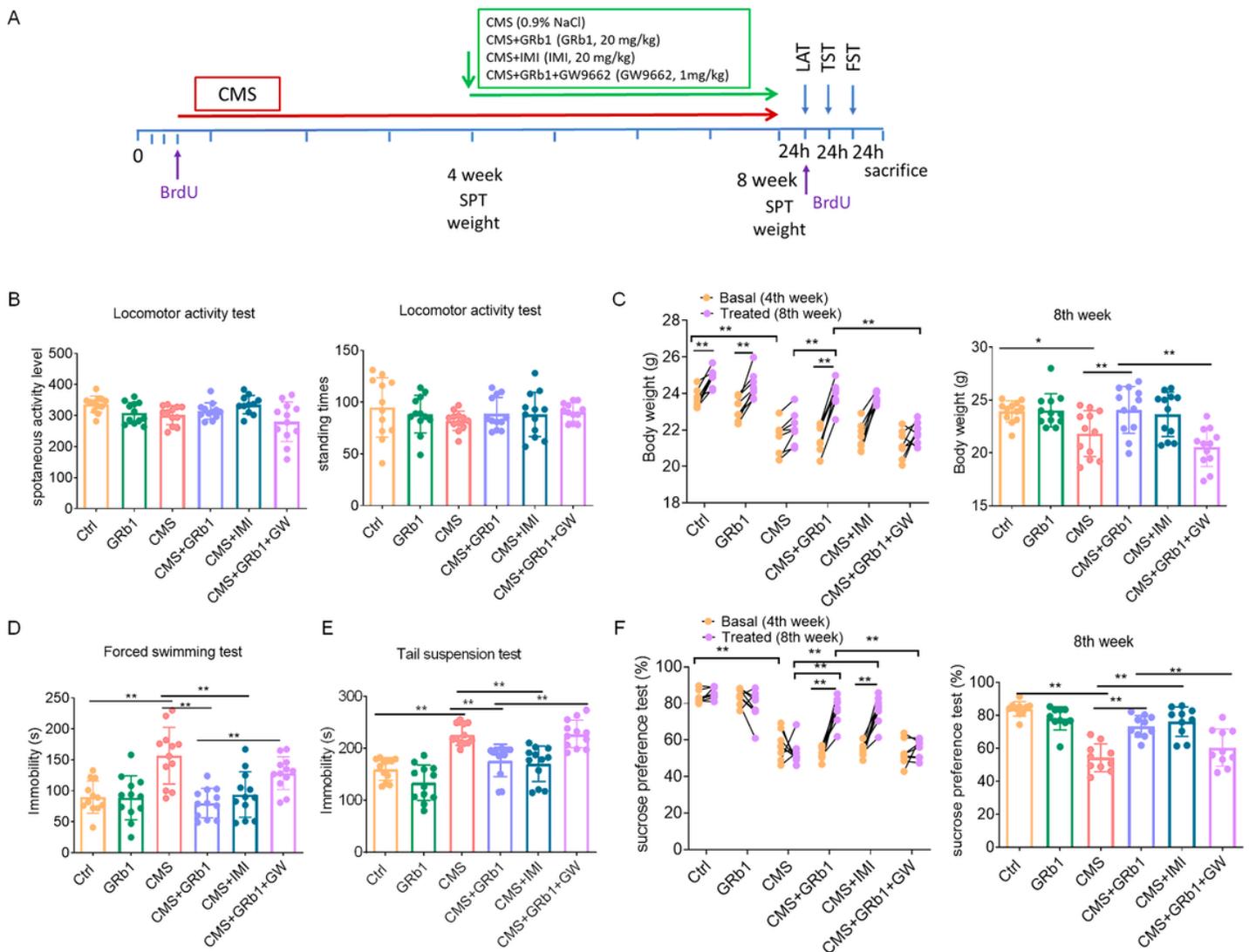
	LPS+GRb1+GW vs. LPS+GRb1	0.6983	
		0.396	
	LPS vs. Ctrl	0.0001	
	GRb1 vs. Ctrl	29.610	0.4226 5
<i>Figure 5J</i>	LPS+GRb1 vs. LPS	< 0.0001	
	LPS+GRb1+GW vs. LPS+GRb1	0.0001	
		5.007	

**Table 6** The F value and P value in multiple comparisons of figure S1

<i>figure</i>	group	F	P	N
<i>Figure 6B-1</i>	LPS vs. Ctrl		0.0454	
	10 µg/ml LPS+GRb1 vs. LPS	18.940	0.4097	6
	20 µg/ml LPS+GRb1 vs. LPS		< 0.0001	
	40 µg/ml LPS+GRb1 vs. LPS		> 0.9999	
<i>Figure 6B-2</i>	LPS vs. Ctrl		0.0100	
	10 µg/ml LPS+GRb1 vs. LPS	6.515	0.0042	6
	20 µg/ml LPS+GRb1 vs. LPS		0.0014	
	40 µg/ml LPS+GRb1 vs. LPS		0.0055	
<i>Figure 6C-1</i>	LPS vs. Ctrl		< 0.0001	
	10 µg/ml LPS+GRb1 vs. LPS	37.831	< 0.0001	6
	20 µg/ml LPS+GRb1 vs. LPS		< 0.0001	
	40 µg/ml LPS+GRb1 vs. LPS		< 0.0001	
<i>Figure 6C-2</i>	LPS vs. Ctrl		0.0133	
	10 µg/ml LPS+GRb1 vs. LPS	8.756	0.0258	6
	20 µg/ml LPS+GRb1 vs. LPS		0.0002	
	40 µg/ml LPS+GRb1 vs. LPS		0.0002	
<i>Figure 6D</i>	LPS vs. Ctrl		0.0357	
	10 µg/ml LPS+GRb1 vs. LPS		0.9795	12
		4.800		
	20 µg/ml LPS+GRb1 vs. LPS		0.0083	
	40 µg/ml LPS+GRb1 vs. LPS		0.9179	
<i>Figure 6E</i>	LPS vs. Ctrl		< 0.0001	
	10 µg/ml LPS+GRb1 vs. LPS		0.9891	12
		25.740		
	20 µg/ml LPS+GRb1 vs. LPS		< 0.0001	
	40 µg/ml LPS+GRb1 vs. LPS		< 0.0001	
<i>Figure 6F</i>	LPS vs. Ctrl		0.0284	
	10 µg/ml LPS+GRb1 vs. LPS	10.021	0.0113	6

Figure 6G	20 µg/ml LPS+GRb1 vs. LPS			< 0.0001
	40 µg/ml LPS+GRb1 vs. LPS			0.0013
	LPS vs. Ctrl			0.0404
	10 µg/ml LPS+GRb1 vs. LPS	10.740	0.9650	10
	20 µg/ml LPS+GRb1 vs. LPS			< 0.0001
	40 µg/ml LPS+GRb1 vs. LPS			0.0105

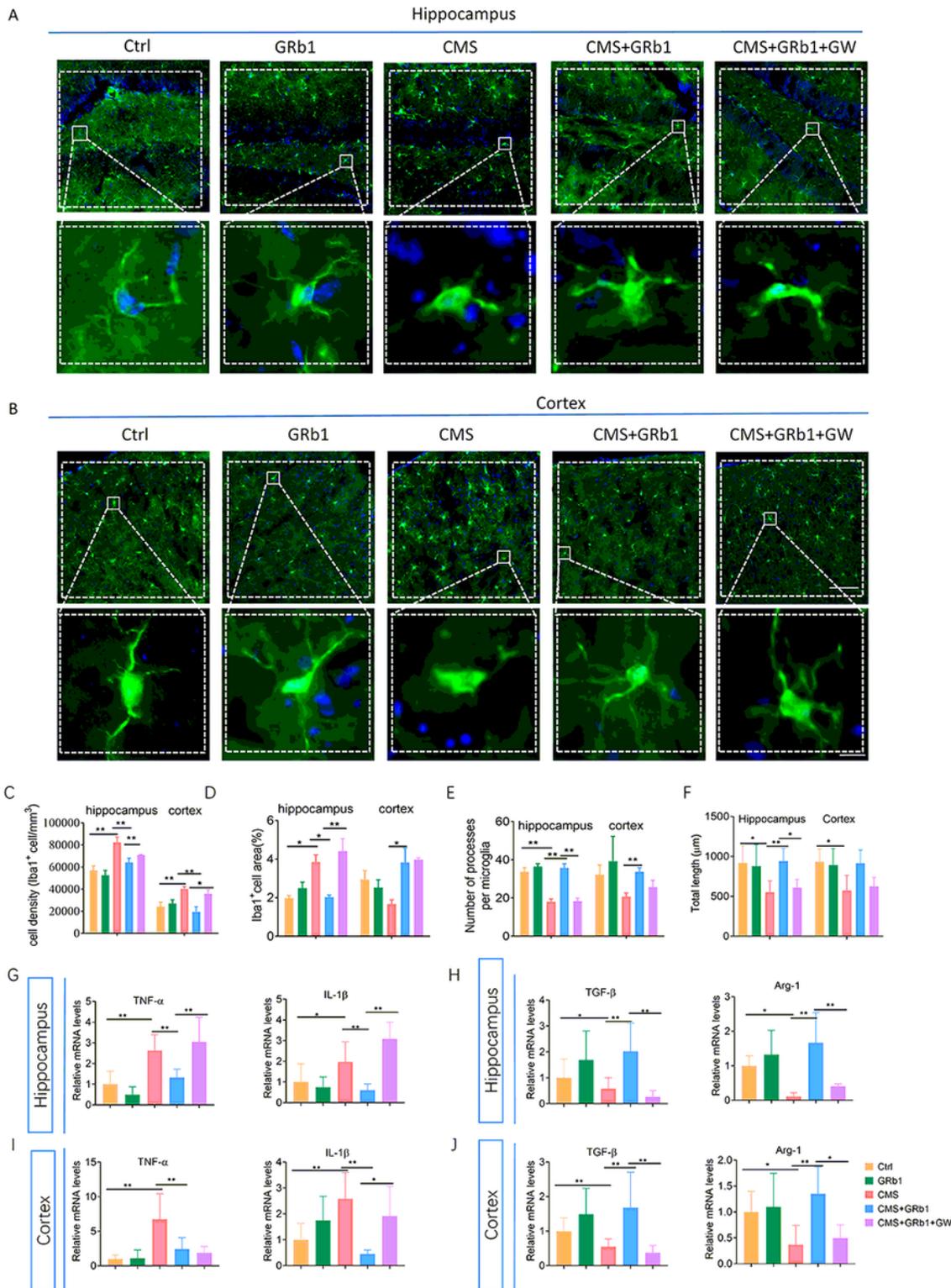
## Figures



**Figure 1**

GRb1 alleviates depressive-like behaviors in male mice exposed to CMS. (a) Drug treatment protocol. (b) Locomotor activity at week 8. (c) Comparison of body weight between week 4 and 8, and comparison of weight at week 8 among the groups. (d–e) Duration of immobility in the forced swimming and tail

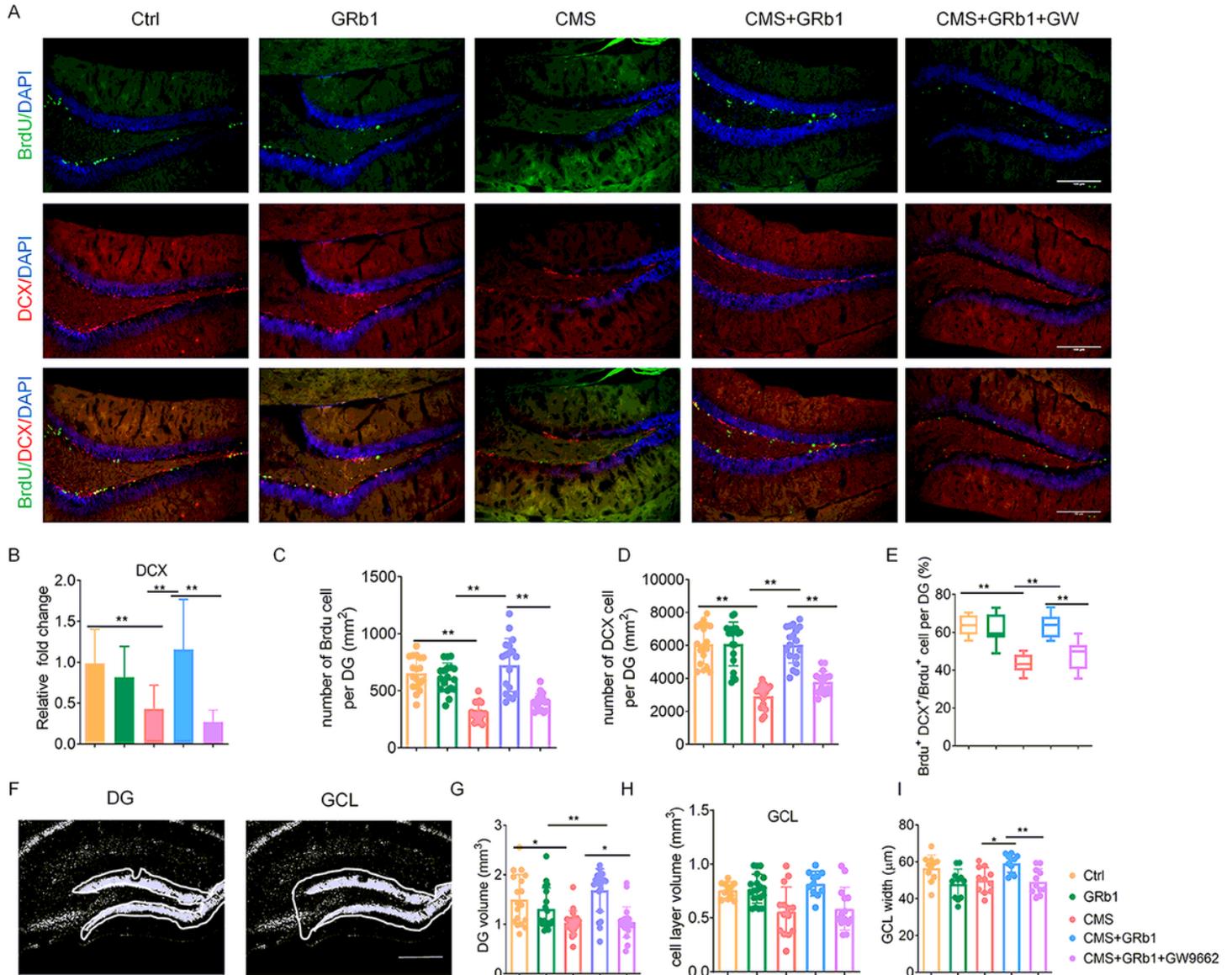
suspension tests at week 8. (f) Comparison of sucrose preference between week 4 and 8, and comparison of preference at week 8 among the groups (n = 12 mice/group). The statistical analysis results are shown in Table 1. \*P < 0.05, \*\* p < 0.01.



**Figure 2**

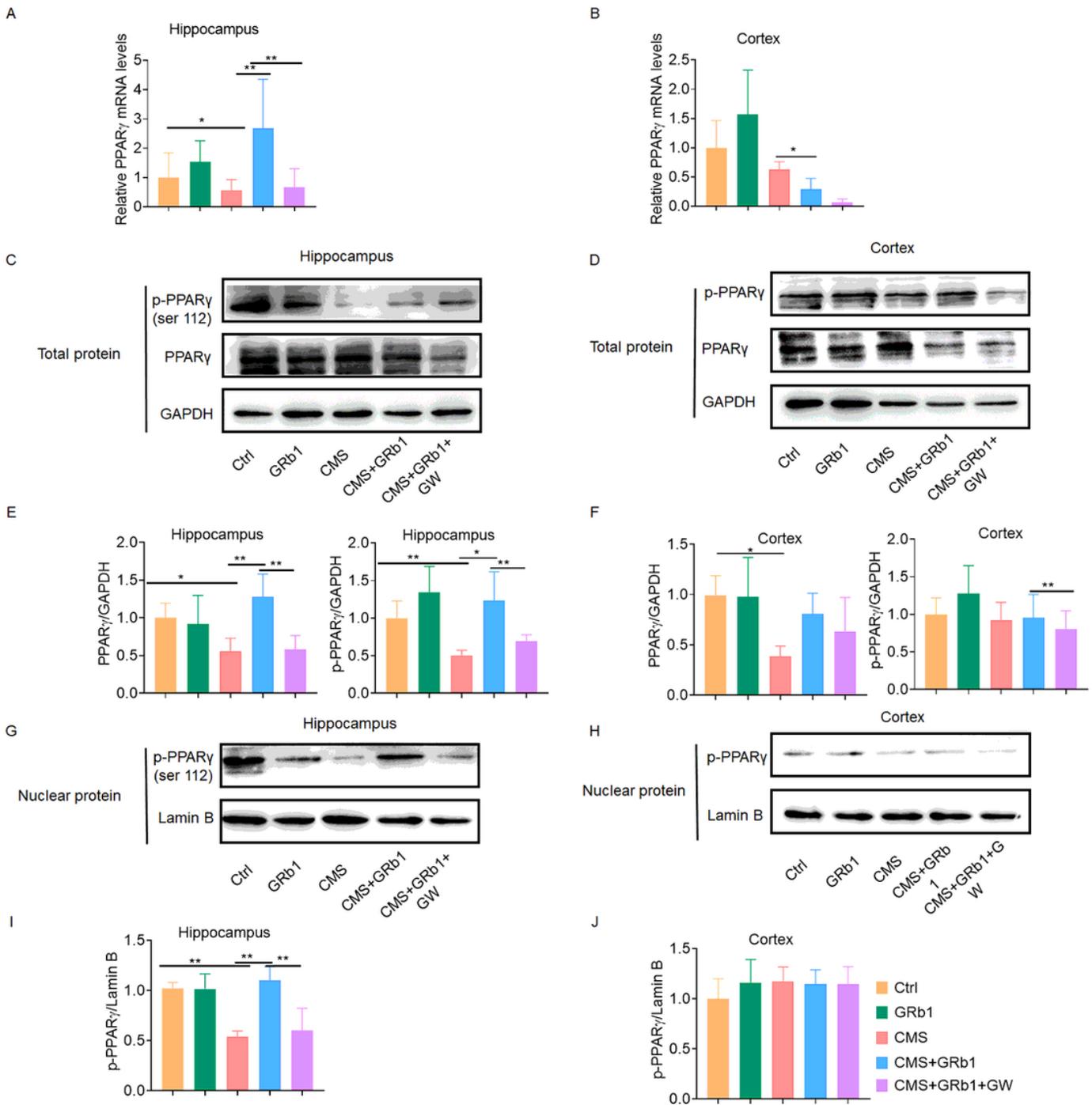
GRb1 reverses the effects of CMS on microglial phenotype and cytokine profile in mice. (a–b) Iba1<sup>+</sup> cells in the hippocampus and cortex were quantified by immunofluorescence. Microglial morphology was

quantified by cell density (c), cell area (d), the number of processes per cell (e), and total length of processes per cell (f) (n = 6 mice/group). (g) Levels of pro-inflammatory microglial markers (TNF- $\alpha$ , IL-1 $\beta$ ) in hippocampus. (h) Levels of anti-inflammatory microglial markers (TGF- $\beta$ , Arg-1) in hippocampus (n = 5 mice/group). (i-j) Changes in levels of pro- and anti-inflammatory mediators in cortex (n = 5 mice/group). Scale bars, 50  $\mu$ m (upper row in each panel), 10  $\mu$ m (lower row in each panel). The statistical analysis results are shown in Table 2. \*P < 0.05, \*\* p < 0.01.



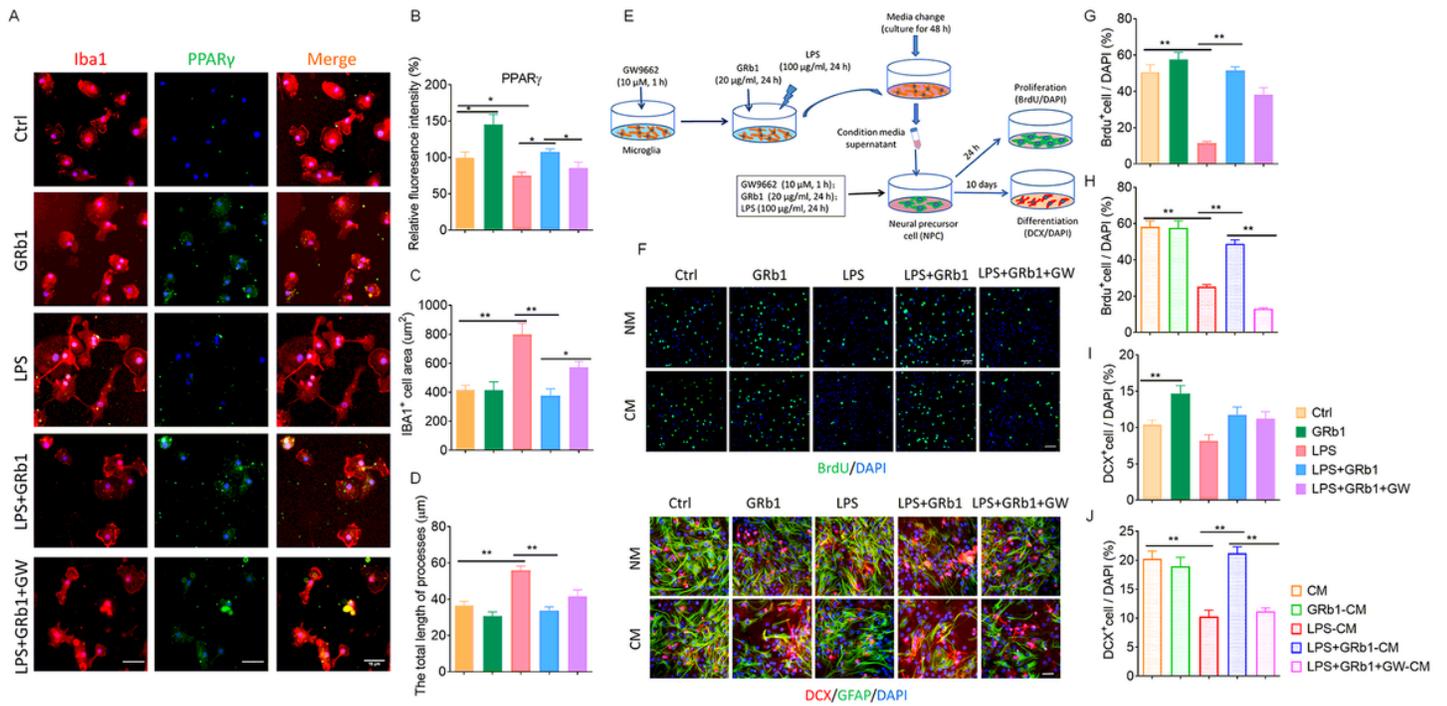
**Figure 3**

GRb1 rescues neurogenesis in the hippocampus of mice exposed to CMS. (a) Immunohistochemistry for DCX and BrdU in DG. (b) Levels of DCX mRNA in DG (n = 5 mice/group). (c-d) Number of BrdU+ or DCX+ cells in DG. (e) Differentiation ratio of proliferative cells in DG. (f) DAPI-stained sections of DG and GCL, which were quantified by measure DG volume (g), GCL volume (h), and GCL thickness (i). Scale bars: 100  $\mu$ m. Data are shown as mean  $\pm$  SEM (n = 6 mice/group). The statistical analysis results are shown in Table 3. \*P < 0.05, \*\* p < 0.01.



**Figure 4**

GRb1 rescues PPAR $\gamma$  expression in mice exposed to CMS. (a–b) Relative PPAR $\gamma$  mRNA levels in the hippocampus and cortex. (c–d) Representative western blot of p-PPAR $\gamma$  and PPAR $\gamma$  in hippocampus and cortex. (e–f) Quantification of PPAR $\gamma$  and p-PPAR $\gamma$  in hippocampus and cortex. (g–h) Representative western blot of p-PPAR $\gamma$  in nuclei of hippocampus and cortex. (i–j) Quantification of p-PPAR $\gamma$  in nuclei of hippocampus and cortex (n = 5 mice/group). The statistical analysis results are shown in Table 4. \*P < 0.05, \*\* p < 0.01.



**Figure 5**

PPAR $\gamma$ -dependent pathways mediate effects of GRb1 in vitro. (a) Representative micrographs after immunostaining against Iba1 display microglial activation phenotype. Fluorescence intensity reflects PPAR $\gamma$  expression. Scale bars: 20  $\mu$ m. (b) Quantitation of PPAR $\gamma$  fluorescence intensity, (c) unbiased stereological quantification of microglial cell area, and (d) total length of processes. (e) Experimental protocol to test the effects of conditioned medium from microglial cultures on neural precursor cell (NPC) cultures. (f) NPCs cultures were maintained in native medium (NM) and directly stimulated by GRb1, LPS, and GW9662; or they were maintained in conditioned medium (CM) of activated microglia. After 1 day or 10 days, the proliferation and differentiation of NPCs were tested, respectively, by immunolabeling BrdU (upper two rows) or DCX and GFAP (lower two rows). Scale bars: 15  $\mu$ m. (g–h) The ratio of BrdU+ to DAPI+ cells in NPCs cultures maintained in native medium or exposed to conditioned medium from microglial cultures. (i–j) The ratio of DCX+ to DAPI+ cells in NPCs cultures maintained in native medium or exposed to conditioned medium. The statistical analysis results are shown in Table 5. \*P < 0.05, \*\* p < 0.01.

## Supplementary Files

This is a list of supplementary files associated with this preprint. Click to download.

- [Onlinefig.S1.png](#)