

Facilitating microglial phagocytosis by which Jiawei Xionggui Decoction alleviates cognitive impairment via TREM2-mediated energy metabolic reprogramming

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Citation: Wen Wen, Jie Chen, Junbao Xiang, Shiqi Zhang, Jingru Liu, Jie Wang, Ping Wang, Shijun Xu, Facilitating microglial phagocytosis by which Jiawei Xionggui Decoction alleviates cognitive impairment via TREM2-mediated energy metabolic reprogramming, *Chinese Journal of Natural Medicines*, 2025, 23(8), 909–919. doi: [10.1016/S1875-5364\(25\)60927-7](https://doi.org/10.1016/S1875-5364(25)60927-7).

View online: [https://doi.org/10.1016/S1875-5364\(25\)60927-7](https://doi.org/10.1016/S1875-5364(25)60927-7)

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Chinese Journal of Natural Medicines

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Original article

Facilitating microglial phagocytosis by which Jiawei Xionggui Decoction alleviates cognitive impairment *via* TREM2-mediated energy metabolic reprogrammingWen Wen^{a,b,c}, Jie Chen^{a,b,c}, Junbao Xiang^{a,b,c}, Shiqi Zhang^{a,b,c}, Jingru Liu^d, Jie Wang^{a,b,c}, Ping Wang^{a,b,c,*}, Shijun Xu^{a,b,c,*}^a State Key Laboratory of Southwestern Chinese Medicine Resources, Chengdu University of Traditional Chinese Medicine, Chengdu 611137, China^b School of Pharmacy, Chengdu University of Traditional Chinese Medicine, Chengdu 611137, China^c Institute of Material Medica Integration and Transformation for Brain Disorders, Chengdu University of Traditional Chinese Medicine, Chengdu 611137, China^d Division of Biosciences, University College London, London WC1E6BT, UK

ARTICLE INFO

Article history:

Received 6 September 2024

Revised 21 December 2024

Accepted 26 February 2025

Available online 20 August 2025

Keywords:

Akt/mTOR/HIF-1 α

Energy metabolic reprogramming

Jiawei Xionggui Decoction

Microglial phagocytosis

Triggering receptor expressed on myeloid cells 2 (TREM2)

ABSTRACT

Triggering receptor expressed on myeloid cells 2 (TREM2)-mediated microglial phagocytosis is an energy-intensive process that plays a crucial role in amyloid beta (A β) clearance in Alzheimer's disease (AD). Energy metabolic reprogramming (EMR) in microglia induced by TREM2 presents therapeutic targets for cognitive impairment in AD. Jiawei Xionggui Decoction (JWXG) has demonstrated effectiveness in enhancing energy supply, protecting microglia, and mitigating cognitive impairment in APP/PS1 mice. However, the mechanism by which JWXG enhances A β phagocytosis through TREM2-mediated EMR in microglia remains unclear. This study investigates how JWXG facilitates microglial phagocytosis and alleviates cognitive deficits in AD through TREM2-mediated EMR. Microglial phagocytosis was evaluated through immunofluorescence staining *in vitro* and *in vivo*. The EMR level of microglia was assessed using high-performance liquid chromatography (HPLC) and enzyme-linked immunosorbent assay (ELISA) kits. The TREM2/protein kinase B (Akt)/mammalian target of rapamycin (mTOR)/hypoxia-inducible factor-1 α (HIF-1 α) signaling pathway was analyzed using Western blotting in BV₂ cells. TREM2^{-/-} BV₂ cells were utilized for reverse validation experiments. The A β burden, neuropathological features, and cognitive ability in APP/PS1 mice were evaluated using ELISA kits, immunohistochemistry (IHC), and the Morris water maze (MWM) test. JWXG enhanced both the phagocytosis of EMR disorder-BV₂ cells (EMRD-BV₂) and increased EMR levels. Notably, these effects were significantly reversed in TREM2^{-/-} BV₂ cells. JWXG elevated TREM2 expression, adenosine triphosphate (ATP) levels, and microglial phagocytosis in APP/PS1 mice. Additionally, JWXG reduced A β -burden, neuropathological lesions, and cognitive deficits in APP/PS1 mice. In conclusion, JWXG promoted TREM2-induced EMR and enhanced microglial phagocytosis, thereby reducing A β deposition, improving neuropathological lesions, and alleviating cognitive deficits.

1. Introduction

Amyloid beta (A β) accumulation and its associated neurotoxicity represent critical factors in the neurological deterioration observed in Alzheimer's disease (AD). The enhancement of A β clearance has emerged as a primary strategy in drug development, with several monoclonal antibodies receiving Food and Drug Administration (FDA) approval for AD treatment, including aducanumab and lecanemab¹⁻³. These developments demonstrate that facilitating A β removal effectively improves cognitive impairment in AD. Microglial phagocytosis, performed by the brain's exclusive immune cells, presents a distinct target for A β clearance^{1,4-7}, though effective therapeutic interventions remain

unavailable.

Triggering receptor expressed on myeloid cells 2 (TREM2), a significant risk gene in AD, is predominantly expressed in brain microglia and plays a crucial role in microglial functions, including phagocytosis, energy metabolism, and survival⁸. TREM2 enhances microglial activation and phagocytosis through mammalian target of rapamycin (mTOR) pathway stimulation⁹. Additionally, A β functions as a TREM2 ligand, binding directly to its extracellular domain. Acute A β administration triggers TREM2-mediated activation of the protein kinase B (Akt)/mTOR/hypoxia-inducible factor-1 α (HIF-1 α) signaling pathway¹⁰. HIF-1 α serves as a key transcriptional regulator of glycolysis by modulating the expression and activity of glucose metabolism-related enzymes¹¹. This process initiates microglial energy metabolic reprogramming (EMR) from oxidative phosphorylation (OXPHOS) to glycolysis, essential for maintaining microglial phagocytosis^{12,13}, a phenomenon known as the Warburg effect. The Akt/mTOR/HIF-

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1 α pathway represents a well-established signaling mechanism for EMR^{12,14}. Prolonged or repeated A β exposure leads to overall microglial energy metabolism deficiency, resulting in reduced or absent Akt/mTOR/HIF-1 α pathway responsiveness, thereby inhibiting microglial EMR, a condition termed microglial immune tolerance^{15,16}. This tolerance exacerbates microglial phagocytic dysfunction and impairs A β clearance, leading to excessive A β accumulation and subsequent cognitive decline¹⁷. Recent research demonstrates that ATV/TREM2 (an agonistic anti-TREM2 antibody) improves energy metabolism and microglial phagocytosis in Alzheimer's disease models *in vivo* and *in vitro*^{18,19}. These findings suggest that enhancing microglial phagocytosis through EMR promotion represents an effective strategy for addressing AD-related cognitive impairment.

Numerous studies have demonstrated that traditional Chinese medicine (TCM) shows significant potential in developing new treatments for AD²⁰⁻²². Research confirms that TCM can mitigate cognitive dysfunction and neuronal damage in AD models by enhancing microglial phagocytosis, as evidenced by studies on *Dactyloctenium aegyptium* stemwood, Juzen Taihoto and Shiquan Dabu Formula²³⁻²⁵. Jiawei Xionggui Decoction (JWXG), a classic formula derived from *Wan Shi Nu Ke*, comprises Chuanxiong Rhizoma (the rhizoma and root of *Ligusticum chuanxiong*), Scutellariae Radix (the radix of *Scutellaria baicalensis*), and Angelica Sinensis Radix (the radix of *Angelica sinensis*). Its composition ratio and application for cognitive impairment improvement have received patent authorization in China (Invention Patent Authorisation No. ZL201310132096.2). Previous studies demonstrate that JWXG enhances cognitive function, decreases A β burden, and inhibits microglial inflammatory response in AD model mice²⁶⁻²⁸. Furthermore, JWXG enhances brain energy metabolism in APP/PS1 mice and multiple infarct dementia rats while modulating lactate content by facilitating monocarboxylate transporter-1 and monocarboxylate transporter-2 expression in multiple infarct dementia rats²⁹⁻³². Additionally, chlorogenic acid, an active constituent of JWXG, activates the mTOR/HIF-1 α signaling pathway³³, the primary pathway regulating EMR. Research indicates that JWXG-derived active ingredients, including baicalin, baicalein, chlorogenic acid, and ferulic acid, effectively reduce cognitive deficits in AD models³⁴⁻³⁹. Notably, baicalin and ferulic acid serve as quality control indices for *Ligusticum chuanxiong*, *Scutellaria baicalensis* and *Angelica sinensis* required in the *Chinese Pharmacopoeia 2020*. Thus, their contents were utilized to assess JWXG quality in this study. This investigation employed 6-month-old (6-mo) transgenic APP/PS1 mice, transgenic TREM2^{-/-}-BV₂ cells, and amyloid- β oligomers (oA β)-induced EMR disorder-BV₂ (EMRD-BV₂) to elucidate JWXG's therapeutic effects and underlying mechanisms in reducing AD cognitive deficits, particularly focusing on TREM2-mediated microglial phagocytosis and EMR.

2. Material and methods

2.1. JWXG preparation

Chuanxiong Rhizoma (Batch No. 2020088) and Scutellariae Radix (Batch No. 2020001) were obtained from Sichuan NEAUTUS Traditional Chinese Medicine Co., Ltd. (Chengdu, China); Angelica Sinensis Radix (Batch No. 202001) was provided by Chengdu Ji'an Kang Medicine Industry Co., Ltd. (Chengdu, China). These herbs were authenticated by Prof. Yuntong Ma (Chengdu University of Traditional Chinese Medicine, Chengdu, China). The JWXG preparation involved boiling 15 g Chuanxiong Rhizoma, 9 g Scutellariae Radix, and 3 g Angelica Sinensis Radix twice in 250 mL of water, followed by filtration and mixing. The decoction was concentrated under vacuum with a rotary evaporator to

0.45 g·mL⁻¹ for content determination. The liquid extract underwent low-temperature vacuum freeze-drying (-80 °C) to produce freeze-dried powder, which was stored in a dryer. The clinical JWXG dosage is 27 g daily for patients. The mice clinical equivalent dosage was calculated based on kg mice/human body weight using a conversion ratio of 9 times, as documented in *Experimental Pharmacology*. Consequently, the daily JWXG dosage for mice was established at 4.05 g·kg⁻¹ in this study.

2.2. High-performance liquid chromatography (HPLC) analysis

After filtration through a 0.22 μ m microporous membrane and dilution to 1:100, an Agilent 1260 liquid chromatograph system (Agilent, USA) and Q Exactive liquid chromatography-tandem mass spectrometry (LC-MS/MS) (Thermo Fisher, USA) were utilized to analyze baicalin, baicalein, ferulic acid and chlorogenic acid in JWXG according to the *Chinese Pharmacopoeia 2020* method for *Ligusticum chuanxiong*, *Scutellaria baicalensis* and *Angelica sinensis*. The compounds were separated using a Poroshell 120 Energy charge (EC)-C₁₈ (4.6 mm \times 100 mm, 2.7 μ m, Agilent, USA) chromatographic column. The elution gradient consisted of purified 0.1% aqueous formic acid (A) and 0.1% acetonitrile formate solution (B). Gradient elution proceeded according to the following program: 0 \rightarrow 20 min, 95% \rightarrow 5% A. The column temperature and injection volume were set to 30 °C and 1 μ L, respectively. Quantification was performed by multiple reaction monitoring mode with transitions of *m/z* 290.988 \rightarrow 123.054 \rightarrow 32.17 for baicalin (+), *m/z* 447.088 \rightarrow 123.071 \rightarrow 54.03 for baicalein (+), *m/z* 193.05 \rightarrow 134.071 \rightarrow 14.86 for ferulic acid (-), *m/z* 353.125 \rightarrow 84.982 \rightarrow 39.63 for chlorogenic acid (-). Other main working parameters were set as usual.

2.3. Cell culture, lentivirus infection and stimulation

The lentivirus target virus [TREM2-short hairpin ribonucleic acid (shRNA)] and an equivalent control virus were used to treat BV₂ cells. After 72-h stimulation, unsuccessfully infected BV₂ cells were eliminated using 5 μ g·mL⁻¹ of puromycin to obtain stable shTREM2-BV₂ and GFP-BV₂ cells. A β ₂₅₋₃₅ peptides (Chinese peptide) were dissolved in HFIP (RHAWN) at a final concentration of 1 mmol·L⁻¹, followed by HFIP evaporation to collect A β peptide and storage at -80 °C. oA β was prepared by dissolving A β ₂₅₋₃₅ peptides to 100 μ mol·L⁻¹ in Dulbecco's modified Eagle medium (DMEM), followed by overnight incubation at 4 °C. BV₂ cells were obtained from the Institute of Basic Medical Sciences, Chinese Academy of Medical Sciences (Identification No. 1101MOU-PUMC000063). BV₂ cells were cultured in basic DMEM containing 10% FBS and 1% P/S and stimulated with either vehicle (DMEM containing 0.5% FBS to prevent cell death), 2 μ mol·L⁻¹ oA β , or 2 μ mol·L⁻¹ oA β + JWXG at varying concentrations for 24 h. In specific experiments, BV₂ cells were pre-treated with 5 mmol·L⁻¹ 2-DG for 1 h to inhibit glycolysis. The experiments employed a tolerant model simulating prolonged conditions, where cells were exposed to vehicle or oA β (2 μ mol·L⁻¹) for 24 h, washed with phosphate-buffered saline (PBS), and further cultured in vehicle DMEM for 3-5 d. Subsequently, the cells were re-stimulated with oA β (2 μ mol·L⁻¹), 2 μ mol·L⁻¹ oA β + JWXG at different concentrations, or incubated with a vehicle for 24 h (Fig. 1).

2.4. Phagocytosis assay

The phagocytosis assay was performed according to previously described methods⁴⁰. Briefly, BV₂ cells were incubated at a density of 30 000/well in 12-well plates. Following re-stimulation with oA β (2 μ mol·L⁻¹) or alternative stimuli, pre-opsonized fluorescent latex beads (50% PBS + 50% FBS, FL-PS-G-01 DaE

Technology, Tianjin, China) were added to the wells at a density of 50 beads/cell and incubated at 37 °C for 4 h. After gentle washing to remove non-phagocytosed beads, the cells were fixed with 4% PFA at room temperature. Images were captured using a fluorescent microscope, and phagocytosed beads were counted.

2.5. Glycolysis measurements

EC assays were conducted according to the previously reported protocol, and the levels of adenosine triphosphate (ATP), adenosine diphosphate (ADP), and adenosine monophosphate (AMP) were determined using the standard curve²⁷. EC was calculated using the following formula:

$$EC = \frac{[ATP] + 0.5 \times [ADP]}{[ATP] + [ADP] + [AMP]}$$

Pyruvate kinase (PK), fructose-6-phosphate kinase (F-6-PK), and hexokinase (HK) activity tests were performed to evaluate the level of EMR in BV₂ cells, following the standard protocols specified in the PK, F-6-PK, and HK activity kit instructions (Suzhou Grace Biotechnology Co., Ltd., Suzhou, China). The intracellular content of lactate and pyruvate in BV₂ cells was measured to determine the ratio between cellular glycolysis and tricarboxylic acid cycle, following the standard protocols specified in the lactate and pyruvate content kit instructions (Nanjing JianCheng Bioengineering Institute Co., Ltd., Nanjing, China).

2.6. Real-time quantitative polymerase chain reaction (RT-qPCR)

Total RNA was extracted using Total RNA Isolation Kits (#R220901, Foregene). cDNA synthesis was performed using a SweScript All-in-One RT SuperMix kit (#G3337-50, ServiceBio). RT-qPCR analysis was conducted using SYBR Green mix (#G3326-05, ServiceBio) with the following primer pairs: *MM-CD68*: 5'-CTC TCT AAG GCT ACA GGC TGC T-3' (forward) and 5'-TCA CGG TTG CAA GAG AAA CA-3' (reverse); *MM-TREM2*: 5'-TAC TGG TGG AGG TGC TGG AG-3' (forward) and 5'-AGG ATG AAA CCT GCC TGG AG-3' (reverse); *MM-β-actin*: 5'-GCT CCG GCA TGT GCA AAG-3' (forward) and 5'-TTC CCA CCA TCA CAC CCT GG-3' (reverse). The relative expression of target gene messenger RNA (mRNA) was analyzed according to a standard procedure¹².

2.7. Western blotting

Western blotting was performed following a standard protocol⁴¹. Briefly, the denatured target proteins were separated using SDS-PAGE, transferred to PVDF membranes, and blocked in a blocking solution (5% skimmed milk dissolved in TBST). Primary antibodies including rabbit anti-HIF-1α (#36169S, CST, MA, USA), rabbit anti-TREM2 (#bs-2723R, Bioss, Beijing, China), rab-

bit anti-phosphorylated (p)-mTOR (Ser 2448, #5536T, CST), rabbit anti-mTOR (#2983S, CST), rabbit anti-p-Akt (Ser473, #4060S, CST), rabbit anti-Akt (#4685S, CST), and rabbit anti-β-actin (#GB12001, Servicebio, Wuhan, China) were applied at a dilution ratio of 1:1000 for 12 h at 4 °C. Subsequently, horseradish peroxidase (HRP)-conjugated secondary antibody was applied for 2 h at room temperature. The membranes were analyzed using a chemiluminescence reagent and quantified using a chemiluminescent detection software system (ChampChemi 610 Plus, Beijing Sage Creation Science Co., Ltd., Beijing, China) and analyzed using quantity one software.

2.8. Animals and treatment

The experimental protocols were approved by the Animal Care and Use Committee of the Institute of Material Medica Integration and Transformation for Brain Disorders (No. IBD 20200010). 3-mo APP/PS1 mice were acquired from Beijing HFK Bio-Technology Co., Ltd., Beijing, China (Certificate No. 1103222011009531), while C57BL/6J mice were obtained from Hunan SJA Laboratory Animal Co., Ltd., Changsha, China (Certificate No. 1107272011006639). All mice received unrestricted access to food and water in a specific pathogen-free (SPF) facility at Chengdu University of TCM. All experimental procedures strictly adhered to the guidelines of the National Institutes of Health (NIH, USA). Following three months of domestication and adaptive feeding, APP/PS1 mice (JWXG group) received daily oral administration of JW for three months, while C57BL/6J mice and the APP/PS1 group received an equal volume of 0.9% saline. Subsequently, mice underwent a Morris water maze (MWM) test following a standard protocol⁴². After the MWM test, animals were anesthetized with 40 mg·kg⁻¹ sodium pentobarbital saline solution (#69020100, i.p., Merck, Darmstadt, Germany). The mouse brain tissue underwent transcranial perfusion with pre-cooled normal saline and was collected after euthanasia. The right hemisphere was preserved in liquid nitrogen for Western blotting, RT-qPCR, and enzyme-linked immunosorbent assay (ELISA) kit analysis, while the left hemisphere was fixed with 4% paraformaldehyde (G1101, Service Bio., Wuhan, China) for morphological analysis (Fig. 1).

2.9. MWM test

The Noldus mice MWM was utilized to assess spatial learning and memory capabilities in mice (Noldus Information Technology, Wageningen, Netherlands). An escape platform (diameter, 4.5 cm) was positioned 1 cm below the water surface. During testing, animals were required to locate the hidden escape platform within 60 s. If unsuccessful, the animal was guided to the platform for 10 s at test completion. The assessment spanned 6 d,

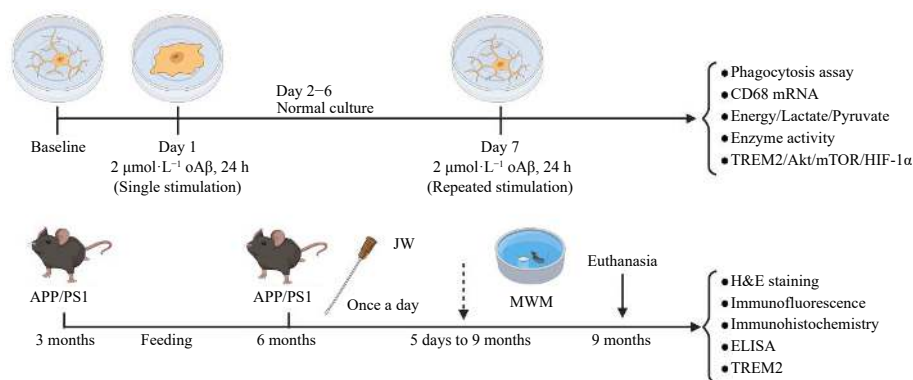


Fig. 1 The experimental flow chart of this work *in vitro* and *in vivo*.

comprising 5 training days and a partial probe trial on day 6 without the platform. On the final day, animals underwent a single probe test without the platform. The escape latency, time to reach the platform or platform zone, duration in the platform or platform zone, and frequency of crossing the platform or platform zone were measured to assess learning and memory capacity.

2.10. Detection of $A\beta_{1-40}/A\beta_{1-42}$ and lactic acid of brain tissue

The ELISA kits of $A\beta_{1-40}$ and $A\beta_{1-42}$ were employed to quantify the levels of $A\beta_{1-40}$ and $A\beta_{1-42}$ in brain tissue. Following the manufacturer's instructions (Elabscience Biotechnology Co., Ltd., Wuhan, China), brain samples were prepared by grinding with PBS. After completing a standard protocol, the OD value was measured at 450 nm following 30 min incubation using the Multiskan spectrum spectrophotometer.

Mouse brain tissue (20 mg) was ground on ice with pure water, and the supernatant was collected after ultrasonic crushing and centrifugation. This procedure was repeated twice. All supernatant fractions were concentrated under vacuum using a rotary evaporator. The samples were prepared, and lactic acid was analyzed according to the previously described protocol⁴³.

2.11. Hematoxylin and eosin (H&E), IHC, and immunofluorescence (IF) staining

H&E staining was performed to evaluate neuronal damage in the cortex and hippocampus of mice, as previously reported³⁹. Brain tissue sections (4 μ m) were dewaxed with xylene and stained, followed by visualization of each stained slice. IHC and IF analyses followed standard protocols^{42,44}. The 4- μ m brain tissue sections underwent baking, rehydration, antigen retrieval, and peroxide blocking before overnight incubation at 4 °C with mouse anti- β -amyloid (1:500, #15126, CST, MA, USA), mouse anti-AT8 (1:500, #MN1020, Thermo Fisher, MA, USA), and rabbit anti-Iba1 (1:500, #GB113502-100, Servicebio, Wuhan, China). Subsequently, sections were treated with HRP-conjugated secondary antibody, Cy3 anti-rabbit secondary antibody (1:300), or 488 anti-mouse secondary antibody (1:300) for 2 h at RT. IHC sections were developed using a DAB kit (#AR1022, BOSTER, Wuhan, China) and counterstained with nuclear hematoxylin. IF sections were counterstained with DAPI solution (2 μ g·mL⁻¹, #G1012, Servicebio, Wuhan, China). IHC sections were analyzed using ImageJ. For microglia quantification around $A\beta$, Iba1 positive cells within 20 μ m of $A\beta$ plaques were manually counted.

2.12. Statistical analysis

ImageJ software was employed for quantitative analysis of H&E, IHC, and IF images, while Quantity One software was used for quantifying Western blotting bands. All results were expressed as mean \pm SEM and analyzed using the R language. Double-factor variance analysis was conducted to evaluate escape latency in the MWM test. Differences between groups were assessed using one-way analysis of variance. Statistical significance was established at $P < 0.05$.

3. Results

3.1. The content of active component in JWXG

LC-MS/MS methodology was utilized to quantify the content of chlorogenic acid, ferulic acid, baicalin, and baicalein, with all four active ingredients detected in JWXG. Based on the standard curve ($R^2 > 0.99$), their respective contents in JWXG were deter-

mined to be 7.96, 1.66, 65.59, and 123.90 μ g·mL⁻¹ (see Supporting information).

3.2. JWXG enhanced microglial phagocytosis in EMRD-BV₂ cells and in APP/PS1 mice

As CD68 serves as a marker of microglial phagocytosis, the expression of *CD68* mRNA was initially measured to assess JWXG's phagocytosis-promoting effect in BV₂ cells under different energy metabolic states. Compared to BV₂ cells, the *CD68* mRNA level in EMR-BV₂ cells (oA β acute stimulation) showed significant elevation (Fig. 2A, $P < 0.01$), while its expression remained unchanged in EMRD-BV₂ cells (repeatedly stimulated by oA β) (Fig. 2B, $P > 0.05$), suggesting that oA β reduplicative stimulation-induced EMRD inhibited microglial phagocytosis. Notably, JWXG significantly enhanced the expression of *CD68* mRNA in EMRD-BV₂ cells in a dose-dependent manner (Fig. 2B, $P < 0.05$). Consistent with *CD68* mRNA expression, JWXG significantly increased the number of fluorescent latex beads within EMRD-BV₂ cells in a dose-dependent pattern (Figs. 2C and 2D, $P < 0.05$ or $P < 0.01$). These observations indicate that JWXG enhances microglial phagocytosis ability in EMRD-BV₂ cells. As anticipated, the ratio of Iba1⁺ microglia to $A\beta$ plaques was significantly higher in JWXG mice compared to APP/PS1 mice, accompanied by substantial microglial accumulation around $A\beta$ plaques (Figs. 2E and 2F, $P < 0.05$). Additionally, the *CD68* mRNA expression in brain tissue showed no significant difference between C57BL/6j and APP/PS1 mice but was markedly elevated following JWXG administration (Fig. 2G, $P < 0.01$). These findings demonstrate that JWXG enhances microglial phagocytosis both *in vitro* and *in vivo*.

3.3. JWXG increased energy production by enhancing EMR *in vitro* and *in vivo*

Brain energy deficiency is a well-documented pathological feature in Alzheimer's disease (AD) and associated animal models, largely due to a metabolic shift from efficient mitochondrial respiration to less efficient glycolysis⁴⁵⁻⁴⁸. To assess the potential of JWXG in restoring cellular energy under energy-restricted conditions, we examined its effects on BV₂ microglial cells subjected to EMRD. No significant difference in EC levels was observed between normal BV₂ and EMRD-BV₂ cells; however, JWXG treatment led to a dose-dependent increase in EC levels in EMRD-BV₂ cells (Fig. 3A, $P < 0.05$). Lactic acid, pyruvate, and the lactic acid/pyruvate ratio—key metabolic indicators of glycolytic activity—also showed no baseline differences between BV₂ and EMRD-BV₂ cells. Nevertheless, all three parameters were significantly elevated following JWXG administration in EMRD-BV₂ cells (Fig. 3B-3D, $P < 0.05$ or $P < 0.01$). The activities of the glycolytic rate-limiting enzymes HK, PFK, and PK did not differ between BV₂ and EMRD-BV₂ cells at baseline but were significantly increased in a dose-dependent manner by JWXG in EMRD-BV₂ cells (Figs. 3E-3G, $P > 0.05$). These enhancements were effectively reversed by 2-DG, a glycolytic inhibitor targeting HK, with the exception of HK activity, which remained elevated. *In vivo*, the brain EC level in APP/PS1 transgenic mice was significantly reduced compared with wild-type C57BL/6j mice, but was markedly restored after JWXG treatment (Fig. 3H, $P < 0.05$ or $P < 0.01$). Correspondingly, brain lactic acid levels were significantly higher in JWXG-treated APP/PS1 mice compared to untreated counterparts (Fig. 3I). These findings suggest that JWXG promotes energy production in both *in vitro* and *in vivo* models by enhancing glycolytic metabolism.

3.4. JWXG promoted microglial phagocytosis through TREM2-mediated Akt/mTOR/HIF-1 α pathway

The Akt/mTOR/HIF-1 α pathway represents a crucial mech-

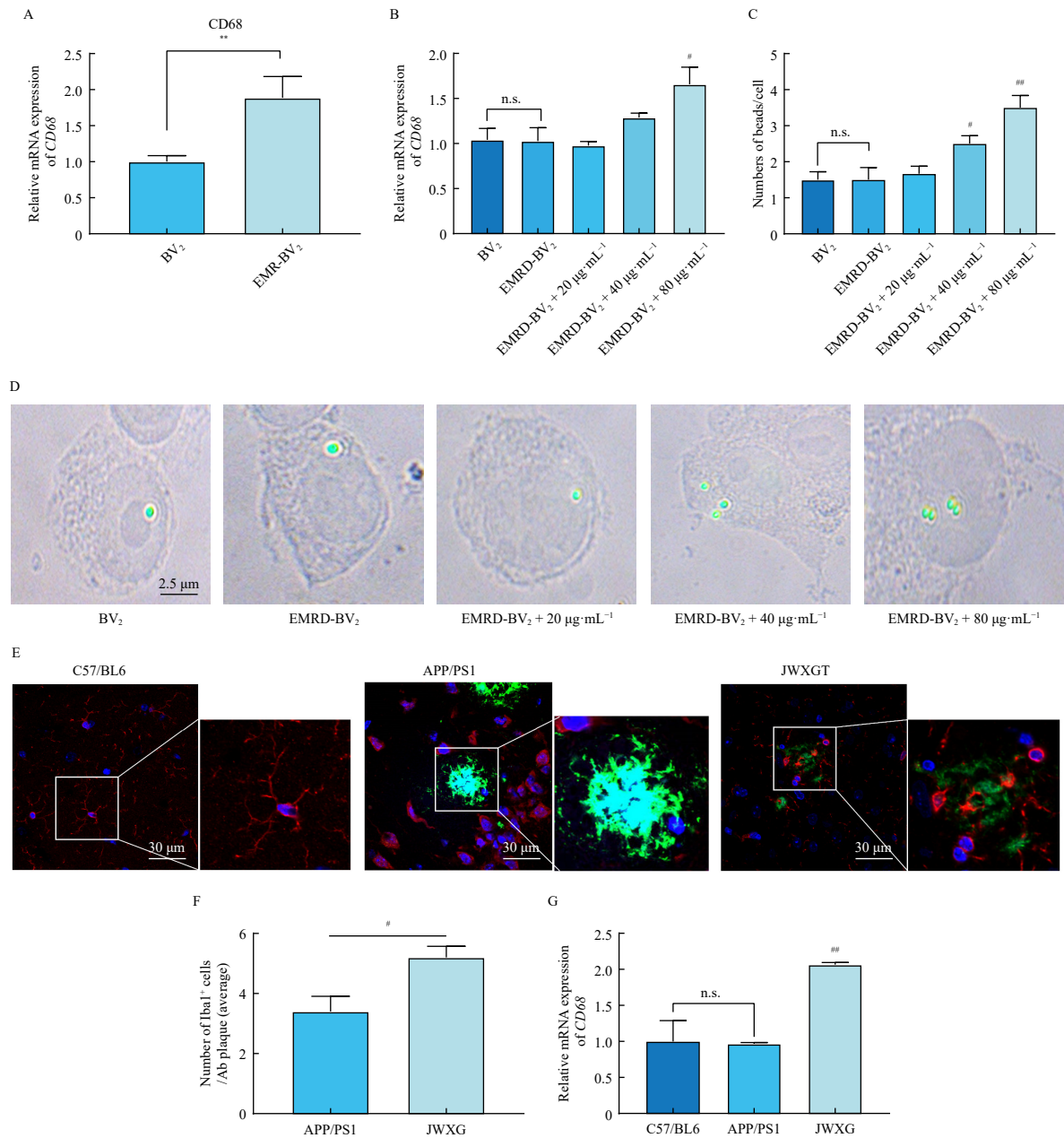


Fig. 2 JWXG enhanced phagocytosis of EMRD-BV₂ and microglia in APP/PS1 mice. (A, B) *CD68* mRNA levels *in vitro* ($n = 6$). (C) Quantification of internalized fluorescent latex beads *in vitro* ($n = 6$). (D) Representative images of phagocytosed fluorescent latex beads *in vitro*. (E) Immunofluorescence images showing colocalization of Iba1⁺ microglia (red), A β plaques (green), and nuclei (DAPI, blue) in the cortex of mice. (F) Quantification of Iba1⁺ microglia colocalized with A β plaques in APP/PS1 mice ($n = 3$). (G) *CD68* mRNA level of mice ($n = 3$). The data are expressed as mean \pm SEM. * $P < 0.05$, ** $P < 0.01$ vs BV₂ cells or C57BL/6j mice; # $P < 0.05$, ## $P < 0.01$ vs EMRD-BV₂ cells or APP/PS1 mice; n.s. means no significant differences.

anism in EMR-associated microglial phagocytosis¹². Initially, the ratios of p-Akt/Akt, p-mTOR/mTOR, and HIF-1 α / β -actin showed no significant differences between EMRD-BV₂ cells and BV₂ cells. However, these ratios significantly increased in EMRD-BV₂ cells following JWXG treatment (Figs. 4A–4D, $P < 0.05$). Additionally, TREM2 expression levels were significantly elevated in JWXG groups compared to EMRD-BV₂ cells and APP/PS1 mice (Figs. 4E–4J, $P < 0.05$ or $P < 0.01$). TREM2, functioning as a transmembrane promotor of microglial phagocytosis located in the microglial membrane, was knocked down in shTREM2-BV₂ cells using lentivirus, with verification through fluorescence and RT-qPCR (Fig. S1). Notably, TREM2 knockdown partially reversed JWXG's effects on microglial phagocytosis enhancement, glycolysis-mediated energy production improvement, and Akt/mTOR/HIF-1 α pathway activation in EMRD-BV₂ cells (Figs. 5A–5N, $P < 0.05$). These findings suggest that JWXG enhances microglial phagocytosis through activation of the TREM2-mediated Akt/

mTOR/HIF-1 α pathway and EMR enhancement.

3.5. JWXG attenuated A β burden and neuronal damage in APP/PS1 mice

Microglial phagocytosis serves a crucial function in eliminating toxic proteins, including A β , from the brain. The A β staining results revealed that A β ⁺ plaques in the cerebral cortex and hippocampus of APP/PS1 mice were significantly elevated compared to C57BL/6j mice but decreased in JWXG-treated mice relative to APP/PS1 mice (Figs. 6A–6C, $P < 0.01$). JWXG's effects on A β ₁₋₄₀ and A β ₁₋₄₂ levels in APP/PS1 mice brain tissues closely paralleled the A β staining findings (Figs. 6D and 6E, $P < 0.05$ or $P < 0.01$). Neuronal impairment represents a characteristic manifestation of A β neurotoxicity⁴⁹. The data demonstrated neuronal injury features in APP/PS1 mice cortex and hippocampus regions, including cortical thinning, neuronal atrophy, and neuron-

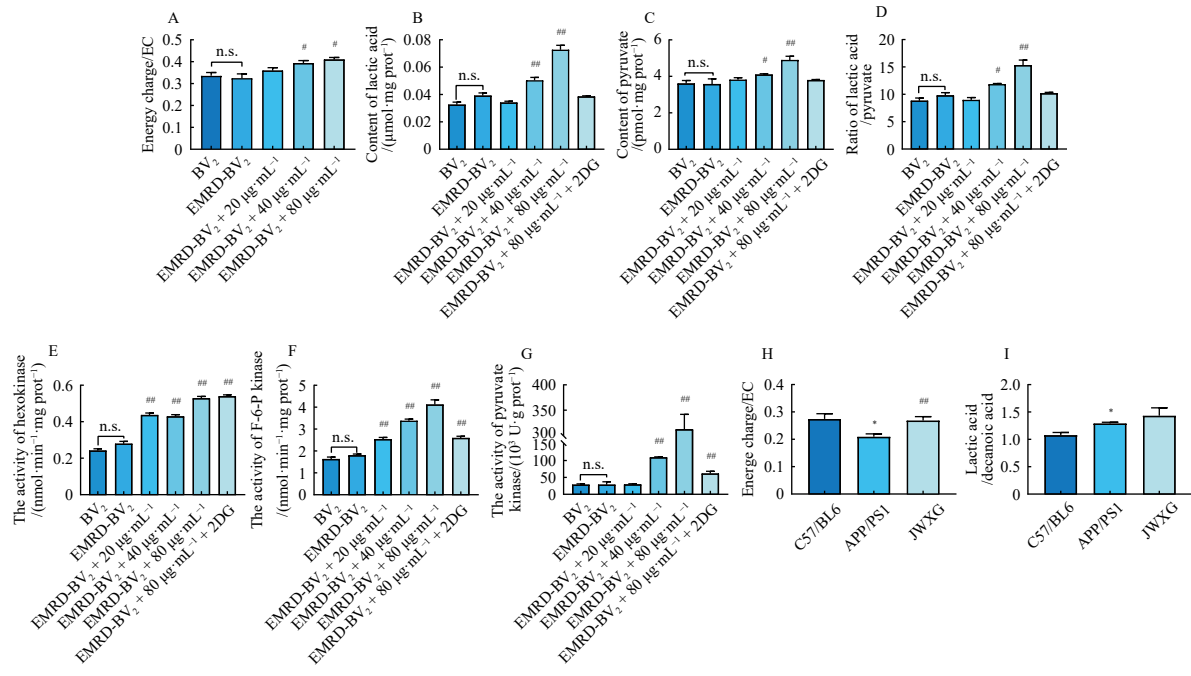


Fig. 3 JWYG increased energy production by enhancing EMR *in vitro* and *in vivo*. (A) The EC level in BV₂ cells (*n* = 6). (B, C) Levels of lactic acid and pyruvate in BV₂ cells (*n* = 6). (D) Lactic acid/pyruvate ratio in BV₂ cells (*n* = 6). (E–G) Activities of hexokinase, fructose-6-phosphate kinase and pyruvate kinase in BV₂ cells (*n* = 6). (H) EC levels of brain tissue in mice (*n* = 3). (I) Brain lactic acid levels in mice (*n* = 3). The data are expressed as mean ± SEM. **P* < 0.05 vs BV₂ cells or C57BL/6J mice; #*P* < 0.05, ##*P* < 0.01 vs EMRD-BV₂ cells or APP/PS1 mice; n.s. means no significant differences.

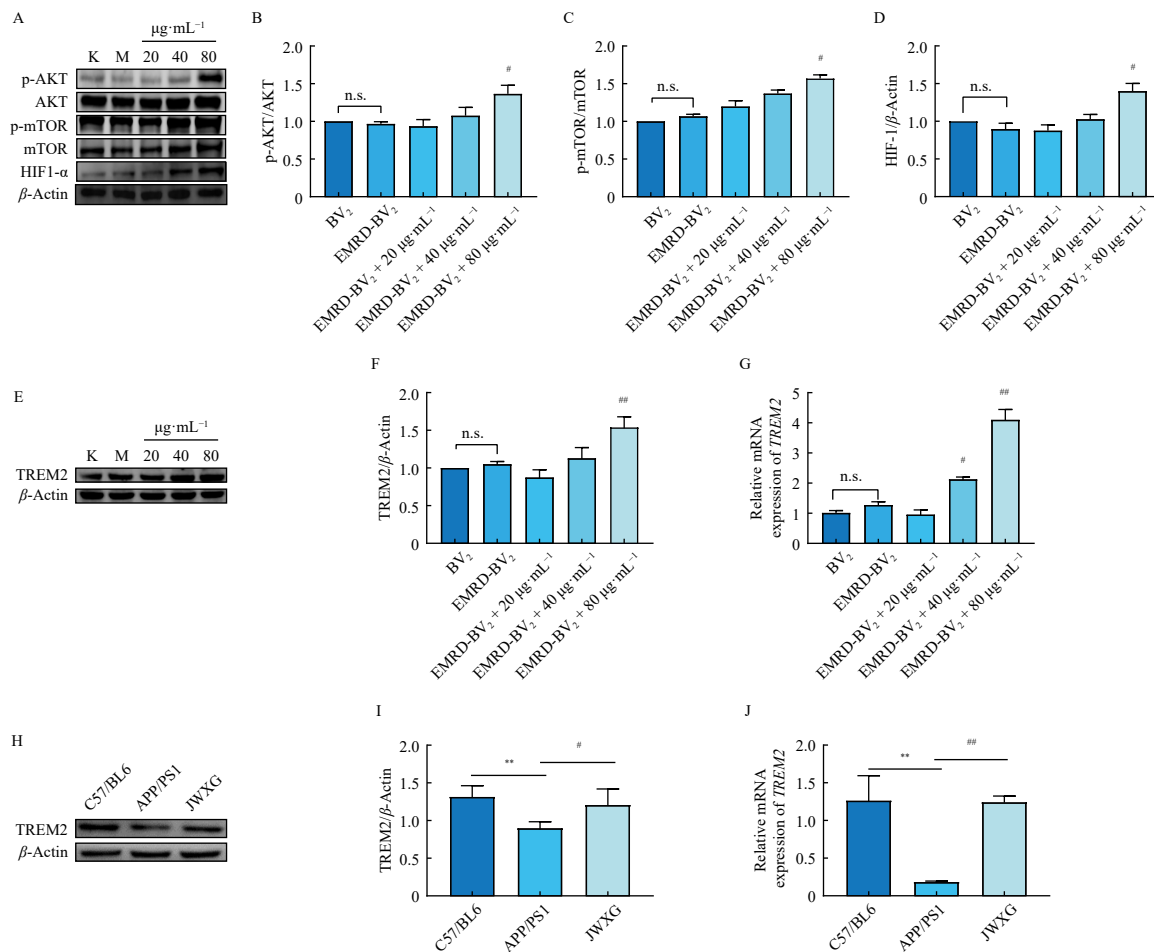


Fig. 4 JWYG activated TREM2-mediated Akt/mTOR/HIF-1α pathway. (A–D) Western blotting results of p-AKT/AKT, p-mTOR/mTOR, HIF-1α/β-actin *in vitro* (*n* = 6). (E, F) Western blotting results of TREM2 *in vitro* (*n* = 6). (G) TREM2 mRNA expression *in vitro* (*n* = 6). (H, I) Western blotting results of TREM2 *in vivo* (*n* = 3). (J) TREM2 mRNA expression *in vivo* (*n* = 3). The data are expressed as mean ± SEM. **P* < 0.01 vs BV₂ cells or C57BL/6J mice; #*P* < 0.05; ##*P* < 0.01 vs EMRD-BV₂ cells or APP/PS1 mice; n.s. means no significant differences.

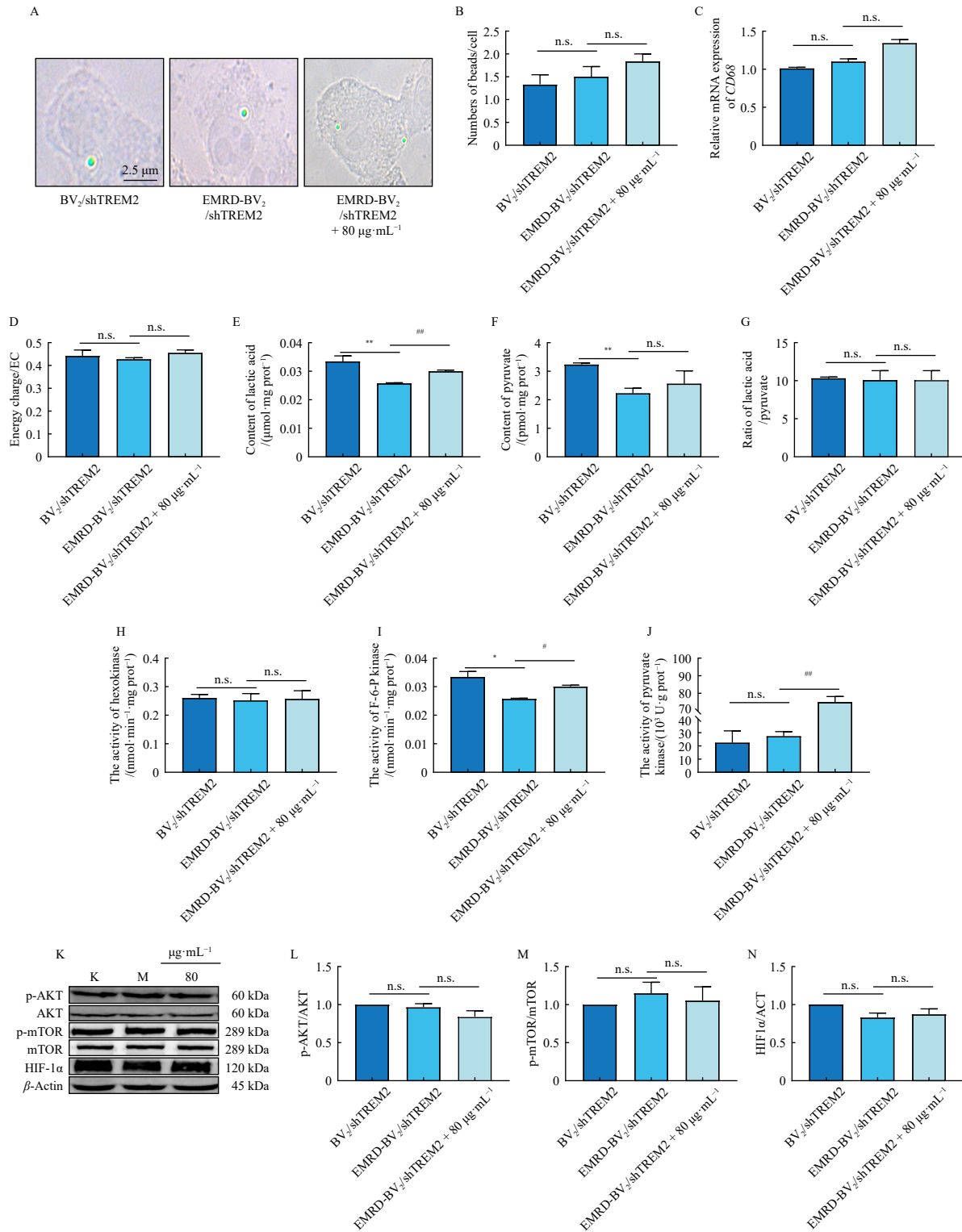


Fig. 5 shTREM2 partially reversed the effects of JWYG *in vitro*. (A) Representative images of the microglial phagocytosis of fluorescent latex beads *in vitro*. (B) Quantification of fluorescent latex beads per cell *in vitro*. (C) *CD68* mRNA levels *in vitro*. (D) EC levels *in vitro*. (E, F) Pyruvate and lactic acid levels *in vitro*. (G) Lactic acid/pyruvate ratio *in vitro*. (H–J). Activities of hexokinase, fructose-6-phosphate (F-6-P) kinase, and pyruvate kinase *in vitro*. (K–N) Representative images and quantitation of p-AKT/AKT, p-mTOR/mTOR, HIF-1 α / β -actin *in vitro*. The data are expressed as mean \pm SEM ($n = 6$). * $P < 0.05$, ** $P < 0.01$ vs BV₂/shTREM2 cells; # $P < 0.05$, ## $P < 0.01$ vs EMRD-BV₂/shTREM2 cells; n.s. means no significant differences.

al reduction and loss; however, JWYG treatment significantly mitigated these neuropathological changes (Fig. 7A). Additionally, JWYG administration significantly ameliorated the reduced NeuN⁺ staining intensity in the CA1 and CA3 regions of APP/PS1 mice (Figs. 7B–Fig. 7D, $P < 0.05$ or $P < 0.01$). These findings further suggest that JWYG reduces A β burden and its neurotoxicity *in vivo* partially through enhanced microglial phagocytosis. The

MWM test evaluated the mice's learning and memory capabilities. Escape latency, a critical MWM test indicator reflecting learning and memory ability, when prolonged, indicates impaired spatial learning and memory capacity⁵⁰. As illustrated in Figs. 8A and 8B, mouse escape latency decreased progressively with increased training time ([4, 20] = 2.038). JWYG significantly reduced mouse escape latency compared to APP/PS1 mice ([4, 20] = 3.723).

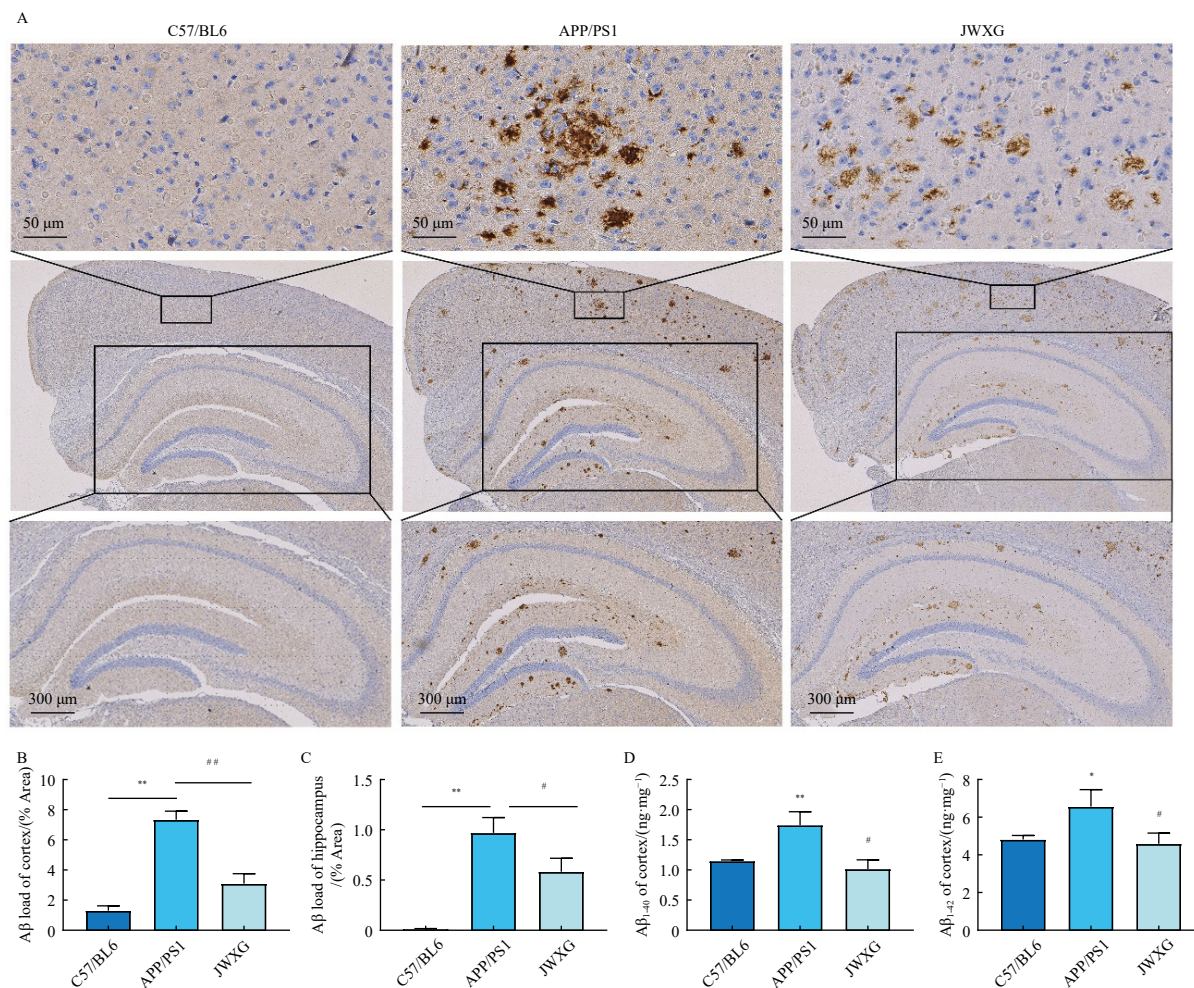


Fig. 6 JWXG reduced Aβ burden in APP/PS1 mice (n = 3). (A–C). Immunohistochemical staining and quantification of Aβ⁺ plaques in the cerebral cortex and hippocampus of mice by immunohistochemistry staining (magnification: 200 × or 33 ×). (D, E). Levels of Aβ₁₋₄₀ and Aβ₁₋₄₂ of brain tissues *in vivo*. The data are expressed as mean ± SEM. *P < 0.05, **P < 0.01 vs C57BL/6j mice; #P < 0.05, ##P < 0.01 vs APP/PS1 mice.

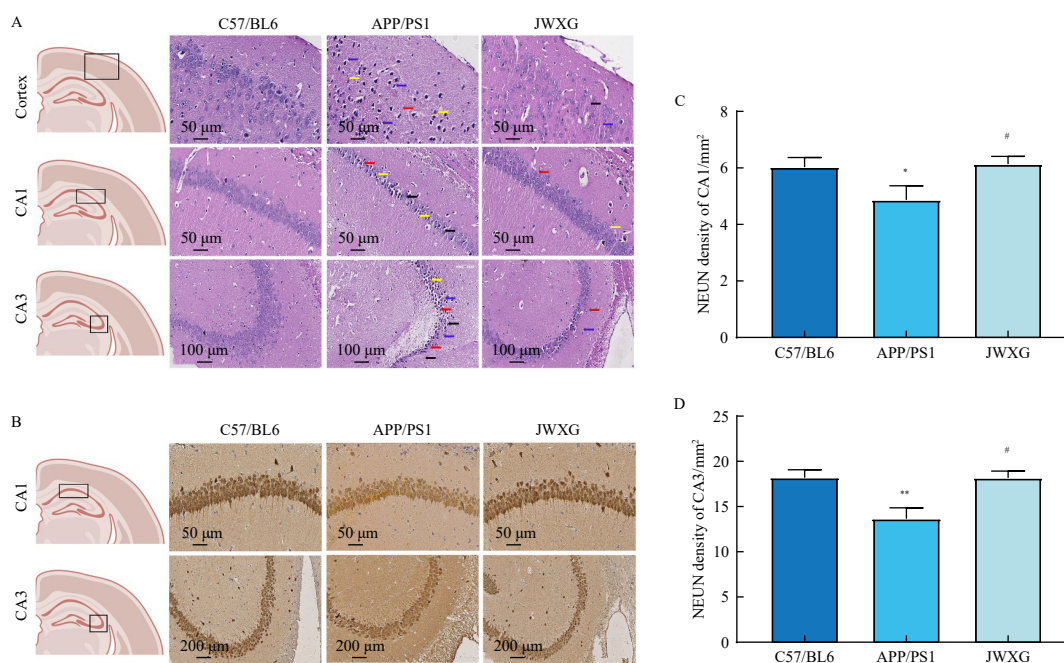


Fig. 7 JWXG attenuated the neuronal injury of APP/PS1 mice. (A) Representative images of H&E staining *in vivo* (magnification: 200 × or 100 ×); nerve cells scattered (red arrow), intercellular space enlarged (black arrow), nucleus staining deepened (yellow arrow), nucleus wrinkled (blue arrow). (B) Representative NeuN⁺ images of immunohistochemical staining (magnification: 200 × or 50 ×). (C, D) NeuN⁺ intensity of each group in CA1 and CA3 regions *in vivo*. The data are expressed as mean ± SEM (n = 3). *P < 0.05, **P < 0.01 vs C57BL/6j mice; #P < 0.05 vs APP/PS1 mice.

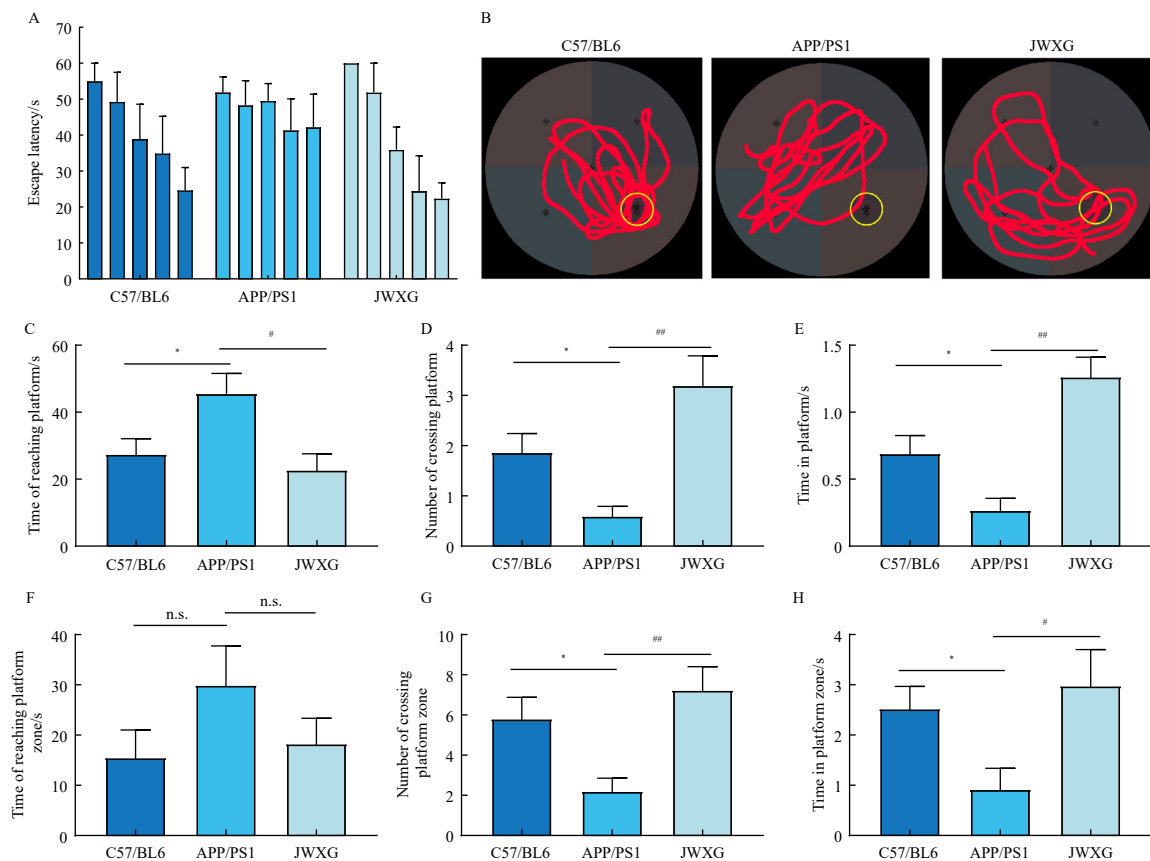


Fig. 8 JWYG ameliorated cognitive deficits in APP/PS1 mice. (A, B) Escape latency of mice and representative images of swimming tracks in the Morris water maze. (C) The time of reaching the platform. (D) Number of platform crossings. (E) Time spent on the platform. (F) Time to reach the platform zone. (G) Number of platform zone crossings. (H) The time spent in platform zone. The data are expressed as mean \pm SEM ($n = 3$). $P < 0.05$ vs C57BL/6j mice; $^{\#}P < 0.05$, $^{\#\#}P < 0.01$ vs APP/PS1 mice.

APP/PS1 mice exhibited significantly diminished spatial learning abilities compared to the C57BL/6j group, demonstrated by increased platform and platform zone reaching time, decreased platform and platform zone crossing frequency, and reduced platform and platform zone dwelling time. However, JWYG treatment significantly reversed these alterations in JWYG mice compared to APP/PS1 mice (Figs. 8C–8H, $P < 0.05$ or $P < 0.01$).

4. Discussion

This study demonstrates that JWYG mitigated cognitive dysfunction and neuronal impairment in AD through activation of the TREM2/Akt/mTOR/HIF-1 α pathway and enhancement of EMR to promote microglial phagocytosis. These findings suggest that enhancing EMR-mediated microglial phagocytosis may represent a promising therapeutic strategy for AD treatment.

A β accumulation and its associated neurotoxicity are considered primary contributors to AD pathogenesis, as they directly compromise cell membrane integrity and disrupt intracellular signaling pathways, leading to neuronal dysfunction or death^{51–53}. Clinical evidence has demonstrated that removal of aggregated A β from brain tissue benefits AD patients³, with FDA approval of two anti-amyloid antibodies, aducanumab and lecanemab, for AD treatment⁵⁴. These developments confirm that facilitating A β clearance represents an effective therapeutic approach. Microglial phagocytosis, autophagy-lysosome pathway, and meningeal lymphatic system serve as crucial mechanisms for A β clearance and maintenance of brain homeostasis^{55–57}. As central nervous system immune cells, microglia monitor the environment during injury response and maintain central homeostasis through phagocytosis and removal of toxic proteins, including A β ^{58–62}. During phagocytosis, microglia transform from a resting state to an amoeboid form, characterized by elevated *CD68* ex-

pression. Consequently, *CD68* mRNA serves as an established biomarker for microglial phagocytosis^{63,64}. Recent research emphasizes oA β for their synaptic toxicity rather than fibrotic amyloid deposits, leading to the use of oA β in EMRD-BV₂ cell studies^{12,65}. Our findings reveal significantly increased *CD68* mRNA expression and phagocytic fluorescent latex bead uptake in oA β acute stimulation-induced BV₂ cells, while no significant differences were observed between immune-tolerant BV₂ cells induced by oA β repeated stimulation (EMRD-BV₂) and unstimulated-BV₂ cells. Similarly, microglial A β phagocytosis showed marked deficits in 9-mo APP/PS1 mice. Notably, JWYG significantly enhanced *CD68* mRNA expression and microglial phagocytosis in both EMRD-BV₂ cells and 9-mo APP/PS1 mice, demonstrating its efficacy in promoting microglial phagocytosis *in vitro* and *in vivo*.

Microglia perform diverse functions that require varying levels of energy, with their metabolic pathways adapting accordingly¹². In a resting state, glucose metabolism primarily occurs through OXPHOS, but when stimulated by neurotoxic substances such as A β or cellular debris, the microglial energy supply shifts from OXPHOS to aerobic glycolysis⁶⁶. Although less efficient in ATP generation compared with mitochondrial respiration, glycolysis enables a faster rate of glucose metabolism, essential for energy-demanding processes like phagocytosis¹⁴. Research indicates that disruption in EMR leads to microglial dysfunction in AD, demonstrating a strong connection between EMR and phagocytosis¹². Recent studies have demonstrated that myeloid cell dysfunction can be reversed by reprogramming myeloid glucose metabolism to restore youthful immune functions and cognitive abilities^{67,68}. This evidence suggests that EMR is crucial for microglial phagocytosis. HK, F-6-PK, and PK serve as three rate-limiting enzymes of glycolysis, while lactic acid levels and the lactic acid/pyruvate ratio typically indicate glucose metabolism status^{69,70}. Our data revealed that JWYG treatment increased EC,

lactic acid content, and lactic acid/pyruvate ratio in EMRD-BV₂ cells, along with a dose-dependent increase in glycolysis rate-limiting enzyme activity. 2-DG, a glucose analog and HK substrate, inhibits glycolysis by reducing HK product rather than HK activity. As anticipated, 2-DG reversed JWXG's effects on glycolysis rate-limiting enzyme activity (except for HK). The enhanced glycolysis levels significantly correlated with increased CD68 expression and phagocytosis both *in vitro* and *in vivo*. These results indicate that JWXG enhances microglial phagocytic function through EMR promotion.

The Akt/mTOR/HIF-1 α pathway serves a critical role in inducing aerobic glycolysis and EMR^{15, 71, 72}. Multiple studies demonstrate that JWXG's active ingredients, including baicalin⁷³, baicalein⁷⁴, ferulic acid⁷⁵, and chlorogenic acid³³, can activate the Akt/mTOR pathway, suggesting JWXG may enhance EMR through increased Akt/mTOR/HIF-1 α pathway activity. Our findings revealed that JWXG significantly activated the Akt/mTOR/HIF-1 α pathway in EMRD-BV₂ cells and 9-mo APP/PS1 mice. These results suggest JWXG's enhancement of microglial phagocytosis closely relates to Akt/mTOR/HIF-1 α pathway activation. TREM2, a transmembrane glycoprotein specifically expressed in CNS microglia, has loss-of-function mutations widely considered a high-risk factor for late-onset AD^{76, 77}. Furthermore, TREM2 is essential for microglial phagocytosis and maintains cellular energetic and biosynthetic metabolism during AD^{9, 78}. TREM2 agonist antibodies are currently under investigation for AD treatment in preclinical trials⁷⁹. A β , a TREM2 ligand, directly binds to TREM2's extracellular domain, resulting in mTOR phosphorylation and HIF-1 α activation, thereby initiating EMR^{17-19, 80, 81}. Additionally, baicalin promotes TREM2 levels in microglia⁸². Our results demonstrated significantly decreased TREM2 gene and protein expression in APP/PS1 mice and EMRD-BV₂ cells, which JWXG treatment notably restored. Notably, shTREM2 blockade of TREM2 in EMRD-BV₂ cells partially reversed JWXG's effects on microglial phagocytosis, glycolysis-mediated energy production, and Akt/mTOR/HIF-1 α pathway activation. Finally, our data demonstrated that JWXG significantly reduced A β burden and neuronal impairment in the cortex and hippocampus while alleviating cognitive deficits in 9-mo APP/PS1 mice. These findings suggest JWXG attenuates cognitive deficits in APP/PS1 mice by activating TREM2-mediated Akt/mTOR/HIF-1 α pathway facilitated microglial phagocytosis.

Funding

This work was supported by the National Natural Science Foundation of China (Nos. 82074150 and 82274240) and the Natural Science Foundation of Sichuan Province (No. 2023NSFC1779).

Declaration of competing interest

These authors have no conflict of interest to declare.

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