

FcγRIIb Exacerbates LPS-Induced Neuroinflammation by Binding with the Bridging Protein DAP12 and Promoting the Activation of PI3K/AKT Signaling Pathway in Microglia

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Introduction: This paper focuses on the expression and role of FcγRIIb in neuroinflammation, exploring the molecular mechanisms by which FcγRIIb interacts with the bridging protein DAP12 to regulate the PI3K-AKT signaling pathway that promote neuroinflammation and aggravate neuronal injury.

Methods: LPS-induced neuroinflammation models in vivo and in vitro were constructed to explore the role and mechanism of FcγRIIb in CNS inflammation. Subsequently, FcγRIIb was knocked down or overexpressed to observe the activation of BV2 cell and the effect on PI3K-AKT pathway. Then the PI3K-AKT pathway was blocked to observe its effect on cell activation and FcγRIIb expression. We analyzed the interaction between FcγRIIb and DAP12 by Immunoprecipitation technique. Then FcγRIIb was overexpressed while knocking down DAP12 to observe its effect on PI3K-AKT pathway. Finally, BV2 cell culture supernatant was co-cultured with neuronal cell HT22 to observe its effect on neuronal apoptosis and cell activity.

Results: In vivo and in vitro, we found that FcγRIIb expression was significantly increased and activated the PI3K-AKT pathway. Contrary to the results of overexpression of FcγRIIb, knockdown of FcγRIIb resulted in a significant low level of relevant inflammatory factors and suppressed the PI3K-AKT pathway. Furthermore, LPS stimulation induced an interaction between FcγRIIb and DAP12. Knockdown of DAP12 suppressed inflammation and activation of the PI3K-AKT pathway in BV2 cells, and meantime overexpression of FcγRIIb suppressed the level of FcγRIIb-induced AKT phosphorylation. Additionally, knockdown of FcγRIIb inhibited microglia activation, which induced neuronal apoptosis.

Discussion: Altogether, our experiments indicate that FcγRIIb interacts with DAP12 to promote microglia activation by activating the PI3K-AKT pathway while leading to neuronal apoptosis and exacerbating brain tissue injury, which may provide a new target for the treatment of inflammatory diseases in the central nervous system.

Keywords: LPS-induced neuroinflammation, FcγRIIb, DAP12, microglia, PI3K-AKT

Introduction

Neuroinflammation is defined as an inflammatory response that occurs within the central nervous system (CNS). The progression of neuroinflammation is characterized by a complex cascade of cellular and biomolecular reactions, with brain triggering a nonspecific immune response upon stimulation.¹⁻³ Common abnormal features in the brains of patients with neuroinflammation include increased expression of pro-inflammatory factors, macrophage activation, peripheral leukocyte infiltration, and damage to neural tissue. Various glial cells are present in the human brain, among which, microglia play a key role in neuroinflammation. In the normal brain environment, microglia are branched and highly motile, and they serve to monitor fluctuations in the brain microenvironment, removing apoptotic neurons and

maintaining the stability of the central nervous system.^{4–6} Microglia are activated and polarized into the proinflammatory phenotype or the anti-inflammatory phenotype;^{7–9} the proinflammatory phenotype secrete large amounts of inflammatory and neurotoxic factors, which exacerbate neuronal damage, while the anti-inflammatory phenotype secrete large amounts of anti-inflammatory and trophic factors upon activation, alleviating inflammation and maintaining the stability of the brain internal environment.^{10,11}

FcγRIIb, a glycoprotein of approximately 40 kDa, consists of two IgG-like extracellular domains, a transmembrane region, and a cytoplasmic tail containing (immunoreceptor tyrosine-based inhibition motif), and it is the only IgG-inhibitory Fc receptor known to be found in both humans and mice. Many of the typical functions of FcγRIIb are based on signaling through the cytoplasmic region ITIM.^{12–15} FcγRIIb is a potent modulator in lupus nephritis, while in arthritis models, it effectively attenuates collagen-induced arthritis symptoms. Although ITIM-mediated inhibition through the cytoplasmic compartment is thought to be the main feature of FcγRIIb-mediated immune regulation, evidence also suggests that inhibition or activation of immune responses can occur independently of ITIM signaling in some specific cases. Following a review of the literature, we found reports indicating a 3.54-fold increase in FcγRIIb expression in RAW264.7 cells after lipopolysaccharide (LPS) treatment.^{16,17} Furthermore, there is evidence to suggest that activation of PI3K is required for FcγR to exert its function, and its activity can be monitored using p-Akt (Ser-473) antibody to determine the change in Akt activity.^{18,19} These reports have opened new ideas about what role FcγRIIb actually plays in a typical model of neuroinflammation. Based on the above logic, we conducted a preliminary exploration of the role of FcγRIIb in LPS-induced neuroinflammation.

DAP12, a member of the type I transmembrane bridging protein family, consists of a short extracellular structural domain, a single transmembrane structural domain, and a cytoplasmic region, (immunoreceptor tyrosine-based activation motif).^{20–22} DAP12 is mainly expressed in microglia in the CNS and macrophages in the periphery, and it does not possess ligand-binding capacity owing to its short extracellular structural domain, resulting in the need to form complexes with specific chaperone membrane receptors to transmit signals downstream.^{23–26} Several signaling pathways involved in DAP12, such as PI3K, PKC, and NF-κB, have been associated with pro-inflammatory responses, phagocytosis, and microglia migration and proliferation.^{27–30} Notably, the induction of pro-inflammatory cytokines and their related molecules by DAP12 in microglia may lead to a phenotypic shift to a toxic phenotype, resulting in neuronal damage.

PI3K-AKT is a highly conserved intracellular signaling pathway that is required for important activity processes in multiple classes of cells, including the synthesis of key proteins, cell proliferation and activation, neuronal activation, and morphological changes.^{31–34} This pathway can be activated by a variety of signals, including growth factors (GF), LPS, and insulin (INS), which produce phospholipids and contribute to protein phosphorylation.^{35–37} Dysregulation of the PI3K-AKT signaling pathway involves multiple pathological changes in multiple systems, such as neurological diseases. Experimental data suggest that activation of the PI3K-AKT signaling pathway in LPS-activated microglia leads to increased levels of proinflammatory factors. Conversely, other data suggest that activation of the PI3K-AKT signaling pathway helps to reduce brain damage and can inhibit inflammatory processes and apoptosis. Aberrant PI3K-AKT signaling is also involved in various diseases, including cancer, neurodegeneration, and neuroinflammatory diseases. In the brain, changes in AKT activity are critical for neuronal development and glial cell activity.^{38–41} In the present study, FcγRIIb promotes microglia inflammatory activation. FcγRIIb promotes microglia activation by activating the PI3K/AKT signaling pathway. FcγRIIb and DAP12 interact to promote microglia activation-mediated LPS-induced neuroinflammation. The “FcγRIIb-PI3K-AKT” positive feedback loop exacerbates neuroinflammation-mediated neuronal injury.

Materials and Methods

Construction of Animal Models

Animal experiments were performed in accordance with the National Institutes of Health Guide for the Care and Use of Laboratory Animals (National Research Council, 1996, USA) and approved by Jiangsu Branch of the National Committee for the Use of Laboratory Animals for Medical Purposes, China. C57BL/6 whole male mice (6–8 weeks, 25–30 g) were used for experiments. The mice were kept in the experimental animal center for 7 days before the induction of the LPS-induced neuroinflammation model to acclimatize and reduce stress. After the mice were stabilized, LPS (*Escherichia coli* O111-B4; Sigma, St. Louis, MO, USA) was injected intraperitoneally into the mice for 6 h, 12 h, 1

d, 3 d, 5 d, and 7 d, while the control mice were injected with an sterile saline (n = 7 groups). After the appropriate length of stimulation, the mice were injected with anesthetic (lidocaine hydrochloride) and then sacrificed, the eyeballs were removed to obtain blood, and brain tissue was removed. The brain tissue samples were stored at -80°C and used for subsequent experiments. During the animal experiments, the pain felt by the mice was minimized as much as possible to ensure the objectivity of the experiments.

Western Blotting

After protein extraction, the protein concentration was determined by Bio-Rad protein assay (BioRad, Hercules, CA, USA), the samples were subjected to sodium polyacrylamide gel electrophoresis (SDS-PAGE), and then the proteins were transferred to polyvinyl fluoride (PVDF) membranes. Closure was performed using 5% skim milk. The membranes were incubated with anti-Fc γ RIIb (anti-rabbit, 1:1000; Signalway Antibody), anti-Fc γ RIIb (anti-rat, 1:500; RnDSystems), anti-DAP12 (anti-rabbit, 1:1000; Abcam), iNOS (anti-rabbit, 1:1000; Abcam), GAPDH (anti-mouse, 1:5000; Proteintech), p-AKT (Ser473) (anti-rabbit, 1:1000; Cell Signaling Technology), AKT (anti-rabbit, 1:1000; Cell Signaling Technology), PI3K (anti-mouse, 1:500; Santa Cruz Biotechnology), and Bax (anti-rabbit, 1:1000; Cell Signaling Technology) overnight at 4°C , followed by incubation with the corresponding secondary antibodies for 2 h at room temperature. Finally, the protein bands were visualized by an enhanced chemiluminescence system (ECL, Pierce Company, Appleton, WI, USA).

Plasmid Transfection

The small interfering and negative control of Fc γ RIIb were purchased from Reebok Biologicals, and the overexpression plasmid and negative control of Fc γ RIIb were purchased from PPL Genetics: si-DAP12#196, 5'-GGGAUUGUUCUGGGUGAVUTT-3'; si-DAP12#104, 5'-UCCUGACUGUGGGAGGAUUTT-3'; si-DAP12#288, 5'-GGAAGGACCCGGAAACAATT-3'. Myc-RNA was transfected into BV2 cells using Lip2000 transfection reagent (Invitrogen Life Technologies, Carlsbad, CA, USA).

Mouse Brain Tissue Processing and Sectioning

The mouse cerebral cortex was removed, fixed in 4% paraformaldehyde, and then treated with 10%, 20%, and 30% sucrose solution for gradient dehydration. The tissues were subsequently sliced with a freezing microtome at a thickness of 6 μm .

Immunohistochemical Fluorescence

The tissues were pre-incubated with 1% bovine serum albumin (1% BSA) for 2 h at room temperature to block non-specific binding of the primary antibody to the tissue membrane, followed by F4/80 (anti-rat, 1:100, Abcam) and a mixture of anti-Fc γ RIIb (anti-rabbit, 1:100; Signalway Antibody) incubated overnight at 4°C in the refrigerator. Subsequently, the tissues were incubated with corresponding secondary antibody for 2 h at room temperature protected from light, shaken dry, and incubated for 30 min in Hoechst dye. A blocking solution (glycerol: ddH $_2$ O= 1:1) was used to block the slices before observing them under a fluorescence microscope (Leica, DM 5000B, Leica CTR 5000, Germany).

Immunocytofluorescence

BV2 cells were spread on 24-well plates and then stimulated with LPS (100 ng/mL LPS, Escherichia coli 055:B5, Sigma Aldrich, L6529). Next, 4% paraformaldehyde was used to fix the cells for 1 h at room temperature. Subsequently, to enhance the permeability of the cell membrane, 0.1% TritonX-100 was added to the 24-well plate containing BV2 cells at 25°C for 15 min before blocking non-specific binding by incubating the cells with 1% bovine serum albumin for 2 h at room temperature. After the blocking solution (1% bovine serum albumin) was discarded, a mixture of Fc γ RIIb and Iba-1 anti-Fc γ RIIb (anti-rat, 1:500; RnDSystems) was added to the 24-well plate and refrigerated overnight at 4°C . Next, the nuclei were incubated with Alexa 488 anti-rabbit IgG (H + L) (Thermo Fisher Scientific) for 2 h, before adding Hoechst dye (Thermo Fisher Scientific) to stain the nuclei. A drop of glycerol was added to the slide, which was then covered with a coverslip, and fluorescence microscopy (Leica, DM 5000B, Leica CTR 5000, Germany) was used for observation.

RNA Extraction and RT-PCR

Total RNA was extracted from the mouse cerebral cortex and LPS-stimulated BV2 cells using a TRIzol kit, while cDNA was synthesized by reverse transcription-polymerase chain reaction using a RevertAid™ RT kit (Thermo Fisher Scientific). The amplified products were subjected to quantitative analysis by q-PCR. This procedure was performed with the following primers (Sunny Shanghai) for DNA sequences: GAPDH Forward primer (5'-3') AGGTCCGGTGTGAACGGATTTG Reverse primer (5'-3') TGTAGACCATGTAGTTGAGGTCA FcγRIIb Forward primer (5'-3') ATGGGAATCCTGCCGTTCCCTA Reverse primer (5'-3') CCGTGAGAACACATGGACAGT TYROBP Forward primer (5'-3') GGTGCCTTCTGTTCCTTCTGTC Reverse primer (5'-3') ATCAGCAGAGTCAACACCAAGTCAC TNFα Forward primer (5'-3') GACGTGGAAGTGGCAGAAGAG Reverse primer (5'-3') TTGGTGGTTTGTGAGTGTGAG IL-6 Forward primer (5'-3') TAGTCCTTCTACCCCAATTTCC Reverse primer (5'-3') TTGGTCCTTAGCCACTCCTTC IL-1β Forward primer (5'-3') GAAATGCCACCTTTTGACAGTG Reverse primer (5'-3') TGGATGCTCTCATCAGGACAG

Enzyme-Linked Immunosorbent Assay (ELISA)

The expression levels of TNF-α, IL-6, and IL-1β in the collected mouse sera and cell supernatants were measured using an ELISA kit (Proteintech or MULTI SCIENCES). The optical density (OD) of each well was measured at 450 nm using an enzyme marker with 630 nm as the calibration wavelength. The standard curve was generated by four-parameter fitting using ELISACalc curve making software and the corresponding concentrations were deduced from the standard curve according to the OD values of the samples. The entire operation was performed according to the manufacturer's instructions.

Cell Culture, Cell Stimulation, and Neuronal Injury Model Construction

BV2 cells (Purchased from Shanghai Zhong Qiao Xin Zhou Biotechnology Co., Ltd; Item No. ZQ0397) and HT22 neurons (Purchased from Shanghai Cell Bank of Chinese Academy of Sciences, respectively) were cultured in Dulbecco's Modified Eagle's Medium (DMEM) (Gibco 8121170) with 10% fetal bovine serum (FBS) and 1% penicillin-streptomycin at 37°C and under 5% CO₂. LPS stimulation was used to induce BV2 cells activation (100 ng/mL LPS, *Escherichia coli* 055:B5, Sigma Aldrich, L6529). Construction of the neuronal injury model: The supernatant of activated BV2 cell culture (MG-CM) was collected, mixed with DMEM complete medium in a 1:1 ratio, and then added to HT22 neurons to generate the neuronal injury model.

Cell Counting Kit 8 Assay (CCK-8)

Briefly, 100 μL of cell suspension was added to each well of a 96-well plate and cultured for 24 h after LPS stimulation for 24 h. Next, 10 μL of CCK-8 reagent (MCE, China) was added to each well and incubated for 2 h protected from light. Finally, the absorbance at 450 nm was measured with an enzyme marker. The data were analyzed according to the manufacturer's instructions.

Immunoprecipitation (IP)

Briefly, the cells were washed three times with phosphate-buffered saline (PBS), before adding 1 mL of PBS to break the cells. Next, the precipitate was centrifuged, before adding 1 mL of RIPA lysate and rotating at 4°C for 1.5 h. The supernatant was centrifuged and 50 μL was aspirated as the input group. The samples were stored at -20°C until required. After that, 25–30 μL of magnetic beads (Protein A/G PLUS-Agarose sc-2003, immunoprecipitation reagent; Santa Cruz Biotechnology) was added to the supernatant to adsorb impurities, and rotated at 4°C for 2–3 h. The samples were centrifuged to settle the beads, before adding FcγRIIb (anti-rat, 4 μL; RnDSystems) antibody to the supernatant and rotating overnight. The next day, 30 μL of magnetic beads was added and continued to rotate for 3 h. Subsequently, the precipitate was centrifuged before adding 500 μL of RIPA lysate to gently explode the beads, before centrifuging and repeating the above operation once more. Finally, 2x loading buffer (Beyotime Biotechnology) was added to the precipitate before immunoblotting.

Statistical Analysis

SPSS 19.0 software was used to calculate the data related to the experiment in this thesis, and the data were expressed using the mean \pm standard deviation. The analysis of differences between two groups of data was done by *t*-test, and all three and more groups of data were subjected to One-way ANOVA analysis was performed on all three and more groups. Statistical plots were plotted by GraphPad Prism. $p < 0.05$ indicates a statistical difference. $p < 0.05$ indicates a statistical difference.

Results

Upregulation of Fc γ RIIb Expression is Involved in Microglia Activation in a Mouse Model of CNS Inflammation

To investigate the role of Fc γ RIIb in LPS-induced neuroinflammation, a mouse CNS inflammation model was constructed by intraperitoneal injection of LPS (9 mg/kg) into C57BL/6 experimental mice. The ELISA results showed that the expression of pro-inflammatory factors TNF- α , IL-6, and IL-1 β in mouse serum was increased sharply after LPS injection and peaked at 6 hours (Figure S1). Additionally, the results of q-PCR showed that the inflammatory factors (TNF- α , IL-6, IL-1 β) in the mouse cerebral cortex peaked at 6 hours, while Fc γ RIIb expression peaked at 12 hours after LPS injection (Figure 1A). The protein expression level of iNOS started to increase at 6 hours and peaked at 3 days, while Fc γ RIIb expression started to increase after 6 hours and peaked at 12 hours (Figure 1B). The immunofluorescence results showed that Fc γ RIIb in the LPS-stimulated group was significantly co-localized with the microglia marker F4/80, while the expression of Fc γ RIIb was significantly increased in the mouse cerebral cortex in the LPS-stimulated group compared to the control group (Figure 1C). The above results suggest that the mouse CNS inflammation model was successfully constructed and that Fc γ RIIb was involved in microglia activation and neuroinflammatory response.

Upregulation of Fc γ RIIb Expression in an in vitro Model of LPS-Induced Neuroinflammation

To further explore the mechanism by which Fc γ RIIb mediates LPS-induced neuroinflammation during the activation of BV2 cells, we stimulated BV2 cells with LPS (100 ng/mL) at different time points to construct an in vitro activation model. The ELISA results showed that the expression levels of TNF- α , IL-6, and IL-1 β were significantly increased in BV2 cells treated with LPS (Figure S2). Additionally, the q-PCR results showed that the expression levels of TNF- α , IL-6, IL-1 β and Fc γ RIIb were significantly increased in BV2 cells treated with LPS (Figure 2A). The results of Western blot showed that the expression of iNOS increased and reached a peak at 12 hours, while the expression of Fc γ RIIb increased after 3 hours and reached a peak at 12 hours (Figure 2B). To further investigate the significance of Fc γ RIIb expression in BV2 cells, we observed the localization of Fc γ RIIb in inflammation-activated BV2 cells. The results of the immunohistochemical fluorescence assay showed that Fc γ RIIb in the LPS-stimulated group was significantly co-localized with the BV2 cell marker F4/80, while the expression of Fc γ RIIb in BV2 cells was found to be significantly increased in the LPS-stimulated group compared to the N group (Figure 2C). The above experimental results indicated that the in vitro activation model of BV2 cells was successfully constructed and that Fc γ RIIb was involved in the process of microglial inflammatory activation.

Fc γ RIIb Promotes Inflammatory Activation of Microglia

The above results showed that Fc γ RIIb expression showed an upregulation trend in inflammation-activated BV2 cells. To confirm the mechanism of action of Fc γ RIIb in the activation of BV2 cells, we knocked down Fc γ RIIb in BV2 cells and observed whether there was an effect on the inflammatory activation of BV2 cells. We transfected plasmids expressing an siRNA targeting Fc γ RIIb into BV2 cells. The Western blot results showed that expression of Fc γ RIIb and iNOS was reduced in BV2 cells (Figure 3A). The q-PCR results also showed that the expression of Fc γ RIIb and inflammatory factors (TNF- α , IL-6, and IL-1 β) was decreased in siRNA-treated BV2 cells (Figure 3B and D). Meanwhile, the results of ELISA showed a downregulated expression trend for inflammatory factors (TNF- α and IL-6) in the cell culture supernatants of siRNA-treated BV2 cells (Figure 3C). The q-PCR assay results were consistent with the ELISA results, in that

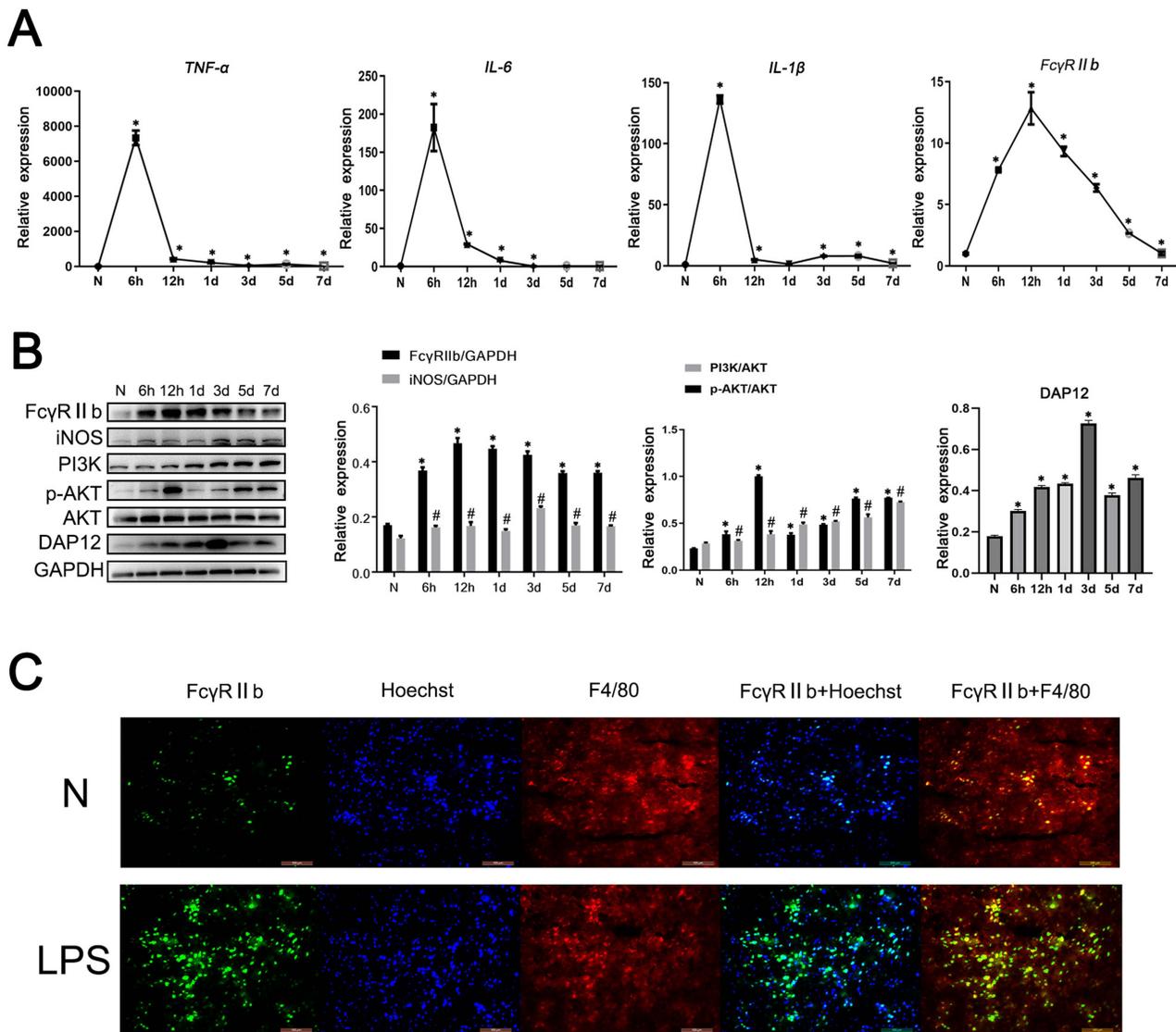


Figure 1 The expression and role of FcγRIIb in a mouse model of CNS inflammation. C57BL/6 mice were injected intraperitoneally with LPS (9 mg/kg), and serum was collected at 6 h, 12 h, 1 d, 3 d, 5 d, and 7 d, while the control group (N) was injected with an equal amount of saline. **(A)** Q-PCR was performed to detect the mRNA expression levels of IL-6, TNF-α, IL-1β, and FcγRIIb in the mouse cerebral cortex. **(B)** Western blot was performed to detect the protein expression levels of FcγRIIb, iNOS, PI3K, p-AKT, AKT and DAPI12 in the cerebral cortex of experimental mice. **(C)** Immunofluorescence staining was used to detect the expression of FcγRIIb in the cerebral cortex tissue of control and LPS mice. FcγRIIb (green), F4/80 (red), Hoechst (blue); scale bar: 100 μm (mean ± SEM represents the resultant values, n = 3, *Indicates compared to the control, #indicates compared to the control, p < 0.05).

expression knocking-down of FcγRIIb in BV2 cells significantly inhibited the expression of inflammatory factors in LPS stimulated group, and the expression levels of TNF-α, IL-6, and IL-1β showed a decreasing trend.

Meanwhile, we transfected FcγRIIb overexpressing plasmid into BV2 cells and then stimulated the cells with LPS (100 ng/mL) for 12 hours. Western blot detection showed that in LPS-stimulated group, the expression levels of iNOS and FcγRIIb were significantly increased in the overexpression group compared to the no-load group (Figure 3E). Moreover, q-PCR results demonstrated that the overexpression group showed increased expression of FcγRIIb, TNF-α, IL-6, and IL-1β in BV2 cells (Figure 3F and H), while ELISA results showed that the expression levels of TNF-α, and IL-6 were significantly upregulated in LPS-stimulated BV2 cells (Figure 3G). The above experimental results suggest that the changes in FcγRIIb expression are positively correlated with microglia activation and that FcγRIIb can promote the inflammatory activation of microglia, thus mediating LPS-induced neuroinflammation.

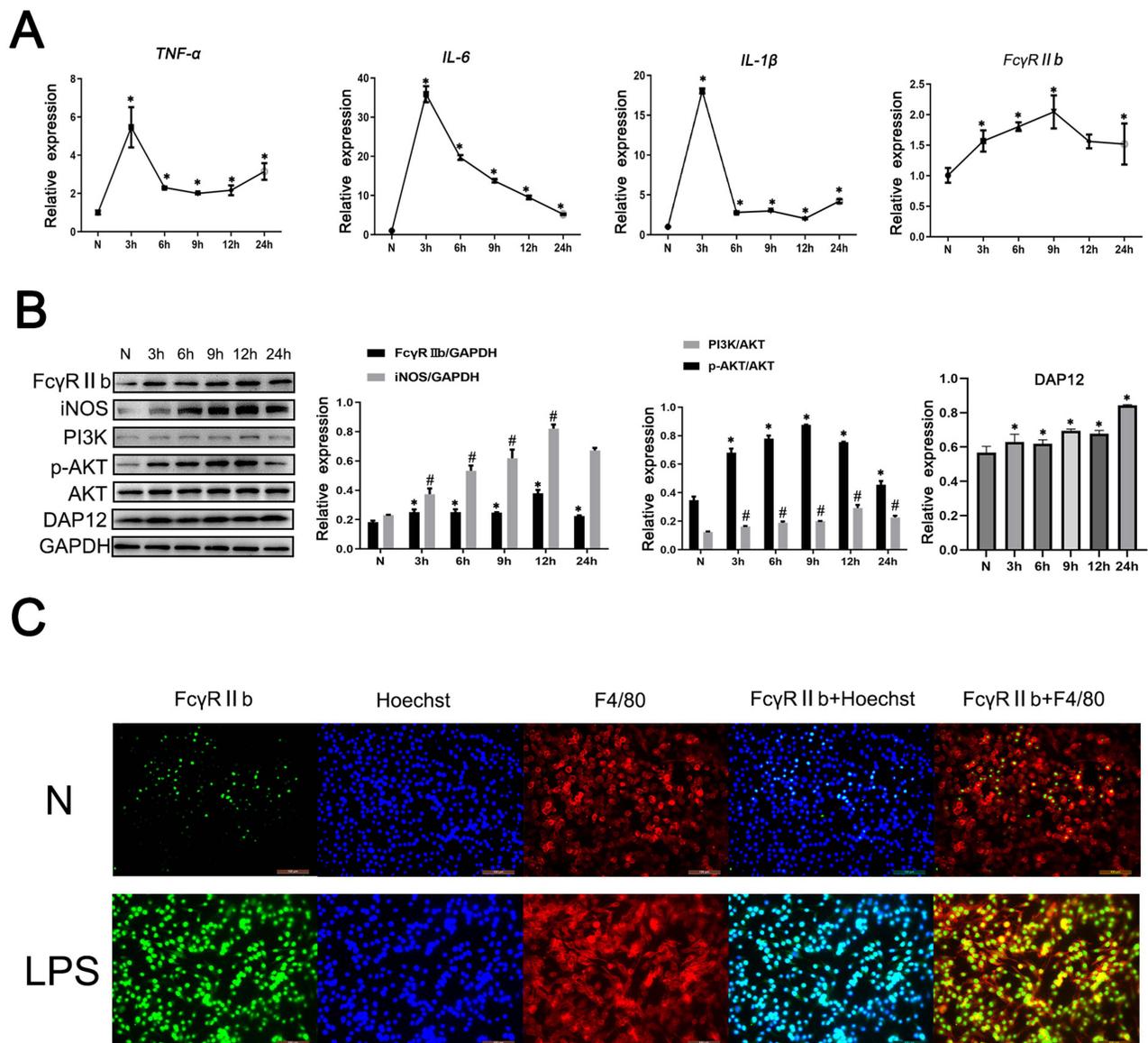


Figure 2 The expression and role of Fc γ RIIb in an in vitro model of LPS-induced neuroinflammation. BV2 cells were stimulated with LPS (100 ng/mL) for 0, 3, 6, 9, 12, and 24 h, and then the cells and the cell culture supernatants were collected. **(A)** Q-PCR was performed to detect the mRNA expression levels of TNF- α , IL-6, IL-1 β , and Fc γ RIIb in BV2 cells. **(B)** Western blot was used to detect the protein expression of Fc γ RIIb, iNOS, PI3K, p-AKT, AKT and DAP12 in BV2 cells. **(C)** Immunofluorescence staining was employed to detect the expression and localization of Fc γ RIIb in the N group and LPS-stimulated group of BV2 cells. Fc γ RIIb (green), F4/80 (red), Hoechst (blue); scale bar: 100 μ m (mean \pm SEM represents the resultant values, n = 3, *Indicates compared to the control, #indicates compared to the control, p < 0.05).

PI3K-AKT Pathway is Involved in the Development of LPS-Induced Neuroinflammation Mediated by Activated Microglia

Many recent studies have found that the PI3K-AKT signaling pathway is involved in various diseases, including neurodegenerative and neuroinflammatory diseases. In BV2 cells treated with LPS, the level of p-AKT in the cerebral cortex of CNS inflammation model mice gradually increased after LPS intraperitoneal injection, reaching the highest level at 12 h, while PI3K expression reached the highest level at 7d (Figure 1C). The results of the cellular immunofluorescence staining assay showed that the expression of p-AKT was increased in the stimulated group compared to the control group (Figure 4A), which is consistent with the results of the Western blot assay. Meanwhile, in vitro stimulation of BV2 cells with LPS for 0, 3, 6, 9, 12, and 24 h showed that the expression of p-AKT increased in BV2 cells and reached the highest level at 9 h, while the expression of PI3K reached the highest level at 12 h (Figure 2C). The results of

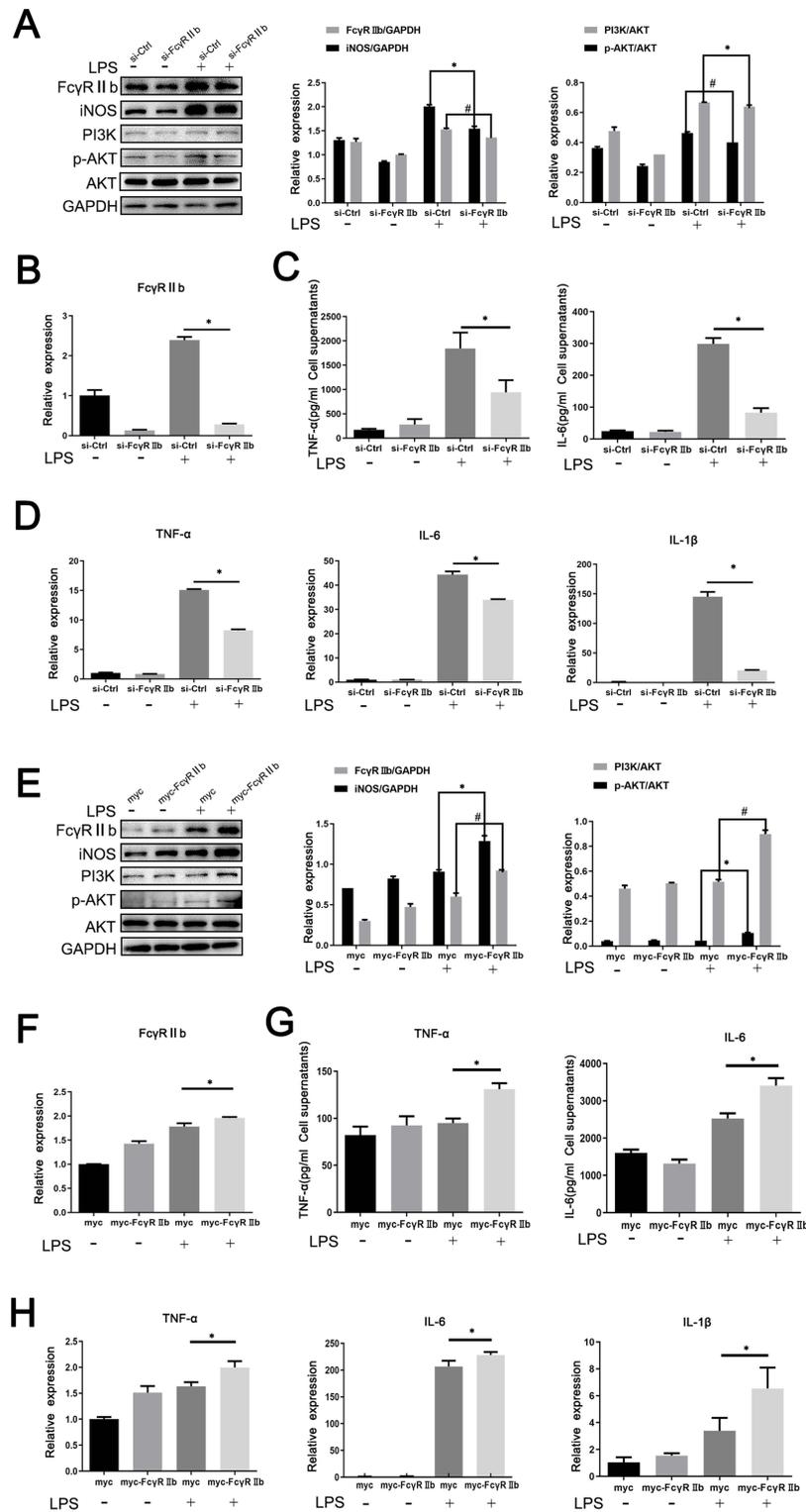


Figure 3 Effect of inhibition or overexpression of FcγRIIb on inflammatory activation in microglia. FcγRIIb interference knocking-down or overexpressing plasmids were transfected into BV2 cells for 48 h and then stimulated with LPS for 12 h. **(A)** Western blot detection of FcγRIIb, iNOS, PI3K, p-AKT and AKT protein expression levels after interference with BV2 cells. **(B)** Q-PCR assay of the mRNA expression levels of the inflammatory factors FcγRIIb in BV2 cells after interference with FcγRIIb. **(C)** ELISA the assay of the expression levels of TNF-α and IL-6 in the cell supernatants collected after interference with BV2 cells. **(D)** Q-PCR assay of the mRNA expression levels of the inflammatory factors TNF-α, IL-6, and IL-1β in BV2 cells after interference with FcγRIIb. **(E)** Western blot assay of the protein expression levels of FcγRIIb, iNOS, PI3K, p-AKT and AKT after overexpression with BV2 cells. **(F)** Q-PCR assay of the mRNA expression levels of FcγRIIb after overexpression with BV2 cells. **(G)** ELISA assay of the expression levels of inflammatory factors TNF-α and IL-6 in BV2 cells. **(H)** Q-PCR assay of the mRNA expression levels of TNF-α, IL-6, and IL-1β after overexpression with BV2 cells. (mean ± SEM represents the result values, n = 3, *Indicates compared to the si-Ctrl group, or myc+LPS group, #Indicates compared to the si-Ctrl+LPS group, or myc+LPS group, p < 0.05).

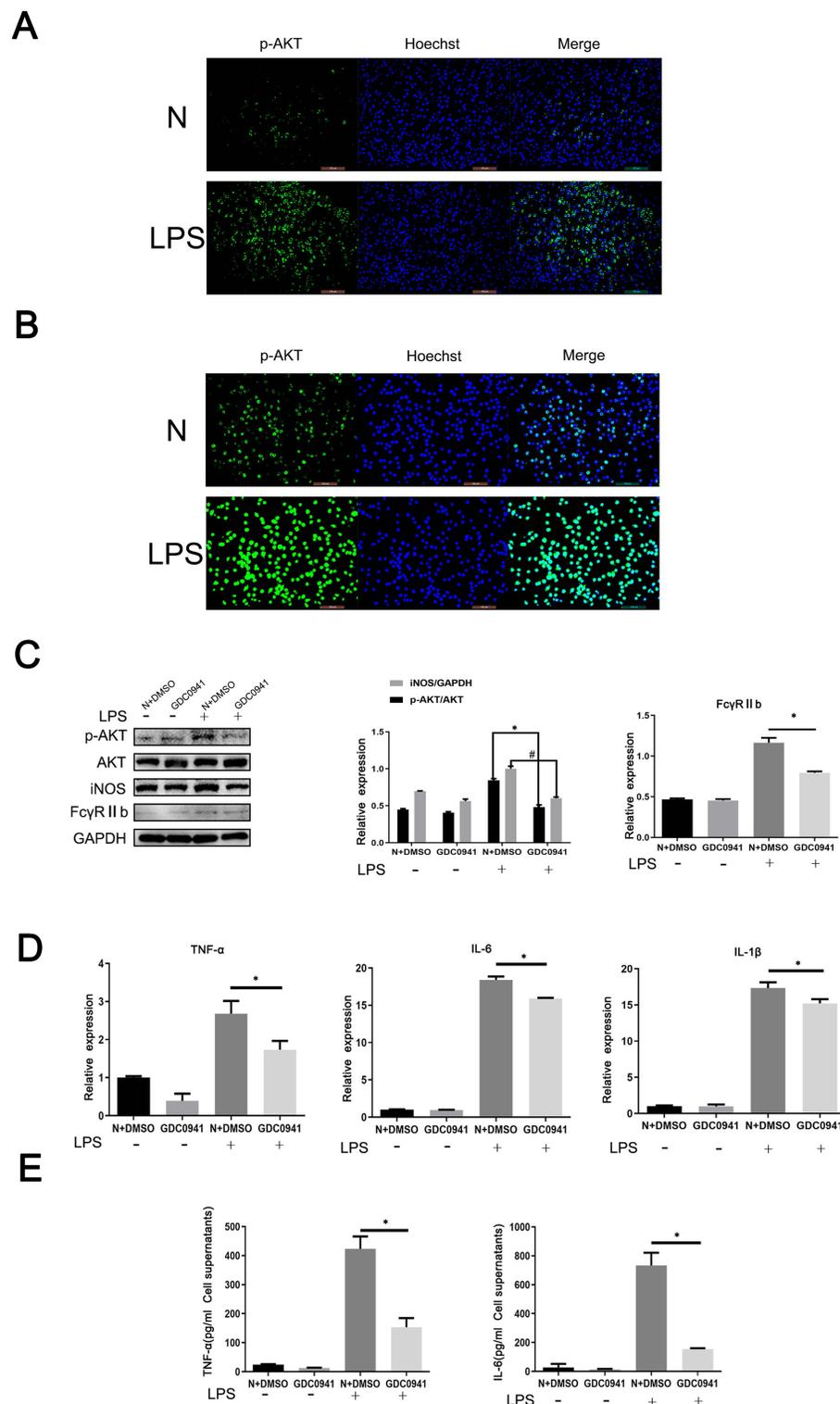


Figure 4 PI3K/AKT signaling pathway is involved in microglia-mediated LPS-induced neuroinflammation. **(A)** Cellular immunofluorescence staining of the expression and localization of p-AKT in control and LPS-stimulated groups in a mouse CNS inflammation model. p-AKT (green), Iba-1 (red), and Hoechst (blue); scale bar: 100 μ m. **(B)** Cellular immunofluorescence staining of the expression and localization of p-AKT in control and LPS-stimulated groups in an in vitro model. p-AKT (green), Iba-1 (red), and Hoechst (blue); scale bar: 100 μ m. **(C)** BV2 cells were pretreated with PI3K-AKT inhibitor GDC0941 for 1 h and then stimulated with LPS (100 ng/mL) for 9 h. Western blotting was performed to detect the protein expression levels of p-AKT, AKT, iNOS and Fc γ RIIb in BV2 cells. **(D)** Q-PCR detection of the mRNA expression levels of TNF- α , IL-6 and IL-1 β in BV2 cells. **(E)** ELISA was performed to detect changes in expression levels of TNF- α and IL-6 in the supernatants of treated BV2 cells. (mean \pm SEM represents the resultant values, n = 3, *Indicates compared to the control, or N+DMSO+LPS group, #Indicates compared to the control, or N+DMSO+LPS group, p < 0.05).

cell immunofluorescence staining also verified that the p-AKT expression in the stimulated group was much higher than that in the control group (Figure 4B). The above experimental data can be used to tentatively verify that the PI3K-AKT signaling pathway is involved in the inflammatory activation of microglia. To further validate the role of the PI3K-AKT pathway in microglia inflammatory activation, we blocked the PI3K-AKT pathway using the pathway blocker GDC0941, the experimental results showed that the expression of iNOS and p-AKT was significantly decreased in BV2 cells treated with LPS (Figure 4C). The results of ELISA and q-PCR assays were consistent with these findings, in that the expression of pro-inflammatory factors was relatively reduced after inhibition of PI3K-AKT pathway activity (Figure 4D and E). The above experimental results confirm that PI3K-AKT pathway activation is involved in the development of LPS-induced neuroinflammation mediated by activated microglia.

FcγRIIb Promotes Microglia Activation Through PI3K-AKT Pathway Activation

The combined experimental data show that the change in FcγRIIb expression was closely associated with PI3K-AKT pathway activation during the inflammatory activation of BV2 cells. Previous works in the literature have reported that FcγRIIb can contribute to disease onset and progression through the activation of the PI3K-AKT signaling pathway during immune regulation. To investigate the mechanism by which FcγRIIb promotes inflammatory activation in BV2 cells, we transfected si-FcγRIIb and si-Ctrl into BV2 cells and observed changes in PI3K-AKT signaling pathway activity in BV2 cells in the LPS-stimulated group after interference with FcγRIIb expression. The results showed that the protein-level of p-AKT and PI3K were significantly reduced after knockdown of FcγRIIb in BV2 cells (Figure 3A). Conversely, the protein expression of p-AKT and PI3K was significantly increased by transfection of the overexpression plasmid (myc-FcγRIIb) compared to the control group (Figure 3E). These results suggest that FcγRIIb mediates microglia inflammatory activation through the positive regulation of PI3K-AKT pathway activity.

Meanwhile, we pretreated BV2 cells with a GDC0941 pathway inhibitor, before stimulating with LPS for 9 h to analyze the effect of blocking the PI3K-AKT pathway on FcγRIIb protein expression in activated BV2 cells. The results revealed that FcγRIIb expression was significantly downregulated in BV2 cells of the LPS-stimulated group after inhibition of the PI3K-AKT signaling pathway (Figure 4C). Therefore, we conclude that PI3K-AKT pathway activity has a positive regulatory effect on the expression of FcγRIIb in inflammation-activated BV2 cells, which can form a positive feedback loop in BV2 cells to mediate the development of LPS-induced neuroinflammation.

FcγRIIb and DAP12 Interact to Promote Microglia Activation-Mediated LPS-Induced Neuroinflammation

The above experimental results suggest that FcγRIIb plays a role in LPS-induced neuroinflammation through the PI3K-AKT signaling pathway. We first confirmed the expression of DAP12 in the cerebral cortex of CNS inflammation model mice and in LPS-activated BV2 cells by Western blot. The results showed that DAP12 was significantly increased in the cerebral cortex of LPS intraperitoneally injected mice. Moreover, the expression of DAP12 was significantly increased in the cerebral cortex of LPS-injected mice (Figure 1C), while the expression in LPS-stimulated BV2 cells also showed an increasing trend (Figure 2C). The above results suggest that DAP12 plays an important role in the inflammatory activation process of BV2 cells. Indeed, numerous studies have reported an irreplaceable role of DAP12 in the development of neuroinflammatory diseases.²⁷ Therefore, we speculated that FcγRIIb has a specific connection with DAP12. To verify this idea, we performed immunoprecipitation experiments on proteins extracted from BV2 cells treated with LPS for 9 h. The experimental results showed that FcγRIIb in LPS-treated BV2 cells interacted with DAP12 (Figure 5A). Taken together, these results suggest that FcγRIIb and DAP12 interact in inflammation-activated microglia.

Although we know that DAP12 interacts with FcγRIIb, we speculate that it is the bridging protein DAP12 that binds to FcγRIIb before inflammation activation of BV2 cells. To test our conjecture, we investigated the association between DAP12 and microglial inflammatory activation by interfering with DAP12 expression in BV2 cells. To this end, we used the cell transfection technique to transfect BV2 cells with siRNA targeting different DAP12 loci (si-

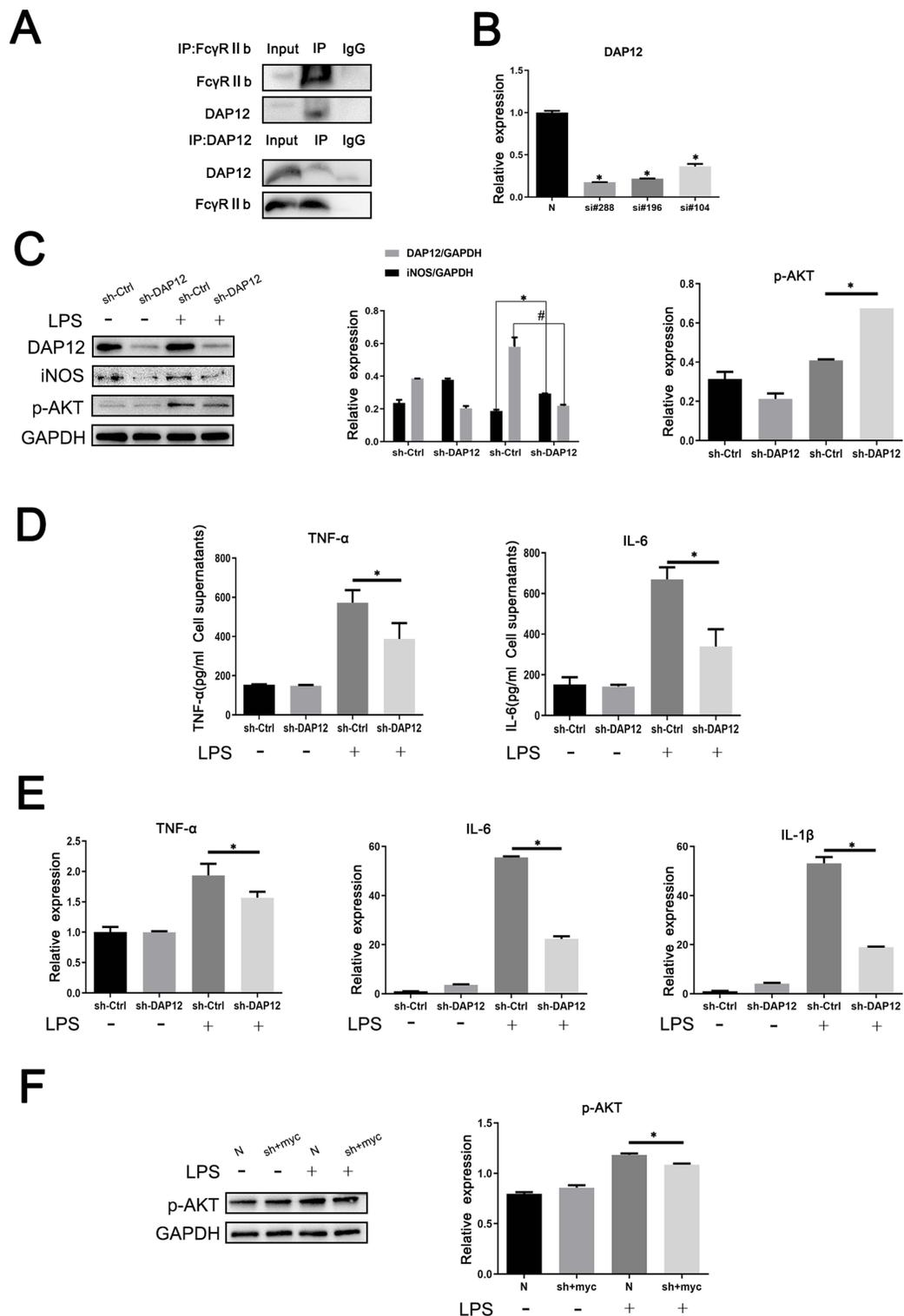


Figure 5 Fc γ RIIb and DAP12 interact to promote microglia activation-mediated LPS-induced neuroinflammation. **(A)** Analysis of the interaction between Fc γ RIIb and DAP12 by immunoprecipitation assay. **(B)** Si-RNA at different loci of DAP12 (si-DAP12#288, si-DAP12#196, and si-DAP12#104) were transfected into BV2 cells, before treating with LPS for 9 h. Q-PCR was performed to detect the transfection efficiency of knocking down si-RNA at different loci of DAP12. **(C)** Western blot detection of the changes in p-AKT, DAP12 and iNOS protein expression levels in BV2 cells after knockdown of DAP12. **(D)** ELISA assay of the expression levels of TNF- α and IL-6 in the supernatants of cells collected by centrifugation. **(E)** Q-PCR assay of the changes in TNF- α , IL-6, and IL-1 β expression in BV2 cells. **(F)** BV2 cells were transfected and then stimulated with LPS. Western blot detected the change in p-AKT protein expression in BV2 cells after overexpressed Fc γ RIIb while interfering with DAP12 (mean \pm SEM represents the result value, n = 3, *Compared to the control, or si-Ctrl+LPS group, or N+LPS group, or si-DAP12+myc-Fc γ RIIb+LPS group, #Indicates compared to the sh-Ctrl+LPS group, p < 0.05).

DAP12#288, si-DAP12#196, and si-DAP12#104) before treating them with LPS for 9 h. The q-PCR results showed that transfection with si-DAP12#288 had the strongest disrupting effect on DAP12 (Figure 5B). After knockdown of DAP12, the Western blot results showed that the expression levels of DAP12 and iNOS proteins were reduced in the LPS-stimulated group (Figure 5C). The ELISA results were consistent with those of Western blot, and showed that interference with DAP12 expression in BV2 cells inhibited the expression of TNF- α and IL-6 (Figure 5D). The q-PCR assay results also clearly showed that knockdown of DAP12 decreased the expression levels of TNF- α , IL-6, and IL-1 β in the stimulated group compared to the control group (Figure 5E). These results indicate that the expression of DAP12 was positively correlated with the degree of inflammatory activation in BV2 cells. Additionally, we found that the phosphorylation level of AKT in the PI3K-AKT signaling pathway was altered after knocking down DAP12; as shown by the Western blot assay, the phosphorylation level of AKT was reduced in the stimulated group compared to the null group (Figure 5C).

To further illustrate the interaction between Fc γ RIIb and DAP12, we overexpressed Fc γ RIIb while interfering with DAP12 to observe the effect on inflammatory activation in BV2 cells. The results of Western blot showed that p-AKT was significantly reduced in the stimulated group compared to the no-load group in BV2 cells after mixing (Figure 5F). Taken together, the results suggest that Fc γ RIIb contributes to microglial inflammatory activation by interacting with the bridging protein DAP12 and relying on DAP12 to deliver activation signals, which in turn activates the PI3K-AKT signaling pathway.

“Fc γ RIIb-PI3K-AKT” Loop Regulates Neuronal Damage Mediated by Microglia Activation-Induced Neuroinflammation

Numerous studies have shown that there are two types of cells in the CNS: neurons and glial cells. Neuroglia are the supporting cells of neurons. When the human brain is damaged, glial cells in the brain are activated to produce large amounts of inflammatory factors that cause neuronal apoptosis; thus, neuroinflammation is closely related to neuronal damage.⁴² To investigate the relationship between the “Fc γ RIIb-PI3K-AKT” and neuronal apoptosis, we constructed a neuronal injury model by first treating BV2 cells according to experimental requirements, collecting cell culture supernatants, and then mixing the collected supernatants with DMEM medium at a ratio of 1:1. Finally, we mixed the collected supernatant with DMEM medium as conditioned medium to culture HT22 neurons for 24 h. The results of Western blot assay showed that the expression of apoptosis-related protein Bax was significantly lower in the si-Fc γ RIIb+LPS group compared to the control group (si-Ctrl+LPS), and when interfering with Fc γ RIIb and blocking the PI3K-AKT pathway (si-Fc γ RIIb+GDC+LPS group), the decrease in Bax expression in neurons was more obvious (Figure 6A). Moreover, the cell viability of the si-Fc γ RIIb+LPS, GDC+LPS, and si-Fc γ RIIb+GDC+LPS groups increased, with that of the si-Fc γ RIIb+GDC+LPS group recovering most significantly (Figure 6B). The above results showed that the positive feedback loop of “Fc γ RIIb-PI3K-AKT” promoted the inflammatory activation of microglia, and that interfering with Fc γ RIIb and blocking the PI3K-AKT pathway could reduce the neuronal damage mediated by LPS-induced neuroinflammation.

Discussion

As human life expectancy increases, neurodegenerative diseases, including Alzheimer’s disease (AD), Parkinson’s disease (PD), and multiple sclerosis (MS), pose an increasing threat to human health.^{42–45} With the increasing mortality rate, the concern for neurodegenerative diseases is growing annually, which has resulted in in-depth exploration. Numerous studies have shown an inextricable relationship between neuroinflammation and neurodegenerative diseases. Innate immune cells, microglia, and astrocytes are important players in neuroinflammation and neurodegenerative diseases. Neuroinflammation is generally an inflammatory response in the CNS caused by various pathological injuries. The onset and progression of neuroinflammation is characterized by a significant increase in the release of pro-inflammatory factors, the release of which leads to loss of synaptic function, neuronal apoptosis, and inhibition of nerve growth.

The CNS contains neurons and glial cells. In contrast to neurons, glial cells are functionally diverse and diverse in type, including astrocytes, oligodendrocytes, and microglia, all of which perform different functions and regulate

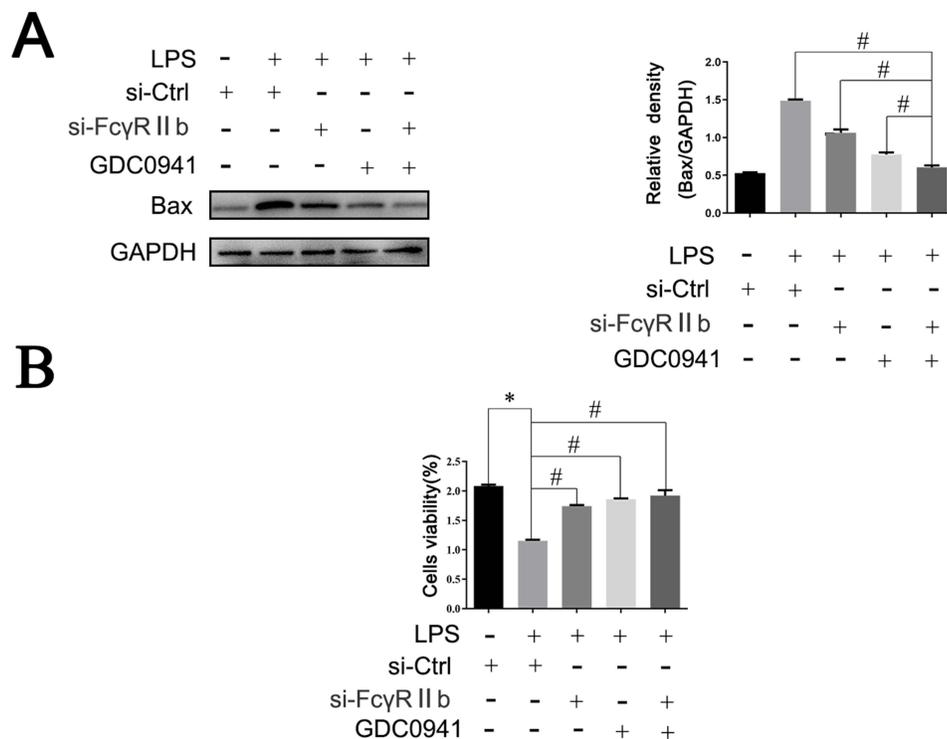


Figure 6 The “F γ RIIb-PI3K-AKT” positive feedback loop exacerbates neuroinflammation-mediated neuronal injury. BV2 cells were transfected with si-Ctrl or si-Fc γ RIIb, pretreated with GDC0941 inhibitor, transfected with si-Fc γ RIIb, treated with pathway inhibitor GDC0941, and then stimulated with LPS. Subsequently, the conditioned medium was prepared with DMEM complete medium and used to culture HT22 neurons for 24 h. **(A)** Western blot to detect the expression level of apoptosis-related protein Bax. (n = 3, # represents p < 0.05 compared to the si-Ctrl+LPS group, si-Fc γ RIIb+LPS group, and GDC+LPS group). **(B)** CCK-8 assay of HT22 cell viability (n = 3, *Compared to the si-Ctrl group, #Compared to the si-Ctrl+LPS group, p < 0.05).

different mechanisms in the brain.^{46,47} Studies have shown that microglia have two distinct phenotypes (neurotoxic and neuroprotective) that keep the brain immune system in dynamic balance. Moreover, microglial phenotypic changes and neuroprotective and neurotoxic functions play different roles depending on the stage of development of neurodegenerative diseases and their severity. Thus, a simple phenotypic classification does not completely separate the various functional areas of microglia. Indeed, the complexity of this process may be the main reason why anti-inflammatory drug trials have not shown significant therapeutic effects.^{48,49}

Fc γ RIIb is available for experimental exploration as an inflammatory factor. All indications suggest that Fc γ RIIb plays different roles and functions in neuroinflammation. In the present study, after exploring the role of Fc γ RIIb in LPS-induced neuroinflammation, we found that in CNS inflammation, Fc γ RIIb binds to the bridging protein DAP12 after LPS stimulation and forms a complex in microglia, thereby activating the PI3K-AKT signaling pathway, which ultimately leads to neuronal apoptosis and brain injury.

We constructed in vivo and in vitro models of LPS-induced neuroinflammation, and we demonstrated that Fc γ RIIb expression was significantly upregulated in microglia activated by inflammation both in vivo and in vitro through several experiments. Additionally, as the PI3K-AKT signaling pathway plays a non-replaceable role in microglia, we investigated and explored this pathway, with the results showing that Fc γ RIIb promotes microglia activation by activating the PI3K-AKT signaling pathway, while PI3K-AKT pathway activity can also inversely affect the expression of Fc γ RIIb. Fc γ RIIb forms a positive feedback loop, which promotes the development of LPS-induced neuroinflammation, and the “Fc γ RIIb-PI3K-AKT” loop exacerbates neuronal damage mediated by LPS-induced neuroinflammation. We now know from the available experimental data that Fc γ RIIb does not play an inhibitory role in LPS-induced neuroinflammation, but rather promotes the development of inflammatory activation. Therefore, we hypothesize that Fc γ RIIb does not play an inhibitory role because it does not stimulate the unique ITIM structure but rather binds to the bridging protein DAP12 to promote the activation of downstream pathways. DAP12 has been reported to act as a co-stimulatory membrane

protein upon LPS stimulation, thereby promoting PI3K-AKT signaling pathway activation, which may be important as a mechanism for the pathogenesis of LPS-induced neuroinflammation. Due to the above speculation, we first confirmed that FcγRIIb interacts with DAP12 by immunoprecipitation assay. Therefore, we next knocked down the bridging protein DAP12, and indirectly verified that the combination of FcγRIIb and DAP12 could promote the activation of microglia. The experimental data showed that AKT phosphorylation was substantially reduced or disappeared after interfering with DAP12. Moreover, by interfering with DAP12 and overexpressing FcγRIIb, we demonstrated that the reduction in AKT phosphorylation was greatly reduced or disappeared. Although the above results widely support our conjecture, given the complexity of the molecular structure, the mechanism of action is unclear and more in-depth exploration is needed to verify our hypothesis.

To investigate the regulatory mechanism of FcγRIIb in LPS-induced neuroinflammation in an *in vitro* model of LPS-induced neuroinflammation, the validation results showed that LPS stimulation activated microglia, increased FcγRIIb expression, increased release of related inflammatory factors, and resulted in neuronal damage. In an *in vivo* model of LPS-induced neuroinflammation, FcγRIIb expression was significantly increased in the brain tissue of mice that had undergone intraperitoneal injection of LPS. The Gene Expression Omnibus (GEO) is an international gene expression repository and a free public database, with datasets relating to the cerebellum, cortex, spinal cord, and hippocampus.^{50,51} Pathway analysis showed that FcγRIIb is enriched in the PI3K-AKT signaling pathway, which is essential for neuronal development and glial cell activity in the brain.^{52,53} Aberrant transduction of the PI3K-AKT signaling pathway can be involved in various diseases, including cancer, neurodegeneration, and neuroinflammatory diseases. Therefore, we speculate that FcγRIIb plays a unique role in CNS inflammation and that FcγRIIb can trigger LPS-induced neuroinflammation by regulating the PI3K-AKT pathway and activating microglia.

To investigate how FcγRIIb regulates the PI3K-AKT pathway and activates microglia to trigger LPS-induced neuroinflammation, we inhibited AKT phosphorylation in microglia using a PI3K inhibitor and showed that the release of inflammatory factors was reduced. Simultaneously, we found that microglia treated with LPS activated AKT phosphorylation, which confirmed the involvement of the PI3K-AKT pathway in microglia activation.^{54–56}

As FcγRIIb is a transmembrane protein, we speculate that it triggers the lower-level signaling pathway by interacting with other receptors or proteins. Therefore, the analysis of the protein network interactions results suggested that FcγRIIb binds to another transmembrane protein DAP12 to form a complex and thus activate intracellular signaling, causing neural damage. To verify this speculation, DAP12 was confirmed to interact with FcγRIIb by immunoprecipitation assay. To further test our hypothesis, we found that AKT phosphorylation was significantly inhibited after disturbing DAP12 in activated microglia, while overexpression of FcγRIIb and DAP12 disruption in activated microglia showed a similar downregulation of AKT phosphorylation. Based on our results, we suggest that FcγRIIb binds to the transmembrane protein DAP12 to form a complex that activates intracellular signaling and causes neural damage. Many typical functions of FcγRIIb have been achieved by signaling through cytoplasmic ITIM, thus playing an inhibitory role in immunomodulatory responses. Our experimental data suggest that the ITIM structural domain is not activated in LPS-stimulated microglia and that FcγRIIb binds to the receptor adapter DAP12 in the cytoplasmic membrane to form a complex, thereby activating the PI3K-AKT pathway and causing brain injury.

It is worth noting that our experimental results only demonstrate that FcγRIIb binds to the bridging protein DAP12 to form a complex in LPS-treated microglia, which activates the PI3K-AKT pathway and causes neuronal damage, but many of the specific mechanisms involved are still unclear and need to be further explored.

In conclusion, We found that FcγRIIb expression was upregulated in microglia activated by inflammation *in vitro* and *in vivo*. FcγRIIb activates the PI3K-AKT signaling pathway by interacting with DAP12 and is involved in microglia inflammatory activation, and the PI3K-AKT pathway activity also affects FcγRIIb expression. The “FcγRIIb-PI3K-AKT” loop exacerbates neuronal damage mediated by neuroinflammation (Figure 7).

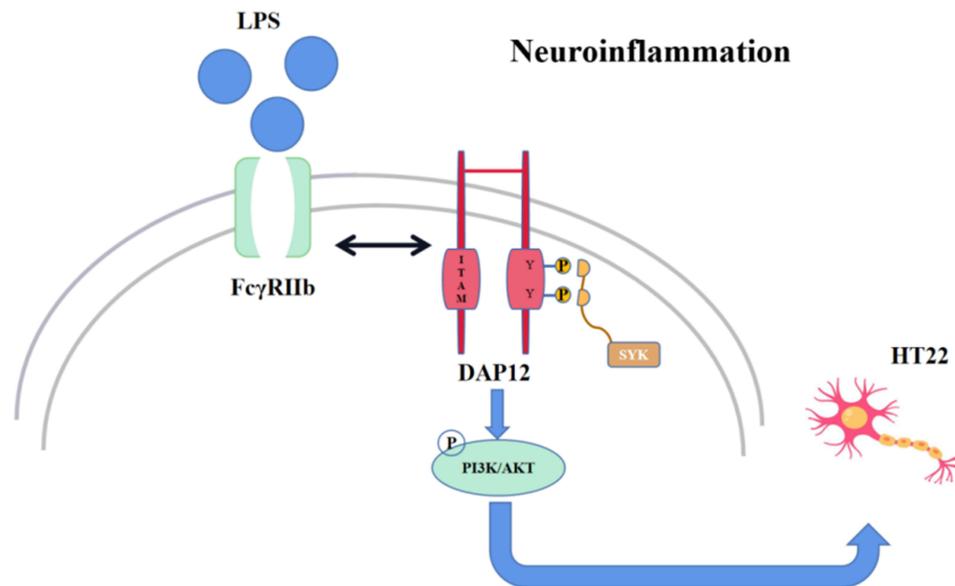


Figure 7 After microglia activation by LPS, FcγRIIb forms a dimer with the bridging protein DAPI2, which activates the downstream PI3K/AKT signaling pathway, thereby exacerbating inflammation and causing neuronal damage.

Data Sharing Statement

The datasets used and/or analyzed during the current study are available from the corresponding author on reasonable request.

Ethics Approval and Consent to Participate

All animal experiments were approved by the Animal Care and Use Committee of Nantong University and the Jiangsu Province Animal Care Ethics Committee.

Author Contributions

All authors made a significant contribution to the work reported, whether that is in the conception, study design, execution, acquisition of data, analysis and interpretation, or in all these areas; took part in drafting, revising or critically reviewing the article; gave final approval of the version to be published; have agreed on the journal to which the article has been submitted; and agree to be accountable for all aspects of the work.

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Disclosure

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