

JQ1 Attenuates Neuroinflammation by Inhibiting Inflammasome-Dependent Canonical Pyroptosis Pathway in Sepsis-Associated Encephalopathy

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Research Article

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Abstract

Sepsis-associated encephalopathy (SAE) is manifested clinically in hyperneuroinflammation and associated with increased morbidity worldwide. Pyroptosis, a novel programmed cell death, has been considered as a causative factor of SAE. Bromodomain-containing protein 4 (BRD4) is a member of the Bromo and Extra-Terminal (BET) family, and promotes inflammatory response in various diseases. Thereby, we examined the effect of JQ1, which is a specific selective inhibitor of BRD4, on the inflammasome-induced pyroptosis of hippocampus in sepsis mice model induced by lipopolysaccharide (LPS) treatment. And we found that JQ1 treatment inhibited the phosphorylation of BRD4, alleviated weight loss and splenomegaly, as well as decreased the serum levels of procalcitonin (PCT) and D-lactate dehydrogenase (D-LDH) induced by LPS injection. Moreover, JQ1 administration decreased the expression of nod-like receptor family protein 1 (NLRP1) or 3 (NLRP3) or the absent in melanoma 2 (Aim2) inflammasomes by blocking nuclear factor kappa B (NFκB) signaling in the hippocampus of sepsis mice. Interestingly, we found that JQ1 selectively attenuated the canonical pyroptosis pathway in SAE mice, indicating by reduced expression of Caspase-1, Caspase-11, gasdermin D (GSDMD) and gasdermin A (GSDMA). JQ1 intervention also suppressed the activation of hippocampal microglia and the release of pro-inflammatory factors, such as IL-1β, IL-18 and IL-6 in SAE mice. In addition, JQ1 treatment protected blood brain barrier (BBB) by up-regulated the expression of tight junction protein occludin and ZO-1 in SAE mice. Furthermore, JQ1 administration remarkably rescued neuronal damage in SAE mice, as enhanced expression of hippocampal NeuN and Doublecortin (DCX). Thus, the protective effects of BRD4 inhibitor JQ1 on SAE were verified in neurons via the inhibition of canonical pyroptosis induced inflammation.

Introduction

Sepsis-associated encephalopathy (SAE) is a diffuse brain dysfunction associated with sepsis, which develops a wide variety of disorders in neural functions [1]. Neuroinflammation [2], loss of blood-brain barrier integrity, microglia activation, and neuronal death [3, 4] have been considered as causative factors of SAE, but the pathological changes of SAE are highly complex and multifactorial and still needed to be clarified. Therefore, it is vital to understand the physiopathology and molecular mechanism of SAE for searching a potential therapeutic strategy.

Pyroptosis, featured by cell membrane perforation and disintegration, is an inflammatory form of programmed cell death [5]. This type of cell death is recently suggested to be dependent on the nod-like receptor family protein 1 (NLRP1) or 3 (NLRP3) or the absent in melanoma 2 (Aim2) inflammasome-dependent, which can further trigger the pyroptosis signaling pathways [6, 7]. Pyroptosis executed by a series of pore-forming proteins called Gasdermins superfamily (GSDMs) [8]. GSDMs have six protein subtypes, including gasdermin A (GSDMA), gasdermin B (GSDMB), gasdermin C (GSDMC), gasdermin D (GSDMD), gasdermin E (GSDME) and pejkakin (PJVK) proteins [9]. Except for PJVK, all members of the GSDM superfamily contain a conserved two-domain structure: N-terminal and C-terminal domains. When the N-terminal domain of these GSDMs is released, it possesses pore-forming activity to cause

inflammatory death associated with the loss of cell membrane integrity and release of inflammatory mediators [9]. Currently, GSDMD [10] and GSDMA [11] are confirmed to be cleaved by Caspase-1/11 and both of which mediate canonical pyroptosis pathway. GSDMC is specifically cleaved by Caspase-8 and mediates non-canonical pyroptosis pathway [12], while GSDME is similarly cleaved by Caspase-3, converting apoptosis to pyroptosis [10]. Accumulating evidence indicates that pyroptosis is induced in central nervous system disease including traumatic brain injury (TBI) [13], multiple sclerosis (MS) [14] and Alzheimer's disease (AD) [15]. Blocking pyroptotic cell death has been proved to be neuro-protective in rodents [16, 17], thus, illuminating the pyroptotic mechanism would benefit for the development of a cure in those diseases.

Bromodomain-containing protein 4 (BRD4) is an important transcriptional regulator of NF- κ B-dependent inflammatory gene expression [18]. Inhibition of BRD4 by small molecules, such as JQ1, suppresses NF- κ B-dependent inflammatory gene expression [19–21]. Mice deficient of BRD4 in myeloid-lineage cells is resistant to LPS-induced sepsis [22]. Phosphorylation of BRD4 (p-BRD4) is critical for its binding to the promoter region of genes [23], and recent study shows that p-BRD4 is involved in neuroplasticity [24]. However, it remains unclear if altered BRD4 or p-BRD4 contributes to pyroptosis of SAE, and whether BRD4 inhibition can reverse this process.

In this study, we found that up-regulated hippocampal expression of BRD4 and p-BRD4 was associated with SAE in mice. Blunting BRD4 by JQ1 alleviated inflammasome-induced pyroptosis, microglia activation and neuroinflammation. Furthermore, JQ1 administration could maintain the blood brain barrier by enhancing tight junction, inhibit neuronal death, as well as protect hippocampal neurogenesis. These results indicate that BRD4 might serve as a promising therapeutic target for SAE.

Materials And Methods

Animals

Adult male C57BL/6 mice weighing 25–30 g (obtained from the Hunan SJA Lab Animal Center of Changsha, Hunan, China) were used in this study. Mice were group-housed (4–8 mice per cage) at a controlled temperature ($22 \pm 2^\circ\text{C}$) under a 12-h light/dark cycle with free access to food and water. The mice adapted to their new environment for 14 days. The experimental protocol was approved by the Animal Care and Use Committee of the University of South China and conformed to the National Institutes of Health Guide for the Care and Use of Laboratory Animals.

Drugs

LPS from *Escherichia coli* 026: B8 (Sigma, USA) was dissolved in 0.9% NaCl. Based on a previous study [25], mice were treated with a single i.p. injection of LPS (10 mg/kg) to establish the animal model of SAE [26], and an equal volume of saline was administered to another cohort of mice as a vehicle control group. BRD4 inhibition was applied by i.p. pretreatment with 50 mg/kg JQ1 (MCE, Shanghai, China) for 1 h before LPS injection, and an equal volume of DMSO was administered as a control group.

ELISA detection

Mice Procalcitonin (PCT) and D-lactate dehydrogenase (D-LDH) ELISA Kits (CUSABIO, WuHan, China) were used to determine the PCT and D-LDH levels, which are the diagnosis and monitoring markers as sepsis. According to the manufacturer's instructions, 24 h after the LPS injection, mice were decapitated and eyeball blood was collected and centrifuged for 20 min at 1500 g at room temperature. The supernatants were collected and used to measure the PCT and D-LDH levels of each sample. The plasma used to test PCT was diluted with Sample Diluent (1:30). Then 90 µl of standard or sample was added to each well and incubate for 2 h at 37°C, followed by 100 µl of biotin-antibody was added to each well and incubated for 1 h at 37°C and 100 µl of HRP-avidin was also added to each well and then incubate for 1 h at 37°C. Aspirate and wash each well after incubation. Wells were then developed with TMB Substrate for 20 min at 37°C in dark. Stop solution was added to each well to terminate the reaction and the absorbance was measured at 450 nm.

Immunohistochemistry

Mice were sacrificed 24 h after LPS injection by administration of an overdose of 10% chloral hydrate (80 mg/kg) and were then perfused transcardially with saline, followed by 100 ml of ice-cold 4% paraformaldehyde solution (pH 7.4). Brains were removed and fixed for 24 h in 4% paraformaldehyde, and cryoprotected with 15% and then 30% phosphate-buffered sucrose (pH 7.4) overnight. Coronal sections (25 µm) of brains were made, and free-floating sections were treated with 3% H₂O₂ to remove endogenous peroxidase, which was then blocked by placing sections in 5% normal sheep serum in 0.01 M phosphate-buffered saline (PBS) containing 0.1% Triton with X-100 for 2 h. Sections were then incubated with rabbit anti-BRD4 antibody (Bethyl, A301), rabbit anti-p-BRD4 (Millipore, ABE1451), rabbit anti-IBA1 (Abcam, ab5076), rabbit anti-NeuN (Millipore, MAB377) and rabbit anti-DCX (CST, 4604S) at room temperature for 2 h, followed by incubation at 4°C overnight. Secondary reagents comprised biotinylated goat anti-rabbit IgG (Proteintech, China). Diaminobenzidine tetrahydrochloride (DAB) (ZSGB-BIO, China) was used as a peroxidase substrate. Washing for all procedures was accomplished in 0.01 M PBS, except for the blocking step. The slices were captured using a digital camera that was attached to the microscope (Olympus, Japan), and the average optical density (OD) of the BRD4, p-BRD4, IBA1, NeuN and DCX immunoreactivity were detected in regions of DG by using Image J software (NIH, USA). In order to avoid any bias, the researcher quantifying the number was blinded to the treatment used.

Western blotting

Mice were anesthetized with an overdose of 10% sodium pentobarbital (80 mg/kg), and the hippocampus (HIP) were rapidly removed following decapitation. The samples were homogenized in a lysis buffer that contained phenylmethansulfonyl fluoride (PMSF) (CW BIO). And the protein concentration was measured using a BCA Assay Kit (CW BIO). The sample extracts were then separated by sodium dodecyl sulfate-polyacrylamide gel electrophoresis (SDS-PAGE) at 90 V for 2 h and transferred onto PVDF membranes (Millipore) for 2 h at 200 mA. Then, the membranes were blocked with 10% nonfat milk for 2 h at room temperature and incubated with the primary antibodies (Table 1) overnight in 5% skim milk solution at

4°C. The next day, after washing three times in 0.02 M Tween-Tris buffered solution, the membranes were incubated with horseradish peroxidase (HRP)-conjugated donkey anti-rabbit (1:3000, Millipore) or donkey anti-mouse (1:2000, CWBIO) polyclonal secondary antibodies for 1.5 h at room temperature. The signals were detected by an enhanced chemiluminescence (ECL) system (CWBIO). The protein levels were quantified by densitometry using NIH ImageJ (NIH, Bethesda, MD, USA).

Table 1
Information for the primary antibodies used in this study

Antibody	Company	Lot number	Molecular weight (KD)	Dilution ratio
BRD4	Abclonal	WH113278	150	1:500
NF-kB	Proteintech	10745-1-AP	65	1:500
NLRP1	Abclonal	A16212	160	1:1000
NLRP3	CST	15101S	110	1:500
AIM2	Abclonal	A3356	47	1:1000
ASC	Santan Cruze	Sc-514414	24	1:500
Caspase-1	Santan Cruze	Sc-56036	45, 20–22	1:500
Caspase-11	Abclonal	A6495	30	1:500
GSDMD	Abclonal	A18281	53	1:1000
GSDMA	Abcam	Ab181027	49	1:1000
Caspase-8	Proteintech	13423-1-AP	32–45/53–57	1:500
GSDMC	Abclonal	A14550	58	1:1000
Caspase-3	CST	9662S	17,19,35	1:1000
GSDME	Proteintech	13075-1-AP	55	1:1000
CD68	Proteintech	28058-1-AP	100	1:500
IL-1 β	Proteintech	16806-1-AP	30–35	1:500
IL-18	Proteintech	10663-1-AP	22	1:500
IL-6	Proteintech	21865-1-AP	24	1:500
occludin	Proteintech	27260-1-AP	59	1:1000
ZO-1	Proteintech	21773-1-AP	220	1:1000
NeuN	CST	24307S	46–55	1:1000
β -Actin	Proteintech	60008-1-Ig	42	1:3000

RNA Isolation and Real Time-PCR

Mice hippocampus total RNA was extracted by Trizol® reagent (CW BIO) according to the manufacturer's instructions. The RNA purity was determined by the A260 nm/A280 nm absorption ratio. cDNA synthesis was performed with the RevertAid™ First Strand cDNA Synthesis Kit (Fermentas) according to the manufacturer's instructions using 2 µg of total RNA. Gene expression was determined by an ABI-7500 real-time PCR system with TB Green™ Premix Ex Taq™ II (Takara). The primers were designed with Primer 3 software (Table 2). A two-step PCR protocol was used according to the manufacturer's instructions. The PCR cycling conditions were 30 s at 95°C followed by 40 cycles at 95°C for 5 s and 60°C for 45 s. The samples were processed in technical duplicates, and a melting analysis was performed for each sample at the end of PCR. The $2^{-\Delta\Delta C_t}$ method was used to determine the relative gene expression as described in our previous study [27].

Table 2
Information for the primers used in this study

Genes	Primers	Sequence 5'-3'
BRD4	<i>forward</i>	CCCTTTTCTGCTCATATTCAGC
	<i>reverse</i>	AATGTATCATAAGCGGAGAGGG
NLRP1	<i>forward</i>	GCCTCACATCCACATACTGCTCAC
	<i>reverse</i>	AGCTCTGCAATCACTTGGTCACTG
NLRP3	<i>forward</i>	CGTTGCAAGCTGGCTCAGTA
	<i>reverse</i>	GGGACTGGGATACAGCCTT
AIM2	<i>forward</i>	CTGCCGCCATGCTTCCTTAACTAG
	<i>reverse</i>	AGCAGTCCAGTTCGGTAGTGTAGG
ASC	<i>forward</i>	ACAATGACTGTGCTTAGAGACA
	<i>reverse</i>	CACAGCTCCAGACTCTTCTTTA
Caspase-1	<i>forward</i>	AGAGGATTTCTTAACGGATGCA
	<i>reverse</i>	TCACAAGACCAGGCATATTCTT
Caspase-11	<i>forward</i>	ATGCTTCTCCAGACATTCTTCA
	<i>reverse</i>	TGGCCTCCTTTATTGGGTAAAT
GSDMD	<i>forward</i>	CTAGCTAAGGCTCTGGAGACAA
	<i>reverse</i>	GATTCTTTTCATCCCAGCAGTC
GSDMA	<i>forward</i>	GAACTTGCACAAGGAGAGGAAA
	<i>reverse</i>	CATCACCACATAGAGGTTCTCC
Caspase-8	<i>forward</i>	ACCAAATGAAGAACAACCTCG
	<i>reverse</i>	CTTCATTTTTTCGGAGTTGGGTT
GSDMC	<i>forward</i>	CCTTTCCAATGAGATTTGTGGG
	<i>reverse</i>	GGAAACTGGAGAACAACACTTC
Caspase-3	<i>forward</i>	GAAACTCTTCATCATT CAGGCC
	<i>reverse</i>	GCGAGTGAGAATGTGCATAAAT
GSDME	<i>forward</i>	GAGAGTCACTCTTCGTTTGGAA
	<i>reverse</i>	CTGAAGTACCAGGTTGTCCATA

Genes	Primers	Sequence 5'-3'
CD68	<i>forward</i>	GAAATGTCACAGTTCACACCAG
	<i>reverse</i>	GGATCTTGGACTAGTAGCAGTG
IL-1 β	<i>forward</i>	GCAGAGCACAAGCCTGTCTTCC
	<i>reverse</i>	ACCTGTCTTGGCCGAGGACTAAG
IL-18	<i>forward</i>	CTGTTGGCCCAATTACTAACAG
	<i>reverse</i>	TCCCGAATTGGAAAGGGAAATA
Il-6	<i>forward</i>	CTCCCAACAGACCTGTCTATAC
	<i>reverse</i>	CCATTGCACAACCTCTTTTCTCA
GAPDH	<i>forward</i>	ACCACCATGGAGAAGGCTGG
	<i>reverse</i>	CTCAGTGTAGCCCAGGATGC

Statistical analyses

All statistical analyses were conducted with GraphPad Prism 5.0 software (GraphPad Software, San Diego, CA, USA). Data are presented as the mean \pm SEM. Significant differences were determined using ANOVAs followed by Bonferroni's post hoc testing for three groups. $p < 0.05$ was considered statistically significant.

Results

JQ1 treatment inhibits the phosphorylation of BRD4 and prevents LPS-induced sepsis-related symptoms in mice

High dose lipopolysaccharide (LPS) could induce sepsis and sepsis-associated encephalopathy [25]. In the present study, we explore the expression of BRD4 and p-BRD4 in the hippocampus of mice, which is a main region susceptibility to sepsis. Real-Time PCR showed that there was significantly differences among the groups in the mRNA level of BRD4 ($F_{(2,15)} = 12.94$, $p = 0.0008$, Figure. 1A). Bonferroni post-test analysis revealed that the mRNA level of BRD4 in DMSO + LPS mice increased when compared with those of the DMSO + NS group ($t = 4.927$, $p < 0.001$), which was reversed by JQ1 treatment ($t = 3.657$, $p < 0.01$). Western blot was used to detect the protein level of BRD4, and one-way ANOVA suggested that there were significantly differences among the groups in the protein level of BRD4 ($F_{(2,11)} = 17.29$, $p < 0.001$, Figure. 1B and 1C). Bonferroni post-test analysis revealed that when compared with the DMSO + NS group, the protein level of BRD4 in DMSO + LPS mice increased significantly ($t = 4.927$, $p < 0.01$), which was reversed by JQ1 treatment ($t = 5.242$, $p < 0.01$). Furthermore, we investigated the expression of BRD4 and p-BRD4 by immunohistochemistry, and the representative images of BRD4 and p-BRD4-positive cells are shown in Figure. 1D. The analysis results show that there were significantly differences

among the groups in the average optical density (OD) of BRD4 ($F_{(2,19)} = 8.053, p = 0.0035$, Figure. 1E) and p-BRD4 ($F_{(2,44)} = 8.952, p = 0.0006$, Figure. 1E). Bonferroni post-test analysis revealed that LPS challenge could increase the expression of hippocampal BRD4 ($t = 3.764, p < 0.001$) and p-BRD4 ($t = 2.794, p < 0.05$), but JQ1 administration could decrease the expression of BRD4 ($t = 2.872, p < 0.05$) and p-BRD4 ($t = 4.096, p < 0.001$). These results suggested that JQ1 treatment not only inhibited the expression of BRD4, but also repressed the phosphorylation of BRD4.

Sepsis exhibited weight loss and splenomegaly. In the present study, the body weight and spleen weight of DMSO + NS, DMSO + LPS and JQ1 + LPS group mice were also determined. Two-way ANOVA with Bonferroni post-test analysis revealed that 24 h after LPS injection, the body weight of mice decreased when compared with the NS group ($t = 3.403, p < 0.01$), but no significant difference was found between the DMSO + NS groups and JQ1 + LPS mice ($t = 1.232, p > 0.05$, Figure. 1F). One-way ANOVA indicated there was a significant difference among the groups in the spleen weight/body weight ($F_{(2,20)} = 25.99, p < 0.001$, Figure. 1G). Bonferroni post-test analysis revealed that the spleen weight/body weight in DMSO + LPS mice increased when compared with those of the DMSO + NS group ($t = 5.806, p < 0.001$), which was rescued by JQ1 treatment ($t = 6.592, p < 0.001$). These results suggested that JQ1 could inhibit weight loss and splenomegaly induced by LPS administration.

Procalcitonin (PCT) and D-lactate dehydrogenase (D-LDH) levels are the diagnosis and monitoring markers in sepsis. So we also detected the levels of serum PCT and D-LDH by using ELISA kits. One-way ANOVA suggested that there were significantly differences among the groups in the levels of PCT ($F_{(2,16)} = 4.046, p = 0.0076$, Figure. 1H) and D-LDH ($F_{(2,18)} = 10.55, p = 0.0012$, Figure. 1I). Bonferroni post-test analysis revealed that LPS caused increased PCT level ($t = 3.694, p < 0.01$) and D-LDH level ($t = 4.349, p < 0.01$) compared to NS mice, these effects were effectively attenuated by JQ1 treatment ((PCT ($t = 2.779, p < 0.05$) and D-LDH ($t = 3.116, p < 0.05$)). These results indicated that JQ1 could prevent the process of sepsis induced by LPS administration.

JQ1 treatment inhibits LPS-induced the activation of NF- κ B and inflammasomes

Inflammasomes, mainly including NLRP1, NLRP3 and Aim2, play an important role in triggering pyroptosis and inflammatory response in SAE [28]. To determine the role of JQ1 treatment on the activation of the inflammasomes, we examined the alterations of mRNA levels of NLRP1, NLRP3, Aim2 and ASC in the hippocampus of mice. One-way ANOVA suggested that there were significantly differences among the groups in the mRNA level of NLRP1 ($F_{(2,16)} = 26.28, p < 0.0001$), NLRP3 ($F_{(2,17)} = 11.86, p = 0.0008$), Aim2 ($F_{(2,16)} = 6.822, p = 0.0085$) and ASC ($F_{(2,17)} = 14.93, p = 0.0003$) (Figure. 2A). Bonferroni post-test analysis revealed that LPS increased the mRNA level of NLRP1 ($t = 5.312, p < 0.001$), NLRP3 ($t = 4.855, p < 0.01$), Aim2 ($t = 3.113, p < 0.05$) and ASC ($t = 4.346, p < 0.01$) compared to NS mice, while JQ1 treatment decreased the levels of NLRP1 ($t = 2.791, p < 0.001$), NLRP3 ($t = 2.758, p < 0.05$), Aim2 ($t = 3.232, p < 0.05$) and ASC ($t = 5.047, p < 0.001$) compared to those LPS mice.

Nuclear factor- κ B (NF- κ B) signaling pathway was found to be involved in LPS-induced inflammatory response [29]. Furthermore, western blotting was used to determine the protein levels of NF- κ B, NLRP1, NLRP3, Aim2 and ASC in the hippocampus of mice. One-way ANOVA suggested that there were significant differences among the groups in the protein level of NF- κ B ($F_{(2,11)} = 12.89, p = 0.0023$), NLRP1 ($F_{(2,11)} = 18.14, p = 0.0007$), NLRP3 ($F_{(2,11)} = 10.21, p = 0.0048$), Aim2 ($F_{(2,11)} = 22.04, p = 0.0003$) and ASC ($F_{(2,11)} = 11.95, p = 0.0029$) (Figure. 2B and 2C). Bonferroni post-test analysis revealed that LPS increased the protein level of NF- κ B ($t = 3.540, p < 0.05$), NLRP1 ($t = 4.468, p < 0.01$), NLRP3 ($t = 4.188, p < 0.01$), Aim2 ($t = 5.807, p < 0.001$) and ASC ($t = 4.582, p < 0.01$) compared to NS mice, while JQ1 treatment decreased the levels of NF- κ B ($t = 4.922, p < 0.01$), NLRP1 ($t = 5.732, p < 0.001$), NLRP3 ($t = 3.566, p < 0.05$), Aim2 ($t = 5.690, p < 0.001$) and ASC ($t = 3.766, p < 0.05$) compared to those LPS mice. These results suggested that JQ1 treatment could inhibit the activation of NF- κ B signaling pathway and NLRP1, NLRP3 and Aim2 inflammasomes induced by LPS injection.

JQ1 treatment ameliorates pyroptosis in the hippocampus of SAE mice exposed to LPS

Accordingly, a feature of pyroptosis is the pore formation in cell membranes, which could be triggered by gasdermins (GSDMs) family protein [30]. Researches showed that GSDMD [10] and GSDMA [11] were cleaved by Caspase-1/11 upon inflammasome activation and mediated canonical pyroptosis. As displayed in the Real-Time PCR analysis, one-way ANOVA suggested that there were significant differences among the groups in the mRNA level of Caspase-1 ($F_{(2,17)} = 20.84, p < 0.0001$), Caspase-11 ($F_{(2,16)} = 7.816, p = 0.0053$), GSDMD ($F_{(2,17)} = 17.86, p = 0.0001$) and GSDMA ($F_{(2,17)} = 20.27, p < 0.0001$) (Figure. 3A). Bonferroni post-test analysis revealed that LPS stimulated up-regulation of the mRNA level of Caspase-1 ($t = 3.994, p < 0.01$), Caspase-11 ($t = 3.590, p < 0.01$), GSDMD ($t = 3.445, p < 0.05$) and GSDMA ($t = 3.410, p < 0.05$) compared to NS mice, and these effects were effectively attenuated by JQ1 treatment ((Caspase-1 ($t = 6.390, p < 0.001$), Caspase-11 ($t = 3.167, p < 0.05$), GSDMD ($t = 5.952, p < 0.001$) and GSDMA ($t = 6.361, p < 0.001$)). Western blotting further confirmed this fluctuating expression pattern of cleaved Caspase-1, cleaved Caspase-11, GSDMD-N and GSDMA-N. One-way ANOVA revealed that there were significant differences among the groups in the protein levels of cleaved Caspase-1 ($F_{(2,11)} = 7.331, p = 0.0129$), cleaved Caspase-11 ($F_{(2,11)} = 7.146, p = 0.0139$), GSDMD-N ($F_{(2,11)} = 20.19, p = 0.0005$) and GSDMA-N ($F_{(2,11)} = 14.74, p = 0.0014$) (Figure.3B and 3C). Bonferroni post-test analysis revealed that LPS stimulated up-regulation of the protein level of cleaved Caspase-1 ($t = 2.985, p < 0.05$), cleaved Caspase-11 ($t = 3.036, p < 0.05$), GSDMD-N ($t = 6.348, p < 0.001$) and GSDMA-N ($t = 3.618, p < 0.05$) compared to NS mice, these effects were effectively attenuated by JQ1 treatment ((cleaved Caspase-1 ($t = 3.569, p < 0.05$), cleaved Caspase-11 ($t = 3.469, p < 0.05$), GSDMD-N ($t = 3.413, p < 0.05$) and GSDMA-N ($t = 5.316, p < 0.01$)). The results indicated JQ1 treatment inhibited LPS-induced the mRNA and protein levels of canonical pyroptosis pathway.

Caspase-8 could cleave GSDMC, and then mediated non-canonical pyroptosis pathway [12]. In order to determine the role of JQ1 treatment on the non-canonical pyroptosis pathway, we examined the

alterations of mRNA and protein levels of Caspase-8 and GSDMC. The results of Real-Time PCR showed the mRNA level of Caspase-8 ($F_{(2,17)} = 0.2848, p = 0.7561$) and GSDMC ($F_{(2,16)} = 0.2408, p = 0.7892$) had no significant difference among DMSO + NS, DMSO + LPS and JQ1 + LPS group (Figure. 3D). And that western blotting further certificated there were no significant difference among these groups in the protein levels of cleaved Caspase-8 ($F_{(2,11)} = 0.04195, p = 0.9591$) and GSDMC-N ($F_{(2,11)} = 1.797, p = 0.2205$) (Figure. 3E and 3F).

GSDME has been demonstrated to be cleaved by Caspase-3, and then converting apoptosis to pyroptosis [10]. In our present study, we found the mRNA level of Caspase-3 ($F_{(2,17)} = 0.3112, p = 0.7372$) and GSDME ($F_{(2,17)} = 0.01089, p = 0.9892$) (Figure. 3G), and the protein level of cleaved Caspase-3 ($F_{(2,11)} = 2.405, p = 0.1456$) and GSDME-N ($F_{(2,11)} = 0.001761, p = 0.9982$) (Figure. 3H and 3I) had no significantly difference among DMSO + NS, DMSO + LPS and JQ1 + LPS group. Because GSDMB wasn't expressed in mice [31], so we didn't detect the expression of GSDMB. All these results indicated JQ1 treatment inhibited LPS-induced the canonical pyroptosis, but not non-canonical pyroptosis or the apoptosis converting to the pyroptosis.

JQ1 suppresses microglia activity and the release of inflammatory cytokines in the hippocampus of SAE mice exposed to LPS

Microglia are innate immune cells in the brain, it could be activated after LPS injection [32]. Here, we investigated the role of JQ1 treatment on the activation of microglia. One way ANOVA showed there was significantly difference among the groups in the mRNA level of CD68 ($F_{(2,17)} = 13.32, p = 0.0005$, Figure. 4A), which was a marker of activated microglia. Bonferroni post-test analysis revealed that LPS stimulated up-regulation of the mRNA level of CD68 ($t = 3.469, p < 0.05$), this effect was effectively attenuated by JQ1 treatment ($t = 5.045, p < 0.001$). Western blotting further demonstrated that the protein level of CD68 ($F_{(2,11)} = 7.869, p = 0.0106$, Figure. 4B and 4C) was significantly different in DMSO + NS, DMSO + LPS and JQ1 + LPS group. The following post-test analysis revealed that when compared with DMSO + NS mice, LPS stimulation up-regulated the protein level of CD68 ($t = 3.625, p < 0.05$), but JQ1 treatment could reverse the expression of CD68 ($t = 3.209, p < 0.05$). Furthermore, we investigated the expression of IBA1, a marker of activated microglia, by using immunohistochemistry, and the representative images of IBA1-positive cells were shown in Figure. 4D. The results showed that there were significantly differences among the groups in the average optical density (OD) of IBA1 ($F_{(2,46)} = 33.89, p < 0.0001$, Figure. 4E). Bonferroni post-test analysis revealed that LPS stimulation could increase the expression of IBA1 ($t = 8.011, p < 0.001$), but JQ1 administration could decrease the expression of IBA1 ($t = 5.582, p < 0.001$). These results indicated that JQ1 could suppress microglia activity induced by LPS injection.

The main inflammatory cytokines that produced by pyroptosis were the IL-1 β and IL-18 [33, 34], which could induce the release of downstream inflammatory factors, such as IL-6. And the inflammatory cytokines over-expression has been associated with impairments in hippocampal-dependent memory [35]. Here, we examined alterations in the mRNA expression levels of IL-1 β , IL-18 and IL-6 in the

hippocampus of SAE. One-way ANOVA suggested that there were significantly differences among the groups in the mRNA levels of IL-1 β ($F_{(2,16)} = 7.729, p = 0.0055$), IL-18 ($F_{(2,17)} = 16.70, p = 0.0002$) and IL-6 ($F_{(2,17)} = 26.13, p < 0.0001$) (Figure. 4F). Bonferroni post-test analysis revealed that expressions of IL-1 β , IL-18 and IL-6 were significantly increased in DMSO + LPS mice compared with the DMSO + NS mice (IL-1 β : $t = 3.457, p < 0.05$; IL-18: $t = 3.051, p < 0.05$; IL-6: $t = 6.783, p < 0.001$), while JQ1 treatment remarkably decreased the expression of IL-1 β , IL-18 and IL-6 (IL-1 β : $t = 3.455, p < 0.05$; IL-18: $t = 5.776, p < 0.001$; IL-6: $t = 5.556, p < 0.001$). We also examined alterations in the protein levels of IL-1 β , IL-18 and IL-6 in the hippocampus of mice. One-way ANOVA revealed that there were significantly differences among the groups in the protein level of IL-1 β ($F_{(2,11)} = 37.66, p < 0.0001$), IL-18 ($F_{(2,11)} = 16.86, p = 0.0009$) and IL-6 ($F_{(2,11)} = 7.486, p = 0.0122$) (Figure. 4G and 4H). Bonferroni post-test analysis revealed that LPS stimulated up-regulation of the protein level of IL-1 β ($t = 3.315, p < 0.05$), IL-18 ($t = 3.614, p < 0.05$) and IL-6 ($t = 2.978, p < 0.05$) compared to NS mice, while JQ1 treatment remarkably decreased expression of IL-1 β , IL-18 and IL-6 (IL-1 β : $t = 8.603, p < 0.01$; IL-18: $t = 5.743, p < 0.001$; IL-6: $t = 3.629, p < 0.05$). These results suggested JQ1 treatment inhibited the upregulated expression of IL-1 β , IL-18 and IL-6 induced by LPS.

JQ1 prevents tight junction and reverses neuronal morphological damage in the hippocampus of SAE mice exposed to LPS

Hippocampus is one of the important region of blood-brain barrier, high dose of LPS could damage the blood brain barrier (BBB) [36]. In our current study, tight junction proteins occludin and ZO-1 were detected to assess the effect of JQ1 on the blood-brain barrier of hippocampus. And the results showed that there were significantly differences among the groups in the protein level of occludin ($F_{(2,11)} = 24.41, p = 0.0002$) and ZO-1 ($F_{(2,11)} = 35.03, p < 0.0001$) (Figure. 5A and 5B). Bonferroni post-test analysis revealed that LPS stimulated down-regulation of the protein level of occludin ($t = 5.776, p < 0.001$) and ZO-1 ($t = 4.964, p < 0.01$) compared to NS mice, while JQ1 treatment remarkably increased the expression of occludin ($t = 6.293, p < 0.001$) and ZO-1 ($t = 8.319, p < 0.001$).

To determine whether neurons were the main cell type that participated in LPS-induced pyroptosis. Western blotting was used to assess the relative expression of the NeuN protein, which is a marker of neuron, and one-way ANOVA revealed that there were significantly differences among the groups in the protein level of NeuN ($F_{(2,11)} = 11.89, p = 0.003$) (Figure. 5A and 5B). Bonferroni post-test analysis revealed that LPS stimulated down-regulation of the protein level of NeuN ($t = 4.674, p < 0.01$) compared to NS mice, while JQ1 treatment remarkably increased the expression of NeuN ($t = 3.540, p < 0.05$). Furthermore, we used immunohistochemistry to estimate the expression of NeuN, and the newborn neurons maker, Doublecortin (DCX). The representative images of NeuN and DCX-positive cells are shown in Figure. 5C. The results showed that there were significantly differences among the average optical density (OD) of NeuN ($F_{(2,22)} = 47.81, p < 0.0001$) and DCX ($F_{(2,50)} = 34.48, p < 0.0001$) (Figure. 5D). Bonferroni post-test analysis revealed that LPS stimulated could increase the expression of NeuN ($t = 6.443, p < 0.001$) and DCX ($t = 6.540, p < 0.001$), but JQ1 administration could decrease the expression of NeuN ($t = 9.659, p < 0.001$) and DCX ($t = 7.606, p < 0.001$). These results suggested that JQ1

administration could enhance tight junction of hippocampus and inhibit neuronal death, as well as protect neurogenesis.

Discussion

In the present study, we showed that LPS enhanced hippocampal BRD4 and p-BRD4, and upregulated NF- κ B-dependent NLRP1, NLRP3 and Aim2 inflammasomes expression, which finally triggered neuronal damage in a canonical pyroptotic manner and promoted the release of inflammatory factors.

Pretreatment with BRD4 inhibitor JQ1 remarkably suppressed the expression of NLRP1, NLRP3 and Aim2 inflammasomes, then inhibited canonical pyroptotic signaling pathway, which mediated by cleaved Caspase-1/11 and GSDMD, GSMDA, and the associated release of the inflammatory cytokines IL-1 β , IL-18 and IL-6 also decreased in the hippocampus of mice. Moreover, JQ1 protected mice from LPS-induced hippocampal microglial activation, tight junction injury of BBB and neuronal damage. These supports the hypothesis that JQ1 inhibits the activation of inflammasomes, alleviates pyroptosis and inflammation, protects BBB and neuronal, and thus protects mice from SAE.

Sepsis, characterized as life-threatening organ dysfunction caused by a dysregulated host response to infection [37], remains a major challenge in medicine with high mortality rates [38, 39]. In response to infection, the activation of inflammasomes becomes an important mediator of pyroptosis, i.e. NLRP3/Caspase-1-mediated pyroptosis [3]. The hippocampus appears to be vulnerable since sepsis survivors develop pronounced hippocampal atrophy [40]. And it has been showed that pyroptosis occurs in the hippocampus of sepsis-induced brain injury [41], but its mechanism and treatment scheme is still to be evaluated.

Increasing studies indicate that BRD4 might be a promising therapeutic target in multiple diseases associated with inflammatory response, such as osteoarthritis [42], rheumatoid arthritis [43] and spinal cord injury [44]. Currently, it is still unclear whether and how BRD4 or p-BRD4 is involved in pyroptosis after SAE induced by LPS, a widely accepted model of endotoxin-mediated inflammation [45]. LPS injected by intraperitoneal could damage blood-brain barrier [46] and then induced SAE [47]. Herein, we first determined that BRD4 and p-BRD4 were up-regulated in the hippocampus of SAE mice induced by LPS injection, which were both reduced by JQ1 treatment. However, whether p-BRD4 is directly related to SAE needs further study. Our study also demonstrated that JQ1 treatment could inhibit weight loss and splenomegaly, and decrease the release of serum PCT and D-LDH following LPS injection to resist sepsis. In addition, the inhibition of BRD4 by its small molecule inhibitor JQ1 could block BBB disruption as manifested by enhancing the expression of tight junction proteins (occludin and ZO-1). To explore whether neuronal damage existed in SAE, we observed the expression of NeuN and DCX in the brains of SAE mice. The results showed that the numbers of both mature neurons and newborn neurons were significantly decreased in the SAE mice compared with control mice, and JQ1 intervention could obviously reverse these changes, indicating JQ1 could protect BBB and inhibit neuronal damage induced by LPS.

The cleavage of GSDMs by inflammasome-activated caspases are critical step in inducing pyroptosis, and the N-terminal domain of GSDMs (GSDMs-N) is commonly recognized as an executor of pyroptosis [48]. In the present study, we explored the mechanism how JQ1 inhibited neuronal damage in SAE. And the results showed that the levels of GSDMD-N and GSDMA-N, which were cleaved by Caspase-1/11, were increased in the brain of SAE mice. While the expression of GSDMC-N and GSDME-N, which were cleaved by Caspase-8 or Caspase3, had no obvious change. These results indicated that SAE could induce neuronal pyroptosis by a canonical manner. Furthermore, the role of BRD4 in neuronal pyroptosis of SAE was further clarified by the BRD4 inhibitor JQ1. Data in our study demonstrated that JQ1 administration could increase the amount of neuron and meanwhile decrease the expression of GSDMD-N, GSDMA-N and cleaved Caspase-1/11, which suggested that JQ1 could protect neuron from LPS-induced damage through inhibiting neuronal pyroptosis.

Pyroptosis is suggested to be inflammasome-dependent. Herein, we investigated the effects of JQ1 on the NLRP1, NLRP3 and Aim2 inflammasomes, and found that the expression of NLRP1, NLRP3, Aim2 and ASC was significantly increased in the hippocampus of SAE, while these effects were effectively attenuated by JQ1 treatment. These results indicated JQ1 inhibited neuronal pyroptosis dependent on the NLRP1, NLRP3, Aim2 inflammasomes. The transcription factor NF- κ B plays a key role in activating the inflammasome signaling pathway [49], and BRD4 is an important transcriptional regulator of NF- κ B [50]. We further determined the expression of NF- κ B in SAE. We found that JQ1 treatment could effectively down-regulate the expression of hippocampal NF- κ B. Microglia could also be activated after LPS injection [32], and a series of inflammatory cytokines such as IL-1 β and IL-18, which were produced during pyroptosis process, were also released by LPS stimulation [33, 34]. Reduced expression of microglia markers (CD68 and IBA1) demonstrated that JQ1 treatment alleviated LPS-induced microglia activation. Additionally, the expression of IL-1 β , IL-18 and the downstream inflammatory factor IL-6 were increased after LPS injection, while JQ1 administration could reverse their high expression, indicating JQ1 could alleviate neuroinflammation in the hippocampus of SAE mice. Thus, we considered that BRD4 mediated NF- κ B activation to initiate the NLRP1, NLRP3 and Aim2 inflammasomes signaling pathway, resulting in pyroptosis, which might be involved in the pathogenic process of SEA.

Conclusion

In conclusion, our results demonstrated that JQ1 could reduce severity of SAE through partial abrogation of neuroinflammation-related pathogenic mechanisms, and JQ1 could be put forward as an ancillary beneficial agent in combination with available drug treatments to slow down SAE pathogenic process and occurrence rate of its clinical signs.

Declarations

Acknowledgments Not applicable.

Author Contributions Xiao Lin Zhong and Zuyao Chen performed PCR and Western Blotting tests, Yajuan Wang and Mingli Mao performed ELISA and Immunohistochemistry tests, Yingcheng Deng, Mengmeng Shi, Yang Xu established the animal model and given JQ1 intervention. Ling Chen performed statistical analysis. Xiao Lin Zhong and Wenyu Cao drafted the manuscript. Xiao Lin Zhong, Ling Chen and Wenyu Cao performed research design and manuscript revision.

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Compliance with Ethical Standards

Conflict of Interest The authors declare that they have no conflict of interest.

Ethics Approval The experimental protocol was approved by the Animal Care and Use Committee of the University of South China and conformed to the National Institutes of Health Guide for the Care and Use of Laboratory Animals.

Consent for Participate Not applicable.

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Figures

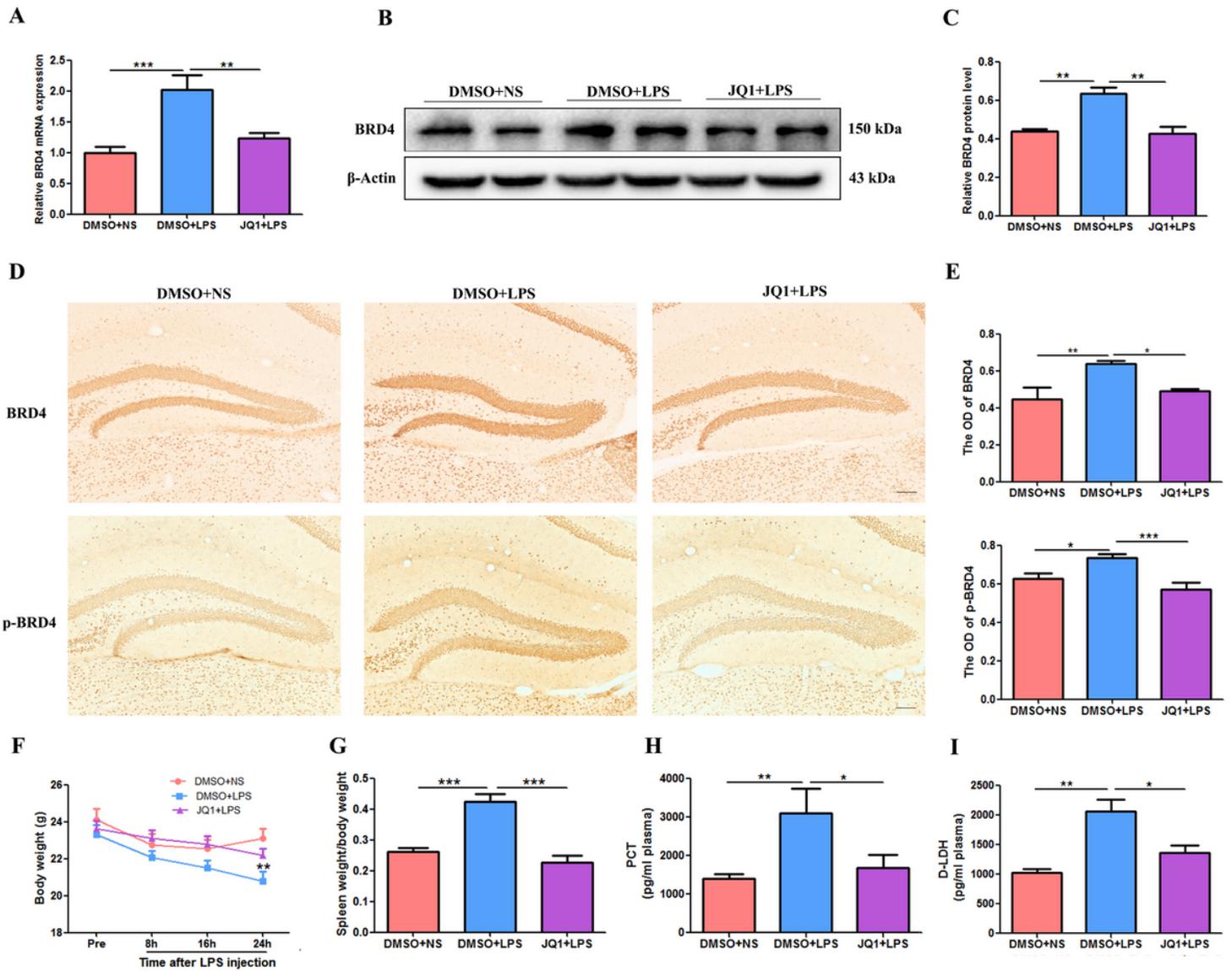


Figure 1

The effect of JQ1 on the expression of BRD4 and its phosphorylation and on the sepsis-related symptoms induced by LPS injection. (A) Representative the mRNA expression level of BRD4. (B) Representative western blotting immunolabeling of BRD4 in the hippocampus and (C) semiquantitative analysis of BRD4 protein relative to β -Actin. (D) Representative immunohistochemistry images of BRD4 and p-BRD4-positive cells and (E) Quantitative analysis of the average optical density (OD) of BRD4 and p-BRD4 in hippocampus for each group. (F) The body weight of mice in pre, 8 h, 16 h and 24 h after LPS stimulation. (G) The spleen weight/body weight of mice 24 h after LPS injection. (H and I) The level of serum Procalcitonin (PCT) and D-lactate dehydrogenase by using ELISA kits. Bar=100 μ m. $p < 0.05$ was considered significant, * $p < 0.05$, ** $p < 0.01$ and *** $p < 0.001$. The results are shown as the mean \pm SEM; $n = 4-8$ animals in each experimental group.

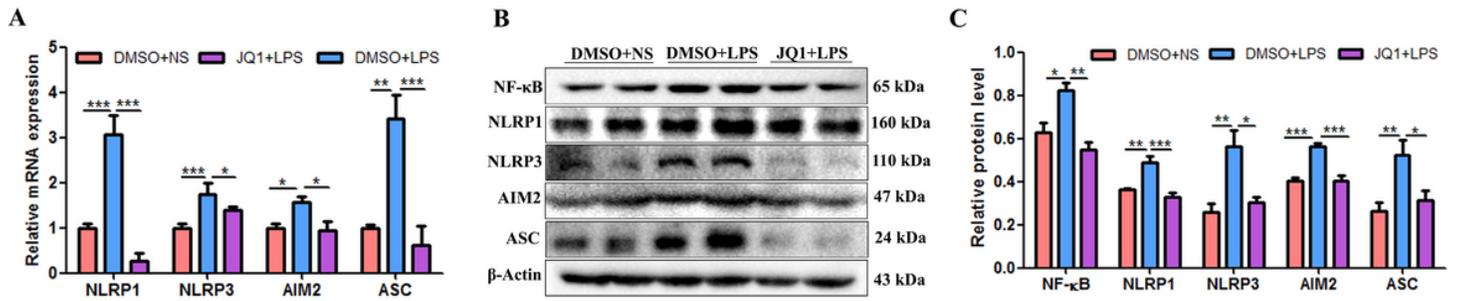


Figure 2

The effect of JQ1 on the expressions of the NF-κB and inflammasomes in the hippocampus induced by LPS injection. (A) Representative the mRNA expression levels of NLRP1, NLRP3, Aim2 and ASC. (B) Representative western blot immunolabeling of NF-κB, NLRP1, NLRP3, Aim2 and ASC and (C) semiquantitative analysis of NF-κB, NLRP1, NLRP3, Aim2 and ASC protein relative to β-Actin. $p < 0.05$ was considered significant, $*p < 0.05$, $**p < 0.01$ and $***p < 0.001$. The results are shown as the mean \pm SEM; $n = 4-8$ animals in each experimental group.

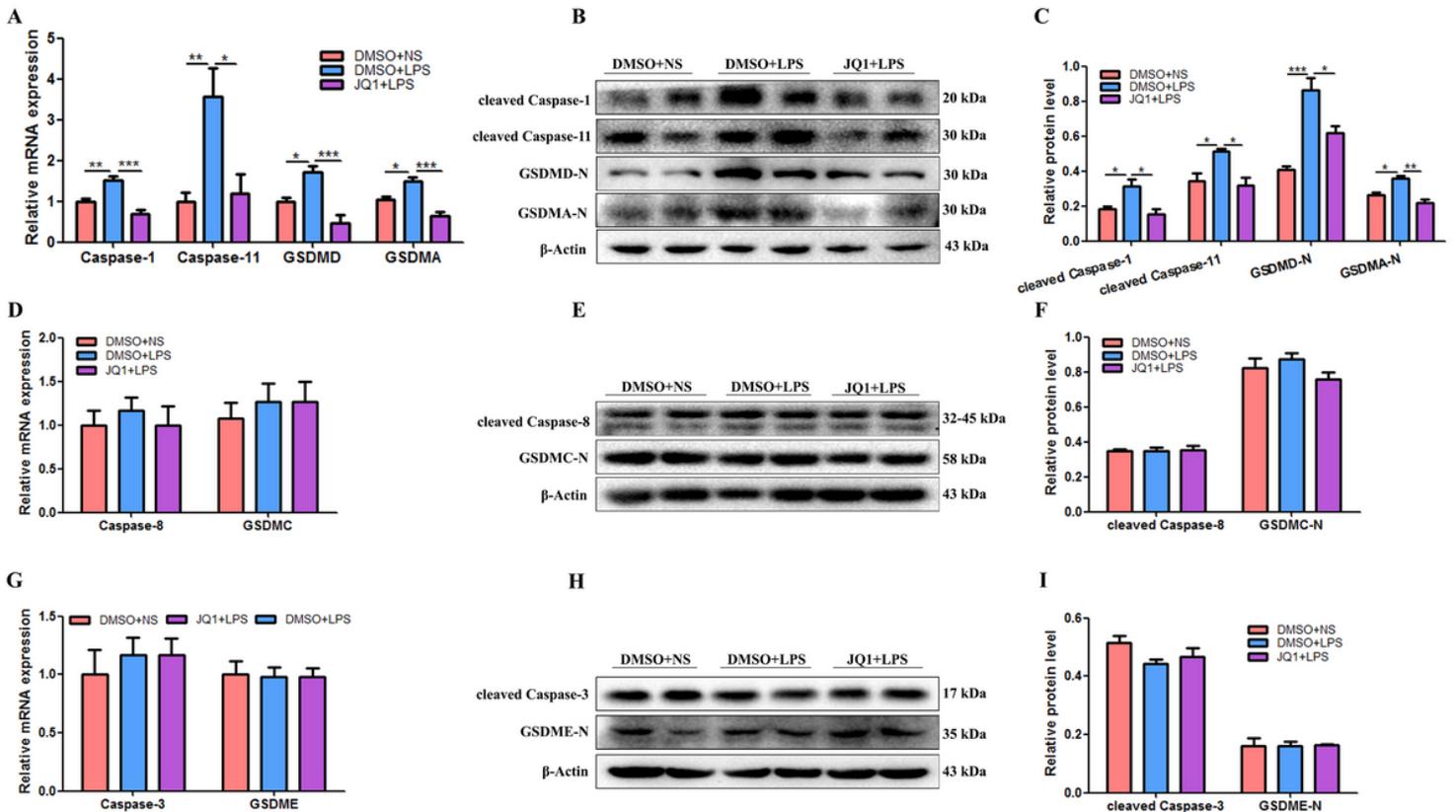


Figure 3

The effect of JQ1 on the expression of gasdermin family protein in the hippocampus induced by LPS injection. (A) Representative the mRNA expression levels of Caspase-1, Caspase-11, GSDMD and GSDMA. (B) Representative western blot immunolabeling of cleaved Caspase-1, cleaved Caspase-11, GSDMD-N and GSDMA-N and (C) semiquantitative analysis of cleaved Caspase-1, cleaved Caspase-11, GSDMD-N

and GSDMA-N protein relative to β -Actin. (D) Representative the mRNA expression levels of Caspase-8 and GSDMC. (E) Representative western blotting immunolabeling of cleaved Caspase-8 and GSDMC-N and (F) semiquantitative analysis of cleaved Caspase-8 and GSDMC-N protein relative to β -Actin. (G) Representative the mRNA expression levels of Caspase-3 and GSDME. (H) Representative western blot immunolabeling of cleaved Caspase-3 and GSDME-N and (I) semiquantitative analysis of cleaved Caspase-3 and GSDME-N protein relative to β -Actin. $p < 0.05$ was considered significant, $*p < 0.05$, $**p < 0.01$ and $***p < 0.001$. The results are shown as the mean \pm SEM; $n = 4-8$ animals in each experimental group.

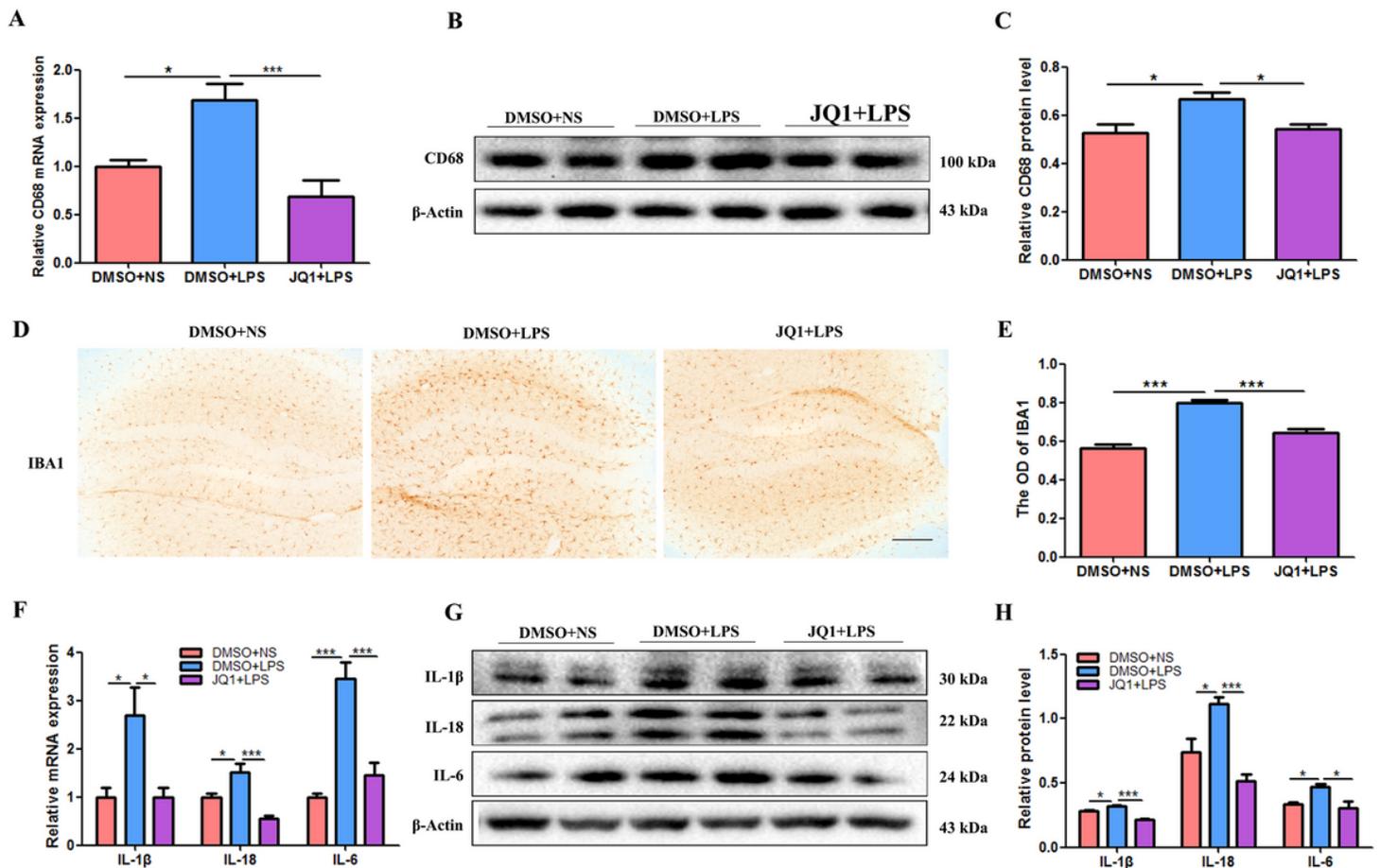


Figure 4

The effect of JQ1 on the activation of microglia and the upregulated expression of inflammatory factors in the hippocampus induced by LPS injection. (A) Representative the mRNA levels of CD68. (B) Representative western blotting immunolabeling of CD68 and (C) semiquantitative analysis of CD68 protein relative to β -Actin. (D) Representative immunohistochemistry images of IBA1 positive cells and (E) quantitative analysis of the average optical density (OD) of IBA1 in hippocampus for each group. (F) Representative the mRNA levels of IL-1 β , IL-18 and IL-6. (G) Representative western blotting immunolabeling of IL-1 β , IL-18 and IL-6 and (H) semiquantitative analysis of IL-1 β , IL-18 and IL-6 protein relative to β -Actin. Bar=100 μ m. $p < 0.05$ was considered significant, $*p < 0.05$ and $***p < 0.001$. The results are shown as the mean \pm SEM; $n = 4-8$ animals in each experimental group.

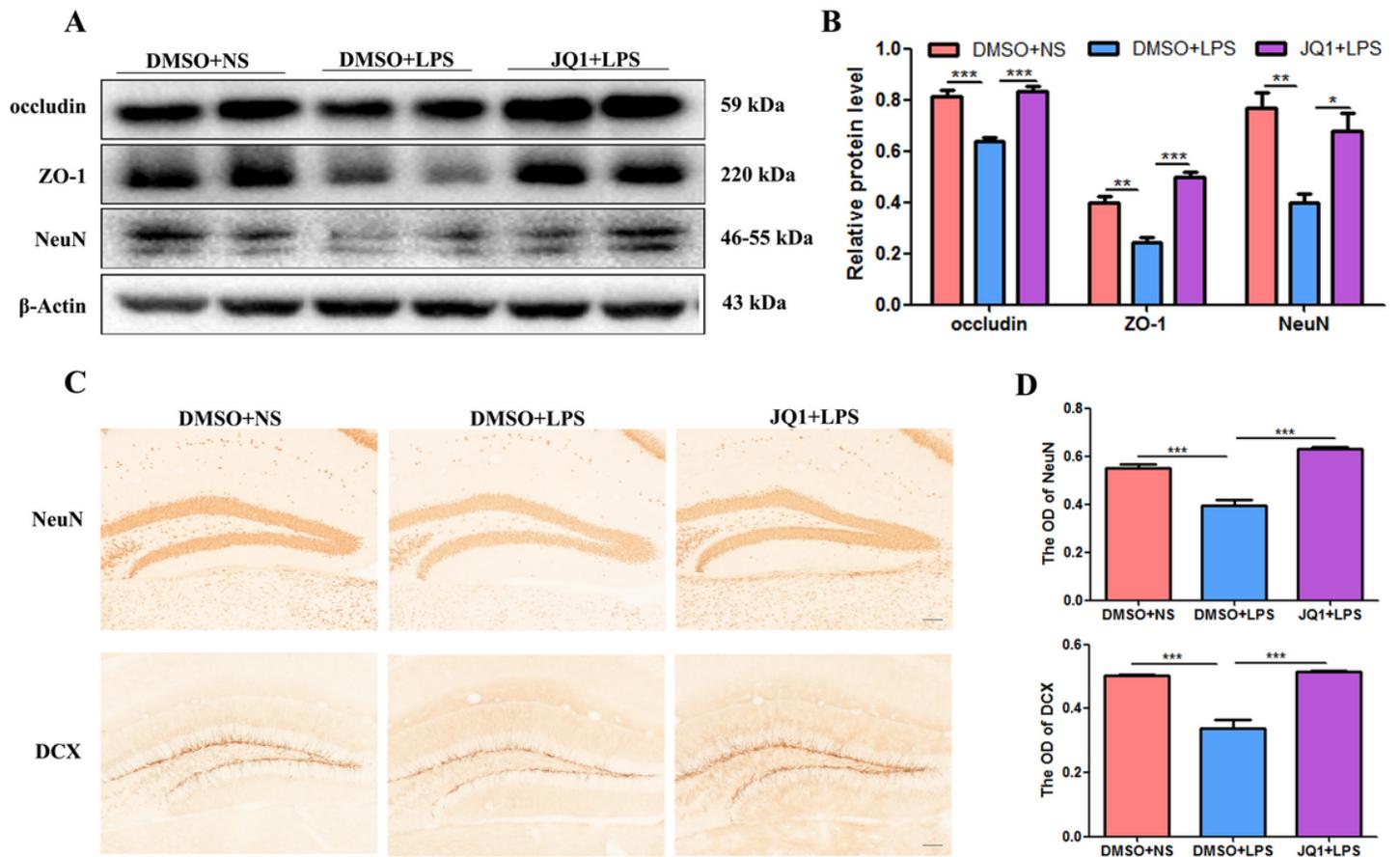


Figure 5

The effect of JQ1 on the neuronal and tight junction proteins in the hippocampus induced by LPS injection. (A) Representative western blotting immunolabeling of occludin, ZO-1 and NeuN and (B) semiquantitative analysis of occludin, ZO-1 and NeuN protein relative to β -Actin. (C) Representative immunohistochemistry images of NeuN and DCX positive cells and (D) quantitative analysis of the average optical density (OD) of NeuN and DCX in hippocampus for each group. Bar=100 μ m. $p < 0.05$ was considered significant, $*p < 0.05$ and $**p < 0.01$. The results are shown as the mean \pm SEM; $n = 4-8$ animals in each experimental group.