

Original Research Article

N-methyl-D-aspartate receptor together with complement C5 accelerated chondrocyte apoptosis and inflammation of osteoarthritis via mediating NLRP3

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Sent for review: 6 November 2022

Revised accepted: 30 March 2023

Abstract

Purpose: NMDAR and complements have been reported to facilitate progression of OA but the correlation between NMDAR and complement C5 in OA is unknown. Therefore, NMDAR antagonist, MK801 and C5 inhibitor, eculizumab were applied to measure functions of NMDAR and C5 in SW1353 cells.

Methods: SW1353 cells were first induced with LPS (5 µg/ml and 10 µg/ml) and later 10 µg/ml LPS induced SW1353 cells were treated by MK801 (100 µM) and eculizumab (100µg/ml). Thereafter, NR1, NR2A and C5 RNA expressions were assessed by RT-qPCR in LPS induced SW1353 cells and NR1 with NR2A was measured under MK801 treatment while C5 was analyzed by eculizumab treatment. Later, apoptosis of SW1353 cells were examined through flow cytometry. Inflammatory cytokines TNF-α, IL-6 and NLRP3 RNA expressions were evaluated by RT-qPCR in SW1353 cells from different groups.

Results: NR1, NR2A and C5 RNA expressions were all significantly upregulated after LPS induction with upregulated apoptosis and increased TNF-α, IL-6 and NLRP3. Treatment of MK801, a NMDAR antagonist, reduced NR1 and NR2A levels. Moreover, apoptosis rate of SW1353 and RNA expressions of inflammatory cytokines were all markedly reduced with MK801. Further, eculizumab treatment significantly downregulated C5 RNA expression in LPS induced SW1353 cells. Added eculizumab also enhanced functions of MK801 in repressing apoptosis rate and inflammatory factors RNA expressions.

Conclusion: MK801 and eculizumab suppressed apoptosis and inflammatory response in SW1353 cells through inhibiting NMDAR and complement C5.

Keywords: NMDAR, complement C5, NLRP3, osteoarthritis

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INTRODUCTION

Osteoarthritis (OA) is the most commonly diagnosed chronic joint disease involving

inflammatory process and other pathological changes, which is the main reason resulting in pain and disability of the aged [1]. OA can be caused by many reasons like age, joint injury,

obesity and so on [2,3] and over a tenth of the population aged older than 60 will develop clinical problems ascribed to osteoarthritis, leading to heavy burden of public health [4]. What's worse, because of aging of population and growing of obesity, occurrence rate of OA will grow gradually [5]. Essentially, OA is a kind of inflammatory disease [6]. During progression of OA, inflammatory response can be attributed to out of balance between inflammatory factors and anti-inflammatory cytokines [7].

Glutamate (Glu) is the main excitatory neurotransmitter through acting on Glu receptor in central nervous system. However, over activation of those receptors can cause damages and even death of nerve cells [8]. The N-methyl-D-aspartate (NMDA) receptor (NMDAR) is a receptor of glutamate that is widely distributed in brain and spinal cord [9]. Based on previous studies, NMDAR is essential to learning and synaptic plasticity and it can mediate death of neurons [10]. Moreover, NMDAR also expressed in other tissues such as renal, cardiovascular system, urinary system and so on [11,12]. In cells, activation of NMDAR caused Na⁺ and Ca²⁺ flowing into cells and overloaded Ca²⁺ resulted in apoptosis or necrosis of cells [13]. NMDAR contains three subunits: NR1, NR2 and NR3 and several different subtypes exists in NR2 (NR2A-D) and NR3 (NR3A and NR3B) [14]. In OA, NMDAR subunits were found to be expressed in OA chondrocytes [15] and NMDAR antagonist, MK801 blocked inflammation of SW1353 cells in a dose dependent manner [15]. Complement system is a representative part of innate immune system which can mediate many kinds of responses like inflammation reaction, cell lysis cessation and pathogen phagocytosis, etc. [16,17] Nevertheless, excessive activation of complements can lead to damages of tissues in diseases [17]. Activation of complements is transited by dissociation of C3 and further activation leads to lysis of C5 to C5a and C5b [18]. C5a is known pro-inflammatory cytokine and C5b induction to C5b-9 can promote inflammation through calcium influx [19,20]. In OA samples, C5 was upregulated, proving that complement C5 participated progression of OA [21]. Moreover, C5a and C5b-9 were both highly expressed in OA patients [22]. For retarding functions of complements, Eculizumab is acknowledged as the inhibitor of complement C5 which can block its activation for preventing cell injury resulted from complements mediated inflammation [23]. Though NMDAR and complement C5 were reported that either of them could accelerate progression of OA, their correlation was seldom mentioned. Therefore, we selected these two cytokines to study the

mechanism beneath them in mediating OA progression and inflammation.

METHODS

Cell culture and treatment

SW1353, human chondrosarcoma cell line [24], was acquired from American Type Culture Collection (ATCC, USA) and incubated in Dulbecco's Modified Eagle Medium (DMEM, Sigma-Aldrich, Germany) replenished with 10% fetal bovine serum (FBS, Sigma-Aldrich, Germany), 2 mM L-Glut and 100 U/ml penicillin and 100 µg/ml streptomycin (Gibco, USA) at 37 °C and 5% CO₂. SW1353 cells were then divided and treated for 12h with different concentrations of LPS (5 µg/ mL and 10µg/ mL) for next experiments [25]. Later, 10 µg/ mL LPS induced SW1353 cells were chosen to be treated with MK801 (100 µM) for 24h [26] and then disposed by 100 µg/ml eculizumab for 4h [18]. After LPS treatment, SW1353 cells in log phase were chosen.

RT-qPCR

After incubation and treatments of SW1353 cells, total RNA was extracted from SW1353 cells strictly according to manufacturer's instruction of TRIzol reagent (Invitrogen, USA). Thereafter, complementary cDNA was compounded through reverse transcription of RNA using SuperScript™ IV First-Strand Synthesis System from Invitrogen (USA). PCR detection then was performed to quantify relative RNA expressions with Fast SYBR™ Green Master Mix (Applied Biosystems, USA) and ProFlex™ PCR System (Applied Biosystems, USA). Sequences of primers were listed as follows: C5: forward, 5'-GGCACAAAGTCCTCCAAATG-3 and reverse, 5'-CCAAACCAAGTCTCCAGTGA-3 [27]; NR1: forward, 5'-GTCCACCAGACTGAAGATTGTGAC-3' and reverse, 5'-CTCCTCCTTGCATGTCCCA-3' [28]; NR2A: forward, 5'-GCTCTTCATCATCAGCAGGG -3' and reverse, 5'-GGATCCCGTCAGATTGAAGTCT -3' [28]; NLRP3: forward, 5'-GGACTATTTCCCAAGATTG-3' and reverse, 5'-ACTCCACCCGATGACAGTT-3' [29] and GAPDH: forward, 5'-GGCATGGACTGTGGTCATGAG -3' and reverse, 5'-TGCACCACCAACTGCTTAGC -3' [28]. For PCR amplification, 2.5 µL diluted cDNA and primers were added into 12.5 µL Fast SYBR™ Green Master Mix and then PCR was performed at ProFlex™ PCR System. Predenaturation was first at 95°C for 10min and later denaturation was at 95°C for 30s. Annealing was at 60°C for 30s and extension was at 72 °C for 1 min, 40 cycles. These three transcripts were

measured in triplicate. Relative expressions of C5, NR1 and NR2A were calculated by $2^{-\Delta\Delta Ct}$ method normalizing to GAPDH.

Flow cytometry

For making apoptosis analysis, SW1353 cells in different groups were first digested with 0.25% trypsin and rinsed by PBS twice. Following, SW1353 cells were resuspended in 195 μ l Annexin V-FITC binding buffer (Beyotime, Shanghai, China) with 5×10^4 cell. Then, 5 μ l Annexin V-FITC (20 μ g/ml) and 10 μ l PI (50 μ g/ml) (Beyotime, Shanghai, China) were added and cultured for 20 min in darkness at room temperature. Attune Flow Cytometer (Invitrogen, USA) was performed to measure distribution of cells. Apoptotic cells were identified as Annexin V-FITC positive and PI negative cells. The detection was repeated three times.

Statistical analysis

All data were displayed as mean \pm SD and analyzed using SPSS 19.0 (USA) and GraphPad Prism 7.0 (USA). Statistical significances of groups were compared with student's t-test and one way ANOVA and $P < 0.05$ was deemed to have statistical meaning.

RESULTS

LPS induced upregulation of NMDAR subunit and complement C5 and promoted apoptosis and inflammatory cytokines of chondrocytes'

To verify involvements of NMDAR and complement C5 in the progression of OA, RNA expressions of NR1, NR2A and complement C5 were measured in different concentrations of LPS induced SW1353 cells. Results showed that NR1, NR2A and C5 RNA levels were all significantly upregulated through LPS induction in a dose dependent manner (Figure 1A). Thereafter, apoptosis of SW1353 under LPS treatment was detected, indicating that growing LPS concentrations resulting in upregulated apoptosis rate of SW1353 cells (Figure 1B). Meanwhile, RNA levels of inflammatory cytokines, TNF- α and IL-6 were checked, revealing that both TNF- α and IL-6 RNA expressions were promoted in a dose dependent manner (Figure 1C). Later, NLRP3 RNA expression was examined in LPS induced SW1353 cells, showing that NLRP3 was also upregulated under LPS treatment (Figure 1D).

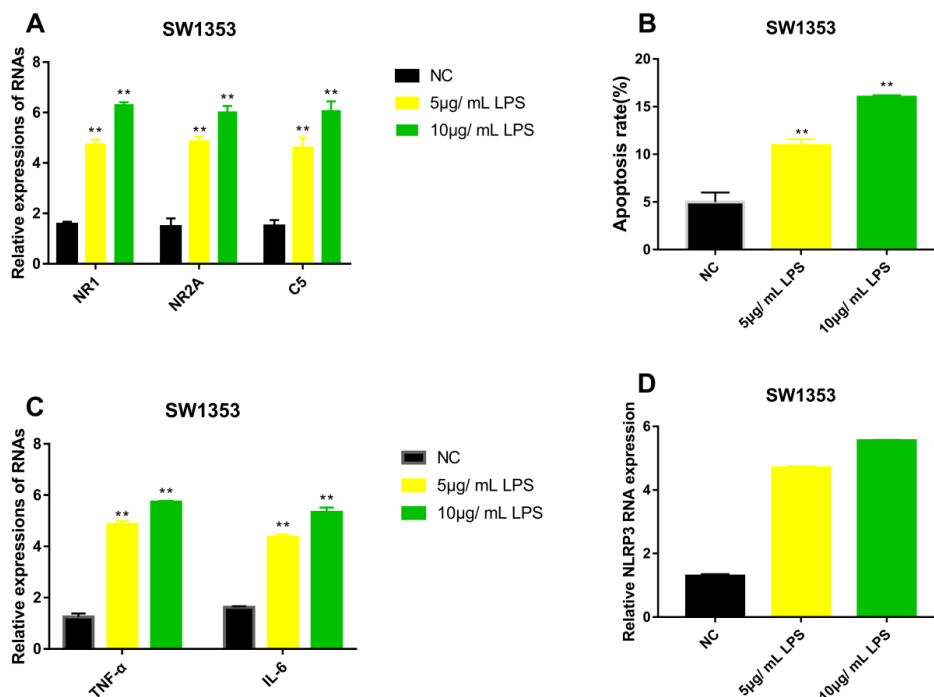


Figure 1: LPS induced upregulation of NMDAR subunit and complement C5 and promoted apoptosis and inflammatory cytokines of chondrocyte A: RNA expressions of NR1, NR2A and C5 were evaluated by RT-qPCR in SW1353 cells under different concentrations of LPS (5 μ g/ml and 10 μ g/ml), $P < 0.05$. B: Apoptosis rate of SW1353 cells treated with different densities of LPS (5 μ g/ml and 10 μ g/ml) were analyzed by flow cytometry, $P < 0.05$. C: TNF- α and IL-6 RNA expressions in LPS (5 μ g/ml and 10 μ g/ml) treated SW1353 cells were assessed with RT-qPCR, $P < 0.05$. D: Relative NLRP3 RNA expressions were validated by RT-qPCR in LPS (5 μ g/ml and 10 μ g/ml) treated SW1353 cells, $P < 0.05$

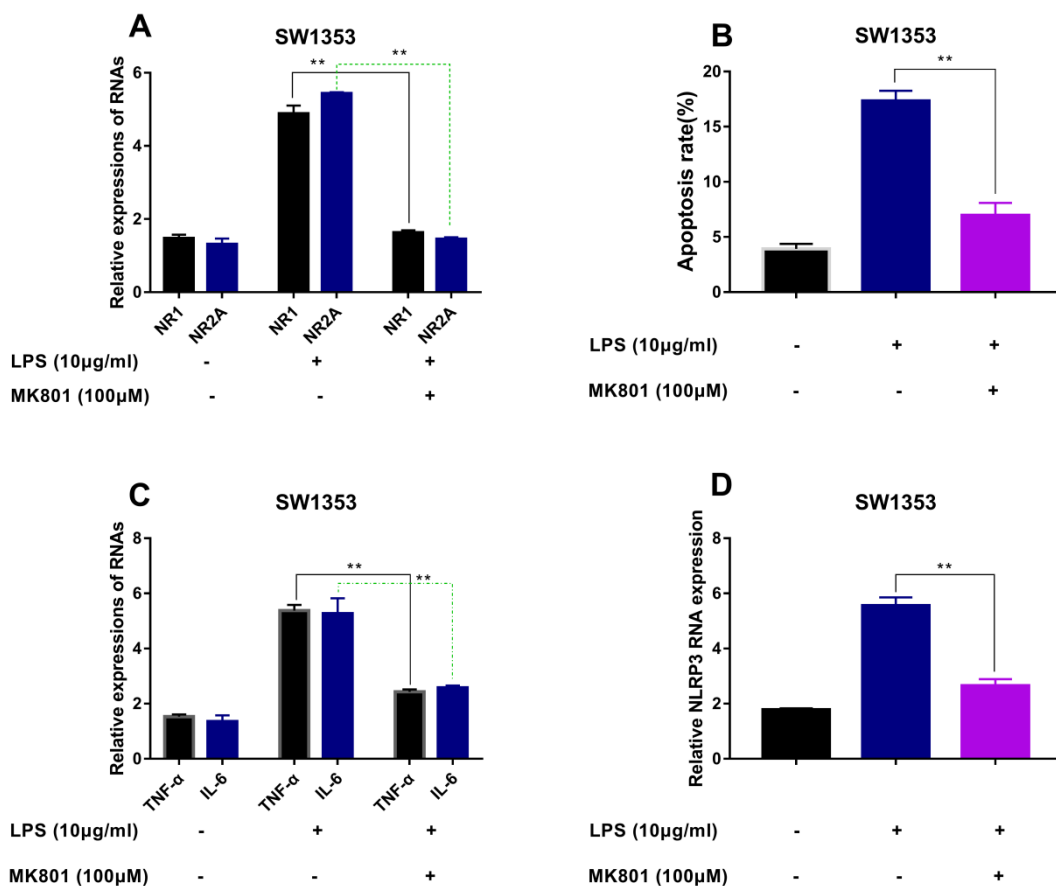


Figure 2: NMDAR antagonist attenuated LPS induced NMDAR subunit and released apoptosis and inflammation in SW1353 cells A: RNA expressions of NR1 and NR2A treated with LPS (10 µg/ml) and MK801 (100µM) were assessed by RT-qPCR in SW1353 cells, $P < 0.05$. B: Apoptosis rate of SW1353 cells under LPS (10µg/ml) and MK801 (100 µM) treatment were checked by flow cytometry, $P < 0.05$. C: TNF-α and IL-6 RNA levels were detected using RT-qPCR in LPS (10µg/ml) and MK801 (100 µM) treated SW1353 cells, $P < 0.05$. D: Relative NLRP3 RNA expression in SW1353 cells after LPS (10µg/ml) and MK801 (100µM) treatment, $P < 0.05$.

NMDAR antagonist attenuated LPS induced NMDAR subunit and released apoptosis and inflammation in SW1353 cells

After RNA expressions of NR1 and NR2A were analyzed in 10µg/ml LPS induced SW1353 cells, mediation of NMDAR antagonist, MK801 (100µM) was measured. After MK801 treatment, both NR1 and NR2A RNA expressions were significantly downregulated compared to untreated SW1353 cells (Figure 2A). Besides that, the apoptosis rate of LPS induced SW1353 cells was markedly reduced after MK801 treatment (Figure 2B). Detections of TNF-α and IL-6 RNA expressions also indicated that added MK801 obviously reduced LPS caused high RNA levels of TNF-α and IL-6 (Figure 2C). Later, MK801 treatment in mediating NLRP3 was detected, indicating that LPS induced upregulated NLRP3 was significantly reversed to low level by MK801 (Figure 2D).

Complement C5 declined LPS induced NMDAR subunit and apoptosis and inflammation of SW1353 cells

After functions of NMDAR in SW1353 cells were examined, role of complement C5 was checked. According to previous experiment, C5 was highly expressed in LPS induced SW1353 cells. Therefore, we used eculizumab, a C5 inhibitor, to figure out mechanism of C5 with NMDAR in progression of OA. As eculizumab (100 µg/ml) applied, LPS caused high expression of C5 was significantly declined (Figure 3A). Thereafter, eculizumab and MK801 treatment resulted in lower apoptosis rate of SW1353 cells compared to SW1353 cells treated by MK801 only (Figure 3B). Following, TNF-α and IL-6 RNA expressions were remarkably decreased under MK801 and eculizumab treatment in comparison with MK801 treatment (Figure 3C). Furthermore, NLRP3 RNA expression was dramatically downregulated after treated by MK801 and added eculizumab led to a much lower level of NLRP3 (Figure 3D).

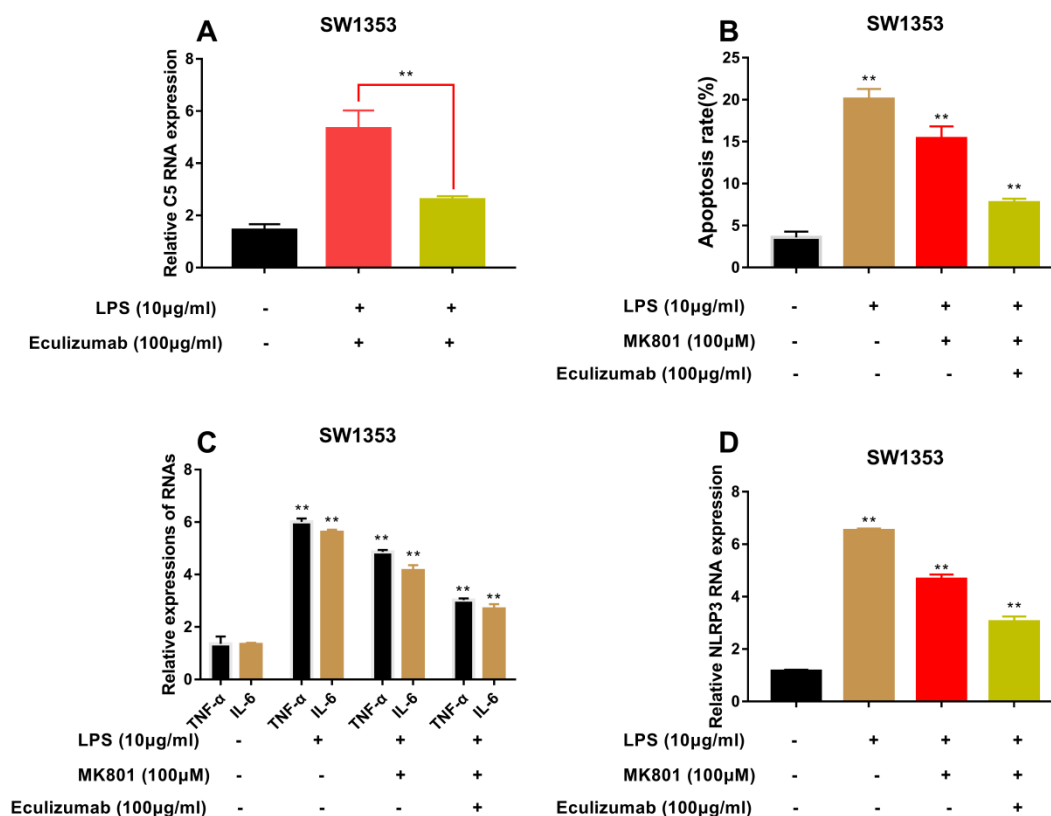


Figure 3: Complement C5 declined LPS induced NMDAR subunit and apoptosis and inflammation of SW1353 cells A: C5 RNA expression in SW1353 cells with LPS (10 µg/ml) and Eculizumab (100 µg/ml) were measured by RT-qPCR, $P < 0.05$. B: Apoptosis rate of SW1353 cells under LPS (10µg/ml), MK801 (100 µM) and Eculizumab (100 µg/ml) were detected by flow cytometry, $P < 0.05$. C: TNF- α and IL-6 RNA expressions were evaluated by RT-qPCR in SW1353 cells with LPS (10µg/ml), MK801 (100 µM) and Eculizumab (100 µg/ml) treatment, $P < 0.05$. D: NLRP3 RNA level in LPS (10 µg/ml), MK801 (100 µM) and Eculizumab (100µg/ml) treated SW1353 cells were analyzed by RT-qPCR, $P < 0.05$

DISCUSSION

According to previous studies, NMDAR worked as a pro-inflammatory factor has been deeply analyzed [30,31]. Normally, NMDAR subunits were represented as objects in progression of OA and suppression of those subunit could retard inflammation in OA [30]. Correspondingly, we also measured functions of NMDAR in SW1353 cells. At the beginning, NR1, NR2A and C5 were apparently upregulated in SW1353 cells after treated by LPS (5 µg/ml and 10 µg/ml) in a dose dependent manner. Moreover, the apoptosis rate of SW1353 was also increased in a dose dependent manner. Furthermore, growing densities of LPS induced upregulation of TNF- α and IL-6 in SW1353 cells were markedly upregulated. Beyond these, NMDA was reported to induce NLRP3 through mediating retinal excitotoxicity in research of Pavlina Tsoka et al. [32]. NLRP3 is the one of the most studied inflammasomes which can activate caspase-1 and resulting in secretions of inflammatory cytokines IL-1 β , IL-18 and TNF- α [33]. In previous studies, inhibition of NLRP3 led to

suppression of pro-inflammatory response in OA samples and cells [34]. Moreover, Icarin attenuated LPS induced inflammation and apoptosis by repressing NLRP3 signaling [35]. NMDA was proven to promote activation of NLRP3 and its activation to caspase-1 needed to be stimulated by activation of C5[32, 36]. Based on these detections, we measured NLRP3 as a biomarker in progression of OA. In our study, NLRP3 RNA expression. Results indicated that LPS simulated OA inflammation could accelerate expressions of NMDAR and complement C5 and upregulated NLRP3 expression. According to our detections, we primarily knew that in LPS induced SW1353 cells, subunits of NMDAR, C5 and NLRP3 have the same tendency.

After that, NMDAR antagonist, MK801 was used to analyze role of NMDAR in LPS induced SW1353 cells. In RNA expressions of NR1 and NR2A detection, MK801 treatment (100 µM) significantly reduced RNA levels of these two subunits which were promoted by LPS (10 µg/ml). Besides that, upregulated LPS induced apoptosis rate of SW1353 cells was dramatically

reduced under MK801 treatment. Furthermore, TNF- α , IL-6 and NLRP3 RNA expression were also declined with MK801 treatment after promoted by LPS. Through these detections, blocking of NMDAR was determined to retard inflammation and apoptosis of SW1353 cells through downregulating subunits NR1 and NR2A. Compared to former studies, we have measured a new way about MK801 suppressing inflammation of OA.

Giuseppina Arbore et al. have declared that assembly of NLRP3 required activation of C5 [36]. Therefore, we measured suppression of C5 in OA to dig out regulation of C5 to NLRP3 in OA. As involvement of NMDAR was examined, role of complement C5 was measured later. Eculizumab was first performed to suppress C5 RNA expression and result of RT-qPCR showed that RNA level of C5 was reversed to low expression in eculizumab (100 μ g/ml) treated LPS-SW1353 cells. Thereafter, correlation between MK801 and eculizumab was checked later, revealing that eculizumab treatment enhanced suppression of apoptosis caused by MK801. Moreover, TNF- α , IL-6 and NLRP3 RNA expressions in LPS induced SW1353 cells with MK801 were also declined after eculizumab treatment. Therefore, C5 played identical functions in facilitating inflammation and apoptosis of SW1353 with NMDAR and eculizumab blocked its role with suppressing NLRP3 and other inflammatory cytokines. Compared to other studies, we have analyzed correlation of C5 and NMDAR in progression of OA and their mediations to NLRP3 inflammasome.

CONCLUSION

NMDAR subunit NR1 and NR2A and complement C5 were all upregulated in LPS induced SW1353 cells and promoted apoptosis and TNF- α , IL-6 and NLRP3 expressions while MK801 inhibited functions of NMDAR and eculizumab suppressed role of complement C5 to retard inflammation and apoptosis of SW1353 cells, indicating that NMDAR and complement C5 could be potential biomarkers to examine inflammatory progression of OA. However, Further studies in vivo and in clinical stage are requested for getting more knowledge of those two factors.

DECLARATIONS

List of Abbreviations

NMDAR: N-methyl-D-aspartate receptor; NLRP3: nucleotide-binding oligomerization domain-

leucine-rich repeat- and pyrin domain-containing 3; OA: osteoarthritis; TNF- α : Tumor Necrosis Factor- α ; IL-6: interleukin 6 NR1: N-methyl-d-aspartate receptor 1; NR2A: N-methyl-d-aspartate receptor 2A; C5: complement C5.

Acknowledgements

None provided.

Funding

This work was supported with funds from Guiyang Science and Technology Contract (2017) No. 30-35 and Guizhou Province Science and Technology Fund Project: Guizhou talents (2018) 5799-50.

Ethical approval

None provided.

Availability of data and materials

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

Conflict of Interest

No conflict of interest associated with this work.

Contribution of Authors

The authors declare that this work was done by the authors named in this article and all liabilities pertaining to claims relating to the content of this article will be borne by them.

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