

Research report

A novel pleiotropic effect of aspirin: Beneficial regulation of pro- and anti-inflammatory mechanisms in microglial cells

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ABSTRACT

Aspirin, one of the most widely used non-steroidal anti-inflammatory drugs, has extensively studied effects on the cardiovascular system. To reveal further pleiotropic, beneficial effects of aspirin on a number of pro- and anti-inflammatory microglial mechanisms, we performed morphometric and functional studies relating to phagocytosis, pro- and anti-inflammatory cytokine production (IL-1 β , tumor necrosis factor- α (TNF- α) and IL-10, respectively) and analyzed the expression of a number of inflammation-related genes, including those related to the above functions, in pure microglial cells. We examined the effects of aspirin (0.1 mM and 1 mM) in unchallenged (control) and bacterial lipopolysaccharide (LPS)-challenged secondary microglial cultures. Aspirin affected microglial morphology and functions in a dose-dependent manner as it inhibited LPS-elicited microglial activation by promoting ramification and the inhibition of phagocytosis in both concentrations. Remarkably, aspirin strongly reduced the pro-inflammatory IL-1 β and TNF- α production, while it increased the anti-inflammatory IL-10 level in LPS-challenged cells. Moreover, aspirin differentially regulated the expression of a number of inflammation-related genes as it downregulated such pro-inflammatory genes as *Nos2*, *Kng1*, *IL1 β* , *Ptgs2* or *Ccr1*, while it upregulated some anti-inflammatory genes such as *IL10*, *Csf2*, *Cxcl1*, *Ccl5* or *Tgfb1*. Thus, the use of aspirin could be beneficial for the prophylaxis of certain neurodegenerative disorders as it effectively ameliorates inflammation in the brain.

1. Introduction

Microglia is the principal immune cell in the central nervous system (CNS). Under pathophysiological conditions such as injury, infection or neurodegeneration (Kettenmann et al., 2011; Kreutzberg, 1996) microglia become activated that proliferate, phagocytose and release pro- and anti-inflammatory cytokines, growth factors or reactive species (Gehrmann et al., 1995; Hanisch, 2002; Kreutzberg, 1996; Luo and Chen, 2012; Smith et al., 2012). Neuroinflammatory

processes are strongly associated with the development of Alzheimer's disease (AD; Akiyama et al., 2000; McGeer and McGeer, 2004), Parkinson's disease (PD; Phani et al., 2012; Shin et al., 2015) and multiple sclerosis (MS; Minagar et al., 2002; Muzio et al., 2007). In AD, for example, β -amyloid accumulation leads to the production of cytokines (IL-1 β , IL-6, TNF- α) and the activation of nitric oxide synthase (Nos) gene through the nuclear transcription factor kappa B (NF- κ B) signaling pathway (Medeiros et al., 2007). IL-1 β in turn increases the expression of amyloid precursor protein (Ge and Lahiri, 2002), while increased expression of TNF- α is reported for both MS

Abbreviations: AD, Alzheimer's disease; Cd, cluster of differentiation; CNS, central nervous system; COX, cyclooxygenase (prostaglandin-endoperoxide synthase, EC 1.14.99.1); DIV, days in vitro; DMEM, Dulbecco's Modified Eagle's Medium; E18, embryonic day 18; ELISA, enzyme-linked immunosorbent assay; ERK1/2, extracellular-regulated kinase 1/2; FBS, fetal bovine serum; GAPDH, glyceraldehyde 3-phosphate dehydrogenase (EC 1.2.1.12); Iba1, ionized calcium binding adaptor molecule 1; IL, interleukin; I κ B, inhibitor of κ B; LPS, bacterial lipopolysaccharide; MS, multiple sclerosis; NF- κ B, nuclear transcription factor kappa B; NO, nitrogen oxide; Nos, nitrogen oxide synthase (EC 1.14.13.39); NSAID, non-steroidal anti-inflammatory drug; p38-MAPK, p38 mitogen-activated protein kinase; PCR, polymerase chain reaction; PD, Parkinson's disease; PBS, phosphate-buffered saline; RT, room temperature; SEM, standard error of mean; SDS, sodium dodecyl sulfate; subDIV, subcloned days in vitro; TBS, Tris-buffered saline; TI, transformation index; TLR4, Toll-like receptor 4; TNF- α , tumor necrosis factor α .

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and PD (Minagar et al., 2002). Strong anti-inflammatory mechanisms can also lead to the development of neurodegenerative diseases (Gehrmann et al., 1995; Ghosh et al., 2013; Gonzalez-Scarano and Baltuch, 1999; Graeber, 2010; Streit, 2002).

Aspirin (acetylsalicylic acid) is one of the most commonly used non-steroidal anti-inflammatory drugs (NSAIDs). By inhibiting the enzyme cyclooxygenase (COX; prostaglandin-endoperoxide synthase, EC 1.14.99.1) it blocks the conversion of arachidonic acid to prostanoids (Vane and Botting, 2003). Apart from its well known beneficial anti-inflammatory effects in the cardiovascular system (Amann and Peskar, 2002) and its role in anti-cancer therapy (Cuzick et al., 2014), it also has widespread effects on immune cell functions such as NO production, cytokine (e.g. IL-1 β , IL-6, IL-8, TNF- α) and adhesion molecule expression (Hussain et al., 2012). Its anti-inflammatory function is based on the inhibition of the NF- κ B signaling pathway (Amann and Peskar, 2002), while its neuroprotective effect in the injured brain is linked to COX-2 inhibition; overexpression of COX-2 is indicative of neuronal damage (Berk et al., 2013; Strauss, 2008). Although aspirin has not been extensively studied in microglia, its anti-inflammatory actions are recently reported (Medeiros et al., 2013; Wang et al., 2011; Yang et al., 2014).

In the present study we investigated the pleiotropic effects of aspirin on secondary, pure microglial cultures derived from mixed primary cultures of 18-day-old embryonic (E18) rat forebrains under control (unchallenged) and bacterial lipopolysaccharide (LPS)-challenged conditions (Gresa-Arribas et al., 2012; Kata et al., 2016; Lund et al., 2006). To reveal the effects of aspirin on pro- and anti-inflammatory mechanisms, we performed quantitative morphometric, functional and gene expression studies relating to phagocytic capability, pro- and anti-inflammatory cytokine production (IL-1 β , TNF- α , and IL-10, respectively) and the expression of various genes related to inflammation. A preliminary report on the effects of aspirin on pure microglial cultures was presented in poster form (Kata and Gulya, 2014).

2. Material and methods

2.1. Animals

All animal experiments were carried out in strict compliance with the European Council Directive (86/609/EEC) and EC regulations (O.J. of EC No. L 358/1, 18/12/1986) regarding the care and use of laboratory animals for experimental procedures, and followed the relevant Hungarian and local legislation requirements. The experimental protocols were approved by the Institutional Animal Welfare Committee of the University of Szeged (I-74-II/2009/MÁB). The pregnant Sprague-Dawley rats (170–190 g) were kept under standard housing conditions and fed ad libitum.

2.2. Antibodies

For a thorough characterization of different microglial phenotypes developed in vitro, an antibody against the ionized calcium binding adaptor molecule 1 (Iba1), an intracellular actin- and Ca²⁺-binding protein expressed in the CNS specifically in macrophages and microglia (Ahmed et al., 2007), was used in our immunocytochemical and Western blot analyses. The anti-glyceraldehyde 3-phosphate dehydrogenase (GAPDH) antibody was used as an internal control in Western blot experiments (Wu et al., 2012). Antibody dilutions, incubation times and blocking conditions were tested for both immunocytochemistry and Western blot analysis. To detect the specificities of the secondary antisera, staining without the primary antibody was performed. In such cases, no fluorescent or Western blot signals were detected.

2.3. Cell cultures

Pure microglial cells were isolated from mixed primary cortical cell cultures of rat embryos of either sex by the method we described earlier (Kata et al., 2016; Szabo and Gulya, 2013). Sibling embryos obtained from the same pregnancy were processed for culturing together; each pregnancy was considered as an independent experiment. Briefly, 10–12 fetal rats (E18) under anesthesia were decapitated and the frontal lobe of the cerebral cortex was removed, minced with scissors, incubated in 9 ml Dulbecco's Modified Eagle's Medium (DMEM; Invitrogen, Carlsbad, USA) containing 1 g/l D-glucose, 110 mg/l Na-pyruvate, 4 mM L-glutamine, 3.7 g/l NaHCO₃, 10,000 U/ml penicillin G, 10 mg/ml streptomycin sulfate, 25 μ g/ml amphotericin B and 0.25% trypsin for 10 min at 37 °C, and then centrifuged at 1000 g for 10 min at room temperature (RT). The pellet was resuspended, washed twice in 5 ml DMEM containing 10% heat-inactivated fetal bovine serum (FBS; Invitrogen) and centrifuged for 10 min at 1000 g at RT. The final pellet was resuspended in 2 ml DMEM/10% FBS, after which the cells were plated in the same medium on a poly-L-lysine-coated culture flask (75 cm², 12 \times 10⁶ cell/flask) and cultured for a number of days in vitro (DIV) at 37 °C in a humidified air atmosphere supplemented with 5% CO₂, in one or other of the following ways: 1) in poly-L-lysine-coated coverslips (15 \times 15 mm; 2 \times 10⁵ cells/coverslip) for immunocytochemistry; 2) in poly-L-lysine-coated Petri dishes (60 mm \times 15 mm; 4 \times 10⁵ cells/dish) for Western blot and enzyme-linked immunosorbent assays (ELISA); or 3) in a poly-L-lysine-coated culture flask (75 cm², 12 \times 10⁶ cells/flask) for the subsequent generation of pure microglial cell cultures.

Secondary microglial cells were subcloned from mixed primary cultures (DIV7) maintained in a poly-L-lysine-coated culture flask (75 cm², 12 \times 10⁶ cells/flask) by shaking the cultures at 100 rpm in a platform shaker for 30 min at 37 °C. Microglia from the supernatant were collected by centrifugation at 3000 g for 8 min at RT and resuspended in 2 ml DMEM/10% FBS. The cells were seeded at a density of 4 \times 10⁵ cells/Petri dish for Western blots or 2 \times 10⁵ cells/coverslip/Petri dish for immunocytochemistry or phagocytosis assays, and cultured in DMEM in a humidified atmosphere supplemented with 5% CO₂ at 37 °C. The medium was changed on the first day after seeding (subDIV1).

2.4. Cell culture treatments

The aspirin concentrations used in our study are comparable to the therapeutic plasma salicylate levels for pain relief, inhibition of platelet aggregation or anti-inflammation (Higgs et al., 1987). On the fourth day of subcloning (subDIV4), DMEM was replaced and the expanded pure microglial cells were treated with either bacterial lipopolysaccharide (LPS; 20 ng/ml final conc., dissolved in DMEM; Sigma, St. Louis, MO, USA) or aspirin (0.1 mM (low) and 1 mM (high) final conc., from \geq 99% acetylsalicylic acid dissolved in sterile, distilled water; Sigma) alone, or with a combination of LPS + aspirin, and the effects were compared in a variety of morphological and functional tests. LPS treatment served as an immunochallenge. Six types of treatment regimens were used: 1) control (unchallenged and untreated) cultures; 2) LPS-challenged cultures received 20 ng/ml LPS; 3) aspirin-treated cultures received 0.1 mM aspirin; 4) aspirin-treated cultures received 1 mM aspirin; 5) LPS-challenged + aspirin-treated cultures were challenged with 20 ng/ml LPS and treated with 0.1 mM aspirin; 6) LPS-challenged + aspirin-treated cultures were challenged with 20 ng/ml LPS and received

1 mM aspirin. Depending on the experiments, the treatments lasted for 6 or 24 h at 37 °C.

2.5. Immunocytochemistry

Pure secondary microglial cultures treated with different treatment regimens were fixed on coverslips with 4% formaldehyde for 5 min and rinsed with 0.05 M phosphate-buffered saline (PBS) for 2 × 5 min. After permeabilization and blocking of the nonspecific sites in 0.05 M PBS solution containing 5% normal goat serum (Sigma), 1% heat-inactivated bovine serum albumin (Sigma) and 0.05% Triton X-100 for 30 min at 37 °C, the cells on the coverslips were incubated overnight in a humidified chamber at 4 °C with rabbit anti-Iba1 polyclonal antibody (1:500 final dilution; Wako, Japan) in the above solution (Kata et al., 2016). The cultured cells were washed for 4 × 10 min at RT in 0.05 M PBS, and then incubated with the Alexa Fluor 568 fluorochrome-conjugated goat anti-rabbit antibody (1:1000 final dilution; Invitrogen) in the dark for 3 h at RT. The cells on the coverslip were washed for 4 × 10 min in 0.05 M PBS at RT, and the nuclei were stained in 0.05 M PBS solution containing 1 mg/ml polyvinylpyrrolidone and 0.5 µl/ml Hoechst 33258 dye (Sigma). The coverslips were rinsed in distilled water for 5 min, air-dried and mounted on microscope slides in Vectashield mounting medium (Vector Laboratories, Burlingame, CA, USA).

2.6. Western blot analysis

Cultured microglial cells (subDIV4) were collected through use of a rubber policeman, homogenized in 50 mM Tris-HCl (pH 7.5) containing 150 mM NaCl, 0.1% Nonidet P40, 0.1% cholic acid, 2 µg/ml leupeptin, 1 µg/ml pepstatin, 2 mM phenylmethylsulfonyl fluoride and 2 mM EDTA, and centrifuged at 10,000g for 10 min (Kata et al., 2016; Szabo and Gulya, 2013). The pellet was discarded and the protein concentration of the supernatant was determined (Lowry et al., 1951). For the Western blot analyses, 5–10 µg of protein was separated on a sodium dodecyl sulfate/polyacrylamide gel (4%/10% stacking gel/resolving gel), transferred onto Hybond-ECL nitrocellulose membrane (Amersham Biosciences, Little Chalfont, Buckinghamshire, England), blocked for 1 h in 5% nonfat dry milk in Tris-buffered saline (TBS) containing 0.1% Tween 20, and incubated overnight with either a rabbit anti-Iba1 polyclonal antibody (1:1000 final dilution; Wako) or a mouse anti-GAPDH monoclonal antibody (clone GAPDH-71.1; 1:20,000 final dilution; Sigma). The membranes were rinsed 5 times in 0.1% TBS-Tween 20 and incubated for 1 h with the peroxidase-conjugated goat anti-rabbit secondary antibody (1:2000 final dilution; Invitrogen) for Iba1 or with the peroxidase-conjugated rabbit anti-mouse secondary antibody (1:2000 final dilution; Sigma) for GAPDH Western blots, then washed 5 times as before. The enhanced chemiluminescence method (ECL Plus Western blotting detection reagents; Amersham Biosciences) was used to reveal immunoreactive bands according to the manufacturer's protocol.

2.7. In vitro phagocytosis

The fluid-phase phagocytic capabilities of the control and variously treated pure microglial cell cultures were determined via the uptake of fluorescent microspheres (carboxylate-modified polystyrene beads, fluorescent yellow-green ($\lambda_{\text{ex}} \sim 470$ nm; $\lambda_{\text{em}} \sim 505$ nm), aqueous suspension, 2.0 µm mean particle size; L4530, Sigma) as we described previously (Szabo and Gulya, 2013; Szabo et al., 2016). Unchallenged (control) and LPS-challenged pure microglial cell cultures (subDIV4) with or without aspirin treatments

were tested for 24 h. At the end of the treatment period, 1 µl of a 2.5% aqueous suspension of fluorescent microspheres per ml was added to the culture, which was then incubated for 60 min at 37 °C. The cells were next washed 5 times with 2 ml of PBS to remove dish- or cell surface-bound residual fluorescent microspheres, and fixed with 4% formaldehyde in PBS. In another setup, we also determined the number of cell membrane-associated but not phagocytosed beads. Such negative controls were treated as above with the exception that microglial cultures with beads were incubated for 60 min at 4 °C. At this temperature, the number of beads associated with cell surface averaged less than 1 bead/100 Iba1-labeled cells, thus the phagocytosis was not considered significant. For measurement of the phagocytotic activity, cells labelled with phagocytosed microbeads and processed for Iba1 immunocytochemistry were counted in 20 random fields in each treatment group (mean ± SEM) under a 20× or 40× objective. Statistically significant differences were determined by two-way ANOVA.

2.8. Determination of IL-1 β , IL-10 and TNF- α

Our protocols for ELISA assays were described previously (Kata et al., 2016). Briefly, the supernatants were collected from each treatment and stored at –20 °C. Concentrations of IL-1 β , IL-10 and TNF- α were measured with rat-specific ELISA kits (eBioscience, Vienna, Austria). The sensitivity of IL-1 β (Cat# BMS630), IL-10 (Cat# BMS629) and TNF- α (Cat#BMS622) assays was 4 pg/ml, 1.5 pg/ml and 11 pg/ml, respectively. As stated by the manufacturer, the overall intra- and inter-assay coefficients of variation were <10% in both cases for IL-1 β and TNF- α , and <5% in both cases for IL-10.

2.9. RNA isolation

Total RNA from control and treated pure microglial cells was purified as described previously (Fabian et al., 2011); columns and wash buffer were from Bioneer (Viral RNA extraction kit; Daejeon, South Korea). Briefly, cells were washed with PBS, incubated in lysis buffer (RA1; Macherey-Nagel, Düren, Germany), then collected and mixed with 70% ethanol in RNase-free water (Bioneer). The mixture was transferred through columns (Bioneer) and washed with 350 µl 80% ethanol in diethylpyrocarbonate-treated water, and then with 600 µl and 300 µl W2 wash buffer (Bioneer). Total RNA was eluted in 50 µl RNase free-water. One µl RNase inhibitor (Applied Biosystems, Foster City, CA, USA) was added to the samples. The quality and quantity of the isolated RNA was measured with NanoDrop1000 Version 3.8.1. (Thermo Scientific, Budapest, Hungary).

2.10. RNA expression

Reverse transcription from 3 µg total RNA in 30 µl was performed with the High Capacity cDNA Archive Kit (Applied Biosystems) according to the manufacturer's protocol as described previously (Kata et al., 2016). cDNA was diluted to 80 µl with nuclease-free water. The instrumentation included the Bravo automatic liquid handling system (Agilent Technologies, Inc., Santa Clara, CA, USA) for polymerase chain reaction (PCR) assay preparation and a LightCycler 1536 System (Roche Diagnostics Corp., Indianapolis, IN, USA) or a Light Cycler Nano Instrument (Roche) for cycling (Woudstra et al., 2013). The expression of 116 inflammation-related genes, together with that of 6 control genes (see below), was measured with Universal Probe Library assays using intron-spanning gene-specific primers (Rat Immune Panel; Avidin Ltd., Szeged, Hungary, www.avidinbiotech.com/custom-pathways/) and the LightCycler 1536 DNA Probe Master kit (Roche). Moreover, the expression of certain

phagocytosis-related genes such as the integrin associated protein or cluster of differentiation 47 (*Cd47*, NM_019195.2), the engulfment or cell motility protein (*Elmo1*, NM_001108415.1), the scavenger receptor class B member 1 (*Scarb1*, NM_031541.1), the plasminogen activator inhibitor-1 (*Serpine1*, NM_012620.1), the signal-regulatory protein α (*Sirpa*, NM_013016.2) and the vesicle-associated membrane protein 7 (*Vamp7*, NM_053531.1) were also analyzed by the Light Cycler Nano Instrument. For the 1536 System, each 2 μ l PCR reaction contained 8 ng cDNA, 0.4 μ l LightCycler DNA Probes Master (5x), the corresponding primer set and UPL probe and the Setup Control (Kata et al., 2016). The PCR cycling protocol was as follows: enzyme activation at 95 °C for 60 s, 50 cycles of denaturation at 95 °C for 0 s, and annealing and extension at 60 °C for 30 s. For the Nano Instrument, each 20 μ l PCR reaction contained 20 ng cDNA, 10 μ l Lightcycler DNA Probes Master (5x), the corresponding primer set and UPL probe and the Setup Control. The PCR protocol was as follows: enzyme activation at 95 °C for 10 min, 50 cycles of denaturation at 95 °C for 15 s, and annealing and extension at 60 °C for 30 s. Gene expression was normalized to the average values of clathrin, heavy chain (*Cltc*, NM_019299.1), *Gapdh* (M17701.1), glucuronidase, beta (*Gusb*, NM_017015.2), hypoxanthine phosphoribosyl-transferase 1 (*Hprt1*, NM_012583.2), phosphoglycerate kinase 1 (*Pgk1*, NM_053291.3), and tubulin, beta 5 class I (*Tubb5*, NM_173102.2) expression as endogenous controls and expressed relative to the unchallenged controls by using the $2^{-\Delta\Delta Ct}$ method.

A total of 122 gene-specific assays were run on 3 independent samples from each condition. Student's *t*-test and two-way ANOVA were applied for the analysis of significance where $p < 0.05$ was considered significant. Gene expression was analyzed by GraphPad Prism 6 (GraphPad Software, Inc., La Jolla, CA, USA). For hierarchical cluster analysis and visualization, the Hierarchical Clustering Explorer (v3.0) software was used (Human-Computer Interaction Lab., University of Maryland, MD, USA; publicly available at <http://www.cs.umd.edu/hcil/multi-cluster/hce3.html>). The complete linkage clustering method was applied with Euclidean distance metric.

2.11. Image analysis and statistics

Digital images were captured by a Nikon Microphot-FXA epifluorescent microscope (Nikon Corp., Tokyo, Japan), using a Spot RT Color CCD camera and the Spot RT software (Spot RT/ke Diagnostic Instruments, Sterling Heights, MI, USA). For the determination of microglial cell purity, Hoechst 33258-labelled cell nuclei that belonged to Iba1-immunopositive cells were counted on coverslip-cultured samples. For each culture, 50–100 randomly selected microscope fields were analyzed. In every case, the cultures had, on average, at least 99 Iba1-positive somata for 100 Hoechst 33258-labelled cell nuclei (>99% purity for microglial cells). Phagocytosed microspheres on 20 randomly sampled microscope fields from 3 coverslips for each treatment regimen were counted with the use of the computer program ImageJ (version 1.47; <http://rsb.info.nih.gov/ij>). For the measurement of area (μm^2), perimeter (μm) and transformation index (TI), Iba1-immunoreactive microglial cell images were converted into binary replicas by using thresholding procedures implemented by ImageJ and Adobe Photoshop CS5.1 software (Adobe Systems, Inc., San Jose, CA, USA) as we published earlier (Kata et al., 2016; Szabo and Gulya, 2013; Szabó et al., 2016). TI was determined (Fujita et al., 1996) according to the following formula: $[\text{perimeter of cell } (\mu\text{m})]^2 / 4\pi [\text{cell area } (\mu\text{m}^2)]$. Color correction and cropping of the light microscopic images were performed when photomicrographs were made for publication and assembled for a panel. Gray-scale digital images of the immunoblots were acquired by scanning the autoradiographic films with a desktop scanner and processed at identi-

cal settings to allow comparisons of the Western blots from different samples. The bands were analyzed through the use of ImageJ. The immunoreactive densities of equally loaded lanes were quantified, and all samples were normalized to the internal GAPDH load controls.

All statistical comparisons were made by using R 3.1.0 for Windows (The R Foundation for Statistical Computing; Vienna University of Economics and Business, Vienna, Austria). Results were analyzed with two-way ANOVA, and the Bonferroni correction was used to establish significance between groups. Values were presented as means \pm SEM; $p < 0.05$ was considered significant; *, ** and *** denote $p < 0.05$, $p < 0.01$ and $p < 0.001$, respectively.

3. Results

3.1. Aspirin regulates microglial morphology in vitro

Microglial morphology was analyzed on pure microglial cultures 4 days after seeding (subDIV4). Immunocytochemistry routinely performed on these secondary cultures consistently detected a >99% incidence of the Iba1 immunopositive microglial cells for the Hoechst 33258 dye-labeled cell nuclei (Fig. 1). The morphological changes elicited by aspirin treatments in unchallenged and LPS-challenged pure microglia cultures (Fig. 2A–F) were quantitatively analyzed on binary silhouettes of individual microglial cells (Fig. 3A–I). Iba1 protein expression was also monitored during treatments (Fig. 2G). The amount of Iba1 immunoreactivity was significantly increased in the aspirin-treated (low: 2.09 ± 0.17 ; high: 2.22 ± 0.26) and the LPS-challenged + aspirin-treated groups (low: 2.30 ± 0.29 ; high: 2.49 ± 0.32) as compared to LPS-challenged values (1.35 ± 0.14 ; Fig. 2G). Most of the unchallenged and untreated (control) microglia displayed slightly amoeboid/ramified morphology with $TI = 6.44 \pm 1.33$; they typically had only a few short processes (Figs. 2A, 3A). LPS challenge resulted in a significant change in morphology (Figs. 2B, 3B, G–I) as amoeboid morphologies with low TI values (1.37 ± 0.03) were more typical; LPS challenge decreased area, perimeter and TI values (by 48%, 65% and 78%, respectively) as compared to the controls. When LPS-challenged cells were treated with aspirin, significantly enlarged and more ramified cells were seen as compared to the values of the LPS-challenged cells (Figs. 2B, E, F, 3B, E–I) indicating that aspirin was able to reverse the morpho-

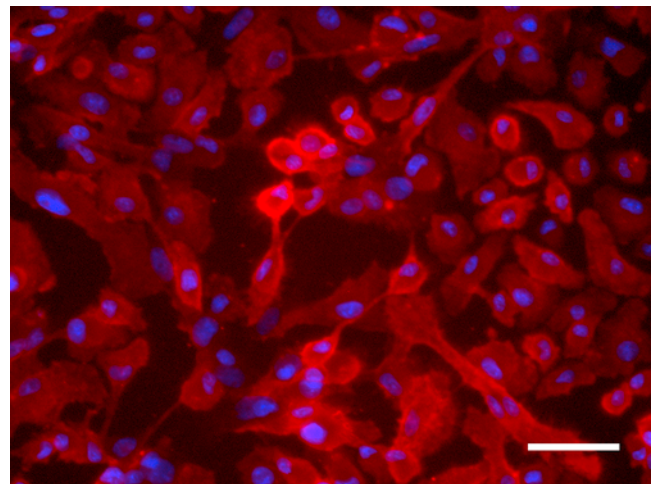


Fig. 1. Localization of Iba1 immunoreactive microglial cells in a pure secondary microglial culture (subDIV4). The typical purity of the culture is >99%; in this representative photomicrograph of a field of view the purity is 100% since every Hoechst 33258-labeled cell nuclei (blue) is surrounded by Iba1 immunopositive cytoplasm (red). Scale bar: 50 μm .

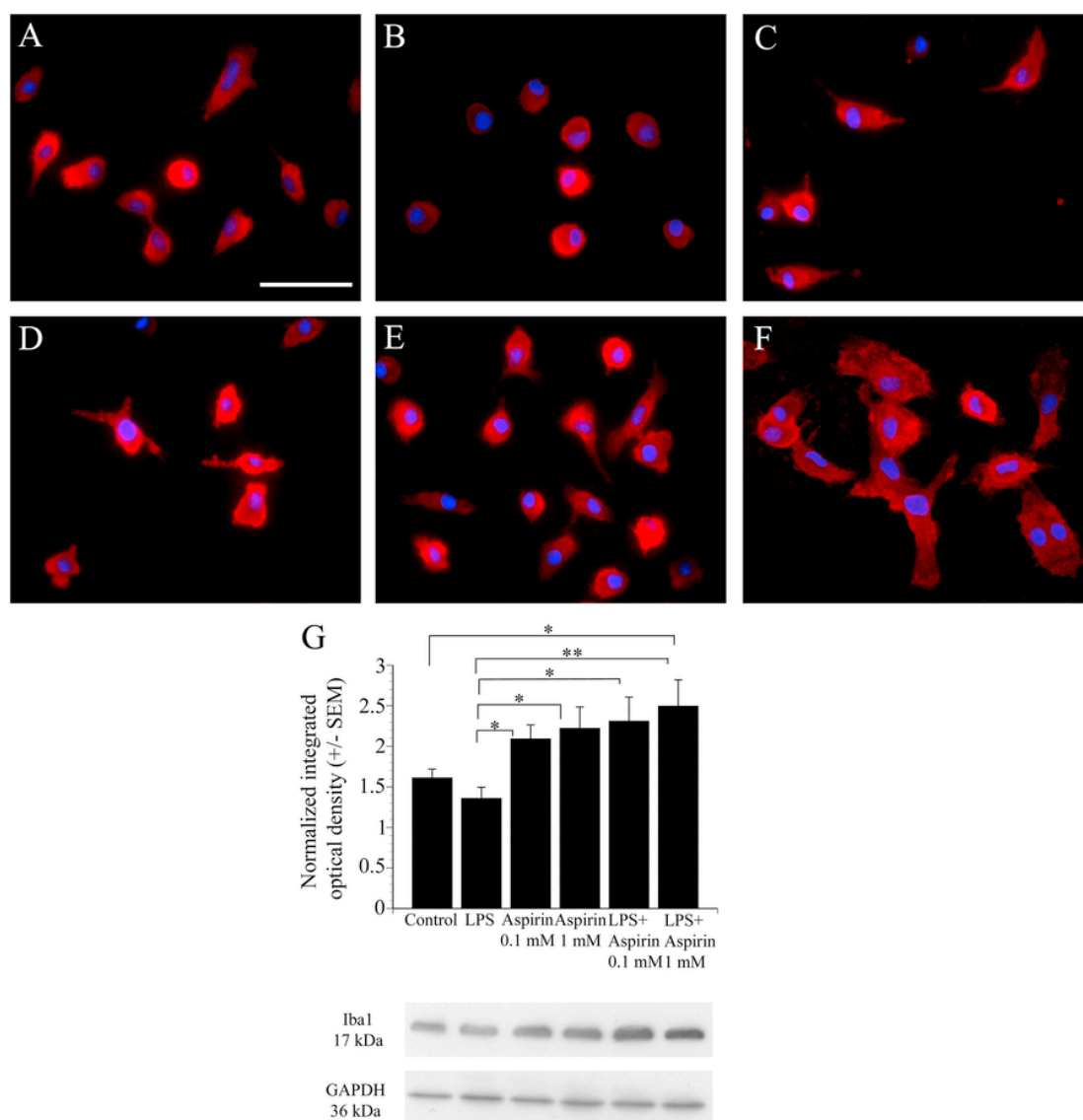


Fig. 2. Aspirin affects microglial morphology and Iba1 immunoreactivity in pure microglial cells. Pure microglial cell cultures (subDIV4) were maintained as described in the Methods section. (A–F) Representative fluorescent immunocytochemical pictures demonstrate the typical cellular distribution of Iba1 immunoreactivity (red) in A) control (unchallenged and untreated), B) LPS-challenged, C) 0.1 mM aspirin treated, D) 1 mM aspirin treated, E) LPS-challenged + 0.1 mM aspirin-treated and F) LPS-challenged + 1 mM aspirin-treated microglial cells. Hoechst 33258-labeled cell nuclei are shown in blue. Scale bar in A (for all pictures): 50 μm. G, Quantitative Western blot analysis of Iba1 and GAPDH immunoreactivities in pure microglial cell cultures. Protein samples from the cultures were separated by gel electrophoresis, transferred to nitrocellulose membranes and probed with either the Iba1 or the GAPDH antibody. Gray scale digital images of the immunoblots were acquired by scanning the autoradiographic films with a desktop scanner. The images were scanned and processed at identical settings to allow comparisons between the Western blots from different samples. Error bars indicate integrated optical density values (means ± SEM) normalized to the internal standard GAPDH. Representative Western blot pictures are beside the graph. Data were analyzed with two-way analysis of variance (ANOVA). **p* < 0.05, ***p* < 0.01.

logical changes induced by LPS-challenge. Both aspirin doses increased the TI values to the control level (low: 5.73 ± 0.56 ; high: 7.36 ± 0.61) and induced ramified morphology with thick processes and microspikes (Figs. 2E, F; 3E, F).

3.2. Aspirin inhibits increases in LPS-induced phagocytosis by microglia

Unchallenged and untreated microglia exhibited a low level of fluid-phase phagocytosis engulfing only 3.63 ± 0.21 beads per cell (*n* = 60; Fig. 4A, G). As expected, microglial activation was associated with robust phagocytic activity as LPS challenge increased phagocytosis significantly to about 350% of the control level (12.87 ± 0.65 ; *n* = 58; Fig. 4A, B, G). Aspirin alone did not affect the phagocytosis appreciably (Fig. 4C, D, G), as the number of

phagocytosed microbeads remained low in both cases (low: 2.72 ± 1.19 , high: 3.30 ± 0.22 ; *n* = 60). However, aspirin inhibited phagocytosis significantly in LPS-challenged microglia (Fig. 4E, F, G). Aspirin (0.1 mM or 1 mM) decreased phagocytosis dose-dependently by about 30% (9.20 ± 0.59 ; *n* = 55; Fig. 4E, G) and 70% (3.86 ± 0.23 microbeads per cell; *n* = 67; Fig. 4F, G), respectively, as compared to LPS treatment.

3.3. Aspirin strongly influences both pro- and anti-inflammatory cytokine levels

Activated microglia express several inflammatory cytokines. When the basal levels of the pro-inflammatory cytokines IL-1β and TNF-α and the anti-inflammatory cytokine IL-10 in unchallenged mi-

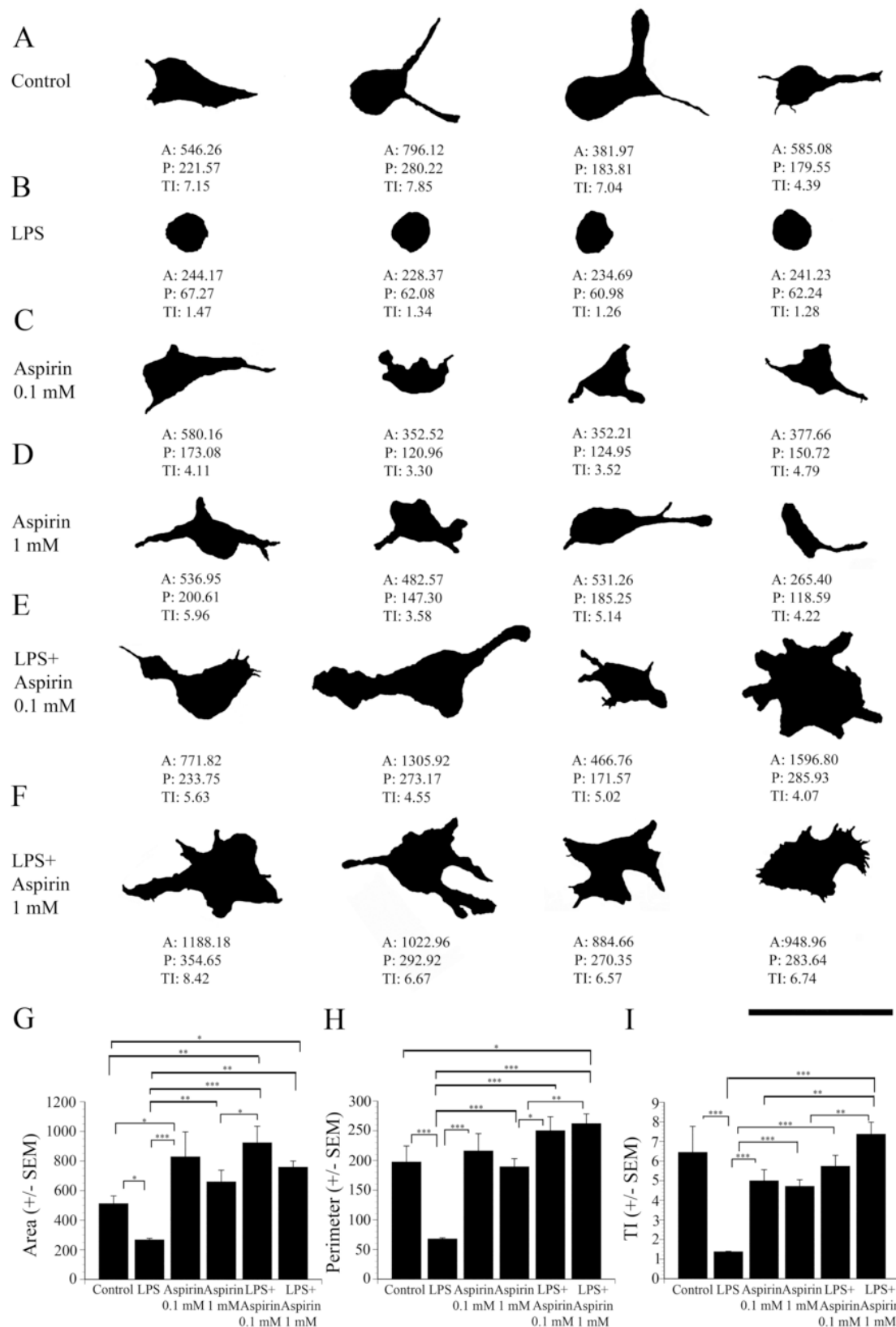


Fig. 3. Quantitative analysis of microglial morphology in pure microglial cell cultures after different treatments. Pure microglial cell cultures (subDIV4) were maintained as described in the Methods section. (A–F) Iba1-positive microglial cells from pure microglial cultures (subDIV4) were photographed, the pictures were digitized and the morphological characteristics were quantitatively analyzed on binary silhouettes of unchallenged (A), LPS-challenged (B), 0.1 mM aspirin treated (C), 1 mM aspirin treated (D), LPS-challenged + 0.1 mM aspirin-treated (E) and LPS-challenged + 1 mM aspirin-treated (F) microglia. Four representative binary silhouettes are shown for each culturing protocol. Scale bar for all silhouettes: 50 μm. Area (G) in μm², perimeter (H) in μm, and TI values (I), calculated as $[\text{perimeter of cell } (\mu\text{m})]^2 / 4\pi [\text{cell area } (\mu\text{m}^2)]$, are

indicated for each digitized cell. Unchallenged and untreated (control) microglia displayed slightly amoeboid/ramified morphology, and neither of the aspirin treatments changed the control values significantly. LPS challenge resulted in amoeboid morphology with the accompanying significant morphometric changes. Both aspirin treatments in LPS-challenged cells resulted in an enlarged and more ramified cell form with a much larger area, perimeter, and TI value as compared with the LPS results (B, E, F, G, H, I). In the control and the aspirin treated cultures, cells displayed short processes or small pseudopodia with a concurrent slight ramification (A, C, D). In the LPS-challenged cultures (B), the microglia became smaller and showed amoeboid morphology. Aspirin treatments in these cells (E, F) resulted in a much larger area (G) but similar perimeter (H) and TI values (I) as compared to the control cultures. (G) Average area (in $\mu\text{m}^2 \pm \text{SD}$) measurements for cultured pure microglial cells. (H) Average perimeter (in $\mu\text{m} \pm \text{SEM}$) measurements for cultured pure microglial cells. (I) Average TI values ($\pm \text{SEM}$) for cultured pure microglial cells. For (G–I), error bars indicate means \pm SEM of 6 replicate measurements from 3 independent culturing. Data were analyzed with two-way analysis of variance (ANOVA). * $p < 0.05$, ** $p < 0.01$, *** $p < 0.001$.

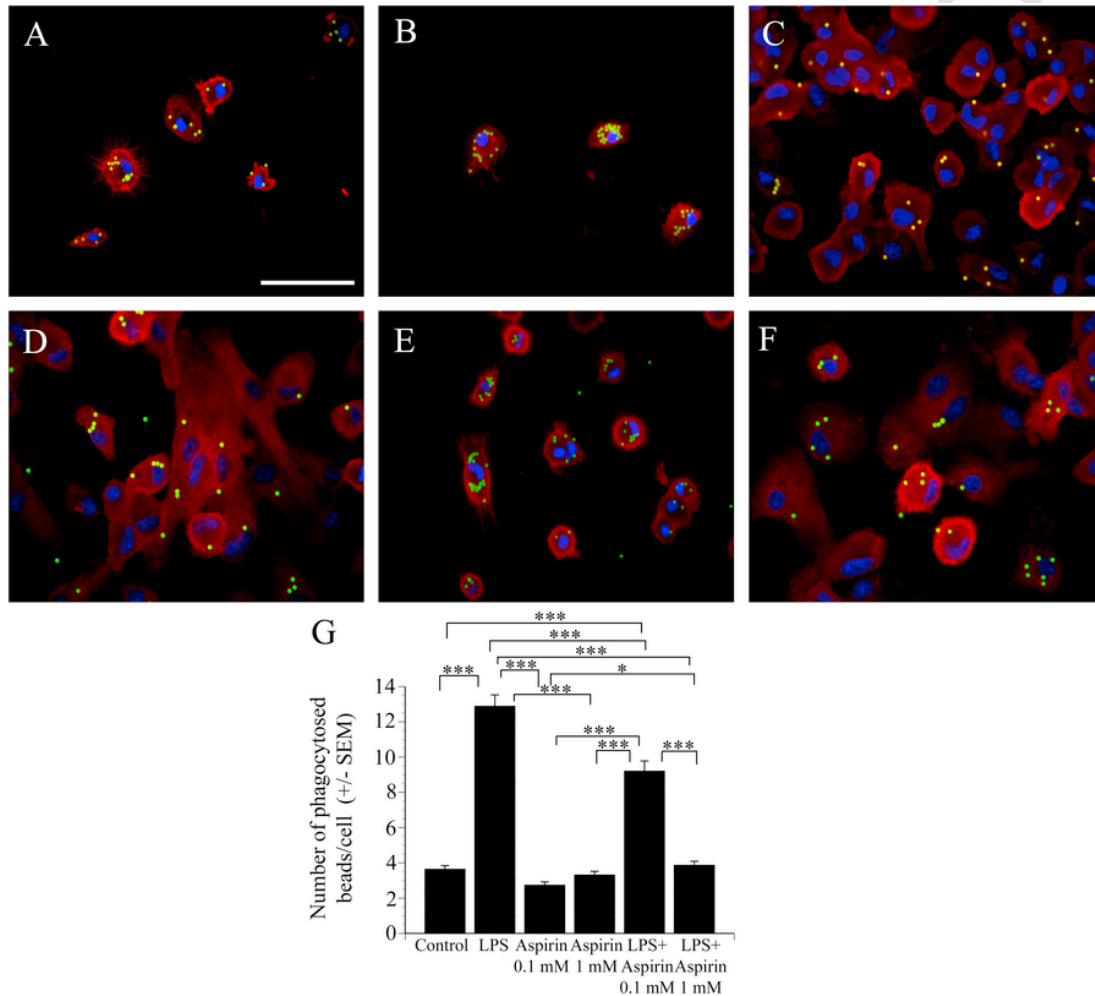


Fig. 4. Aspirin strongly inhibits phagocytosis. Pure microglial cultures (subDIV4) were maintained and treated with fluorescent microbeads (2 μm in diameter) as described in the Methods section. Iba1-specific fluorescent immunocytochemistry (red: microglia; blue: nucleus; green: microspheres) on unchallenged (A), LPS-challenged (B), 0.1 mM aspirin treated (C), 1 mM aspirin treated (D), LPS-challenged + 0.1 mM aspirin-treated (E), and LPS-challenged + 1 mM aspirin-treated (F) microglia. Scale bar in A (for all pictures) = 50 μm . (G) Quantitative analysis of the number of phagocytosed microbeads revealed that LPS dramatically activated phagocytosis, while aspirin significantly decreased this microglial function. Error bars indicate means \pm SEM of 6 replicate measurements from 3 independent culturing. Data were analyzed with two-way ANOVA. * $p < 0.05$, *** $p < 0.001$.

croglia were compared with the levels from aspirin-treated LPS-challenged cells, a unique regulatory pattern emerged (Fig. 5). In the case of IL-1 β the basal level in unchallenged (control) microglia was 22.45 ± 2.87 pg/ml (Fig. 5A). Both aspirin treatments were able to change this level (low: 10.26 ± 2.62 pg/ml; high: 10.19 ± 2.23 pg/ml). As expected, LPS challenge significantly elevated the IL-1 β level to 336.15 ± 51.72 pg/ml. However, co-incubation of LPS and 1 mM aspirin significantly inhibited IL-1 β level by about 50%, to 169.6 ± 19.42 pg/ml, while 0.1 mM aspirin decreased it by about 16.5%, to 280.05 ± 25.0 pg/ml (Fig. 5A).

A similarly strong effect of aspirin was demonstrated on the level of TNF- α , another pro-inflammatory cytokine, in LPS-challenged microglial cells (Fig. 5C, D). Two different treatment periods (6 h and 24 h) were used as the TNF- α production responded quickly to the

LPS challenge. The level of TNF- α in the control group could not be detected, but its level quickly rose, to 1187 ± 94.12 pg/ml in the LPS-challenged cells after 6 h, and the level was still elevated after 24 h (189.92 ± 27.4 pg/ml). Aspirin treatments (0.1 mM and 1 mM) resulted in small, detectable TNF- α levels for both doses (27.69 ± 2.89 pg/ml and 29.23 ± 1.73 pg/ml after 6 h, while 8.24 ± 1.81 pg/ml and 9.09 ± 0.6 pg/ml after 24 h). When aspirin was co-administered to LPS-challenged cells for either 6 h or 24 h, it inhibited the production of TNF- α in both cases, but only the high aspirin concentration decreased the TNF- α level significantly by about 25% (to 844.76 ± 67.11 pg/ml after 6 h) and 50% (96.79 ± 6.02 pg/ml after 24 h) (Fig. 5C, D).

Aspirin also affected the production of IL-10 (Fig. 5B). LPS challenge increased the basal IL-10 level (24.98 ± 3.47 pg/ml) signifi-

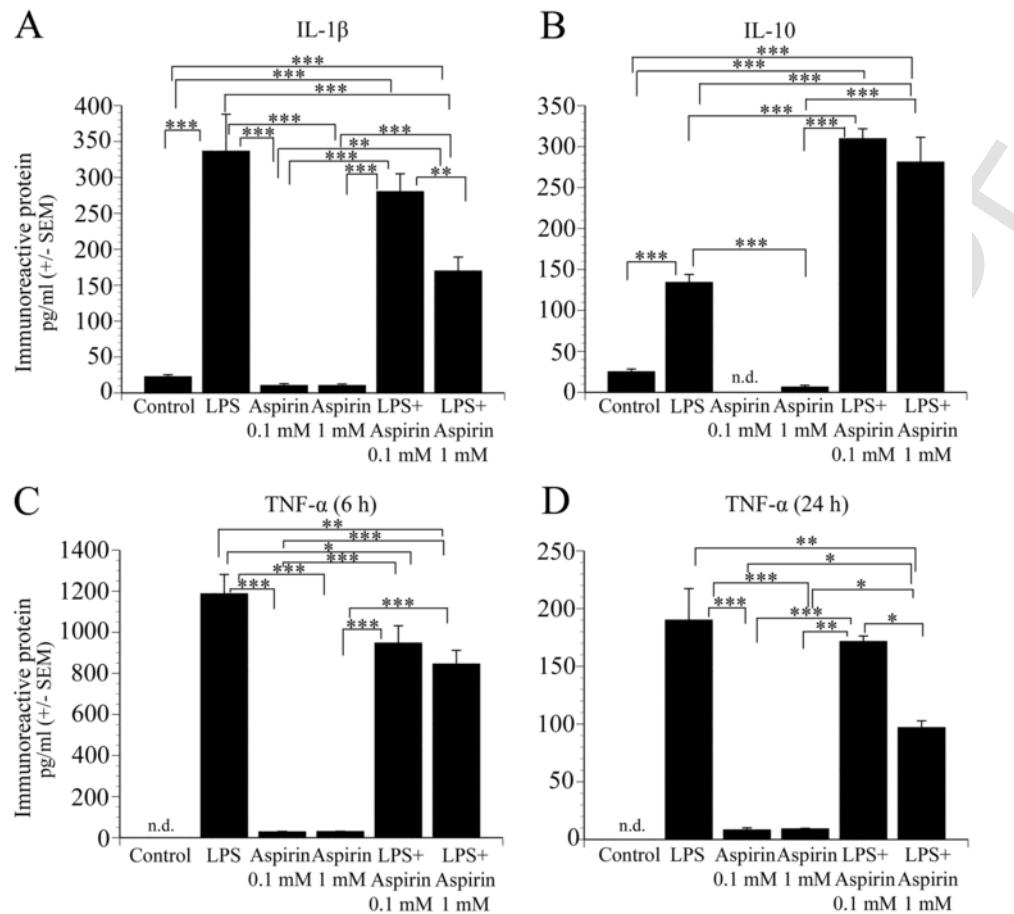


Fig. 5. Aspirin reduces the pro-inflammatory cytokine IL-1 β and TNF- α levels and increases the anti-inflammatory IL-10 production. Pure microglial cultures (subDIV4) were maintained and treated as described in the Methods section. After treatments for 6 or 24 h, immunoreactive protein levels (pg/ml \pm SD) for IL-1 β (A), TNF- α (C, D) and IL-10 (B) were detected by ELISA. TNF- α production was measured after 6 h and 24 h (C, D). As expected, pro-inflammatory cytokine production was significantly increased in the LPS-challenged cells (A, C, D). Aspirin (1 mM) inhibited this effect for both IL-1 β (A) and TNF- α (C, D). The level of the anti-inflammatory cytokine IL-10 was measured after 24 h (B). Aspirin (0.1 mM and 1 mM) did not increase the level of IL-10 in unchallenged cultures but strongly increased it in LPS-challenged cultures (to 115% and 130% of the control values, respectively) (B). Error bars indicate means \pm SEM of 6 replicate measurements from 3 independent culturing. Data were analyzed with two-way ANOVA. n.d. = not detected. * p < 0.05; ** p < 0.01, *** p < 0.001.

cantly (134.0 ± 9.9 pg/ml). When aspirin was co-administered with LPS, the IL-10 protein expression was further increased as compared to the LPS challenged value (low: 309.47 ± 12.29 pg/ml, high: 280.95 ± 30.3 pg/ml). Aspirin administered alone to unchallenged cells had no significant effects on the microglial IL-10 levels.

3.4. Aspirin affects the expression of inflammation-related genes

We analyzed the effects of aspirin on the expression of 116 inflammation-related genes in unchallenged and LPS-challenged pure microglial cells with or without aspirin treatment. The hierarchical cluster analysis of 46 such genes is summarized in Fig. 6, and those with the most noteworthy and significant expression changes in response to the treatments are listed in Table 1. The genes upregulated by the LPS challenge included those coding for chemokine ligands 1, 2, 4, 5, 7, 9 (*Cxcl1* = 84.30-fold, *Ccl2* = 16.95-fold, *Ccl4* = 9.80-fold, *Ccl5* = 38.26-fold, *Cxcl5* = 33.19, *Cxcl9* = 73.44-fold,) IL-1 β (*Il1 β* = 24.70-fold), IL-6 receptor (*Il6* = 7.57-fold) and nitric oxide synthase (*Nos2* = 369.22-fold). Only a few genes were downregulated by the LPS challenge, including *Tgfb2* (−1.73-fold), *Sirpa* (−2.42-fold), *Tlr5* (−3.42-fold) and *Myl2* (−4.46-fold).

The levels of expression of some inflammation-related genes strongly downregulated by aspirin are highlighted in Fig. 7. Aspirin treatment in unchallenged cells affected fewer, but similarly impor-

tant microglial genes involved in pro- and anti-inflammatory processes. In unchallenged cells the genes downregulated by aspirin (0.1 mM or 1 mM) included the interleukin 1 receptor antagonist (*Il1rn* = −1.37-fold; −1.44-fold), the anti-inflammatory DNA damage inducible transcript 3 (*Ddit* = −1.43-fold, and −1.79-fold), the pro-inflammatory IL6 (*Il6* = −1.23-fold; −1.55-fold) and the pro-inflammatory chemokine (C-C motif) receptor 1 (*Ccr1* = −1.42-fold, −1.44-fold). When aspirin was co-administered to LPS-challenged cultures (Table 1), a more complex picture emerged. Some of the LPS-upregulated genes were inhibited by aspirin, as seen in the case of *Ccr1* (Fig. 7A), *Il1rn* (Fig. 7B) or *Nos2* (Fig. 7D). *Kng1* (Fig. 7C), a pro-inflammatory gene was downregulated by both aspirin treatments following LPS challenge. Other genes were further upregulated by aspirin when applied to LPS-challenged microglia such as *Ccl5* or *Csf2*, or the pro-inflammatory *Ccl4* (Table 1).

4. Discussion

Microglial activation results in profound morphological, functional and gene expression changes that activate both pro- and anti-inflammatory mechanisms (Kroner et al., 2014) that in turn not only protect the nervous tissue but can elicit chronic inflammation that could lead to the development of neuropathological conditions (Jones and Thomsen, 2013; Long-Smith et al., 2009). Aspirin, one of

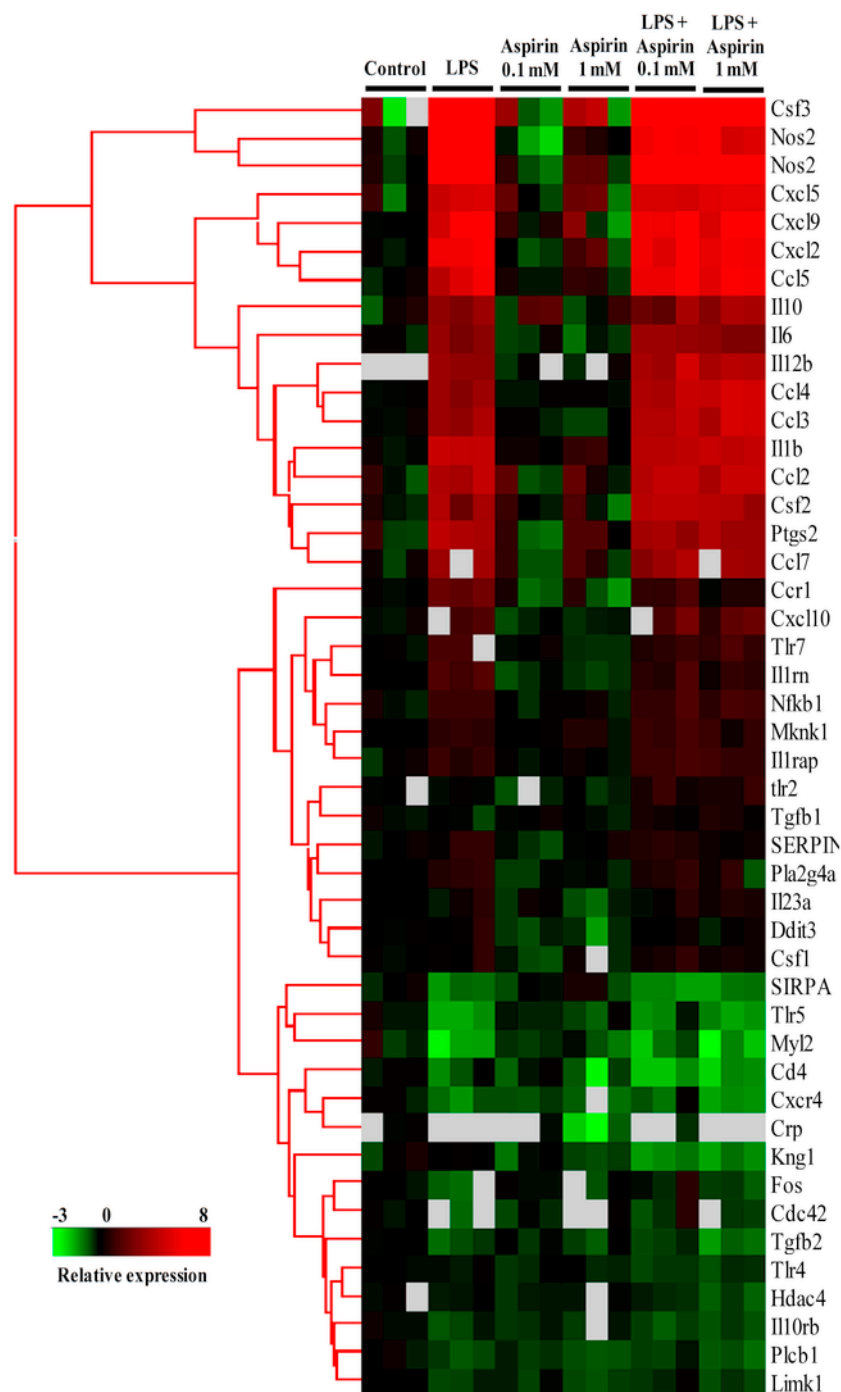


Fig. 6. Hierarchical cluster analysis of inflammation related genes. Pure microglial cultures (subDIV4) were maintained and treated as described in the Methods section. Cells (subDIV4) were cultured with or without LPS for 24 h in the presence or absence of aspirin. Aspirin: 0.1 mM or 1 mM; LPS: 20 ng/ml; LPS + aspirin: LPS (20 ng/ml) + aspirin (0.1 mM or 1 mM). For hierarchical cluster analysis and visualization, the Hierarchical Clustering Explorer (v3.0) software was used. The complete linkage clustering method was applied with Euclidean distance metric. The heat map depicts expression values relative to control samples on a log₂ scale (overexpression: red, repression: green and no change: black). Missing values are indicated in grey.

the most commonly used drugs, is noted for its strong anti-inflammatory actions. It acetylates the active site of the enzyme COX, and irreversibly blocks the conversion of arachidonic acid to prostanoïd (Vane and Botting, 2003). Apart from its classical beneficial role in cardiovascular diseases, however, aspirin may be beneficial in neuroinflammation-related disorders given that appropriate target cells, the microglia, are present in the nervous tissue.

In the present study we investigated the complex pleiotropic, beneficial effects of aspirin on the modulation of inflammation by quan-

titatively analyzing numerous morphological, functional (phagocytosis, cytokine release) and gene expression parameters in pure, secondary microglial cultures. As our cultures were >99% pure, they provided a unique opportunity to study these functional and expression parameters without the significant influence of any contaminating cell types. This is a crucial factor when, for example, the levels of secreted pro- and anti-inflammatory peptides or gene expression levels are measured, as other cell types in the CNS are also capable of expressing such genes (Gruol et al., 2014).

Table 1
Differentially expressed transcripts in aspirin-treated pure microglial cells with or without LPS challenge.

Gene	Name NCBI Reference Sequence ID RGD ID	LPS	Aspirin 0.1 mM	Aspirin 1 mM	LPS + Aspirin 0.1 mM	LPS + Aspirin 1 mM
Ccl2	Chemokine ligand 2 NM_031530.1 RGD ID 3645	+ 16.95	+ 1.14	+ 1.70	+ 20.92	+ 21.35
Ccl3	Chemokine ligand 3 NM_013025.2 RGD ID 3647	+ 11.33	− 1.01	− 1.31	+ 18.41	+ 24.79
Ccl4	Chemokine ligand 4 NM_053858.1 RGD ID 620441	+ 9.80	− 1.05	+ 1.06	+ 17.68	+ 29.80
Ccl5	Chemokine ligand 5 NM_031116.3 RGD ID 69069	+ 38.26	+ 1.02	+ 1.35	+ 84.12	+ 62.69
Ccl7	Chemokine ligand 7 NM_001007612.1 RGD ID 1359152	+ 11.48	− 1.14	+ 1.50	+ 10.82	+ 11.82
Ccr1	Chemokine receptor 1 NM_020542.2 RGD ID 708446	+ 4.51	− 1.42	− 1.44	+ 2.16	+ 1.29
Cxcl1	Chemokine ligand 1 (melanoma growth-stimulating activity, alpha) NM_030845.1 RGD ID 619869	+ 84.30	− 1.51	+ 4.11	+ 43.46	+ 246.78
Cxcl2	Chemokine ligand 2 NM_053647.1 RGD ID 70069	+ 69.02	− 1.37	+ 1.73	+ 63.70	+ 60.07
Cxcl5	Chemokine ligand 5 (Cxcl6) NM_022214.1 RGD ID 708540	+ 33.19	+ 1.41	+ 2.11	+ 34.89	+ 46.37
Cxcl9	Chemokine ligand 9 NM_145672.4 RGD ID 628798	+ 73.44	+ 1.40	+ 1.22	+ 86.22	+ 68.51
Cxcl10	Chemokine ligand 10 NM_139089.1 RGD ID 620209	+ 2.83	− 1.30	− 1.24	+ 3.93	+ 3.21
Csf1	Colony stimulating factor-1 AF515736.1 RGD ID 621063	+ 1.31	− 1.49	− 1.03	+ 1.48	+ 1.30
Csf2	Colony stimulating factor-2 XM_001074265.1(old) NM_053852.1 RGD ID 621065	+ 10.01	+ 1.20	− 1.00	+ 20.66	+ 14.84
Csf3	Colony stimulating factor 3 NM_017104.1 (old), NM_017104.2 RGD ID 2426	+ 305.58	+ 1.23	+ 4.96	+ 282.90	+ 278.46
Ddit3	DNA-damage inducible transcript 3 NM_001109986.1 RGD ID 62391	+ 1.25	− 1.43	− 1.79	+ 1.14	+ 1.04
Il1β	Interleukin 1 beta NM_031512.2 RGD ID 2891	+ 24.70	+ 1.18	+ 1.62	+ 19.88	+ 22.98
Il1rap	Interleukin 1 receptor accessory protein NM_012968.1 RGD ID 2893	+ 1.97	− 1.02	+ 1.06	+ 2.30	+ 2.08
Il1rn	Interleukin 1 receptor antagonist NM_022194.2 RGD ID 621159	+ 2.83	− 1.37	− 1.44	+ 2.02	+ 1.64
Il6	Interleukin 6 M26744.1 RGD ID 2901	+ 7.57	− 1.23	− 1.55	+ 10.82	+ 7.22
Il10	Interleukin 10 NM_012854.2 RGD ID 2886	+ 8.47	+ 2.01	+ 1.04	+ 6.28	+ 11.87
Il18	IFN-gamma-inducing factor IL-18 AY077842.1 RGD ID 2889	+ 2.24	+ 1.63	− 1.29	+ 1.90	+ 2.03
Kngr1	Kininogen 1 NM_012696.2 RGD ID 2980	+ 1.04	− 1.36	− 1.55	− 2.75	− 2.82
Nos2	Nitric oxide synthase U26686.1 RGD ID 3185	+ 369.22	− 1.28	+ 1.78	+ 73.06	+ 51.21

Table 1 (Continued)

Gene	Name NCBI Reference Sequence ID RGD ID	LPS	Aspirin 0.1 mM	Aspirin 1 mM	LPS + Aspirin 0.1 mM	LPS + Aspirin 1 mM
Pla2g4a	Phospholipase A2, group IVA (cytosolic, calcium-dependent) NM_133551.2 RGD ID 67366	+1.73	-1.33	-1.16	+1.66	+1.07
Ptgs2	Prostaglandin endoperoxide synthase 2 NM_017232.3 RGD ID 620349	+16.83	-1.33	+2.14	+11.76	+12.55
Sirpa	The signal-regulatory protein a NM_013016.2 RGD ID 3449	-2.42	-1.22	+1.04	-2.84	-2.63
Tgfb1	Transforming growth factor, beta 1 NM_021578.2 RGD ID 69051	-1.21	+1.05	-1.10	+1.25	+1.34
Tgfb2	Transforming growth factor, beta 2 NM_031131.1 RGD ID 70491	-1.73	-1.15	-1.43	-1.47	-2.41
Tlr2	Toll-like receptor 2 NM_198769.2 RGD ID 735138	-1.03	-1.49	-1.22	+1.63	+1.63
Tlr7	Toll-like receptor 7 EF032637.1 RGD ID 1563357	+2.71	+1.05	-1.36	+1.79	+2.21

The National Center for Biotechnology Information (NCBI) Reference Sequence Database can be retrieved at <http://www.ncbi.nlm.nih.gov/refseq/>. A description of a gene function (RGD ID) can be found in The Rat Genome Database 2015 (Shimoyama et al., 2015) at <http://rgd.mcw.edu/>. Data are expressed as fold-change over the expression of unchallenged and untreated (control) microglial cells. Treatments had different effects on inflammation-related genes. Most of the genes were upregulated by LPS challenge. Aspirin, either alone or in combination with LPS, had differential effects on the expression of certain genes. For example, aspirin treatment downregulated *Tgfb2*, *Kng1* or *Ddit3*, while upregulated other genes such as *Csf3*, *IL10* and *Cxcl1* in unchallenged cells. In LPS-challenged microglia, aspirin inhibited *IL1rn*, *Ccr1*, *Kng1*, *Nos2*, while it had a synergistic effect with LPS on others (e.g. *Cxcl5*, *Ccl4*, *Csf2*).

Aspirin reversed the morphological effects of LPS-induced microglia activation through the increase of the area, perimeter and TI values; the combined treatment with LPS also inhibited this activation through the development of microspikes and inducing the cells to become more ramified. Aspirin also affected the protein synthesis of Iba1, a Ca^{2+} -binding protein that is implicated in actin cytoskeleton remodeling (Ohsawa et al., 2004; Szabo et al., 2016).

LPS is a potent activator of microglial phagocytosis (Kata et al., 2016; Lund et al., 2006; Szabo and Gulya, 2013; Szabo et al., 2016), a crucial factor both in normal and pathologic functions of the CNS (Kettenmann et al., 2011). Alteration of this function could be harmful (Hickman and El Khoury, 2014; Lue et al., 2015). Pro-inflammatory phenotypes are associated with increased phagocytic activity, and blocking of phagocytosis may prevent some forms of inflammatory neurodegeneration; this may be beneficial under different neuropathological conditions (He et al., 2014; Neher et al., 2011). Aspirin inhibited phagocytosis in the LPS-challenged microglia only, while it did not inhibit the basal phagocytotic activity in unchallenged cells, a physiological microglial function crucial for the healthy CNS.

LPS is the main ligand for Toll-like receptor 4 (TLR4). TLR4-induced signalling activates the NF- κ B and MAPK pathways leading to the production of pro-inflammatory cytokines (Fuentes et al., 2016; Korneev et al., 2017). Aspirin inhibits the production of pro-inflammatory cytokines through the suppression of LPS-induced NF- κ B and MAPK pathways in microglial cells (Cianciulli et al., 2016; Sun et al., 2017; Yang et al., 2014). NF- κ B activity is suppressed in the cytoplasm with an inhibitory I κ B protein. The phosphorylation and degradation of I κ B activates NF- κ B and allows to the NF- κ B dimer to translocate to the nucleus and activate target genes such as *IL1 β* and *TNF- α* (Sun et al., 2017; Yang et al., 2014). LPS-stimulated degradation of I κ B could be strongly inhibited by aspirin (Yang et al., 2014). Anti-inflammatory property is mediated by inhibition of several members of MAPK family, e.g., p38 MAPK and ERK1/2 (Jung et al., 2009; Li et al., 2017); according to recent data, aspirin enhances the

inhibition of p38-MAPK and ERK phosphorylation in a microglial cell line (Yang et al., 2014).

Cytokines are important to the normal and pathologic immunomodulatory functions of the microglia. Previous studies demonstrated a strong relationship between cytokines and neurodegenerative diseases (McGeer and McGeer, 2010; Smith et al., 2012). Microglial cells activated by β -amyloid deposits produce toxic mediators and pro-inflammatory cytokines such as IL-1 β , IL-6, TNF- α and iNos that could result in chronic inflammation (Benveniste et al., 2001). High level of IL-1 β could be observed in the vicinity of amyloid plaques of AD patients (McGeer et al., 1993) that could be responsible for plaque accumulation (Akiyama et al., 2000). COX-2 is also increased in AD patients (Akiyama et al., 2000). Recent attempts have been made to ameliorate neuroinflammation through the suppression of microglial activation both in vivo and in vitro (Lim et al., 2015; Wang et al., 2011; Yang et al., 2016). Similarly to the findings for AD, previous studies also suggested that activated microglia could be associated with neuronal loss during PD due to the activation of cytokines, COX-2 and iNos expression (Orr et al., 2002; Phani et al., 2012) or increased levels of TNF- α and nitrogen oxide could cause demyelination in MS (Minagar et al., 2002).

As expected, both pro- (IL-1 β , TNF- α) and anti-inflammatory (IL-10) cytokines were increased in LPS-challenged microglia as compared with the unchallenged cells. Aspirin did not significantly affect the basal cytokine levels in unchallenged cells, but strongly reduced the levels of the pro-inflammatory cytokines IL-1 β and TNF- α when tested in LPS-challenged microglia. Besides the inhibition of these pro-inflammatory agents, aspirin was also able to elevate the anti-inflammatory IL-10 in LPS-challenged cells. Thus, aspirin is remarkably beneficial as it concomitantly decreased pro-inflammation-related cytokines but increased anti-inflammation-related cytokine peptides in immunochallenged cells.

We also demonstrated that aspirin, LPS and their combination had differential effects on the expression of several inflammation-related genes in pure microglial cells. Some of these genes encode

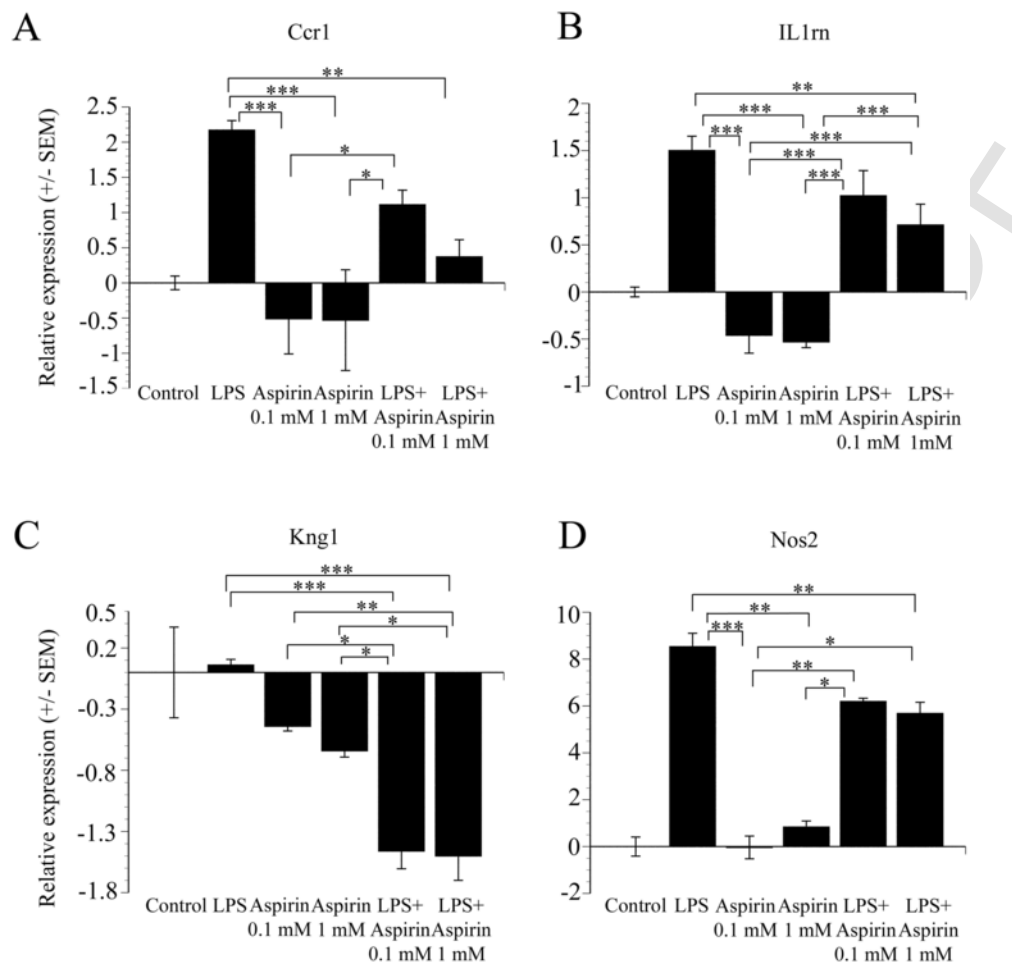


Fig. 7. Relative expression levels of various inflammation-related genes identified by real-time PCR in pure microglia cells. Pure microglial cultures (subDIV4) were maintained and treated as described in the Methods section. The transcription levels of the chemokine receptor 1 (*Ccr1*, A), the interleukin 1 receptor antagonist (*IL1rn*, B), the kininogen 1 (*Kng1*, C) and the nitric oxide synthase (*Nos2*, D) genes are shown in unchallenged (control), LPS-challenged and LPS-challenged + aspirin-treated cells. Except for *Kng1*, these pro-inflammatory genes were upregulated in LPS-challenged cells but were all strongly inhibited by both doses of aspirin. Relative expression levels (on a \log_2 scale) \pm SEM from at least 3 separate experiments are shown for each condition. Data were analyzed with two-way ANOVA. * $p < 0.05$, ** $p < 0.01$, *** $p < 0.001$.

markers that can be related to different microglia subtypes proposed as M1 or M2 phenotypes (Moehle and West, 2015; Orihuela et al., 2016). In this context, for example, the pro-inflammatory *IL1 β* , *IL6*, *IL18*, *Cxcl9*, *TNF- α* , *Cxcl10*, and the chemokine *Ccl5*, genes encode M1 markers, while the anti-inflammatory *IL10* and the pro-inflammatory *Tgfb2* genes are M2 markers (Franco and Fernández-Suárez, 2015). In an LPS-based animal model for Parkinson's disease, a Rho kinase inhibitor reversed the inflammatory M1 to anti-inflammatory M2 microglia, decreased NF- κ B activation, and inhibited IL-12 and TNF- α generation (He et al., 2016), demonstrating a protective effect against LPS-mediated dopaminergic degeneration. It must be noted, however, that the concept of M1/M2 polarization of microglia cells is not universally accepted and needs reassessment (Martinez and Gordon, 2014; Ransohoff, 2016).

Our study demonstrated that several genes were upregulated after LPS challenge, confirming its strong pro-inflammatory effects on microglia. Aspirin, both alone and in combination with LPS, displayed different effects on microglial gene expression. It suppressed the pro-inflammatory effects of LPS on some genes such as *Ccr1* and *Nos2*. It may be of importance that *Ccr1* was previously associated with both MS and experimental autoimmune encephalomyelitis (Rottman et al., 2000) and *Ccr1* protein was localized around the demyelinating plaques (Eltayeb et al., 2007). In our experiment (Fig. 7A) both aspirin doses decreased *Ccr1* expression. Multiple studies suggested

that *Nos2* has a central role in inflammatory reactions and pathogenesis of certain neurodegenerative diseases (Akiyama et al., 2000; Block and Hong, 2005; Galea and Feinstein, 1999). Thus, downregulating *Nos2* expression could be crucial for possible future therapies. We showed that aspirin inhibited *Nos2* expression significantly in LPS-challenged microglia. Kininogen (*Kng1*) is well known for its pro-inflammatory properties (Sharma and Yusof, 1998; Stewart et al., 1997) as it increases the release of inflammatory cytokines and affects the formation of inflammatory exudate and pain (Ueno and Oh-ishi, 2003).

Aspirin also increased the expression of genes that were predominantly anti-inflammatory e.g. *IL10*, *Tgfb1* or *Ccl5*. IL-10 is a well known anti-inflammatory cytokine that limits the production of pro-inflammatory cytokines including IL-1 α and - β , IL-6, IL-12, IL-18, and TNF- α (Couper et al., 2008). Previous studies showed that *Tgfb1* had a protective effect against excitotoxicity and the loss of this protein resulted in neurodegeneration (Brionne et al., 2003). *Ccl5* was also upregulated in our studies. Although *Ccl5* was demonstrated to induce pro-inflammatory mechanisms (Skuljec et al., 2011), it was also shown to ameliorate AD-like pathology by recruiting microglia to β -amyloid deposits (Lee et al., 2012) and to protect against neuronal injury (Campbell et al., 2013; Lim and Mocchetti, 2015). Upregulation of such anti-inflammatory genes in neurodegenerative disorders could be beneficial.

As activated microglia often damage neuronal tissues (Hong et al., 2016; Lui et al., 2016) or cause chronic inflammation by excessive cytokine production combined with high level of phagocytosis, effective inhibition of pro-inflammatory actions by aspirin could be an important prophylactic therapy in preventing neuroinflammation and thus neurodegeneration. Our data indicate that aspirin beneficially regulates microglia, cells that could be targets in treating or preventing neurodegenerative disorders. With respect to the large population who receives aspirin for pain medication or prevention of a number of pathologic conditions, to study the interactions of this drug with the nervous system is even more important. Early observations revealed that NSAIDs could reduce the risk of AD (McGeer et al., 1996; Stewart et al., 1997), a notion that has not been proven in randomised controlled trials assessing the efficacies of aspirin, steroidal and non-steroidal anti-inflammatory drugs in AD (Jaturapatporn et al., 2012). However, we recently demonstrated that rosuvastatin, a new generation 3-hydroxy-3-methylglutaryl coenzyme A reductase inhibitor regulated microglial pro- and anti-inflammatory functions very favorably (Kata et al., 2016). As the effects of aspirin on selected morphological, functional and gene expression characteristics of microglia are very similar to that of rosuvastatin, and these drugs are frequently prescribed together for a number of cardiovascular illnesses, their extended use may have contributed to the lower than expected incidence of AD (Schrijvers et al., 2012). Thus, the individual and combined effects of aspirin and rosuvastatin on microglial functions should be further analyzed for a prophylactic therapeutic approach in neurodegenerative disorders.

5. Conclusion

Aspirin elicits strong responses to microglial functions in LPS-challenged pure microglial cells. The most important pleiotropic beneficial effects of aspirin are 1) the robust inhibition of phagocytosis in activated microglia, 2) the inhibition of synthesis of pro-inflammatory cytokines combined with a very strong stimulation of anti-inflammatory cytokine production, and 3) the beneficial differential expression of a number of inflammation-related genes. Such beneficial regulation of microglial functions could point to the possible benefits of aspirin therapy in certain neuroinflammatory and neurodegenerative diseases.

Author contributions

This study is based on an original idea of KG. Designed the experiments: LH, LGP and KG. Performed the experiments: DK, IF, LZ and LH. Contributed unpublished reagents/analytic tools: LGP. Analyzed the data: DK, IF and LH. Wrote the paper: DK and KG. All authors have read and approved the final manuscript.

Conflict of interests

LGP holds an equity position in Avidin Ltd. that produces a high-throughput QPCR product (Rat immune panel) and has a service using the same kit (<http://avidinbiotech.com/custom-pathways/>).

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lish, or the preparation of the manuscript. At the time of the experiments, D.K. was a Ph.D. student at the University of Szeged.

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