



# The Opening of Connexin 43 Hemichannels Alters Hippocampal Astrocyte Function and Neuronal Survival in Prenatally LPS-Exposed Adult Offspring

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Clinical evidence has revealed that children born from mothers exposed to viral and bacterial pathogens during pregnancy are more likely to suffer various neurological disorders including schizophrenia, autism bipolar disorder, major depression, epilepsy, and cerebral palsy. Despite that most research has centered on the impact of prenatal inflammation in neurons and microglia, the potential modifications of astrocytes and neuron-astrocyte communication have received less scrutiny. Here, we evaluated whether prenatally LPS-exposed offspring display alterations in the opening of astrocyte hemichannels and pannexons in the hippocampus, together with changes in neuroinflammation, intracellular Ca<sup>2+</sup> and nitric oxide (NO) signaling, gliotransmitter release, cell arborization, and neuronal survival. Ethidium uptake recordings revealed that prenatal LPS exposure enhances the opening of astrocyte Cx43 hemichannels and Panx1 channels in the hippocampus of adult offspring mice. This enhanced channel activity occurred by a mechanism involving a microglia-dependent production of IL- $1\beta$ /TNF- $\alpha$  and the stimulation of p38 MAP kinase/iNOS/[Ca<sup>2+</sup>]<sub>i</sub>-mediated signaling and purinergic/glutamatergic pathways. Noteworthy, the activity of Cx43 hemichannels affected the release of glutamate, [Ca<sup>2+</sup>], handling, and morphology of astrocytes, whereas also disturbed neuronal function, including the dendritic arbor and spine density, as well as survival. We speculate that excitotoxic levels of glutamate triggered by the activation of Cx43 hemichannels may contribute to hippocampal neurotoxicity and damage in prenatally LPS-exposed offspring. Therefore, the understanding of how astrocyte-neuron crosstalk is an auspicious avenue toward the development of broad treatments for several neurological disorders observed in children born to women who had a severe infection during gestation.

Keywords: neuroinflammation, hemichannel, connexin, glia, pannexin

# INTRODUCTION

Environmental factors during early development have a crucial impact on brain function, causing individual differences that could lead to behavioral alteration and increased risk for neurological diseases over the lifetime (Faa et al., 2016). One of the early experiences that affect the brain outcome is the maternal infection, which impairs the complex immune harmony between the maternal and fetal environments, leading to a disrupted immune profile in the developing brain (Gumusoglu and Stevens, 2019). Indeed, clinical evidence has revealed that children born from mothers exposed to viral and bacterial pathogens during pregnancy are more likely to suffer various neurological disorders such as schizophrenia, autism bipolar disorder, major depression, epilepsy, and cerebral palsy (Bergdolt and Dunaevsky, 2018). Most of these epidemiological data have been reproduced in rodent models linked to the administration of lipopolysaccharide (LPS) during gestation (Boksa, 2010).

Although the offspring from LPS-exposed pregnant rodents displays a wide spectrum of brain abnormalities, including behavioral and cognitive changes, anatomical abnormalities, altered synaptic transmission, and immune disturbances (Golan et al., 2005; Rousset et al., 2006; Escobar et al., 2011), the involved mechanisms remain unknown. Moreover, most research has centered on the impact of prenatal inflammation in neurons and microglia, however, the potential modifications of astrocytes and neuron-astrocyte communication have received less scrutiny. Astrocytes encompass the most ubiquitous glial cell type and are endowed with the ability to sense neuronal function and react to it by releasing biomolecules termed "gliotransmitters" (e.g., glutamate, ATP, and D-serine) (Perea et al., 2009). Brain function is highly dependent on astrocytes, as they govern the energy supply to neurons (lactate) along with controlling the homeostatic balance of extracellular pH, neurotransmitters and ions, as well as modulating the redox response and intracellular  $Ca^{2+}$  concentration ( $[Ca^{2+}]_i$ ) signaling (Santello et al., 2019). During brain disease, astrocytes undergo a protective cell reaction called "reactive astrogliosis," however, when damage turns persistent, this response could disturb astrocyte-to-neuron communication and facilitate the recruitment of the innate immune response (Pekny and Pekna, 2014).

Despite that is known that prenatal LPS exposure triggers reactive astrogliosis (Hao et al., 2010; Zager et al., 2015), the signaling that shed light on this phenomenon and whether other astrocytic properties (e.g., gliotransmitter release, [Ca<sup>2+</sup>]<sub>i</sub> dynamics, NO production) are disturbed remain poorly understood. Current studies suggest that cellular cascades associated to hemichannels and pannexons are pivotal for astroglial function and dysfunction (Abudara et al., 2018). Hemichannels are plasma membrane channels composed by the oligomerization of six connexin monomers around a central pore that permit the diffusion of ions and small molecules, acting as a signaling route for communication between the cytoplasmic and extracellular compartments (Abudara et al., 2018). On the other hand, pannexins channels or pannexons are made of the oligomerization of pannexins, a three-member protein family with similar secondary and tertiary structures

than connexins and with the capacity to form plasma membrane channels (Dahl, 2018).

In astrocytes, hemichannels and pannexons permit the release of gliotransmitters that have been found crucial for synaptic transmission and plasticity, as well as behavior and memory (Stehberg et al., 2012; Ardiles et al., 2014; Chever et al., 2014; Walrave et al., 2016; Meunier et al., 2017). However, during pathological conditions, the permanent activity of these channels has been linked to the homeostatic disturbances occurring in the pathogenesis and progression of multiple diseases (Salameh et al., 2013; Orellana et al., 2016; Leybaert et al., 2017). In a previous study, using neonatal primary cell cultures, we demonstrated that astrocytes obtained from the offspring of LPS-exposed dams show an elevated opening of hemichannels and pannexons (Avendano et al., 2015). Nevertheless, whether prenatal LPS exposure affects the opening of these channels in the adult offspring and the possible impact of this on astrocyte function and neuronal survival is still ignored. Here, by performing studies in the stratum radiatum of the hippocampus, we found that prenatal LPS exposure increases the activity of astrocyte connexin 43 (Cx43) hemichannels and pannexin-1 (Panx1) channels ex vivo in acute brain slices from adult offspring mice. Relevantly, the opening of Cx43 hemichannels affected the release of glutamate,  $[Ca^{2+}]_i$  handling, and morphology of astrocytes, whereas also impaired the dendritic arbor and spine density, as well as neuronal survival.

#### MATERIALS AND METHODS

#### **Reagents and Antibodies**

The mimetic peptides gap19 (KQIEIKKFK, intracellular loop domain of Cx43), gap19<sup>I130A</sup> (KQAEIKKFK, negative control), Tat-gap19 (YGRKKRRQRRR-KQIEIKKFK, intracellular loop domain of Cx43), Tat-gap19<sup>I130A</sup> (YGRKKRRQRRR-KQAEIKK FK, negative control), Tat-L2 (YGRKKRRQRRR-DGANVDM HLKQIEIKKFKYGIEEHGK, second intracellular loop domain of Cx43), Tat-L2<sup>H126K/I130N</sup> (YGRKKRRQRRR-DGANVD MKLKQNEIKKFKYGIEEHGK, negative control), <sup>10</sup>panx1 (WRQAAFVDSY, first extracellular loop domain of Panx1) and <sup>10</sup>panx1<sup>src</sup> (FSVYWAQADR, scrambled peptide) were obtained from Genscript (New Jersey, United States). HEPES, water (W3500), minocycline, SB203580, polyclonal anti-Cx43 antibody, anti-glial fibrillary acidic protein (GFAP) monoclonal antibody, minocycline, oATP, ethidium (Etd) bromide (Ex-Max 493 nm/Em-Max 620 nm), sulforhodamine 101 (SR101) (Ex-Max 586 nm/Em-Max 605 nm) and probenecid (Prob) were purchased from Sigma-Aldrich (St. Louis, MO, United States). A740003, U-73122, 2-APB, MTEP, SIB-1757, LN-6, and A740003 were obtained from Tocris (Bristol, United Kingdom). Fluo-4-AM (Ex-Max 494 nm/Em-Max 506 nm), DAF-FM diacetate (Ex-Max 495 nm/Em-Max 515 nm), monoclonal anti-Iba-1 antibody, BAPTA-AM, diamidino-2-phenylindole (DAPI) (Ex-Max 359 nm/Em-Max 461 nm), goat anti-mouse Alexa Fluor 488 (Ex-Max 495 nm/Em-Max 519 nm) were obtained from Thermofisher (Waltham, MA, United States). Fluoro-Jade C (F-Jade) (Ex-Max 485 nm/Em-Max 525 nm) were obtained

from Chemicon (Martinsried/Munich, Germany). A soluble form of the TNF- $\alpha$  receptor (sTNF- $\alpha$ R1) and a recombinant receptor antagonist for IL-1 $\beta$  (IL-1ra) were from R&D Systems (Minneapolis, MN, United States).

#### Animals

The animals were treated and handled according to the National Institutes of Health guidelines (NIH, Baltimore, MD, United States). The experimental procedures were approved by the Bioethical and Biosafety Committee of the Faculty of Biological Sciences at the Pontificia Universidad Católica de Chile. Mice of 8–9 weeks of age were housed in cages in a temperature-controlled ( $24^{\circ}$ C) and humidity-controlled vivarium under a 12 h light/dark cycle (lights on 8:00 a.m.), with *ad libitum* access to food and water. The experiments performed in this study involved the following number of offspring animals per group of treatment: control (54), prenatal LPS (78), prenatal LPS + Tat-gap19 (6) and prenatal LPS + Tat-gap19<sup>1130A</sup> (6) (see below for details).

### Prenatal LPS Exposure Protocol and Mimetic Peptide *in vivo* Administration

The protocol of inflammatory stimulation was applied on gestation day 17. Pregnant mice were randomly assigned to one of two groups: (1) control (0.9% NaCl, i.p. injection) and (2) prenatal LPS (0.01 µg/g E. Coli LPS, i.p. injection). Given that prenatal LPS inflammation may induce sex-specific brain effects in the offspring (Makinson et al., 2017), we used only male offspring in our studies. Following full term delivery male offspring were used to prepare acute brain slices. In some in vivo experiments, we used the gap19 mimetic peptide coupled to the TAT membrane translocation motif (Tat-gap19), which is known to cross the blood-brain barrier (Abudara et al., 2014). Although gap19 contains the KKFK sequence that is a known cellmembrane translocation motif that facilitates plasma membrane permeability (Carrigan and Imperiali, 2005), we used the TAT version of this peptide in order to increase its cell membrane permeability and chances to interact with its union site: the intracellular C-terminal tail of Cx43 (Abudara et al., 2014). An I130A-modified gap19 analog (Tat-gap19<sup>I130A</sup>) was employed as a negative control peptide because amino acid I130 is implicated in the formation of hydrogen bonds and thereby crucial for gap19 activity (Wang et al., 2013). Accordingly, Tat-Gap19<sup>I130A</sup> exerts no inhibitory effects on Cx43 hemichannels (Wang et al., 2013). Tat-gap19 (23 mg/kg), Tat-gap19<sup>I130A</sup> (23 mg/kg) or saline solution were administrated via intraperitoneal (i.p.) injections beginning on PND 30, as has been previously described to be useful in acute and long-lasting administration in rodents (Crespo Yanguas et al., 2018; Chen et al., 2019; Maatouk et al., 2019). A second dose was given on PND45 followed by injections in PND 60, 75, 90, and 105.

#### **Acute Brain Slices**

Mice were anesthetized under isoflurane, decapitated and brains were extracted and cut into coronal slices (300  $\mu$ m) using a vibratome (Leica, VT1000GS; Leica, Wetzlar, Germany) filled

with ice-cold slicing solution containing (in mM): sucrose (222); KCl (2.6); NaHCO<sub>3</sub> (27); NaHPO<sub>4</sub> (1.5); glucose (10); MgSO<sub>4</sub> (7); CaCl<sub>2</sub> (0.5) and ascorbate (0.1), bubbled with 95%  $O_2/5\%$  CO<sub>2</sub>, pH 7.4. Then, the slices were transferred at room temperature (20–22°C) to a holding chamber in ice-cold artificial cerebral spinal fluid (ACSF) containing (in mM): NaCl (125), KCl (2.5), glucose (25), NaHCO<sub>3</sub> (25), NaH<sub>2</sub>PO<sub>4</sub> (1.25), CaCl<sub>2</sub> (2), and MgCl<sub>2</sub> (1), bubbled with 95%  $O_2/5\%$  CO<sub>2</sub>, pH 7.4, for a stabilization period of 60 min before dye uptake experiments (see below).

#### Treatments

Some acute brain slices were pre-incubated for 15 min before and during experiments with the following agents: mimetic peptides against Cx43 hemichannels (Tat-L2 and gap19, 100  $\mu$ M) and pannexin1 (Panx1) channels (<sup>10</sup>panx1, 100  $\mu$ M), Prob (pannexin channel blocker, 500  $\mu$ M), minocycline (inhibitor of microglial activation, 50 nM), sTNF- $\alpha$ R1 (soluble form of the receptor that binds TNF- $\alpha$ , 300 ng/ml), IL-1ra (IL-1 $\beta$  receptor endogenous blocker, 300 ng/ml), SB203580 (p38 MAP kinase inhibitor, 1  $\mu$ M), L-N6 (iNOS inhibitor, 1  $\mu$ M), A740003 (P2X<sub>7</sub> receptor blocker, 200 nM), oATP (general P2X receptor blocker, 200  $\mu$ M), BAPTA-AM (intracellular Ca<sup>2+</sup> chelator, 10  $\mu$ M), MTEP (selective mGluR<sub>5</sub> antagonist, 50 nM), SIB-1757 (selective mGluR<sub>5</sub> antagonist, 5  $\mu$ M), U-73122 (selective phospholipase C (PLC) inhibitor, 5  $\mu$ M), 2-APB (inhibitor of IP<sub>3</sub> receptor antagonist, 50  $\mu$ M), tetrodotoxin (TTX, 0.5  $\mu$ M).

# Dye Uptake in Acute Brain Slices and Confocal Microscopy

For dye uptake and ex vivo "snapshot" experiments, acute brain slices were incubated with 5  $\mu$ M ethidium (Etd) for 10 min in a chamber filled with ACSF and bubbled with 95% O<sub>2</sub>/5% CO<sub>2</sub>, pH 7.4. Afterward, the slices were washed three times (5 min each) with ACSF, and fixed at room temperature with 4% paraformaldehyde for 60 min, rinsed once with 0.1 mM glycine in phosphate buffered saline (PBS) for 5 min and then twice with PBS for 10 min with gentle agitation. Then, the slices were incubated two times for 30 min each with a blocking solution (PBS, gelatin 0.2%, Triton-X 100 1%) at room temperature. Afterward, the slices were incubated overnight at 4°C with anti-GFAP monoclonal antibody (1:500, Sigma) to detect astrocytes. Additionally, some slices not previously subjected to Etd uptake were incubated overnight at 4°C with anti-Iba-1 monoclonal antibody (1:500, Thermofisher) to detect microglia or antipolyclonal Cx43 antibody (1:400, SIGMA) to detect Cx43. Later, the slices were washed three times (10 min each) with blocking solution and then incubated for 2 h at room temperature with goat anti-mouse Alexa Fluor 488 (1:1000) antibody and Hoechst 33342. Further, the slices were washed three times (10 min each) in PBS and then mounted in Fluoromount, coverslipped and examined in a confocal laser-scanning microscope (Eclipse Ti-E C2, Nikon, Japan). Stacks of consecutive confocal images were taken with 40X objective at 100 nm intervals were acquired sequentially with three lasers (in nm: 408, 488, and 543), and Z projections were reconstructed using Nikon confocal

software (NIS-elements) and ImageJ software. At least six cells per field were selected from at least three fields in each brain slice. To assess the fluorescent intensity and distribution of Cx43 in astrocytes, stacks of consecutive confocal images were taken with the same confocal microscope, but with a 60X oil immersion objective (1.4 NA) at 200 nm intervals. Images were acquired sequentially with three lasers (in nm: 488 and 543), and Z projections were reconstructed using Nikon confocal software (NIS-elements). Image analysis of Z projections was then performed with ImageJ software. Cx43 intensity in areas close to the plasma membrane and cytoplasm was modeled by using the Otsu plugin for automatic image thresholding and the "enlarge" function of ImageJ. With the latter, we created a 10-pixel extension from the contour of the intracellular GFAP signal of each selected astrocyte to obtain an approximation of the plasma membrane area. Dye uptake or Cx43 fluorescence was calculated with the following formula: Corrected fluorescence = Integrated Density - ([Area of selected cell] x [Mean fluorescence of background readings]).

# Enzyme-Linked Immunosorbent Assay (ELISA)

ELISA assays were performed to determine the amount of TNF- $\alpha$  and IL-1 $\beta$  in the hippocampus. Mice were anesthetized with ketamine/xylazine (10:1 mg/kg of body weight, i.p.) and then perfused and decapitated. Afterward the hippocampus was removed and homogenized with an Ultra-Turrax homogenizer in buffer containing Tris-HCl 100 mM pH 7.4, EDTA 5 mM, SDS 1%, PMSF 1 µM and the protease inhibitor cocktail (ratio: 0.1 g hippocampus tissue: 1 ml lysis buffer) (Pierce, Rockford, IL, United States). Protein concentrations were determined by using a detergent-compatible Bio-Rad protein assay kit (Bio-Rad, Richmond, CA, United States). Then, the samples were centrifuged at 14,000 g for 10 min. Supernatants were collected and protein content assayed by BCA method. Cytokine levels were determined by sandwich ELISA, according to the manufacturer's protocol (IL-1 $\beta$  and TNF- $\alpha$  EIA kit, Enzo Life Science, United States). For the assay, 100 µl of samples were added per ELISA plate well and incubated 4°C overnight. A calibration curve with recombinant cytokine was included. Detection antibody was incubated at room temperature for 2 h and the reaction developed with avidin-HRP and substrate solution. Absorbance was measured at 450 nm with reference to 570 nm with the microplate reader Synergy HT (Biotek Instruments). The results were normalized by protein amount.

# [Ca<sup>2+</sup>]<sub>i</sub> and NO Imaging

Acute brain slices were incubated for 20 min at 34°C in ACSF solution containing 1  $\mu$ M SR101, washed and processed for Fluo-4 AM (Ca<sup>2+</sup> indicator) or DAF-FM (NO indicator) loading. For that, acute slices were incubated for 60 min at 37°C in ACSF containing 0.02% Pluronic F-127 and 5  $\mu$ M Fluo-4 AM or 5  $\mu$ M DAF-FM. Then, slices were transferred on the stage of a confocal laser scanning microscope and Ca<sup>2+</sup> or NO measurements were carried out for 20 min. Fluo-4 or DAF-FM were excited with an argon laser (488 nm) and emission

was filtered with a 515  $\pm$  15 nm filter, whereas SR101 was excited with a HeNe green laser (543 nm) and emission was filtered with a 605  $\pm$  75 nm filter. Acquisitions were carried out in the frame-scanning mode at 1 frame every 2 s with a 60x objective (NA 0.95; Nikon, Tokyo, Japan) on an Eclipse microscope (Nikon Instruments, Tokyo, Japan) equipped with and confocal head (confocal C2 head, Nikon) and controlled by the NIS-element software. The NO/Ca<sup>2+</sup> imaging data was analyzed using FIJI-IMAGE-J programs. Images with obvious motion were excluded for analysis. ROIs in astrocytes, including somata and processes, were manually identified on the basis of morphology. Fluorescence intensity was calculated with the following formula: Corrected total cell fluorescence = Integrated Density - ([Area of selected cell] × [Mean fluorescence of background readings]). At least four cells per field were selected from at least three fields in each brain slice. For spontaneous  $[Ca^{2+}]_i$  oscillations, the peaks were detected using the algorithm developed by Igor Pro from WaveMetrics. The frequency and amplitude were calculated and measured.

# Measurement of Extracellular ATP and Glutamate Concentration

Acute hippocampal slices were immersed in oxygenated ACSF (as above), pH 7.4, at room temperature (20-22°C) for 30 min either under control conditions or exposed to different agents. Then, extracellular ATP was measured using a luciferin/luciferase bioluminescence assay kit (Sigma-Aldrich, St. Louis, MO, United States), while extracellular levels of glutamate were determined using an enzyme-linked fluorimetric assay (Sigma-Aldrich, St. Louis, MO, United States). The amount of glutamate and ATP in each sample was inferred from standard curves. Briefly, after the experiments, the slices were washed twice with ACSF solution and sonicated in icecold PBS containing 5 µM EDTA, Halt (78440) and T-PER protein extraction cocktail (78510) according to manufacturer instructions (Pierce, Rockford, IL, United States). Total proteins from tissue homogenates were measured using the Bio-Rad protein assay.

# Golgi Staining

Mice were deeply anesthetized with isoflurane (4%) before euthanizing by decapitation. Brains were removed quickly from the skull to avoid any damage to the tissue. After rinsing, the tissue was sliced in approximately 10 mm thick blocks. The blocks were stained with the FD Rapid GolgiStain<sup>TM</sup> kit (FD NeuroTechnologies, Ellicott City, MD, United States). They were first immersed in the impregnation solution (A and B) which was replaced after 6-12 h and then kept in dark for 15-16 days. Afterward, the blocks were put in Solution C which was replaced after 24 h and kept in dark for the next 48-60 h. Cryomicrotome (Microm Thermo Scientific, Walldorf, Germany) was used to cut 200 µm thick slices. Slices were mounted on a gelatin-coated microscope slides, stained, and dehydrated and coverslipped with Permount. Tissue preparation and staining were all done by the same person following the FD Rapid GolgiStain<sup>TM</sup> kit manufacturer's protocol. Neuronal dendritic arbors and

spines were imaged using motorized microscope-computer based system and the MFB Stereo Investigator software version 11 (MBF–Bioscience, Williston, ND, United States). System was composed of z-axis motorized Olympus BX51 microscope equipped with x-y motorized stage guided by MAC5000 stage controller (Ludl Electronic Products Ltd., Hawthorne, NY, United States).

#### **Morphometry and Sholl Analysis**

Image processing of slices labeled for microglia (Iba-1 immunostaining), astrocytes (GFAP immunostaining) or neurons (golgi staining) was performed using the Fiji-ImageJ software (Schindelin et al., 2012). All samples were coded and analyzed randomly by a researcher blinded to animal number and condition. A minimum of 10 cells from each animal where chosen for analysis and their image data were imported using the BioFormats plugin and then channels separated with the Split channels tool. Later, the Iba-1, GFAP or golgi channel were selected and Z-axis projection of the sum of planes was performed using the Z projection tool. Afterward, microglia, astrocytes or neurons were selected and cut with the crop tool to facilitate their analysis when they fulfilled the following criteria: (i) presence of untruncated processes, (ii) consistent and strong staining along the entire arborization field, and (iii) relative isolation from neighboring cells to avoid overlap. Afterward, signal was segmented with the threshold tool and converted to binary mask before its skeletonization with the skeletonize tool. The latter tool allowed to obtain segment length and any possible bifurcation of the skeletonized image analyzed with the Fiji-ImageJ software. Due their complexity, drawings of neurons were done before skeletonization by using the Neuromantic software (Myatt et al., 2012), which allow the semi-manual or semi-automatic reconstruction of neurons from single images or image stacks. Then various features were measured including maximum and total branch length of cell processes, number of terminals, maximum path distance (maximum length of a path between the soma and terminal dendrites), as well as the number of branches were measured with the AnalizeSkeleton plugin of Fiji-ImageJ and/or the Neuromantic software. Further, the plugin Sholl analysis of Fiji-ImageJ was used to place concentric circles around the cell starting from the soma and radiating outward at increasing radial increments of 5 µm (Sholl, 1953). Different parameters were measured including the numbers of intersections (points where the cellular processes cross concentric rings), area under the Sholl curve, the maximum number of intersections, the radius of highest count of intersections (maximum intersect. radius) and the sum of intersections divided by intersecting radii (mean of intersections).

### **Spine Density Estimation**

Dendrite spine counting was conducted blind to the experimental condition. Measurements were obtained from the CA1 area of the dorsal hippocampus, whereas secondary or tertiary dendrite branches from the apical part (stratum radiatum) and from the basal part (stratum oriens) of the pyramidal cells were analyzed. Dendrite fragments chosen for analysis had to meet the following criteria: (i) good staining and impregnation without breaks, (ii) location about 150  $\mu$ m (apical part) or about 40  $\mu$ m (basal part) from the soma, (ii) branch fragments must be in the same focus plane and have a length about 30  $\mu$ m (20–50  $\mu$ m), and (iv) the branch fragment must be relatively straight to minimize errors connected with length measuring. About 8 fragments per brain were analyzed and spines were counted in the Fiji-ImageJ software. Afterward, signal was segmented with the threshold tool and converted to binary mask before its skeletonization with the skeletonize tool. The latter tool allowed to obtain segment length and the number of spines using the semiautomatic counting plugin of the Fiji-ImageJ software.

### **Neuronal Death Quantification**

Acute brain slices were fixed in 40% ethanol at 4°C for 5 min, treated with 0.1% Triton X-100 in PBS for 10 min and rinsed twice with distilled water. Preparations were incubated with 0.001% F-Jade in distilled water and gently shaken for 30 min in the dark. Later, F-Jade was removed and slices were incubated with anti-GFAP monoclonal antibody (Sigma, 1:400) diluted in 0.1% PBS-Triton X-100 with 2% NGS at 4°C overnight. After five rinses in 0.1% PBS-Triton X-100, slices were incubated with goat anti-mouse IgG Alexa Fluor 488 (1:1000) at room temperature for 50 min. After several washes, coverslips were mounted in DAKO fluorescent mounting medium and examined with a confocal laser-scanning microscope (Olympus, Fluoview FV1000, Tokyo, Japan).

#### **Statistical Analysis**

For each data group, results were expressed as mean  $\pm$  standard error (SEM); n refers to the number of independent experiments. Detailed statistical results were included in the figure legends. Statistical analyses were performed using GraphPad Prism (version 7, GraphPad Software, La Jolla, CA, United States). Normality and equal variances were assessed by the Shapiro-Wilk normality test and Brown-Forsythe test, respectively. Unless otherwise stated, data that passed these tests were analyzed by unpaired *t*-test in case of comparing two groups, whereas in case of multiple comparisons, data were analyzed by one or two-way analysis of variance (ANOVA) followed, in case of significance, by a Tukey's post hoc test. When data were heteroscedastic as well as not normal and with unequal variances, we used Mann-Whitney test in case of comparing two groups, whereas in case of multiple comparisons data are analyzed by Kruskal-Wallis test followed, in case of significance, by Dunn's post hoc test. A probability of P < 0.05 was considered statistically significant.

# RESULTS

# Prenatal LPS Exposure Enhances the Opening of Cx43 Hemichannels and Panx1 Channels in the Hippocampus of Adult Offspring Mice

The offspring of LPS-exposed dams exhibit alterations in hippocampal-dependent synaptic plasticity and memory

(Hao et al., 2010; Kelley et al., 2017), as well as increased neuronal death and astrogliosis (Ling et al., 2002; Zager et al., 2015). Because the exacerbated activity of astrocyte hemichannels and pannexons impact synaptic impairment, neuronal loss and astrogliosis in the hippocampus (Abudara et al., 2015; Yi et al., 2016; Gomez et al., 2018), we examined whether prenatal LPS exposure modulates the functional activity of these channels in the hippocampal CA1 region of the offspring. For that reason, we investigated hemichannel and pannexon activity by measuring ethidium (Etd) uptake in acute brain slices from the offspring mice following different months after birth. Etd enters to the cytosol of activated cells through selective large-pore channels, including hemichannels and pannexons. After its intercalation with base pairs of DNA and RNA, Etd becomes fluorescent, denoting channel activity (Johnson et al., 2016). Etd uptake by GFAP-positive astrocytes on acute brain slices was studied taking "snapshot" images in the stratum radiatum of the hippocampal CA1 region.

Astrocytes analyzed in acute brain slices from control offspring displayed a weak Etd uptake in the stratum radiatum (Figure 1A). Nonetheless, 4 months old offspring mice from LPS-exposed dams exhibited hippocampal astrocytes with increased Etd uptake compared to control conditions (~900%, Figures 1A-C). Temporal analysis of these responses showed that astroglial Etd uptake rapidly increased 1 month after birth in prenatally LPS-exposed offspring and reached a maximum in 4 months old offspring (Figure 1C). Thus, hereinafter and unless otherwise stated, this postnatal period was used in all further experiments throughout this study. Because Cx43 hemichannels and Panx1 channels are pivotal pathways for dye passage in astrocytes (Contreras et al., 2002; Iglesias et al., 2009), the potential involvement of these channels in the prenatal LPS-induced astroglial Etd uptake was investigated. Accordingly, acute brain slices were pre-incubated for 15 min before and during Etd uptake recordings with a battery of diverse pharmacological molecules. Tat-L2 (100  $\mu$ M) and gap19 (100  $\mu$ M); two mimetic peptides that inhibit Cx43 hemichannels by biding the intracellular L2 loop of Cx43 (Iyyathurai et al., 2013); strongly blunted the prenatal LPS-evoked Etd uptake in hippocampal astrocytes to  $\sim 30$ and 26% compared to 100% of the maximum response, respectively (Figure 1D). Moreover, an adapted Tat-L2 (Tat-L2<sup>H126K/I130N</sup>), in which 2 aa essential for binding of L2 to the CT tail of Cx43 are mutated, did not cause an equivalent inhibitory response (Figure 1D). Equivalently, we noted that an inactive structure of gap19 containing the I130A modification (gap19<sup>I130A</sup>), failed to reduce the prenatal LPS-dependent Etd uptake in astrocytes (Figure 1D). To elucidate the participation of Panx1 channels to the prenatal LPS-induced Etd uptake in hippocampal astrocytes, we employed the mimetic peptide <sup>10</sup>panx1 with an amino acid sequence homologous to the first extracellular loop domain of Panx1 (Pelegrin and Surprenant, 2006), as well as probenecid.  $^{10}\text{panx1}$  (100  $\mu\text{M})$  and probenecid (500  $\mu$ M) but not a scrambled peptide for <sup>10</sup>panx1 partially





counteracted the prenatal LPS-mediated astrocyte Etd uptake (**Figure 1D**). Collectively, this evidence suggests that prenatal LPS exposure augments the opening of astrocyte Cx43 hemichannels and Panx1 channels in the hippocampus from the adult offspring.

# Activation of Microglia and IL-1β/TNF-α/p38 MAP Kinase/iNOS Signaling Contribute to the Opening of Astrocyte Cx43 Hemichannels in Prenatally LPS-Exposed Adult Offspring

Given that release of inflammatory cytokines is critical for modulating molecular, morphological and functional properties of astrocytes during pathological conditions (Agulhon et al., 2012), we evaluated whether prenatal LPS exposure could modulate the hippocampal levels of these cytokines in the offspring. During the 1 month period after birth, the hippocampus of prenatally LPS-exposed offspring showed a strong  $\sim$ 5.5-fold increase in IL-1 $\beta$  levels compared to control that then dropped progressively in the following months (Figure 2A). Likewise, prenatal LPS exposure triggered a prominent 2.5-fold increase in hippocampal TNF-a levels of 1 month old offspring, which was slightly decreasing over time (Figure 2B). One of the major sources of cytokine production in the brain is the microglia and its activation has been observed along with neuroinflammation in prenatally LPS-exposed offspring (Schaafsma et al., 2017). Given that activation of microglia occurs along with changes in their morphology (Kettenmann et al., 2011), we measured the total branch length and branch points of microglial processes at the stratum radiatum. Analysis starting at the cell body throughout the end of each process, permit us to calculate the sum of all branch lengths and number of branch points of each microglia arbor. We found that prenatal LPS exposure reduced the branch points and the total length of microglial processes in the 4 months old offspring hippocampus (Figures 2C,D). Relevantly, microglia processes from prenatally LPS-exposed offspring showed similar branch points and length than their control counterparts when brain slices were treated with 50 nM minocycline (Figures 2C,D), a molecule that attenuates microglial activation (Kim and Suh, 2009).

To explore deeper the arbor complexity of microglia in the prenatally LPS-exposed offspring, we employed a Sholl analysis, which consists in place concentric rings at established intervals from the soma to then count branch intersections at each ring. We observed that hippocampal microglia of the offspring of LPS-exposed dams are significantly different from those of control offspring (Figures 2E-K). Particularly, during 4 months after birth, a dramatical reduction in the number of intersections between branches and Sholl rings was detected in hippocampal microglia of prenatally LPSexposed offspring (Figure 2E). Prenatal LPS exposure also reduced microglial branch complexity as measured by the area under the Sholl curve for the total number of branch intersections at 5–60  $\mu$ m from the soma (Figure 2L). Furthermore, hippocampal microglia of the offspring of LPS-exposed dams also exhibited decreased values in the maximum number of intersections, the radius of highest count of intersections (maximum intersect. radius) and the

sum of intersections divided by intersecting radii (mean of intersections) (**Figures 2M–O**). Relevantly, minocycline treatment greatly prevented not only the arbor reduction and altered morphology observed in hippocampal microglia from prenatally LPS-exposed offspring (**Figures 2E–O**), but also the increased production of IL-1 $\beta$  and TNF- $\alpha$  occurring in these conditions (**Figure 2P**).

On the other hand, we found that minocycline prominently blunted the prenatal LPS-induced Etd uptake in hippocampal astrocytes, whereas pretreatment with a soluble form of TNF-a receptor that binds TNF-a (sTNF-aR1) and a recombinant antagonist for IL-1 $\beta$  receptor (IL-1ra) caused equivalent responses (Figure 2Q). It is known that IL-1 $\beta$  and TNF- $\alpha$ along with p38 MAP kinase activation, lead to NO-dependent S-nitrosylation of astrocytic Cx43 hemichannels, increasing their activity (Retamal et al., 2007). In this line, we found that the prenatal LPS-induced Etd uptake in hippocampal astrocytes was prominently tackled by blocking p38 MAP kinase with 10 µM SB202190 or of iNOS by 5 µM L-N6 (Figure 2Q). Altogether, these observations reveal that TNF- $\alpha$ /IL-1 $\beta$  and activation of iNOS/p38 MAP kinase pathways appear to be pivotal for the prenatal LPS-evoked opening of astrocyte Cx43 hemichannels but not Panx1 channels in the hippocampus. Accordingly, the Cx43 hemichannel blocker gap19 failed to trigger any additive inhibition in the prenatal LPS-induced Etd uptake when slices were treated with minocycline (Figure 2Q). By contrast, the Panx1 channel blocker <sup>10</sup>panx1 caused an additive inhibition when slices were stimulated with minocycline (Figure 2Q), suggesting that pannexon activity is not linked to the release of cytokines from activated microglia.

Prior studies have described that opening of Panx1 channels relies on direct protein-protein interactions with P2X<sub>7</sub> receptors (P2X<sub>7</sub>Rs) (Iglesias et al., 2008). According with this evidence, we found that 200 nM A740003, a selective P2X<sub>7</sub>R antagonist, caused a partial reduction in the prenatal LPS-induced Etd uptake in hippocampal astrocytes (**Figure 2Q**), which was close to the inhibitory effect induced by Panx1 channel blockers (**Figure 1D**). Relevantly, <sup>10</sup>panx1 did not evoke any additive inhibition on Etd uptake of that caused by A740003 (**Figure 2Q**), underscoring the possibility that prenatal LPS-induced opening of Panx1 channels could take place via the activation of P2X<sub>7</sub>Rs.

### Prenatal LPS Exposure Increases the Astrocyte Production of NO and the Release of ATP via Panx1 Channels in the Offspring Hippocampus

Given that NO opens Cx43 hemichannels (Retamal et al., 2006) and because inhibition of iNOS with LN-6 greatly reduced the Etd uptake caused by prenatal LPS exposure in hippocampal astrocytes (**Figure 2Q**), we tested if this condition could affect NO production in the offspring hippocampus. DAF-FM (NO indicator) and SR101 (astrocyte marker) fluorescence imaging revealed that hippocampal astrocytes from prenatally LPS-exposed offspring showed a  $\sim$ 2.5-fold augment in basal NO production compared to control conditions (**Figures 3A–C**). The fact that LN-6 totally suppressed the prenatal-LPS-induced



FIGURE 2 | Microglia and IL-1β/TNF-α/p38 MAP kinase/iNOS signaling participate in the prenatal LPS-induced opening of astrocyte Cx43 hemichannels on offspring hippocampus. Averaged data of hippocampal levels of IL-1β (A) and TNF-α (B) from control offspring (white bars) or prenatally LPS-exposed offspring (black bars) following different postnatal periods. \*\*\* p < 0.0001, \*\* p < 0.005, \* p < 0.05 versus control, two-way ANOVA Bonferroni's post hoc test, mean ± S.E.M., n = 3. (C,D) Averaged data of total branch length (C) and number of branches (D) by hippocampal microglia in acute slices from control offspring (white bars) or prenatally LPS-exposed offspring (black bars) of 4 months old. Also shown are the effects of treatment with 50 nM minocycline for 2 h in acute slices prenatally LPS-exposed offspring of 4 months old (gray bars). \*p < 0.0001 versus LPS, one-way ANOVA Dunnett's post hoc test, mean ± S.E.M., n = 3. (E) Averaged data of Sholl analysis by hippocampal microglia from control offspring (white circles) or prenatally LPS-exposed offspring of 4 months old alone (black circles) or plus treatment with 50 nM minocycline (gray circles). \*p < 0.001 versus LPS, two-way ANOVA Tukey's post hoc test, mean ± S.E.M., n = 3. (F-K) Representative Iba-1 (black) positive hippocampal microglia in acute slices from control offspring (F,G) or prenatally LPS-exposed offspring of 4 months old alone (H,I) or plus treatment with 50 nM minocycline (J,K). Insets of microglia (G,I,K) were taken from the area depicted within the red squares in (F,H,J). (L-O) Averaged data of area under the curve of Sholl analysis (L), maximum intersection (M), maximum intersection radius (N), and mean of intersections (O) by hippocampal microglia from control offspring (white bars) or prenatally LPS-exposed offspring of 4 months old alone (black bars) or plus treatment with 50 nM minocycline (gray bars). \*p < 0.0001 versus LPS, one-way ANOVA Dunnett's post hoc test, mean ± S.E.M., n = 3. (P) Averaged data of hippocampal levels of IL-1β and TNF-α from control offspring (white bars) or prenatally LPS-exposed offspring of 4 months old alone (black bars) or plus treatment with 50 nM minocycline (gray bars). \*\*\* p < 0.0001, versus control, two-way ANOVA Bonferroni's post hoc test, mean ± S.E.M., n = 3. (Q) Averaged data normalized to the maximal effect (dashed line) induced by prenatal LPS exposure on Etd uptake by hippocampal astrocytes in acute slices from 4 months old offspring exposed to the following pharmacological agents: 50 nM minocycline, 50 nM minocycline + 100 μM gap19, sTNF-αR1 + IL-1ra (300 ng/ml each), 1 μM SB203580, 1 μM L-N6, 50 nM minocycline + 100 μM  $^{10}$ panx1, 200 nM A740003 or 100  $\mu$ M  $^{10}$ panx1 + 200 nM A740003. \*\*p < 0.0001, \*p < 0.005 versus LPS, \*p < 0.0001 versus minocycline, one-way ANOVA Dunnett's post hoc test, mean  $\pm$  S.E.M., n = 3. Calibration bars: black bar = 180  $\mu$ m; yellow bar: 80  $\mu$ m.



FIGURE 3 | Prenatal LPS exposure increases the production of NO by astrocytes and the Panx1 channel-dependent release of ATP on offspring hippocampus. Representative images showing SR101 (red) and DAF-FM (green) staining by hippocampal astrocytes in acute slices from control offspring (A) or prenatally LPS-exposed offspring (B) of 4 months old. Insets of astrocytes were taken from the area depicted within the red squares in (A,B). (C) Averaged of DAF-FM signal fluorescence by astrocytes in acute slices from control offspring (white bar) or prenatally LPS-exposed offspring of 4 months old alone (black bars) or in combination with the following pharmacological agents: 100  $\mu M$  gap19, 100  $\mu M$   $^{10} panx1$  and 1  $\mu M$  L-N6. \*p < 0.0001 versus LPS, one-way ANOVA Tukey's post hoc test, mean  $\pm$  S.E.M., n = 3. (D) Averaged data of ATP release by acute hippocampal slices from control offspring (white bar) or prenatally LPS-exposed offspring of 4 months old alone (black bars) or in combination with the following blockers: 100 µM gap19, 100 µM <sup>10</sup>panx1 and 1 µM L-N6. \*p < 0.0001 versus LPS, \*p < 0.0001 versus control, one-way ANOVA Tukey's post hoc test, mean  $\pm$  S.E.M., n = 3. Calibration bar = 85  $\mu$ m.

production of NO indicates that iNOS is the major contributor to this response (**Figure 3C**). A previous study has related the opening of Panx1 channels with the production of NO (Orellana et al., 2013). In opposition to this finding, we observed that neither <sup>10</sup>panx1 nor gap19 were effective in to prevent the prenatal-LPS-induced production of NO (**Figure 3C**), suggesting that Panx1 channels and Cx43 hemichannels and are not involved in this response.

iNOS stimulation is crucial for the hemichannel/pannexondependent release of ATP from astrocytes occurring after LPS treatment (Avendano et al., 2015). With this in mind and given that A740003, a selective antagonist of P2X<sub>7</sub>Rs, counteracted the astrocyte Etd uptake triggered by prenatal LPS exposure (**Figure 2Q**), we investigated if this condition could impact the release of ATP in the offspring hippocampus. Measurements of extracellular ATP levels with the luciferin/luciferase bioluminescence assay showed that prenatal LPS exposure dramatically augmented the release of ATP in ~7-fold in the offspring hippocampus compared to control conditions (**Figure 3D**). Importantly, probenecid or <sup>10</sup>panx1 but not gap19 markedly reduced the release of ATP caused by prenatal LPS exposure (from ~73 pmol/mg to ~30 pmol/mg and ~31 pmol/mg, respectively) (**Figure 3D**). Similarly, 200  $\mu$ M oATP or 200 nM A740003 prominently reduced the prenatal LPS-evoked release of ATP (**Figure 3B**).

# Cx43 Hemichannel Opening Evoked by Prenatal LPS Exposure Contributes to [Ca<sup>2+</sup>]<sub>i</sub> and Glutamatergic Signaling on Offspring Hippocampus

At the next step, we decided to analyze the effect of hemichannel/pannexon blockers on astroglial  $[Ca^{2+}]_{i}$ in prenatally LPS-exposed offspring. As indicated by the assessment of Fluo-4 (Ca<sup>2+</sup> indicator) and SR101, hippocampal astrocytes from offspring of LPS-exposed dams showed a 3-fold augment in basal levels of Ca2+ signal compared to control astrocytes (Figures 4A-C). Importantly, blockade of Cx43 hemichannels with Tat-L2 or gap19 dramatically suppressed the prenatal LPS-mediated increase in astroglial  $[Ca^{2+}]_i$  in the hippocampus (Figure 4C). Similar observations were obtained upon treatment with minocycline but not with probenecid or <sup>10</sup>panx1 (Figure 4C). Altogether these findings indicate that microglial-dependent opening of Cx43 hemichannels but not Panx1 channels participate in the prenatal LPS-induced increase in astroglial  $[Ca^{2+}]_i$  in the hippocampus.

Glutamate modulates  $[Ca^{2+}]_i$  dynamics in astrocytes, particularly through the stimulation of metabotropic glutamate receptors (mGluRs) and subsequent release of Ca<sup>2+</sup> from intracellular stores (Bradley and Challiss, 2012). Here, we found that prenatal LPS-evoked  $[Ca^{2+}]_i$  response was prominently blunted by 50 nM MTEP or 5 µM SIB-1757, being the latter two selective antagonists of mGluR<sub>5</sub> (Figure 4C). Noteworthy, selective blockade of phospholipase C (PLC) or IP<sub>3</sub> receptors with 5 µM U73122 or 50 µM APB, respectively, as well as chelation of [Ca<sup>2+</sup>]<sub>i</sub> with 10 µM BAPTA-AM, strikingly counteracted the increase in astroglial  $[Ca^{2+}]_i$  triggered by prenatal LPS exposure (Figure 4C). On the other hand, inhibition of P2X<sub>7</sub>Rs with oATP or A740003 caused a slight reduction in the prenatal LPS-induced astroglial  $[Ca^{2+}]_i$  signal (Figure 4C). This could imply that astroglial [Ca<sup>2+</sup>]<sub>i</sub> response resulting from prenatal LPS exposure is a consequence of the Cx43 hemichanneldependent release of glutamate and further stimulation of mGluR<sub>5</sub> rather than signaling via Panx1 channels. Consistent with this notion, we saw that prenatal LPS exposure induced a 7-fold increase in the release of glutamate in the offspring hippocampus, a response that was totally blunted by Tat-L2 or gap19 but not with probenecid or <sup>10</sup>panx1 (Figure 4D). Further, we tested whether [Ca<sup>2+</sup>]<sub>i</sub> and mGluR<sub>5</sub> signaling were implicated in the Etd uptake observed in hippocampal astrocytes from prenatally LPS-exposed offspring. BAPTA-AM, but not inhibition of PLC, IP<sub>3</sub> receptors or mGluR<sub>5</sub>, significantly reduced the prenatal LPS-induced Etd uptake in hippocampal astrocytes (Figure 4E). In this scenario, we further performed the analysis of spontaneous [Ca<sup>2+</sup>]<sub>i</sub> oscillations in astrocytes. We found that prenatal LPS exposure increase the number of spontaneous astroglial [Ca<sup>2+</sup>]<sub>i</sub> oscillations and their amplitude in the offspring hippocampus, a response that was totally blunted by BAPTA-AM but not by gap19,



Tat-L2, 100  $\mu$ M <sup>10</sup>panx1, 500  $\mu$ M Probenecid (Prob), 10  $\mu$ M Bapta-AM, 50 nM MTEP or 5  $\mu$ M SIB-1757. \*\*p < 0.005 versus LPS, one-way ANOVA Tukey's *post hoc* test, mean  $\pm$  S.E.M., n = 3. Calibration bar = 85  $\mu$ m.

Tat-L2, <sup>10</sup>panx1, probenecid, MTEP or SIB-1757 (**Figure 4F**). Similar responses were observed when the peak amplitude of spontaneous astroglial  $[Ca^{2+}]_i$  oscillations was analyzed (**Figure 4G**). To figure out the possible contribution of neurons to these responses, we performed the above experiments in presence of 0.5 µM TTX. We found that TTX did not affect the prenatal LPS-induced changes in NO and  $[Ca^{2+}]_i$  levels, as well as the release of glutamate, suggesting that neurons do not participate in these processes under these conditions (**Supplementary Figures S1A–E**). Collectively, these data suggest that spontaneous  $[Ca^{2+}]_i$  oscillations evoked by prenatal LPS exposure are likely necessary for the opening of astroglialCx43 hemichannels, whereas the subsequent release of glutamate through them is needed for the increase in basal  $[Ca^{2+}]_i$  via the activation of mGluR<sub>5</sub> receptors. Although previous studies have associated the channel-dependent Etd uptake with changes in the distribution of Cx43 in astrocytes (Avendano et al., 2015), we found that prenatal LPS exposure did not alter the total amount or distribution of Cx43 in hippocampal astrocytes (**Supplementary Figures S2A–G**).



bars) or prenatally LPS-exposed offspring of 4 months old alone (black bars) or plus the *in vivo* administration of 23 mg/kg Tat-gap19 (gray bars) or 23 mg/kg Tat-gap19<sup>1130A</sup> (red bars). \*p < 0.0001 versus LPS, one-way ANOVA Dunnett's *post hoc* test, mean  $\pm$  S.E.M., n = 3. Calibration bar = 40  $\mu$ m.

# Activation of Cx43 Hemichannels Contributes to Increased Arborization of Hippocampal Astrocytes in Prenatally LPS-Exposed Offspring

One of the crucial aspects of reactive astrogliosis is the hypertrophy of cellular processes accompanied by crucial changes

in the arborization and morphology of astrocytes (Pekny and Pekna, 2014). To understand whether the above described prenatal LPS-induced changes in astrocytes are accompanied by alterations in their arborization, we analyzed the maximum and total branch length of astroglial processes at the stratum radiatum (**Figures 5A–E**). Similar to the previous measurements made in microglia, we calculated the longest branch and the sum of all



hemichannels. (A–C) Representative golgi (black) staining by CA1 pyramidal neurons from control offspring (A), prenatally LPS-exposed offspring of 4 month old alone (B) or in combination with the *in vivo* administration of 23 mg/kg Tat-gap19 (C). (D–G) Averaged data of total branch length (D), number of branches (E), number of terminals (F), and maximum path distance (G) by CA1 pyramidal neurons from control offspring (white bars) or prenatally LPS-exposed offspring of 4 months old alone (black bars) or in combination with the *in vivo* administration of 23 mg/kg Tat-gap19 (gray bars). \*p < 0.05, \*\*p < 0.05, \*\*p < 0.05, versus LPS, one-way ANOVA Dunnett's *post hoc* test, mean ± S.E.M., n = 3. (H) Averaged data of Sholl analysis by apical (upper panel) and basal (bottom panel) dendritic arbor of CA1 pyramidal neurons from control offspring (white circles) or prenatally LPS-exposed offspring of 4 months old alone (black circles) or plus the *in vivo* administration of 23 mg/kg Tat-gap19 (gray circles). \*p < 0.05 versus LPS, two-way ANOVA Tukey's *post hoc* test, mean ± S.E.M., n = 3. (H) and maximum intersection (K) by apical (upper panel) and basal (bottom panel) dendritic arbor of CA1 pyramidal neurons from control offspring (white bars) or prenatally LPS-exposed offspring of 4 months old alone (black circles) or plus the *in vivo* administration of 23 mg/kg Tat-gap19 (gray circles). \*p < 0.05 versus LPS, two-way ANOVA Tukey's *post hoc* test, mean ± S.E.M., n = 3. (H-K) Averaged data of area under the curve of Sholl analysis (I), mean of intersections (J) and maximum intersection (K) by apical (upper panel) and basal (bottom panel) dendritic arbor of CA1 pyramidal neurons from control offspring (white bars) or prenatally LPS-exposed offspring of 4 months old alone (black bars) or plus the *in vivo* administration of 23 mg/kg Tat-gap19 (gray bars). \*p < 0.05, \*\*p < 0.05, \*\*p < 0.01 versus LPS, one-way ANOVA Tukey's *post hoc* test, mean ± S.E.M., n = 3. Calibration bar = 45 µm.

branch lengths of each astrocyte arbor, which were depicted as maximum and total branch length, respectively. Measurements of cell arbor disclosed that maximum branch length remained without alterations between hippocampal astrocytes from control and prenatally LPS-exposed offspring (Figure 5D). Nonetheless, astrocytes from prenatally LPS-exposed offspring exhibited a ~2fold increase in both total branch length (Figure 5E) and the number of branches (Figure 5F). Sholl analysis demonstrated that hippocampal astrocytes from the offspring of LPS-exposed dams showed more complex arbors than in control animals (Figures 5A-C,G). Specifically, in these astrocytes, both the number of intersections between branches and Sholl rings, as well as the area under the Sholl curve were increased (Figures 5G,H). Prenatal LPS exposure triggered equivalent augmented values in the maximum number of intersections, the radius of the highest count of intersections and mean of intersections (Figures 5I-K). These findings suggest that prenatal LPS enhance the complexity

of astrocyte branch arbors in the hippocampus of prenatally LPS-exposed offspring.

To unveil the contribution of Cx43 hemichannel activity in prenatal LPS-induced increase in branch arbor complexity, we injected prenatally LPS-exposed offspring mice during postnatal months with the gap19 mimetic peptide containing the cell-penetrating TAT linker (Tat-gap19; 23 mg/kg, see section Materials and methods), which crosses the blood-brain barrier (BBB) (Abudara et al., 2014). Notably, Tat-gap19 completely prevented the prenatal LPS-induced increase in diverse arbor parameters, including total branch length (**Figure 5E**), number of branches (**Figure 5F**), arbor complexity (**Figure 5G**), the area under the Sholl curve (**Figure 5H**), mean of intersections (**Figure 5I**), maximum number of intersections (**Figure 5J**), and the radius of highest count of intersections (**Figure 5K**). Relevantly, the inactive form of Tat-gap19 containing the I130A modification (TAT-gap19<sup>I130A</sup>) induced no effect on the prenatal

LPS-mediated increase on astrocyte arborization (**Figures 5D–K**). These findings suggest that opening of Cx43 hemichannels is crucial for the prenatal LPS-mediated increment in astrocyte arbor branch complexity in the offspring hippocampus.

# The Opening of Cx43 Hemichannels Is Required for the Prenatal LPS-Mediated Reduction in Arbor Branch Complexity and Dendritic Spine Density of Hippocampal Pyramidal Neurons in the Offspring

Hippocampal synaptic dysfunction has been linked with retraction of dendrites pyramidal neurons, as well as loss of synapses in diverse neurological disorders (Luo and O'Leary, 2005; Riccomagno and Kolodkin, 2015). Nonetheless, whether prenatal LPS exposure causes dendritic retraction and spine density reduction in the hippocampus have not been studied in detail (Fernandez de Cossio et al., 2017). Here, analysis of dendritic arbor (Figures 6A-C) showed that prenatal LPS exposure caused a  $\sim$ 2-fold decrease in total branch length (Figure 6D), number of branches (Figure 6E) and the number of terminals (Figure 6F) in CA1 pyramidal neurons. These neurons also exhibited a decrease in the maximum path distance between the soma and terminal dendrites when compared with their control counterparts (Figure 6G). A precise Sholl analysis underscored that CA1 pyramidal neurons from the offspring of LPS-exposed dams displayed a ~2-fold decline in arbor complexity in both basal and apical dendrites (Figure 6H). Indeed, prenatal LPS exposure diminished the area under the Sholl curve (Figure 6I), the mean of intersections between branches and Sholl rings (Figure 6J), as well as the maximum number of intersections (Figure 6K). Of note, dendritic retraction evoked by prenatal LPS exposure was accompanied by decreased spine density in apical but not basal dendrites of CA1 pyramidal neurons (Figures 7A-D).

Consistent with what occurred in hippocampal astrocytes, *in vivo* treatment with Tat-gap19 strongly counteracted the prenatal LPS-induced decrease in arborization, although this inhibitory response was more effective in apical rather than basal dendritic arbor (**Figures 6H–K**). In addition, Tat-gap19 also totally blunted the prenatal LPS-induced decrease in spine density in apical dendrites of CA1 pyramidal neurons (**Figure 7D**). TAT-gap19<sup>I130A</sup>, the inactive form of Tat-gap19, did not change the prenatal LPS-mediated reduction in the neuronal pyramidal arbor or spine density (not shown). Altogether these findings argue for a crucial role of Cx43 hemichannels in the prenatal LPS-mediated reduction of dendritic arbor and spine density of CA1 pyramidal neurons.

# Cx43 Hemichannels Participate in the Prenatal LPS-Induced Neuronal Death in Hippocampal Slices

It is well established that prenatal-LPS exposure triggers neuronal death in the offspring (Ling et al., 2002; Hao et al., 2010) and diverse studies have proposed that uncontrolled release of



substances via opening of astrocyte Cx43 hemichannels could be toxic for neighboring neurons (Orellana et al., 2011; Yi et al., 2016). With this in mind, we evaluated if prenatal LPS exposure could induce cell death in CA1 pyramidal neurons and whether Cx43 hemichannels are involved in this process. In control offspring, most pyramidal neurons were negative for F-Jade staining (~3 neurons/field) and most astrocytes displayed a normal grade of GFAP expression (Figures 8A-C,J,K). However, prenatally LPS-exposed offspring exhibited a ~10-fold augment in CA1 pyramidal neurons displaying F-Jade staining, which was accompanied by a qualitative augment in GFAP reactivity (Figures 8D-F,J,K). Relevantly, in vivo treatment with Tat-gap19 greatly tackled both the prenatal LPS-induced F-Jade staining of CA1 pyramidal neurons (to ~8 neurons/field) and the enchanced reactivity of GFAP (in ~40%) in hippocampal slices (Figures 8G-K). In the presence of TAT-gap $19^{1130A}$ , the F-Jade staining by CA1 pyramidal neurons remained unaltered in prenatally LPSexposed offspring (Figures 8J,K). The above data suggest that



prenatal LPS exposure on offspring hippocampus. (A-I) Representative images depicting Fluoro-Jade (F-Jade, green), GFAP (red) and DAPI (blue) staining in acute slices from control offspring (A-C) or prenatally LPS-exposed offspring of 4 months old alone (D-F) or in combination with the in vivo administration of 23 mg/kg Tat-gap19 (G-I). Insets of gray scale GFAP staining for astrocytes are shown from the area depicted within the white squares in (B,E,H). (J) Averaged data of GFAP fluorescence per field in acute slices from control offspring (white bars) or prenatally LPS-exposed offspring of 4 months old alone (black bars) or in combination with the in vivo administration of 23 mg/kg Tat-gap19 (gray bars). \*p < 0.01 versus LPS, one-way ANOVA Tukey's post hoc test, mean  $\pm$  S.E.M., n = 3. (K) Averaged number of F-Jade-positive CA1 pyramidal neurons per field in acute slices from control offspring (white bars) or prenatally LPS-exposed offspring of 4 months old alone (black bars) or in combination with the in vivo administration of 23 mg/kg Tat-gap19 (gray bars). \*p < 0.001 versus LPS, one-way ANOVA Tukey's post hoc test, mean  $\pm$  S.E.M., n = 3. Calibration bar = 100  $\mu$ m.

Cx43 hemichannels are main contributors to the hippocampal neuronal death caused by prenatal LPS exposure in the offspring.

# DISCUSSION

In this work, we reported that prenatal LPS exposure augments the activity of astrocyte Cx43 hemichannels and Panx1 channels in the hippocampus of adult offspring mice. This enhanced channel activity occurred by a mechanism involving a microglia-dependent production of IL-1 $\beta$ /TNF- $\alpha$  and the stimulation of p38 MAPK/iNOS/[Ca<sup>2+</sup>]<sub>i</sub>-mediated pathways and

purinergic/glutamatergic signaling. Noteworthy, the opening of Cx43 hemichannels affected the release of glutamate,  $[Ca^{2+}]_i$  handling, and morphology of astrocytes, whereas also disturbed neuronal function, including the dendritic arbor and spine density, as well as survival.

Previous evidence indicates that prenatal LPS exposure triggers diverse disturbances in the offspring brain, including alterations in hippocampal-dependent synaptic plasticity and memory (Golan et al., 2005; Escobar et al., 2011; Kelley et al., 2017), as well as increased neuronal death and astrogliosis (Ling et al., 2002; Hao et al., 2010; Zager et al., 2015; Cho et al., 2018). This study suggests that part of the above-mentioned abnormalities induced by prenatal LPS exposure could take place by the persistent opening of astrocyte Cx43 hemichannels and/or Panx1 channels within the hippocampus. As assayed by Etd uptake in acute brain slices, we found that a single LPS injection during pregnancy increases the opening of Cx43 hemichannels and Panx1 channels in hippocampal astrocytes from the stratum radiatum in the offspring. These responses were prominently inhibited by Tat-L2 or gap19, whereas probenecid or <sup>10</sup>panx1 showed a partial inhibitory effect. Thus, Cx43 hemichannels rather than Panx1 channels were the major responsible for the prenatal LPS-induced Etd uptake in astrocytes. The latter is consistent with the enhanced activity reported for both channels in astrocyte cultures obtained from prenatally LPS-exposed neonates (Avendano et al., 2015), as well as astrocytes from different animal pathological models such as neuropathic pain (Tonkin et al., 2018), Alzheimer's disease (Yi et al., 2016), epileptic seizures (Santiago et al., 2011), spinal cord injury (Garre et al., 2016), and acute brain infection (Karpuk et al., 2011).

How does prenatal LPS exposure trigger the opening of Cx43 hemichannels and Panx1 channels in hippocampal astrocytes ex vivo? Multiple lines of evidence indicate that environmental factors during early development impact the future inflammatory balance and immunity response of the offspring (Boksa, 2010). Here, we observed that prenatal LPS exposure induced a microglia-dependent long-lasting production of both IL-1ß and TNF- $\alpha$  on offspring hippocampus. The latter response was accompanied by a profound retraction of microglia cellular processes compatible with amoeboid features typical of activated microglia. This is in agreement with the presence of activated microglia as well as with upregulated levels of IL-1ß and TNF- $\alpha$  in the offspring's brain of LPS-exposed dams (Boksa, 2010). In addition, our experiments showed that minocycline, a molecule that attenuates microglial activation, or inhibition of IL- $1\beta$ /TNF- $\alpha$  signaling, dramatically suppressed the prenatal LPSinduced opening of astrocytic Cx43 hemichannels but not Panx1 channels on offspring hippocampus. These findings harmonize with prior studies showing that release of IL-1 $\beta$  and TNF- $\alpha$ from activated microglia causes the activation of astroglial Cx43 hemichannels (Retamal et al., 2007). Thereby, in our system, the opening of astroglial Cx43 hemichannels likely resulted from the IL-1β/TNF-α-mediated activation of p38 MAP kinase and further iNOS-dependent S-nitrosylation of Cx43, as has been previously described (Retamal et al., 2006; Retamal et al., 2007) (Figure 9). In accord with this hypothesis, we detected that prenatal LPS-induced astroglial Etd uptake was strongly



opening of Panx1 channels and further release of Ca<sup>-+</sup> stored in the endoplasmic reticulum. In parallel, the activation of astroglial P2x7 receptors lead to the opening of Panx1 channels and further release of ATP, possibly through direct protein-to-protein interactions. Relevantly, the modulation of [Ca<sup>2+</sup>]<sub>i</sub> dynamics evoked by Cx43 hemichannels may alter astroglial morphology (not depicted), whereas the excitotoxic release of glutamate through Cx43 hemichannels may affect neuronal arborization and survival by unknown mechanisms.

blunted by blocking p38 MAP kinase or iNOS. In the same line, hippocampal astrocytes from prenatally LPS-exposed offspring showed increased levels of NO production, as measured by DAF-FM signal. Of note, this response was totally blunted by the inhibition of iNOS but not by blockers of Cx43 hemichannels or Panx1 channels, suggesting that these channels do not participate in astrocyte NO production caused by prenatal LPS exposure.

A cornerstone underlying the opening of Panx1 channels came from their close association with P2X<sub>7</sub>Rs (Dahl, 2018). Indeed, Panx1 co-immunoprecipitates with P2X<sub>7</sub>Rs (Pelegrin and Surprenant, 2006; Poornima et al., 2012), and proline 451 in the C-terminal tails of these receptors has been found crucial in this interaction (Iglesias et al., 2008; Sorge et al., 2012). In this context, our experiments showed that activation of P2X<sub>7</sub>Rs and opening of astrocytic Panx1 channels are

part of the mechanism involved in the prenatal LPS-induced release of ATP on offspring hippocampus. In concordance with these findings, previous studies have described that ATP triggers its own release via P2X<sub>7</sub>Rs and further activation of Panx1 channels (Iglesias et al., 2009; Garre et al., 2016). The fact that Panx1 channels contribute to the prenatal LPS-induced Etd uptake and release of ATP but not glutamate is puzzling. One possibility is that despite that both channels may be permeable to Etd, they differ in their contribution to the release of ATP and glutamate in our system. Although this seems paradoxical, recent studies have demonstrated that hemichannels and pannexons do not act as freely permeable non-selective pores, but they select permeants in an isoform-specific form (Hansen et al., 2014a,b; Nielsen et al., 2017). Thus, fluorescent dye uptake cannot be employed as an indicator of

permeability to ions or small biologically relevant molecules. Alternatively, other explanation to these findings may imply that another cell type, expressing functional Panx1 channels may be responsible for the Panx1-dependent release of ATP and Etd uptake found in astrocytes in the hippocampus. Regardless of its source, we can speculate that ATP may activate distant astrocytes and/or neurons in a paracrine manner, triggering Ca<sup>2+</sup> responses that could rely on the reactive profile of astrocytes (Butt, 2011). If so, the stimulation of purinergic receptors could be shut down by diffusion of ATP to distant areas as well as by desensitization of P2Y receptors and degradation of extracellular ATP by exonucleases. Simultaneously, a negative feedback loop is the counteracting effect that could be evoked by ATP on Panx1 channels (Qiu and Dahl, 2009).

How does Cx43 hemichannel opening affect Ca<sup>2+</sup> signaling? Hemichannels are permeable to  $Ca^{2+}$  (De Bock et al., 2012; Fiori et al., 2012). In this scenario, alterations on  $[Ca^{2+}]_i$ handling linked to astrocyte hemichannel opening could be pivotal in the potential vicious cycle underpinning astrocyte dysfunction in the prenatally LPS-exposed offspring. Supporting this idea, we found that prenatal LPS exposure increased the basal  $[Ca^{2+}]_i$  and the number and amplitude of spontaneous oscillations by astrocytes on offspring hippocampus. Notably, the increase of spontaneous astroglial  $[Ca^{2+}]_i$  oscillations and their amplitude was totally blunted by BAPTA-AM but not by gap19, Tat-L2, <sup>10</sup>panx1, probenecid, MTEP or SIB-1757. Moreover, the increase in basal  $[Ca^{2+}]_i$  evoked by prenatal LPS exposure was dependent on the microglia-mediated opening of Cx43 hemichannels and subsequent stimulation of mGluR<sub>5</sub>, but not activation of Panx1 channels. Collectively, these data suggest that spontaneous [Ca<sup>2+</sup>]<sub>i</sub> oscillations evoked by prenatal LPS exposure are necessary for the opening of astroglialCx43 hemichannels, which result in the release of glutamate and the subsequent downstream increase of basal [Ca<sup>2+</sup>]<sub>i</sub> via intracellular stores (Figure 9). Consistent with this, selective blockade of PLC or IP<sub>3</sub> receptors, as well as chelation of  $[Ca^{2+}]_i$ , strongly blunted the increase in basal astroglial  $[Ca^{2+}]_i$  triggered by prenatal LPS exposure. Furthermore, prenatal LPS exposure enhanced the release of glutamate on offspring hippocampus, a response completely dependent on the opening of Cx43 hemichannels but not Panx1 channels. These data concord with previous studies showing the release of glutamate via activation of Cx43 hemichannels (Ye et al., 2003) and with the fact that mGluR<sub>5</sub> controls [Ca<sup>2+</sup>]<sub>i</sub> responses in astrocytes (Panatier and Robitaille, 2016). Previous studies have described that opening of Cx43 hemichannels is regulated by  $[Ca^{2+}]_i$  (De Bock et al., 2012; Meunier et al., 2017). In this line, we noted that chelation of  $[Ca^{2+}]_i$  reduced the prenatal LPS-induced Etd uptake by astrocytes on offspring hippocampus, whereas inhibition of mGluR<sub>5</sub>, PLC or IP<sub>3</sub> receptor did not affect this response.

In the inflamed brain, among other disturbances, astrocytes and neurons undergo the remodeling of their dendritic arbor as well as several morphological changes (Luo and O'Leary, 2005; Pekny and Pekna, 2014; Riccomagno and Kolodkin, 2015). In this work, we detected for the first time that prenatal LPS exposure augments the complexity of astrocyte branch arbors on offspring hippocampus, whereas in neurons occurred the opposite. This evidence harmonizes with studies reporting that neuropathological conditions augment the arborization of hippocampal astrocytes (Beauquis et al., 2013; Chun et al., 2018) and induce dendritic retraction of CA1 pyramidal neurons (Christian et al., 2011; Burak et al., 2018). Usually, the reduction in neurite arborization lined to the loss of branching and a decline in total neurite lengths is accompanied by a decrease in synaptic number and retraction of dendritic spines (Luo and O'Leary, 2005; Rossi and Volterra, 2009). In this context, our experiments revealed that dendritic retraction of CA1 pyramidal neurons triggered by prenatal LPS occurred in parallel with a reduction in the number of apical but not basal dendritic spines in these neurons. It is not clear why the dendritic spine density of basal dendrites is not altered by prenatal LPS exposure. However, this evidence are supported by prior studies describing the layerspecific spine density of CA1 pyramidal neurons (Spruston, 2008) and the opposite regulation of this feature between apical and basal dendrites during different pathological conditions (Hyer and Glasper, 2017; Maynard et al., 2017).

Remarkably, the prenatal LPS-induced alterations on arborization and morphology of astrocytes and neurons were dramatically counteracted with the administration of a specific blocker of Cx43 hemichannels that crosses the BBB. Thus, prenatal LPS exposure elicits divergent morphological effects on offspring astrocytes and neurons that likely reflect their inflammatory status, a phenomenon in which the opening of Cx43 hemichannels seems to be crucial. Substantial levels of glutamate at the synaptic cleft could be neurotoxic under pathological conditions (Lau and Tymianski, 2010). In this context, recent evidence indicates that glutamate released by a mechanism implicating the opening of astroglial Cx43 hemichannels could reduce neuronal survival (Orellana et al., 2011; Yi et al., 2016). Here, we observed that selective inhibition of Cx43 hemichannels greatly prevents the prenatal LPS-induced death of CA1 pyramidal neurons on offspring hippocampus. The latter suggests that Cx43 hemichannels likely contribute to neuronal damage either by altering astrocyte functions (e.g.,  $[Ca^{2+}]_i$  handling) and/or through the release of excitotoxic amounts of glutamate. Supporting this idea, the prenatal LPSinduced overexpression of GFAP, [Ca<sup>2+</sup>]<sub>i</sub> increase and release of glutamate was strongly blunted by blocking Cx43 hemichannels. Excitotoxic levels of glutamate along with a dysfunctional astroglial partnership plausibly could cause neuronal damage via osmotic and  $[Ca^{2+}]_i$  imbalance, as well as caspase activation (Orellana et al., 2011; Moidunny et al., 2016) (Figure 9).

Our findings suggest that opening of astrocyte Cx43 hemichannels occurs at early phases of postnatal life in prenatally LPS-exposed offspring and is accompanied by hippocampal neuroinflammation, as well as diverse astrocyte and neuronal alterations in function and morphology. Future studies are needed in order to elucidate whether astroglial hemichannel/pannexon opening evoked by prenatal LPS exposure may also take place at fetal stages. We speculate that excitotoxic levels of glutamate triggered by the activation of Cx43 hemichannels may contribute to the hippocampal neurotoxicity and damage in prenatally LPS-exposed offspring. Therefore, the understanding of how astrocyte-neuron crosstalk is affected in prenatally LPS-exposed offspring is a promising avenue toward the development of common therapies for several neurological disorders observed in children born to women who had a severe infection during pregnancy.

### DATA AVAILABILITY STATEMENT

The datasets generated for this study are available on request to the corresponding author.

### **ETHICS STATEMENT**

This study was carried out in accordance with the recommendations of "Protocolo de Cuidado y Uso Animal del Comité Ético Cienífico para el Cuidado de Animales y Ambiente." The protocol was approved by the "Comité Ético Científico para el Cuidado de Animales y Ambiente" of the Ponificia Universidad Católica de Chile.

# **AUTHOR CONTRIBUTIONS**

CC, JEO, BA, LM, CI, TA, and JAO conceived, performed, and analyzed the experiments. JAO wrote and edited the manuscript. All authors read and approved the final manuscript.

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# SUPPLEMENTARY MATERIAL

The Supplementary Material for this article can be found online at: https://www.frontiersin.org/articles/10.3389/fncel. 2019.00460/full#supplementary-material

FIGURE S1 | Neurons do not contribute to the prenatal LPS-induced changes in astroglial function observed in the offspring hippocampus. (A) Averaged of DAF-FM signal fluorescence by astrocytes in acute slices from control offspring (white bar) or prenatally LPS-exposed offspring of 4 months old alone (black bars) or in combination with 0.5  $\mu$ M TTX and the following pharmacological agents: 100 μM gap19, 100 μM <sup>10</sup>panx1 and 1 μM L-N6. \*p < 0.0001 versus LPS, one-way ANOVA Tukey's post hoc test, mean  $\pm$  S.E.M., n = 3. (B) Averaged of basal Fluo-4 signal fluorescence by hippocampal astrocytes in acute slices from control offspring (white bar) or prenatally LPS-exposed offspring of 4 months old alone (black bars) or in combination with 0.5  $\mu$ M TTX and the following pharmacological agents: 50 nM minocycline, 100 µM gap19, 100 µM Tat-L2, 100 μM <sup>10</sup>panx1, 500 μM Probenecid (Prob), 10 μM Bapta-AM, 5 μM U-73122, 50 μM 2-APB, 50 nM MTEP, 5 μM SIB-1757, 200 μM oATP, and 200 nM A740003. \*\*p < 0.0001, \*p < 0.005 versus LPS, one-way ANOVA Tukey's post hoc test, mean  $\pm$  S.E.M., n = 3. (C) Averaged data of glutamate release by acute hippocampal slices from control offspring (white bar) or prenatally LPS-exposed offspring of 4 months old alone (black bars) or in combination with 0.5 µM TTX and the following blockers: 100 µM gap19, 100 µM Tat-L2, 100 µM <sup>10</sup>panx1, 500 μM Probenecid (Prob). \*p < 0.0001 versus LPS, one-way ANOVA Tukey's post hoc test, mean  $\pm$  S.E.M., n = 3. (D) Averaged of spontaneous [Ca<sup>2+</sup>]; oscillations by hippocampal astrocytes in acute slices from control offspring (white bar) or prenatally LPS-exposed offspring of 4 months old alone (black bars) or in combination with 0.5  $\mu$ M TTX and the following pharmacological agents: 100 μM gap19, 100 μM Tat-L2, 100 μM <sup>10</sup>panx1, 500 μM Probenecid (Prob), 10  $\mu$ M Bapta-AM, 50 nM MTEP or 5  $\mu$ M SIB-1757. \*\*p < 0.005 versus LPS, one-way ANOVA Tukey's post hoc test, mean  $\pm$  S.E.M., n = 3. (E) Averaged of peak amplitude of spontaneous [Ca2+]; oscillations by hippocampal astrocytes in acute slices from control offspring (white bar) or prenatally LPS-exposed offspring of 4 months old alone (black bars) or in combination with 0.5 µM TTX and the following pharmacological agents: 100 µM gap19, 100 µM Tat-L2, 100  $\mu M$   $^{10} panx1,$  500  $\mu M$  Probenecid (Prob), 10  $\mu M$  Bapta-AM, 50 nM MTEP or 5 μM SIB-1757. \*\*p < 0.005 versus LPS, one-way ANOVA Tukey's post hoc test, mean  $\pm$  S.E.M., n = 3.

**FIGURE S2** | Prenatal LPS exposure does not affect total level and distribution of Cx43 in astrocytes. (**A**,**B**) Representative confocal images depicting GFAP (green) and Cx43 (red) staining by astrocytes in acute slices from control offspring (**A**) or prenatally LPS-exposed offspring of 4-month old (**B**). (**C**-**F**) Insets of astrocytes and examples of draws for staining analysis were taken from the area depicted within the white squares in (**A**,**B**). Calibration bars: white = 120  $\mu$ m; yellow = 20  $\mu$ m. (**G**) Quantification of membrane, intracellular and total staining of Cx43 by astrocytes in acute slices from control offspring (**W**) or prenatally LPS-exposed offspring of 4 months old (black bars). Data were obtained from at least three independent experiments with three or more repeats each one (≥20 cells analyzed for each repeat).

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**Conflict of Interest:** The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

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