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Pannexin Protein Expression in the Rat Middle Cerebral Artery

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Key Words

 Cell-to-cell communication - Cerebrovasculature -Hemichannels - Middle cerebral artery - Pannexins - Rat

Abstract

Background: Connexin proteins are well known to participate in cell-to-cell communication within the cerebral vasculature. Pannexins are a recently discovered family of proteins that could potentially be involved in cell-to-cell communication. Herein, we sought to determine whether pannexins are expressed in rat middle cerebral artery (MCA). *Methods:* A combination of RT-PCR, immunoblotting and immunohistochemistry techniques was used to characterize the expression pattern of pannexins in rat MCA. A fluorescent dye uptake approach in cultured smooth muscle cells was used to determine whether these cells have functional hemichannels. *Results:* We report for the first time that pannexins are expressed in the cerebral vasculature. We reveal that pannexin 1 is expressed in smooth muscle but not in endothelium and pannexin 2 is expressed in both endothelium and smooth muscle. Fluorescent dye entered cultured smooth muscle cells in the absence of extracellular calcium or when the cells were depolarized, which was prevented by the putative hemichannel blocker carbenoxolone. *Conclusions:* The identification of pannexins in rat MCA indicates that

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 Accessible online at: www.karger.com/jvr pannexin expression is not restricted to neuronal cells. Dye uptake in cultured smooth muscle cells exhibited properties similar to those of connexin and pannexin hemichannels, which may represent another form of cell-to-cell communication within the vasculature. Copyright © 2012 S. Karger AG, Basel

Introduction

 Within the cerebral vasculature, cell-to-cell communication is critical for modulating tone and thus blood flow. Large arteries in particular, such as the middle cerebral artery (MCA), play a major role in the regulation of cerebrovascular resistance [1] . Cells can communicate in two different ways, namely by gap junctional communication and paracrine signaling. In gap junctional communication, a direct exchange of signaling molecules via gap junctions (junctional channels) enables intercellular communication. Paracrine intercellular communication, on the other hand, does not involve direct intercellular signaling but rather the release of diffusible signaling molecules (such as calcium and ATP) that act on neighboring cells.

 Historically, connexins were considered the exclusive structural protein of junctional and nonjunctional chan-

 Dr. Elke M. Sokoya Department of Human Physiology, Flinders University Room 6E136, Flinders Medical Centre Flinders Drive, Bedford Park, SA 5042 (Australia) Tel. +61 8 8204 5983, E-Mail elke.sokoya @ flinders.edu.au nels. However, 10 years ago, a new family of proteins called pannexins was identified [2]. Rather than forming gap junctions, pannexins appear to form hemichannels within the plasma membrane [3] and Ca^{2+} -permeable channels in the endoplasmic reticulum [4]. To date, 3 pannexins have been described in the rodent and human genomes, namely pannexin 1 (Panx1), pannexin 2 (Panx2) and pannexin 3 (Panx3). Panx1 is ubiquitously expressed in many organs including the eye, thyroid, kidney, liver and central nervous system [5-7], while Panx2 appears to be more confined to the central nervous system [8, 9]. Panx3 is localized to the skin, osteoblasts and chondrocytes [5]. Within the central nervous system, Panx1 has been reported in neurons and Purkinje cells [3] while Panx2 has been described in neurons [6, 8], and curiously its expression can be induced in astrocytes after ischemia/reperfusion injury [10].

 Local changes in intracellular calcium can be propagated and regenerated along the vasculature, a phenomenon known as calcium wave propagation. Calcium waves have been recorded in both endothelial cells [11, 12] and smooth muscle cells [13, 14]. The propagation of calcium waves involves both the release of ATP through hemichannels or 'nonjunctional channels' (paracrine signaling pathway) and the ensuing local activation of purinergic receptors by ATP to cause an increase in inositol triphosphate, which can then move directly through gap junctions to adjacent cells (direct gap junction pathway) [15]. While gap junctions comprising connexins have been implicated in calcium wave propagation within the vasculature [16], less is known regarding the composition of hemichannels. Pannexin hemichannels are prime candidates for the ATP release channel since, unlike connexin hemichannels, they can open at physiological calcium concentrations [17] and are highly permeable to ATP [18].

 While connexins have been widely reported in cerebral arteries [19–21] and attributed to cerebral vascular function [20, 22, 23], it is unknown whether their expression is shared with pannexins. In the present study, we sought to characterize the expression profile of Panx1 and Panx2 in the rat MCA. Our results indicate that Panx1 is present in smooth muscle alone while Panx2 is expressed in both endothelium and smooth muscle of rat MCA. Moreover, dye uptake studies in cell culture suggest that smooth muscle cells may have functional hemichannels that are comprised of connexins and possibly pannexins. The presence of pannexins within the cerebrovasculature suggests that they may contribute to the regulation of vascular tone in the brain.

Materials and Methods

 Experiments were carried out in accordance with the National Institutes of Health guidelines for the care and use of laboratory animals and were approved by the Animal Protocol Review Committee at Baylor College of Medicine. Male Long-Evans rats (275–325 g) were housed under a 12-hour light/12-hour dark cycle with unrestricted access to food and water.

MCAs were removed from the rat brain, cleaned of connective tissue and placed in a digestion buffer containing (in mmol/l) NaCl (135), KCl (5), $MgCl_2$ (1.5), Na₂HPO₄ (0.42), NaH₂PO₄ (0.44), NaHCO₃ (4.2), HEPES (10) and 1 mg/ml BSA (pH 7.25). MCAs were cut into pieces and enzymatically dissociated with 18 U/ml papain and 1 mg/ml dithioerythritol (35 min at 37° C), 1.2 mg/ml collagenase II, 0.8 mg/ml soybean trypsin inhibitor and 60 U/ml elastase (10 min at 37° C). The tissue was washed several times with digestion buffer between enzymatic incubations. Single smooth muscle cells or small endothelial cell clusters were visually identified with a microscope and isolated by micropipette.

Extraction of Total RNA and cDNA Synthesis from MCAs Two MCAs were harvested from the brain of a male rat under sterile conditions and snap-frozen in liquid nitrogen. Frozen vessels were pulverized with a pestle, and total RNA was extracted using the RNeasy® Micro Kit (Qiagen) according to the manufacturer's instructions. mRNA was reverse transcribed to generate first-strand cDNA using random hexamers. Synthesis of cDNA was performed with SuperScript II (Invitrogen) with the following incubation times: 5 min at 65° C, 12 min at 25° C, 50 min at 42°C and 15 min at 70°C. RNase H was added and incubated for 20 min at 37° C to degrade any remaining RNA.

Extraction of RNA and cDNA Synthesis from Single Cells

 MCAs were enzymatically digested (see Cell Digestion Protocol above), and the cells were viewed on the stage of an inverted microscope. With the aid of suction, a micropipette [prefilled with RNAlater (Qiagen)] was used to specifically draw either 50– 100 endothelial cells or smooth muscle cells into the tip. The cells were then expelled into a PCR tube and snap-frozen in liquid N_2 . cDNA was synthesized with a one-step method. The following reagents were added directly to the tube containing $2 \mu l$ of cells: 2μ l of 5 \times 1st strand buffer, 0.5 μ l of 10 mmol/l deoxynucleotide triphosphate, 1 μ l of 50 μ mol/l random hexamer, 1 μ l of 0.1 mmol/l DTT, 1 µl of 10 U/µl RNase Inhibitor and 1.5 µl of diethylpyrocarbonate-treated water. The cells were lysed by exposing them to 2 freeze-thaw cycles, and then 1μ l of 50 U/ μ l Super Script II Reverse Transcriptase (Invitrogen) was added. cDNA was synthesized during a 1-hour incubation at 42° C.

Polymerase Chain Reaction

PCR amplification was performed with 0.2μ l of Platinum Taq DNA polymerase, $1 \times$ PCR Buffer, 1.5 mmol/l MgCl₂, 0.2 mmol/l deoxynucleotide triphosphate mixture and 1μ mol/l of each primer for 40 cycles. Primer pairs were as follows: 5'-TAAAC-CCCA-GCTATGGAGCCA-3' (forward) and 5'-GGCGTCAGT-AAAATCCCGTTC-3' (reverse) for Panx1; 5'-GGCCCCGTA-AGAAAATGAAGT-3 (forward) and 5 -TGTATCCGTGGCT-

GT-CTTCCT-3 (reverse) for Panx2; 5 -AGCTGTCACATCA-CCCCACT-GT-3' (forward) and 5'-TGTGCGAGTGACATGA-TGCTG-3' (reverse) for Panx3, and 5'-GGCTGCCTTCTCT-TGTGACAA-3' (forward) and 5'-CGCTC-CTGGAAGATGG-TGAT-3 (reverse) for GAPDH. All primer pairs were designed using Primer Express (Applied Biosystems, Calif., USA), with the exception of GAPDH [24]. The annealing temperature was 55° C for Panx1, 60°C for Panx2 and 56°C for Panx3. For the negative control, reverse transcriptase was omitted. The RT-PCR products were analyzed by electrophoresis on a 1.8% agarose gel containing ethidium bromide, in parallel with a 50-base pair DNA ladder (Invitrogen). The DNA fragments were visualized and photographed using a UV light table/camera documentation system. The predicted sizes of the PCR amplification products specific to Panx1, Panx2, Panx3 and GAPDH are 125, 140, 101 and 180 base pairs, respectively. Sequencing was performed at the Child Health Research Center Core Laboratory (Baylor College of Medicine).

Protein Isolation **Protein Isolation**

 The MCAs were harvested from the brain, cleaned of connective tissue and snap-frozen in liquid nitrogen. Pooled frozen MCAs were pulverized with a pestle, homogenized in a modified RIPA buffer and centrifuged (8 min at 14,000 rpm). A portion of the supernatant was used for total protein analysis (modified Lowry assay) and a portion was loaded onto the gel.

Western Gel and Transfer

 MCA homogenate was combined with Laemmli buffer and boiled for 5 min. Then, 2.5-20 µg of protein was loaded onto each lane of the gel (7.5% SDS-PAGE gel). The MCA homogenate was run in parallel with a standard (Precision Plus Protein Dual Color Standard, Invitrogen). The gel was run at room temperature at 90 V for 1.5 h and then transferred onto an Immobilon-P membrane (Millipore) at 4°C (1.5 h at 250 mA with stirring). The membrane was blocked in 5% Bio-Rad Blotting Grade Blocker containing 0.1% Tween-20 in $1 \times$ PBS (BB) for 1 h. This was followed by incubation overnight at 4°C in primary antibody diluted in 50% BB (1:20,000 Panx1 dilution or 1:50,000 Panx2 dilution) against each pannexin protein. The membrane was washed in PBS and incubated in secondary antibody (antirabbit IgG, 1:10,000 dilution with 50% BB) conjugated to horseradish peroxidase. The membrane was then placed in Pierce SuperSignal® West Femto Maximum Sensitivity Substrate followed by exposure to film. To confirm that each lane contained equivalent amounts of protein, the membrane was stripped with RestoreTM Western Blot Stripping Buffer (Pierce) and then probed for anti-GAPDH (Calbiochem; 1:100,000 dilution). Rat brain was used as a positive control [5, 6]. For negative controls, we (1) omitted the primary antibody, (2) incubated the homogenate with nonimmune rabbit IgG and (3) incubated the homogenate with the appropriate immunogenic peptide.

Generation of Pannexin Antibodies

 For the generation of a rat Panx1 (rPanx1) and a rat Panx2 (rPanx2) antibody, two amino acid peptide sequences, CMSLQT-KGEDQGSQR (corresponding to amino acids 383–396) and CSAEPPVVKRPRKKM (corresponding to amino acids 408– 421), respectively, were selected for their immunogenicity using PCGENE. BLAST database searches indicated that these sequences were unique to their respective protein. Each peptide was synthesized and conjugated via a cysteine residue added at the N terminus during synthesis. Two rabbits were immunized to each peptide, and the sera were affinity purified. Affinity purification yielded final antibody concentrations of 0.5 mg/ml (rPanx1) and 1.5 mg/ml (rPanx2). Synthesis of peptides, rabbit immunization and affinity purification of the antibodies were performed by Sigma-Genosys (The Woodlands, Tex., USA).

Preparation of MCA Cross-Sections

 Rats were anesthetized with pentobarbital (50 mg/kg, i.p.). The vascular system was perfused by a transcardial approach with heparinized saline solution (20 units/ml) at a pressure of 80 mm Hg. The brain was removed and a wedge containing the MCA was harvested and snap-frozen in dry ice-chilled 2-methylbutane. Cryosections (10 μ m thick, -20°C) were mounted onto Superfrost Plus slides (Fisher Scientific), air dried and stored at –80° C.

Immunofluorescence Light Microscopy

 MCA cross-sections were washed in PBS and then fixed with 4% paraformaldehyde in PBS for 30 min at 4°C. Following a wash step in PBS, sections were permeabilized (0.1% Tween-20) and blocked (10% goat serum, 0.5% BSA). Sections were incubated with either Panx1 primary antibody (1:35 dilution), Panx2 primary antibody (1:300 dilution) or rabbit IgG (Jackson ImmunoResearch Labs) overnight at 4°C. Immunohistochemical labeling was demonstrated by incubation with Alexa Fluor 594-conjugated goat antirabbit IgG (1:500 dilution) for 30 min at room temperature. Specimens were incubated (10 min at room temperature) in 1 μ mol/l DAPI (Molecular Probes) for nuclei detection, rinsed in PBS, mounted using Airvol and allowed to dry in air overnight. Pannexin immunolabeling was captured with an epifluorescence microscope (Nikon TE2000U) and deconvolved using AutoDeblur software (Media Cybernetics Inc).

Smooth muscle cells were cultured from the rat thoracic aorta and used after the first passage. The rat aorta was dissected and transferred to a sterile dish containing cold DMEM and antibiotics (100 U/ml penicillin, 100 mg/ml streptomycin). The connective tissue was removed and the aorta pinned onto a silicone-covered preparation dish. The vessel was cut open longitudinally and the endothelium was gently removed with curved forceps. The medial layer was then stripped off in small pieces and deposited in a T25 culture flask which was set upright in a tissue incubator. After 24–48 h, the flask was placed horizontally, and culture medium (DMEM/F12 containing 10% FBS, 100 U/ml penicillin and 100 mg/ml streptomycin) was changed every 48 h. The media explants were removed between days 7 and 10 before the cells were passaged and plated onto 35-mm petri dishes at low densities so that single cells predominated.

 Dye Uptake Experiments Cultured rat aorta smooth muscle cells were used at passage 1. Cells were exposed to either HEPES buffer with 1 mmol/l $CaCl₂$ (normal calcium), HEPES buffer with no $CaCl₂$ and 2 mmol/l EGTA (zero calcium) or high-potassium buffer. For the high-potassium experiments, the culture medium was replaced with either HEPES buffer (normal K^+) containing (in mmol/l) NaCl

Fig. 1. RT-PCR products for Panx1 (125 bp), Panx2 (140 bp) and Panx3 (101 bp) were found in rat brain and rat MCA. GAPDH (180 bp) was used as an internal control. Lane 1: 50-bp ladder (size marker). No product was detected if reverse transcriptase was omitted (data not shown). Data are representative of 3 separate preparations.

(120), potassium gluconate (4.2), $\text{Na}_2\text{HPO}_4(0.9)$, sodium pyruvate (1.2), glucose (10), HEPES (20) and CaCl₂ (1) or isotonic high-K⁺ buffer containing (in mmol/l) NaCl (4.2), potassium gluconate (120), Na_2HPO_4 (0.9), sodium pyruvate (1.2), glucose (10) and HEPES (20). Lucifer Yellow was added to the bath at a concentration of 0.5% for 5 min. After a brief wash, Lucifer Yellow was imaged with an epifluorescence microscope. In some experiments, hemichannels were blocked by adding 100μ mol/l carbenoxolone. The integrity of cell membranes was confirmed by adding Lucifer Yellow in conjunction with 0.4% rhodamine B isothiocyanatedextran.

Results

 Expression of Pannexin Message in Rat MCA Transcripts for Panx1 and Panx2 were detected in both rat brain and rat MCA (fig. 1). While Panx3 was also detected, this was only marginal. PCR products were not detected when template cDNA was omitted from the PCR or when reverse transcription reactions were performed in the absence of reverse transcriptase (data not shown). Sequencing of the pannexin products indicated that the bands corresponded to Panx1 (NM_199397), Panx2 (NM_199409) and Panx3 (NM_199398). In freshly dissociated endothelial cells, message encoding Panx2 but not Panx1 was detected (fig. 2a). RT-PCR products indicating the presence of endothelial nitric oxide synthase and absence of $SM22\alpha$ confirmed the purity of the endothelial cell sample. Freshly dissociated smooth muscle cells demonstrated PCR product for both Panx1 and Panx2 message (fig. 2b). The purity of the smooth muscle cell preparation was confirmed by the presence of PCR amplicons for SM22 α but not for endothelial nitric oxide synthase.

To determine whether pannexin protein was expressed in rat MCA, we used immunoblotting and generated our own rabbit polyclonal anti-Panx1 and anti-Panx2 antibodies. Since our RT-PCR data showed very weak message for Panx3, we decided to focus on the expression profile of Panx1 and Panx2.

 To investigate Panx1 protein expression in the rat MCA, protein extracts were assayed for Panx1 expression by Western blot. Panx1 was weakly expressed in rat MCA, although GAPDH protein expression was confirmed in our sample (fig. 3a). No band was detected in either rat brain or rat MCA when the primary antibody was incubated with the control peptide (fig. 3a), when the primary antibody was replaced with nonimmune rabbit IgG or when the primary antibody was omitted (data not shown). By contrast, affinity-purified rPanx1 antisera recognized a single band at 58 kDa in rat brain, tissue known to express high levels of Panx1 [5, 6] (fig. 3a), but not in lung (online suppl. fig. S1; for all online suppl. material, see www.karger.com/doi/10.1159/000332329), tissue known to lack Panx1 [5]. These results are consistent with previous studies in rat brain showing a 58-kDa isoform that is associated with the plasma membrane [9, 25] . The 58-kDa band is 11 kDa larger than the predicted size but has been shown to represent the glycosylated protein expressed on the cell surface [26] .

 To investigate Panx2 protein expression in the rat MCA, protein extracts were assayed for Panx2 expression by Western blot. Panx2 was shown to be highly expressed in rat MCA (fig. 3b). Each band was absent when the primary antibody was incubated with the control peptide (fig. 3b), when the primary antibody was replaced with

Fig. 2. RT-PCR product for Panx2 (140 bp) was found in freshly dissociated endothelial cells (ECs; **a** , left panel) and RT-PCR products for Panx1 (125 bp) and Panx2 (140 bp) were found in freshly dissociated smooth muscle cells (SMCs; **b**, left panel) from rat MCA. The right panels show RT-PCR products indicating the absence of

SM22 α (325 bp) and the presence of endothelial nitric oxide synthase (eNOS; 356 bp) in dissociated endothelial cells (a, right panel) and vice versa in dissociated smooth muscle cells (**b**, right panel). No product was detected if reverse transcriptase was omitted (data not shown). Data are representative of 3 separate preparations.

50 37 Brain MCA Brain MCA Panx2 Panx2 antibody Control peptide 75 kDa kDa $GAPDH$ 37 **b**

Fig. 3. a Tissue extracts from rat brain $(10 \mu g)$ and MCA $(20 \mu g)$ were run on 10% SDS-PAGE gels and probed with Panx1 antibody in the absence and presence of the control peptide. **b** Tissue extracts from rat brain (10 μ g) and MCA (2.5 μ g) were run on 10% SDS-PAGE gels and probed with Panx2 antibody in the absence

and presence of the control peptide. Each band disappeared when the primary antibody was incubated with the control peptide. Each membrane was stripped and reprobed for GAPDH. The molecular mass is indicated. Data are representative of 4 separate preparations.

Fig. 4. Pannexin immunofluorescence in frozen cross-sections of rat MCA. **a** Panx1 fluorescence (red) showed weak staining in smooth muscle (arrow) and was undetected in endothelial cells. **b** Red fluorescence was absent in sections in which the control rabbit IgG was substituted for the primary antibody. **c** Panx2 fluorescence (red) was detected in both endothelial cells (bottom two arrows) and smooth muscle cells (top arrow). **d** Red fluorescence was absent in sections incubated with a control rabbit IgG. Green fluorescence is the autofluorescence of the internal elastic lamina that separates the endothelium from smooth muscle. Blue fluorescence identifies nuclei labeled with DAPI. Data are representative of 3 separate preparations. Scale bar = 20 μ m. (Color only in online version).

nonimmune rabbit IgG or when the primary antibody was omitted (data not shown). Affinity-purified rPanx2 antisera recognized a single band at 47 kDa in rat brain, tissue known to express Panx2 at high levels [5, 6], but not in lung (online suppl. fig. S1), tissue known to lack Panx2 [5].

To characterize the localization of Panx1 within the cells of the vascular wall, immunohistochemical staining

was performed on frozen MCA cross-sections. Panx1 immunofluorescence showed a weak signal in the smooth muscle of MCA cross-sections and no signal in the endothelium (fig. 4a). These findings are consistent with the RT-PCR data in freshly dissociated vascular cells (fig. 2).

 In frozen cross-sections, Panx2 fluorescence was detected in both endothelial cells and smooth muscle cells of rat MCA (fig. 4c). Again this is consistent with the detection of Panx2 message in endothelial and smooth muscle cell populations (fig. 2).

Primary cultured rat smooth muscle cells were confirmed as smooth muscle cells based on positive staining for both smooth muscle cell α -actin (online suppl. fig. S2) and myosin heavy chain and absence of staining for von Willebrand factor (data not shown). Note that fibroblasts are positive for smooth muscle cell α -actin but negative for myosin heavy chain. These data underscore the high purity of smooth muscle cells in our preparation. These cells also showed positive staining for connexin (Cx) 43 (online suppl. fig. S2), and Western blot analysis confirmed that they expressed Panx2 (online suppl. fig. S3).

 The functional state of hemichannels in cultured smooth muscle cells was assessed by the cellular uptake of Lucifer Yellow from the extracellular media. When cells were exposed to zero extracellular calcium solution, they showed a significant increase in Lucifer Yellow fluorescence which was prevented in the presence of the hemichannel blocker carbenoxolone (fig. 5a). Similar findings were also seen in response to depolarization with high potassium (fig. 5b). Lucifer Yellow fluorescence was absent when dextran-rhodamine was added to the bath, indicating that cell injury does not account for the dye uptake since dextran-rhodamine (3,000 Da) is too large to move through hemichannels.

Discussion

 Within the cerebral vasculature, cell-to-cell communication is critical for modulating tone and thus blood flow. There is abundant literature implicating a role for connexin gap junctions in promoting endothelial calcium changes in response to smooth muscle-dependent agonists [27], conducted vasomotor responses [22, 28], myogenic responses [23, 29], cerebral vasomotion [20] and endothelium-dependent dilations [21, 30]. However, it is currently not known whether pannexin hemichannels are present within the cerebral vasculature.

Color version available online

 The present study provides an initial characterization of pannexin expression in the rat MCA. We provide the following new data: (1) Panx1 protein is weakly expressed in smooth muscle but not in endothelium; (2) Panx2 protein is expressed in both endothelium and smooth muscle, and (3) cultured smooth muscle cells exhibit functional properties similar to those of connexin and pannexin hemichannels.

 To our knowledge, this is the first report of pannexin expression within the vasculature of the brain. As this paper was being reviewed, a report was published demonstrating that Panx1, but not Panx2, is expressed in thoracodorsal resistance arteries [31] . Interestingly, evidence was also presented suggesting an association between Panx1 and the α_1 -adrenergic receptor where Panx1 opens to release purines to enhance phenylephrine-induced vasoconstriction.

 Previous studies from our laboratory have shown that the endothelium of rat MCA expresses Cx40 and Cx43 while the smooth muscle expresses Cx37 and Cx40 [21]. Other cell types have also been found to coexpress connexins and pannexins. For example, cortical astrocytes express both Cx43 [32] and Panx1 [33] . However, erythrocytes express Panx1 but not connexins [34] . Connexin expression is known to be dependent upon the vascular bed and even species studied. Therefore, it will be interesting to see whether this also holds true for pannexin expression.

 It is currently difficult to unequivocally demonstrate the presence of functional hemichannels. Unfortunately, there are no pharmacological agents that can discriminate between connexins and pannexins, and even the more specific mimetic peptides have been shown to be promiscuous [35]. Carbenoxolone is a derivative of glycyrrhetinic acid that blocks both connexin and pannexin hemichannels [36]. It has been shown to block dye uptake and ATP release in pannexin-expressing cells [34] . Carbenoxolone has also been shown to block volume-regulated anion channels [37] and $P2X_7$ receptors [38]; however, Pelegrin and Surprenant [39] showed that the latter was due to the functional coupling between Panx1 and $P2X₇$ receptors and that carbenoxolone had no effect in single astrocytes transfected with $P2X_7$.

 Our fluorescent dye uptake results in cultured smooth muscle cells are consistent with the presence of functional hemichannels. Note that these cells were plated at low densities, thereby ruling out a potential gap junction-mediated transfer of dye between cells. Connexin, but not pannexin, hemichannels are sensitive to changes in extracellular calcium [17, 39]. Therefore, the increase in Lucifer Yellow fluorescence seen in smooth muscle cells after depletion of extracellular calcium suggests connexin hemichannel opening (fig. 5a). This is most likely mediated by Cx43 since (1) this is the predominant connexin protein expressed in rat aorta smooth muscle [40] (online suppl. fig. S2) and (2) Lucifer Yellow dye uptake was blocked in the presence of carbenoxolone.

 Under normal extracellular calcium conditions, depolarization of cultured smooth muscle cells with high potassium caused an increase in cellular Lucifer Yellow fluorescence which was blocked by carbenoxolone (fig. 5b). Since Cx43-expressing cells have failed to show hemichannel currents in response to depolarizing voltages [41, 42], our data are consistent with the activation of pannexin hemichannels.

 During pathological insults such as ischemia, vascular connexin hemichannels could be activated during episodes of significantly reduced extracellular calcium. On the other hand, pannexin hemichannels could be activated during conditions of cellular depolarization such as traumatic brain injury [43] , stroke [44] and subarachnoid hemorrhage [45]. Under normal conditions, Panx1 could contribute to the propagation of calcium waves within the vasculature by mediating the release of ATP [18] and as such play an important role in mediating vascular tone. In addition to forming hemichannels on the plasma membrane, recent findings have also indicated that pannexins form calcium-permeable channels within the endoplasmic reticulum [4, 46]. Given the fundamental role of calcium homeostasis in controlling vascular tone and cerebral blood flow [47, 48], pannexins may play a vital role in the regulation of vasodilatation.

 The identification of pannexins in the rat MCA indicates that pannexin expression is not limited to neuronal/ astrocytic cell populations within the central nervous system. Further studies are required to understand the role of pannexin hemichannels within the vasculature and their functional differences compared to connexin hemichannels. The possibility that pannexins may be an important therapeutic target during pathologies such as stroke and traumatic brain injury warrants further investigation.

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