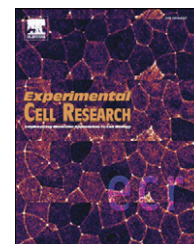


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Review

Cell membrane permeabilization via connexin hemichannels in living and dying cells

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ABSTRACT

Vertebrate cells that express connexins likely express connexin hemichannels (Cx HCs) at their surface. In diverse cell types, surface Cx HCs can open to serve as a diffusional exchange pathway for ions and small molecules across the cell membrane. Most cells, if not all, also express pannexins that form hemichannels and increase the cell membrane permeability but are not addressed in this review. To date, most characterizations of Cx HCs have utilized cultured cells under resting conditions and revealed low open probability and unitary conductance close to double that of the corresponding gap junction channels. In addition, the cell membrane permeability through Cx HCs can be markedly affected within seconds to minutes by various changes in the intra and/or extracellular microenvironment (i.e., pH, pCa, redox state, transmembrane voltage and intracellular regulatory proteins) that affect levels, open probability and/or (single channel) permeability of Cx HC. Net increase or decrease in membrane permeability could result from the simultaneous interaction of different mechanisms that affect hemichannels. The permeability of Cx HCs is controlled by complex signaling cascades showing connexin, cell and cell stage dependency. Changes in membrane permeability via hemichannels can have positive consequences in some cells (mainly in healthy cells), whereas in others (mainly in cells affected by acquired and/or genetic diseases) hemichannel activation can be detrimental.

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Introduction

Connexin hemichannels (Cx HCs)

Connexin hemichannels are hexameric oligomers of transmembrane proteins termed connexins (Fig. 1), and docking of two hemichannels in apposed membranes forms intercellular gap junction channels. When open at the unapposed cell surface, non-junctional hemichannels are aqueous pores (Fig. 1) permeable to ions and small molecules that allow diffusional exchange between the intra and extracellular compartments and also constitute a route for autocrine/paracrine cellular communication [1,2]. Cx HCs and gap junction channels frequently are oppositely influenced by various experimental and physiological conditions [3–8].

Although synthesis and trafficking pathways of Cx HCs might differ from those of intercellular channels, hemichannels formed

by most studied connexins (all but Cx26 [9]) oligomerize in the Golgi/trans-Golgi network [10–13]. Upon assembly, hemichannels (except Cx26 HCs [9]) are transported to the non-junctional plasma membrane via a cytoskeleton-mediated route [14–16]. Once inserted in the plasma membrane, they diffuse laterally to join the external aspect of gap junctional plaques [17,18], where they dock with hemichannels from a neighboring cell to form intercellular channels. However, Cx43 HCs may also be directly targeted to the region where gap junctional plaques are found via a microtubule/dynein/ β -catenin/N-cadherin-dependent pathway [19]. It might be possible that hemichannels in non-junctional membrane and those in gap junctions are derived from different precursor pools. In support of this notion, fluorescent-tagged Cx43 can be delivered to cell protrusions or to cell surface domains that lack a contacting cell [20].

In cells expressing wild-type connexins, changes in surface hemichannel levels and/or intrinsic properties of hemichannels

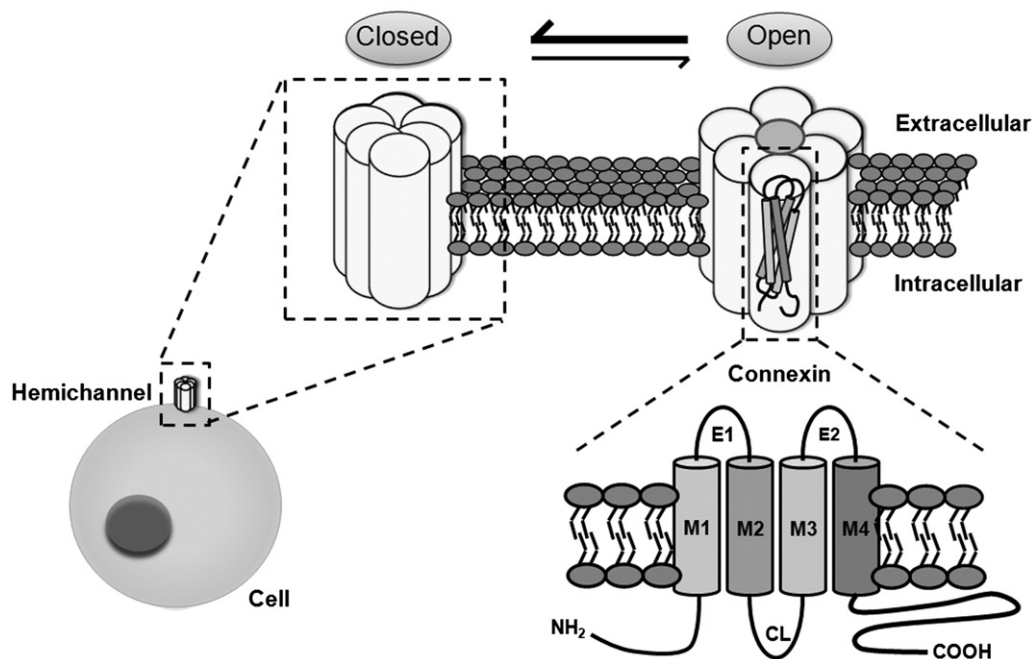


Fig. 1 – Diagram illustrating basic structures of connexins and undocked hemichannels present at the cell surface. The membrane topology of a connexin consists of 4 membrane-spanning domains (M1–M4), 2 extracellular loops (E1 and E2) and 1 cytoplasmic loop (CL). The amino ($-\text{NH}_2$) and carboxy ($-\text{COOH}$) terminal tail are intracellular. A hemichannel is formed by six connexins that oligomerize laterally leaving a central pore. In cultured cells under resting conditions hemichannels remain preferentially closed, but they can be activated by diverse physiological and pathological conditions, offering a diffusional transmembrane route between the intra and extracellular milieus.

found at the cell surface (e.g., open probability and permeability) can significantly affect the cell membrane permeability. Increases in hemichannel levels have been clearly associated with rises in intracellular-free Ca^{2+} concentration ($[\text{Ca}^{2+}]_i$) [6,21]. Variations in hemichannel open probability can result from changes in covalent modifications of connexin subunits (e.g., redox and phosphorylation state) or variations in transmembrane physicochemical conditions (e.g., transmembrane voltage, pH and concentration of monovalent cations) that affect the pore forming proteins. Although changes in hemichannels unitary conductance upon application of specific stimuli have not yet been reported, they are likely to occur in Cx43 HCs after changes in phosphorylation state that reduce their size cutoff [22]. Additionally, several connexin mutations affect the open probability and/or permeability of hemichannels [23–31] and in most cases possible permeability changes have not been yet evaluated.

Cultured cells (primary cultures or cell lines) under “resting” conditions show low hemichannel activity defined as low frequency of unitary current events and/or low membrane permeability to small molecules (e.g., ATP and small fluorescent dyes). However, cells within organisms are exposed to diverse stimuli, which may transiently increase the hemichannel activity. Moreover, there may also be endogenous hemichannel inhibitors that contribute to control transient increases in hemichannel activity. With regard to positive regulators, increased hemichannel activity has been recorded in cells exposed to endogenous ligands associated with cell proliferation or differentiation, such as lipophilic molecules and growth factors [4,6,32]. Thus, the activity of hemichannels in cultured cells under “resting” conditions might not reflect the importance of these transmembrane pathways in cells *in vivo*.

Activation of hemichannels has also been associated with cell death under diverse pathological conditions (for review see [33]). In acquired diseases, the primary cause of cell death is directly related to the pleiotropic effects of the injury, and a high level of hemichannel activity makes the cells more susceptible to the injury (e.g., generation of reactive oxygen species, ATP depletion and increase in $[\text{Ca}^{2+}]_i$ as occurs after ischemia-reperfusion). In contrast, sustained and uncompensated high hemichannel activity found in human genetic diseases associated with point connexin mutations and gain of hemichannel function could be the primary cause of death, as observed in cells transfected with mutant connexins [23–25,27–29,31]. Under these conditions, the deleterious effect of hemichannel activation is possibly related to the long lasting and unbalanced Ca^{2+} influx leading to activation of intracellular Ca^{2+} -dependent hydrolases (e.g., proteases, phospholipases and endonucleases) and cellular depletion of vital small solutes and metabolites such as ATP and glutathione.

Do cells in tridimensional tissues express functional hemichannels in their surface? To our knowledge, demonstrations of hemichannel activity *in vivo* or in tissue slices bathed in extracellular solution containing divalent cations have not been reported. However, ATP is released through hemichannels in whole-mount retinal pigment epithelium [34], and high Cx43 HC activity occurs in colonocytes during infectious diarrhea where it may mediate water release [35].

Structure–function relationship

Connexin proteins are encoded by at least 21 different genes in humans and are named according to their predicted molecular

mass in kDa from Cx23 to Cx58 [36]. All studied connexins share a common membrane topology with four transmembrane domains, two extracellular loops, one intracellular loop, and both N- and C-terminus in the cytosolic side of the cell membrane (for a review, see [37]). The main divergences in length and amino acid sequence among connexins are located in the C-terminal tail and cytoplasmic loop. Several recent studies have focused on how each connexin protein domain affects the functional state of hemichannels (see below).

Post-translational modifications of connexins affect preferentially amino acid residues located in the C-terminus and include palmitoylation, myristoylation, nitrosylation, protease cleavage and phosphorylations [38–42]. But the possible role of most of these covalent modifications on hemichannels remains unknown. Phosphorylation is the most extensively studied covalent modification of connexins (for reviews, see [43,44]). Electrophysiological studies of Cx43 HCs using PKC inhibitors suggested that activation of a novel PKC isoform completely inhibits hemichannels, whereas activation of a conventional isoform (βII) only partially reduces the hemichannel-mediated currents [45]. Moreover, MAP kinase or PKC-mediated phosphorylation of Cx43 HCs reconstituted in liposomes reduces their permeability to molecules [22,46], and PKC-mediated phosphorylation of Cx43 at serine 368 is required to decrease the hemichannel-mediated permeabilization in this model [47]. In addition, opening of Cx46 HCs expressed in *Xenopus* oocytes is also decreased by PKC-mediated protein phosphorylation [48].

In astrocytes and exogenous expression systems, S-nitrosylation of Cx43 and Cx46, i.e., covalent binding of NO to Cys, affects the activity of hemichannels [39,49]. In cortical rat astrocytes, treatment with the NO donors, GSNO or Nor-1 or metabolic inhibition increases hemichannel-mediated dye uptake [39]. NO is also involved in Cx43 HC-mediated cell permeabilization induced by proinflammatory cytokines in astrocytes [5] or Cx43 transfected C6 glioma cells [4]. Although NO does not affect the open probably of Cx46 HCs under control condition, it changes the opening and closing kinetics at positive voltages as well as the permeability to large molecules [49]; the possible relevance of NO 46 HCs expressed by lens fibers remains to be demonstrated. Proteolysis of Cx46 is another putative post-translational modification that might affect Cx46 HCs. In support to this possibility, it has been shown that proteolysis shifts the pK_a of gap junction channels formed by rat Cx46 from 6.8 to 6.6 [50], but its functional consequences on Cx46 HCs remains to be studied.

In gap junction channels formed by Cxs 43 or 45, the C-terminal tail may participate in channel gating induced by intracellular acidification or membrane potential through a “ball and chain mechanism” [51,52]. Moreover, the Cx43 C-terminus binds to the second half of the cytoplasmic loop in a pH-dependent action [54]. It remains unknown whether similar conformational changes account for the hemichannel sensitivity to intracellular pH variations. In support of this possibility, intracellular acidification rapidly reduces Cx46 HC current in *Xenopus* oocytes [53]. In addition, nuclear resonance magnetic experiments reveal that acidification induces α -helical structure and protonation of histidine residues in a Cx43 peptide corresponding to the second half of the intracellular loop, without mayor structural changes in the C-terminus.

In ^{15}N -HSQC NMR analyses, the carboxy terminal (CT) domain of Cx43 was shown to be highly disorganized with two α -helical

regions one at Ala³¹¹–Ser³²⁵ and the other at Asp³³⁹–Lys³⁴⁵ [55]. The CT can interact with several intracellular and membrane proteins, such as ZO-1 [56], ZO-2 and ZO-3 [57], ubiquitin protein ligase, Nedd4 [58], several protein kinases (for reviews, see [43,44]), aquaporin-0 [59] and NOV, a member of the CCN gene family [60]. The interaction between connexins and other proteins has been supported by co-localization and pull-down experiments [61,62]. However, it has not been determined whether connexins forming surface hemichannels interact with any proteins other than protein kinases. Truncation of Cx43 CT changes the sub-cellular localization, number and size of Cx43 gap junctions but does not prevent their assembly into hemichannels [6,63]. Accordingly, mutant Cxs 32, 43, 46 and 50 lacking most of their CT form functional hemichannels and gap junction channels [6,39,64,65], showing that the CT is not essential for channel function. Hemichannels and gap junction channels formed by most CT mutants studied show relevant differences in their electrophysiological and permeability properties [6,39,64,65]. Truncation of Cx43 C-terminus affects the location, decreases the number and increases the size of gap junction plaques in cardiomyocytes, suggesting its involvement in the regulation of critical steps in hemichannels trafficking [63]. In addition, the Cx43 CT alone seems to play a regulatory function in glioma cell invasion [66], inhibition of cell growth [67], determination of the infarct size and susceptibility to ventricular tachyarrhythmias after acute myocardial infarction [68] and neuroprotection during cerebral ischemia [69]. To our knowledge, similar studies addressing the possible role of surface hemichannels have not been reported and it remains to be determined the extent to which many biological responses are mediated by gap junction channels or hemichannels or both.

Much less is known about the functional role of the N-terminal tail of connexins. Mutation of the N-terminus of different connexins affects some hemichannel properties including gating polarity [70], charge selectivity [71] and perm-selectivity [72] as well as formation of functional hemichannels [73]. Cx43 tagged with enhanced green fluorescent protein (EGFP) on its N-terminus shows formation of gap junction plaques but fails to form functional gap junction channels or hemichannels in HeLa cells transfectants [74]. The latter might be due to a pore blocking action of the altered N-terminal tail, as has been proposed for channels composed of wild-type Cx26, in which the N-terminus forms a plug that physically closes the pore [75]. The N-terminus has α -helical region [76,77], which at least in Cx32 has a flexible turn around glycine 12 allowing the N-terminus to extend into the channel vestibule [78]. Together, these data indicate that the N-terminus is important in formation of Cx HCs as well as in control of their functional state.

The intracellular loop (IL) of connexins has been less studied. However, it plays a major role in pH sensitivity of gap junction channels composed of Cx38 [79]. Functionally, the IL has been shown to participate in the modulation of permeability, conductance and voltage gating of Cx26 and Cx30 based gap junction channels [80]. Probably, the IL affect functions (but not formation) of gap junction channels [81] because it interacts directly with the connexin C-terminus [82,83] and other proteins such as aquaporin-0, calmodulin, beta-tubulin and possibly cadherins [84–87]. This intramolecular interactions (IL–C-terminus) can be modulated by changes in intracellular pH [88]. Moreover, the IL of fish Cx35 and chicken Cx56 can be phosphorylated [89,90], suggesting

that channel functions can be modulated through changes in post-translational covalent modifications of the IL. Since phosphate can be protonated, which changes the net charge of the phosphorylated amino acid residues, this covalent modification may affect other hemichannels properties such as pH and voltage sensitivity.

Each extracellular loop (EL) has three conserved cysteine residues (Cys), which can form disulfide bridges with Cys located in the same or other loops [91]. The ELs, are implicated in slow hemichannel gating (also call loop gating) [92], which is modulated by extracellular divalent cations [93]. Loop gating involves movement of the TM1-E1 region resulting in a reduction of diameter of the hemichannel lumen [94], mainly due to a rotation and inward tilt of all six M1 segments [95]. Interestingly, different peptides sharing the EL sequence block Cx HCs in diverse models [5,96–98] and prevent the formation of new gap junction channels [99,100]. However, a nonspecific allosteric effect was proposed as the mode of action of these peptides [101]. More selective peptides for pannexin hemichannels are now available [102], and future studies might provide similar reagents for Cx HCs together with more knowledge on their mechanism of action.

The transmembrane domains (TM) are believed to form the hemichannel aqueous pore, but little clarity exists regarding the specific amino acid residues lining it. Important advances have been made using SCAM (substituted Cys accessibility method) experiments. Prominent reduction in the hemichannel activity occurred after binding of the sulfhydryl-specific methanethiosulfonate MTS to the M1 domain of Cx46 [103] and M3 domain of Cx32 [104], suggesting that these TMs line the pore of hemichannels formed by those connexins. However, the recently reported 3.5 Å resolution map of wild-type human Cx26 [77], indicate that M3 and M4 are facing the hydrophobic membrane environment, while M1 and M2 line the pore interior.

Permeability

Most studies of hemichannel permeability have measured the uptake of synthetic molecules with different mass, shape and net charge such as Lucifer yellow [46,105], ethidium [3,5,6,74], calcein [106], 5(6)-carboxyfluorescein [47,107] and DAPI [108]. Release or uptake measurements of biologically relevant molecules such as ATP [109,110], glutamate [111], NAD⁺ [112], glutathione [113], prostaglandin E₂ [114], 2-NBDG [5] and ascorbate [115] have also provided information on permeability of hemichannels. For paracrine/autocrine actions, transient opening of hemichannels may allow release of enough signaling molecule to reach effective concentrations in a confined discrete extracellular space. There is still the question of whether Cx HCs provide a diffusional pathway for passive uptake of metabolites, generally thought to be mediated by relatively specific membrane transporters. An answer may be found through calculation of apparent affinity constants and maximal transport velocity of hemichannels for particular permeant molecules, which could be compared to those of transporters for the same molecules (e.g., glucose).

Is permeability a fixed or a plastic feature of hemichannels? Recent evidence indicates that hemichannel permeability can be affected by covalent modification of the hemichannel subunits, which likely cause conformational changes in the hemichannel structure. For example, in *Xenopus* oocytes the permeability of Cx46 HCs to Lucifer yellow and ethidium is slightly lowered (~15%) after treatment with nitric oxide (NO) without changes in

5(6)-carboxyfluorescein uptake, suggesting that NO reduces the size cutoff of Cx46 HCs, possibly through S-nitrosylation of connexin subunits [49]. Moreover, the permeability of Cx43 HCs reconstituted in liposomes to charged organic dyes is reduced by PKC phosphorylation of Cx43, which presumably reduces the pore diameter [22]. Interestingly, when all six connexin subunits are phosphorylated, the sucrose permeability becomes close to zero, but hemichannels remain permeable to ethyleneglycol (MW 62) [22]. Thus, even when impermeable to larger molecules, phosphorylated Cx43 HCs remain permeable to small molecules and most likely to ions. Therefore, in addition to changes in number of surface hemichannels and/or open probability of hemichannels, changes in cell membrane permeability via hemichannels could result from changes in intrinsic hemichannel permeability features. To date, reduction in hemichannel permeability as consequence of putative covalent modifications have been described but increases in hemichannel permeability may also occur and further research will be required to validate or disapprove this possibility.

Regulation of the activity of hemichannels at the cell surface

Number of hemichannels in the cell membrane

Cell membrane permeability via hemichannels is affected by relatively rapid changes in the amount of surface hemichannels, and biotinylation of surface membrane proteins with subsequent “pull down” has proved a valuable tool to study the magnitude and velocity of these changes [21,33,39]. Cortical astrocytes or Cx32 HeLa transfectants exposed to metabolic inhibition for only 15 min show an abrupt increase of ~2-fold in their surface Cx43 and Cx32 levels, respectively [21,39]. Moreover, Cx43 transfected HeLa cells treated with a Ca^{2+} ionophore (4Br-23187) or FGF-1 require ~30 min and ~7 h, respectively, to reach a ~2-fold increase in surface hemichannel levels [6].

Rises in $[Ca^{2+}]_i$ appear to be a key regulator of surface hemichannel amount. Increased hemichannel levels in Cx43 HeLa transfectants after FGF-1 requires sustained rises in $[Ca^{2+}]_i$ and p38 MAPK activation [6]. Accordingly, blockade of intracellular Ca^{2+} rises with BAPTA (cells preloaded with BAPTA-AM) or application of SB202190, a p38 MAPK inhibitor, similarly prevent the increase in surface Cx43 HC levels, suggesting that $[Ca^{2+}]_i$ rises and p38 MAPK activation are in the same activation pathway. In Cx32 HeLa transfectants, the level of surface hemichannels increase after metabolic inhibition, which requires Cx HC-mediated Ca^{2+} influx [21]. Interestingly, the increase in surface hemichannel levels of cells treated with metabolic inhibitor, FGF-1 or a Ca^{2+} ionophore is not paralleled by proportional rises in total connexin levels, suggesting that the mechanism involves connexin redistribution and not synthesis *de novo* [6,21]. In addition, cytosolic stress induced by different stimuli reduces Cx43 HC internalization/degradation and increases the number of surface hemichannels in rat kidney fibroblasts [116]. Different forms of metabolic stress are associated with increased cytosolic Ca^{2+} levels. It is likely that amount of hemichannels at the cell membrane can be modulated by increases in $[Ca^{2+}]_i$ that affect insertion and/or removal of hemichannels from the cell surface. Thus, increase in $[Ca^{2+}]_i$ and consequently the number of surface hemichannels could possibly be mediated by numerous membrane receptors and ion channels; receptors linked to G proteins that lead to IP_3 generation and release of Ca^{2+} from intracellular stores or activation of several TRP channels, P2X receptors and pannexin hemichannels (Fig. 2).

In addition to the above mechanisms, which take place over a time course of several, the cell membrane permeability could change much more quickly via gating of Cx HCs.

Gating mechanisms

Transmembrane voltage

In cultured cells under resting conditions, hemichannels have a low open probability at negative membrane potentials and open probability is increased at positive potentials (for review see [117]). Two gating mechanisms have been described for rat Cx46 HCs, termed fast and slow gating (slow gating is also known as loop gating) [92]. Normally, positive membrane potentials activate fast gating which corresponds to fast transitions between the fully open state and a substate. At negative membrane potentials, the loop gating activates slow transitions, perhaps involving multiple substates, between the fully open state, substates and fully closed state [117]. Recently, Verselis and Srinivas [93] showed that slow but not fast gating of Cx46 HCs is modulated by extracellular divalent cations. Negative cell membrane voltages cause closure by loop gating, and divalent cations help to stabilize the closed conformation. This phenomenon may be due to an interaction between divalent cations and the extracellular loops, which induces a narrowing of the hemichannel pore [94]. In general, the biological importance of Cx HC activation by voltage remains to be demonstrated.

Extracellular cations

Increasing extracellular Ca^{2+} concentration ($[Ca^{2+}]_o$) reduces the open probability of all Cx HCs examined (for a review, see [118]). In Cx32 HCs, Ca^{2+} ions bind to a 12 Asp ring at the external vestibule of the hemichannel pore [119]. Alternatively, Cx37 HCs may be directly blocked by extracellular Ca^{2+} going through the pore and reaching binding sites located at the cytoplasmic side [120]. Using atomic force microscopy, a 1.8-nm external pore diameter was visualized in Cx43 HCs reconstituted in lipid bilayers under control conditions. Moreover, removal of extracellular Ca^{2+} increased the outer hemichannel pore diameter to 2.5 nm [121,122]. In addition, Ca^{2+} -induced inhibition of Cx50 HCs is modulated by extracellular concentration of monovalent cations [123] and Ca^{2+} probably acts through the stabilization of the closed configuration of the loop gate [93]. Mg^{2+} , the other main extracellular divalent cation, also affect the functional state of hemichannels [124]. Increasing extracellular Mg^{2+} shifts the voltage dependence of activation, i.e., opening, to more positive potentials, decreases the maximum conductance, accelerates deactivation and slows activation [124]. These effects of Mg^{2+} on Cx46 HCs may result from sequential activation with two independent divalent cation binding sites. In addition, a mutation in the first extracellular domain of hCx46 (hCx46*N63S) results in hemichannels with increased sensitivity to Mg^{2+} -induced blockade [124], which may provide further insight on the mechanism of action of Mg^{2+} . The reduction in hemichannel activation caused by extracellular divalent cations can be overcome by other agents. For example, nitric oxide donors and inflammatory cytokines increase the hemichannel activity in astrocytes, in the presence of divalent cations [5,39]. Under these conditions, the number of hemichannels in the cell surface is not elevated, and the increase in hemichannel activity is rapidly blocked by DTT, a cysteine reducing agent, suggesting that nitrosylation of Cx43 reduces the sensitivity of Cx43 HCs to extracellular divalent cations.

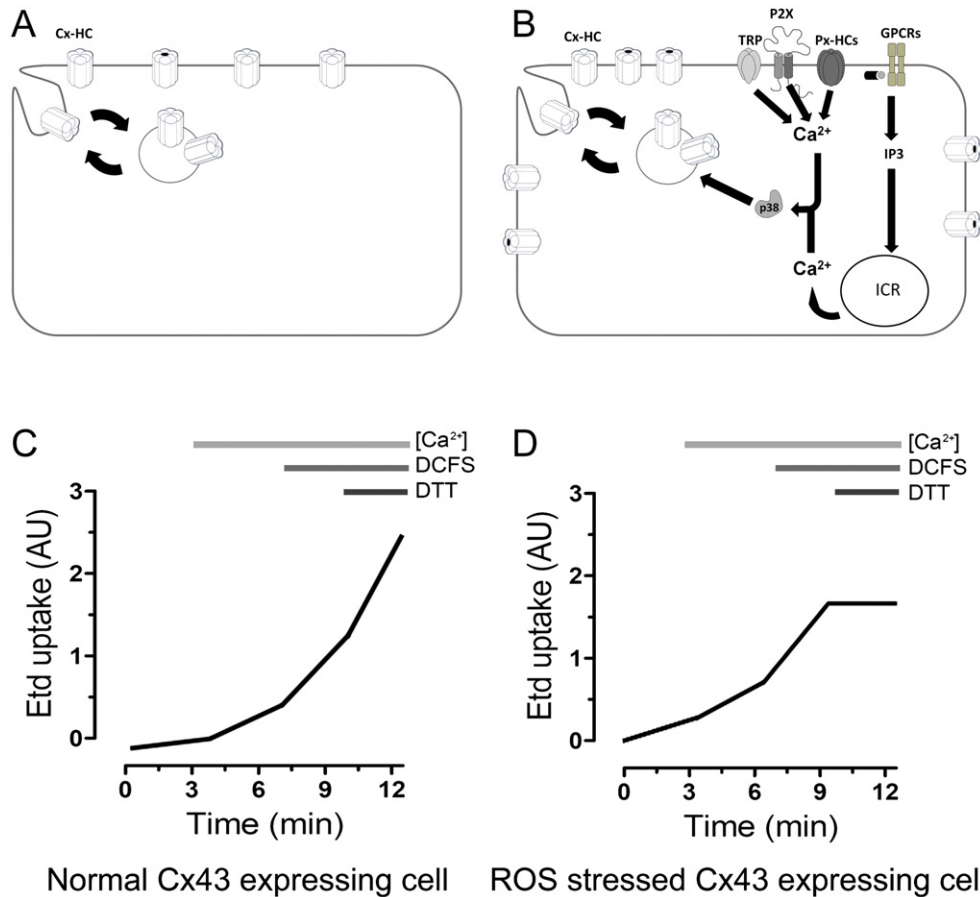


Fig. 2 – Scheme of several conditions that affect membrane permeability via Cx43 HCs. (A) Few Cx43 HCs at the cell membrane of a normal single cell under resting condition. Open hemichannels (black dot at the center) are few. Arrows indicate insertion and removal of Cx43 HCs contained in transport vesicles. (B) Single cell under metabolic stress, where diverse membrane channels [Ca^{2+} permeable TRP channels (TRPs), purinergic P2X receptors (P2Xs), pannexin hemichannels (Px HCs) known to be activated by reactive oxygen species lead to Ca^{2+} influx]. Also, G-protein-coupled receptors (GPCRs) can increase [Ca^{2+}]_i by generation of IP₃, which acts on intracellular Ca^{2+} stores (ICS) to release Ca^{2+} into the cytoplasm. Under both conditions, the increase in [Ca^{2+}]_i leads to activation of p38 MAP kinase (p38) that either reduces (–) removal from or enhances (+) insertion of Cx43 HCs into the cell membrane. Under both conditions, the net number of surface Cx43 HCs is augmented. (C and D) Graphs illustrate time lapse measurements of Etd uptake in Cx43 expressing cells under control conditions (left) or conditions that generate an oxidant stress due to excessive generation of reactive oxygen species (ROS) (right). In both cases, Etd uptake is first shown (~3 min) under control conditions and then treatment with agents that increase [Ca^{2+}]_i (i.e., FGF-1 in normal cell and metabolic inhibitors in ROS stressed cell), followed by change of the extracellular medium to divalent cation-free solution (DCFS) followed at about 10 min of recording with 10 mM DTT. DTT induced opposite effects in normoxic and ROS stressed cells.

Hemichannels formed by Cx50, a main lens connexin, are sensitive to extracellular monovalent cations [125]. Substitution of extracellular Na^+ with K^+ , Cs^+ , Rb^+ or NH_4^+ , in the presence of physiological extracellular Ca^{2+} concentration, results in >10-fold potentiation of macroscopic Cx50 HC currents, an effect reversed upon returning to Na^+ . Replacement of Na^+ with Li^+ , choline or TEA does not potentiate the current. The magnitude of potentiation of Cx50 HC currents is progressively decreased as external Ca^{2+} is reduced. The potentiation is reduced in low Ca^{2+} solution, and the main effect of K^+ appears to be a reduction in the ability of divalent cations to close Cx50 HCs. Cx46 HCs exhibit only a modest increase in open probability upon replacing Na^+ with K^+ . Studies using reciprocal chimeric hemichannels demonstrate that the difference in regulation by monovalent ions in Cx46 and Cx50 HCs resides in the region from the NH_2 -terminal through the

cytoplasmic loop [125]. A possible biological relevance of hemichannel regulation by extracellular monovalent ions is in ephaptic communication as in the retina [126].

Other divalent cations that affect hemichannel activity are Zn^{2+} , Co^{2+} , Cd^{2+} , Mg^{2+} and Ba^{2+} [127]. In bass retinal hemichannels, currents generated by depolarization in Ca^{2+} -free saline and in saline containing 1 mM Ca^{2+} medium are markedly inhibited by extracellular Zn^{2+} . The Zn^{2+} effect on hemichannel currents is concentration dependent (half-maximum inhibitory concentration = 37 μM) and is more potent as compared to other divalent cations ($\text{Zn}^{2+} > \text{Cd}^{2+}$, $\text{Co}^{2+} > \text{Ca}^{2+} > \text{Ba}^{2+} > \text{Mg}^{2+}$). Interestingly, Zn^{2+} and Ca^{2+} modulate hemichannels independently in additive experiments. Cell treatment with *N*-bromosuccinimide to modify histidine residues suppresses the inhibitory action of Zn^{2+} , and Zn^{2+} may act through histidine residues on the extracellular domain

of hemichannels independent of $[Ca^{2+}]_o$ [127]. The effect of Zn^{2+} on Cx38- or Cx35-HCs is biphasic; low $[Zn^{2+}]$ (10 μ M) enhances the currents generated by depolarizing voltages 40 mV, and higher Zn^{2+} (1 mM) reduces the currents [128]. This differential effect of Zn^{2+} on hemichannel activity may be explained by differences in connexins as well as by differences in the cell system used to evaluate its effects.

pH

Hemichannels composed of different connexin types are sensitive to intracellular acidification [53,129,130]. A direct effect of H^+ on the connexin subunit was suggested in Cx46 HCs, since the response is very fast and is observed in the excised patch configuration [53]. Chemical gating of Cx46 HCs seems to affect a pH sensor located close to the channel pore [53] and pH reduction shifts the voltage activation to more positive voltages [131]. However, intracellular acidification might rapidly affect $[Ca^{2+}]_i$ in intact cells, which could affect activation of Cx32 and Cx43 HCs.

Redox potential

S-nitrosylation of Cx43 increases dye uptake in cortical astrocytes [39] and is probably associated with changes in the voltage sensitivity of Cx46 HCs [49]. The open probability of Cx43 HCs increases several fold after treatment of normal cells with dithiothreitol (DTT) [41], suggesting that both reduction and oxidation can modulate hemichannel gating, possibly depending on the cell type and its metabolic state. In agreement with this possibility, the increase in Cx43 HC activity observed in cells under oxidative stress, such as that induced by NO donors [39], metabolic inhibition [39], inflammatory cytokines [5] and hypoxia-reoxygenation [98], is completely inhibited by DTT.

The increase in Cx43 HC activity induced in normal cells by DTT (Fig. 2) or nitric oxide donors occurs in seconds and without changes in levels of surface Cx43 [39,41], suggesting that both result from increase in open probability of existing surface hemichannels. The effect of DTT is not abolished by buffering the $[Ca^{2+}]_i$ and the effect of both NO donors and DTT occurs in the presence of extracellular divalent cations. The increase in Cx43 HC activity induced by DTT is not reproduced by extracellular application of reduced glutathione, but application of glutathione in the recording pipette increases the open probability [41]. Moreover, the recovery in membrane permeability induced by DTT in cells during metabolic inhibition is reproduced by application of glutathione ethyl ester, which is membrane permeable, but not by extracellular reduced glutathione [39]. These data indicate that intracellular but not extracellular Cys residues mediate the changes in Cx43 HC activity induced by alterations in redox potential. Changes in redox potential may not affect all Cx HCs, because not all connexins have cysteine residues in intracellular domains, e.g., Cx45.

Mechanosensitivity

Mechanical stimulation induces ATP release through Cx43 HCs in osteoblastic cells and chondrocytes [132,133]. Moreover, mechanical stimulation enhances the activity of Cx46 HCs at negative transmembrane potentials but inhibits the increase in hemichannel activity induced by positive membrane potentials [134]. The mechanosensitivity of lens hemichannels may assist accommodation by providing a transient path for volume flow as the lens changes shape [134]. Cx30 HCs are also mechanosensitive and may

be involved in regulation of pressure-induced natriuresis by releasing ATP into the tubular fluid [135]. Additionally, Cx43 HCs in osteocytes activate after shear stress, accelerating PGE_2 release [136]. The mechanism of action of mechanical stress on Cx HC activity remains unknown but clearly occurs in the presence of extracellular divalent cations and most likely without changes in intracellular redox potential.

Interaction between different hemichannel-mediated membrane permeabilization mechanisms

Cx46 HCs expressed in *Xenopus* oocytes show a progressive current facilitation after repetitive positive pulses to potentials greater than +20 mV, which depends on the time between pulses (<20 s) [137,138]. The mechanism of hemichannel current facilitation by repetitive depolarizing voltages remains unknown. However, it is unlikely to occur as a consequence of increased $[Ca^{2+}]_i$ because increases in $[Ca^{2+}]_i$ require >15 min to significantly increase the hemichannel-mediated cell permeabilization to dyes [3,6,21,39]. Facilitation of hemichannel currents is also observed in oocytes expressing Cx46 with a truncated C-terminus; nominal removal of divalent cations from the extracellular side causes maximal current activation of these truncated hemichannels and also prevents facilitation, indicating that Cx46 HCs show activation independent of their C-terminal domain [138]. Similarly, application of consecutive electrical pulses to Cx46 expressing oocytes in the absence of extracellular divalent cations does not affect the maximal current, which is ~3-fold higher than the maximal current generated by repetitive voltage steps at physiological $[Ca^{2+}]_o$ (Fig. 3). Moreover, repetitive application of depolarizing voltages from a holding potential of -60 mV to +80 mV in the presence of extracellular divalent cations induces current facilitation to a maximal value, which is lower than that recorded in the absence of divalent cations (Fig. 3). However, the current facilitation depends on the amplitude of the depolarization, reaching a maximum at approximately +60 mV [138], suggesting that repetitive voltages open a smaller fraction of Cx46 HCs than does divalent cation-free solution. These results also suggest that Cx46 HC currents would increase when repetitive voltage steps are applied to cells bathed with saline solution containing physiological concentrations of Ca^{2+} and Mg^{2+} .

Potentiation occurs after sequential stimulation of cells with different treatments known to enhance hemichannel activity. Incubation of Cx43 HeLa transfectants with 20 ng/ml FGF-1 induces a ~2-fold increase in the dye uptake rate [6]. The increase in hemichannel activity induced by FGF-1 requires elevation of intracellular $[Ca^{2+}]_i$ and is associated with increase in Cx levels at the cell surface [6]. Application of DTT to Cx43 HeLa transfectants treated with FGF-1 further increases Cx HC activity (Figs. 4A and B) and to a level higher than that induced by DTT alone (Fig. 4B), effects that are not observed in parental cells (Fig. 4B). The effect induced by FGF-1 + DTT is greater than the sum of the effects of the two stimuli by themselves and is completely blocked by La^{3+} , indicating the participation of Cx HCs and absence of pannexin hemichannels. As expected, a similar response occurs after adding DTT to cells treated with the Ca^{2+} ionophore 4-Br-A23187 (Figs. 4C and D). Again, the response is higher than with either compound alone and was also prominently inhibited by La^{3+} . Permeabilization by multiple stimuli may well be greater than that induced by any of them singly, particularly if there is synergistic action. Similarly, exposure of cortical astrocytes to a divalent cation-free

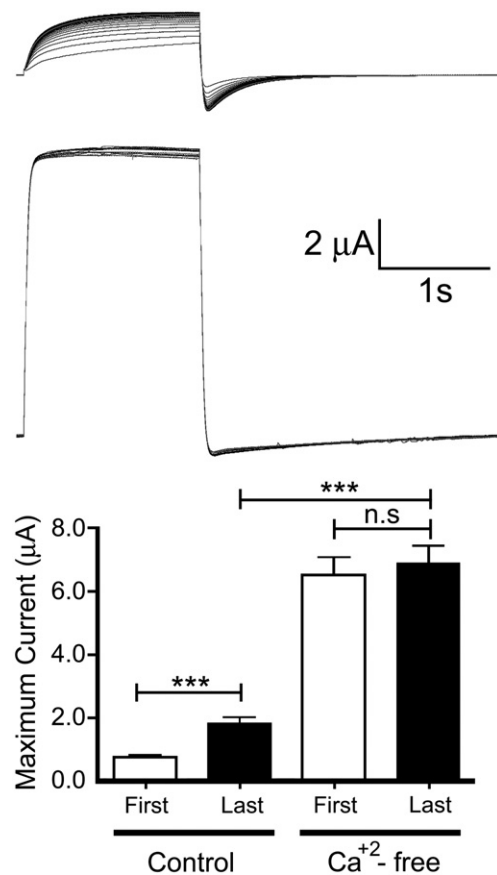


Fig. 3 – Facilitation of Cx46 HC current induced by successive depolarizations might not result in activation of all hemichannels available at the plasma membrane. *Xenopus laevis* oocytes were injected with cRNA of rat Cx46 and 24 h later electrophysiology recordings were performed. The pulse protocol was 2.5 s at -60 mV following by 1.5 s at $+80$ mV for 25 repetitions. Two current records are shown; the top record is superimposed traces showing progressive increase in outward and tail currents induced by the pulses under control conditions (in ND96 bath solution, plus 1.8 Ca^{+2} and 1.0 Mg^{+2}). The bottom record was obtained with the same protocol as in (A), but the cell was bathed in ND96 recording solution without Ca^{+2} and Mg^{+2} (Ca^{+2} free). Under this condition the initial current was ~ 4.0 times bigger than the maximal current in control medium ($n = 3$, $***p < 0.05$) and there was little variation in Cx46 HC currents during successive pulses. The bar graph shows mean \pm SEM of several experiments, pointing out the difference between the first and the last pulse under control conditions ($***p < 0.05$). However, no difference was found between the first and last current in oocytes in Ca^{+2} -free solution (n.s., $p > 0.05$).

solution plus DTT induces a higher rate of ethidium uptake than the sum of each stimulus acting alone, and the effect is abrogated by β -glycyrrhetic acid (β -GA), a known hemichannel blocker (Fig. 5). In summary, permeabilization responses *in vivo* may involve multiple hemichannel activating mechanisms, and may induce an unexpectedly great increase in cell permeability (Fig. 2). Increased insertion and increase in P_o of inserted channels would be multiplicative and could account for these observations.

Cell permeabilization through hemichannels induced by metabolic inhibition (iodoacetic acid and antimycin A) in HeLa-Cx43 and HeLa-Cx32 transfectants exposed to metabolic inhibitors show a higher (La^{3+} sensitive) ethidium uptake rate increase after exposure to a divalent cation-free solution than without metabolic inhibition [21]. However, under this condition, the total response is not higher than the sum of each stimulus by itself, indicating that potentiation was less than by summation. Moreover, application of metabolic inhibitors after divalent cation-free solution does not further affected the cell permeabilization, suggesting a requirement for Ca^{2+} influx in the metabolic inhibition induced hemichannel response or saturation of available Cx32 HCs. Therefore, not all activating stimuli induce potentiation of the Cx HC activity, possibly depending on the cell state, stimulus nature and molecular mechanism.

Concluding remarks

Historically, hemichannels were regarded simply as precursors of gap junction channels, since there was limited evidence that hemichannels were not open to the external milieu prior to gap junction channel formation. Moreover, their permeability properties predicted from that of gap junction channel was so high that their opening was thought to be deleterious. A possible error has been the extrapolation of gap junction channel permeability to that of the constituent hemichannels. The human genome encodes 21 connexins and most cell types express two or more, which frequently offers the formation of heteromeric hemichannels instead. Moreover, most permeability studies have been carried out in homomeric gap junction channels formed by no more than 40% of connexins and the most relevant information obtained in those studies is that gap junction channels could show charge selectivity. Nonetheless, the size cutoff of the few homo and heteromeric gap junction channels so far characterized remain unknown and numerous reports still quote that gap junctions are permeable to molecules of up to 1–1.2 kDa based on studies obtained in insect cells which do not express connexins. Now it is clear that hemichannels should be viewed as intercellular communication pathways independent of gap junctions and in parallel with their role as precursors of gap junction channels. Clarification of their role as a pathway between cytoplasm and extracellular space will require further work. For example, it is unknown if cellular sorting mechanisms for delivery of hemichannels that will be functional as hemichannels differ from those hemichannels that are destined to form gap junction channels. In agreement with this possibility, in epithelial and endothelial cells, gap junctions are often located in the baso-lateral membrane below the apical tight junctions. Hence, in these tissues it is clear that if there are hemichannels in the apical membrane, they do not form gap junction channels.

There is little doubt that surface hemichannels not forming gap junctions will be found interacting with other membrane or cytoplasmic proteins in the same way as gap junction channels, but at this point identity of binding partners is uncertain. As discussed here, intercellular communication through Cx HCs is a highly regulated process, and cell signaling that upregulates Cx HC activity frequently down regulates gap junction based intercellular communication. Although these differences could be explained by spatial separation of regulatory pathways not yet described, they

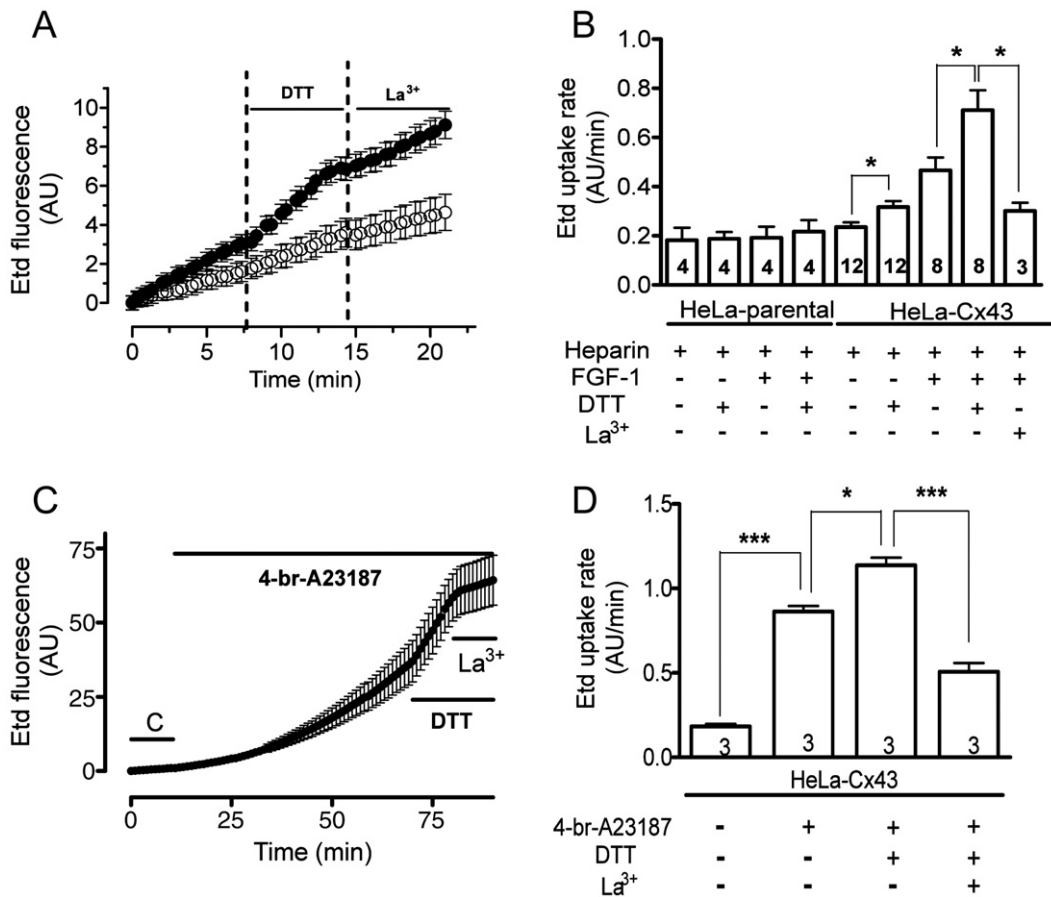


Fig. 4 – DTT induces additional increase in the Cx HC-mediated dye uptake of HeLa-Cx43 cells treated with FGF-1 or a Ca²⁺ ionophore. (A) Time lapse recording showing the Etd uptake of HeLa-Cx43 cells or parental HeLa cells incubated for 7 h with 20 ng/ml FGF-1 plus heparin (filled circles) or heparin alone (white circles), after addition of 10 mM DTT and after treatment with 200 μM La³⁺. Representative experiment showing mean ± SEM of 20 cells. (B) Average Etd uptake rates of parental HeLa cells or HeLa-Cx43 cells in the control condition (heparin alone), after addition of 10 mM DTT or incubated for 7 h with 20 ng/ml FGF-1 plus heparin and treated with DTT; and after addition of La³⁺. (C) Time lapse recording showing Etd uptake by HeLa-Cx43 cells in the control condition (c) and after addition of 2.5 μM 4-Br-A23187, 10 mM DTT and La³⁺ (150 μM). Representative experiment showing mean ± SEM of 20 cells. (D) Average Etd uptake rates of HeLa-Cx43 cells in the control condition, after addition of 4 Br-A23187, then DTT and then La³⁺. The number of independent experiments is indicated within each bar. 20 cells were recorded in each experiment. **p* < 0.05; ****p* < 0.001.

could also be explained by differences in post-translational hemichannel modifications before or during delivery to different membrane domains.

We know that the activity of Cx HCs is modulated through changes in their surface levels, permeability and electrophysiological properties. Stimuli activating Cx HCs through different mechanisms induce responses that can be connexin specific. In addition, healthy cells are likely capable of adapting to transient activation of Cx HCs through active compensatory mechanisms, whereas damaged cells (metabolically compromised, under proinflammatory environments and/or carrying hemichannel defective Cx mutations) may not be able to overcome the metabolic consequences of hemichannel activation. The increase in number of hemichannels in the cell surface membrane induced by rises in [Ca²⁺]_i is a response common to healthy and sick cells. In both cases, hemichannels are a pathway that contributes to increase in [Ca²⁺]_i, but sick cells may be unable to restore their [Ca²⁺]_i. Under the same thought, hemichannels might be part of

responses generated by different drugs or biologically relevant compounds acting on molecular targets that lead to increases in [Ca²⁺]_i and/or changes in intra and/or extracellular compartments including redox state, pCa, pH, membrane potential and activity of protein phosphatases and kinases. Thus, a component of cell responses elicited by compounds that increase [Ca²⁺]_i may be due to changes in hemichannel activity rather than changes mediated exclusively by the primary target. An example of this possibility is the rise in membrane permeability induced by FGF-1, which also leads to a delayed increase in intracellular [Ca²⁺]_i in HeLa transfectants [6]. This Cx HC transactivation is connexin dependent since it occurs in Cx43 and Cx45 but not in Cx26 HeLa transfectants. Thus, conditions that potentiate hemichannel activity may differ in cells expressing different connexins.

To date, the demonstration that functional hemichannels exist at the cell surface and appear relevant to numerous cellular responses has utilized mainly *in vitro* studies. Now, it is important to acquire evidence from *in vivo* experimental models. Although *in*

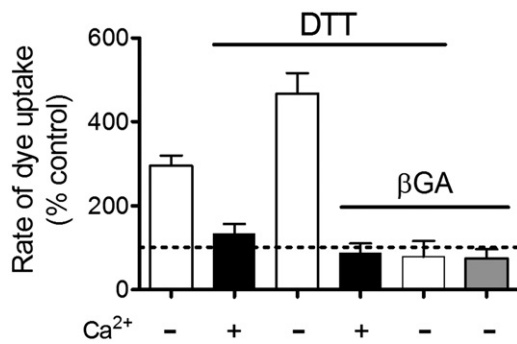


Fig. 5 – Treatment with divalent cation-free solution and DTT causes a synergistic potentiation of the dye uptake response mediated by Cx43 HCs in HeLa transfectants. Time lapse measurements of Etd uptake rate were performed in physiological concentrations of divalent cations (Ca²⁺ +, filled bars) or in Ca²⁺ and Mg²⁺-free solution (Ca²⁺ -, white bars), without (gray bar) or with addition of 100 mM DTT followed by application of 18β-glycyrrhetic acid (βGA, 150 μM). Uptake rates were normalized to the control value of each experiment. The dotted line indicates control dye uptake rate expressed as 100%. Each plotted number corresponds to the mean ± SEM of 4 independent experiments including 20 cells in each of them.

in vitro preparations are convenient for analysis of the effects of single variables, *in vivo* data where cells are under the influence of multiple regulatory mechanisms will be required to establish functional relevance of hemichannels in both physiological and pathophysiological conditions.

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REFERENCES

- [1] W.H. Evans, E. De Vuyst, L. Leybaert, The gap junction cellular internet: connexin hemichannels enter the signalling limelight, *Biochem. J.* 397 (2006) 1–14.
- [2] M.V. Bennett, J.E. Contreras, F.F. Bukauskas, J.C. Sáez, New roles for astrocytes: gap junction hemichannels have something to communicate, *Trends Neurosci.* 11 (2003) 610–617.
- [3] J.E. Contreras, H.A. Sánchez, E.A. Eugénin, D. Speidel, M. Theis, K. Willecke, F.F. Bukauskas, M.V. Bennett, J.C. Sáez, Metabolic inhibition induces opening of unapposed connexin 43 gap junction hemichannels and reduces gap junctional communication in cortical astrocytes in culture, *Proc. Nat. Acad. Sci. U.S.A.* 99 (2002) 495–500.
- [4] E. De Vuyst, E. Decrock, M. De Bock, H. Yamasaki, C.C. Naus, W.H. Evans, L. Leybaert, Connexin hemichannels and gap junction channels are differentially influenced by lipopolysaccharide and basic fibroblast growth factor, *Mol. Biol. Cell* 18 (2007) 34–46.
- [5] M.A. Retamal, N. Froger, N. Palacios-Prado, P. Ezan, P.J. Sáez, J.C. Sáez, C. Giaume, Cx43 hemichannels and gap junction channels in astrocytes are regulated oppositely by proinflammatory cytokines released from activated microglia, *J. Neurosci.* 27 (2007) 13781–13792.
- [6] K.A. Schalper, N. Palacios-Prado, M.A. Retamal, K.F. Shoji, A.D. Martínez, J.C. Sáez, Connexin hemichannels composition determines the FGF-1-induced membrane permeability and free [Ca²⁺]_i responses, *Mol. Biol. Cell* 19 (2008) 3501–3513.
- [7] J.A. Orellana, P.J. Sáez, K.F. Shoji, K.A. Schalper, N. Palacios-Prado, V. Velarde, C. Giaume, M.V. Bennett, J.C. Sáez, Modulation of brain hemichannels and gap junction channels by pro-inflammatory agents and their possible role in neurodegeneration, *Antioxid. Redox Signal.* 11 (2009) 369–399.
- [8] E. De Vuyst, N. Wang, E. Decrock, M. De Bock, M. Vinken, M. Van Moorhem, C. Lai, M. Culot, V. Rogiers, R. Cecchelli, C.C. Naus, W.H. Evans, L. Leybaert, Ca²⁺ regulation of connexin 43 hemichannels in C6 glioma and glial cells, *Cell Calcium* 46 (2009) 176–187.
- [9] J.A. Diez, S. Ahmad, W.H. Evans, Assembly of heteromeric connexons in guinea-pig liver en route to the Golgi apparatus, plasma membrane and gap junctions, *Eur. J. Biochem.* 262 (1999) 142–148.
- [10] N.M. Kumar, D.S. Friend, N.B. Gilula, Synthesis and assembly of human beta 1 gap junctions in BHK cells by DNA transfection with the human beta 1 cDNA, *J. Cell Sci.* 108 (1995) 3725–3734.
- [11] C.H. George, J.M. Kendall, W.H. Evans, Intracellular trafficking pathways in the assembly of connexins into gap junctions, *J. Biol. Chem.* 274 (1999) 8678–8685.
- [12] L.S. Musil, D.A., Good enough biochemical analysis of connexin43 intracellular transport, phosphorylation, and assembly into gap junctional plaques, *J. Cell Biol.* 5 (1991) 1357–1374.
- [13] M. Koval, J.E. Harley, E. Hick, T.H. Steinberg, Connexin46 is retained as monomers in a trans-Golgi compartment of osteoblastic cells, *J. Cell Biol.* 137 (1997) 847–857.
- [14] K. Jordan, J.L. Solan, M. Dominguez, M. Sia, A. Hand, P.D. Lampe, D.W. Laird, Trafficking, assembly, and function of a connexin43-green fluorescent protein chimera in live mammalian cells, *Mol. Biol. Cell* 10 (1999) 2033–2050.
- [15] P.E. Martin, R.J. Errington, W.H. Evans, Gap junction assembly: multiple connexin fluorophores identify complex trafficking pathways, *Cell Commun. Adhes.* 8 (2001) 243–248.
- [16] B.N. Giepmans, I. Verlaan, T. Hengeveld, H. Janssen, J. Calafat, M.M. Falk, W.H. Moolenaar, Gap junction protein connexin-43 interacts directly with microtubules, *Curr. Biol.* 11 (2001) 1364–1368.
- [17] U. Lauf, B.N. Giepmans, P. Lopez, S. Braconnot, S.C. Chen, M.M. Falk, Dynamic trafficking and delivery of connexins to the plasma membrane and accretion to gap junctions in living cells, *Proc. Nat. Acad. Sci. U.S.A.* 99 (2002) 10446–10451.
- [18] G. Gaietta, T.J. Deerinck, S.R. Adams, J. Bouwer, O. Tour, D.W. Laird, G.E. Sosinsky, R.Y. Tsien, M.H. Ellisman, Multicolor and electron microscopic imaging of connexin trafficking, *Science* 296 (2002) 503–507.
- [19] R.M. Shaw, A.J. Fay, M.A. Puthenveedu, M. von Zastrow, Y.N. Jan, L.Y. Jan, Microtubule plus-end-tracking proteins target gap junctions directly from the cell interior to adherens junctions, *Cell* 128 (2007) 547–560.
- [20] J. Simek, J. Churko, Q. Shao, D.W. Laird, Cx43 has distinct mobility within plasma-membrane domains, indicative of progressive formation of gap-junction plaques, *J. Cell Sci.* 122 (2009) 554–562.
- [21] H.A. Sánchez, J.A. Orellana, V.K. Verselis, J.C. Sáez, Metabolic inhibition increases activity of connexin-32 hemichannels permeable to Ca²⁺ in transfected HeLa cells, *Am. J. Physiol. Cell Physiol.* 297 (2009) 665–678.
- [22] X. Bao, S.C. Lee, L. Reuss, G.A. Altenberg, Change in permeant size selectivity by phosphorylation of connexin 43 gap-junctional hemichannels by PKC, *Proc. Nat. Acad. Sci. U.S.A.* 104 (2007) 4919–4924.
- [23] G.M. Essenfelder, R. Bruzzone, J. Lamartine, A. Charollais, C. Blanchet-Bardon, M.T. Barbe, P. Meda, G. Waksman, Connexin30 mutations responsible for hidrotic ectodermal dysplasia cause

- abnormal hemichannel activity, *Hum. Mol. Genet.* 13 (2004) 1703–1714.
- [24] B.C. Stong, Q. Chang, S. Ahmad, X. Lin, A novel mechanism for connexin 26 mutation linked deafness: cell death caused by leaky gap junction hemichannels, *Laryngoscope* 116 (2006) 2205–2210.
- [25] D.A. Gerido, A.M. DeRosa, G. Richard, T.W. White, Aberrant hemichannel properties of Cx26 mutations causing skin disease and deafness, *Am. J. Physiol. Cell Physiol.* 293 (2007) 337–345.
- [26] R. Dobrowolski, A. Sommershof, K. Willecke, Some oculodentodigital dysplasia-associated Cx43 mutations cause increased hemichannel activity in addition to deficient gap junction channels, *J. Membr. Biol.* 219 (2007) 9–17.
- [27] T.D. Matos, H. Caria, H. Simoes-Teixeira, T. Aasen, O. Dias, M. Andrea, D.P. Kelsell, G. Fialho, A novel M163L mutation in connexin 26 causing cell death and associated with autosomal dominant hearing loss, *Hear. Res.* 240 (2008) 87–92.
- [28] J.R. Lee, A.M. Derosa, T.W. White, Connexin mutations causing skin disease and deafness increase hemichannel activity and cell death when expressed in *Xenopus* oocytes, *J. Invest. Dermatol.* 129 (2009) 870–878.
- [29] P.J. Minogue, J.J. Tong, A. Arora, I. Russell-Eggitt, D.M. Hunt, A.T. Moore, L. Ebihara, E.C. Beyer, V.M. Berthoud, A mutant connexin50 with enhanced hemichannel function leads to cell death, *Invest. Ophthalmol. Vis. Sci.* 50 (2009) 5837–5845.
- [30] C.K. Abrams, M.V. Bennett, V.K. Verselis, T.A. Bargiello, Voltage opens unopposed gap junction hemichannels formed by a connexin 32 mutant associated with X-linked Charcot-Marie-Tooth disease, *Proc. Nat. Acad. Sci. U.S.A.* 99 (2002) 3980–3984.
- [31] G.S. Liang, M. de Miguel, J.M. Gómez-Hernández, J.D. Glass, S.S. Scherer, M. Mintz, L.C. Barrio, K.H. Fischbeck, Severe neuropathy with leaky connexin32 hemichannels, *Ann. Neurol.* 57 (2005) 749–754.
- [32] D.J. Belliveau, M. Bani-Yaghoub, B. McGirr, C.C. Naus, W.J. Rushlow, Enhanced neurite outgrowth in PC12 cells mediated by connexin hemichannels and ATP, *J. Biol. Chem.* 281 (2006) 20920–20931.
- [33] K.A. Schalper, J.A. Orellana, V.M. Berthoud, J.C. Sáez, Dysfunctions of the diffusional membrane pathways mediated by hemichannels in inherited and acquired human diseases, *Curr. Vasc. Pharmacol.* 7 (2009) 486–505.
- [34] R.A. Pearson, N. Dale, E. Llaudet, P. Mobbs, ATP released via gap junction hemichannels from the pigment epithelium regulates neural retinal progenitor proliferation, *Neuron* 46 (2005) 731–744.
- [35] J.A. Guttman, A.E. Lin, Y. Li, J. Bechberger, C.C. Naus, A.W. Vogl, B.B. Finlay, Gap junction hemichannels contribute to the generation of diarrhea during infectious enteric disease, *Gut* (2009).
- [36] K. Willecke, J. Eiberger, J. Degen, D. Eckardt, A. Romualdi, M. Guldenagel, U. Deutsch, G. Sohl, Structural and functional diversity of connexin genes in the mouse and human genome, *Biol. Chem.* 383 (2002) 725–737.
- [37] M. Yeager, N.B. Gilula, Membrane topology and quaternary structure of cardiac gap junction ion channels, *J. Mol. Biol.* 223 (1992) 929–948.
- [38] D. Locke, I.V. Koreen, A.L. Harris, Isoelectric points and post-translational modifications of connexin26 and connexin32, *FASEB J.* 20 (2006) 1221–1223.
- [39] M.A. Retamal, C.J. Cortes, L. Reuss, M.V. Bennett, J.C. Sáez, S-nitrosylation and permeation through connexin 43 hemichannels in astrocytes: induction by oxidant stress and reversal by reducing agents, *Proc. Nat. Acad. Sci. U.S.A.* 103 (2006) 4475–4480.
- [40] J.L. Solan, P.D. Lampe, Connexin43 phosphorylation: structural changes and biological effects, *Biochem. J.* 419 (2009) 261–272.
- [41] M.A. Retamal, K.A. Schalper, K.F. Shoji, M.V. Bennett, J.C. Sáez, Opening of connexin 43 hemichannels is increased by lowering intracellular redox potential, *Proc. Nat. Acad. Sci. U.S.A.* 104 (2007) 8322–8327.
- [42] O. Traub, J. Look, R. Dermietzel, F. Brümmer, D. Hülser, K. Willecke, Comparative characterization of the 21-kD and 26-kD gap junction proteins in murine liver and cultured hepatocytes, *J. Cell Biol.* 108 (1989) 1039–1051.
- [43] P.D. Lampe, A.F. Lau, Regulation of gap junctions by phosphorylation of connexins, *Arch. Biochem. Biophys.* 384 (2000) 205–215.
- [44] A.P. Moreno, A.F. Lau, Gap junction channel gating modulated through protein phosphorylation, *Prog. Biophys. Mol. Biol.* 94 (2007) 107–119.
- [45] G. Hawat, G. Baroudi, Differential modulation of unapposed connexin 43 hemichannel electrical conductance by protein kinase C isoforms, *Pflugers Arch.* 456 (2008) 519–527.
- [46] D.Y. Kim, Y. Kam, S.K. Koo, C.O. Joe, Gating connexin 43 channels reconstituted in lipid vesicles by mitogen-activated protein kinase phosphorylation, *J. Biol. Chem.* 274 (1999) 5581–5587.
- [47] X. Bao, L. Reuss, G.A. Altenberg, Regulation of purified and reconstituted connexin 43 hemichannels by protein kinase C-mediated phosphorylation of Serine 368, *J. Biol. Chem.* 279 (2004) 20058–20066.
- [48] A. Ngezahayo, C. Zeilinger, I.I. Todt, I.I. Marten, H. Kolb, Inactivation of expressed and conducting rCx46 hemichannels by phosphorylation, *Pflugers Arch.* 436 (1998) 627–629.
- [49] M.A. Retamal, S. Yin, G.A. Altenberg, L. Reuss, Modulation of Cx46 hemichannels by nitric oxide, *Am. J. Physiol. Cell Physiol.* 296 (2009) 1356–1363.
- [50] R. Eckert, pH gating of lens fibre connexins, *Pflügers Arch.* 443 (2002) 843–851.
- [51] M. Delmar, W. Coombs, P. Sorgen, H.S. Duffy, S.M. Taffet, Structural bases for the chemical regulation of Connexin43 channels, *Cardiovasc. Res.* 62 (2004) 268–275.
- [52] F. Liu, F.T. Arce, S. Ramachandran, R. Lal, Nanomechanics of hemichannel conformations: connexin flexibility, *J. Biol. Chem.* 281 (2006) 23207–23217.
- [53] E.B. Trexler, F.F. Bukauskas, M.V. Bennett, T.A. Bargiello, V.K. Verselis, Rapid and direct effects of pH on connexins revealed by the connexin46 hemichannel preparation, *J. Gen. Physiol.* 113 (1999) 721–742.
- [54] H.S. Duffy, P.L. Sorgen, M.E. Girvin, P. O'Donnell, W. Coombs, S.M. Taffet, M. Delmar, D.C. Spray, pH-dependent intramolecular binding and structure involving Cx43 cytoplasmic domains, *J. Biol. Chem.* 277 (2002) 36706–36714.
- [55] P.L. Sorgen, H.S. Duffy, S.M. Cahill, W. Coombs, D.C. Spray, M. Delmar, M.E. Girvin, Sequence-specific resonance assignment of the carboxyl terminal domain of Connexin43, *J. Biomol.* 23 (2002) 245–246 NMR.
- [56] T. Toyofuku, Y. Akamatsu, H. Zhang, T. Kuzuya, M. Tada, M. Hori, c-Src regulates the interaction between connexin-43 and ZO-1 in cardiac myocytes, *J. Biol. Chem.* 276 (2001) 1780–1788.
- [57] X. Li, S. Lu, J.I. Nagy, Direct association of connexin36 with zonula occludens-2 and zonula occludens-3, *Neurochem. Int.* 54 (2009) 393–402.
- [58] K. Leykauf, M. Salek, J. Bomke, M. Frech, W.D. Lehmann, M. Durst, A. Alonso, Ubiquitin protein ligase Nedd4 binds to connexin43 by a phosphorylation-modulated process, *J. Cell Sci.* 119 (2006) 3634–3642.
- [59] X.S. Yu, J.X. Jiang, Interaction of major intrinsic protein (aquaporin-0) with fiber connexins in lens development, *J. Cell Sci.* 117 (2004) 871–880.
- [60] A. Gellhaus, X. Dong, S. Propson, K. Maass, L. Klein-Hitpass, M. Kibschull, O. Traub, K. Willecke, B. Perbal, S.J. Lye, E. Winterhager, Connexin43 interacts with NOV: a possible mechanism for negative regulation of cell growth in choriocarcinoma cells, *J. Biol. Chem.* 279 (2004) 36931–36942.
- [61] B.N. Giepmans, W.H. Moolenaar, The gap junction protein connexin43 interacts with the second PDZ domain of the zona occludens-1 protein, *Curr. Biol.* 8 (1998) 931–934.

- [62] J.G. Laing, R.N. Manley-Markowski, M. Koval, R. Civitelli, T.H. Steinberg, Connexin45 interacts with zonula occludens-1 and connexin43 in osteoblastic cells, *J. Biol. Chem.* 276 (2001) 23051–23055.
- [63] K. Maass, J. Shibayama, S.E. Chase, K. Willecke, M. Delmar, C-terminal truncation of connexin43 changes number, size, and localization of cardiac gap junction plaques, *Circ. Res.* 101 (2007) 1283–1291.
- [64] A. Revilla, C. Castro, L.C. Barrio, Molecular dissection of transjunctional voltage dependence in the connexin-32 and connexin-43 junctions, *Biophys. J.* 77 (1999) 1374–1383.
- [65] A.M. DeRosa, R. Mui, M. Srinivas, T.W. White, Functional characterization of a naturally occurring Cx50 truncation, *Invest. Ophthalmol. Vis. Sci.* 47 (2006) 4474–4481.
- [66] D.C. Bates, W.C. Sin, Q. Aftab, C.C. Naus, Connexin43 enhances glioma invasion by a mechanism involving the carboxy terminus, *Glia* 55 (2007) 1554–1564.
- [67] X. Dang, B.W. Doble, E. Kardami, The carboxy-tail of connexin-43 localizes to the nucleus and inhibits cell growth, *Mol. Cell. Biochem.* 242 (2003) 35–38.
- [68] K. Maass, S.E. Chase, X. Lin, M. Delmar, Cx43 CT domain influences infarct size and susceptibility to ventricular tachyarrhythmias in acute myocardial infarction, *Cardiovasc. Res.* 84 (2009) 361–367.
- [69] M.G. Kozoriz, J.F. Bechberger, G.R. Bechberger, M.W. Suen, A.P. Moreno, K. Maass, K. Willecke, C.C. Naus, The connexin43 C-terminal region mediates neuroprotection during stroke, *J. Neuropathol. Exp. Neurol.* 69 (2010) 196–206.
- [70] V.K. Verselis, C.S. Ginter, T.A. Bargiello, Opposite voltage gating polarities of two closely related connexins, *Nature* 368 (1994) 348–351.
- [71] S. Oh, V.K. Verselis, T.A. Bargiello, Charges dispersed over the permeation pathway determine the charge selectivity and conductance of a Cx32 chimeric hemichannel, *J. Physiol.* 586 (2008) 2445–2461.
- [72] L. Dong, X. Liu, H. Li, B.M. Vertel, L. Ebihara, Role of the N-terminus in permeability of chicken connexin45.6 gap junctional channels, *J. Physiol.* 576 (2006) 787–799.
- [73] J.W. Kyle, P.J. Minogue, B.C. Thomas, D.A. Domowicz, V.M. Berthoud, D.A. Hanck, E.C. Beyer, An intact connexin N-terminus is required for function but not gap junction formation, *J. Cell Sci.* 121 (2008) 2744–2750.
- [74] J.E. Contreras, J.C. Sáez, F.F. Bukauskas, M.V. Bennett, Gating and regulation of connexin 43 (Cx43) hemichannels, *Proc. Nat. Acad. Sci. U.S.A.* 100 (2003) 11388–11393.
- [75] A. Oshima, K. Tani, Y. Hiroaki, Y. Fujiyoshi, G.E. Sosinsky, Three-dimensional structure of a human connexin26 gap junction channel reveals a plug in the vestibule, *Proc. Nat. Acad. Sci. U.S.A.* 104 (2007) 10034–10039.
- [76] J.W. Kyle, V.M. Berthoud, J. Kurutz, P.J. Minogue, M. Greenspan, D.A. Hanck, E.C. Beyer, The N terminus of connexin37 contains an alpha-helix that is required for channel function, *J. Biol. Chem.* 284 (2009) 20418–20427.
- [77] S. Maeda, S. Nakagawa, M. Suga, E. Yamashita, A. Oshima, Y. Fujiyoshi, T. Tsukihara, Structure of the connexin 26 gap junction channel at 3.5 Å resolution, *Nature* 458 (2009) 597–602.
- [78] B.D. Kalmatsky, S. Bhagan, Q. Tang, T.A. Bargiello, T.L. Dowd, Structural studies of the N-terminus of Connexin 32 using 1H NMR spectroscopy, *Arch. Biochem. Biophys.* 490 (2009) 9–16.
- [79] X. Wang, L. Li, L.L. Peracchia, C. Peracchia, Chimeric evidence for a role of the connexin cytoplasmic loop in gap junction channel gating, *Pflugers Arch.* 431 (1996) 844–852.
- [80] D. Manthey, K. Banach, T. Desplantez, C.G. Lee, C.A. Kozak, O. Traub, R. Weingart, K. Willecke, Intracellular domains of mouse connexin26 and -30 affect diffusional and electrical properties of gap junction channels, *J. Membr. Biol.* 181 (2001) 137–148.
- [81] A. Seki, W. Coombs, S.M. Taffet, M. Delmar, Loss of electrical communication, but not plaque formation, after mutations in the cytoplasmic loop of connexin43, *Heart Rhythm* 1 (2004) 227–233.
- [82] A. Seki, H.S. Duffy, W. Coombs, D.C. Spray, S.M. Taffet, M. Delmar, Modifications in the biophysical properties of connexin43 channels by a peptide of the cytoplasmic loop region, *Circ. Res.* 95 (2004) 22–28.
- [83] X. Zhang, Y. Qi, Role of intramolecular interaction in connexin50: mediating the Ca²⁺-dependent binding of calmodulin to gap junction, *Arch. Biochem. Biophys.* 440 (2005) 111–117.
- [84] X.S. Yu, X. Yin, E.M. Lafer, J.X. Jiang, Developmental regulation of the direct interaction between the intracellular loop of connexin 45.6 and the C terminus of major intrinsic protein (aquaporin-0), *J. Biol. Chem.* 280 (2005) 22081–22090.
- [85] Y. Zhou, W. Yang, M.M. Lurtz, Y. Ye, Y. Huang, H.W. Lee, Y. Chen, C.F. Louis, J.J. Yang, Identification of the calmodulin binding domain of connexin 43, *J. Biol. Chem.* 282 (2007) 35005–35017.
- [86] E.Y. Kang, M. Ponzio, P.P. Gupta, F. Liu, A. Butensky, D.E. Gutstein, Identification of binding partners for the cytoplasmic loop of connexin43: a novel interaction with beta-tubulin, *Cell Commun. Adhes.* (2009) 1–10.
- [87] C. Nambara, Y. Kawasaki, H. Yamasaki, Role of the cytoplasmic loop domain of Cx43 in its intracellular localization and function: possible interaction with cadherin, *J. Membr. Biol.* 217 (2007) 63–69.
- [88] B.J. Hirst-Jensen, P. Sahoo, F. Kieken, M. Delmar, P.L. Sorgen, Characterization of the pH-dependent interaction between the gap junction protein connexin43 carboxyl terminus and cytoplasmic loop domains, *J. Biol. Chem.* 282 (2007) 5801–5813.
- [89] X. Ouyang, V.M. Winbow, L.S. Patel, G.S. Burr, C.K. Mitchell, J. O'Brien, Protein kinase A mediates regulation of gap junctions containing connexin35 through a complex pathway, *Brain Res. Mol. Brain Res.* 135 (2005) 1–11.
- [90] V.M. Berthoud, E.C. Beyer, W.E. Kurata, A.F. Lau, P.D. Lampe, The gap-junction protein connexin 56 is phosphorylated in the intracellular loop and the carboxy-terminal region, *Eur. J. Biochem.* 244 (1997) 89–97.
- [91] S. Rahman, W.H. Evans, Topography of connexin32 in rat liver gap junctions. Evidence for an intramolecular disulphide linkage connecting the two extracellular peptide loops, *J. Cell Sci.* 100 (1991) 567–578.
- [92] E.B. Trexler, M.V. Bennett, T.A. Bargiello, V.K. Verselis, Voltage gating and permeation in a gap junction hemichannel, *Proc. Nat. Acad. Sci. U.S.A.* 93 (1996) 5836–5841.
- [93] V.K. Verselis, M. Srinivas, Divalent cations regulate connexin hemichannels by modulating intrinsic voltage-dependent gating, *J. Gen. Physiol.* 132 (2008) 315–327.
- [94] V.K. Verselis, M.P. Trelles, C. Rubinos, T.A. Bargiello, M. Srinivas, Loop gating of connexin hemichannels involves movement of pore-lining residues in the first extracellular loop domain, *J. Biol. Chem.* 284 (2009) 4484–4493.
- [95] Q. Tang, T.L. Dowd, V.K. Verselis, T.A. Bargiello, Conformational changes in a pore-forming region underlie voltage-dependent “loop gating” of an unapposed connexin hemichannel, *J. Gen. Physiol.* 133 (2009) 555–570.
- [96] P. Gomes, S.P. Srinivas, W. Van Driessche, J. Vereecke, B. Himpens, ATP release through connexin hemichannels in corneal endothelial cells, *Invest. Ophthalmol. Vis. Sci.* 46 (2005) 1208–1218.
- [97] E. De Vuyst, E. Decrock, L. Cabooter, G.R. Dubyak, C.C. Naus, W.H. Evans, L. Leybaert, Intracellular calcium changes trigger connexin 32 hemichannel opening, *EMBO J.* 25 (2006) 34–44.
- [98] J.A. Orellana, D.E. Hernández, P. Ezan, V. Velarde, M.V. Bennett, C. Giaume, J.C. Sáez, Hypoxia in high glucose followed by reoxygenation in normal glucose reduces the viability of cortical astrocytes through increased permeability of connexin 43 hemichannels, *Glia* 58 (2010) 329–343.
- [99] K.A. Dora, P.E. Martin, A.T. Chaytor, W.H. Evans, C.J. Garland, T.M. Griffith, Role of heterocellular Gap junctional communication in endothelium-dependent smooth muscle

- hyperpolarization: inhibition by a connexin-mimetic peptide, *Biochem. Biophys. Res. Commun.* 254 (1999) 27–31.
- [100] S. Boitano, W.H. Evans, Connexin mimetic peptides reversibly inhibit Ca(2+) signaling through gap junctions in airway cells, *Am. J. Physiol. Lung Cell. Mol. Physiol.* 279 (2000) 623–630.
- [101] J. Wang, M. Ma, S. Locovei, R.W. Keane, G. Dahl, Modulation of membrane channel currents by gap junction protein mimetic peptides: size matters, *Am. J. Physiol. Cell Physiol.* 293 (2007) 1112–1119.
- [102] J.P. Reyes, C.Y. Hernandez-Carballo, G. Perez-Flores, P. Perez-Cornejo, J. Arreola, Lack of coupling between membrane stretching and pannexin-1 hemichannels, *Biochem. Biophys. Res. Commun.* 380 (2009) 50–53.
- [103] J. Kronengold, E.B. Trexler, F.F. Bukauskas, T.A. Bargiello, V.K. Verselis, Single-channel SCAM identifies pore-lining residues in the first extracellular loop and first transmembrane domains of Cx46 hemichannels, *J. Gen. Physiol.* 122 (2003) 389–405.
- [104] I.M. Skerrett, J. Aronowitz, J.H. Shin, G. Cymes, E. Kasperek, F.L. Cao, B.J. Nicholson, Identification of amino acid residues lining the pore of a gap junction channel, *J. Cell Biol.* 159 (2002) 349–360.
- [105] T.F. Liu, H.Y. Li, M.M. Atkinson, R.G. Johnson, Intracellular lucifer yellow leakage from Novikoff cells in the presence of ATP or low extracellular Ca: evidence for hemi-gap junction channels, *Meth. Find. Exp. Clin. Pharmacol.* 17 (1995) 23–28.
- [106] R.P. Kondo, S.Y. Wang, S.A. John, J.N. Weiss, J.I. Goldhaber, Metabolic inhibition activates a non-selective current through connexin hemichannels in isolated ventricular myocytes, *J. Mol. Cell. Cardiol.* 32 (2000) 1859–1872.
- [107] X. Bao, G.A. Altenberg, L. Reuss, Mechanism of regulation of the gap junction protein connexin 43 by protein kinase C-mediated phosphorylation, *Am. J. Physiol. Cell Physiol.* 286 (2004) 647–654.
- [108] F.F. Bukauskas, M.M. Kreuzberg, M. Rackauskas, A. Bukauskiene, M.V. Bennett, V.K. Verselis, K. Willecke, Properties of mouse connexin 30.2 and human connexin 31.9 hemichannels: implications for atrioventricular conduction in the heart, *Proc. Nat. Acad. Sci. U.S.A.* 103 (2006) 9726–9731.
- [109] C.E. Stout, J.L. Costantin, C.C. Naus, A.C. Charles, Intercellular calcium signaling in astrocytes via ATP release through connexin hemichannels, *J. Biol. Chem.* 277 (2002) 10482–10488.
- [110] J. Kang, N. Kang, D. Lovatt, A. Torres, Z. Zhao, J. Lin, M. Nedergaard, Connexin 43 hemichannels are permeable to ATP, *J. Neurosci.* 28 (2008) 4702–4711.
- [111] Z.C. Ye, M.S. Wyeth, S. Baltan-Tekkok, B.R. Ransom, Functional hemichannels in astrocytes: a novel mechanism of glutamate release, *J. Neurosci.* 23 (2003) 3588–3596.
- [112] S. Bruzzone, L. Guida, E. Zocchi, L. Franco, A. De Flora, Connexin 43 hemi channels mediate Ca2+-regulated transmembrane NAD+ fluxes in intact cells, *FASEB J.* 15 (2001) 10–12.
- [113] S. Rana, R. Dringen, Gap junction hemichannel-mediated release of glutathione from cultured rat astrocytes, *Neurosci. Lett.* 415 (2007) 45–48.
- [114] P.P. Cherian, A.J. Siller-Jackson, S. Gu, X. Wang, L.F. Bonewald, E. Sprague, J.X. Jiang, Mechanical strain opens connexin 43 hemichannels in osteocytes: a novel mechanism for the release of prostaglandin, *Mol. Biol. Cell* 16 (2005) 3100–3106.
- [115] S. Ahmad, W.H. Evans, Post-translational integration and oligomerization of connexin 26 in plasma membranes and evidence of formation of membrane pores: implications for the assembly of gap junctions, *Biochem. J.* 365 (2002) 693–699.
- [116] J.K. VanSlyke, L.S. Musil, Cytosolic stress reduces degradation of connexin43 internalized from the cell surface and enhances gap junction formation and function, *Mol. Biol. Cell* 16 (2005) 5247–5257.
- [117] F.F. Bukauskas, V.K. Verselis, Gap junction channel gating, *Biochim. Biophys. Acta* 1662 (2004) 42–60.
- [118] J.C. Sáez, M.A. Retamal, D. Basilio, F.F. Bukauskas, M.V. Bennett, Connexin-based gap junction hemichannels: gating mechanisms, *Biochim. Biophys. Acta* 1711 (2005) 215–224.
- [119] J.M. Gómez-Hernandez, M. de Miguel, B. Larrosa, D. González, L.C. Barrio, Molecular basis of calcium regulation in connexin-32 hemichannels, *Proc. Nat. Acad. Sci. U.S.A.* 100 (2003) 16030–16035.
- [120] M.C. Puljung, V.M. Berthoud, E.C. Beyer, D.A. Hanck, Polyvalent cations constitute the voltage gating particle in human connexin37 hemichannels, *J. Gen. Physiol.* 124 (2004) 587–603.
- [121] J. Thimm, A. Mechler, H. Lin, S. Rhee, R. Lal, Calcium-dependent open/closed conformations and interfacial energy maps of reconstituted hemichannels, *J. Biol. Chem.* 280 (2005) 10646–10654.
- [122] D.J. Müller, G.M. Hand, A. Engel, G.E. Sosinsky, Conformational changes in surface structures of isolated connexin 26 gap junctions, *EMBO J.* 21 (2002) 3598–3607.
- [123] M. Srinivas, D.P. Calderon, J. Kronengold, V.K. Verselis, Regulation of connexin hemichannels by monovalent cations, *J. Gen. Physiol.* 127 (2006) 67–75.
- [124] L. Ebihara, X. Liu, J.D. Pal, Effect of external magnesium and calcium on human connexin46 hemichannels, *Biophys. J.* 84 (2003) 277–286.
- [125] M. Srinivas, D.P. Calderon, J. Kronengold, V.K. Verselis, Regulation of connexin hemichannels by monovalent cations, *J. Gen. Physiol.* 127 (2006) 67–75.
- [126] M. Kamermans, L. Fahrenfort, Ephaptic interactions within a chemical synapse: hemichannel-mediated ephaptic inhibition in the retina, *Curr. Opin. Neurobiol.* 14 (2004) 531–541.
- [127] Z. Sun, D.Q. Zhang, D.G. McMahon, Zinc modulation of hemi-gap-junction channel currents in retinal horizontal cells, *J. Neurophysiol.* 101 (2009) 1774–1780.
- [128] R.L. Chappell, J. Zakevicius, H. Ripps, Zinc modulation of hemichannel currents in *Xenopus* oocytes, *Biol. Bull.* 205 (2003) 209–211.
- [129] D. Francis, K. Stergiopoulos, J.F. Ek-Vitorin, F.L. Cao, S.M. Taffet, M. Delmar, Connexin diversity and gap junction regulation by pHi, *Dev. Genet.* 24 (1999) 123–136.
- [130] H. Ripps, H. Qian, J. Zakevicius, Properties of connexin26 hemichannels expressed in *Xenopus* oocytes, *Cell. Mol. Neurobiol.* 24 (2004) 647–665.
- [131] B. Jedamzik, I. Marten, A. Ngezahayo, A. Ernst, H.A. Kolb, Regulation of lens rCx46-formed hemichannels by activation of protein kinase C, external Ca(2+) and protons, *J. Membr. Biol.* 173 (2000) 39–46.
- [132] M. Romanello, B. Pani, M. Bicego, P. D'Andrea, Mechanically induced ATP release from human osteoblastic cells, *Biochem. Biophys. Res. Commun.* 289 (2001) 1275–1281.
- [133] M. Garcia, M.M. Knight, Cyclic loading opens hemichannels to release ATP as part of a chondrocyte mechanotransduction pathway, *J. Orthop. Res.* (2009).
- [134] L. Bao, F. Sachs, G. Dahl, Connexins are mechanosensitive, *Am. J. Physiol.* 287 (2004) 1389–1395.
- [135] A. Sipos, S.L. Vargas, I. Toma, F. Hanner, K. Willecke, J. Peti-Peterdi, Connexin 30 deficiency impairs renal tubular ATP release and pressure natriuresis, *J. Am. Soc. Nephrol.* 20 (2009) 1724–1732.
- [136] A.J. Siller-Jackson, S. Burra, S. Gu, X. Xia, L.F. Bonewald, E. Sprague, J.X. Jiang, Adaptation of connexin 43-hemichannel prostaglandin release to mechanical loading, *J. Biol. Chem.* 283 (2008) 26374–26382.
- [137] L. Ebihara, E. Steiner, Properties of a nonjunctional current expressed from a rat connexin46 cDNA in *Xenopus* oocytes, *J. Gen. Physiol.* 102 (1993) 59–74.
- [138] M.A. Retamal, S. Yin, G.A. Altenberg, L. Reuss, Voltage-dependent facilitation of Cx46 hemichannels, *Am. J. Physiol. Cell Physiol.* 298 (2010) 132–139.