

## Deletion of a single allele of Cx43 is associated with a reduction in the gap junctional intercellular communication and increased cell proliferation of mouse lung pneumocytes type II

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**Abstract.** *Objectives:* Connexins (Cx) are proteins that form the gap junctional channels at neighbouring plasma membranes between adjacent cells. Cxs are involved in cell communication, which is reportedly correlated with cell proliferation and differentiation. Alterations in connexin expression and/or gap junctional intercellular communication (GJIC) capacity have long been postulated to be important in a number of pathological conditions including cancer. This study was performed to determine the consequences of the deletion of a single allele of *Gjal* (Cx43 gene) in Alveolar Type II cells (APTII), and its impact on GJIC and cell proliferation. *Material and methods:* In order to do so, APTII from wild type (Cx43<sup>+/+</sup>) and heterozygous (Cx43<sup>+/-</sup>) mice were harvested and cultured for 4 days. The GJIC capacity was evaluated by scrape-loading method, with the transfer of lucifer yellow dye. The expression of Cx43 was evaluated by immunofluorescence method and Western blotting. Cell proliferation was evaluated by 3-(4,5-dimethylthazol-2-yl)-2,5-diphenyltetrazolium bromide assay. *Results:* It was observed that GJIC capacity was significantly reduced and cell proliferation index was significantly higher in Cx43<sup>+/-</sup> cells compared to Cx43<sup>+/+</sup> cells. *Conclusions:* These results show that knocking out one allele of Cx43 leads to a lower cell to cell communication capacity, and consequently induces a higher cell proliferation. Because chemically induced lung adenomas in mice are known to originate from APTII, these alterations may play a critical role in their susceptibility to lung carcinogenesis.

### INTRODUCTION

Gap junctions are clusters of channels formed by special type of proteins named the connexins (Cx) that form the gap junctions. They allow the direct passage of small molecules less than

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1 kDa between the cytoplasmic compartments of two adjacent cells (Goodenough *et al.* 1996). Several molecules can easily diffuse through gap junctions, such as secondary messengers,  $\text{Ca}^{2+}$  and  $\text{IP}_3$  (Saez *et al.* 1989; Charles *et al.* 1991; Finkbeiner 1992). About 21 different connexins with different patterns of tissue distribution and developmental expression have been reported.

Gap junctional intercellular communication (GJIC) is known to play an important role in maintaining homeostasis, development and differentiation of many tissues. Impairment of GJIC and decreased Cx expression has been reported to be associated with carcinogenic process (Trosko & Ruch 1998).

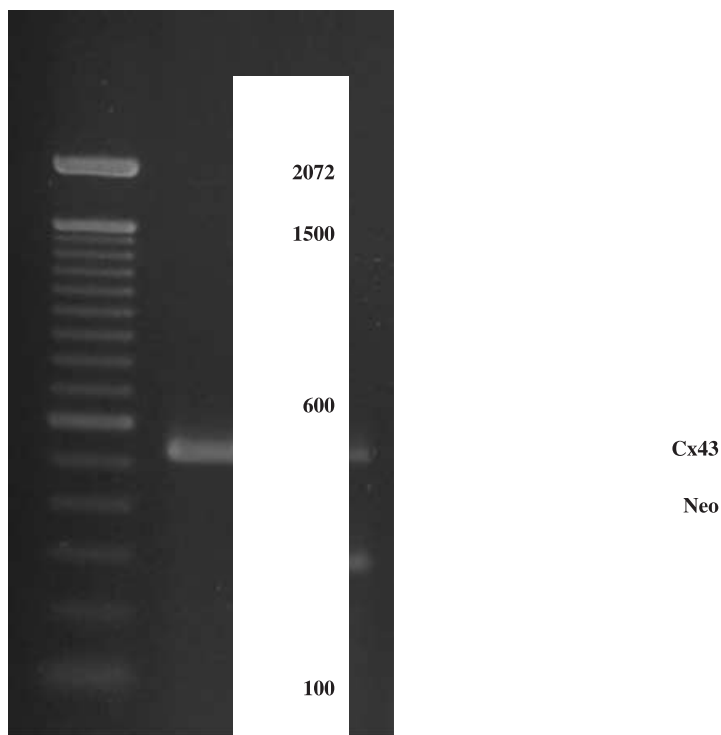
Furthermore, studies on reintroduction of connexins expression in cancer cells showed that these proteins have a role in tumour suppression. For example, expression of Cx26 gene has been reported to reduce growth of HeLa cells after transfection (Mesnil *et al.* 1995). In addition, regenerative growth, tumorigenesis and tumour progression are associated with a reduction in the levels of GJIC (Fehrenbach 2001). Similarly, reduction in gap junction function and expression of Cx genes has been reported using several growth factors, oncogenes and tumour promoters (Ohshima *et al.* 1985). This reduction or complete loss of GJIC appears to be providing an advantage to cancer cells to escape tissue-specific growth control signals, and helps in abnormal cellular proliferation. Abnormalities in the expressing GJIC have been reported in several solid tumours namely liver, breast, bladder and skin (Rehm *et al.* 1988).

There is supporting evidence that vast majority of chemically induced lung tumours and/or spontaneous pulmonary neoplasms originate from alveolar type II epithelial cells (APTII) (Hennemann *et al.* 1992; Traub *et al.* 1994; Mason *et al.* 2000). Normally, in lung tissue Cx26, Cx32, Cx43 and Cx46 are expressed (Kasper *et al.* 1996; Lee *et al.* 1997; Abraham *et al.* 1999; Avanzo *et al.* 2004; King & Lampe 2004), whereas APTII cells in particular express high levels Cx43 (Lee *et al.* 1997; Abraham *et al.* 1999). However, their functional role in normal lung as well as during lung tumour development is not clear and still needs to be investigated. Furthermore, recent reports suggests that both *Gjal* (Cx43 gene) and Cx32 genes function as tumour suppressors in lung (Avanzo *et al.* 2004; King & Lampe 2004). Accordingly, in the present study, in order to understand further role of Cx43, we have made an attempt to study the impact of deletion of a single Cx43 allele on GJIC, Cx43 expression patterns and cell proliferation. We wish to report in this study that the deletion of a single Cx43 allele results in the reduction of GJIC capacity in APTII cells, which, is associated with an increase in cell proliferation. Because chemically induced lung adenomas in mice are known to originate from APTII, these alterations may play a critical role in their susceptibility to lung carcinogenesis.

## MATERIALS AND METHODS

### Mice

Cx43<sup>+/-</sup> mice were obtained from the International Agency for Research on Cancer (IARC, Lyon, France). These mice were generated by replacing the exon-2 of the Cx43 gene by the neomycin resistance gene (Reaume *et al.* 1995). They were originally produced in the C57BL/6 strain, but their background was subsequently changed to CD1 by serial breeding at the IARC animal facility. Due to the postnatal lethality of Cx43-knockout mice, only heterozygous (Cx43<sup>+/-</sup>) and wild type (Cx43<sup>+/+</sup>) APTII from male mice were used in this study. Cx43 expression has been well characterized by real-time polymerase chain reaction (PCR) and Western blot in these Cx43<sup>+/-</sup> mice in comparison to Cx43<sup>+/+</sup> mice (Avanzo *et al.* 2004). Heterozygous mice indeed show reduced mRNA amount for Cx43 and lower Cx43 levels.



**Figure 1.** Electrophoresis agarose gel of PCR-amplified Cx43 and neo-sequences of DNA extracted of tails from Cx43<sup>+/-</sup> and Cx43<sup>+/+</sup> mice. Bands corresponding to the Cx43 and neo-sequences are clearly visible after ethidium bromide staining. MW, molecular weight; +/+, Cx43 wild-type; +/-, Cx43 heterozygous.

### Genotyping By PCR

DNA from each mouse was obtained from tail biopsies. Genotyping was performed by PCR, as described by Yamakage *et al.* (1998). The following Cx43 primers were used: sense 5'-CCCCACTCTCACCTATGTCTCC-3' and antisense 5'-ACTTTTGCCGCCTAGCTATCCC-3' observed at 520 bp; neo-sense 5'-GGCCACA GTCGATGAATCCAG-3' and antisense 5'-TATC-CATCATGGCTGATGCAA-3' observed at 294 bp (Fig. 1). After genotyping, the rodents were kept under controlled conditions, from 6 to 10 weeks of age.

### Isolation and culture of APTIS

Raw cell suspensions were prepared from female or male Cx43<sup>+/+</sup> and Cx43<sup>+/-</sup> mouse lungs, using a modification of recently published methods (Corti *et al.* 1996; Dobbs & Gonzalez 2002; Rice *et al.* 2002). Briefly, the mice were anaesthetized with Avertin (250–300 µl/10 g of body weight, intraperitoneal injection). The trachea was isolated and cannulated with a 20-gauge luer stub adapter. The diaphragm was cut open, and the chest plate and thymus were removed. The lungs were perfused with approximately 10 ml of warm saline solution (37 °C) via the right ventricle with a 21-gauge needle fitted on a 10-ml syringe. Dispase (DP, 3 ml at 4 °C) (Becton Dickinson Bioscience, Le Pont de Claix, France) and agarose (0.5 ml, 45 °C) were quickly instilled in the trachea. The lungs were immediately covered with ice for 2 min to solidify the agarose gel, and were then removed and incubated in 1 ml of DP for 45 min at room temperature. The lungs were subsequently transferred to a 60-mm culture dish containing 7 ml of

4-(2-hydroxyethyl)-1-piperazineethanesulfonic acid-buffered Dulbecco's modified Eagle's medium (DMEM) and 100 U/ml DNase I at 4 °C. The lung tissue was gently detached from the bronchi. The cell suspension was filtered through progressively smaller cell strainers (100 µm and 40 µm; BD Bioscience) and nylon gauze (20 µm, Millipore, Molsheim, France). The cells were collected by centrifugation at 130 *g* for 8 min (4 °C) and re-suspended in 2 ml of DMEM without foetal bovine serum.

### **Purification of APTIIs from raw cell suspensions**

The cells were incubated in 500 µg immunoglobulin G (IgG) (Sigma-Aldrich, St. Louis, MO, USA) for 1 h, at 37 °C and 5% CO<sub>2</sub>. After incubation, the cells were re-centrifuged. The resultant pellet was re-suspended in DMEM with 10% foetal bovine serum, 36.6 mg/ml of L-glutamine (Sigma), 100 µg/ml of streptomycin (Sigma), 100 U/ml of penicillin (Sigma) and 10 ng/ml of keratinocyte growth factor (KGF) (Sigma). This suspension was pre-plated for a period not shorter than 4 h in order to eliminate fibroblasts. The purified APTIIs were plated in Matrigel (Sigma)/collagen (70 : 30 v/v) at 37 °C and 5% CO<sub>2</sub> according to Rice *et al.* (2002). The medium was changed after 24 h, and every 48 h thereafter. After the enzymatic digestion with DP and the mechanical dissociation of the lungs, around  $10.7 \pm 5.5 \times 10^6$  cells were recovered per mouse ( $n = 97$ ). IgG purification reduced the final concentration to  $6.06 \pm 2.3 \times 10^6$  cells, with a recovery of  $91.04 \pm 1.95\%$ .

### **Preparation of IgG plates**

Non irradiated plastic plates (for bacterial growth) were added with 100 µl IgG diluted in phosphate-buffered saline (PBS), pH 7.3, with 0.1% bovine serum albumin. The plates were kept for 3 h at 25 °C. This solution was removed and the plates were washed five times with PBS (1 ml), and once with DMEM.

### **Collagen preparation**

A solution-containing type IV collagen was prepared according to Michalopoulos & Pitot (1975). Approximately 1 g of collagen fibres were obtained from two Wistar rats. The fibres were irradiated by ultraviolet light for 24 h and dissolved in 300 ml of a 1 : 1000 acetic acid solution in sterile water, and stirred for 24 h at 4 °C. The suspension was allowed to rest for 24 h at 4 °C, after which the supernatant was recovered.

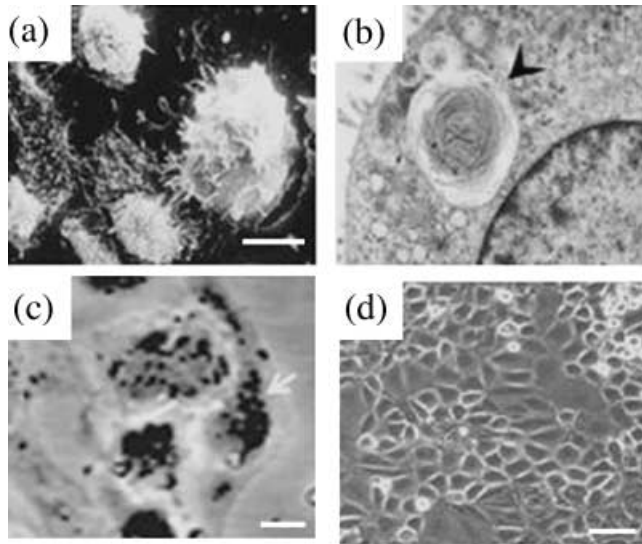
A 10 ml collagen solution was vigorously mixed with DMEM (10%) and the pH was adjusted to 7.0 with NaOH. Seven milliliters of Matrigel were added to each 3 ml of the collagen solution. A volume of 250 µl was used in each plate. Collagen polymerization was carried out for 30 min. The supernatant was aspirated and the plates were dried immediately before use.

### **Cell viability**

The cellular viability was determined by the dye exclusion method using the vital dye trypan blue at 0.4% in PBS, pH 7.4.

### **Osmium tetroxide/tannic acid staining**

Cells cultured for 2 days were fixed for 15 min in 1.5% glutaraldehyde, washed with PBS and post-fixed with 1% osmium tetroxide for 1 h, followed by enough volume of tannic acid (1%, pH 6.8) to cover the fixed cultures overnight. APTIIs were visually checked for morphology and stained for lamellar bodies, an APTII marker, in order to determine the cell type in culture (Fig. 2c). APTIIs were identified in cultures by the presence of dark, lamellate cytoplasmic inclusions. For quantification purposes, about 500 cells were counted and the results were expressed as percentage. Only attached and spread cells were quantified.



**Figure 2. Morphological appearance of Alveolar Type II (APTII) cells *in vitro*.** APTII cells were isolated from mouse lungs, and grown in culture to determine phenotype, as described in MATERIALS AND METHODS. (a) Scanning electron micrograph of APTII at day 2 of isolation. (b) Transmission electronmicrograph of APTII cells from Cx43<sup>+/-</sup> male mice, showing lamellar body in the cytoplasm ( $\times 25\,600$ ) (arrowhead). (c) Cells stained with tannic acid on day 2 of culture, APTII were densely packed and displayed distinct and numerous lamellar bodies (large dark granules) in  $> 95\%$  of the cells in culture (arrowheads) in cells from Cx43<sup>+/-</sup> mice. (d) Phase contrast micrographs on day 4 representing cultured Cx43<sup>+/-</sup> and Cx43<sup>+/+</sup> APTIIs. Bars, 10  $\mu\text{m}$  (c) and 50  $\mu\text{m}$  (d).

### Electron microscopy

Cultures of APTII cells on 0.4  $\mu\text{m}$  Millicell-PC (Millipore) membranes were washed with PBS, and observed after 12 h of plating (Hasegawa *et al.* 2001). The cells were then fixed with glutaraldehyde 2.5% in PBS, post-fixed in osmium tetroxide 1% in PBS, dehydrated in a graded ethanol series, and embedded in Epon. Blocks of interest were cut from the dishes and visualized by phase contrast. Microscopic sections for ultra-structure analysis were contrasted with uranyl acetate for 10 min and observed in an electron microscope.

### Rna extraction, cDNA construction and real-time PCR analysis of Cx43 expression in lung

Total RNA was extracted from tissues by a single-step technique with TRIzol Reagent (Invitrogen Life Technologies, Carlsbad, CA, USA) according to the manufacturer's protocol. The quality of the RNA samples was determined by electrophoresis in 2% agarose gels and staining with ethidium bromide. 18S and 28S bands were visualized under ultraviolet light (not shown). Total RNA was then treated with DNase I (Invitrogen Life Technologies) before processing further.

Oligo DT (1  $\mu\text{l}$ ) and dNTPs (1  $\mu\text{l}$ ) were added to the total RNA sample (500  $\mu\text{g}$ ) and incubated at 65  $^{\circ}\text{C}$  for 5 min. After that, Buffer 5X (Superscript II) (4  $\mu\text{l}$ ), DTT (1 M, 2  $\mu\text{l}$ ) and RNase OUT (1  $\mu\text{l}$ ) were added, and the mixture was incubated at 42  $^{\circ}\text{C}$  for 2 min. Again Superscript II (1  $\mu\text{l}$ ) was added, followed by incubation at 42  $^{\circ}\text{C}$  for 50 min and subsequent incubation was performed at 70  $^{\circ}\text{C}$  for 15 min. For the removal of remnant RNA, 1  $\mu\text{l}$  RNase H was added, and the mixture incubated at 37  $^{\circ}\text{C}$  for 20 min. All reagents were purchased from Invitrogen Life Technologies.

Real-time PCR analysis was performed in ABI Prism<sup>®</sup>. 7000 Sequence Detection Systems (Applied Biosystems, Foster City, CA, USA) using TaqMan Universal Master Mix (Applied Biosystems, Part number 4304437). PCR primers and the TaqMan probes for Cx43 and

glyceraldehyde-3-phosphate dehydrogenase (GAPDH) quantification were selected using the Primer Express software programmes (Applied Biosystems) and checked by a basic local alignment search tool search of GenBank. The Cx43 forwards primer, 5'-GTGCCGGCTTCACTTTCA TTAAG-3', and the Cx43 reverse primer, 3'-CCAAGGCGCTCCAGTCA-5', were chosen to amplify a 65-bp fragment. Primers to Cx43 were designed from separate exons (GenBank accession numbers L\_10387 and L\_10388 for forwards and reverse primers, respectively). The internal Cx43 TaqMan probe (FAM-3'-TCTGGGCACCTCTCTTT-5'-NFQ) was designed following the general rules outlined by the manufacturer. The GAPDH primers, 5'-CATGGCCTTCCGTGTTCCCTA-3' forward primer and 3'-GCGGCACGTCAGAT CCA-5' reverse primer (55-bp) (GenBank NM\_008084), and the TaqMan probe VIC-5'-CCCCCAATGTGTCCGTC-3'-MGBNFQ were used as a housekeeping control. The TaqMan probes carried a 5'-reporter dye 6-carboxy-fluorescein (FAM to Cxs) or (VIC for GAPDH), a 3'-non fluorescent quencher dye (NFQ) and a minor groove binding (MGB). The primers and probes were used with 100% efficiency at the final concentrations of 0.9  $\mu\text{M}$  and 0.25  $\mu\text{M}$ , respectively.

The thermal cycling conditions to cDNA quantification assays were established according to ABI Prism® 7000 sequence detection systems parameters (Applied Biosystems). Analysis of relative gene expression data was performed according to the  $2^{-\Delta\Delta\text{CT}}$  method (Livak & Schmittgen 2001).

### **Cx43 immunostaining**

For *in vitro* assays,  $5 \times 10^6/\text{ml}$  cells were seeded in eight-well permanox slide (Laboratory-Tek Brand Products, Naperville, IL, USA) and cultured for 4 days. The medium was removed and the cells were washed three times with PBS, pH 7.3 at 37 °C. Cells were permeabilized for 2 min with methanol : acetone 1 : 1 at room temperature. Then, slides were washed five times in PBS and blocked in block solution (PBS, 0.5% Triton X-100, 2% heat-inactivated goat serum) for 45 min. Monoclonal anti-Cx43 diluted 1 : 100 (Zymed, San Francisco, CA, USA) was used.

### **Immunoblotting**

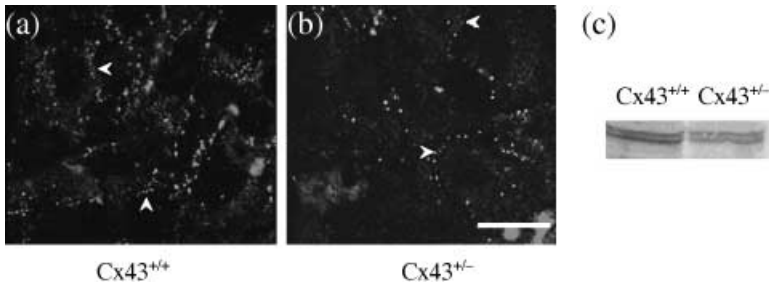
Confluent cultures of APTII cells in 35-mm dishes were washed with PBS and directly lysed in sodium dodecyl sulfate (SDS) sample buffer (60 mM Tris-HCl, pH 6.8, 0.5% SDS). Protein extracts were subjected to SDS-polyacrylamide gel electrophoresis and transferred to a nitrocellulose membrane. Non-specific staining was blocked by 3% bovine serum albumin overnight and the blots were incubated with polyclonal rabbit anti-Cx43 antibody (1/1000, 2 h) followed by incubation with a secondary peroxidase-conjugated antirabbit antibody (1/500). A 3,3'-diaminobenzidine solution was used to reveal Cx43 protein.

### **Studies of cell-cell communication capacity by scrape-loading technique**

Gap junctional intercellular communication capacity was assessed by the technique of scrape loading described by El-Fouly & colleges (1987). Briefly, GJIC capacity was evaluated by the intercellular transfer of 5% Lucifer yellow (Sigma), in PBS after 5 min. The cells were fixed in 4% paraformaldehyde and photographs were taken with a 40X objective. Stained cells were counted in each side of rhodamine albumin in triplicate experiments.

### **Cell proliferation, measured by the MTT assay**

APTII cells were plated in 35 mm dishes and the cell proliferation index was measured by reduction of the yellow dye 3-(4,5-dimethylthazol-2-yl)-2,5-diphenyltetrazolium bromide (MTT, Sigma), to a blue formazan product. Optical density of the blue formazan product was measured at 570 nm in a Shimadzu spectrophotometer. Triplicate experiments were performed.



**Figure 3. Localization of Cx43 in Alveolar Type II (APTII) cells by indirect immunofluorescence method on day 4 of culture.** For the identification of Cx43-expression in APTII cells from Cx43<sup>+/-</sup> and Cx43<sup>+/+</sup>, APTIIs were collected with Dispase, cultivated on glass slides for 4 days, permeabilized, immunostained with anti-Cx43 (a and b) antiserum followed by rhodamine-conjugated goat antirabbit immunoglobulin G (IgG) secondary antibody, and imaged in fluorescence microscopy. Negative control (APTII cells submitted to immunocytochemistry but omitting the primary antibody) not showed. Note that Cx43 was found in cell-to-cell region in both Cx43<sup>+/+</sup> (a) and Cx43<sup>+/-</sup> (b) mice (arrowheads). Cx43 was higher in the membrane area of Cx43<sup>+/+</sup> mice in comparison to Cx43<sup>+/-</sup> mice. These experiments were repeated twice for each plate with similar results. Bar, 10  $\mu$ m. (c) Immunoblots of Cx43 in APTII cells. APTII cells cultured were solubilized, resolved by sodium dodecyl sulfate-polyacrylamide gel electrophoresis, transferred to polyvinylidene difluoride, and blotted with anti-Cx43. Kaleidoscope pre-stained standards were used as loading control (not showed). For positive controls, HeLa cells transfected with the Cx43 gene was used (results not showed). Cx43 clones were kindly supplied by Dr Mesnil. These experiments were repeated twice for each Cx with similar results.

### Statistical analysis

All results were analyzed with a two-tail Student's *t*-test. Values of  $P < 0.05$  were considered significant.

## RESULTS

### Characterization of APTII cells in culture

Morphology and presence of lamellar body were investigated to confirm the identity of APTIIs used in this study (Fig. 2a,b). About  $95.9 \pm 0.2\%$  of isolated cells were typical APTIIs after osmium tetroxide staining (Fig. 2c). A total of  $92.9 \pm 4.0\%$  of cells were viable by the trypan blue exclusion test. APTIIs used in this study showed characteristic morphology in a 4-day culture (Fig. 2d).

### Connexin localization in APTIIs

Immunostaining for Cx43 resulted in a punctuate pattern on the plasma membranes of both Cx43<sup>+/+</sup> and Cx43<sup>+/-</sup> cells (Fig. 3a,b). However, the intensity was much higher in Cx43<sup>+/+</sup> cells in comparison to Cx43<sup>+/-</sup> cells.

### Expression of connexins by real-time PCR and immunoblot analysis

Western blot was performed in 4-day cultures of APTII. Cx43 is less expressed in Cx43<sup>+/-</sup> APTIIs than in Cx43<sup>+/+</sup> APTIIs and showed a difference of 50% in relation to heterozygous mice (Table 1 and Fig. 3).

**Table 1.** Levels of normalized Connexins mRNA in mouse lungs determined by quantitative RT-PCR<sup>a</sup>

Groups	Difference in cycles between Cx43 and GAPDH $\Delta$ CT	Normalized Cx43 amount $2^{-\Delta\Delta$ CT
Cx43 <sup>+/+</sup>	6.7 $\pm$ 0.2	1.2 $\pm$ 0.03
Cx43 <sup>+/-</sup>	8.1 $\pm$ 0.7	0.5 $\pm$ 0.04*

<sup>a</sup>Using reverse transcriptase, cDNA was synthesized from 500  $\mu$ g of total RNA. Aliquots of cDNA were used as templates for real-time PCR reactions containing primers and probes for Cx43, or primers and probe for GAPDH. Each reaction contained cDNA derived from 20 ng of total RNA. Eight different lungs from each group were analyzed, and four replicates of each reaction were performed by RT-PCR.

\* $P < 0.05$ , Tukey-Kramer test.

**Table 2.** Growth curves representing MTT-test of APTII cells from Cx43<sup>+/-</sup> and Cx43<sup>+/+</sup> mice in primary culture

Groups	Optical density		
	4 Days	6 Days	8 Days
Cx43 <sup>+/+</sup>	0.660 $\pm$ 0.043	1.113 $\pm$ 0.042	0.964 $\pm$ 0.011
Cx43 <sup>+/-</sup>	0.901 $\pm$ 0.010*	1.588 $\pm$ 0.015*	1.096 $\pm$ 0.018*

Cells ( $5 \times 10^4$ /ml) were plated in 35 mm dishes and cultured in conditioned medium, as described in METHODS. Results obtained from Cx43<sup>+/-</sup> mice were statistically significant in each culture periods. These results are represented as means from triplicate experiments. \*Significant results ( $P < 0.05$ ).

### Evaluation of APTII proliferation

Table 2 shows the proliferation of APTII cells at two culture periods. At both the time-points, APTIIs from Cx43<sup>+/+</sup> mice showed a lower proliferation rate in comparison to Cx43<sup>+/-</sup> cells ( $P < 0.05$ ), quantified by the MTT-test. At 8 days in culture, APTII cells reached confluence and started dying.

### GJIC capacity in cultured APTII cells by the scrape-loading technique

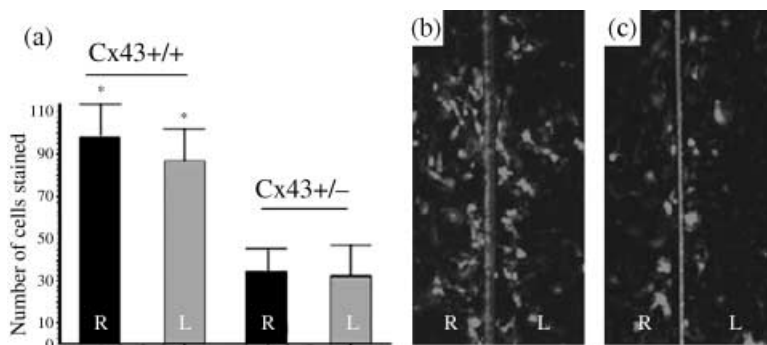
This technique was performed on the fourth day after APTII seeding. The amplitude of lucifer yellow diffusion after 5 min was  $90 \pm 8$  in APTIIs from Cx43<sup>+/+</sup> groups and  $38 \pm 6$  in APTIIs from Cx43<sup>+/-</sup> groups (Fig. 4), representing a statistically significant difference ( $P < 0.05$ ).

## DISCUSSION

It has been reported that chemically induced pulmonary adenomas originate from APTII cells (Mason *et al.* 2000). We have earlier reported the higher incidence of urethane-induced lung adenomas in Cx43<sup>+/-</sup> mice compared to Cx43<sup>+/+</sup> mice (Avanzo *et al.* 2004). In this study, we have examined the consequences of the deletion of one allele of *Gjal* (Cx43 gene) for GJIC capacity and the associated alterations in the proliferation of APTIIs.

In order to accomplish this goal, we have first standardized the procedure to isolate these cells and maintain them in culture. It is known that maintenance and differentiation of alveolar cells in culture can be profoundly altered by both the soluble and insoluble macromolecules that





**Figure 4.** Gap junctional intercellular communication (GJIC) detected by the scrape loading and dye transfer (SLDT) method in Alveolar Type II (APTII) cells on day 4 of culture. Confluent monolayers were scraped with a sharp scalpel and loaded with Lucifer Yellow (LY, 457.25 kDa) (5% in PBS) for 5 min at room temperature. Observations were performed under epifluorescence. (a) Number of APTIIs from Cx43<sup>+/+</sup> mice stained in green was counted in each side of the rhodamine-albumin line (red, 10 kDa) used as a negative control. (b) APTIIs from Cx43<sup>+/+</sup> mice. (c) APTIIs from Cx43<sup>+/-</sup> mice. Note in the centre lane that rhodamine-albumin remains on the scraped area. These experiments were repeated three times for each plate with similar results. For each experiment, 10<sup>7</sup> APTII cells were plated initially. On day 4 of culture the cells reached more than 90% of confluence and the cells were scraped in region with complete confluence. \*Represent significant results ( $P < 0.05$ ).

compose the extracellular matrix, and the developmental fate of APTII cells can be altered by changing matrix or growth factor components (Isakson *et al.* 2001). We used a combination of three different methods to establish the primary culture from APTII mouse cells (Corti *et al.* 1996; Dobbs & Gonzalez 2002; Rice *et al.* 2002). The matrigel : collagen type IV matrix (70 : 30 v/v) plus Dulbecco's Modified Eagle's Medium (DMEM) 10/KGF allowed us to keep APTIIs in culture with an abundant presence of lamellar bodies, compared to a previous study that utilized type I collagen/fibronectin matrix supplemented with KGF (Isakson *et al.* 2001).

We also determined the presence of Cx43 in gap junction plaques in culture. On the basis of immunofluorescence studies and real-time PCR analysis, Cx43 of APTIIs showed lower levels of expression in Cx43<sup>+/-</sup> compared to Cx43<sup>+/+</sup> cells. These results were confirmed by immunoblotting and were in agreement with GJIC results. Our scrape-loading study with lucifer yellow dye-coupling showed that the GJIC capacity was lower in Cx43<sup>+/-</sup> derived APTII cells. This study of communication capacity is important, because it allows the quantification of functional parameters of gap junctions. Impaired GJIC has been shown to correlate with neoplastic transformation in several human tissues, such as breast (Laird *et al.* 1999), brain (Soroceanu *et al.* 2001), endometrium (Saito *et al.* 2001), prostate (Hossain *et al.* 1999) and lung (Ruch *et al.* 2001). This result is in accordance with our previous study showing that Cx43<sup>+/-</sup> mice are more susceptible to lung carcinogenesis. During Cx43 deficiency, adenoma cells from male or female Cx43<sup>+/-</sup> mice presented higher numbers of S-phase cells than those from Cx43<sup>+/+</sup> mice, suggesting that APTIIs are implicated in this phenomenon (Mason & Shannon 1997; Avanzo *et al.* 2004).

Several molecules can pass through gap junction channels and could directly influence the cell growth. After cell proliferation analyses using MTT-test, we observed that isolated APTIIs cultured from Cx43<sup>+/-</sup> mice showed a higher proliferation rate in comparison to Cx43<sup>+/+</sup> APTIIs. According to Loewenstein (1979) and Trosko & Ruch (1998), the communication-deficient cells may not exchange neither inhibitory nor stimulatory factors, being unable to obey the homeostatic mechanisms present in its respective tissue and accordingly this may lead to deregulated cell proliferation.

In conclusion, this study shows, for the first time, that the deletion of a single Cx43 allele results in the reduction of GJIC capacity in APTIIs, which is associated with an increase in cell growth rate. Because chemically induced lung adenomas in mice are reportedly originate from APTIIs, these alterations may play a critical role in their susceptibility to lung carcinogenesis.

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