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Stat3 is a positive regulator of gap junctional intercellular communication in cultured, human lung carcinoma cells

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Abstract

Background: Neoplastic transformation of cultured cells by a number of oncogenes such as *src* suppresses gap junctional, intercellular communication (GJIC); however, the role of Src and its effector Signal transducer and activator of transcription-3 (Stat3) upon GJIC in non small cell lung cancer (NSCLC) has not been defined. Immunohistochemical analysis revealed high Src activity in NSCLC biopsy samples compared to normal tissues. Here we explored the potential effect of Src and Stat3 upon GJIC, by assessing the levels of tyr418-phosphorylated Src and tyr705-phosphorylated Stat3, respectively, in a panel of NSCLC cell lines.

Methods: Gap junctional communication was examined by electroporating the fluorescent dye Lucifer yellow into cells grown on a transparent electrode, followed by observation of the migration of the dye to the adjacent, non-electroporated cells under fluorescence illumination.

Results: An inverse relationship between Src activity levels and GJIC was noted; in five lines with high Src activity GJIC was absent, while two lines with extensive GJIC (QU-DB and SK-LuCi6) had low Src levels, similar to a non-transformed, immortalised lung epithelial cell line. Interestingly, examination of the mechanism indicated that Stat3 inhibition in any of the NSCLC lines expressing high endogenous Src activity levels, or in cells where Src was exogenously transduced, did not restore GJIC. On the contrary, Stat3 downregulation in immortalised lung epithelial cells or in the NSCLC lines displaying extensive GJIC actually suppressed junctional permeability.

Conclusions: Our findings demonstrate that although Stat3 is generally growth promoting and in an activated form it can act as an oncogene, it is actually *required* for gap junctional communication both in nontransformed lung epithelial cells and in certain lung cancer lines that retain extensive GJIC.

Keywords: Stat3, Electroporation, Indium-Tin oxide, Gap junctions, Src, Cell to cell adhesion, Lung cancer

Background

Gap junctions are plasma membrane channels that provide a path of direct communication between the interiors of neighboring cells and are formed by the connexin (Cx) family of proteins. An increase in cell proliferation correlates with a reduction in gap junctional, intercellular communication (GJIC) [1]. In fact, a number of oncogene products such as v-Src [2], the polyoma virus middle Tumor antigen, an oncogene which acts by activating Src

family kinases (mT [3,4]), the chaperone Hsp90N [5], vRas [6,7] and others have been shown to interrupt junctional communication.

Extensive evidence has indicated that expression of the Src tyrosine kinase leads to gap junction closure, through phosphorylation of the ubiquitous connexin, Cx43. Src exerts its effect either through direct tyrosine phosphorylation of Cx43, or indirectly, through activation of the serine/threonine, Erk1/2 or protein kinase C family kinases [8]. Examination of levels of tyr-418 phosphorylated, ie activated Src in a number of Non Small Cell Lung Cancer (NSCLC) biopsies previously showed the presence of higher Src activity than the surrounding, non-tumor lung tissue

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[9,10]. However, Src's contribution to GJIC suppression in NSCLC lines and primary cells which may express other oncogenes in addition to Src, or different levels of Src effectors, remains to be determined.

The Signal Transducer and Activator of transcription-3 (Stat3), an important Src downstream effector, is a cytoplasmic transcription factor. Following phosphorylation on tyr-705 by Src, as well as by growth factor or cytokine receptors such as the IL6 family, Stat3 normally dimerises through a reciprocal SH2 domain-phosphotyrosine interaction and translocates to the nucleus, where it induces the transcription of specific genes [11]. The effect of Src upon Stat3 activation in NSCLC lines is at present unclear. Examination of Stat3 levels in certain NSCLC lines demonstrated that Src is a major Stat3 activator, transducing signals from EGFR and IL6 that lead to apoptosis inhibition [12], while in another report [13] Src inhibition in different NSCLC lines was found to actually *increase* Stat3-ptyr705. However, we and others previously demonstrated that cell-to-cell adhesion, as observed at confluence of cultured cells, causes a dramatic increase in Stat3 activity levels in a number of cellular systems ([14-16] reviewed in [17]); for this reason, cell density must be taken into account in the examination of the effect of different factors such as Src upon Stat3 activity levels. In the present report this was achieved by measuring Stat3-ptyr705 phosphorylation and activity levels at a range of densities.

We previously assessed GJIC in a number of lung cancer lines [18]. In the present work GJIC was examined using an apparatus where cells were grown on a glass slide, half of which was coated with electrically conductive, optically transparent, indium-tin oxide. An electrode was placed on top of the cells and an electrical pulse, which opens transient pores on the plasma membrane, was applied in the presence of the fluorescent dye, Lucifer yellow. Although this technique is adequate for a number of lines, the turbulence generated as the electrode is removed can cause cell detachment, which makes GJIC examination problematic. Here, we revisited the question of GJIC levels in lung cancer lines using an improved technique, where the upper electrode is eliminated. This approach is valuable for the electroporation of tumor-derived lines especially at high densities, where cell adhesion to the substratum may be weak. Interestingly, the results revealed that cell density *per se* triggers a dramatic increase in both Cx43 levels and GJIC. Two NSCLC lines, QU-DB and SK-LuCi6 were found to have extensive GJIC, similar to control, nontransformed lung epithelial cells, while GJIC in five other lines was very low or undetectable. Investigation of the mechanism of gap junction closure revealed an inverse relation between Src activity levels and GJIC. Further studies led to the discovery that, unlike Ras inhibition in Src-transformed fibroblasts [19], Stat3 inhibition in NSCLC lines with high

Src activity does not restore GJIC. On the contrary, Stat3 inhibition in lines displaying extensive GJIC (QU-DB, SK-LuCi6) suppressed junctional permeability, indicating that Stat3 activity is actually *required* for the maintenance of gap junction function in these lung cancer lines.

Results

Cell density upregulates GJIC and connexin-43 protein levels

A number of reports showed that gap junction function is dependent upon cell to cell contact and the assembly of adherens junctions [20,21]. Since the opportunity for engagement of cadherins, key components of adherens junctions, is expected to increase with cell density, we examined the effect of cell density upon GJIC. To this effect, we took advantage of the nontransformed mouse lung epithelial type II line, E10 that has extensive GJIC, an even and flat morphology and good adhesion to the substratum even at high densities [22] (Figure 1A). In addition, unlike nontransformed human lung lines such as NL-20 [23], E10 cells can be grown in the absence of growth factors that could affect GJIC. Cells were plated in electroporation chambers and when 90% confluent or at 3 days post-confluence Lucifer yellow was electroporated and the movement of the dye through gap junctions observed under fluorescence and phase contrast illumination (see Methods). The results are presented as the average number of cells where dye transferred, per cell loaded with the dye by electroporation (GJIC). As shown in Figure 1B, *a-c*, although cells at 90% confluence did display some gap junction transfer (GJIC ~1.5), GJIC increased to ~6 at 3 days post-confluence (Figure 1B, *d-f*), indicating that cell density causes a dramatic increase in GJIC (Table 1,A).

We next examined the levels of Cx43, a widely expressed gap junction protein, at different cell densities. Cells were plated in plastic petri dishes at a confluence of 80% and at different times up to 5 days post confluence, total protein extracts were probed for Cx43 by Western blotting. As shown in Figure 1C, cell density caused a dramatic increase in Cx43 levels, which plateaued at ~3 days post-confluence (lane 1 vs 5).

GJIC and connexin-43 in NSCLC lines and freshly explanted tumor cells

In light of the above findings, we examined GJIC levels at different densities up to 4 days post-confluence in a panel of human lung cancer lines [18]. Two NSCLC lines, QU-DB (Figure 2A, *a-c*) and SK-LuCi6 (Table 1,A) displayed extensive GJIC at their peak density, while five NSCLC lines had very low GJIC (e.g. A549, Figure 2B, *a-c*, and Table 1,B). In addition, primary cells explanted and cultured from a moderately differentiated adenosquamous carcinoma

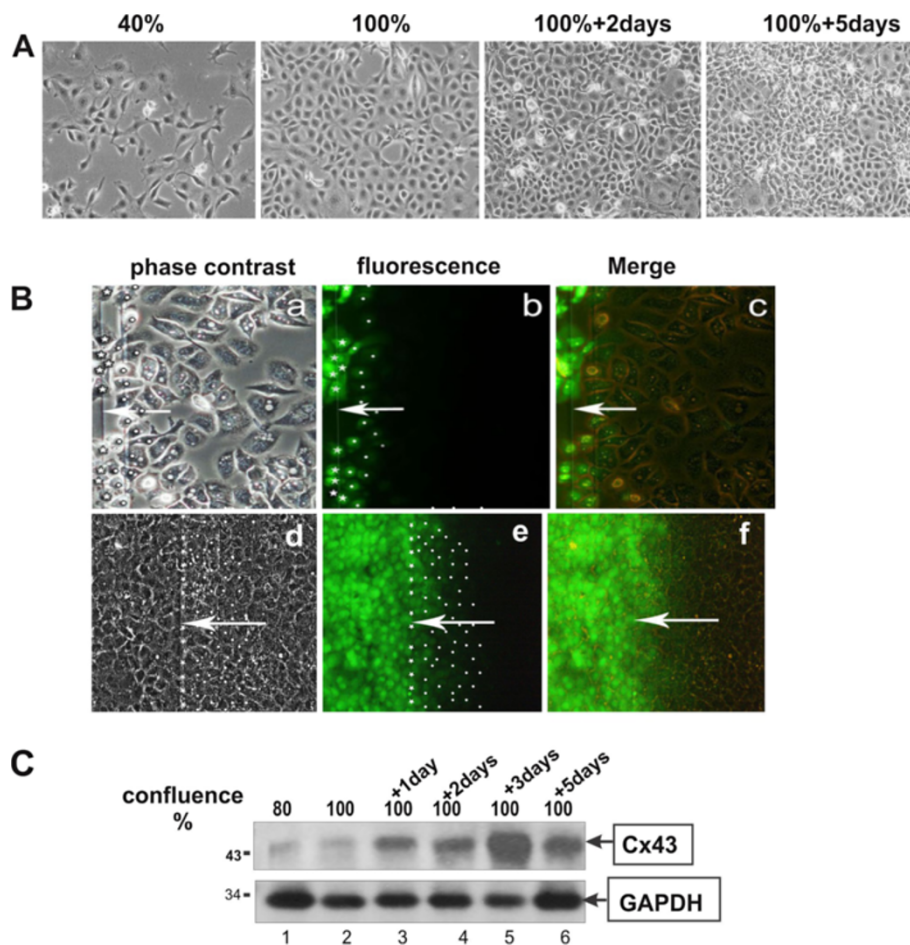


Figure 1 Cell density increases GJIC and Cx43 levels. **A.** Immortalised lung epithelial E10 cells were plated in 3 cm plastic petri dishes, grown to different densities and photographed under phase-contrast illumination. Magnification: 240x. **B.** E10 cells were plated in electroporation chambers and subjected to a pulse in the presence of Lucifer yellow when 90% confluent (**a-c**) or 3 days after confluency (**d-f**) and photographed under phase-contrast (**a, d**), fluorescence (**b, e**) or combined (**c, f**) illumination (see Methods, Figure 7). Arrows point to the position of the edge of the electroporated area. In **a, b, d** and **e**, stars mark cells loaded with the dye at the edge of the electroporated area and dots mark cells into which the dye was transferred through gap junctions. Magnification: 240x. **C.** E10 cells were seeded in plastic petri dishes and when they reached the indicated densities, detergent cell extracts were probed for Cx43 (top) or GAPDH (bottom) as a control.

(Figure 3, **a-b**), a poorly differentiated adenocarcinoma, and an adenocarcinoma (Table 2) had no GJIC.

Examination of Cx43 levels showed that QU-DB cells had levels similar to E10, which increased dramatically with cell density, while Cx43 levels in A549 cells were almost undetectable, at any cell density (Figure 4A). SK-LuCi6 cells had levels similar to QU-DB, while all other NSCLC lines examined had very low Cx43 levels at all densities tested (not shown). The above data taken together indicate that, besides nontransformed epithelial cells, cell density causes a dramatic increase in GJIC and Cx43 protein levels in two lung carcinoma lines which display extensive GJIC. Nevertheless, the majority of lung cancer lines examined (5/7) had very low or no detectable gap junctional communication, even at high cell densities (Table 1,B).

Src activity and GJIC suppression in NSCLC lines

We next examined Src-tyr418 phosphorylation, as an indication of Src activity. As shown in Figure 5A and C, A549 cells displayed high Src-ptyr418 levels, similar to the levels in SK-LuCi6 or E10 cells expressing activated Src by retroviral transduction (lines SK-LuCi6-*Src*, E10-*Src*, respectively, Figure 5C, lanes 1 vs 3 and Table 1,B), while Src-ptyr418 levels in QU-DB cells were low (Figure 5A, lanes 5-8), similar to E10 (Figure 5B, lanes 5 and 6). Lines CALU-1, SW-900, CALU-6 and SK-Lu1 had Src-ptyr418 levels comparable to SK-LuCi6-*Src* (Figure 5B, lanes 1-3 vs 7 and Table 1,B), while SK-LuCi6 had low levels, similar to QU-DB (Figure 5B, lanes 4-5). Examination of gap junctional communication revealed that five lines with high Src-ptyr418 (A549, CALU-1, SW-900, CALU-6, LuCi-1) had very low or no detectable GJIC (Figure 2B,

Table 1 Effect of Stat3 downregulation upon GJIC

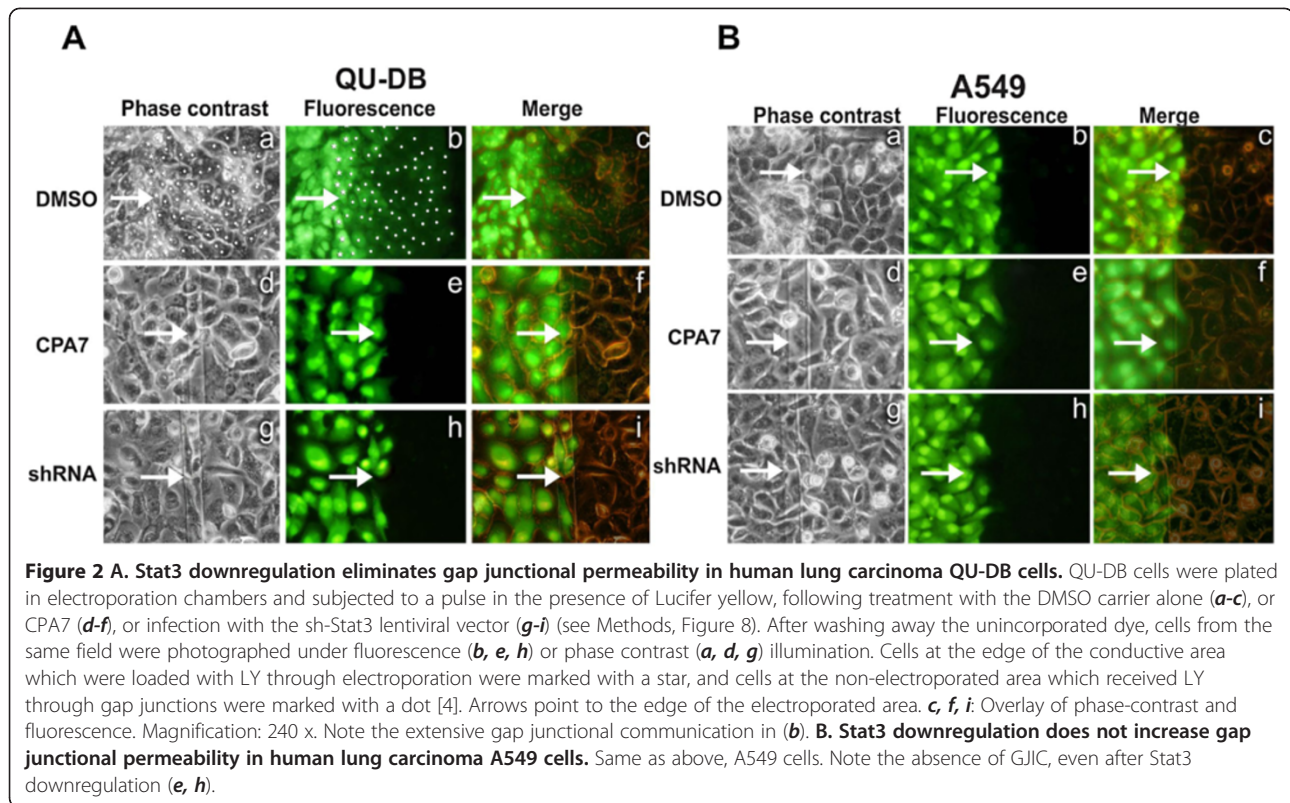
A. Cells with extensive junctional communication						
Cell line	Treatment ^a	Src ^b (%)	Stat3 ^b (%)		GJIC ^c	
			50%	100+3d	90%	100+3d
E10	-	6±1	9±3	26±9	1.5±0.5	6.0±1
"	DMSO	6±1	9±3	30±8	-	6.0±1
"	CPA7	5±1	2±1.1	3±1	-	0.2±0.1
"	sh-Stat3	N/A	6±1.1	8±2	-	1.0±0.2
QU-DB	-	7±1	10±2	20±4	1±0.2	6.3±1
"	DMSO	7±1	10±2	22±3	-	6.3±1
"	CPA7	5±1	2±0.5	2±0.5	-	0.2±0.1
"	sh-Stat3	N/A	5±3	8±2	-	0.8±0.2
SK-LuCi6	-	5±1	8±2	21±4	1.8±0.2	6.5±1
"	DMSO	5±1	8±2	20±5	-	6.5±1
"	CPA7	5±1.2	2.8±1.25	3±1	-	0.3±0.1
"	sh-Stat3	N/A	6±2	4±0.5	-	1±0.2
"	Jak inhib.1	5.2±0.3	4.2±1.1	5±0.5	-	0.5±0.2
"	Stat3C	5.1±1	22±9	97±10	-	8±1
B. Cells expressing activated Src						
A549	-	95±11	93±12	320±32	0.1 ±0.1	0.3 ±0.1
"	DMSO	95±11	93±12	320±32	-	0.3 ±0.1
"	CPA7	93±10	8±1	12±2	-	0.1 ±0.1
"	sh-Stat3	N/A	12±3	11±4	-	0.1±0.1
E10-Src	DMSO	98±12	98±15	350±28	-	0.4 ±0.2
"	CPA7	95±11	5±1	15±4	-	0.1 ±0.1
"	sh-Stat3	N/A	9±3	20±3	-	0.1±0.1
SK-LuCi6-Src	DMSO	100±12	100±12	420±33	-	0.2±0.1
"	CPA7	98±10	3±1	9±1	-	0.1 ±0.1
"	sh-Stat3	N/A	11±3	17±3	-	0.3±0.1
"	sh-Stat3+Das.	4±1	6±1.1	14±3	-	0.2±0.1
SK-Lu-1	DMSO	85±5	90±11	311±23	-	1±0.2
"	CPA7	82±4	6±1	8±3	-	0.1 ±0.1
CALU-1	DMSO	96±9	100±10	290±12	-	0.1 ±0.1
"	CPA7	90±12	8±2	6±1	-	0.1 ±0.1
SW-900	DMSO	100±13	100±12	405±21	-	0.1 ±0.1
"	CPA7	96±11	12±1	11±2	-	0.1 ±0.1
CALU-6	DMSO	95±11	93±10	300±18	-	0.1 ±0.1
"	CPA7	93±10	10±1	15±5	-	0.1 ±0.1

^aFor Stat3 inhibition, cells were treated with 50 μM CPA7, or the DMSO carrier for 24 hrs, or infected with a lentivirus vector expressing a Stat3-specific, shRNA [37]. For Stat3 upregulation, cells were infected with a retroviral vector containing Stat3C. Jak inhibitor-1 was used at 5 μM [16].

^b Stat3-tyr705 or Src-ptyr418 levels were measured by Western blotting. Numbers represent relative values obtained by quantitation analysis. Averages of at least three experiments ±SEM are shown. For Stat3, data from cells grown to 50% confluence or 3 days after confluence are presented [15], with the average of the values for DMSO-treated, Src-transduced, SK-LuCi6-Src cells grown to 50% confluence taken as 100%. The transcriptional activity values obtained paralleled the Stat3-705 phosphorylation levels indicated (Figure 4C and D, see Methods).

^cGJIC was assessed by *in situ* electroporation at the indicated confluences (see Methods, Figure 8). Quantitation was achieved by dividing the number of cells into which the dye had transferred through gap junctions (denoted by dots, Figure 1B and 2A), by the number of cells at the edge of the electroporated area (denoted by stars). Numbers are averages ±SEM of at least three experiments, where transfer from more than 200 cells was examined.

N/A: Not applicable.



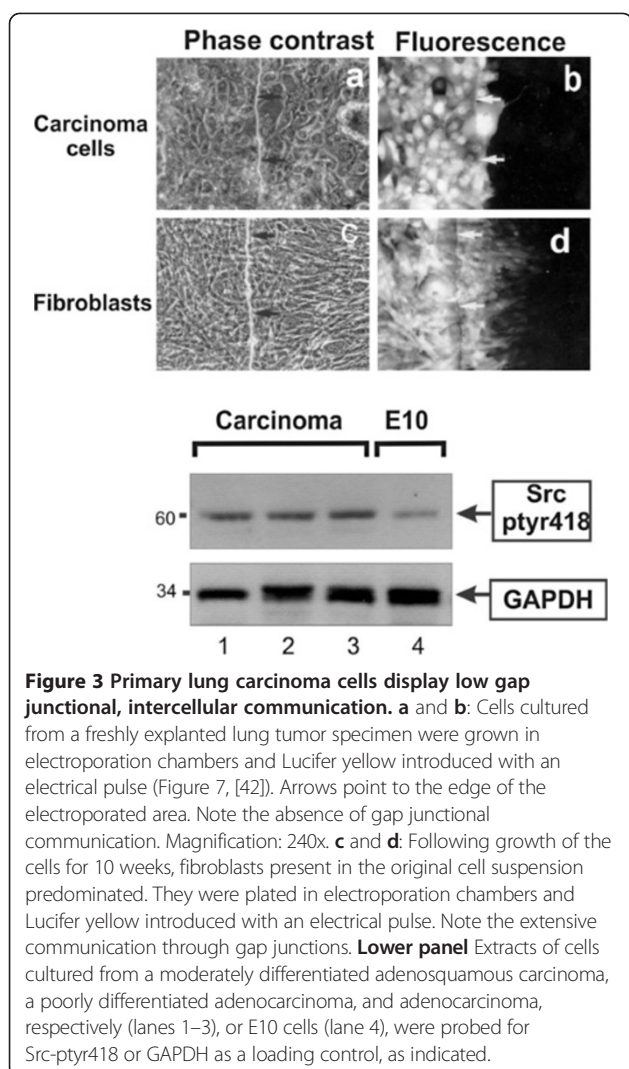
a-c and Table 1,B). In addition, Src expression in SK-LuCi6 or E10 cells eliminated junctional permeability (E10-Src and SK-LuCi6-Src, Table 1, B), in agreement with the known Src effect of GJIC suppression. Conversely, the two lines with low Src-ptyr418 levels (QU-DB and SK-LuCi6), had high GJIC, especially at high densities (Figure 2A and Table 1, A). Primary cells from the three tumor specimens were found to have higher Src activity than the E10, consistent with previous results from biopsy tissues (Figure 3, bottom panel). Taken together, these data point to an inverse relationship between Src activity levels and GJIC in these NSCLC lines.

Stat3 is a positive regulator of GJIC in NSCLC lines

Stat3 is a prominent effector of the non-receptor tyrosine kinase Src [24]. However, Stat3 can be activated by cytokine and membrane tyrosine kinase receptors, which can act in a Src-independent manner [11]. Therefore, to assess the specific contribution of Src to Stat3 activation in the lung cancer lines, we at first examined the correlation between Src-ptyr418 and Stat3-ptyr705 levels. As shown before for a number of cell types (reviewed in [17]), high cell density caused an increase in Stat3-ptyr705 levels in all lines (e.g. Figure 5A, lanes 1-4 and 5-8), therefore Stat3-ptyr705 levels were assessed at a confluence of 50% for this experiment (see Methods). The results showed elevated Stat3-ptyr705 levels in the five lines with

high Src-ptyr418 at all cell densities, comparable to SK-LuCi6-Src cells (e.g. A549 vs QU-DB, Figure 5A, lanes 1-4 vs 5-8 and Figure 5B and Table 1,B). At the same time, QU-DB and SK-LuCi6 cells had low levels of both Src-ptyr418 and Stat3-ptyr705 (Figure 5B). The above data point to a correlation between Src and Stat3 activity levels in the NSCLC lines. We next examined the effect of Src inhibition upon Stat3-ptyr705 in the lines found to have high Src-ptyr418. The results showed that treatment with the Src inhibitor Dasatinib caused a dramatic reduction in Stat3-ptyr705 (e.g. line A549, Figure 5C, and Additional file 1: Additional data, Table Add-1). Similar results were obtained with the PD180970 and SU6656 Src inhibitors (see Methods). These findings indicate that Src may, in fact, be an important Stat3 activator in these cells.

The effect of Stat3 inhibition upon GJIC in the 5 lines with high Src activity was examined next. As shown in Figure 3C and D treatment with the Stat3 inhibitor, CPA7 for 15 hrs [25], or knockdown with a Stat3-specific, shRNA, essentially eliminated Stat3, tyr705 phosphorylation and activity in A549 cells. However, CPA7 treatment (Figure 2B, d-f), or Stat3 knockdown (Figure 2B, g-i) did not increase junctional permeability in A549 cells. Similar results were obtained with SK-Lu1, CALU-1, SW-900 and CALU-6 lines (Table 1,B). The above data taken together indicate that the high Stat3 activity, which could be, at least in part, due to high Src activity in these lines, cannot be responsible



for the lack of junctional communication in the lung carcinoma lines examined.

Since the lung cancer lines might express other oncogenes besides Src, we examined the role of Stat3 in the Src-triggered GJIC suppression specifically, using

Table 2 GJIC in primary lung carcinoma cells^a

	Cells ^b	GJIC ^a
Adenosquamous carcinoma, moderately differentiated	carcinoma cells	0.1 ± 0.1
	fibroblasts	5.8 ± 1.2
Adenocarcinoma, poorly differentiated	carcinoma cells	0.1 ± 0.1
Adenocarcinoma	carcinoma cells	0.1 ± 0.1

^aImmediately after surgery, cells were placed in culture and GJIC examined (see Methods, Figure 7). After 8-10 weeks in culture, most of the tumor cells had died while the fibroblasts present in the initial suspension predominated. These cells did not express cytokeratins, contrary to tumor cells [18]. The fibroblasts shown were derived from the moderately differentiated adenocarcinoma tumor above (Figure 3, c-d). GJIC was examined as in Table 1, at 3 days after confluence.

the Src-transduced, SK-LuCi6-*Src* line. As expected, Src expression disrupted gap junctional permeability. Interestingly, subsequent Stat3 inhibition with CPA7 or shRNA did not restore GJIC (Table 1,B). Taken together, the above findings indicate that Stat3 cannot be part of a pathway leading to Src-induced, gap junction closure in SK-LuCi6-*Src* cells.

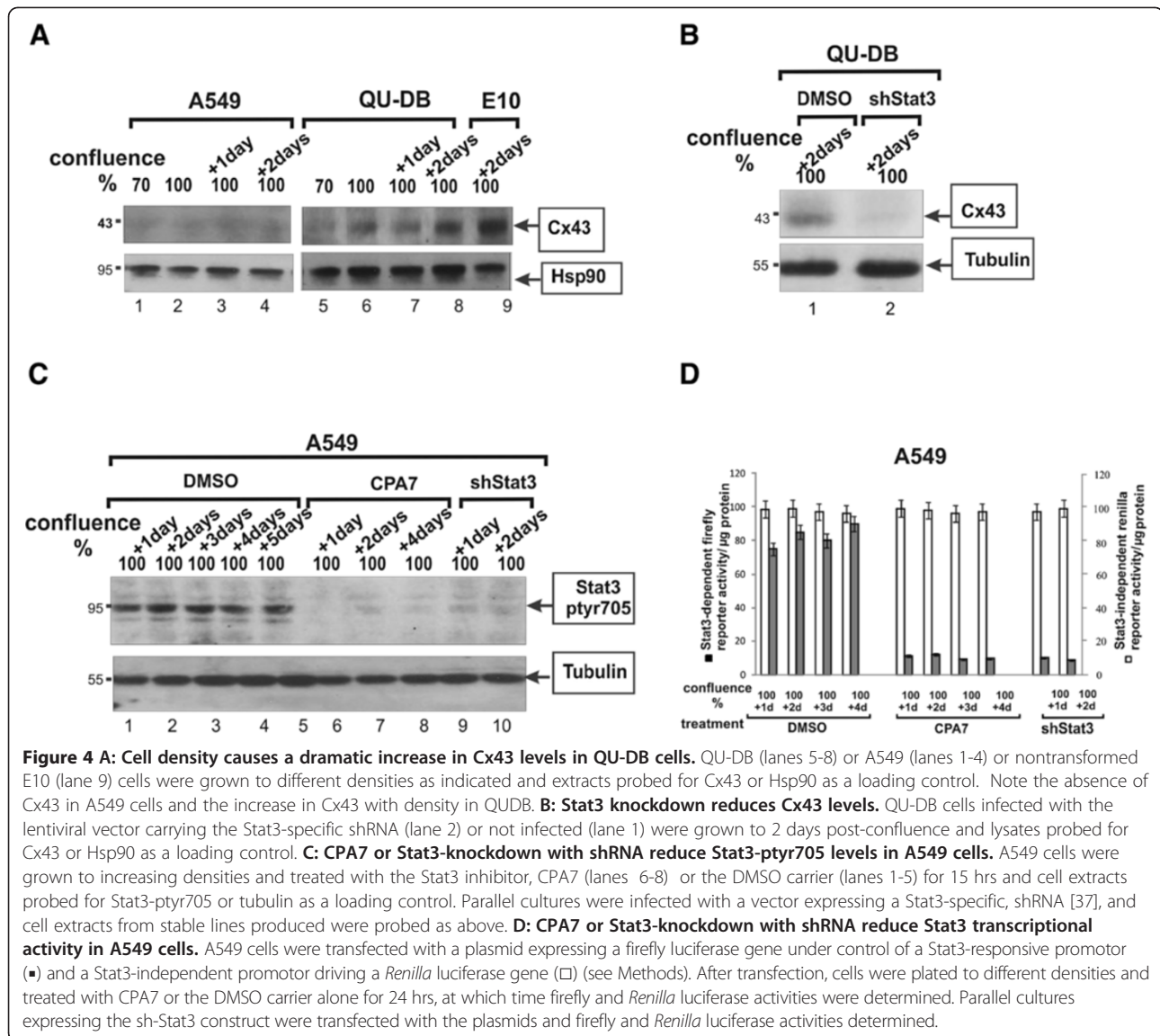
We then examined the possibility that Stat3 might play a *positive* role in the maintenance of gap junctional permeability, by assessing the effect of Stat3 inhibition upon GJIC levels in QU-DB cells which have low Src activity and extensive GJIC. As shown in Figure 2A (d-f), Stat3 downregulation through CPA7 treatment essentially *abolished* GJIC in QU-DB cells. Reduction of Stat3 levels through infection with the sh-Stat3 lentivirus vector gave similar results (Figure 2A, g-i). Similarly, Stat3 downregulation in SK-LuCi6 or E10 cells caused a dramatic decrease in GJIC (Table 1,A). Conversely, expression of the constitutively active form of Stat3, Stat3C [26], increased the already extensive gap junctional communication in SK-LuCi6 cells (Table 1,A).

Examination of Cx43 levels following sh-Stat3 expression revealed a dramatic reduction (Figure 4B), indicating that Stat3 is required for the maintenance of Cx43 protein levels. TUNEL staining revealed that Stat3 inhibition by CPA7 treatment caused an increase in apoptosis in SK-LuCi6 cells (Figure 6A). In addition, CPA7 treatment caused an increase in PARP cleavage in these cells, even at a confluence of 50% (Figure 6B, lane 2). At 3 days post confluence, the time of GJIC examination, PARP cleavage was greater (lane 4), in agreement with previous results indicating that Stat3 inhibition causes apoptosis which is more pronounced in confluent cultures [27]. This finding hints at a link between GJIC reduction and apoptosis induced by Stat3 inhibition.

We next examined whether Stat3 inhibition might also affect Cx43 mRNA levels, through quantitative RT-PCR analysis [28]. The results showed that Stat3 inhibition by CPA7 treatment, or downregulation through shRNA expression brought about a substantial reduction in Cx43 mRNA levels, indicating an effect of Stat3 upon Cx43 gene transcription as well. In any event, taken together, our data reveal that, rather than increasing junctional permeability as might have been expected based on the well documented ability of Stat3 to act as a Src effector, Stat3 inhibition eliminates GJIC, indicating that Stat3 activity is actually *required* for gap junction function in two cultured lung carcinoma lines which display extensive GJIC.

Discussion

Extensive data from our group and others demonstrated that oncogenes such as mT, Src or Ras can suppress gap junctional, intercellular communication [3,6]. Moreover,



it was shown that lower levels of these gene products were sufficient to eliminate gap junction function than the levels necessary for full transformation [4,29], indicating that a decrease in GJIC may be an early event in neoplastic conversion. In this communication we used an improved procedure to examine GJIC in lung cancer lines as well as in primary lung tumor cells. All cell lines had been established from NSCLC tumors which were known to be metastatic [18], except QU-DB, which was derived from a patient that was a long term survivor [30]. Our results reveal that GJIC was low in the majority of cases, except in the QU-DB and SK-LuCi6 lines. Assuming that the establishment process did not bring about an *increase* in GJIC, the existence of extensive GJIC in line SK-LuCi6 which was established from a rapidly metastatic tumor [31] indicates that intercellular

communication does not necessarily inhibit metastasis; other factors may supercede potential growth inhibitory effects of intercellular communication and may be responsible for tumor growth and metastasis.

We next examined the mechanism of GJIC suppression by assessing the role of Src and its effector Stat3. Our results revealed an inverse relationship between Src-tyr418 phosphorylation levels and GJIC in a number of lines. Since Src is known to suppress gap junctional communication in cultured cells such as rodent fibroblasts and epithelial cells, it is tempting to speculate that Src may be responsible, at least in part, for gap junction closure in these lines. However, repeated attempts to reinstate GJIC by reducing Src activity levels through treatment with the Src kinase family-selective, pharmacological inhibitors

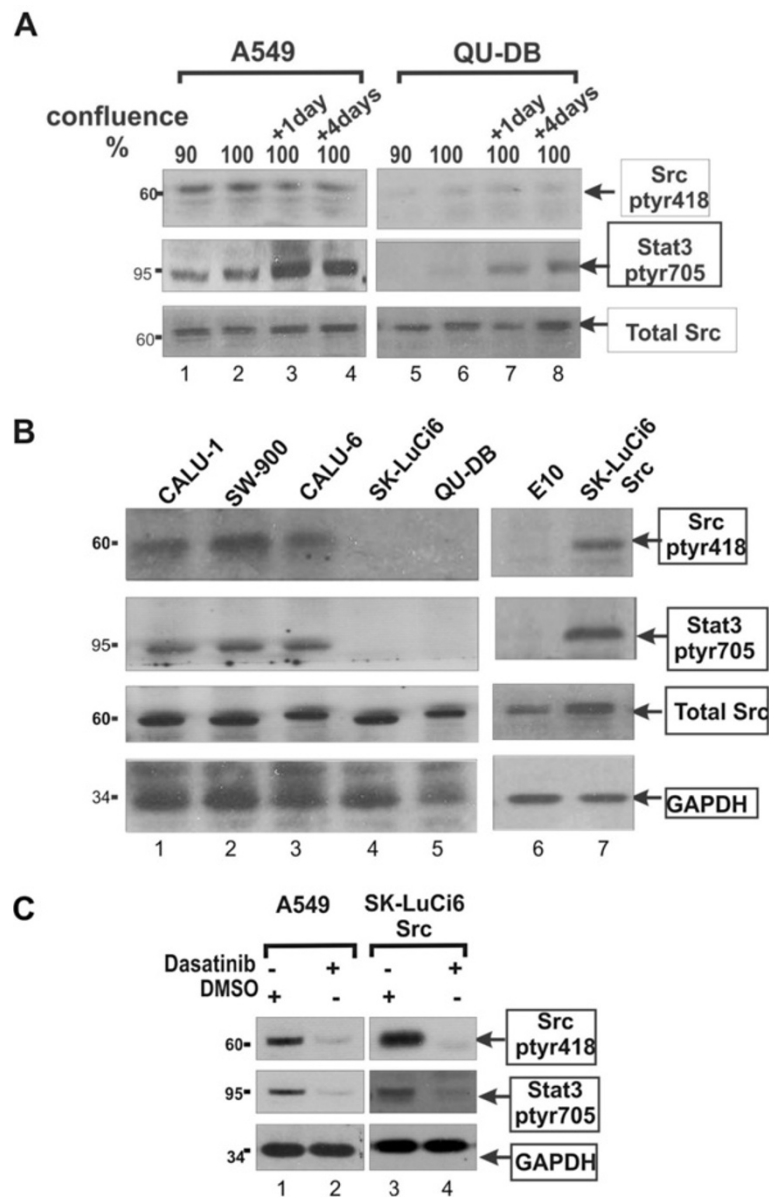
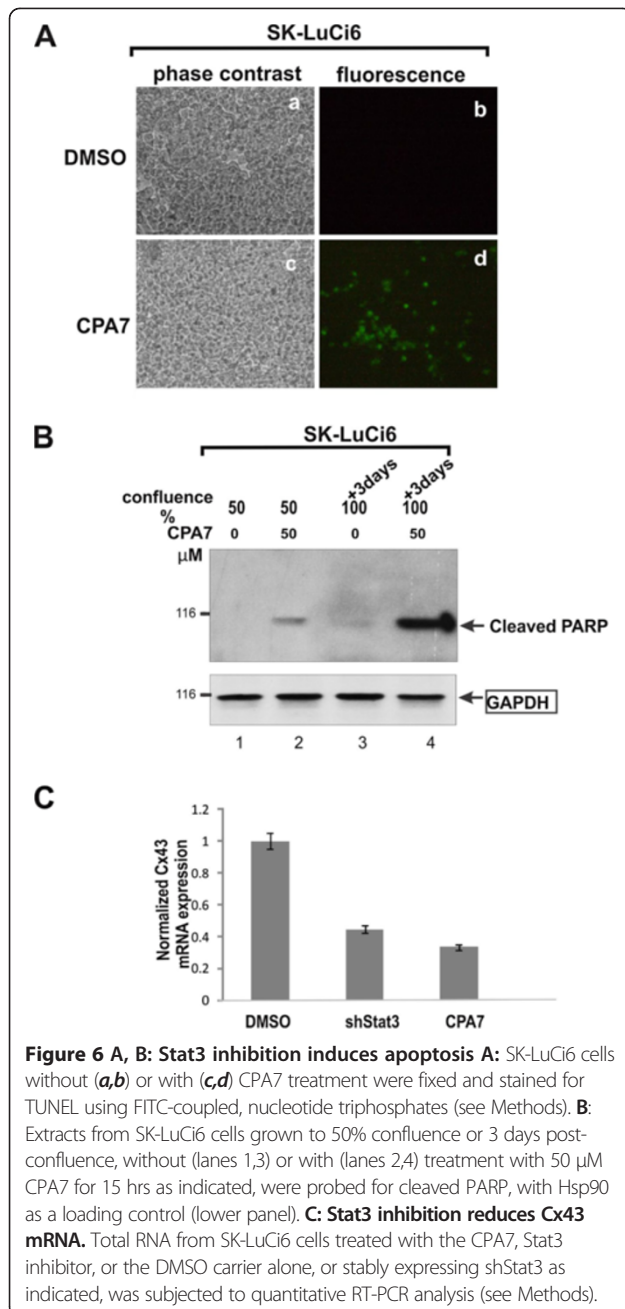


Figure 5 A: A549 cells have high Src-ptyr418 levels. QU-DB (lanes 5-8) or A549 (lanes 1-4) cells were grown to different densities as indicated and extracts probed for Src-ptyr418, Stat3-ptyr705 or total Src. Note the low levels of Src-ptyr418 in QU-DB cells. **B: Src-ptyr418 and Stat3-ptyr705 in NSCLC lines.** The indicated cell lines were grown to 50% confluence and extracts probed for Src-ptyr418, Stat3-ptyr705, total Src or GAPDH as a loading control. **C: Dasatinib reduces Stat3-ptyr705 levels in A459 cells:** A549 or SK-LuCi6-Src cells were grown to subconfluence and treated with the Src-selective inhibitor, Dasatinib (1 μ M) or the DMSO carrier alone and cell extracts probed for Src-ptyr418, Stat3-ptyr705 or GAPDH as a loading control, as indicated.

Dasatinib, PD180970 or SU6656, or infection with Adenoviral vectors expressing a Src dominant-negative mutant or c-Src kinase [15] in A549 cells which have high Src-ptyr418 were unsuccessful (not shown). Possibly other oncoproteins besides Src, or other factors may be important contributors to GJIC suppression in these lines. Alternatively, since low levels of activated Src were previously shown to be sufficient for GJIC suppression in mouse fibroblasts

[3,4], the possibility that the residual Src activity in treated cells might be sufficient to interrupt gap junctional communication cannot be excluded. Dasatinib treatment of SK-LuCi6-Src cells did cause a partial restoration of GJIC, although the high levels of SK-LuCi6 were not attained, possibly due to the high Src activity levels in this line.

We also examined GJIC in freshly explanted, primary cells from 3 NSCLC specimens. Since the



senescence process can reduce GJIC [32], cells were plated in electroporation chambers immediately after surgery at densities of ~80%, so that they would reach confluence within 1–2 days, and GJIC examined every day for up to 10 days. No gap junctional communication was ever detected in any of the preparations, although fibroblasts from the same tissue had extensive GJIC (Figure 3, c-d). Src-418 levels were relatively high in cells from all three tumor specimens, indicating that Src may have played a role in GJIC suppression. However, the possibility that the initiation of the senescence process even a

day after surgery may have affected GJIC cannot be excluded.

Stat3 does not transmit Src signals to gap junction closure

Several signal transducers besides Stat3 are known to be downstream effectors of the Src kinase such as Ras/Raf/Erk, PI3k/Akt, the Crk-associated substrate (Cas) and others [33]. Constitutively active Ras is neoplastically transforming and can suppress GJIC [6,29]. Examination of the mechanism of Src-mediated, GJIC suppression previously indicated that inhibition of Ras in Src-transformed, rat fibroblasts reinstated gap junctional communication [19]. Conversely, mT expression in Ras-deficient cells did not suppress GJIC [34]. These data taken together underline the importance of the Ras pathway in GJIC reduction by activated Src. It was also shown later that Cas is required for the Src-induced, reduction in gap junctional communication [35]. In sharp contrast, our present data with Src-transduced, SK-LuCi6-Src cells demonstrate that Stat3 inhibition does not restore GJIC, indicating that a role of Stat3 in the Src-induced, GJIC suppression in these cells is unlikely, despite the fact that constitutively active Stat3 can act as an oncogene and transform established lines [36].

Stat3 plays a positive role in gap junctional communication

The fact that cell density upregulates Stat3 concomitant with an increase in both Cx43 and GJIC prompted us to explore a potential positive role of Stat3 upon GJIC. Interestingly, Stat3 inhibition in two NSCLC lines which exhibit extensive junctional communication (QU-DB, SK-LuCi6) abolished GJIC, indicating that Stat3 does in fact play a positive role in the maintenance of gap junction function. This conclusion is in agreement with a previous report indicating that Stat3 inhibition eliminated GJIC in nontransformed rat liver epithelial cells as well [37].

Results from a number of labs demonstrated that Stat3 activates a number of anti-apoptotic genes, such as Bcl-xL, Mcl1 and Akt1 [11]. Global induction of apoptosis with etoposide, cycloheximide or puromycin was shown to lead to a loss of cell coupling, probably due to caspase-3-mediated degradation of Cx43, in primary bovine lens epithelial and mouse NIH3T3 fibroblasts [38]. Interestingly, we previously demonstrated that Stat3 inhibition in cells transformed by Src or the Large Tumor antigen of Simian Virus 40 leads to apoptosis [15,39], possibly due to activation of the transcription factor E2F family, potent apoptosis inducers, by these oncogenes. Therefore, apoptosis induced by Stat3 downregulation in cells with high Src may have triggered gap junction closure.

We previously demonstrated that while Stat3 inhibition in sparsely growing, normal mouse fibroblasts causes

growth retardation, at high densities, such as needed for optimal gap junction formation, Stat3 inhibition leads to apoptosis [27]. Therefore, apoptosis induction through a reduction in Stat3 levels or activity could explain the dramatic reduction in Cx43 and GJIC upon Stat3 pharmacological or genetic inhibition, in lines with low Src activity. Still, our results also demonstrate a substantial reduction in Cx43 mRNA levels upon Stat3 inhibition, pointing to a transcriptional effect of Stat3 upon the Cx43 promoter in these NSCLC lines, as previously demonstrated in other cell types [28,40,41].

Conclusions

Our results demonstrate that Stat3 is not transmitting Src signals leading to gap junction closure in the NSCLC cell lines examined. In the contrary, although Stat3 is generally growth promoting and in an activated form it can act as an oncogene, we show for the first time that Stat3 is actually *required* for gap junctional communication both in normal epithelial cells and in certain tumor cell lines that retain GJIC. This novel role of Stat3 in gap junction function may be an important regulatory step in progression of tumours that exploit such a pathway.

Methods

Examination of gap junctional communication

To examine gap junctional communication by *in situ* electroporation, it is important to be able to reliably distinguish cells that were loaded with Lucifer yellow directly by electroporation, from cells that received the dye from neighbouring cells by diffusion through gap junctions. This was achieved using a slide where a 3 mm-wide strip of ITO had been removed by etching with acids, leaving two co-planar electrodes, supported by the same glass slide substrate (Figure 7) [42].

In a further improvement (Figure 8), the coating was removed from the glass surface in ~20 µm wide lines, to define electrode and non-conducting regions. Etching was done using a laser beam, so that the nonconductive glass underneath is exposed. It was important to ensure that only the 800Å coating was removed, without affecting the glass, so that cell growth would be unaffected across the line. This was achieved with a UV laser operating at a 355 nm wavelength using approximately 1 Watt of output power with 60% of the energy delivered to the surface of the glass. The beam was manipulated by mirrors on a pair of galvanometers to produce the desired pattern.

To form the two electrodes, the coating was removed in a straight line in the middle (2). A dam of nonconductive plastic (3) was bonded onto this line, to divert the current upwards, thus creating a sharp transition in electric field intensity between electroporated and non-electroporated sections. To provide areas where the cells are not electroporated, the ITO was also removed in two parallel lines

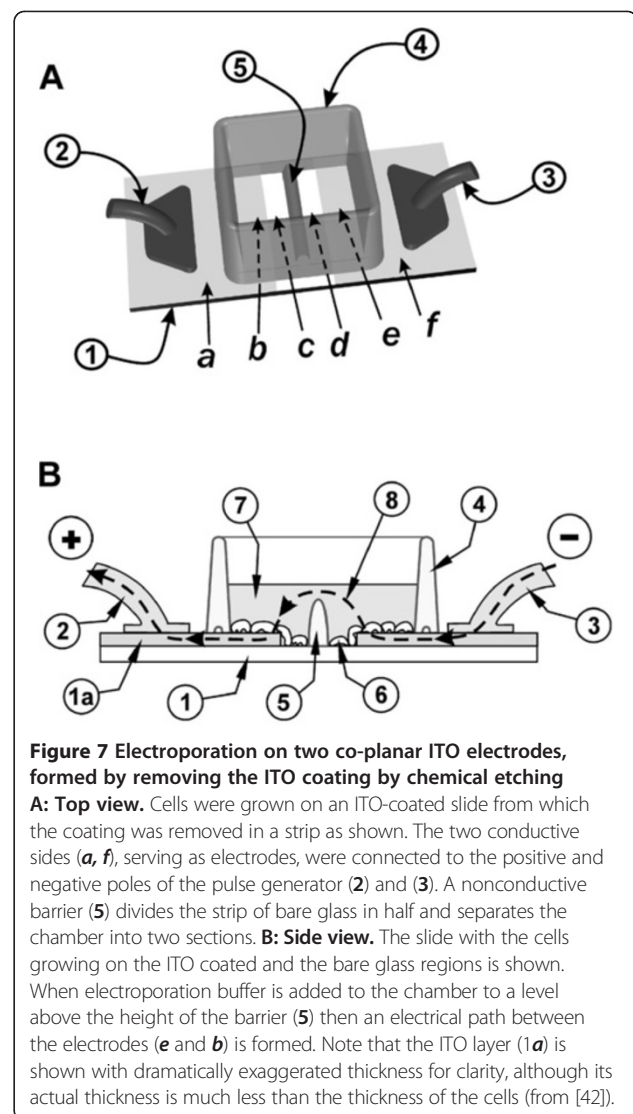
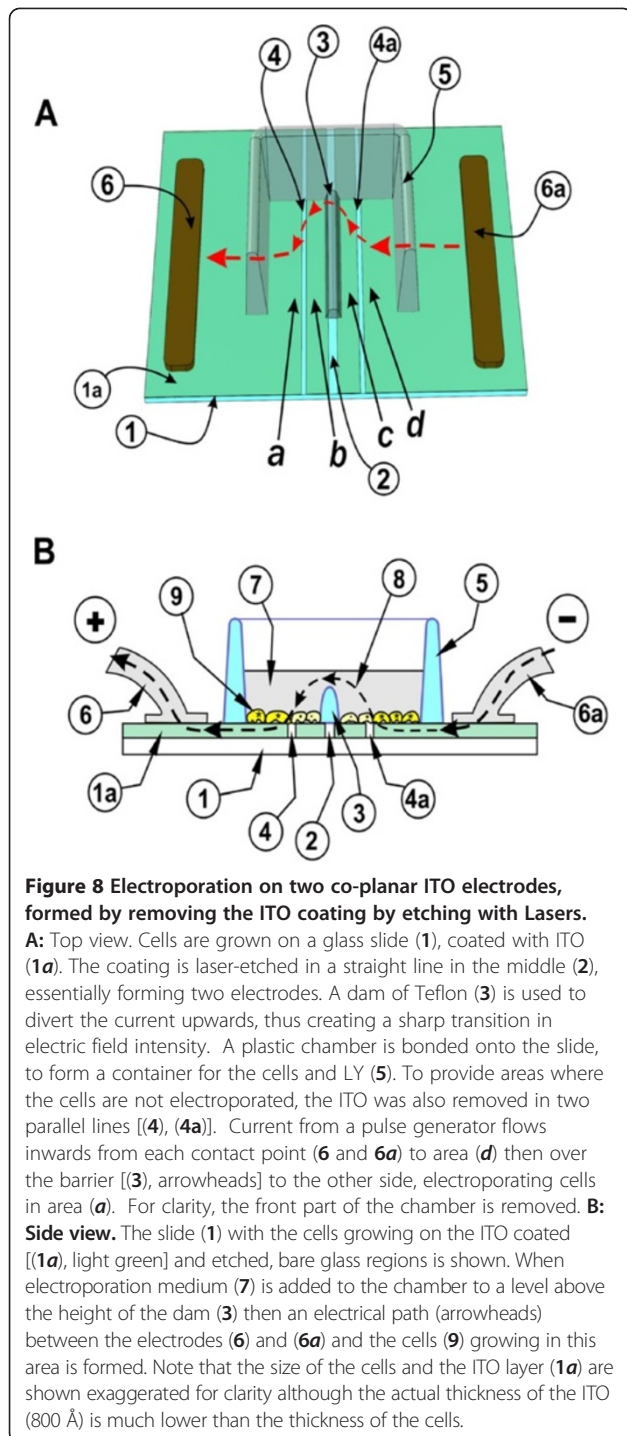


Figure 7 Electroporation on two co-planar ITO electrodes, formed by removing the ITO coating by chemical etching
A: Top view. Cells were grown on an ITO-coated slide from which the coating was removed in a strip as shown. The two conductive sides (a, f), serving as electrodes, were connected to the positive and negative poles of the pulse generator (2) and (3). A nonconductive barrier (5) divides the strip of bare glass in half and separates the chamber into two sections. **B: Side view.** The slide with the cells growing on the ITO coated and the bare glass regions is shown. When electroporation buffer is added to the chamber to a level above the height of the barrier (5) then an electrical path between the electrodes (e and b) is formed. Note that the ITO layer (1a) is shown with dramatically exaggerated thickness for clarity, although its actual thickness is much less than the thickness of the cells (from [42]).

[(4) and (4a)]. A plastic chamber was bonded onto the slide, to form a container for the cells and electroporation solutions (5). Current flows inwards from each contact point (6 and 6a), via a conductive highway under the well (5) electroporating cells in area (d) then over the barrier [(8), arrowheads] to the other side, in area (a). In this configuration, cells which acquired LY by electroporation [growing in (a) and (d)] and cells into which LY traveled through gap junctions [(b) and (c)] both grow on ITO, separated only by a laser-etched line of ~20 µm. Extensive experimentation showed that in this setup the electroporation intensity is uniform across the electroporated area (see Figure 1B and Figure 2).

Cells were plated in the chamber and when they reached the appropriate density (90% confluence, to 5 days post-confluence), the growth medium was replaced with Calcium-free DMEM supplemented with 5 mg/ml Lucifer yellow (7). The slide/chamber was placed into a



holder where electrical contacts were established and a set of electrical pulses delivered to the cells. Extensive experimentation indicated that 10 pulse pairs, each pulse of 18 Volts peak value, 100 μ s length and spaced 0.5 seconds apart, with one of each pair having a polarity opposite to that of its partner gave optimal results. Following a 5 min incubation at 37°C, the unincorporated dye was washed away with Calcium-free DMEM supplemented with

10% dialysed fetal calf serum and cells observed and photographed under fluorescence and phase contrast illumination. Communication is expressed as the number of cells into which the dye has transferred per cell loaded with the dye by electroporation at the edge of the electroporated area. All experiments were conducted at least three times, with at least 5 slides each time, and the results are presented as average GJIC \pm SEM where the transfer from at least 200 cells is assessed.

The equipment (*In situ* Porator) was supplied by Cell Projects Ltd UK.

Cell lines, culture techniques and Stat3 activity measurement

All cells were grown in DMEM with 10% fetal calf serum. Extra care was taken to ensure that cell seeding was uniform, by passing cells at subconfluence, when cell to cell adhesion was low. Confluence was estimated visually and quantitated by imaging analysis of live cells under phase contrast [14]. To ensure that the growth medium was not depleted of nutrients, it was changed every day.

Cells were cultured from surgically explanted tumors as previously described [18].

Stat3C and activated Src were expressed in SK-LuCi-6 cells through infection with the culture supernatant from a Phoenix amphotropic packaging line transfected with a pBabe-puro-Stat3C plasmid [26]. shStat3 was expressed by retroviral vector infection as described [16].

Stat3 transcriptional activity was measured as described, by transient transfection of the pLucTKS3 construct [24]. As a control, cells were co-transfected with the reporter pRLSRE, which contains two copies of the serum response element (SRE) of the c-fos promoter, subcloned into the *Renilla* luciferase reporter, pRL-null (Promega) [15]. Following transfection, cells were plated to different densities and luciferase activity determined.

For Stat3 immunostaining, SK-LuCi6 cells were fixed with 4% paraformaldehyde, permeabilized in 0.2% Triton-X100 and probed with a Stat3 antibody (Cell Signalling, #9132 diluted at 1:100) followed by AlexaFluor-coupled, goat anti-rabbit IgG (Invitrogen #A11008, used at 1:400).

Inhibitors

Stat3 was inactivated using two approaches: (1). Treatment with 50 μ M CPA7 [PtCl₃(NO₂)(NH₃)₂] [25] overnight, or (2). Expression of shRNA, delivered with a lentivirus vector as described [37]. Jak inhibitor-1 was from EMD Biosciences (5 μ M [16]).

Src was inactivated using 3 pharmacological inhibitors: Dasatinib (0.5 or 1 μ M, up to 72h), PD180970 (0.2 μ M with redosing every 12h for a total of 24h), or SU6656 (5 μ M for 24h) [15].

Western blotting

It was conducted on proteins extracted from cell pellets [43], using antibodies to Cx43 (Cell Signalling, #3512, used at a 1:500 dilution), Stat3-ptyr705 (Cell Signalling, #9131, 1:1,000), Src-ptyr418 (Invitrogen, #44-660G, 1:1,000) or total Src (rabbit monoclonal 36D10, Cell Signalling, #2109, 1:1,000), followed by secondary antibodies and ECL reagents (Biosource). Alpha-Tubulin (Cell Signalling #2125, 1:5,000), GAPDH (BD Transduction, #14C10, 1:5,000) or Hsp90 (Assay designs, #SPA-830, 1:5,000) served as loading controls.

qRT-PCR

SK-LuCi6 cells were treated with CPA7 for 15 hrs. The RNeasy Mini Kit (Qiagen, Hilden Germany, cat. #74104) was used for the purification of RNA. cDNA synthesis from 1 µg of total RNA was performed using the iScript cDNA synthesis kit (Bio-RAD Laboratories, Hercules, CA). qRT-PCR was performed using iQ SYBR Green Supermix (Bio-RAD Laboratories, Hercules, CA) and 20 µM primer. Primer sequences were [28]:

Cx43: Forward: 5'-GCCTGAACTTGCCTTTTCAT-3',
Reverse: 5'-CTCCAGTCACCCATGTTGC-3', [28] to generate a product of 500 bp [40].

As internal reference genes we used GAPDH: (forward: 5'-AATGCATCCTGCACCACCAA-3', Reverse: 5'-GTAGCCATATTCATTGTCATA-3') [40] and 18S RNA [16]. mRNA from SK-LuCi6 cells where Stat3 was downregulated with sh-Stat3 was analysed in a similar manner. Results from 3 independent experiments, each conducted in triplicate were averaged out and SEM calculated.

Additional file

Additional file 1: Additional data.

Abbreviations

Cx43: Connexin-43; GJIC: Gap junctional, intercellular communication; NSCLC: Non-small cell lung cancer.

Competing interests

The corresponding author received a royalty from PARTEQ, the intellectual property arm of Queen's University, for a patent where she is a co-inventor of the apparatus used in this work. PARTEQ had no involvement whatsoever in the content of this paper.

Authors' contributions

MG did the bulk of the benchwork. SG and RA conducted some of the initial experiments. Aaron Trotman-Grant did the Stat3C experiments. ET obtained the results on the primary lung carcinoma cells. LR conceived of the study, designed and coordinated it and drafted the manuscript. All authors read and approve of the manuscript.

Authors' information

MG is a postdoctoral fellow, funded by the US Army breast cancer program. RA and ET were graduate students. ATG is currently doing a project. LR is a professor at Queen's University, Kingston, Canada.

Acknowledgements

We would like to thank Dr. Mike Baird and Shallyn Littlefield for CPA7 synthesis, Dr. Jackie Bromberg for the pBabe-puro-Stat3C plasmid, Kevin Firth, P.Eng., of Ask Sciences products, Kingston, Ontario for engineering design, Dr. Christopher Nicol for advice and help with qRT-PCR assays, Drs Bruce Banfield and Renee Finnen for fluorescent microscopy and Dr. Andrew Craig for the NL20 line. We are grateful to Dr. Elissavet Kardami (University of Manitoba, Canada) for a generous gift of connexin-43 antibody. The financial assistance of the Canadian Institutes of Health Research (CIHR), the Canadian Breast Cancer Foundation (Ontario Chapter), the Natural Sciences and Engineering Research Council of Canada (NSERC), the Canadian Breast Cancer Research Alliance, the Ontario Centers of Excellence, the Breast Cancer Action Kingston and the Clare Nelson bequest fund through grants to LR is gratefully acknowledged. RA was supported by a Canada Graduate Scholarships Doctoral award from CIHR, the Ontario Women's Health Scholars Award from the Ontario Council on Graduate Studies and a Queen's University Graduate Award. ET was the recipient of an OGSST award. SG was the recipient of an NSERC summer studentship. MG was supported by a postdoctoral fellowship from the US army Breast Cancer Program, the Ministry of Research and Innovation of the Province of Ontario and the Advisory Research Committee of Queen's University.

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Received: 12 April 2012 Accepted: 13 December 2012

Published: 18 December 2012

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doi:10.1186/1471-2407-12-605

Cite this article as: Geletu et al.: Stat3 is a positive regulator of gap junctional intercellular communication in cultured, human lung carcinoma cells. *BMC Cancer* 2012 **12**:605.

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