Connexin expression and gap-junctional intercellular communication in ES cells and iPS cells

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INTRODUCTION

Gap junctions are cell–cell communicating junctions that consist of multimeric proteins called connexins and mediate the exchange of low-molecular-weight metabolites and ions between contacting cells (Oyamada et al., 2013). Gap-junctional intercellular communication (GJIC) has long been hypothesized to play a crucial role in the maintenance of homeostasis, morphogenesis, cell differentiation, and growth control in multicellular organisms. Discoveries of human genetic disorders due to mutations in connexin genes and experimental data on connexin knockout mice provide direct evidence that gap junctional intercellular communication is essential for tissue functions and organ development and that its dysfunction causes diseases. Connexin-related signaling also involves extracellular signaling (hemichannels) and non-channel intracellular signaling.

GJIC during embryonal development has been demonstrated by using microelectrode impalements to monitor the cell-to-cell movement of ions (ionic coupling) and by microinjection of small-molecular-weight fluorescent dyes such as Lucifer yellow into a single cell and observation of the subsequent dye spread into the surrounding cells (dye coupling) (Lo and Gilula, 1979; Kalimi and Lo, 1988, 1989). It has been revealed that in many instances, GJIC is established within the first few cleavages and results in the entire embryo becoming interconnected as a syncytium. As development progresses, however, dye coupling delineates boundaries defining restrictions in GJIC that effectively segregate the developing embryo or tissue into a number of “communication compartment” domains. Thus, cells lying within a communication compartment are well coupled, exhibiting both ionic and dye coupling, whereas there is little or no coupling between cells situated across a compartment border. Such restriction of GJIC and the segregation of cells into communication compartment domains are almost always associated with embryogenesis and development.

Pluripotent stem cells, i.e., embryonic stem (ES) and induced pluripotent stem (iPS) cells, can indefinitely proliferate without commitment and differentiate into all cell lineages. ES cells are derived from the inner cell mass of the preimplantation blastocyst, whereas iPS cells are generated from somatic cells by overexpression of a few transcription factors. Many studies have demonstrated that mouse and human iPS cells are highly similar but not identical to their respective ES cell counterparts. The potential to generate basically any differentiated cell types from these cells offers the possibility to establish new models of mammalian development and to create new sources of cells for regenerative medicine. ES cells and iPS cells also provide useful models to study connexin expression and gap-junctional intercellular communication (GJIC) during cell differentiation and reprogramming. In 1996, we reported connexin expression and GJIC in mouse ES cells. Because a substantial number of papers on these subjects have been published since our report, this Mini Review summarizes currently available data on connexin expression and GJIC in ES cells and iPS cells during undifferentiated state, differentiation, and reprogramming.

Keywords: connexins, gap-junctional intercellular communication, ES cells, iPS cells, differentiation, reprogramming, pluripotency

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In 1996, we first reported the expression of connexin genes and GJIC during in vitro cardiomyocyte differentiation of mouse ES cells (Oyamada et al., 1996). Because a substantial number of papers on these subjects have been published since our first report, this Mini Review summarizes currently available data on connexin expression and GJIC in ES cells and iPS cells during undifferentiated state, differentiation, and reprogramming.

**QUESTIONS ABOUT CONNEXIN EXPRESSION AND GAP-JUNCTIONAL INTERCELLULAR COMMUNICATION IN ES/iPS CELLS**

Main questions about connexin expression and GJIC in ES/iPS cells that have been addressed thus far can be summarized as below:

1. What kinds of connexins are expressed in undifferentiated ES/iPS cells?
2. To what extent do undifferentiated ES/iPS cells communicate with each other via gap junctions?
3. What changes in connexin expression and GJIC occur during differentiation of ES/iPS cells?
4. What roles do connexin expression and/or GJIC play in maintenance of pluripotency in ES/iPS cells?
5. What changes in connexin expression and GJIC occur during induction of pluripotency in somatic cells (reprogramming)?
6. What roles do connexin expression and/or GJIC play in reprogramming?

**CURRENTLY AVAILABLE DATA ON CONNEXIN EXPRESSION AND GAP-JUNCTIONAL COMMUNICATION IN ES CELLS**

Table 1 summarizes results of published papers concerning connexin expression and GJIC in ES cells.

**CONNEXIN EXPRESSION AND GAP-JUNCTIONAL INTERCELLULAR COMMUNICATION IN iPS CELLS**

Table 2 summarizes results of published papers concerning connexin expression and GJIC in iPS cells.

Using human iPS cells, Sharovskaya et al. (2012) reported that GJIC is re-established during reprogramming to pluripotency: GJIC in incompletely reprogrammed cells was markedly decreased compared with that in the parental somatic cells, but GJIC in completely reprogrammed cells exceeded that in the parental somatic cells and was comparable to that in human ES cells. They drew an analogy between dramatic reduction of GJIC in incompletely reprogrammed iPS cells, cells lacked characteristics of parental HUVEC Cx37 expression, whereas Cx43 expression increased three- to five-fold.

Ke et al. (2013) demonstrated that Cx43 is specifically and highly enriched in undifferentiated human iPS cell lines during and after the reprogramming process. They also showed that iPS cells display functional GJIC and that Cx43 expression is gradually upregulated (~4.5-fold increase) during the reprogramming process. They observed that the Cx43 protein level increased gradually along with the expression of the pluripotency marker NANOG. Because Cx43 has been identified as a downstream target of the key pluripotency transcription factors OCT4, SOX2 and NANOG (Boyer et al., 2005), Cx43 expression might be upregulated by the key factors during reprogramming. They also found that the ectopic expression of Cx43 enhances the reprogramming efficiency (~3-fold increase), whereas the knockdown of endogenous Cx43 expression by RNAi reduces the efficiency, possibly by affecting the MET process, as reported by changes in E-cadherin expression. In addition, they showed that pharmacological GJIC inhibitors, CBX, 18-a-GA and the Cx43 mimetic peptide GAP27, did not affect the efficiency of iPS cell generation, suggesting that the effect of Cx43 on the efficiency of iPS cell generation may be attributed to the Cx43 protein itself but not to the function of GJIC, i.e., through a GJIC-independent pathway.

Taken together, these results suggest that Cx43 may represent a pluripotency marker of iPS cells and may play an important role in the reprogramming process.

Lundy et al. (2013) recently have developed a cell culture protocol capable of generating and maintaining highly purified human ES cell- and iPS cell-derived cardiomyocytes for several months in vitro. They have shown that these human ES cell- and iPS cell-derived cardiomyocytes are capable of maturing to a phenotype that more closely resembles adult cardiomyocytes in both structure and function. A robust induction of key cardiac structural markers including Cx43 has been demonstrated in late-stage ES cell- and iPS cell-derived cardiomyocytes. These findings suggest that ES cell- and iPS cell-derived cardiomyocytes are capable of slowly maturing to more closely resemble the phenotype of adult cardiomyocytes and may eventually possess the potential to regenerate the lost myocardium with robust de novo force-producing tissue.
| ES cell lines | Connexin expression in undifferentiated cells | GJIC in undifferentiated cells | Differentiation from ES cells | Connexin expression during differentiation | GJIC during differentiation | Methods used to determine the final phenotype of differentiated cells | References |
|---------------|---------------------------------------------|-----------------------------|-------------------------------|------------------------------------------|---------------------------|------------------------------------------------|-------------|
| Mouse ES cells (J1) | Cx43\(^1\), Cx45\(^1\) Not detected: Cx40\(^1\) | Present\(^3\) | Cardiomyocytes | Cx40\(^1\), Cx43\(^{1,2}\), Cx45\(^1\) | Present\(^3\). Restricted to neighboring beating cells | Contraction, Ca\(^{2+}\)-imaging, cardiac-specific gene expression | Oyamada et al., 1996 |
| Mouse ES cells (D3) | Cx43\(^2\) | Present\(^3\) | Cardiomyocytes | Cx43\(^2\) | Contraction, EM | Westfall et al., 1997 |
| Mouse Cx43\(^{-/-}\) ES cells (R1) | Cx45\(^1\), No compensatory upregulation of Cx40\(^1\) and Cx45\(^1\) | Very low GJIC\(^3\) | Cardiomyocytes. Cx43 knockout did not significantly change either the time course, frequency of cardiomyocytic differentiation, or expression of cardiac-specific genes | Upregulation of Cx40\(^1\) | Very low GJIC\(^3\) | Contraction, cardiac-specific gene expression | Oyamada et al., 2000 |
| Mouse ES cells (D3) | Cx43\(^{1,2}\) | Cardiomyocytes | Increases in Cx40\(^2\) and Cx43\(^2\) during cardiac differentiation | Contraction, cardiac-specific gene expression, electrophysiology | | Van Kempen et al., 2003 |
| Mouse ES cells (HM1) | | Cardiomyocytes | Upregulation of Cx40\(^2\) at a peak around day 3 (hanging drop period) + 14 | Cardiac-specific gene expression, ANEPPS fluorescence, electrophysiology | | Fijnvandraat et al., 2003 |
| Mouse ES cells (CCE) | Cx43\(^{1,2}\), Cx45\(^{1,2}\) No or very low expression: Cx37\(^1\), Cx40\(^1\) | Cardiomyocytes (irregular contractions in Cx45\(^{-/-}\) cells) | Cx37\(^1\), Cx40\(^1\), Cx43\(^{1,2}\), Cx45\(^1\) | Contraction, Ca\(^{2+}\)-imaging, multielectrode array, cardiac-specific gene expression, EM | | Egashira et al., 2004 |
| Human ES cells (H1, H2, H9, H14) | Cx43\(^{1,2}\), Cx45\(^1\) | Present\(^3\) | | | | Carpenter et al., 2004 |
| Human ES cells (GE01, GE09, BG01, BG02, TE06) | Cx43\(^1\), Cx45\(^1\) | | | | | Bhattacharya et al., 2004 |

(Continued)
Table 1 | Continued

| ES cell lines | Connexin expression in undifferentiated cells | GJIC in undifferentiated cells | Differentiation from ES cells | Connexin expression during differentiation | GJIC during differentiation | Methods used to determine the final phenotype of differentiated cells | References |
|---------------|---------------------------------------------|-------------------------------|-------------------------------|------------------------------------------|----------------------------|--------------------------------------------------------------------------------|------------|
| Human ES cells (HES-3, HES-4) | Cx43<sup>1</sup> (As one of the candidate human ES marker genes) | Present<sup>3</sup> | | | | | Richards et al., 2004 |
| Human ES cells (HES-3, HES-4) | Cx43<sup>1</sup>, Cx45<sup>1,2</sup> | Present<sup>3</sup> | | | | | Wong et al., 2004, 2006 |
| Mouse ES cells (Royan B1) | Cardiomyocytes | Presence of gap junctions in 21-day cardiomyocytes by EM | Cardiac-specific gene expression, EM, pharmacological reagents | Baharvand et al., 2005 |
| Mouse ES cells (DBA/1LacJ) | Cardiomyocytes | Cx43<sup>1</sup>, Cx45<sup>1</sup> | Contraction, Ca<sup>2+</sup>-imaging, cardiac-specific gene expression, EM | Chaudhary et al., 2006 |
| Human ES cells (BG01, H1) | Cx43<sup>1</sup>, Cx40<sup>1,2</sup>, Cx45<sup>1,2</sup>, Cx26<sup>1</sup>, Cx30<sup>1</sup>, Cx30.2<sup>1</sup>, Cx30.3<sup>1</sup>, Cx31<sup>1</sup>, Cx31.1<sup>1</sup>, Cx31.9<sup>1</sup>, Cx32<sup>1</sup>, Cx36<sup>1</sup>, Cx37<sup>1</sup>, Cx46<sup>1</sup>, Cx47<sup>1</sup>, Cx53<sup>1</sup>, Cx60<sup>1</sup>, Cx62<sup>1</sup> Not detected: Cx40.1<sup>1</sup>, Cx50<sup>1</sup> | Presence of GJIC<sup>3,5</sup> and hemichannels Extremely rare dye coupling between ES cells and feeder cells | Contraction, electrophysiology, cardiac-specific gene expression, EM, pharmacological reagents | Huetter et al., 2006 |
| Cynomolgus monkey ES cells (CMK-6) | Cx43<sup>1</sup> | Embryoid bodies (EBs) | Suppression of Cx43 mRNA expression during EB differentiation | Yamamoto et al., 2007 |
| Human ES cells (HES2, HES-3, ENY) | Cx43<sup>2</sup> | Presence of GJIC mediated transport of shRNA | | | | | Wolvetang et al., 2007 |
| Mouse ES cells (D3) | Cx43<sup>1,2</sup> | Present<sup>3</sup>, Cx43 silencing inhibited GJC, induced a loss of pluripotent state, and decreased in the proliferation rate | EBs GJC blockers and Cx43-siRNA inhibited the formation of EBs from ES cells | Todorova et al., 2008 |
Table 1 | Continued

| ES cell lines | Connexin expression in undifferentiated cells | GJIC in undifferentiated cells | Differentiation from ES cells | Connexin expression during differentiation | GJIC during differentiation | Methods used to determine the final phenotype of differentiated cells | References |
|---------------|-----------------------------------------------|-------------------------------|-------------------------------|------------------------------------------|----------------------------|---------------------------------------------------------------|------------|
| Mouse ES cells (HM1) | Cx43<sup>1</sup>, Cx43<sup>2</sup>, Cx43<sup>3</sup>, Cx31<sup>1</sup>, Cx31<sup>2</sup>, Cx32<sup>1</sup>, Cx32<sup>2</sup>, Cx37<sup>1</sup>, Cx37<sup>2</sup>, Cx30<sup>3</sup>, Cx30<sup>3</sup>, Cx45<sup>1</sup>, Cx45<sup>2</sup>, Cx46<sup>1</sup>, Cx46<sup>2</sup>, Cx47<sup>1</sup>, Cx50<sup>1</sup>, Cx50<sup>2</sup>, Cx57<sup>1</sup> | Present<sup>3</sup>, <sup>4</sup> Reduction of GJIC by decreased expression of Cx31 or Cx45 via RNA interference in Cx43<sup>−/−</sup> ES cells did not lead to apoptosis | Present<sup>3</sup> | Not characterized | Present but attenuated<sup>3</sup> Restricted to differentiated cells Absence of GJIC between pluripotent and differentiating cells | Worsdorfer et al., 2008 |
| Mouse ES cells (D3): Sox1-promoter-GFP + ES cells and Cx43<sup>−/−</sup> ES cells | Cx43<sup>1</sup> | Neuroectodermal cells Cx43<sup>−/−</sup> ES cells showed a failure of oligodendrocyte development and an amplification of astrocytic cells | Wild-type ES cells showed “two-tailed” Cx43 expression with a maximum at day 7 | Sox1-promoter-GFP, neuronal lineage-specific gene expression | Parekkadan et al., 2008 |
| Human ES cells (hESM01) | Present<sup>3</sup> | Not characterized | Present but attenuated<sup>3</sup> | Sharovskaya et al., 2009 |
| Mouse ES cells (R1) | Cx43<sup>2</sup> expression in cell sheets of mouse ES cell-derived cardiomyocytes | Cardiomyocytes | Cardiac-specific gene expression, multielectrode array | Matsuura et al., 2011 |

<sup>1</sup>mRNA level; <sup>2</sup>Protein level; <sup>3</sup>dye coupling (Lucifer yellow etc.); <sup>4</sup>neurobiotin tracer coupling; <sup>5</sup>electrical coupling; EM, electron microscopy.
Table 2 | Connexin expression and GJIC in iPS cells.

| iPS cell lines | Connexin expression in undifferentiated cells | GJIC in undifferentiated cells | Differentiation from iPS cells | Connexin expression during differentiation | GJIC during differentiation | Methods used to determine the final phenotype of differentiated cells | References |
|----------------|---------------------------------------------|-----------------------------|--------------------------------|------------------------------------------|----------------------------|-----------------------------------------------------------------|------------|
| Mouse iPS cells (O9), Mouse ES cells (E14.1) | Cx43 in iPS cell- and ES cell-derived cardiomyocytes on day 22 | Cardiomyocytes differentiated from iPS and ES cells with the use of a standard EB-based protocol | Cx43 in iPS cell and ES cell-derived cardiomyocytes | Contraction, cardiac-specific gene expression, Gπ2 staining, multielectrode array | Mauritz et al., 2008 |
| Human iPS cells reprogrammed from primary keratinocytes | Cx43 1 | Cx43 in iPS cell- and ES cell-derived cardiomyocytes | Contraction, cardiac-specific gene expression, multielectrode array, electrophysiology, pharmacological reagents | Pfannkuche et al., 2009 |
| Mouse iPS cells (O9, N10), Mouse ES cells (R1, D3) | Cx43 in iPS cell- and ES cell-derived cardiomyocytes | Cardiomyocytes | Cx43 in iPS cell- and ES cell-derived cardiomyocytes | Contraction, cardiac-specific gene expression, multielectrode array, electrophysiology, pharmacological reagents | Martinez-Fernandez et al., 2009 |
| Mouse iPS cells reprogrammed without c-MYC | Cx431, Cx431 | Cx43 in iPS cell-derived cardiomyocytes in vivo and in vitro | In vivo 3 germ layer differentiation, i.e., endoderm, ectoderm, and mesoderm. In vitro cardiomyocyte differentiation | Contraction, cardiac-specific gene expression, multielectrode array, electrophysiology, EM | Sharovskaya et al., 2012 |
| Human iPS cells reprogrammed from HUVECs | Cx431, Cx431 | GJIC is re-established during reprogramming to pluripotency | Cx43 in iPS cell-derived cardiomyocytes | HUVECs express Cx431, Cx371, and Cx451 | Low GJIC in HUVECs | Ke et al., 2013 |
| Human iPS cells reprogrammed from human embryonic fibroblasts | Cx431, 2, Cx251, Cx261, Cx301, Cx30.21, Cx30.31, Cx311, Cx31.11, Cx31.91, Cx371, Cx401, Cx451, Cx461, Cx471, Cx591, Cx611, Cx621, Cx43 increases during reprogramming | Human embryonic fibroblasts express Cx431, 2 at low levels | Present3 | Human embryonic fibroblasts express Cx431, 2 at low levels | Lundy et al., 2013 |
| Human iPS cell line (iMR90) | Cx431, 2 expression in late-stage (80–120 days) cardiomyocytes vs. early stage (20–40 days) counterparts | Cardiomyocytes differentiated from iPS and ES cells using a long-term culture protocol | Significant increase in Cx431, 2 expression in late-stage (80–120 days) cardiomyocytes vs. early stage (20–40 days) counterparts | Optical contraction analysis, electrophysiology, Gπ2 staining, cardiac-specific gene expression, EM | Lundy et al., 2013 |

1 mRNA level; 2 protein level; 3 dye coupling (Lucifer yellow etc.); EM, electron microscopy.
CONCLUSIONS: CURRENT ANSWERS TO THE QUESTIONS ON CONNEXIN EXPRESSION AND GAP-JUNCTIONAL INTERCELLULAR COMMUNICATION IN ES/iPS CELLS

It seems reasonable to conclude that mRNAs encoding almost all of the connexins are expressed in ES/iPS cells. At protein level, however, expression of only a few connexins, such as Cx43, Cx45, Cx31, and Cx40, has been confirmed. Many studies have shown that undifferentiated ES/iPS cells communicate with each other via gap junctions at a high level. Several studies using Cx43 RNAi demonstrated that Cx43 contributes substantially to a high level of GJIC in undifferentiated ES/iPS cells.

Concerning changes in connexin expression and GJIC during differentiation of ES/iPS cells, it has been shown that expression of tissue-related connexins, such as Cx40, Cx43, Cx45, and Cx37 in the cardiomyocyte, is upregulated and that GJIC between pluripotent and differentiated cells is disrupted, resulting in formation of “communication compartment.” Regarding changes in connexin expression and GJIC during induction of pluripotency in somatic cells, the studies mentioned here have demonstrated that GJIC is re-established and Cx43 expression is upregulated during reprogramming to pluripotency.

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Cx43-mediated GJIC has been shown to play an important role in maintenance of pluripotency. In fact, pharmacological blockers of GJC and Cx43 downregulation by siRNA have been shown to induce a loss of their pluripotent state in mouse ES cells (Todorova et al., 2008). Cx43 has also been shown to play an important role in reprogramming, possibly by GJIC-independent mechanism including effects on the MET process (Sharovskaya et al., 2012; Ke et al., 2013).

Because the literature on connexin expression and GJIC in iPS cells is limited, it is difficult to conclude whether there are differences between ES and iPS cells at present. However, available data suggest that human ES and iPS cells share a similar feature concerning expression and function of connexins.

Although important roles of connexin expression and/or GJIC in ES/iPS cells can be currently perceived, many critical questions including precise mechanisms by which connexin expression influences pluripotency and reprogramming remain to be clarified.

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