INTRODUCTION

Human adult progenitor stem cells exhibiting pluripotent properties are a potential therapy for inducing cardiac repair following damage caused by ischemia (Martin-Rendon et al. 2008; Murry et al. 2005; Psaltis et al. 2008; Segers and Lee 2008; Yip et al. 2008). The clinical outcomes of injecting/perfusing autologous adult bone marrow stem cells into heart muscle or into coronary blood vessels have been variable and controversial and the underlying mechanisms by which bone marrow stem cells can repair heart tissue are unclear (Balsam et al. 2004; Passier et al. 2008; Roell et al. 2007).

One mechanism by which progenitor bone marrow (BM)-derived stem cells, defined by CD34 antigen expression (Sitnicka et al. 2003; Zhang et al. 2007), improve heart function could involve their participation in direct local intercellular signaling. Cell interactions are complex and include intercellular signaling across gap junctions as well as paracrine signaling that may implicate the component connexin hemichannels between stem cells and cardiac cells. We examined therefore whether connexin proteins that provide membrane-traversing channels underpinning functional coupling are present in bone marrow cells administered in cardiac repair therapy. Connexins are also involved in cell adhesion processes (Cotrina et al. 2008) and their importance, together with other proteins, e.g., cadherins (Zhang et al. 1998) in cell adhesion and cohesion, has long been appreciated. Encouraging these studies is the fact that cardiac tissue is a rich depository of connexin (Cx)43, a protein that plays important roles in human cardiogenesis (Moore et al. 2008) and is a determinant of myocardial infarct size (Kanno et al. 2003).

To determine whether progenitor stem cells, especially CD34+ cells, or myeloid subpopulations purified from BM or cord blood (CB) express connexins and engage in communication with resident cardiac cells via membrane channels, we carried out a comprehensive study of connexin expression in adult BM and CB cells. We also investigated their intercellular coupling competence using dye transfer approaches. Connexins are a 22 member highly conserved family of proteins in humans (Willecke
et al. 2002). They oligomerize in the endoplasmic reticulum into hemichannels that are delivered to the plasma membrane where they dock with partner hemichannels exposed on closely aligned neighboring cells and generate gap junction intercellular channels (Evans et al. 2006). Flow-cytometric studies using specific connexin channel inhibitors were carried out to establish whether any intercellular coupling detected was mediated by gap junctions. The results show that adult BM and CB progenitor stem cells have a very low capacity to communicate in a connexin-dependent manner and support an emerging view that autologous bone marrow cells may contribute to improved cardiac output following a cardiac infarction not by direct intercellular communication but rather by other mechanisms that may include secretion of growth factors and cytokines that promote proangiogenic effects.

**METHODS**

RNA expression of 20 human connexins and N-cadherin was carried out using standard reverse transcriptase–polymerase chain reaction (RT-PCR) techniques as previously described (Oviedo-Orta et al. 2000). The primers to each of the Cxs used are listed in Table I. Briefly, total RNA was extracted using the RNeasy mini kit (Qiagen, Germany) and contaminating genomic DNA removed by treatment with RNase-free DNase I (Qiagen). First-strand cDNA was synthesized with 1 μg total RNA using the protoscriptII RT-PCR kit (New England Biolabs). Since most connexins are encoded within a single exon, a reverse transcriptase–free reaction was performed to demonstrate absence of genomic DNA contamination. Human placenta transcriptase–free reaction was performed to demonstrate absence of genomic DNA contamination. Human placenta DNA (Sigma) was used as a control. The PCR programme was 94°C for 5 min, followed by 35 cycles at 94°C for 30 s, 58°C for 30 s, and 72°C for 30 s, with a final step of 72°C for 10 min. Cord blood and normal marrow were obtained with informed consent and approval from the South East Wales Research Ethics Committee and were performed in accordance with the ethical standards laid down in the 1964 Declaration of Helsinki. BM mononuclear cells were pooled from patients and CD34+ cells purified using MiniMACS columns (Miltenyi Biotec., Surrey, UK). Cell purity was determined by flow cytometry (FACS Calibur) and data processed using the programme WinMDI2.8 (Purdue University Cytometry Labs). CD34+ cells were cultured in standard (RPMI-1640) medium supplemented with stem cell factor (SCF), interleukin (IL)-3, IL-6, granulocyte-macrophage colony-stimulating factor (GM-CSF), G-CSF, and FMS-like tyrosine kinase 3 ligand (FLT3). HeLa cells transfected with Cx43 (Martin et al. 2001) and HL-1 cells, a mouse atrial cardiomyocyte tumor cell line, were grown in Dulbecco’s modified Eagle’s medium (DMEM) or Claycombe medium (JRH Biosciences, Hertfordshire, UK), respectively, as described (Verma et al. 2008; White et al. 2004).

Western blotting of cell proteins extracted at 4°C in sodium dodecyl sulfate with added proteolytic inhibitors (1 mg/ml leupeptin, 1 mg/ml aprotinin, and 0.5 mM phenylmethyl sulfonyl fluoride) was carried out in 4–12% (w/v) polyacrylamide gels. Separated proteins were electrophoretically transferred to nitrocellulose filters and nonspecific protein binding sites blocked before exposure to anti-connexin antibodies. After treatment with horseradish peroxide–conjugated secondary antibodies, the blots were visualized using chemiluminescence. Western blots were quantified using NIH image software. Western blotting of cell proteins extracted at 4°C in sodium dodecyl sulfate with added proteolytic inhibitors (1 mg/ml leupeptin, 1 mg/ml aprotinin, and 0.5 mM phenylmethyl sulfonyl fluoride) was carried out in 4–12% (w/v) polyacrylamide gels. Separated proteins were electrophoretically transferred to nitrocellulose filters and nonspecific protein binding sites blocked before exposure to anti-connexin antibodies. After treatment with horseradish peroxide–conjugated secondary antibodies, the blots were visualized using chemiluminescence. Western blots were quantified using NIH image software.

### Table 1. PCR primers used to analyze 20 human connexins and N-cadherin.

| Connexins | Accession numbers | Forward & reverse primers | Product size (bp) |
|-----------|-------------------|---------------------------|------------------|
| Cx25      | NM_198568         | ggggcacagcagggaaatc tgcgttccagcttagtgg | 192 |
| Cx26      | NM_004004         | cctgaagttggtgctcaaaagt gcggcagaaagagaatca | 150 |
| Cx30      | NM_006783         | ttgcattcgagcagatttg ggtttggttccttgga | 244 |
| Cx30.2    | NM_181538         | gccactggaattatcagag ccggcagatgctcaatgt | 250 |
| Cx30.3    | NM_153212         | tgttggatggttctgc gcacacatacccaactct | 236 |
| Cx31      | NM_024009         | ttctggaagcttcaatag tggtgtcagagtgtagcag | 186 |
| Cx31.1    | NM_005268         | tccaagccctcagaagagag ctaggccgaaaaatgcag | 229 |
| Cx31.9    | NM_152219         | caagaggtcagttgc gatgtcagagatcaggtglag | 150 |
| Cx32      | NC_000023         | tcccttgctcactctagt cctctagtggtgcctctgt | 156 |
| Cx36      | NM_020660         | ttctcagctcgcagagaga gatgtcagttcagcagag | 218 |
| Cx37      | BC027889          | aagatctcgtgcggcagag ttcaccccaacatagtcg | 244 |
| Cx40      | BC013313          | tagagcaagttgcctcact gtagtccgacaccaggcagta | 186 |
| Cx40.1    | NM_153368         | aagagctgtggtgctcaca tgctcagagatcaggtggta | 229 |
| Cx43      | BC026329          | atgcagcagttgcctcctg tgtcttgatgctcatgtctg | 249 |
| Cx45      | NM_005497         | cagctgtaagccagcagaaag tttcatgagcccatctcc | 217 |
| Cx46      | NM_021945         | catcttaagcagctttcc gcggctcaatagtaagagag | 186 |
| Cx47      | NM_020435         | gatccacaaacactcaccct aagggcgctagcagctc | 229 |
| Cx50      | NM_005267         | gagagaagctcggctctcagg cggggtggctacttccttcc | 175 |
| Cx59      | NM_030772         | agtgcacacccggcagct ggacccctccttcggtgtagtg | 205 |
| Cx62      | NM_032602         | ggccacagttgcctggtg tcctcagctgatgcttcctg | 193 |
| GAPDH     | NM_002046         | tgccttgcgacccacacact tgccttgcacccaccct | 329 |
| N-cadherin| NM_001792         | ccagagatggtgtaggtgtaag | 201 |
antibodies, signals were amplified using an enhanced chemiluminescence (ECL) solution (Amersham Biosciences, UK). Connexin antibodies were generated to a range of intracellular peptides linked to keyhole limpet haemocyanin (Oviedo-Orta et al. 2000) or were purchased from Zymed or Chemicon laboratories (USA). These antibodies bind to rodent and human connexins.

Coupling was measured by detection of dye transfer between cells. Monolayer cells were grown to confluence in 25-cm² diameter flasks. Donor cells were loaded with 5 mM calcein (Molecular Probes), a fluorescent probe that permeates Cx37 and Cx43 gap junction channels (Veitch et al. 2004) and recipient cells with 5 μg/ml DiI C18 (Molecular Probes). After incubation in 5% CO₂ at 37°C for 30 min, the dye-loaded cells were washed with phosphate-buffered saline (pH 7.4) and then harvested after treatment with trypsin. Cells were resuspended in culture medium and 2 × 10^5 donor and recipient cells in a 1:1 ratio were cultured at 37°C in 5% CO₂ for 4 h. As controls, non-dye-loaded cells of each category were used. Dye transfer was evaluated by flow cytometry and repeated 3 to 4 times. Cells grown in suspension were treated as with confluent monolayers with omission of trypsin treatment. To study the involvement of gap junctional coupling, cells were treated for 30 min with the following gap junction inhibitors: 18\(\alpha\)-glycyrrhetinic acid (18GA) or Gap 27 (sequence SRPTEKTIFII: residues 204–214 of Cx43) as stated in the figure legends. In some experiments, Gap 27 was substituted by a second Cx mimetic peptide Gap 26 (sequence VCYDKSFPISH-VR; residues 63–75 of Cx43) that, as previously shown (Evans and Leybaert 2007), also inhibits gap junctional communication.

RESULTS

Adult BM and CB cells were fractionated into subpopulations of stated purity and RNA expression of 20 human connexins was examined by RT-PCR (Table 2). Cx37 expression was detected in bone marrow and cord blood CD34⁺ cells and in cord blood CD14⁺ monocyte cell populations. Cx43 was also detected in CB and BM derived CD34⁺ cells as well as in CB CD14⁺ cells. A signal was repeatedly observed with Cx26 (a connexin found in skin and the ear; Willecke et al. 2002) in CD14⁺ cells in CB but not in BM and is probably an artefact. Cx26 was not detected in CD34⁺ cells purified from cord blood or bone marrow. mRNA expression of N-cadherin, an adhesion protein expressed at low levels, provided a positive control in CD34⁺ cells from both sources. Freshly isolated CD34⁺

| Cxs | CD34⁺ (93%) | CD14⁺ (95%) | CD15⁺ (92%) | CD34⁺ cultured for 10 days | CD34⁺ (86%) | CD14⁺ (80%) | CD15⁺ (98%) |
|-----|-------------|-------------|-------------|----------------------------|-------------|-------------|-------------|
| Cx25 | –           | –           | –           | –                          | –           | –           | –           |
| Cx26 | –           | +           | –           | –                          | –           | –           | –           |
| Cx30 | –           | –           | –           | –                          | –           | –           | –           |
| Cx30.2 | –       | –           | –           | –                          | –           | –           | –           |
| Cx30.3 | –         | –           | –           | –                          | –           | –           | –           |
| Cx31 | –           | –           | –           | –                          | –           | –           | –           |
| Cx31.1 | –         | –           | –           | –                          | –           | –           | –           |
| Cx31.9 | –         | –           | –           | –                          | –           | –           | –           |
| Cx32 | –           | –           | –           | –                          | –           | –           | –           |
| Cx36 | –           | –           | –           | –                          | –           | –           | –           |
| Cx37 | +           | +           | –           | –                          | +           | –           | –           |
| Cx40 | –           | –           | –           | –                          | –           | –           | –           |
| Cx40.1 | –         | –           | –           | –                          | –           | –           | –           |
| Cx43 | +           | +           | –           | +                          | +           | –           | –           |
| Cx45 | –           | –           | –           | –                          | –           | –           | –           |
| Cx46 | –           | –           | –           | –                          | –           | –           | –           |
| Cx47 | –           | –           | –           | –                          | –           | –           | –           |
| Cx50 | –           | –           | –           | –                          | –           | –           | –           |
| Cx59 | –           | –           | –           | –                          | –           | –           | –           |
| Cx62 | –           | –           | –           | –                          | –           | –           | –           |
| N-Cad | +          | –           | –           | –                          | –           | –           | –           |

Note. Numbers in parentheses indicate purity of subpopulation analyzed.
cells are a largely quiescent population; to determine whether the cell cycle status affected connexin expression, we repeated the analysis on CD34\(^+\) cells cultured in the presence of growth factors. Culturing of these cells for 13 days did not promote connexin mRNA expression.

Cx protein expression was examined by Western blotting. Since antibodies to the full range of Cxs are unavailable, we confined our attention to Cx32, Cx37, Cx40, and Cx43 using appropriate tissue controls expressing these connexins. Figure 1 shows that Cx32, Cx37, Cx40, and Cx43 could not be detected in BM and CB stem cell progenitor populations.

We next examined whether CD34\(^+\) cells communicated via gap junctions with each other, as well as with other mammalian model cells expressing Cx43, by following intercellular transfer of calcein, a small fluorescent dye loaded into donor cells. To validate this assay of cell communication and the efficacy of the two gap junction inhibitors used, we demonstrated first that HeLa cells expressing Cx43 (previously shown to be communication competent on the basis of intracellular Ca\(^{2+}\) transfer and electrical coupling; Paemeleire et al. 2000) transferred dye (Figure 2A). Also, HL-1 cells that display cardiac-type properties and express Cx43 and Cx37 (Verma et al. 2008) were coupled but to a lesser extent. In both instances, dye transfer was inhibited by the gap junction inhibitor 18GA and connexin mimetic peptide Gap 27. Dye transfer between freshly prepared BM or CB CD34\(^+\) was absent (Figure 2B).

Low-level transfer was observed in cells that were cultured for 13 days. However, two peptide inhibitors of gap-junctional coupling had little effect on the low-level dye transfer detected, indicating that this was unlikely to be mediated by gap junctions. We also analyzed the capacity of the CD34\(^+\) cells to communicate across gap junctions by their coculture with HL-1 cells or with HeLa cells expressing Cx43 (Figure 2B). No dye coupling was observed between CD34\(^+\) cells and each of the Cx43-expressing cells used in these studies. The data are summarized in Figure 3.

**DISCUSSION**

Several reports have claimed that adult human stem cells, especially autologous BM CD34\(^+\) subpopulations, administered directly into hearts of patients with ischemic damage, enhanced cardiac functioning (Balsam et al. 2004; Martin-Rendon et al. 2008; Murry et al. 2005; Psalitis et al. 2008; Roell et al. 2007; Segers and Lee 2008; Yip et al. 2008; Zhang et al. 2007). For such treatment to be effective, one likely scenario is that BM stem cells introduced into the heart become functionally integrated and are retained in cardiac tissue, possibly transdifferentiating into cardiac-like cells (Breitbach et al. 2007). Here, we addressed the thesis that key proteins enabling the functional integration of stem cells into heart may be connexins, especially Cx43, a protein that is expressed at high levels in myocardial tissue.

Connexins are proteins that oligomerize into dodecameric gap junction channels that facilitate direct intercellular cell communication, allowing metabolic, ionic, and electrical coordination of cell assemblies. Connexins are also present in the plasma membrane as unopposed hexameric hemichannels and these, when in open channel configuration, connect the cell interior to the external environments, allowing passage through the channels of, for example, ATP and calcium (Evans et al. 2006; Verma et al. 2008). Connexin hemichannels in the plasma membrane confer the independent property of cell-cell adhesion in addition to intercellular communication (Cotrina et al. 2008; Wong et al. 2006).

The present results show that CD34\(^+\) cells that display stem cell progenitor properties (Kawamoto et al. 2006) express very low levels of the 20 connexin isoforms examined. Only Cx37 and Cx43 were detected at the mRNA level. Specifically, Cx32, present in primitive hematopoietic progenitor cells in mice (Hirabayashi et al. 2007), was not found. At the functional level, flow cytometry studies were carried out in the presence of specific inhibitors of gap junctions and these showed that CD34\(^+\) cells derived from adult BM and CB showed little capacity to communicate with each other, if at all, in a connexin-dependent manner; these cells also did not communicate with other cell types that express Cx43, including a cell line that has retained many cardiodependent properties. The use of two gap junction inhibitors, 18GA and the connexin mimetic peptides Gap 27 and Gap 26 (Evans and Leybaert 2007), reinforces the data. Taken together, therefore, the compositional and functional data suggest that gap-junctional communication occurring between the introduced adult progenitor stem cells and resident heart cells is unlikely. Since Cx43 and Cx37 enhance cell adhesion (Wong et al. 2006), the extremely low expression of these proteins also does not encourage any general hypo-
Figure 2. (A) Validation of dye transfer assay showing HeLaCx43 cells co-cultures with HL-1 cells and inhibition by 18GA (75 μM) and Gap 27 (150 μg/ml). (B) Transfer of calcein between BM or CB CD34+ cells cultured for 5, 10, or 13 days as well as in co-cultured with HL-1 and HeLa Cx43. The data represent a compromise between voltage and compensation settings.
Possible mechanisms. Tunneling nanotubes extending players in improving the performance of hearts damaged proteins to assemble into gap junction channels. Cx43 and Cx37 recorded may be sufficient to allow a few out because the extremely low levels of mRNA encoding Cx-mediated communication cannot be entirely ruled of many adult stem cells. However, a possible role for or a low connexin expression is a characteristic property tissue. Indeed, they reinforce a broad view that a lack of the integration of bone marrow CD34+ cells or myocytes in hearts damaged by hypoxia (Drexler and Wollert 2009). Finally, in attempts to identify optimum cell types to drive robust cardiac myogenesis, cardiac progenitor cells have been identified in embryonic/adult hearts and these could also be key players in inducing cardiac repair (Nahrendorf et al. 2007).

In conclusion, the results show that connexin proteins, assembled into hemichannels and gap junctions in the plasma membrane where they underpin intercellular communication, are expressed at such low levels by progenitor CD34+ cells and myeloid lineage cell types that they are unlikely to provide a mono-causal explanation for clinical data demonstrating beneficial outcomes of application of autologous adult BM stem cells in cardiac repair. Deeper knowledge of adult stem cell plasticity (Chen et al. 2006; Jones and Wagers 2008; Raff 2003) and a better understanding of the complex biochemical mechanisms underpinning cell differentiation and interactions, and especially paracrine factors released by BM cells, is required to unravel the variable clinical trial results.

Declaration of interest: The authors report no conflicts of interest. The authors alone are responsible for the content and writing of the paper.

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