Cellular Bioenergetics is an Important Determinant of the Molecular Imaging Signal Derived from Luciferase and the Sodium-Iodide Symporter

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Running title: Bioenergetics Influences Cell Imaging Signal

Subject codes:
[124] Cardiovascular imaging agents/Techniques
[32] Nuclear cardiology and PET
[150] Imaging
[27] Other Treatment
[33] Other nuclear testing
[140] Energy metabolism
[142] Gene expression

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This manuscript was sent to Mark Alan Sussman, Consulting Editor, for review by expert referees, editorial decision, and final disposition.

In November 2012, the average time from submission to first decision for all original research papers submitted to Circulation Research was 15.8 days.

DOI: 10/1161/CIRCRESAHA/112/273375
ABSTRACT

Rationale: Molecular imaging is useful for longitudinal assessment of engraftment. However, it is not known which factors, other than cell number can influence the molecular imaging signal obtained from reporter genes.

Objective: The effects of cell dissociation/suspension on cellular bioenergetics and the signal obtained by firefly luciferase (Fluc) and human Na-I symporter (hNIS) labeling of cardiosphere-derived cells (CDCs) was investigated.

Methods and Results: $^{18}$FDG uptake, ATP levels, $^{99m}$Tc-pertechnetate uptake and bioluminescence were measured in vitro, in adherent and suspended CDCs. In vivo dual isotope SPECT-CT imaging or bioluminescence imaging (BLI) were performed 1hr and 24hrs following CDC transplantation. SPECT quantification was performed using a phantom for signal calibration. Cell loss between 1hr & 24hrs post-transplantation was quantified by qPCR and ex vivo luciferase assay.

Cell dissociation followed by suspension for 1hr resulted in decreased glucose uptake, cellular ATP, $^{99m}$Tc uptake and BLI signal by 82%, 43%, 42%, and 44% respectively, when compared to adherent cells, in vitro. In vivo $^{99m}$Tc uptake was significantly lower at 1hr, when compared to 24hrs following cell transplantation in the non-infarct ($p<0.001$, n=3) and infarct ($p<0.001$, n =4) model, despite significant cell loss during this period. The in vivo BLI signal was significantly higher at 1hr than at 24hrs ($p<0.01$), with the BLI signal being higher when CDCs were suspended in glucose-containing medium compared to saline (PBS).

Conclusion: Adhesion is an important determinant of cellular bioenergetics, $^{99m}$Tc-pertechnetate uptake and BLI signal. BLI and NIS imaging may be useful for in vivo optimization of bioenergetics in transplanted cells.

Keywords:
Molecular imaging, metabolism, stem cell

Non-standard Abbreviations:
CDCs Cardiosphere-derived cells
HSV-tk Herpes simplex virus thymidine kinase
Fluc Firefly luciferase
hNIS Human sodium-iodide symporter
polyHEMA poly-2hydroxyethyl methacrylate
INTRODUCTION

Cellular cardiomyoplasty is plagued by low engraftment and small functional benefit. Molecular imaging facilitates the study of in vivo stem cell biology and optimization of cell therapies by permitting in vivo quantification of stem cell engraftment. For in vivo visualization, transplanted cells need to express a reporter gene or they must be labeled ex vivo.\(^1\) Reporter gene strategies (e.g. bioluminescence imaging of firefly luciferase gene expression\(^2\) and PET/SPECT imaging of herpes simplex virus thymidine kinase (HSV-tk)\(^3\) or human sodium-iodide symporter (hNIS)\(^4\) gene expression) are superior to ex vivo labeling of cells with tracers (e.g. \(^18\)FDG \(^5,\)\(^6\) or Indium\(^7\)) or nano-particles for longitudinal assessment of engraftment,\(^8\) because transplanted cell viability is a prerequisite for reporter gene expression. However, it is not known which factors, other than cell number can influence the molecular imaging signal obtained from reporter genes.

Cell metabolism is an important determinant of cell survival, proliferation and function.\(^9,\)\(^11\) Studies in cancer cells reveal that they primarily utilize glucose via glycolysis, rather than oxidative phosphorylation for ATP generation, despite the availability of oxygen – this phenomenon is referred to as the \textit{Warburg effect}.\(^12\) However, very little is known about the determinants of metabolism and its relationship to function in stem cells.

In this study, we examined the effect of cell dissociation/suspension on cellular bioenergetics and the molecular imaging signal obtained by hNIS (SPECT) and firefly luciferase (fluc) labeling (bioluminescence imaging or BLI) of cardiosphere-derived cells (CDCs). CDCs are adherent cells composed of a mixture of cardiac-derived progenitor and supporting cells, and have recently completed a phase1 clinical trial.\(^13\) NIS promotes cellular uptake of iodide or \(^99\)mTc-pertechnetate, driven by the trans-
membrane sodium gradient, which is maintained by Na\(^+\)-K\(^+\)-ATPase, whereas BLI is based on oxidation of the injected substrate D-luciferin by luciferase, a reaction that requires oxygen, Mg\(^{2+}\), and ATP, and results in emission of photons. Since signal generation by luciferase and NIS is linked to cellular ATP, we hypothesized that cellular bioenergetics would influence the molecular imaging signal obtained from \(h\)NIS and \(f\)luc-labeled cells. We studied \(h\)NIS labeling, because it is a human gene, that could be useful for longitudinal follow up of cell engraftment in small and large animal models and patients; and luciferase labeling because it is widely used for quantification of engraftment in rodent models of cell transplantation. Our results illustrate that cell dissociation which is an essential first step in most clinical and experimental studies of cell transplantation impairs cellular bioenergetics and the molecular imaging signal derived from NIS and luciferase.

METHODS

**Cell isolation and culture.**
Cardiosphere-derived cells (rCDCs) were isolated from syngeneic, male Wistar Kyoto rat hearts as previously described. rCDCs were cultured in IMDM medium (Invitrogen) containing 10% FBS, 10% glutamine and 0.1mM mercaptoethanol, and expanded to 3-5 passages prior to experiments. Please see supplemental data section for details.

**Lentivirus synthesis.**
A third-generation lentiviral vector system (kindly supplied by Professor Inder Verma, Salk Institute, USA) was used to label the rCDCs. Please see supplemental data section for details.

**In vitro glucose uptake (\(^{18}\)FDG) uptake.**
One day before the experiment, rCDCs were plated at a density of 1\(\times\)10\(^5\) cells/well. Prior to labeling, cells were washed twice with PBS and the medium was changed to glucose free-DMEM for 1hr. \(^{18}\)FDG (74 kBq/ml) was added to half the plated cells for 30min to measure glucose uptake in adherent cells; the remainder of the cells were trypsinized and suspended in medium containing \(^{18}\)FDG (74 kBq/ml) for 30min in order to measure glucose uptake in cell suspension. Subsequently, cells were washed twice with cold PBS to remove any remaining free \(^{18}\)FDG, suspended in 1ml 20% sodium dodecylsulfate to lyse the cells, and transferred to 5ml vials. Counts were recorded in a gamma-counter (LKB Wallac, Turku, Finland). After the gamma counting, 4ml of ice cold acetone was added to each sample, and samples were kept at 4\(^\circ\)C to allow for radiotracer decay, prior to performing the Bradford protein assay.
ATP measurements.

ATP estimation was performed using the ATP Determination Kit (A22066, Molecular Probes) using a luminometer (Turner BioSystem Veritas, Madison, WI, USA). All experiments were performed using 1 × 10^4 rCDCs (non-transduced) per well in a 96 well plate following 18-24 hrs of culture for the adherent condition, and following trypsinization and 1hr of suspension for the suspension condition. For suspension conditions, the wells were coated with poly-2hydroxyethyl methacrylate (polyHEMA), overnight, prior to cell plating. PolyHEMA was chosen because it is known to prevent cell attachment and spreading^{18,19}. Cell lysis Buffer (Cell Signaling Technology) was used to lyse the cells in each well for 20min; standard reaction solution was injected through the automated injector. The signal was normalized to protein content using the Bradford Assay. ADP/ATP ratio was assessed using ApoSENSOR ADP/ATP Ratio Assay Kit.

Flow cytometry.

Annexin V and Propidium iodide (Invitrogen) were used to quantify apoptotic and dead cells, respectively immediately after cell dissociation and after suspension in cell culture medium for 1hr and 6hrs. Please see supplemental data section for detailed methods.

Measurements of cellular metabolism.

Seahorse Bioscience XF96 instrument^{20} was used to measure the rate of change of dissolved O_2 in each well (termed OCR or oxygen consumption rate) which reflects oxidative phosphorylation, and change in pH in the media which reflects glycolysis, the 2 main sources of ATP in cells. Oxygen concentration and pH were measured over 5 min periods with a mixing time of 2min in each cycle with three cycles in total for adherent CDCs (plated for 18-24hrs) and rCDCs suspended for 1hr in Seahorse medium containing glucose (25mM) in a specialized 96 well plate (n=3). Restoration of metabolism following cell suspension was assessed by re-plating viable rCDCs in a specialized 96 well plate (Seahorse biosciences) following suspension for 0, 3hrs and 6hrs in cell culture medium; metabolic measurements (OCR and ECAR) were performed at 24hrs following cell dissociation.

The effect of Oligomycin (2µM), an inhibitor of the mitochondrial ATP synthase was used to assess dependence of CDCs on oxidative phosphorylation for ATP synthesis. Iodoacetate (600µM), an inhibitor of the glycolytic enzyme, glyceraldehyde-3-phosphate dehydrogenase (GAPDH) was used to assess the contribution of glycolysis to ATP synthesis. Cells were treated with these compounds for 30min prior to measurements.

The respiratory rates in each well were normalized to protein content using Bradford Assay. These experiments were performed at least in triplicate and repeated 2-3 times.
In vitro $^{99m}$Tc-Pertechnetate uptake.

rCDCs were transduced with a 3rd generation lentivirus expressing the human sodium-iodide symporter ($Lv$-CMV-hNIS) at an MOI of 20. In vitro $^{99m}$Tc-pertechnetate uptake was measured by incubating $h$NIS$^+$ rCDCs cultured for 18-24hrs in a 6 well plate at a density of $10^5$ cells/well for the adherent condition, or trypsinized and suspended in PBS (phosphate buffered saline) or IMDM medium (Invitrogen); $^{99m}$Tc-pertechnetate (11.1 kBq/mL) was added for 1hr, immediately after generation of cell suspensions. The effect of perchlorate (100 μM), a specific NIS blocker on $^{99m}$Tc-pertechnetate uptake was measured by adding perchlorate to some wells prior to the addition of $^{99m}$Tc-pertechnetate. At the end of 1hr, cells were rinsed twice with ice cold PBS and lysed with 20% sodium dodecylsulfate. Counts were recorded in a gamma-counter (LKB Wallac, Turku, Finland) and the Bradford protein assay was performed to normalize $^{99m}$Tc uptake by protein content. We investigated cell suspension in PBS (phosphate buffered saline) and IMDM medium because PBS/saline, which lacks substrates, has been extensively used in experimental$^{16,21}$ and clinical studies$^{13}$ of CDC transplantation; IMDM medium (Invitrogen) contains metabolic substrates like D-glucose, Ca$^{2+}$, Mg$^{2+}$ and is used for CDC culture.

In order to investigate the influence of trypsin on $^{99m}$Tc-pertechnetate uptake, $h$NIS$^+$ rCDCs were dissociated either using 0.05% trypsin (Invitrogen), which usually takes ~2min, or non-enzymatic cell dissociation solution ( Sigma-Aldrich; contains EDTA and other proprietary reagents) which takes ~ 20-30min (for complete dissociation into single cells), following which they were suspended in IMDM medium containing $^{99m}$Tc-pertechnetate (11.1 kBq/ml) for 1hr. Subsequently, cells were rinsed twice using ice cold PBS, lysed, and analyzed using the gamma counter; Bradford assay was performed to normalize uptake results to protein content. These experiments were performed in triplicate and repeated twice.

All in vitro studies were performed at 37°C.

In vivo $^{99m}$Tc uptake (SPECT/CT Imaging).

Dual isotope imaging was performed. Data was acquired in list mode and post-processed by applying two energy windows ("75 keV +10%/-10%" and "140 keV +10%/-10%") to obtain $^{201}$Tl and $^{99m}$Tc projections separately. Please see supplemental methods section for details of animal surgery and imaging.

Three million NIS$^+$ rCDCs suspended in 100µl of IMDM (Invitrogen) were injected directly into the myocardium at three sites in the anterior wall of the left ventricle using a 30G needle. In vivo dual isotope SPECT imaging was performed 1hr after injection of $^{99m}$Tc-pertechnetate (555-740 MBq) and $^{201}$Tl (37-74 MBq);$^4$ CT imaging was performed prior to SPECT imaging. Animals were allowed to recover in their
cages after completion of imaging on day 0. After 24hrs, the same rats were re-injected with $^{99m}$Tc-pertechnetate (555-740 MBq) and $^{201}$Tl (37-74 MBq) via the tail vein and \textit{in vivo} dual isotope SPECT-CT imaging was performed. The rats were euthanized after completing the 24hr imaging protocol.

\textbf{Ex-vivo SPECT imaging.}

In order to validate the results obtained by \textit{in vivo} imaging and to confirm the origin of the \textit{in vivo} signal, a high resolution \textit{ex vivo} SPECT scan was performed in hearts excised from a separate set of animals at 1hr (n=2) and 24hrs (n=2) following transplantation with NIS$^+$ rCDCs. Please see supplemental data section for details.

\textbf{Image quantification.}

For absolute quantification of \textit{in vivo} $^{99m}$Tc-pertechnetate uptake, we generated a dose response plot by dual isotope SPECT imaging of a rat size phantom containing several concentrations of $^{99m}$Tc-pertechnetate and $^{201}$Tl. Calibration factor (MBq/i.i.) was defined as the quotient of the known activity concentration (MBq/ml) within the radioactive sphere in the phantom divided by the measured mean image intensity (i.i/cm$^3$) within a ROI drawn over the small sphere in the SPECT image of the phantom. For absolute quantification, regions of interest (ROI) were manually defined, image intensity within the ROI was multiplied by the calibration factor (CF) to give the radioactivity (MBq), and further divided by the ROI volume to get the uptake concentration (MBq/ml). Please see supplemental data section for details.

\textbf{In vivo bioluminescence imaging.}

For \textit{in vivo} BLI, rCDCs were labeled with a 3$^{rd}$ generation lentivirus expressing firefly luciferase (\textit{Lv-CMV-fLuc}) at an MOI of 20, which did not affect CDC survival, proliferation or differentiation. One million rCDCs transduced with \textit{Lv-CMV-fLuc} were trypsinized, suspended in 100µl of PBS/saline (n=4) or IMDM (n=4) and then injected intra-myocardially into 3 sites in the anterior wall of the left ventricle of non-infarcted male WK rats using a 30G needle, following a left sided thoracotomy. \textit{In vivo} BLI was performed using the Xenogen IVIS 200 system 1hr and 24hrs after cell transplantation. We studied non-infarcted rats in order to reduce signal variability resulting from the location of the transplanted cells (infarct versus border-zone) given the small size of the left ventricle and known dependence of the BLI signal upon the availability of oxygen.

\textbf{In-vitro BLI.}

\textit{In vitro} BLI was performed in a 96 well plate using a luminometer (Turner VERITAS Microplate, Promega, Madison, WI, USA) using \textit{fluc}-transduced rCDCs. Ten thousand cells ($10^4$ cells/well) were plated for 18-24 hrs for the \textit{adherent} cell condition and for 1hr on polyHema coated wells (\textit{suspension
condition) in PBS/saline or PBS/saline containing Ca^{2+}/Mg^{2+} (1mM) and D-glucose (5.6mM). We used Ca^{2+}/Mg^{2+}/glucose-containing PBS rather than IMDM in order to avoid the possible effects of phenol red, a component of IMDM medium, on the bioluminescence signal. The signal (RLU) was measured in suspended and adherent cells after injection of 30µg/ml of D-Luciferin (sodium salt, Gold Biotechnology, Saint Lois, MO, USA) into each well using the automated injector. Cells were lysed using the lysis buffer contained in the Dual-Luciferase® Reporter Assay System (E1910). The signal was normalized by protein content using the Bradford assay. All experiments were performed at least in triplicate and repeated twice.

**Quantification of engraftment.**

We used the ex vivo luciferase assay to quantify in vivo engraftment of fluc⁺ rCDCs, and quantitative PCR for the male rat-specific SRY gene to quantify engraftment of NIS⁺ rCDCs in the first 24hrs following transplantation. Please see supplemental data section for detailed methods and rationale for the 2 methods.

**Statistical analysis.**

For matched comparisons of continuous variables at different time points, the paired t-test was used (or repeated measures ANOVA, in case of more than 2 groups). For comparisons of continuous variables between two independent groups the Student’s t-test was used. A p<0.05 was chosen for statistical significance. Values are reported as mean ± SD.

**RESULTS**

**Bioenergetics in CDCs.**

Since cell trypsinization/dissociation is an important step in most experimental and clinical studies of cell transplantation in the heart, we initially compared glucose uptake, ATP levels and cellular metabolism in non-transduced rCDCs that were trypsinized and suspended in culture media for 1hr, and CDCs that were plated on an adherent surface (tissue culture treated wells) for 18-24hrs.

Cell viability was consistently >96% immediately post-dissociation, when assessed by trypan blue staining and flow cytometry (Fig 1A, B). However, CDC viability was reduced to 83±4% and 74±3% (n=3), respectively, following 1hr and 6hrs of suspension (Fig 1C).

Cellular glucose (¹⁸FDG) uptake and ATP levels were decreased by 82% and 43% respectively, following 1hr of cell suspension, when compared to adherent cells. (Fig 2A: ¹⁸FDG: 3.03±1.06 × 10⁴ vs. 1.74±0.23 × 10⁵ counts/mg protein in suspended and adherent cells respectively, p<0.001; Fig 2B: ATP:...
2.19±0.27 vs. 3.87±0.65 nmol/mg protein in suspended and adherent cells respectively; \( p<0.005 \). The ADP/ATP ratio increased from 0.27 in adherent rCDCs, to 1.1 and 1.8 in suspended rCDCs at 1hr and 3hrs respectively, indicating progressive bioenergetic deficits with increasing periods of cell suspension.

Oxygen consumption rate (OCR) which reflects oxidative phosphorylation was significantly reduced following 1hr of cell suspension (25±3 pmol/min in adherent rCDCs vs. 12±5 pmol/min in suspended rCDCs; \( n=3 \), \( p<0.01 \)). Extracellular acidification rate (ECAR), which reflects glycolysis was also significantly reduced upon 1hr of cell suspension (16±2 mpH in adherent rCDCs vs. 10±3mpH in suspended rCDCs; \( n=3 \); \( p<0.05 \)), indicating that cell suspension impairs cellular bioenergetics (Fig 2C).

Oxygen consumption rate and extracellular acidification rate returned to normal at 24hrs, following re-plating of viable cells suspended for 3hrs and 6hrs, indicating that the metabolic derangements following cell suspension are reversible and linked to cell adhesion (Fig 2D).

Oligomycin, an inhibitor of mitochondrial F1-F0 ATP synthase, reduced OCR by 59 ± 2% (\( p<0.001; n=3 \)) in adherent CDCs (Fig 2E), without affecting cellular ATP levels (Fig 2F); in contrast, iodoacetate, an inhibitor of glycolysis reduced ATP levels by 88±3% (\( p<0.001; n=6 \)) despite the presence of glucose (Fig 2F), indicating dependence of CDCs on glycolysis for ATP synthesis, evidence for the Warburg effect.

**In vitro studies.**

We measured the effect of cell dissociation/suspension on the signal derived from \( h \)NIS and firefly luciferase labeling of rCDCs in vitro, in the presence of either IMDM medium (Invitrogen) or PBS containing glucose/Ca\(^{2+}\)/Mg\(^{2+}\) and PBS/saline.

Trypsinization followed by suspension of NIS\(^+\) CDCs for 1hr in regular culture medium (IMDM) decreased \( ^{99m}\)Tc-pertechnetate uptake by 42% (1.10±0.21 × 10\(^4\) vs. 1.89±0.15 × 10\(^4\) counts/mg protein in suspended and adherent rCDCs respectively; \( p<0.001 \)) (Fig 3A); suspension in PBS resulted in an additional decrease in \( ^{99m}\)Tc-pertechnetate uptake by 43% (6.23±3.14 × 10\(^3\) counts/mg protein; \( p<0.001 \) vs. rCDCs suspended in IMDM, Fig 3A). Since \( h \)NIS is a surface protein that may be degraded by trypsin, we measured \( ^{99m}\)Tc-pertechnetate uptake in suspended \( h \)NIS\(^+\) rCDCs dissociated using trypsin or non-enzymatic dissociation solution that contains EDTA, but no enzymes. We found that \( ^{99m}\)Tc-pertechnetate in suspended rCDCs dissociated using trypsin was similar to that obtained by non-enzymatic dissociation (1.08±0.14 × 10\(^4\) vs. 1.07±0.14 × 10\(^4\) counts/mg protein, \( p=0.45 \)), indicating that
trypsin-mediated NIS protein degradation does not underlie the decreased $^{99m}$Tc-pertechnetate uptake observed in NIS$^+$ rCDCs following suspension (Fig 3B).

In vitro BLI revealed a 44% reduction in signal (compared to adherent rCDCs) following suspension of fluc$^+$ rCDCs for 1hr in saline/PBS containing glucose, Ca$^{2+}$ and Mg$^{2+}$ ($1.35\pm0.008 \times 10^4$ vs. $2.41\pm0.16 \times 10^4$ RLU/mg protein in suspended and adherent cells respectively; $p<0.01$), and further reduction by 95% when suspended in PBS/saline ($6.21\pm0.23 \times 10^2$ RLU/mg protein, $p<0.01$) (Fig 3C).

**In vivo studies.**

Since cell dissociation/suspension decreased $^{99m}$Tc uptake and BLI in vitro, we investigated the effect of cell dissociation/suspension on in vivo $^{99m}$Tc uptake and BLI, acutely following cell transplantation, when the transplanted cells are likely to still be in suspension, and at 24hrs following transplantation, when live cells are expected to have attached to the cardiac extracellular matrix, and non-adherent cells have undergone anoikis (and hence are unable to generate a molecular imaging signal).

The in vivo BLI signal was higher at 1hr than at 24hrs. Importantly, at both time points, the in vivo BLI signal was lower when rCDCs were suspended in PBS/saline, when compared to the glucose-containing medium IMDM ($4.09 \pm 1.73 \times 10^5$ vs. $1.05 \pm 0.34 \times 10^6$ RLU at 1hr for PBS and IMDM respectively, $p<0.05$; and $5.04 \pm 2.40 \times 10^4$ vs. $1.73 \pm 0.37 \times 10^5$ RLU for PBS and IMDM respectively at 24hrs; $p<0.05$) (Fig 4 A, B, C), indicating that presence of substrates in the cell suspension medium improves the sensitivity of in vivo BLI.

NIS$^+$ rCDCs were suspended in IMDM to optimize their metabolism in suspension. In vivo dual isotope SPECT-CT imaging of intra-myocardially-injected NIS$^+$ rCDCs suspended in IMDM revealed that NIS$^+$ rCDCs were clearly visible 24hrs post-transplantation, but not 1hr post-transplantation in infarct and non-infarct models of cell transplantation (Fig 5A); perfusion imaging confirmed good perfusion at the site of cell injection in the non-infarcted animals (Fig 5B) and ex vivo SPECT imaging confirmed the presence of cells at the 1hr time point (Fig 5C). Calibration factor was calculated to be $3.37\times10^{-5}$ MBq/i.i. from the dose response plot (Fig 5D). Signal quantification revealed that the in vivo $^{99m}$Tc-pertechnetate uptake ratio was significantly greater at 24hrs, compared to 1hr in both non-infarct ($0.05\pm0.01$MBq at 1hr vs. $0.23\pm0.05$MBq at 24 hrs) and infarct ($0.09\pm0.02$MBq at 1hr vs. $0.33\pm0.12$MBq at 24 hrs) rat models ($p<0.005$; Fig 5E). Similarly, the ex-vivo $^{99m}$Tc-pertechnetate uptake ratio was higher at 24hrs ($0.01$MBq at 1hr vs. $0.12\pm0.001$MBq at 24 hrs).

Longitudinal, in vivo SPECT/CT imaging of NIS$^+$ rCDCs and BLI of fluc$^+$ rCDCs over 7days.
revealed progressive cell loss from day 1 to 7 post-transplantation (Supplemental Fig I). Based on our in vitro studies which reveal that cellular metabolism is stable after adhesion for 24 hrs (Supplemental Fig II), we believe that factors other than cell adhesion play a role in cell loss at later time points following transplantation.

**Confirmation of Engraftment.**

Ex vivo luciferase assay (n=12) using fluc’rCDCs, and quantitative PCR for the male-specific SRY gene using NIS’ rCDCs (n=8) in separate sets of animals revealed a significant reduction in cell number (-72.39%±14.10% for fluc’rCDCs and -74.77%±13.13% for NIS’rCDCs) between 1 hr and 24 hrs post-transplantation (Fig 6A, B), confirming similar amounts of in vivo cell loss in both groups in the first 24 hrs.

**DISCUSSION**

This is the first study to report a link between stem cell adhesion, bioenergetics and the molecular imaging signal. The novel results of this study are as follows: 1) CDCs exhibit the Warburg effect (aerobic glycolysis); 2) cell adhesion is an important regulator of cellular metabolism and the molecular imaging signal derived from hNIS and luciferase; 3) in vivo ⁹⁹ᵐTc-pertechnetate uptake is higher at 24 hrs following transplantation of cell suspensions (when compared to 1 hr), despite fewer cells at this time point.

**NIS imaging.**

Reporter gene strategies are useful in the longitudinal assessment of engraftment because transplanted cell viability is required for gene expression.¹ We used a self-inactivating, 3rd generation lentivirus and the constitutively active cytomegalovirus virus (CMV) promoter which results in integration of the expression cassette into the host genome and high levels of transgene expression. Previous studies by our group have shown that expression of firefly luciferase or the human NIS gene in CDCs at an MOI of 20, did not affect CDC viability or function.⁴, ²¹ NIS promotes cellular uptake of iodide or ⁹⁹ᵐTc-pertechnetate and Na⁺, driven by the trans-membrane sodium gradient,¹⁴ which is generated by the Na⁺-K⁺- ATPase. Remarkably, after ectopic NIS expression, only cells over-expressing NIS will transport ⁹⁹ᵐTc-pertechnetate or iodine-124 (¹²⁴I) after intravenous injection of these tracers, permitting noninvasive, longitudinal monitoring of stem cell engraftment by single-photon emission computed tomography (SPECT) and positron emission tomography (PET), respectively.⁴, ²⁵, ²⁶
In vivo $^{99m}$Tc-pertechnetate uptake was low at 1hr following cell transplantation and increased at 24hrs, despite significant cell loss during this period. This finding is not due to impaired perfusion at the injection site at 1hr or due to cell proliferation, based on our perfusion results in non-infarcted animals at 1hr following transplantation (Fig 5B) and quantitative PCR for the SRY gene using NIS+ CDCs (Fig 6). We attribute this result to impairment of bioenergetics in suspended CDCs based on our in vitro studies in adherent and suspended cells.

For in vivo studies, hNIS+ rCDCs were dissociated using trypsin, suspended in IMDM and injected intra-myocardially. We used IMDM rather than PBS/saline as the vehicle for our in vivo experiments because in vitro studies revealed that $^{99m}$Tc-pertechnetate uptake was higher when cells are suspended in IMDM, when compared to PBS/saline (Fig 3A). Our in vitro studies also confirmed that $^{99m}$Tc-pertechnetate uptake was similar in CDCs dissociated using trypsin or non-enzymatic dissociation solution (Fig 3B), indicating that impairment of cellular bioenergetics, rather than trypsin-mediated degradation of NIS is the dominant mechanism underlying this phenomenon.

Cell dissociation and suspension resulted in significant down-regulation of glucose uptake, cellular metabolism and ATP levels (Fig 2A, B). Our in vitro studies also indicate a progressive increase in the cellular ADP/ATP ratio from 1hr to 3hrs in suspension, which would result in a lower Gibbs free energy availability from ATP hydrolysis, limiting Na$^+$-K$^+$ ATPase function, that in turn reduces transmembrane Na$^+$ gradient and consequently $^{99m}$Tc-pertechnetate uptake – this could explain the very low in vivo $^{99m}$Tc-uptake in the immediate post-transplantation period (the time interval between cell dissociation and completion of in vivo SPECT/CT (NIS) imaging was ~2-3hrs). Transplanted cells would be expected to adhere to the cardiac extracellular matrix (ECM) over the 24hr period post-transplantation, resulting in improved bioenergetics which would translate into improved $^{99m}$Tc-pertechnetate uptake- in vitro studies confirmed increased $^{99m}$Tc-pertechnetate uptake in NIS+CDCs, cultured on adhesive surfaces for 18-24hrs, when compared to suspended cells. Cells that do not attach in vivo would undergo anoikis, which would explain some of the cell loss in the first 24hrs. Other mechanisms underlying cell loss could be progressive cell egress from the transplantation site via lymphatics and coronary veins (an effect that is more pronounced when cells are injected into contractile, non-infarcted myocardium) and cell death due to oxidative stress, inflammation or other mechanisms. Increase in the in vivo SPECT signal despite significant cell loss probably occurred because the magnitude of the increase in $^{99m}$Tc-pertechnetate uptake of CDCs surviving at 24hrs greatly exceeded the effects of cell loss over the same time period.

Bioluminescence imaging.

BLI-fluc imaging is based on the oxidation of the substrate D-luciferin by luciferase, a reaction that requires oxygen, magnesium, and ATP, and results in a red-shifted light emission (wavelength: 500–
700 nm), which can be detected using a camera. Since the luciferase reaction is directly dependent on ATP and Mg\(^{2+}\), composition of the cell suspension solution had a profound influence on the BLI signal. We found that the BLI signal generated by cells suspended in IMDM was higher than when suspended in PBS/saline at 1hr and 24hrs following transplantation. These differences could be related to differences in the \textit{in vivo} cell survival and/or metabolism between the 2 groups. Further studies are needed to test whether cell suspension in solutions that contain substrates, Mg\(^{2+}\) and Ca\(^{2+}\) are superior to PBS/saline with respect to cell engraftment.

As in the case of \(^{99m}\)Tc-pertechnetate uptake, the \textit{in vitro} BLI signal was also higher in \textit{fluc}+ rCDCs that were adherent for 18-24hrs, compared to suspended cells in our \textit{in vitro} studies. Hence, transplanted \textit{fluc}− rCDCs would also be expected to attach to cardiac ECM and improve cellular bioenergetics and light generation by luciferase over the first 24hrs following transplantation. However, the BLI signal was higher at 1hr, compared to 24hrs post-transplantation. This discrepancy in the \textit{in vivo} molecular imaging signal at 24hrs, between \textit{in vivo} SPECT and BLI is not due to differences in the \textit{in vivo} survival of \textit{fluc}− CDCs and NIS− CDCs because cell loss in the first 24hrs was similar in the 2 groups (Fig 6A). We attribute the different results at 24hrs, using NIS and luciferase, to the dependence on O\(_2\) for generation of the luciferase signal, but not the NIS signal. Our \textit{in vitro} metabolism studies reveal that stem cells rely primarily on glycolysis for ATP generation, a phenomenon referred to as the Warburg effect. Furthermore, inhibition of the mitochondrial \(F_1-F_0\) ATP synthase did not affect ATP levels (Fig 2F), in contrast to cell dissociation which rapidly depressed ATP levels (Fig 2B). Since signal derived from NIS relies on the trans-membrane Na\(^+\) gradient, which is dependent on cellular ATP levels, it would be expected to increase with cell adhesion at 24hrs, following \textit{in vivo} cell transplantation. However, the luciferase reaction is dependent on O\(_2\) as well as ATP. Low O\(_2\) tension at the transplantation site due to tissue damage from injection combined with cell loss during this interval would be expected to impair recovery of the BLI signal but not the NIS signal at 24hrs.

### Future implications for stem cell imaging.

We chose 1hr as the first time point for our \textit{in vitro} and \textit{in vivo} studies because 1hr post-transplantation is often chosen as the baseline time point for longitudinal assessment of engraftment in experimental models of cell transplantation. We picked 24hrs as the 2\(^{nd}\) time point, because it is unclear from the literature whether the signal at 1hr or 24hrs should be used as the baseline signal for longitudinal follow up of engraftment using molecular imaging.

With \textit{in vivo} BLI, the BLI signal was higher at 1hr, compared to 24hrs post-transplantation and hence, more accurately represented cell fate in the first 24hrs following transplantation. However, the NIS-derived signal, using suspended cells was higher at 24hrs when compared to 1hr following cell
transplantation despite significant cell loss. This result has important implications if NIS is used for longitudinal assessment of engraftment. Based on our results, the NIS-derived signal immediately post-transplantation, using suspended cells, is more reflective of cellular bioenergetics than actual cell number, when compared to the signal at 24hrs. Hence, it could confound quantification of engraftment in the first 24hrs, using SPECT-CT imaging of NIS⁺ rCDCs. This phenomenon may be harnessed to monitor the effects of small molecules and tissue engineered scaffolds on transplanted cell bioenergetics in the immediate post-transplantation period.

**Study limitations.**

This is the first report of impaired bioenergetics by cell dissociation/suspension- further in depth studies are needed to understand the signaling pathways underpinning this phenomenon. Tissue engineered matrices containing Arg-Gly-Asp (RGD) motifs that promote encapsulated cell attachment to the matrix, or cell-cell adhesion by generation of cell aggregates may abrogate the down-regulation of cellular metabolism in stem cell suspensions, and need to be tested.

We limited this study to rat CDCs because we have extensive experience with stem cell isolation, gene transduction and molecular imaging using this cell type. However, we believe that these results are not specific to CDCs and will be reproducible in other adherent cell types, e.g. mesenchymal stem cells (MSCs), embryonic stem cells and endothelial progenitor cells. Unfortunately, we were unable to transduce commercially available human MSCs using this 3rd generation lentivirus and hence were unable to test this clinically important cell type.

Technical limitations precluded cell sorting to exclude dead cells, in cell suspensions, prior to in vitro experiments at 1hr and measurement of transplanted cell bioenergetics in vivo, to verify our in vitro results. However, using flow cytometry, we consistently observe an ~15% reduction in cell viability which is much less than the reduction of cellular ATP levels, respiratory measurements, ⁹⁹ᵐาะTc uptake and in vitro BLI, indicating that cell suspension rather than decrease in cell viability is the primary cause of the observed depression of cellular bioenergetics.

**CONCLUSION**

Cell dissociation impairs cellular bioenergetics, resulting in reduced ⁹⁹ᵐาะTc-pertechnetate uptake and BLI signal. Since cell survival, proliferation and function are intimately linked to metabolism, BLI and NIS imaging could be useful tools for in vivo optimization of bioenergetics and engraftment of transplanted cells.
ACKNOWLEDGEMENTS
We are grateful to Jianhua Yu, Catherine Foss, PhD and Miguel Aon, PhD for helpful advice, James Fox and Gilbert Green for technical assistance.

SOURCES OF FUNDING
This study was funded by AHA-BGIA and NIH RO1 HL092985. Dr. Angel Chan was supported by NIH T32HL07227 Training Grant. Dr. Xiaoping Lin was partially supported by the China Scholarship Council. Dr. Styliani Vakrou was supported by a grant from the Hellenic Society of Cardiology. Kirubel Woldemichael was supported by an NIH diversity fellowship.

DISCLOSURES
None

REFERENCES


FIGURE LEGENDS

**Fig 1. Cell viability following suspension.** (A) CDC viability: representative flow plot and bargraph (B) immediately following cell dissociation reveals high CDC viability (97.5 ± 1.1; n=7). (C) Cell viability is reduced following suspension for 1hr and 6hrs (n=3)

**Fig 2. Cell dissociation impairs bioenergetics.** (A) 18FDG uptake and (B) ATP levels in 18-24hr adherent and 1hr suspended CDCs following dissociation using trypsin(C) OCR and ECAR are significantly reduced in suspended CDCs (D) Re-plating after suspension for 1hr and 6hrs results in restoration of cellular metabolism, when assessed at 24hrs (n=6) (E) Oligomycin reduces OCR without affecting ATP levels in adherent CDCs (n=3) (F) Iodoacetate but not Oligomycin reduced ATP levels in adherent CDCs, indicating dependence on glycolysis for ATP generation (n=6)

**Fig 3. In vitro 99mTc-pertechnetate uptake and BLI.** (A) 99mTc-pertechnetate uptake is reduced by cell dissociation/suspension and inhibited by perchlorate (B) 99mTc-pertechnetate uptake was similar in suspended cells dissociated using trypsin and non-enzymatic dissociation solution (C) In vitro BLI signal is significantly higher in adherent cells, when compared to suspended cells. Suspension in PBS containing Ca²⁺, Mg²⁺, glucose results in a significantly higher signal than suspension in PBS.

**Fig 4. Bioluminescence imaging (BLI) following intra-myocardial transplantation of fLuc⁺ CDCs.** Representative in vivo BLI images in rats reveal higher signal when CDCs are suspended in IMDM medium (A) when compared to PBS (B) at 1hr and 24hrs following cell transplantation (AP view). (C) Bar graphs summarizing in vivo BLI results at 1hr and 24hrs following transplantation reveal that signal is significantly higher at 1hr, when compared to 24hrs. Suspension in IMDM medium results in a significantly higher signal at both time points

**Fig 5. In vivo dual isotope SPECT/CT imaging following intra-myocardial transplantation of NIS⁺ CDCs.** (A) Representative coronal images of in vivo imaging (perfusion/²⁰¹Tl in red and ⁹⁹mTc-pertechnetate in green), 1hr and 24hrs following cell transplantation into normal myocardium reveals distinct ⁹⁹mTc-pertechnetate uptake signal from transplanted cells at 24hrs, but not at 1hr. (B) Corresponding polar plot of ²⁰¹Tl perfusion reveals normal perfusion 1hr post-transplantation into normal myocardium. (C) Representative SPECT/CT co-registered image of explanted rat heart at 1hr following transplantation reveals an area of ⁹⁹mTc-pertechnetate uptake (indicated by arrow). (D) Dose response plot for ⁹⁹mTc was obtained by dual isotope SPECT scanning, using a rat size phantom containing several doses of ⁹⁹mTc-pertechnetate and ²⁰¹Tl. (E) Summary of in vivo ⁹⁹mTc-pertechnetate uptake in infarct and non-infarct animals, imaged at 1hr and 24hrs following cell transplantation confirms

DOI: 10/1161/CIRCRESAHA/112/273375 19
significantly increased signal at 24hrs (0.07±0.03MBq at 1hr vs. 0.28±0.18MBq at 24h). Cell-derived signal at 1hr is similar to the background signal from myocardium (0.05±0.02MBq).

**Fig 6.** Confirmation of engraftment following intra-myocardial transplantation of \( f\text{Luc}^+ \) and \( \text{NIS}^+ \) rCDCs. (A) Similar amounts of cell loss were observed between 1hr and 24hrs following intra-myocardial transplantation of \( f\text{Luc}^+ \) and \( \text{NIS}^+ \) rCDCs \( (P<0.005) \). (B) Standard curve correlating ex vivo luciferase activity in rat heart homogenates with known numbers of \( f\text{Luc}^- \) rCDCs.
Novelty and Significance

What Is Known?

- Stem cell engraftment in the heart is low and results in small functional benefit.
- Molecular imaging is useful to study in vivo stem cell biology and optimize engraftment.
- Cell survival, proliferation and function are intimately linked to metabolism.

What New Information Does This Contribute?

- Stem cell dissociation and suspension impairs cellular bioenergetics, resulting in reduction of the molecular imaging signal obtained by Single photon emission computer tomography (SPECT) and bioluminescence imaging (BLI) of cardiosphere-derived cells (CDCs) labeled with the sodium-iodide symporter (NIS) and luciferase, respectively.
- Impairment of cellular bioenergetics in dissociated NIS+ cells can confound quantification of engraftment by in vivo SPECT imaging in the first 24h after transplantation.
- SPECT imaging of NIS-labeled cells may be useful for in vivo optimization of bioenergetics and engraftment in transplanted stem cells.

Cell transplantation in the heart is limited by low engraftment. We used in vivo molecular imaging by SPECT and BLI to examine engraftment of CDCs in the first 24h after transplant. Using a combination of in vitro studies of cell viability, metabolism and function, and in vivo longitudinal imaging, we demonstrate a link between CDC dissociation/suspension, bioenergetics and the molecular imaging signal obtained by NIS (SPECT imaging) and firefly luciferase labeling (BLI) of transplanted cells. Specifically, we found that cell dissociation/suspension reversibly impairs cellular metabolism and ATP levels, resulting in decreased $^{99m}$Tc-pertechnetate uptake and BLI signal in suspended cells, when compared with adherent cells. This translated into increased $^{99m}$Tc-pertechnetate uptake (SPECT signal) at 24h, compared with 1h post-transplantation, despite considerable cell loss during this interval. However, the BLI signal decreased between 1h and 24h post-transplantation, and was reflective the cell loss during this period. Because the BLI signal depends on ATP, Mg$^{2+}$ and O$_2$, while the NIS-derived signal depends only on ATP, we propose that SPECT imaging of NIS+ cells may be useful for in vivo optimization of cellular bioenergetics and engraftment in small/large animal models and humans.
Figure 1

Panel A: Flow cytometry graph showing Annexin V-FITC and Propidium Iodide staining.

Panel B: Bar graph showing % viability compared to adherent cells for IMDM 1Hour and IMDM 6Hour conditions.
**Figure 3**

**A**

- Adherent Suspension in Medium
- Suspension in PBS

Tc-99m Uptake ($10^3$ Cts/mg Protein)

- No drugs
- + PERCHLORATE

$p<0.001$

**B**

- Trypsin
- Non-enzymatic dissociation

Tc-99m Uptake ($10^3$ Cts/mg protein)

- $P=0.45$

**C**

- Adherent Suspension in PBS with Ca/Mg/Glucose
- Suspension in PBS

Bioluminescence Activity ($10^3$ RLU/mg protein)

- $N=4$
- $N=3$
- $N=4$

$p<0.001$

$p<0.001$
Figure 4
Figure 5

Coronal view

Polar map

Figure 5

Transverse view
Figure 6
Cellular Bioenergetics is an Important Determinant of the Molecular Imaging Signal Derived from Luciferase and the Sodium-Iodide Symporter

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*Circ Res.* published online December 19, 2012;
*Circulation Research* is published by the American Heart Association, 7272 Greenville Avenue, Dallas, TX 75231
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Print ISSN: 0009-7330. Online ISSN: 1524-4571

The online version of this article, along with updated information and services, is located on the World Wide Web at:
http://circres.ahajournals.org/content/early/2012/12/11/CIRCRESAHA.112.273375

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**Supplemental Material**

**Detailed Methods**

**Cell isolation and culture**
Briefly, small pieces of myocardial tissue (explants) derived from male rats were placed on fibronectin-coated dishes. In the following days, cells exited the explants and formed an adherent monolayer on the dish surface with phase bright cells on top. These cells are harvested using mild enzymatic digestion and transferred to D-poly-lysine coated dishes, where they form three dimensional structures called cardiospheres which are enriched in cardiac progenitors. Cardiospheres are subsequently harvested and grown as monolayers in fibronectin-coated flasks – these cells are called cardiosphere-derived cells (CDCs). CDCs were cultured in IMDM medium (Invitrogen) containing 10% FBS, 10% glutamine and 0.1mM mercaptoethanol, and expanded to 3-5 passages prior to lentiviral transduction.

**Lentivirus synthesis**
The cDNA encoding the hNIS gene or the cDNA pGL4.10[luc2] encoding firefly luciferase (Promega, Madison, WI, USA) was sub-cloned in place of eGFP into the vector RRLsin18.cPPT.CMV.eGFP.Wpre, resulting in plasmids designated cpPPT.CMV.hNIS or pPPT.CMV.fLuc. Viral vectors were produced by Lipofectamine 2000 (Invitrogen) transfection of 4 lentiviral vector plasmids into HEK293T cells (ATCC, Manassas, VA, USA). Vector-containing supernatant was collected 48 and 72 hours after transfection, filtered, and concentrated (Centricon Plus-70, Millipore, Billerica, MA, USA). Viral titer was assigned on concentrated supernatant by HIV-1 p24 ELISA (Dupont, Wilmington, DE, USA). For genetic labeling, rCDCs were transduced at a multiplicity of infection of 20 yielding transduction efficiencies of >70% for hNIS expression and >90% for luc expression. NIS expression was confirmed by immunostaining using a monoclonal mouse anti-hNIS antibody (Abcam, Cambridge, MA, USA) and by in-vitro $^{99m}$Tc-pertechnetate uptake, while luciferase expression was examined by immunostaining using a polyclonal goat anti-luciferase antibody (Promega) and by an in vitro bioluminescence assay. We have previously demonstrated that transduction of CDCs with firefly luciferase or hNIS at an MOI (multiplicity of infection) of 20 does not affect cell proliferation, using in vitro studies. $^{1,2}$ Percentage of transduced cells was calculated by immunostaining prior to cell transplantation.

**Flow Cytometry**
Annexin V and Propidium iodide were used to identify apoptotic and dead cells, respectively. Annexin V was diluted at a concentration of 1 mg/ml in binding buffer and cells ($1 \times 10^6$ cells) were re-suspended in 1ml of this freshly made solution. Cells were incubated for 10 min in the dark at room temperature then PI solution (0.1ml to give final conc. of 1mg/ml) was added to this solution 5 minutes prior to the analysis with flow cytometry equipment (with 10,000 events collected per sample). Cell debris was excluded by scatter gating and 10,000 gated events were collected per sample using BD Accuri C6 flow cytometer. For measurements of cell viability after dissociation, CDCs were trypsinized, counted and used immediately after suspension in media. Trypan blue measurements were compared to measurements by flow cytometry. For measurements of cell viability following suspension for 1hr and 6hrs, CDCs were trypsinized, counted and suspended in cell culture medium for 1hrs and 6hrs respectively, in the incubator at 37ºC prior to measurements by flow cytometry.

**In vivo SPECT/CT imaging**
Male WKY rats underwent left thoracotomy in the 4th or 5th intercostal space under general anesthesia (isoflurane inhalation, 4% for induction and 2% for maintenance). In the myocardial infarction group, the heart was exposed and the left anterior descending coronary artery was
ligated using a 5-0 silk suture. Three million NIS\(^+\) rCDCs suspended in 100\(\mu\)l of IMDM (Invitrogen) were injected directly into the myocardium at three sites in the anterior wall of the left ventricle using a 30G needle. In the non-infarct group, the same procedure was followed as the myocardial infarction group with the exception of the ligation of the left anterior descending coronary artery. Subsequently, the chest was closed with a 3-0 silk suture. \(^{99}\text{TcO}_4^-\) (\(^{99}\text{Tc}\) labeled technetium-pertechnetate; 555-740 MBq) and \(^{201}\text{TlCl}\) (\(^{201}\text{Tl}\) labeled thallous chloride; 37-74 MBq) were injected intravenously via the tail vein immediately after intra-myocardial cell transplantation, to determine stem cell retention and myocardial perfusion. The isoflurane was turned off and the animal was monitored for spontaneous breathing and allowed to fully recover prior to imaging.

*In vivo* dual isotope SPECT imaging was performed 1hr after injection of \(^{99}\text{Tc}\)-pertechnetate and \(^{201}\text{TlCl}\). CT imaging was performed prior to SPECT imaging. Both scans were performed on a small animal SPECT/CT system (X-SPECT-CT from Gamma Medica Inc., Northridge, CA) using inhalational isoflurane as the anesthetic agent, administered via a nose cone. Animals were allowed to recover in their cages after completion of imaging on day 0. After 24hrs, the same rats were re-injected with \(^{99}\text{Tc}\)-pertechnetate (555-740 MBq) and \(^{201}\text{TlCl}\) (37-74 MBq) via the tail vein and *in vivo* dual isotope SPECT-CT imaging was performed. The rats were euthanized after completing the 24hr imaging protocol.

In a second study, rats were serially imaged at 1d, 3-4d and 7d following intra-myocardial transplantation of NIS\(^+\) or \(\text{fluc}\(^+\)\) rCDCs.

**SPECT/CT image acquisition and processing:** The SPECT module X-SPECT-CT system is composed of two gamma camera heads each consisting of pixelated NaI(Tl) with a total area of 125 mm \(\times\) 125 mm, divided into 80 \(\times\) 80 number of pixels with 1.56 mm pitch. Low-energy knife-edge pinhole collimators were used with a pinhole aperture of 1 mm diameter and a focal length of 9cm; a radius-of-rotation of 5.42 cm was used. Each camera head acquired 128 projections over a 180-degree range, with an acquisition time of 30s for each projection for all scans except for the ex vivo heart scans which was changed to 40s per projection.

In the dual isotope SPECT imaging, data were acquired in listmode and were subsequently re-binned into two energy windows (“75 keV \(+10\%/-10\%\)” and “140 keV \(+10\%/-10\%\)”) to obtain separate sets of \(^{201}\text{Tl}\) and \(^{99}\text{Tc}\) projections. The \(^{99}\text{Tc}\) and \(^{201}\text{Tl}\) projection datasets were reconstructed using a 3D pinhole ordered-subset expectation-maximization (OS-EM) imaging reconstruction algorithm with 8 and 4 updates, respectively with an isotropic reconstructed image voxel size of 0.7 mm.

X-ray computed tomography (CT) was performed on the microCT module with an X-ray tube voltage of 75 kVp. A total of 512 projections were acquired over a 360-degree range. The projections with 1,184 \(\times\) 1,120 isotropic pixels (100 \(\mu\)m) were reconstructed into a CT volume of 512\(^3\) isotropic voxels with 170\(\mu\)m pixel size. The SPECT and CT were then registered using rigid body transform, with pre-set parameters specific to the system.

**SPECT image quantification:** For absolute quantification, a calibration factor (CF) was calculated from an experimental study by inserting a small hollow sphere filled with a known amount of radioactivity of \(^{99}\text{Tc}\) or \(^{201}\text{Tl}\) in water in an average rat-size water-filled cylindrical phantom to simulate a rat scan. Dose-response plot for \(^{99}\text{Tc}\)-pertechnetate was obtained by dual isotope SPECT imaging of varying doses of \(^{99}\text{Tc}\)-pertechnetate and \(^{201}\text{Tl}\) and then used to calculate the calibration factor. SPECT data for the phantom were acquired using exactly the same acquisition settings as those used in the animal experiments. In this case, CF (MBq/i.i.) was defined as the quotient of the known activity concentration (MBq/ml) within the radioactive sphere in the phantom divided by the measured mean image intensity (i.i/cm\(^3\)) within a ROI drawn over the small sphere in the SPECT image of the phantom.
To quantify the tracer uptake in vivo, regions-of-interest (ROI) were manually defined on a region of increased focal tracer uptake and on a contra-lateral normal region of a mid-myocardial section. In the case of no observable increased focal myocardial tracer accumulation, an ROI was placed on the distal anterior wall. The total radioactivity at the region of interest was calculated by the image intensity within the ROI multiplied by the CF. The radioactivity concentration (MBq/mL) within the ROI was calculated by the total activity divided by the volume of the ROI. The background activity was calculated by placing an ROI on the baso-lateral wall of the heart.

Myocardial perfusion polar maps were generated by combining the short-axis image slices through the left ventricular myocardium of the $^{201}$TI SPECT dataset. A perfusion defect was extracted from a threshold of 60% of the maximum intensity of the myocardium.

**Ex-Vivo SPECT imaging**

The same procedure was adopted as for the in vivo imaging group, except that only $^{99m}$Tc-pertechnetate was injected. For the 1hr ex-vivo group, the rats were injected with $^{99m}$Tc-pertechnetate (555-740 MBq) immediately after injection of the NIS$^+$ rCDCs and imaged 1hr later. Immediately after the SPECT scan, the rat was sacrificed and the heart was rinsed with PBS and washed thoroughly to remove any remaining blood before ex vivo scanning. Imaging parameters were identical to the ones used in the in vivo acquisitions, with the exception of the time per projection which was decreased to 40s per projection. For the second group of rats, the same procedure was performed as the 1hr ex vivo rats with the exception that the rats were injected with $^{99m}$Tc-pertechnetate (555-740 MBq), imaged and sacrificed 24hrs following cell transplantation.

**Quantification of engraftment by ex vivo luciferase assay:** We performed the ex vivo luciferase assay in a separate set of animals, to quantify engraftment of fluc$^+$ rCDCs in the first 24hrs following transplantation. This assay has high sensitivity and is not affected by conditions such as ischemia/hypoxia, cellular metabolism and diffusion of luciferin to the cell transplantation site, which can all affect the in vivo signal, thus permitting reliable quantification of engraftment. For this purpose, we performed intra-myocardial injection of 1million fluc$^+$ rCDCs in 12 animals (6 animals without myocardial infarction and 6 animals with infarction). Three animals from each group were sacrificed at 1hr and 24hrs after cell transplantation. A standard curve was constructed to calculate cell number.

Hearts were harvested and cut into 200mg pieces, from apex to base (4-5 pieces/heart). The tissue samples were mixed with 1ml of lysis buffer (Promega) and 0.5ml of 10% BSA (as a non-specific protease inhibitor). Tissue samples were homogenized by manually grinding (Duall 24 Glass homogenizer, Kontes, Vineland, NJ, USA), the homogenates were collected in 50ml conical tubes (VWR, West Chester, PA, USA) and centrifuged at 25,000 g for 45min, at 4°C. The supernatants were collected and a luciferase assay was performed using a Monolight 2010 luminometer (Analytical Luminescence Laboratories, USA). Twenty microliters of each sample was mixed with 100µl of luciferase assay reagent (Promega) in 75mm glass tubes (VWR) and placed in the instrument (2s measurements). Results were reported as relative light units (RLUs) and converted to cell numbers using the regression equation of the corresponding standard curve.

**Standard curve preparation:** Luciferase over-expressing rCDCS (from the same isolates that were used for the in vivo experiments) were also used for the standard curve preparation. Different numbers of rCDCs (starting from 500 up to 100,000 in duplicates) were pelleted. Hearts from normal male WKY rats were harvested and cut into small pieces. Cell pellets were lysed (Luciferase lysis buffer, Promega) and 200mg of rat heart tissue was added to the cell lysate. Cell lysates and tissue samples were processed as described above and values
obtained by the luciferase assay were used to construct the standard curve. Luciferase activity demonstrated excellent linear correlation to cell numbers.

**Quantification of engraftment by quantitative polymerase chain reaction (qPCR):**
Genomic DNA was isolated from aliquots of the homogenate corresponding to 12.5mg of myocardial tissue, according to the manufacturer's instructions (Qiagen). Real time PCR was performed using the TaqMan® chemistry (Applied Biosystems), with the rat SRY gene as target (forward primer: 5'-GGA GAG AGG CAC AAG TTG GC-3', reverse primer: 5'-TCC CAG CTT GCT GAT C-3', TaqMan probe: 6FAM CAA CAG AAT CCC AGC ATG AAT TCA G TAMRA). For absolute quantification of gene copy number, a standard curve was constructed with samples derived from multiple log dilutions of genomic DNA isolated from male rat CDCs. All samples were spiked with 50ng of female genomic DNA to control for any effects this may have on reaction efficiency in the actual samples. The copy number of the SRY gene at each point of the standard curve is calculated based on the amount of DNA in each sample and total mass of the rat genome per diploid cell. (http:www.cbs.dtu.dk/databases/DOGS/index.html). All samples were tested in triplicate. The qPCR assay was repeated twice with DNA samples isolated on 2 occasions, from each heart. For each reaction, 50ng of template DNA was used. Real-time PCR was performed in an ABI PRISM 7700 instrument. The result from each reaction, i.e. copies of the SRY gene in 50ng of genomic DNA, was expressed as the number of engrafted cells/heart, by first calculating the copy number of the SRY gene in the total amount of DNA corresponding to 12.5mg of myocardium and then extrapolating to the total weight of each heart.

**Supplemental Figures and Figure Legends**

**A**

Supplemental Figure 1: Longitudinal in vivo SPECT/CT imaging of NIS⁺ rCDCs (A) and BLI of fluc⁺ rCDCs (B) over 7 days revealed progressive cell loss from day 1 to 7 post-transplantation (n=3).
Supplemental Figure 2: In vitro studies reveal that cellular metabolism is stable after adhesion for 24hrs.
Supplemental Figure 3: In vitro studies: Ouabain (100 µM), a blocker of Na⁺-K⁺-ATPase reduced ⁹⁹mTc-pertechnetate uptake by ~17% in adherent CDCs (p<0.01) and in CDCs suspended in glucose-containing media (p=0.01), but had no effect when CDCs were suspended in PBS (p=0.76), confirming a link between cellular Na⁺-K⁺-ATPase activity and NIS-mediated ⁹⁹mTc-pertechnetate uptake.

Supplemental References
