Serial Noninvasive In Vivo Positron Emission Tomographic Tracking of Percutaneously Intramyocardially Injected Autologous Porcine Mesenchymal Stem Cells Modified for Transgene Reporter Gene Expression

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Background—Porcine bone marrow-derived mesenchymal stem cells (MSCs) were stably transfected with a lentiviral vector for transgene expression of the trifusion protein renilla luciferase, red fluorescent protein and herpes simplex truncated thymidine kinase (LV-RL-RFP-tTK; positron emission tomography [PET] reporter gene) for in vivo noninvasive tracking of the intramyocardially delivered MSC fate.

Methods and Results—A closed-chest, reperfused myocardial infarction was created in farm pigs. Sixteen days after myocardial infarction, LV-RL-RFP-tTK-MSCs were injected intramyocardially using electromechanical mapping guidance in the infarct border zone (n=110057). PET-computed tomographic metabolic and perfusion imaging was performed after an intravenous injection of 10 mCi [18F]-FHBG and 13N-ammonia PET at 30±2 hours and 7 days after LV-RL-RFP-tTK-MSC treatment. Fusion imaging of the [18F]-FHBG PET-computed tomography with MRI was used to determine the myocardial location of the injected LV-RL-RFP-tTK-MSCs. Seven days after injections, [18F]-FHBG PET showed a decreased cardiac uptake with a mild increased pericardial and pleura uptake in the treated animals, which was confirmed by the measurement of luciferase activity. At 10 days, infarct size by MRI in the LV-RL-RFP-tTK-MSC-treated animals was smaller than controls (n=7) (23.3±1.5% versus 30.2±3.5%, P<0.005). The presence of the LV-RL-RFP-tTK-MSCs (5.8±1.1% of the injected cells) in the myocardium 10 days after intramyocardial delivery was confirmed histologically.

Conclusions—Reporter gene imaging enables the tracking of the persistence of viable LV-RL-RFP-tTK-MSC in the peri-infarcted porcine myocardium at 10 days after delivery using clinical PET scanners. (Circ Cardiovasc Imaging. 2008;1:94-103.)

Key Words: cells ■ imaging ■ infarction ■ mapping ■ revascularization

Cardiac transplantation of stem cells (SCs) has been shown to improve regional perfusion and systolic function of the failing heart after myocardial infarction (MI).1,2 However, once delivered to the heart, unlabeled cells cannot be visualized or tracked in vivo. Although iron oxide-labeled SCs can be detected by MRI, this method is insensitive to a small number of cells, is hindered by the label dilution because of cell division and migration, and cannot distinguish live from dead cells.3–5

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The reporter gene approach is another method to track SC fate noninvasively.6 Because the reporters are expressed only in living cells and are passed to daughter cells on cell division without dilution, the sensitivity for in vivo detection is enhanced.6,7 Using a reporter system, multimodality (including bioluminescence, fluorescence, and positron emission tomography [PET]) imaging permits longitudinal monitoring
of the cell survival, homing, and quantification over an extended period, if the reporter gene is stably expressed.

Coupling of the reporter gene to tissue specific gene promoters could also allow monitoring of cell differentiation, eg, detection of angiogenic gene expression would be possible.6 Because PET reporter gene imaging in a large animal model of MI on a clinical PET scanner has not been demonstrated, the aim of the present study was to adapt and validate the reporter gene method for in vivo monitoring of cardiac SC therapy in a large animal MI model that would facilitate translational research.

Methods

Cell Culture
Porcine bone marrow (BM)-derived mesenchymal stem cells (MSCs) were selected from 100 mL BM mononuclear cell fraction using a Ficoll-Paque gradient (Ficoll-Paque, Amersham Biosciences, Uppsala, Sweden) and stored at 4°C in Baxter bag (Baxter Healthcare, Ltd, Thetford, Norfolk, UK) in seven pigs (MSC group). Buffy coats were plated at 50 000 cells/cm² in alpha modified Eagle’s medium without nucleotides but also containing 10% FCS and 2 mmol/L glutamine, penicillin or streptomycin supplemented with 1 ng/mL Fibroblast Growth Factor 2. Cells were harvested by trypsinization when 75% confluent and replated at a cell density of 1000 cells/cm². Cells were tested for absence of CD45 expression. In vitro cell differentiation testing was performed in CD45⁻ cultures.

Differentiation and Characterization of the MSCs Stemness
The differentiation media for adipogenesis consisted of DMEM (low glucose), 20% FCS, 0.5 mmol/L isobutylmethylxantine, 60 μmol/L indomethacin, and 10 to 6 mol/L dexamethasone. For osteoblastic differentiation, the medium consisted of DMEM (high glucose), 10% FCS, 10 mmol/L β-glycerophosphate, 50 μg/mL L-ascorbic acid, and 10 to 7 mol/L dexamethasone. For chondrogenic differentiation, a micropellet system was used. Cells were suspended in DMEM, containing 0.1 μmol/L dexamethasone, 1 mmol/L sodium pyruvate, 0.17 mmol/L ascorbic acid, 0.35 mmol/L proline, and 1:500 stock from Cambrex of insulin-transferrin-selenium. The medium was supplemented with bone morphogenic protein-2 at 100 ng/mL or transforming growth factor-β at 10 ng/mL (Figure 1).

Cell Transfection
MSCs were transfected by incubation with cell culture supernatants from 293 T lentiviral packaging cells, previously transfected, using the calcium phosphate method, with the envelope plasmid pMD-G-VSVG, the packaging plasmid pCMV-DR8.2, and an expression plasmid construct containing the renilla luciferase (RL)-red fluorescence protein (RFP)-herpes simplex truncated thymidine kinase (tTK) (LV-RL-RFP-tTK). Fluorescence-activated cell sorting was used to select the highest RFP⁺ cells (10% of the total transfected MSC) for in vivo delivery (Figure 1).

Effect of Reporter Genes on LV-RL-RFP-tTK-MSCs Cell Viability and Proliferation
For cell viability and proliferation assay, the cells were serially washed with PBS (Invitrogen, Carlsbad, Calif), collected at different
Chemical purity of [18F]-FHBG. The radiochemical yield was 10% to 12% (corrected for decay) in five runs, and the synthesis time was 120 minutes. The specific radioactivity of the radiotracer was 44.4 to 74 GBq/minute from the end of bombardment. The average specific radioactivity of LV-RL-RFP-tTK-MSCs, the cells (0.5 × 10^6/mL) were incubated at 36°C in PBS containing 5 mmol/L D-glucose and 15 mmol/L NaCl for 20 to 180 minutes. The unbound tracer was washed. The cells were resuspended in 1 mL PBS and loaded in 1-cm diameter tubes after which the activity was counted in a gamma counter (Canberra Packard, Dreieich, Germany) for 1 minute within the 18F-sensitive energy window to determine isotope incorporation.

Animal Protocol

Fourteen domestic pigs (25 to 30 kg) underwent induction of MI (closed-chest, reperfused model, seven of them had previous BM-harvesting) (Figure 2). The pigs were sedated with 12 mg/kg ketamine hydrochloride, 1 mg/kg xylazine, and 0.04 mg/kg atropine intramuscularly after overnight fasting. Following endotracheal intubation, a 6F introducer sheath (Terumo Medical Corporation, Somerset, NJ) was inserted in the femoral artery. After the administration of 200 IU/kg of heparin, selective angiography of the left coronary arteries was performed, and a balloon catheter (3.0 mm in diameter, 15 mm long) (Maverick, Boston Natick, Mass) was advanced into the left anterior descending coronary artery after the origin of the first major diagonal branch. The mid left anterior descending coronary artery was then occluded by inflation of the balloon at 5 atm for 60 minutes, followed by reperfusion via balloon deflation. After hemodynamic stabilization, the pig was allowed to recover.

Sixteen days after acute MI, after baseline cardiac MRI, seven pigs received 3D NOGA-guided intramyocardial injections of LV-RL-RFP-tTK-MSC (Group LV-RL-RFP-tTK-MSC), whereas seven pigs served as controls (Group Control). For in vivo tracking of the injected cells, [18F]-FHBG PET-computer tomography (CT) imaging was performed 30 ± 2 hours and 7 days after injections.

Ten days after delivery of the LV-RL-RFP-tTK-MSC, the pigs underwent another MRI and then were humanely euthanized and the hearts excised. The endocardial injection sites were localized and sectioned. Tissue samples of the heart (ie, injection sites, infarcted-noninjected sites, and normal myocardium) and different organs (ie, pericardium, pleura, lung, mediastinal lymphatic nodes, liver, spleen, bone marrow) were collected. A part of all pigs and subjected to either histopathologic cell counting or luciferase assay (Figure 2).

Intramyocardial Delivery of Transfected Cells Using 3D NOGA Guidance

After placement of 8F sheath (Terumo Medical Corporation), the diagnostic electromechanical NOGA mapping catheter (Cordis, a Johnson & Johnson, Miami Lakes, Fla) was introduced into the LV cavity. Detailed descriptions of the endocardial mapping system components are previously published.9–11

After the diagnostic NOGA endocardial mapping, the LV-RL-RFP-tTK-MSCs were injected into the peri-infarct myocardium in 11.3 ± 1.2 sites with 6.3 ± 1.1 × 10^3 cells/injections using the Myostar injection catheter (Cordis, Johnson & Johnson). The injections (0.3 mL cell suspension each) were given slowly (30 to 40 seconds) and only to areas with a unipolar voltage above 5 mV, using the quality control criteria.9

Cardiac PET Imaging

For determination of the myocardial perfusion and to facilitate the localization of the injected LV-RL-RFP-tTK-MSC, 13N-ammonia PET was performed 1 hour before [18F]-FHBG PET. After the transmission CT scan, dynamic image acquisition was started simultaneously with the intravenous injection of 500 MBq 13N-ammonia (12 frames with 10 seconds, three frames with 60 seconds, three frames with 300 seconds). After physical decay of 13N-ammonia to nearly undetectable activity levels, 10 mCi (~0.33 mCi/kg) [18F]-FHBG were injected intravenously, and PET scans were completed 1 hour after tracer injection.

Emission images were summed and analyzed with the transmission PET data by using an ECAT EXACT 921 (CTI/Siemens) scanner over 60 minutes in three-dimensional acquisition mode. For attenuation correction, 15 minutes transmission scans were acquired after completing the dynamic scan. Random-, scatter-, and attenuation-corrected transaxial images were reconstructed by an ordered subset EM (OSEM) method (four iteration, eight subset) supplied by the manufacturer. The uptake of radioactivity was decay corrected and expressed as standard uptake value, normalizing the accumulation to the injected activity and the body weight. Standard uptake value was calculated as activity per milliliter/injected activity/body weight. Emission images between 40 to 60 minutes were summed and analyzed together with the transmission data by using MATLAB 7.1.

Regional myocardial concentrations (percentage dose per milliliter) were estimated from maximum pixels within the regions of interest (ROIs).12 The count ratio of [18F]-FHBG PET relative to remote myocardium was calculated.12

Cardiac MRI of the Pigs

Cardiac MRI was performed using a 1.5-T clinical scanner (Avanto, Siemens, Erlangen, Germany) by using a phased array coil and a vector ECG system. Cine MR images were acquired using a retrospectively ECG-gated, steady-state free precession cine MRI technique in short-axis and long-axis views of the heart using 1.2 ms
Histological Detection of Intramyocardially Injected MSCs

Tissue samples from the noninfarcted, infarcted-treated, and infarcted-nontreated myocardium were fixed in 4% buffered paraformaldehyde for 3 days at 4°C, and embedded in paraffin for routine histological procedures. Five-micrometer sections were counterstained using Hoechst stain (Sigma Aldrich H6024) at 0.5 μg/mL, incubated for 10 minutes.

For counting of the RFP^+ cells, 10 myocardial tissue blocks (injected and remote areas, 8×8×8 mm of size) were sectioned, and the sections with the appropriate depths (determined by confocal microscopy) with the RFP^+ cells were further cut into final 15-μm slices. The total numbers of RFP^+ cells were counted in 200 slices of each block. The fluorescence images were acquired on a Leica TCS-SP2-AOBS confocal microscope (Leica TCS-SP2-AOBS), with a 63×/1.30 glycerol-immersion objective. An UV/HeNe laser with filter settings for DAPI/TRITC fluorescence was used for simultaneous two channels detection.

Localization of the LV-RL-RFP-tTK-MSCs Injections

For confirmation of the location and viability of the injected cells, six pigs underwent the same protocol as the animals in Group LV-RL-RFP-tTK-MSC, but received the LV-RL-RFP-tTK-MSC intramyocardially using NOGA guidance in two sites (0.72±0.25×10^6 cells/injection site) with an additional injection of nontransfected MSCs in the opposite posterior wall. 13N-ammonia PET followed by [18F]-FHBG PET-CT scans was performed 8 hours (n=3) or 7 days (n=3) after MSC delivery (Figure 2).

Additionally, transmission PET scans were registered to the corresponding MRI scans, using landmark-based registration for further confirmation of the injection sites with PET-MRI fusion software. The homologous anatomical landmarks were delineated on the contours of lung, thorax and the heart apex. The PET images were resliced by the evaluated rigid body transformations and the MRI and PET images were visualized by BrainCAD image fusion software (www.pet.dote.hu/braincad).

Luciferase Assay

Tissue samples (1.52±0.09 g) collected from the infarcted (injected and noninjected) and normal myocardium and different organs of the animals were mechanically disrupted at high speed for 15 to 20 seconds in ice-cold standard relaxing solution: Na-ATP, MgCl2, ethylene glycolbis (β-aminoethyl ether)-N,N,N,N-tetraacetic acid (EGTA), KCl, imidazole at 4:1:2:100:10 mmol/L/L, respectively (1 g tissue/0.5 mL solution).13 The homogenized tissue was centrifuged at 800 rpm for 1 minute at 4°C. The supernatant was discarded and the tissue was resuspended in standard relaxing solution containing 0.3% Triton X-100 for 6 minutes to remove membranous structures. After washing twice with the relaxing solution, the cells and cell fragments were resuspended in standard relaxing solution. Luciferase was determined using components of the Dual-Luciferase Reporter Assay System (Promega, Madison, Wisconsin, USA). Briefly, an equal volume of passive lysis buffer was added to the homogenized tissue samples, and incubated 15 minutes at room temperature. Luciferase was measured in a Luminoscan Ascent luminometer (Labsystems) and expressed in relative light units (RLU) per μg protein.

Statistical Analysis

Parameters in the LV-RL-RFP-tTK-MSCs treated animals and controls were expressed as means±standard deviation. The analysis of the quantitative MRI data was performed blinded to treatment arm. Changes in MRI parameters (in Groups LV-RL-RFP-tTK-MSC and Control) and the cell doubling data were tested using an analysis of variance. The probability values yielded by the multiple comparisons were corrected for multiplicity by using the method of Bonferroni-Holm. A P<0.005 was considered statistically significant.

The authors had full access to and take full responsibility for the integrity of the data. All authors have read and agree to the manuscript as written.
Results

In Vitro Controls of the Transfected Cells

In vitro assays of osteogenic, chondrogenic and adipogenic differentiation were not altered by transfecting the MSCs (Figure 1). Both cell viability and proliferation assays showed no significant difference between the nontransfected MSCs and LV-RL-RFP-tTK-MSC with a cell doubling time of 3.67 and 4.05 days, respectively (Figure 1). The LV-RL-RFP-tTK-MSC stably expressed both RFP and RL for five passages as examined by fluorescence-activated cell sorting and enzyme assays, respectively.

In vitro studies of [18F]-FHBG uptake demonstrated an increasing uptake of the tracer with extended incubation (Figure 3A) in a minimum detectable concentration of $1 \times 10^5$ MSCs/mL in vitro (Figure 3B).

In Vivo Cell Tracking of the Intramyocardially Delivered Transfected MSCs

PET imaging demonstrated focal [18F]-FHBG tracer uptake of the anterior myocardial wall accompanied by a pattern of intense tracer foci at the local injections of the LV-RL-RFP-tTK-MSC when injected in two sites with a distance of at least 2 to 3 cm 8 hours after delivery (Figure 4). Fusion imaging of PET with MRI or CT confirmed the myocardial localization of the tracer activity presumably corresponding to the RL-RFP-tTK modified MSCs (Figure 4). Seven days after LV-RL-RFP-tTK-MSC delivery, no specific [18F]-FHBG accumulation was detected in the two injection sites. When MSCs were injected in 11.3 ± 1.2 sites (Group LV-RL-RFP-tTK-MSC) [18F]-FHBG imaging showed diffuse distribution of the tracer 30 hours after injections, but not in the controls. The count ratio of [18F]-FHBG in the entire anterior wall relative to remote myocardium was 5.3 ± 1.8. Placement of ROI on the intense tracer foci of the [18F]-
FHBG uptakes, the count ratio of the [18F]-FHBG of ROI relative to remote myocardium was 11.3±3.5. Comparing the [18F]-FHBG tracer uptakes between the LV-RL-RFP-tTK-MSC-treated animals and controls, the count of the entire anterior wall and injection foci were 5.8±1.6 and 10.9±2.7 times greater in the treated animals relative to the controls, respectively. Seven days after cell transplantation, PET images displayed faint focal activity in the myocardium, and a similar activity pattern in the pericardium and pleura, indicating the spreading of wandering living cells to the surrounding tissues (Figure 5). The count ratio of [18F]-FHBG of the tracer foci relative to remote myocardium was 2.1±0.3, and relative to the heart of control animals 1.8±0.2 on day 7. Placement of ROIs on the pleura and pericardial locations of [18F]-FHBG uptakes, the counts were 1.6±1.1 times higher in the treated animals compared with the control animals.

**Effect of Cell-Based Gene Therapy on Infarct Size and Myocardial Function**

One pig in each group died during MI induction. MRI at 10 days revealed a trend of a decreased end-diastolic volume (EDV) \(P=0.07\), an increased ejection fraction \(P=0.06\), and a statistically significantly smaller infarct size \(P<0.005\) in the Group LV-RL-RFP-tTK-MSC compared with control (Table 1, Figure 6).

**Number of Engrafted LV-RL-RFP-tTK-MSC**

Ten days after LV-RL-RFP-tTK-MSC implantation, fluorescence confocal microscopy of the myocardium showed the presence of RFP\(^+\) cells distributed through an elongated track in the area surrounding the intramyocardial injections in Group LV-RL-RFP-tTK-MSC (Figure 7) but not in samples from control pigs. The mean RFP\(^+\) cells/pig in the cardiac tissue samples was \(3.46±0.72\times10^5\), ie, \(5.8±1.1\%\) of the injected cells (Figure 7).

**Luciferase Activity**

Analysis of luciferase enzyme activities revealed high level of expression of the RL gene in the myocardial injection sites 10 days after delivery and some RL expression in the remote myocardial site, pericardium, pleura, liver, spleen, mediasti-

![Figure 5. A, NOGA endocardial mapping of pig left ventricle 16 days after MI with 12 injections (black points) of LV-RL-RFP-tTK-MSC at the border zone of anterior wall MI. B, PET image of a pig 30 hours after injection of LV-RL-RFP-tTK-MSC demonstrates diffuse high tracer activity in the myocardium, corresponding with the 12 injection sites of the transfected cells. The emission and transmission slices are represented by hot color bar and grayscale with standard uptake values. C, No uptake is detected in the control animal. D, PET images seven days after injection of LV-RL-RFP-tTK-MSCs with decreased but still detectable scattered tracer activity in the myocardium in the MSC-treated animal, and in the surrounding tissues indicating wandering of the living cells probably through the lymphatic routes. E, No tracer accumulation in the control pig at 23 days after MI.](http://circimaging.ahajournals.org/)

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Histological analysis confirmed the presence of the formation. In contrast, our PET study with MSCs revealed a activity of murine embryonic stem cells transfected with Previously, Cao et al.15 showed increased thymidine kinase cells and their progeny in large animal model of MI.14 only clinically translatable method for in vivo tracking of the gene method is able to detect biodistribution of transfected cells to distant organs that may have unwanted side effects from the cells engineered for angiogenesis, longer survival, and transdifferentiation.

We used the intramyocardial delivery of the cells, as Hofmann et al.17 demonstrated that intracoronary delivery resulted in <3% retention of SCs in the infarcted heart within a few hours. Similar to our results, the intramyocardial delivery of embryonic cardiomyoblasts into rat hearts using in vivo dual-isotope imaging detected about 20% engrafted cells for 96 hours.15,16 However, cell retention may vary based on not only the method of delivery but also the stem cell type.

The stable transfection of the cells with lentiviral or nonviral vector carrying reporter genes might influence the cell biology and physiology. Wu et al.19 demonstrated, that embryonic stem cells transfected with the lentiviral vector of trifusion reporter gene had no significant effects on embryonic stem cell viability, proliferation, and differentiation capability in vitro and in vivo.

Even if the monitoring of cell fate is of clinical importance in cell-based regeneration therapies, the usage of reporter genes in human studies has some limitations, such as extensive molecular manipulation and genetic modification of the stem cells. The remote homing of the living cells necessitates further preclinical investigations from several aspects: (1) even if the ex vivo viral gene transfer uses a replication-
defective retroviral vector, a low titer of contaminating infectious lentiviral vectors could be detected by Bloëmer et al, which might carry a potential hazard of undesirable viral vector shuttle18; (2) cell transfection with therapeutic gene might lead to an unwanted vascularization of remote organs, hampered by the resistance of lentiviral vectors to gene silencing. Further refinement of this method will be the elimination of viral vector use for transfection, increasing the safety of cell-based gene therapy in general.

Limitations
Although both groups included small number of animals, the results were consistent between animals and between in vivo imaging and postmortem histological analysis.

The RFP$^+$ cells were counted by confocal microscopy, a method that is limited through autofluorescence and the sectioning of the relatively large pig heart into 10 blocks. However, the number of counted cells corresponded well with the cell number-dependent luciferase activity presented in Figure 7. An alternative and more reliable approach would be a sex-mismatch transplant and TaqMan polymerase chain reaction on Y-chromosomes, if we did not use autologous stem cells for delivery.

The presented study was a prospective study, with its limitation of a nonblinded study design. However, the analysis of the quantitative MRI data was performed blinded to the treatment arm, and thus, the functional improvement of the infarct size is not biased.

In conclusion, we have demonstrated the feasibility of PET and optical imaging of the stable expression of the trifusion gene protein in a relevant large acute ischemic animal model. Thus, this technique demonstrates promise as an approach to translational research of cell-based gene therapy in cardiovascular disease.

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CLINICAL PERSPECTIVE

The potential clinical impact of the presented study is the demonstration of the feasibility of serial noninvasive in vivo tracking of cardially delivered stem cells transfected with the reporter gene by using multimodality optical imaging with clinical PET, CT, and MRI in a relevant experimental model of myocardial ischemia (closed-chest reperfused infarction). The model and percutaneous application of cells by a clinical system support the translatability of this study. The insertion of reporter genes into stem cells allowed the monitoring of cell fate, because the reporter is expressed as long as the cells are alive and passes to the daughter cells on cell division. This study has also confirmed the efficacy of cell-based gene therapy, showing a decrease in infarct size by percutaneous intramyocardial delivery of the cells. The study has highlighted the migratory itinerary of the cardially delivered cells, as living transplanted stem cells have been detected in remote organs. The clinical relevance of this finding is that the reporter gene method is able to detect the biodistribution of transfected cells to distant organs, where they may have unwanted side-effects from the cells engineered for angiogenesis, longer survival and transdifferentiation. The further refinement of this method will be the coupling of the imaging reporters with therapeutic genes with controlled release of angiogenic factors, and elimination of viral vector use for transfection. This method has the potential of increasing the safety and efficacy of cell-based gene therapy in human regenerative medicine in ischemic or degenerative organ disorders.
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