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Improved Cellular Uptake of Perfluorocarbon Nanoparticles for In Vivo

Murine Cardiac ¹⁹F MRS/MRI and Temporal Tracking of Progenitor Cells

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Abstract

Herein, we maximize the labeling efficiency of cardiac progenitor cells (CPCs) using perfluorocarbon nanoparticles (PFCE-NP) and ¹⁹F MRI detectability, determine the temporal dynamics of single-cell label uptake, quantify the temporal viability/fluorescence persistence of labeled CPCs in vitro, and implement in vivo, murine cardiac CPC MRI/tracking that could be translatable to humans.

FuGENE^{HD}-mediated CPC PFCE-NP uptake is confirmed with flow cytometry/confocal microscopy. Epifluorescence imaging assessed temporal viability/fluorescence (up to 7 days [D]). Nonlocalized murine ¹⁹F MRS and cardiac MRI studied label localization in terminal/longitudinal tracking studies at 9.4T (D1–D8).

A 4–8 fold ¹⁹F concentration increase is evidenced in CPCs for FuGENE vs. directly labeled cells. Cardiac ¹⁹F signals post-CPC injections diminished in vivo to ~31% of their values on D1 by D7/D8. Histology confirmed CPC retention, dispersion, and macrophage-induced infiltration. Intra-cardiac injections of PFCE-NP-labeled CPCs with FuGENE can be visualized/tracked in vivo for the first time with ¹⁹F MRI.

Keywords: Fluorine MRI, perfluorocarbon nanoparticles, cardiac stem cells, macrophages, tracking.

(Graphical Abstract: Modified Figure 7)

1. Introduction

Implantation of stem cells (SCs)¹ promises cardiac tissue regeneration and structural and functional improvements following injury [1]. While the feasibility of SC therapies has been proven, efficacy has yet to be demonstrated conclusively [2].

In recent years, perfluorocarbon nanoparticles (PFCE-NPs) have been applied in numerous therapeutic applications [3] and have led to direct tracking and quantification of labeled SCs and other cell populations using ¹⁹F-MRI [4–8], benefitting from fluorine's MR visibility, the lack of in vivo tissue/organ content, and clinical relevance. However, to-this-date, there are significant challenges and limitations with this approach [9], such as the efficiency of label uptake, compromizing cellular fate and cardiac signal detectability in vivo [2].

Different agents commonly used for DNA transfection (FuGENE^{HD}, lipofectamine, others), herein referred to as "labeling agents", have been employed [10–14] to increase the efficiency of cellular uptake of ¹⁹F compounds, targeting immune-based therapies [6, 7, 15], and work unrelated to the direct administration of cardiac SCs [16, 17, 18]. Alternative strategies using

¹ ANOVA: Analysis of variance; BW: Bandwidth; CPCs: Cardiac progenitor cells; CDCs: Cardiosphere-derived cells; CT: Collagenase/Trypsin; DAPI: 4',6-diamidino-2-phenylindole; ECG: Electrocardiographic; FLASH: Fast low-angle shot; FOV: Field-of-view; FSC: Forward scatter; GFP: Green fluorescent protein; IMDM: Iscove's Modified Dulbecco's Medium; ISO: Isoflurane; LV: Left-ventricular; MPIOs: Iron oxide particles; MRI: Magnetic Resonance Imaging; MRS: Magnetic Resonance Spectroscopy; NPs: Nanoparticles; NEX: Number of excitations; PFCE-NPs: Perfluoro-crown-ether nanoparticles; PFCE: Perfluoro-crown-ethers; PBS: Phosphate buffer solution; PW: Pulse width; RF: Radiofrequency; SSC: Side scatter; SCs: Stem cells; SNR: Signal-to-noise-ratios; SPIO: Superparamagnetic iron oxide; ST: Slice thickness; SPGR: Spoiled gradient echo; TE: Echo time; TR: Repetition time; TFA: Trifluoroacetic acid; 2D: Two-dimensional; 3D: Three-dimensional.

electroporation techniques have led to moderate label payload increases (¹⁹F or other) [19, 20, 21, 18] at the expense of increased risks of cellular toxicity, difficulties in optimization, imbalances of cell homeostasis, and possible permanent permeabilization of the plasma membrane [21, 18].

Despite prior ¹⁹F MRI advances in the visualization of pre-labeled SCs with PFCE-NPs or other MRI contrast agents [5, 15], there have been no prior reports on murine cardiac ¹⁹F imaging of exogenously administered cardiac SCs following direct, intra-cardiac injections. To-this-date, human or in vivo murine studies are lacking, while prior reported imaging attempts in rats have been prohibitively lengthy, costly (multiple millions of cells/animal), and practically complex to reproduce in mice [20, 22].

Invariably, low-label uptake is prohibitive for the visualization and tracking of labeled SCs in longitudinal cardiac applications in vivo.

In this proof-of-principle study, we show significant enhancements of PFCE-NP label uptake in two types of cardiac progenitor cells (CPCs) [23] following labeling with the agent FuGENE [24], thereby accomplishing in vivo cardiac ¹⁹F MRI visualization and tracking with minimal cellular toxicity. We report significant improvements of ¹⁹F MRI signal detection in the labeled CPCs (cardiosphere-derived (CDCs) and collagenase-trypsin (CT) cells) using FuGENE compared to directly labeled or electroporated cells. CDCs have been shown to improve cardiac function in models of MI, whereby administered cells were retained—albeit in low numbers—and expressed cardiac proteins [23, 26, 27]. CT cells have been isolated by enzymatic digestion and slow adhesion and have a comparable gene expression and differentiation profile to CDCs [28].

The purposes of the study are to a) maximize the labeling efficiency of CPCs using PFCE-NPs and the ¹⁹F MRI detectability, b) determine the temporal dynamics of single-cell label uptake, c) quantify the temporal viability/fluorescence persistence of labeled CPCs in vitro, and d) implement an in vivo, murine cardiac CPC administration protocol, ¹⁹F MRI and tracking, that could be translatable to human applications.

2. Methods

Animal Ethics

All experimental procedures in this prospective, proof-of-principle animal study were approved by the Home Office, UK and International regulatory bodies, and adhered to local institutional guidelines. Every effort was expended to ensure humane care of the animals in all animal studies.

Cell Isolation and Labeling

Cell isolation, labeling, labeling efficiency, cell viability, and fixation studies were conducted in accordance to standard procedures. Methodological details are listed as Supplementary Materials.

In Vitro and in Vivo Imaging

Label Tracking - Confocal microscopy

Temporal Labeling Kinetics: Fluorescent NPs and/or nuclear green fluorescent protein (GFP) stained cells were imaged [excitation wavelengths: λ_{green} =488 nm, λ_{red} =633 nm, emission ranges: 500–550 nm (green) and 650–700 nm (red)] using a Leica TCS SP8 confocal microscope (Leica-

Microsystems, Mainhem, UK) with HyD detectors (objective NA=1.4, 63X) at different times post-seeding (1–6 h), and at two FuGENE concentrations (25, 100 μ l in 2 ml IMDM). Image post-processing was conducted using the Imaris software (Bitplane, Zurich, Switzerland) that allowed generation of three-dimensional (3D) renditions and movies showing the temporally dependent label uptake up to 6 h post-labeling.

High-content Epifluorescence Microscopy Imaging

Temporal Viability and Fluorescence: Live cells were stained with Calcein (CellTraceTM Calcein Red-Orange, Thermo Fisher Scientific, UK) for high-content imaging in 6-well plates. Independent cell cohorts (n=3, ~10–20 k cells/cohort) were maintained in culture up to 7 d (D), and a time course study (D1–D7) of live cells was conducted (at low/high proliferative rates, given initial low/high cell densities) to assess cell survival (Calcein/Hoechst), labeling efficiency (Calcein/Atto647), and fluorescence (Atto647), using a high-content imaging system (Operetta, Perkin-Elmer, Llantrisant, UK). The Operetta's Harmony software was used for image analyses.

In Vivo Cell Administration using Direct Injections

Surgical Procedure: C57BL/6 mice (adult female, 20-30 g, n=6) were anesthetized with 4% isoflurane (ISO) and placed on a heating pad, intubated, and ventilated with 100% oxygen (MiniVent, Type 845, Hugo Sachs Elektronik, Harvard Apparatus, March-Hugstetten, stroke rate of 175 strokes/min and a stroke volume of 250 μ l). Analgesic was administered (80 μ l, Vetergesic). A left thoracotomy was performed, and the FuGENE-labeled cells (~1.7–2×10⁶ CT cells in ~50 μ l of IMDM media) were injected at left-ventricular (LV) myocardial sites using a

0.5 ml syringe. The chest was sutured and the intubation tubing removed to allow spontaneous breathing. Mice were monitored for 30–60 min for recovery.

MRI/MRS

All experiments were conducted on a 9.4 T Agilent scanner (Agilent Technologies, Santa Clara, CA, USA). Relevant methodological details pertaining to radiofrequency coils and in vitro studies are listed as Supplementary Materials.

In Vivo MRI

In Vivo Cardiac ¹⁹*F MRI and Cellular Tracking:* C57BL/6 mice (n=6) were anesthetized with 4% ISO and were maintained for up to 95 min with 1.5–2.0% ISO in 100% oxygen.

All animals were placed in a specially constructed cradle and were their body temperature was maintained at ~37°C. Electrocardiographic and breathing rates were monitored using a gating system. Heart rates were maintained between 350–500 beats/min.

Six animal studies were completed in total. Three successful studies were terminal at D1, and another three were tracking studies (two mice were rescanned at D4 and D7, and one mouse at D5 and D8). Poor visualization of ¹⁹F signal was evidenced in one of the tracking studies (on D7), attributed to partial mis-localization of injected cells.

Details on ¹H and ¹⁹F image (spoiled grass) and spectroscopy (saturation recovery) acquisitions are listed as Supplementary Materials.

Image Processing

Details on image/spectral analyses are listed in the Supplementary Material section. Briefly, ¹⁹F images were interpolated, colored coded, and overlaid on the anatomical ¹H MRI. For MRS, spectral areas were quantified.

Histology and Immunofluorescence Staining

Histology

Post-mortem histological evaluation was performed in mouse hearts on the same day following cellular implantation (terminal-D1) or at subsequent days (tracking-D7, D8) to assess the CPC injection and retention sites. The excised hearts were dehydrated and fixed (in a 4% methanol-free formaldehyde solution), processed, embedded in paraffin, and stored. Serial transverse paraffin sections of 10–17 μ m were cut from base to apex, processed, and stained using the nuclear stain 4',6-diamidino-2-phenylindole (DAPI, Sigma-Aldrich, UK). Imaging and analyses were performed on a Leica bright-field optical, and on the confocal microscope, by two blinded investigators (DAPI, Atto647).

Immunofluorescence Staining

Hearts from particle-injected mice were fixed in 4% methanol-free formaldehyde solution before being transferred to 70% ethanol, followed by paraffin embedding. Hearts were sectioned (10-17 µm) and blocked for 2 h in DAKO® Protein Block (Thermo Fisher Scientific, UK). Thereafter, the sections were incubated for 2 h with goat polyclonal anti-mouse Galectin 3 antibody (RnD Systems AF1197, Oxford, UK) followed by incubation with the secondary antibody donkey anti-goat Alexa Fluor 488 (A11055, Invitrogen, Thermo Fisher Scientific, UK). Sections were washed and mounted with Fluoromount G mounting medium (Southern Biotech 0100-01, USA) with DAPI. Images were obtained with the confocal microscope.

Statistical Analyses

All results are reported as mean \pm standard deviation (SD). Two-tailed Student's t-tests (XLSTAT, Addinsoft, New York, USA), and two-way analysis of variance (ANOVA) with repeated measures (SPSS, Version 24, IBM, USA) were also used to assess mean comparisons (α =5%) (control, labeled, FuGENE-labeled, electroporated), and the effects of the labeling regime (control, labeling, FuGENE labeling), time (D1, D4, D7), and their interactions, for in vitro CDC/CT cell assays. Power calculation for the tracking studies of control, injected mice (exposure of controls p₀=0.99, relative risk of disease associated with exposure RR=0, α =5%) yielded a required minimum sample size of n=3.

3. Results

Cell Isolation and Labeling

FuGENE Optimization: Figure 1 shows sequentially gated, flow cytometric results (CT cells) (without/with FuGENE) using Atto647-fluorinated-PFCE-NPs. The labeling efficiency was > 99% (in both labeling instances) with noted increases in the total label load (D1) upon FuGENE use (>4-fold fluorescence increases with FuGENE). Successful persistence of cellular label (labeling efficiency was > 99%) was confirmed with independent FuGENE-labeled CT cell cohorts post-labeling (D8) using flow cytometry (results not shown). For the FuGENE optimization studies, the overall cell viability was >88%.

In Vitro and in Vivo Imaging

Labeling Efficiency: There is an innate heterogeneity of labeling in the studied CPCs, confounded by the label methodology (endocytosis-based internalization), cellular uptake mechanism, and the heterogeneous content of isolated cells (containing cardiac progenitors mixed with endothelial cells, fibroblasts, and others). Correspondingly, given the sensitivity disparities for flow (single cells, nm to tens of μ m), epifluorescence imaging (multiple thousands of cells, tens to hundreds of μ m), and MRI (cell pellets containing > 10k cells per voxel, hundreds of μ m), there are expected disparities in the detection of labeled cells.

High-content Imaging

Temporal Viability and Fluorescence: Figure 2 shows quantified results for the temporally dependent proliferation and fluorescence using optical imaging of (a) FuGENE-labeled CDC and (b) CT cells at days 1 (D1), D4, and D7, at high seeding densities in vitro. The increased density of the initially seeded cells promoted enhanced proliferative activity at D4–D7 and label dilution, as reflected by the decreased levels of fluorescent intensity at D4/D7 compared to D1.

Histograms of the temporally varying cell proliferation, and mean fluorescence signal/cell, are also shown for both types of studied cells (Figure 2). The mean labeling efficiency increased to \sim 30–80% upon FuGENE use vs. \sim 5–10% following labeling (no FuGENE) (p<0.003/p<0.0042; CDC/CT) at day 1 (D1). However, the mean fluorescence signal decayed in labeled cells (CDC/CT) (without/with FuGENE) by D2/D4 owing to the possible cleavage of the fluorophore and cytosolic destruction of NPs (D1–D3) [Supplementary Figure S2] and cell proliferation (D4-D7) (p<0.007/p<0.001 for labeled, and p<0.0002/p<0.0001 FuGENE-labeled cases). Indicated is also the improved performance of CDCs (labeled/FuGENE-labeled) that elicit increased

fluorescent signals vs. CTs at D1 (p<0.04/p<0.006 CDC/CT). Two-way ANOVA with repeated measures yielded significant variability effects in viability with respect to time (CDC cells, p= 8.8×10^{-7}), and time (p= 1×10^{-7}), labeling scheme (p=0.01), and their interactions (p=0.002) (CT cells). Correspondingly, significant variability effects in fluorescence were noted for both cell types (CDC/CT) in terms of time (p= $0.002/1.8\times10^{-5}$), labeling scheme (p= $1.4\times10^{-7}/4.4\times10^{-11}$), and their interactions (p= $5.6\times10^{-6}/5.2\times10^{-10}$).

Electroporation: Similar viability/fluorescence trends to those in Figure 2 were observed for FuGENE-labeled compared to electroporated cells seeded at low densities [Figure 3 (A, B)]. By D7, viability was higher for electroporated cells vs. FuGENE-labeled cells (p<0.04), but the initial fluorescence response was better for FuGENE-labeled cells (p<0.002, D1). However, the mean fluorescence signal decayed significantly in both cell types (electroporated and with FuGENE) by D4 as cells proliferated (p<0.025, D1 vs. D4 for electroporated, and p<0.002, D1 vs. D4 FuGENE-labeled cases).

Label Tracking - Confocal microscopy

Temporal Labeling Kinetics: Figure 4(A-F) shows phase contrast confocal microscopy images of GFP positive CPCs, and (B, C) enlarged views of CPCs post-FUGENE labeling with low (25 μ l), and high doses (100 μ l) (red color). Figure 4(D–F) depicts 3D renditions of cells, (D) label localization (no FuGENE), and (E, F) labeled cells with 25 and 100 μ l of FuGENE. As indicated, cluster formation prohibits label uptake at high FuGENE doses [Figure 4(C, F)]. This problem is ameliorated at lower/optimal dose levels (Figure 4(B, E)). Supplementary Figure S2 contains three movies that confirm the temporal labeling kinetics using confocal live cell imaging (Movie

1: cellular labeling at 1.2 h post-label administration, Movie 2: cellular label with FuGENE with 25 μ l per million cells at 5.7 h, and Movie 3: high FuGENE dose with 100 μ l per million cells at 4.15 h). Significant label uptake was confirmed within 2–3 h post-label administration in all cases.

MRI/MRS

¹⁹*F-MRI/MRS:* Figure 4 shows unlocalized ¹⁹F MRS of TFA and labeled cells without/with FuGENE indicating signal increases from NPs (G–I), label localization at low doses (H), and an increased signal when the label is predominantly extracellular (I). Figure 4(J, K) shows axial ¹⁹F MRI from labeled CPCs in pellet suspensions at low/high FuGENE doses (butterfly coil). The results from MRS/MRI (Figure 4(G–K)) confirm confocal findings. Specifically, low label uptake is manifested by simple labeling (Figure 4(G)) that significantly enhances upon FuGENE dose increases (Figure 4(H, I)). The localization of FuGENE-encapsulated NPs in the extracellular space at nonoptimal FuGENE doses justifies the increased signal.

Confirmation of the label uptake mechanism and label localization attributed to FuGENE is provided by separate control experiments (Supplementary Figure S3(A–D)). In this figure, ¹⁹F MRS control phantom tests are summarized, confirming the source of the fluorine signal increase due to FuGENE. Independent in vitro labeling/FuGENE experiments were conducted (solenoid coil with a homogeneous B_1 response) for the quantification of ¹⁹F concentrations with respect to the concentration of a TFA standard (50 mM). Table 1 summarizes the SNR, cell seeding density, labeling efficiency, and ¹⁹F concentrations. As shown, indicative 4-8-fold concentration improvements (0.3 vs. 1.4–2.5 mM) are achieved with FuGENE, yielding corresponding labeling efficiency increases (10 vs. 44–75%).

Furthermore, comparison of the unlocalized, in vitro ¹⁹F spectra from electroporated and FuGENE-labeled CT cells (butterfly coil) indicates a 125-fold signal (D1) increase with FuGENE labeling compared to electroporation (Supplementary Figure S4 (A, B)). FuGENE-labeled cells following electroporation yielded 10% higher signal compared to FuGENE-labeled cells (Supplementary Figure S4 (C, D)).

In Vivo Cardiac ¹⁹*F MRI, Tracking, and Quantification:* Figure 5 shows overlaid ¹H MRI with ¹⁹F for terminal studies and bright field histology, confirming the injection locations. High-resolution, gated ¹H MRI are shown in Figure 6(A–F), and a sequence of overlaid ¹H/¹⁹F MRI of ~1.7–2 million FuGENE-labeled cells injected in the antero-lateral ventricular myocardium (D1, D4, and D8). Localization of the enhanced ¹⁹F signal regions was confirmed with histological staining, as described below.

Quantification: Temporal ¹⁹F voxel signal decreases were noted in vivo at increasing time intervals post-CPC administration, whereby the evoked ¹⁹F signal progressively diminished to \sim 31% of its value after injections, over a period of 7-8 days (Figures 6–7).

Histology: Typical bright field, and confocal microscopy imaging (nuclear [DAPI], label [Atto648], macrophages [Gal3]) are shown for a terminal (D1), and tracking studies (D7 or D8) confirming retention/localization of injected cells (D1), and infiltration of macrophages (Gal3) in fixed hearts at D7/D8 (Figure 8).

4. Discussion

Hitherto direct labeling methods led to an overall low and heterogeneous label uptake that prohibited cost-efficient, fast, and practical ¹⁹F MRI visualization and tracking of labeled SCs in cardiac applications.

Our choice of FuGENE versus protamine-sulphate, electroporation, or other transfection agents (e.g., lipofectamine) was based on the increased CPC viability and label uptake compared to those observed in preliminary experimental protamine-sulphate and lipofectamine studies (data not shown). Labeling was measured by flow cytometry (indicating label persistence at D8) and optical microscopy.

The observed discrepancy of the labeling efficiency of flow cytometry and epifluorescence imaging is attributed to the detection sensitivity differences of the two modalities. Flow represents a highly sensitive, single-cell-based optical method, while epifluorescence microscopy relies on volume-averaged optical signal responses from hundreds of cells. Epifluorescence imaging and MRI exhibited higher cellular sensitivity detection thresholds than flow cytometry, as reported below. This suggests that the level of labeling in many of the cells detected by flow cytometry would be too low to be detected using MRI.

Temporal kinetic studies in single cells indicate that NPs localize within FuGENE vesicles within the first 3-4 h, however, high doses lead to aggregate formations in the extracellular space, resulting in a low-label uptake, whereas the low-label dose achieved increased intracellular uptake, as confirmed by confocal microscopy. Based on recent MRI work [29], the in vitro load detection threshold for ¹⁹F PFCE labels (using cells and TFA) was 0.5 mM (butterfly). Additionally, the efficiency of labeling with FuGENE increased by 4.4–7.5 times vs. simple labeling yielding corresponding ¹⁹F concentration improvements in the order of 1.4–2.5 mM. Correspondingly, SNR increases in the order of 4.4–14 times were noted in ¹⁹F MR images

in vitro compared to the simple labeling case. Imaging can thus be executed within a few min using conventional gradient echo pulse sequences [29]. Signal enhancements may be highly beneficial in minimizing cell label load and potential toxicity in applications of cardiac pathology in vivo.

The fluorescent intensity decreased within 2-4 days in cultures in vitro primarily as a result of fluorophore cleavage, secondary to subsequent proliferation and label dilution (following day 3). Despite the slightly reduced viability rate following labeling with FuGENE (compared to labeling without it, or electroporation), the proliferative capacity of cells recovered. Notably, the persistence of in vivo MRI signal relies on the long-term fates of the NPs. The migratory paths of endosome-packaged NPs within cells involved lysosomal destruction and exocytosis, the temporal evolution of this process was faster than the timescale of cell division.

Several primary mechanisms may cause changes to cellular status (and possibly to the evoked signal) with time including possible: a) cellular migration/dispersion and clearance through the microcirculation, b) cell death, c) macrophage infiltration and label/cellular debris scavenging, hypoxic-induced proliferation Ultimately, and e) (no signal change). physiologically/pathologically relevant mechanisms supersede. For example, during cultures in vitro, cells proliferate and dilute label (as indicated herein). Under ischemic conditions, cells may undergo an initial, hypoxia-induced proliferation, causing label dilution but minimal overall ¹⁹F signal changes. Alternatively, cells may disperse and migrate from the injection site, ultimately entering the microcirculation, or lyse, and release the label into the extracellular space, thereby causing fluorescence, or possible ¹⁹F MRI signal decreases. The infiltration of macrophages at later times post-injection will also initiate cellular and label scavenging. While the end-effect is a complex interplay of the proposed mechanisms, in almost all cases, temporally dependent, voxelinduced signal decreases are expected, justified by in vivo findings.

In terms of quantification, the evoked signal on days subsequent to the day of injection may arise from both viable cells, macrophage-endocytosed NPs that remain at the injection site, and extracellularly released NPs from lysed CPCs. While noninvasive, single voxel ³¹P metabolic MRS may be able to assess the viability of injected cells [30], its potential applicability is limited by spatial resolution since CPCs will likely disperse, migrate, or lyse, post-injection. Combination of bioluminence techniques with MRI may address such issues, however, preclinical acquisition protocols become complex.

While the consistency of localization/detection of injected cells in the myocardium using ¹⁹F MRI has been confirmed with histology at the end of D1 post-CPC injection, the major limitations of this work include the limited animal sample size in tracking studies, and the lack of additional immunohistochemical analyses required for the detailed investigation of the spatio-temporal evolution of signal changes, including the confirmation of label localization within the interstitial space, within recruited macrophages, and/or the microvasculature.

Despite these shortcomings, we have demonstrated fast, in vivo cardiac ¹⁹F MRI post-SC injections in the in vivo mouse. More importantly, the protocol for cell preparation and injection has potential to be directly translatable to the clinic. The cell types employed herein have been used before with success in animal/human studies of cardiac pathology [26, 27]. However, additional issues ought to be resolved first (cell density, administration route, optimization of cell preparation methodologies for large-scale use) drawing on knowledge from successful strides based on localized injections of dendritic cells in humans [31, 32]. No potential problems are envisaged in terms of ¹⁹F label release and the safety profile of NPs, since the doses are small

and since all of these components have been previously used in humans with no reported adverse toxic effects [32]. PFCEs are inert, they are not metabolized in vivo, and will be excreted through exhalation in the lungs, and through the liver, spleen, kidneys, and blood, as has been previously been reported [33, 31]. It is understood that a long-clearance time is unacceptable in clinical studies, as has been observed in prior murine studies using high-dose PFCE emulsions [34]. Nevertheless, additional work needs to be expended to address clearance times for the trace amounts of PFCEs used herein in labeled cells, as it is unrealistic to extrapolate prior inferences based on prior dendritic cell studies.

The methodology may be further developed clinically as a noninvasive tool for determining the successful CPC administration type and route [35] and the temporally dependent retention in the acute phases of myocardial infarction.

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Author contributions

Guarantors of integrity of entire study: CC, MS, EMcN, CAC, JP, RSU, RC, AA, SPP, KK Study concepts/study design: CC, MS, EMcN, CAC, SPP, JP, RSU Data acquisition: CC, EMcN, AA, AS, RSU Data analysis/interpretation: CC, EMcN, RSU, SPP, CAC, MS, RC, AA, AS, JP, KK, AH, LP, RA Manuscript drafting: CC Manuscript revision for important intellectual content: CC, CAC, MS, EMcN, RC, RSU, AS, JP, RSU, SPP, AA, KK, AH, LP, RA

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Competing Financial Interests

None.

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Tables

Table 1: SNR of ¹⁹F MRI from labeled and transfected CT cells (at an optimal FuGENE dose), and mean fluorine concentration per cell (based on MRS) using the solenoid coil. Concentrations of labeled samples were estimated using an external reference TFA solution (50 mM), and ranged between 0.3–2.5 mM, respectively.

Biomarker Sample	Relaxation Times (T ₁ /T ₂) (ms)	¹⁹ F MRI Signal-to- noise-ratio (SNR)	Cell Seeding Density	Labeling Efficiency (%)	¹⁹ F Concentration (mM)
TFA (50 mM)	2728/2365¥				50
NP (solution) *	773/360				3.3
Labeled CT	1324/NA	27	$\sim 10^{6}$	10	0.3
FuGENE- labeled CT [†]	1360/533	103–373	~ 10 ⁶	44–75	1.4–2.5

^{*}Results denote an average of two independent measurements

[†]Improvements in SNR in the order of 4–14 times are achieved using FuGENE-labeling compared to simple labeling

[¥]Results reproduced from Constantinides et al. [29]

Figure Captions

- Figure 1: Flow cytometric results of CDCs. Side scatter (SSC) vs. forward side scatter (FSC) (CT cells) and side scatter vs. side scatter plots (singlets vs. doublets) in justification of the successful implementation of the labeling protocol without and with FuGENE (on day 1 [D1] post-labeling). The cells were labeled with PFCE-NPs containing Atto647. Successful persistence of cellular label was confirmed with independent CT cell cohorts on D8 post-labeling using flow cytometry (results not shown).
- Figure 2: Temporal quantification of control, labeled and FUGENE-labeled CDCs. Temporally dependent proliferation, and fluorescence response (Calcein, Atto647) using optical-epifluorescence imaging of FuGENE-labeled cells at days 1 (D1), D4, and D7, at a high seeding density. Increased cell density promotes proliferative activity at D4–D7 and label dilution, as reflected by the decreased levels of fluorescent intensity. Histograms of temporally dependent proliferation, and mean fluorescence intensity for (A, B) CDC (n=3), and (C, D) CT cells (D1– D7) (n=3). Epifluorescence studies of unlabeled control cells conducted in (B, D) did not elicit fluorescent signals (detected signal values were only attributed to autofluorescence and noise – not shown). Mean control fluorescence signals (unlabeled cells) matched the elicited mean fluorescence signals from labeled cells at D4–D7.
- **Figure 3: Quantification of FUGENE-labeled and electroporated CDCs.** (A, B) Histograms of temporally dependent proliferation and mean fluorescence intensity

(Calcein, Hoechst, Atto647) for FuGENE-labeled and electroporated-labeled CT cells (D1-D7) plated at low-seed densities in correlation to epifluorescence imaging (n=3). Increased proliferative activity promoted label dilution at D4-D7, as reflected by the decreased levels of fluorescent intensity. Mean control fluorescence signals (unlabeled cells) matched elicited mean fluorescence signals from labeled (FuGENE, electroporated) cells at D4-D7.

- Figure 4: Validation of label localization in CDCs using confocal microscopy and ¹⁹F MRS/MRI. (A) Phase contrast (PC) confocal microscopy of cardiac progenitor cells (CPCs), (B, C) CPCs post-labeling with 25 (intracellular white arrows) and 100 μl (extracellular grey arrows) of FuGENE, (D-F) corresponding label localization in (D) labeled cells, (E, F) and labeled cells with 25 and 100 μl of FuGENE. Green and red colors respectively indicate nuclear GFP and perfluorocrown (PFCE) NP label fluorescence. (G-I) Unlocalized ¹⁹F magnetic resonance spectroscopy (MRS) of trifluoro-acetic acid (TFA) and labeled cells with and without and with transfection FuGENE indicating signal increases from NPs post-transfection (H, I) label localization at low doses (H), and a spectral shift when the label is predominantly extracellular (I); (J, K) Corresponding axial ¹⁹F MRI from labeled CPCs in pellet suspensions at low/high transfection FuGENE doses (butterfly coil).
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injection sites (window/level settings for ¹⁹F images are different for different views).

- Figure 6: In vivo cardiac ¹⁹F MRI tracking and temporal signal quantification. (A, F) In vivo, gated ¹H and ungated oblique ¹H/¹⁹F MRI of the heart following a surgical procedure of cell injection (butterfly coil). The sequence of cardiac ¹⁹F MRI depicts views with narrow receiver bandwidth (1–3 kHz) acquisitions identifying the prolonged and contributory effects of ISO anesthesia, as reported earlier by Constantinides et al. [25] (window and level settings for ¹⁹F are different for different views). (G, H) Normalized histograms of the spectral areas of ISO and labeled cells (gated/ungated) at D1 (n=3, 4–6 h post-surgery), D4–D5 (n=3), and D7–D8 (n=2), and (I) corresponding normalized histogram of ¹⁹F signal from the entire volume of the detected, hyperenhanced regions at the injection sites with respect to the total fluorine signal on D1 [Note that three mice were imaged in the D7–D8 group but signal was only detected in two].
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Figure 8: Nuclear, bright field histological and immunostaining imaging. (A–E) Histological unstained and stained sections of murine hearts in justification of the retention of injected labeled cells, based on nuclear (DAPI), and surface marker (Atto647) staining (bright spots) using confocal imaging. (C) Histological/stained short-axis, mid-apical views of three murine hearts ([A–D] terminal experiments in two mice at (A, B) 4 and (C, D) 6 h post-cell injection on day 1 (D1), and [E–H] on D7/D8) indicating the sites of injected CPCs using bright field (C, D, E, H), nuclear (A), surface (B, E, Atto647), and immunostaining (F-G, Galectin3 (Gal3)), using confocal imaging.

Figures



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CDC (n=3), and (C, D) CT cells (D1–D7) (n=3). Epifluorescence studies of unlabeled control cells conducted in (B, D) did not elicit fluorescent signals (detected signal values were only attributed to autofluorescence and noise – not shown). Mean control fluorescence signals (unlabeled cells) matched the elicited mean fluorescence signals from labeled cells at D4–D7. Statistical significance bars are not shown in A and C (CDC: Control-D1 vs. D7, p<0.007, D4 vs. D7, p<0.014; Labelled-D1 vs. D4, p<0.021, D4 vs. D7, p<0.019, D1 vs. D7, p<0.012; FuGENE-labeled: D1 vs. D7, p<0.001, D4 vs. D7, p<0.006; and CT: Control-D1 vs. D4, p<0.002, D4 vs. D7, p<0.002, D1 vs. D7, p<0.001; Labeled-D1 vs. D4, p<0.009, D4 vs. D7, p<0.019, D1 vs. D7, p<0.012; FuGENE-labeled: D1 vs. D4, p<0.009, D4 vs. D7, p<0.019, D1 vs. D7, p<0.012; FuGENE-labeled: D1 vs. D4, p<0.009, D4 vs. D7, p<0.019, D1 vs. D7, p<0.012; FuGENE-labeled: D1 vs. D4, p<0.009, D4 vs. D7, p<0.019, D1 vs. D7, p<0.012; FuGENE-labeled: D1 vs. D4, p<0.009, D4 vs. D7, p<0.019, D1 vs. D7, p<0.012; FuGENE-labeled: D1 vs. D4, p<0.009, D4 vs. D7, p<0.019, D1 vs. D7, p<0.012; FuGENE-labeled: D1 vs. D4, p<0.009, D4 vs. D7, p<0.019, D1 vs. D7, p<0.012; FuGENE-labeled: D1 vs. D4, p<0.009, D4 vs. D7, p<0.019, D1 vs. D7, p<0.012; FuGENE-labeled: D1 vs. D7, p<0.009, D4 vs. D7, p<0.022, (D7), p<0.06; Controlled vs. FuGENE-labeled: D1 vs. D7, p<0.047, (D7), p<0.030).



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Supplementary Material

Cell Isolation and Labeling

Cell Isolation: Cardiac progenitor cells (CPCs) were isolated from adult, C57BL/6, or C57BL/6-Tg(CAG-EGFP)1Osb/J (The Jackson Laboratory, Bar Harbor, ME, USA) mouse atria. Cardiosphere-derived cells (CDCs) were expanded as described previously [1]. Collagenase/Trypsin (CT) cells were isolated by digestion of atrial tissue with 0.05% trypsin-EDTA and adhered on fibronectin-coated petri dishes over periods of 72 h [2].

Labeling-Viability: Cells were plated in Iscove's Modified Dulbecco's Medium (IMDM, Thermo Fisher Scientific, UK) and incubated in culture with fluorescent PFCE-NPs (Atto647, ATTO-TEC, GmbH, Germany) (10 mg/million cells) [3], for ~24 h before trypsinization, isolation, and pelleting. Cell viability was assessed using the Trypan Blue exclusion assay.

FuGENE Optimization: FuGENE (Promega, Madison, WI, USA) was used to label the cells with the NPs. To optimize the dose, three different FuGENE doses were used in two separate experiments (25, 100, and 200 μ l of FuGENE in ~10⁶cells). FuGENE was pre-mixed and incubated with the NPs for ~20 min before cell labeling. Cells were transfected overnight.

Electroporation: Cells were resuspended in 400µl of IMDM media and were electroporated using a BIORAD Gene PulserX cell electroporator unit (BIORAD, Hertfordshire, UK) using a single rectangular pulse (amplitude 250 V, duration 5 ms). They were then resuspended in a larger volume of IMDM media for epifluorescence and MR imaging.

Labeling efficiency: Labeling efficiency was determined with a CyAn ADP flow cytometer (Beckman Coulter, Indianapolis, IN, USA). Processing was conducted offline using FlowJo (FlowJo LLC, Version 10, Ashland, OR, USA). Cells were determined to be positively labeled (with 95% confidence) by comparison to the fluorescence of unlabeled cell samples.

Cell Fixation: Cells destined for flow cytometry were fixed in a 2% methanol-free formaldehyde solution (Thermo Scientific Pierce, UK), mixed with phosphate buffer solution (PBS) (1:7 v/v) (Sigma-Aldrich, UK), in final solutions containing 0.5-1 ml.

Transverse Electron Microscopy: For transverse electron microscopy (TEM), pellets of cell extracts were blocked in agar, fixed in 2.5% glutaraldehyde, washed in 0.1 M phosphate buffer (Sigma-Aldrich, UK), fixed in 1% osmium tetroxide, dehydrated in increasing concentrations of ethanol, and embedded in an epon araldite/resin mixture. The resin was polymerized at 60 °C overnight and ultrathin sections were cut using a Reichert Jung ultramicrotome (Vienna, Austria) with thicknesses in the range of 80-100 nm. Sections were stained with lead citrate and uranyl acetate, before being examined with a JEM 1010 TEM operating at 80 kV (JEOL, Tokyo, Japan).

MRI/MRS

The 9.4T scanner was equipped with a Direct Drive console and a 1000 mT/m actively shielded gradient set (internal diameter=60 mm) (Agilent Technologies, Santa Clara, CA, USA).

Radiofrequency Coils: For MRI studies, a 4×4 cm² single-turn, transmit/receive butterfly (on a 28 mm diameter plastic former) and a custom-made solenoid coil, were used as described

previously [2, 4]. The broad frequency response of the coils allowed intermittent ${}^{1}H/{}^{19}F$ MRI/MRS. To minimize the B₁ effects, all samples used for in vitro and quantification studies were consistently placed on the coils.

¹⁹*F MRI-Phantom Studies:* a) Aqueous cylindrical phantoms containing trifluoroacetic acid (TFA) (100 and 200mM) and NaF solutions (50–200 mM) [Sigma-Aldrich, UK] were used to optimize the radiofrequency (RF) coil response using the conventional spoiled gradient echo (SPGR) ¹⁹*F* MRI sequences as reported previously [4].

Separate phantoms containing PFCE-NPs (8 mg/ml) sonicated in deionized water and mixed in cell media (IMDM, Thermo Fisher Scientific, UK) to an Eppendorf volume of 1.7 ml comprising a) separate TFA and FuGENE vials (200 μ l), b) separate TFA, NPs, and FuGENE vials, and c) a separate TFA and a pre-mixed NP/FuGENE vial (200 μ l), were used for control phantom validations using ¹⁹F MRS (repetition time (TR)=20s, pw=32–80 μ s, bandwidth (BW)=19.84 kHz, number of excitations (NEX)=16, 512–1024 points), using the solenoid coil [2].

The acquisition parameters for all other in vitro studies were: TR=5.71 ms, echo time (TE)=2.87 ms, flip angle=50°, NEX=256, matrix= 32×32 , dummy scans=100, slice thickness (ST)=15 mm, BW)=6.01 kHz, field-of-view (FOV)= $40-60\times40-60$ mm². For MRS, fully relaxed spectra were acquired for cell studies.

In Vivo Cardiac ¹⁹F MRI and Cellular Tracking

¹*H Imaging:* Double-gated, two-dimensional (2D) multislice, and ungated 2D/3D spoiled grass (SPGR) cardiac ¹*H* images (to match the corresponding ungated ¹⁹*F* MRI) of the mouse thorax were acquired in vivo (2D: TR/TE=1.9-2.13/1-1.1 ms, flip angle= $20-50^{\circ}$, NEX=2-6,

FOV= 40×40 mm², matrix= 128×128 , acquisition time=3-5 min;3D: TR/TE=1.9-2.13/1-1.1 ms, flip angle= $20-50^{\circ}$, NEX=2-6, FOV= 40×40 mm², matrix= $128 \times 128 \times 128$, acquisition time=3-5 min).

¹⁹*F MRI/MRS*: Ungated ¹⁹*F* SPGR images matched ¹H MRI (2D: TR/TE=16.5/8.3 ms, flip angle=50°, NEX=768, FOV=40×40 mm², 1 slice, ST=5 mm, matrix=32×32, BW=2–4 kHz, acquisition time=2.42 min; 3D:TR/TE=16.4/8.3 ms, flip angle=20–30°, NEX=12–72, FOV=40×40 mm², matrix=32×32×32, BW=1–3 kHz, acquisition time=3–19 min). The excitation frequency for ¹⁹F acquisitions was determined based on nonlocalized saturation recovery, gated ¹⁹F MRS of the labeled CPCs acquired a priori (TR=800–1200 ms, 512 points, BW=20kHz, NEX=128–256).

Image Processing

Image and Spectral Analyses: ¹⁹F images were imported and interpolated (bicubic splines) in ImageJ (NIH, Bethesda, MD, USA) to match the ¹H matrix size. Thoracic ¹H and ¹⁹F MRI were overlaid in ImageJ (opacity=30–70%). SNR values were estimated using standard methodologies. For in vitro studies, fluorine concentrations were evaluated by estimating the areas of fully relaxed spectra with respect to a TFA reference standard of known concentration. For tracking studies, the mean signal was estimated as the average ¹⁹F signal from all images in the 3D stack in ImageJ. Spectra were read and processed in CSX (P. Barker, Johns Hopkins, Baltimore, MD, USA) or using custom-written IDL tools (Harris Geospatial Solutions, Broomfield, CO, USA). *Epifluorescence Image Analyses:* No localized fluorescence signals were detected from control (unlabeled) cells. The mean fluorescent intensities quantified in control studies were attributed to noise and auto-fluorescence and matched the mean signal intensities from labeled cells at D4-D7. Labeled cell detection (D1, D4, or D7) was based on fluorescence threshold intensity detection at values above the background intensity (as determined from background non-cellular regions at D1, D4, or D7).

References (Supplementary Material)

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Supplementary Figure Captions

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Supplementary Figure S1: Temporally dependent cellular responses following labeling and

FuGENE labeling. Transverse electron microscopy (TEM) images from control, labeled, and FuGENE-labeled cells. (A-C) Typical EM images from control CT cells harvested on D1 after plating. Sequence of EM images (D1-D3) following (D-F) labeling and (G-I) FuGENE-labeling.

- **Supplementary Figure S2: Study of the temporal label kinetics using dynamic confocal imaging.** Study of the temporal label kinetics using dynamic confocal imaging (labeling [**Movie 1**: acquired at 1.2 h post-label administration], optimal [**Movie 2**: 25 μl per million cells acquired at 5.7 h post-label administration], and high FuGENE doses [**Movie 3**: 100 μl per million cells acquired at 4.15 h post-label administration]) confirming increased (Movie 2), decreased (Movie 1) intracellular label uptake and cluster formations (Movie 3) at high FuGENE doses. Significant label uptake was confirmed within 2-3 h post-label administration. Label uptake was distinguished from transient particle adherence on the extracellular membrane based on 3D visualizations and cross-correlations of single-cell phase contrast and confocal images post-labeling. Indicative 2D examples are depicted in Figure 4.
- Supplementary Figure S3: Control validation experiments using MRS. (A-D) Control MRS experiment confirming that the source of ¹⁹F signal is attributed to FuGENElabeled NPs. In sequence shown are magnitude spectra (drawn to the same scale) of (A) TFA, (B) TFA and FuGENE (separate vials), (C) TFA, FuGENE and NP solution (separate vials), and (D) a real spectrum of TFA and FuGENE-NPs (mixed, same vial) (birdcage coil).

Supplementary Figure S4: FUGENE-labeling and electroporation studies with CDCs using

MRS. (A-D) Pairwise comparisons (A and B, and C and D, drawn to the same scale) of nonlocalized, fully relaxed ¹⁹F MRS responses of electroporated-labeled and FuGENE-labeled, and FuGENE-labeled and electroporated-FuGENE labeled CT cells (butterfly coil). Pairwise comparisons were conducted on separate occasions using single, separate samples in Eppendorf vials positioned at the same location on the butterfly coil.

Supplementary Figures



Supplementary Figure S1: Temporally dependent cellular responses following labeling and FuGENE labeling. Transverse electron microscopy (TEM) images from control, labeled, and FuGENE-labeled cells. (A-C) Typical EM images from control CT cells harvested on D1 after plating. Sequence of EM images (D1-D3) following (D-F) labeling and (G-I) FuGENE-labeling.



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