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In vivo tracking of adenoviral-transduced iron oxide-labeled bone marrow-derived dendritic cells using magnetic particle imaging

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Abstract

Background Despite widespread study of dendritic cell (DC)-based cancer immunotherapies, the *in vivo* postinjection fate of DC remains largely unknown. Due in part to a lack of quantifiable imaging modalities, this is troubling as the amount of DC migration to secondary lymphoid organs correlates with therapeutic efficacy. Magnetic particle imaging (MPI) has emerged as a suitable modality to quantify *in vivo* migration of superparamagnetic iron oxide (SPIO)-labeled DC. Herein, we describe a popliteal lymph node (pLN)-focused MPI scan to quantify DC *in vivo* migration accurately and consistently.

Methods Adenovirus (Ad)-transduced SPIO⁺ (Ad SPIO⁺) and SPIO⁺ C57BL/6 bone marrow-derived DC were generated and assessed for viability and phenotype, then fluorescently labeled and injected into mouse hind footpads (n=6). Two days later, *in vivo* DC migration was quantified using whole animal, pLN-focused, and *ex vivo* pLN MPI scans.

Results No significant differences in viability, phenotype and *in vivo* pLN migration were noted for Ad SPIO⁺ and SPIO⁺ DC. Day 2 pLN-focused MPI quantified DC migration in all instances while whole animal MPI only quantified pLN migration in 75% of cases. *Ex vivo* MPI and fluorescence microscopy confirmed that pLN MPI signal was due to originally injected Ad SPIO⁺ DC.

Conclusion We overcame a reported limitation of MPI by using a pLN-focused MPI scan to quantify pLN-migrated Ad SPIO⁺ and SPIO⁺ DC in 100% of cases and detected as few as 1000 DC (4.4 ng Fe) *in vivo*. MPI is a suitable preclinical imaging modality to assess DC-based cancer immunotherapeutic efficacy.

Relevance statement Tracking the *in vivo* fate of DC using noninvasive quantifiable magnetic particle imaging can potentially serve as a surrogate marker of therapeutic effectiveness.

Key points

- Adenoviral-transduced and iron oxide-labeled dendritic cells are in vivo migration competent.
- Magnetic particle imaging is a suitable modality to quantify *in vivo* dendritic cell migration.
- Magnetic particle imaging focused field of view overcomes dynamic range limitation.

Keywords Adenoviridae, Cell tracking, Immunotherapy, Magnetic iron oxide nanoparticles, Mice (inbred C57BL/6)

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Graphical Abstract

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In vivo tracking of adenoviral-transduced iron oxide-labeled bone marrow-derived dendritic cells using magnetic particle imaging



Mouse dendritic cells (DC) were iron oxide-labeled (SPIO+) or transduced with an adenovirus and then SPIO-labeled (Ad SPIO+). Magnetic particle imaging using a 12-cm whole animal field of view (FOV) identified SPIO+ DC (grey) and Ad SPIO+ DC (green) at the hind footpad injection site on day 0 (a) and day 2 (b). Injection site signal prevented lymph nodemigrated SPIO+ DC quantification (b, red). A 2-cm FOV permitted quantification of lymph nodemigrated SPIO⁺ and Ad SPIO⁺ DC (c, grey and green) and was confirmed through ex vivo magnetic particle imaging of excised popliteal lymph nodes (d)

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Magnetic particle imaging is a noninvasive modality suitable for in vivo tracking of dendritic cells. It has the potential to serve as a surrogate marker of therapeutic effectiveness.

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Background

For cancer patients who are nonresponsive to or metastatically progress after standard of care treatment options, cancer immunotherapy remains an available treatment option [1]. Cancer immunotherapy initiates the enhancement of one's own immune system to combat cancer. As such, ex vivo-generated autologous dendritic cells (DC) can be exploited as professional antigenpresenting cells (APC) to express tumor-associated antigens (TAA) in conjunction with costimulatory molecules required to initiate a potent TAA-specific immune response upon injection into the host [2]. Delivery of TAA to DC can occur via different mechanisms, but viral transduction of a TAA-expressing vector remains an efficient delivery system [3]. The immune response elicited by this DC-based cancer vaccine strategy includes the direct activation of TAA-specific CD4⁺ and CD8⁺ T cells in conjunction with indirect activation of innate immune cells, namely natural killer (NK) cells, NK T cells, and macrophages [4]. Despite the potential of DC to orchestrate robust TAA-specific immune responses [5], their treatment effectiveness remains suboptimal [6] and the in vivo fate of therapeutic DC following administration remains largely unknown.

The paucity of knowledge on in vivo DC localization is troubling. First, DC migration to secondary lymphoid organs post injection and subsequent presentation of TAA to prime T cells post injection is required and secondly, quantification of this DC migration correlates with the rapeutic potency [7, 8]. When considering that a limitation of treatment effectiveness stems from < 5% of injected DC reaching secondary lymphoid organs [6, 9, 10], the necessity for noninvasive quantifiable DC in vivo tracking is needed to both assess and improve upon vaccine outcomes.

European Radiology

EXPERIMENTAL

We and others have employed magnetic resonance imaging (MRI) as a noninvasive imaging modality to track both human and mouse DC in vivo in preclinical and clinical studies [11-15]. By labeling DC ex vivo with superparamagnetic iron oxide nanoparticles (SPIO), SPIO⁺ DC are identifiable in vivo as regions of signal hypointensities resulting from the effect of SPIO⁺ DC on surrounding proton relaxation. Although highly sensitive [16], quantification of SPIO⁺ DC with MRI remains semiquantitative [17] and low in specificity as signal hypointensities inherent in certain tissues cannot be discriminated from SPIO⁺ DC-induced hypointensities [18]. Magnetic particle imaging (MPI) has now emerged as a

preclinical small animal alternative to MRI cell tracking of SPIO⁺ cells [19]. Unlike MRI, MPI signal is quantifiable as it is linearly proportional to iron mass. Tissue/bone hypointensities that complicate MRI are also nonexistent with MPI [20]. Low resolution and biocompatibility of SPIO are noted limitations of MPI [21, 22]; yet hardware upgrades [23] and the development of new MPI-tailored SPIO formulations are ongoing [24]. We previously tracked in vivo SPIO⁺ bone marrow-derived DC (BMDC) migration with MPI [14]. However, a major challenge has been the limited dynamic range of MPI when there are two signals of interest in the field of view (FOV) with different iron concentrations. In our preclinical model, the high iron concentration from administered BMDC at the injection site shadows the lower iron concentration from BMDC in the draining lymph node (LN), reducing the ability to separate and accurately quantify signals. Due to its broad applicability across a range of disease states and with the first human MPI scanner designs increasingly becoming a reality [25], preclinical studies optimizing the tracking of SPIO⁺ BMDC is of timely importance.

Herein, we designed a combined protocol to generate adenoviral-transduced SPIO-labeled BMDC without affecting viability, phenotype or *in vivo* migration to popliteal LN (pLN) detected by MPI. Additionally, by combining the use of a focused FOV centered on pLN with a multichannel joint image reconstruction algorithm, we are the first group to successfully overcome the dynamic range limitation of MPI to permit accurate quantification of pLN-migrated BMDC. MPI is a suitable preclinical imaging modality to noninvasively assess DC-based cancer vaccine efficacy.

Methods

Animals, reagents, and antibodies

C57BL/6 mice were purchased from Charles River Laboratories, Inc. (Senneville, CAN). All animal studies were performed in accordance with institutional and national guidelines. Supplemental Table S1 contains a complete list of reagents and antibodies used in this study.

Mouse bone marrow-derived dendritic cell (BMDC) generation

BMDC were generated from C57BL/6 mice femurs and tibias as previously detailed [17, 26] with the following modifications. Briefly, BMDC progenitors were cultured in complete RPMI media containing interleukin (IL)-4 and granulocyte-macrophage colony-stimulating factor (GM-CSF) at 10 ng/mL and 4 ng/mL, respectively for four days at 37°C and 5% CO₂. After four days, immature BMDC were enriched from heterogenous cell cultures by HistodenzTM gradient centrifugation.

Adenoviral transduction

Immature day 4 BMDC were transduced with a replication-defective recombinant human adenovirus (Ad) type 5 expressing enhanced green fluorescent protein (eGFP) as a pseudo-TAA [27] at a multiplicity of infection (MOI) of 30 for 2 h at 37 °C and 5% CO₂. BMDC incubated in media lacking Ad served as the untransduced control BMDC.

SPIO labeling and BMDC maturation

Following Ad transduction, Ad BMDC and untransduced BMDC were labeled as previously described with the SPIO FeREX[®] (used in Fig. 1 only) and Synomag[®]-D. Synomag[®]-D is a 50 nm dextran-encapsulated cell labeling agent with a nanoflower-shaped iron oxide core and is supplied as a suspension in water [13, 14, 28]. Briefly, FeREX[®] alone or Synomag[®]-D SPIO was complexed with protamine sulfate (0.24 mg/mL) and heparin (8 USP units/mL) in incomplete RPMI and coincubated with BMDC at a final concentration of 200 µg Fe/mL for 4 h at 37 °C and 5% CO₂. IL-4- and GM-CSF-supplemented complete RPMI was then added to immature day 4 BMDC culture and incubated overnight at 37 °C and 5% CO₂. On day 5, BMDC were further matured [12] in culture for 24 h.

BMDC magnetic column enrichment

On the sixth and final culture day, SPIO⁺ BMDC were enriched via magnetic column separation [14]. After collection and repeated washing in PBS to remove free SPIO, BMDC were resuspended in PBS in a 12×75 mm polypropylene tube and placed in a EasySepTM magnet (Stemcell Technologies, Vancouver, CAN) for 5 min. Both tube and magnet were inverted to collect BMDC not labeled with SPIO (unlabeled, UL) that remained in the flow through. Conversely, both Ad-transduced and untransduced BMDC that incorporated SPIO were retained in the tube and were subsequently collected for downstream application as Ad SPIO⁺ and SPIO⁺ BMDC, respectively.

Immunophenotyping and viability assessment

A previously used protocol from our group was employed with modifications to assess day 6 mature BMDC viability and phenotype [26] using the fluorescent antibodies and dyes listed in Supplemental Table S1. Zombie NIRTM fixable vital dye was added to BMDC at room temperature for 20 min in PBS to determine viability. Next, BMDC underwent surface immunofluorescence staining in the presence of TruStain FcXTM anti-mouse CD16/CD32 block at 4 °C for 25 min. Extensive washing in HBSS and 0.1% bovine serum albumin ensued prior to fixation in



Fig. 1 Ad-transduced and SPIO-labeled BMDC are *in vivo* migration competent. CD11c⁺ bone marrow-derived dendritic cells (BMDC) were either adenovirus- (Ad)-transduced and then superparamagnetic iron oxide nanoparticle- (SPIO)-labeled (**a**, Ad SPIO⁺, green histogram) or labeled with SPIO and then Ad-transduced (**b**, SPIO⁺ Ad, gray histogram). Black histograms in **a** and **b** depict minimal fluorescence observed with untransduced SPIO⁺ BMDC. Ad SPIO⁺ transduced at a multiplicity of infection (MOI) = 30 and MOI = 60 both resulted in a significantly higher percentage of eGFP⁺ BMDC compared to MOI = 10. No further increases in eGFP⁺ BMDC percentage were noted at MOI = 60 when compared to MOI = 30 (**c**). Increasing the MOI did not significantly improve eGFP⁺ BMDC percentage in SPIO⁺ Ad condition (**d**). Therefore, the Ad SPIO⁺ (MOI = 30) (**c**, middle bar) culture condition was employed for all subsequent experiments. Data are shown as mean ± SEM for *n* = 3 independent experiments (**c**, two-way ANOVA, * *p* = 0.0053, ** *p* = 0.0084, **c** and **d**, ns-no statistical significance). One million Ad SPIO⁺ BMDC were fluorescently labeled with PKH26 immediately before hind footpad adoptive transfer. Two days later, excised popliteal lymph nodes (pLN) were processed for digital morphometric analysis of PKH26 (**e**, red bar) and eGFP (**e**, green bar) fluorescence, with representative fluorescence microscopy images shown in Supplemental Figure S1. The percentage of total BMDC that were eGFP⁺ at the time of injection (**c**, MOI = 30) is comparable to the percentage of pLN migration competent BMDC that were eGFP⁺ **e** Data is representative of *n* = 2 independent experiments with four mice per group (**e**, *t*-test, ns-no statistical significance).

4% paraformaldehyde and storage at 4 °C. When necessary, eGFP expression due to Ad transduction and fluorescent membrane-intercalating dye incorporation (detailed below) was assessed using flow cytometry and compared to an aliquot of nonfluorescent BMDC. Data was acquired on a LSRII analytical flow cytometer (BD Biosciences, San Jose, USA) and analyzed using FlowJo Software (v10, Tree Star, Inc., Ashland, USA).

Adoptive cell transfer (ACT)

Following magnetic enrichment and immediately prior to ACT, SPIO⁺ and Ad SPIO⁺ BMDC were fluorescently labeled with Tag-It VioletTM (Violet) cell tracking dye (5 μ M) using an established protocol for membraneintercalating dyes [29]. Low (3×10⁵ cell) and high (5×10⁵ cell) injection doses of Violet⁺ SPIO⁺ BMDC were prepared in 40 μ L PBS and subcutaneously injected into the right hind footpad of C57BL/6 mice. An identical number of Violet⁺ Ad SPIO⁺ BMDC were injected into the contralateral hind footpad (*n*=3 mice per injection dose).

In a separate experiment, Ad SPIO⁺ BMDC were labeled with PKH26 as described previously [13] prior to hind footpad ACT of 1×10^6 PKH26⁺ Ad SPIO⁺ BMDC (*n*=2 independent experiments with 8 total mice).

Magnetic particle imaging (MPI)

Mice were imaged as previously described [14] using a MomentumTM MPI scanner (Magnetic Insight Inc., Alameda, USA) on days 0 and 2 post-injection using imaging

parameters described below. Prior to imaging, mice were fasted for 12 h with water, a laxative, and corn bedding to mitigate gastrointestinal (GI) iron signal in MPI images. Mice were anesthetized initially with 2% isoflurane and maintained with 1% isoflurane during imaging. Afterwards, mice were returned to cages with food *ad libitum*.

Day 0

Post-injection images were acquired in 3D using 3.0 Tesla/m (T/m) selection field gradient and drive field strengths of 20 mT and 23 mT in the X and Z axes, respectively, for isotropic imaging. The MPI signal was collected from 35 projection angles equally distributed between 0° and 180°. A 12 cm (Z)×6 cm (X) field of view (FOV) was centered on the mouse body and encompassed the entire imaging area without any signal being present at the edges of the FOV. These images were reconstructed using the prescribed native reconstruction algorithm equipped in the "regular user interface". The acquisition time for full 12 cm (Z) FOV images was 40 min.

Day 2

Images were acquired using two different imaging sequences. Initially, mice were imaged using the same 3D image parameters and the reconstruction described above. Subsequently, mice were imaged again in 3D using a 3.0 T/m selection field gradient and drive field strengths of 20 mT and 23 mT in the X and Z axes, respectively,

with 35 projection angles. Single channel imaging was used to collect information along the Z axis only. The MPI signal was collected from 35 projection angles equally distributed between 0° and 180°. A smaller 2 cm $(Z) \times 6$ cm (X) FOV was used to only image low signal from BMDC that migrated to pLN and excluded impeding stronger signal from the injection site. These images were reconstructed using a multichannel joint algorithm where an inverse combiner was enabled to prevent image artifacts due to MPI signal being present at the FOV edges [30]. The acquisition time for focused 2 cm (Z) FOV images was 15 min. These customized image parameters were implemented using the "advanced user interface" which allowed the user access to a configuration editor where changes were made to prescribed sequences equipped on the MPI system. Excised pLN were then imaged using a full 12 cm (Z) FOV.

Analysis and quantification

Two samples containing 1×10^6 SPIO⁺ and 1×10^6 Ad SPIO⁺ BMDC were singly imaged using the 12 cm (Z) and 2 cm (Z) FOV 3D image parameters described above. Iron loading per cell was measured from these samples by converting the measured MPI signal to iron mass using calibration lines unique to the image sequence, as described [14]. This iron per cell measurement was subsequently used for all in vivo and ex vivo image analysis to estimate cell numbers from MPI signal. All MPI images were imported into Horos[™], an open-source clinically relevant image analysis software (version 3.3.6, Annapolis, USA). Images were viewed using a custom MPI color look-up table. MPI signal was measured within a specific region of interest (ROI) using a semiautomatic segmentation tool. Total MPI signal for an ROI was calculated by multiplying the ROI volume by the mean signal. The signal to noise ratio (SNR) was calculated by dividing the mean signal for a ROI by the standard deviation of the background noise. The SNR had to be greater than 5 for the MPI signal to be considered detectable and for images to be further quantified. Iron content was calculated by dividing the total MPI signal by the slope of the respective calibration line for those image parameters. All MPI images were delineated and analyzed in the same way to ensure consistency. Cell numbers were estimated by dividing the iron quantified within a given ROI by the respective iron/cell measurements.

Popliteal lymph node (pLN) histology

Upon final *in vivo* MPI imaging session completion, pLN were excised from euthanized mice, fixed with 4% paraformaldehyde, and promptly underwent *ex vivo* MPI. Next, pLN were cryoprotected in sucrose and processed into 16 µm tissue sections [26] for imaging using an EVOSTM M7000 Imaging System (Thermo Fisher Scientific, Burlington, CAN). For quantitative image analysis, pLN sections were imaged for Violet fluorescence. Digital morphometric analysis of imaged pLN sections was performed using Image Pro Premier 9.2 software (Media Cybernetics, Rockville, USA) and represented as area of Violet fluorescence/area of lymph node.

Similarly to above, pLN were excised and processed from mice that received hind footpad injections of PKH26⁺ Ad SPIO⁺ BMDC two days prior. A Leica DMIRE2 fluorescence microscope (Leica, Wetzlar, GER) and Image Pro software were used to perform digital morphometric analysis of pLN sections [17] to determine the area of PKH26⁺ and eGFP⁺ fluorescence per area of lymph node.

Statistical analysis

All data is presented as means \pm standard error of the mean (SEM). A paired *t*-test, two-way ANOVA with Tukey's multiple comparison test, and simple linear regression analysis (Graph Pad Prism, Version 9.4.1, La Jolla, USA) were used. Statistical significance was defined when $p \leq 0.05$.

Results

SPIO labeling and BMDC viability are unaffected by prior adenoviral transduction

Although Ad transduction [3] as well as SPIO labeling of dendritic cells [18] are both widely reported on in the context of cancer immunotherapy, the outcome of Ad transduction combined with SPIO labeling within one BMDC generation protocol was unknown. Preliminary experiments demonstrated that BMDC transduction prior to SPIO labeling (Ad⁺ SPIO, MOI=30) yielded a significantly higher transduction percentage $(40.9 \pm 4.4\%)$ than observed at MOI=10 $(23.7 \pm 4.8\%)$ and further MOI increases did not improve the transduction percentage (Fig. 1a, c). SPIO labeling prior to transduction (SPIO⁺ Ad) was associated with inferior transduction efficiencies compared to Ad⁺ SPIO BMDC (Fig. 1b, d). Importantly, the percentage of Ad-transduced BMDC (PKH26⁺ eGFP⁺) of total pLN-migrated Ad SPIO⁺ BMDC (PKH26⁺) was comparable to the percentage of Ad-transduced BMDC at the time of injection (Fig. 1c and Supplemental Figure S1). This indicated that Adtransduction does not impair in vivo BMDC migration. Next, the viability of singlet, CD45⁺CD11c⁺ BMDC was measured using flow cytometry and Zombie $\text{NIR}^{^{\text{TM}}}$ fixable vital dye (Fig. 2a-d). SPIO⁺ and Ad SPIO⁺ BMDC were associated with a small but significant decrease in viability compared to UL BMDC (Fig. 2e, gray and green

bars compared to white bar, p < 0.001 and p = 0.002); however, BMDC viability was consistently > 90% across all conditions. More importantly, cell viability was equivalently high for Ad SPIO⁺ BMDC (93.5 ± 0.82%) and for untransduced SPIO⁺ BMDC (93.2 ± 0.74%, mean ± standard error of the mean (Fig. 2e, p = 0.823). As expected, UL and untransduced SPIO⁺ BMDC expressed negligible eGFP (Fig. 2f, open and gray histogram, respectively) compared to ~ 30% of Ad SPIO⁺ BMDC that expressed eGFP (Fig. 2g, green histogram). By comparing cell counts prior to and immediately after magnetic column enrichment, the SPIO labeling percentage was calculated and did not significantly differ between SPIO⁺ (78 ± 8.1%) and Ad SPIO⁺ (67 ± 8.6%) BMDC (Fig. 2h, p=0.135), suggesting that our currently outlined Ad transduction and SPIO labeling protocol is suitable for continued use.



Fig. 2 BMDC maintained high viability after Ad transduction and efficiently labeled with SPIO. The viability of singlet (**a**), CD45⁺ (**b**), CD11c⁺ (**c**) BMDC was measured by Zombie NIRTM fixable viability dye and flow cytometry (**d**). No significant differences in viability were measured between superparamagnetic iron oxide nanoparticle-labeled (SPIO⁺) BMDC (**e**, gray bar) and adenovirus- (Ad)-transduced SPIO⁺ (**e**, green bar), although the viability of both populations exhibited a small but significant decrease compared to untransduced and unlabeled (UL) BMDC (**e**, white bar). The viability was > 93% for all conditions. Unlabeled (UL) (**f**, open histogram) and SPIO⁺ (**f**, gray histogram) BMDC have negligible enhanced green fluorescent protein (eGFP) expression while Ad SPIO⁺ BMDC express eGFP following transduction with an Ad encoding eGFP (**g**, green histogram). Prior transduction did not affect the percentage of BMDC that successfully labeled with SPIO (**h**). Data are shown as means ± SEM for *n* = 5 independent experiments (**h**, paired *t* test, ns–no statistical significance, *p*=0.135)

An important characteristic of a cell labeling agents is efficient incorporation into cells of interest without impacting cell phenotype. Within our set-up, mature BMDC were defined as $CD11c^+CD86^+$ (Fig. 3a) and their percent expression did not significantly differ between UL (68.9 ± 4.5%), SPIO⁺ (72.8 ± 2.8%) and Ad SPIO⁺ (74.6 ± 3.2%) BMDC (Fig. 3b, white, gray and green bars, respectively). This trend persisted across a range of cell surface markers critical for therapeutic BMDC function and included markers of BMDC migration (Fig. 3c, CCR7), activation (Fig. 3d, CD40), antigen presentation (Fig. 3e, I-A^b) and T cell stimulation (Fig. 3f/g, ICOSL



Fig. 3 BMDC phenotype is unaffected by SPIO labeling or Ad transduction. Representative flow cytometric dot plots to identify singlet BMDC expressing both CD11c and CD86 (CD11c⁺CD86⁺) are shown in (**a**). The percentage of mature BMDC (identified as CD11c⁺CD86⁺) did not differ between untransduced and unlabeled (UL) BMDC (**b**, white bar), superparamagnetic iron oxide nanoparticle-labeled (SPIO⁺) BMDC (**b**, gray bar) and adenovirus- (Ad)-transduced SPIO⁺ BMDC (**b**, green bar). Neither SPIO labeling or Ad transduction altered the phenotype of CD11c⁺CD86⁺ mature BMDC as assessed by markers of migration (**c**, CCR7), activation (**d**, CD40), MHC Class II antigen presentation (**e**, I-A^b) and T cell stimulation (**f** and **g**, ICOSL and OX40L, respectively). Data are shown as means ±SEM for *n* = 5 independent experiments (**b**–**g**, two-way ANOVA, * *p* < 0.05)

and OX40L). In addition to no phenotypic changes due to Ad transduction or SPIO incorporation, >92% of CD11c⁺CD86⁺ mature BMDC remarkably expressed all phenotypic markers in question, furthering support for our combined Ad transduction and SPIO labeling protocol as we progressed to *in vivo* MPI studies.

In vivo detection of SPIO⁺BMDC and Ad SPIO⁺BMDC migration using 3D MPI

We next sought to measure SPIO⁺ and Ad SPIO⁺ BMDC *in vivo* migration with MPI. This longitudinal MPI study is outlined schematically in Fig. 4 with quantified data that is representative of 5 independent experiments presented in Fig. 5.

On day 0, scanning was performed immediately after ACT using a 12-cm FOV (Fig. 4a) with 3D MPI signal shown in full dynamic range. MPI signal was localized to the right and left hind footpad injection sites due to SPIO⁺ BMDC and Ad SPIO⁺ BMDC, respectively

(Fig. 5a, white and green bracket, respectively, 3×10^5 cell injection dose each). Two days later, the same FOV (Fig. 4b) and imaging parameters identified a distinct region of interest (ROI) at the left pLN region quantified as 1.9×10^3 Ad SPIO⁺ (green bracket) but the signal at the right pLN region (white bracket and red asterisk) could not be separated from the high signal emanating from the footpad injection sites (Fig. 5b). The 3D MPI signal window leveled to the pLN region results in oversaturation of both GI-localized and persistent injection site signal (Fig. 5b, orange arrow, gray and green brackets). Despite animals being fasted prior to imaging, GI signal was visible due to residual food and iron contained within consumed bedding. To combat the difficulty of resolving pLN signal near a source of higher signal intensity [14], a second day 2 scan using a 2-cm focused FOV (Fig. 4c) centered on the pLN was conducted. MPI signal was clearly present in the right and left pLN regions and guantified as 1.0×10^3 SPIO⁺ and 2.0×10^3 Ad SPIO⁺ BMDC, respectively (Fig. 5c, white and green brackets,



Fig. 4 Comparison of FOV size to detect and quantify *in vivo* SPIO⁺ BMDC pLN migration. On day 0, SPIO⁺ and Ad SPIO⁺ BMDC were adoptively transferred into the hind footpads of C57BL/6 mice immediately prior to MPI using a 12-cm whole animal FOV (**a**, pink box). The mouse orientation within the scanner is shown and overlaid with MPI signal. Imaging with a 12-cm FOV (**b**, pink box) was then repeated two days later and immediately followed by MPI of the pLN region using a 2-cm FOV (**c**, pink box). Mice were euthanized to excise pLN that were then placed in a 0.5 mL polypropylene tube before a final *ex vivo* MPI session using a 12-cm FOV (**d**, pink box). **Note that MPI signal was enhanced in all panels for the purpose of creating a schematic. All other MPI signal presented in this manuscript is not enhanced



Fig. 5 MPI detection and quantification of *in vivo* SPIO⁺ and Ad SPIO⁺ BMDC migration to pLN. On day 0, SPIO⁺ BMDC (**a**, gray outline) and Ad SPIO⁺ BMDC (**a**, green outline) were detected at the injection site (hind footpads, 3×10^5 BMDC injection dose) immediately following adoptive cell transfer. Two days later, MPI using a 12-cm FOV was unable to quantify SPIO⁺ BMDC migration to the pLN (**b**, gray bracket and red asterisk) but quantified pLN-migrated Ad SPIO⁺ BMDC (**b**, green brackets). GI signal is denoted by the orange arrow in **b**. In contrast, a 2-cm FOV MPI scan detected and quantified both SPIO⁺ BMDC and Ad SPIO⁺ BMDC migration to the pLN (**c**, gray and green brackets, respectively). pLN were then excised from animals and underwent *ex vivo* MPI to detect pLN-migrated SPIO⁺ BMDC (**d**, gray bracket) and Ad SPIO⁺ BMDC (**d**, green bracket). MPI images are slices selected from 3D imaging using 35 projections. Data is representative of *n*=5 independent experiments

respectively). Postmortem MPI using a 12-cm FOV (Fig. 4d) on excised pLN validated *in vivo* signal (Fig. 5d, white and green bracket, respectively).

Focused small FOV MPI with multichannel joint image reconstruction is superior to whole animal FOV MPI for quantifiable detection of pLN-migrated BMDC

For the longitudinal MPI experiment detailed above, day 2 MPI images are shown for all animals that received either a low (Fig. 6a, 3×10^5 cells) or high (Fig. 6b, 5×10^5 cells) injection dose. *In vivo* pLN-localized MPI signal is shown for all animals following acquisition using a 12-cm and focused 2-cm FOV, respectively (Fig. 6a/b, left and middle panels, respectively). Excised pLN were also imaged using a 12-cm FOV for *in vivo* validation purposes (Fig. 6a/b, right panels). Using a 12-cm FOV, quantifiable signal was detected in only 4/6 right pLN and 5/6 left pLN as summarized for SPIO⁺ and Ad SPIO⁺ BMDC (Fig. 6c, gray and green bars). This is in stark contrast to

our ability to confidently detect quantifiable signal in 6/6 right pLN and 6/6 left pLN using a focused 2-cm FOV with multichannel joint reconstruction (Fig. 6d) and from excised pLN set-up (Fig. 6e) that is summarized for SPIO⁺ and Ad SPIO⁺ BMDC (Fig. 6d/e, gray and green bars). Irrespective of the FOV employed or the injection dose, there was no significant difference between the number of pLN-migrated SPIO⁺ BMDC and Ad SPIO⁺ BMDC, although the data does trend that increased injection dose leads to increased pLN-migrated BMDC (Fig. 6c-e). In summary, we have demonstrated that a focused 2-cm FOV with multichannel joint reconstruction results in quantifiable signal detection in 100% of pLN and is a noted improvement to 75% of pLN with quantifiable signal detected using a 12-cm FOV. It is also important to note that signal quantification using either a 12-cm or 2-cm FOV remains strongly linearly correlated to ex vivo pLN quantification (Fig. 6f/g, $R^2 = 0.91$ and $R^2 = 0.88$).

MPI pLN signal detection is due to originally injected SPIO⁺ and Ad SPIO⁺ BMDC

By rendering both SPIO⁺ and Ad SPIO⁺ BMDC Violet fluorescent (Violet⁺) immediately prior to injection, excised pLN can be imaged after terminal MPI to identify Violet⁺ BMDC and confirm that *in vivo* pLN MPI signal is the result of originally injected BMDC. Compared to an aliquot of BMDC removed prior to labeling, 100% of SPIO⁺ and Ad SPIO⁺ BMDC were Violet⁺. Moreover, the Violet mean fluorescence intensity was consistent across both cell populations and thus, is a suitable readout for comparing pLN migration (Fig. 7a, b). Upon completion of day 2 MPI, representative pLN cryosections are shown for Violet⁺ SPIO⁺ and Violet⁺ Ad SPIO⁺ BMDC from both low and high injection doses (Fig. 7c, e and Fig. 7d, f, respectively). In all instances, Violet⁺ BMDC have localized to the central and paracortical T cellrich areas of the pLN (Fig. 7c-f). Digital morphometry then revealed that SPIO⁺ BMDC and Ad SPIO⁺ BMDC migrate equivalently to the pLN two days post ACT for both low (Fig. 7g, p=0.470) and high (Fig. 7h, p=0.539) injection doses, which is in direct agreement with in vivo migration quantified using MPI (Fig. 6).

Discussion

For DC-based immunotherapies, the lack of standardization and scientific consensus surrounding administration route, injection dose, and optimal DC source [31, 32] means that MPI must be robust enough to detect SPIOlabeled DC regardless of progenitor cell origin. Our prior studies highlighted a limitation of MPI observed when attempting to resolve a strong and weak ROI near each other [14]. To optimize therapeutic DC tracking with MPI, this study describes the efficient labeling of differently generated (Ad-transduced and untransduced) SPIO⁺ BMDC and the use of an image reconstruction algorithm that allows for the choice of a focused FOV to detect and quantify SPIO⁺ DC migration to the pLN 100% of the time despite its nearness to high injection site hind footpad signal.

We have previously reported efficient MRI cell labeling agent uptake by immature DC in the absence of transfection agents [26]. However, transfection agents have been employed to enhance SPIO label uptake and thus, improve MRI/MPI sensitivity to detect therapeutically relevant number of SPIO⁺ BMDC in vivo [33]. Despite reports of transfection agents resulting in cytotoxicity [34], protamine sulfate and heparin were successfully used in this study as CD11c⁺ BMDC viability was>90% for SPIO⁺ and Ad SPIO⁺ BMDC (Fig. 2e). SPIO nanoparticles for MPI are continuing to be developed with a focus on improving biocompatibility, for example, through surface nanoparticle modifications or liposomal delivery [34, 35] that will improve image resolution and may circumvent the need for transfection agents altogether. Also, we used a naked/dextran-coated SPIO [28] to label DC and is most likely a contributing factor to the reported 70-80% of DC that incorporated SPIO (Fig. 2h). Magnetic column enrichment controlled for this reduced labeling percentage by ensuring that only SPIO⁺ and Ad SPIO⁺ DC were characterized and adoptively transferred. Biocompatible SPIO nanoparticles, in addition to improving image resolution [36], will not only improve cell loading but also will improve the percentage of cells that are labeled and is a focus of our future studies.

An important aspect of this study was to demonstrate that BMDC can be dually labeled by Ad and incorporate SPIO nanoparticles without compromising their subsequent ability to migrate *in vivo*. Ad are extensively used for ex vivo and in vivo DC transduction as they are nonreplicating, nonintegrating vectors associated with efficient and sustained gene delivery [10, 37] that influences the effectiveness of DC-based cancer vaccines [38]. While eGFP served as a pseudo TAA in this study, Ad transgenes have consisted of a single TAA [39], multiple TAAs to combat antigen loss variants [10] as well as immunostimulatory molecules like CD40, CD137L, 4-1BBL, and OX40L [39-42] to activate DC and TAAspecific T cells, respectively. Despite their widespread use, Ad DC transduction still yields conflicting results [43], potentially due to the lack of standardization in generating and the source of therapeutic DC. This may explain why the percentage of Ad-transduced CD11c⁺

(See figure on next page.)

Fig. 6 Focused small FOV MPI to accurately quantify *in vivo* BMDC signal. *In vivo* MPI images of the pLN region acquired using a 12-cm FOV (**a**, left panels) and 2-cm FOV (**a**, middle panels) are shown for three mice that received low dose $(3 \times 10^5 \text{ SPIO}^+ \text{ and Ad SPIO}^+ \text{ BMDC})$ hind footpad injections. Corresponding *ex vivo* pLN MPI images are also shown (**a**, right panels). MPI images of three mice that received high dose $(5 \times 10^5 \text{ SPIO}^+ \text{ and Ad SPIO}^+ \text{ BMDC})$ injections are displayed in **b** using the same layout described in **a**. Red asterisks indicate pLN in which BMDC migration could not be quantified using a 12-cm FOV (**a** and **b**, left panels) but were quantified using a 2-cm FOV and *ex vivo* MPI set-up (**a** and **b**, middle and right panels, respectively). Images are shown window-leveled to the dynamic range of the pLN. MPI quantification of pLN-migrated BMDC determined using 12-cm FOV (**c**), 2-cm FOV (**d**) and *ex vivo* MPI (**e**) are summarized for low and high dose SPIO⁺ (**c**-**e**, gray bars) and Ad SPIO⁺ (**c**-**e**, green bars) BMDC injection conditions. Only signal that was ≥ 5 times the standard deviation above background signal was used for quantification. Linear regression analysis revealed a strong correlation between 12-cm FOV and *ex vivo* pLN MPI quantification (**f**, R^2 = 0.91) for pLN that were able to be quantified using a 12-cm FOV. In contrast, BMDC migration to all pLN was successfully quantified with a 2-cm FOV and strongly correlated with *ex vivo* pLN MPI quantification (**g**, R^2 = 0.88). Data is representative of n = 5 independent experiments



Fig. 6 (See legend on previous page.)

BMDC presented here and reported by others [44, 45] is lower than what has been reported in the literature for DC in general [3]. Discrepancies in transduction efficiency between BMDC and DC derived from other sources is primarily attributed to low expression of the coxsackie and adenovirus entry receptors on BMDC [46]. Moreover, we presented a dual Ad transduction SPIO labeling protocol and this combination constrains the modifications that we can make to improve transduction. Ad transduction prior to SPIO labeling was performed based on our earlier studies (Fig. 1) and importantly, the presence of heparin during SPIO labeling likely inhibits further transduction after the two-hour initial incubation has elapsed [47]. Extending the transduction incubation time would likely result in increased transduction efficiency but also Ad-induced DC maturation [46] that would lead to maturing DC subsequently incorporating less SPIO. Although optimizations are necessary (i.e., increasing MOI to improve transduction efficiency [43]), this combined protocol is a crucial proof-of-principle study that differently matured DC can efficiently label with SPIO and be tracked in vivo with MPI.

Successful DC tracking is contingent on sufficient cell labeling agent incorporation to permit in vivo detection without impacting mature DC viability, phenotype, and function [48]. Analysis of BMDC phenotype revealed that neither Ad transduction or SPIO labeling affected CCR7 expression and by extension, BMDC ability to migrate in vivo, as nearly 100% of SPIO⁺ mature CD11c⁺CD86⁺ BMDC were CCR7⁺ (Fig. 3c). Although additional CCR7-independent markers of DC migration such as CX3CL1 and CXCL12 have been described [49], CCR7 remains the most important DC migration marker. For this reason, prostaglandin E_2 (PGE₂) is included in the day 5 maturation cocktail despite its negative effect on IL-12 secretion, which along with IFN-y, promotes a $T_{\rm H}$ 1-mediated immune response [32, 50]. In stark contrast to a previous BMDC study [51], we report a remarkably high (near 100%) surface expression of CD40 on SPIO⁺ and Ad SPIO⁺ BMDC (Fig. 3d) which should elicit IL-12 production in secondary lymphoid organs following interaction with CD40L on CD4⁺ T_{H} cells [48]. In addition to the CD40:CD40L and I-A^b:T cell receptor (TCR) interactions between DC and T cells, >92% of BMDC coexpress OX40L and ICOSL (Fig. 3e-g) to interact with OX40 and ICOS, respectively, on activated T cells [52, 53]. This was expected as TNF- α is a component of the maturation cocktail and induces OX40L and ICOSL upregulation [1, 42]. It is imperative that DC highly express OX40L as interaction with OX40 leads to IFN- γ production [1, 54]. The mature and primed phenotype described above suggests that in a DC-based cancer vaccine setting, both SPIO⁺ and Ad SPIO⁺ BMDC should elicit a strong and sustained T_H1-mediate response. In conjunction with broad innate immune cell activation conferred by professional APC like DC, this would establish a proinflammatory environment conducive for TAA-specific CD8⁺ T cell activation and proliferation [42, 52, 53] that is an absolute requirement for immunotherapeutic effectiveness.

Irrespective of DC source, generation, TAA loading or dose, knowledge of where and in what quantities DC migrate to post injection is required to evaluate DC-based cancer immunotherapy effectiveness [8]. Iron oxide-based MRI has been the most successful noninvasive imaging modality to date for DC tracking owing to the extensive anatomical information it provides alongside its high sensitivity of detection. SPIO⁺ DC in vivo produce T2-weighted decreases to SNR capable of detecting as little as ~ 2×10^3 SPIO⁺ cells under preclinical ¹H MRI conditions and permitting comparison of migration differences associated with differently matured DC [12, 18]. Despite its popularity, disadvantages have been widely reported and generally fit into two categories. First, SPIO⁺ DC-induced signal void is not specific and cannot be distinguished from tissues whose chemical shift artifacts generate signal loss [1, 18]. Secondly, as SPIO⁺ DC signal void is not linearly related to iron concentration, it is at best a semiguantitative imaging modality [17] and is limited in its ability to accurately predict treatment outcome.

Recently, MPI has emerged as an exciting preclinical imaging modality capable of directly quantifying the amount of iron, and by extension, the amount of SPIO⁺ cells, within a ROI [14]. This permits for "hot spot" imaging similarly to what is seen for ¹⁹F MRI [26, 29, 55] and overcomes one of the main limitations of ¹H MRI detailed above. MPI does not provide any anatomical information. However, it is feasible to combine with a second imaging modality to provide anatomical context

⁽See figure on next page.)

Fig. 7 Adoptively transferred SPIO⁺ and Ad SPIO⁺ BMDC are source of MPI signal detected in pLN. Magnetically enriched superparamagnetic iron oxide-labeled (SPIO⁺) and adenovirus- (Ad)-transduced SPIO⁺ BMDC were labeled with Tag-It Violet (Violet⁺, **a** and **b**, respectively) and formulated into low (3×10^5) and high (5×10^5) hind footpad injections. Two days later, pLN were excised and cryosectioned to reveal Violet fluorescence for low (**c**) and high (**e**) SPIO⁺ BMDC and low (**d**) and high (**f**) Ad SPIO⁺ injection doses, respectively. Violet⁺ pLN fluorescence was quantified and did not significantly differ between SPIO⁺ BMDC and Ad SPIO⁺ BMDC for both low (**g**) and high (**h**) injection doses. Data are shown as means ±SEM for *n*=3 mice per dose (**g** and **h**, paired *t* test, *p*=0.470 and *p*=0.539). The experimenter was blinded prior to image acquisition and images were taken at × 100 magnification (scale bar=500 µm). Data is representative of *n*=5 independent experiments



Fig. 7 (See legend on previous page.)

to the observed MPI signal [19]. Indeed, a prior study by our group used ¹H MRI to confirm that SPIO⁺ BMDC detected by MPI were localized to pLN [14] 72 h following ACT into the hind footpad. This study additionally brought attention to the difficulty of resolving strong and weak MPI signal near each other. We perceive this to be a persistent issue for DC tracking as 3-5% of injected DC reach the pLN (weak signal) [11], which is in close proximity to the hind footpad injection site (strong signal). Even after fasting and housing animals with corn bedding that contains minimal iron prior to MPI, GI signal was still evident (Figs. 4b and 5b) and could similarly interfere with weak pLN signal quantification. GI signal will also be an issue for human MPI. Therefore, we made use of a focused FOV to quantify pLN signal and exclude areas of high iron concentration that limit the dynamic range capable of being resolved [56].

Two days following ACT, we first employed a standard FOV to measure pLN-migrated SPIO⁺ and Ad SPIO⁺ BMDC. pLN signal was detected in 9 of 12 pLN (75%), which is an improvement to MPI signal detection in 50% of pLN observed in a previous and similarly designed experiment [14]. Strong injection site signal, presumably due to nonmigrating or apoptotic BMDC [26], that is unable to be resolved from pLN signal precluded quantification in 3 of 12 pLN. By switching to an advanced image reconstruction algorithm allowing a focused 2-cm FOV that largely excludes injection site MPI signal, pLN signal was detected and quantified in 12 of 12 pLN (100%), demonstrating that MPI is capable of consistent detection and quantification of in vivo migration for two distinct DC populations (SPIO⁺ and Ad SPIO⁺ DC). Importantly, a smaller FOV also drastically reduced image acquisition time from 40 to 15 min, improving efficiency. We also acknowledge that a focused FOV may not be applicable for imaging studies in which the location of in vivo migration is unknown.

Excised pLN MPI was performed in earlier experiments to confirm that MPI signal was originating in the pLN. To remain consistent, we also included excised pLN MPI in this study and as expected, we detected signal in 100% of pLN. Further studies were performed to ensure that pLN-localized signal was due to injected BMDC as opposed to false positive signal resulting from resident macrophage uptake of apoptotic SPIO-labeled cells [57]. By labeling SPIO⁺ and Ad SPIO⁺ BMDC with Violet fluorescence prior to ACT, digital morphometric analysis of pLN confirmed that originally injected migrationcompetent Violet⁺ DC were the source of MPI signal and that no difference in *in vivo* migration capacity was noted between SPIO⁺ and Ad SPIO⁺ BMDC. In all pLN, SPIO⁺ and Ad SPIO⁺ BMDC penetrated the central and paracortical regions. This suggests that DC were viable upon reaching the pLN as nonviable DC phagocytosed by macrophages would remain in the cortex. Furthermore, this observation has immunological implications as TAA-presenting DC engage with and activate CD4⁺ and CD8⁺ TAA-specific T cells within the paracortex [48, 49] as we demonstrated in a prior BMDC-based cancer vaccine study [26].

Previously, the use of a focused FOV to isolate low signal regions was challenging, if not impossible, with the prescribed native reconstruction algorithm equipped on the Momentum[™] MPI scanner (Magnetic Insight Inc., Alameda, USA) and the only option for a user limited to accessing the regular user interface. Native image reconstruction assumes that there is no signal along the edges of the FOV; if there is signal present, these values are set to zero for each line along the transmit axis. This assumption causes an inverted negative image artifact which can prevent signal detection and quantification. Thus, in previous experiments, a full 12-cm FOV was used to ensure all signal was encompassed within a large area. For these experiments, our MPI system was upgraded to allow the user access to an advanced user interface where image sequences could be changed through the configuration editor. Most importantly, this permitted the use of a novel reconstruction algorithm referred to as multichannel joint image reconstruction [30], recently implemented as a reconstruction option on the Momentum[™] scanner (Magnetic Insight Inc., Alameda, USA). The multichannel joint reconstruction method uses an iterative reconstruction technique to recover edge information using information from an orthogonal axis, preventing this artifact, and allowing the user to prescribe a small FOV on the region of interest. Additionally, with this improved method, as few as 1,000 pLN-migrated DC (4.4 ng Fe) were detected in vivo following footpad injection. This is the lowest amount of Fe detected in vivo thus far and the lowest number of cells detected that were labeled with a commercially available SPIO. Previously, Song et al. [58] reported on the *in vivo* detection of 250 HeLa cells (7.8 ng Fe) labeled with a custom synthesized Janus particles and implanted subcutaneously on the back of a mouse. This study demonstrates the feasibility of using this type of reconstruction to resolve and quantify discrete sources of signal without interference from proximate, varying iron concentrations that are not of interest.

In conclusion, this is the first *in vivo* MPI study that employed multichannel joint image reconstruction with a focused FOV for image acquisition and represents an important step forward in establishing MPI as a noninvasive imaging modality capable of tracking the fate of injected DC to serve as a surrogate marker of therapeutic effectiveness. Recently, DC-based therapies have garnered much interest within combination cancer immunotherapy in autoimmune and transplantation settings [59] as well as in infectious disease settings such as SARS-CoV-2 [60]. Regardless of disease setting, noninvasive and quantifiable imaging is required for cell-based immunotherapies to be successful.

Abbreviations

Adoptive cell transfer
Adenovirus
Antigen presenting cell
Bone marrow-derived dendritic cell
Dendritic cell
Enhanced green fluorescent protein
Field of view
Gastrointestinal
Granulocyte-macrophage colony-stimulating factor
Lymph node
Multiplicity of infection
Magnetic particle imaging
Magnetic resonance imaging
Popliteal lymph node
Region of interest
Signal to noise ratio
Superparamagnetic iron oxide
Tumor-associated antigen

Supplementary Information

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Additional file 1: Supplemental Table S1. Antibodies and reagents. Supplemental Fig. S1. Ad SPIO⁺ BMDC migrate to the popliteal lymph node following injection. One million Ad SPIO⁺ BMDC were fluorescently labeled with PKH26 immediately before hind footpad adoptive transfer. Two days later, excised popliteal lymph nodes were processed into 16 µm cryosections to observe PKH26 (a) and enhanced green fluorescent protein (b) fluorescence at × 100 magnification. Image inset outlined by a white square in (a and b) is shown at × 400 magnification for PKH26 (c) and eGFP fluorescence (d) as well as a PKH26 and eGFP overlay (e) to identify Ad SPIO⁺ BMDC in pLN. Scale bars = 100 µm (a-b) and 20 µm (c-e). Data is representative of n = 2 independent experiments with four mice per group.

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Authors' contributions

CF wrote the manuscript with contributions from JG. CF, JG, and JWB performed experiments. CF, JG, PJF, and GAD planned and designed experiments. All authors read and approved the final manuscript.

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Availability of data and materials

The datasets used and/or analyzed during the current study are available from the corresponding author on reasonable request.

Declarations

Ethics approval and consent to participate

All animal studies were performed in accordance with institutional and national guidelines.

Consent for publication

Not applicable.

Competing interests

The authors declare that they have no competing interests.

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