

NIH Public Access

Author Manuscript

J Magn Reson. Author manuscript; available in PMC 2013 October 10

Published in final edited form as:

J Magn Reson. 2012 April ; 217: 86–91. doi:10.1016/j.jmr.2012.02.014.

Detection of Iron-labeled Single Cells by MR Imaging Based on Intermolecular Double Quantum Coherences at 14 T

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Abstract

To evaluate the efficiency and feasibility of intermolecular multiple quantum coherence (iMQC) magnetic resonance (MR) imaging for single cell detection, we obtained intermolecular double quantum coherence (iDQC) and conventional gradient echo (GE) images of macrophage cells labeled by contrast agents in gel. The iDQC images obtained with echo-planar readout visualized the labeled cells effectively and with a higher contrast than seen in conventional GE images, especially at low planar resolutions and with thick slices. This implies that iDQC imaging with contrast agents could be a good alternative to conventional MR imaging for detecting labeled single cells or cell tracking under favorable conditions.

1. Introduction

Over the past decade, cellular and molecular imaging (including positron emission tomography (PET) [1], single-photon emission computed tomography (SPECT) [2], optical imaging with fluorescence or bioluminescence imaging [1, 3], and magnetic resonance imaging (MRI) [4]) has attracted attention owing to its ability to monitor biological processes in living organisms. This involves not only direct imaging of individual cells and molecules but also diagnosis and management of disease processes. MR applications include stem cell tracking [5–7], gene expression [8, 9], and cancer detection [10]. Conventional MRI techniques enable *in vivo* long-term investigation at relatively high resolution, but they are limited by their low sensitivity in the detection of a specific cell and/or molecule.

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One major class of modern MR contrast agents uses iron oxide particles, either by themselves [11, 12] or conjugated to attach to specific receptors [13–15]. Thus, many groups have been developing methods to improve the cellular uptake of iron oxide particles [16, 17] or have been using micron-sized iron oxide particles to increase the cell labeling efficiency [18]. Many studies have visualized iron-labeled single cells at a high planar resolution (100 μ m) and thin slice thickness (200 μ m) using 3D images or a strong gradient [19–22]. In general, MR detection of iron-labeled single cells requires a sensitive RF coil to achieve a high signal-to-noise ratio (SNR) and a strong gradient to achieve high resolution. In addition, various MRI techniques have been proposed to visualize the local dipolar field more effectively. For example, the 3D fully refocused (steady-state free precession) gradient-echo sequence (known as FIESTA on GE scanners) [20-22] can enhance the effects of the local dipoles and contrast between the local dipole and background. Positive contrast with iron oxide particles is possible by several approaches: spectrally selective RF pulses can excite and refocus the off-resonant water surrounding the labeled cells [23], a mismatched slice gradient [24] or a mismatched coherence-selection gradient pulse pair [25] can dephase the background signal, or an echo-dephased steady state free precession method can be used [26].

In this study, we used intermolecular multiple quantum coherences (iMQCs) to enhance the visibility of iron-labeled single cells at high field (14 T). Because of the intrinsic sensitivity to changes in magnetization and susceptibility structures, iMQC imaging signals provide a fundamentally different contrast mechanism from conventional MRI contrast [27–31], and are particularly well suited to imaging local anisotropy produced by magnetic nanoparticles [32, 33]. Our previous study showed that intermolecular double quantum coherence (iDQC) imaging produced unique contrast that reflected susceptibility structures of small (~ 100 μ m) air bubbles formed on the walls of capillary tubes [34]. This susceptibility variation can create resonance frequency differences between distant spins that are coupled by intermolecular dipolar interactions. The iMQC image intensity around the bubbles is then reduced by additional dephasing effects and phase distortions [25, 34, 35].

iMQC imaging has been demonstrated at field strengths from 0.016 T [36] to 17.6 T [37–40]. In the present study, we compared images of iron-labeled macrophage cells in gel obtained with iDQC with EPI detection (iDQC_EPI) with those obtained through gradient echo (GE) imaging, at different planar resolutions and slice thicknesses, to evaluate the efficiency and feasibility of iMQC imaging for single cell detection at 14 T. To reveal the effects of the excitation and readout methods, we also compared the iDQC_EPI image with the conventional EPI and iDQC images with spin-echo detection (iDQC_SE) at the spatial resolution of 100 μ m \times 100 μ m \times 1 mm.

2. Experimental methods

2.1. Cell sample preparation

Macrophage cells (Raw 264.7) with an average diameter of 11 μ m were obtained from a mouse monocytic cell line. The cells were maintained at 37°C and 5% CO₂ in Dulbecco's modified Eagle's medium (DMEM; Gibco BRL, Gaithersburg, MD, USA) supplemented with 1% fetal bovine serum (FBS; Gibco BRL). Cell labeling was accomplished by adding a ferumoxide injectable solution (Feridex IV; Advanced Magnetics, Inc., Cambridge, MA, USA) with an iron concentration of 400 μ g Fe/mL to 10⁶ cells in the culture medium. Labeled cells were incubated for 4 h, washed three times with PBS in an adherent state, and then detached. The cells were suspended in 25 mL of DMEM and centrifuged for 5 min at 1000 rpm. Cells were counted with a hemocytometer.

For MRI studies, labeled and unlabeled cells were suspended in agarose (1%, w/w) at cellular concentrations of 5000 cells/mL and placed in 5-mm NMR tubes. The cellular iron content was assessed by an inductively coupled plasma-atomic emission spectrometer (ICP-AES).

2.2. Magnetic resonance imaging

All experiments were performed on a 14.1 T Bruker Avance DMX600 NMR spectrometer equipped with a triple gradient system for microscopic imaging. The maximum gradient strength was 200 Gauss/cm, and a 5-mm saddle-type RF coil was used. Imaging data were obtained and analyzed using Bruker's ParaVision 3.0.2 imaging software.

The iDQC EPI and iDQC SE images were acquired with a CRAZED-type pulse sequence (90°- -gradient-120°-[double-area gradient]-2 -TE/2-180°-TE/2) with EPI and SE readouts at the end of the sequence, respectively [34]. Four-step phase cycling (for the first RF pulse and the receiver phases) was introduced to suppress unwanted coherences such as singlequantum contaminations. To acquire the iDQC_EPI images, we set echo time (TE) = 90 ms, matrix size = 64×64 , bandwidth (BW) = 50 kHz, = 10 ms, correlation distance (d_c) = 100 μ m, NS = 32, and scan time = 5 min (NS = 128 and scan time = 21 min for the images of the 0.1-mm-thick slice). We also set TR = 10 s to minimize contamination from stimulated or interscan-stimulated echoes. Slice thickness was also varied to study the resolution dependence of the cell detection. The iDQC_SE image with 1-mm slice thickness was also acquired with the same parameters, but the total scan time was 42 min due to the difference in readout technique. The GE images were acquired with TE = 90 ms, TR = 500 ms, and total scan time = 5 min (hence, NS = 10). In acquiring most of these images, we have fixed TE, the bandwidth, and the total scan time in order to facilitate comparisons. Conventional spin-echo images with EPI detection (SE_EPI) were also acquired to evaluate T_2 -contrast effects during TE under the same parameters as those in the iDQC experiments, except during multiple quantum excitation/selection.

The fractional signal loss (S/S) was used to assess the amount of contrast, namely single cell detectability, for a given signal void generated by an iron-labeled cell in the images. In our experiments, S/S was calculated by taking the difference (S) between the signal from the central voxel of the cell region (S_{cell}) and the signal from the background gel containing no labeled cells (S) and then dividing the difference by the background gel signal (S) [21, 22]. For each of the images, a mean S/S value was calculated, averaging over six cells. To evaluate each performance, in addition to the overall data quality of the iDQC and GE images, we also calculated the contrast-to-noise ratio (CNR = (S/S) · SNR) because it reflects both the contrast and the SNR [21, 22]. In general, the CNR indicates the detectability of individual cells and clinical lesions in real systems. However, when an image has a sufficient SNR value, the S/S value of the image may better represent the single cell detectability than the CNR because the S/S value is not significantly altered by the SNR [21], while the CNR takes considerable impact from the SNR.

3. Results and discussion

3.1. Detection of iron-labeled single cells with different spatial resolutions

For low-resolution images, a small inhomogeneity such as an iron-labeled cell (which is smaller than a voxel) is ordinarily blurred by averaging all the magnetizations in a voxel [18]. Therefore, spatial resolution comparable to the size of the cell, along with an acceptable SNR, is normally required for detecting single cells. However, as discussed elsewhere [34], iMQC MR imaging is known to be capable of magnifying small variations in a voxel.

To evaluate the feasibility and efficiency of iMQC MR imaging for single cell detection, the iDQC_EPI and GE images of the iron-labeled (~20 pg Fe/cell) macrophage cells in gel were acquired at various spatial resolutions. As shown in Figs. 1 and 2, single cell detection was generally easier with thin slices or high planar resolution. At a slice thickness of 100 μ m, the images showed distinct signal loss that contrasted against the high background signal produced by the agarose medium. The large magnetic susceptibility of the iron oxide particles affected a region 10 times larger than the actual size of a single cell (~11 μ m) in the iDQC_EPI and GE images. At a slice thickness of 0.1 mm or 0.5 mm with a planar resolution of 100 μ m × 100 μ m, the cells were easily detected in both the GE and iDQC_EPI images. However, at a slice thickness of 1 mm or thicker, the iDQC_EPI images provided higher cell contrast than did the GE images. In particular, at the 2-mm slice thickness in images with a planar resolution of 100 μ m × 100 μ m, some labeled single cells remained clearly distinguishable in the iDQC_EPI image.

The *S/S* and CNR values, representing averages over the six most distinct cells selected in the GE image with 0.1-mm slice thickness (see the magnified images and arrows in Fig. 1b), were measured to visualize the dependence of image contrast on the spatial resolution. As seen in Fig. 1, the S/S values in the iDQC_EPI and GE images seemed to show a somewhat different tendency. At the 0.1-mm slice thickness, the iDQC EPI image had a lower cell-contrast (S/S) value than the GE image, but the S/S value decreased with increasing slice thickness slower than that of the GE image. This could be explained by the intrinsic property of iMQCs that arise from long-range dipole-dipole couplings. Consequently, at a slice thickness of 1 mm or higher, the iDQC_EPI image had larger S/Svalues than the GE images. This result implies that under certain favorable conditions, iDQC_EPI images may offer better contrast than GE images for iron-labeled single cell detection. The mean *S/S* values in both the iDQC EPI and GE images at the spatial resolution of 100μ m \times 100 μ m \times 500 μ m were ~ 0.5. This value was similar to the value (~ 0.45) obtained from the 3D FIESTA pulse sequence at the spatial resolution of 100 μ m \times $100 \ \mu m \times 200 \ \mu m at 1.5 \ T \ [22]$. The CNR curve showed a different pattern from the S/Scurve because the CNR is the product of contrast (S/S) and SNR. The SNR increases with the slice thickness, hence the CNR curve showed a maximum at the 1-mm (or 0.5-mm) slice thickness. Generally, compared with conventional MR images, the SNR of the iDQC image is expected to be low because of the nature of its signal source. However, since we have fixed TE (= 90 ms) and total scan time (= 5 min, except for the iDQC_EPI image at the 0.1mm slice thickness) for image acquisition, the SNR values of the GE images became similar to those of the iDQC images. The long TE in the GE detection could cause enhanced image contrast but reduced SNR. The experimental results showed that the iDQC EPI images at more than 0.5-mm slice thickness had sufficiently large SNR and high S/S values to detect the iron-labeled single cells. Thus, when iDQC_EPI images include a sufficiently strong signal (for example, at low resolution, such as in clinical images), iDQC imaging could be a good alternative for cellular or molecular MR imaging experiments.

Figure 2a shows iDQC_EPI and GE images at the various planar resolutions with 1-mm slice thickness. At all resolutions, the iDQC_EPI images showed greater cell detectability than the GE images. In particular, at the planar resolution of 150 μ m × 150 μ m, some iron-labeled single cells were readily detected in the iDQC_EPI image but were somewhat indistinct in the GE image. A resolution of 150 μ m × 150 μ m has often been used to obtain GE images of the brains of small animals such as mice. With this planar resolution, a 3D GE image may be needed to detect labeled cells (<0.1 pixel) clearly, because the labeled cells are not easily detectable in a conventional 2D image with a slice thickness of 1 mm. As shown in Figs. 1 and 2, however, the iDQC_EPI image showed relatively good detectability for the labeled cells, even at the slice thickness of 1 mm or higher. The mean S/S and CNR values for the same six labeled cells in the iDQC_EPI and GE images were also measured

(see Fig. 2b). Both the mean *S/S* and CNR values of the iDQC_EPI images were higher than those of the GE images. The higher CNR values were attributed to the good signal of the iDQC_EPI image with 1-mm slice thickness, as discussed earlier. These mean *S/S* and CNR values further confirmed that labeled cells could be more detectable in the iDQC_EPI images than in the GE images under some favorable conditions. In particular, for a thick slice and/or a low planar resolution, the iDQC_EPI imaging method could be a more useful tool than the conventional GE imaging method for detecting single cells.

3.2. Comparison of images from the different imaging methods

We also acquired SE_EPI and iDQC_SE images to evaluate the contributions of excitation and readout methods in image contrast. Figure 3 shows the GE, iDQC_EPI, SE_EPI, and iDQC_SE images at the spatial resolution of 100 μ m × 100 μ m × 1 mm for the same sample. The mean *S/S* and CNR values for the same six labeled cells of the four images were also measured, as well as the SNR of the images. The *S/S* value of the iDQC_EPI image was the biggest, while those of the other images were similar to each other (60 ~ 70 % of the iDQC_EPI).

As would be expected, the conventional SE image (not shown) does not have enough contrast to detect iron-labeled single cells. However, the adoption of EPI readout (SE_EPI) significantly enhanced the cell detectability since the EPI readout includes additional T_2^* -weighted contrast as a result of the long readout window [41]. The iDQC_EPI and SE_EPI images were acquired using the same EPI acquisition method with the same readout parameters, except during the iDQC excitation. In principle, therefore, T_2 contrast effects during the TE period should be the same in both experiments, but the iDQC_EPI image showed enhanced contrast because the signals intrinsically originated from the multispin coherences. Although both signals of iDQC_EPI and iDQC_SE images originated from intermolecular multispin coherences, they showed different cell contrast due to the additional T_2^* contrast effects as in the previous case of the conventional SE and SE_EPI images.

Consequently these experimental results show that the large enhancement of cell contrast in the iDQC_EPI image arises from both the intrinsic iMQC property and the EPI-readout method. It was reported that the iDQC images generally reflected their enhanced T_2^* -weighted sensitivity due to the particular iMQC properties [32, 34, 38, 39]. Rose defined a kind of threshold CNR that dictates the performance of human signal detection, and he determined that the threshold value was ~5 [21, 42]. Based on this criterion, the CNR values of all images were sufficient to detect the iron-labeled cells. Hence, the relative cell detectability will depend mainly on the cell contrast (S/S).

3.3. Contrast sources in iDQC MR images

To a greater or lesser extent, all four of the images in Fig. 3 reflect T_2 *-weighted contrasts. The GE images reflect mainly the T_2 *-weighted contrast, while the others reflect the T_2 -weighted contrast as a result of the long TE as well as some T_2 *-weighted contrast. The iDQC excitation with EPI detection seemed to be particularly suitable for detecting iron-labeled single cells.

In general, the combination of the iMQC MR imaging technique with contrast agents may offer two major benefits for enhancing the contrast for single cell detection. First, iDQC detections magnify the effects of the short relaxation time (mainly T_2 in this experiment) because iDQCs arise from the two-spin correlations (exponentially decaying twice as fast as T_2) over a correlation distance (depending on the correlation gradient), while the conventional GE and/or SE signals arise from the summation of the single spin

magnetizations (exponentially decaying with T_2 *or T_2) in a voxel. In addition, the iDQC signals of the iron-labeled cells and nearby coupled gel spins were significantly decreased owing to the additional dephasing effects induced by the chemical shift variations of these cells. These additional enhancements might facilitate reliable contrast in MR images even with relatively small amounts of contrast agents and low spatial resolutions. Figure 4 graphically presents the enhanced contrast of the iDQC image when coupled with contrast agents. As discussed in our previous study [34], the intervoxel effects in iMQC microscopic imaging could increase the detectability (by the signal loss around the labeled cells) of small-sized single cells. Combining the effects of the iDQC method and the EPI readout reflecting its T_2 *-weighted contrast might significantly enhance the detectability of iron-labeled single cells.

4. Conclusion

We have demonstrated the feasibility of combining iDQC imaging with contrast agents for single cell detection. To enhance contrast and/or detectability, a combination of iDQC excitation and EPI readout was used. As compared to conventional GE imaging, contrast-enhanced images were obtained with iDQC_EPI imaging even at low spatial resolutions. Experimental results show that iDQC_EPI imaging could be an alternative (and possibly a better alternative under certain favorable conditions) to conventional GE imaging for cellular and molecular imaging of paramagnetic and ferromagnetic contrast agents.

Acknowledgments

This work was supported by a Korea Basic Science Institute grant (T31404), NAP of the Korea Research Council of Fundamental Science & Technology (PGB042), a Korean Research Foundation Grant (KRF-2007-331-C00137), the Seoul R&BD program (10543), and an NIH grant (EB02122).

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> Feasibility verification of iDQC MR imaging method for single cell detection. > Comparison of iDQC and GE images at various experimental conditions. > iDQC images show higher cell detectability than GE images under some favorable conditions. > iDQC imaging could be an alternative to GE imaging for cellular/molecular imaging.



Figure 1.

MR images of iron-labeled cell samples according to (a) slice thickness variations (0.1, 0.5, 1.0, and 2.0 mm from left to right) at 100 μ m × 100 μ m planar resolution. (b) A more magnified image (top) showing the cells selected for the calculation of the fractional signal loss (*S/S*) value and the resulting mean *S/S*, SNR, and mean CNR graphs (bottom). The error bars represent the standard deviation of the mean value from the six cells.



Figure 2.

(a) MR images of iron-labeled cell samples ordered by planar resolution variations (78 μ m × 78 μ m, 100 μ m × 100 μ m, and 150 μ m × 150 μ m from left to right) with 1-mm slice thickness and (b) graphs of calculated mean fractional signal loss (*S/S*) SNR, and mean CNR values for the six cells in Fig. 1b. The error bars represent the standard deviation of the mean value from the six cells.



Figure 3.

(a) Comparison of MR images from different imaging methods and (b) graphs of calculated mean fractional signal loss (S/S), SNR, and mean CNR values for the six cells in Fig. 1b. The spatial resolution of the images was 100 μ m × 100 μ m× 1 mm. The error bars represent the standard deviation of the mean value from the six cells.



Figure 4.

Signal sources and corresponding pixel images of conventional single quantum coherence (SQC, a and b) and iDQC (c and d) imaging, considering effects of contrast agent (CA) and distant dipolar field (DDF; correlation distance of 100 μ m). The pixel dimension and single cell size were set to 100 μ m × 100 μ m and 10 μ m, respectively. Expansion of the effects to a 3D voxel could be straightforward.