

Published in final edited form as:

Magn Reson Imaging. 2014 April ; 32(3): 245–249. doi:10.1016/j.mri.2013.10.013.

Assessing reproducibility of diffusion-weighted magnetic resonance imaging studies in a murine model of HER2+ breast cancer

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Abstract

Background and purpose—The use of diffusion-weighted magnetic resonance imaging (DW-MRI) as a surrogate biomarker of response in preclinical studies is increasing. However, before a biomarker can be reliably employed to assess treatment response, the reproducibility of the technique must be established. There is a paucity of literature that quantifies the reproducibility of DW-MRI in preclinical studies; thus, the purpose of this study was to investigate DW-MRI reproducibility in a murine model of HER2+ breast cancer.

Materials and methods—Test–Retest DW-MRI scans separated by approximately six hours were acquired from eleven athymic female mice with HER2+ xenografts using a pulsed gradient spin echo diffusion-weighted sequence with three *b* values [150, 500, and 800 s/mm²]. Reproducibility was assessed for the mean apparent diffusion coefficient (ADC) from tumor and muscle tissue regions.

Results—The threshold to reflect a change in tumor physiology in a cohort of mice is defined by the 95% confidence interval (CI), which was $\pm 0.0972 \times 10^{-3} \text{ mm}^2/\text{s}$ ($\pm 11.8\%$) for mean tumor ADC. The repeatability coefficient defines this threshold for an individual mouse, which was $\pm 0.273 \times 10^{-3} \text{ mm}^2/\text{s}$. The 95% CI and repeatability coefficient for mean ADC of muscle tissue were $\pm 0.0949 \times 10^{-3} \text{ mm}^2/\text{s}$ ($\pm 8.30\%$) and $\pm 0.266 \times 10^{-3} \text{ mm}^2/\text{s}$, respectively.

Conclusions—Mean ADC of tumors is reproducible and appropriate for detecting treatment-induced changes on both an individual and mouse cohort basis.

Keywords

Reproducibility; Diffusion-weighted MRI; Apparent diffusion coefficient; Mouse

1. Introduction

Current radiographic analysis of treatment response is based on the Response Evaluation Criteria in Solid Tumors (RECIST), which uses one-dimensional changes in tumor size to determine response [1]. However, it is well recognized that RECIST may not provide the most sensitive response assessment to novel and emerging therapies that target the molecular or cellular aspects of the tumor itself or, for example, its microenvironment [2, 3]. Development of imaging methods that can provide an earlier assessment of disease response is an active area of research, as it would allow for the initiation of alternative, potentially more effective, treatments while avoiding unnecessary toxicities associated with ineffective therapy. Diffusion-weighted magnetic resonance imaging (DW-MRI) has the potential to offer earlier assessments of disease response, as parameters derived from quantitative analyses of such data can be used to characterize treatment-induced alterations in tumor cell density [4–7].

The microscopic, thermally-induced behavior of molecules moving in a random pattern is referred to as self-diffusion or Brownian motion. The rate of diffusion in cellular tissues is described by an apparent diffusion coefficient (ADC), which is influenced by the number and separation of barriers that a diffusing water molecule encounters. DW-MRI maps the ADC, and in well-controlled situations the variations in ADC have been shown to correlate inversely with tissue cellularity [8]. It has been shown, in preclinical [9–12] and clinical [13–15] settings, that exposure of tumors to both chemotherapy and radiotherapy consistently leads to measurable increases in water diffusion in cases of favorable treatment response.

In their 2009 DW-MRI review, Padhani et al. recommended that each imaging center determine the reproducibility of DW-MRI data at their institution to allow for proper study design and to assess the significance of treatment-induced changes [16]. Reproducibility of clinical DW-MRI data have been previously assessed in normal and diseased tissues with a variety of image acquisition methods and data analyses including differences in the number and strength of diffusion-weighted images, as well as the reproducibility statistics themselves [17–20]. However, there is a paucity of such data in the preclinical setting. With the increasing use of DW-MRI as a surrogate biomarker of response in preclinical studies [9, 10, 12], investigating reproducibility is imperative to interpret whether parameter changes during therapeutic interventions reflect tumor cellularity instead of measurement error in mouse models of disease. Thus, the objective of this study was to quantify DW-MRI reproducibility in a murine model of HER2+ breast cancer in order to assist in the interpretation of results collected from future longitudinal treatment response studies.

2. Materials and methods

2.1. Animal and tumor xenograft model

Trastuzumab-resistant breast cancer cells, HR6, were generously gifted from Dr. Carlos Arteaga, M.D. at Vanderbilt University. This cell line was developed by harvesting cells from BT474 xenografts that initially responded but recurred in the presence of maintained trastuzumab [details are provided in Ref. 21]. (This cell line was selected as it is part of an

ongoing investigation of trastuzumab-resistance in breast tumors.) Female athymic nude mice ($n = 11$, 4–6 weeks, Harlan, Indianapolis, IN) were implanted with 0.72 mg, 60-day release, 17β -estradiol pellets (Innovative Research of America, Sarasota, FL). Twenty-four hours later, approximately 10^7 HR6 cells were injected subcutaneously into the right flank. Mice were anesthetized with 2% isoflurane in pure oxygen for both procedures. Tumor volumes were measured weekly after cell injection via calipers, and the mean tumor volume at the start of imaging was 298 mm^3 (range: 136 mm^3 – 526 mm^3). All animal procedures were approved by our Institution's Animal Care and Use Committee.

2.2. Image acquisition

Test–retest DW-MRI sessions were performed on 11 mice using a 7 T MRI scanner (Agilent Technologies (formally Varian), Palo Alto, CA) equipped with a 38-mm quadrature RF coil (Doty Scientific, Columbia, SC). Anesthesia was induced and maintained for each imaging session via 2% isoflurane in pure oxygen. Animal respiration rate was monitored, and animal body temperature was maintained at an external temperature of 32°C by means of a flow of warm air directly into the bore of the magnet. Each animal was placed in a custom built restraint, and the tumor region was first localized via 3D gradient echo scout images. Fifteen 1 mm thick slices with a 1 mm gap were acquired in order to cover the entire tumor region. A standard pulsed gradient spin echo sequence was used to acquire diffusion-weighted images with three b values (150, 500, and 800 s/mm^2) and gradients applied simultaneously along the three orthogonal directions (x , y , and z). Scan acquisition parameters were: TR/TE = 2000/30 ms, gradient duration $\delta = 3 \text{ ms}$, gradient interval $\Delta = 20 \text{ ms}$, two signal excitations, and an acquisition matrix of 128×128 over a $28 \times 28 \text{ mm}^2$ field of view. Image acquisition was triggered with respiration and navigator corrected [22] to reduce image artifacts due to bulk motion. Animals were removed from the scanner, allowed to recover completely from anesthesia, and given free access to food and water between test–retest imaging sessions. Repeat DW-MRI scan acquisitions were separated by a median of 5.88 hours (range: 1.50–7.82).

2.3. Image analysis

To construct an ADC parametric map, signal intensities from images acquired at three b values were fit for each image voxel using a nonlinear least squares optimization method to Eq. (1):

$$S(b) = S_0 \cdot \exp(-\text{ADC} \cdot b), \quad (1)$$

where S_0 and $S(b)$ are the signal intensities before and after application of diffusion gradients, respectively. As we did not acquire images without application of diffusion gradients (i.e., $b = 0$), S_0 was a free parameter in our optimization routine. Both image volumes were viewed simultaneously to ensure that a similar tissue section was analyzed, and the $b = 150 \text{ s/mm}^2$ image was utilized to define tumor boundaries. Care was taken during setup so that animal position and orientation in the magnet were similar between repeat acquisitions; however, image co-registration was not performed and thus separate regions of interest (ROIs) were manually drawn around the tumor for each scan. ROIs were also drawn within the skeletal muscle on the same slice as the tumor ROIs. Mean ADC values from each ROI were calculated and compared between repeated measurements. To ensure data integrity, a water phantom was imaged simultaneously with the animal in each imaging session. Additionally, the reproducibility of the mean ADC from the water phantom was computed. All data analysis methods were performed in MATLAB® (The MathWorks, Natick, MA).

2.4. Reproducibility statistics

Reproducibility statistics employed in this study follow the methods previously described by Bland and Altman [23], and are similar to what was previously implemented with imaging data by Galbraith et al. [24]. For each data set, the difference between repeat measurements, d was calculated. The distribution of the differences d was tested for normality using a two-sided Shapiro–Wilk test. A Kendall's tau test was used to estimate the correlation between the magnitude of the difference values and overall mean parameter value for the repeated measurements. A Wilcoxon signed-rank test was performed with the original data to test the null hypothesis of no bias (i.e., average difference is zero) between repeated measurements. The statistical measurements of reproducibility were then calculated as follows:

1. The root-mean-square deviation (rMSD) is computed using the differences between repeat measurements d :

$$\text{rMSD} = \sqrt{\frac{\sum d^2}{n}} \quad (2)$$

2. The 95% confidence interval (CI) for change which might occur in a group of n subjects:

$$\text{CI} = \pm \frac{1.96 \cdot \text{std}(d)}{\sqrt{n}}, \quad (3)$$

where $\text{std}(d)$ is the standard deviation of the difference between repeated measurements. Any change in a group of size n greater than this value would be significant at the 5% level.

3. The within-subject standard deviation (wSD) is:

$$\text{wSD} = \frac{\text{rMSD}}{\sqrt{2}}. \quad (4)$$

4. The repeatability coefficient (r) is:

$$r = 2.77 \cdot \text{wSD}, \quad (5)$$

or, equivalently:

$$r = 1.96 \cdot \text{rMSD}. \quad (6)$$

The repeatability coefficient defines the magnitude of the maximum difference expected in 95% of paired observations; i.e., an observed difference greater than this value between scans for an individual would be significant at the 5% level.

Due to our moderate sample sizes, we replaced 1.96 in Eq. (3) with the appropriate t-statistic for our sample size, which for 11 data sets is 2.228. Statistical analyses were performed using the statistical toolbox in MATLAB®. A significance value of $p < 0.05$ was used for all statistical tests.

In addition to the above reproducibility statistics, we also quantified the intraclass correlation coefficient (ICC). As defined by Landis and Koch for interpreting the strength of agreement of categorical data, an ICC greater than 0.61 is considered substantial agreement between repeated measurements [25].

3. Results

Examples of ADC parametric maps from repeated scans are depicted in Fig. 1 annotated with labels for tumor ('T'), muscle ('M'), and water phantom ('W'). It can be noted from the figure that the animal orientation is slightly different between repeated scans, thus showing the need for drawing separate ROIs for each imaging session. The average volume (\pm standard error) of the tumor ROIs was $46.02 \text{ mm}^3 (\pm 5.2004)$ and $44.5 \text{ mm}^3 (\pm 4.86)$ from scan 1 and 2, respectively. Statistically, there is no difference in tumor volume ROIs between repeated scans (Wilcoxon, $p > 0.05$). The reproducibility statistics for the tumor, muscle, and water phantom are listed in Table 1. Neither ROI had an average difference significantly different from normal as determined from the Wilcoxon signed-rank test. Additionally, the difference between repeat measurements d was independent of the mean for each ROI (thus, a logarithmic (or other) transformation was not required). Fig. 2 displays Bland–Altman plots for all ROIs, with each panel plotting the differences in ADC values between the repeated scans against the mean ADC value. The mean difference and 95% CIs are displayed as solid and dotted lines, respectively. The 95% CIs, which define the required change to surpass the expected measurement variability for a group of mice, are $\pm 0.0972 \times 10^{-3} \text{ mm}^2/\text{s} (\pm 11.8\%)$, $\pm 0.0949 \times 10^{-3} \text{ mm}^2/\text{s} (\pm 8.30\%)$, and $\pm 0.200 \times 10^{-3} \text{ mm}^2/\text{s} (\pm 7.78\%)$ for mean tumor, muscle, and water phantom ADC, respectively. Repeatability ranges (dashed lines), which provide thresholds of significance for individuals, are $\pm 0.273 \times 10^{-3} \text{ mm}^2/\text{s}$, $\pm 0.266 \times 10^{-3} \text{ mm}^2/\text{s}$, and $\pm 0.574 \times 10^{-3} \text{ mm}^2/\text{s}$ for mean tumor, muscle, and water phantom ADC, respectively.

We calculated the ICC for each ROI (please see Table 1). The ICCs for the tumor and muscle ROIs were greater than 0.61; thus, mean ADC for each ROI had substantial agreement between repeated measurements [25]. The ICC of the water phantom is zero, suggesting that the measurement system is producing 100% of the variability because the variability between subjects is (essentially) zero.

4. Discussion

We chose to quantify reproducibility for individuals and groups in a manner similar to Galbraith et al. [24], which is based on the limits of agreement methods described by Bland and Altman [23]. This analysis provides objective statistical thresholds that define the range of repeatability by quantifying the maximum difference expected to be observed between two repeat observations for an individual. Additionally, the 95% CI for the mean difference provides a measure of spontaneous change that is expected in groups of similar size to our cohort. Both the 95% CI and repeatability coefficient are useful in, for example, a longitudinal treatment response study where the objective is to quantify changes in parameters after a specific intervention. For example, mean tumor ADC would need to increase or decrease by $0.0972 \times 10^{-3} \text{ mm}^2/\text{s}$ to reflect a treatment-induced change in a group analysis, or change by $\pm 0.273 \times 10^{-3} \text{ mm}^2/\text{s}$ for an individual animal. The 95% CI when expressed as a percentage of the mean is $\pm 11.8\%$, which is not unreasonable as treatment-induced changes greater than $\pm 20\%$ have been reported when evaluating DW-MRI as a biomarker of response in several preclinical mouse models of cancer [12, 26], including a mouse model of HER2+ human breast cancer [9]. The reproducibility for muscle ADC was also expressed as a 95% CI and repeatability coefficient, which were $\pm 0.0949 \times 10^{-3} \text{ mm}^2/\text{s}$ and $\pm 0.266 \times 10^{-3} \text{ mm}^2/\text{s}$, respectively. The 95% CI expressed as a percentage of the mean is $\pm 8.30\%$, so that changes greater than this value would reflect a significant change in muscle for a cohort of mice of similar size as in our study.

Reproducibility in terms of agreement between repeat measurements (defined by the ICC) was worse for muscle (ICC = 0.63) compared to tumor (ICC = 0.80). This result is not

surprising, considering that muscle ADC values in this study are calculated from a diffusion-weighted pulse sequence that encodes along a single diffusion direction, instead of the trace, for each b value. Since diffusion in muscle tissue is anisotropic [27–29], ADC values measured along only one direction, acquired by applying diffusion gradients simultaneously on all three axes, would be dependent on animal orientation in the scanner. Although the orientation was kept as consistent as possible, one might expect that slight differences between scans would have a greater impact on reproducibility in muscle tissue compared to most tumor models where water diffusion is nearly isotropic [16].

The ICC for tumor and muscle from our study were slightly lower than a previous reproducibility study by O’Flynn et al. who observed an ICC of 0.91 for mean ADC of normal human breast tissue [20]. It is worth noting that, while we observed lower ICCs than O’Flynn et al., our main objective was to quantify a threshold by which changes greater than that value would represent therapeutic interventions. The 95% CI and repeatability coefficient define these thresholds in a population or an individual, respectively, and thus are more appropriate than the ICC when interpreting results from a longitudinal treatment response study [19, 30].

The reproducibility data provided in this manuscript can also be used to assist in prospectively powering, for example, a study designed to test the therapeutic response of a given drug regimen. If a quality estimate of the error in an individual ADC measurement is known, then a quality estimate on the number of animals required in each cohort to achieve a desired power can be determined. We take the standard deviation of the difference between the repeated measurements to be a reasonable estimate of the standard deviation of the difference in the response of matched pairs (i.e., two measurements on the same animal separated by time). If we set the type 1 error (i.e., α) to be 0.05 and assume a (reasonable) difference in ADC between the experimental and control groups of 20%, then 10 animals per group is sufficient to reject the null hypothesis that the population means of the two groups are equal with a probability (power) of 0.90.

In our study, we attempted to be as consistent as possible when determining the imaging FOVs between repeated acquisitions. However, differences in the measured ADC values from the ROIs exist. A possible explanation for this is that we are not comparing the exact sections of tissue from each scan. We chose to remove the animal from the scanner between repeat acquisitions to more closely mimic a study in which an animal is scanned multiple times over multiple days, thereby allowing us to quantify the variability that we might expect during a treatment study. Additionally, allowing the animal to recover between repeat scans minimized any physiological stress that may occur during long periods ($> \sim 2.5$ hours) of anesthesia. Removing the animal and allowing for recovery would mean that there might be slight differences in ROIs (both tumor and muscle), and thus possible repositioning errors between acquisitions. We expect that errors associated with differences in animal position would negatively impact reproducibility, thus resulting in larger 95% CIs and repeatability coefficients.

5. Conclusion

The development of imaging biomarkers with the sensitivity to detect treatment-induced changes in tumor physiology before tumor volume change is an active area of interest. Before an imaging biomarker can be reliably employed in a treatment response study, the reproducibility must be assessed. In this study, we assessed reproducibility of DW-MRI data in conjunction with an ADC measurement. These test–retest analyses of mean tumor ADC measurements were very reproducible, and thus DW-MRI can be used in serial studies to

measure treatment-induced changes of tumor physiology in our mouse model of HER2+ human breast cancer.

Acknowledgments

We thank the National Institutes of Health for funding through R01 CA138599, P50 CA98131, P30 CA68485, IR21CA169387, and 1 S10 RR17858. We also thank Dr. Henry Ong, Ph.D. and Dr. Jack Skinner, Ph.D. for image acquisition/analysis advice, as well as Dr. Zoe Yu, M.D. and Ms. Cammie Rinehart Sutton for cell culture and animal assistance.

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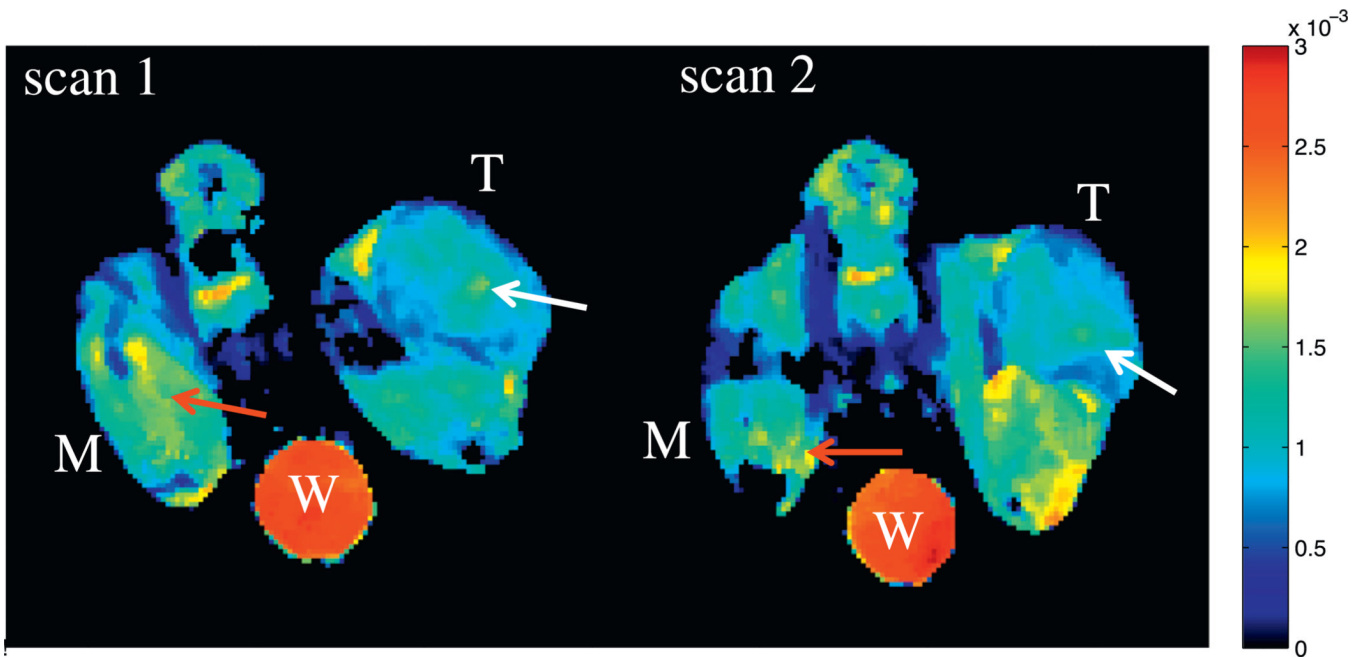


Fig. 1. Test-retest example of ADC parametric maps. Three regions of interest are indicated: tumor (denoted with a 'T'), skeletal muscle (denoted with 'M'), and a water phantom (denoted with a 'W'). Take note of how the animal orientation is slightly different between repeat scans, which might affect ADC reproducibility as the same tissue sections from each scan might not be analyzed. This variation in positioning is somewhat noticeable in the tumor, as the location and size of higher ADC regions (white arrows) are different; however, the variability of regions with higher ADC values (red arrows) is much larger in the muscle. Units of ADC are in mm^2/s .

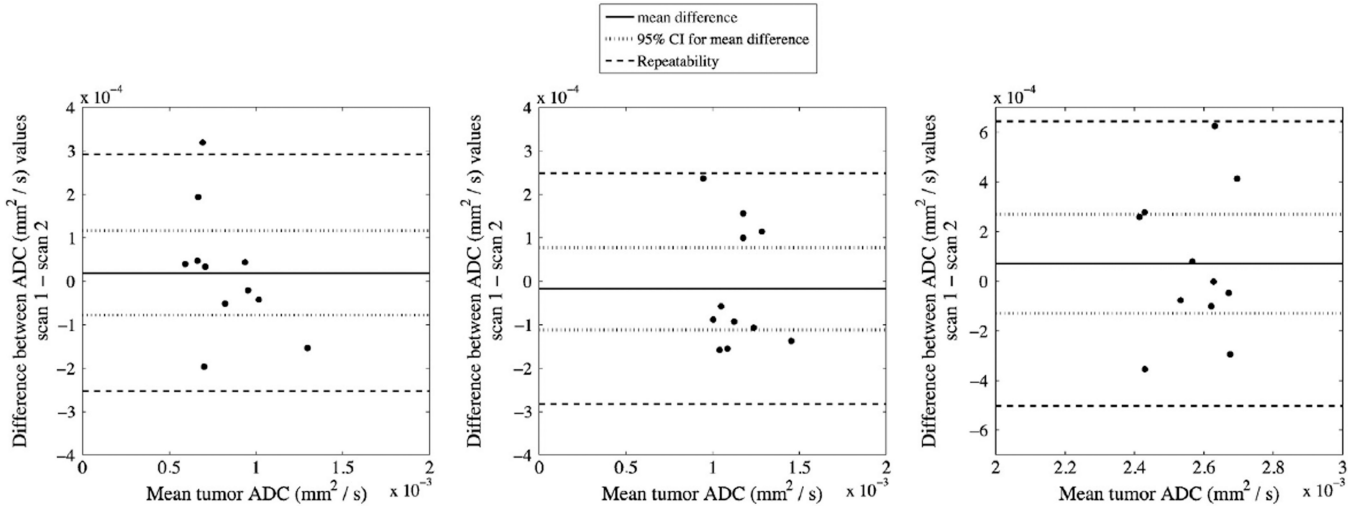


Fig. 2. Bland–Altman plots displaying the difference in ADC between scans plotted against mean ADC for both tumor (left panel) and muscle (right panel) ROIs. The mean difference (solid line) is shown with 95% confidence intervals (dotted lines), which defines the significant threshold for the population. Repeatability is also shown (dashed lines), and represents the threshold required to guarantee a statistically significant change in an individual mouse.

Table 1

ADC (mm^2/s) reproducibility analysis for the tumor, muscle, and water phantom.

ROI	Mean ($\times 10^{-3}$)	Mean difference ($\times 10^{-3}$)	95% CI for mean difference ($\times 10^{-3}$)	wSD ($\times 10^{-3}$)	Repeatability ($\times 10^{-3}$)	ICC
Tumor	0.822	0.0193	± 0.0972 (11.8%)	0.0985	0.273	0.80
Muscle	1.14	-0.0173	± 0.0949 (8.30%)	0.0960	0.266	0.63
Water phantom	2.57	0.0707	± 0.200 (7.78%)	0.207	0.574	0.00

ADC, apparent diffusion coefficient; ROI, region of interest; CI, confidence interval; wSD, within-subject standard deviation; ICC, intraclass correlation coefficient.