Early assessment of breast cancer response to neoadjuvant chemotherapy by semi-quantitative analysis of high-temporal resolution DCE-MRI: Preliminary results

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\textbf{ABSTRACT}

\textbf{Purpose:} To evaluate whether semi-quantitative analysis of high temporal resolution dynamic contrast-enhanced MRI (DCE-MRI) acquired early in treatment can predict the response of locally advanced breast cancer (LABC) to neoadjuvant chemotherapy (NAC).

\textbf{Materials and Methods:} As part of an IRB-approved prospective study, 21 patients with LABC provided informed consent and underwent high temporal resolution 3 T DCE-MRI before and after 1 cycle of NAC. Using measurements performed by two radiologists, the following parameters were extracted for lesions at both examinations: lesion size (short and long axes, in both early and late phases of enhancement), radiologist’s subjective assessment of lesion enhancement, and percentages of voxels within the lesion demonstrating progressive, plateau, or washout kinetics. The latter data were calculated using two filters, one selecting for voxels enhancing \( \geq 50\% \) over baseline and one for voxels enhancing \( \geq 100\% \) over baseline.

Pretreatment imaging parameters and parameter changes following cycle 1 of NAC were evaluated for their ability to discriminate patients with an eventual pathological complete response (pCR).

\textbf{Results:} All 21 patients completed NAC followed by surgery, with 9 patients achieving a pCR. No pretreatment imaging parameters were predictive of pCR. However, change after cycle 1 of NAC in percentage of voxels demonstrating washout kinetics with a 100\% enhancement filter discriminated patients with an eventual pCR with an area under the receiver operating characteristic curve (AUC) of 0.77. Changes in other parameters, including lesion size, did not predict pCR.

\textbf{Conclusion:} Semi-quantitative analysis of high temporal resolution DCE-MRI in patients with LABC can discriminate patients with an eventual pCR after one cycle of NAC.

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\section{1. Introduction}

Neoadjuvant chemotherapy (NAC) is offered to selected patients with locally advanced breast cancer (LABC) to reduce tumor burden before surgery and to allow for possible earlier treatment of occult micrometastatic disease [1]. Studies have confirmed that patients undergoing NAC have a lower risk of requiring mastectomy (i.e., are more likely to qualify for breast conservation therapy) and have equivalent survival to patients undergoing adjuvant chemotherapy [2,3].

With more breast cancer patients receiving NAC as a component of therapeutic management, the need for noninvasive assessment of treatment response has emerged as an important challenge for imaging. Because NAC is typically given in multiple cycles over several weeks, the ability to identify patients early in treatment who are not responding to a particular chemotherapy would allow the treating oncologist to discontinue an ineffective treatment (with

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potential short-term and long-term toxicities) and substitute an alternative regimen. Additionally, since pathological response is correlated with disease-free and overall survival [4,5], early noninvasive response assessment may have similar prognostic significance [6].

Prospective comparisons of different noninvasive imaging modalities have identified magnetic resonance imaging (MRI) as a useful technique in the setting of NAC for breast cancer [7,8]. Among different MRI approaches, dynamic contrast enhanced MRI (DCE-MRI) is considered especially promising due to its ability to assess changes in tumor vascularity in addition to changes in gross tumor size. To the extent that gross tumor shrinkage may lag behind changes in tumor vascularity in the context of a biological treatment response, DCE-MRI may be able to predict response earlier than techniques oriented exclusively toward tumor size.

DCE-MRI is an umbrella term used to describe a spectrum of MRI techniques and analytic approaches including both quantitative and semi-quantitative methods applied to data acquired via high and low temporal resolution sampling [9]. The literature on DCE-MRI for assessment of breast cancer response to NAC has evolved along two main avenues of investigation: fully quantitative approaches using different tracer pharmacokinetic models applied to high temporal resolution acquisitions [10–14], and a variety of alternative approaches employing semi-quantitative analyses of low temporal resolution/high spatial resolution images [15–18]. Both approaches have their strengths: theoretical considerations and empirical data suggest that diagnostic performance of DCE-MRI may improve with increasing temporal resolution [19], but semi-quantitative approaches may offer greater reproducibility by virtue of their simplicity, especially when deployed across multiple sites in a large-scale clinical trial [20].

Given the relative advantages of these two approaches, and out of consideration that a blended approach might offer enhanced reproducibility while retaining the ability to characterize changes in tumor vascularity with precision, we undertook this study to investigate whether a semi-quantitative analysis of high temporal resolution DCE-MRI data could provide useful early information regarding breast cancer response to NAC. We employed a semi-quantitative approach for kinetic curve type categorization similar to one previously applied by other investigators for initial characterization of lesions as benign or malignant [21]. While others have studied changes in tumor washout kinetics during preoperative therapy [15], this is the first attempt to our knowledge to evaluate whether a semi-quantitative analysis of high temporal resolution DCE-MRI data can be used to predict pathological response after one cycle of NAC.

2. Materials and methods

2.1. Patients and clinical protocol

Patients with pathologically proven LABC who were scheduled to receive NAC were eligible for this IRB-approved prospective study. After providing informed consent, patients underwent DCE-MRI before and after one cycle of NAC. Human epidermal growth factor receptor 2 (HER2) positive patients received paclitaxel, carboplatin, and trastuzumab every three weeks for six cycles. Most patients with HER2 negative tumors received doxorubicin and cyclophosphamide administered every two weeks for four cycles followed by twelve weekly cycles of paclitaxel, although a subset of patients with “triple negative” disease (i.e., negative for estrogen receptor (ER), progesterone receptor (PR), and HER2 overexpression) received weekly cisplatin and paclitaxel combined with either everolimus or placebo for twelve weeks as part of a separate clinical trial. After NAC, patients underwent either mastectomy or breast conservation therapy.

2.2. Pathological analysis

After surgery, specimens were evaluated for pathological treatment response. A patient was classified as having had a pathological complete response (pCR) to NAC if she had complete absence of residual disease at the primary tumor site and complete absence of disease in any resected lymph nodes. A patient was considered as not having achieved a pCR if she had any residual disease at the primary tumor site and/or residual lymph node disease.

2.3. MRI methods

Patients were screened prior to imaging to ensure adequate renal function before administration of intravenous gadolinium contrast. The study protocol specified a minimum estimated glomerular filtration rate (eGFR) of 90 ml/min, with this value obtained within 30 days of imaging.

MRI was performed on a Philips 3 T Achieva MR scanner (Philips Healthcare, Best, The Netherlands) using a 4-channel receive-doubledistance coil (Invivo Inc., Gainesville, FL). High temporal resolution DCE-MRI data were acquired using an RF-spoiled 3D gradient echo multi-flip angle acquisition with TR = 7.9 ms, TE = 4.6 ms, a flip angle of 20°, one signal acquisition, and a sensitivity encoding (i.e., SENSE) factor of 2 applied in the anterior–posterior direction. Twenty sagittal slices were acquired with a slice thickness of 5 mm, in-plane field of view (FOV) of 22 cm², acquisition matrix of 192 × 192, and no interslice spacing. Dynamic scanning was performed over 25 acquisitions with a temporal resolution of 16 s per acquisition, for a total scan time of 400 s. The first three acquisitions were unenhanced baseline scans; after the third baseline scan, 0.1 mmol/kg (9–15 mL) of gadopentetate dimeglumine (Gd-DTPA, Magnevist, Wayne, NJ) was administered through an antecubital vein catheter via a power injector (Medrad, Inc., Warrendale, PA) at 2 mL/s, followed by a saline flush. An investigator with over 10 years of experience in medical image registration (X.L.) monitored cine loops of the dynamic frames to ensure that images were not corrupted by patient motion.

2.4. Image analysis

Graphical user interface (GUI) software was constructed using Matlab 2010a (MathWorks, Natick, MA) to display acquired images and to facilitate data extraction (Fig. 1). Using the GUI software, two board-certified radiologists (R.A. and T.H.), each with more than five years of breast MRI experience, measured all lesions and drew regions of interest (ROIs) around lesion volumes on the pretreatment and post-cycle 1 image sets.

Data extracted for analysis included lesion size, perceived enhancement, and semi-quantitative enhancement kinetics parameters. For lesion size, tumor measurements were made in long-axis and short-axis using the slice on which the lesion appeared largest. Following the methodology described by Loo et al. [17], measurements were performed both during the initial phase of enhancement (before 120 s) and during the late phase of enhancement (at 400 s).

For perceived enhancement, each radiologist performed a subjective assessment of the percentage of lesion volume demonstrating enhancement at each timepoint. Enhancement was graded in quartiles as follows: less than 25% enhancement, 25% to 50% enhancement, 50% to 75% enhancement, and greater than 75% enhancement.

For semi-quantitative analysis of enhancement kinetics, the GUI first filtered for voxels within each ROI demonstrating signal
Fig. 1. Screen capture from the graphical user interface (GUI). The top row displays postcontrast MRI images from all three timepoints (baseline, after one cycle of NAC, and at completion of NAC). The graphs depict time–signal intensity curves for single voxels, chosen by the user by placing cross-hairs within the images. The middle row displays subtraction images corresponding to the slices in the top row. The bottom row contains parametric color maps representing the shape of the enhancement curve for each voxel within the slice, with one map using a 50% enhancement filter and the other map using a 100% enhancement filter. Blue = progressive (type I) kinetics, yellow = plateau (type II) kinetics, red = washout (type III) kinetics.
intensity increases of either $\geq 50\%$ or $\geq 100\%$ over baseline following contrast administration, thus removing any nonenhancing voxels or voxels with low-level enhancement. Then, using the $\geq 50\%$ and $\geq 100\%$ enhancement filters, the GUI automatically calculated the percentage of voxels within each ROI exhibiting progressive (Type I), plateau (Type II), or washout (Type III) enhancement kinetics [22]. Enhancement type for each voxel was designated with reference to $S_{\text{slope}}$ for that voxel, defined as

$$S_{\text{slope}} = \left( \frac{S_{\text{tail}} - S_{\text{peak}}}{S_{\text{peak}}} \right) \times 100\%$$

where $S_{\text{peak}}$ was the peak signal intensity during the first 120 s following contrast injection and $S_{\text{tail}}$ was the average signal intensity over the final two dynamic acquisitions. Voxels were designated as having progressive (Type I) enhancement kinetics if $S_{\text{slope}}$ was $\geq 10\%$ or greater, plateau (Type II) if $S_{\text{slope}}$ was between $-10\%$ and $+10\%$, and washout (Type III) if $S_{\text{slope}}$ was $-10\%$ or less. In order to minimize anomalies from spurious signal intensity fluctuations over the course of a dynamic contrast run, the entire time–signal intensity curve for each voxel was smoothed by a moving average filter of length three before entering into the enhancement kinetics analysis.

2.5. Statistical analysis

A non-parametric Spearman rank correlation analysis was used to estimate the correlation between the two radiologists with respect to observed imaging parameters. Logistic regression analysis was then performed in two steps. First, as a preliminary step to select covariates, a Mann–Whitney U test was employed to examine the median difference of continuous variables (e.g., size measurements) between pathological responders and non-responders, and a Fisher’s exact test was used to determine the association between categorical variables (e.g., perceived enhancement) and pathological response or non-response. For variables found to discriminate patients with a pCR at a p-value of less than equal to 0.05, receiver operating characteristic (ROC) analysis was then performed using a logistic regression model. 95% confidence intervals for the area under the curve (AUC) were generated based on 2000 bootstrap samples. All data were analyzed using publicly available R version 2.11.0 statistical software.

3. Results

3.1. Patient characteristics and pathological response

A total of 21 patients completed the study. Table 1 presents an overview of patient and tumor characteristics. Patients underwent post-cycle 1 scanning at a median of 14 days following baseline imaging (range: 7–28 days). At pathological analysis, 9 patients were classified as having achieved a pCR. Table 2 provides a summary of tumor type, scan timing, and pathological response for all patients.

3.2. Interobserver variability

The non-parametric Spearman rank correlation analysis demonstrated statistically significant correlation between both radiologists for all continuous variable measurements with exception of short-axis lesion diameter during initial phase of enhancement ($\rho$ for short-axis lesion diameter during initial phase of enhancement = 0.33, $p = 0.14$; mean $\rho$ for all other measurements = 0.77, median 0.75, $p = 0.00–0.02$). The correlation between the two radiologists was deemed sufficient to allow observations from both radiologists to be averaged together for all subsequent analyses.

### Table 1

<table>
<thead>
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<th>Variable</th>
<th>Value</th>
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<td>Range</td>
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<tr>
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</tr>
<tr>
<td>Triple negative</td>
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</tr>
<tr>
<td>Tumor grade</td>
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<tr>
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<tr>
<td>High</td>
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<td>Mastectomy</td>
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<tr>
<td>Breast conservation therapy</td>
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### Table 2

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<th>Pathological response</th>
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<td>pCR</td>
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<td>non-pCR</td>
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<td>HER2 +</td>
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</tr>
<tr>
<td>21</td>
<td>Triple negative</td>
<td>9</td>
<td>non-pCR</td>
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</tbody>
</table>


predictive ability, although percentage change in long-axis diameter measured early in enhancement did approach statistical significance (p = 0.08). Fig. 3 summarizes all of the parameter change data.

4. Discussion

In recent years, an extensive literature has emerged around advanced imaging techniques for the prediction of breast cancer response to NAC. Techniques currently under investigation include volumetric ultrasonography, positron emission tomography, and magnetic resonance imaging comprising both conventional anatomic and advanced functional imaging methods [23–35]. This is an important area of investigation because current trials are investigating the use of different NAC regimens in multiple patient groups, including patients with smaller tumors that are less amenable to reliable evaluation by palpation. An imaging technique with the ability to predict pathological response early in treatment could have important ramifications for patient care, including allowing the treating oncologist to discontinue an ineffective treatment and substitute an alternative regimen [36].

The literature around DCE-MRI in this setting is especially rich and varied. Investigations into DCE-MRI for the prediction of breast cancer response to NAC have employed different DCE-MRI temporal resolutions, data analysis methods, imaging timing relative to NAC, and outcome variables. In general, the literature segregates into two broad categories: studies employing fully quantitative analysis of high temporal resolution DCE-MRI data using complex pharmacodynamic modeling [10–14,37,38], and studies employing a variety of semi-quantitative analyses of low temporal resolution data [15–18]. The variation in approaches reflects the challenges of DCE-MRI protocol design in the breast, where there are competing imperatives for high spatial-resolution imaging (to depict lesion morphology and to maximize sensitivity for small disease foci) and high temporal-resolution imaging (to model subtle changes in vascular permeability). It also reflects a possible disconnection between highly specialized centers, which may possess the expertise and dedicated analytic resources to perform rigorous and complex pharmacokinetic modeling, and nonspecialized clinical sites, which may have access to such techniques only through commercially available computer assisted detection (CAD) software. At the present time, it is not clear how DCE-MRI should evolve and be translated into broad clinical use for assessing breast cancer response to NAC. Advocates of quantitative analysis of high temporal resolution data might highlight promising data from pharmacokinetic modeling as well as studies employing a variety of semi-quantitative approaches as potentially problematic, especially when trying to standardize image acquisition and analysis across multiple sites in a large-scale clinical trial; for example, a simple, semi-quantitative, three-point signal enhancement ratio (SER) approach [16,39] was chosen over more complex methods for use in the multi-site American College of Radiology Imaging Network (ACRIN) 6657/Investigation of Serial Studies to Predict Your Therapeutic Response with Imaging And molecular Analysis (1-SPY) trial [20].

This study was performed out of consideration that a blended technique, employing a semi-quantitative analysis of high temporal resolution DCE-MRI data, might offer useful early information on important outcome variables. In general, the literature segregates into two broad categories: studies employing fully quantitative analysis of high temporal resolution DCE-MRI data using complex pharmacodynamic modeling [10–14,37,38], and studies employing a variety of semi-quantitative analyses of low temporal resolution data [15–18]. The variation in approaches reflects the challenges of DCE-MRI protocol design in the breast, where there are competing imperatives for high spatial-resolution imaging (to depict lesion morphology and to maximize sensitivity for small disease foci) and high temporal-resolution imaging (to model subtle changes in vascular permeability). It also reflects a possible disconnection between highly specialized centers, which may possess the expertise and dedicated analytic resources to perform rigorous and complex pharmacokinetic modeling, and nonspecialized clinical sites, which may have access to such techniques only through commercially available computer assisted detection (CAD) software. At the present time, it is not clear how DCE-MRI should evolve and be translated into broad clinical use for assessing breast cancer response to NAC. Advocates of quantitative analysis of high temporal resolution data might highlight promising data from pharmacokinetic modeling as well as studies employing a variety of semi-quantitative approaches as potentially problematic, especially when trying to standardize image acquisition and analysis across multiple sites in a large-scale clinical trial; for example, a simple, semi-quantitative, three-point signal enhancement ratio (SER) approach [16,39] was chosen over more complex methods for use in the multi-site American College of Radiology Imaging Network (ACRIN) 6657/Investigation of Serial Studies to Predict Your Therapeutic Response with Imaging And molecular Analysis (1-SPY) trial [20].

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This study was performed out of consideration that a blended technique, employing a semi-quantitative analysis of high temporal resolution DCE-MRI data, might offer useful early information on
breast cancer response to NAC. We adapted a straightforward method used previously by other investigators [21] to categorize individual voxels as having progressive (Type I), plateau (Type II), or washout (Type III) enhancement kinetic curves, and we found that after a single cycle of NAC, a reduction in the percentage of voxels demonstrating washout (Type III) kinetics with a $\geq 100\%$ enhancement filter was significantly associated with pCR. Our AUC of 0.77 for this single predictor variable compares favorably with the AUC of 0.73 found in the ACRIN/I-SPY trial for a four-predictor variable model [20]. Other imaging parameters in our study, including lesion size measurement and subjective assessment of lesion enhancement, did not discriminate between pathological responders and non-responders.

Fig 1 illustrates imaging results for a patient who achieved a pCR. In this example, there was a subjectively appreciable decrease after one cycle of NAC in the proportion of intralesional voxels exhibiting washout kinetics (coded red), although the lesion itself had not changed significantly in size. At the completion of NAC, however, the lesion had disappeared. The findings suggest that in this patient, a biological treatment response manifested as a change in tumor vascularity that was detectable earlier than a change in gross tumor size.

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There are a number of potential advantages to our approach. First, the technique is simple and requires significantly less effort, resources, and expertise than a formal quantitative analysis. Traditional quantitative DCE-MRI analysis typically requires a pre-contrast T1 map, accurate estimation of the arterial input function (i.e., the time rate of change of the concentration of contrast agent in a feeding artery), and pharmacokinetic modeling; none of those data or analyses was employed to arrive at the results presented here. Second, by virtue of its simplicity, this method may be more robust to variations in scanner settings, platforms, and imaging protocols, especially when deployed across multiple sites in a large-scale clinical trial, although this would have to be proved systematically. Third, a voxelwise analysis may offer the advantage of improved depiction of tumor heterogeneity when compared to some other semiquantitative approaches that average parameter values across an entire lesion ROI.

We emphasize that this is a preliminary study with several limitations owing to our small sample size. First, our study population was heterogeneous, including patients with different breast tumor subtypes and different treatment regimens that also affected scan timing; our sample size was not large enough to perform subanalyses on response prediction performance within these subgroups. Second, our study was not powered to allow for comparison of our semi-quantitative technique against a fully quantitative pharmacodynamic modeling analysis, and we therefore make no claim that one technique is superior to the other. Third, although reduction in the percentage of voxels demonstrating washout kinetics was able to discriminate patients with pCR from patients without pCR, there remained some overlap between the two groups; indeed, we note that the greatest percentage decrease in intravoxel washout voxels was observed in a patient who did not achieve a pCR (Fig. 3), a finding that will require an explanation before this approach can be considered ready for clinical translation. We anticipate the results of the present study being most useful for motivating other investigators to conduct retrospective analysis of similar datasets and for guiding future studies.

In conclusion, we have shown that semi-quantitative analysis of enhancement kinetics data from a high temporal resolution DCE-MRI acquisition can predict pathological response of LABC to a single cycle of NAC. Our results demonstrate the feasibility of incorporating a relatively simple semi-quantitative DCE-MRI analysis for early response assessment in settings that lack the expertise and resources for dedicated, formal quantitative analysis. These results suggest that important early treatment response information can be obtained from high temporal resolution DCE-MRI without the use of rigorous pharmacokinetic modeling.

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References


