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RESEARCH ARTICLE

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# Development of a Non-invasive Assessment of Hypoxia and Neovascularization with Magnetic Resonance Imaging in Benign and Malignant Breast Tumors: Initial Results

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## Abstract

**Purpose:** To develop a novel magnetic resonance imaging (MRI) approach for the noninvasive assessment of hypoxia and neovascularization in breast tumors.

**Procedures:** In this IRB-approved prospective study, 20 patients with suspicious breast lesions (BI-RADS 4/5) underwent multiparametric breast MRI including quantitative BOLD (qBOLD) and vascular architecture mapping (VAM). Custom-made in-house MatLab software was used for qBOLD and VAM data postprocessing and calculation of quantitative MRI biomarker maps of oxygen extraction fraction (OEF), metabolic rate of oxygen (MRO<sub>2</sub>), and mitochondrial oxygen tension (mitoPO<sub>2</sub>) to measure tissue hypoxia and neovascularization including vascular architecture including microvessel radius (VSI), density (MVD), and type (MTI). Histopathology was used as standard of reference. Appropriate statistics were performed to assess and compare correlations between MRI biomarkers for hypoxia and neovascularization.

**Results:** qBOLD and VAM data with good quality were obtained from all patients with 13 invasive ductal carcinoma (IDC) and 7 benign breast tumors with a lesion diameter of at least 10 mm in all spatial directions. MRI biomarker maps of oxygen metabolism and neovascularization demonstrated intratumoral spatial heterogeneity with a broad range of biomarker values. Bulk tumor neovasculature consisted of draining venous microvasculature with slow flowing blood. High OEF and low mitoPO<sub>2</sub> were associated with low MVD and *vice versa*. The heterogeneous pattern of MRO<sub>2</sub> values showed spatial congruence with VSI. IDCs showed significantly higher MRO<sub>2</sub> ( $P=0.007$ ), lower mitoPO<sub>2</sub> ( $P=0.021$ ), higher MVD ( $P=0.005$ ), and lower (*i.e.*, more pathologic) MTI ( $P=0.001$ ) compared with benign breast tumors. These results indicate that IDCs consume more oxygen and are more hypoxic and neovascularized than benign tumors.

**Conclusions:** We developed a novel MRI approach for the noninvasive assessment of hypoxia and neovascularization in benign and malignant breast tumors that can be easily integrated in a diagnostic MRI protocol and provides insight into intratumoral heterogeneity.

**Key words:** MRI, Imaging biomarkers, Tumor hypoxia, Oxygen metabolism, Neovascularization, Breast cancer

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## Introduction

Breast cancer (BC) is a complex disease with remarkable intratumoral heterogeneity which leads to varied genetic, phenotypic, and behavioral characteristics; clinical presentations; and treatment responses [1–6]. Tumor hypoxia has been recognized as an important feature and a key driver of BC heterogeneity [7] that leads to the development of cell clones and an aggressive and treatment-resistant phenotype characterized by rapid progression and poor prognosis [8]. In addition, the presence of tumor hypoxia has a profound influence on the efficacy of cancer therapies and is a strong prognostic factor for disease progression, metastases, and survival [7, 9]. To survive and grow in hypoxic conditions, tumor cells co-opt adaptive mechanisms, with a key mechanism being tumor neovascularization [10]. Increased tumor neovascularization has been associated with poorer recurrence-free, cancer-specific, and overall survival, as well as failure to respond to chemotherapy [11–14].

It has to be noted that the distribution of hypoxia and induced neovascularization are not uniform within a tumor and significantly contribute to BC heterogeneity [8]. Therefore, tumor hypoxia and the induced neovascularization may provide powerful physiological stimuli that can be exploited as a tumor-specific condition, enabling the design of hypoxia-based imaging biomarkers and hypoxia-activated anti-cancer drugs. Nevertheless, to date, the assessment of tumor hypoxia, neovascularization, and consequently BC heterogeneity, and the ability to stratify patients based on the tumor's hypoxia status is limited in its clinical application due to invasiveness [15] and poor reproducibility of the technique [16, 17], high costs (positron emission tomography, PET) [18–20], or low spatial resolution (near-infrared spectroscopy, NIRS) [21–23].

New individual magnetic resonance imaging (MRI) techniques which allow a non-invasive, quantitative measurement of hypoxia as well as microvascular vessel caliber and architecture with hitherto unparalleled accuracy have been developed and pioneered in patients with brain cancers [24–26]. Advanced quantitative blood oxygenation level-dependent (qBOLD) imaging allows the absolute quantification of the tissue oxygen extraction fraction (OEF), the oxygen metabolic rate (MRO<sub>2</sub>), and the mitochondrial oxygen tension (mitoPO<sub>2</sub>) [27]. Through these, calculation of the actual tissue oxygen tension and thereby a direct measurement of tumor hypoxia becomes possible [28]. Vascular architectural mapping (VAM) can measure and quantify microvascular vessel caliber and architecture, thus depicting the development of aberrant tumor vessels indicative of neovascularization and response to antiangiogenic therapy in brain cancer [29].

In breast cancer, a technique that not only depicts tumor hypoxia but also the induced neovascularization and BC heterogeneity is necessary yet currently does not exist. We hypothesized that the non-invasive assessment of hypoxia and neovascularization with MRI is feasible in benign and malignant breast tumors and provides insights into intratumoral heterogeneity. The aim of this study was to develop a novel MRI approach for the non-invasive synergistic assessment of hypoxia and neovascularization in benign and malignant breast tumors, which can be integrated in a diagnostic MRI protocol.

## Material and Methods

The institutional review board approved this prospective, single-institution study, and all patients gave written, informed consent.

### *Patients*

Between January and March 2017, 20 consecutive female patients (mean age 49 years, range 21–84 years) who met the following inclusion criteria were enrolled and underwent multiparametric MRI of the breast at 3 Tesla: 18 years or older; not pregnant; not breastfeeding; suspicious imaging finding at mammography or breast ultrasound (Breast Imaging Reporting and Data System [BI-RADS] assessment category 4–5); lesion size  $\geq 1$  cm<sup>3</sup>; no previous treatment, *i.e.*, breast biopsy before MRI or neoadjuvant chemotherapy; and no contraindications to MRI or MR contrast agents.

### *MRI Data Acquisition*

All MRI examinations were performed on a 3 Tesla clinical scanner (Tim Trio, Siemens, Erlangen, Germany) equipped with a standard 16-channel breast coil (Sentinelle, Invivo, FL). In addition to a state-of-the-art MRI protocol [30] with T2-weighted, diffusion-weighted, and dynamic contrast enhanced (DCE) MR imaging, the following sequences were performed:

1. qBOLD imaging: (i) a multi-echo gradient echo (GE) sequence for T2\*-mapping (8 echoes; TR: 750 ms; TE: 5–40 ms) and (ii) a multi-echo spin echo (SE) sequence for T2-mapping (8 echoes; TR: 2000 ms; TE: 15–120 ms).

2. VAM: (i) a separate DWI sequence (*b* values: 0 and 850 s/mm<sup>2</sup>; TR/TE: 3000/53 ms; TA 50 s) and (ii) a dynamic susceptibility contrast (DSC) bolus-tracking perfusion MRI sequence combined with a hybrid single-shot

gradient echo spin echo (GESE) EPI readout [31] (TR: 1360 ms; TE(GE): 25 ms; TE(SE): 93 ms).

Geometric parameters were identically chosen for the four qBOLD and VAM sequences: coronal slice orientation; field-of-view (FoV):  $320 \times 240$  mm; in-plane resolution:  $2.5 \times 2.5$  mm, slice thickness: 6 mm; 8 slices. DSC GESE perfusion examinations were performed with 60 dynamic measurements during administration of a single dose (0.1 mmol/kg-bodyweight) of gadolinium-based contrast agents (GBCA; gadoterate-meglumine, Dotarem, Guerbet, France) at a rate of 4 ml/s followed by a 20-ml bolus of saline using an MR-compatible injector (Spectris, Medrad). This resulted in 60 dynamic volumes of both GE-EPI and SE-EPI for tracking the first-pass peak contrast media bolus dynamics. The additional acquisition time (TA) for the qBOLD (R2\* and R2-mapping: TA, 1.5 and 2.5 min, respectively) and VAM sequences (DSC GESE perfusion and DW-EPI: TA, 1.4 min and 50 s, respectively) was less than 7 min.

Details of the MRI sequence protocol are shown in the top part of Fig. 1. In short, the diagnostic MRI sequences (T2w TSE, axial DWI, not shown in Fig. 1) were followed by the T2- and T2\*-mapping sequences for qBOLD, the coronal DWI for VAM, and the native T1w FLASH sequence for DCE perfusion. Injection of contrast media was performed during DSC GESE perfusion sequence immediately followed by the five diagnostic dynamic T1w FLASH sequences.

Hence, we obtained three different perfusion MRI data (GE-DSC, SE-DSC, and DCE perfusion) with injection of only one single dose of contrast media. Note: coronal slice orientation was chosen in order to (i) reduce FoV and motion artifacts (*e.g.*, berating) compared to axial orientation and to (ii) facilitate AIF determination (*i.e.*, using the inflow effect of the contrast media bolus and reducing blood flow parallel to the slices).

### *MRI Data Processing and Calculation of MRI Biomarker*

Processing of qBOLD and VAM data as well as calculation of MRI biomarker maps for oxygen metabolism and neovascularization were performed with custom-made MatLab (MathWorks, Natick, MA) software.

The qBOLD data processing consisted of three steps: (i) corrections for background fields of the T2\*-mapping data [32] and for stimulated echos of the T2-mapping data [33]; (ii) calculation of relaxivity maps (R2\* and R2) from the T2\*- and T2-mapping data; and (iii) calculation of absolute blood volume (BV) and flow (BF) maps from the GE data of the DSC GESE perfusion data *via* manual selection of a local arterial input function (AIF) [34] (Fig. 1, left part of “a MRI Data Acquisition”) in GE-DSC by an experienced radiologist (K.P. or T.T.H.). These maps (R2\*, R2, BV,

BF) were used for calculation of MRI biomarker maps of oxygen metabolism, including oxygen extraction fraction (OEF), metabolic rate of oxygen (MRO<sub>2</sub>) [27], and the average mitochondrial oxygen tension (mitoPO<sub>2</sub>) [35, 36], using the equations shown in Fig. 1, left part of panel “c Calculation of MRI biomarker” highlighted with “qBOLD”. OEF describes the percent of the oxygen removed from the blood for tissue consumption. MRO<sub>2</sub> is the rate of oxygen consumed by the tissue in  $\mu\text{mol}/100 \text{ g}\cdot\text{minute}$ , and mitoPO<sub>2</sub> (in mmHg) reflects the balance between the delivery and consumption of oxygen, *i.e.*, the oxygen tension in the tissue.

The VAM data processing consisted of three steps: (i) normalization to the baseline (pre-bolus) signal which was determined as the mean of the signals from the 4th to the 15th dynamic volume, (ii) fitting of the first pass bolus curves for each voxel of the DSC GE and SE perfusion data with a previously described gamma-variate function [37], and (iii) calculation of the  $\Delta R_{2,GE}$  versus  $(\Delta R_{2,SE})^{3/2}$  diagram [31], which is the so-called vascular hysteresis loop (VHL) (Fig. 1, right part of “b MRI Data Processing”). These data were used for calculation of MRI biomarker maps of neovascularization, including microvessel density (MVD; in units of  $\text{mm}^{-2}$ ), vessel size index (VSI, *i.e.*, microvessel radius; in units of  $\mu\text{m}$ ) [38], and the microvessel type indicator (MTI; in units of  $\text{m}^{-5/2}$ ) [24]. Maps of MVD and VSI were calculated using the equations shown in Fig. 1, right part of panel “c Calculation of MRI biomarker” highlighted with “VAM”. Calculation of the MTI map which was defined as the area of the vascular hysteresis loop (VHL) signed with the rotational direction of the VHL, *i.e.*, a clockwise VHL direction, was identified with a plus sign, and a counter-clockwise VHL direction was identified with a minus sign. Based on the findings of Emblem et al. [39], if in a voxel of the VAM dataset in the vascular system venule- and capillary-like components are present, the VHL in this voxel transverses counterclockwise. If there are arterioles and capillaries, the VHL transverses clockwise.

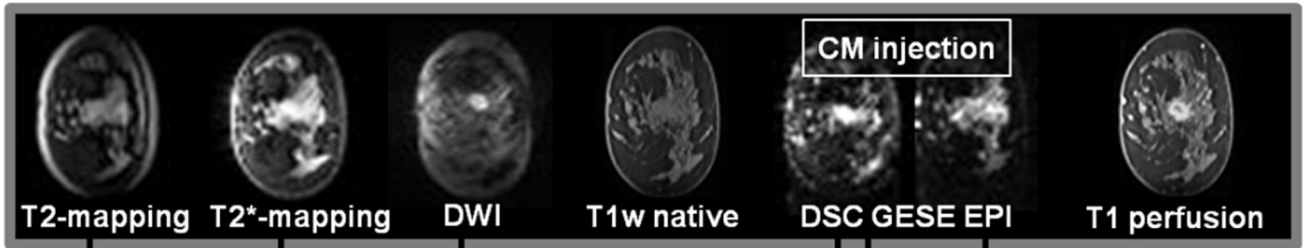
### *Quantitative Analysis*

For quantitative analysis, regions of interest (ROIs) were manually defined based on features seen in the MVD maps for high, medium, and low neovascularization. MRI biomarker values for oxygen metabolism (OEF, MRO<sub>2</sub>, mitoPO<sub>2</sub>) and for neovascularization (MVD, VSI, MTI) were averaged for each ROI. Additionally, the total tumor volumes were calculated on DCE-MR images.

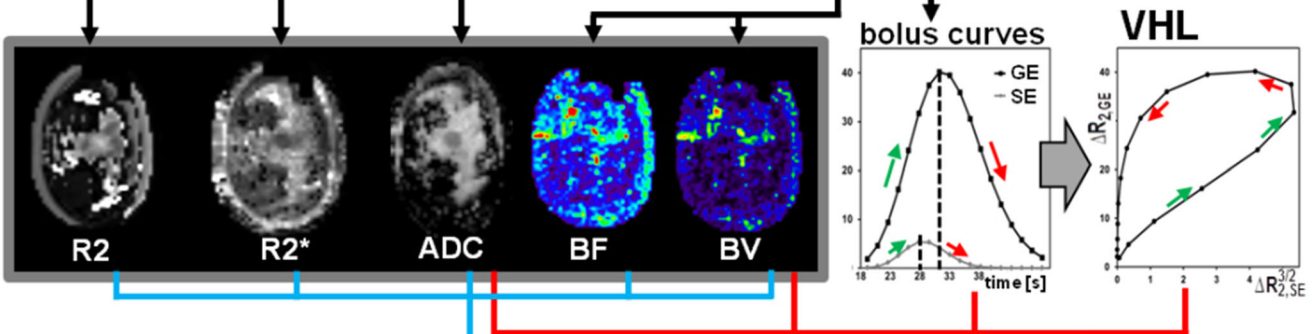
### *Histopathologic Analysis*

Histopathology was used as the standard of reference in all lesions by means of either image-guided needle biopsy or

**a MRI Data Acquisition**



**b MRI Data Processing**



**c Calculation of MRI Biomarker**

**qBOLD**

$$OEF = \frac{R2^* - R2}{k \cdot BV}$$

$$MRO_2 = \frac{C_a \cdot BF}{k \cdot BV} \cdot (R2^* - R2)$$

$$mitoPO_2 = P_{50} \cdot \left( \frac{2}{OEF} - 1 \right)^{1/h} - \frac{MRO_2}{L}$$

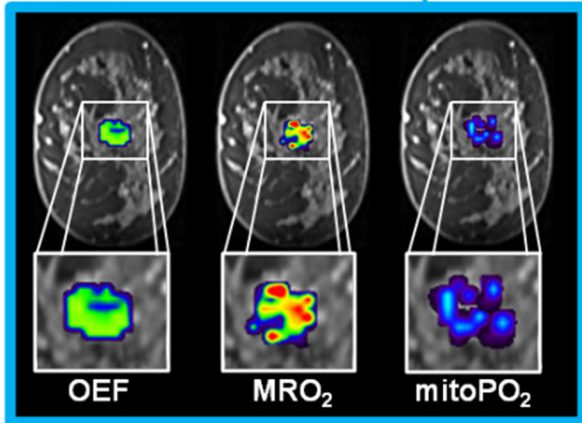
**VAM**

$$MVD = \frac{Q_{max}}{b} \cdot \left( \frac{BV^2}{4\pi \cdot ADC \cdot \bar{R}^2} \right)^{1/3}$$

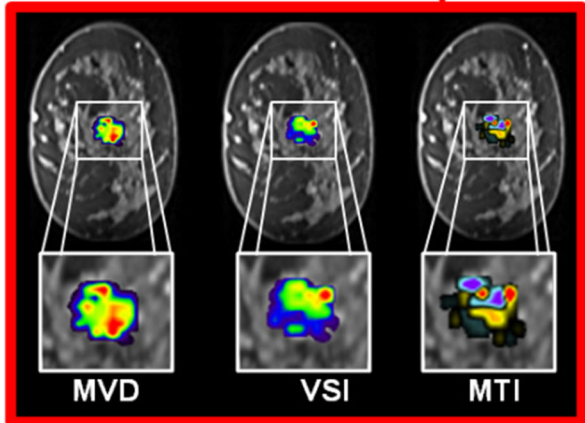
$$VSI = \left( \frac{BV \cdot ADC \cdot b^3}{2\pi \cdot Q_{max}^3} \right)^{1/2}$$

$$MTI = A_{VHL}^{asc} - A_{VHL}^{desc}$$

**d Oxygen Metabolism**



**Neovascularization**



surgery, regardless of the imaging results. All cases were read by an experienced breast pathologist. For a benign diagnosis at image-guided needle biopsy, the final diagnosis

was benign (seven of the 20 patients). For a high-risk lesion, which had an uncertain potential for malignancy, the final diagnosis was established with open surgery ( $n = 5$ ).



◀ **Fig. 1.** MRI data processing pipeline for combined non-invasive *in situ* characterization of neovascularization, oxygen metabolism, and hypoxia, respectively, of human breast tumors. Firstly, an MRI data acquisition protocol was used which was adapted from neuroimaging research to be compatible with clinical routine breast imaging purposes. Secondly, data processing of MRI raw data included calculation of relativity maps ( $R_2$  and  $R_2^*$ ), apparent diffusion coefficient (ADC), blood flow and volume (BF and BV), and vascular hysteresis loops (VHL), respectively. Finally, we used the equations shown by gray box for calculation of MRI biomarker maps of oxygen metabolism and hypoxia using the qBOLD approach and neovascularization using the VAM approach. Note:  $k = 4/3 \cdot \pi \cdot \gamma \cdot \Delta\chi \cdot \text{Hct} \cdot B_0$  ( $\gamma = 2.67502 \cdot 10^8$  rad/s/T is the nuclear gyromagnetic ratio;  $\Delta\chi = 0.264 \cdot 10^{-6}$  is the difference between the magnetic susceptibilities of fully oxygenated and fully deoxygenated hemoglobin;  $\text{Hct} = 0.42 \times 0.85$  is the microvascular hematocrit fraction);  $C_a = 8.68$  mmol/ml is the arterial blood oxygen content;  $P_{50}$  is the hemoglobin half-saturation tension of oxygen (27 mmHg),  $h$  is the Hill coefficient of oxygen binding to hemoglobin (2.7), and  $L$  (4.4  $\mu\text{mol}/\text{mmHg}$  per min) is the tissue oxygen conductivity;  $Q_{\text{max}} = \max[\Delta R_{2, \text{GE}}] / \max[(\Delta R_{2, \text{GE}})^{3/2}]$ ;  $R^- \approx 3.0$   $\mu\text{m}$  is the mean vessel lumen radius; and  $b$  is a numerical constant ( $b = 1.6781$ ). For guidance of the interpretation of MTI maps: negative MTI values were assigned to cool colors and positive to warm colors. Consequently, maps of MTI enabled differentiation between supplying arterial (areas with warm colors) and draining venous microvasculature (areas with cool colors). More specifically, a voxel with high-arterial blood volume, for example, shows a high positive MTI value and an orange to red color in the MTI maps; a capillary voxel shows a very low MTI value around 0 and a black color in the MTI maps; and a voxel with high venous blood volume shows a high, negative MTI value and a purple color in the MTI maps.

## Statistical Analysis

Software (SPSS 21, IBM, Chicago, IL, USA) was used for statistical evaluation. A Mann-Whitney test was used for nonparametric comparison of metric biomarker values. Linear regression was performed and Spearman's rank correlation coefficient was calculated to investigate the association between MRI biomarkers for oxygen metabolism for hypoxia measurement and neovascularization.  $P$  values less than 0.05 were considered to indicate significance.

## Results

qBOLD and VAM data of good quality were obtained from 20 patients (mean age 47 years, range 21–84 years). A total of 20 breast lesions were assessed. In 13 lesions, the histopathology revealed an invasive ductal carcinoma (IDC) and seven had a benign histological finding: five fibroadenoma (FA), one recurrent mastitis, and one intraductal papilloma. The mean lesion volume for these 20 patients was 9507  $\text{mm}^3$  (881–34,523  $\text{mm}^3$ ) and the lesion diameter was at least 1 cm in all three directions of space. Details on patient and lesion characteristics are summarized in Table 1.

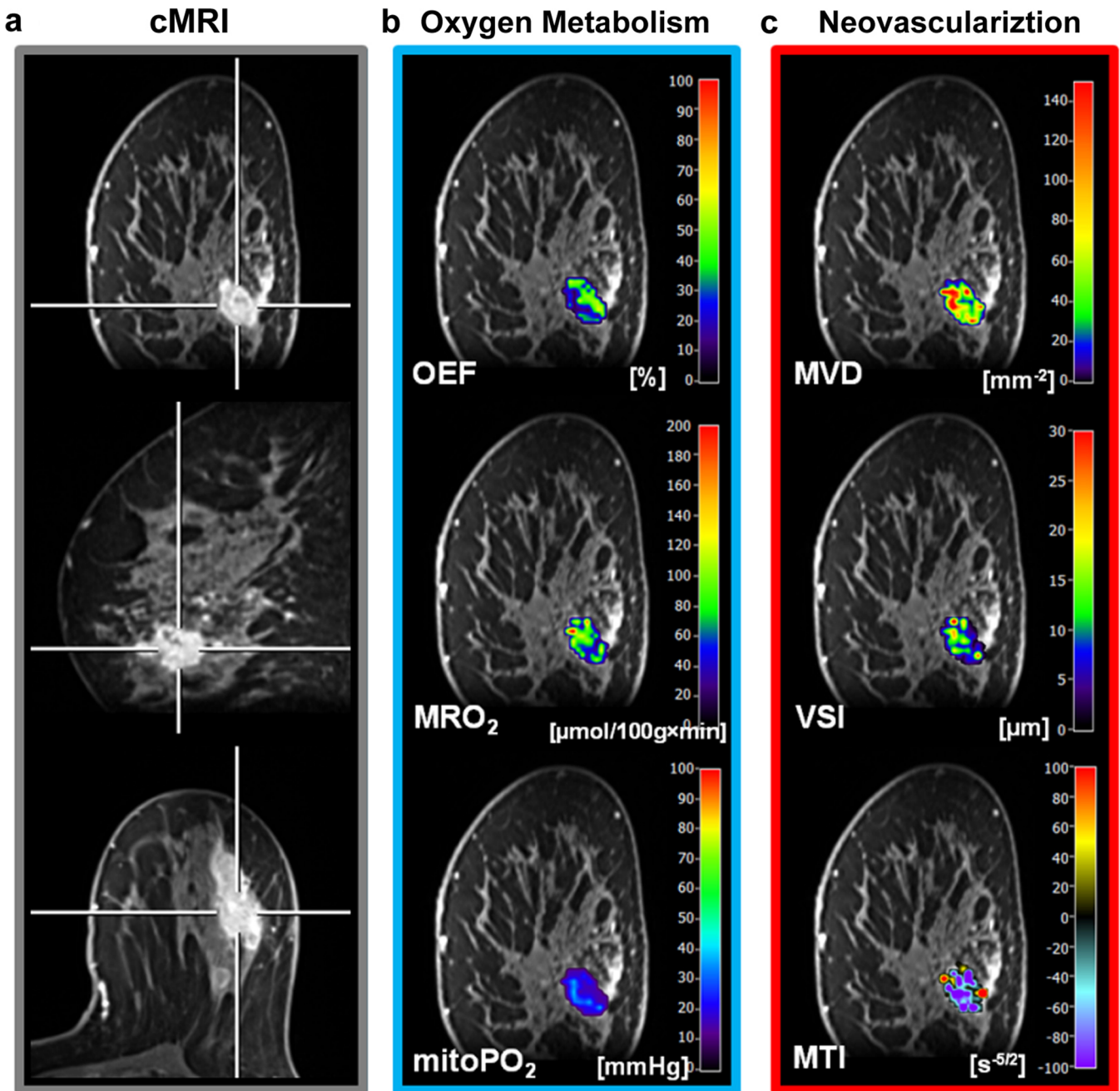
## Qualitative Analysis of MRI Biomarker Maps

A representative case for MRI-based combined depiction of tumor hypoxia through quantitative assessment of oxygen metabolism and neovascularization in a patient with IDC is depicted in Fig. 2. MRI biomarker maps of both oxygen metabolism and neovascularization

**Table 1.** Patient and tumor characteristics

ID	Age [years]	Histopathology	Location	Tumor volume [ $\text{mm}^3$ ]	BI-RADS
1	39	IDC	11:00 axis, left breast	5636	5b
2	60	IDC	9:00 axis, left breast	5817	5b
3	48	IDC	4:00 axis, left breast	1877	5b
4	62	intraductal papilloma	8:00 axis, right breast	919	3
5	35	IDC	8:00 axis, right breast	6004	5b
6	48	IDC	3:00 axis, left breast	3760	5b
7	37	IDC	3:00 axis, left breast	3940	5b
8	21	FA	9:00 axis, left breast	34,174	2
9	56	IDC	9:00 axis, right breast	8181	5b
10	84	IDC	9:00 axis, left breast	8508	5b
11	50	IDC	3:00 axis, left breast	10,325	5b
12	58	IDC	3:00 axis, right breast	2368	5b
13	39	FA	2:00 axis, left breast	1649	2
14	53	IDC	4:00 axis, left breast	11,289	5b
15	53	IDC	central, left breast	15,482	5b
16	31	FA	9:00 axis, right breast	6589	2
17	22	FA	2:00 axis, left breast	31,656	2
18	55	recurrent mastitis	central, right breast	20,473	2
19	65	IDC	2:00 axis, right breast	7592	5b
20	26	FA	11:00 axis, right breast	3969	2

IDC invasive ductal carcinoma, FA fibroadenoma, BI-RADS Breast Imaging Reporting and Data System



**Fig. 2.** Non-invasive synergistic assessment of neovascularization, oxygen metabolism, and hypoxia, respectively, in a 53-year-old patient with an invasive ductal carcinoma (IDC, patient 14 in Table 1). **a** Conventional MRI (cMRI) using dynamic contrast-enhanced (DCE) T1-weighted perfusion MR images in coronal, sagittal, and axial orientation (top-down) show lesion size and position. The white line indicates the slice intersection. **b** Imaging biomarker maps of oxygen extraction fraction (OEF), metabolic rate of oxygen (MRO<sub>2</sub>), and mitochondrial oxygen tension (mitoPO<sub>2</sub>) in coronal orientation (top-down) as well as **c** microvessel density (MVD), vessel size index (VSI), and micro vessel type indicator (MTI) in coronal orientation (top-down) demonstrates intratumoral spatial heterogeneity. The IDC showed high MRO<sub>2</sub>, low mitoPO<sub>2</sub>, high MVD, and low (*i.e.*, more pathologic) MTI.

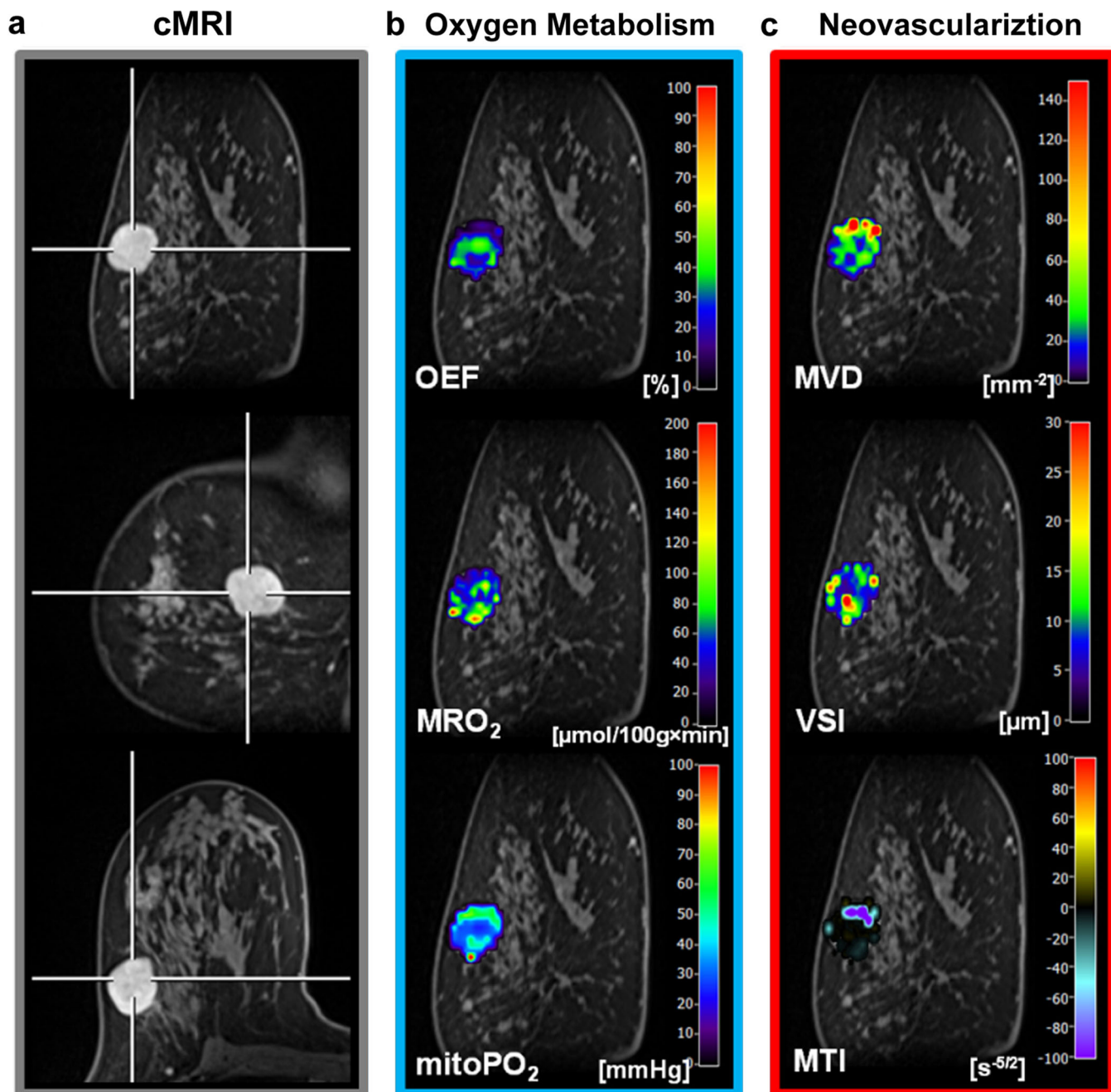
demonstrated intratumoral spatial heterogeneity with a broad range of biomarker values. High OEF and low mitoPO<sub>2</sub> (*i.e.*, hypoxia) were associated with low MVD and *vice versa*. The heterogeneous pattern of MRO<sub>2</sub> values, however, showed spatial congruence with VSI.

The MTI map revealed that the bulk of the tumor neovasculture consisted of draining venous microvasculature with slow flowing blood (purple regions). Additionally, the MTI map allowed for the detection of supplying arterial vessel structures which were

identifiable as red and orange areas. These qualitative observations were confirmed by the remaining patients including those with benign lesions. A representative case of a patient with fibroadenoma is described in Fig. 3.

### Quantitative Analysis

To investigate the intratumoral spatial heterogeneity of neovascularization, oxygen metabolism, and hypoxia, we performed an ROI-based quantitative analysis of the MRI



**Fig. 3.** Non-invasive synergistic assessment of neovascularization, oxygen metabolism, and hypoxia, respectively, in a 31-year-old patient with a benign fibroadenoma (FA, patient 16 in Table 1). **a** Conventional MRI (cMRI) using dynamic contrast-enhanced (DCE) T1-weighted perfusion MR images in coronal, sagittal, and axial orientation (top-down) show lesion size and position. The white line indicates the slice intersection. **b** Imaging biomarker maps of oxygen extraction fraction (OEF), metabolic rate of oxygen (MRO<sub>2</sub>), and mitochondrial oxygen tension (mitoPO<sub>2</sub>) in coronal orientation (top-down) as well as **c** microvessel density (MVD), vessel size index (VSI), and micro vessel indicator (MTI) in coronal orientation (top-down) demonstrate intratumoral spatial heterogeneity with lower MRO<sub>2</sub>, higher mitoPO<sub>2</sub>, lower MVD, and higher MTI as compared to malignant tumors, e.g., Fig. 2.



biomarker maps. For this purpose, ROIs were manually defined in areas showing high, moderate, and low neovascularization—as identified on the MVD maps—with high, moderate, and low MVD values, respectively. MVD were chosen for two reasons: (i) MVD is a common parameter for neovascularization, and (ii) it was demonstrated in a previous study that MVD is best suited for detection of neovascularization in brain tumors [26].

A histogram analysis of the MRI tumor hypoxia biomarkers through quantitative assessment of oxygen metabolism and neovascularization for benign (BI-RADS 2/3) and malignant breast lesions (BI-RADS 5) is shown in Fig. 4. Breast carcinoma (IDC) showed significantly higher  $MRO_2$  ( $P=0.007$ , Fig. 4b) and lower  $mitoPO_2$  ( $P=0.021$ , Fig. 4c) as well as higher MVD ( $P=0.005$ , Fig. 4d) and lower (*i.e.*, more pathologic) MTI ( $P=0.001$ , Fig. 4f) compared with benign breast lesions. In other words, IDC lesions that consumed more oxygen were more hypoxic and neovascularized compared with benign lesions.

The results of correlation analyses between MRI biomarkers of neovascularization, oxygen metabolism, and hypoxia for benign and malignant breast lesions, respectively, are shown in Fig. 5. In both benign and malignant breast lesions, OEF and MVD showed a significant negative correlation (BI-RADS 2/3:  $R=-0.506$ ,  $P=0.019$  and BI-RADS 5:  $R=-0.550$ ,  $P<0.001$ ) whereas  $MRO_2$  and VSI revealed a significant positive correlation (BI-RADS 2/3:  $R=0.583$ ,  $P=0.006$  and BI-RADS 5:  $R=0.599$ ,  $P<0.001$ ) with neovascularization, oxygen metabolism, and hypoxia, respectively (Fig. 5a, b, d, e).

In other words, the higher the MVD, the lower the necessary extraction of oxygen from the vasculature (*i.e.*, OEF) to satisfy the oxygen demands. On the other hand, tumor regions with predominantly thicker microvessels (higher VSI) consumed more oxygen (higher  $MRO_2$ ). Furthermore,  $mitoPO_2$  showed significantly positive correlation with MVD only for breast cancer (IDC;  $R=0.514$ ,  $P=0.001$ ; Fig. 5f) but not for benign breast lesions ( $R=0.336$ ,  $P=0.136$ ; Fig. 5c). Hence, the lower the MVD in IDC, the lower the oxygen tension in the tissue. All other correlations between MRI biomarkers for oxygen metabolism and neovascularization were not statistically significant.

## Discussion

In breast cancer, tumor hypoxia is one of the key drivers of the development of an aggressive and treatment-resistant tumor phenotype and is a strong prognostic factor for disease progression, metastases, and survival [7–9]. In addition, several studies have reported an association between MVD and poorer recurrence-free, cancer-specific, and overall survival [11–13] as well as an association with clinical response to chemotherapy [14]. The distribution of hypoxia and induced neovascularization are not uniform within a tumor and therefore significantly contribute to BC heterogeneity [8].

To date, the assessment of tumor hypoxia, neovascularization, and BC heterogeneity is limited in their clinical application and there is a need for the development of an accurate, reliable, and non-invasive imaging technique that can assess tumor hypoxia, neovascularization, and BC heterogeneity to achieve improved tumor characterization and tailored treatment.

In this study, we developed a novel MRI approach for the non-invasive assessment of hypoxia and neovascularization in benign and malignant breast tumors. The MRI sequences can be easily integrated into a clinical routine MRI protocol, requiring less than 7 min of additional scan time and no additional injection of gadolinium-based contrast agents (GBCAs).

The proposed approach was tested in 20 consecutive patients with breast tumors and achieved good data quality in lesion with a diameter of at least 10 mm in all three spatial directions. MRI biomarker maps of neovascularization and oxygen metabolism for direct hypoxia measurement were successfully calculated.

Our approach demonstrated that breast cancer showed significantly higher  $MRO_2$  ( $P=0.007$ ), lower  $mitoPO_2$  ( $P=0.021$ ), higher MVD ( $P=0.005$ ), and lower (*i.e.*, more pathologic) MTI ( $P=0.001$ ) compared with benign breast tumors. This indicates that breast cancer consumes more oxygen and is more hypoxic and neovascularized than benign tumors. These findings confirm the importance of tumor hypoxia and neovascularization as powerful physiological stimuli that can be exploited as a tumor-specific condition and can be used to design of hypoxia-based imaging biomarkers and hypoxia-activated anti-cancer drugs [40].

To date, only a few studies have used GE-based DSC perfusion sequences as part of a DCE MRI protocol. In most of these studies [41–44], two separate perfusion sequences in combination with two injections of contrast media resulting in up to a triple dose of contrast media were necessary. As it has been recently shown that the *i.v.* application of GBCAs may cause MRI signal changes in the brain, there is an ongoing debate regarding the safety and use of GBCAs. There is international consensus that the application of GBCAs should only be considered when necessary diagnostic information cannot otherwise be obtained and therefore administration of higher doses of the contrast agent are not acceptable within the recommendation [45–47].

In line with this recommendation, we developed a technique that allows consistent and direct measurement of neovascularization, oxygen metabolism, and hypoxia with the application of the single dose of GBCA already required in a standard imaging breast MRI protocol. This is a significant advancement over prior approaches.

Wang, et al. [48] developed a dual-echo GE sequence for simultaneous measurements of T1w DCE MRI and T2\*w DSC MRI parameters in the breast that allowed the calculation of semi-quantitative parameters such as blood flow and blood volume. They showed that their approach



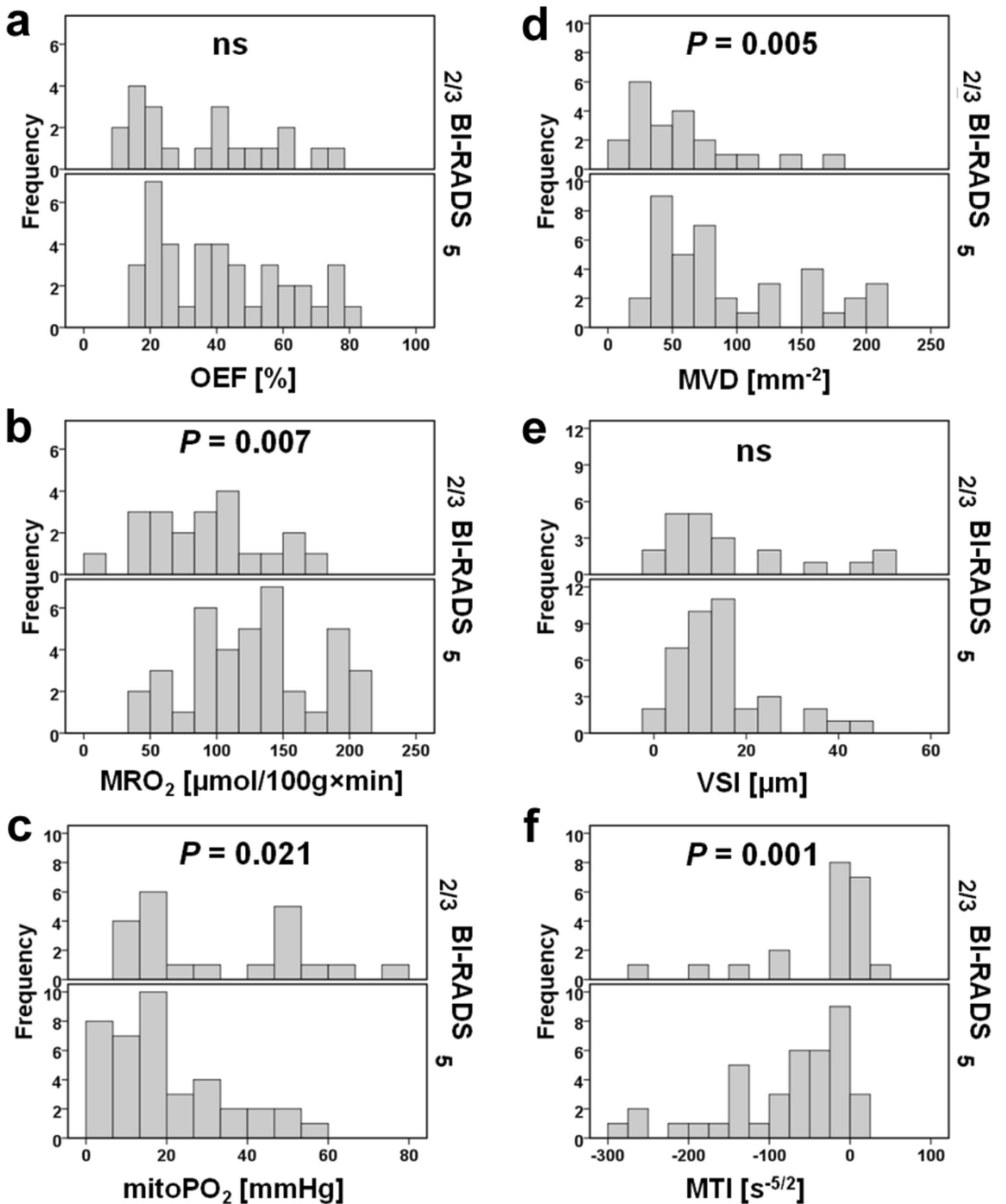


Fig. 4. Histogram analysis of **a** oxygen extraction fraction (OEF), **b** metabolic rate of oxygen ( $MRO_2$ ), **c** mitochondrial oxygen tension ( $\text{mitoPO}_2$ ), **d** microvessel density (MVD), **e** vessel size index (VSI), and **f** microvessel type indicator (MTI) for benign (BI-RADS 2/3) and malignant breast lesions (BI-RADS 5).  $P$  values show significance of difference between the histograms as calculated with a Mann-Whitney test.

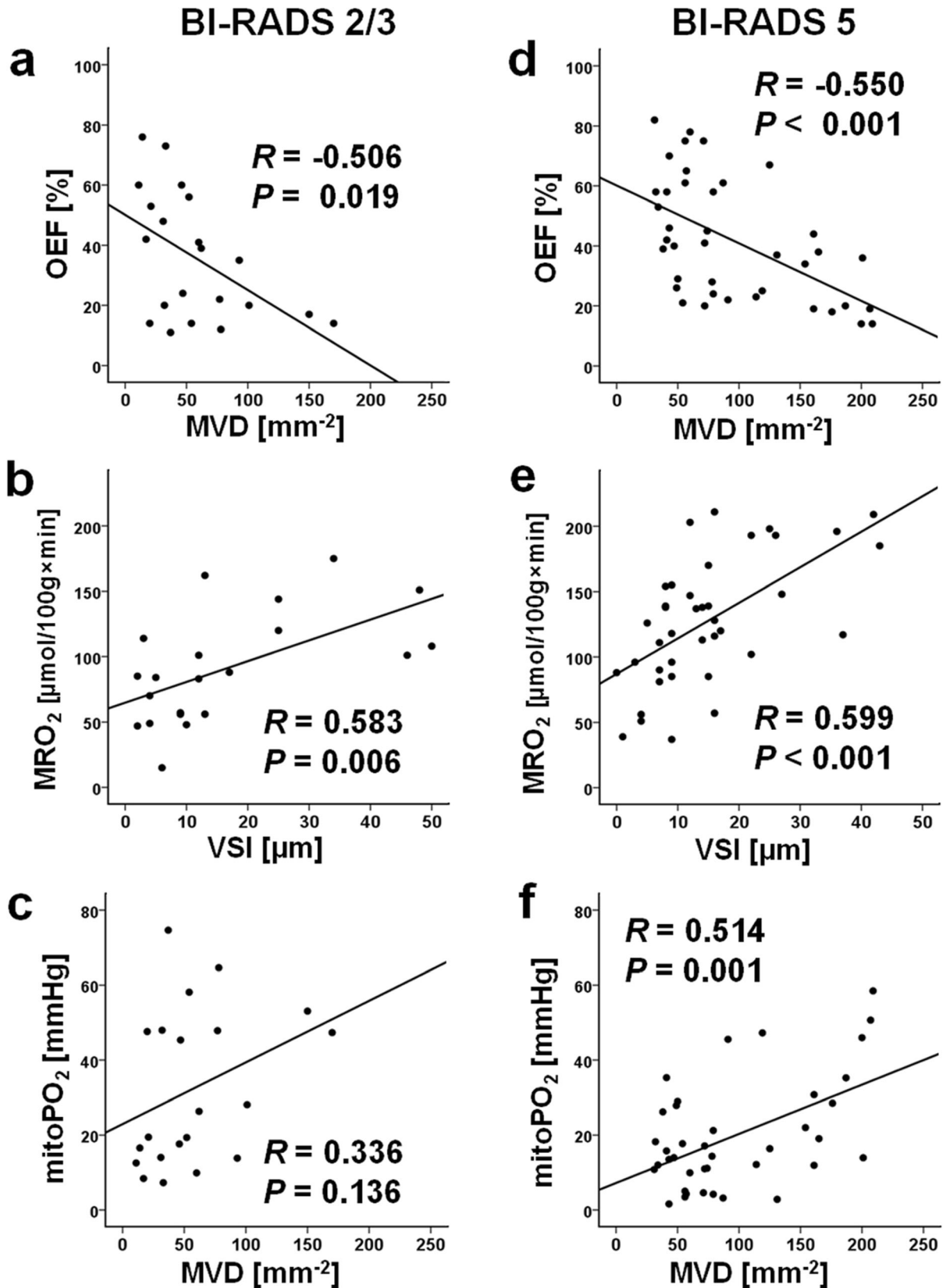


Fig. 5. Linear regression and Spearman's correlation analysis between MRI biomarker for neovascularization, oxygen metabolism, and hypoxia, respectively. Benign breast lesions (BI-RADS 2/3) revealed significant correlation between **a** extraction fraction (OEF) and microvessel density (MVD), and between **b** metabolic rate of oxygen ( $\text{MRO}_2$ ) and vessel size index (VSI), but not between **c** mitochondrial oxygen tension (mito $\text{PO}_2$ ) and MVD. Breast cancer (BI-RADS 5), however, showed significant correlation between **d** OEF versus MVD, **e**  $\text{MRO}_2$  versus VSI, and **f** mito $\text{PO}_2$  versus MVD, respectively.

improved the diagnostic performance of breast MRI for the differentiation of breast cancer from benign fibroadenomas. However, in contrast to our synergistic approach, no information about microvascular architecture, *i.e.*, MVD and VSI, was obtained as no SE-DSC perfusion MRI measurements were included. Kim et al. [49] used GE and SE steady-state susceptibility contrast MRI in an orthotopic human breast cancer xenograft model and computed MRI biomarker maps of MVD and VSI, which were validated with *ex vivo* whole tumor 3D micro-CT angiography. They found that the MRI biomarkers significantly correlated with their micro-CT analogs. The findings of the current study for MVD and VSI are in good agreement with these previous results. It has to be noted that none of the previous studies included examinations of oxygen metabolism and subsequently hypoxia with MRI or another technique. To date, invasive sampling with needle electrodes or PET using radioactive tracers such as  $^{15}\text{O}-\text{O}_2$  has been mainly used for the assessment of oxygen metabolism. Our results for tissue oxygen tension (mitoPO<sub>2</sub>) and MRO<sub>2</sub> are in good agreement with these results [50–54] yet were obtained non-invasively and without ionizing radiation, which is an inherent benefit of our approach.

In breast cancer, tumor hypoxia has been recognized as an important driver of intratumoral heterogeneity which in turn leads to the development of cell clones with an aggressive and treatment-resistant phenotype characterized by rapid progression, treatment failures, and patient deaths [1, 2, 6–9]. Prior studies that investigated tumor hypoxia in breast cancer using the PET/CT imaging with the radio-tracer [ $^{18}\text{F}$ ]fluoromisonidazole ([ $^{18}\text{F}$ ]FMISO) have shown that higher [ $^{18}\text{F}$ ]FMISO-tumor-to-background-ratio (TBR) and estrogen receptor negativity were independent predictors of shorter disease-free survival, where higher [ $^{18}\text{F}$ ]FMISO-TBR was associated with higher plasma levels of angiogenic hypoxic markers [55]. Cheng, et al. also showed that [ $^{18}\text{F}$ ]FMISO PET/CT can be used to predict primary endocrine resistance in ER-positive breast cancer [20]. A limitation of the assessment of tumor hypoxia with PET/CT is the associated radiation dose and long up-take and examination times. In addition, in comparison to MRI, the spatial resolution is much lower limiting the detailed depiction of intratumoral heterogeneity. In the current study, MRI biomarker maps of neovascularization, oxygen metabolism, and hypoxia, respectively, showed intratumoral spatial heterogeneity with a broad range of MRI biomarker values. We found that OEF inversely and mitoPO<sub>2</sub> directly correlated with MVD. This might be explained by the fact that in a less dense tumor microcirculation (low MVD), higher extraction of oxygen from the vasculature (high OEF) is necessary to satisfy the oxygen demands of the tumor. Additionally, the tissue oxygen tension is lower in these regions compared with areas with higher MVD. Consequently, neovascularization is initiated to sustain oxygen and nutrient supply and enable the further progression of

the tumor. The further the development of the tumor microcirculation as driven by ongoing neovascularization, the higher the VSI and the possible oxygen consumption (MRO<sub>2</sub>) of the tumor region. In this study, we expand on the current knowledge by providing non-invasive insight into the distinct intratumoral heterogeneity of breast cancer, which in the future may be used to develop hypoxia-based imaging biomarkers and guide targeted treatment with hypoxia-activated anticancer drugs.

A current limitation of the study is that due to the lower signal-to-noise ratio of the employed sequence, in particular regarding the SE-DSC data compared with a standard T1w DCE perfusion sequence, only tumors with a diameter of at least 10 mm in all three spatial directions were included. However, there is active research to improve spatial resolution of qBOLD and VAM imaging. It can therefore be expected that in the future, smaller lesions can also be assessed and this current limitation will be overcome. In this study, we performed no direct histopathologic correlation for MRI biomarkers. It has to be noted that previous studies that assessed MVD by immunohistochemistry of biopsy samples using antibodies against different factors such as VIII, CD31, and CD34 revealed very heterogeneous results with MVD ranging between less than 10 and few hundred mm<sup>-2</sup> [56–58]. These discrepancies, apart from differences in MVD assessment methodology and patient selection criteria, might be explained by intratumoral heterogeneity which was demonstrated in this study. Therefore, histopathological evaluation of whole tumor sections precisely registered to the pre-operative MRI data which mandates sophisticated planning is required for validation and was beyond the scope of this study. The aim of this feasibility study was the development of the MRI approach, and direct histopathologic correlation with derived MRI biomarkers is part of an on-going larger-scale study.

## Conclusions

In conclusion, we developed a novel MRI approach for the non-invasive assessment of hypoxia and neovascularization in benign and malignant breast tumors and provide insight into the distinct intratumoral heterogeneity of breast cancer. Our non-invasive approach can be easily integrated in a diagnostic MRI protocol and does not require additional application of GBCAs and has the potential to improve tumor characterization and facilitate tailored breast cancer treatment. However promising, the performance, and the fashion of implementation, of this novel MRI approach for the non-invasive assessment of hypoxia and neovascularization in benign and malignant breast tumors remains to be further investigated.

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**Compliance with Ethical Standards.** The institutional review board approved this prospective, single-institution study, and all patients gave written, informed consent.

#### Conflict of Interest

The authors declare that they have no conflict of interest.

#### Ethical Approval

All procedures performed in studies involving human participants were in accordance with the ethical standards of the institutional and/or national research committee and with the 1964 Helsinki declaration and its later amendments or comparable ethical standards.

#### Informed Consent

Informed consent was obtained from all individual participants included in the study.

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