



# Clinical experience of tensor-valued diffusion encoding for microstructure imaging by diffusional variance decomposition in patients with breast cancer

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**Background:** Diffusion-weighted imaging plays a key role in magnetic resonance imaging (MRI) of breast tumors. However, it remains unclear how to interpret single diffusion encoding with respect to its link with tissue microstructure. The purpose of this retrospective cross-sectional study was to use tensor-valued diffusion encoding to investigate the underlying microstructure of invasive ductal carcinoma (IDC) and evaluate its potential value in a clinical setting.

**Methods:** We retrospectively reviewed biopsy-proven breast cancer patients who underwent preoperative breast MRI examination from July 2020 to March 2021. We reviewed the MRI of 29 patients with 30 IDCs, including analysis by diffusional variance decomposition enabled by tensor-valued diffusion encoding. The diffusion parameters of mean diffusivity (MD), total mean kurtosis ( $MK_T$ ), anisotropic mean kurtosis ( $MK_A$ ), isotropic mean kurtosis ( $MK_I$ ), macroscopic fractional anisotropy (FA), and microscopic fractional anisotropy ( $\mu FA$ ) were estimated. The parameter differences were compared between IDC and normal fibroglandular breast tissue (FGBT), as well as the association between the diffusion parameters and histopathologic items.

**Results:** The mean value of MD in IDCs was significantly lower than that of normal FGBT ( $1.07 \pm 0.27$  vs.  $1.34 \pm 0.29$ ,  $P < 0.001$ ); however,  $MK_T$ ,  $MK_A$ ,  $MK_I$ , FA, and  $\mu FA$  were significantly higher ( $P < 0.005$ ). Among all the diffusion parameters,  $MK_I$  was positively correlated with the tumor size on both MRI and pathological specimen ( $r_s = 0.38$ ,  $P < 0.05$  vs.  $r_s = 0.54$ ,  $P < 0.01$ ), whereas  $MK_T$  had a positive correlation with the tumor size in the pathological specimen only ( $r_s = 0.47$ ,  $P < 0.02$ ). In addition, the lymph node (LN) metastasis group had significantly higher  $MK_T$ ,  $MK_A$ , and  $\mu FA$  compared to the metastasis negative group ( $P < 0.05$ ).

**Conclusions:** Tensor-valued diffusion encoding enables a useful non-invasive method for characterizing

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breast cancers with information on tissue microstructures. Particularly,  $\mu$ FA could be a potential imaging biomarker for evaluating breast cancers prior to surgery or chemotherapy.

**Keywords:** Magnetic resonance imaging (MRI); diffusion-weighted imaging (DWI); tensor-valued diffusion encoding; breast cancer; invasive ductal carcinoma (IDC); breast

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## Introduction

Magnetic resonance imaging (MRI) is widely used for evaluating breast cancer because of its non-invasive nature and excellent soft tissue contrast (1-4). Particularly, single diffusion encoding along a single direction per shot has been incorporated as a key imaging technique into routine breast MRI examination to complement dynamic contrast-enhanced (DCE) MRI, since single diffusion encoding can produce contrast in tissues without using gadolinium contrast medium injections and images can be acquired rapidly (5-8).

Single diffusion encoding is a well-known, non-invasive functional MRI technique based on quantitative measure of water diffusion in tissues to provide apparent diffusion coefficient (ADC) which reflects tissue microstructure and tumor cellularity of the breast for lesion characterization; however, previous studies reported inconsistent ADC values with large variability (9-16). One possible explanation for this issue is that the ADC is an average metric that does not capture heterogeneity within individual voxels, meaning that it cannot distinguish variable tumor cellularity, extent of infiltration, and unavoidable overlap between tumorous and non-tumorous tissues (17-20). Therefore, there have been attempts to apply fitting the models of diffusion-tensor invariants based on the diffusion-weighted imaging—such as diffusion tensor imaging (DTI) and diffusion kurtosis imaging (DKI)—to the evaluation of breast tumors (21-25).

Recently, there has been an increasing effort to understand the relation between diffusion parameters and tissue heterogeneity by considering the ‘kurtosis of the diffusion propagator’ or ‘variance of the distribution of diffusivities’ in systems that contain multiple Gaussian components (25). For example, the mean kurtosis (MK) from DKI can quantify the total non-Gaussianity, which may be caused by tissue heterogeneity (26), but it cannot distinguish the specific source of the kurtosis or estimate the microscopic diffusion anisotropy. Recently, tensor-valued

diffusion encoding (27), also known as multidimensional diffusion encoding (28), has been made possible at a wide range of clinical MRI systems by efficient experimental design (29). The term ‘tensor-valued’ indicates that diffusion encoding is applied along multiple directions per signal acquisition (27,28), unlike single diffusion encoding where diffusion encoding is applied along one direction at a time. Q-space trajectory imaging (QTI) is an analysis method that uses data encoded by multiple  $b$ -values and  $b$ -tensor shapes, to decompose the total diffusional variance into two components (27,30,31), namely the ‘anisotropic variance’ caused by microscopic diffusion anisotropy related to eccentric cells and tissue structures (30,32,33), and the ‘isotropic variance’ caused by heterogeneous isotropic diffusivity due to variable cell density or tissue mixtures (30).

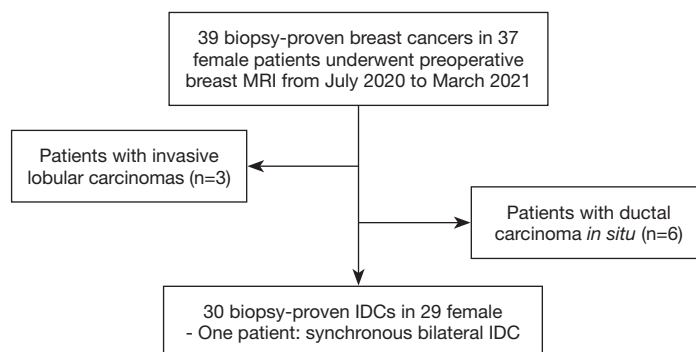
To the best of our knowledge, only one recent study has introduced the clinical potential of tensor-valued diffusion sequence in limited number of study patients with breast cancers (34). In the present study, we aimed to investigate the clinical utility of the tensor-valued diffusion sequence by assessing the microstructural detail of breast cancers in a larger number of homogeneous study patients and in a different scanning environment. Also, we present the following article in accordance with the STROBE reporting checklist (available at <https://dx.doi.org/10.21037/qims-21-870>).

## Methods

### *Study population*

The study was conducted in accordance with the Declaration of Helsinki (as revised in 2013). The study was approved by Institutional Review Board of Gyeongsang National University Changwon Hospital (No.: GNUCH 2021-06-013, retrospectively registered on June 30, 2021), and individual consent for this retrospective analysis was waived.

We searched picture archiving and communication system reports and retrospectively reviewed biopsy-



**Figure 1** Flow chart for patients' inclusion and exclusion. MRI, magnetic resonance imaging; IDC, invasive ductal carcinoma.

proven breast cancer patients who underwent preoperative breast MRI examination from July 2020 to March 2021 at Gyeongsang National University Changwon Hospital (GNUCH). We found 37 female patients with 39 breast cancers. Of them, we excluded 3 patients with invasive lobular carcinomas and 6 patients with ductal carcinoma in situ, because the number of patients was too small to provide appropriate characteristics on MRI. The final analysis included 29 female patients with 30 breast cancers that were pathologically proven as invasive ductal carcinoma (IDC) (mean age,  $49 \pm 10$  years; interval 39–72 years), and one patient had synchronous bilateral breast cancer (Figure 1). Among the 30 lesions, 26 lesions were surgically removed at GNUCH. We also reviewed prognostic cancer stages based on histopathological reports according to the 8<sup>th</sup> edition of the American Joint Committee on Cancer (AJCC) and the histologic grade with the Nottingham modification of the Scarff-Bloom-Richardson Grading System (35,36).

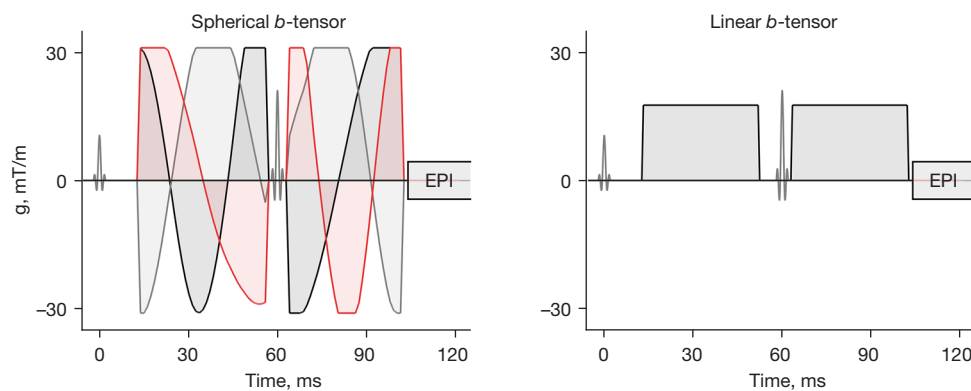
### MRI acquisition

Breast MRI was performed using the 3T system (Signa™ Architect; GE Healthcare, Waukesha, WI, USA) with an 8-channel breast coil in prone position. The examination included two morphological sequences: a 3D Dixon-based fat-suppressed T2-weighted sequence with repetition time (TR) = 2,000 ms, echo time (TE) = 90 ms, axial slices at resolution =  $0.7 \times 0.7 \times 1.2$  mm<sup>3</sup>, and an acquisition time of 3 min; and a DCE 3D T1-weighted sequence with high temporal and spatial resolution using a dual-echo 3D spoiled gradient echo sequence with Dixon fat-water separation with TR = 5.0 ms, TE = 1.7 ms, flip angle = 10 degrees, resolution =  $0.7 \times 0.7 \times 0.6$  mm<sup>3</sup>, and an acquisition time of 8 min.

The examination also included single diffusion encoding to acquire data for the regular ADC analysis, and tensor-valued diffusion encoding intended for QTI analysis. The single diffusion encoding was a spin-echo sequence with echo-planar readout with  $b$ -values 0 and 1,500 s/mm<sup>2</sup>, TR = 2,970 ms, TE = 82 ms, number of excitations = 6, number of slices of each  $b$ -value = 40, axial slices at resolution =  $1.3 \times 1.3 \times 5.0$  mm<sup>3</sup>, and an acquisition time of 3:27 min. Tensor-valued diffusion encoding was performed by a prototype pulse sequence which is named multidimensional diffusion provided by the vendor. It used gradient waveforms optimized for the MRI hardware, including compensation for concomitant gradients (37,38). Two waveforms were used to yield linear and spherical  $b$ -tensor encoding (LTE and STE) at three  $b$ -values 100, 1,000, and 2,000 s/mm<sup>2</sup>, where the LTE was directed along [4 10 15] directions and the STE was repeated [6 10 10] times for each  $b$ -value, respectively. We used TR = 9,370 ms, TE = 120 ms, resolution =  $4.3 \times 4.3 \times 5.0$  mm<sup>3</sup> with a total acquisition time of 9:47 min. The two waveforms are depicted in Figure 2. Detailed technical information on the tensor-valued diffusion sequence, sampling scheme, and gradient waveforms are available at [https://github.com/filip-szczepankiewicz/fwf\\_seq\\_resources](https://github.com/filip-szczepankiewicz/fwf_seq_resources). In addition, the following GitHub repository also contains auxiliary routines for setup of acquisition protocols and image visualization: <https://github.com/markus-nilsson/md-dmri>.

### Image processing and analysis

Motion correction and eddy-current correction were not performed, but data was inspected visually to exclude any data with gross errors. At the stage of the visual inspection, there was no patient to exclude due to gross error. We



**Figure 2** Schematic gradient waveforms producing linear and spherical tensor encodings used for obtaining tensor-valued diffusion images. EPI, echo-planar imaging.

performed QTI according to the method described in Westin *et al.* (27). We adopted the parameter nomenclature in Szczepankiewicz *et al.* (30), including the mean diffusivity (MD), the anisotropic and isotropic mean kurtosis ( $MK_A$  and  $MK_I$ ), and total mean kurtosis ( $MK_T = MK_A + MK_I$ ), as well as the fractional anisotropy (FA), which is equivalent to FA from DTI, and microscopic fractional anisotropy ( $\mu$ FA) which reflects the diffusion anisotropy without the effects of orientation dispersion. These parameters are similar to the ones employed by Naranjo *et al.* (34), however, they are derived from a cumulant expansion signal representation rather than an inverse Laplace transform. The main benefit of this is that the fitting is markedly faster and robust to the signal sampling scheme; however, it does not allow estimation of specific compartment signal fractions. The parameter estimation was performed using the open-source toolbox for multidimensional diffusion MRI (39,40).

The analysis was performed in MATLAB (R2019a, The Mathworks, Natick, MA, USA), and detail information for data analysis using MATLAB software is available at <https://github.com/markus-nilsson/md-dmri>. The regions-of-interest (ROIs) of the tumor were manually drawn on S0 maps calculated by QTI parameters, and the DCE findings were used to confirm the delineation of the tumor. All ROIs were delineated by one breast radiologist with 6 years of experience in breast MRI. Note that we avoided the inclusion of peritumoral edema, intratumoral necrosis, and normal breast parenchymal tissue. The mean values for each QTI parameter across all voxels in the ROI were calculated for each tumor. The same values were also calculated for normal fibroglandular breast tissue (FGBT) at the contralateral healthy breast.

### Histopathological analysis

A board-certified pathologist with 7 years of experience retrospectively reviewed the results of the immunohistochemical analysis of the hematoxylin-and-eosin-stained surgical specimens. These results were used to evaluate tumor size, grade, and pathologic type; estrogen receptor, progesterone receptor, c-erb-B2, and Ki-67 expression; and axillary lymph node (LN) metastasis, which was assessed through fine needle aspiration biopsy, sentinel LN biopsy, or axillary LN dissection. We used a cutoff value of 14% to classify the participants into low Ki-67 expression and high Ki-67 expression groups according to the St Gallen International Expert Consensus (41). We also classified the patients into groups based on hormonal receptors, c-erb-B2 expression (negative or positive), histologic grade (1, 2, or 3), and AJCC prognostic stage (1, 2, or 3).

### Statistical analysis

The distribution of the data was tested using the Kolmogorov-Smirnov test. For parameters with a non-normal distribution, the Mann-Whitney U-test was used to compare diffusion metrics between the IDC and FGBT and those between the IDC subgroups, and the Kruskal-Wallis test was used to compare diffusion metrics among the subgroups according to the AJCC prognostic cancer stage and IDC histologic grade. Cohen's *d* effect sizes (ES) were calculated to determine the standardized mean differences between the groups (42,43). We analyzed inter-method reliability for tumor size measurement between breast MRI and pathological examination using intraclass correlation

**Table 1** Characteristics of the 30 invasive ductal carcinomas

Variables	Total (n=30)
Age at diagnosis (year)	49±10
Amount of fibroglandular tissue	
Almost entirely fatty	1
Scattered fibroglandular tissue	10
Heterogeneous fibroglandular tissue	12
Extreme fibroglandular tissue	7
Background parenchymal enhancement	
Minimal enhancement	15
Mild enhancement	7
Moderate enhancement	5
Marked enhancement	3
Tumor size on MRI (mm) <sup>†</sup>	
≤10	4
>10–20	14
>20–40	8
>40	4
AJCC prognostic cancer stage <sup>‡</sup>	
Stage I	18
Stage II	3
Stage III	5
Lesion type	
Mass	25
Non-mass enhancement	5
Hormonal subtypes	
Luminal type	25
C-erb-B2 positive type	3
Triple negative type	2

<sup>†</sup>, tumor size defined as longest diameter on dynamic contrast-enhanced images; <sup>‡</sup>, pathologic cancer stage was determined only in the 26 patients underwent surgery at Gyeongsang National University Changwon Hospital. MRI, magnetic resonance imaging; AJCC, American Joint Committee on Cancer.

coefficient (ICC) analysis. Spearman's correlation analysis was also used to evaluate the correlations between the diffusion metrics and tumor size. All statistical analyses were conducted using SPSS, version 24.0 (IBM Corp., Armonk, NY, USA), and  $P < 0.05$  was considered significant.

## Results

### *Clinicopathological characteristics of patients and tumors*

We have summarized the clinicopathological characteristics of the 30 IDC patients in *Table 1*. The mean tumor size was  $28 \pm 23$  mm on DCE T1-weighted MRI (range, 7–87 mm) and  $25 \pm 16$  mm on pathological examination (range, 10–54 mm). Additionally, we observed good reliability between the tumor size measured on MRI and that on pathological examination (ICC = 0.80, 95% confidence interval: 0.54–0.91,  $P < 0.001$ ).

### *QTI parameters in IDC and normal FGBT*

The QTI parameters and ADC value obtained from ROI analysis of IDC and normal FGBT are summarized in *Table 2*, and a representative example of diffusion parameter maps obtained through postprocessing is depicted in *Figures 3–5*. All diffusion parameters showed significant differences between IDCs and normal FGBT. The MD and the mean ADC of IDC were significantly lower than those of normal FGBT ( $P < 0.001$ , respectively), and there was a strong correlation between the MD and the mean ADC of IDC ( $r_s = 0.75$ ,  $P < 0.001$ ). However, all other QTI parameters, including  $MK_T$ ,  $MK_A$ ,  $MK_I$ , FA, and  $\mu FA$ , were significantly higher in IDC than in normal FGBT (ES range, 0.4–2.2, all  $P < 0.05$ ). Among the parameters,  $MK_T$  showed a higher difference in mean value than MD between IDC and FGBT, and, there was a greater elevation of  $MK_T$ , which consisted of a high  $MK_A$  and low  $MK_I$ , in IDC than in FGBT. On qualitative histopathological review, IDC tended to have a higher cell density and lower cell density variance with eccentric growth and replacement along the radiating ducts and lobular structures with segmental distribution than FGBT (*Figures 3K, 3L, 4K, 4L, 5K, 5L*). Additionally,



**Table 2** Summary of QTI parameters in IDCs and normal FGBT. Cohen's *d* ES are provided to document the standardized mean differences between IDCs and FGBT. The MD and the mean ADC are given in units of  $\mu\text{m}^2/\text{ms}$ , remaining parameters are unitless

Variables	MD	MK <sub>T</sub>	MK <sub>A</sub>	MK <sub>I</sub>	FA	$\mu\text{FA}$	ADC
IDCs	1.07±0.27	1.07±0.45	0.84±0.46	0.26±0.28	0.26±0.07	0.69±0.14	0.94±0.20
FGBT	1.34±0.29	0.38±0.36	0.22±0.21	0.16±0.19	0.21±0.08	0.40±0.13	1.38±0.30
P value	<0.001	<0.001	<0.001	0.018	0.008	<0.001	<0.001
ES	1.0	1.7	1.8	0.4	0.7	2.2	1.7

Data are mean  $\pm$  standard deviation. P values were calculated using Mann-Whitney's U test. QTI, Q-space trajectory imaging; IDC, invasive ductal carcinoma; FGBT, fibroglandular breast tissue; ES, effect sizes; MD, mean diffusivity; ADC, apparent diffusion coefficient; MK<sub>T</sub>, total mean kurtosis; MK<sub>A</sub>, anisotropic mean kurtosis; MK<sub>I</sub>, isotropic mean kurtosis; FA, macroscopic fractional anisotropy;  $\mu\text{FA}$ , microscopic fractional anisotropy.

the largest ES between IDC and FGBT was seen for  $\mu\text{FA}$  (Table 2), which suggests that it outperforms conventional diffusion parameters in distinguishing the two tissue types.

### Correlation between QTI parameters and clinicopathological characteristics of IDC

Table 3 demonstrates the correlation between tumor size and diffusion parameters. MK<sub>I</sub> was positively correlated with tumor size, both on MRI and pathological examination ( $r_s=0.38$ ,  $P=0.04$  vs.  $r_s=0.54$ ,  $P=0.005$ , respectively), whereas MK<sub>T</sub> and ADC only demonstrated moderate correlations with the tumor size on pathological examination ( $r_s=0.46$ ,  $P=0.02$  vs.  $r_s=-0.48$ ,  $P=0.02$ ).

Table 4 demonstrates the comparison of diffusion parameters in IDC according to prognostic factors. IDCs with LN metastasis showed significantly higher MK<sub>T</sub>, MK<sub>A</sub>, and  $\mu\text{FA}$  than those without LN metastasis (ES: 0.99, 0.96, and 0.81, respectively). Although no significant difference was seen in any diffusion parameter between the groups according to hormonal receptor, c-erb-B2, and Ki-67 expression, the positive c-erb-B2 expression group had a higher  $\mu\text{FA}$ , with a large ES, than the negative c-erb-B2 expression group (ES: 0.83,  $P=0.23$ ). According to the histologic grades, QTI parameters—MD, MK<sub>T</sub>, MK<sub>A</sub>, MK<sub>I</sub>, FA, and  $\mu\text{FA}$ —showed no significant differences among the grade 1, grade 2, and grade 3 groups. Furthermore, there were no significant differences in the six QTI parameters among the AJCC stage 1, stage 2, and stage 3 groups.

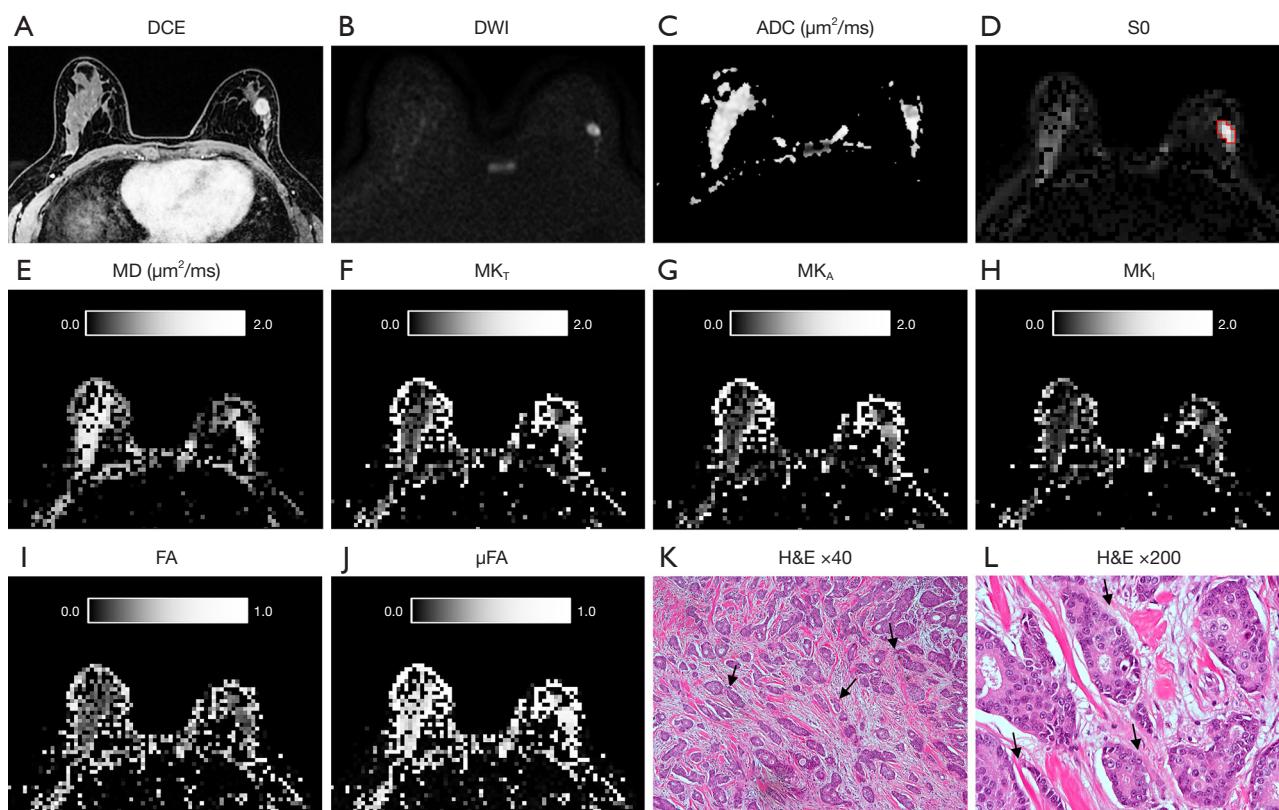
## Discussion

In this study on patients with pathologically proven IDC, we observed that IDC had a significantly lower mean value

of MD than normal FGBT. Furthermore, the mean values of QTI parameters were significantly higher in IDC than in normal FGBT. The mean MK<sub>A</sub> was significantly higher than the mean MK<sub>I</sub> in IDC, which contributed to a high MK<sub>T</sub>. Among all QTI parameters,  $\mu\text{FA}$  had the largest ES for differentiating IDC from FGBT. Additionally, IDCs with LN metastasis had higher MK<sub>T</sub>, MK<sub>A</sub>, and  $\mu\text{FA}$  than those without LN metastasis.

In real-world practice, single diffusion encoding plays a key role in differentiating between benign and malignant lesions on breast MRI by using ADC as a common biomarker to indicate the average cell density in each imaging voxel (44). However, single diffusion encoding cannot provide detailed information on tumor microstructure. Therefore, DTI is performed with conventional breast MRI, and this enables the estimation of MD and FA, which is used to determine whether a breast lesion is benign or malignant (44-46). Furthermore, MK can be estimated using DKI. MK can provide information on non-Gaussian diffusion, which may be related to microstructural tissue heterogeneity (26,47,48). Also, MK has been found to improve the accuracy of differentiation of breast lesions and have a higher diagnostic potential than ADC (49,50). However, the interpretation of MK could not be linked to the two components of diffusional variance, which decreased the specificity. To overcome this limitation, tensor-valued diffusion encoding, i.e., *b*-tensors with different shapes, may be used, as it provides detailed microstructural information by decomposing MK into its anisotropic and isotropic components (30,31).

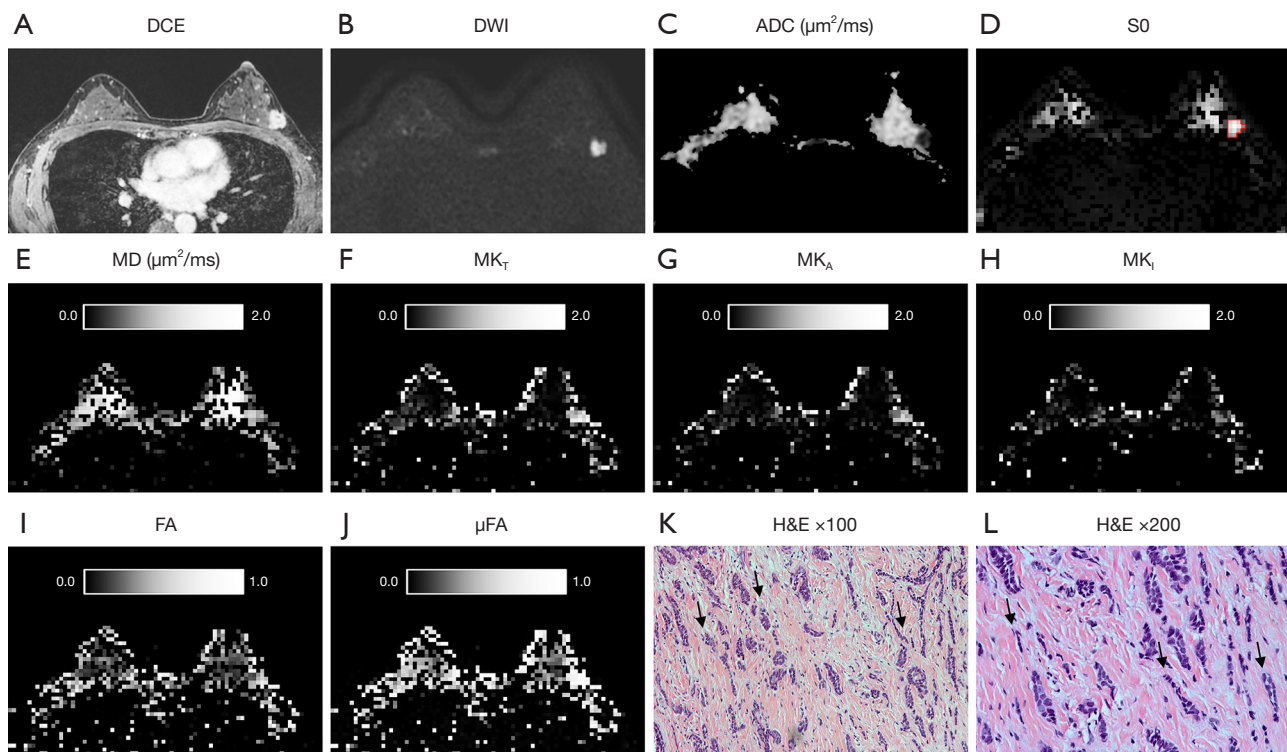
In the current study, the MD values in IDC were significantly lower than those in FGBT, which is consistent with the results of previous studies that have demonstrated a negative correlation of MD with tissue cellularity (44).



**Figure 3** A 50-year-old woman with stage I invasive ductal carcinoma in the left breast. (A) Axial DCE T1-weighted fat-suppressed image reveals a round shaped mass with heterogeneous enhancement in the left breast at the 2 o'clock position. (B,C) Single diffusion encoding image ( $b$ -value  $1,500 \text{ s/mm}^2$ ) (B) and ADC map (C) show the prominent diffusion restriction of the tumor (mean ADC,  $0.92 \mu\text{m}^2/\text{ms}$ ). (D) Region-of-interest of the tumor is defined on S0 map, referring to the DCE image and single diffusion encoding image. (E,F) MD (E) is slightly decreased (mean MD,  $1.10 \mu\text{m}^2/\text{ms}$ ) whereas  $\text{MK}_T$  (F) is elevated in the tumor. The isotropic heterogeneity and the microscopic anisotropic component cannot be distinguished on single diffusion encoding image and  $\text{MK}_T$  maps. (G,H) On the  $\text{MK}_A$  (G) map and  $\text{MK}_I$  (H) map, the tumor exhibits elevation of  $\text{MK}_A$  and relatively low  $\text{MK}_I$ , indicating that dominance of the anisotropic component in the IDC. (I,J) Macroscopic FA (I) map reveals insignificant elevation of FA in the tumor. Conversely,  $\mu\text{FA}$  (J) is increased in the tumor. (K,L) Histopathological examinations of a specimen from the breast conserving surgery shows the IDC is composed of multiple pleomorphic tumor cells with large nuclei along the radiating ductal structures. The tumor cells originated from mammary duct epithelium infiltrate and replace the surrounding duct and lobular structures with fibrosis. The tumoral and underlying structures appear well organized over longer distances along the radiating direction, that is, elongated structure resembling mammary duct (black arrows), explains the increased  $\text{MK}_A$ . Areas of higher cell density and fibrosis with lower cell density are mixed, which were reflected by increased  $\text{MK}_I$ . Also, some ducts filled with tumor cells appear to be cut in cross sections rather than in the entire longitudinal direction according to the cutting direction of the pathological specimens [ $\times 40$  (K) and  $\times 200$  (L) magnification; hematoxylin and eosin stain]. DCE, dynamic contrast-enhanced; DWI, diffusion-weighted imaging; ADC, apparent diffusion coefficient; MD, mean diffusivity;  $\text{MK}_T$ , total mean kurtosis;  $\text{MK}_A$ , anisotropic mean kurtosis;  $\text{MK}_I$ , isotropic mean kurtosis; FA, fractional anisotropy;  $\mu\text{FA}$ , microscopic fractional anisotropy; IDC, invasive ductal carcinoma.

Additionally, the MD value of IDC in our study was  $1.07 \mu\text{m}^2/\text{ms}$ , which was within the range reported ( $0.71$ – $1.62 \mu\text{m}^2/\text{ms}$ ) in previous studies (24). In addition, there was a strong correlation between the MD and mean ADC of IDC which could be interpreted that the two

parameters being influenced by cellular density within the voxel as expected (51). Although these two parameters reflect the average diffusivity, the MD has the major difference and advantage because it is free of bias from non-Gaussian diffusion and  $b$ -value dependence (25).

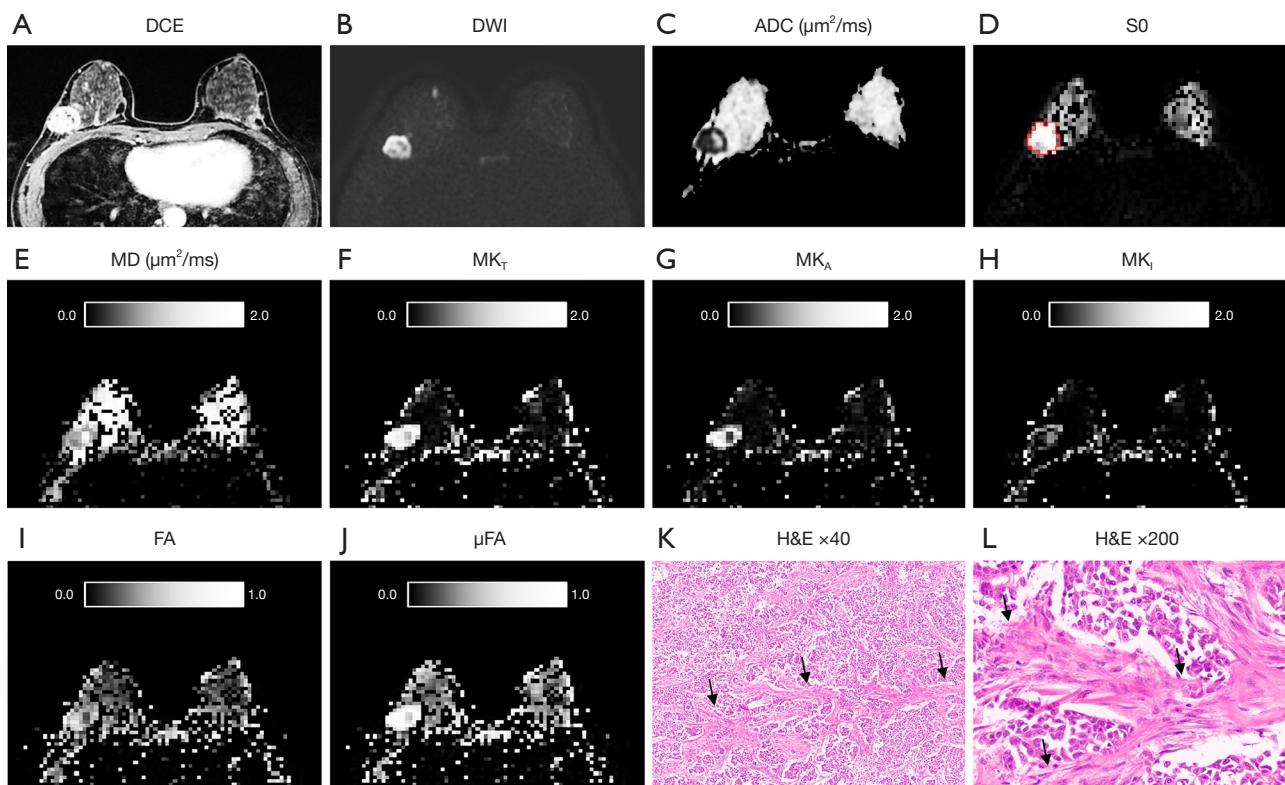


**Figure 4** A 45-year-old woman with stage II invasive ductal carcinoma in the left breast. (A) Axial DCE T1-weighted fat-suppressed image reveals an irregular mass with focal skin retraction and heterogeneous enhancement in the left breast at the 2 o'clock position. (B,C) Single diffusion encoding image ( $b$ -value  $1,500 \text{ s/mm}^2$ ) (B) and ADC map (C) show the prominent diffusion restriction of the tumor (mean ADC,  $0.58 \mu\text{m}^2/\text{ms}$ ). (D) Region-of-interest of the tumor is drawn on S0 map, referring to DCE image and single diffusion encoding image. (E,F) MD (E) is mildly decreased (mean MD,  $0.86 \mu\text{m}^2/\text{ms}$ ) and  $\text{MK}_T$  (F) is absolutely increased in the tumor. (G,H) The tumor has high  $\text{MK}_A$  and relatively low  $\text{MK}_I$  on the  $\text{MK}_A$  (G) map and  $\text{MK}_I$  (H) map, interpreted as dominance of the anisotropic component in the IDC. (I,J) FA (I) map shows slight elevation of FA in the tumor. On the other hand,  $\mu\text{FA}$  (J) is markedly increased in the tumor. (K,L) Histopathological examinations of a specimen from the breast conserving surgery shows the IDC is composed of elongated ductal structures, resembling mammary ducts that are consisted of large nuclei-contained tumor cells. This finding reflects variable cell density and dominant anisotropic diffusion effect, resulting in changes of QTI parameters. The pleomorphic tumor cells infiltrate and replace the surrounding duct and lobular structures with resembled structure. The tumoral and underlying structures appear well organized along the radiating direction, that is, ductal structure with fibrosis (black arrows), explains the higher  $\text{MK}_A$ . Also, some ducts filled with tumor cells appear to be cut in cross sections rather than in the entire longitudinal direction according to the cutting direction of the pathological specimens [ $\times 100$  (K) and  $\times 200$  (L) magnification; hematoxylin and eosin stain]. DCE, dynamic contrast-enhanced; DWI, diffusion-weighted imaging; ADC, apparent diffusion coefficient; MD, mean diffusivity;  $\text{MK}_T$ , total mean kurtosis;  $\text{MK}_A$ , anisotropic mean kurtosis;  $\text{MK}_I$ , isotropic mean kurtosis; FA, fractional anisotropy;  $\mu\text{FA}$ , microscopic fractional anisotropy; IDC, invasive ductal carcinoma; QTI, Q-space trajectory imaging.

We also found that microscopic anisotropy was a dominant component of diffusion restriction in IDC, which was consistent with the findings of a recent study (34). Interestingly, our findings were similar to those of a previous study that used tensor-valued diffusion encoding in brain tumors and found high  $\text{MK}_A$  values in meningiomas due to an abundance of eccentric cells and cell structures (30). Based on the histopathological correlation, we speculate

that, regardless of the organ studied, the consistency of the result may be related to the similar tumor growth patterns of IDCs and meningiomas; IDC has a growth pattern that extends along the ductal system, with penetration into the adjacent stroma and high cellularity, prominent nucleoli, and numerous mitoses (52). These histopathological features can indicate directionality and eccentricity based on the anatomical characteristics of tumor growth. IDCs may have





**Figure 5** A 45-year-old woman with stage III invasive ductal carcinoma in the right breast. (A) Axial DCE T1-weighted fat-suppressed image reveals an irregular mass with heterogeneous enhancement in the right breast at the 8 o'clock position. (B,C) Single diffusion encoding ( $b$ -value  $1,500 \text{ s/mm}^2$ ), (B) and ADC map (C) show the diffusion restriction of the tumor (mean ADC,  $0.77 \mu\text{m}^2/\text{ms}$ ). (D) Region-of-interest of the tumor were defined on S0 map, referring to the dynamic contrast enhancement images and single diffusion encoding image. (E,F) MD (E) is markedly decreased (mean MD,  $1.08 \mu\text{m}^2/\text{ms}$ ) whereas  $\text{MK}_T$  (F) is elevated in the tumor. The isotropic heterogeneity and the microscopic anisotropic component cannot be distinguished on single diffusion encoding and  $\text{MK}_T$  maps. (G,H) On the  $\text{MK}_A$  (G) map and  $\text{MK}_I$  (H) map, the tumor exhibits high  $\text{MK}_A$  and relatively low  $\text{MK}_I$ , indicating that the anisotropic component is dominant in the IDC. (I,J) Macroscopic FA (I) map reveals mildly elevated FA in the tumor. Conversely,  $\mu\text{FA}$  (J) is prominently increased in the tumor. (K,L) Histopathological examinations of a specimen from the breast conserving surgery shows the IDC is composed of multiple oval and pleomorphic tumor cells with large nuclei along the radiating the ductal and lobular structures, reflecting higher cell density and anisotropic diffusion effect to lead change in QTI parameters. The tumor cells originated from mammary duct epithelium infiltrate and replace the surrounding duct and lobular structures with fibrosis. The tumoral and underlying structures appear well organized over longer distances along the radiating direction, that is, fascicular (black arrows), explains the higher  $\text{MK}_A$ . Areas of higher cell density and fibrosis with lower cell density are mixed, which were reflected by  $\text{MK}_I$ . Also, some ducts filled with tumor cells appear to be cut in cross sections rather than in the entire longitudinal direction according to the cutting direction of the pathological specimens [ $\times 40$  (K) and  $\times 200$  (L) magnification; hematoxylin and eosin stain]. DCE, dynamic contrast-enhanced; DWI, diffusion-weighted imaging; ADC, apparent diffusion coefficient; MD, mean diffusivity;  $\text{MK}_T$ , total mean kurtosis;  $\text{MK}_A$ , anisotropic mean kurtosis;  $\text{MK}_I$ , isotropic mean kurtosis; FA, fractional anisotropy;  $\mu\text{FA}$ , microscopic fractional anisotropy; IDC, invasive ductal carcinoma; QTI, Q-space trajectory imaging.

features similar to meningiomas that grow along the dura mater and show hypercellularity, relatively uniform tumor cells, and a tendency to encircle adjacent tumor cells (53). Our results were also consistent to the recent study in prostate cancer that demonstrated higher  $\text{MK}_I$  and  $\text{MK}_A$

values in the tumor (54). Interestingly, this recent study also showed these two QTI parameters had significant differences within the different location of normal prostatic tissues that might be related to an increase in within-voxel diffusion heterogeneity according to its zonal anatomy,

**Table 3** Correlation between tumor size and QTI parameters.  $MK_I$  is the only parameter that demonstrates positive correlation with the tumor size measured on both MRI and the pathological specimen. The MD and the mean ADC are given in units of  $\mu\text{m}^2/\text{ms}$ , remaining parameters are unitless

Variables	MD	$MK_T$	$MK_A$	$MK_I$	FA	$\mu\text{FA}$	ADC
MRI tumor size							
$r_s$	-0.23	0.36	0.33	0.38	-0.02	0.32	-0.35
P value	0.23	0.05	0.07	0.039	0.92	0.09	0.056
Pathological tumor size							
$r_s$	-0.31	0.46	0.38	0.54	0.06	0.37	-0.48
P value	0.12	0.019	0.06	0.005	0.78	0.07	0.015

QTI, Q-space trajectory imaging;  $MK_I$ , isotropic mean kurtosis; MD, mean diffusivity; ADC, apparent diffusion coefficient;  $MK_T$ , total mean kurtosis;  $MK_A$ , anisotropic mean kurtosis; FA, macroscopic fractional anisotropy;  $\mu\text{FA}$ , microscopic fractional anisotropy;  $r_s$ , Spearman's rank correlation coefficient. P values were calculated using Spearman's correlation analysis.

reflecting the different compositions of various tissues (54). Unlike the prostate, the breast is not an organ with a detailed anatomical division, and the non-glandular tissues such stroma or fat tissue also inherently exist along the driving direction of the ductal distribution. Therefore, we could not find these differences in QTI parameters according to the detailed anatomical division.

In this study, we evaluated the correlation between QTI parameters and tumor size. However, only isotropic heterogeneity ( $MK_I$ ) showed a moderate positive correlation with tumor size on both MRI and pathological examination. The interpretation of this observation remains unclear. A potential explanation based on our histopathologic findings is that large tumors are more likely to contain poorly perfused regions with variable cell density than small tumors due to aggressive cell growth or necrosis. It is well-known that intratumoral necrosis is contributed to intratumoral heterogeneity of breast cancer which indicates its ability to adapt to a new microenvironment, leading to chemotherapy resistance or tumor progression (55-59). Considering this issue,  $MK_I$  may be a useful non-invasive parameter to estimate the intratumoral heterogeneity of breast cancer. Further studies of QTI parameters with histopathological correlation are required to clarify the relationship between intratumoral heterogeneity and treatment response in different types of breast cancers.

In this study, we also observed higher mean values of FA and  $\mu\text{FA}$  due to the disorganized growth and increased cellularity of tumors in IDC than in FGBT, and these findings are consistent with the results of previous studies of brain tumors and breast tumors (30,34,60). However, our findings are inconsistent with those of other DTI studies that

have reported lower or similar values of FA in breast cancer compared with FGBT (21,44,46,61-63). These discordant results suggest that FA has a limited role as a parameter for evaluating breast cancers. Interestingly, we observed that  $\mu\text{FA}$  was better than FA at differentiating IDC from FGBT, therefore, we believe that  $\mu\text{FA}$  can be a more reliable parameter than FA for assessing breast cancers because of the following reasons: (I)  $\mu\text{FA}$  is a straightforward microscopic analogue to FA and is closely related to  $MK_A$  (30); therefore, this parameter could be a useful biomarker for demonstrating the dominant microstructural changes in breast cancer, and (II) brain imaging studies have shown that FA as a parameter of voxel scale anisotropy is confounded by the orientation coherence of the tissue (i.e., cell elongation and cell orientation) and voxel size (60,64-66).

In addition, we found that the LN metastasis group had higher  $MK_T$ ,  $MK_A$ , and  $\mu\text{FA}$  than the metastasis negative group. These results suggest that QTI parameters could be useful non-invasive biomarkers for evaluating LN metastasis in breast cancers, because preoperative identification of LN metastasis is important for determining pretreatment decisions, such as surgical planning and adjuvant therapy (67,68).

Despite of the promising results, this study has several limitations which should be considered when interpreting the results. First, tensor-valued diffusion encoding tends to have a longer acquisition time than single diffusion encoding, and a reduced resolution may be required to maintain a high signal-to-noise ratio. This may introduce biases from dissimilar partial volume effects, especially for small lesions. Therefore, careful evaluation using this novel sequence with conventional breast MRI, including high-

**Table 4** QTI parameters in IDCs according to the prognostic factors. The MD and the mean ADC are given in units of  $\mu\text{m}^2/\text{ms}$ , remaining parameters are unitless

Factor	No. of tumors	MD	MK <sub>T</sub>	MK <sub>A</sub>	MK <sub>I</sub>	FA	$\mu\text{FA}$	ADC
Hormone receptor								
Positive	25	1.09±0.30	1.05±0.50	0.80±0.47	0.28±0.26	0.26±0.08	0.67±0.15	0.94±0.21
Negative	5	1.05±0.12	0.98±0.26	0.91±0.49	0.06±0.35	0.24±0.05	0.73±0.14	0.92±0.12
P value		0.96	0.83	0.55	0.23	0.83	0.48	0.87
ES		0.17	0.18	0.23	0.71	0.30	0.41	0.12
C-erb-B2								
Positive	3	0.97±0.06	1.13±0.16	1.10±0.55	0.03±0.50	0.23±0.06	0.79±0.15	0.84±0.02
Negative	27	1.10±0.29	1.03±0.49	0.78±0.45	0.27±0.25	0.26±0.08	0.67±0.14	0.95±0.20
P value		0.28	0.60	0.25	0.70	0.65	0.23	0.25
ES		0.62	0.27	0.63	0.61	0.42	0.83	0.77
Ki-67 <sup>†</sup>								
<14%	7	1.14±0.36	0.89±0.58	0.68±0.53	0.21±0.14	0.27±0.09	0.63±0.18	1.07±0.16
≥14%	19	1.09±0.27	1.08±0.46	0.77±0.39	0.34±0.24	0.26±0.07	0.67±0.12	0.90±0.20
P value		0.82	0.36	0.53	0.23	0.96	0.69	0.05
ES		0.16	0.36	0.19	0.66	0.12	0.26	0.94
Histologic grades <sup>‡</sup>								
Grade 1	10	1.03±0.26	1.19±0.55	0.85±0.50	0.36±0.28	0.28±0.10	0.70±0.13	0.95±0.21
Grade 2	8	1.23±0.39	0.81±0.47	0.69±0.54	0.12±0.22	0.24±0.06	0.63±0.20	1.05±0.19
Grade 3	8	1.02±0.18	1.10±0.33	0.87±0.42	0.28±0.36	0.23±0.07	0.70±0.13	0.81±0.10
P value		0.46	0.27	0.61	0.14	0.61	0.80	0.09
ES		0.28	0.34	0.42	0.58	0.42	0.53	0.23
AJCC prognostic stages <sup>§</sup>								
Stage I	18	1.12±0.27	1.00±0.53	0.76±0.50	0.25±0.27	0.26±0.09	0.67±0.16	0.98±0.21
Stage II	3	0.91±0.46	1.29±0.17	1.26±0.35	0.03±0.49	0.27±0.10	0.84±0.08	1.29±0.18
Stage III	5	1.04±0.16	1.11±0.41	0.77±0.38	0.42±0.24	0.24±0.04	0.67±0.11	0.90±0.13
P value		0.29	0.37	0.15	0.28	0.93	0.11	0.20
ES		0.29	0.03	0.58	0.30	0.59	0.68	0.29
Lymph node metastasis								
Positive	11	0.98±0.24	1.30±0.37	1.08±0.49	0.28±0.30	0.26±0.09	0.75±0.12	0.90±0.08
Negative	19	1.15±0.28	0.89±0.45	0.66±0.38	0.23±0.28	0.25±0.07	0.64±0.15	0.97±0.22
P value		0.27	0.033	0.026	0.49	0.55	0.047	0.31
ES		0.26	0.99	0.96	0.17	0.12	0.81	0.42

P values for tests for equal mean values from Mann-Whitney's U test or Kruskal-Wallis test. ES are provided to document the standardized mean differences between the groups. <sup>†</sup>, Ki-67 index was measured only in the patients underwent surgery at our hospital; <sup>‡</sup>, <sup>§</sup>, histologic grade and AJCC cancer stage were determined in the only 26 patients who underwent surgery at Gyeongsang National University Changwon Hospital. QTI, Q-space trajectory imaging; ES, effect size; MD, mean diffusivity; MK<sub>T</sub>, total mean kurtosis; MK<sub>A</sub>, anisotropic mean kurtosis; MK<sub>I</sub>, isotropic mean kurtosis; FA, macroscopic fractional anisotropy;  $\mu\text{FA}$ , microscopic fractional anisotropy; ADC, apparent diffusion coefficient.

resolution single diffusion encoding, should be performed in clinical practice. Alternatively, tensor-valued diffusion encoding can be performed using MRI systems with high gradient performance (29) or by exploiting methods, such as simultaneous multi-slice imaging, to decrease scan time and yield high-resolution images in a limited amount of time. Second, we did not apply correction for motion, eddy-currents, or geometric distortion. We expect that such post-processing may improve spatial sharpness, and parameter accuracy, especially in regions of high contrast, however, implementing such methods was outside the scope of the present study. Third, the interpretation of  $MK_I$  and  $MK_A$  in terms of diffusional variance caused by anisotropic and isotropic variance is based on the assumption that the diffusion in each voxel has an approximately multi-Gaussian distribution (30). Since the experiment may also encode time-dependent diffusion (69), the metrics may be confounded by the size of the restrictions, intra-compartment kurtosis, and exchange (69,70). In addition, we did not perform quantitative microscopy, instead, radiological and pathological correlation was qualitative in the same manner of the recent previous studies (54,71). Therefore, further studies should corroborate our interpretation of the parameters by evaluating the association of QTI parameters with quantitative histopathology.

In conclusion, we found that QTI parameters reflected changes in tissue microstructures in IDC. The predominant histopathological features of IDC were high  $MK_A$  and  $\mu FA$ , which represent the eccentricity of tumor cells and cell structures.  $\mu FA$  was superior to other parameters for differentiating IDC from FGBT; therefore, it could be a potential imaging biomarker for evaluating breast cancers. QTI is not affected by tissue organization as DTI is, hence, tensor-valued diffusion encoding can provide additional information about microstructural compartments in the breast cancer where DTI is limited (72,73). Therefore, QTI parameters based on tensor-valued diffusion encoding may facilitate a deeper understanding of tissue microstructure and intratumoral heterogeneity in patients with breast cancer who are prior to undergo surgery or received chemotherapy.

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*Ethical Statement:* The authors are accountable for all aspects of the work in ensuring that questions related to the accuracy or integrity of any part of the work are appropriately investigated and resolved. The study was conducted in accordance with the Declaration of Helsinki (as revised in 2013). The study was approved by Institutional Review Board of Gyeongsang National University Changwon Hospital (No. GNUCH 2021-06-013, retrospectively registered on June 30, 2021), and individual consent for this retrospective analysis was waived.

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