Evaluation of intratumoral heterogeneity by using diffusion kurtosis imaging and stretched exponential diffusion-weighted imaging in an orthotopic hepatocellular carcinoma xenograft model

Ran Guo1,2, Shuo-Hui Yang3#, Fang Lu3, Zhi-Hong Han4, Xu Yan5, Cai-Xia Fu6, Meng-Long Zhao1,2, Jiang Lin1,2#

1Department of Radiology, Zhongshan Hospital, Fudan University, Shanghai 200032, China; 2Shanghai Institute of Medical Imaging, Shanghai 200032, China; 3Department of Radiology, 4Department of Pathology, Shuguang Hospital, Shanghai University of Traditional Chinese Medicine, Shanghai 200021, China; 5MR Scientific Marketing, Siemens Healthcare, Shanghai 201318, China; 6Siemens Shenzhen Magnetic Resonance Ltd., Shenzhen 518057, China

#These authors contributed equally to this work.

Correspondence to: Shuo-Hui Yang, MD, PhD. Department of Radiology, Shuguang Hospital, Shanghai University of Traditional Chinese Medicine, 185 Pu An Road, Shanghai, 200021, China. Email: caddie_yang1980@aliyun.com; Jiang Lin, MD, PhD. Department of Radiology, Zhongshan Hospital, Fudan University, 180 Fenglin Road, Shanghai 200032, China. Email: lin.jiang@zs-hospital.sh.cn.

Background: To investigate the value of diffusion kurtosis imaging (DKI) and diffusion-weighted imaging (DWI) with a stretched exponential model (SEM) in the evaluation of tumor heterogeneity in an orthotopic hepatocellular carcinoma (HCC) xenograft model.

Methods: Thirty orthotopic HCC xenograft nude mice models were established and randomly divided into two groups, the sorafenib induction group (n=15) and control group (n=15). Every mouse in each group underwent MRI with DKI and SEM on a 1.5T MR scanner at 7, 14, and 21 days after sorafenib intervention. DKI and SEM parameters including mean kurtosis (MK), mean diffusivity (MD), α, and distributed diffusion coefficient (DDC) were measured, calculated, and compared between the two groups and among different time points. Sequential correlations between histopathological results including necrotic fraction (NF), micro-vessel density (MVD), Ki-67 index, standard deviation (SD), and kurtosis from hematoxylin-eosin staining, and DKI and SEM parameters were analyzed.

Results: MK, MD, and DDC of HCC in the sorafenib induction group were significantly higher than those in the control group at each time point (P<0.05), while α was significantly lower (P<0.05). Significantly positive correlations were found between MK and NF (r=0.693, P=0.010), SD (r =0.785, P=0.003), kurtosis (r=0.779, P=0.003), between MD and NF (r=0.794, P=0.003), SD (r=0.629, P=0.020), kurtosis (r=0.645, P=0.018), and between DDC and NF (r=0.800, P=0.003), SD (r=0.636, P=0.020), kurtosis (r=0.664, P=0.016), and significantly negative correlations were observed between α and NF (r=−0.704, P=0.009), SD (r=−0.754, P=0.003), and kurtosis (r=−0.792, P=0.003) in the sorafenib induction group.

Conclusions: DKI and SEM parameters may be potentially useful for evaluating intratumoral heterogeneity in HCC.

Keywords: Tumor heterogeneity; magnetic resonance imaging; diffusion kurtosis imaging (DKI); stretched exponential model (SEM); hepatocellular carcinoma (HCC)

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Introduction

Heterogeneity, arising from the genetic instability of tumors, leads to an apparently chaotic growth of tumor structures including highcellularity, necrosis, immature angiogenesis, hemorrhage, and myxoid change, etc. (1). It poses a critical challenge for clinicians to accurately predict the physiological behavior and clinical outcome of a tumor (2). Intratumoral heterogeneity is associated with tumor development, progression, and response to treatment (3).

Diffusion-weighted imaging (DWI) can be utilized to assess tumor cell proliferation, and tissue necrosis (4). However, the conventional mono-exponential diffusion model is not suitable for evaluating tumor heterogeneity because of the non-Gaussian behavior of water in the tumor (5). This non-Gaussian behavior of diffusion which reflects tissue heterogeneity and irregularity, however, can be explored by using DWI with high b values and advanced modeling of DWI data (6,7). Recently, several studies have reported that diffusion kurtosis imaging (DKI) and stretched exponential model (SEM) derived from DWI can provide valuable information on tumor microstructural complexity, which could be used for improved characterization of tumors, evaluation of prognosis, and monitoring of their therapeutic efficacy (5,8-10).

Hepatocellular carcinoma (HCC) is one of the most frequently diagnosed cancers worldwide (11). Despite endeavors to improve prognosis, the overall survival rate of HCC is still rather dismal since it is a heterogeneous disease with wide variations in outcomes and response to therapy (12). Although a few studies have tentatively investigated the values of DKI and SEM in demonstration of the heterogeneity of HCC (13-15), a directly quantitative correlation between DKI and SEM parameters with the histopathological results of HCC heterogeneity, is scarce.

Sorafenib has antiangiogenic and antiproliferative effects on HCC (16). Using a low dose of 10 mg/kg body weight, a prior study reported that sorafenib could induce tumor ischemia, hypoxia, and necrosis, and partially inhibit tumor proliferation and angiogenesis (16). As a result, a low dose of sorafenib could presumably more rapidly induce a heterogeneous HCC compared with its natural growth status. Hence, using a sorafenib-induced HCC xenograft animal model, we aimed to sequentially correlate quantitative DWI parameters from DKI and SEM with quantitative histopathologic tumor tissue composition for the purpose of evaluating HCC heterogeneity.

Methods

Orthotopic HCC xenograft animal model and sorafenib induction

This experiment was approved by the Institutional Animal Care and Use Committee of our hospital. The human HCC-LM3 cell line and 4–6-week-old male BALB/c nude mice (Shanghai Institute of Materia Medica, Chinese Academy of Sciences, Shanghai, China) weighing between 23 and 25 g were used to establish the orthotopic HCC xenograft model (4). Briefly, HCC-LM3 cells (5x10^6/0.2 mL/site) were inoculated subcutaneously in the left axilla of a nude mouse. When HCC tumor grew to 1 cm in diameter, it was removed and cut into the volume of 1 mm^3 blocks. A piece of tumor block was implanted into the left lobe of the liver of the nude mouse. At the 14th day after implantation, the mice were randomly divided into the sorafenib induction group (n=15) and control group (n=15). The mice in the induction group were administered daily with sorafenib solution at a dose of 10 mg/kg body weight by oral gavage (16). The mice in the control group were administered with 0.2 mL of 0.9% saline once a day. At the 7th, 14th, and 21st day after the intervention, 5 mice were taken randomly from the induction group and control group for MRI examinations and the following histopathological correlations.

MRI techniques

All HCC nude mice models were examined using a 1.5T MRI scanner (MAGNETOM Aera; Siemens Healthcare, Erlangen, Germany) with a 16-channel wrist coil. All mice were anesthetized with 1.5% sodium pentobarbital at a dose of 40 mg/kg body weight by intraperitoneal injection. The mouse was then placed in the prone position in an animal cradle with a belt holding the abdomen to limit respiratory motion.

Coronal and transverse T2-weighted imaging (T2WI) and transverse T1-weighted imaging (T1WI) were done using turbo spin echo with a section thickness of 2 mm, an intersection gap of 0.2 mm, and a field of view (FOV) of 100x100 mm^2. Other parameters of coronal T2WI sequence were as follows: repetition time (TR) =4,000.0 ms, echo time (TE) =74.0 ms, bandwidth =150.0 Hz/pixel, flip angle (FA) =150°, matrix size =272x320, acquisition time = 1 minute 48 s. Other parameters of transverse T1WI were...
time =1 minute 48 s. For DKI-SEM measurement, a free-breathing single-shot echo planar DWI sequence was applied in the transverse plane using three orthogonal diffusion directions with 6 b values [average number] of 0 [1], 500 [2], 800 [3], 1,000 [4], 1,500 [6], and 2,000 [6] s/mm$^2$ (7,13). Parameters of DKI-SEM sequences were as follows: section thickness =2 mm, intersection gap =0.4 mm, FOV =176×295 mm$^2$, matrix size =118×198, TR =5,000.0 ms, TE =75.0 ms, bandwidth =1,405 Hz/pixel. Parallel imaging with an acceleration factor of 2 was used to shorten the scan time and reduce image distortion.

**Image postprocessing and quantitative measurement**

The original DWI images were imported to an in-house developed postprocessing program based on MATLAB (Mathworks, Natick, MA, USA) for offline DKI and SEM calculation, which was described in a previous study (17). The noise of the diffusion images was first suppressed by applying a Gaussian filter with a full width at half maximum of 3 mm. A voxel-by-voxel fitting was then performed by using the following equations, from which metrics of DKI and SEM were derived: $S(b)/S(0) = \exp \left(-b \cdot MD + \frac{1}{6}b^2 \cdot MD^2 \cdot MK\right)$; $S(b)/S(0) = \exp \left[-(b \cdot DDC)\alpha\right]$ (5). $S(b)$ is the signal intensity at a particular b value, and $S(0)$ is the signal intensity without diffusion weighting. MD is mean diffusivity, a corrected apparent diffusion coefficient (ADC) without non-Gaussian bias. MK is mean kurtosis and quantifies the deviation of water moving from the Gaussian distribution. DDC represents distributed diffusion coefficient reflecting the average rate of diffusion. The heterogeneity index $\alpha$, with a range of 0–1, describes the heterogeneity of water diffusion (5). ADC of a mono-exponential model is automatically generated by the postprocessing software. The pixel-based fitted curve was also drawn by using this software (Figure 1).

Diffusion images were analyzed and measured by two independent radiologists with 12- and 25-year experience in liver MR image interpretation. Both of them were blinded to the grouping of mice and the histopathological results. Tumor size defined by the greatest dimension on the transverse T2W images was measured (4). Both transverse T1W and T2W images were reviewed to exclude hemorrhage in the tumor. If there was hemorrhage found on this greatest dimension slice, the most adjacent slice would be analyzed. Regions of interest (ROIs) were manually drawn along the tumor border on the slice showing the greatest dimension of the tumor on ADC images (4,17). Then, the ROIs were automatically

![Figure 1](image-url)
copied onto the DKI and SEM maps by the postprocessing software (Figure 2). Each observer measured the tumor twice on the same tumor slice, and the results were averaged. After a 4-week interval, these measurements were repeated to minimize any learning bias (18). The observers were aware of the goal of the study but were unaware of other information regarding the group and the time point of each model and the histopathological results. The final averaged values from the observer with 25 years of experience were used for subsequent statistical analysis.

**Histopathological analysis**

All nude mice were sacrificed after MRI examinations, and the tumors were removed and fixed in the 10% buffered formalin solution for at least 24 hours. With reference to the slice thickness of DKI-SEM sequences, each tumor was cut into the section of 2 mm thickness in its greatest dimensional transverse plane. The section was embedded in paraffin and then cut into a slice with a thickness of 3 μm.

Aperio ScanScope and Leica SCN400 (Leica biosystems, Buffalo Grove, IL, USA) were used to perform the scanning of all the histopathological slices including hematoxylin-eosin (HE), anti-CD31, and anti-Ki-67 by a pathologist with 10 years of experience in histopathological analysis. The scanning images of the whole slices were imported to a PC with a screen resolution of 1,377×768 pixels and post-processed with SlidePath.

**Figure 2** DKI and SEM images and corresponding histopathological images of a hepatocellular carcinoma in the sorafenib induction group on the 21st day. (A) Transverse T2WI shows the greatest dimension of the tumor. (B) ADC map for ROI outlining the tumor. (C) MK map (MK =1.320). (D) MD map (MD =0.761×10⁻³ mm²/s). (E) α map (α=0.735). (F) DDC map (DDC =0.602×10⁻³ mm²/s). (G) Hematoxylin-eosin staining image shows patchy and irregular necrosis (black arrows, 10x) in the tumor. (H) Anti-CD31 immunohistochemistry image shows sparsely intratumoral micro-vessels (black arrows, 20x). (I) Anti-Ki-67 immunohistochemistry image shows heterogeneously stained Ki-67 positive cells (black arrows, 20x). DKI, diffusion kurtosis imaging; SEM, stretched exponential model; T2WI, T2-weighted imaging; ADC, apparent diffusion coefficient; ROI, region of interest; MK, mean kurtosis; MD, mean diffusivity; DDC, distributed diffusion coefficient.
Gateway Client software (Leica biosystems, Buffalo, IL, USA) for magnification, observation, and final image capturing. The histopathological measurements were performed on the captured images by using ImageJ software (v.1.48; National Institutes of Health, Bethesda, MD, USA) and Image-Pro Plus 6.0 software (Media Cybernetics, Rockville, MD, USA).

HE staining was performed to demonstrate the morphology and structure of HCC cells and mesenchymal tissues. The histogram analysis completed on the whole slice of the HE staining image with ImageJ software to obtain the parameters that reflect tumor heterogeneity: standard deviation (SD) and kurtosis. An ROI was manually drawn along the tumor border on this HE slice after conversion from color to grayscale (Figure 3) (19-23).

The measurement of tumor necrotic fraction (NF) was described in detail by a previous report (24). Briefly, five areas with four in each corner and one in the center on each HE whole slice were selected at lower power (4× and 10× magnification). Then, necrotic areas were obtained with Image-Pro Plus 6.0 in three higher-power FOVs (20× magnification) in each of the aforementioned five areas. Finally, NF was calculated according to the following equation: \[ NF = \frac{\text{Area}_{\text{necrosis in FOV}}}{\text{Area}_{\text{tumor in FOV}}} \]

Anti-CD 31 staining (Abcam, Cambridge, UK) was performed to obtain the micro-vessel density (MVD). Three regions with the densest CD 31-positive vessels of anti-CD 31 staining images were found at 4× magnification. These regions were magnified by 20×, and then these corresponding images were captured (4). MVD was obtained on these captured images with ImageJ software.

Anti-Ki-67 staining (Abcam, Cambridge, UK) was done to show the cell proliferation of the tumor. Ki-67 index was obtained similarly according to a previous report (25).

Every aforementioned histopathological index was measured twice, and the results were averaged for further analysis.

**Statistical analysis**

Mann-Whitney U-test was performed to analyze the differences between DKI and SEM parameters along with the differences of histopathological results between the induction and control groups at each time point. Differences in DKI and SEM parameters among different time points in both the induction and control groups were evaluated by using the Kruskal-Wallis H-test and Mann-Whitney U-test. The Spearman rank correlation test was used to investigate the correlations between DKI and SEM parameters and histopathological results, and the Benjamini-Hochberg method was performed to control for a false discovery rate. The correlation coefficient rho (\(r\)) was obtained to evaluate the degree of correlations: \(0.0<|r|<0.2\) represented a poor or no relationship; \(0.2 \leq |r| < 0.4\) was weak; \(0.4 \leq |r| < 0.6\) was moderate; \(0.6 \leq |r| < 0.8\) was good; \(|r| > 0.8\) was excellent (4). Two-way analysis of variance (ANOVA) was performed to analyze the effects of three-time points and the corresponding accumulated effects of sorafenib on DKI and SEM parameters. All the intra- and inter-observer agreement of DKI and SEM parameters and intra-observer agreement of histopathological measurements were evaluated by using intraclass correlation coefficients (ICCs) in the induction group. The inter-observer agreement was defined as the agreement between the two observers with the averaged values from two measurements at a 4-week interval. ICCs were defined as follows: \(\leq 0.40\), poor to fair reliability; \(0.41-0.60\), moderate reliability; \(0.61-0.80\), good reliability; \(>0.80\), excellent reliability (26). Measurement data are expressed as (mean ± SD). P<0.05 was considered statistically significant. Statistical analysis was performed using the SPSS software (v.18.0; IBM, Armonk, NY, USA).

**Results**

The DKI-SEM acquisition and analysis were successfully performed in all 30 mice, and all the tumors were clearly displayed on diffusion images. Intratumoral hemorrhage was observed in two mice in the induction group (one at the 14th day and one at the 21st day after sorafenib intervention) and in one mouse in the control group (at the 21st day after saline administration), but none of them were found on the measuring slice of the tumor.

**Comparisons of DKI and SEM parameters between the sorafenib induction group and control group**

MK, MD, and DDC of HCC in the sorafenib induction group were significantly higher than those in the control group at each time point, whereas \(\alpha\) was significantly lower (Table 1, Figure 4).

**Comparisons of DKI and SEM parameters at different time points in the sorafenib induction group and control group**

Kruskal-Wallis H-test showed that DKI and SEM...
parameters of the induction group at different time points were significantly different (MK: $\chi^2=12.150$, $P=0.002$; MD: $\chi^2=11.786$, $P=0.003$; $\alpha$: $\chi^2=11.180$, $P=0.004$; DDC: $\chi^2=12.020$, $P=0.002$). In the induction group, MK, MD, and DDC were significantly higher on the 14th and 21st days than those on the 7th day (MK: $Z=-2.440$, $P=0.015$ and $Z=-2.627$, $P=0.009$; MD: $Z=-2.402$, $P=0.016$ and $Z=-2.611$, $P=0.009$; DDC: $Z=-2.402$, $P=0.016$ and $Z=-2.611$, $P=0.009$). Significant increases of these parameters were also found on the 21st day vs. 14th day (MK: $Z=-2.627$, $P=0.009$; MD: $Z=-2.514$, $P=0.012$; DDC: $Z=-2.611$, $P=0.009$). Meanwhile, $\alpha$ was significantly lower on the 14th and 21st days compared with the 7th day ($Z=-2.207$, $P=0.027$ and $Z=-2.627$, $P=0.009$), and lower on the 21st day when compared with the 14th day ($Z=-2.440$, $P=0.015$) (Figure 4). In the control group, MK increased slightly on the 14th and 21st days, while $\alpha$ displayed a decreasing trend from the 7th to the 21st day (P>0.05). MD and DDC declined slightly on the 14th day, and then increased on the 21st day (P>0.05). No significant differences were found in any of these DKI and SEM parameters at three time points in the control group (MK: $\chi^2=4.973$, $P=0.083$; MD: $\chi^2=4.340$, $P=0.114$; $\alpha$: $\chi^2=12.020$, $P=0.002$).

Figure 3 Grayscale images from two whole slice hematoxylin-eosin staining HCCs and their frequency histograms of the pixel distribution on the 21st day after intervention. (A) Grayscale image of a tumor in the sorafenib induction group (SD =20.645, kurtosis =5.451); (B) grayscale image of a tumor in the control group (SD =16.161, kurtosis =2.850); (C) frequency histograms of the pixel distribution in the grayscale images of the tumors in sorafenib induction group (blue) and control group (red). HCC, hepatocellular carcinoma; SD, standard deviation.
Comparisons of tumor size and histogram parameters in hematoxylin-eosin staining between the sorafenib induction group and control group

At each time point, the tumor size in the induction group was slightly smaller than that in the control group with no significant difference observed ($7^{\text{th}}$ day: $1.220\pm0.264$ vs. $1.240\pm0.136$ cm, $P=0.750$; $14^{\text{th}}$ day: $1.640\pm0.215$ vs. $1.660\pm0.080$ cm, $P=0.595$; $21^{\text{st}}$ day: $1.680\pm0.117$ vs. $1.940\pm0.301$ cm, $P=0.240$).

SD and kurtosis of histogram parameters in the induction group were significantly higher than those in the control group ($P<0.05$) ($\chi^{2}=3.438$, $P=0.179$; DDC: $\chi^{2}=1.758$, $P=0.415$).

### Table 1 Comparisons of DKI and SEM parameters of HCC between the sorafenib induction group and control group

<table>
<thead>
<tr>
<th>Group</th>
<th>MK ($\times10^{-3}$ mm$^2$/s)</th>
<th>MD ($\times10^{-3}$ mm$^2$/s)</th>
<th>$\alpha$</th>
<th>DDC ($\times10^{-3}$ mm$^2$/s)</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>7 days</td>
<td>14 days</td>
<td>21 days</td>
<td>7 days</td>
</tr>
<tr>
<td>Induction</td>
<td>1.247±0.012</td>
<td>1.279±0.021</td>
<td>1.352±0.048</td>
<td>0.705±0.015</td>
</tr>
<tr>
<td>Control</td>
<td>1.182±0.049</td>
<td>1.200±0.030</td>
<td>1.240±0.023</td>
<td>0.677±0.015</td>
</tr>
<tr>
<td>Z</td>
<td>−2.214</td>
<td>−2.619</td>
<td>−2.619</td>
<td>−2.193</td>
</tr>
<tr>
<td>P</td>
<td>0.027</td>
<td>0.009</td>
<td>0.009</td>
<td>0.028</td>
</tr>
</tbody>
</table>

DKI, diffusion kurtosis imaging; SEM, stretched exponential model; HCC, hepatocellular carcinoma; MK, mean kurtosis; MD, mean diffusivity; DDC, distributed diffusion coefficient.

**Figure 4** Sequential comparisons of DKI and SEM parameters of HCC in the sorafenib induction group and control group at different time points. Box-plots show sequential changes of MK (A), MD (B), $\alpha$ (C), and DDC (D) in these two groups. All parameters were tested with Mann-Whitney $U$-test (*$P<0.05$, **$P<0.01$). Center line = median; upper and lower margins of box = upper and lower quartiles. DKI, diffusion kurtosis imaging; SEM, stretched exponential model; HCC, hepatocellular carcinoma; MK, mean kurtosis; MD, mean diffusivity; DDC, distributed diffusion coefficient.

Comparisons of tumor size and histogram parameters in hematoxylin-eosin staining between the sorafenib induction group and control group

At each time point, the tumor size in the induction group was slightly smaller than that in the control group with no significant difference observed ($7^{\text{th}}$ day: $1.220\pm0.264$ vs. $1.240\pm0.136$ cm, $P=0.750$; $14^{\text{th}}$ day: $1.640\pm0.215$ vs. $1.660\pm0.080$ cm, $P=0.595$; $21^{\text{st}}$ day: $1.680\pm0.117$ vs. $1.940\pm0.301$ cm, $P=0.240$). SD and kurtosis of histogram parameters in the induction group were significantly higher than those in the control group ($P<0.05$) (Table 2, Figure 3).
Table 2 Comparisons of SD and kurtosis of HCC in hematoxylin-eosin staining between the sorafenib induction group and control group

<table>
<thead>
<tr>
<th>Group</th>
<th>7 days</th>
<th>14 days</th>
<th>21 days</th>
<th>7 days</th>
<th>14 days</th>
<th>21 days</th>
</tr>
</thead>
<tbody>
<tr>
<td>Induction group</td>
<td>16.903±0.531</td>
<td>18.727±1.500</td>
<td>20.449±0.421</td>
<td>−4.890±2.259</td>
<td>−1.070±1.301</td>
<td>6.250±1.718</td>
</tr>
<tr>
<td>Control group</td>
<td>15.804±0.699</td>
<td>16.153±0.758</td>
<td>17.119±1.110</td>
<td>−9.346±2.158</td>
<td>−6.256±2.366</td>
<td>−1.350±4.487</td>
</tr>
<tr>
<td>Z</td>
<td>−2.193</td>
<td>−2.402</td>
<td>−2.611</td>
<td>−2.193</td>
<td>−2.611</td>
<td>−2.611</td>
</tr>
<tr>
<td>P</td>
<td>0.028</td>
<td>0.016</td>
<td>0.009</td>
<td>0.028</td>
<td>0.009</td>
<td>0.009</td>
</tr>
</tbody>
</table>

SD, standard deviation; HCC, hepatocellular carcinoma

Comparisons of histopathological results (NF, MVD and Ki-67 index) between the sorafenib induction group and control group

At each time point, NF of the induction group was significantly greater than that of the control group, and significantly lower MVD and Ki-67 index were observed in the induction group than those in the control group (P<0.05) (Table 3).

Correlations between DKI and SEM parameters and histopathological results in the sorafenib induction group and control group

In the induction group, MK, MD, and DDC showed significantly positive correlations with NF, SD, and kurtosis. α was significantly and negatively correlated with NF, SD, and kurtosis. No correlations were found between DKI and SEM parameters and MVD and Ki-67 index (Table 4). In the control group, there was no correlation between DKI and SEM parameters and NF, MVD, and Ki-67 index, and SD and kurtosis (Table 5).

Comparisons of the effects of three time points and the corresponding accumulated effects of sorafenib on DKI and SEM parameters in the HCC models

Two-way ANOVA revealed significantly accumulated effects of sorafenib from 7 to 21 days on all DKI and SEM parameters (MK: F=13.713, P<0.001; MD: F=43.904, P<0.001; α: F=15.426, P<0.001; and DDC: F=38.050, P<0.001). However, no significant effects from these three time points on DKI and SEM parameters were observed (MK: F=3.151, P=0.061; MD: F=2.820, P=0.079; α: F=0.963, P=0.396 and DDC: F=1.840, P=0.180).

Intra- and interobserver agreement on DKI and SEM parameters in the sorafenib induction group

The intra- and interobserver agreements for measuring all DKI and SEM parameters were excellent (Table 6).

Intra-examiner agreement on histopathologic SD, kurtosis, NF, MVD, and Ki-67 index were 0.979 [95% confidence interval (CI): 0.941–0.993], 0.976 (95% CI: 0.932–0.992), 0.973 (95% CI: 0.922–0.991), 0.923 (95% CI: 0.790–0.974), and 0.965 (95% CI: 0.898–0.988) (P<0.001), respectively.

Discussion

The heterogeneity of a tumor is related to the presence and changes of different components within the tumor during its formation, progression, and treatment (3,27,28). Patients with higher heterogeneity are likely to have a poorer prognosis, which may be secondary to more aggressive biology or treatment resistance (3,29). Prior knowledge of tumor heterogeneity, especially in a non-invasive way, is critical both before and after treatment. This study demonstrated that parameters from DKI and SEM could largely reflect the intratumoral heterogeneity of HCC. We also observed that the tumor heterogeneity changed over time, which was correlated with sequentially obtained histopathological results. Therefore, DKI and SEM parameters may be used as valuable imaging biomarkers for evaluating the heterogeneity in HCC.

In this experiment, at each time point after 10 mg/kg sorafenib intervention, NF in the induction group was significantly larger than that in the control group, while MVD and Ki-67 index was significantly lower. These results indicate that 10 mg/kg sorafenib did have destructive effects on HCC. We further found that SD and kurtosis
Table 3 Comparisons of histopathological results of HCC between the sorafenib induction group and control group

<table>
<thead>
<tr>
<th>Group</th>
<th>NF (%)</th>
<th>MVD</th>
<th>Ki-67 index (%)</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>7 days</td>
<td>14 days</td>
<td>21 days</td>
</tr>
<tr>
<td>Induction group</td>
<td>20.549±2.149</td>
<td>25.244±2.338</td>
<td>31.748±3.374</td>
</tr>
<tr>
<td>Control group</td>
<td>11.643±1.201</td>
<td>12.991±1.397</td>
<td>19.272±1.616</td>
</tr>
<tr>
<td>Z</td>
<td>−2.611</td>
<td>−2.611</td>
<td>−2.611</td>
</tr>
<tr>
<td>P</td>
<td>0.009</td>
<td>0.009</td>
<td>0.009</td>
</tr>
</tbody>
</table>

HCC, hepatocellular carcinoma; NF, necrotic fraction; MVD, micro-vessel density

Table 4 Correlations between DKI and SEM parameters and histopathological results of HCC in the sorafenib induction group

<table>
<thead>
<tr>
<th>Parameters</th>
<th>NF</th>
<th>MVD</th>
<th>Ki-67 index</th>
<th>SD</th>
<th>Kurtosis</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>r</td>
<td>95% CI</td>
<td>P*</td>
<td>r</td>
<td>95% CI</td>
</tr>
<tr>
<td>MK</td>
<td>0.693</td>
<td>0.315–0.870</td>
<td>0.010</td>
<td>0.239</td>
<td>−0.369–0.761</td>
</tr>
<tr>
<td>MD</td>
<td>0.794</td>
<td>0.495–0.930</td>
<td>0.003</td>
<td>0.403</td>
<td>−0.183–0.772</td>
</tr>
<tr>
<td>α</td>
<td>−0.704</td>
<td>−0.868–(−0.386)</td>
<td>0.009</td>
<td>−0.391</td>
<td>−0.821–0.194</td>
</tr>
<tr>
<td>DDC</td>
<td>0.800</td>
<td>0.514–0.938</td>
<td>0.003</td>
<td>0.405</td>
<td>−0.151–0.798</td>
</tr>
</tbody>
</table>

DKI, diffusion kurtosis imaging; SEM, stretched exponential model; HCC, hepatocellular carcinoma; SD, standard deviation; NF, necrotic fraction; MVD, micro-vessel density; MK, mean kurtosis; MD, mean diffusivity; DDC, distributed diffusion coefficient; CI, confidence interval; P*, Benjamini-Hochberg P value.

Table 5 Correlations between DKI, SEM parameters and histopathological results of HCC in the control group

<table>
<thead>
<tr>
<th>Parameters</th>
<th>NF</th>
<th>MVD</th>
<th>Ki-67 index</th>
<th>SD</th>
<th>Kurtosis</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>r</td>
<td>95% CI</td>
<td>P*</td>
<td>r</td>
<td>95% CI</td>
</tr>
<tr>
<td>MK</td>
<td>0.340</td>
<td>−0.205–0.728</td>
<td>0.578</td>
<td>0.309</td>
<td>−0.290–0.842</td>
</tr>
<tr>
<td>MD</td>
<td>−0.129</td>
<td>−0.574–0.435</td>
<td>0.762</td>
<td>0.134</td>
<td>−0.477–0.701</td>
</tr>
<tr>
<td>α</td>
<td>−0.280</td>
<td>−0.793–0.255</td>
<td>0.578</td>
<td>−0.269</td>
<td>−0.819–0.402</td>
</tr>
<tr>
<td>DDC</td>
<td>0.002</td>
<td>−0.525–0.499</td>
<td>0.995</td>
<td>−0.439</td>
<td>−0.891–0.131</td>
</tr>
</tbody>
</table>

DKI, diffusion kurtosis imaging; SEM, stretched exponential model; HCC, hepatocellular carcinoma; SD, standard deviation; NF, necrotic fraction; MVD, micro-vessel density; MK, mean kurtosis; MD, mean diffusivity; DDC, distributed diffusion coefficient; CI, confidence interval; P*, Benjamini-Hochberg P value.
based on the histogram of HE staining in the induction group were significantly higher than those of the control group. According to previous reports (20,30,31), SD (or variance or coefficients of variation) and kurtosis derived from the histopathological image were positively associated with frequency and distribution of pixels, which was presumably related to tissue heterogeneity. Therefore, our histopathological results confirmed that a low dose of sorafenib could induce a more heterogeneous HCC compared with HCC without intervention.

Higher MK and lower α were shown to be associated with the complexity of various cancers, such as rectal and prostate cancers (7,32). Significantly increased MK and slightly decreased α were found within 24 hours after sorafenib treatment on a subcutaneous HCC model, which were caused by many intra- and intercellular factors including necrosis in the tumor (15). Jansen et al. also found that metastatic nodes in head and neck squamous cell carcinoma had higher heterogeneity when they were necrotic (33). In our study, increased MK and decreased α in the sorafenib induction group were correlated with SD and kurtosis, two histopathological markers reflecting the intratumoral heterogeneity of the tumor (20,30). Meanwhile, they were also correlated with NF. As a result, the non-uniformly distributed necrosis within the tumor was presumably one of the most important causes of increased tumor heterogeneity when a low dose of sorafenib was used. However, no significant difference was found in tumor size between the two groups at our study time points. This implies the significance of functional parameters MK and α which changed earlier than the morphological index of HCC. Similar results were also shown after irradiation of nasopharyngeal carcinoma in a xenograft model (34). However, there were conflicting results regarding the changing trends of MK and α in studies evaluating neoadjuvant chemoradiotherapy of rectal cancer (8,35). The discrepancy might be explained by complete tumor cell death, liquefaction, or massive necrosis, as well as therapy-induced fibrosis, which could reduce the tumor heterogeneity in these studies. To avoid complete or massive necrosis, which is likely when applied with a full dosage (4), we used a low dose of sorafenib to observe the tumor heterogeneity more closely at multiple time points.

In our study, MD and DDC exhibited significant increases after sorafenib treatment, which might be explained by decreased tumor cell density, increased intratumoral necrosis, and increased extracellular spaces. These values again showed positive correlations with histopathological SD, kurtosis, and NF, which indicate their potential to describe the complexity of tumor microstructure, particularly after treatment.

However, no correlations were found between DKI and SEM parameters, and MVD and Ki-67 metrics in the induction group. There were several possible explanations. First, DKI and SEM parameters might not have been sensitive to alterations in tumor cell proliferation and angiogenesis in our observation period. Second, tumor cell death and tissue necrosis might be the major contributing factors leading to the increased heterogeneity after sorafenib intervention (15). Third, the positioning of ROI which covered the necrotic areas might impair the demonstration of the real effects of cell proliferation and angiogenesis on these DKI and SEM parameters.

Differing from the sorafenib induction group, DKI and SEM parameters in the control group was not correlated with any histopathological markers. This might be due to the balance maintained between various tissue compositions in the natural course of HCC growth within our observational time points. Cell proliferation and tissue necrosis, which may have opposite effects on these DWI parameters, are two major counteracting factors leading
to intratumoral complexity (4), and we speculate that the balance between them is broken by sorafenib.

Our study has several limitations. First, the measurement on the slice showing the greatest dimension of the tumor may not have adequately represented the heterogeneity of the whole tumor (36). Second, although great efforts were made to match the location of the tumors between DWI and histopathological images, no exact co-registration method was available, which may have led to some errors. Furthermore, until now, there has been no well-established histopathological standard for evaluation of tumor heterogeneity. However, with digital whole slide scanning technology and quantitative image analysis such as histogram, and grayscale values from HE staining representing different tissue components, their distribution, and even spatial interrelationships may be analyzed more reliably and objectively with gray-level co-occurrence matrix methods, such as entropy, contrast, and correlation (22,23,37,38). Third, although hemorrhage was excluded from ROIs, micro-hemorrhage, which was invisible on MR images, may have influenced the accuracy of the DKI and SEM results. Fourth, the animal sample size was relatively small and might have led to a wider 95% CI for correlations between imaging parameters and histopathological results of HCC in the sorafenib induction group. Further studies with more animal models are warranted to verify our results before translation to clinical practice. Fifth, multiple time points might be a confounder for studying the correlations between the imaging and histopathological parameters. However, the two-way ANOVA test revealed no significant effects from these time points on DKI and SEM parameters. Last, this study focused on DKI and SEM parameters, which may only reflect some aspects of HCC heterogeneity. A recent report found that histogram analysis based on blood-oxygenation-level-dependent (BOLD) and dynamic contrast-enhanced (DCE)-MRI may provide a quantification of HCC heterogeneity as well (39). However, the latter techniques are time-consuming and more complicated.

In conclusion, DKI and SEM could reflect intratumoral heterogeneity of HCC and monitor its dynamic changes after application of low-dose sorafenib. Parameters of DKI and SEM may be potentially used for evaluating tumor heterogeneity of HCC.

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Footnote

Conflicts of Interest: The authors have no conflicts of interest to declare.

Ethical Statement: The investigation was approved by the Institutional Animal Care and Use Committee of Zhongshan Hospital, Fudan University.

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