A Comparative Study of Two-Compartment Exchange Models for Dynamic Contrast-Enhanced MRI in Characterizing Uterine Cervical Carcinoma

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A variety of tracer kinetic methods have been employed to assess tumor angiogenesis. The Standard two-Compartment model (SC) used in cervix carcinoma was less frequent, and Adiabatic Approximation to the Tissue Homogeneity (AATH) and Distributed Parameter (DP) model are lacking. This study compares two-compartment exchange models (2CXM) (AATH, SC, and DP) for determining dynamic contrast-enhanced magnetic resonance imaging (DCE-MRI) parameters in cervical cancer, with the aim of investigating the potential of various parameters derived from 2CXM for tumor diagnosis and exploring the possible relationship between these parameters in patients with cervix cancer. Parameters (tissue blood flow, \( F_p \); tissue blood volume, \( V_p \); interstitial volume, \( V_e \); and vascular permeability, \( PS \)) for regions of interest (ROI) of cervix lesions and normal cervix tissue were estimated by AATH, SC, and DP models in 36 patients with cervix cancer and 17 healthy subjects. All parameters showed significant differences between lesions and normal tissue with a \( P \) value less than 0.05, except for \( PS \) from the AATH model, \( F_p \) from the SC model, and \( V_p \) from the DP model. Parameter \( V_e \) from the AATH model had the largest AUC (\( r = 0.85 \)). Parameters \( F_p \) and \( V_p \) from SC and DP models and \( V_e \) and \( PS \) from AATH and DP models were highly correlated, respectively, (\( r > 0.8 \)) in cervix lesions. Cervix cancer was found to have a very unusual microcirculation pattern, with over-growth of cancer cells but without evident development of angiogenesis. \( V_e \) has the best performance in identifying cervix cancer. Most physiological parameters derived from AATH, SC, and DP models are linearly correlated in cervix cancer.

1. Introduction

Neovascularization plays a fundamental role in the growth of solid tumors [1]. Various experimental models as well as clinicopathological observations have shown that solid tumors (e.g., breast [2], lung [3], and cervical carcinoma [4, 5]) cannot attain diameters > 2-3 mm without their own vascular supply. Dynamic contrast-enhanced magnetic resonance imaging (DCE-MRI) is commonly used for the assessment of tumor angiogenesis. Analysis of DCE-MRI data can be performed using tracer kinetic models to derive quantitative parameters of tissue microcirculation. A variety of tracer...
kinetic methods have been employed to characterize various tumors and to assess the effects of antiangiogenic and antivascular drugs in clinical trials [6–15].

To date, the Generalized Kinetic (GK or Tofts) model and the Extended Generalized Kinetic (EGK or extended Tofts) model are frequently used for analysis of DCE-MRI data in oncology and drug trials. It is commonly assumed that \( K^{\text{trans}} \) yielded from the two models encompasses the effects of both blood flow and vessel permeability, such that \( K^{\text{trans}} \) reflects permeability when blood flow is much larger than permeability and \( K^{\text{trans}} \) reflects blood flow when permeability dominates [16]. However, novel vascular targeting drugs could reduce both permeability and blood flow and might exert different pharmacokinetic effects on blood flow and permeability [17, 18]. Thus, a method that can separately estimate tissue blood (plasma) flow (\( F_p \)) and vessel permeability (PS) is more valuable for the assessment of drug effects. Improvements in the temporal resolution of DCE-MRI have been reported. However, these models (AATH, SC, and DP) with respect to tumor microcirculation would be clinically desirable. Moreover, to facilitate the consistent interpretation of treatment effects for multicenter clinical trials employing different tracer kinetic models, it would be of interest to understand the relationships between parameters of these tracer kinetic models, so that if one kinetic model is used in a certain trial, one could get a sense of what would be the likely changes in another kinetic model. In this study, three two-compartment models (AATH, SC, and DP) were applied in cervix carcinoma to investigate the potential of various parameters in tumor diagnosis and to examine the possible relationship between parameters of these tracer kinetic models.

2. Materials and Methods

2.1. Patients. This retrospective study was approved by the institutional research ethics review board, and informed consent was obtained from all patients. Sixty-eight consecutive female patients (mean age, 50.4 years; age range, 41–75 years) clinically suspected of cervix cancer presented to our department in the period of April 2016 to July 2018. Patients were excluded for the following reasons: (1) poor image quality of DCE-MRI such as significant motion artifacts (\( n = 1 \)) or incomplete images (\( n = 2 \)); (2) patients with a history of targeted chemotherapy or radiation therapy before examination (\( n = 4 \)); (3) patients diagnosed of other cervical lesions, such as submucous myoma of uterus (\( n = 2 \)) and the endometrial carcinoma (\( n = 1 \)); and (4) no mass was identified for patients with stage Ia on DCE and other MRI sequences (\( n = 5 \)). In the end, 36 patients with cervix cancer and 17 healthy subjects were included in this retrospective study. Cervix cancer was clinically staged according to the International Federation of Gynecology and Obstetrics classifications [26]. These 36 patients were classified into stage Ib (\( n = 17 \)), Iia (\( n = 14 \)), and Iib (\( n = 5 \)). All patients were confirmed histopathologically. Hysterectomy was performed for cervix cancer patients and biopsy for healthy subjects. Patient characteristics are summarized in Table 1.

2.2. MR Imaging Protocol. All scans were performed on a 3.0 T scanner (Discovery™ MR750w, General Electric, USA) using an 8-channel torso phased-array coil. Routine clinical MRI scan sequences included a transverse fast spin-echo T1-weighted sequence (repetition time/echo time = 550–700/7–10 ms; field of view = 320 × 320 mm; matrix size = 512 × 512; slice thickness = 5.0 mm; intersection gap = 6 mm), a short time inversion recovery (STIR) T2-weighted sequence (repetition time/echo time = 3000–4000/70–80 ms; field of view = 256–320 × 256–320 mm; matrix size = 512 × 512; slice thickness = 5.0 mm; intersection gap = 6–7 mm), and diffusion-weighted imaging (DWI) (repetition time/echo time = 2400/60 ms, field of view = 320 × 320 mm, matrix size = 256 × 256 mm; slice thickness = 5.0 mm, intersection gap = 7 mm, b value = 0, 1000 s/mm²).

DCE-MRI was performed using a three-dimensional T1-weighted spoiled gradient echo sequence (LAVA, repetition time/echo time = 3/1 ms, flip angle = 4°, 8° and 11°, field of view = 360 × 360 mm, matrix size 256 × 256, slice thickness = 5 mm, 6 slices per slab). Ten precontrast scans of each flip angle (4°, 8°, and 11°) were acquired in the axial plane under quiet respiration. Dynamic postcontrast scans were acquired using the same sequence and a flip angle of 11°, with the intravenous injection of gadopentetated imeglimine (Magnevit; Bayer Healthcare Pharmaceuticals Inc., NJ) at an injection rate of 2 mL/second standard dose of 0.1 mmol/kg. A total of 180 consecutive scans were acquired for the dynamic series with a temporal resolution 2 s. Subsequently, a routine late contrast-enhanced T1-weighted scan (repetition time/echo time = 4/2 ms, flip angle = 13°, field of view = 280 × 280 mm, matrix size 512 × 512, slice thickness = 3 mm) was acquired in the sagittal plane.

2.3. Tracer Kinetic Models. Details of the four tracer kinetic models used in this study can be found in several review papers [21, 22]. Here, we would only list the essential operational equations for these models which specify the
To avoid possible effects of inflow for more details of the three tracer kinetic models.

cross-referencing to confirm the location and size of the T1-weighted, T2-weighted, and DW images were used for 10 years of experience in gynecological radiology. Routine Figure 1(a)) by two experienced radiologists with more than lineated on the central four slices (as illustrated in slices from the imaging volume (of 6 slices) were selected for and inhomogeneity near boundaries, only the central four roths when contrast-enhanced scans were evaluated. The size of ROI is no less than 10 voxels to ensure the robustness of measurement. The normal ROIs were selected in the normal cervical tissue away from the lesions. The areas of necrotic, cystic, and hemorrhages were avoided when drawing the lesion ROIs. Finally, 107 ROIs for cervix cancer were obtained from 36 patients. 103 ROIs for normal tissues were obtained from 17 healthy subjects and 24 patients with smaller masses which can be delineated accurately. AIF was sampled from a voxel that clearly resided within the iliac artery on one of the four central slices as shown in Figure 1(b). Desirable features for AIF selection included an early bolus arrival time and high peak value and signal-to-noise ratio. Fitting of voxel-level tissue concentration-time curves $C_{tiss}(t)$ and generation of parametric maps were performed using a commercially available software (Miitalsytics, Fitpu Healthcare, Singapore). The software allows for the selection of an individual AIF for each patient case and employs a constrained nonlinear optimization algorithm in fitting the various models.

2.4. Image Postprocessing. To avoid possible effects of inflow and inhomogeneity near boundaries, only the central four slices from the imaging volume (of 6 slices) were selected for processing. For each patient, regions of interest (ROIs) for tumor lesions and normal cervix tissue were manually delineated on the central four slices (as illustrated in Figure 1(a)) by two experienced radiologists with more than 10 years of experience in gynecological radiology. Routine T1-weighted, T2-weighted, and DW images were used for cross-referencing to confirm the location and size of the lesions.

2.5. Statistical Analysis. For each patient, the median parameter value of all voxels within the tumor ROIs on multiple slices is taken as a representative statistic of the parameter in the tumor. The median values of the fitted

| Table 1: Patient characteristics. |
|----------------------------------|

| Parameters                  | No. of patients |
|-----------------------------|-----------------|
| Age; mean (range) years     | 42 (42–75)      |
| Histological subtype        |                 |
| Adenocarcinoma (AC)         | 2               |
| Squamous cell carcinoma (SCC)| 34              |
| Tumor grade                 |                 |
| Well                        | 6               |
| Moderate                    | 30              |
| FIGO* stage                 |                 |
| Ib                          | 17              |
| IIa                         | 14              |
| IIb                         | 5               |

Abbreviation: FIGO International Federation of Gynecology and Obstetrics. *According to FIGO 2009 staging criteria.

Dependence of tissue tracer concentration $C_{tiss}(t)$ (as a function of time $t$) on the arterial input function (AIF) and relevant physiological parameters:

Standard Compartment (SC) model

$$C_{tiss}(t) = AIF \otimes F_p \{ A \exp(\alpha t) + (1 - A) \exp(\beta t) \}, \quad (1a)$$

where

$$\begin{align*}
(\alpha & \beta) = \\
\frac{1}{2} \left[ -\left( \frac{PS}{V_p + \frac{PS}{F_p}} \right) \right. \quad (1b) \\
& \pm \sqrt{\left( \frac{PS}{V_p + \frac{PS}{F_p}} \right)^2 - 4 \left( \frac{PS}{V_p} \right)}
\end{align*}$$

and $\otimes$ denotes the convolution operator.

Distributed Parameter (DP) model

$$A = \frac{\alpha + (PS/V_p) + (PS/V_e)}{\alpha - \beta}, \quad (1c)$$

where $u(t)$ denotes the Heaviside unit-step function and $I_1$ is the modified Bessel function.

Adiabatic Approximation to the Tissue Homogeneity (AATH) model

$$C_{tiss}(t) = AIF \otimes F_p \left\{ 1 - \exp \left( \frac{PS}{V_e} \right) \right\} \left[ 1 - \exp \left( \frac{PS}{F_p} \right) \right] \left[ 1 - \exp \left( -\frac{PS}{F_p} \right) \right] \left( t - \frac{V_p}{F_p} \right) \left\{ 1 + \int_0^{t-(V_p/F_p)} \exp \left( -\frac{PS}{V_e} \right) \left( \frac{FS}{V_e} \right)^{1/2} I_1 \left( \frac{PS}{V_e} \right) \right\}, \quad (2)$$

Interested readers can refer to recent review papers [21, 22] for more details of the three tracer kinetic models.

$$ \begin{align*}
C_{tiss}(t) &= AIF \otimes F_p \{ A \exp(\alpha t) + (1 - A) \exp(\beta t) \}, \\
\text{where} & \quad (1a) \\
(\alpha & \beta) = \\
\frac{1}{2} \left[ -\left( \frac{PS}{V_p + \frac{PS}{F_p}} \right) \right. \quad (1b) \\
& \pm \sqrt{\left( \frac{PS}{V_p + \frac{PS}{F_p}} \right)^2 - 4 \left( \frac{PS}{V_p} \right)} \\
A &= \frac{\alpha + (PS/V_p) + (PS/V_e)}{\alpha - \beta}, \quad (1c) \\
\text{and} & \quad \otimes \text{ denotes the convolution operator.} \\
\text{Distributed Parameter (DP) model} \\
C_{tiss}(t) &= AIF \otimes F_p \left\{ 1 - \exp \left( \frac{PS}{V_e} \right) \right\} \left[ 1 - \exp \left( \frac{PS}{F_p} \right) \right] \left[ 1 - \exp \left( -\frac{PS}{F_p} \right) \right] \left( t - \frac{V_p}{F_p} \right) \left\{ 1 + \int_0^{t-(V_p/F_p)} \exp \left( -\frac{PS}{V_e} \right) \left( \frac{FS}{V_e} \right)^{1/2} I_1 \left( \frac{PS}{V_e} \right) \right\}, \quad (2) \end{align*}$$
Figure 1: Continued.
Figure 1: Example of a patient with stage IIb cervix cancer. (a) ROIs for cervix carcinoma (blue) and normal cervix tissue (red) are shown for the central four slices of the DCE-MRI dataset, and the location within the iliac artery where the AIF was sampled was marked with a red dot. (b) Sampled AIF used in model fitting. (c) Parameter maps generated using the three models (AATH, SC, and DP) for tumor and the normal tissue ROIs. (d) Examples of curve fittings for a tumor voxel. In the legend, the four numbers within square brackets beside each model are their respective parameter values: \( F_p \) (mL/min/100 mL), \( V_p \) (mL/100 mL), PS (mL/min/100 mL), \( V_e \) (mL/100 mL)).
parameters were used because the median is more robust to outliers (that could occur during data fitting) than the mean. A two-way model average measure, intraclass correlation coefficient (ICC), was used to test the interobserver consistency. Then, all the measurements from the two observers were averaged for further comparison. Agreement was interpreted according to the ICC as follows: >0.80, excellent; 0.61–0.80, good; 0.41–0.60, moderate; 0.21–0.40, fair; and <0.2, poor agreement [27].

The normality of the distribution of all parameters was analysed by the Kolmogorov–Smirnov test. The receiver operating characteristic (ROC) curves of all parameters were obtained and the areas under the curves (AUC) were evaluated to determine the discriminating power of DCE parameters between the lesion and normal tissues. Interpretation of AUC values is application dependent, and in general, it is appropriate that values ≥0.9 would be “excellent,” ≥0.8 “good,” ≥0.7 “fair,” and <0.7 “poor” [28].

The Pearson correlation coefficient r was used to explore possible relationship between median parameter values of the three models. A strong correlation was assumed for 0.8 < r ≤ 1, a moderate correlation for 0.5 < r ≤ 0.8, a weak correlation for 0.3 < r ≤ 0.5, and no correlation for r ≤ 0.3 [29]. The Bland–Altman plot was used to show agreements between parameter values of the three models. All statistical analysis were performed using SPSS software 18.0 (Chicago, IL, USA), and P < 0.05 was considered statistically significant.

3. Results

For the case shown in Figure 1(a), the parameter maps of one slice generated using the three methods (AATH, SC, and DP) are shown in Figure 1(c). It is evident that three methods attain similar results in V_e and PS, and both estimates are smaller in tumor ROI than in normal tissue ROI. For F_p, the estimate by AATH is smaller in tumor ROI and the estimates by SC and DP are apparently close between tumor and normal tissue ROI. For V_p, the estimates vary from close between tumor and normal tissue ROI (AATH) to smaller in tumor ROI (SC and DP).

Table 2 shows the ICC values for the measured parameters, where most ICC values are greater than 0.9, indicating very good agreement between measurements from two observers. Thus, the parameter values as measured by two observers are averaged and utilized in the analysis as follows.

Table 3 shows the nonparametric statistical analysis of parameters between cervix cancer and normal tissue ROIs. The parameters F_p, V_e and PS of the three DCE models (AATH, SC, and DP models) were smaller in cervix cancer lesions than in normal cervix tissue, and the differences between lesions and normal tissue were mostly significant except for PS from the AATH model and F_p from the SC model. V_p by AATH and SC was significantly larger in cervix cancer lesions than in normal cervix tissue, and DP yielded contradictory results, though the difference between lesions and normal tissue was not statistically significant. Parameter V_e attained fair to good AUC values (>0.83 for AATH and DP models and >0.75 for the SC model). Other parameters, e.g., V_p from the AATH model and PS from the SC model also showed fair AUC values.

Results of the Pearson correlation between the same parameters estimated by different models are shown in Table 4. Comparing parameter estimates of the three models (AATH, SC, and DP models) for cervix cancer lesion, it was found that F_p and V_e from SC and DP models and V_e and PS from AATH and DP models were highly correlated, respectively (r > 0.8). Moderate correlations were observed between V_p from AATH and V_e from SC and DP models, V_e and PS from AATH and SC models, and V_e and PS from SC and DP models (0.5 < r < 0.8). No correlation was observed between F_p from AATH and F_p from SC and DP models, respectively (r < 0.3). Comparing parameter estimates of the three models (AATH, SC, and DP models) for normal cervix tissues, it was found that good agreement existed between the estimates of F_p, V_e from the three models, and PS from AATH and DP models (r > 0.8). Moderate correlations were observed between PS from AATH and SC models and V_p from SC and DP models (0.5 < r < 0.8). A weak correlation was observed between PS from SC and DP models (0.3 < r < 0.5). No correlation was observed between V_p from AATH and V_p from SC and DP models, respectively (r < 0.3). Results of the Bland–Altman test for comparison of the parameter with the same biophysical meaning obtained using the three models are shown in Figure 2. Bland–Altman plots demonstrated good agreements between the estimates of V_e from the three models for both normal cervix tissue and cervix lesion because their y-coordinate values were centered at zero (Figures 2(a) and 2(b)). Good agreements were observed between V_e from SC and DP models and PS from AATH and SC models in normal cervix tissue (Figure 2(a)). The y-coordinate value of other parameters estimated by AATH, SC, and DP modes stayed away from zero, indicating less agreement between them in normal cervix tissue and cervix lesion.

4. Discussion

This preliminary study evaluated the performance of physiological parameters derived from AATH, SC, and DP models with respect to tumor microcirculation in cervix cancer. The angiogenic activity of cervix cancer assessed by various DCE-MRI models improved in vivo understanding of the fundamental processes involved in tumor angiogenesis. Parameter V_e has the best performance in identifying cervix cancer among all parameters. Most physiological parameters derived from the three DCE-MRI models are linearly correlated in cervix cancer.

The parameter V_e of all three DCE models showed that the V_e value of cervix cancer tissue was significantly smaller than that of normal cervix tissue. The corresponding AUC values were greater than 0.75, and V_e of the AATH model had the largest AUC (0.85) among all parameters, indicating a fairly good performance in terms of diagnostic value. In tracer kinetic modeling, V_e stands for the fractional volume of extravascular extracellular space, which is closely pertaining to the degree of cell growth. The more the cells grow,
| Lesion | AATH | SC | DP |
|--------|------|----|----|
| ICC    | 0.922| 0.975| 0.959| 0.939| 0.935| 0.980| 0.968| 0.857| 0.965| 0.989| 0.958| 0.958|
| 95% CI | 0.886–0.947| 0.963–0.983| 0.940–0.972| 0.911–0.958| 0.905–0.956| 0.971–0.987| 0.953–0.978| 0.790–0.903| 0.949–0.976| 0.984–0.993| 0.938–0.971| 0.938–0.971|

| Normal | ICC | 0.934| 0.661| 0.908| 0.948| 0.978| 0.945| 0.910| 0.887| 0.906| 0.812| 0.943| 0.892|
| 95% CI | 0.902–0.955| 0.499–0.770| 0.864–0.938| 0.924–0.965| 0.967–0.985| 0.919–0.963| 0.867–0.939| 0.833–0.923| 0.861–0.936| 0.702–0.882| 0.916–0.961| 0.841–0.927|

ICC, intraclass correlation coefficient. CI, confidence interval of difference.
cellular space. The first two processes are of interest in DCE transportation process for the tracer molecule: the flowing process within the vascular space, the exchange process between the vascular and extravascular extracellular space, and the diffusion process within the extravascular extracellular space. The first two processes are of interest in DCE modeling, and the associated parameters are $F_p$ for the flowing process and $PS$ for the exchange process, both of which are linked to nutrition supply to cell growth. In most solid tumors, both parameters usually increase due to the over-growth of tumor cells. From Table 2, it is observed that both parameters as estimated by the three models are linked to nutrition supply to cell growth. In other words, the smaller the value of $V_p$ is, the less the $V_e$ value is. Tumor is typically characterized by over-growth of cells. Thus, the observed smaller value of $V_e$ in cervix cancer tissue suggests the over-growth of cells in cervix cancer tissue in comparison with normal cervix tissue, which is consistent with the feature of other cancer tissue.

In the tissue region of interest, there are three types of transportation process for the tracer molecule: the flowing process within the vascular space, the exchange process between the vascular and extravascular extracellular space, and the diffusion process within the extravascular extracellular space. The first two processes are of interest in DCE modeling, and the associated parameters are $F_p$ for the flowing process and $PS$ for the exchange process, both of which are linked to nutrition supply to cell growth. In most solid tumors, both parameters usually increase due to the over-growth of tumor cells. From Table 2, it is observed that both parameters as estimated by the three models are linked to nutrition supply to cell growth. In other words, the smaller the value of $V_p$ is, the less the $V_e$ value is. Tumor is typically characterized by over-growth of cells. Thus, the observed smaller value of $V_e$ in cervix cancer tissue suggests the over-growth of cells in cervix cancer tissue in comparison with normal cervix tissue, which is consistent with the feature of other cancer tissue.

A table comparing model parameters between cervix carcinoma and normal cervix tissue is provided. The table includes columns for $F_p$, $V_p$, $V_e$, and $PS$, with values given for AATH-normal cervix, AATH-cervix cancer, SC-normal cervix, SC-cervix cancer, DP-normal cervix, and DP-cervix cancer for AATH, CC, and DP models. The table also includes $P$ values for statistical comparison using the Mann-Whitney U test, with $a$ indicating $P < 0.05$.

Table 4: Results of the Pearson correlation between parameters of the three models (AATH, SC, and DP) in cervix carcinoma and normal cervix tissue. $r > 0.5$ and $P < 0.05$ are indicated by $^*$. 

| Lesion       | $F_p$ (mL/min/100 mL) | $V_p$ (mL/100 mL) | $V_e$ (mL/100 mL) | $PS$ (mL/min/100 mL) |
|--------------|----------------------|------------------|------------------|---------------------|
| AATH-CC      | 0.121 ($P = 0.214$)  | 0.710 ($P < 0.001$) | 0.572 ($P < 0.001$) | 0.667 ($P < 0.001$) |
| AATH-DP      | 0.095 ($P = 0.329$)  | 0.732 ($P < 0.001$) | 0.899 ($P < 0.001$) | 0.921 ($P < 0.001$) |
| CC-DP        | 0.813 ($P < 0.001$)  | 0.850 ($P < 0.001$) | 0.565 ($P < 0.001$) | 0.749 ($P < 0.001$) |
| Normal       | 0.829 ($P < 0.001$)  | 0.261 ($P = 0.008$) | 0.824 ($P < 0.001$) | 0.554 ($P < 0.001$) |
| AATH-CC      | 0.815 ($P < 0.001$)  | 0.247 ($P = 0.012$) | 0.912 ($P < 0.001$) | 0.890 ($P < 0.001$) |
| AATH-DP      | 0.981 ($P < 0.001$)  | 0.696 ($P < 0.001$) | 0.886 ($P < 0.001$) | 0.385 ($P < 0.001$) |

The unusual observation in the microcirculation pattern of cervix cancer tissue is also evident in Figure 3, which exemplifies the change of tissue intensity after venous injection of contrast media. The baseline corresponds to the period the contrast media has not flowed into the tissue of interest. Here, the intensity of cervix cancer tissue is slightly brighter than that of normal cervix tissue, indicating that cervix cancer tissue has shorter spin-lattice relaxation time. With the arrival of contrast media, tissue intensity rapidly increases till a "peak" value, followed by a part with intensity slowly decreasing, which stands for the wash-in phase and wash-out phase of contrast media, respectively. The speed of uptake in the wash-in phase depends on the value of blood flow, and the larger the value, the sharper the slope of the uptake curve. The sharper slope of normal cervix tissue suggests a lower blood flow for the cervix cancer tissue, which is well in line with the observation in DCE modeling. The "peak" value is primarily related to the degree of microvasculature. If the tissue develops more microvessels, there would accumulate more tracer molecules, hence to observe brighter intensity value. The lower "peak" value of cervix cancer tissue suggests that cervix cancer tissue might develop fewer vessels. This is consistent with the finding by the DP model, but not consistent with either that by AATH and CC models or previous pathohistological experiment. This apparent contradiction might be due to the dysfunction...
Figure 2: Continued.
of vessels newly developed in cancer tissue [33], which may accumulate less tracer molecules, resulting in the lower “peak” value of cervix cancer.

The wash-out phase involves two processes: one process where the contrast media flows out of the tissue of interest and another process where the contrast media leaks out of the tissue of interest. The flow-out process would lead to lower tissue intensity, and the exchange process would determine the speed of intensity decreasing. The higher the blood flow is, the faster the intensity decreases. The more leakier the vessel wall is, the slower the intensity decreases. The trend of wash-out phase would be determined by both factors. If the blood flow is large and the effect of flow-out process is dominant, a downward trend would be observed. If the blood flow is small and the vessel wall is leaky, a slowly upward or flat trend could be observed. As there is an evident downward trend in the wash-out phase of normal cervix tissue, it demonstrates the higher blood flow rate in normal cervix tissue. Consequently, the intensity difference between cervix cancer tissue and normal cervix tissue is decreasing in the early stage of wash-out phase. In spite of this, it is clear that the intensity difference becomes stable towards the tail stage of the observed wash-out phase, which
is more influenced by the exchange process. As aforementioned, the measured permeability \( PS \) in DCE modeling is larger in normal cervix tissue; thus, the downward trend in normal cervix tissue encounters larger resistance, which helps precisely to explain the observed phenomena in the wash-out phase.

From the aforementioned analysis, cervix cancer exhibits very unusual pattern in microcirculation, with over-growth of cells, decreased blood flow, decreased vessel wall permeability, and without well development of increased and functional microvascularity. One immediately notices what is the growing mechanism for tumor cells without additional nutrition supply through the blood transportation system. It is known that microvascular proliferation is not the only factor of cancer cell growth and some other factors like interstitial fluid pressure, oxygen, and human papilloma virus (HPV) [9,34,35] but may also promote tumor growth. But, which of these pathophysiological mechanisms is the dominating factor for the growth of cervix cancer cells is not clear so far. It would require a histopathologic and etiologic study to uncover the difference of growth condition between cervix carcinoma and other tumors. For clinicians to provide individual cancer patients with optimal treatment and precise prognosis, further knowledge on the tumor microenvironment would be of great value.

In the situation where two different kinetic models were employed in a multicenter clinical trial for the purposes of assessment and monitoring of treatment effect using DCE-MRI, it would be desirable that the kinetic parameters estimated by the two models can be related, i.e., that they were at least linear correlated. Parameter \( V_e \) derived from all three models, \( V_p \) estimated by SC and DP models, and PS from AATH and SC models in cervix lesion were highly correlated as well as in good agreement, indicating a fairly good performance in terms of evaluation of treatment effect in multicenters. In most situations, parameters derived from different models were not quantitatively equivalent but highly correlated. For example, if the SC and DP models were employed in different centers and one center observed a decrease in perfusion (tissue blood flow) of cervix cancer estimated by SC-\( F_p \) due to a particular treatment, then another center employing the DP model should also observe a similar percentage decrease in DP-\( F_p \). Because SC-\( F_p \) and DP-\( F_p \) were highly correlated but not quantitatively equivalent, we could only compare their relative difference (percentage change) in \( F_p \) before and after treatment and not their absolute values. In a DCE CT study of brain tumors [36], parameters \( F_p, V_p, V_e, \) and PS from the SC model were found to be highly correlated with those from the DP model. These correlation results for these parameters in meningiomas derived from DCE CT are very similar to the present results derived from DCE-MRI for cervix cancer (\( r > 0.5 \)). However, comparing parameter estimates of the three models (AATH, SC, and DP models) for cervix in this study, it was found that the Pearson correlation between parameters of the three models in cervix cancer lesion and normal cervix tissue yield inconsistent conclusion, i.e., \( F_p \) from AATH model and \( F_p \) from SC and DP models were highly correlated, respectively, in normal cervix tissue, but very weak correlation in cervix cancer lesions. \( V_p \) from the AATH model and \( V_p \) from SC and DP models were highly correlated, respectively, in cervix cancer lesions, but very weak correlation in normal cervix tissue. (Table 4). Thus, the parameter correlations between kinetic models of tumors.
may be different from normal tissue, and it should not be
generalized without validation for different tumors char-
acterized with different perfusion. Parameter relationships
under various tracer kinetic models in other tumors need
further investigation.

This study has a few limitations. There was a lack of
pathohistological experiment, and the previous pathohisto-
logical results by other researchers were employed for com-
parison in this study. The present DCE-MRI data were acquired
using a protocol with high temporal resolution (2 s) in order to
capture rapid changes in arterial and tumor concentration with
time, which led to a tradeoff in organ coverage and image spatial
resolution. As a result, effects of lesion heterogeneity might not
be fully accounted for. Possible error sources that could affect
contrast concentration estimation in this study include the lack
of correction for effects of (i) B1 field inhomogeneity and (ii)
T2* relaxation. (i) Concentration estimation based on variable
flip angle T1 mapping would be sensitive to spatial non-
uniformity in the radio-frequency transmit field (B1) which
causes variations in the prescribed flip angles. (ii) At high-
contrast concentrations, effects of T2* relaxation become in-
creasingly substantial in higher field strengths and could not be
ignored during concentration estimation. Previous studies have
demonstrated that failure to account for T2* effects could result in
underestimation of the AIF, especially for concentrations around the AIF peak [37, 38]. Both effects (i) and (ii) can result in
effects of lesion heterogeneity, which further translate to errors in the estimation of kinetic
parameters [37, 38]. A major limitation of this study is therefore
the lack of quantification of such errors and their propagation to
the estimated kinetic parameters, which would be the subject of
further study in our future work.

In conclusion, two-compartment exchange models
turned out to be very promising tools in analyzing DCE data
and assessing the microcirculation pattern in cervix cancer
tissue, hence with great value to assist cervix cancer di-
agnosis and prognosis. Parameter Vn has the best perform-
ance in identifying cervix cancer tissue among all parameters.

Data Availability

The data of this study are already presented in this paper.

Ethical Approval

All procedures performed involving human participants
were in accordance with the ethical standards of the Second
Affiliated Hospital and Yuying Children’s Hospital of
Wenzhou Medical University.

Consent

All patients signed informed consent forms, allowing their
data to be used in this study.

Conflicts of Interest

The authors declare that there are no conflicts of interest
regarding the publication of this paper.

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