

Tumor Habitat Analysis Using Longitudinal Physiological MRI to Predict Tumor Recurrence After Stereotactic Radiosurgery for Brain Metastasis

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Objective: It is difficult to predict the treatment response of tissue after stereotactic radiosurgery (SRS) because radiation necrosis (RN) and tumor recurrence can coexist. Our study aimed to predict tumor recurrence, including the recurrence site, after SRS of brain metastasis by performing a longitudinal tumor habitat analysis.

Materials and Methods: Two consecutive multiparametric MRI examinations were performed for 83 adults (mean age, 59.0 years; range, 27–82 years; 44 male and 39 female) with 103 SRS-treated brain metastases. Tumor habitats based on contrastenhanced T1- and T2-weighted images (structural habitats) and those based on the apparent diffusion coefficient (ADC) and cerebral blood volume (CBV) images (physiological habitats) were defined using k-means voxel-wise clustering. The reference standard was based on the pathology or Response Assessment in Neuro-Oncologycriteria for brain metastases (RANO-BM). The association between parameters of single-time or longitudinal tumor habitat and the time to recurrence and the site of recurrence were evaluated using the Cox proportional hazards regression analysis and Dice similarity coefficient, respectively.

Results: The mean interval between the two MRI examinations was 99 days. The longitudinal analysis showed that an increase in the hypovascular cellular habitat (low ADC and low CBV) was associated with the risk of recurrence (hazard ratio [HR], 2.68; 95% confidence interval [CI], 1.46–4.91; P = 0.001). During the single-time analysis, a solid low-enhancing habitat (low T2 and low contrast-enhanced T1 signal) was associated with the risk of recurrence (HR, 1.54; 95% CI, 1.01–2.35; P = 0.045). A hypovascular cellular habitat was indicative of the future recurrence site (Dice similarity coefficient = 0.423).

Conclusion: After SRS of brain metastases, an increased hypovascular cellular habitat observed using a longitudinal MRI analysis was associated with the risk of recurrence (i.e., treatment resistance) and was indicative of recurrence site. A tumor habitat analysis may help guide future treatments for patients with brain metastases.

Keywords: Radiosurgery; Brain metastasis; Response assessment; Tumor habitat; Radiation necrosis

INTRODUCTION

Stereotactic radiosurgery (SRS) is becoming a popular therapeutic option for brain metastases [1,2]; however,

distinguishing radiation necrosis (RN) from tumor recurrence remains challenging. RN and tumor recurrence have common imaging findings, such as contrast enhancement, peritumoral edema, and a mass effect

Received: July 21, 2022 Revised: November 8, 2022 Accepted: December 11, 2022

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[3]. There is no definitive imaging technique that can accurately differentiate between these two entities [4,5]. Because both RN and tumor recurrence may coexist in SRS-treated tissue comprising a viable tumor, capillary damage, ischemia, and chemotherapy damage [4,6-8], a combination of diffusion- and perfusion-weighted MRI can provide useful information. Diffusion- and perfusion-weighted MRI have been confirmed as effective tools that can increase the diagnostic confidence [4,6,9,10], depict angiogenesis and necrotic tissue, and improve the diagnostic accuracy better than conventional MRI alone [11-13].

A tumor habitat analysis can distinguish subregions within a heterogeneous tumor by identifying similar voxels with common tumor biology [14,15]. A recent habitat analysis using structural MRI [16] showed that a habitat defined by low T2 and low T1 contrast-enhancing signal intensity was associated with viable tumors after SRS for brain metastases. Such a guantitative tumor habitat analysis may be helpful for assessing the residual tumor burden of metastatic tumors after treatment and determining the target for additional SRS. Nonetheless, this study was limited to cross-sectional anatomical MRI data. A previous radiological-pathological correlation analysis of 11 brain metastases treated with SRS revealed that a single MRI examination could not sufficiently determine pathological failure, but that serial MRI was practical for predicting the treatment response [17]. A longitudinal study using diffusion-weighted and dynamic susceptibility contrast (DSC) perfusion-weighted imaging, which reflect tumor cellularity and vascularity, may reveal effective tools that can predict treatment resistance and local recurrence sites.

We hypothesized that an evaluation of brain metastases after SRS using serial diffusion- and perfusion-weighted physiological MRI would be useful for predicting the treatment response by identifying subregions of recurrent tumors and treatment-induced changes, and that a spatial tumor habitat analysis would enable the identification of the recurrence site by clustering similar apparent diffusion coefficient (ADC) and cerebral blood volume (CBV) values. This study aimed to predict tumor recurrence (i.e., the treatment response) after SRS of brain metastases by performing a longitudinal tumor habitat analysis using MRI.

MATERIALS AND METHODS

Patient Population

After receiving approval from the Institutional Review Board

of Asan Medical Center (IRB No. 2021-4701), this singlecenter, retrospective, clinical study was conducted according to the United States Health Insurance Portability and Accountability Act (HIPAA) regulations and the Declaration of Helsinki. The need for written informed consent was waived. Additionally, it was conducted in accordance with the Strengthening the Reporting of Observational Studies in Epidemiology (STROBE) statement [18].

Using our center's Redcap database of brain metastases. we identified 301 patients treated during January 1, 2014 to November 30, 2020; of those 301 patients, 212 who fulfilled the following criteria were first identified and evaluated: age older than 18 years; initially diagnosed with brain metastasis and underwent subsequent SRS; brain metastasis with a size of at least $1 \times 1 \text{ cm}^2$; and a contrast-enhancing lesion (CEL) in the brain metastasis that could be analyzed. Their eligibility for study inclusion was further evaluated using the following criteria: the tumor was enlarged by more than 20% or at least 5 mm after SRS, resulting in clinical suspicion of tumor recurrence according to the response assessment in neuro-oncology brain metastases (RANO-BM) criteria; underwent follow-up MRI including diffusion-weighted imaging (DWI) and DSC perfusion-weighted imaging at least twice; the first followup MRI was performed within 6 months after SRS; and adequate follow-up examination results were available for the assessment of the treatment response. We excluded 129 patients for the following reasons: 39 patients had brain metastases that did not show a sufficient increase in size; 43 patients underwent the first follow-up MRI more than 6 months after SRS; 36 patients did not undergo follow-up MRI including DWI and DSC perfusion-weighted imaging at least twice; and 11 patients did not have adequate followup examination results to allow an assessment of the treatment response. Finally, 83 patients (mean age, 59.0 years; range, 27-82 years; 44 male and 39 female) with a total of 103 brain metastases were included. A flowchart of the patient inclusion process is shown in Figure 1. Clinical data were collected from the database of the Asan Medical Center. When multiple SRS-treated metastases were present, up to three of the largest lesions of each patient were included.

Definition of Progression: Reference Standard

At our institution, patients with brain metastases underwent follow-up assessments involving brain MRI every 2 or 3 months after the initial SRS for 1 or 2 years





Fig. 1. Flowchart of the patient inclusion process. DSC = dynamic susceptibility contrast, DWI = diffusion-weighted imaging, SRS = stereotactic radiosurgery

during regular outpatient visits under the supervision of a multidisciplinary team [1,19]. If a patient developed a new symptom or if the neurological symptoms deteriorated, then MRI was performed regardless of the scheduled followup period. When clinically indicated, surgery was performed to confirm the final diagnosis of a viable tumor or RN. If surgery was not possible, then the viable tumor was determined using MRI in accordance with the RANO-BM criteria and serial follow-up examinations with intervals of at least 3 months were performed. Supplemental imaging examinations, such as 3,4-dihydroxy-6-[18F]fluoro-Lphenylalanine) (¹⁸F-DOPA) PET/CT, ¹¹C-methionine PET/ CT, and ¹⁸F-fluorodeoxyglucose (¹⁸F-FDG) PET/CT, were also performed for individuals with equivocal MRI results. The clinicoradiological diagnosis was determined by a consensus reached during a multidisciplinary meeting involving three neurosurgeons (all with at least 15 years of experience with neuro-oncology) and two neuroradiologists (with 21 and 9 years of experience with neuro-oncologic imaging, respectively) who reviewed all imaging and medical records. When a CEL exhibited a steady increase in size during two or more successive follow-up MRI examinations within a 2- to 3-month interval and necessitated a change in treatment, the patient was classified as having tumor recurrence. The date of recurrence was recorded and the MRI examination was considered the confirmatory scan. In contrast, when a CEL subsequently regressed or became

stable without a change in treatment within 6 months of the index imaging, the patient was categorized as having RN. The time to recurrence was defined as the time from SRS to the first local recurrence event.

MRI Acquisition

A 3T scanner was used for all MRI examinations (Ingenia 3.0 CX; Philips Healthcare). The following structural and physiological MRI sequences were performed: T2-weighted imaging (T2WI); fluid-attenuated inversion recovery (FLAIR) imaging; T1-weighted imaging (T1WI); DWI; DSC perfusion-weighted imaging; and three-dimensional (3D) contrast-enhanced (CE) T1WI.

The DWI parameters included the following: repetition time (TR)/echo time (TE), 3000/56 ms; diffusion gradient, $b = 0 \text{ s/mm}^2$ and $b = 1000 \text{ s/mm}^2$; field of view (FOV), 250 x 250 mm²; matrix, 256 x 256; and slice thickness/ gap, 5 mm/2 mm. The ADC values were calculated using DWI images with $b = 1000 \text{ s/mm}^2$ and $b = 0 \text{ s/mm}^2$. DSC perfusion-weighted imaging was performed using a gradient-echo echo-planar imaging protocol. A preload of 0.01 mmol/kg gadoterate meglumine was administered, followed by a dynamic bolus of a standard dose of 0.1 mmol/kg gadoterate meglumine (Dotarem; Guerbet) delivered at a rate of 4 mL/s using an MRI-compatible power injector (Spectris; Medrad). The contrast bolus was followed by 20 mL of saline at the same injection rate. The



imaging parameters were as follows: TR/TE, 1808/40 ms; flip angle, 35°; FOV, 240 x 240 mm²; slice thickness/gap, 5 mm/2 mm; matrix, 128 x 128; and total acquisition time, 1 minute and 54 seconds. Dynamic acquisition was performed at a temporal resolution of 1.5 seconds, and 60 dynamics were acquired.

Deep Learning-Based Segmentation of Contrast-Enhancing Lesions and Image Processing

Brain extraction was performed for 3D CE-T1WI and FLAIR imaging using an algorithm (https://github.com/MIC-DKFZ/ HD-BET). Lesion segmentation was performed based on the images obtained using 3D CE-T1WI and FLAIR imaging and the 3D nnUNet-based algorithm (https://github.com/MIC-DKFZ/nnUNet) [20] of the PyTorch package version 1.1 of Python 3.7 (https://www.python.org). Because the CEL, necrosis, and peritumoral edema were segmented, only the CEL was included in the subsequent analysis. Hemorrhagic lesions were automatically excluded from the analysis if they showed similar hyperintensities on T1WI and CE-T1WI. The process was validated by an experienced neuroradiologist.

During T2WI and CE-T1WI, the signal intensity was normalized using kernel density estimation-based white matter segmentation [21] in R (version 4.1.1; Institute for Statistics and Mathematics; https://www.r-project.org/). To perform the DSC analysis, a pharmacokinetic map was computed using Nordic ICE (NordicNeuroLab). The quantity of blood (mL per 100 mL of tissue) was determined using the integrated DSC module, which combines a leakage correction algorithm of the relative CBV (rCBV) and manual noise thresholding. The Weisskoff–Boxerman method, which is based on the time-dependent deviation of the pixel-wise concentration-time curve from a reference curve, indicated that the leakage did not affect the estimations [22]. Normalized CBV (nCBV) maps were created by normalizing the rCBV maps in accordance with normal white matter.

To assess the changes in the follow-up examination results, the 3D CE-T1WI images of each patient in the dataset were co-registered and resampled into isometric voxels. Next, using rigid transformations with six degrees of freedom in the SPM package (version 12; https://www.fil. ion.ucl.ac.uk/spm/), the T2WI, nCBV, and ADC images were co-registered and resampled as isovoxel CE-T1WI images.

Tumor Habitat Analysis

The k-means clustering algorithm in scikit-learn (https://github.com/scikit-learn/scikit-learn) in Python

3.7 (https://www.python.org) was applied to aggregate the voxel clusters based on the signal intensities of T2WI and CE-T1WI or the values of nCBV and ADC reflecting functionally coherent subregions of the CEL. Using k-means clustering, samples from the dataset were classified as a given number of clusters with equal variances. Because the optimal number of clusters in a dataset is crucial to k-means clustering, three, four, and five clusters were initially evaluated (Supplementary Fig. 1).

Regarding the structural habitats, three clusters showed differences considering both T2WI and CE-T1WI results, whereas four and five clusters were identified by emphasizing the T2WI and CE-T1WI results, respectively. The four and five clusters displayed narrow ranges. Similarly, regarding the physiological habitats, three clusters demonstrated differences in both ADC and nCBV results, whereas four and five clusters with constrained ranges were identified by emphasizing the ADC and nCBV results, respectively. Therefore, based on the voxel-wise differences, three clusters were chosen to avoid overly parameterized models [16,23].

Structural habitats were defined as follows: an enhancing tissue habitat with high CE-T1 signal intensity irrespective of T2 signal intensity; a solid low-enhancing habitat with low T2 and CE-T1 signal intensity; and a nonviable tissue habitat with high T2 and low CE-T1 signal intensity. Low and high values were interpreted using the results of a data-driven analysis of k-means clustering without a specific threshold.

The ADC and CBV feature maps were used to define the following physiological habitats: a hypervascular cellular habitat with relatively low ADC and relatively high CBV values compared with other habitats; a hypovascular cellular habitat with relatively low ADC and CBV values; and a nonviable tissue habitat with relatively high ADC and relatively low CBV values.

The establishment of the tumor habitat is shown in Figure 2. For the longitudinal tumor habitat analysis, the differences in voxels in each habitat observed during the first and second examinations were calculated.

Analysis of the Recurrence Site

The recurrence site was analyzed using the tumor habitats established by the second follow-up examination and the confirmatory scan (MRI on the date of recurrence). The CEL at the recurrence site shown by the confirmatory scan was compared with the tumor habitats defined during





Fig. 2. Tumor habitats observed using structural MRI and physiological MRI to evaluate brain metastasis. To construct habitats using structural MRI, k-means clustering was applied for T2-weighted and contrast-enhanced T1-weighted images. To construct habitats using physiological MRI, k-means clustering was applied for ADC and CBV images. These images are of a 60-year-old male with colon cancer who was treated with stereotactic radiosurgery for metastasis in the left occipital lobe. Enhancing, solid low-enhancing, and nonviable tissue habitats are indicated by red, green, and blue subregions in the structural habitats, respectively. Similarly, hypervascular cellular, hypovascular cellular, and nonviable tissue habitats are represented by red, green, and blue subregions in the physiological habitats, respectively. ADC = apparent diffusion coefficient, CBV = cerebral blood volume

the second follow-up examination. The images obtained during the confirmatory scan were registered on the images obtained during the second follow-up examination. The region of interest of the CEL on the image obtained during the confirmatory scan was transferred to the tumor habitat observed during the second follow-up examination. The overlap of each habitat with the CEL at the time of recurrence was calculated using the Dice similarity coefficient $\binom{2|H\cap R|}{|H|+|R|}$ [24], where H indicates each tumor habitat and R indicates the CEL at the time of recurrence. Briefly, the calculation is performed as follows: 2 (overlapped area between each tumor habitat at the second follow-up and CEL at the time of recurrence)/(sum of the tumor habitat at the second follow-up + CEL at the time of recurrence). The Dice similarity coefficient ranged from 0 (no overlap) to 1 (perfect agreement).



Statistical Analysis

The demographic characteristics of the patients were compared using the chi-square test or Fisher's exact test for discrete variables and the Student's *t* test for continuous variables.

A univariable analysis using Cox regression or the Kaplan-Meier method (log-rank test) was performed to analyze the associations of tumor habitats with the time to recurrence. The hazard ratios (HRs) indicate the relative change in the hazard incurred by a 1-unit increase in each parameter; during this study, 5000 voxels were considered one unit. A multivariable analysis was not performed because each tumor habitat was derived from a given CEL volume, resulting in mutual dependency among the tumor habitats and leading to multicollinearity in the regression analysis.

For the significant predictors identified by the univariable Cox regression, an optimal cutoff for stratifying groups at low risk and high risk for recurrence was estimated using maxstat in R (version 4.1.1; Institute for Statistics and Mathematics) with 10-fold cross-validation, which ensured unbiased prediction within the sample [25]. Because the z-score is the ratio of each regression coefficient to its standard error, the Wald statistic is the asymptotically standard normal when it is hypothesized that the corresponding β is zero [26]. These data were used to demonstrate the statistical significance of each spatiotemporal habitat type.

The R Statistical Package (version 4.1.1; Institute for Statistics and Mathematics) was used for all statistical analyses. Statistical significance was defined as P < 0.05.

RESULTS

Patient Characteristics

Eighty-three patients (mean age, 59.0 years; range, 27–82 years; 44 male and 39 female) with a total of 103 brain metastases who underwent longitudinal followup including MRI after SRS were enrolled in the study. Table 1 lists the characteristics of the patients and their brain metastases. The tumor recurrence and RN subgroups were not statistically significantly different in terms of age, sex, primary cancer origin, time interval between the detection of brain metastasis and SRS, time from SRS to the first and second imaging follow-up examinations, additional radiation therapy, combined chemotherapy, targeted therapy, steroid use, mean number of times SRS was performed, and metastasis size before and at the time **Table 1.** Characteristics of the Patients and Their Metastatic

 Lesions

	Tumor	Radiation	Р	
	Recurrence	Necrosis	P	
Number of patients	34	49		
Age, years	60.0 ± 12.6	58.3 ± 10.5	0.508	
Male	18 (53)	26 (53)	> 0.999	
Primary cancer			0.199	
Lung	20 (59)	28 (57)		
Breast	5 (15)	7 (14)		
Other	9 (26)	14 (29)		
Time from detection to SRS, days	19.5 ± 10.5	15.6 ± 9.7	0.090	
SRS to first imaging follow-up, days	86.5 ± 34.1	83.0 ± 36.6	0.661	
SRS to second imaging follow-up, days	178.7 ± 54.7	189.8 ± 79.3	0.482	
SRS to confirmatory scan, days	350.9 ± 150.9	508.2 ± 349.4	0.016	
Adjuvant therapy			0.223	
Additional RT	0 (0)	1 (2)		
Combined with chemotherapy	4 (12)	1 (2)		
Targeted therapy	12 (35)	6 (12)		
Steroid usage	23 (68)	15 (28)	0.317	
Number of metastatic lesions	37	66		
Mean dose of SRS, Gy*	28.4 ± 14.6	24.6 ± 15.2	0.223	
Size prior to SRS, cm*	2.7 ± 1.1	2.8 ± 0.9	0.699	
Size at time of SRS, cm*	2.8 ± 1.0	2.9 ± 1.0	0.730	

Results are reported as the number (percent) or as mean \pm standard deviation. *Data for metastatic lesions. RT = radiation therapy, SRS = stereotactic radiosurgery

of SRS. The time from SRS to the confirmatory scan was longer in the RN subgroup (mean \pm standard deviation [SD], 508.2 \pm 349.4 days) than in the tumor recurrence group (mean \pm SD, 350.9 \pm 150.9 days) (*P* = 0.016).

Single-Time Analysis of the Tumor Habitat

The results of the univariable analysis performed to evaluate the associations between structural and physiological habitats and time to recurrence are summarized in Table 2. At the time of the first MRI examination, a solid low-enhancing habitat (low T2 and CE-T1 signal intensity) was associated with recurrence. Furthermore, a high number of voxels showing a solid lowenhancing structural habitat (HR, 1.54; 95% confidence interval [CI], 1.01–2.35; P = 0.045) were associated with recurrence. At the time of the second MRI examination, a



Table 2. Single Time Point Tumor Habitats and Their Associations

 with Recurrence of Brain Metastases After Stereotactic

 Radiosurgery

	Hazard Ratio	95% CI	Р			
First MRI						
Structural MRI habitats based on	T2WI and	CE-T1WI				
Enhancing tissue habitat	0.83	0.42-1.63	0.590			
Solid low-enhancing habitat	1.54	1.01-2.35	0.045			
Nonviable tissue habitat	0.45	0.14-1.50	0.195			
Physiologic MRI habitats based on ADC and CBV						
Hypervascular cellular habitat	1.21	0.35-4.13	0.764			
Hypovascular cellular habitat	0.70	0.30-1.61	0.401			
Nonviable tissue habitat	1.06	0.67-1.69	0.799			
Second MRI						
Structural MRI habitats based on	T2WI and	CE-T1WI				
Enhancing tissue habitat	0.97	0.50-1.89	0.927			
Solid low-enhancing habitat	1.95	1.23-3.08	0.004			
Nonviable tissue habitat	1.01	0.44-2.30	0.987			
Physiologic MRI habitats based o	n ADC and	CBV				
Hypervascular cellular habitat	20.1	2.90-139	0.002			
Hypovascular cellular habitat	1.56	0.92-2.66	0.100			
Nonviable tissue habitat	1.02	0.59-1.77	0.932			
Harard ratios reported have are for a 1 unit (E000 vevels) increase						

Hazard ratios reported here are for a 1 unit (5000 voxels) increase in each imaging parameter. ADC = apparent diffusion coefficient, CBV = cerebral blood volume, CE = contrast-enhanced, CI = confidence interval, T1WI = T1-weighted imaging, T2WI = T2weighted imaging

high number of voxels showing either a solid low-enhancing structural habitat (HR, 1.95; 95% CI, 1.23–3.08; P = 0.004) or a hypervascular cellular physiological habitat (HR, 20.1; 95% CI, 2.90–139; P = 0.002) were associated with recurrence. The concordance index (C index) was 0.65 (95% CI, 0.55–0.75) for all statistically significant tumor habitats at a single time point.

Longitudinal Analysis of the Tumor Habitat: Recurrence Risk

The results of the longitudinal analysis of the tumor habitats are presented in Table 3. During the longitudinal analysis, an increase in the hypovascular cellular habitat was strongly associated with recurrence (HR, 2.68; 95% CI, 1.46–4.91; P < 0.001). The C index was 0.71 (95% CI, 0.62– 0.80) after considering the changes in the hypovascular hypercellular habitat and all statistically significant tumor habitats at a single time point. Representative habitats of patients with SRS-treated metastasis with recurrence evaluated using longitudinal physiological MRI are shown in Figure 3 and Supplementary Figure 2. **Table 3.** Longitudinal MRI Tumor Habitat Analysis andAssociations with Recurrence of Brain Metastasis After StereotacticRadiosurgery

Change in Tumor Habitat	Hazard Ratio	95% CI	Р			
Structural MRI habitats based on T2WI and CE-T1WI						
Enhancing tissue habitat	3.08	0.52-18.3	0.216			
Solid low-enhancing habitat	1.64	0.75-3.61	0.215			
Nonviable tissue habitat	1.92	0.82-4.48	0.133			
Physiologic MRI habitats based on ADC and CBV						
Hypervascular cellular habitat	4.02	0.42-38.4	0.227			
Hypovascular cellular habitat	2.68	1.46-4.91	< 0.001			
Nonviable tissue habitat	0.70	0.18-2.76	0.607			

Hazard ratios reported here are for a 1 unit (5000 voxels) increase in each imaging parameter. ADC = apparent diffusion coefficient, CBV = cerebral blood volume, CE = contrast-enhanced, CI = confidence interval, T1WI = T1-weighted imaging, T2WI = T2weighted imaging

The optimal cutoff for the longitudinal physiological MRI evaluations of the habitats of groups at low risk and high risk for recurrence was an increase of more than 1345 voxels in the hypovascular cellular habitat. This cutoff separated the low- and high-risk groups, which were significantly different according to the log-rank test (P < 0.001). Supplementary Figure 3 shows the Kaplan–Meier survival curves of the groups at low risk and high risk for recurrence based on hypovascular cellular habitats.

Longitudinal Analysis of Tumor Habitat: Recurrence Sites

Thirty-seven lesions (35.9%; 37/103) were classified as tumor recurrence in 34 patients (41.0%; 34/83). The mean Dice similarity coefficient for the hypovascular cellular habitat observed using physiological MRI was the highest (0.423; range, 0.004–0.969; SD, 0.259), followed by the mean Dice similarity coefficient for the solid low-enhancing habitat observed using structural MRI (0.298; range, 0.015– 0.942; SD, 0.247). Figure 4 shows the relationship between temporal changes in the hypovascular cellular habitats and the recurrence site.

DISCUSSION

We demonstrated that brain metastasis recurrence after SRS could be predicted by performing a longitudinal analysis involving diffusion- and perfusion-weighted physiological MRI of the tumor habitat. An increase in the hypovascular cellular habitat, which exhibits both low ADC and CBV values, was significantly associated with the time





Fig. 3. Prediction of recurrence using a hypovascular cellular habitat. A 51-year-old male with brain metastasis attributable to lung cancer (adenocarcinoma) underwent whole brain radiotherapy and SRS of a lesion in the right frontal lobe. **A, B.** The longitudinal physiological MRI examination shows an increased hypovascular cellular habitat at the anterior aspect of the lesion and mild contrast enhancement. **C.** The confirmatory scan shows that the tumor progressed at the anterior aspect of the lesion. The time to recurrence was 16 months after SRS. ADC = apparent diffusion coefficient, CBV = cerebral blood volume, SRS = stereotactic radiosurgery

to recurrence of SRS-treated metastasis.

A tumor habitat analysis can identify distinct tumor subregions and cell populations that can be correlated with the biological state of the tissue by co-registering the images with histologic specimens [14,27]. We longitudinally analyzed tumor habitats using both physiological MRI and structural MRI for brain metastasis after SRS and compared their predictive values for recurrence. Structural MRI was found to be useful for the initial follow-up examination after SRS, during which a solid low-enhancing habitat (low T2 and CE-T1 signal intensity) was associated with recurrence. However, physiological MRI was helpful for predicting recurrence during the follow-up examinations and identified the tumor habitat that most closely matched the recurrence site. It was suggested that defining tumor subregions is helpful in radiation therapy planning, and the understanding of the spatial distribution of physiological tumor subregions could help optimize local radiotherapy [27,28]. Therefore, the hypovascular cellular subregion on images obtained with physiological MRI should be considered a target for further SRS or other local therapies.

When brain metastasis regrows after SRS, it is important to identify viable tumors that require follow-up imaging. Our study identified tumor habitats that could have viable tumors based on the findings of structural MRI and physiological MRI. These tumor habitats were solid





Fig. 4. Relationship between the increase in the hypovascular cellular habitat and recurrence site. A 64-year-old female with brain metastasis attributable to breast cancer. **A**, **B**. The longitudinal physiological MRI analysis shows that the hypovascular cellular habitat increased (from 1581 voxels to 3217 voxels) whereas the other habitats decreased. Spatial mapping shows an increase in the hypovascular cellular habitat at the medial and upper lateral aspects of the enhancing masses (arrows). **C**. Subsequently, the confirmatory scan performed during the 9-month follow-up examination shows that the hypovascular cellular habitat became the recurrence site (arrows). ADC = apparent diffusion coefficient, CBV = cerebral blood volume

low-enhancing with enhanced tissue on structural MRI and hypervascular cellular and hypovascular cellular on physiological MRI. Among them, the following habitats were associated with tumor recurrence at a single time point: solid low-enhancing habitat on structural MRI during the first and second follow-up examinations and hypervascular cellular on physiological MRI during the second follow-up examination. This suggests that viable tumors have inherent cellularity and a certain degree of vascularity. Accordingly, the aforementioned habitats may be potential prognostic factors for tumor recurrence. However, the predictive power for tumor recurrence cannot be guaranteed [29]. From the perspective of the longitudinal analysis, only changes in the hypovascular cellular habitat were associated with tumor recurrence, indicating that the cellularity of the viable tumor is an important factor that can determine tumor recurrence. Moreover, viable tumors can capture existing vessels or generate new ones [30,31]. However, the vasculature remains relatively low until vessel invasion [32]. Consequently, an increase in the hypovascular hypercellular habitat during longitudinal follow-up examinations might indicate a surge in viable tumors before vascular invasion.

Korean Journal of Radiology

Therefore, it could be used as a predictive biomarker to help determine the optimal additional therapy before overt tumor recurrence.

Importantly, we performed a habitat analysis of the CEL with both a solid low-enhancing habitat and a hypovascular cellular habitat within it. When the CEL and the solid portion and/or cellular portion with low T2 signal intensity were matched, tumor recurrence was indicated; however, when the CEL and solid portion and/or cellular portion were not matched, RN was indicated. Previous studies that used this interpretation found similar results. Using structural MRI, a T1/T2 mismatch, defined as nonmatching lesion boundaries on T2WI and CE-T1WI, showed 83.3% sensitivity and 91.1% specificity for RN [33]. Using physiological MRI, a layered appearance (without matching) consisting of outer, middle, and inner layers of high CBV, low ADC, and high ADC values, respectively, was suggestive of RN [34], with sensitivity and specificity rates of approximately 100% for 16 patients. Nonetheless, the visual analysis of enhancement patterns comprising T1/T2 signal intensity mismatch [35] or a layered appearance of high CBV, low ADC, and high ADC values with brain metastasis [36] is limited because it is a subjective assessment. In contrast, a quantitative tumor habitat analysis using k-means clustering applied to co-registered isometric voxels during our study provided objective and robust measurements that could be used to assess RN and viable tumors.

The hypovascular cellular habitat observed with physiological MRI had the highest mean Dice similarity coefficient (0.423; range, 0.004–0.969; SD, 0.259). The number of voxels of the CEL tended to change between the second follow-up examination and recurrence. Parametric response mapping [37], which monitors changes in a single voxel, provided an evaluation of the Dice similarity coefficient of the second follow-up examination and recurrence. Furthermore, parametric response mapping can be used to monitor each tumor habitat and determine the region matching the CEL at the time of recurrence, despite changes in tumor volume. As a result, rather than focusing on the actual number, a mean Dice similarity coefficient of 0.423 was utilized to determine the best tumor habitat corresponding to the recurrence site.

The number of patients with recurrent brain metastasis after SRS is likely to increase with improvements in systemic therapy [38]. The recent guidelines of the American Society of Clinical Oncology and the European Society of Clinical Oncology [1,38] recommend follow-up at 3-month intervals to diagnose brain metastasis recurrence, and that treatment should be continued according to the patient's performance status, neurological function, and prior treatment. A longitudinal analysis using a combination of diffusionand perfusion-weighted MRI can predict [29] the timing of recurrence and stratify patients into groups at low risk and high risk for recurrence by defining hypovascular cellular habitats. The results of this assessment could be helpful in planning the next treatment option, especially localized therapy, surgery, or further SRS.

This study has several limitations. First, the strict pathological confirmation of image-based segmentation is difficult. Only a few patients underwent surgery for the recurrence of brain metastasis. Accurate localization of brain metastasis recurrence is limited because of the surgical difficulty. Nonetheless, we attempted to demonstrate that the recurrence site matched the tumor habitat by analyzing serial MRI examination results. It is essential to establish biologically validated habitat imaging as a promising research field [28,39,40] by continuing to perform research and applying those results to solid tumors [41,42]. Therefore, future studies are necessary to confirm the radiological and pathological correlations. Second, regarding structural MRI, the assignment of voxels to the enhancing habitat, solid low-enhancing habitat, or nonviable tissue habitat is based on logical assumptions. A tumor habitat analysis is a data-driven analysis that groups similar voxels within the CEL; however, this type of colocalization could be difficult to determine in clinical practice.

In conclusion, an increase in the hypovascular cellular habitat observed using longitudinal physiological MRI was associated with the risk of recurrence (i.e., treatment resistance) and was indicative of the recurrence site. These results may be helpful for monitoring patients after SRS and determining future localized therapeutic targets.

Supplement

The Supplement is available with this article at https://doi.org/10.3348/kjr.2022.0492.

Availability of Data and Material

The datasets generated or analyzed during the study are available from the corresponding author on reasonable request.



Conflicts of Interest

Ji Eun Park, Seo Young Park, and Ho Sung Kim who is on the editorial board of the *Korean Journal of Radiology* was not involved in the editorial evaluation or decision to publish this article. All remaining authors have declared no conflicts of interest.

Author Contributions

Conceptualization: Ji Eun Park. Data curation: Young-Hoon Kim, Young Hyun Cho, Jeong Hoon Kim. Formal analysis: Seo Young Park. Funding acquisition: Ji Eun Park. Investigation: Da Hyun Lee, Ji Eun Park, Young-Hoon Kim. Methodology: Ji Eun Park, NakYoung Kim. Project administration: Ji Eun Park. Resources: Da Hyun Lee, Young-Hoon Kim. Software: NakYoung Kim. Supervision: Ji Eun Park, Ho Sung Kim. Validation: Young Hyun Cho, Jeong Hoon Kim, Ho Sung Kim. Visualization: Da Hyun Lee, Ji Eun Park. Writing—original draft: Da Hyun Lee. Writing—review & editing: Ji Eun Park.

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Funding Statement

This research was supported by the Ministry of Health& Welfare, Republic of Korea (HI21C1161) and by a grant (2021IP0078) from the Asan Institute for Life Sciences, Asan Medical Center, Seoul, Korea.

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