ORIGINAL ARTICLE

3'-Deoxy-3'-[¹⁸F]-fluorothymidine ([¹⁸F]-FLT) transport in newly diagnosed glioma: correlation with nucleoside transporter expression, vascularization, and blood-brain barrier permeability

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Abstract 3'-Deoxy-3'- $[^{18}F]$ -fluorothymidine ($[^{18}F]$ -FLT), a marker of cellular proliferation, has been used in positron emission tomography (PET) examination of gliomas. The aim of this study was to investigate whether the uptake of ¹⁸F]-FLT in glioma correlates with messenger RNA (mRNA) levels of the equilibrative nucleoside transporter 1 (ENT1), microvascular density (assessed by CD34 immunohistochemistry), and the blood-brain barrier (BBB) breakdown. A total of 21 patients with newly diagnosed glioma were examined with [¹⁸F]-FLT PET. Tumor lesions were identified as areas of focally increased [¹⁸F]-FLT uptake, exceeding that of surrounding normal tissue. Dynamic analysis of [¹⁸F]-FLT PET revealed correlations between the phosphorylation rate constant k_3 and ENT1 expression; however there was no correlation between the kinetic parameters and CD34 score. There was a good correlation between the gadolinium (Gd) enhancement score (evaluating BBB breakdown) and ENT1 expression, CD34 score, and Ki-67 index. This preliminary study

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Department of Cell Physiology, Kagawa University Faculty of Medicine, Miki, Kagawa, Japan suggests that ENT1 expression might not reflect accumulation of [¹⁸F]-FLT in vivo due to BBB permeability in glioma.

Keywords $[^{18}F]$ -FLT · Nucleoside transporter · CD34 · Blood-brain barrier · Glioma

Introduction

3'-deoxy-3'-[¹⁸F]-fluorothymidine ([¹⁸F]-FLT), a radiolabeled fluorinated thymidine analog, has been attracting attention as a promising positron emission tomography (PET) tracer for evaluating cell proliferation activity reflecting the ability to synthesize nucleic acid in various malignant tumors [1–4]. [¹⁸F]-FLT PET has been shown to be useful for preoperative grading and assessing cell proliferation activity in human glioma [4, 5]. In addition, [¹⁸F]-FLT PET imaging is under investigation to potentially monitor reprogression and response to chemoradiotherapy in postoperative human glioma.

The mechanism of [¹⁸F]-FLT uptake into cells is a based on thymidine metabolism. [¹⁸F]-FLT and can be phosphorylated by thymidine kinase 1 (TK1), a principle enzyme in the salvage pathway of DNA synthesis, and trapped inside proliferative cells [1]. However, [¹⁸F]-FLT must cross plasma membranes before it can interact with TK1. To interpret [¹⁸F]-FLT PET images as measures of cell proliferation, we need to understand the mechanisms for its transport, accumulation, and retention in tissues. Efficient cellular uptake requires the presence of specialized plasma membrane nucleoside transporter proteins [6, 7]. The equilibrative nucleoside transporters (ENT) are bidirectional and broadly selective. They catalyze the facilitated diffusion of nucleosides down their concentration gradients

[8]. ENT1 is a ubiquitous protein and is the most abundant and widely distributed plasma membrane nucleoside transporter in the human central nervous system [9]. After intravenous injection, FLT distributes rapidly in the extracellular fluid and is transported into the cytosol by nucleoside transporters, probably mainly by the ENT1 [10, 11]. Recent studies report that ENT1 expression might be important for drug toxicity (gemcitabine and capecitabine), and [¹⁸F]-FLT uptake also might predict a sensitivity to drug-induced inhibition of nucleic acid synthesis [11–13]. In addition, our recent study demonstrated that [¹⁸F]-FLT uptake in the tumor could be significantly related to transport through the disrupted blood-brain barrier (BBB) but not through phosphorylation [14]. Generally, magnetic resonance (MR) imaging after contrast enhancement can be used to evaluate BBB breakdown in the tumor.

On the other hand, angiogenesis is fundamental in tumor growth and progression [15]. A recent study reported a correlation between the imaging of cell proliferation and tumor angiogenesis as reflected by microvessel density (MVD) in lung cancer [16]. Immunohistochemistry for CD34 can be used to determine MVD in malignant tumors.

The aim of our study was to investigate [¹⁸F]-FLT uptake and evaluate whether it correlates with ENT1 mRNA expression, microvascular density (CD34 score), or BBB breakdown in human brain glioma.

Materials and methods

Patients

Between April 2009 and March 2012, 21 consecutive patients with newly diagnosed gliomas (14 men, 7 women; mean age 45.2 \pm 17.2 years, range 21–72 years) were investigated. The use of [¹⁸F]-FLT as a PET tracer was approved by Kagawa University Faculty of Medicine, Human Subjects Ethics Committee. Informed consent for ¹⁸F]-FLT PET with arterial blood sampling and use of tumor samples for further histological and biochemical analysis was obtained from all patients before the examination. All patients underwent preoperative [¹⁸F]-FLT PET kinetic analysis and MR imaging with contrast enhancement usually within 1 week. No patients had had previous treatments except steroid therapy before the PET examination. Histopathology was confirmed in tissue specimens obtained by stereotactic biopsy or resection using a navigation system (StealthStation; Medtronic-Sofamor Danek, Memphis, TN, USA) from the region of interest (ROI) in which [¹⁸F]-FLT kinetic parameters were evaluated. All tumors were graded according to the World Health Organization (WHO) grading system (malignancy scale) for central nervous system tumors. Tumor types and grades were distributed as follows and are listed in Table 1: WHO grade II diffuse astrocytoma (n = 3), WHO grade II oligoastrocytoma (n = 1), WHO grade III anaplastic astrocytoma (n = 5), WHO grade III anaplastic oligodendroglioma (n = 1), WHO grade III anaplastic oligoastrocytoma (n = 1), and WHO grade IV glioblastoma multiforme (n = 10).

Positron emission tomography scanning procedure

¹⁸Fl-FLT was synthesized using an HM-18 cyclotron (Sumitomo Heavy Industries Co., Tokyo, Japan) using a previously reported procedure [12]. The radiochemical purity of the $[^{18}F]$ -FLT produced was >95 %. The specific activity of [¹⁸F]-FLT at the time of patient injection was >7.4 GBg/µmol (>0.2 Ci/µmol). Dynamic [¹⁸F]-FLT PET scanning was performed using an ECAT EXACT HR + Scanner (Siemens/CTI, Knoxville, TN, USA) in the three-dimensional acquisition mode (collimating septa retracted). The imaging system enables 51 simultaneous transverse acquisitions per field of view (FOV), with intersection spacing of 3 mm, for a total axial FOV of 15 cm. The in-plane reconstructed resolution was 4.7 mm full width at half maximum. [¹⁸F]-FLT PET examination was performed 2–14 (mean 8.5 \pm 3.8) days before surgery. No special dietary instructions were given to patients before examination. Nine patients received steroid treatment for brain edema at the time of PET examination. Images were acquired with patients in the supine position, resting, with their eyes closed. First, a 5-min transmission scan for attenuation correction was obtained using an external source of ⁶⁸Ge rod sources rotating around the head. Dynamic emission data were acquired for 60 min $(4 \times 20 \text{ s}, 4 \times 40 \text{ s}, 4 \times 60 \text{ s}, 4 \times 180 \text{ s}, 8 \times 180 \text{ s})$ 300 s) starting immediately after intravenous bolus injection of $[^{18}F]$ -FLT (mean 279 \pm 72 MBq). Arterial blood samples were drawn from the brachial artery at 15-s intervals for the first 3 min, followed by increasingly longer intervals to 60 min, to measure arterial plasma radioactivity using an automatic well-type gamma counter (ARC-400, Aloka, Tokyo, Japan). Dynamic PET acquisition data were reconstructed by the filtered back-projection method using a Hanning filter (cutoff frequency, 0.4 cycles per pixel) to generate 128×128 matrices.

Static and semiquantitative [18 F]-FLT uptake in the tumor was measured by determining the standardized uptake value (SUV). An ROI was set manually by an observer around the hottest area of each lesion or its center located on the MR image if increased [18 F]-FLT uptake was absent. The maximum SUV (SUV_{max}) was regarded as a representative value of [18 F]-FLT uptake in each tumor. Alternatively, the tumor to normal brain tissue (T/N) ratio was calculated for evaluating [18 F]-FLT uptake in the

Table 1 Summary of 21 cases with clinical and in vitro data

No.	Age/sex (year)	WHO grade	Dig.	FLT-kinetic			SUV _{max}	T/N	MRI + grade	Ki-67	CD34 (IHC)	ENT1
				$\frac{K_1 \text{ (ml/min/g)}$	k ₃ (/min)	<i>K_i</i> (ml/ min/g)		ratio	(score)	index (%)	(score)	(PCR)
1	27/F	4	GBM	0.110	0.090	0.025	1.9	17.27	3	80	2	157
2	26/M	4	GBM	0.039	0.072	0.017	1.88	11.75	3	80	3	570
3	37/M	4	GBM	0.071	0.055	0.012	1.41	8.81	3	20	3	195
4	61/F	4	GBM	0.087	0.151	0.016	1.51	9.44	3	70	2	782
5	28/F	4	GBM	0.090	0.292	0.025	1.53	6.65	2	30	2	112
6	53/M	4	GBM	0.085	0.106	0.027	1.70	9.44	3	25	2	175
7	62/M	4	GBM	0.058	0.093	0.018	1.53	9.00	3	30	2	82
8	67/M	4	GBM	0.061	0.092	0.013	1.45	7.63	3	30	3	279
9	54/M	4	GBM	0.063	0.302	0.024	2.06	9.81	3	40	3	1,414
10	72/M	4	GBM	0.081	0.093	0.019	1.62	9.53	3	50	3	410
1	70/F	3	AA	0.038	0.076	0.007	1.11	6.53	0	30	1	69
2	45/M	3	AA	0.040	0.000	0.000	0.89	6.85	1	10	3	152
3	31/M	3	AA	0.043	0.094	0.015	1.19	10.81	1	20	2	312
4	33/M	3	AA	0.057	0.265	0.020	2.09	13.06	1	70	3	106
5	66/F	3	AA	0.037	0.060	0.006	2.98	10.64	2	15	2	206
6	64/M	3	AO	0.129	0.069	0.013	2.76	19.71	2	50	2	263
7	28/F	3	AOA	0.027	0.047	0.014	0.96	7.38	0	5	1	303
1	31/M	2	DA	0	0	0	0.52	2.36	0	3	2	60
2	36/M	2	OA	0	0	0	0.36	2.57	0	8	1	68
3	37/F	2	DA	0	0	0	0.79	2.72	1	1	2	100
4	21/M	2	DA	0	0	0	0.78	3.12	1	2	2	249

Dig. diagnosis, *FLT* fluorothymidine, *GBM* glioblastoma multiforme, *AA* anaplastic astrocytoma, *AO* anaplastic oligoastrocytoma, *OG* anaplastic oligoastrocytoma, *DA* diffuse astrocytoma, *SUV* standard uptake value, *MRI* magnetic resonance imaging, *IHC* immunohistochemistry, *ENT1* equilibrative nucleoside transporter 1, *PCR* polymerase chain reaction

lesion. The T/N ratio is calculated by dividing the tumor SUV_{max} by the mean SUV of contralateral normal gray matter.

Kinetic analysis

Kinetic analysis of [¹⁸F]-FLT PET data was conducted using a two-tissue compartment four-rate parameter model involving blood volume fraction, as previously described [17]. This model is similar to the two-tissue compartmental model for fluorodeoxyglucose (FDG) [18]. The transfer from blood into tissue across the BBB is represented by K_1 (ml/min per gram), whereas the return of [¹⁸F]-FLT from tissue representing nonphosphorylated [¹⁸F]-FLT back to the blood is represented by k_2 (/min). K_1 represents transport of [¹⁸F]-FLT by the nucleoside transporters across the intact BBB in unenhanced tumors, whereas K_1 represents mainly passive [¹⁸F]-FLT influx through the disrupted BBB in well-enhanced tumors. The metabolic trapping of $[^{18}F]$ -FLT through phosphorylation is represented by k_3 (/min) and is the rate-limiting step for its intracellular trapping in tissue. Loss of image signal through these processes is described by k_4 (/min), which represents dephosphorylation and subsequent efflux via transporters or diffusion. Metabolic flux of [¹⁸F]-FLT is estimated from parameters derived by fitting the [¹⁸F]-FLT input function and the total blood activity curve to tissue time-activity curve data. In the calculation, it appears that k_4 is very small compared with k_3 , because the dephosphorylation rate is very low in tumor tissue [17, 18], thus, k_4 was assumed to be zero. Parametric images were calculated using the Lawson–Hanson nonnegative least-squares algorithm. The total metabolic flux constant of [¹⁸F]-FLT (K_i) was estimated from these parameters as:

 $K_i = K_1 \times k_3/(k_2 + k_3)[\text{ml/min/g}].$

The rate constant values were obtained from the calculated parametric images using ROIs placed, as described above. A parametric image-based method was used for retrieving the [¹⁸F]-FLT PET kinetic parameters, as the quality of the parametric images was sufficient for evaluation with acceptable noise. This area always contained the maximum [¹⁸F]-FLT uptake area set for SUV_{max} measurement.

Magnetic resonance imaging

MR images after contrast enhancement were used to evaluate BBB breakdown in the tumor and to identify exact lesion location. T_1 -weighted MR images with 0.1 mmol/kg gadolinium diethylenetriaminepentaacetic acid (Gd-DTPA) were obtained using a 1.5-T whole-body MR imaging unit (Signa Advantage, GE Medical Systems, Milwaukee, WI, USA, or VISART, Toshiba, Tokyo, Japan). The degree of Gd enhancement in the tumor on T_1 -weighted MR images was evaluated visually by comparing the intensity of lesion enhancement with that of the normal choroid plexus [19]. Gd enhancement in tumors was graded as score 0 (none), score 1 (less than that of choroid plexus), score 2 (equal to choroid plexus), and score 3 (more than choroid plexus).

Real-time polymerase chain reaction (PCR) assay

Total RNA from frozen tumor tissue (5 mm³ samples) was extracted using Isogen (Nippon Gene, Toyama, Japan), according to the manufacturer's instructions, and quantified by measuring absorbance at 260 nm. Total RNA (1 µg) was reverse-transcribed into complementary DNA (cDNA) using the Prime Script II 1st strand cDNA synthesis Kit (TAKARA Bio Inc, Tokyo, Japan). Quantitative real-time (RT)-PCR was performed using an ABI Prism 7300 RT-PCR system (Applied Biosystems, Carlsbad, CA, USA) with TAKARA Premix Ex Tag Perfect real time (TAKARA Bio Inc, Tokyo, Japan). Primer probes for human ENT1 (hENT1) (SLC29A1) and human β -actin as an internal control were purchased from Applied Biosystems (Assay ID: Hs01085706 m1 and Hs99999903 m1, respectively). PCR conditions were as follows: initial denaturing at 95 °C for 10 s, followed by 40 cycles of 95 °C for 10 s and 60 °C for 31 s. Reactions were run in triplicate in 96-well plates using standard default settings. Reactive quantification of RNA was determined using the delta-delta Ct method.

Immunohistochemistry

A representative standard formalin-fixed, paraffin-embedded section (4-µm thick) from each brain tumor specimen was immunohistochemically stained with each primary antibody using the avidin–biotin–peroxidase complex technique and Histofine Simple Stain MAX Peroxidase detection reagent kit designed specifically for immunohistochemical staining on formalin-fixed paraffin-embedded human tissue sections (Nichirei Biochemicals Co, Tokyo, Japan). Primary antibodies were rabbit anti-CD34 (monoclonal; 1:150, Abcam Inc. CA, USA) and mouse anti-Ki-67 (monoclonal; 1:50, Dako, Glostrup, Denmark). Normal rabbit and mouse immunoglobulin G (IgG) was used as negative controls. We performed semiquantitative immunohistochemistry analysis of CD34 in microvascular endothelial cells on the basis of both the percentage of positive microvessels and the intensity of staining (score 0–3) according to methods described previously [20, 21]. Briefly, in area of the most intense CD34-positive neovascularization, individual microvessel counts (MVCs) were made on a 400× magnification field. Any endothelial cell or endothelial cluster was considered a single countable microvessel. MVC was expressed as an absolute number of microvessels (NMVs) per 400× field for each case. Cases in which NMVs \leq 5 were considered score 0, $5 < NMV \leq 15$ were score 1, $15 < NMV \leq 35$ were score 2, and NMV > 35 were score 3. This grading procedure was performed by a blinded observer without prior knowledge of data in all cases (Fig. 1).

Statistical analysis

Correlation of the T/N ratio with the Ki-67 index and the $[^{18}F]$ -FLT kinetic parameters (K_1 , k_3 , and K_i) were calculated using linear regression analysis. Linear regression analysis was also used to examine whether the [¹⁸F]-FLT kinetic parameters $(K_1, k_3, \text{ and } K_i)$ were related to ENT1 expression determined by RT-PCR. Correlation of the Gdenhancement score with K_1 , K_i , T/N ratio, Ki-67, ENT1 (RT-PCR), and CD34 score were analyzed using nonparametric Spearman's rank test. The relationship between ¹⁸F]-FLT kinetic parameters and the grading of immunohistochemical (IHC) staining for CD34 (CD34 score) were also analyzed using the nonparametric Spearman's rank test. All statistical analyses were conducted using statistical analysis software for Excel 2010 (Ekuseru-Toukei 2010; Social Survey Research Information Co., Tokyo). P values <0.05 were considered statistically significant.

Results

Values for kinetic parameters, SUV_{max} and T/N ratio, MR imaging and CD34 scores, Ki-67 index and ENT1 mRNA levels are given in Table 1. First, there was a good correlation between the T/N ratio for [¹⁸F]-FLT and the Ki-67 index, a measure of proliferation (r = 0.733, p = 0.0002; Fig. 2a). To understand the basis of the increased T/N ratio, we examined the relationship between the composite K_i with T/N and found good correlation (i = 0.61, p = 0.0034; Fig. 2b). Changes in K_i may reflect changes in K_1 (influx across the BBB) and k_3 (phosphorylation of [¹⁸F]-FLT). There was good correlation between K_1 and T/N ratio (r = 0.8042, p < 0.0001; Fig. 2c). There was no significant correlation between k_3 and T/N ratio (Fig. 2d).

Next, BBB disruption may cause an increase in K_1 and thus K_i . Indeed K_1 and K_i were both significantly correlated with

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Fig. 1 a A three-compartment model for [¹⁸F]-fluorothymidine ([¹⁸F]-FLT) kinetics used in this study. Microphotographs shows immunohistochemistry [equilibrative nucleoside transporter 1 (ENT1), CD34, and Ki-67]. **b** Immunohistochemical analysis (score 1-3) of CD34 expression. Gliomatosis showing CD34 expression in microvascular endothelial cells. Number of microvessels (NMVs) per ×400 field for each case. The cases in which NMV < 5 were considered score 0, $5 < NMV \le 15$ were score 1, 15 < NMV < 35were score 2, and NMV > 35were score 3. Insets higher magnification



BBB disruption (Gd-enhancement score) (p = 0.0004, Fig. 3a; p = 0.0032; Fig. 3b). In addition, BBB disruption correlated with both T/N ratio and Ki-67 proliferative index (p = 0.0111, Fig. 3c; p = 0.0024, Fig. 3d). The Gd-enhancement score also showed significant correlations with ENT1 mRNA levels and CD34 scores (p = 0.016, Fig. 3e; p = 0.007, Fig. 3f).

Finally, in contrast to the correlation of K_1 and K_i with BBB disruption, there was no significant correlation of these two parameters with ENT1 mRNA levels (Fig. 4a, b), although there was a significant relationship between ENT1 and k_3 (r = 0.4778, p = 0.0285, Fig. 4c). There was also no significant correlation of the CD34 score and either K_1 , K_i or k_3 (Fig. 5a–c).

Discussion

The present study, including 21 patients with glioma, investigated the relationship between ¹⁸F]-fluorothymidine

([¹⁸F]-FLT) positron emission tomography (PET) kinetic parameters and expression of ENT1 by RT-PCR, microvascular density, and BBB disruption. FLT is an antiretroviral drug that inhibits DNA synthesis [22]. Shields et al. developed [¹⁸F]-FLT as a tracer to image proliferation in vivo. They also established [¹⁸F]-FLT as an analog substrate of thymidine, which is intracellularly phosphorylated during the S-phase [1].

[¹⁸F]-FLT transport in gliomas

This study addressed the hypothesis that the increased [¹⁸F]-FLT accumulation in gliomas might reflect changes in ENT1 expression [10, 11]. However, whereas analysis of [¹⁸F]-FLT kinetics with PET revealed a correlation between the phosphorylation rate constant k_3 and ENT1 expression, there was no correlation between the influx rate constant K_1 and ENT1 expression. These results suggest that the influx rate constant K_1 might reflect not only [¹⁸F]-

Fig. 2 Linear regression analysis between T/N ratio [¹⁸F]-fluorothymidine ([¹⁸F]-FLT) and the Ki-67 index, a measure of proliferation, and the [¹⁸F]-FLT positron emission tomography (PET) kinetic parameters $(K_1, k_3 \text{ and } K_i)$. **a** There was a significant correlation between T/N ratio and Ki-67 index (r = 0.733, p = 0.0002). **b** There was a correlation between T/N ratio and K_i (r = 0.61, p = 0.0034) and **c** K_1 (r = 0.8042, p < 0.0001), **d** but not k_3

Fig. 3 Correlation the bloodbrain barrier (BBB) breakdown in tumors [gadolinium (Gd) enhancement score] with kinetic parameter (K_1 and K_i), T/N ratio, Ki-67, equilibrative nucleoside transporter 1 (ENT1) [real-time polymerase chain reaction (RT-PCR)], and CD34 score using the nonparametric Spearman's rank test. There was a significant correlation of the Gd-enhancement score with a $K_1 (p = 0.0004), \mathbf{b} \ K_i$ (p = 0.0032,), c T/N ratio(p = 0.0111), **d** Ki-67 index (p = 0.0024), e ENT1 (RT-PCR) (p = 0.016), and **f** CD34 score (p = 0.007)





Fig. 4 Linear regression analysis between equilibrative nucleoside transporter 1 (ENT1) messenger RNA (mRNA) expression by realtime polymerase chain reaction (RT-PCR) and the [18 F]-fluorothymidine ([18 F]-FLT) positron emission tomography (PET) kinetic

parameters (K_1 , K_i and k_3). There was no correlation between **a** K_1 and **b** K_i , but **c** there was a correlation between ENT1 (RT-PCR) and k_3 (p = 0.0285



Fig. 5 Spearman's rank correlation test was used between CD34 score expression by immunohistochemistry and [18 F]-FLT PET kinetic parameters (K_1 , K_i and k_3). There were no correlations between CD34 score and kinetic parameters

FLT accumulation in tumor cells but leakage to extracellular space across the BBB, and the effect of [¹⁸F]-FLT uptake via nucleoside transporters on the transport rate constant K_1 might be very small compared with the effect of influx through the disrupted BBB. It might suggest the reason why k_3 relates to ENT1 expression. These results suggest that FLT accumulation might reflect inflammation and postoperative changes.

Ullrich et al. [23] reported that kinetic analysis models with [¹⁸F]-FLT PET enable discrimination of increased [¹⁸F]-FLT uptake due to increased transport through the BBB. The long half-life of [¹⁸F allows for kinetic analysis, providing differentiation between metabolized and nonmetabolized [¹⁸F]-FLT in tissue. Kinetic analysis of [¹⁸F]-FLT PET imaging was performed to estimate the extent to which [¹⁸F-FLT uptake was due to influx and transport effects across the BBB versus tracer retention related to phosphorylation and tumor growth. Previous studies reported that kinetic analysis of [¹⁸F]-FLT PET could clarify whether transport or metabolic trapping largely contributes to the increased accumulation of FLT in tumors [24]. The mathematical aspects of the $[^{18}F]$ -FLT model and the boundary conditions were adopted in our implementation of the previous [¹⁸F]-FLT kinetic model [17, 25].

[¹⁸F]-FLT and angiogenesis in gliomas

In lung cancer, Yang et al. [16] reported a correlation between [¹⁸F]-FLT imaging of cell proliferation and tumor angiogenesis, as reflected by MVD. There are no reports about [¹⁸F]-FLT PET and MVD in gliomas. We evaluated whether [¹⁸F]-FLT PET kinetics reflected MVD (CD34 score) in gliomas. However, there was no correlation between any of the kinetic parameters and CD34 score. It is possible that CD34 might not be suitable as an MVD marker, as CD34 staining is seen in both mature and immature vessels [16]. Other markers should be examined in the future.

[¹⁸F]-FLT and BBB permeability

A previous study showed that the major portion of $[^{18}F]$ -FLT uptake is similar to the low levels observed in unenhanced lesions on MR imaging and demonstrates inside the volume of increased permeability defined by MR imaging T₁-weighted Gd enhancement [26]. In this study, gliomas with no MR imaging T₁-weighted Gd enhancement had low accumulation on $[^{18}F]$ -FLT PET images compared with those with enhancement, and there was a good correlation between [¹⁸F]-FLT PET images (SUV_{max}, T/N ratio) and Gd-enhanced MR imaging. In general, Gd contrast enhancement on T₁-weighted MR images of brain lesions, such as tumor, necrotic tissue, and inflammation, is due to BBB disruption and increased blood vessel permeability [27]. It should be noted that there was some increase in T/N ratio for [¹⁸F]-FLT in tumors without Gd enhancement (Table 1), and this remains to be understood. [¹⁸F]-FLT can cross the BBB and plasma membranes; however, the mechanisms of transport and influx to tumor cells across the BBB in vivo have not been fully elucidated. To interpret [¹⁸F]-FLT PET images as a measure of cell proliferation, we need to understand the mechanisms that account for its transport, accumulation, and retention in tissues.

Conclusions

This study demonstrates that $[^{18}F]$ -FLT uptake in newly diagnosed human brain gliomas is related to transport through the disrupted BBB but not to phosphorylation activity. Although tissue ENT1 expression reflects the phosphorylation rate constant k_3 , the transport rate constant K_1 determined by $[^{18}F]$ -FLT PET kinetic analysis did not accurately reflect ENT1 expression and should not be used as a surrogate biomarker of transport activity in human brain gliomas. Several MVD markers should be examined in future studies.

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Conflict of interest None.

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