Kinetics of 3'-Deoxy-3'-¹⁸F-Fluorothymidine During Treatment Monitoring of Recurrent High-Grade Glioma

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3'-deoxy-3'-18F-fluorothymidine (18F-FLT) is used as a biomarker of cell proliferation. We investigated the kinetics of ¹⁸F-FLT during treatment of malignant glioma with bevacizumab and irinotecan. Methods: Fifteen patients with recurrent high-grade brain tumors (2 grade III, 13 grade IV) were studied at baseline (study 1 [S1]), after 1 course of therapy (2 wk, study 2 [S2]), and at the end of therapy (6 wk, study 3 [S3]). ¹⁸F-FLT (1.5 MBq/kg) was administered intravenously, and dynamic PET was performed for 1 h. Curves representing blood clearance and tumor uptake were derived from factor images and summed frames with thresholding techniques or with a fixed cube. The standard ¹⁸F-FLT model was used to estimate the rate constants. ¹⁸F-FLT uptake was measured at 2 time points (early standardized uptake value [SUV_{early}] and late SUV [SUV_{late}]). Results: Parameters appeared similar for curves derived from factor images and summed frames; the steepest drop occurred between S1 and S2 for transport, influx, SUV_{early} , and SUV_{late} . Three groups were distinguished on the basis of clinical outcome: patients who died within 6 mo (group 1 [G1], n = 4), survived 6–12 mo (group 2 [G2], n = 6), and survived more than 1 y (group 3 [G3], n = 5). None of the rate constants was significantly different between the groups. Long-term survivors (G3) showed a significantly different SUV change (in percentage) between S1 and S3, whereas short-term survivors (G1 and G2) did not. Conclusion: Overall, the relative SUV change from S1 to S3 predicted a favorable clinical outcome, whereas the SUV change from S1 to S2 did not. Long-term survivors (G3) showed a significant drop in SUV from S1 to S2 and from S1 to S3. Significant correlations were found between SUV and both the rate constant and the influx rate. The correlation coefficient between SUV_{late} and influx rate was 0.91, permitting response monitoring by the measurement of ¹⁸F-FLT uptake changes.

Key Words: ¹⁸F-FLT (3'-deoxy-3'-¹⁸F-fluorothymidine); positron emission tomography; malignant brain tumor; kinetic modeling; factor analysis; therapy monitoring

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he imaging of cell proliferation has emerged as an avenue of monitoring neoplastic disease. 3'-deoxy-3'fluorothymidine (FLT) labeled with ¹⁸F has provided a noninvasive way of studying cell proliferation. Although ¹⁸F-FLT is not incorporated into DNA, it is phosphorylated by thymidine kinase-1 and trapped inside the cytosol. Phosphorylated ¹⁸F-FLT appears suitable for imaging with PET (*1*,*2*) and represents thymidine incorporation into DNA, as has been demonstrated in various tumor models including 2 glioma cell lines (*3*). The clinical application of ¹⁸F-FLT in brain gliomas has been reported (*4*,*5*). The kinetic model has been described elsewhere (*6*).

Various tracers have been used to monitor therapy. For PET and glioma, results have been reported for ¹⁸F-FDG, ¹¹C-methionine, ¹⁸F-tyrosine, and others (7–11).

The National Cancer Institute trial on recurrent glioma using PET and ¹⁸F-FLT showed that the influx rate K_i and phosphorylation rate k_3 reached statistical significance in separating recurrent glioma (n = 15) from radiation necrosis (n = 4) (12). It also reported that estimates of the standardized uptake value (SUV) or visual analysis were not able to distinguish these entities. In addition, the authors of this report found that ¹⁸F-FLT performed better than ¹⁸F-FDG.

In a previous study (5), the percentage change in ¹⁸F-FLT uptake was used to categorize patients as responders (SUV decrease > 25%) or nonresponders. In the current study, we investigated the kinetics of ¹⁸F-FLT of recurrent high-grade brain tumors during treatment. These parameters provide more information than did the ¹⁸F-FLT uptake alone, and our hypothesis is that they contain predictive information. In addition, we evaluated the effect of different analysis methods and compared their results. A volume of interest (VOI) was defined to generate the input curve and another to derive the output curve. Two types of VOI were used of variable or fixed size and defined on parametric or composite images. These methods can be automated easily and are suitable quantification techniques for routine clinical applications.

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The tracer kinetics were investigated with compartmental modeling. In this study, the absolute value of parameters, and their relative change, was evaluated, and the different analysis techniques were compared. In addition, ¹⁸F-FLT uptake was measured at early and late time points and expressed as SUV. The final results were correlated to clinical outcome and survival.

MATERIALS AND METHODS

Patients

Twenty patients with recurrent glioma were prospectively enrolled. Nine were included from our previous study (5), and 11 were newly enrolled. All patients underwent dynamic ¹⁸F-FLT studies at baseline and during therapy, after which, 4 patients had to be removed because of technical issues (2 had severe head motion during the study, 1 had an incomplete study, and the other had an infiltrated dose [inadequate bolus with activity remaining in the arm]). One patient had a complete response to therapy; no tumor could be delineated in the ¹⁸F-FLT PET study after treatment, and kinetic modeling failed.

The final study population consisted of 15 patients (6 men and 9 women; average age, 53 y; age range, 26–77 y). The study was approved by our Institutional Review Board, and all patients signed an informed consent form. Histopathology revealed 13 grade IV tumors (glioblastoma multiforme) and 2 grade III tumors (anaplastic astrocytoma and anaplastic mixed glioma). Each patient underwent 3 studies: at baseline (study 1 [S1]), after 1 course of therapy (average interval, 16 ± 6 d; study 2 [S2]), and after 3 courses of therapy (average interval, 30 ± 8 d; study 3 [S3]). For every patient, the most active tumor (i.e., highest ¹⁸F-FLT uptake) at baseline was selected. Therapy consisted of biweekly cycles of bevacizumab (10 mg/kg) and irinotecan (125 or 350 mg/m²). Table 1 shows the population data, grade, number of treatments, and pathology of the malignant gliomas.

Image Acquisition

¹⁸F-FLT was synthesized by a modification of a previously reported procedure (13). A 1.5 MBq/kg (0.04 mCi/kg) dose of ¹⁸F-labeled FLT was administered. PET was performed using an ECAT HR+ system (CTI/Siemens). A transmission scan (2dimensional mode) of 5-min duration was acquired first. ¹⁸F-FLT was administered intravenously, and a dynamic acquisition (3-dimensional mode) was started simultaneously. Twenty-three frames $(8 \times 15, 2 \times 30, 2 \times 60, \text{ and } 11 \times 300 \text{ s})$ were acquired. The images were reconstructed with iterative techniques as described elsewhere (14). The following parameters were used: maximum a posteriori for the transmission scan and ordered-subset expectation maximization for the emission scan, corrections for attenuation and scatter, and a gaussian kernel of 5 mm in full width at half maximum as a postreconstruction smoothing filter. The final reconstructed volume set had a matrix size of 128×128 and consisted of 63 planes. The voxel size was $2.06 \times 2.06 \times 2.425$ mm, using a zoom of 2.5 in the axial planes.

Processing

Factor analysis was performed on the volume set of reconstructed images as described elsewhere (15). Briefly, factor analysis evaluates temporal changes in dynamic datasets and generates functional images of the vessels, tumor, brain structures, bone marrow, and the like. Rebinning produced a dataset with $4.1 \times 4.1 \times 4.9$ mm voxels that was used for further processing. For our application, factor analysis generated 3 factors and corresponding factor images, which are the associated spatial distribution of a factor. Factor 1 represents the large vessels and factor 2 the tumors in the brain. For every patient, studies were available at 3 time points (S1, S2, and S3). For the blood curve, VOIs were generated using a 50% threshold of the maximum voxel in the transverse sinuses. These VOIs for the input function were generated completely automatically with our processing modules. For the brain tumor, VOIs were derived from factor images (F2) or summation of frames between 45 and 60 min

					World Health				World Health		
			Age	Initial	Organization			Pathology	Organization grade	Prior	Survival
Group	Patient	Sex	(y)	pathology	grade	Initial therapy	(d)	recurrence	recurrence	treatments	(d)
1	1	Μ	69	GBM	4	Chemotherapy/XRT	76	GBM	4	1	81
	2	F	65	GBM	4	Chemotherapy/XRT	578	GBM	4	2	107
	3	Μ	50	AA	3	Chemotherapy/XRT	332	AA	3	2	59
	4	F	59	GBM	4	Chemotherapy/XRT	322	GBM	4	1	169
2	5	F	37	AMG	3	Chemotherapy/XRT	160	AMG	3	1	260
	6	Μ	64	GBM	4	Chemotherapy/XRT	178	GBM	4	1	234
	7	Μ	26	AA	3	Chemotherapy	462	GBM	4	3	364
	8	F	35	AA	3	XRT	2,804	GBM	4	3	318
	9	Μ	45	GBM	4	Chemotherapy/XRT	132	GBM	4	1	344
	10	F	54	GBM	4	Chemotherapy/XRT	91	GBM	4	1	343
3	11	F	61	GBM	4	Chemotherapy/XRT	719	GBM	4	2	400
	12	F	62	GBM	4	Chemotherapy/XRT	316	GBM	4	1	709
	13	F	47	GBM	4	Chemotherapy/XRT	160	GBM	4	1	374
	14	Μ	76	GBM	4	Chemotherapy/XRT	134	GBM	4	1	978
	15	F	40	GBM	4	Chemotherapy/XRT	462	GBM	4	1	366
						art of treatment in cu					

(Sum), using simple thresholding techniques. The effect of varying isocontours (55%-75%) was investigated. In this threshold method, both the shape and the size of the VOI varied between studies. In another analysis, a cube of $5 \times 5 \times 5$ voxels ($10 \times 10 \times 12 \text{ mm} = 1.2 \text{ cm}^3$) was generated and positioned around the voxel with maximum tumor uptake on the factor images (F2-cube) or summed frames (Sum-cube). In this cube analysis, the shape of the VOI is constant across studies, but the location is related to the maximum count in the tumor.

Compartmental Model

The kinetic model for ¹⁸F-FLT is based on the original work of Shields et al. (1). Our implementation for the brain was published elsewhere (6). Briefly, the model assumes a transport step of ¹⁸F-FLT from the vascular to the tissue space. There is no partitioning in the vascular space, and plasma and whole-blood activity concentrations are the same (16). In the tissue space, there is an exchangeable compartment (with intact ¹⁸F-FLT) and a compartment with phosphorylated ¹⁸F-FLT (mono-, di-, and triphosphate ¹⁸F-FLT). ¹⁸F-FLT is metabolized by the liver to ¹⁸F-FLTglucoronide, which is present only in the vascular space-that is, not transported into the brain-tissue compartment. Metabolites were corrected by subtracting a metabolite fraction from the vascular curve. This fraction was calculated as $0.42 \times (1 \exp[-0.029 \times \text{time}]$) as explained in our earlier publication (6). Kinetic modeling yields the 4 rate constants $(k_1, k_2, k_3, and k_4)$ between compartments. The blood volume fraction in tissue V_b was estimated as a fifth parameter. The influx rate constant K_i was calculated as $k_1k_3(k_2 + k_3)^{-1}$ and the volume of distribution V_d as $k_1(k_2 + k_3)^{-1}$. Metabolite correction of the blood curve to yield the input function was performed as described previously (6). A recovery coefficient of 0.7 was chosen to convert the image-based blood time-activity curve to the input function. Because the tumors, compared with the image resolution, are relatively large the recovery coefficient of the output function was set to unity.

Outcome Analysis

Of the 15 patients studied, 4 died within 4 mo (group 1 [G1]) and 6 died between 6 and 12 mo (group 2 [G2]). The remaining 5 survived for more than 1 y (group 3 [G3]). The overall survival time and time between initial diagnosis and current study treatment are given in Table 1. The median survival was 94 d for G1, 331 d for G2, and 542 d for G3. The patient with the complete response (no ¹⁸F-FLT uptake after treatment) had an overall survival of 1,054 d.

Statistical Analysis

Results are presented as the mean \pm SD. The Student *t* test was used to compare the kinetic parameters and ¹⁸F-FLT uptake. For small sample sizes (*n* < 10), the Wilcoxon Mann–Whitney *U* test

(2-sample rank test) was applied. Pearson correlation was used to study the relationship between subgroups.

RESULTS

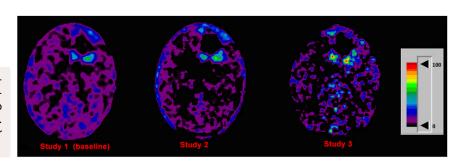
In Figure 1, for each of the 3 studies a representative factor slice is shown for a short-term survivor (G1). Figure 2 shows an example for a long-term survivor (G3).

Comparison of the varying thresholds of the tumor VOI delineated on the factor images (F2, threshold 55% - 75%) showed similar results for the parameter values of k_1 , K_i, V_b , early SUV (SUV_{early}), and late SUV (SUV_{late}). The correlation coefficients of these parameters between thresholding methods were high and ranged from 0.91 to 0.99 both for the entire set (n = 45) and for the 3 study sets (S1, S2, and S3; n = 15 each). Intermediate correlations (0.7 < r < 0.9) were found for k_2 and V_d , whereas k_3 and k_4 showed poor correlations (r < 0.6). Figure 3 shows the average results of all patients for the 3 sets of studies, using tumor VOIs delineated on summed frames (Sum-75%). Results in Figure 3 suggest a trend for k_1 , but no significantly different results were obtained for the absolute value of the parameters tested. The study after the first course of treatment (S2) generally shows lower parameter values, with a tendency to rebound at S3, except for k_4 . Considering the absolute values of the entire population, no statistical differences were found for the micro- or macroparameters between time points S1, S2, and S3 for any of the analysis methods (F2-55%-75%, Sum-75%, F2-cube, or Sum-cube).

Subsequently, the relative parameter change defined as (S1-S2)/S1 or (S1-S3)/S1 (i.e., fraction of change) was evaluated. In Figure 4, box plots are shown depicting the relative changes between S1 and S2 (left panel) and between S1 and S3 (right panel). Figure 4 shows narrow plots for k_1 at an early time point, with large variability later for G1 and G2. G3, on the other hand, shows consistently small variability between S1 and S3 for all 3 parameters. The variability of k_3 is large (except of G3 for $S1 \rightarrow S3$), signifying large fluctuations in phosphorylation rate during therapy.

Table 2 shows the population mean using the factor images with varying threshold and the summed images with 75% threshold. Inspection of the cube VOIs showed a similar pattern. None of the parameter values showed significant differences between the 4 methods analyzed:

FIGURE 1. Representative factor images from patient 2 of G1. Note increasing intensity (proportional to ¹⁸F-FLT influx rate) from S1 to S3. Images are displayed using same color scale (shown in arbitrary units).



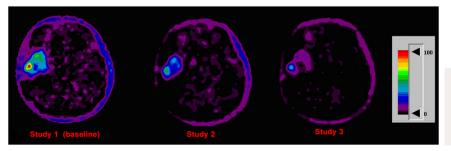


FIGURE 2. Representative factor images from patient 13 of G3. Note decreasing ¹⁸F-FLT influx rate from S1 to S3. Images are displayed using same color scale (in arbitrary units).

threshold VOI or cube VOI on factor or summed images. When we compared the tumor volume (in voxels), VOIs obtained with a threshold of 70% on the factor images (F2-70%) fitted best with a threshold of 75% on the summed images (Sum-75%), and linear regression analysis furnished a slope of 1.050 and intercept of 0.077 (r = 0.87). Figure 5 shows the scatter diagram for K_i for a variable VOI (F2-70%) versus a fixed VOI (Sum-cube).

The same techniques were used to measure the SUV, either threshold VOIs or a fixed cube, on the factor or summed images. ¹⁸F-FLT uptake reaches a maximum around 10–20 min (4,6); therefore, we calculated SUV_{early} (from 10 to 25 min) and SUV_{late} (from 50 to 60 min). The average value for SUV_{early} was 0.83 and for SUV_{late} 0.85. Both have a high correlation coefficient, compared with K_i (0.86 and 0.90, respectively).

A scatter plot of SUV_{late} versus K_i for the 3 studies is shown in Figure 6. The parameters at each time point, S1–

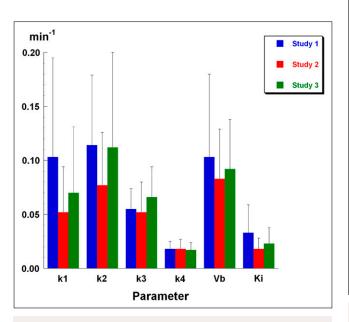


FIGURE 3. Average results per study (S1–S3) for rate constants (k_1 – k_4), V_b, and K_i. Bars denote SD. VOIs are delineated with threshold of 75% on summed images from 45 to 60 min. n = 15 for each column. None of shown parameters demonstrates significant differences for mean between S1, S2, or S3. Dimension of k_1 – k_4 and K_i is min⁻¹, whereas V_b is fraction (unitless).

S3, show an excellent correlation and similar slopes; none of the regression parameters was significantly different. The range of the parameters shows a decrease, both for K_i and SUV_{late} , when proceeding from the baseline study S1 to either S2 or S3.

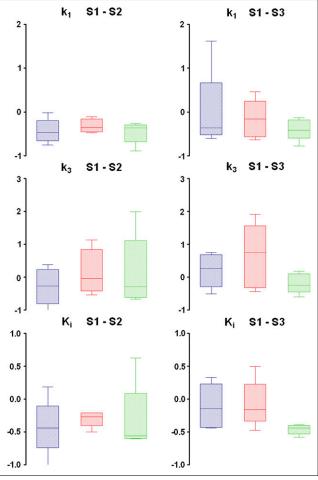


FIGURE 4. Box plots of relative changes for parameters k_1 , k_3 , and K_i . On left, relative changes between S1 and S2 are shown; on right, fractional changes between S1 and S3 are shown. Note small variability in k_1 for S1 \rightarrow S2 for all 3 groups. Variability for G3 (green) remains small for S1 \rightarrow S3 for all 3 parameters shown. For k_3 , wide variability in relative changes exists, except for G3 between baseline and end of therapy. blue = G1; red = G2; green = G3.

Image threshold	<i>k</i> ₁	k ₂	k ₃	k_4	V _b	¥	۷d	SUV _{early}	SUV _{late}
Factor image									
75%	0.071 ± 0.055	0.091 ± 0.068	0.047 ± 0.026	0.018 ± 0.007	0.086 ± 0.062	0.023 ± 0.017	0.52 ± 0.30	0.86 ± 0.44	0.88 ± 0.48
20%	0.069 ± 0.055	0.090 ± 0.064	0.046 ± 0.024	0.017 ± 0.007	0.086 ± 0.056	0.022 ± 0.017	0.51 ± 0.28	0.83 ± 0.41	0.85 ± 0.46
65%	0.064 ± 0.046	0.086 ± 0.040	0.048 ± 0.029	0.017 ± 0.007	0.086 ± 0.056	0.021 ± 0.016	0.47 ± 0.24	0.80 ± 0.39	0.81 ± 0.43
60%	0.062 ± 0.047	0.102 ± 0.084	0.062 ± 0.083	0.018 ± 0.007	0.086 ± 0.051	0.021 ± 0.015	0.44 ± 0.24	0.77 ± 0.38	0.78 ± 0.40
55%	0.060 ± 0.043	0.094 ± 0.047	0.053 ± 0.031	0.019 ± 0.008	0.085 ± 0.048	0.020 ± 0.014	0.41 ± 0.24	0.74 ± 0.36	0.74 ± 0.38
Cube	0.063 ± 0.051	0.090 ± 0.060	0.045 ± 0.028	0.017 ± 0.009	0.083 ± 0.055	0.019 ± 0.015	0.50 ± 0.31	0.77 ± 0.37	0.76 ± 0.38
Summed image									
75%	0.075 ± 0.070	0.101 ± 0.070	0.058 ± 0.026	0.018 ± 0.007	0.093 ± 0.057	0.025 ± 0.019	0.44 ± 0.25	0.85 ± 0.45	0.90 ± 0.49
Cube	0.068 ± 0.057	0.115 ± 0.074	0.061 ± 0.038	0.018 ± 0.009	0.087 ± 0.052	0.022 ± 0.016	0.39 ± 0.23	0.76 ± 0.38	0.78 ± 0.40
Donlad data for	Dooled data for all natients and all studies $(n = 45)$	studies $(n = 45)$							
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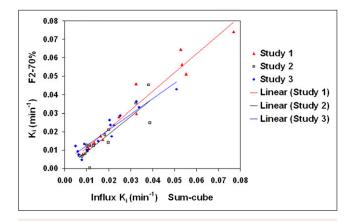


FIGURE 5. Comparison of K_i obtained with method F2-70% (*y*-axis) vs. Sum-cube (cube around maximum tumor voxel of frames summed from 45 to 60 min, *x*-axis).

Tumor Kinetics and Survival

The kinetic parameters were compared with clinical outcome, and patients were grouped arbitrarily into 6-mo survival periods. Four patients with glioblastoma died within 4 mo of the ¹⁸F-FLT study (G1), 6 patients survived 6–12 mo after therapy (G2), and the remaining 5 survived more than 1 y (G3). Detailed inspection of the absolute values of the various parameters did not reveal any statistically significant differences between patient groups or studies. However, analyzing the relative or fractional change between studies—that is, from S1 to S2 (Δ S1 \rightarrow S2) or from S1 to S3 (Δ S1 \rightarrow S3)—revealed several significant differences. For the factor-derived tumor VOI (F2-70%), significant differences were found for k_1 , K_i, SUV_{early}, and

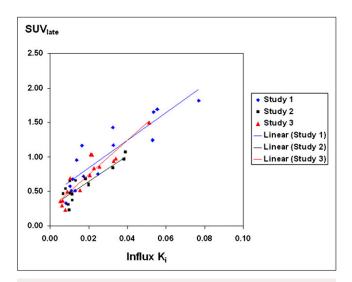


FIGURE 6. Scatter diagram of influx rate constant K_i vs. ¹⁸F-FLT uptake SUV_{late} for 3 studies analyzed with cube VOI around tumor on 45- to 60-min summed image. Correlation coefficients were 0.89 for S1 (P < 0.001), 0.88 for S2 (P < 0.001), and 0.92 for S3 (P < 0.001).

SUV_{late} in G2 for Δ S1 \rightarrow S2 and in G2 and G3 for both Δ S1 \rightarrow S2 and Δ S1 \rightarrow S3. For the Sum-75%, tumor VOI significant differences in K_i and SUV were found only for G3 for both $\Delta S1 \rightarrow S2$ and $\Delta S1 \rightarrow S3$. More significant differences were found for the cubes positioned around the maximum tumor uptake than for the variable VOIs. For F2cube, parameters V_b, K_i, V_d, and SUV showed a significant relative change in G3 for both $\Delta S1 \rightarrow S2$ and $\Delta S1 \rightarrow S3$. The same was true for Sum-cube. In addition, this latter cube VOI found significant differences in G2 for K_i and V_d both for $\Delta S1 \rightarrow S2$ and for $\Delta S1 \rightarrow S3$. Combining G1 and G2 (n = 10) and comparing these parameters with G3 (n =5) yielded the same statistically significant differences as before. Thus, most significant relative changes between studies were found with a cube of $5 \times 5 \times 5$ voxels around the tumor maximum. Using a variable tumor VOI reduced the number of significant changes, and in general, only relative changes in K_i and SUV remained statistically significant. SUV changes were always significantly different for G3, both for early and for late time points (Fig. 7).

Results with regard to SUV changes of more than 25%, which was the criterion used to define a responder in our earlier study (5), revealed that G3 had a 100% sensitivity both for $\Delta S1 \rightarrow S2$ and for $\Delta S1 \rightarrow S3$. These relative changes were statistically significant (P < 0.01).

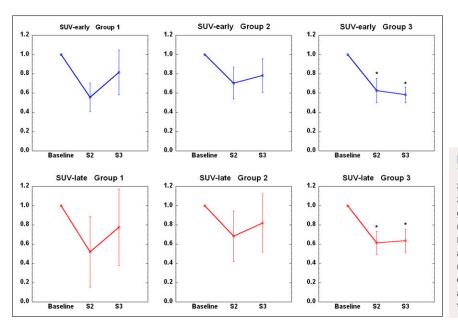
DISCUSSION

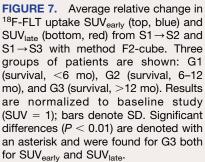
In this study, the kinetics of ¹⁸F-FLT in malignant brain tumors were investigated at baseline and during treatment and stratified to clinical outcome. Our analysis is completely image-based and virtually operator-independent. The input function was derived automatically from factor images of the transverse sinuses; no user interaction was required. The tumor VOIs were derived with thresholding techniques or defined as a cube around the voxel with maximum ¹⁸F-FLT uptake. In earlier work, we have demonstrated that factor analysis can reliably generate input and output functions (*17–22*). In another study, we have shown that a general 3-compartment model is adequate for describing the ¹⁸F-FLT kinetics of malignant brain tumors. As shown by Shields et al., the ¹⁸F-FLT model is a noninvasive way for evaluating cellular proliferation (2). ¹⁸F-FLT influx rate is a representation of thymidine flux or incorporation rate into DNA (*I*), as was demonstrated by Chen et al. (*4*,*5*) and by Ullrich et al., who found that ¹⁸F-FLT uptake correlated with in vivo proliferation rate in newly diagnosed high-grade gliomas (*11*).

In this study, we used a total dynamic acquisition time of 1 h. This acquisition time is 15 min shorter than that in our study of 2007 (16). The mathematic studies of Muzi et al. (16) have shown that the errors (defined as SEE/mean) increase when the acquisition duration is decreased from 120 to 60 min. For k_1 , the increase is from 13% to 15% and for K_i from 4% to 9%, which are still acceptable. The error for k_4 is 42%, whereas the other parameters have already more than 55% errors for the 2-h scan.

Another approach would be to set k_4 to zero to minimize errors in the parameter estimates. Although k_4 is small, it cannot be neglected as we have shown in earlier work (2,16). Muzi et al. came to a similar conclusion in their mathematic studies (16).

The subdivision into groups G1, G2, and G3 was based on survival—that is, average survival of 3.4, 10.2, and 18.5 mo, respectively. In general, the largest relative change in parameters was seen between baseline and 2 wk of therapy (Δ S1 \rightarrow S2), as shown in Figures 3 and 7. Thereafter, the trend reveals that the changes persist for G3, whereas G1 returns to normal. These changes between baseline and end of therapy monitoring (Δ S1 \rightarrow S3) revealed signif-





icant differences for V_b and V_d , and for K_i , SUV_{early} , and SUV_{late} , for the analysis methods using a cube (fixed-volume VOI). When thresholding techniques with variable-volume tumor VOIs were used, the number of significant changes decreased for K_i , SUV_{early} , and SUV_{late} and mainly for G3, suggesting that a persistently decreased ¹⁸F-FLT uptake in the tumor was a predictor for longer survival (Fig. 7).

Other reported studies (2,16) have shown that transport to the cells is fast and primarily determined by perfusion, whereas the phosphorylation reaction (k_3) is the ratelimiting step, determining uptake and retention of ¹⁸F-FLT in somatic tissue (16). This finding suggests that simple semiquantitative analysis (SUV) is sufficient for clinical applications, as was shown here (Fig. 6) for the different patient groups (G1, G2, and G3) and time points (S1, S2, and S3). This observation is not supported by the work of the Cologne group, who claims that the association between K_i and uptake is tumor grade-dependent and does not apply within 1 World Health Organization grade (11). In an earlier study from this group (8), a significant correlation was found between ¹⁸F-FLT uptake and both k_1 and k_3 , suggesting that the ¹⁸F-FLT uptake is related to a combination of factors. As shown in Figure 6, we found an excellent correlation between ¹⁸F-FLT uptake and influx rate K_i (containing both k_1 and k_3). Contrary to the studies of the Cologne group, we corrected the input function for metabolites. In addition, we enrolled only patients with recurrent glioma. The correlations between k_1 and SUV were always high (r > 0.7), but between k_3 and SUV, we found a correlation coefficient of 0.6-0.7 for the baseline study (S1) only. The later studies S2 and S3 did not show any relationship (r < 0.2).

Other investigators have used ¹¹C-methionine for evaluating response to therapy. Galldiks et al. (23) reported that ¹¹C-methionine was able to measure a response after 3 cycles of temozolomide, offering a method to measure biologic activity. Jacobs et al. (8) compared ¹¹C-methionine directly with ¹⁸F-FLT and indicated that the 2 tracers have an uptake that is not related. ¹¹C-methionine has a high sensitivity and specificity for glioma. These authors suggested that ¹⁸F-FLT helps differentiate low- from high-grade tumors and vields complementary information. Ceyssens et al. (24), on the other hand, argue that interindividual ¹¹C-methionine uptake does not permit individual grading. They conclude that the clinical use of ¹¹C-methionine is limited to recurrence detection, biopsy guidance, and radiation therapy target delineation. In our study, we found significant changes in ¹⁸F-FLT uptake and influx rate in a subgroup of long-term survivors after 1 course of therapy.

In the National Cancer Institute trial, ¹⁸F-FLT was able to separate recurrence from radionecrosis (*12*). Parameters K_i and k_3 reached statistical significance. These authors emphasized that simple blood-brain barrier breakdown might increase transport without affecting the influx (retention in the salvage pathway). This finding is not consistent with what we reported earlier (*5*). ¹⁸F-FLT uptake was measurable in stable patients with recurrent glioma, who had an intact blood-brain barrier, because they had no contrast enhancement on gadolinium-enhanced T1-weighted MRI (4).

Limitations of this study are the small numbers of patients per subgroup. On the other hand, all tumors were recurrent and high-grade, minimizing heterogeneity. Despite the small number of patients, robust effects were found for the relative change of parameters between S1 and S2 (Δ S1 \rightarrow S2) that were statistically significant. If the decreased uptake of ¹⁸F-FLT persists between S2 and S3, the patient is likely to have a good prognosis and will survive longer (G3 in Fig. 7).

CONCLUSION

¹⁸F-FLT kinetics were investigated during treatment of recurrent high-grade brain tumors. The standard 3-compartment model with corrections for V_b, metabolites, and partial volume was used to estimate kinetic parameters. Several analysis methods, including variable versus fixed tumor VOI (defined on factor vs. summed images), were compared. A cube with a fixed volume of 1.2 cm³ defined on the summed images from 45 to 60 min after ¹⁸F-FLT injection yielded the highest number of significant differences between the subgroups. The largest change of kinetic parameters occurred between baseline and 2 wk of treatment. Significant changes were found for V_b, K_i, V_d, SUV_{early}, and SUV_{late}. After stratification to overall survival, the patients with the best prognosis showed a change at 2 wk of treatment that persisted at 6 wk, whereas the patients with short survival times (G1) returned to baseline values at 6 wk. High correlations were found between ¹⁸F-FLT uptake in tumor (SUV) and influx rate constant (K_i), indicating that in clinical practice simple uptake measurements are sufficient for therapy monitoring and predicting short- and long-term survival.

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