#### **CLINICAL STUDY**



# Rate of change in maximum <sup>18</sup>F-FDOPA PET uptake and non-enhancing tumor volume predict malignant transformation and overall survival in low-grade gliomas

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#### Abstract

**Purpose** To examine whether the rate of change in maximum <sup>18</sup>F-FDOPA PET uptake and the rate of change in nonenhancing tumor volume could predict malignant transformation and residual overall survival (OS) in low grade glioma (LGG) patients who received serial <sup>18</sup>F-FDOPA PET and MRI scans.

**Methods** 27 LGG patients with  $\ge 2^{18}$ F-FDOPA PET and MRI scans between 2003 and 2016 were included. The rate of change in FLAIR volume (uL/day) and maximum normalized <sup>18</sup>F-FDOPA specific uptake value (nSUV<sub>max</sub>/month), were compared between histological and molecular subtypes. General linear models (GLMs) were used to integrate clinical information with MR-PET measurements to predict malignant transformation. Cox univariate and multivariable regression analyses were performed to identify imaging and clinical risk factors related to OS.

**Results** A GLM using patient age, treatment, the rate of change in FLAIR and <sup>18</sup>F-FDOPA nSUV<sub>max</sub> could predict malignant transformation with > 67% sensitivity and specificity (AUC = 0.7556, P = 0.0248). A significant association was observed between OS and continuous rates of change in PET uptake (HR = 1.0212, P = 0.0034). Cox multivariable analysis confirmed that continuous measures of the rate of change in PET uptake was an independent predictor of OS (HR = 1.0242, P = 0.0033); however, stratification of patients based on increasing or decreasing rate of change in FLAIR (HR = 2.220, P = 0.025), PET uptake (HR = 2.148, P = 0.0311), or both FLAIR and PET (HR = 2.354, P = 0.0135) predicted OS.

**Conclusions** The change in maximum normalized <sup>18</sup>F-FDOPA PET uptake, with or without clinical information and rate of change in tumor volume, may be useful for predicting the risk of malignant transformation and estimating residual survival in patients with LGG.

Keywords <sup>18</sup>F-FDOPA PET · Biomarker · Low grade glioma · MRI

## Introduction

The lack of effective treatments for patients with gliomas remains a significant health problem and highlights the need for novel and innovative therapeutic approaches. While maximum resection followed by chemoradiation is the standard of care for glioblastoma, there is no standard of care for

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low-grade gliomas (LGGs), and management of low-grade gliomas remains one of the most controversial areas in clinical neuro-oncology [1]. Patients with low-grade, IDH mutant gliomas tend to be young (mean age 35–40 [2]) and can have a relatively long survival (5–13 years from diagnosis [2–5]). Aggressive chemoradiation in these patients can result in significant morbidity (~98% of patients in RTOG-9802) [5], leading some oncologists to adopt a "watch and wait" strategy [6, 7] prior to aggressive surgery and/or chemoradiation [6, 8]. Since low-grade IDH mutant tumors grow very slowly and patients may have a relatively long survival, testing of new, often non-beneficial therapies, in large clinical trials based solely on survival outcomes can take more than 20 years (e.g. RTOG-9802) [3]. Thus, there

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is a significant demand for non-invasive imaging biomarkers that can quickly identify transformation of low-grade into higher grade gliomas, as well as predict overall survival in low-grade gliomas.

Although magnetic resonance imaging (MRI) is routinely used for serial evaluation of tumor burden in patients with LGGs [9], amino acid positron emission tomography (PET) is often used for confirming the presence of metabolically active tumors. Unlike normal brain tissue, tumor cells use amino acids for fuel and a variety of enzymatic processes. Thus, amino acid PET has higher tumor-to-background uptake (i.e. contrast to noise) compared with traditional FDG PET and better delineates hypermetabolic regions from less aggressive areas of the tumor [10-17]. While [<sup>18</sup>F]-fluoro-ethyl-tyrosine (<sup>18</sup>F-FET) PET is the most frequently used amino acid PET tracer in neuro-oncology due to its long half-life and simple synthesis [17], other tracers including methyl-L-[<sup>11</sup>C]-methionine (<sup>11</sup>C-MET) and 3,4-dihydroxy-6-[<sup>18</sup>F]-fluoro-L-phenylalanine (<sup>18</sup>F-FDOPA) provide similar information [18, 19], as they share a common transport mechanism (i.e. system L amino acid transporters like LAT1-4). <sup>18</sup>F-FDOPA PET is regularly used for diagnosis and management of movement disorders including Parkinson's disease [20, 21], so its dual use in neuro-oncology is economically attractive. <sup>18</sup>F-FDOPA PET combined with MRI has been shown to significantly improve diagnosis [22], identify the degree of malignancy, and differentiate pseudoprogression from true tumor growth [23–25]. Thus, we hypothesized serial changes in <sup>18</sup>F-FDOPA PET combined with changes in MRI may be useful as a unique tool for predicting both malignant transformation as well as predicting residual survival in patients with LGG. In the current study, we examined a series of 27 patients with LGGs who received two or more <sup>18</sup>F-FDOPA PET scans and corresponding MRI scans to determine whether a general model consisting of clinical information, the time rate of change in maximum <sup>18</sup>F-FDOPA PET uptake, and the time rate of change in non-enhancing tumor volume could be used to predict eventual malignant transformation and residual OS from the time of the last PET scan.

#### Methods

#### **Patient selection**

The UCLA Medical Center institutional review board approved this study (IRB IRB#15-000467). Patients from 2003 to 2016 with histology confirmed LGG (WHO II) who underwent at least two <sup>18</sup>F-FDOPA PET scans and MRI scans within 3 months of the PET scans, and had no surgical intervention between scans were eligible for inclusion. A total of 27 patients were identified as having two or more <sup>18</sup>F-FDOPA PET scans, MRI scans consisting of at least pre- and post-contrast T<sub>1</sub>-weighted images and either T<sub>2</sub>-weighted turbo spin echo or T<sub>2</sub>-weighted fluidattenuated inversion recovery (FLAIR) images with sufficient quality, lack of contrast enhancement, and adequate clinical information including survival. When available, isocitrate dehydrogenase (IDH) 1 mutational status, 1p19q codeletion status, and O<sup>6</sup>-methylguanine–DNA methyltransferase (MGMT) promoter methylation results were obtained from the patient's clinical chart. IDH mutation status was determined by genomic sequencing analysis using polymerase chain reaction (PCR) and/or through immunohistochemistry (IHC) of resected tumor tissue. 1p/19 co-deletion status was identified using fluorescence in situ hybridization (FISH) or Foundation medicine. MGMT methylation status was determined by PCR analysis. Malignant transformation was determined either by histology or radiological changes consistent with transformation (e.g. emergence of contrast enhancement suggestive of transformation to higher grade as indicated in the clinical radiology report). OS was measured from the time of the 2nd PET scan until death. Table 1 summarizes these patient demographics and molecular information.

## Magnetic resonance imaging acquisition and analysis

Anatomic MRI consisted of at least standard anatomic  $T_1$ -weighted pre- and post-contrast images (2D axial turbo spin echo with 3 mm slice thickness and no interslice gap or 3D inversion-prepared gradient echo images with 1–1.5 mm isotropic voxel size), and 2D axial  $T_2$ -weighted or  $T_2$ -weighted FLAIR images acquired at 3 mm slice thickness with no interslice gap. The presence or absence of nodular contrast enhancement (CE) on post-contrast  $T_1$ -weighted images within the lesion was determined by the official clinical radiology report. All 27 patients included in the current study lacked CE on post-contrast  $T_1$ -weighted images.

A single volume of interest (VOI) was segmented based on regions of  $T_2$  hyperintensity on  $T_2$ -weighted or  $T_2$ -weighted FLAIR images by two investigators (T.O. and J.Y.) and confirmed by another investigator (B.E.) using a semi-automatic procedure documented previously [26–28] and the Analysis of Functional NeuroImages (AFNI) software (NIMH Scientific and Statistical Computing Core; Bethesda, MD, USA). All volumes are reported in microliters ( $\mu$ L) and the time rate of change in tumor volume was quantified by calculating the change in volume divided by the time interval between the two MRI scans (units of  $\mu$ L/day).

WHO 2007												
Patient ID	Sex ,	Age	(Histo- logical) diagnosis	IDH status	1p19q status	MGMT status	Date of initial diag- nosis	Date of 1st PET scan	Date of 2nd PET scan	Date of PET scans	Therapy between PET scans	Transformed < 6 months after 2nd PET scan
-	W N	45	ΓO	MUT	Codel	М	1/16/2009	4/7/2009	5/2/2013	1486	None	Yes
2	Е	62	LA	ΜT	Unknown	Unknown	2/28/2012	10/26/2010	1/17/2012	448	None	Yes
3	7 M	49	OA	ΜT	Non-codel	М	1/15/2009	7/21/2009	6/8/2010	322	None	Yes
4	X	34	ΓO	MUT	Codel	М	7/2/2009	7/27/2010	3/6/2012	588	None	Yes
5	F	48	LA	Unknown	Unknown	Unknown	6/8/1995	4/28/2009	4/27/2010	364	TMZ	No
9	M	34	LA	MUT	Non-codel	U	6/12/1997	2/8/2011	2/6/2014	1094	TMZ	Yes
7	7 M	40	ΓO	MUT	Unknown	U	1/12/1998	9/12/2007	4/20/2010	951	CCNU	No
8	F	4	OA	MUT	Codel	М	4/12/2000	3/14/2007	8/31/2010	1266	None	Yes
6	X	31	ΓO	MUT	Codel	М	4/24/2007	10/30/2008	10/24/2011	1089	None	Yes
10	M	34	ΓO	MUT	Codel	М	11/14/2001	2/21/2007	6/19/2008	484	RT	Yes
11	X	50	OA	Unknown	Unknown	Unknown	8/22/2003	2/9/2010	11/2/2010	266	CCNU	No
12	M	54	LA	Unknown	Unknown	Unknown	6/10/2005	6/19/2008	12/8/2009	537	CCNU	Yes
13	N	36	LA	ΜT	Non-codel	U	3/25/2005	6/6/2007	6/21/2011	1111	None	No
14	Ш	31	OA	Unknown	Codel	Unknown	1/12/2009	9/23/2009	2/21/2012	881	None	No
15	Z	30	LO	ΜT	Non-codel	Unknown	11/4/1998	5/9/2007	11/14/2007	189	None	No
16	X	32	OA	MUT	Non-codel	Unknown	7/12/2007	2/6/2008	7/28/2009	538	TMZ	Yes
17	۲ W	48	OA	Unknown	Codel	Unknown	11/5/2007	3/31/2009	6/20/2013	1542	None	No
18	Z	35	OA	MUT	Codel	U	3/20/2008	9/9/2008	9/8/2009	364	TMZ	No
19	Ň	71	OA	Unknown	Unknown	Unknown	2/19/2001	2/24/2009	7/31/2009	157	TMZ	No
20	Ц.	34	OA	MUT	Non-codel	М	5/7/2008	10/21/2008	5/30/2013	1682	RT	No
21	N	33	LO	MUT	Codel	U	3/2/2009	3/2/2010	10/30/2014	1703	None	Yes
22	Z	39	LA	Unknown	Unknown	Unknown	7/10/2009	11/16/2010	6/21/2011	217	TMZ	No
23	Ц	24	LA	MUT	Unknown	М	2/10/2009	10/30/2009	3/1/2011	487	TMZ	No
$24^{a}$	X	24	LO	MUT	Codel	Unknown	11/24/2010	1/4/2011	1/5/2012	366	TMZ	No
25	ц.	53	LA	MUT	Non-codel	U	1/31/2011	9/5/2013	4/10/2014	217	TMZ+RT	No
26	с,	37	OA	MUT	Non-codel	U	11/21/2011	10/2/2012	10/24/2013	387	None	Yes
27	~ W	87	LA	WT	Non-codel	М	12/14/2011	4/3/2014	8/7/2014	126	None	No

Table 1 Patient demographics

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 $^{\rm a} \rm Cerebellar \, \rm IDH^{\rm MUT}$  , 1p19q codeleted patient with unknown MGMT status

## <sup>18</sup>F FDOPA PET image acquisition and analysis

<sup>18</sup>F-FDOPA PET scans were acquired using a high-resolution full-ring PET scanner (ECAT-HR; CTI/MIMVista) after subjects fasted for more than 4 h. <sup>18</sup>F-FDOPA was synthesized and injected intravenously according to previously established procedures [29, 30]. A CT scan was acquired prior to PET for attenuation correction. Three-dimensional <sup>18</sup>F-FDOPA emission data were acquired for a total of 30 min, and data were integrated between 10 and 30 min from injection to obtain 20-min static <sup>18</sup>F-FDOPA images following reconstruction. PET images were reconstructed using an ordered-subset expectation maximization (OSEM) iterative reconstruction algorithm consisting of six iterations with eight subsets [31, 32]. Lastly, a Gaussian filter with a full width at half maximum of 4 mm was applied. The resulting voxel sizes were 1.34 mm × 1.34 mm × 3 mm for <sup>18</sup>F-FDOPA PET images. These images were then fused to the MRI scans using a 12-degree of freedom affine registration transformation with a normalized mutual information cost function using FSL (FLIRT; FMRIB Software Library, Oxford, England; https://www.fmrib.ox.ac.uk/fsl/).

Standardized uptake volume (SUV) maps were calculated based on the radioactive activity divided by the decay-corrected injected dose per body mass [33]. The resulting SUV images were then normalized to the basal ganglia (nSUV) as described previously [13, 34, 35]. Maximum nSUV was quantified within T<sub>2</sub> hyperintense VOIs at each respective time point (nSUV<sub>max</sub>) and the rate of change in nSUV<sub>max</sub> divided by the time interval between the two PET scans (units of nSUV<sub>max</sub>/month).

## General linear model (GLM) integrating clinical and MR-PET imaging information

Multiple linear regression was used to create a series of general linear models (GLMs) integrating known clinical information and MR-PET imaging information for the purposes of predicting the risk of malignant transformation (WHO II  $\rightarrow$  III or II  $\rightarrow$  IV) within 6 months of the 2nd PET scan.

#### Statistical analyses

Statistical analysis was performed in MATLAB (R2018a; MathWorks, Natick, MA) or GraphPad Prism (Version 7.0c; GraphPad Software, La Jolla, CA). The receiver operating characteristic (ROC) curve was used to determine whether a combination of clinical information (e.g. age, MGMT status, on treatment during scans, etc.) combined with the rate of change in non-enhancing tumor volume and rate of change in nSUV<sub>max</sub> could discriminate between patients who demonstrated malignant transformation within 6 months of the 2nd PET scan and those who did not transform. The area under the curve (AUC), along with sensitivity and specificity for detecting malignant transformation, was evaluated as a measure of model performance.

The Shapiro–Wilk test was used to test for normally-distributed data. For normally-distributed data, Student's t-test or analysis of variance (ANOVA) was performed. For nonnormally-distributed data, a Mann–Whitney or Wilcoxon rank-sum analysis of medians was performed. Univariate and multivariate cox proportional hazards regression analysis was performed to investigate the association between OS and one or more predictor variables including clinical information such as age and treatment along with imaging metrics like rate of change in non-enhancing tumor volume and rate of change in nSUV<sub>max</sub>. Kaplan–Meier curves were used to display differences in OS, and the log-rank test was used to compare survival when appropriate. Statistical significance was defined as alpha=0.05 and no correction for multiple comparisons was performed.

## Results

A total of 27 patients met the inclusion criteria for the current study. The mean age for these patients was  $50.9 \pm 13.4$  s.d. years and 29% were female. The average time interval between serial <sup>18</sup>F-FDOPA PET exams was 1.3 years (range 125 days to 4.7 years) and matched MRI scans used in the current study was 1.3 years (range 125 days to 4.6 years). According to WHO 2007 histological criteria, 9 patients had low-grade diffuse astrocytomas (LA), 8 patients had low-grade oligodendrogliomas (LO), and 10 patients had low-grade mixed oligoastrocytomas (OA). A total of 15 patients were IDH mutant gliomas, 5 patients were IDH wild type, and 7 patients did not have known IDH status. Ten patients were 1p19q codeleted, 9 patients were non-codeleted, and 8 patients did not have known 1p19q status. A total of 9 patients were MGMT methylated, 7 were unmethylated, and the remaining 11 did not have MGMT status available. Approximately 48% of patients (13 of 27) were not on active therapy between the two PET and MRI exams and about 44% of patients (12 of 27) demonstrated histological or radiographic evidence of malignant transformation within 6 months of the 2nd MR-PET exam. Within 6 months of the 2nd PET scan, about 52% (14 of 27) patients accepted chemotherapy and/or radiation treatment. Of the patients who accepted therapy, about 43% (6 of 14) had malignant transformation.

Figure 1a–c illustrates MR-PET changes in a 24-yearold female patient with a residual survival of around 1 year after the 2nd PET scan. This patient had a WHO II, IDH1 mutant, MGMT methylated diffuse glioma with multi-parametric MR-PET imaging showing increasing tumor extent



**Fig. 1 a** Fluid attenuated inversion recover (FLAIR) MRI images, **b** normalized <sup>18</sup>F-FDOPA PET uptake, and **c** MR-PET fused images in a 24-year-old female patient with a WHO II, IDH1 mutant, MGMT methylated diffuse glioma with multi-parametric MR-PET imaging showing increasing tumor extent crossing the midline and an increasing focus of <sup>18</sup>F-FDOPA PET uptake over a period of around 1.3 years while on temozolomide. This patient expired approximately

1 year after the 2nd PET scan. **d** Fluid attenuated inversion recover (FLAIR) MRI images, **e** normalized <sup>18</sup>F-FDOPA PET uptake, and **f** MR-PET fused images in a 35-year-old male patient with a WHO II, IDH1 mutant, unmethylated, 1p19q codeleted oligodendroglioma with a small tumor burden and minimal PET uptake that is relatively stable over a year of evaluation while on temozolomide. This patient expired more than 9.7 years from the time of the last PET scan

crossing the midline and an increasing focus of <sup>18</sup>F-FDOPA PET uptake over a period of about 1.3 years while on temozolomide. Conversely, Fig. 1d–f illustrates a 35-year-old male patient with a residual survival of more than 9.7 years from the 2nd PET scan. This patient had a WHO II, IDH1 mutant, unmethylated, 1p19q codeleted oligodendroglioma with a small tumor burden and minimal PET uptake that was relatively stable over 1 year of evaluation while on temozolomide.

There was a weak but significant correlation between the rate of change in  $^{18}\mbox{F-FDOPA}$  PET  $\mbox{nSUV}_{\rm max}$  and the rate of change in non-enhancing tumor volume using FLAIR (Fig. 2a; Pearson's correlation,  $R^2 = 0.1769$ , P = 0.0289). We did not observe any notable differences in the rate of change in FLAIR volume or rate of change in PET uptake between tumor subtypes using the WHO 2007 histological criteria (Fig. 2b; Kruskal–Wallis, FLAIR volume, P = 0.7075, PET uptake, P = 0.5226). Similarly, we did not observe any significant differences between IDH mutants and wild type gliomas (Fig. 2c; FLAIR, P = 0.6262, PET, P = 0.9623), patients with 1p19q co-deletion and those without co-deletion (Fig. 2d; FLAIR, P=0.4668, PET, P=0.9154), patients with MGMT methylated or unmethylated tumors (Fig. 2e; PET, P = 0.6065, FLAIR, P = 0.2441). There was also no difference in the rate of change in FLAIR volume between patients on active treatment compared with those who were off active treatment during the time of the MRI and PET

scans (Fig. 2f; Mann–Whitney, P = 0.9051); however, there was a slight, insignificant trend toward lower rate of change in PET uptake in patients who were on active treatment compared to those off treatment (Fig. 2f; Mann–Whitney, P = 0.1546).

#### **Predicting malignant transformation**

Next, we tested whether the rate of change in FLAIR volume or rate of change in PET  $nSUV_{max}$  could be used to predict malignant transformation within 6 months of the 2nd MRI or PET examination. Results suggested no difference between the rate of change in FLAIR volume and PET uptake between transformers compared with those who did not transform within 6 months (Fig. 3a; Mann–Whitney, MRI, P = 0.6141, PET, P = 0.3473). Creation of a new metric from the linear combination of both the rate of change in FLAIR volume and the rate of change in PET uptake via a general linear model (GLM) of the form:

$$Risk = -4.53 \times 10^{-3} \cdot (\Delta MRI \ FLAIR \ Volume[\mu L/day]) + 3.38 \cdot (\Delta^{18} F-FDOPA \ Uptake[nSUV_{max}/month])$$

$$Intercept$$

$$+ 0.44$$
(1)



**Fig. 2** Comparison of rate of change in FLAIR volume and maximum normalized <sup>18</sup>F-FDOPA PET Uptake (nSUV<sub>max</sub>) in Histological and Molecular Subtypes of Low-Grade Gliomas (LGGs). **a** Correlation between rate of change in FLAIR volume and normalized maximum <sup>18</sup>F-FDOPA PET uptake in WHO II low grade gliomas ( $R^2$ =0.1769, P=0.0289). **b** Rate of change in FLAIR volume and <sup>18</sup>F-FDOPA nSUV<sub>max</sub> for WHO 2007 (histological) tumor subtypes. **c** Rate of change in FLAIR volume and <sup>18</sup>F-FDOPA nSUV<sub>max</sub> for isoci-

trate dehydrogenase (IDH)-1 mutant and wild type LGGs. **d** Rate of change in FLAIR volume and <sup>18</sup>F-FDOPA nSUV<sub>max</sub> for 1p19q codeleted and non-codeleted LGGs. **e** Rate of change in FLAIR volume and <sup>18</sup>F-FDOPA nSUV<sub>max</sub> for MGMT methylated and unmethylated LGGs. **f** Rate of change in FLAIR volume and <sup>18</sup>F-FDOPA nSUV<sub>max</sub> for LGG patients on active treatment compared with those not on active treated during the PET and MRI examinations

resulted in a trend toward detecting a difference between transformed and non-transformed LGGs (Fig. 3b; Mann–Whitney, P = 0.1712). Expanding this model to include patient age and whether the patient was on active treatment during the MRI and PET exams resulted in a GLM of the form:

$$Risk = -0.159 \times 10^{-4} \cdot (Age) + 0.3 \cdot (On Tx)$$

$$+ \underbrace{0.61}^{Intercept} -3.94 \times 10^{-3} \cdot (\Delta MRI FLAIRVolume[\mu L/day])$$

$$+ 1.94 (\Delta^{18}F-FDOPA Uptake[nSUV_{max}/month]) (2)$$

In the above linear model, "Age" is an integer in years and "On Tx" is a binomial with zero representing no treatment between serial PET scans and one for patients with treatment between scans. This model demonstrated a significant difference between transformed and non-transformed LGG patients (Fig. 3c; Mann–Whitney, P = 0.0396). Receiveroperator characteristic curves (ROCs) for predicting malignant transformation of LGGs within 6 months of the 2nd PET or MRI exam (Fig. 3d) suggested that the rate of change in FLAIR volume (Fig. 3e; AUC = 0.5611, P = 0.5914) and rate of change in PET uptake (Fig. 3e; AUC = 0.6111, P = 0.3291) could not predict transformation. Similarly, a GLM using only the rate of change in FLAIR and PET



**Fig. 3** Prediction of malignant transformation in LGGs using serial changes in MR-PET measurements. **a** Rate of change in FLAIR volume and <sup>18</sup>F-FDOPA nSUV<sub>max</sub> for LGGs undergoing malignant transformation within 6 months of the last MR-PET scan. **b** Relative risk of malignant transformation estimated using a general linear model (GLM) consisting of only the rate of change in FLAIR volume and the rate of change in maximum normalized <sup>18</sup>F-FDOPA PET uptake. **c** Relative risk of malignant transformation estimated using a GLM composed of patient age, whether they were on active treatment at the time of the MR-PET exams, the rate of change in FLAIR

volume, and the rate of change in maximum normalized <sup>18</sup>F-FDOPA PET uptake. **d** Receiver-operator characteristic (ROC) curves for **a** the rate of change in FLAIR volume, **b** the rate of change in maximum normalized <sup>18</sup>F-FDOPA PET uptake, **c** the GLM using MRI and PET only, and **d** the GLM using age, treatment, MRI, and PET. **e** Comparison between area under the ROC curves (AUC) showing no significant difference between each technique (P=0.6238), although the GLM using both clinical and imaging characteristics demonstrated AUCs significantly higher than chance. \*P < 0.05

uptake could not reliably predict transformation (Fig. 3e; AUC = 6722, P = 0.1304). However, a GLM using the rate of change in FLAIR volume and the rate of change in PET uptake combined with patient age and whether they were on active treatment could predict transformation (Fig. 3e; AUC = 0.7556, P = 0.0248) with approximately 87% sensitivity and 67% specificity using a risk threshold of 0.5907.

#### Survival analysis

Lastly, we performed Cox regression to estimate residual OS from the 2nd PET scan using clinical factors (age, on treatment between scans) and imaging predictor variables (rate of change in non-enhancing tumor volume and rate of change in PET nSUV<sub>max</sub>), both individually (univariate) and in a combined (multivariable) model. Univariate analysis suggested both static measurements of continuous FLAIR volume at the 2nd time point (HR=1.0178, P=0.0212) and continuous normalized PET uptake at the 2nd time point (HR=2.4050, P=0.0270) were associated with OS. In patients with IDH status, MGMT methylation status, or 1p19q codeletion status available, no significant association was observed with OS (IDH, P=0.9986; MGMT, P=1.000; 1p19q, P=0.9990). Univariate analysis demonstrated a trend toward increased

risk of death with increasing age (Table 2; Hazard Ratio (HR) = 1.027, Confidence Interval (CI) = [0.9939, 1.0613], P = 0.1107), a trend toward shorter OS with increasing rate of change FLAIR tumor volume (Table 2; HR = 1.0111, CI [0.9974, 1.0250], P = 0.1142), and a significantly shorter OS in patients with increasing rate of change in PET nSUV<sub>max</sub> (Table 2; HR = 1.0212, CI [1.0070, 1.0356], P = 0.0034). No real association was observed between treatment status and OS (Table 2; HR = 1.1013, CI [0.5037, 2.4078], P = 0.8090). When considering all four variables, Cox multivariable regression analysis suggested only the continuous measure of the time rate of change in PET nSUV<sub>max</sub> was an independent predictor of OS in LGGs (Table 2; HR = 1.0242, CI [1.0080, 1.0407], P = 0.0033).

Although continuous measures of the change in FLAIR volume was not a predictor of OS in LGGs, grouping patients into those with increasing versus decreasing FLAIR volume stratified patients into short and long-term residual OS (Fig. 4a; Log-rank, P = 0.025, HR = 2.220). Similarly, patients with an increasing PET uptake had worse residual OS (Fig. 4b; Log-rank P = 0.0311, HR = 2.148). Additionally, patients with increasing rate of both FLAIR volume and PET uptake had significantly shorter OS compared to patients with decreasing FLAIR volume or PET uptake

 Table 2
 Cox proportional-hazards model analysis

Characteristic (unit)	Overall survival (univariate)		Overall survival (multivariable)	
	<i>P</i> -value	Hazard ratio (95% CI)	<i>P</i> -value	Hazard ratio (95% CI)
Age (years)	0.1107	1.0270 (0.9939–1.0613)	0.1525	1.0246 (0.9910–1.0594)
Treatment status (on vs. off)	0.809	1.1013 (0.5037-2.4078)	0.188	1.7590 (0.7588-4.0779)
Rate of change in FLAIR volume (uL/day)	0.1142	1.0111 (0.9974–1.0250)	0.9304	1.0007 (0.9853-1.0163)
Rate of change in normalized PET SUV ( $\times 1000$ nSUV <sub>max</sub> /month)	0.0034**	1.0212 (1.0070–1.0356)	0.0033**	1.0242 (1.0080–1.0407)

*Note* Continuous values of the rate of change in PET uptake were scaled by  $\times 1000$  for regression  $^{**}P < 0.01$ 



**Fig. 4** Kaplan–Meier plots showing significant differences in residual OS from the 2nd PET scan in patients with **a** a positive or negative rate of change in FLAIR volume (HR=2.220, P=0.025); **b** a positive or negative rate of change in PET uptake (HR=2.148, P=0.0311); and **c** patients with a positive rate of change in both FLAIR vol-

ume and PET uptake compared to those with either a negative rate in FLAIR volume or PET uptake changes (HR=2.354, P=0.0135). Black=positive rate of change or high risk. Gray=negative rate of change or low risk

(Fig. 4c; Log-rank, P = 0.0135, HR = 2.354), suggesting the combination of both change in FLAIR volume and PET uptake may be meaningful for identifying patients at highest risk.

## Discussion

<sup>18</sup>F-FDOPA PET imaging and MRI are routinely used in the clinic to monitor treatment response in patients with LGGs. However, most patients only receive a single PET and corresponding MRI scan when clinical questions arise. While <sup>18</sup>F-FDOPA PET combined with MRI has been shown to improve diagnosis [22], identify the degree of malignancy, and differentiate pseudoprogression from true tumor growth [23–25], few studies have explored the value of serial PET and MRI scans in predicting response assessment or long-term outcome [34, 36, 37] and no studies to our knowledge have examined the use of both serial PET and MRI to predict malignant transformation or survival in LGGs.

Although the current study did not find any significant differences between histological or molecular subtypes of LGGs, results suggest a linear combination of imaging features with or without clinical factors may be beneficial for predicting malignant transformation and long-term survival. In particular, a simple linear combination containing age, treatment, the rate of change in T2 hyperintense lesion volume on FLAIR and rate of change in <sup>18</sup>F-FDOPA PET nSUV<sub>max</sub> were able to predict subsequent malignant transformation with > 67% sensitivity and specificity. When controlling for clinical factors and tumor growth dynamics on MRI, the change in maximum tumor metabolism using PET appears significantly associated with survival. This suggests metabolic information may compliment traditional measures of tumor growth and may be important for understanding aspects of long-term disease control in LGGs.

Although we found an association between static measurements of FLAIR volume and PET uptake with survival in LGGs, the rate of change in PET uptake appeared to provide additional information in terms of predicting transformation and survival. This was not completely surprising, considering there was a correlation between nSUV<sub>max</sub> on the second PET scan and change in <sup>18</sup>F-FDOPA uptake (data not shown, P=0.0023,  $R^2=0.3160$ ); however, there were some patients with favorable outcome that had high PET uptake on the 2nd PET scan but exhibited decreasing PET uptake over time. For LGGs, serial <sup>18</sup>F-FDOPA PET and MRI exams may provide a more accurate illustration of tumor progression and prediction of patient outcome.

The current study had a number of important limitations that should be addressed. First, the current study was a small, retrospective study involving only 27 patients over a period of more than 10 years. Because this was a retrospective study spanning a long period of time, there was a significant amount of missing clinical and molecular information on these limited number of patients. Therefore, the lack of significant findings between histological and molecular subtypes may be premature and may require further investigation. Also, patients did not have their PET and MRI scans at a specific controlled interval, but instead had PET scans based on clinical or radiographic (e.g. MRI) changes, which may have biased the results because these patients represent a subset of patients with specific interval changes requiring evaluation. Additionally, due to this being a retrospective study, we were unable to acquire multiple baseline PET scans to correct for noise and resulting intrasubject scan variations. A more systematic study in a large cohort of LGG patients with PET and MRI scans at controlled intervals, even when lacking clinical or radiographic changes, is warranted.

#### Conclusion

The rate of change in non-enhancing tumor volume on FLAIR combined with the rate of change in maximum normalized <sup>18</sup>F-FDOPA PET uptake, with or without clinical information including age and treatment, may be useful for predicting the risk of malignant transformation and residual survival in patients with LGGs.

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#### **Compliance with ethical standards**

**Conflict of interest** All authors declare that they have no conflict of interest.

**Ethical approval** All procedures performed in studies involving human participants were in accordance with the ethical standards of the institutional and/or national research committee and with the 1964 Helsinki declaration and its later amendments or comparable ethical standards. Written informed consent was obtained from all individual participants included in the study.

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