# Hybrid MR-PET of Brain Tumours using Amino Acid PET and Chemical Exchange Saturation Transfer MRI

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## Abstract

#### Purpose

PET using radiolabelled amino acids has become a promising tool in the diagnostics of gliomas and brain metastasis. Currently, amide proton transfer (APT) chemical exchange saturation transfer (CEST) MR imaging is evaluated for brain tumour imaging. In this hybrid MR-PET study, we compared in brain tumours with 3D data derived from APT-CEST MRI and amino acid PET using O-(2-<sup>18</sup>F-fluoroethyl)-L-tyrosine (<sup>18</sup>F-FET).

#### **Methods**

Eight patients with gliomas were investigated simultaneously with <sup>18</sup>F-FET PET and APT-CEST MRI using a 3T MR-BrainPET scanner. CEST imaging was based on a steady-state approach using a  $B_1$  average power of 1µT.  $B_0$  field inhomogeneities were corrected and parametric images of magnetisation transfer ratio asymmetry (MTR<sub>asym</sub>) and differences to the extrapolated semi-solid magnetisation transfer reference method, APT# and nuclear Overhauser effect (NOE#), were calculated. Statistical analysis of the tumour-to-brain ratio of the CEST data was performed against PET data using the non-parametric Wilcoxon test.

#### Results

A tumour-to-brain ratio derived from APT# and <sup>18</sup>F-FET presented no significant differences and no correlation was found between APT# and <sup>18</sup>F-FET PET data. Distance between local hot spots APT# and <sup>18</sup>F-FET were different (average  $20 \pm 13$  mm, range 4 - 45 mm).

#### Conclusion

For the first time CEST images were compared with <sup>18</sup>F-FET in a simultaneous MR-PET measurement. Imaging findings derived from<sup>18</sup>F-FET PET and APT CEST MRI seems to provide different biological information. The validation of imaging findings by histological confirmation is necessary, ideally using stereotactic biopsy.

Keywords: CEST; APT; <sup>18</sup>F-FET PET; hybrid MR-PET; glioma;

## Introduction

Diagnostics of brain tumours remains unsatisfactory despite high-resolution morphological imaging with conventional magnetic resonance imaging (MRI), in particular in pre-treated patients [1]. Multiple imaging modalities may be combined in order to improve diagnostic accuracy, such as advanced MRI and positron emission tomography (PET). In brain tumours, information about amino acid transport has become an important additional tool to standard MRI and has shown potential to define biological tumour volume for treatment planning, treatment monitoring and recurrence evaluation [2]–[8]. Several amino acid tracers are available, such as L-[methyl-<sup>11</sup>C]-methionine (<sup>11</sup>C-MET), O-(2-<sup>18</sup>F- fluoroethyl)-L-tyrosine (<sup>18</sup>F-FET) or 3,4-dihydroxy- 6-[<sup>18</sup>F]fluoro-L-phenylalanine (<sup>18</sup>F-FDOPA). Fluorine-18 labelled radiotracers are preferred due to the logistical advantages of the longer half-life. Furthermore, <sup>18</sup>F-FET exhibits different tracer kinetics depending on the tumour grade when compared to <sup>18</sup>F-FDOPA or <sup>11</sup>C-MET [4].

Chemical exchange saturation transfer (CEST) is an MRI technique that enables the indirect detection of metabolites using the MR free water signal [10]. CEST is based on the phenomenon of transferring saturation from a solute pool (metabolites) to the water pool resulting in contrast that is dependent on metabolite concentration and its proton exchange rate [11], [12]. Repeating the measurements at different saturation frequencies allows the acquisition of the Z-spectrum, where the contrast of multiple metabolites is encoded. This technique has many applications such as measurements of glycogen (GlycoCEST) [13], glutamate (GluCEST) [14], glucose (GlucoCEST) [15] and amide proton transfer (APT) [16]. Moreover, CEST contrast is also pH dependent [16]–[19]. APT-CEST, which provides information at 3.5 ppm downfield from water, has been shown to allow differentiation between low- and high-grade gliomas [20], [21]. In addition, APT-CEST has also shown its potential to distinguish between recurrent tumour and radiation necrosis [22], [23]. Furthermore, the signals from the upfield nuclear Overhauser effect (NOE) at -3.5 ppm were also presented as a unique contrast in gliomas at 7 T [24]–[26].

The introduction of hybrid MR-PET scanners promotes the synergy between both modalities by allowing the simultaneous acquisition of human MRI and PET data [27]. However, to the best of our knowledge, no study in the literature has compared amino acid PET of brain tumours with APT-CEST using simultaneous MR-PET. Initial attempts to compare CEST signals from amine and <sup>18</sup>F-FDOPA using non-simultaneous measurements have recently been reported [28], [29]. This is the first study to investigate simultaneous MR-PET brain tumour imaging using APT-CEST and <sup>18</sup>F-FET PET.

Here, we present a methodology to obtain volumetric APT-CEST and <sup>18</sup>F-FET PET to study brain tumours using simultaneous MR-PET. We compare CEST imaging with the more clinically established <sup>18</sup>F-FET PET to explore similarities between both imaging modalities.

## **Materials and Methods**

#### Measurement details

#### **Subjects**

Eight patients with suspected cerebral glioma were included in this feasibility study (four females, four males; median age: 56 years; age range: 20 - 69 years). All subjects presented increased <sup>18</sup>F-FET uptake in the tumour, defined as <sup>18</sup>F-FET uptake of at least 1.6 times the uptake of healthy brain tissue of the same subject [30]. Seven subjects presented a single <sup>18</sup>F-FET positive region and one subject presented two <sup>18</sup>F-FET positive regions. Detailed information on the patient group is given in Table 1. The university ethics committee and the relevant federal authorities approved the study. All subjects provided prior written, informed consent for their participation in the study.

#### Instrumentation

Measurements were performed in a hybrid 3 T MR-BrainPET scanner (MAGNETOM Trio, Siemens Healthineers, Erlangen, Germany) equipped with a dedicated BrainPET with a spatial resolution of approximately 3 mm full-width at half maximum (FWHM) in the centre [31]. The BrainPET is inserted in a MAGNETOM Trio 3 T MR machine. A dedicated MR head coil was used, which is composed of an outer birdcage coil for transmission and an inner eight-channel receive array coil for signal reception.

#### **MR** Acquisitions

CEST measurements were performed before MR contrast agent injection in the clinical protocol. The clinical protocol includes a  $T_1$ -weighted magnetization-prepared rapid gradient echo (MPRAGE) sequence and a  $T_2$ -weighted fluid-attenuated inversion recovery sequence (FLAIR). A contrast-enhanced MPRAGE sequence (CE-MPRAGE) was conducted after intravenous injection of the contrast agent gadoteric acid (Dotarem, Guerbet, France).

CEST acquisition was based on the steady-state approach proposed by Ref. [32]. In our implementation, a segmented 3D EPI readout with three shots per partition was used [33]. In order to suppress the fat signal, a rectangular excitation pulse with a duration of 2.4 ms was used as proposed by Ref. [34]. The TR interval, consisting of a100 ms Gaussian RF pulse, crusher gradients and a partial EPI readout (echo train length = 26ms), amounted to TR = 136ms and an average B<sub>1</sub> power of 1  $\mu$ T

[35]. Further sequence parameters were: flip angle = 11 degrees; TE/readout duration = 12.5/26 ms; matrix size =  $80 \times 80 \times 48$ ; voxel size =  $3.0 \times 3.0 \times 3.0 \text{ mm}^3$ ; sagittal slice orientation (readout superior-inferior); slice partial Fourier = 6/8; phase partial Fourier = 7/8 and slice oversampling of 25%. The Z-spectrum was unevenly sampled at 69 different frequencies ranging from 14 to -14 ppm (±14.0, ±13.5, ±13.0, ±12.5, ±12.0, ±11.5, ±11.0, ±10.5, ±10.0, ±9.5, ±9.0, ±8.5, ±8.0, ±7.5, ±7.0, ±5.0, ±4.0, ±3.8, ±3.7, ±3.6, ±3.5, ±3.4, ±3.3, ±3.2, ±3.1, ±3.0, ±2.9, ±2.8, ±2.7, ±2.0, ±1.0, ±0.5, ±0.2, ±0.1, 0.0) and additionally at 300 ppm for data normalisation. Each image was acquired in 18 seconds (135 TRs; k-space centre acquired with the 45<sup>th</sup> TR). Seamless acquisition of all images amounted to total measurement time of 21 minutes.

In addition, a 3D multi-echo gradient echo (GRE) sequence was also acquired in order to compute B<sub>0</sub> maps. Measurement details were as follows: FA = 7°; TR/TE<sub>1</sub> = 50/2.32 ms;  $\Delta$ TE = 2.85 ms; 12 echoes; monopolar readout; matrix size =  $128 \times 104 \times 72$ ; voxel size =  $1.7 \times 1.7 \times 2.0$  mm<sup>3</sup>.

#### **PET** Acquisition

<sup>18</sup>F-FET was produced via nucleophilic <sup>18</sup>F-fluorination with a radiochemical purity of greater than 98 %, a specific radioactivity greater than 200 GBq/µmol and a radiochemical yield of about 60 %, as described previously [36]. According to the German guidelines for brain tumour imaging using labelled amino acid analogue, all patients fasted for at least 4 h before the PET acquisition [37]. Dynamic PET scans from 0 to 50 min post-injection were performed after intravenous injection of approximately 250 MBq of <sup>18</sup>F-FET. Attenuation correction was performed by a template-based approach using MRI [38]. PET data were reconstructed using an OP-OSEM3D algorithm with 4 subsets and 32 iterations where normalization and attenuation, scatter, dead time and random corrections were included [39]. The reconstructed dynamic data set consisted of 16 time frames (5 × 1 min; 5 × 3 min; 6 × 5 min), image matrix of  $256 \times 256 \times 153$  and voxels of  $1.25 \times 1.25 \times 1.25 \text{ mm}^3$ .

#### Data Processing

All data processing and analyses were performed using in-house developed MATLAB code (The MathWorks, Inc., Natick, Massachusetts, United States), Python code (Python Software Foundation, version 2.7, available at <u>http://www.python.org</u>) and the software PMOD (Version 3.7, PMOD Technologies Ltd., Switzerland). Data co-registration was performed using the FLIRT tool

from FSL [40], [41] using as reference the APT-CEST image at 3.5 ppm [42].

#### Field maps

Field maps for  $B_0$  correction of the CEST data were calculated from the multi-echo GRE data. Images were masked based on the magnitude data and field maps were unwrapped using the algorithm from Ref. [43].

#### **CEST Data Processing**

 $B_0$  field maps were used to correct the Z-spectrum for field shifts using a smoothing spline interpolation on a voxel-by-voxel basis. After correction, the data from the offsets between 14 and 7 ppm were fitted to the Henkelman's two-pool magnetisation transfer (MT) model as suggested for the extrapolated semi-solid magnetisation transfer reference (EMR) method [43]–[46].

In order to calculate the CEST-weighted images two approaches were considered: magnetisation transfer ratio asymmetry ( $MTR_{asym}$ ) and differences to the EMR (named using #). The  $MTR_{asym}$  at 3.3 to 3.7 ppm was used, while EMR for the APT# and NOE# were calculated at 3.3 to 3.7 ppm and at -3.7 to -3.3 ppm, respectively.

#### Data Analysis

A volume-of-interest (VOI) analysis was performed to compare the different CEST metrics with <sup>18</sup>F-FET PET data. Three VOIs were considered to evaluate the tumour region. One PET-based tumour VOI (FET-VOI) was defined as follows: a spherical background VOI was positioned in the hemisphere contralateral to the lesion in healthy brain tissue (volume of background VOI, 14.1 ml; 7220 voxels, 30 mm diameter). The tumour VOI was defined in the summed <sup>18</sup>F-FET PET image from 20 to 40 min post-injection by a 3D auto-contouring process using a tumour-to-brain ratio (TBR) of 1.6 or more. This metric is widely used in <sup>18</sup>F-FET PET and the cut-off value is based on a biopsy controlled study of cerebral gliomas [7]-[9], [30], [47]. Two MR-based VOIs were manually defined: FLAIR-VOI, based on the image hyperintensities; CE-MPRAGE-VOI, based on contrast-enhanced regions (Fig. 1).

Statistical analysis of the TBR of the CEST metrics was performed against <sup>18</sup>F-FET data using a non-parametric Wilcoxon test using SPSS (IBM Corp. Released 2013. IBM SPSS Statistics, Version 22.0. Armonk, NY: IBM Corp). P-values of 0.05 or less were considered statistically significant. In addition, the hot spots in the tumour area were localized in <sup>18</sup>F-FET PET and APT# and their distance calculated.

## Results

Figure 2 presents Z-spectra for a pixel in the tumour and for a pixel in normal appearing white matter. The signal at 3.5 ppm is increased in the tumour voxel compared to the healthy tissue pixel. This is also visible in the difference to its corresponding EMR curve. In contrast, no peak was found in the downfield area of the Z-spectrum in healthy appearing brain tissue. In this figure it can also be seen that the EMR follows the conventional macromolecular pool for wide frequencies offsets in the range of 14 to 7 ppm.

Figures 1, 3 and 4 show the clinical and parametric images obtained for two subjects. Figure 1 shows a case with a tumour where the contrast-enhanced region did not correspond with the metabolically active area in PET. The contrast-enhanced regions in the  $T_1$ -weighted MRI also did not correspond with the FLAIR hyperintensities. In the MTR<sub>asym</sub> (Fig. 1g), the tumour showed an increased value compared to healthy appearing brain parenchyma. In the downfield APT# images, an increased intensity was observed in the metabolically active tumour area as defined by <sup>18</sup>F-FET PET while the NOE# showed no differences between the tumour area and healthy appearing brain tissue. In Fig. 3, a case of a tumour with a necrotic area is presented. Here, the clinical data showed a contrast enhancement around the necrotic area, and a hyperintense signal in the FLAIR image. In the PET data the metabolically active area surrounds the necrotic region. For the CEST data, increased MTR<sub>asym</sub> values were observed in the necrotic region, while in the EMR parametric images (NOE# and APT#) the necrotic region presented values near zero, suggesting a reduced effect of direct water and MT contributions. In Fig. 4, a case of a tumour in the frontal lobe is presented. Here, the clinical data showed a contrast enhancement in the tumour region, and a hyperintense signal in the FLAIR image. In the PET data the metabolically active area has a larger extension when compared to the standard clinical data. For the CEST data, increased MTR<sub>asym</sub> values were observed in the tumour region. In the EMR parametric images, NOE# presented no signal changes while APT# presented an increased signal in the tumour region with a different pattern than <sup>18</sup>F-FET. This example also illustrates the advantage of using 3D CEST where the full tumour extent could be evaluated.

Table 2 presents the VOI analysis for the various metrics. The different CEST contrasts provided different information when compared with <sup>18</sup>F-FET PET (p<0.05, Wilcoxon test). TBR varied across the different VOI definitions. Additionally, the location of the maxima of APT# and <sup>18</sup>F-FET PET differ (Tab. 3).

## Discussion

The aim of this study was to compare different CEST metrics with amino acid PET in cerebral gliomas using the clinically established PET tracer, <sup>18</sup>F-FET, to examine the similarities / differences between the two methods. Each modality has already shown its diagnostic potential, for example, for tumour grading or discrimination of radiation necrosis against tumour recurrence [9], [47], [48]. While <sup>18</sup>F-FET PET achieves a moderate accuracy of 77 % for tumour grading [49], APT has shown more promising data with an area under curve in ROC analysis of 81 – 88 % [22]. To the best of our knowledge, no study in the literature has ever compared <sup>18</sup>F-FET PET and APT. Thus, for the first time this study investigates the relation between <sup>18</sup>F-FET PET and CEST related metrics in brain tumours using simultaneous MR-PET at 3T and three-dimensional imaging.

In principle, <sup>18</sup>F-FET PET and APT are based on different biochemical and physiological mechanisms. <sup>18</sup>F-FET PET provides information about the expression of large amino acid transporter systems (L-type amino acid transporters 1 and 2 (LAT-1 and LAT-2)) that are more strongly expressed in brain tumours than in healthy tissue [50]. Consequently, an increased <sup>18</sup>F-FET signal is not necessarily related to increased protein synthesis or increased protein content but to an increased expression of LAT transporters. In contrast, the APT signal is related to the peptides and mobile proteins and not to the total protein content [51], as well as local modification of pH [52]. Our results present no relation between APT-CEST and <sup>18</sup>F-FET PET in the tumour region (Tab. 2), suggesting that the expression of LAT transporters and transfer of peptides and mobile proteins/change of pH is not strictly coupled. In addition, hot-spot locations in <sup>18</sup>F-FET and APT differ (Tab. 3), which suggests that both modalities could be used to further understand the increasingly important issue of tumour heterogeneity.

Several CEST metrics (MTR<sub>asym</sub>, NOE# and APT#) were calculated in this work using an irradiation power of 1  $\mu$ T and compared with <sup>18</sup>F-FET PET. In the literature, most studies performed at 3 T use an effective power of 2  $\mu$ T or higher [16]-[18], [22], [23], [48], [53]. The value of 2  $\mu$ T is chosen in order to obtain a null contrast in the normal appearing matter using MTR<sub>asym</sub> [53], once the contribution of NOE and APT signals cancel out in healthy tissue. However, at higher powers the influence of the macromolecular pool also increases [54], resulting in APT-weighted images (MTR<sub>asym</sub>) including MT contributions. Thus, in this work an irradiation power of 1  $\mu$ T was used in order to obtain nearly the maximum APT contrast and reduce the influence of the macromolecular pool that

was further corrected using the EMR model. Moreover, this methodology proved to be stable regarding the applied power and allowed NOE# contrast to be additionally obtained [46]. The origin of this signal is still under discussion [24]–[26] and it has also been suggested that it could be linked to protein folding and structure [56]. Moreover, studies investigating NOE effects are often performed at higher field strengths (>3 T) [24]-[26], which also influences the contrast.

In addition to various CEST metrics, different radiotracers are also available to measure amino-acid transport such as <sup>18</sup>F-FDOPA. This radiotracer was used non-simultaneously with two-dimensional CEST in Refs [28], [29]. In these studies, the authors explored the CEST signal from amines at 3 ppm as a measure of pH using a saturation power of 6  $\mu$ T. They found elevated MTR<sub>asym</sub> signal in regions with elevated uptake of <sup>18</sup>F-FDOPA but it was not quantified.

Tissue pH and T<sub>1</sub> are a potential confounding factors in the present work, as it influences the CEST contrast [16], [24], [52], [57], [58]. Regarding pH, an increased CEST signal can be associated to an alkalinisation of the tissue or an increase protein concentration that could not be disentangled in this work. However, imaging pH exclusively is beyond the scope of the presented work that focused on evaluating different CEST metrics in the metabolically active tumour area defined by <sup>18</sup>F-FET PET. Regarding T<sub>1</sub>, this is also a factor that can also influence CEST contrast [16], [24]. In the tumour region, the increasing of T<sub>1</sub> occurs usually in simultaneous with an increase of water content that may reduce the influence of this factor in CEST signal [16]. Further studies are required to understand the relationship between CEST, <sup>18</sup>F-FET, pH, tumour extent, and hot-spot locations.

Defining tumour extent is an important issue for patient assessment and treatment planning. With conventional MRI, this is usually based on the contrast-enhanced regions of T<sub>1</sub> weighted images [59]–[61]. Contrast-enhancement in MRI, however, indicates areas with a disrupted blood-brain barrier (BBB) that are, on the one hand, not specific for tumour tissue and, on the other hand, may not detect the full extent of the tumour. In contrast, <sup>18</sup>F-FET uptake is independent of the BBB integrity [62] and allows a more specific delineation of tumour extent than contrast-enhanced MRI [30]. Thus, the results cannot be compared directly and often show discrepancies. For example, in the case of low-grade gliomas where the BBB is mostly intact, often no contrast enhancement is observed, while there is still an increased uptake of <sup>18</sup>F-FET [30]. Thus, the present study also considers the <sup>18</sup>F-FET positive regions to define the metabolically tumours region. This approach is supported by the biopsy-controlled study of Ref. [30]. In the CEST literature, PET data is usually unavailable for comparison and consequently different studies present different strategies to delineate the tumour region that may or may not be accurate. In Ref. [18] ROIs were placed over the CE-MPRAGE or in FLAIR data when no contrast enhancement was found. In Ref. [21], [23] circular ROIs were only placed in the MTR<sub>asym</sub> with higher intensities in order to avoid the effect of averaging over larger ROIs. However, those tumour delineations do not necessarily reflect the solid and metabolic active area of the tumours and VOI values can change across different VOIs (Tab. 2). By using <sup>18</sup>F-FET PET, the solid and metabolically active tumour tissue can be defined with high reliability; this presents an enormous advantage of the present study compared to previous studies.

Three-dimensional acquisition is required in clinical applications in order to obtain whole tumour imaging. In CEST studies, two-dimensional imaging is often used to avoid long acquisition times [24], [26], [63]. In this study a steady-state approach based on segmented 3D EPI was used [32] to obtain fast full brain coverage. It was possible to acquire a single image in 18 seconds, without the use of parallel imaging, due to hardware limitation (reduced number of receive channels). In our approach, we sampled the Z-spectrum with 70 points in 21 min, which is still rather long but this could easily be reduced to about 10 mins by employing parallel imaging. Nevertheless, if a simplified sampling scheme with less sampling offsets is considered, as suggested in [21], the measurement time could be reduced to 6 min.

The CEST imaging technique benefits from higher static magnetic fields (> 3 T) due to the increased spectral dispersion and increased signal-to-noise ratio. For that reason, the majority of the studies are performed at 7 T [20], [24], [32], [64]. However, no human MR-PET machine is currently available at 7T.

The obtained results need to be confirmed in a larger and diversified population of patients. Nevertheless, the presented methodology can be combined with high field methods [65], pointing towards human MR-PET experiments up to 9.4 T to allow further insights into tumour metabolism [66].

### Conclusions

For the first time CEST imaging was compared with amino acid PET using <sup>18</sup>F-FET in a simultaneous MR-PET measurement. Based on the well-founded assumption that high <sup>18</sup>F-FET uptake reflects metabolically active tumour our data suggests that APT CEST provides substantially different

information.

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## Compliance with ethical standards

## **Conflict of interest**

The 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.

## **Informed consent**

Informed written consent was obtained from all individual participants included in the study.

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## Legends

**Figure 1.** A representative patient case (glioblastoma) with FLAIR (a), CE-MPRAGE (b), <sup>18</sup>F-FET PET (c), MTR<sub>asym</sub> (g), NOE# (h) and APT# (j) images. The respective VOI definitions are shown in subfigures (d-f). PET shows prominent <sup>18</sup>F-FET uptake in the tumour area in the frontal lobe (c) while APT# (j) exhibits minor differences compared with the normal brain. When compared with the normal brain tissue, NOE# (i) presented reduced contrast and MTR<sub>asym</sub> (g) presented high contrast.

**Figure 2.** Z-Spectrum and respective EMR fitting for a tumour pixel (a) and healthy tissue pixel (b). The difference between both curves are presented in the bottom plots from where the NOE# and APT# can be calculated (c-d).

**Figure 3.** A patient case with a tumour (glioblastoma) with a necrotic region. Clinical FLAIR (a), CE-MPRAGE (b), <sup>18</sup>F-FET PET (c) and CEST metrics (MTR<sub>asym</sub> (d), NOE# (e), APT# (f)) are also presented. The tumour shows a cystic lesion in MRI (a,b) and corresponding strong signal in MTR<sub>asym</sub> (d), which do not correspond to the active tumour depicted in <sup>18</sup>F-FET PET. NOE# (e) and APT# (f)) also presented different pattern when compared with <sup>18</sup>F-FET which could not depict the rim of the lesion.

**Figure 4.** A patient case with a tumour (oligoastrocytoma) in the frontal lobe. Clinical FLAIR (a), CE-MPRAGE (b), <sup>18</sup>F-FET PET (c) and CEST metrics (MTR<sub>asym</sub> (d), NOE# (e), APT# (f)) are also presented. <sup>18</sup>F-FET PET (c) detects clearly the metabolically active tumour tissue in suspicious areas in CE-MPRAGE (b) and FLAIR (a). CEST metrics (d-f) present different information when compared with <sup>18</sup>F-FET.











## Tables

Patient ID	Diagnosis	Treatment	
1	Glioblastoma/Reactive changes	Radio/chemotherapy	
2 ( lesion 1)	Glioblastoma Radio/chemotherapy		
2 (lesion 2)	Glioblastoma	Radio/chemotherapy	
3	Anaplastic Ependymoma	Surgery, radiotherapy	
4	Glioblastoma Surgery, radio/chemotl		
5	Anaplastic Astrocytoma	None	
6	Glioblastomas None		
7	Glioblastoma Surgery, radio/chemother		
8	Anaplastic Oligoastrocytoma Surgery, chemothera		

 Table 1. Diagnostic and treatment data for all the patients.

**Table 2.** VOIs analysis of CEST data. The MR-related metrics and <sup>18</sup>F-FET PET encode differentinformation (p<0.05, Wilkoxon test). Ranges: FLAIR-VOI (APT#:2.93 – 4.92; NOE#:4.04–5.66;</td>MTR<sub>asym</sub>: -1.89 – -0.75), CE-MPRAGE-VOI (APT#:2.43 – 4.93; NOE#: 3.17–5.12; MTR<sub>asym</sub>: -0.75 –0.72) and: FET-VOI (APT#:3.25 – 5.93; NOE#: 3.86–6.04; MTR<sub>asym</sub>: -1.85 – 0.65)

	FLAIR-VOI		CE-MPRAGE-VOI		FET-VOI	
	Value	TBR	Value	TBR	Value	TBR
APT#	3.71±0.71	0.86±0.21	3.89±0.91	0.99±0.29	4.44±0.73	1.13±0.24
NOE#	4.98±0.54	1.30±1.0	4.13±0.80	0.71±0.15	5.13±0.68	0.89±0.12
MTR <sub>asym</sub>	-1.27±0.39	0.76±0.39	-0.24±0.50	0.16±0.27	-0.70±0.41	0.41±0.41

	Hotspot distances (mm)
Patient ID	APT# vs. <sup>18</sup> F-FET
1	23.4
2(1)	45.0
2(2)	27.0
3	6.61
4	27.17
5	4.24
6	4.24
7	21.63
8	24.19

Table 3. Distances between local hotspots defined by APT# and  $^{18}\mbox{F-FET}$  PET.