#### **ORIGINAL ARTICLE**



# Multiparametric PET/MR imaging biomarkers are associated with overall survival in patients with pancreatic cancer

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#### Abstract

**Purpose** To correlate the overall survival (OS) with the imaging biomarkers of dynamic contrast-enhanced magnetic resonance imaging (DCE-MRI), diffusion-weighted imaging (DWI), magnetic resonance spectroscopy, and glucose metabolic activity derived from integrated fluorine 18 fluorodeoxyglucose positron emission tomography (<sup>18</sup>F–FDG PET)/MRI in patients with pancreatic cancer.

**Methods** This prospective study was approved by the institutional review board and informed consent was obtained from all participants. Sixty-three consecutive patients (mean age,  $62.7 \pm 12$  y; men/women, 40/23) with pancreatic cancer underwent PET/MRI before treatment. The imaging biomarkers were comprised of DCE-MRI parameters (*peak, IAUC*<sub>60</sub>, *K*<sup>trans</sup>, *k*<sub>ep</sub>, *v*<sub>e</sub>), the minimum apparent diffusion coefficient (ADC<sub>min</sub>), choline level, standardized uptake values, metabolic tumor volume, and total lesion glycolysis (TLG) of the tumors. The relationships between these imaging biomarkers with OS were evaluated with the Kaplan-Meier and Cox proportional hazard models.

**Results** Seventeen (27%) patients received curative surgery, with the median follow-up duration being 638 days. Univariate analysis showed that patients at a low TNM stage ( $\leq 3$ , P = 0.041), high *peak* (P = 0.006), high ADC<sub>min</sub> (P = 0.002) and low TLG (P = 0.01) had better OS. Moreover, high TLG/*peak* ratio was associated with poor OS (P = 0.016). Multivariate analysis indicated that ADC<sub>min</sub> (P = 0.011) and TLG/*peak* ratio (P = 0.006) were independent predictors of OS after adjustment for age, gender, tumor size, and TNM stage. The TLG/*peak* ratio was an independent predictor of OS in a subgroup of patients who did not receive curative surgery (P = 0.013).

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**Conclusion** The flow-metabolism mismatch reflected by the TLG/*peak* ratio may better predict OS than other imaging biomarkers from PET/MRI in pancreatic cancer patients.

 $\label{eq:keywords} \ \text{PET/MR} \cdot \text{Pancreatic cancer} \cdot \text{Dynamic contrast-enhanced} \ \text{MRI} \cdot \text{Diffusion} \ \text{weighted} \ \text{MRI} \cdot \text{MR} \ \text{spectroscopy} \cdot \text{Overall} \ \text{survival}$ 

### Introduction

Pancreatic ductal adenocarcinoma (PDAC) is the fourth-most common cause of cancer death [1]. Currently, radical resection of the primary tumor and regional lymphadenectomy are the only potentially curative treatments. Despite recent improvements in adjuvant chemotherapy, patients with PDAC continue to have dismal prognoses: the 5-year overall survival (OS) rate is about 8% [2]. Therefore, identification of novel prognostic biomarkers in patients with PDAC is crucial and can improve the selection of patients for adjuvant therapy and thus, survival outcome.

Because noninvasive imaging biomarkers are used extensively in cancer research [3], positron emission tomography/ magnetic resonance imaging (PET/MRI) has shown great potential in tumor diagnosis, treatment planning, surveillance, and follow-up in oncology patients [4–7]. Multiparametric PET/MRI offers various imaging biomarkers for tumor characteristics including blood flow by dynamic contrastenhanced MRI (DCE-MRI), cellularity by diffusionweighted imaging (DWI), metabolites by magnetic resonance spectroscopy (MRS), and metabolism by fluorine 18 fluorodeoxyglucose (<sup>18</sup>F–FDG) in a single examination [6]. Furthermore, these imaging biomarkers can provide novel insights about tumor characteristics and aggressiveness in PDAC.

Among these imaging biomarkers, associations between tumor blood flow-metabolism mismatch and adverse tumor biology have been illustrated in various cancers [8]. Using <sup>15</sup>O–H<sub>2</sub>O and <sup>18</sup>F–FDG PET/computed tomography (CT) to quantify blood flow and metabolic activity in pancreatic tumors, a previous study found that high tumor metabolism and low blood flow were linked with an aggressive cancer phenotype and potentially reflect treatment resistance [9]. The flowmetabolism mismatch indicates tumor adaptation to hypoxic stress, and this adaptation could be associated with poorer patient outcomes for PDAC [10–12]. DCE-MRI is a noninvasive technique that can detect tumor blood flow, vascularity, and permeability in vivo and has been used as a functional imaging tool in an integrated PET/MRI examination [6]. Thus, we assume that the correlation between flow and metabolism from DCE-MRI and PET may illustrate tumor aggressiveness and predict outcome in PDAC.

In addition, a positive correlation between high cellularity, reflected by a low apparent diffusion coefficient (ADC) from DWI, and high metabolic activity from <sup>18</sup>F–FDG PET has

been investigated in many cancers [13-15]. For example, it has recently been reported that in PDAC and periampullary cancer, advanced-stage tumors had lower minimum ADC (ADC<sub>min</sub>) values and higher glucose metabolic activities than did early-stage tumors. And, a high ratio of metabolic tumor volume (MTV)/ADC<sub>min</sub> was independently associated with shorter progression-free survival (PFS) [16]. Thus, the combined information from multiparametric PET/MRI biomarkers can provide more accurate prognostication about the survival of patients with PDAC.

Our hypothesis is that imaging biomarkers from multiparametric PET/MRI before treatment can be used to predict OS in patients with PDAC. The aim of this study was to correlate the OS with the imaging biomarkers of DCE-MRI, DWI, MRS, and glucose metabolic activity derived from integrated PET/MRI in patients with pancreatic cancer.

### **Materials and methods**

#### Patients

This prospective study was approved by the institutional review board in our hospital, and informed consent was obtained from all patients. From August 2014 to July 2016, 106 consecutive patients were admitted for examination by PET/ MRI before treatment. Inclusion criteria were suspicious pancreatic cancer on sonography or CT and no local or systemic treatments. Exclusion criteria were pregnancy and contraindications for 3-T MRI. All patients were pathology-proven by surgery, endoscopic ultrasound, or CT-guided biopsy. Tumor size, histological grade, and lymph node metastasis were recorded. Finally, 63 patients (mean age,  $62.7 \pm 12$  y; range, 34– 83 y; 40 men, 23 women) with PDAC comprised our study population after excluding 43 patients with chronic pancreatitis (n = 15) or other malignancy (n = 28) (Fig. 1). All patients tolerated this examination. Patient characteristics are presented in Table 1.

Some patients participated in another study [16] that analyzed associations between imaging biomarkers from DWI, MRS, and PET with tumor grade, clinical stage, and PFS in patients with PDAC or periampullary cancer. In this study, we focus primarily on associations between imaging biomarkers and OS in patients with PDAC, and we include imaging





biomarkers from DCE-MRI. Neither DCE-MRI nor OS data have been reported previously.

### DCE-MRI

### **PET/MRI protocol**

Whole-body PET/MRI was performed using an integrated PET/MRI system (Biograph mMR; Siemens Healthcare, Erlangen, Germany). Approximately 60 min after injection of FDG (5.18 MBq/kg) and bladder evacuation, PET was performed from the head to the mid-thighs in five bed positions (acquisition time, 4 min/position) with the patient in a supine arm-down position. Images were reconstructed using an ordered-subsets expectation maximization iterative algorithm (two iterations, 21 subsets) with a 5-mm post-reconstruction Gaussian filter and a  $172 \times 172$  image matrix. Attenuation correction of the PET data was performed using a 4-tissue-class (air, lung, fat, and soft tissue) segmented attenuation map acquired using a 2-point Dixon MRI sequence. Simultaneous MRI with coronal and axial T2-weighted half-Fourier single-shot turbo spin-echo sequences was performed while acquiring PET data at each bed position. Beyond MRI acquisition protocol, PET information was strictly linked to the employed beta-emitting tracer.

Following simultaneous PET and MRI acquisition, axial T1-weighted imaging with a 2-D fast low-angle shot and axial T2-weighted imaging with fat suppression were performed on the abdomen (Table 2).

DCE-MRI was performed using a 3-D gradient-echo sequence with isotropic resolution in three dimensions. A T1 map of the pancreas was initially created using various flip angles (5°, 10°, 15°, and 20°) to establish baseline T1 values. A MR pulse sequence with a T1-weighted volumetric interpolated breath-hold examination (3.17-s temporal resolution per volume) was started 50 s prior to injection of gadobutrol (0.05 mmol/kg, Gadovist® 1.0; Schering, Berlin, Germany) at a flow of 3 mL/s. Baseline images were acquired 10 s (10 phases) before initiating contrast agent injection. All patients were instructed to suspend breathing for as long as possible, then breathe slowly and smoothly during imaging. Imaging time was 3 min, 57 s to obtain 90 volumes (1260 images).

### DWI

Pulse sequences included an axial diffusion-weighted echoplanar imaging sequence with free breathing. The ADC map was calculated using a monoexponential function (b-values, 0, 600, and 1000 s/mm<sup>2</sup>) [16].

#### MRS

Single-voxel MRS data were acquired using a point-resolved selective spectroscopy sequence and standard parameters

|   | o o patients with patieneatic cancers |
|---|---------------------------------------|
| Age (years) *                             | 62.7±12 (34~83)                       |
| Gender (M/F) *                            | 40/23                                 |
| Tumor size (cm) *                         | $3.4 \pm 1.5$                         |
| Glucose (mg/dL) *                         | $117 \pm 41$                          |
| FDG (MBq) *                               | $371\pm86.6$                          |
| CEA (ng/ml) *                             | 6.2 (0.4~5783)                        |
| CA19–9 (U/ml) *                           | 221 (1.5~26,000)                      |
| Tumor location <sup>†</sup>               |                                       |
| Head                                      | 33 (52)                               |
| Neck                                      | 8 (13)                                |
| Body                                      | 16 (25)                               |
| Tail                                      | 6 (10)                                |
| Surgery method <sup>†</sup> $(n = 21)$    |                                       |
| Whipple operation                         | 14                                    |
| Distal pancreatectomy                     | 3                                     |
| Exploratory laparotomy and biopsy         | 2                                     |
| Bypass and biopsy                         | 2                                     |
| TNM staging <sup>†</sup>                  |                                       |
| Ι   | 4 (6)                                 |
| П   | 13 (21)                               |
| III                                       | 9 (14)                                |
| IV  | 37 (59)                               |
| Histology grade <sup>†</sup> ( $n = 22$ ) |                                       |
| Well-differentiated                       | 3                                     |
| Moderately differentiated                 | 9                                     |
| Poorly differentiated                     | 10                                    |

 Table 1
 Clinical information of the 63 patients with pancreatic cancers

\*Data are means  $\pm$  standard deviations or median values, with range in parentheses

†Data are numbers of patients, with percentages in parentheses

FDG: Fluorine 18 fluorodeoxyglucose

CEA: serum carcinoembryonic antigen

CA19–9: carbohydrate antigen 19–9

(1000/30 ms TR/TE;  $90^{\circ}$  flip angle) during free-breathing. Automated optimization of transmitter pulse power, localized shimming, gradient tuning, and water suppression were used. Data were acquired at a spectral bandwidth of 1200 Hz, and 200 signals were averaged for each water-suppressed spectrum [16].

#### Post-contrast multiphasic and whole-body MRI

Finally, multiphasic (arterial, portovenous, and equilibrium phases) T1-weighted imaging of the pancreas was performed using gadobutrol (0.05 mmol/kg), and contrast-enhanced imaging of the whole body was performed using an axial volumetric interpolated breath-hold examination sequence. Magnetic resonance cholangiopancreatography was also performed. The total length of the PET/MRI examination was approximately 60 min per participant (Figs. 2 and 3).

#### Image analysis

One radiologist (B.B.C., with 12 years of experience in abdominal imaging) measured the DCE-MRI, DWI, MRS, and PET parameters of the pancreatic tumor. One nuclear medicine physician (M.F.C., with 14 years of experience) evaluated the lymph nodes and distant metastases on PET/MRI. The radiologist and nuclear medicine physician independently performed their parts on the workstation.

The TNM stages (American Joint Committee on Cancer, 7th edition) of study patients were determined by a multidisciplinary team for pancreatic cancer at our hospital. In those who did not receive curative surgery, stage was determined by biopsy and all available image results.

The DCE-MRI data were analyzed using a commercial software tool (MIStars; Apollo Medical Imaging, Melbourne, Australia) [17]. Motion correction was performed using a 2-D rigid body registration. Two semi-quantitative parameters (peak and IAUC<sub>60</sub>) were derived. The former was defined as (SImax - SIbase)/SIbase \* 100, where SIbase represents the average baseline signal before the arrival of contrast agent in the arteries, and SI<sub>max</sub> represents the maximum value of the first pass of the time-signal intensity curve. The latter was the area under the curve for the first 60 s after contrast material arrived in the region of interest (ROI). Three quantitative parameters (volume transfer constant  $[K^{trans}]$ , reflux constant  $[k_{ep}]$ , and extra-vascular extra-cellular volume fraction  $[v_e]$ ) were derived using a bicompartmental model [18] and nonlinear fitting of the individual timeintensity curve. Regions of interest were manually drawn on the abdominal aorta at the level of the celiac axis for the arterial input function. These parameters were automatically calculated pixel-by-pixel using a constrained nonlinear leastsquares fitting algorithm with adjustable delay time [17]. The mean ROI size was  $4.9 \pm 4.6$  cm<sup>2</sup> (range, 0.6–25.6 cm<sup>2</sup>).

To measure the ADC, ROIs were manually drawn on the ADC map along the contour of the tumor on the single slice containing the largest area of the tumor. The lowest ADC value in an ROI,  $ADC_{min}$ , represented the greatest tumor cellularity. The mean ROI size was  $8.7 \pm 6.6$  cm<sup>2</sup> (range, 0.6–28.4 cm<sup>2</sup>).

In MRS, post-processing, zero filling, Fourier transformation, and phase correction were implemented. The volumes of interest were  $3.6 \pm 1.3$  cm<sup>3</sup> (range, 3.4-12.5 cm<sup>3</sup>). The area of the choline peak at 3.2 ppm was calculated using Syngovia software. MRS was successful in 56 patients.

Relevant PET-related parameters were  $SUV_{max}$ , which reflects the maximum standardized uptake value (adjusted for body weight);  $SUV_{mean}$ , which reflects average SUVs; MTV, expressed as the tumor volume with FDG uptake, which was segmented using a fixed-percentage threshold method at 50% of the  $SUV_{max}$ ; and total lesion glycolysis (TLG), representing the product of MTV and the average SUVs of the included

**Table 2** Parameters of PET/MRIexamination

| Sequence   | Repetition<br>time<br>(msec) | Echo<br>time<br>(msec) | Flip<br>angle<br>(°) | Matrix      | Field of<br>view<br>(mm) | Slice<br>thickness<br>/gap (mm) | NEX | Acquisition<br>time<br>(min:sec) |
|--|------------------------------|------------------------|----------------------|-------------|--------------------------|---------------------------------|-----|----------------------------------|
| During PET   |                              |                        |                      |             |                          |                                 |     |                                  |
| Coronal T2WI   | 1240                         | 87                     | 90                   | 384×<br>258 | 450×<br>302              | 6/0.6                           | 1   | 2:37                             |
| Axial T2WI   | 1000                         | 85                     | 90                   | 320×<br>260 | 350×<br>284              | 6/0                             | 1   | 3:14                             |
| After PET  |                              |                        |                      |             |                          |                                 |     |                                  |
| Axial T1WI   | 138                          | 2.5                    | 70                   | 640×<br>240 | 300×<br>253              | 5/0                             | 1   | 0:42                             |
| Axial T2WI   | 2000                         | 97                     | 121                  | 640×<br>496 | 300×<br>233              | 5/0                             | 1   | 1:30                             |
| DCE-MRI  | 4.7                          | 2.5                    | 15                   | 192×<br>168 | 340×<br>298              | 5/0                             | 1   | 3:57                             |
| DWI  | 7500                         | 83                     | 90                   | 312×<br>216 | 350×<br>242              | 5                               | 3   | 3:00                             |
| MRCP   | 1500                         | 85                     | 90                   | 512×<br>496 | 260×<br>252              | 4/0                             | 1   | 1:18                             |
| Dynamic axial<br>T1WI<br>post                            | 3.3                          | 1.2                    | 15                   | 265×<br>224 | 320×<br>280              | 1.8/0                           | 1   | 0:22 (3 sets)                    |
| contrast<br>Whole-body<br>axial T1WI<br>post<br>contrast | 3.3                          | 1.2                    | 15                   | 256×<br>192 | 370×<br>298              | 3/0                             | 1   | 2:24                             |

T2WI: T2-weighted images; FLAIR: fluid-attenuated inversion recovery images; T1WI: T1-weighted images; DCE-MRI: dynamic contrast-enhanced magnetic resonance image; DWI: diffusion weighted imaging; MRCP: magnetic resonance cholangiopancreatography; NEX: Number of excitations



**Fig. 2** Images of a 56-year-old man with pancreatic head cancer (arrow) (T4N1M1). (a) Pre-contrast T1-weighted MR (b) PET (c) fused PET/ venous phase CE T1-weighted VIBE MR. (d–f) DCE-MRI color maps:

(d)  $IAUC_{60}$  (e) peak (f)  $K^{trans}$  (g) diffusion-weighted image (b = 1000) (h) ADC map (i) MR spectroscopy. This patient underwent chemotherapy and overall survival was 199 days



**Fig. 3** Images of a 76-year-old man with pancreatic head cancer (arrow) (T1N1M0). (a) Venous phase CE T1-weighted MR (b) axial PET (c) fused PET/venous phase CE T1-weighted MR. (d–f) DCE-MRI color maps: (d)  $IAUC_{60}$  (e) peak (f)  $K^{trans}$  (g) diffusion-weighted image (b = 1000) (h)

Table 3Pearson correlationsamong PET/MR imaging

biomarkers

coronal PET (i) MR cholangiopancreatography shows marked dilatation of common bile duct and main pancreatic duct. This patient underwent Whipple operation and overall survival was 638 days

| Parameters          |                   | peak    | IAUC <sub>60</sub> | K <sup>trans</sup> | k <sub>ep</sub> | Ve     |
|---------------------|-------------------|---------|--------------------|--------------------|-----------------|--------|
| ADC <sub>min</sub>  | Correlation (r)   | 0.258   | -0.002             | 0.032              | 0.179           | -0.138 |
|                     | P value           | 0.024*  | 0.495              | 0.405              | 0.087           | 0.148  |
| Choline             | Correlation $(r)$ | 0.053   | 0.008              | 0.368              | 0.357           | 0.245  |
|                     | P value           | 0.350   | 0.475              | 0.003**            | 0.003**         | 0.034* |
| SUV <sub>max</sub>  | Correlation (r)   | -0.003  | 0.183              | -0.298             | -0.139          | -0.235 |
|                     | P value           | 0.491   | 0.076              | 0.009**            | 0.139           | 0.032* |
| SUV <sub>mean</sub> | Correlation (r)   | -0.02   | 0.02               | -0.076             | 0.137           | -0.108 |
|                     | P value           | 0.437   | 0.438              | 0.277              | 0.142           | 0.2    |
| MTV                 | Correlation (r)   | -0.331  | -0.33              | -0.085             | -0.094          | -0.094 |
|                     | P value           | 0.004** | 0.004**            | 0.254              | 0.232           | 0.232  |
| TLG                 | Correlation (r)   | -0.302  | -0.213             | -0.1               | 0.007           | -0.120 |
|                     | P value           | 0.008** | 0.047*             | 0.219              | 0.478           | 0.174  |
| Size                | Correlation (r)   | -0.395  | -0.081             | -0.270             | -0.174          | -0.209 |
|                     | P value           | 0.001** | 0.265              | 0.016*             | 0.086           | 0.05*  |
|                     |                   |         |                    |                    |                 |        |

 $peak = (maximum signal - baseline signal) / baseline signal; IAUC_{60} = initial area under the curve for the first 60 s after contrast material arrival; K<sup>trans</sup> = forward volume transfer constant; <math>k_{ep}$  = reverse volume transfer constant;  $v_e$  = extravascular extracellular space volume per unit volume of tissue; ADC<sub>min</sub> = minimum apparent diffusion coefficient; SUV<sub>max</sub> = maximum standardized uptake value; SUV<sub>mean</sub> = mean standardized uptake value; MTV = metabolic tumor volume; TLG = total lesion glycolysis

\* indicates P < 0.05

\*\* indicates P < 0.01

| Table 4 | Differences in DCE-MRI | parameters and TLG/ | <i>peak</i> ratio relative to | pathological | grades and TNM stag | ze |
|---------|------------------------|---------------------|-------------------------------|--------------|---------------------|----|
|         |                        |                     |                               |              | /                   |    |

| Parameter                        |                          | peak (%)    | <i>IAUC<sub>60</sub></i> (/100, sec) | $K^{trans}$ (min <sup>-1</sup> / 1000) | $k_{ep} ({\rm min}^{-1}/1000)$ | $v_e(\%/10)$ | TLG/<br>peak |
|----------------------------------|--------------------------|-------------|--------------------------------------|--|--------------------------------|--------------|--------------|
| Tumor differentiation $(n = 22)$ | Well-moderate $(n = 12)$ | 333 ± 42    | $402\pm216$                          | $61\pm95$                              | $417\pm491$                    | 106±<br>128  | 9.2±7.6      |
|                                  | Poor $(n = 10)$          | $327\pm96$  | $426\pm241$                          | $14\pm19$                              | $201\pm254$                    | $32\pm31$    | $12\pm11$    |
|                                  | P value                  | 0.582       | 0.771                                | 0.093                                  | 0.159                          | 0.059        | 0.674        |
| T-stage ( $N = 63$ )             | $\leq 3 \ (n = 29)$      | $325\pm71$  | $436\pm223$                          | $43\pm70$                              | $358\pm354$                    | $85\pm101$   | $12\pm9$     |
|                                  | 4 ( <i>n</i> = 34)       | $287\pm 64$ | $393\pm226$                          | $43\pm80$                              | $348\pm567$                    | $78\pm105$   | $37\pm75$    |
|                                  | P value                  | 0.039*      | 0.473                                | 0.199                                  | 0.253                          | 0.336        | 0.002*       |
| N-stage $(N = 63)$               | 0 (n = 15)               | $324\pm59$  | $390\pm268$                          | $51\pm85$                              | $508\pm771$                    | $90\pm119$   | $9\pm7$      |
|                                  | 1 (n = 48)               | $298\pm72$  | $420\pm211$                          | $41\pm72$                              | $304\pm336$                    | $78\pm98$    | $30\pm 64$   |
|                                  | P value                  | 0.076       | 0.561                                | 0.656                                  | 0.453                          | 0.695        | 0.007*       |
| N-stage† $(n = 17)$              | 0 ( <i>n</i> = 8)        | $350\pm30$  | $363 \pm 230$                        | $75 \pm 110$                           | $435\pm362$                    | 135±<br>149  | $5\pm3$      |
|                                  | 1 (n = 9)                | $334\pm87$  | $509\pm147$                          | $25\pm42$                              | $326\pm518$                    | $43\pm33$    | $9\pm 5$     |
|                                  | P value                  | 0.2         | 0.2                                  | 0.0236*                                | 0.277                          | 0.2          | 0.114        |
| M-stage ( $N = 63$ )             | 0 (n = 26)               | $323\pm74$  | $418\pm215$                          | $42\pm71$                              | $372\pm410$                    | $76\pm95$    | $14\pm21$    |
|                                  | 1 ( <i>n</i> = 37)       | $292\pm 64$ | $417\pm233$                          | $44\pm78$                              | $340\pm524$                    | $84\pm109$   | $33\pm71$    |
|                                  | P value                  | 0.058       | 0.955                                | 0.726                                  | 0.463                          | 0.582        | 0.002*       |
| TNM stage $(N = 63)$             | $\leq 2 (n = 17)$        | $339\pm76$  | $444\pm201$                          | $49\pm82$                              | $402\pm424$                    | $89\pm110$   | $9\pm 8$     |
|                                  | >2 $(n = 46)$            | $292\pm 64$ | $402\pm233$                          | $41\pm73$                              | $334\pm499$                    | $78\pm101$   | $31\pm 65$   |
|                                  | P value                  | 0.01*       | 0.394                                | 0.241                                  | 0.239                          | 0.202        | <0.001*      |

 $peak = (maximum signal - baseline signal) / baseline signal; IAUC_{60} = initial area under the curve for the first 60 s after contrast material arrival; K<sup>trans</sup> = forward volume transfer constant; <math>k_{ep}$  = reverse volume transfer constant;  $v_e$  = extravascular extracellular space volume per unit volume of tissue †A subgroup of patients (n = 17) who received curative surgery with lymph node dissection. The pathological findings are used as the gold standard \* indicates P < 0.05

voxels. The fixed-threshold MTV and TLG were automatically derived from these tumor delineations by the software (available as a predefined Syngovia tool) [16].

Excellent inter-observer reliability has been reported for  $ADC_{min}$ ,  $SUV_{max}$ ,  $SUV_{mean}$ , TLG, and MTV [16].

### **Statistical analysis**

Summary statistics are presented as the mean ( $\pm$  standard deviation [SD]) for continuous variables, or frequency and percentage for categorical variables. Pearson correlation coefficients (*r*) were calculated to evaluate the relationships between imaging biomarkers and tumor size. Differences in the DCE-MRI imaging biomarkers relative to tumor grade and TNM stages were compared using the Mann–Whitney test. The TNM stages were based on surgical or imaging findings. For the nodal stage, a subgroup of patients (*n* = 17) who received curative surgery with lymph node dissection was also analyzed, and pathological findings were used as the gold standard.

OS was measured from the date of the first MR exam to the date of death or study completion (June 30, 2017). Time to

progression (TTP) was defined as the time from the first MR exam to the time of tumor progression. For survival analysis, each variable was dichotomized as either high or low based on the median value of the variable. The Kaplan-Meier method was used to plot survival curves, and the two-sided log-rank test was used to assess differences in OS and TTP between patient groups. Multivariable analysis was investigated using the step-wise forward Cox proportional hazard model with the variables age, tumor size, TNM stage, and imaging biomarkers. A subgroup analysis of patients without curative surgery was also performed. Data were analyzed using SPSS software (SPSS for Windows 22; IBM Corp., Armonk, NY, USA). Significance was recognized when P < 0.05.

### Results

#### **Clinical treatment and follow-up**

Of 63 patients, 39 (62%) died during this study. All patients died of pancreatic cancer, so the disease-specific OS and general survival were the same in our study cohort. Median

Fig. 4 Kaplan–Meier survival curves show that patients with low values of (a) *peak* or (b) minimum ADC (ADC<sub>min</sub>) have shorter overall survival than those with high values. Patients with high values of (c) total lesion glycolysis (TLG) or (d) TLG/ *peak* ratio have shorter overall survival than those with low values



survival was 234 days (range, 6–966 days), and median follow-up was 638 days. Fifty-four patients (85.7%) had disease progression during follow-up and median TTP was 89 days (range, 6–966 days). Seventeen (17/63, 27%) patients received curative surgery (Whipple operation, n = 14; distal pancreatectomy, n = 3), and 15 received adjuvant chemotherapy. In 46 patients who did not receive curative surgery, 39 received chemotherapy, and seven received conservative treatment. Those who did not receive surgery had a much shorter OS than those who did (median survival 244 vs. 635 days, P = 0.012).

# Correlations between PET/MRI biomarkers and tumor size

Values for *peak* were positively correlated with ADC<sub>min</sub> (r = 0.258, P = 0.024). Choline was positively correlated with  $K^{trans}$  (r = 0.368, P = 0.003),  $k_{ep}$  (r = 0.357, P = 0.003), and  $v_e$  (r = 0.234, P = 0.034). In contrast, SUV<sub>max</sub> was negatively correlated with  $K^{trans}$  (r = -0.298, P = 0.009) and  $v_e$  (r = -0.235, P = 0.032). Both *peak* and *IAUC*<sub>60</sub> were negatively

correlated with MTV and TLG (all P < 0.05). Large tumors had lower values for *peak*,  $K^{trans}$ , and  $v_e$  than small tumors (all P < 0.05; Table 3).

# Association between imaging biomarkers and pathologic grade, TNM stage

Among DCE-MRI biomarkers, *peak* was lower in tumors with T4 (T4 vs. T $\leq$ 3, *P* = 0.039) and higher TNM stage (stage >2 vs.  $\leq$ 2, *P* = 0.01) than in lower-stage tumors. The ratio TLG/*peak* was higher for tumors with T4 (T4 vs. T $\leq$ 3, *P* = 0.002), N1 (N1 vs. N0, *P* = 0.007), M1 (M1 vs. M0, *P* = 0.002), or higher TNM stage (stage >2 vs.  $\leq$ 2, *P* < 0.001) than in lower-stage tumors (Table 4).

## Relationships between imaging biomarkers and OS in all patients

Univariate analyses revealed that advanced stage (P = 0.041; hazard ratio [HR] = 2.041, 95% confidence interval [CI] = 1.029–4.048), low *peak* (P = 0.006; HR = 2.559, 95% CI =

**Table 5** Correlation of imagingbiomarkers with overall survivalin all 63 patients

**Table 6**Correlation of imagingbiomarkers with overall survivalin a subgroup of 46 patientswithout curative surgery

|   |        | Univari | Univariate  |             |       | riate        |           |  |
|---|--------|---------|-------------|-------------|-------|--------------|-----------|--|
| Parameters                                  | Cutoff | HR      | 95% CI      | P value     | HR    | 95% CI       | P value   |  |
| Age (years)                                 | 64     | 1.13    | 0.6-2.127   | 0.706       |       |              |           |  |
| Gender (female vs. male)                    |        | 0.629   | 0.322-1.23  | 0.175       |       |              |           |  |
| Size (cm) (≦3.3 vs. >3.3)                   | 3.3    | 0.777   | 0.413-1.46  | 0.433       |       |              |           |  |
| ΓNM stage (4 vs.≦3)                         |        | 2.041   | 1.029-4.048 | 0.041*      |       |              |           |  |
| maging biomarkers                           |        |         |             |             |       |              |           |  |
| veak (%)                                    | 297    | 2.559   | 1.315-4.981 | $0.006^{*}$ |       |              |           |  |
| AUC <sub>60</sub> (/100, sec)               | 399    | 1.279   | 0.68-2.405  | 0.445       |       |              |           |  |
| $K^{trans}$ (min <sup>-1</sup> /1000)       | 7      | 1.392   | 0.735-2.638 | 0.310       |       |              |           |  |
| $x_{ep} ({\rm min}^{-1}/1000)$              | 213    | 1.288   | 0.680-2.439 | 0.437       |       |              |           |  |
| v <sub>e</sub> (%/10)                       | 27     | 1.066   | 0.564-2.015 | 0.844       |       |              |           |  |
| $ADC_{min} (10^{-6} \text{ mm}^2/\text{s})$ | 844    | 2.861   | 1.477-5.543 | $0.002^*$   | 0.999 | 0.998–1      | $0.011^*$ |  |
| Choline                                     | 3.33   | 1.210   | 0.615-2.378 | 0.581       |       |              |           |  |
| SUV <sub>max</sub> (g/mL)                   | 7.2    | 0.985   | 0.525-1.846 | 0.961       |       |              |           |  |
| SUV <sub>mean</sub> (g/mL)                  | 4.46   | 0.985   | 0.525-1.846 | 0.961       |       |              |           |  |
| MTV (cm <sup>3</sup> )                      | 8.54   | 0.577   | 0.305-1.09  | 0.09        |       |              |           |  |
| ГLG (g)                                     | 33     | 0.421   | 0.217-0.816 | $0.01^{*}$  |       |              |           |  |
| ГLG/peak                                    | 11.81  | 0.444   | 0.230-0.858 | 0.016*      | 4.610 | 1.565-13.582 | $0.006^*$ |  |

HR = hazard ratio; CI = confidence interval; peak = (maximum signal - baseline signal) / baseline signal; $IAUC_{60} =$  initial area under the curve for the first 60 s after contrast material arrival;  $K^{trcans} =$  forward volume transfer constant;  $k_{ep}$  = reverse volume transfer constant;  $v_e$  = extravascular extracellular space volume per unit volume of tissue; ADC<sub>min</sub> = minimum apparent diffusion coefficient; SUV<sub>max</sub> = maximum standardized uptake value; SUV<sub>mean</sub> = mean standardized uptake value; MTV = metabolic tumor volume; TLG = total lesion glycolysis

The multivariate analysis includes continuous variables (age, tumor size, *peak*, ADC<sub>min</sub>, TLG, TLG/*peak*) and categorical variables (gender and TNM stage)

\* indicates P value is significant < 0.05

| OS                  | Univaria | ite          |             | Multivariate |              |         |  |
|---------------------|----------|--------------|-------------|--------------|--------------|---------|--|
| Parameters          | HR       | 95% CI       | P value     | HR           | 95% CI       | P value |  |
| Age                 | 0.993    | 0.964-1.023  | 0.654       |              |              |         |  |
| Gender              | 0.502    | 0.242-1.042  | 0.064       |              |              |         |  |
| Size                | 1.019    | 0.786-1.32   | 0.887       |              |              |         |  |
| TNM stage (4 vs.≦3) | 1.061    | 0.477-2.359  | 0.885       |              |              |         |  |
| Image biomarkers    |          |              |             |              |              |         |  |
| peak                | 0.993    | 0.988-0.999  | 0.021*      |              |              |         |  |
| ADC <sub>min</sub>  | 0.999    | 0.998-1      | $0.019^{*}$ |              |              |         |  |
| TLG                 | 1.005    | 1.002-1.009  | $0.004^{*}$ |              |              |         |  |
| TLG/peak            | 4.682    | 1.426-15.374 | 0.011*      | 4.573        | 1.373-15.236 | 0.013*  |  |

HR = hazard ratio; CI = confidence interval; *peak* = (maximum signal – baseline signal) / baseline signal; ADC<sub>min</sub> = minimum apparent diffusion coefficient; TLG = total lesion glycolysis

The continuous variables of imaging biomarkers are used in the univariate analysis. The multivariate analysis includes continuous variables (age, tumor size, *peak*, ADC<sub>min</sub>, TLG, TLG/*peak*) and categorical variables (gender and TNM stage)

\* indicates P value is significant < 0.05

**Table 7** Correlation of imagingbiomarkers with time toprogression in all 63 patients

|   |        | Univariate |             | Multiva     | riate |             |             |
|---|--------|------------|-------------|-------------|-------|-------------|-------------|
| Parameters                                  | Cutoff | HR         | 95% CI      | P value     | HR    | 95% CI      | P value     |
| Age (years)                                 | 64     | 1.06       | 0.62-1.812  | 0.832       |       |             |             |
| Gender (female vs. male)                    |        | 1.165      | 0.671-2.022 | 0.587       |       |             |             |
| Size (cm) (≦3.3 vs. >3.3)                   | 3.3    | 0.671      | 0.392-1.148 | 0.145       |       |             |             |
| ΓNM stage (4 vs.≦3)                         |        | 1.748      | 1.011-3.023 | $0.045^{*}$ |       |             |             |
| maging biomarkers                           |        |            |             |             |       |             |             |
| veak (%)                                    | 297    | 1.697      | 0.987-2.916 | 0.056       |       |             |             |
| AUC <sub>60</sub> (/100, sec)               | 399    | 1.185      | 0.694-2.203 | 0.533       |       |             |             |
| $K^{trans}$ (min <sup>-1</sup> /1000)       | 7      | 1.492      | 0.871-2.555 | 0.145       |       |             |             |
| $k_{ep} ({\rm min}^{-1}/1000)$              | 213    | 1.043      | 0.611-1.780 | 0.878       |       |             |             |
| $v_e  (\%/10)$                              | 27     | 1.143      | 0.669-1.955 | 0.624       |       |             |             |
| $ADC_{min} (10^{-6} \text{ mm}^2/\text{s})$ | 844    | 1.694      | 0.988-2.906 | 0.056       |       |             |             |
| Choline                                     | 3.33   | 1.432      | 0.811-2.527 | 0.216       |       |             |             |
| SUV <sub>max</sub> (g/mL)                   | 7.2    | 0.825      | 0.482-1.408 | 0.48        |       |             |             |
| SUV <sub>mean</sub> (g/mL)                  | 4.46   | 0.825      | 0.482-1.408 | 0.48        |       |             |             |
| MTV (cm <sup>3</sup> )                      | 8.54   | 0.6        | 0.350-1.027 | 0.062       |       |             |             |
| ГLG (g)                                     | 33     | 0.454      | 0.263-0.781 | $0.004^{*}$ | 1.004 | 1.001-1.007 | $0.012^{*}$ |
| TLG/peak                                    | 11.81  | 0.402      | 0.233-0.694 | $0.001^*$   | 2.130 | 1.158-3.915 | $0.015^{*}$ |

HR = hazard ratio; CI = confidence interval; *peak* = (maximum signal – baseline signal) / baseline signal;  $IAUC_{60}$  = initial area under the curve for the first 60 s after contrast material arrival;  $K^{trans}$  = forward volume transfer constant;  $k_{ep}$  = reverse volume transfer constant;  $v_e$  = extravascular extracellular space volume per unit volume of tissue; ADC<sub>min</sub> = minimum apparent diffusion coefficient; SUV<sub>max</sub> = maximum standardized uptake value; SUV<sub>mean</sub> = mean standardized uptake value; MTV = metabolic tumor volume; TLG = total lesion glycolysis

The multivariate analysis includes continuous variables (age, tumor size, *peak*, ADC<sub>min</sub>, TLG, TLG/*peak*) and categorical variables (gender and TNM stage)

\* indicates P value is significant < 0.05

1.315–4.981, Fig. 4a), low ADC<sub>min</sub> (P = 0.002; HR = 2.861, 95% CI = 1.477–5.543, Fig. 4b), and high TLG (P = 0.01; HR = 3.511, 95% CI = 1.182–10.429, Fig. 4c) were prognostic factors for poor OS. In addition, a high metabolism-to-flow ratio (TLG/*peak* [P = 0.016; HR = 3.511, 95% CI = 1.182–10.429]) was a prognostic factor for poor OS (Fig. 4d). Cox multivariate analysis revealed that TLG/*peak* (P = 0.006; HR = 4.61, 95% CI = 1.565–13.582) and ADC<sub>min</sub> (P = 0.011; HR = 0.999, 95% CI = 0.998–1) remained independent predictors of OS after adjusting for age, gender, tumor size, TNM stage, and other imaging biomarkers (Table 5).

# Relationships between imaging biomarkers and OS in patients without curative surgery

Because curative surgery was an important prognostic factor for OS, we further performed subgroup analyses using the 46 patients who did not receive curative surgery to determine the prognostic values for these imaging biomarkers. Similarly, univariate analyses revealed that *peak*, ADC<sub>min</sub>, TLG, and TLG/*peak* were prognostic factors (all P < 0.05). Only the final factor was significant in multivariate analyses after adjustment for age, gender, tumor size, TNM stage, and other imaging biomarkers (P = 0.013; HR = 4.573, 95% CI = 1.373–15.236) (Table 6). Therefore, TLG/*peak* was the best predictor for OS in patients who did not receive curative surgery.

## Relationships between imaging biomarkers and TTP in all patients

Univariate analyses revealed that advanced stage (P = 0.045; HR = 1.748, 95% CI = 1.011–3.023), high TLG (P = 0.004; HR = 2.205, 95% CI = 1.280–3.797), high TLG/*peak* (P = 0.001; HR = 2.489, 95% CI = 1.441–4.3) were prognostic factors for shorter TTP. Cox multivariate analysis revealed that TLG (P = 0.012) and TLG/*peak* (P = 0.015) remained independent predictors of TTP after adjusting for age, gender, tumor size, TNM stage, and other imaging biomarkers (Table 7).

#### Discussion

In this study, we found that high *peak*, high  $ADC_{min}$ , and low TLG were good prognostic predictors of OS in patients with PDAC. The TLG/*peak ratio* and  $ADC_{min}$  were independent predictors after controlling for age, gender, size, and stage. In patients who did not receive curative surgery, the former remained an independent predictor for OS. Besides, both high TLG and TLG/*peak* ratio were independent predictors for shorter TTP as well.

The parameters of DCE-MRI have been found to correlate with tumor angiogenesis, degree of tumor fibrosis, expression of vascular endothelial growth factor Ki67, and microvascular density in PDAC [19, 20]. Compared to normal pancreas, those affected by PDAC were found to have lower values for IAUC,  $K^{trans}$ , and  $k_{ep}$ , and larger values for  $v_e$  [21, 22], suggesting decreased blood flow and increased interstitial space within the tumors. In this study, tumors with low peak values were associated with advanced TNM stage and poor OS. This variable is a semi-quantitative parameter and represents a combination of blood flow, interstitial volume, and tumor permeability [17]. We assume that the abundant fibrous stroma in more aggressive tumors impede blood flow, leading to intratumoral hypoxia and impaired drug delivery. Therefore, tumors with low peak values were associated with poor prognoses. Also, a previous study found that tumors with low values for  $K^{trans}$ and  $k_{ep}$  were associated with poor responses after antiangiogenic therapy [23]. Based on these results, it seems that high tumor angiogenesis before treatment is associated with better therapy response and survival outcome in PDAC.

DWI can potentially depict and characterize a variety of focal pancreatic lesions [24], particularly for staging in PDAC [25]. The ADC is significantly lower in tumors with dense fibroses and can serve as a biomarker of fibrosis architecture [26, 27]. Previous studies showed that a low pretreatment ADC was associated with shorter PFS after chemotherapy [28] and shorter OS in patients with resected PDAC [29]. We found that a low pretreatment ADC main independently predicted a poor outcome in all patients and also in the subgroup of patients without curative surgery. We also found a positive correlation between ADC<sub>min</sub> and *peak*, suggesting reduced blood flow in tumors with high cellularity or abundant fibroses. Further studies are needed to evaluate the usefulness of DWI for monitoring treatment.

Using <sup>1</sup>H–MRS, we found that choline was positively correlated with  $K^{trans}$ ,  $k_{ep}$ , and  $v_{e}$ . In PDAC, increased levels of choline suggest increased membrane activity, particularly enhanced cellular proliferation. In vivo, poor perfusion can limit substrate delivery, thus decreasing metabolite concentrations in poorly vascularized regions of the tumor [30]. Although a previous study found that low choline was associated with longer survival in PDAC [31], we did not find it was associated with OS, likely because the patient sample in this study was small.

Several studies have shown the potential role of PET in determining therapy response, recurrence detection, and survival prediction [32]. This study confirms that high TLG indicates poor OS and TTP. Tumor blood flow and glucose metabolism are correlated with each other because when blood flow decreases, tumor hypoxia promotes gene expression via the transcription factor hypoxia-inducible factor  $1\alpha$ . which leads to accelerated glycolysis [33]. When more highly expressed, it might promote inflammation and fibrosis in PDAC [34], and it has been shown to correlate with poor prognoses in these patients [35]. Therefore, the combined flow-metabolism imaging biomarker, TLG/peak, outperforms other imaging biomarkers in predicting OS, even in patients without curative surgery. This biomarker could be obtained in a single study via integrated PET/MRI in clinical practice, and it is potentially a biomarker for monitoring therapy response.

Several studies have found correlations between imaging biomarkers and tumor histologic findings such as microvascular density, fibrosis, and nuclear grade [16, 27, 36]. In this study,  $v_e$  seems to be lower in poorly differentiated PDAC compared to well- to moderately differentiated tumors (P = 0.059), but further studies with larger sample sizes are needed.

We acknowledge some limitations in this study. First, the number of patients is limited, especially among those who received curative surgery. Because PDAC is extremely aggressive, very few patients are treated with curative surgery. Second, further studies should take into account additional elements that affect patient survival, such as performance status. Third, a comparative effectiveness study should be performed to evaluate serum markers versus imaging biomarkers. Fourth,  $ADC_{min}$  and  $SUV_{max}$  are essentially single data points and could be influenced by image noise. Our findings should be validated with multicenter studies. Another limitation of this study is the long duration of the PET/MRI acquisition protocol for multiparametric imaging. According to our results, MRS could be omitted to reduce examination time in clinical practice because choline was not correlated with survival outcome.

In conclusion, multiparametric PET/MRI biomarkers are associated with OS and TTP in patients with PDAC. These prognostic biomarkers might be useful for the design and development of future trials or the selection of personalized therapeutic options, possibly changing the management and improving the prognosis for patients with PDAC.

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

Conflict of interest None.

**Ethical approval** All procedures performed in studies involving human participants were in accordance with the ethical standards of the

institutional and national research committee and with the 1964 Helsinki declaration and its later amendments or comparable ethical standards.

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

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