Epithelial-to-Mesenchymal Transition in Pancreatic Cancer is associated with Restricted Water Diffusion in Diffusion-Weighted Magnetic Resonance Imaging

Philipp Mayer¹, Anne Kraft², Wilfried Roth³, Thilo Hackert³, Frank Bergmann⁴,⁵, Hans-Ulrich Kauczor¹, Verena Steinle¹, Ekaterina Khristenko¹, Miriam Klauss¹, Matthias M. Gaida²,⁶,⁷

¹ Clinic for Diagnostic and Interventional Radiology, Heidelberg University Hospital, 69120 Heidelberg, Germany.
² Institute of Pathology, University Medical Center Mainz, JGU-Mainz, 55131 Mainz, Germany.
³ Department of General, Visceral, and Transplantation Surgery, Heidelberg University Hospital, 69120 Heidelberg, Germany.
⁴ Institute of Pathology, Heidelberg University Hospital, 69120 Heidelberg, Germany.
⁵ Clinical Pathology, Klinikum Darmstadt GmbH, 64283 Darmstadt, Germany.
⁶ Research Center for Immunotherapy, University Medical Center Mainz, JGU-Mainz, 55131 Mainz, Germany.
⁷ Joint Unit Immunopathology, Institute of Pathology, University Medical Center, JGU-Mainz and TRON, Translational Oncology at the University Medical Center, JGU-Mainz, 55131 Mainz, Germany.

Corresponding authors: Dr. med. Philipp Mayer; Clinic for Diagnostic and Interventional Radiology, Heidelberg University Hospital, Im Neuenheimer Feld 420, 69120 Heidelberg, Germany. Email: Philipp.Mayer@med.uni-heidelberg.de, Telephone: +49(0)6221/56-37345, Fax: +49(0)6221/56-5730. Prof. Dr. med. Matthias M. Gaida; Institute of Pathology, University Medical Center Mainz, Langenbeckstr. 1, 55131 Mainz, Germany. Email: Matthias.Gaida@unimedizin-mainz.de, Telephone: +49(0)6131/17-3269, Fax: +49(0)6131/17-6604.

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Abstract

**Purpose:** This study aimed to evaluate the potential of diffusion-weighted magnetic resonance imaging (DW-MRI) as imaging biomarker for epithelial-to-mesenchymal transition (EMT) in pancreatic ductal adenocarcinoma (PDAC).

**Methods:** In forty-two patients, preoperative apparent diffusion coefficient (ADC) values of therapy-naive PDAC were compared with immunohistochemical expression profiles of the epithelial marker E-cadherin as well as mesenchymal transcription factors Runx-related transcription factor 2 (Runx2) and Zinc finger E-box-binding homeobox 1 (Zeb1), as determined by Allred immunoreactivity score.

**Results:** We observed a significant positive rank correlation between the ADC and the E-cadherin Allred score ($\rho = 0.553, p < 0.001$) and significant negative rank correlations between the ADC and the Runx2 Allred score ($\rho = -0.526, p < 0.001$) as well as the Zeb1 Allred score ($\rho = -0.710, p < 0.001$). Compared to tumors with low ADC values < 1.3 $\mu$m²/s, tumors with ADC values ≥ 1.3 $\mu$m²/s had significantly higher Allred scores for E-cadherin (median, 4 versus 5; $p < 0.001$) and significantly lower Allred scores for Runx2 (median, 3 versus 2; $p = 0.003$) as well as Zeb1 (median, 4 versus 0; $p < 0.001$).

**Conclusion:** In PDAC, tumor plasticity in terms of EMT is well reflected by ADC values from DW-MRI. In the near future, DW-MRI could be beneficial for identification of PDAC patients that might profit from personalized EMT-targeted therapies.

Key words: Pancreatic cancer, epithelial-to-mesenchymal transition, diffusion-weighted magnetic resonance imaging.

Introduction

Pancreatic ductal adenocarcinoma (PDAC) is a highly aggressive tumor, which is predicted to achieve the second rank in cancer-associated deaths in 2030 [1]. Since most early-stage tumors are
asymptomatic, the majority of PDACs are detected at a progressed stage and only a minor portion of patients qualify for potentially curative resection [2]. Other factors contributing to the poor prognosis of PDAC include early vascular invasion, the rapid development of distant metastases, and its notoriously poor response to (radio)chemotherapy [3].

Clinical imaging plays a key role not only for the initial diagnosis of PDAC, but also for the assessment of the local tumor mass, detection of distant metastases, and monitoring of disease in patients undergoing (neo-)adjuvant chemotherapy as well [4]. Computed tomography (CT) is the most widely used imaging modality to assess the local tumor spread and particularly the involvement of major abdominal vessels which determines tumor resectability [5,6] as well as extrapancreatic perineural invasion which is a significant predictor of recurrence [7]. A couple of studies reported conventional contrast-enhanced CT to be useful for characterization of some micro structural features of PDAC, such as cellular density [8] and stroma content [9]. However, the major strength of conventional contrast-enhanced CT is rather the accurate depiction of the macroanatomy than the probing of the tissue microstructure [5].

Magnetic resonance imaging (MRI) is often seen as a problem-solving tool for detection and characterization of PDACs [10]. Its superior soft tissue contrast is beneficial for detection of small non-contour deforming tumors, and for detection of a minor portion of PDACs that are isoattenuating on CT without any visible tumor-pancreas contrast [10,11]. In particular, diffusion-weighted (DW)-MRI, which quantitates the random motion of water molecules in biological tissues, has a very high diagnostic performance for detection of PDACs with reported sensitivity and specificity values well above 90% [12]. In recent years, several studies have reported encouraging results for (sub)cellular characterization of PDAC lesions by DW-MRI [12], including their cell density [13], growth pattern [14], stromal content [15], immune cell infiltration [16], as well as tumor hypoxia [17]. The ultrastructural findings in PDAC show a heterogeneous picture with tumor cells sealed by an extensive fibrotic stroma with interspersed inflammatory cells [18]. PDAC shows usually a glandular growth pattern of tumor cells, however also (micro-) papillary, solid-nest-like, cribriform, or single-cell dissociated tumor growth can be detected, reflecting the high plasticity of this cancer [14]. Various molecular factors, such as the loss of epithelial markers and the gain of a mesenchymal tumor phenotype (epithelial-to mesenchymal-transition; EMT) or dense infiltrates of neutrophils, are associated with changes in the microarchitecture and tumor growth pattern [19,20]. In PDAC, as in a variety of other tumors, EMT is associated with an aggressive, invasive phenotype, where tumor cells disseminate from the main tumor mass and change their morphology and molecular phenotype, towards anti-apoptotic, pro-metastatic, and chemoresistant behavior [21]. The process of EMT in tumor cells can be induced by a variety of environmental factors, for example numerous cytokines, such as transforming growth factor (TGF)-β, by loss of cell-cell contact, by proteases, matrix stiffness, or by tumor hypoxia [20,22–24]. Typical features of EMT besides the change of cell morphology and loss of polarity, are the downregulation of the epithelial cell adhesion molecule E-cadherin and different cytokeratins, in parallel to the upregulation of mesenchymal markers such as vimentin or fibronectin [22]. This process is induced by transcription factors such as the Runt-related transcription factor 2 (Runx2), as an early regulator of EMT, or the Zinc finger E-box-binding homeobox 1 (Zeb1), as a later occurring key EMT regulator [25,26]. In PDAC undergoing EMT, microarchitectural changes are described, such as the separation of tumor cells from the main tumor mass and disseminated tumor growth in small clusters or as single cells, but also the loss of cytokeratins and expression of EMT-promoting transcription factors Zeb1 or nuclear β-Catenin [19,20,27]. Of note is that EMT can confer resistance to radiochemotherapy. Targeted inhibition of Zeb1 is known to increase radiosensitivity of cancer cells through an EMT-dependent mechanism [28]. In a study by Arumugam et al., high levels of Zeb1 and low levels of E-cadherin in PDAC cell lines were correlated with resistance to gemcitabine, 5-fluorouracil (5-FU), and cisplatin [29].

Several strategies targeting EMT in PDAC are being investigated in preclinical studies [30] and targeting EMT could increase PDAC’s sensitivity to conventional radiochemotherapy in the future.

To date, there are no clinically established imaging biomarkers for prediction of EMT in vivo. While contrast-enhanced CT is widely used to assess the macroscopic tumor extent, DW-MRI can probe the tissue composition at a more granular scale [12]. DW-MRI was proven to well reflect tissue properties of PDAC that are closely linked to EMT, including tumor growth pattern [14], stroma content and properties [15], as well as tumor hypoxia [17], and previous studies reported an association of DW-MRI and EMT in xenograft tumor models [31,32]. Therefore, we hypothesized that DW-MRI could represent a biomarker for EMT in PDAC.
Material and Methods

Patients

The study protocol was approved by the local Ethics Committee (S-044/2012). Forty-two PDAC patients from a previously well-characterized cohort [17] were retrospectively included in the present study. All patients had surgical resection of therapy-naive PDAC and had a DW-MRI of the pancreas on the day before surgical tumor resection at the University Hospital of Heidelberg, Germany. Written informed consent was obtained from all participants in the study. The detailed clinical parameters are listed in Table 1. The workflow of the study is presented in Figure 1.

Table 1. Clinical and pathological patient characteristics

| Median Age (IQR) | 67.5 years (42 – 85 years) |
|------------------|---------------------------|
| **Sex**          |                           |
| Female           | 19 (45.2 %)               |
| Male             | 23 (54.8 %)               |
| **Neoadjuvant (radio)chemotherapy** |                     |
| No               | 42 (100.0 %)              |
| Yes              | 0 (0.0 %)                 |
| **T stage**      |                           |
| T1               | 0 (0.0 %)                 |
| T2               | 19 (45.2 %)               |
| T3               | 23 (54.8 %)               |
| T4               | 0 (0.0 %)                 |
| **N stage**      |                           |
| N0               | 8 (19.0 %)                |
| N1               | 8 (19.0 %)                |
| N2               | 26 (61.9 %)               |
| **M stage**      |                           |
| M0/Mx            | 40 (95.2 %)               |
| M1               | 2 (4.8 %); retroperitoneal lymph node metastasis in 1 patient, hepatic metastasis in 1 patient |
| **Grading**      |                           |
| G1               | 0 (0.0 %)                 |
| G2               | 25 (59.5 %)               |
| G3               | 17 (40.5 %)               |
| **Tumor localization** |                     |
| Pancreatic head  | 33 (78.6 %)               |
| Pancreatic body and/or tail | 9 (21.4 %)               |
| **Type of pancreatectomy** |                 |
| ppWhipple        | 18 (42.9 %)               |
| cWhipple         | 9 (21.4 %)                |
| Left resection   | 8 (19.0 %)                |
| Total pancreatectomy | 7 (16.7 %)               |

Abbreviations: cWhipple, conventional Whipple procedure; IQR, interquartile range; ppWhipple, pylorus-preserving Whipple procedure.

Diffusion-weighted MRI

DW-MRI was performed on a clinical 1.5 Tesla scanner (Magnetom Aera, Siemens Healthcare, Erlangen, Germany) using a 6-element body-phased array coil and a 24-channel spine array coil. DW-sequence was a single-shot echo-planar imaging sequence in axial orientation. Acquisition parameters were as follows: Number of slices per b-value, 14; slice thickness, 5.0 mm; distance factor, 5 %; phase encoding direction, anterior-posterior; repetition time (TR), 2200 ms; time to echo (TE), 58.0 ms; acquisition matrix, 130 * 92; spectral fat saturation; bandwidth, 2262 Hz/pixel; GRAPPA (generalized autocalibrating partially parallel acquisitions) acceleration factor, 2. All DW-MR examinations included at least the b-values $b_0 = 0 \text{s/mm}^2$, $b_{50} = 50 \text{s/mm}^2$, and $b_{800} = 800 \text{s/mm}^2$. Opposed to our previous study where apparent diffusion coefficients had been calculated from $b_50$ and $b_{800}$ (ADC$_{50,800}$) [17], ADC values were now calculated from $b_0$ and $b_{800}$ (ADC$_{0,800}$; in the following referred to as ADC).

Immunohistochemistry

Human tissue samples were provided by the local tissue bank in agreement with the regulations of the tissue bank and the approval of the local Ethics committee (no. 206/2005). All patients gave written informed consent. Paraffin sections of the respective tumor specimens were used for

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immunohistochemistry, which was performed at the Institute of Pathology, University Medical Center Mainz, Germany. Prior to incubation with the primary antibody, heat-induced antigen retrieval was conducted using buffer (pH 6.0 for Runx2; pH 9.0 for E-cadherin and Zeb1; Dako EnVision, Glostrup, Denmark). The following primary antibodies were used: E-cadherin (ready to use; Dako), Runx2 (1:50; Cell Signaling Technology, Leiden, Netherlands), and Zeb1 (1:500; Novus Biological, Abingdon, UK).

Analyses were performed on whole tumor sections to evaluate membranous, of Runx2 and Zeb1 nuclear. The staining was semi-quantitative and widely established using a chromogen (Dako) and analyzed using the Gryphax Subra camera system (Jenoptik, Jena, Germany). For confirmation of results, quantification was cross-checked software-based by ImageJ (US National Institutes of Health, Bethesda, MD, USA).

Statistical analyses

Statistical analyses were performed using the statistical software package MedCalc (version 20, MedCalc Software Ltd, Ostend, Belgium). Data were expressed as absolute and relative frequencies for categorical data and as median with interquartile range (IQR) for continuous variables. The Mann–Whitney U test was used for comparison of continuous variables between groups. Correlation analyses were performed using Spearman’s rank correlation coefficient. Inter-rater reliability was assessed with the intra-class correlation coefficient (ICC) (two-way ICC, random raters’ assumption reproducibility). As proposed by Song et al. [34], ICC values were interpreted as poor correlation (0.00 – 0.20), fair correlation (0.21 – 0.40), moderate correlation (0.41 – 0.60), good correlation (0.61 – 0.80), or excellent correlation (0.81 – 1.00). P < 0.05 were considered to be statistically significant.

Results

Diffusion-weighted imaging analyses

Tumor ADC values could be determined for all patients. Agreement for ADC values between both raters was excellent with ICC = 0.856 (95% confidence interval (CI), 0.748 – 0.920). In the following, given ADC values represent mean values for both raters, if not stated otherwise. ADC values of tumors ranged from 0.959 µm²/s to 1.631 µm²/s (median, 1.297 µm²/s; IQR, 1.166 – 1.419 µm²/s) and were similar to those from previous studies on DW-MRI in treatment naïve PDAC patients [35–38].

Immunohistochemical analyses

E-cadherin showed a membranous staining pattern. E-cadherin Allred scores ranged from 0 to 8 (median, 5; IQR, 4 – 6). Runx2 and Zeb1 showed a nuclear staining pattern with Allred scores from 0 to 7 for Runx2 (median, 2; IQR, 2 – 4) and from 0 to 5 for Zeb1 (median, 2.5; IQR, 0 – 4). There was a significant negative correlation between the E-cadherin Allred score and the Zeb1 Allred score (ρ = -0.604, p < 0.001) and a significant positive correlation between the Runx2 Allred score and the Zeb1 Allred score (ρ = 0.507, p < 0.001). The E-cadherin Allred score was only weakly negatively correlated with the Runx2 Allred score (ρ = -0.278, p = 0.074). E-cadherin, Runx2, and Zeb1 Allred scores were neither significantly different between T2 tumors and T3 tumors (p ≥ 0.294) nor between moderately differentiated (G2) and poorly differentiated tumors (G3, p ≥ 0.422) or between lymph node negative (N0) and positive tumors (N1 or N2, p ≥ 0.704).

Radiopathological correlation

We observed a significant positive correlation between the ADC and the E-cadherin Allred score (ρ = 0.553, p < 0.001). There were significant negative correlations between the ADC and the Runx2 Allred score (ρ = -0.526, p < 0.001) as well as between the ADC and the Zeb1 Allred score (ρ = -0.710, p < 0.001). ADC values were significantly lower in tumors with low E-cadherin expression (E-cadherin Allred < 5; median ADC, 1.170 µm²/s; IQR, 1.119 – 1.286 µm²/s) than in tumors with high E-cadherin expression (E-cadherin Allred ≥ 5; median ADC, 1.403 µm²/s; IQR, 1.261 – 1.522; p = 0.001) (Figure 2A). ADC values were significantly higher in tumors with Runx2 Allred scores ≥ 3 (median ADC, 1.378 µm²/s; IQR, 1.231 – 1.498 µm²/s) than in tumors with Runx2 Allred scores < 3 (median ADC, 1.172 µm²/s; IQR, 1.090 – 1.324 µm²/s; p = 0.003) (Figure 2B). Tumors with Zeb1 Allred scores < 3 had significantly higher ADC values (median ADC, 1.403 µm²/s; IQR, 1.288 – 1.532 µm²/s) than tumors with Zeb1 Allred scores ≥ 3 (median ADC, 1.166 µm²/s; IQR, 1.094 – 1.300 µm²/s; p < 0.001) (Figure 2C). Vice versa, compared to tumors with ADC values < 1.3 µm²/s, tumors with ADC values ≥ 1.3 µm²/s had significantly higher Allred scores for E-cadherin (median Allred, 4; IQR, 3 – 5; versus median Allred, 5; IQR, 5 – 7.25; p < 0.001) and significantly lower Allred scores for Runx2 (median Allred, 3; IQR 2 – 4; versus median Allred, 2; IQR, 0 – 3; p = 0.003) as well as Zeb1 (median Allred, 4; IQR, 2.75 – 4.25; versus median Allred, 0; IQR, 0 – 2.25; p < 0.001).
ADC values were neither significantly different between T2 tumors and T3 tumors (p = 0.950) nor between moderately differentiated (G2) and poorly differentiated tumors (G3, p = 0.654) or between lymph node negative (N0) and positive tumors (N1 or N2, p = 0.353). This is in line with the results from a study by Rosenkrantz et al. [39] but in contradiction to the results from a study by Wang et al. [40].

DW images and immunohistochemical stainings from a patient with high diffusivity and epithelial tumor phenotype are presented in Figure 3. DW images and immunohistochemical stainings from a patient with low diffusivity and mesenchymal tumor phenotype are presented in Figure 4. A schematic representation of the dependency of the diffusivity on EMT in PDAC is presented in Figure 5.
Figure 4: Images from a patient with low diffusivity. The T2-HASTE MR image shows the tumor in the pancreatic head (arrows). A VOI in red encompasses the tumor on the DW-MR images. Please note that the tumor is markedly hyperintense in the b800 image and markedly hypointense in the ADC map. The ADC value of the tumor was 1.048 µm²/s. In immunohistochemistry (lower row), tumor cells showed only focal membranous E-cadherin expression (arrows in the left image) but showed strong nuclear positivity for Runx2 (arrows in the middle image) and Zeb1 (arrows in the right image). Please note the Zeb1-positive stroma cells (arrowheads in the right image).

Figure 5: Schematic representation of the dependency of the MRI diffusivity on EMT. Epithelial type PDAC with high expression of E-cadherin exhibit significantly higher diffusivity (arrows) in DW-MRI than mesenchymal type PDAC with low expression of E-cadherin and higher expression of Runx2 and Zeb1.
Discussion

Pancreatic cancer is characterized by high tumor plasticity, which is responsible for its aggressive behavior and therapeutic resistance [3,41]. In this vein, multiple studies described molecular features being associated with this particular tumor aggressiveness. A well-established concept is epithelial to mesenchymal transition in PDAC, where tumor cells lose their epithelial features and de-differentiate into a mesenchymal, aggressive cellular phenotype, which is characterized by anti-apoptotic, pro-invasive, pro-metastatic, and chemoresistant features [21]. EMT in pancreatic cancer can be visualized by disseminated tumor growth, loss of cytokeratins, or expression of the mesenchymal transcription factors Zeb1 or nuclear β-Catenin [19,20,27]. We decided to use the established epithelial marker E-cadherin, and two mesenchymal markers, Runx2 as early-stage marker for EMT, and Zeb1 as later occurring marker. Epithelial features could be shown in the majority of tumors and were negatively correlated with the expression of the mesenchymal markers Runx2 and Zeb1, indicating a transient and dynamic EMT process in the tissues. The inverse correlation of E-Cadherin and Zeb1 is well documented for EMT in pancreatic cancer, resulting in consequent resistance to the clinically widely used chemotherapies [29]. In line with these findings, also the expression of the second investigated EMT marker, Runx2, in pancreatic cancer is associated with chemoresistance [42].

To date, prediction of EMT and other molecular markers indicating EMT-mediated aggressiveness or chemoresistance of PDAC is only possible using a tissue biopsy, because of the lack of established imaging biomarkers in vivo. We hypothesized that DW-MRI, which is widely used for pancreatic imaging, could represent an imaging biomarker for EMT since it was previously proven to reflect histological tissue characteristics of PDAC which are closely linked to EMT, tumor growth pattern [14], immune cell infiltrates [14], desmoplastic stroma characteristics [15], and tumor hypoxia [17], although the direct prove of EMT and imaging markers have not been established in PDAC. In the present study, we found a significant positive correlation between the ADC and E-cadherin expression in tumor cells. On the other hand, the ADC was significantly negatively correlated with Runx2 as well as Zeb1 expression. ADC values were significantly lower in tumors with low E-cadherin expression compared to high E-cadherin expression. Supporting that notion of an association with intratumor EMT, the ADC values were significantly higher in tumors with high Allred immunoreactivity scores for Runx2 and Zeb1 than in tumors with low Runx2 and Zeb1 expression. Vice versa, compared to tumors with low ADC values, tumors with high ADC values had significantly higher expression of E-cadherin and significantly lower expression of Runx2 as well as Zeb1. Our data clearly show that intratumor EMT is closely linked to restricted diffusion in MRI. A possible explanation is that the larger glandular lumina and stable glandular structures of epithelial phenotype PDAC enable a relatively unimpeded diffusion of water molecules while the mesenchymal phenotype PDAC exhibits a more clustered growth pattern, sometimes even micro- or non-glandular and single cellular [27] which had been linked to more restricted water diffusivity [14]. In line with this, Zhang et al. also reported a positive, albeit weak, correlation between the epithelial marker E-cadherin and the ADC in a chronic pancreatitis rat model [43]. Similarly, in a study on EMT in a colorectal carcinoma xenografts model, tumors with ongoing EMT had lower diffusivity values than tumors without EMT [32]. In seeming contradiction, Chen et al. found less restricted diffusion in mesenchymal-like mouse xenograft tumors than in their epithelial-like counterparts, using hepatocellular carcinoma (HCC) cell lines [31]. They hypothesized that the tighter cell-cell junctions in the epithelial-like tumors lead to more densely packed tumor cells with decreased extracellular space, and thus more restricted diffusion than in the mesenchymal-like tumors [31]. From the histological side, this effect is explainable because injected xenograft models form a solid tumor mass, without a desmoplastic stroma or glandular tumor formation, as in human (pancreatic) cancer. With intact epithelial adhesion molecules, a solid tumor with stable cell-cell contacts, but without significant intercellular spaces is outgrown, compared to the mesenchymal xenograft, which due to loss of intercellular contacts, creates small spaces, where water molecules can diffuse freely. This effect is obviously less prominent in the human PDAC in the present study: Cancer areas with high expression of epithelial markers form glandular structures with large intercellular spaces, surrounded by desmoplastic stroma. Mesenchymal cells from small or single cellular clusters, without larger empty spaces, and are therefore sealed in desmoplastic stroma.

Since EMT induction can be associated with increased glucose uptake of tumor cells [44], several studies focused on 18F-fluorodeoxyglucose (18F-FDG) uptake in positron emission tomography (PET) as an imaging biomarker for EMT in different tumor types, including esophageal cancer [45,46], HCC [47], oral
squamous cell carcinoma [48], and non-small cell lung cancer [44]. However, high costs, the required logistics, and limited availability are obvious drawbacks of $^{18}$FDG-PET [49,50], potentially restraining its widespread clinical use. On the other hand, DW-MRI is not limited by these drawbacks of $^{18}$FDG-PET and might therefore represent a more practicable imaging biomarker for the prediction of intratumor EMT.

We provide first-time evidence for a human pancreatic cancer cohort that tumor plasticity in terms of EMT is directly correlated to ADC values in DW-MRI. Non-invasive estimation of EMT in PDAC by DW-MRI could help to identify aggressive PDAC lesions with high risk for chemotherapy failure. In the future, DW-MRI may also be a helpful tool to select patients for personalized (experimental) therapies, targeting EMT [30].

Progress in (personalized) treatment methods for PDAC is of urgent need due to the still devastating prognosis of the disease. While recent advances in personalized medicine have led to significant improvements in overall survival in other solid malignancies, there have been only modest survival improvements for PDAC patients [51]. This failure cannot be attributed to a scarcity of research in personalized medicine in PDAC [51]. While previously used targeted therapies were limited by an insufficient understanding of the molecular aspects of the disease, better preclinical models mimicking human disease have led to a rapid increase in knowledge on PDAC’s genetics and biology [52]. New therapeutic strategies include exploitation of DNA repair abnormalities, targeting cancer-cell metabolism and EMT as well as components of the stroma or the tumor micromilieu [52–54]. However, although up to one fourth of PDAC harbour clinically actionable molecular alterations and molecularly matched therapies may improve overall survival in these patients, the percentage of PDAC patients who ultimately receive targeted therapies is very low [51,55,56]. This shows that the challenge for personalized medicine in PDAC might not only be the dearth of targeted and other novel therapy options, but rather improving patient stratification and making novel therapies more accessible to individual patients and treating physicians [51,52]. Sole reliance on molecular markers for treatment stratification can be problematic in PDAC since tiny biopsy specimens often do not contain enough tumor cells, depending on the exact biopsy technique and the abundance of the stroma, and their results are affected by the biopsy site due to the notorious tumor heterogeneity of PDAC [51]. Radio-pathological correlation studies, such as the present study, lead to the understanding that many microscopic and molecular alterations of the tumor tissue are accompanied by inter- and intraindividual qualitative and quantitative changes in radiological imaging [57]. Therefore, radiological imaging, especially quantitative imaging, which allows non-invasive analysis of the whole tumor tissue, in combination with pathological and molecular tissue analysis, could serve as a valuable vein for treatment stratification in the era of personalized medicine [51].

In summary, the present study is an example that microarchitectural changes, seen in histopathological correlates, are directly reflected by MRI. The clinical impact is an upfront estimation, whether the tumor belongs to an aggressive, chemoresistant phenotype, which can be generated in the initial tumor staging and where potentially in the future PDAC patients may benefit from EMT-targeted therapies.

**Abbreviations**

- 5-FU: 5-fluorouracil; $^{18}$FDG: $^{18}$F-fluorodeoxyglucose; ADC: apparent diffusion coefficient; CI: confidence interval; cWhipple: conventional Whipple procedure; CT: computed tomography; DKFZ: German Cancer Research Center; DW-MRI: diffusion-weighted magnetic resonance imaging; EMT: epithelial-to-mesenchymal transition; GRAPPA: generalized autocalibrating partially parallel acquisitions; HCC: hepatocellular carcinoma; ICC: intra-class correlation coefficient; IQR: interquartile range; MITK: Medical Imaging Interaction Toolkit; MRI: magnetic resonance imaging; PDAC: pancreatic ductal adenocarcinoma; PET: positron emission tomography; ppWhipple: pylorus-preserving Whipple procedure; Runx2: Runt-related transcription factor 2; TGF-β: transforming growth factor-β; VOI: volume of interest; Zeb1: Zinc finger E-box-binding homeobox 1.

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**Competing Interests**

The authors have declared that no competing interest exists.

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