

Review

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Current state of radiomics in hepatobiliary and pancreatic malignancies

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Abstract

Rising in incidence, hepatobiliary and pancreatic (HPB) cancers continue to exhibit dismal long-term survival. The overall poor prognosis of HPB cancers is reflective of the advanced stage at which most patients are diagnosed. Late diagnosis is driven by the often-asymptomatic nature of these diseases, as well as a dearth of screening modalities. Additionally, standard imaging modalities fall short of providing accurate and detailed information regarding specific tumor characteristics, which can better inform surgical planning and sequencing of systemic therapy. Therefore, precise therapeutic planning must be delayed until histopathological examination is performed at the time of resection. Given the current shortcomings in the management of HPB cancers, investigations of numerous noninvasive biomarkers, including circulating tumor cells and DNA, proteomics, immunomics, and radiomics, are underway. Radiomics encompasses the extraction and analysis of quantitative imaging features. Along with summarizing the general framework of radiomics, this review synthesizes the state of radiomics in HPB cancers, outlining its role in various aspects of management, present limitations, and future applications for clinical integration. Current literature underscores the utility of radiomics in early detection, tumor characterization, therapeutic selection, and prognostication for HPB cancers. Seeing as single-center, small studies constitute the majority of radiomics literature, there is considerable heterogeneity with respect to steps of the radiomics workflow such as segmentation, or delineation of the region of interest on a scan. Nonetheless, the introduction of the radiomics quality score (RQS) demonstrates a step towards greater standardization and reproducibility in the young field of radiomics. Altogether, in the setting of continually improving artificial



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intelligence algorithms, radiomics represents a promising biomarker avenue for promoting enhanced and tailored management of HPB cancers, with the potential to improve long-term outcomes for patients.

Keywords: Pancreatic neoplasms, pancreatic ductal adenocarcinoma, hepatocellular carcinoma, cholangiocarcinoma, radiomics, artificial intelligence

INTRODUCTION

Hepatobiliary and pancreatic malignancies comprise a heterogeneous group of diseases that rank amongst the leading causes of cancer-related deaths worldwide^[1]. Despite improvements in cancer surveillance, imaging, and treatment, the prognosis of these patients remains poor, with the 5-year survival rate of pancreatic cancer [pancreatic ductal adenocarcinoma (PDAC)] and biliary tract carcinomas (BTCs) reported of only 10% and 5%-18%, respectively^[2,3]. The prognosis of localized hepatocellular carcinoma (HCC) is marginally better at 33%; however, it drops off to 2% in the context of metastatic disease^[4].

Poor outcomes are predominantly driven by a delay in the diagnosis of these diseases^[1]. Diagnostic challenges of hepatobiliary and pancreatic (HPB) malignancies are multifaceted. Firstly, due to the asymptomatic nature of the disease, these patients are diagnosed with more advanced diseases, with only 43% of hepatic cancer and 20% of pancreatic cancers being diagnosed at an early stage^[5,6]. Secondly, there is a lack of screening modalities (PDAC) or the current screening strategies are not very accurate (HCC), e.g., the ultrasound-based protocol for HCC has a sensitivity of 47%-63% for detection of early-stage disease^[7]. Biomarkers for screening of these diseases are lacking^[8,9]. Additionally, pancreatic cancer often mimics benign lesions on imaging, which can result in misdiagnosis and delay in the delivery of appropriate care^[6]. It is estimated that nearly 5%-11% of all patients undergoing pancreaticoduodenectomies for pancreatic cancer turn out to have benign lesions^[10]. Altogether, there is a critical need for noninvasive biomarkers to facilitate earlier diagnosis of hepatobiliary and pancreatic malignancies.

Once a diagnosis is established, tumor characterization (tumor grade, presence of nodal disease, extent of local invasion, and molecular profile) is essential in determining appropriate care. This is vital for sequencing of systemic therapy and surgical planning^[11,12]. While helpful, current imaging modalities are limited by their accuracy in determining these features. Furthermore, the selection of appropriate systemic therapies presents a challenge in the management of these diseases. Currently, clinical tools to predict treatment response are absent, and it is not determined until the patient undergoes resection and a histopathological examination of the specimen is performed. If the disease was resistant to the administered chemotherapeutics, unfortunately, the patient gained no benefit from this therapy, while allowing time for resistant clones to proliferate and result in the progression of disease. For instance, in the setting of metastatic PDAC, Nab-paclitaxel combined with gemcitabine represents the standard of care; however, there is considerable heterogeneity among patients with respect to duration of treatment (0.1-21.9 months), secondary to the significant toxicities of these therapies, as demonstrated by the MPACT trial, as well as treatment response, with the SIEGE trial reporting almost a fifth of patients failing to reach their first treatment response assessment^[13,14]. As such, there is a critical need to robustly validate biomarkers to both spare patients from toxicities associated with treatments that they are unlikely to respond to and to identify new avenues for targeted treatments.

Recently, multiple biomarkers have shown promise in hepatobiliary and pancreatic malignancies, including circulating tumor cells, circulating tumor DNA, proteins, and radiomics^[9,15-17]. The identification and validation of biomarkers could aid in patient stratification, treatment planning, and prediction of response

to therapy and risk of recurrence, thereby improving survival^[18]. This narrative review focuses on the current literature on radiomics as a biomarker for hepatobiliary and pancreatic malignancies. We discuss how radiomics could help guide the management of these diseases, the current limitations of radiomics, and future applications and integration in the clinical setting.

METHODS

To identify and synthesize literature for a narrative review regarding the utility and applications of radiomics in the management of HPB tumors, PubMed and Embase, as well as Google Scholar, were queried. These platforms were searched from inception until July 2023. Given that a qualitative review was planned, all article types published in English were considered eligible for inclusion. Terms such as “radiomics” and “segmentation” were combined with the term, “biomarker”, and various iterations of aspects of management, such as “prognostication” and “treatment response” and specific HPB tumors (e.g., “HCC” “PDAC”, *etc.*).

DEFINING RADIOMICS

Recently, the field of radiomics has burgeoned and shown promise as a potential tool for early diagnosis, tumor characterization, and prognostication^[19]. This involves extraction of high-dimensional data from images and providing feature data for quantitative description of lesions^[20]. Radiographic images contain a number of quantifiable features that may be mined and analyzed to offer insights into disease processes^[19]. The general radiomics workflow can be broadly summarized into four main steps: image acquisition, region of interest (ROI) segmentation, feature extraction, and analysis [Figure 1]^[21]. Image segmentation consists of the delineation of the boundaries of a ROI, such as a tumor and adjacent anatomical structures. This step can be done manually, semi-automatically, or fully automatically with deep learning algorithms. Manual segmentation and semi-automatic segmentation, though time-consuming, have been the most encountered methods of segmentation in current literature^[22]. While these methods of segmentation are prone to introducing inter-observer biases stemming from inconsistencies in the delineation of the boundaries of ROIs, prior studies have uncovered conflicting results on the influence this actually has on radiomics features^[21,23]. Once ROIs have been segmented, the radiomic features are extracted via the conversion of imaging data into quantifiable features, such as signal intensity, shape, texture, and higher-order features. Signal intensity features are obtained through analysis of histograms of individual voxel signal intensities and offer insights into the central tendency of the distribution. Shape and texture features are both calculated in three dimensions by considering the correlation of signal intensities of surrounding voxels. Higher-order features may also be extracted following the application of secondary wavelet or Gaussian filters^[22]. The number of features extracted through these processes can significantly vary depending on filter and software specifications. Overfitting of the model may occur in cases where a high number of features are coupled with a low number of cases in a classification task. The next step of the workflow, feature selection, mitigates this risk by selecting relevant features using techniques such as random forest algorithms and by excluding non-reproducible features. Relevant features may then undergo subsequent analysis using machine learning algorithms.

APPLICATION OF RADIOMICS IN PANCREATIC CANCER

Pancreatic cancer is one of the leading causes of cancer-related deaths globally, with an estimated 5-year survival of approximately 12%^[24]. As compared to other malignancies, the incidence of PC is on the rise while the outcomes remain poor. Over the last four decades, minimal improvement in survival has been observed, the only major development being the introduction of multiagent systemic therapies in the last decade^[25]. The two key challenges that are faced in the management of these patients are the lack of screening tools, resulting in a delay in diagnosis and limited biomarkers that can help guide management in

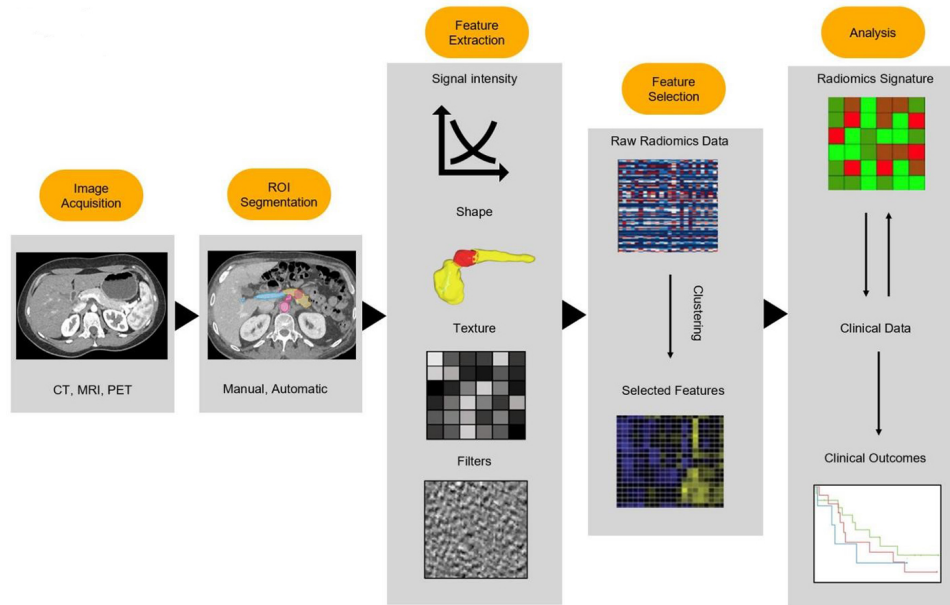


Figure 1. Current approach to radiomic analysis. CT: Computed tomography; MRI: magnetic resonance imaging; ROI: region of interest.

terms of selection, duration, and sequencing of therapeutics^[26]. Oncological resection in conjunction with systemic therapy provides the best chance at cure.

A majority of patients are asymptomatic or present with non-specific symptoms such as abdominal pain and weight loss. Implementation of screening via magnetic resonance imaging (MRI) and endoscopic ultrasound (EUS) has shown promise in high-risk populations; however, implementing them across the entire population is not feasible due to the relatively low incidence of disease. Upon suspicion of disease, computed tomography (CT) coupled with EUS with fine needle biopsy can diagnose the disease. Given current approaches, only 20% of patients are diagnosed with resectable disease, and therefore, better screening tools are required. In terms of therapeutic selection, the only clinically available biomarker is carbohydrate antigen 19-9 (CA19-9), but it has several limitations. Firstly, it can be elevated in non-neoplastic diseases such as pancreatitis and biliary obstruction. Second, approximately 20% of patients are non-producers, rendering its use to estimate tumor biology infeasible. Third, biomarkers to predict sensitivity or resistance to systemic therapies are not available. As a result, treatment response can only be estimated several months into the administration of these therapies when the disease has typically progressed to an incurable stage.

As described earlier, radiomic features represent tumor characteristics and can be applied to various aspects of the management of pancreatic cancer, which are described as follows.

Diagnosis and surgical resectability

Early diagnosis of disease presents one of the greatest challenges in the management of pancreatic cancer. Radiomics has shown to be a promising tool for the differentiation of normal pancreatic tissue from pancreatic cancer. Application of radiomics has demonstrated strong discriminatory ability with the reported AUC [area under the receiver operating characteristic (ROC) curve] of 0.79-0.99^[27-29]. Beyond normal pancreatic parenchyma, radiomics has also been employed in differentiating pancreatic cancer from various mimicking lesions, most notably autoimmune pancreatitis (AIP). Studies of CT-based radiomic

features have demonstrated an accuracy of 85%-95.2% in differentiating AIP from PC^[30-34]. Similar studies have been performed to differentiate PNETs from PDACs and SPT, achieving AUCs ranging from 0.86-0.99^[35-38]. Of note, these studies included atypical hypovascular NF-PNETs, which more closely mimic PDAC than typical PNETs, and radiomics analysis outperformed clinic-radiological factors^[36,37,39,40]. Radiomics combined with machine learning methods would potentially result in the development of tools that will allow for radiomics-based screening for asymptomatic pancreatic cancer. Though the sensitivity of CT in the detection of PDAC ranges from 76%-96%, early CT findings of PDAC, such as tumoral heterogeneity and loss of fatty marbling, can be particularly subtle and may be missed even by experienced radiologists^[28]. Radiomics poses an avenue for quantitative analysis and detection of these changes. Radiomics analysis may also autonomously run in the background of scans and automate the process of screening^[41]. This can effectively enable every abdominal CT scan, regardless of indication, to be used to screen for PDAC. The improved quantitative analysis of images combined with an increase in the sheer volume of scans being screened through this approach makes radiomics a suitable tool for screening for PDAC. Through this, it may be possible to detect disease at an earlier stage when a larger proportion of patients are amenable to surgical resection.

Pancreatic cysts present a diagnostic challenge and comprise a heterogeneous group of lesions with biological behavior varying between benign indolent lesions and a propensity for progression to invasive cancer. Radiomics has been applied to differentiate cyst types. In one of these studies, the authors were able to differentiate serous cystadenomas and pancreatic cystic lesions with an AUC of 0.77 based on 22 radiomic features, which outperformed clinical and standard imaging features (AUC: 0.71)^[42]. In another study, Xie *et al.* were able to differentiate macrocystic SCNs and MCNs with an AUC of 0.99^[43]. Available literature on studies applying radiomics to pancreatic cystic lesions was summarized by Machicado *et al.*, who reported AUCs between 0.77 and 0.99 in differentiating different cyst types^[44]. Mucinous cystic lesions of the pancreas are precursors to pancreatic cancer, but not all of these patients will go on to develop cancer^[10]. Therefore, risk stratification is essential if we are to resect lesions prior to their progression to cancer while avoiding surgery in patients with benign lesions given the high morbidity and mortality associated with these procedures. Radiomics has been applied to the characterization of these cysts and has shown an accuracy of 84% in distinguishing between common types of pancreatic cysts^[45].

Currently, work is being performed on assessing the utility of radiomics in detecting high-grade dysplasia in these lesions and predicting the risk of progression to pancreatic cancer. Tobaly *et al.* trained a radiomics model based on preoperative CT to differentiate low-grade dysplasia, high-grade dysplasia, and invasive cancer in patients with IPMN and demonstrated an AUC of 0.84 and 0.71 on internal and external validation, respectively^[46]. Similarly, Polk *et al.* combined radiomics with conventional variables (thickened and enhanced cyst wall and enhanced mural nodule) and reported an AUC of 0.93 (95%CI: 0.85-1.0) in differentiating low-grade dysplasia and high-grade dysplasia or invasive cancer^[47]. If these tools are developed, we will be able to accurately screen high-risk patients and recommend appropriate care.

Radiomic-based analysis could also improve surgical planning. Determination of resectability and the likelihood of margin negative resections, particularly in the setting of neoadjuvant therapy, is challenging, and the ability of conventional pancreas protocol CT (PPCT) to determine this remains low^[48,49]. This is driven by the fact that it is difficult to differentiate dead tissue and viable tumors on a PPCT. Since radiomics provides a greater deal of information regarding the tissue in the region of interest, studies have applied radiomics to detect the presence of vascular invasion and predict positive resection margins and demonstrated improved prediction compared to conventional PPCT^[50-52]. Recently, Schlanger *et al.* systematically reviewed studies employing artificial intelligence and machine learning across two categories:

preoperative diagnosis and patient evolution; five studies focused on the prediction of complications^[53]. Two of these studies developed imaging-based models, utilizing CT texture features and MRI-based features, respectively, to predict postoperative pancreatic fistula^[54,55]. Capretti *et al.* similarly harnessed 100 preoperative CT scans to develop a model to predict clinically relevant postoperative pancreatic fistula with an AUC of 0.807, which was also able to predict postoperative overall complications and length of stays with AUCs of 0.690 and 0.709^[56].

Characterizing tumor biology

One of the most promising applications of radiomics has been its ability to noninvasively assess and characterize tumor pathology. Considerable work has been done in correlating tumor grade with CT and MRI radiomic features. The majority of current work has been conducted on radiomics-guided grading in pancreatic neuroendocrine tumors and has found a remarkable predictive ability for radiomic analysis to preoperatively distinguish WHO grade 1 from WHO grade 2 or 3 lesions (AUC: 0.736-0.902)^[57-61]. Similar results have been reported for pancreatic cancer (AUC: 0.91-0.994)^[62,63]. Radiomic features have been shown to be able to accurately determine the preoperative risk of lymph node and liver metastases and superior mesenteric vein and portal vein invasion in pancreatic cancer with an AUC of 0.841-0.912^[51,64-67]. Molecular analysis can allow for personalized approaches to the management of pancreatic cancer; however, it is currently limited by cost and the need for invasive procedures to acquire tissue^[68]. Radiogenomics is an area of great interest where radiomic features can be used to determine the mutational profile of these lesions. Current studies on radiogenomics in pancreatic cancer have identified key radiomic features that are highly predictive of genetic alterations in SMAD4, KRAS and p53, HNF1a, and KRT81 status^[69-72].

Predicting treatment response and prognostication

As discussed earlier, real-time assessment of treatment response is essential in the management of pancreatic cancer. However, currently available clinical tools fail to provide accurate assessment, resulting in delays in tailoring of therapy that enables us to derive maximum benefit from the administered therapies while minimizing toxicity. In this regard, studies have assessed radiomic features in a longitudinal manner across chemotherapy, known as delta radiomics^[73]. These studies have reported a strong association between delta radiomics and treatment response gauged by changes in CA19-9, RECIST criteria, and grade of response observed on histopathological examination of the surgical specimen^[74-79]. However, current studies are limited by the heterogeneity of the radiomic features found to be significantly associated with treatment response, the small sample size of the study population, and the single institution nature of these studies^[73]. Future studies must consolidate current evidence by analyzing the various radiomic signatures used across the studies to create a core set of validated radiomics signatures that can be broadly translated to clinical practice. Lastly, radiomics has shown promise in prognostication. Radiomic models based on features extracted from pretreatment scans have been shown to be highly predictive of overall survival, local control, and recurrence and have outperformed current clinical prediction models with an AUC of 0.78-0.92^[80-83].

APPLICATION OF RADIOMICS IN HEPATOBILIARY MALIGNANCIES

Diagnosis and surgical resectability

Unlike most solid cancers, the diagnosis of HCC & intrahepatic cholangiocarcinoma (ICC) can be established without biopsy confirmation^[84]. As a result, the role of noninvasive imaging is of particular importance in the diagnostic workflow of these malignancies. While HCC most often presents with highly characteristic and specific imaging findings, it can occasionally be challenging to discern definitively on imaging from other mimicking hypervascular lesions such as hemangiomas, adenomas, and focal nodular hyperplasia (FNH)^[85,86]. Radiomic-guided studies have demonstrated efficacy in differentiating benign liver lesions from malignant ones with an AUC of differentiating FNH from HCC ranging from 0.917 to 0.971, an AUC for differentiating hemangiomas from HCC of 0.83-0.96 and a sensitivity and specificity for

accurately classifying adenoma 0.80-0.93/0.78-0.93^[87-91]. Another study distinguished combined hepatocellular cholangiocarcinoma (C-HCC) from HCC with an AUC of 0.79-0.81 and outperformed current diagnostic criteria when used in conjunction to differentiate intrahepatic cholangiocarcinoma from HCC^[92]. Radiomic features have also been applied to distinguish primary and metastatic liver tumors, with one study reporting an accuracy of 83% in differentiating metastatic liver disease from primary tumors^[93]. Differentiating neoplastic portal vein involvement and thrombosis from benign portal vein thrombosis is also essential in determining the underlying HCC's resectability. Canellas *et al.* were able to reliably distinguish neoplastic and bland thrombi using radiomics signatures based on thrombus density values^[94]. Moreover, microinvasion, encompassing infiltration of portal vein, hepatic vein, or bile duct, is considered a marker of poor prognosis after hepatic resection and transplantation^[95]. Several studies have also demonstrated promising results of select CT and U/S-based radiomics signatures in accurately predicting the presence of microvascular tumor invasion^[96-99]. For ICC, in a study of 203 cases, radiomics-based models demonstrated the ability to predict futile resections with greater AUC (0.804) than traditional clinical risk factor-based models (AUC: 0.590)^[100].

Characterization of tumor biology

Traditional biopsies are restricted to the small sample of tumor that is biopsied, which may not be representative of the tumor at large^[101,102]. With radiomics-based tumor grading, the entirety of the tumor is evaluated when being graded, thus providing physicians with greater certainty regarding tumor characteristics^[103]. Applications of radiomics in this regard have demonstrated promise, with studies reporting that radiomics signatures strongly correlate with pathological grade of HCCs, allowing for rapid, noninvasive tumor grade determination^[104]. Subsequent research has since further correlated radiomic signatures with tumor Ki-67 status, demonstrating the utility of radiomics in assessing HCC proliferation indices entirely noninvasively^[105].

Treatment selection

Liver transplantation and liver resection represent potentially curative options for early-stage HCC; however, most patients present with later-stage HCC, rendering them ineligible for these options. As such, physicians and patients must choose from a variety of locoregional therapies, including radiofrequency ablation (RFA), transarterial chemoembolization (TACE), and transarterial radioembolization (TARE), all of which have particular advantages and disadvantages^[106,107]. In recent years, there have been investigational efforts to harness radiomics to aid in this challenging clinical decision-making process. For instance, one group employed a DL-based radiomics strategy to develop nomograms for predicting progression-free survival among HCC patients undergoing liver resection or RFA (c-index = 0.726 for RFA, 0.741 for resection)^[108]. To identify future candidates for RFA, another group calculated a radiomics signature based on textural features extracted from patients with significantly longer progression-free survival after RFA ($P = 0.008$)^[109]. Suh *et al.* demonstrated the utility of a radiomics-based signature as a feasible barometer for stratifying and determining patient suitability for liver resection vs TACE^[110]. In unresectable cases, radiomics has also demonstrated efficacy in predicting the development of progressive disease following TACE^[111,112]. In this non-surgical cohort of patients, radiomics analysis has also demonstrated the ability to accurately predict tumor immunoscores and immune phenotypes^[113,114]. This data has been validated to tumor responsiveness to novel immunotherapy and can help make treatment decisions^[114].

Prognostication

Multiple studies have reported the value of applying radiomics to predict outcomes in hepatobiliary diseases. These studies demonstrated that the addition of radiomics to conventional clinical variables increases the accuracy of predicting early recurrence of disease, disease-specific recurrence, and long-term mortality as compared to clinical variables alone with AUCs ranging from 0.59 to 0.91^[107,115,116]. For

comparison, a prognostic nomogram incorporating age, N stage, M stage, tumor size, and surgery yielded AUCs of 0.909 and 0.890 for predicting 1-year and 5-year survival, respectively^[117]. Similar studies have demonstrated certain radiomics features to accurately and independently predict post-surgical overall survival^[118,119]. Seeing as HCC recurrence rates range from 25% after liver transplantation to over 70% after liver resection or RFA^[120], numerous studies investigating the potential of radiomics in predicting recurrence risk after various treatments are also underway. In a multi-institutional study, based on contrast-enhanced computed tomography (CECT) analysis, preoperative and postoperative ML-based radiomics models were developed to predict HCC recurrence after liver resection, and both exhibited better prognostic performance compared to the current staging system (c-index = 0.733-0.801)^[121]. Another study used CECT images from 184 patients to derive radiomics signatures of risk-straying patients based on their recurrence-free survival after RFA (c-index = 0.755)^[122]. Other groups have also utilized DCE-MRI to develop radiomics models; for instance, one study compared the prognostic performance of radiomic scores for the tumor region, perilesional region, non-tumoral region, which independently had the highest prognostic score (c-index = 0.72); the combined score of the three regions yielded a c-index of 0.83^[123]. Focusing on patients with unresectable disease, a separate study group used preoperative multiparameter MRI (mp-MRI) to create a radiomics model to predict early progression after TACE (AUC: 0.800)^[106]. Furthermore, studies suggest that models combining radiomic and clinical input may provide the best performance. For instance, one study established a pretreatment CT-based radiomics model for overall survival after TACE and reported that a composite model was more predictive than a radiomics or clinical signature alone (HR: 19.88, 95%CI: 6.37-92.02, $P < 0.001$)^[106].

Similar efforts to estimate treatment response and survival for CCA, which remains a highly lethal hepatobiliary malignancy, have been conducted. A number of studies have radiomics-based models to predict lymph node metastasis (LNM) in ICC patients with validation cohort AUCs ranging from 0.80 to 0.89^[124-126]. Two studies have developed radiomics-based models for preoperative prediction of microvascular invasion (MVI)^[127,128]. In another study focusing on LNM in ECC, an MRI-based radiomics model achieved an AUC of 0.9 in predicting LNM^[129]. Other researchers have explored recurrence after surgery with implications for informing surveillance schedules and any adjuvant therapy. For example, one group employed random forest and logistic algorithms to develop clinical, radiomic, and combined models; the radiomics model had higher AUCs (sensitivity 0.846, specificity 0.771 in the validation cohort) than the clinical model and comparable AUCs to the combined model^[100].

Altogether, current literature underscores that the incorporation of radiomics may enhance models to predict response to various treatments, thereby guiding appropriate treatment selection and sequencing. In facilitating more informed clinical decision-making, radiomics models will hopefully help improve patient outcomes for hepatobiliary cancers.

CURRENT PITFALLS

Despite promising results, the field of radiomics is still in its relative infancy. As of now, there are still certain critical limitations across the majority of literature on radiomics in hepatobiliary and pancreatic diseases, which need to be addressed prior to its clinical integration [Table 1].

First, a vast majority of existing literature on radiomics comprises retrospective single-institution series with relatively small sample sizes. Currently, a majority of these radiomics-based models have not been validated in other datasets and are therefore not generalizable. Second, prior to the extraction of radiomic features from images, the areas of interest within those images must be delineated, a process known as segmentation^[22]. Current segmentation methods are either manual, semi-automated or fully automated.

Table 1. Current limitations of radiomics

No.	Limitations
1	Limited validation of radiomics models
2	Segmentation required with most literature based on manual segmentation techniques (1) Operator dependent (2) Time intensive (3) Introduces inherent variation that radiomic features may be sensitive to
3	Lack of standardization in the workflow of radiomics
4	Radiomics features are sensitive to variations in imaging protocols, which limits multicentric generalization

Most literature on radiomics is based on manually segmented datasets. Manual segmentation introduces human error and is highly operator-dependent^[22]. In the case of hepatobiliary and pancreatic malignancies, particularly pancreatic cancer, these tumors have irregular boundaries, which makes segmentation even more difficult. Although variations in segmentation methods have been linked to introducing specific differences in radiomic signatures, no current consensus exists on the segmentation techniques. Deep-learning-based segmentation has shown promise in bridging this gap; however, it depends on the accumulation of large datasets to train these models. Work is underway; however, this has not yet been achieved. Additionally, manual segmentation is time-intensive and requires oversight by trained radiologists, thus hampering the feasibility of using it in the clinical setting. Third, in order to eliminate bias, mitigate confounding factors, and encourage reproducibility, the radiomics workflow of studies should be scrutinized and entirely standardized. Many current studies have ambiguous descriptions regarding their process of feature extraction algorithms, mathematical definitions, inconsistencies in feature nomenclature and pre-processing methodology^[130]. This has been the source of skepticism from clinicians who perceive the radiomics model as a black box that generates satisfying clinical prediction results for any given clinical outcome^[130]. Moving forwards, a clear and transparent description of all these processes, alongside any feature reduction or exclusion, should be attempted. Lastly, radiomic features have been demonstrated to be sensitive to variations in image acquisition protocols^[22,131]. Even variations in the time at which contrast-enhanced images are captured following the administration of contrast have been shown to significantly affect the radiomic features of acquired images^[132]. Due to this, imaging protocols must be standardized across all centers evaluating radiomics. Efforts are underway to address these shortcomings, one example of which is the introduction of the radiomics quality score (RQS), a generalizable tool to assess the quality of radiomic studies, which is now being integrated into a majority of studies in the field^[133].

FUTURE DIRECTIONS AND CLINICAL INTEGRATION

Despite limitations, current results of radiomics-based studies are encouraging and denote a promising future for the field. As the radiomics workflow becomes more standardized and the radiomics process becomes more easily reproducible, significant clinical implications can be expected in the following domains.

As a screening tool, radiomics has already demonstrated significant potential in differentiating normal tissue from malignancy. With currently ongoing work on automated deep learning-based radiomics workflow, it can be expected that tools that can accurately screen for hepatobiliary and pancreatic malignancies will be developed. Upon validation, these can then be made available for widespread clinical integration. This tool could be run on radiographic images to autonomously highlight suspicious areas of interest and bring them to the attention of a reading radiologist, providing a probability assessment of potential early malignancy.

For diagnostics, current diagnostic criteria consist of varying combinations of pathology- and radiology-based input to reach a definitive diagnosis. Both these methods, however, are limited by their current accuracy in differentiating lesions and human error in the workflow. Radiomics has already demonstrated the potential to accurately diagnose malignancies through their unique radiomic footprints. This is of particular importance in pancreatic lesions where there is a significant overlap in radiological features between these lesions. Additionally, radiomics has demonstrated the ability to noninvasively predict tumor characteristics with high accuracy, including tumor grade and the presence of nodal disease. In particular, for tumors that demonstrated intertumoral heterogeneity, such as pancreatic neuroendocrine tumors, pathological assessment is limited to the tissue obtained on biopsies and might fail to provide information about the whole lesion^[134,135]. Radiomics allows for “virtual biopsies” that can overcome this by capturing the spatial heterogeneity of the entire tumor, enabling optimal characterization. Furthermore, the noninvasive nature of radiomics can result in repeat assessment at varying intervals for patients who are on surveillance. Another interesting application in tumor characterization is the emerging field of radiogenomics. Albeit limited, the literature available on this has suggested that radiomics can be used to genetically characterize these tumors. As this field develops, radiomics has the potential to allow for accurate diagnostics and tumor characterization that is superior to the current diagnostic modalities.

Radiomics has also demonstrated potential in predicting response to systemic therapies noninvasively. Current biomarkers fail to accurately predict response to administered therapies, resulting in delays in tailoring of therapies in a timely manner. If radiomics-based assessment of treatment response were possible, it would allow for timely modification in systemic therapy, thus maximizing the chance for complete or near complete treatment response and improved survival. In terms of surgical planning, radiomics has demonstrated the ability to predict the presence of vascular invasion more accurately than experienced radiologists. Lastly, radiomics has shown promise in prognostication via accurate prediction of postoperative complications, and recurrence-free and overall survival. Accurate prognostication is essential not only in guiding management, as previously discussed, but also in managing patient expectations through the course of their care. Radiomics-based prognostication has outperformed a variety of existing prognostic factors. In studies where radiomics has not independently outperformed existing factors, the addition of radiomics to established factors enhanced the accuracy of the existing models. While it is unlikely that radiomics models will ever independently be used to prognosticate patients, it is very likely that models combining radiomic features and clinical factors will be used across all metrics of prognostication in the future.

CONCLUSIONS

In conclusion, the field of radiomics is advancing rapidly and has shown promise as a tool for early detection, tumor characterization, therapeutic selection, and prognostication in hepatobiliary and pancreatic malignancies. Despite its shortcomings, we believe that with improvements and automation of segmentation techniques, optimization of radiomic analyses, and introduction of standardized guidelines for research on radiomics, the tools developed using this technology will become more robust. Clinical application of these tools will provide precision in the management of these patients, resulting in improvement in patient outcomes.

DECLARATIONS

Authors' contributions

Made substantial contributions to the conception and design of the study and performed data analysis and interpretation: Grewal M, Ahmed T, Javed AA

Performed data acquisition, as well as providing administrative, technical, and material support: Grewal M, Ahmed T, Javed AA

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Conflicts of interest

All authors declared that there are no conflicts of interest.

Ethical approval and consent to participate

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Consent for publication

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