

Review

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Review

Artificial Intelligence and Rectal Cancer: Beyond Images

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Abstract: *Introduction.* Cancer variability plus medical big data can be handled by artificial intelligence. Its models can have different input types: images and many others (as numbers, predefined categories, free texts). The non image elements are as important as images, as per clinical practice and literature. This article reviews such models, with non image component, applied as use case to rectal cancer. *Results and Discussion.* Secondary literature analysis shows all reviews focusing on image inputs only. Primary literature models of interest include pure (only non image) and combined (both non image and image) inputs. Non image models show significant performance. Combined models frequently exhibit better behavior than their unimodal parts. *Conclusion.* To the best of our knowledge, no reviews focus on non image inputs, either alone or combined with images. Non image components require instead substantial attention, in optimal clinical practice and research. Multimodality –beyond images– is important, in rectal cancer and possibly other pathologies. *Methods.* Literature search was performed on PubMed, without temporal limits, in English, using ample keywords; for secondary literature, appropriate filters were employed.

Keywords: artificial intelligence; machine learning; deep learning; images; unstructured data; structured data; electronic health records; real-world data; combined models; big data; multivariate models; predictive models; digital medicine; personalized medicine; precision medicine; rectal cancer

1. Introduction

Rectal cancer is a world wide leading cause of morbidity and mortality. It is the 8th highest cancer type as per incidence ($\approx 700,000$ new cases per year) and has significant mortality ($\approx 300,000$ deaths per year) [1].

Cancer, in general, is still a fundamental burden in human society, despite significant technological and clinical progresses. Globally, it is the second leading cause of death [2]; furthermore, its trend is increasing [1]. One of the main reasons for cancer burden is its heterogeneity. Variability spans multiple levels: disease level (100+ cancer types), inter patient level (each patient is unique, as per history and current status) and even intra patient level (cancer evolves dynamically, particularly as per its genome). This daunting complexity hinders both its biological understanding as well as its clinical management (prevention, diagnosis, therapy and prognosis).

Fortunately, a very promising paradigmatic shift is ongoing in medicine (particularly, in oncology): evidence based medicine is gradually being complemented by personalized medicine³. Evidence based medicine –more traditional and established, based on clinical trials– is fundamental in the development of medical knowledge, having generated the standard of care in the last three decades [3,4]. Nonetheless, its methodology generates evidence for an ‘average patient’ (‘one size fits all’ approach), with application to the actual, unique patient being problematic [3,5]. Personalized medicine –more modern, with its precision and individualized perspective– can usefully complement

evidence based medicine, particularly in oncology. Certain personalized medicine enablers –big data (particularly real world data), and artificial intelligence (particularly machine learning)– hold great promise [3].

Big data are digital data created in enormous amount ($\approx 10^{12}$ terabyte/year, with growing trend). Big data, to be so defined, should have not only high ‘volume’ (quantity); but also high ‘variety’ (multiple variables and sources), ‘velocity’ (acquisition speed) as well as ‘veracity’ (quality-related parameter). Medical big data have enormous potential to improve research and clinical practice [6]; but they need to be elaborated by proper mathematical and computational methodologies. Particularly, AI –when rigorously designed, developed and validated [7]– can successfully handle them [8].

Artificial intelligence is the imitation of human intelligence by machine computing [9]. An artificial intelligence model is a mathematical relationship between an input (the data, with one or more independent variables) and one or more outputs (the dependent, pursued variables); in the typical situation, independent variables (also known as features, predictors or covariates) are many (with cardinality in the order of magnitude of tens or even hundreds), with only one dependent variable (also known as outcome, target or label). Inputs can mainly be regrouped into ‘structured’ and ‘unstructured’. Structured data –including numbers and pre-defined categories– have higher standardization and usually data quality. Unstructured data –including images and uncategorized (freely typed) text– usually need to be mined (e.g. radiomics for images and natural language processing for texts). Particularly, radiomics is an approach which aims to extract structured quantitative information (consisting in even several hundreds of predictors) from unstructured medical images [11].

Machine learning is a notable and intensively researched subset of artificial intelligence. Machine learning consists in mimicking human learning by computer algorithms [10]. These algorithms are able to ‘learn’ features and patterns directly from the input data without being explicitly programmed. Thus, machine learning –both the model structure and its subsequent performance– is strongly influenced by the data, those given as input (training) and those employed to validate (evaluation). Generally speaking, increasing the quantity of such data (i.e. the sample size) tends to provide more reliable results, as in statistics, an advantage of big data over ‘traditional’ data. But, simultaneously, big data are also required to have sufficient data quality, in order to give reliable and realistic results; as such, they need curated and standardized data collection [11,12].

Machine learning is mainly categorized into supervised and unsupervised learning (a hybrid methodology –semi supervised– is also possible). Supervised learning deals with labeled data (data for which the outcome has given, known values, called ‘labels’); unsupervised one with unlabeled data (data for which either outcome labels are too difficult/expensive to obtain or even it is complicated to identify an actual outcome in the data set). Supervised machine learning can be further divided into ‘classification’ (when the outcome is qualitative, i.e. categorical) and ‘regression’ (outcome is quantitative, i.e. numerical): in classification, the model predicts to which outcome class (category or group) each data item (in general object or entity, in medicine patient) belongs; in regression, it estimates, for each of them, a number, which usually has the mathematical meaning of a probability. Some algorithms can also perform both classification and regression tasks. Such models include decision trees (and random forests, which are their ensemble version) and support vector machines. Other notable model classes include general linear models (which can be employed for regression, as linear regression, or classification, as logistic regression, whose name is misleading) and Cox regression (which handles temporal data). Least absolute shrinkage and selection operator (LASSO) involves linear regression with feature selection (regularization).

A noteworthy subset of machine learning is deep learning, which employs artificial neural networks with more than one hidden layer. Artificial neural networks are machine learning algorithms inspired by human brain: they are composed of relatively simple units (called ‘artificial neurons’, imitating biological neurons), which are networked together in a complex fashion (Figure 1).

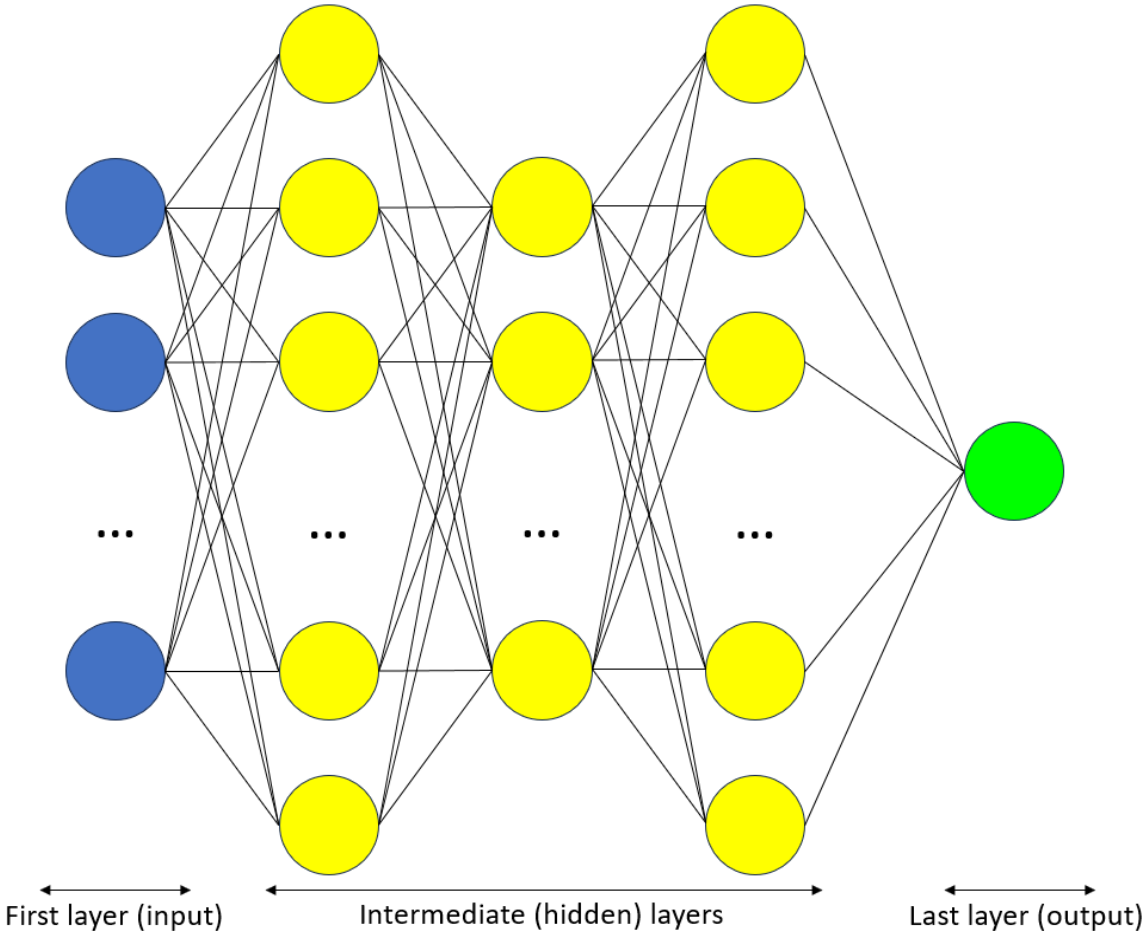


Figure 1. Schematic representation of a typical artificial neural network , with multiple hidden layers (i.e. deep learning). Circles represent perceptrons.

Perceptrons are usually grouped into ‘layers’, each performing a specific task; ‘hidden’ layers are all besides the first and the last ones, which are the input and the input, respectively. Because of its architectural connectivity, deep learning models can accomplish complex tasks, as classification and generation. In medicine, it is mainly applied to images.

Medical images –such as imaging, histology and endoscopy– are fundamental, for both diagnostic and therapeutic purposes [13]. But images are not the only data type employable in medicine, to support decision making: structured data, as well as unstructured data different from images, are also essential6. Real world data [14,15] –defined as all data generated outside clinical trials[3,16]– are an important source for evidence generation. Particularly, electronic health records (or electronic medical records) –as medical reports and nurse diaries– can be a mine of patient information [17]. Medical big data are hence intrinsically multimodal, i.e. are of heterogeneous types and come from a plethora of sources.

Multimodality is also a property of oncology, particularly for RC [3,5]. Multimodal integration can be beneficial for the whole medical workflow, in both clinical practice and research: disease understanding, prevention, diagnosis, therapy, prognosis and follow-up. The integration of different roles –including but not limited to physicians, nurses, biomedical technicians and data scientists– is of paramount importance for advancing knowledge and for improving healthcare. Medicine is a multi-source, synthetic, holistic process: it cannot be reduced to images only.

Nonetheless, to the best of our knowledge, all reviews on artificial intelligence applied to rectal cancer focus on models whose input are based solely on images (herein called ‘image models’). This seems to contradict not only the above mentioned medical holism, but also the literature. Indeed, models with only non image input (herein termed ‘non image models’) can show high performance,

e.g. in pan cancer application, entirely based on RWD/EHRs [18], or in RC specifically, as presented in the *Results and Discussion* section. Moreover, there are also ‘multimodal’ models, which are multivariate models (analyzing multiple variables) with multiple variable types, i.e. artificial intelligence models whose input is simultaneously based on images and non image components. Such models (in the following named ‘combined models’; in the literature also known as ‘hybrid models’ or ‘holistic models’) can exhibit higher performances than their unimodal, constituting models [6,11,19], highlighting even more the importance of multimodality.

While three of them [11,12,20] only briefly mention the existence of combined models, all reviews found [11,12,20–36] focus solely on image models. Since non image inputs –integrated (combined models) or not (non image models) with images– are as essential as image inputs, reviewing the existing primary literature about them is needed.

This Review focuses on original articles reporting artificial intelligence models (particularly machine learning algorithms) that have some non image input and that are applied to rectal cancer, without restrictions on study outcome.

2. Results and Discussion

Tables 1 and 2 present primary literature ($N = 23$ articles, for a total of 11,941 patients) dealing with artificial intelligence applied to rectal cancer, with different outcomes, with at least one non image input, so both non image models and combined models.

Table 1 contains general information, while Table 2 provides more in-depth analysis.

Table 1. Literature about non image artificial intelligence with application to rectal cancer.

Reference	First author	Year	Journal	Aim	Number of patients			TRIPOD	AI model(s)
					Total	Training	Validation		
NIMs									
37	Peng J	2014	PLOS One	Predict OS, DMs (and LR)	917	833	84	2b	Cox regression
38	Sun Y	2017	Journal of Surgical Oncology	Predict DMs after nCRT	522	425	97	2b	Cox regression
39	Valentini V	2011	Journal of Clinical Oncology	Predict OS, LR, DMs	2795	2242	553	3	Cox regression
CMs									
40	Chen LD	2020	European Radiology	Predict TDs	127	87	40	2b	ANN
41	Cheng Y	2021	Abdominal Radiology	Predict response (particularly, pCR) to nCRT	193	128	65	2a	Logistic regression
42	Cui Y	2019	European Radiology	Predict response (particularly, pCR) to nCRT	186	131	55	2a	Logistic regression
43	Dinapoli N	2018	International Journal of Radiation Oncology	Predict response (particularly, pCR) to nCRT	221	162	59	3	GLM
44	Ding L	2020	Cancer Medicine	Predict preoperative LN metastases	545	362	183	2a	Logistic regression, DL
45	Huang YQ	2016	Journal of Clinical Oncology	Predict preoperative LN metastases	526	326	200	2b	Logistic regression
46	Jin C	2021	Nature Communications	Predict response (particularly, pCR) to nCRT	622	321	301	3	DL
47	Kleppe A	2022	Lancet Oncology	Optimize adjuvant therapy	2072	997	1075	3	Cox regression, DL
48	Li M	2021	World Journal of Gastroenterology	Predict PNI	303	242	61	2a	Logistic regression
49	Liu H	2022	Journal of Magnetic Resonance Imaging	Evaluate KRAS mutation	376	288	88	2a	Logistic regression, DL
50	Liu S	2021	Frontiers in Oncology	Detect preoperative EMVI	281	198	83	2b	Logistic regression
51	Liu X	2021	Lancet EBioMedicine	Predict DMs after nCRT	235	170	65	3	DL
52	Mao Y	2022	Frontiers in Oncology	Predict response (particularly, pCR) to nCRT	216	151	65	2a	Logistic regression
53	Peterson KJ	2023	Journal of Gastrointestinal Surgery	Predict response (particularly, pCR) to nCRT	131	111	20	2a	Logistic regression
54	van Stiphout RGPM	2011	Radiotherapy & Oncology	Predict response (particularly, pCR) to nCRT	953	Various groupings		3	SVM

55	van Stiphout RGPM	2014	Radiotherapy & Oncology	Predict response (particularly, pCR) to nCRT	190	112	78	3	Logistic regression
56	Wan L	2019	Abdominal Radiology	Predict response (particularly, pCR) to nCRT	120	84	36	2b	Logistic regression
57	Wei Q	2023	European Radiology	Predict response (particularly, pCR) to nCRT	151	100	51	3 (plus 2a)	RF
58	Wei Q	2023	Abdominal Radiology	Predict preoperative LN metastases	125	80	45	2a	Logistic regression
59	Yi X	2019	Frontiers in Oncology	Predict response (particularly, pCR) to nCRT	134	101	33	2a	SVM, RF, LASSO

Legend. DM = Distant metastasis. EMVI = Extra mural venous invasion. GLM = General linear model. LASSO = Least absolute shrinkage and selection operator. LN = Lymph node. LR = Local recurrence. nCRT = Neoadjuvant chemoradiotherapy. OS = Overall survival. pCR = Pathological complete response. PNI = Peri neural invasion. RF = Random forest. SVM = Support vector machine. TD = Tumor deposit. TRIPOD = Transparent Reporting of a multivariable prediction model for Individual Prognosis Or Diagnosis.

In Table 1, as per aim, most articles address the prediction of the response to neoadjuvant therapy (neoadjuvant chemoradiotherapy, part of the standard treatment of locally advanced rectal cancer); additional purposes include prediction of metastases and other rectal cancer features. Particularly, response prediction is analyzed in 11 (47.8 %) papers; prediction of clinical outcomes (overall survival, local recurrence and distant metastases) and of risk factors (KRAS mutation, tumor deposit, peri neural invasion, extra mural venous invasion) are the purpose of 4 (17.4 %) articles for each subset; prediction of lymph node staging in 3 (13.0 %) papers; prediction of histological examination in 1 (4.4 %) article.

Sample size is between hundreds and thousands of patients per study.

Classification based on TRIPOD (Transparent Reporting of a multivariable prediction model for Individual Prognosis Or Diagnosis) statement [62] is reported. Generally, based on it, one prediction model (particularly, based on artificial intelligence) can be regrouped in this manner: *1a* training only; *1b* training plus validation based on resampling (cross validation, bootstrapping, ...); *2a* training plus validation based on random split (between training and validation data sets); *2b* training plus validation based on non random split (e.g. based on temporal flow of the patients); *3* training plus external validation (i.e. training and validation are implemented upon different populations, centers, institutions, even countries and continents); *4* (external) validation only (this applies to an already published prediction model).

Most artificial intelligence models for rectal cancer that were found belong to the combined model category ($N = 20$), compared to non image models ($N = 3$). Variability among chosen artificial intelligence model (i.e. reason why so many models exist and continue to be used, instead of choosing always an optimal, ideal model) is linked to the so called 'no free lunch theorem' [64]. This fundamental theorem holds (in its original form or in analogous formulations) for many artificial intelligence tools (supervised models, unsupervised models, feature selection techniques, ...). It declares that no artificial intelligence model performance is the absolute best across all scenarios, contexts and data sets. This explain the variety of methodologies found in artificial intelligence. The commonest type of model found (69.6 %), including both non image and and combined models, is regression (either logistic or Cox ones); particularly, most models (52.2 %) have logistic regression type, which could be due to its interpretability, as model coefficients are directly related to the importance of each predictor; if models are ranked by decreasing patient number, the first three models are instead all Cox regression type. Another notable technique is support vector machine (8.7 %).

Table 2. Details about studies listed in Table 1.

Reference	First author	Year	Model input(s)			Performance	Number of centers	External validation(s)?	Note
			Non images?	Images?	CM better?				
NIMs									
[37]	Peng J	2014	Yes (demographic and clinicopathological)			C-Index = 0.73–0.76	1	No	
[38]	Sun Y	2017	Yes (clinicopathological)			C-Index = 0.71	1	No	
[39]	Valentini V	2011	Yes			C-Index = 0.68–0.73	5	Yes	Subsequently re-validated by Reference [60]
CMs									
[40]	Chen LD	2020	Yes (clinical)	Yes (radiomics, ANN, US)	Yes	AUC = 0.80	1	No	
[41]	Cheng Y	2021	Yes	Yes (radiomics, MRI)	Yes	AUC = 0.91–0.94	1	No	
[42]	Cui Y	2019	Yes (from EHRs)	Yes (radiomics, MRI)	Yes	AUC = 0.97	1	No	
[43]	Dinapoli N	2018	Yes (clinical)	Yes (radiomics, MRI)		AUC = 0.75	3	Yes	Subsequently re-validated by Reference [61]
[44]	Ding L	2020	Yes	Yes (DL, MRI)		AUC = 0.89–0.92	1	No	
[45]	Huang YQ	2016	Yes (clinicopathological)	Yes (radiomics, CT)		C-Index = 0.78	1	No	
[46]	Jin C	2021	Yes (blood tumor markers)	Yes (DL, MRI)	Yes	AUC = 0.97	3	Yes	
[47]	Kleppe A	2022	Yes (markers)	Yes (DL, histopathology)	Yes	HR = 3.06–10.71	3	Yes	
[48]	Li M	2021	Yes (clinical)	Yes (radiomics, CT)	Yes	AUC = 0.80	1	No	
[49]	Liu H	2022	Yes (clinical)	Yes (DL, MRI)	Yes	AUC = 0.84	1	No	
[50]	Liu S	2021	Yes (clinical)	Yes (radiomics, MRI)	Yes	AUC = 0.86	1	No	
[51]	Liu X	2021	Yes (clinicopathological)	Yes (radiomics, DL, MRI)	Yes	C-Index = 0.78	3	Yes	

[52]	Mao Y	2022	Yes (clinicopathological)	Yes (radiomics, CT)	Yes	AUC = 0.87	1	No
[53]	Peterson KJ	2023	Yes (clinical [from EHRs])	Yes (radiomics, MRI)	Yes	AUC = 0.73	1	No
[54]	van Stiphout RGPM	2011	Yes (clinical)	Yes (PET-CT)	Yes	AUC = 0.86	4	Yes
[55]	van Stiphout RGPM	2014	Yes (clinical)	Yes (PET-CT)		AUC = 0.70	2	Yes
[56]	Wan L	2019	Yes (clinical)	Yes (MRI)	Yes	AUC = 0.84	1	No
[57]	Wei Q	2023	Yes (clinical)	Yes (radiomics, MRI)	Yes	AUC = 0.87	2	Yes
[58]	Wei Q	2023	Yes (clinical)	Yes (radiomics, MRI)	Yes	AUC = 0.85	1	No
[59]	Yi X	2019	Yes (radiological, clinicopathological)	Yes (radiomics, MRI)		AUC = 0.90–0.93	1	No

Legend. AUC = Area under the ROC curve. C Index = Concordance index. CT = Computerized tomography. HR = Hazard ratio. MRI = Magnetic resonance imaging. PET = Positron emission tomography. ROC = Receiver operating characteristic. US = Ultra sound.

Table 2 shows additional information about articles listed in Table 1, particularly details about artificial intelligence model in terms of its input(s) and its validation.

For those models having an image component (combined models), the most common source of images is imaging techniques, mainly magnetic resonance imaging (65.0 %) and computerized tomography (25.0 %). An intensively employed methodology to deal with images is radiomics; as mentioned above, deep learning is also frequently applied to images.

Performance is expressed through either area under the receiver operating characteristic (AUROC or simply AUC) curve, concordance index (C index, a generalization of AUC [39]) or hazard ratio. Both non image models and combined models show relatively good performances (ranges of AUC and C index of 0.70–0.97 and 0.68–0.78, respectively). Interestingly, in combined models, an integrated model (e.g. deep learning plus pathological staging markers [47]) very often (75.0 %) shows higher performance than its individual components. Both these aspects suggest that multimodality is highly desirable, in agreement with the literature [64]. This can be more general, as it also holds even for images alone, i.e. without any non image input (e.g. where multiple imaging modalities are combined together, within the same model) [65].

Some notes about how performance was reported. If the model is externally validated, then the performance refers to external validation; otherwise, to the internal one. If more than one model is present for the same outcome, the best performing model is presented. If more than one outcome is pursued in the study, performance will be shown as a range.

It can be noted that too many studies (65.2 %) are still monocenter. Multicenter projects tend to be better than the latter, as they tend to provide both better input data, as per both quantity (number of patients, also pooled [39]) and quality (diversity and heterogeneity of populations, lowering overfitting and promoting generalizability), and consequent better performance and clinical applicability of the model. Particularly, although externally validated designs are encouragingly rising, compared to a decade ago, their presence should continue to increase, for the reasons previously stated. To summarize, more multicenter, externally validated, prospective studies are needed [66].

Regarding data preprocessing, most (82.6 %) articles report at least one methodology.

The most reported methodology type (60.9 %) is feature selection, which seems reasonable, given the well known importance of the ‘dimensionality curse’. The latter refers to the need of empirically finding an optimal balance between underfitting (where the proposed model is too simple, hence incapable to accurately describe the data and produce reliable predictions) and overfitting (the model is instead excessively complex, as it copies exactly the training data patterns but it is incapable of properly generalizing them to new, unseen data). This trade off is reached by choosing an appropriate number of model predictors; this number should be related to the number of data items (e.g. quantities as events per variable).

Other reported methods include data normalization and standardization (39.1 %), imputation (13.0%), augmentation (4.3%) and binning (4.3%). Data normalization consists in mathematically ‘compacting’ one variable data to the [0, 1] interval; data standardization involves bringing them to a standard Gaussian distribution (with mean 0 and standard deviation 1). Data imputation is a set of techniques whose aim is to handle missing values, estimating them based on various mathematical methods (the simplest being substituting a missing value in a feature with the feature median or mean). Data augmentation is the synthetic generation of data, which can be particularly useful in the case of class imbalance: in a binary outcome scenario, this would mean that the proportion of data items with a certain outcome value (label) is significantly different compared to that with the other, complementary label.

As previously mentioned, the rationale to explore and analyze the above presented primary articles is that all ($N = 19$) found secondary literature papers [11,12,20–36] focus solely on image models; at maximum, few of them briefly mention the existence of the combined models; but none of them report about non image models. Thus, a detailed review of artificial intelligence models whose input has at least one non image element was needed. This provided the motivation for the present

study. We now present more details about this evaluation of the reviews status: the secondary literature analysis is summarized in Table 3. As a note, almost half of the found reviews (42.1 %) deal with radiomics.

Table 3. Review articles, dealing with artificial intelligence models of rectal cancer, found in our initial secondary literature evaluation.

Reference	First author	Year	Journal	IMs	CMs	NIMs	Notes
[11]	Lambin P	2017	<i>Nature Reviews Clinical Oncology</i>	Main focus	Briefly mentioned		Radiomics
[12]	Meldolesi E	2016	<i>Future Oncology</i>				
[20]	Jia LL	2022	<i>Frontiers in Oncology</i>				Systematic review (with meta-analysis)
[21]	Bedrikovetski S	2021	<i>BMC Cancer</i>				Radiomics
[22]	Coppola F	2021	<i>Diagnostics</i>				Deep learning
[23]	Kalantar R	2021	<i>Diagnostics</i>		Not present		Deep learning; Systematic review
[24]	Kuntz S	2021	<i>European Journal of Cancer</i>				
[25]	Kwok HC	2022	<i>Abdominal Radiology</i>				Radiomics
[26]	Miranda J	2022	<i>Clinical Imaging</i>				
[27]	Namikawa K	2020	<i>Expert Review of Gastroenterology & Hepatology</i>				Deep learning
[28]	Pacal I	2020	<i>Computers in Biology and Medicine</i>				
[29]	Qin Y	2022	<i>Frontiers in Oncology</i>				Radiomics
[30]	Reginelli A	2021	<i>Diagnostics</i>				
[31]	Staal FCR	2021	<i>Clinical Colorectal Cancer</i>				Radiomics; Systematic review
[32]	Stanzione A	2021	<i>World Journal of Gastroenterology</i>				Radiomics
[33]	Tabari A	2022	<i>Cancers</i>				
[34]	Wang PP	2021	<i>World Journal of Gastroenterology</i>				
[35]	Wong C	2023	<i>Journal of Magnetic Resonance Imaging</i>				
[36]	Xu Q	2021	<i>Cancer Management and Research</i>				

Legend. CM = Combined model. IM = Image model. NIM = Non image model.

3. Conclusion

While there are no reviews dealing with artificial intelligence models applied to rectal cancer with non image inputs, the latter –alone (non image models) or together with image inputs (combined models)– can significantly and reliably describe medical data sets, predicting clinical outcomes with high performance, comparable to that of image models. Moreover, combined models usually exhibit better behavior compared with their constituting parts; this points to promote the synergic, holistic usage of multimodal data sources, to more comprehensively describe reality –also complex as human cancer– and find actionable features. Hence, multimodality –in both artificial intelligence and oncological management– is a fundamental approach to benefit patients.

As per clinical perspective on obtained results, rectal cancer is well suited as pathological model to be employed with artificial intelligence methodologies: necessity of multimodal therapies (radiotherapy, chemotherapy, surgery –all three are involved); opportunity to intensify therapies based on risk (adjuvant chemotherapy yes/no, demolitive surgery yes/no, radiotherapy dose escalation yes/no); good healing possibility, hence need to identify patients with either high risk to cure them or low risk to avoid unneeded toxicity. Because of those reasons, the literature has chosen to investigate –as clinical question for artificial intelligence techniques– stratification of outcomes, prediction of risk factors present at histological exam, prediction of lymph node staging and response prediction.

To conclude, within artificial intelligence applied to rectal cancer, it is needed to integrate different aspects: multiple data types (image and non image inputs); several medical procedures (neoadjuvant chemoradiotherapy and surgery); complementary knowledge branches (multidisciplinary clinical board and research team). This can be true, more generally, also for other oncological and non diseases. In the currently ever growing patient-centric viewpoint, it is ultimately beneficial to contemplate sources and features beyond images.

4. Methods

The articles, regarding artificial intelligence applied to rectal cancer, were searched on PubMed website (comprising MEDLINE database plus others), without temporal period limitations, written in English language only, through the following comprehensive search string: *rectal-cancer AND (artificial-intelligence OR machine-learning OR deep-learning OR predictive-model)*. Regarding secondary literature, found articles were internally filtered according to all these three literature types: reviews, systematic reviews and meta analyses.

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