

A comprehensive assessment of artifcial intelligence applications for cancer diagnosis

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Abstract

Artifcial intelligence (AI) is being used increasingly to detect fatal diseases such as cancer. The potential reduction in human error, rapid diagnosis, and consistency of judgment are the primary motives for using these applications. Artificial Neural Networks and Convolution Neural Networks are popular AI techniques being increasingly used in diagnosis. Numerous academics have explored and evaluated AI methods used in the detection of var‑ ious cancer types for comparison and analysis. This study presents a thorough evaluation of the AI techniques used in cancer detection based on extensively researched studies and research trials published on the subject. The manuscript offers a thorough evaluation and comparison of the AI methods applied to the detection of fve primary cancer types: breast cancer, lung cancer, colorectal cancer, prostate cancer, skin cancer, and digestive cancer. To determine how well these models compare with medical professionals' judgments, the opinions of developed models and of experts are compared and provided in this paper.

Keywords Artifcial intelligence · Cancer detection · Cancer diagnosis · Prediction · Deep learning · Machine learning · Image Processing

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1 Introduction

The recent advancement of AI applications in the medical sector has demonstrated success in rapid and precise diagnosis. The assessment of AI approaches used in cancer detection is considered essential since the disease is one of the most deadly and has a signifcant need for diagnosis. Many academics have conducted studies and developed methods to assess the efectiveness of a particular cancer detection technology.

By analyzing and contrasting the six primary cancer types provided in this paper, the primary objective of this study is to provide a broader view of various methodologies and malignancies in a single manuscript. This article is helpful to readers to get a comprehensive idea of cutting-edge AI methodologies, which further stimulates constructive research.

AI has made advances in many medical felds, including diagnosis, treatment, drug development, patient care, etc. Several studies on AI in cancer diagnosis have been undertaken. Cancer is still a leading cause of worldwide fatalities, and hence the error-free diag– nosis of cancer is crucial. AI has proven to be significantly accurate in the imaging diagnosis of tumors (i.e., pathological diagnosis and endoscopic diagnosis). AI assists doctors in providing more accurate and effective medical care to patients (Dong et al. [2020](#page-50-0)).

Correct identifcation necessitates the mining of quantitative information for an informed diagnosis. However, human error is possible, and intricate faults make the issue worse. To efectively treat patients, AI must be used in therapeutic imaging assessment. Image segmentation, image registration, and image visualization are its three major pillars. In advanced countries, the Multidisciplinary Group (MDT), a team of specialists and medi‑ cal professionals, is used to determine the most efective course of therapy for a condition. AI gathers data from numerous sources for a comprehensive analysis and treatment plan.

It is often expected (Shastry and Sanjay [2022](#page-51-0)) that a health professional's diagnosis will be more precise with more information they have about the patient. The quantity of information that is currently available to health practitioners, however, is sometimes seen as overwhelming. Because it may be used to fnd pertinent patterns in complicated data, machine learning provides a solution to this issue. ML is applicable to "precision cancer treatment" for treating cancer patients. Based on the unique genetic profles of each patient's tumors, "precision cancer treatment" seeks to reliably forecast the appropriate drug treatments for a given patient (Injadat et al. [2021\)](#page-50-1).

Deep learning (DL), a branch of machine learning, is employed in drug discovery, identifcation, and diagnostics to simulate human intelligence. It makes use of an Artifcial Neural Network (ANN) to mimic how input is processed by artifcial neurons, how output is generated, and the working of other hidden layers in the network. DL has been used in mechanical surgical treatments for gynecological disorders and heart valve replacement and is expected to play a signifcant role in the fght against cancer (Shastry and Sanjay [2022\)](#page-51-0).

The literature is primarily derived from relevant past studies conducted on AI in cancer diagnosis. This survey consist of 36 research and review-based papers detailing AI applications and their comparisons across the diagnosis of major cancers. This review is created by thoroughly examining the relevant prior research and compiling the data in an organized manner. The primary objectives are to collect information about currently used and leading AI applications, their performance analysis, comparison of diferent study experiments, and understanding of the future scope in a single manuscript. This paper reviews heterogeneous set of studies for diverse cancer types to further analyze AI approaches, indicating a clear conclusion on the usage and infuence of AI techniques in cancer detection.

Furthermore, the utilization of numerous studies enables a clear comparison of methodologies to identify the best-performing models hence stimulating future research.

The studies conducted in the past on the most recent developments in AI for cancer detection primarily emphasized the evaluation of a particular method or experiment that led to the development of a more efective model. Other research experiments compared various approaches to see which ones are more efective against specifc malignancies. This review adopts a different strategy by offering a thorough analysis of 6 significant tumors. The comparison of research is described, tabulated and potential new methods resulting from study experiments have been discussed. The primary medical modalities and AI techniques discussed in the paper are as shown in the Fig. [1](#page-2-0).

In the fight against cancer, AI techniques are powerful tools. AI is gradually being utilized for cancer screening. Automated approaches for detecting cancer are being implemented with the help of AI. Computer algorithms are being used to analyze magnetic resonance imaging (MRI) images, leaving little room for error. AI is also showing an immense

Fig. 1 Medical and AI techniques discussed in the paper for six major cancer types

contribution to drug discovery. Better cancer surveillance is taking place due to the advanced technology ofered by the development of AI (Saba [2020](#page-51-1)). There are a variety of AI models for each cancer type discussed in this paper. Every method uses a different data– set to train and test the model. The training process with each AI technique may use biased input data. In this case, the generalized approach of AI for cancer detection needs to be presented. Data Preprocessing is an important strategy that needs to be performed in such a way that the training data fts the diverse groups. Another strategy is use bias detection techniques during the development of the model and deployment of the model. This important strategy is analysis of the data and the output of the model to identify potential biases.

This paper is organized as follows: First section gives the introduction. This section pre‑ sents an extensive overview of AI applications in five primary malignancies. Section [2](#page-3-0) presents a thorough analysis of the artifcial intelligence methods utilized in the detection of breast cancer. Section [3](#page-8-0) describes lung cancer detection techniques with detailed analysis. Section [4](#page-13-0) presents AI techniques and their analysis for colorectal cancer type. Section [5](#page-21-0) presents prostate cancer detection techniques using AI. Section [6](#page-27-0) describes skin cancer and Sect. [7](#page-35-0) describes digestive cancer detection techniques. Section [8](#page-43-0) presents describes the ML approaches used for personalized cancer treatment using genotypic features. Section [9](#page-48-0) presents a discussion on the all the techniques studied. The authors discusses here the problems with AI techniques and the confdentiality of data. Section [10](#page-49-0) presents conclusion and future research directions.

2 AI techniques for breast cancer

AI-based breast cancer detection techniques leverage the power of artifcial intelligence and machine learning algorithms to analyze medical data including mammograms, MRIs, and ultrasounds for early detection and diagnosis of breast cancer. These algo– rithms recognize the patterns and anomalies within the images that may indicate the presence of tumors or suspicious lesions, often with high accuracy. By continuously learning from vast amounts of data, AI systems can improve their performance over time. AI-based solutions help radiotherapists to become more skilled investigators and aid in the early diagnosis of breast cancer. Mammography, tomography, breast ultrasound, MRI, CT scans, and PET scans are the most frequently utilized tools for the diagnosis of breast cancer (Shastry and Sanjay [2022\)](#page-51-0). Mammography is also known as breast screening. It is highly possible to detect breast cancer during screening requiring a minimal amount of time. Ultrasound technique delivers sound waves inside the body to examine the internal structure of the body. A transducer is used in this technique. This provides sound waves. It is located on the skin. They treat tissues as obstacles and react to them and echo. These echoes are recorded. The echo values are then trans‑ formed to grayscale for digital analysis. The positron emission tomography technique employs F-fuorodeoxyglucose. Imaging the body allows physicians to locate a tumor. The technique has its basis in detecting radiolabel cancer cells. These cells are specifc tracers. MRIs are employed for detection. Elastography is a method that makes it possible to remove breast cancer tissue larger than normal parenchyma. Elastography is used for imaging by sound waves bouncing off the tissues. These waves are recorded, and the values are converted to grayscale for digital analysis. Elastography can differentiate between malignant and benign tumors (Sharif [2021](#page-51-2)). Active microwave imaging is

a newly discovered technique for early detection in breast detection. When microwaves are bombarded with cancer tissues, they exhibit properties very diferent from healthy tissues (Bindu et al. [2006](#page-50-2)).

Mammography has been proven as one of the most useful screening tools for breast cancer. In previous investigations, half of the mammographically identifed breast tumors were evident. Double reading of mammograms have been proven more potent in comparison to a single reading. In double reading, the sensitivity rate is enhanced by 5–15% (Watanabe et al. [2019\)](#page-51-3). 10–30% of breast cancers are missed on mammography, which is ascribed to thick parenchyma hiding lesions, inadequate placement, perception mistakes, and interpretation errors.

The goal of the study by researchers was to create an artifcial intelligence algorithm for mammography breast cancer diagnosis and observe its improved diagnostic accuracy. An AI algorithm was developed and validated. This experiment was carried out using 170,230 mammography examinations from fve institutions in the United States, South Korea, and the United Kingdom. For the multicentre, observer-blinded, reader study, 320 mammograms were obtained independently from two institutions, with 160 being cancer-positive, 64 benign, and 96 normal. 14 radiologists acted as readers, assessing each mammogram for the probability of malignancy, likelihood of malignancy (LOM), location of malignancy, and the need to remember the patient, frst without and then with the AI algorithm's assistance. The performance of AI and radiologists was evaluated using the LOM-based area under the receiver operating characteristic curve (AUROC), as well as recall-based specifcity and sensitivity.

To diagnose breast cancer, an AI system was built that employs the biggest breast cancer data set among existing AI algorithms. The algorithm was able to exhibit comparable performance in validation data sets from diferent nations since it was trained with data from multiple institutions. With the use of large-scale mammography data, the AI program outperformed doctors in terms of diagnosis, notably in early-stage invasive breast tumors. Mammographic characteristics of tumors found by the AI system were analyzed through a comparative study with radiologists to gain a better understanding of AI behavior. This comparison is shown in Fig. [2](#page-4-0).

Fig. 2 Comparison of radiologist observations and AI performance

When compared to radiologists, AI algorithm constructed using large-scale mammography data performed better in breast cancer detection. The considerable increase in radiologists' performance was observed when assisted by AI justifes using AI as a diagnostic assistance tool for mammography. According to this study, AI has the potential to improve early-stage breast cancer diagnosis in mammography. The performance of radiologists was greatly enhanced when assisted by AI, particularly in thick breast regions on a mammogram, which represents one of the primary challenges in screening. Such advancements leads to an increase in screen-detected malignancies and a decrease in interval cancers, improving mammography screening efectiveness. (Kim [2020\)](#page-50-3)

The radiologists' accuracy is compared with AI systems using a non-inferiority null hypothesis (margin $= 0.05$) based on differences in the area under the receiver operating characteristic (AUROC) curve. The sensitivities of the radiologist and the AI system are compared with HMRC using standard analysis of variance (ANOVA) of two multi-reader multi-case (MRMC) modalities with the same level of specificity. The AUC of the AI system (0.840) is statistically non-inferior to that of the 101 radiologists (0.814) with a difference of 0.026 resulting in slightly higher for the AI system for low and mid-specifcity.

Although the AI system has shown better results than the collective performance of the radiologists, it consistently performs low compared to the best radiologist. An ideal AI system should be able to function to the limits of the imaging modality itself, in other words, not be able to detect mammographic occult cancers while minimizing false positives. These systems are suitable for mass population screening to provide better and consistent results in a short time. It is benefcial in countries that lack experienced breast radiologists. The tested AI system based on deep learning algorithms has a similar performance to an average radiologist for detecting breast cancer in mammography (Rodriguez-Ruiz et al. [2019\)](#page-51-4).

From previous mammograms, the radiologists studied and classifed the retrospective data into three categories:

- 1. Actionable: This mammography was found in 155 previous mammograms from 90 individuals.
- 2. Non-actionable: In this category, because the lesion that was subsequently biopsied fell below the recollection threshold, it is not possible to be utilized.
- 3. Excluded: In this category,prior lumpectomy on the ipsilateral side and synthetic or three-dimensional tomography pictures are excluded (Watanabe et al. [2019\)](#page-51-3).

cmAssist is a newly invented AI-CAD for mammography that uses deep learning, a prominent kind of artifcial intelligence. The cmAssist method uses a combination of unique deep learning-based networks to accomplish great sensitivity without losing specifcity. In addition, to enrich the varied appearances of cancer and benign structures in training dataset, the algorithms are trained using a proprietary, patent-pending data enrichment approach. The dataset consisted of 2D Full-Field Digital Mammograms (FFDM) collected from a community healthcare facility in Southern California for a retrospective study. All patients in the collected dataset were females, aged 40–90 years, who had a biopsy performed between October 2011 and March 2017. Of 1393 patients, 499 had a cancer biopsy, 973 had a benign biopsy, and 79 had both cancer and benign biopsies. The training set is made up of photos taken using a variety of mammography machines of diferent kinds.

Radiologist accuracy enhanced substantially as a result of the usage of cmAssist, as shown by a 7.2% improvement in the area-under-the-curve (AUC) of the receiver operating characteristic (ROC) curve for the reader group with a two-sided p-value of 0.01.

With the usage of cmAssist, all radiologists observed substantial increase in cancer detection rate (CDR) with two-sided p-value of 0.030 and a confdence interval of 95%.

When radiologists employed cmAssist, their accuracy and sensitivity for identifying cancers that had previously gone undetected improved signifcantly. After utilizing cmAssist, the percentage improved drastically in CDR for the radiologists on the reader panel which went from 6 to 64% (mean 27%), while there is no increase in the number of false-positive recalls.

On this data set of missed malignancies, cmAssist has a maximum feasible sensi‑ tivity of 98% when used in stand-alone mode. Future work in AI-assisted false-positive reduction is underway, leading to even greater gains in cancer diagnosis accuracy (Watanabe et al. [2019](#page-51-3)).

2.1 Breast cancer detection using artifcial neural networks

Basic data mining problems like classifcation and regression are efective in solving the breast cancer detection problems. While Artifcial Neural Networks are frequently used to identify breast cancer with good amount of accuracy. ANN based design consists of a collection of linked neurons organized into three layers: input, hidden, and output. This kind of network learns to perform assigned tasks by taking into consideration a sufficient number of instances. The input layer consists of several neurons which is equal to the number of characteristics in the dataset. The hidden layer is another component of ANN with all hidden layers counted as a single one. In the past experiments with ANN, 31 neurons are connected to 9 of the hidden layers. There is a $9-9$ mapping of connection between the frst two hidden layers. As the problem is of the binary classifcation there is only a single neuron at the output.

The cross-validation was executed using a ten-fold technique, which meant that the dataset was divided into ten equal groups. The dataset used is the breast cancer dataset acquired from the University of California Irvine (UCI) machine learning repository. This dataset consists of 699 instances classifed as benign or malignant, with 458 instances (65.50%) being benign and 241 instances (34.50%) being malignant. The model was calibrated for 100 epochs with fve batch sizes, and the activation function is employed in the hidden layers while the sigmoid is employed at the output layer. The ANNs outperformed and computed 98.24% accuracy. The model produced by ANN is more robust and accurate than any other method, and it has the scope to make major advancements in breast cancer prediction.

However, ANN does not account for the training time and False Negative (FN) outcome when training the model. The researchers proposes a new strategy and the usage of the convolutional auto-encoder for breast cancer detection. The proposed hybrid convolutional network uses two models of convolutional autoencoders. First for picking up features and the second for categorizing those features. The fully connected layer is used for the output of the convolutional layers for categorizing input pictures as benign or malignant only after acquiring the most critical attributes. With a shorter training time, the suggested model achieves superior performance. The model's sensitivity is found as 93.50% which is better than the previous studies (Sharif [2021\)](#page-51-2).

Research has proven ANN to be highly accurate in case the of a breast cancer diagnosis. However, this method has some limitations:

- 1. ANN has some parameters to be tuned at the beginning of the training process such as hidden layers and hidden nodes, learning rates, and activation function.
- 2. It takes a long time for the training process due to the complex architecture and param‑ eters update process in each iteration that needs expensive computational costs.
- 3. It can be trapped to local minima so that the optimal performance cannot be guaranteed.

Several attempts have been made to fnd answers to the constraints of neural networks. Huang and Babri showed that Single Hidden Layer Neural Networks (SFLN) can tackle those problems using a three-step extreme learning procedure termed ELM. ELM was compared to traditional Gradient-Based Back Propagation Artifcial Neural Networks in terms of performance (BP-ANN). Performance is measured using sensitivity, specifcity, and accuracy. It is observed that Extreme Learning Machine Neural Networks (ELM ANN) gives superior results in general than BP-ANN. Doctors may benefit from intelligent classifcation algorithms, particularly in reducing errors caused by inexperienced practitioners.

Three main diferences between BP-ANN and ELM ANN are observed based on the defnition:

- 1. We need to adjust several parameters like the number of hidden nodes, momentum, rate of learning, and termination criteria for BP ANN. Whereas, ELM ANN is an easy algorithm that only requires defning the number of nodes and no tuning.
- 2. In hidden and output nodes, ELM ANN can utilize both differentiable and non-differentiable activation functions whereas BP ANN can only employ diferentiable activation functions.
- 3. The BP ANN is trained with a model which has a low training error to terminate at a local minimum. Whereas, ELM ANN is trained with a model that has minimal training error and weight norm, allowing it to develop better-generalized models and reach global minima.

The performance measures used for classifcation problems were accuracy, specifcity, and sensitivity. Similar experiments were conducted with BP ANN for comparison. Results showed that generally ELM ANN was better than BP ANN.

ELM ANN surpassed BP ANN in breast cancer diagnosis. Even though the specificity rate found slightly lower, sensitivity and accuracy rates are signifcantly better in ELM ANN. The researchers concluded that ELM ANN has an enhanced generalization model than BP ANN based on these outcomes. The dataset used for these fndings is the Breast Cancer Wisconsin Dataset obtained from the University of Wisconsin Hospital, Madison by Dr. William H. Wolberg. The dataset consists of 699 instances with 10 attributes plus the class attributes, with a class distribution of 65.5% (458 instances) for benign and 34.5% (241 instances) for malignant (Prasetyo et al. [2014](#page-51-5)).

2.2 Breast cancer detection using active microwave imaging

Microwave imaging is an upcoming technique that offers therapeutic applications. The interactions with the tissues are studied. The malignant cancer tissues show visibly more scattering than the normal tissues. This can help in the detection of the tissues in the early stages where cancer could be cured. In this technique, malignant tissues are estimated to have higher water content than normal ones.

Methods in Active Microwave Imaging are sated below:

- 1. Confocal microwave technique: It is a non-ionizing technique. This method takes advan tage of the fact of the see-through properties of the breast to create a dielectric contrast of tissues based on water content. Time shifting and the summing of the signals are used for the detection of cancer tissues. Acquiring data: There is an antenna that is used as a trans-receiver. The antenna is set around the sample to observe it. An observation is taken every 10 degrees. At last, signals are recorded at every step added to get the fnal signal. The data is validated using time-domain analysis based on the fnite diference method.
- 2. 2D-microwave tomographic imaging: This is the traditional method used in breast cancer diagnosis. Acquiring data: The breast sample is brought into view using the bow-tie antenna. A frequency of 3000MHz is used for the sample. The antenna behavior is the same as the confocal microwave technique. The sample readings are taken in steps of 10 degrees. With the experimentation on active microwave imaging, following conclusions are made:
	- Active microwave imaging is very useful in the detection of cancer in its early stages.
	- Studies show that microwave tomographic imaging could effectively distinguish and image the tissues showing the signifcant diference in dielectric-permittivity.
	- Tumor location can be determined using Confocal Imaging. This is achieved by var iations seen in signal strength due to water content (Bindu et al. [2006\)](#page-50-2).

Table [1](#page-9-0) represents the observations among AI-enabled Breast Cancer detection techniques.

3 AI enabled lung cancer detection techniques

The AI techniques share similarities in their application to breast cancer as well as the lung cancer. Each cancer type mentioned here presents unique challenges and considerations. Therefore, AI solutions are tailored to address the specifc characteristics and complexities of each cancer type, as well as the clinical context in which they are deployed. The safety of AI-based approaches in cancer diagnosis and treatment for breast cancer and the lung cancer is measure with the similar set of tools.

AI-based lung cancer detection techniques use advanced machine learning algorithms to analyze medical data including chest X-rays, CT scans, and PET scans for the early detection and diagnosis of lung cancer. These algorithms are trained on large datasets of annotated images to recognize subtle patterns and abnormalities indicative of lung tumors, often achieving high levels of accuracy in classifcation tasks. AI systems can assist radiologists by fagging suspicious areas for further examination, reducing interpretation time, and potentially improving diagnostic accuracy. The most popular ways of detecting lung cancer are lung radiography and computed tomography (CT) scans (Mustafa and Azizi [2016](#page-50-4)).

3.1 CT lung cancer detection

CT lung cancer detection is one such AI-backed solution that is used to aid physicians, lowering their workload, improving hospital operational processes and giving them more time to create a high-quality doctor-patient connection. Computer-aided detection (CAD)

employing CT scans relieves clinicians' workload and increases efficiency by discovering previously undetected lung cancer nodules (Sathykumar and Munoz [2020\)](#page-51-6).

Small-cell lung carcinoma (SCLC) and non-small-cell lung carcinoma (NSCLC) are the two common types of lung cancer used for therapeutic reasons. Lung cancer staging is a method of determining the extent to which cancer has progressed from its initial site and it is one of the elements that afect lung cancer diagnosis and possible treatments (Mustafa and Azizi [2016\)](#page-50-4).

Radiologists employ X-ray imaging of the lungs to diagnose early lung cancer. When existent tissue is to be separated from the absence of an early tumor, radiology physician monitoring becomes more challenging. This accurate detection result may aid radiologists in more properly assessing and diagnosing patients' ailments. The study's purpose is to develop smart application software or intelligent machines that can identify and categorize early tumor types by utilizing an artifcial neural network. This precise detection result may assist radiologists to analyze and diagnose patients' diseases more efectively so that early lung tumor test fndings might beneft patients in treatment by avoiding danger or no risk (Pandiangan et al. [2019](#page-50-6)).

The authors compared and assessed the results of four separate research experiments. Sensitivity, accuracy, the area under the curve (AUC), specificity, and the receiver operation characteristics (ROC) curve are some of the metrics that the models examined. High specificity number suggests a low rate of lung cancer misdiagnosis whereas a low specificity value indicates a high probability of false positives. The model's accuracy is the per‑ centage of data that was properly categorized. Finally, both the ROC and AUC curves are employed in the various research group model performance measurements.

In a research experiment done by $ToAYaA\Sar$ et al. the authors used lung CT images for lung nodule cancer detection. The use of image augmentation, Minimum Redundancy Maximum Relevance (MRMR), principal component analysis (PCA), and suitable feature selection resulted in the model performing very accurately. Out of the various iterations, the use of deep features with KNN and MRMR gives the best result with an accuracy of 99.51%. Three other research studies considered got a less accurate result in comparison due to the lack of image augmentation and feature selection techniques (Sathykumar and Munoz [2020](#page-51-6)).

3.2 AI based lung segmentation techniques

The researchers devised a lung segmentation technique to improve segmentation accuracy and separate and remove the trachea from the lungs. Digital image processing techniques have been utilized to improve quality and accuracy, as well as to demonstrate the evolution of the feld.

The approach used in research is focused on meeting the goals of the study. Traditional lung X-ray image processing and ANN procedures are used to identify and classify early cancers. These fndings are utilized to help enhance visual observation of the target item. The image processing system includes pre-processing, image noise reduction,image enhancement, lung organ segmentation, object edge detection, and tumor boundary identifcation. In X-ray pictures, low contrast is used to distinguish between malignant and nonmalignant tumors.

The back-propagation technique is used in this study. The beneft of this technique is that it properly reduces mistakes caused by the discrepancy between the actual output and the predicted outcomes. The performance of artifcial neural networks with graphics is demonstrated.

Image processing techniques are used on 25 lung nodule samples and 25 non-lung nodule samples from each of the 50 standard lung X-ray samples of parents. An artifcial intelligence (AI) machine has been developed that can identify 10 samples correctly. The machine works by combining two-dimensional (2D) X-ray images with previously studied tumor characteristics. After training with a large dataset, it should be able to achieve close to 100% detection performance (Pandiangan et al. [2019](#page-50-6)).

Recently, Cengil and Cinar proposed a CNN-based model for the prediction of lung cancer. This technique makes use of the TensorFlow library for the detection mechanism. The authors used SPIE-AAPM-LungX dataset which contains images of 60 patients. The model supports an accuracy of 70% which is less than the standards required.

Another CNN-based approach is presented by Sasikala et al. This approach is applied to Lung Images Dataset Consortium and Image Dataset Resource Initiative (LIDC-IDRI) dataset. The pre-processing layers consisted of median flters which removed any unwanted features from the images. The accuracy achieved was 96%. A comparison study is presented by Gunaydin et al. They compared models like KNN, SVM, and decision trees to detect lung cancer. Data set used in this study is the Standard Digital Image Dataset from the Japanese Society of Radiology Technology. All techniques shows good results with the accuracy of 95.05%. Asuntha and Srinivasan proposed a diferent model with new approaches using deep learning techniques. They have used highly efficient feature extraction techniques such as wavelet transform, HoG nodes, Zernike moment, SIFT, and LBP. After the primary feature has been extracted, a fuzzy particle swarm optimization is used to select the most evident features. The model is tested on several data sets and has shown the accuracy of 97% (Patel [2022\)](#page-51-7).

Chemotherapy, radiation, and surgery are the methods for treating lung cancer, depend‑ ing on the kind of cancer. One of the most commonly used surgical treatments for the initial stage of lung cancer is the elimination of a lung lobe. The chemotherapy treatment is determined by the kind of tumor. In advanced instances, chemotherapy increases overall survival as well as the quality of life when compared to supportive treatment alone. For patients who are unable to have surgery, radiotherapy is frequently combined with chemotherapy. Smoking cessation and prevention are both efective methods for avoiding the development of lung cancer. Long-term vitamin A, D, or E supplementation has not been shown to lessen the incidence of lung cancer. The larger consumption of fruits and vegetables appears to reduce the risk, though the fact is that there is no established link between food and lung cancer (Mustafa and Azizi [2016](#page-50-4)).

Despite the many diverse types of cancer, lung cancer, with its unique development and spreading processes, can infuence normal cells and disturb the cell signaling process, which modifes the function of cell division. To date, an enormous number of studies have been conducted in several aspects of diagnosis of cancer or pre-cancer stages by using arti– fcial intelligent systems-based algorithms. Some algorithms are supervised and some set of algorithms are un-supervised along with diferent features extracted from pathological images. To the best of the author's knowledge, there have been no systematic review and meta-analysis studies to evaluate the performance as well as to estimate the current status of existing approaches to lung cancer. The well-known databases are explored in this systematic review and meta-analysis based solely on a Boolean query for lung cancer and the accompanying artifcial intelligent systems. Preferred Reporting Items for Systematic Reviews and Meta-Analyses Diagnostic Test Accuracy (PRISMA-DTA) is used to conduct the systematic review, The English-language papers that use various sorts of prediction

models to distinguish between healthy and malignant cell pictures are taken from the databases. The relevant publications that had the necessary data, such as the true positive, true negative, total sample size, false positive, and false negative values, are then chosen from the search results. The studies with insufficient data are excluded from further examination.

The subsequent research employed a variety of artifcial intelligent systems, including support vector machines (SVMs), artificial neural networks (ANNs) with diverse training strategies, and statistical techniques, despite the different retrieved features from lung cancer images. Artificially intelligent systems are used in each of them to make decisions similar to those of a clinical practitioner when diagnosing lung cancer. I-squared parameters generated by Meta-DiSc software enable a detailed assessment of the heterogeneity in a meta-analysis study and the potential impact of the included studies on it. I-squared indices are classifed as low, moderate, or high depending on whether they fall within the range of 0 to 25%, 25 to 50%, or above 75%. Additionally, the SROC curve and the estimated values for the combined diagnostic odds ratio, AUC, sensitivity, and specifcity are 0.77, 0.74, 17.22, and 0.872, respectively. Since the AUC value is higher, it is concluded that artif‑ cial intelligence systems are efective in diferentiating between types of lung cancer. The fndings for the development trends in the quantity of success in the performance of the artificially intelligent systems have been presented, taking lung cancer diagnosis and conducting a meta-analysis on the papers into consideration. Eventually, two publication bias tests have shown that the possibility of publication bias exists. Additionally, while sensitivity and specifcity trends are moderate, the diagnostic odds ratio and AUC values shows enormously high trends (Sokouti et al. [2019](#page-51-8)).

Table [2](#page-14-0) represents the critical observations of AI techniques for lung cancer detection.

4 AI techniques for colorectal cancer

The lung cancer detection techniques discussed in the previous section share some commonalities with colorectal cancer in terms of diagnostic and treatment challenges. These two cancer types are distinct in terms of anatomical notation. Clinical decision support systems are used in both the cases of lung cancer as well as colorectal cancer. Artificial intelligence has seen signifcant growth in tumor evaluation for CRC over the past years. Various techniques and steps involved in tumor recognition are the evaluation of some categorical as well as numeric data. Obtaining a correct diagnosis from clinical data during the initial steps of the medical assessment might reduce the chance of human error while also saving time. Colonoscopy, one of the most widely used computer-aided diagnosis techniques, is the most commonly used method for detecting and screening colorectal cancer (Lorenzo– vici and Dulf [2021](#page-50-7)).

4.1 Prediction of colorectal cancer using AI

The high occurrence and fatality rate of CRC presents the question of artificial intelligence's usage in colorectal cancer epidemiology. As broad approaches to understanding research and data have started, several stages of using deep learning in epidemiological studies have started. However, they do experience some difficulty because of factors like data access, classification and accuracy. Various facets of artificial intelligence (AI) utilized for colorectal cancer prediction include:

Table 2 Observations on lung cancer detection techniques

- 1. Geo AI: It is a subset of AI that was initially used in environmental healthcare and now being used by healthcare professionals. It functions by using information from a given region to collect more specialized data using artifcial intelligence. Information is read‑ ily available on a wide range of topics, including geographic (terrain) information, food consumption information, healthcare information etc. The idea behind GeoAI is that particular regions that have a higher rate of colorectal cancer can gain the most from this kind of data collection.
- 2. Digital epidemiology: It is a promising but possibly divisive area of AI. This technique focuses on obtaining data from digital sources including social media and digital devices, and this data is utilized for the early identifcation of SARS and real-time surveillance of Covid-19. This provides access to a vast body of knowledge that was previously unavailable to doctors and can help with early illness identifcation and public health surveillance. The predicted information might not, however, be what the doctors require.
- 3. Data mining: It is another application that serves as a practical tool. It analyzes massive amounts of data gathered from several sources and extracts facts, patterns, and data from them. When used to diagnose colorectal cancer, data mining tools helps in fnding connections, linkages, and a variety of other potentially hidden characteristics that are never explored and may contain a signifcant piece of the information for what causes colorectal cancer (Yu and Helwig [2022](#page-51-9)).

4.2 AI techniques in colorectal cancer screening

- 1. Colonoscopy: It is a highly rated method due to its attributes like high sensitivity as well as decision-making power on lesions. However, small-size or fat polyps have a high chance of being missed by the naked eye (Yu and Helwig [2022\)](#page-51-9). A fexible tube (endoscope) with a small video camera at the tip is used to do colonoscopy explora‑ tion. When polyps bigger than 10 mm are detected, traditional colonoscopy is efective. Although DCE (dye-based chromoendoscopy) may identify microscopic or fat polyps, it has signifcant limitations, such as interobserver and intraobserver variability. White light alone is used in an AI-based method to determine whether polyps are present in a video frame. Polyp categorization is the process of cataloging the type of polyp found after it has been discovered. Using three well-defned methods, depending on whether the polyp seems to be benign, pre-malignant, or malignant, the doctor can make a better judgment about whether or not to remove it. Colorectal polyps are characterized as adenomatous, hyperplastic, or serrated and infammatory based on their anatomical pathogenesis. AI might help medics distinguish polyps in the future, which is a promising development. There is a form of serrated polyp known as a sessile serrated adenoma that is not generally neoplastic (Viscaino and Bustos [2021](#page-51-10)). An experiment study of colonoscopy with 1058 patients was conducted, which resulted in the conclusion that colonoscopy reduces the mortality rate.
- 2. Blood tests: Studies have found that certain blood test data and some features could be used to identify patients carrying the risk of CRC. One such test is the Complete Blood Count. Slow bleeding from cancer in CRC causes iron defciency anemia, which is accounted for by this test. A machine learning model, MeScore, is developed and tested rigorously for calculating the risk factors of CRC.
- 3. CT colonography: A non-invasive imaging test for CRC is CT colonography (CTC). The ability of CTC to diagnose and distinguish between various lesions can be reduced, and the capacity for detection can be increased by CAD.
- 4. Colon capsule endoscopy: It is an invasive procedure. It is to be noted that the invasion is minimal. It required more laxatives. It removes the capsule from the GI tract by flush– ing it out. It required manual reading, which increases the error rate. AI techniques are applied to these images to interpret the results.
- 5. Polyp characterization: This is a critical step. The characterization of polyps carry a lot of weight. AI comes into play here for predicting malignant or benign lesions. Most colorectal polyps are hyperplastic. Even in that case, it is absolutely important to have a high-precision diagnosis. A retrospective study was carried out at three hospitals, using methods like conventional white light endoscopy, etc., to construct and evaluate deep learning models for the automatic classifcation of colorectal lesions histologically in white-light colonoscopy images. This study contained a total of 3828 pictures from 1339 participants. The results showed a promising future for deep learning models.
- 6. Magnifying narrow-band imaging: It is a less bandwidth-intensive form of endoscopy that uses image enhancement. It is an optical flter. These flters are used for green and blue light illumination in a sequence. It comes in handy for polyp characterization. A computer-based method is tested for classification. All 214 study participants underwent a zoomed NBI (narrow band imaging) colonoscopy and had a total of 434 polyps measuring 10 mm or less in size. The diagnostic agreement between two specialists and computer-aided classifer systems is 98.7%, with no discernible diference.
- 7. Magnifying chromoendoscopy: It is a technique to amp up the visualization of pit pat‑ terns on the surfaces of polyps to distinctly identify benign and neoplastic polyps. An analysis is done on an automated computer system named HuPAS, which could outline the boundaries of the pits. The aided system is in 100% compliance with the expert diagnosis for type I and II pits. For types IIIL and IV, it shows values of 96.6 and 96.7, respectively.
- 8. Endocytoscopy: This technique involves magnifying real-time images by 380–500 folds. This allows for clear visuals. In esophageal cancer (EC), a contact light microscopy system is added to the colonoscope's distal tip, which permits instantaneous evaluation of nuclei and cytological structures. CAD-EC is developed to automate export-dependent diagnosis. The database consists of images from 242 patients. CAD-EC shows a sensitivity of 89.4%, a specifcity of 98.9%, an accuracy of 94.1%, a PPV of 98.8%, and an NPV of 90.1% in diferentiating invasive cancer from adenoma.
	- 1. Confocal endomicroscopy/confocal laser endomicroscopy: It allows for real-time \times 1000 image magnification. This technique requires a lot of training and is hence performed only by experts. To distinguish between neoplastic and non-neoplastic polyps, a study of the diagnosis capacity of computer-based automated pCLE classifcation was conducted, and it was contrasted with experienced endoscopists who made a diagnosis based solely on pCLE recordings. For computer-based automated pCLE classifcation and expert performance in diferentiating neoplastic and nonneoplastic lesions, the results showed a sensitivity, specifcity, and accuracy of 92.5 vs. 91.4%, 83.3 vs. 84.7%, and 89.6 vs. 89.6%, respectively. These diferences were not statistically signifcant.
	- 2. Laser-induced fluorescence spectroscopy: This technique gives us a real-time auto– mated distinction between benign and neoplastic polyps. It includes an optical fber device called WavSTAT. This component emits laser waves onto targeted tissue and

then releases the light to give the results. A new version of LIFS is studied which makes use of WavSTAT4. The accuracy of LIFS using WavSTAT4 in predicting polyp histology was 84.7%. For distal colorectal diminutive polyps only, the NPV for excluding adenomatous histology increased to 100%. The dataset used was a combination of electronic medical record data from two unrelated patient popula‑ tions in Israel and the UK to develop an ML-based prediction model for identifying individuals at high risk of colorectal cancer based on attributes such as complete blood count (CBC), age, and sex (Goyal and Mann [2020\)](#page-50-8).

4.3 AI for colorectal cancer diagnosis

One of the numerous sectors in which machine learning techniques are gaining prominence is the development of computer-aided diagnosis systems. The most widely used traditional machine-learning algorithms in medical applications for data analysis are decision trees. Important factor when working with CAD (computer-aided diagnosis) in tumor recognition is confdence analysis (Lorenzovici and Dulf [2021](#page-50-7)). CAD tries to locate aberrant or suspicious areas in order to improve detection rates while lowering false-negative rates (FNR). (Viscaino and Bustos [2021\)](#page-51-10)

- 1. Endoscopy and MRI/CT imaging: One of the most efective methods for diagnosing colorectal cancer makes use of convolution neural networks (CNN), an ANN type, and computer vision, endoscopy, and MRI/CT imaging. It analyses a vast number of images to fnd patterns and objects. The frst steps in improving the endoscope have been made possible by segmentation technology, which derives from computer vision's capacity to distinguish between things. The important thing to remember is that optimal segmen– tation would allow for the separation of anatomy and malignant and normal masses. Pattern recognition is used to distinguish between normal and abnormal conditions. The detection, segmentation and classifcation of pictures are currently the most signifcant areas where AI has been able to infuence this kind of imaging. Additionally, general initiatives to enhance picture quality for better reading and advancements in segmenta– tion technology have been observed. Along with the obvious advantages of improved diagnostic accuracy, using deep learning in colonoscopies also has the added beneft of reducing any variability in detection rates. One of the major challenges was dealing with bigger artifacts, which might lead to misunderstanding of data or false-positive results. However, the lack of relevant datasets is the main issue. When attempting to employ CNNs for image or object identifcation, there are generally certain restrictions or problems. Pixel appearance anomalies in images can cause misinterpretations owing to missing or irregular pixelation. Because anatomy is not always the same in all people, it might be mistakenly perceived. Anatomical variations or patient posture practices can affect how an image is taken, causing positional alterations that could result in misunderstanding. Additionally, it might be difficult to distinguish between useful information and information that is only an artifact when segmenting data. The limitations of endoscopes and CT/MRI imaging serve as a reminder that AI is still being improved and is not yet ready for full application and they also highlight the continued need for technicians in the interpretation of imaging.
- 2. Genetic and pathological diagnosis: Some progress has been made in training AI to categorize cancers based on histology. When infammatory tissue are present in data sets, the systems, however, has identifcation problems. The inability of AI to detect

pictures may not be its fault; instead, CNNs requires higher-quality training data, and stain normalization in hematoxylin and eosin pictures are able to aid AI by increasing its accuracy. By paying more attention to characteristics like textures, spatial connections, and morphology, the accuracy may be increased even further. Fuzzy systems have also proven efective since they allow for more thorough information interpretation. Any type of pathology can use this scope to fnd abnormalities or forecast the possibility of malignancy based on predetermined criteria. (Yu and Helwig [2022\)](#page-51-9)

4.4 AI for colorectal cancer treatment

- 1. Therapeutic assessment: Rectal cancer treatment using an AI model enables the earlyonset diferentiation between therapeutic full response (CR) and not-respond to therapy (NR). AI makes it possible to comprehend some metabolic processes and drug-induced changes that are directly related to the development of colorectal cancer. AI has signifcantly enhanced the processing of complicated networks of biological information. Additionally, the accuracy of AI's colorectal cancer ANN algorithm prediction is rising in importance. ANN is distinguished by nonlinear models that are adaptable to medical and clinical research. The following are a few advantages observed with this technique:
	- 1. It has the potential to improve the optimization process, resulting in adaptable nonlinear models that are cost-efective, particularly in huge data scenarios.
	- 2. It predicts clinical outcomes with high accuracy and reliability.
	- 3. Improves academic dialogue and knowledge dissemination.
- 2. IBM Watson for oncology: Clinical decision-support systems (CDSSs), a new AI trend for therapeutic suggestions, shows enormous potential for therapeutic management in cancer given the increasing growth of clinical information. By imitating human think– ing, it makes it possible to acquire and analyze knowledge in a way that is superior to the traditional human touch. The selection of cancer therapies has been expedited by AI with a cognitive computing system called IBM Watson for Oncology (WFO). WFO is gaining recognition in the feld of cancer therapy. It formulates recommendations using natural language processing and clinical data from many sources (treatment guidelines, professional opinions, works of literature, and medical records). The datasets used in the study included 10,000 to 40,000 CT images for studies on CT imaging and AI. These images were gathered by the research parties themselves. AI training data is frequently labeled and supervised. The known input and output values have been connected clearly. However, doctors' notes and other unstructured, person-specifc information make up the majority of a case record, making it challenging to feed WFO (Yu and Helwig [2022\)](#page-51-9).

Colorectal surgery is another promising area in which AI has been used. According to research, AI greatly decreased the need for extra surgery after T1 CRC endoscopic resection. In particular, a support vector machine for supervised machine learning was developed to identify patients who required extra surgical intervention. As a result of surgery's difficult, time-consuming, and non-scalable nature, the use of AI in surgery is still limited. The only purpose of AI's presentable results is to expedite the decision-making process (Yu and Helwig [2022\)](#page-51-9).

An experiment was performed by NoA©mi Lorenzovici et al. to maximize the effectiveness of the colorectal cancer detection system by using an innovative dataset. For every patient, the dataset contains 33 blood and urine samples. Also, when determining the diagnosis, the patient's living environment was taken into consideration. Various machine learning approaches, such as classification and shallow and deep neural net– works are used to create the intelligent computer-aided colorectal cancer diagnosis system. The problem is solved using two approaches, traditional machine learning algo– rithms, and the regression problem solved using artifcial neural networks. The initial step in the frst approach is pre-processing and it is done by classifying each record from the dataset as unhealthy (denoted by 1) or healthy (denoted by 0). If there is the presence of other diseases or risk factors, then it is denoted as 1, otherwise 0. The response variable is obtained using two approaches; the frst, absolute deviation; and the second, by fnding the weight of each predictor. In the frst approach, the weights are calculated on the assumption that each variable had roughly the same efect on the diagnosis. The second method for determining the proper weights is to expand individual variables' healthy and unhealthy ranges. But even though the second technique appeared to be more rational than the frst, the results did not support these assumptions in the theory.

The first goal of the experiment is to solve the binary classification problem. There– fore, for the training of the model, a labeled response variable is needed to represent the answer of the model to the input data. After obtaining the predictors and the response variable, various models are trained using the Classifcation Learner toolbox in MAT‑ LAB. This toolbox allows to train models, and each model is trained using two separate validation strategies, each with three alternative approaches. The frst validation technique used is k-fold cross-validation and it had various approaches such as 5-Fold Cross-Validation, 25-Fold Cross-Validation, and 50-Fold Cross-Validation. Regarding the percentage of data held out, three diferent methods are tested, and those are 5% of data held out for validation, 15% of data held out for validation, and 25% of data held out for validation.

In the second approach, taking into consideration the fact that deep neural networks perform better on large datasets than basic machine learning algorithms, solving a regression problem was not performed using the regression learner but rather with the Neural Network Toolbox in MATLAB. The first three types of neural networks are shallow neural networks, having only one hidden layer and a varying number of neurons on that hidden layer. In the frst case, when the hidden layer has only three neurons, an MSE(mean squared error) of 1.94 is obtained after training the network. In the second case, a network with 10 neurons on one hidden layer is constructed, and in the third case, a shallow neural network having 20 neurons was made. The next three networks are built on a 5-hidden layer architecture and, therefore are called deep neural networks. Evaluating the results obtained after training the frst network with fve hidden layers, it is concluded that this network has the worst performance. After training the classification algorithms, the result– ing models' performances are compared. Traditional machine learning techniques achieves a maximum accuracy of 77.8% while solving the binary classifcation problem. In terms of mean squared error minimization, however, the regression issue addressed with deep neu– ral networks performed substantially better, with a value of 0.0000529. In conclusion, deep neural networks are both more reliable and more efficient in colorectal cancer detection than traditional machine learning algorithms (Lorenzovici and Dulf [2021](#page-50-7)).

Although evaluation of slides by pathologists is the standard for cancer detection, with the increase in workload the authors have seen the development of computerized screening. This is done by whole slide images (WSI), which are a digital form of glass slides. computer-aided diagnosis (CAD) is possible through the application of medical image analysis, CNN, and AI. With increasing cases of CRC, the number needed to diagnose increases as a study shows only 7.4% of cancers are detected as positive.

Unique AI based deep learning model is trained for screening colonic malignancies in colorectal specimens for better detection and classifcation. This unique algorithm developed by Qritive consists of two models, frst to detect the high-risk regions in Whole Slide Images (WSIs) by gland segmentation and second to classify the WSIs into "low risk" (benign, inflammation) and "high risk" (dysplasia, malignancy) categories. The WSIs of 294 colorectal specimens are collected from the archives of Singapore General Hospital's pathology and The Cancer Genome Atlas (TCGA). The attributes selected from the data included the histological features present in the WSIs, such as glandular structures, dysplastic regions, blood vessels, necrosis, mucin, and inflamma tion, which were used to train the AI models for accurate detection and classifcation of high-risk regions indicative of malignancy. Slides containing poor-quality images and other malignancies are excluded. The gland segmentation model is a deep learn– ing model that uses Faster Region-Based Convolutional Neural Network (Faster-RCNN) and Resnet-101 for feature extraction. The second model is a gradient-boosted decision tree classifer, which classifes WSIs as high risk and low risk. The model is initially trained based on 39 WSIs annotated by pathologists. The gland segmentation model is trained by dividing the WSIs into tiles. These tiles are separated into categories if tis‑ sue is found and others are discarded. The model is thoroughly trained by using iterative training to reduce the loss function. Data augmentation by rotation, mirroring, contrasting, changing brightness, and adding Gaussian noise and Gaussian blur is done to increase the training set and make the model robust.

The trained segmentation model was applied to the 105 WSIs from the Singapore General Hospital to number them according to the following features:

- 1. Area classifed as dysplasia or adenocarcinoma with a minimum of 70% certainty.
- 2. Area-weighted average prediction confdence for adenocarcinoma or dysplasia objects.
- 3. The boolean fag is set to 1 if the slide has a prediction certainty of more than 85%.
- 4. Predictions for cancer or dysplasia are in the bottom 1 percentile.

After the numeric classifcation, a fvefold cross-validation is carried out to train, tune, and compare several models. The best-performing model is selected. This model is tested on 150 WSIs and the results are compared to those of pathologists. It is found that the model could classify 119 out of 150 slides correctly, thus resulting in a high sensitivity of 97.4%, a specifcity of 60.3%, and an accuracy of 91.7%.

The confusion matrix generated from the predicted results of biopsies is depicted in table [3](#page-20-0) below.

On evaluating the data from the testing dataset, it is observed that its high accuracy of 91.7 shows uniformity between the AI model and the pathologist. With a sensitiv‑ ity of 97.4%, the model assures the minimization of false negatives (Goyal and Mann [2020](#page-50-8)). This model guarantees the identifcation of high-risk CRCs. These identifed specimens are then be sent to a pathologist for fnal verifcation. Thus, the AI model is

used as a screening tool to help reduce the diagnostic burden of pathologists by identi-fying significant cases (Ho and Zhao [2022](#page-50-9); Viscaino and Bustos [2021\)](#page-51-10).

Table [4](#page-22-0) lists the observations in literature for colorectal cancer.

5 Prostate cancer

Though the risk factors, diagnostic methods, and treatment modalities are diferent for prostate cancer with respect to breast cancer, lung cancer, and colorectal cancer, there are common AI methods to deal with all four types of cancers. Medical Image analysis is the common technique used among all the mentioned types of cancers. AI techniques for colorectal cancer involve analyzing medical imaging data like CT scans. Similarly, AI techniques for skin cancer, especially melanoma, use deep learning to analyze dermatoscopic images, identifying malignant lesions by visual patterns. By utilizing imaging data and machine learning, AI enhances early detection and treatment strategies, highlighting its transformative potential in cancer care. The second most prevalent cancer to cause death in males is prostate cancer, which is also the most frequently diagnosed non-skin malignancy in men. The volume of prostate biopsies has increased, and there is a dearth of urological pathologists, which strains the ability to diagnose prostate cancers. These difculties are encountered during prostate therapeutic interventions (Booven and Kuchakulla [2021](#page-50-10)). Prostate cancer should be found early and located while it is still curable since excellent cancer-specifc survival is anticipated for the majority of locally contained illnesses. Radia‑ tion therapy and radical prostatectomy are frequent therapies for individuals with locally advanced, high-risk illnesses. The Gleason grading system's pathologic grade is the main factor in clinical risk evaluation. Generally speaking, cancers with just Gleason pattern 3 are regarded as low risk. The risk of developing metastatic illness and dying from cancer is increased in tumors with dominant Gleason patterns 4 and 5 (Harmon and Tuncer [2019\)](#page-50-11).

It is anticipated that the development and acceptance of digital pathology technology will increase the precision of anatomic pathology diagnosis. Machine-learning algorithms have shown some accuracy in recognizing disease phenotypes and characteristics when applied to digital images (Perincheri and Levi [2021](#page-51-11)).

There is widespread knowledge of the value of multiparametric magnetic resonance imaging (mpMRI) as a method for detecting prostate cancer. The prostate and pelvic areas can be seen in high-contrast, high-resolution anatomical pictures thanks to mpMRI. Radiologists must possess a high degree of competence in order to assess and interpret all mpMRI sequences. Literature has been developed to describe the complicated heterogene– ity of localized prostate cancer using the functional imaging properties of mpMRI. In the past ten years, there has been a significant rise in the usage of AI applications in both radiology and pathology (Harmon and Tuncer [2019](#page-50-11)).

5.1 AI in localized prostate cancer

Contributions of AI to prostate mpMRI are particularly expected to increase the sensitivity of prostate cancer diagnosis and lower inter-reader variability. The TZ shows the largest advantage, enabling readers with average competence to reach 83.8% sensitivity with automatic detection as opposed to 66.9% with mpMRI alone. Litjens et al. are able to show that integrating AI-based prediction with the Prostate Imaging Reporting and Data Sys‑ tem (PI-RADS) enhances both cancer detection and clinically relevant (aggressive) disease

Table 4 Observations in literature for colorectal cancer

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classifcation. While there haven't been as many studies on the automated identifcation of prostate cancer using mpMRI using deep learning. Early research suggests increased detection rates compared to earlier studies. Here, PI-RADS v2 classifcation in conjunction with automated detection enhanced the diagnosis of clinically relevant malignancy over radiol ogist-only detection. Biopsy intra-lesion spatial targeting of aggressive spots on mpMRI is not best suited for AI algorithms. The algorithm should be educated using voxel-based classifers against a system that classifes each tumor "patch" according to its regional Gleason score in order to enable more precise spatial learning (Harmon and Tuncer [2019\)](#page-50-11).

Machine learning algorithms used in cancer therapy often include semantic segmenta– tion of stromal, epithelial, and lumen components. Due to how time-consuming it is to geographically annotate at the full resolution of digital pathology pictures, this degree of spatial annotation is not viable. Theoretically, more thorough pathologic details of each lesion would be necessary as input to enhance spatial labeling at the mpMRI level. The use of AI in pre-processing stages to enable more intricate histopathologic spatial analysis with radiologic imaging has shown potential. According to research by Kwak et al., the tissue component density is substantially correlated with the MRI signal characteristics. The differences between tumors with Gleason patterns 3, 4, and 5 were shown by Chatterjee et al. to be bigger for the gland component volumes than for cellularity measures. The detection and classifcation of intermediate prostate tumors are now constrained by pathology and imaging. Most models are built using weakly labeled data, and there aren't many highquality annotations available. The foundation for high-resolution radio-logic labeling can be improved through pathological classifer performance (Harmon and Tuncer [2019](#page-50-11)).

Digital pathological evaluation using automated Gleason grading has been the main focus of prostate cancer research. Building classifiers from manually created characteristics obtained in relatively uniform environments has been the foundation of traditional machine-learning techniques. Prostate cancer grading systems may now perform bet– ter thanks to more recent developments in deep learning applications. Deep learning and machine learning were used to create a novel network for prostate cancer diagnosis and grading. According to the results, the accuracy of epithelial identifcation was 99%, while the accuracy of low- and high-grade illness was between 71 and 79%. New possibilities for image-based jobs at full-scale resolution will open up as technology advances in processing power (Harmon and Tuncer [2019\)](#page-50-11).

5.2 AI methods in prostate cancer diagnosis

- 1. ANN and Histopathologic Diagnosis of Prostate Cancer: A prostate cancer diagnosis frequently depends on the histological recognition of prostatic adenocarcinoma. Numer– ous investigations were undertaken, and it was shown that many of them demonstrate the potential of AI ANNs to provide more precise patient counseling and circumvent histopathologic variabilities.
- 2. ANN and Magnetic Resonance Imaging (MRI) diagnosis of prostate cancer: Detection and aggressiveness assessment using magnetic resonance imaging (MRI) have both been studied as treatment options. Although MR spectroscopy, T2-weighted MR imaging, and apparent diffusion coefficient (ADC) have been useful methods to evaluate prostate cancer, there is still disagreement about how to best apply them. These AI classifer techniques have been proposed as potential methods for non-invasive tumor progression detection and decision support for active monitoring programs.
- 3. Artifcial Neural Network in Biomarker Diagnosis and Risk Stratifcation: PSA level testing has helped to inform prostate cancer diagnosis and prognosis. A deluge of biomarkers has been discovered and incorporated into clinical assays over the past ten years. Analyzing and verifying the biomarkers can be greatly aided by ANNs. In several investigations, clinical prediction models were created using machine learning techniques. The fndings demonstrate that novel non-invasive biomarkers can be found using computationally directed proteomics.
- 4. ANN for patient interaction and patient centered care Many prostate cancer patients are still uncertain about their potential treatment options after receiving a diagnosis. Understanding how specifc therapies are implemented might therefore help patients feel more at ease and enjoy their treatment more. Numerous research shows the potential of AI ANNs to enable the creation of useful patient-centric tools to aid in informing patients about their treatment options and illness progress monitoring.
- 5. Classification system for prostate cancer risk stratification using ANN: Numerous criteria work in conjunction with ANN to classify patients into risk groups that are classifed into 7 categories ranging from extremely low to very high risk. Although these risk categories provide a solid framework for stratifcation, none of them takes the possibility of repetition into account. Therefore, a deeper understanding of the mechanisms underlying recurrence could equip us with the knowledge necessary to modify risk variables and, consequently, select the most appropriate therapies (Booven and Kuchakulla [2021\)](#page-50-10).

5.3 Study experiments

A study experiment is presented in this section to evaluate Paige Prostate's performance on a prostate core biopsy. In this study, Sudhir Perincheri et al. evaluated Paige Prostate's performance on a prostate core biopsy using the Whole Slide Images dataset from Yale Medicine, which the algorithm has never seen before. Paige Prostate is a machine-learning algorithm that classifes a whole-slide image as "suspicious" if any prostatic lesions are found or "not suspicious" if none are found. It is developed at Memorial Sloan Kettering Cancer Center (MSKCC) in New York using data from its digital slide collection.

The primary objective of the study is to investigate two possible use case scenarios. The frst is its use as a pre-screening tool to spot negative cores that don't need a pathologist to manually evaluate them, and the second is its use as a second read tool to spot cancer foci the pathologist missed. 1876 prostate core samples from 118 consecutive individuals are included in the analysis. The primary diagnostic is a board-certifed pathologist's clinically recorded unique diagnosis. The Paige Prostate algorithm is run on scanned, identifer-free stained samples.

Except for High-Grade Prostatic Intraepithelial Neoplasia (HG-PIN), the algorithm classifed each core biopsy as suspicious if it discovered abnormalities.

Additionally, if the distribution of the core-needle biopsy slides difered noticeably from the thumbnail, it fagged the slides as being out of distribution. Digital images and/or glass slides were inspected to confrm the presence of diagnostic lesion tissue at the scanned level. Without being aware of the final diagnosis or algorithm classification, two genitourinary pathologists manually inspected the entire slides. Thirty randomly chosen core biopsy photos are combined with erratic biopsy images. When one of the two reviewers observed atypia in a core biopsy, the algorithm classifed the sample as "suspicious," and when neither of the two reviewers did, the sample is classifed as "discrepant." The same

pathologists manually evaluated the pictures of any core biopsies that had discrepancies following blinded assessment while emphasizing the area of concern.

5.3.1 Results of the study experiment

Of the 118 patients represented by the 1876 core biopsies in the research, 86 are found to have prostatic adenocarcinoma, and 32 are found to be cancer-free. According to Paige Prostate's analysis, at least one core is deemed questionable in 84 of the 86 individuals with adenocarcinoma, whereas none is in 26 of the 32 patients who did not have carcinoma or glandular atypia.

Among the non-suspicious labels, 16 of the 46 discrepant cores are not interpretable manually, resulting in the fnal diagnosis of adenocarcinoma. Diagnostic lesional tissue is not found in the scanned picture of 19 additional discrepant cores. Taking away the 16 faulty scans and considering the 19 cores with no diagnostic tissue in the scanned picture as concordant, leaves 11 discrepant cores.

Thirty-four (34) of the 477 core biopsies classifed as "suspicious" by Paige Prostate resulted in a benign prostatic tissue diagnosis. In 16 instances, at least one of the reviewers identifed unusual glands during the manual review. The system classifed two cores as out of distribution because they lacked prostatic glandular tissue. With the exception of one core that developed granulomatous prostatitis, the remaining cores were completely benign.

In suspected circumstances, Paige Prostate includes a tool that shows regions of concern. At least one of the two manual reviewers classifed six more cores as odd after reexamining suspected discrepant cores with Paige Prostate annotation. To avoid bias, 30 randomly chosen core biopsy images were combined with discrepant cores for manual reviewers, which included 22 benign cores. The manual readings and the fnal results are identical, with the exception of one core having adenocarcinoma. In the data set, Paige Prostate shows a positive predictive value of 97.9%, a negative predictive value of 99.2%, sensitivity of 97.7%, and specifcity of 99.3%.

The study by Sudhir Perincheri et al. carefully studies the model's performance on unseen data and identifes areas where the algorithm might be improved. Through their fndings, the false negative rate might be decreased, for instance, by better out-of-focus scan detection. On the other hand, disregarding photos with a single focus smaller than 0.25 mm would reduce the algorithm's false positive rate. Paige Prostate has the potential to be a powerful tool with a wide range of use-case applications in anatomic pathology clinical settings (Perincheri and Levi [2021\)](#page-51-11).

In a study experiment conducted by Derek J Van Booven, et al. the authors analyzed the advancements in AI-based artifcial neural networks (ANN) and their current role in prostate cancer diagnosis and management. For this systematic review, the authors explored PubMed, Embase, and the Web of Science to fnd and evaluate publications about the use of active surveillance and machine learning in the detection, staging, and management of prostate cancer. To make decisions during the diagnosis and treatment of prostate cancer, evidence from several modalities must be interpreted. MRI-guided biopsies, genomic biomarkers, Gleason, and PSA levels grading are just a few exam– ples of the tools that are utilized to diagnose, risk stratify, and subsequently monitor patients during corresponding follow-ups. A process that is extremely challenging and time-consuming for humans can be made possible by artifcial intelligence (AI), allowing therapists to recognize complex correlations and manage vast data volumes.

Utilizing artifcial intelligence-based ANN tools that could be seamlessly integrated with some of the crucial instruments frequently used in prostate cancer detection and lowering the degree of subjectivity, it is possible to use limited resources while enhancing the overall efficiency and accuracy in prostate cancer diagnosis and treat– ment (Booven and Kuchakulla [2021\)](#page-50-10). An exceptional possibility for AI applications to enhance detection, classification and overall prognostication arises from the close rela tionship between histology and functional imaging parameters. Improvement in systematic assessment and characterization of pathological and radiologic interpretation is clinically necessary for prostate cancer. To continue making progress in this area, mature data sets with top-notch annotations are required (Harmon and Tuncer [2019](#page-50-11)).

Observations in literature for prostate cancer have been shown in the table [5.](#page-28-0)

6 AI techniques for skin cancer detection

In most of the available literature, Gray Level Co-occurrence Matrix1 (GLCM) is a go-to technique for feature extraction and GoogleNet for classification. Such models had an average accuracy of around 85%. Out of all the literature's, Esteva et al. achieved a notable accuracy of 96% using GoogleNet.

Stanley et al. uses fuzzy-based histogram analysis for diferentiating benign skin lesions from melanomas in clinical images. The model is trained upon 129 melanoma and 129 benign clinical images. Results shows that the model is able to distinguish between both of them with true positive rates between 89.17 and 93.30% and true negative rates between 84.33 and 86.04%.

Al Nazi and Abir proposes a model where augmentation techniques are used to overcome the issue of having fewer images in the dataset. ISIC 2018 and *PH*² data set is used for this purpose. The model achieves an impressive 92% accuracy despite having fewer images in the dataset.

Han et al. proposes a Region-based CNN model for SC prediction. 11,06,886 images are used to train the region-based CNN. The model is trained with two approaches, one with 80% sensitivity to malignant modules with an F1 score of 72.1% and the second with 90% sensitivity to malignant nodules with an F1 score of 76.1% . The model provides better accuracy than dermatologists (Patel [2022\)](#page-51-7).

The computer-aided systems can easily extract features like color variation, asymmetry, texture, etc. There have been many diferent methods, like the 7-point checklist, ABCD rule and Menzies method to improve the diagnosis of skin cancer. The major steps in the computer-aided diagnosis of melanoma are the acquisition of skin lesions, segmentation of lesions from the skin region (to detect areas of interest), extraction of geometric features (accountable for increasing accuracy) and feature classifcation. The feature extraction in CAD depends majorly on the ABCD rule *i.e.* asymmetry, bor– der irregularity, color, and diameter (Kanimozhi and Murthi [2016](#page-50-12)). In treating suspicious cases, physicians have been assisted by system-aided technology. They can also be used by less experienced doctors as an extra tool to obtain a preliminary estimate and enhance patient follow-up processes. These techniques are often divided into two categories: the key category is related to attribute mining from skin images, and the second category uses picture properties like texture and color to recognize patterns.

Table 5 Observations in literature for prostate cancer

6.1 AI‑based melanoma detection

The technique that is used for melanoma detection is computerized detection using Arti– fcial Neural Networks, and this technique proves to be more helpful and powerful than dermoscopy. MATLAB has been extensively used for image processing and developing diagnostic algorithms. The methodology for the detection of melanoma using image pro‑ cessing is depicted below:

- 1. Image pre-processing: An image of the lesion is captured by any lens and in any lighting condition. This image thus needs to be pre-processed.
	- (a) Image scaling: The size of the image is enlarged or reduced to make the lesion more clearly visible.
	- (b) RGB to grayscale: The colored image is then converted into a grayscale image using the rgb2gray function.
	- (c) Grayscale to binary image: This grayscale image is then converted into a binary image using the im2bw function. The output image performs thresholding and converts all the pixels having a value above the threshold to 1(white) and all other pixels to 0 (black).
- 2. Segmentation: This is used to divide the image into several parts, identifying the different objects in the image.
	- (a) Background subtraction: This technique helps in extracting the foreground of the image (objects of interest) for further processing. It is also known as blob detection.
	- (b) Edge detection: It segregates the objects in an image by boundaries by detecting discontinuities in brightness.
	- (c) Masking: It sets the pixel values of the background to 0 or some other value and highlights only the lesion.
- 3. Feature extraction: Melanoma skin lesions are mainly identifed by their geometric features. The features that are most signifcant for melanoma detection are:
	- (a) Area
	- (b) Perimeter
	- (c) Major Axis Length
	- (d) Minor Axis Length
	- (e) Circularity Index
	- (f) Irregularity Indices (A, B, C, D)
- 4. Classifcation: ANN is used in the classifcation stage according to the ABCD rule. The neural networks are constructed in three layers with a feed-forward architecture.
	- (a) The input layer contains information from the features extracted in the previous stage.
	- (b) These input units are integrated with the hidden units in the hidden layer. These hidden units are then linked to the output layer

(c) The output layer gives a result that indicates if cancer is present or not.

 Once the layers are developed, the neural network is then trained with data having the results already present. The backpropagation algorithm is used to train the ANN.

5. Results: The values received from the network are input into a confusion matrix. Once the algorithm is completely trained, the unknown images are given to the algorithm for classifcation as cancerous or non-cancerous. The trained algorithm gives an accuracy of more than 96%. This tool is made more feasible and robust since it can detect cancer in an image taken in any condition and hence can be used in rural areas or where an expert diagnosis is not possible (Kanimozhi and Murthi [2016\)](#page-50-12).

6.2 Skin lesion analysis on gray level profle

Some of the visible features of melanoma in its early stages are skin dryness, fungus, and allergic symptoms. Currently, the captured skin images are analyzed in the frequency domain, but this doesn't prove to be an efficient way as the skin color in these images does not vary much. Hence, this study aims at analyzing images in the gray level profle. This serves as the input parameter for our system. This method analyzes the variations of the RGB spectrum of the skin images using novel six-sigma threshold and region connectivity concepts to detect the border of a lesion and identify the incidence and propagation. An automated method for the diagnosis of skin cancer based on dermoscopic images has been developed. The retrieved features are based on a gray level co-occurrence matrix (GLCM) and a multilayer perceptron classifier (MLP) to differentiate Melanocytic Nevi from malignant melanoma.

It is important to recognize the texture in an image by its contrast, correlation, homogeneity, and energy properties. In gray images, texture can be identifed using 2-D histograms and co-occurrence matrices.

The images captured by the dermatoscope are converted from RGB to grayscale images. The following steps are followed for extracting the gray level co-occurrence matrix:

- 1. The image data is quantized into a given number of gray levels.
- 2. An *NxN* GLCM matrix is created by considering a window of pixels surrounding the given pixel. Then the number of times the given intensities occur in the given matrix is calculated. The total number of times the given spatial relationship occurs in W will equal the sum of all the GLCM elements.
- 3. The GLCM is then made symmetric.
- 4. Then the probability of each pixel is obtained by dividing it by the sum of all intensities in W.

Then, characteristics like homogeneity, entropy, energy, contrast, and correlation can be quantifed.

Then the area and perimeter of the region of interest can be calculated by using segmentation(using K-means clustering) and boundary extraction.

- 1. Convert RGB image to a $L^*a^*b^*$ image which helps in segmentation using k-means clustering.
- 2. Convert L*a*b* image to a grayscale image.
- 3. Convert gray image to a binary image by using the Otsu algorithm.

Based on the features extracted, the researchers trained a model of Back-Propagation net– work. The algorithm uses 50 images, 10 input neurons, and 4 training classes. Researchers have used around 20 images for testing purposes.

The results shows that the developed classifer has 100% accuracy for classifying noncancerous images. The accuracy for the classifcation of skin cancer of type 1 is 95%, while that of skin cancer of types 2 and 3 is 95.8% and 95%, respectively. Thus, the authors have developed a model with high accuracy for the classifcation of cancerous images into 3 types and non-cancerous images (Goel and Singh [2015\)](#page-50-13).

6.3 Skin cancer classifcation with convolutional neural network and transfer learning

With present AI systems, in the detection of skin cancer, the following challenges are observed. A freely accessible dataset called HAM10000 containing 10,015 dermoscopy images is acquired from patients belonging to Australia and Austria. 6705 images are benign, 1113 are malignant, and 2197 belong to unknown lesions:

- 1. The high variances in the shape and size of images make it difcult to give an accurate prediction.
- 2. Pre-processing is essential for good accuracy.
- 3. The contrast from neighboring skin cells plays a part in the analysis. So if contrast is low, detection becomes tricky.
- 4. Colour illumination hinders detection capabilities due to properties like texture, light rays, and refections.
- 5. The presence of moles on the human body, which is unrelated to skin cancer, can throw the system off track.

In this classifcation technique, transfer learning is used. It is a type of machine learning where a pre-existing model is applied to related problems. Training a DNN model from scratch requires a lot of images. These types of datasets with sufficient quality are sparse. The proposed model is a Deep Convolutional Neural Network (DCNN) model with the Adam optimizer as a gradient descent algorithm. An augmentation step is applied in the end to reduce the drawbacks of images. The dataset is divided into two ways as 70% training, 20% validation, 10% testing set, and 80% training and 20% validation. The results are put up against existing DNN models like AlexNet, DenseNet, and ResNet which are implemented using Python due to support of powerful libraries like Keras, Tensorfow and PyTorch. The best results are with 100 epochs and an accuracy value of 0.847 (Ali and Miah [2021\)](#page-50-14). Performance comparison with AlexNet, ResNet, VGG-16, DenseNet and MobileNet is presented below:

- Precision: The proposed DCNN model has a precision of 94.63, which is only lower than AlexNet, which stands above it with an accuracy of 96.89.
- Recall: The proposed model has a recall value of 93.91, with BesNet, VGG-16, and MobileNet besting it with recall values of 97.50, 95.18, and 94.57, respectively.
- F1 score: The proposed model has the highest F1 score with a value of 94.27.
- Training accuracy: The proposed model had a training accuracy of 92.69, just behind BesNet and MobileNet, which has higher values of 92.78 and 92.93, respectively.

• Testing accuracy: The proposed model has the highest training accuracy of 90.16 among the models.

In the following section the performance of the CNN model is presented with 80% training set against the above models.

- Precision: The proposed DCNN model has a high precision of 96.57, which is only slightly lower than AlexNet. Alexnet has a precision of 97.88.
- Recall: The suggested model has a recall value of 93.66, with DenseNet and MobileNet besting it with recall values of 94.83 and 94.15, respectively. BesNet has a very close value of 93.63.
- F1 score: The proposed model has the highest F1 score with a value of 95.09.
- Training accuracy: Training accuracy for the suggested model is 93.16, just behind AlexNet's higher value of 93.82.
- Testing accuracy: With a training accuracy of 91.43, the presented model outperformed the others.

The comparison of computation time is as given below:

- Time per epoch: Among all the models, the proposed model required the least amount of computation. It shows a tie per epoch value of $9-10$ s which is the lowest when compared to others. The only close model performance is by MobileNet, with 11 to 12 secs. The rest of the models take an average time of 14–17 secs for time per epoch value.
- Total time: The total time taken to compute images is 16 min, which is 3 min lower than the next highest performance of MobileNet. This is a great achievement. The rest of the models averaged around 24 min.

The proposed model is most suitable for CAD systems which will classify lesions at an early stage. The proposed model has a better classifcation rate than existing models and transfer learning models. The other models were trained on the same data set while com-paring them (Ali and Miah [2021](#page-50-14)).

6.4 Improved skin cancer classifcation using a combination of human and artifcial intelligence

With the advancement of artifcial intelligence in healthcare, various studies conducted in the field have often highlighted the comparison of computer-aided diagnosis with physicians to evaluate which is the more efficient of the two. However, the researchers combined both diagnostic methods in an efort to improve skin cancer detection. The authors fused the systems by frst obtaining an independent classifcation of the lesion and then merging the procured results to form an overall classifer, taking both systems into account. The task is to correctly classify the most common skin cancer lesions. The three systems compared includes dermatologists from German University Hospitals, a convolution neural network(CNN), and a combination of both.

The study utilizes the International Skin Imaging Collaboration (ISIC) archive, which contains dermoscopic images. The majority of the images used in the study came from the HAM10000 Dataset, which is a subset of the ISIC archive. The dataset was supplemented with an additional 4291 images from the ISIC archive, resulting in a

total of 11,444 images used in the study out of which 6390 were verifed for biopsy. To avoid bias for the 300 test images, a random generator is programmed using Python. An online questionnaire is circulated with the test images to 13 skin cancer experts at 13 university hospitals in Germany.

As a concentrated evaluation of 300 samples is impractical, the test set is splited into 6 questionnaires with 50 images each and randomly assigned to each clinic. The dermatologists verifes the skin lesions from the 50 test samples and answered on the identifed type of lesion along with the certainty of their response on a scale of 10 with 0 being least certain, 10 being very certain and 5 being most uncertain. To avoid careless responses, the authors have established statistical criteria for the detection of outliers.

The CNN Model is trained on the same training set which included 585 images of class a, 910 of class b, 3101 of class k, 4219 of class n, and 3521 images of class m accounting for a total of 12,336 samples of diferent classes of lesions. Considering the reliable classifcation of the ResNet50 architecture, the authors opted for the same model for this study.

To obtain the best results, the classifcation from dermatologists have to be fused with the CNN Model. The specification of confidence obtained from the dermatologist is also used as an input to the fusion algorithm as a probability metric of the selected class. XGBoost, an efficient tree algorithm with gradient boosting, is used iteratively to avoid bias in the results. RandomizeSearchCV is used for hyperparameter tuning to obtain the largest possible test set.

The results from the three classifers are evaluated, considering the diferential diagnosis task and how well benign lesions are distinguished from malignant ones. From the results obtained, the mean accuracy of physicians' classifcation is noted as 42.94%, while the CNN model accounts for 81.59% accuracy, and the fusion method shows the best results by displaying accuracy of 82.95%. The sensitivity and specificity of the three systems are also evaluated, and the results obtained are as follows.

The classifcation by dermatologists displayes a mean sensitivity of 66% and a mean specifcity of 62%. Those of CNN are 86.1% and 89.2%, and those of the fusion method are 89% and 84%, respectively.

From the evaluation, it is observed that the combination of man and machine increases the average sensitivity from 86.1% solely obtained through AI to greater than 89% by including human evaluation. Thus, the study concludes that the combination of human decisions with those of artificial intelligence achieves higher accuracy and better results and could be further utilized not only for skin cancer but also for prospective assessment of oncological transformations (Hekler [2019](#page-50-15)).

Artifcial intelligence is gaining ground swiftly in the feld of dermatology. It has the potential to revolutionize patient care, particularly by improving the sensitivity and precision of screening for skin lesions, including cancer. However, clinical and photographic data of all skin types are needed for AI research, and this can only be acquired through increased global skin imaging collaboration. It is necessary to record the sensitivities, specifcities, and performance in future research and in actual environments. AI is not a danger to dermatologists' knowledge; rather, in the next few years, it may be employed to enhance clinical practice. If practicing dermatologists had a greater understanding of AI concepts, they would be able to deliver better skin care. The review of the techniques displays an ever-increasing scope of AI and Computer-aided diagnosis in skin cancer detection (Das [2021\)](#page-50-16).

6.5 Deep learning techniques for skin lesion analysis and melanoma cancer detection

This section describes segmentation Techniques for skin lesions analysis.

- 1. U-Net architecture: Fully Convolutional Network (FCN) was the one to develop this. It is made up of a convolutional layer-based encoder-decoder network. Decoder layers are made up of upsampling layers, while encoder levels are made up of pooling layers. Due to the loss of image features caused by this approach, some important information may be lost.
- 2. FCN: It is a band of the pool and convolutional layers. When tested at the International Symposium on Biomedical Imaging (ISBI), a multi-stage FCN that was created with parallel integration produced a system with a state-of-the-art performance of 95.51% segmentation accuracy and 91.18% dice coefficient score.
- 3. Deep residual network: It is an ANN subtype. It uses skip connections to skip across convolutional layers as it rises in a pyramidal pattern. To extract features, a system with more than 50 layers was used. Because it consumes a lot of computational resources, this can limit its practical application.
- 4. Convolutional DE-Convolutional Neural Networks (CDCNN): Both convolutional and deconvolutional networks make up this architecture. Deconvolutional networks are CNNs that function in the opposite direction. For feature extraction, both networks are applied. These systems demand a lot of computing power. Due to the intensive adjusting of several factors, this method is not particularly practical (Adegun and Viriri [2021](#page-50-17)).

Skin lesion segmentation algorithms and techniques in ISIC 2018 database are given below. Major metrics for performance evaluation of the segmentation and classifcation deep learning models include:

- 1. Dice coefficient: This gauges the degree to which the actual result and the projected result are comparable and overlap.
- 2. Sensitivity: This is the proportion of positive outcomes(prediction) among those who are positive.
- 3. Specifcity: This is the proportion of negative outcomes(prediction) among those who tested negative.
- 4. Accuracy: In relation to the total number of cases investigated, it calculates the percent‑ age of true results (including true positives and true negatives).

Some of the best-performing cutting-edge methods used in ISIC 2018 have been investigated, and their performance has been documented.

- 1. A model developed using an encoder-decoder technique with Deeplab and PSPNET shows an accuracy of 94.2%, specifcity of 96.3%, and sensitivity of 90.6%.
- 2. The model developed using Ensemble VGG16, U-net, DenseNet and Inception v3 with Conditional Random Field (CRF) post-processing shows an accuracy of 94.5%, specificity of 95.2%, and sensitivity of 93.4%.
- 3. A model developed using a Fully convolutional encoder-decoder network with CRF shows an accuracy of 94.3%, specifcity of 96.4%, and sensitivity of 91.8%.
- 4. The model developed using a Fully convolutional encoder-decoder network with CRF shows 94.5%, 94.2%, and 94.0% accuracy, specifcity, and sensitivity respectively.
- 5. A model developed using UNet with DeepLabV3 shows 94.7%, 94.5%, and 94.5% accuracy, specifcity, and sensitivity respectively.
- 6. The model developed using FCN with a region-proposal network (RPN) and CRF shows 93.7%, 91.8%, and 92.1% accuracy, specifcity, and sensitivity respectively.
- 7. A model developed using Mask-RCNN shows 93.9%, 96.8%, and 90.4% accuracy, specificity, and sensitivity respectively.
- 8. The model developed using the Mask R-CNN model with ResNet50 shows 93.7%, 94.3%, and 93.6% accuracy, specifcity, and sensitivity respectively.
- 9. Model developed using U-Net variants shows 93.8%, 93.9%, and 94.5% accuracy, specificity, and sensitivity respectively.
- 10. Model developed using U-Net shows 93.7%, 97.9%, and 87.9% accuracy, specifcity and sensitivity respectively.

Similarly, models are also prepared for ISIC 2019 but results shows ISIC 2018 models performed better. The reason for better performance is found to be pre-processing images before the application of learning algorithms. Generally, ensemble models with state-ofthe-art models such as Resnext, NASNet, SENet, DenseNet121, InceptionV3, Inception– ResNetV2, Xception, EfficientNetB1, EfficientNetB2, and EfficientNetB3 seem to be more successful and perform at a high level. However, developing this model takes a significant amount of time and resources. Performance-wise deep learning algorithms are recom-mended for usage (Adegun and Viriri [2021](#page-50-17)).

Table [6](#page-36-0) represents the observations of AI techniques in the literature for skin cancer.

7 AI based techniques for digestive cancer detection

White Light Imaging (WLI) and Narrow Band Imaging (NBI) are widely used methods for gastric cancer detection. It has been discovered that narrow-band imaging (NBI) magnification endoscopy is more effective than white light imaging (WLI) for the early identification of gastric cancer.

AI methods are proposed for gastric cancer and stomach diseases. A high-perfor‑ mance computer, deep learning and a significant amount of digital data have been determined to be three components that make artifcial intelligence (AI) valuable to doctors in the field of picture identification in recent times. The very first CNN-based AI system for identifying gastric and esophageal malignancies has been developed by Japanese endoscopists in the feld of gastrointestinal endoscopy. Numerous researchers have noted the value of AI in separating cancerous from non-cancerous tissue using magnified NBIs. The system is created to forecast the depth of invasion of stomach cancer using artificial intelligence, and its specificity, sensitivity, and accuracy are noticeably better than those of experienced endoscopists. By utilizing its great sensitivity and speed, AI may be employed for picture verifcation following endoscopy to ensure that malignancies are not missed by accident. The decision to assist AI in the treatment of gastric cancer still needs a more thorough categorization than stage classifcation. Although AI proved successful in identifying cancerous lesions, the diagnostic precision was insufficient since it was unable to distinguish between malignant and benign lesions with appropriate accuracy. In order to address this problem, AI is utilized to

Table 6 Performance observations for skin cancer

identify gastric ulcers, a frequent condition that serves as a non-cancerous gastrointes– tinal lesion that must be separated from gastric cancer. Furthermore, the stomach's ana– tomical components can be recognized by AI. An artifcial intelligence (AI) system is created to anatomically categorize the upper gastrointestinal tract, locate the stomach in real-time, and assess how comprehensively the stomach is covered (Suzuki and Yoshitaka [2021](#page-51-12)).

AI plays active role in Pharyngeal And Esophageal Cancer detection. High-precision AI systems have been proven in reports to employ both still and moving pictures. To detect early esophageal cancer in a clinical scenario, AI systems still need to be improved, especially during screening endoscopies, in which the endoscope is swiftly moved through the esophagus. In terms of anatomy, it can be very difficult for nonexperts to see the cervical esophagus and the esophageal mucosa close to the esophagogastric junction. Additionally, many reports don't include validation flms with images that are of low quality due to defocus, halation, mucus, blur, and inadequate air insufflation. So, rather than using the actual speed in clinical practice, the researchers believe that high-speed videos should be used to test the AI system. It would also be useful if AI could predict the cancer risk with normal background esophagus in NBI or WLI, as the presence of several unstained areas after iodine stain indicates the possibility of esophageal squamous cell carcinomas (ESCCs). Such a prediction would help to detect cancers in the daily practice of endoscopic examination. With these developments, AI systems will be able to help endoscopists more successfully identify early esophageal and pharyngeal tumors, increasing survival rates. (Suzuki and Yoshitaka [2021](#page-51-12))

Methods exists in literature for endoscopic detection of early Esophageal Cancer. Current CAD systems are incapable of detecting Barrett's esophagus-related early neoplastic lesions as well as selecting a biopsy site. To solve this, a CAD system was developed using deep learning. The sensitivity and specifcity were higher than those of the existing ones. Ebigbo et al. developed a model using CNN with modalities of High Defnition- White Light Endoscopy (HD-WLE) and NBI. The authors used a dataset of 296 images, with 148 in each training and test. The results showed that HD-WLE had a sensitivity of 97% and a specifcity of 88%, while NBI had a sensitivity of 94% and a specifcity of 80%. De Groof et al. developed an HD-WLE model using SVM. He used 120 images, with 60 for training and testing datasets. It resulted in 95% sensitivity and 85% specifcity (Niu and Zhao [2020](#page-50-18)).

AI based Endoscopic Optical Coherence Tomography and Confocal Laser Endomicroscopy identifes lesions using the esophageal, mucosal, and submucosal structures. The primary challenges are complex imaging technology and high time consumption while reading images. After the application of Volumetric Laser Endomicroscopy(VLE) by Swager et al., the system becomes superior to VLE experts. Ebigbo et al. performed real-time diagnosis on 62 images captured with a CAD system, yielding a sensitivity of 83.7% and a specifcity of 100%. Veronese et al. developed a model using CLE images and support vector machine (SVM). This system could distinguish between gastric metaplasia (GM), intestinal metaplasia (IM), and neoplasia. It achieved a sensitivity of 96%, 95%, and 100% for GM, IM, and neoplasia, respectively (Niu and Zhao [2020\)](#page-50-18). Following are three important methods of AI in digestive cancer (Niu and Zhao [2020](#page-50-18)):

1. Artifcial Intelligence based on White Light Endoscopy and Narrow Band Imaging: To increase the accuracy of WLE, Cai et al. developed a CAD system using DL. This system is tested with WLE images. There is no signifcant diference between senior endoscopists. Several models are developed, but none of them gave any signifcant results. A few models managed to get better results than senior endoscopists, but the specificity dropped down.

- 2. Artifcial Intelligence in the pathological diagnosis of early Esophageal Cancer: When it comes to EC detection, pathological diagnosis is the way to go. Thus, many have tried to use AI in combination with pathological diagnosis. The results did not outclass the experts. The performance of the system as a whole was not far from an expert. A slight inferiority was observed.
- 3. Models developed using Deep Convolutional Neural Network for Gastric Cancer Detection: Using a backpropagation algorithm, deep learning allows us to break down images and extract relevant clinical features from them. The model trains itself on the accumulated images and then diagnoses new images. Currently, neural networks developed using CNN are best for image recognition(Hirasawa [2018](#page-50-19)).

Sabo et al. developed two models to distinguish low-grade and high-grade dysplasia. He made use of HE-stained (hematoxylin and eosin) image features. This model performed well in borderline lesions, which were not distinguishable before. Overall, HE-stained pathological image features may pave the way toward AI pathological diagnosis (Niu and Zhao [2020\)](#page-50-18). Model developed by Dehua Tanga, et al. is white light imaging and is one method for detecting EGC, but its sensitivity rate is only 40–60%. A system has been developed by the authors for the detection of EGC under WLI using AI based on deep convolutional neural networks (DCNN).

A study is conducted on 1568 patients belonging to 4 institutions in China. Around 45,000 images were gathered after examination by expert endoscopists, and they are used for temporal validation of the DCNN system. Another dataset containing 26 oesophago gastroduo denoscopy (OGD) videos are used for assessing the performance of the DCNN system. The external validation dataset contained around 1514 images. The testing dataset contained around 300 cancerous images and 300 control images (non- malignant). The boundaries of the cancerous lesion are marked by 4 expert endoscopists, and these boundaries are then circumscribed as annotation boxes by the computer.

The parameters of the neurons of the DCNN system are set to random values and then trained according to the images, with every image undergoing the same process multiple times. The architecture of deep networks has 2 parts: backbone structure for image feature extraction and detection and decision layers for detection of the location of the lesion. The backbone structure chosen for this system is the Darknet-53 model, which has 53 layers of neurons. The four coordinates of the surrounding box for the decision of location are predicted by the decision layer. These boxes are then classifed into three categories: small, medium, or large. The average accuracy of the system for the four institutions was found to be 88.2%. The sensitivity and negative predictive value was found to be greater than 85%. The specifcity of this system was found to be between 81.7 to 90.3%. The positive predictive value ranged from 80.5 to 90.5%. The AUC values were found to be quite high between 0.887 and 0.940, resulting in excellent performance of the system. The accuracy of the developed system(95.3%) was found to be higher than expert endoscopists (87.3%)and trainees (73.6%). The results clearly indicate that this DCNN system has much better results compared to expert endoscopists. The system can identify only EGC and pre-cancerous lesions under WLI but not NBI (Tang [2020](#page-51-13)).

The model developed by Toshiaki Hirasawa, et al. is presented in this section.

- 1. Preparation and data set of images: The dataset for this study comprises 13,584 images of gastric cancer lesions used for training and 2296 stomach images designated for independent testing. These images were obtained from two hospitals and two clinics in Japan during routine clinical procedures through standard endoscopic methods using conventional equipment. Exclusion criteria involved fltering out images of inferior quality and those acquired through specialized techniques like chromoendoscopy or NBI. An expert manually delineated cancerous areas in the training images, excluding magnifed and subpar-quality ones.
- 2. Construction of CNN: To construct the model deep neural network architecture known as Single Multishot box Detector is used (SSD, [https://arxiv.org/abs/1512.02325\)](https://arxiv.org/abs/1512.02325). No alterations to the algorithm are made. The SSD consists of 16 layers or more. Every CNN layer is fne tuned using stochastic learning gradients. It has a global learning rate of 0.0001. The images are resized to appropriate size along with the bounding box. For evaluation of accuracy an independent dataset is considered. This dataset consist of 69 patients. This dataset includes images which has 77 gastric cancer lesions. 62 patients has a single lesion, 6 patients has a couple lesions and a single patient has 3 lesions. The fnal dataset consist of 2296 images in total. Each case has 18 to 69 images taken.
- 3. Results: The model takes 47 s to analyse the 2296 submitted test images. It gives a sensitivity(detected number of correct gastric cancer lesions/actual number of gastric cancer lesions) of 92.2% and PPV (detected number of correct gastric cancer lesions/ number of lesions that are diagnosed as gastric cancer by the CNN) of 30.6%. There are two prime reasons for misdiagnosis, with the first one being the presence of gastritis, which induced a change in color tone, and the second being normal anatomical structures of the cardia, angulus, and pylorus. The missed six lesions are even difficult to diagnose even for experts.
- 4. Study limitations
	- (a) Use of high quality images only.
	- (b) Large training data set
	- (c) Use of only gastric cancer images. Endoscopic surveys shows low cases of gastric cancer.
	- (d) Occult cancer could be present in false positives.
	- (e) A single person marked the training and set images.
	- (f) Accuracy of CNN is not compared with that of endoscopists.

This CNN model detects early stage cancer. Despite all of its limitations, it achieves an incredible processing speed and have good accuracy. CNNs have an aptitude for diagnostics, and this could be leveraged further (Hirasawa [2018](#page-50-19)).

It would be possible to detect cancer early by using AI-based endoscopes. When it comes to early gastrointestinal malignancies, endoscopists' diagnostic skills and accuracy might be inconsistent, so AI technology's improved diagnostic capabilities may be helpful. Endoscopic diagnosis would be more accurate if AI and endoscopists' knowledge were combined (Suzuki and Yoshitaka [2021\)](#page-51-12).

Table [7](#page-41-0) shows the observations on digestive cancer detection techniques.

8 Discussions

After thoroughly evaluating the AI techniques for diferent cancer types, the following discussions are made and presented with the critical evaluation. The visual representations employed contribute to the efectiveness of conveying the comparative analyses. With a detailed review of each category of cancer, the authors present the following discussions. These discussions lead to the future development of AI in cancer detection and prediction:

For breast cancer, the evaluated techniques as shown in Fig. [3](#page-43-1) encompassed Convolutional Auto-Encoder, Artifcial Neural Network (ANN), AI-CAD, and ANN with extreme learning. Notably, the fndings underscored ANN's outstanding performance, securing an accuracy rate of 98.24%. Subsequently, ANN with Extreme Learning demonstrated the next-best accuracy at 96.4%. AI-CAD and Convolutional Auto-Encoder yielded an accuracy of 95% and 93%, respectively. Domain-specifc libraries such as Mahotas and sci-kitimage cater to the unique requirements of medical image analysis, contributing to the accu racy and reliability of breast cancer detection models. Based on these results, the study fnds the utilization of ANN as the optimal technique for breast cancer detection.

In case of lung cancer detection, prominent techniques such as Artificial Neural Net– work (ANN), k-Nearest Neighbors (KNN), Convolutional Neural Network (CNN), and back propagation-based Artifcial Neural Network (bp-ANN) have been evaluated. ANN achieved a remarkable accuracy of 100%, an achievement tempered by the limitation of a relatively small dataset used for testing. Following closely as it can be seen in Fig. [4](#page-44-0), KNN emerged as the top performer, boasting an accuracy of 99.51%. CNN demonstrated a commendable accuracy of 97%, leveraging its inherent fexibility. Concluding the set of techniques, bp-ANN exhibited a noteworthy accuracy of 96.6%. These results collectively underscore the efectiveness of these methodologies in lung cancer detection, with KNN standing out as the most accurate performer among the assessed techniques.

Artifcial Neural Networks (ANNs) are frequently developed using Python, and alterna‑ tive implementations can be pursued in Java or R. The implementation of ANNs is prominently facilitated through widely adopted frameworks such as TensorFlow, PyTorch, and Keras. It is crucial to highlight that these frameworks offer a sophisticated high-level interface, streamlining the process of constructing intricate neural network architectures.

Technique

Fig. 3 Performance evaluation of breast cancer detection techniques

Fig. 4 Performance evaluation of lung cancer detection techniques

Convolutional Neural Networks (CNNs) are conventionally constructed using program‑ ming languages such as Python, with a focus on leveraging the robust ecosystems of deep learning frameworks. Commonly utilized frameworks, such as TensorFlow and PyTorch, form the foundation for CNN development. The high-level abstraction provided by Keras, often integrated with TensorFlow, facilitates streamlined design and implementation of convolutional operations crucial for image-based data processing.

The k-Nearest Neighbors (KNN) algorithm is typically instantiated using languages suitable for numerical computations, with Python frequently chosen for its versatility. Implementations commonly employ specialized libraries, such as scikit-learn, renowned for its concise interface and efectiveness in the construction and training of KNN models.

The reviewed model was implemented using Matlab NNTOOL. However, Back Propagation-based Artifcial Neural Networks (bp-ANN) share an implementation approach with general Artifcial Neural Networks. They are often developed using languages like Python, with a focus on leveraging deep learning frameworks, including TensorFlow, PyTorch, and Keras. These frameworks provide built-in support for the backpropagation algorithm, optimizing neural network weights during the training process.

In order to detect and classify colorectal polyps, artifcial intelligence (AI) and machine learning algorithms are used. These include support vector machines (SVM) for medical imaging and surgical performance classifcation, random forests, decision trees, CNN, and deep neural networks. The CNN and deep neural networks are implemented using MAT– LAB's machine learning and neural network toolboxes to support the algorithms for cancer detection. Utilizing scientifc computer libraries and programming languages like MAT‑ LAB and Python, biophysics-inspired models were employed for numerical simulations and modeling.

In the context of colorectal cancer, the available range of techniques presented a diverse array of options, as illustrated above in Fig. [5](#page-45-0). The best performing models are Random Forest CNN Bayesnet and SVM, KNN, Decision Tree and Random Forest with accuracy of 97% each. Despite exhibiting equal performance Random Forest CNN BayesNet is preferred due to its neural network nature rather than being a conventional algorithm. This characteristic afords Random Forest CNN BayesNet greater potential for growth and adaptability.

Fig. 5 Performance evaluation of colorectal cancer detection techniques

In the evaluation of Prostate Cancer detection techniques, the examined methodologies comprised the Paige Prostate Algorithm, Multi-parametric Magnetic Resonance Imaging using AI, and Artifcial Neural Network (ANN). Notably, Multi-parametric Magnetic Resonance Imaging using AI exhibited the highest performance, boasting a 99% accuracy, closely trailed by the Paige Prostate Algorithm at 98%, while ANN demonstrated an accuracy of 93% as illustrated in Fig. [6](#page-45-1).

The Paige Prostate algorithm, a notable tool in prostate-related analyses, is typically implemented using Python, although Java or R can be considered as alternative languages. The preference for Python stems from its widespread use and robust support within the medical image processing and machine learning domains.

Multiparametric magnetic resonance imaging (mp-MRI) is implemented using dedicated libraries and frameworks tailored for medical image analysis. Python, with its versatile ecosystem, is frequently chosen, with libraries like SimpleITK or Pydicom

Fig. 6 Performance evaluation of prostate cancer detection techniques

playing crucial roles. Further integration with specialized image processing libraries such as scikit-image ensures comprehensive analysis of the multiparametric data.

The findings of this study unequivocally designate Multi-parametric Magnetic Resonance Imaging using AI as the superior technique. For scenarios necessitating the segregation of high-risk and low-risk cancer tissues, the Paige Prostate Algorithm is a more favorable choice due to its high accuracy.

In Fig. [7](#page-46-0) of Skin cancer detection the top models evaluated are GoogleNet, Transfer Learning, CNN, Image Processing with ANN and Back Propagation Neural Network using gray-level co-occurrence matrix.

The models like GoogleNet, Transfer Learning and CNN are implemented using Python with the availability of powerful libraries like Keras, Tensorflow and PyTorch. For implementing Image Processing and GLCM, we can use libraries like OpenCV or scikit-image in Python, or functions provided by MATLAB's Image Processing Toolbox.

BPNN can be implemented using neural network libraries like TensorFlow, PyTorch, or Keras in Python, or MATLAB's Neural Network Toolbox.

A notable observation is the reliance of models on CNN as their base. The illustrative fndings unequivocally highlight the capabilities inherent in neural networks, particularly exemplifed by two standout performers, Image processing with an impressive accuracy of 96.9% and GoogleNet closely following with a commendable accuracy of 96%. This clearly points out that neural networks have a high utility ceiling for cancer detection.

We fnd usage of ANN with image processing here as the all over best technique here but GoogleNet is a close second choice for skin cancer detection.

In Fig. [8](#page-47-0) evaluation of digestive cancer detection techniques such as Convolutional Neu– ral Network (CNN), Deep Convolutional Neural Network (DCNN), and Deep Neural Network (DNN) revealed that DCNN emerged as the leading performer, achieving an accuracy of 95.3%. These networks are commonly deployed in Python, utilizing libraries like Tensorfow, Keras, PyTorch, among others. This outcome not only signifes the notable utility of DCNN in the context of digestive cancer detection but also highlights potential of neural networks. The current accuracy of Deep Convolutional Neural Network (DCNN) holds promise for substantial improvement over time by training it on new samples. This indicates a high growth potential for DCNN, leading us to find this technique as particularly well-suited for digestive cancer detection. Specialized CNN implementation such as

Fig. 7 Performance evaluation of skin cancer detection techniques

Fig. 8 Performance evaluation of digestive cancer detection techniques

the Single Shot MultiBox Detector (SSD), Hirasawa (2018) (2018) , further supports the efficacy of tailored approaches in this domain.

Following a thorough comparison of various AI techniques, it is evident that deep learn– ing techniques such as Convolutional Neural Network (CNN) with further fne-tuning shows the best performance in detecting malignancies across a variety of cancers and can be a suitable choice for further research for detection.

A noticeable point is that CNN by itself wasn't up to par with the task. But the custom version of CNN performed a lot better than traditional CNN. CNN by itself is versatile in giving decent accuracies across any type of cancer but to bring out its full potential it needs to be tailored. The pairing of CNN with professional cancer specialists has the potential to signifcantly enhance the early diagnosis of various cancer types.

A series of prediction techniques were also studied in review, and following a thor‑ ough evaluation, it was found that an algorithm created by Qritive, an AI-powered platform had an accuracy of 91.7% in predicting Colorectal Cancer, whereas the accuracy of AI's Colorectal Cancer ANN algorithm prediction is rising in importance. Although, ANN prediction approaches for Breast Cancer showed an accuracy of 98.24%. CNN was used to predict Lung Cancer with a 70% accuracy rate and Skin Cancer using a region-based CNN. Paige Prostate an AI software has been created to identify signifcant area in prostate biopsy images, highlighting regions with highest probability of having cancerous tissue helping in the prediction of prostate cancer at an early stage. ANN produced the best outcome with the highest accuracy out of all the prediction approaches covered in this review.

8.1 Challenges with AI in cancer diagnosis

The following are the major challenges associated with using AI techniques for the detection as well as the prediction of cancer:

- 1. Bias in results caused by bias in training data or design decisions is a signifcant and well-observed issue for machine learning systems.
- 2. Transferability is a signifcant concern for AI; successful algorithms frequently fail when applied to various contexts.
- 3. Because of the dependency of AI on computational requirements for information management, AI approaches becomes expensive.
- 4. Sensitizing people to the ethical dilemma of using patient data without their agreement in some circumstances is crucial.
- 5. Confdentiality of Data- Large amounts of high-quality data are needed for training and validating artifcial intelligence (AI) systems. Critical problems include who owns the data, who permits its usage, and how it is protected. Despite the concerns, implementation is now moving forward with signifcant enthusiasm. Strong motivations for the quick adoption of AI are created by the social, cultural, and economic systems in healthcare. It is very challenging to adopt traditional opt-in or opt-out consent methods. A few AI developers are working on novel methods of data security. Any health system that wants to use deep learning applications must frst deal with data privacy as a fun‑ damental first step.

AI prevents emotional problems, social and ethical norms, and tiredness. However, radiotherapists and other medical professionals will not be replaced by AI in the medical feld. AI is not completely self-dependent and cannot overrule human participation.

9 Conclusion

The majority of developments have been in the field of "narrow AI", which are now capable of performing a wide range of specialized tasks, such as playing board games, translating between languages, listening to and acting on human commands, or spotting specifc patterns in visual data (such as recognizing faces in CCTV images or suspicious areas). However, lately AI has made a signifcant impact on the healthcare industry, particularly in the feld of cancer prediction. An in-depth study of the machine and deep learning models used in cancer early detection utilizing medical imaging is provided, which ofers a critical and analytical assessment of the most cutting-edge cancer diagnostic and detection methodologies now available.

In the clinical and translational fields of oncology, the power of AI regarding therapeutic advice in cancer shows great potential, which translates to better and more individual– ized therapies for people in need. AI-based computer-assisted systems can help physicians with essential aspects like detecting and classifying various types of cancers. To overcome AI's shortcomings, physicians and technicians must work together to create clinically relevant solutions.

Deep learning technologies significantly improves the way we acquire data, make diagnoses, and treat cancer. Though methods for obtaining medical information may be dis‑ puted, there is little doubt that cancer early detection will have a major impact as soon as acceptable data collection methods are discovered. In the numerous felds of study that AI-based cancer prediction algorithms are applied to using machine and deep learning approaches for extracting and categorizing disease characteristics, AI techniques play a vital role in early cancer prediction and diagnosis. With more active efforts toward developing novel biomarkers, creating tailored treatments, and accumulating big datasets, the prospective advantages and reach of AI in cancer detection are only going to improve. It further draws information from digital health tools and help physicians make accurate medical decisions.

The outcomes of AI-based models are promising and provide a competitive edge over traditional techniques. Future research will have to overcome several restrictions to develop clinically relevant techniques. Overcoming real-time limits, as well as the capacity to identify additional components like folds and blood veins, and the capacity to get uncer– tainty estimates from machine learning and artifcial intelligence model predictions. In this instance, the promise of AI-powered technologies is that they will revolutionize the delivery and consumption of healthcare, signifcantly improve cancer prevention, and open up enormous fnancial opportunities.

10 Future scope

This review promotes future research opportunities. Hyperparameter tuning techniques may be used to further improve the performance of current models. The use of ensemble models, which may combine machine learning classifers might improve performance and robustness. Identifying cancer may beneft from the inclusion of additional deep learning models and architectures, such as the combination of CNN-RNN models. Improved feature extraction from digital images may be used as a reliable input to models, enhancing their performance even further.

AI implementation may alter the options available to patients in more or less predictable ways. By using personal health and non-health data in addition to research information, for instance, the range of therapy options may be modifed and restricted, giving the patient fewer alternatives than are now available. AI certainly has the ability to produce both good and bad outcomes. However, every algorithm will encode values, either overtly or, increas– ingly frequently in the 'new AI' age, implicitly.

The authors also present here the future research direction of AI in Colorectal Cancer. It is observed that deep learning technologies significantly improve data acquisition, diagnosis, and treatment of colorectal cancer. Though methods for obtaining medical information may be disputed, there is little question that colorectal cancer early detection will have a major impact as soon as acceptable data collection methods are discovered. Modes for related patterns also increase significantly. In the clinical and translational fields of oncology, the power of AI regarding therapeutic advice in colorectal cancer shows great poten– tial, which translates to better and more individualized therapies for people in need. The outcomes of AI-based models are promising and provide a competitive edge over tradi‑ tional techniques. Future research will have to overcome several restrictions to develop clinically relevant techniques. Overcoming real-time limits, as well as the capacity to identify additional components like folds and blood veins, and the capacity to get uncertainty estimates from machine learning/artifcial intelligence model predictions.

AI-based computer-assisted systems can help physicians with essential aspects like detecting and classifying colorectal polyps. To overcome AI's shortcomings, physicians and technicians must work together to create clinically relevant solutions.

Declarations

Conflict of interest All the authors certify that they have no affiliations with or involvement in any organization or entity with any fnancial interest (such as honoraria; educational grants; participation in speakers' bureaus; membership, employment, consultancies, stock ownership, or other equity interest; and expert testimony or patent-licensing arrangements) or non-financial interest (such as personal or professional relationships, afliations, knowledge, or beliefs) in the subject matter or materials discussed in this manuscript.

 Research involving human participants and/or animals No human or animal was harmed while performing this review.

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