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Blood Cancer Detection from Blood Smear Images using Machine Learning Algorithms

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Abstract

Artificial intelligence techniques in computer vision have made a substantial contribution to the development of imaging analysis in medicine by improving the accuracy of predictions, which has resulted in more suitable treatment and diagnostics. By offering a second view, these techniques can help hematologists and other medical professionals make better diagnoses in the area of automated leukemidical imaging and cancer in the blood diagnosis. A thorough examination of the existing DM and DL image processing algorithms is provided in this study, with a particular emphasis on how to identify of leukocytes in smear blood imaging alongside various clinical imaging regions. The primary aim of the suggested investigation is to identify the best DM and DL techniques for clinical imaging, particularly for identifying the types of lymphocytes from smear blood data. This review article delves closely into the sophisticated algorithms for DL, especially the emerging models built around CNNs that operate in the clinical computational imaging area. According to a review of associated research, white blood cell identification using micro smear captures is a common application of standard AI technologies. They aid in the diagnosis of numerous illnesses, including blood cancer, and give medical professionals important information. As the researchers and professionals assigned to the processing of medical images, we construct recommendations for future investigations based on the extensive analysis of white blood cell associated research study that is laid out through the current research.

Keywords: Blood Cancer; Leukemidical Image Processing; Medical Image Processing; Smear Blood Images; AI; CNN; Deep Learning; Data Mining; White Blood Cells Classification.

1 | Introduction

Artificial intelligence (AI) approaches have become commonplace for many purposes, including the interpretation of images from hospitals [1]. Medical imaging evaluation has become a crucial component of contemporary healthcare systems, providing intelligent support to medical specialists. Image analysis in medicine processes many image modalities, including CT scans, MRIs, blood smear pictures, and ultrasounds, and is essential in the diagnosis of numerous disorders [2]. Regarding diagnosis and academic objectives, imaging paradigms are essential in medical imaging evaluation because they help identify and categorize both

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soft and hard tissue of various human organs [3]. Professionals in computer vision have a lot to contribute to the analysis of medical images. AI is important for leukocyte identification of cancer, healthcare data commentary, and retrieving images into computer-assisted diagnostics. Since the efficacy of AI algorithms immediately impacts medical diagnosis and therapy processes, the use of computers in detection depends on them [4, 5]. It facilitates how doctors use conventional working methods and helps them even more with diagnosis and therapy. Computer-aided design is greatly impacted by contemporary technological advancements in device fashion, capacity for storage, and high rates of processing power. Leukocyte characterization using MRI scans was previously an important application for the computer-aided design system using AI. This gives medical professionals crucial knowledge that aids in the diagnosis of many hematic issues, including blood-related cancers. The primary goal of image processing is to support clinicians efficiently during the diagnosis procedure. Within the medical community, leukocytes or WBCs are seen to be a good indicator of most human disorders. Diverse individual illnesses can be identified by variations within the dimensions, form, and hue of leukocytes as well as by changes in their geometrical structure, as seen in smear imaging. Different blood cell types exist, including leukocytes, which are further classified into five subtypes. Different AI methods have been developed for recognizing and classifying white blood cells in images of micro-blood smears. Traditional methods rely on the laborious, difficult, and error-prone manual examination of WBCs in images of smear blood [6-9]. Diagnostic procedures, along with suitable therapy, are greatly aided by automatic systems [10]. As a result, automated Leukocyte detection through microscopically scanned blood sample data has become increasingly prevalent since it may minimize physicians' load and offer precise findings that help professionals during the procedure of diagnosis [11]. Computerized Leukocyte classification in blood sampling imaging could be accomplished primarily through two methods: data mining (DM) and deep learning (DL) approaches. Both offer the possibility of helping produce computerized systems that can improve the efficiency of clinical hematologics [11-13]. Figure 1 displays an extensive outline of the DM versus DL Approaches for the classification of white blood cells. By examining leucocytes, several computer-aided diagnosis techniques may automatically identify a variety of hematic conditions, including cancer of the blood [12]. DM involves a series of interrelated stages, like ROI segmentation, the extraction of features, and appropriate classifications. For determining the ROI from an image, a range of methods for data mining can be employed [13]. Using the DM techniques, the extraction of features is an additional process.

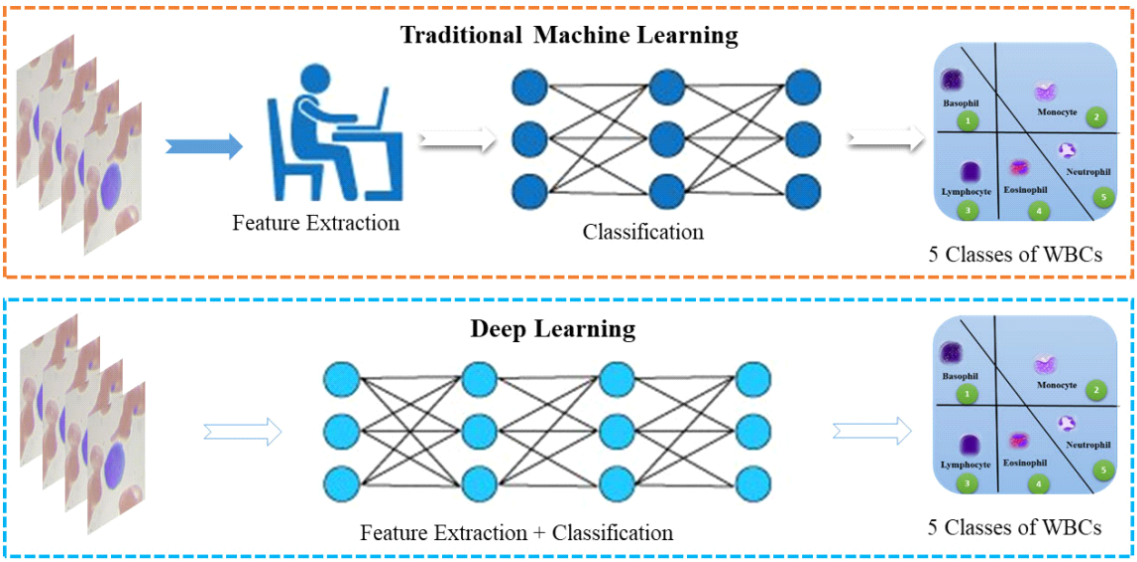

Figure 1. Summary of traditional ML and DL models for the classification of leukocytes.

Nonetheless, since feature dynamics are different, choosing the best feature extractor might be difficult. Exceptional image processing approaches are now available because of the widespread adoption of DL methods, particularly in clinical hematology when blood pictures are used [14-24]. This work aims to classify

WBCs in blood pictures and offers an extensive overview of the various deep-learning algorithms that are currently offered and their medical implications. Microscopic blood smears remain an issue that is rarely addressed, even though there have been numerous studies on image processing employing methods such as DL along with future trends aimed at CT, MRI, or a chest X [14, 25]. Thus, by examining cutting-edge AI techniques—specifically, leucocyte categorization methods—our study aims to close this gap. The main goal of the research topic is to present an extensive analysis of the application of DL.

The most popular DM and DL techniques are identified in this suggested study using unconventional classifying methods, which are then evaluated in different groups based on the approach and research topic. This work also contributes to the direction of future studies by using DL methods for the classification of leucocytes in images from blood smears. Among the important contributions of the suggested review work are the following:

- This paper explores the numerous possibilities and uses of employing AI algorithms in blood cancer classifications.
- It also identifies the DL methods that currently exist for leukocyte identification and evaluates their accuracy.
- We discuss the main obstacles and needs for AI algorithms, then we discuss their prospects and possibilities for computational imaging research.

The rest of the article is organized in the following manner: Part two goes into great length about the work examination process and reviewing technique. A short overview of the processing of medical images is provided in Part Three. The comprehensive overview, along with uses regarding deep learning in leucocyte identification based on blood images are presented throughout the fourth part. The prerequisites and present concerns are covered by part five. The sixth part outlines the planned review study's prospects for the future. The seventh part concludes with a discussion of the latest advances in models of DL and the recommendations for further research.

2 | The Research Methodology

A thorough explanation of the digital libraries employed in the planned review study's formal research process is given in this section. To identify the literature that meets the search requirements and make deliberate use of the accessible digital resources, a planned search process is necessary [15]. Considering the Recommended Items for Reporting over Systematic Reviewing, we used both human and automated searches within the suggested study to find the most pertinent research papers [16]. We used both automatic and manual search functions to retrieve the most pertinent content. Our search approach starts with a machine-generated search across digital records to find pertinent information. Then, image processing and leucocyte classification specialists verify the results. We look for papers published between 2014 and 2024 for the suggested survey. Given the selected keywords, both a computerized and customized search is conducted across all the available sources. The authors choose "Medical Imaging processing," “Classification of Leucocytes," OR "WBC's Identification in Bloody Smears Images" as some of these keywords. These steps are used for establishing search keywords:

- Research questions serve as the basis for the main terms.
- For the main terms, alternate equivalents or spelling are found.
- Publications and academic papers that are pertinent are used to find keywords.
- To switch up equivalents, use the Boolean operator's OR.
- The AND operator links the important phrases, and the query is created when the key phrases are analyzed to obtain pertinent data within the sources.

The aforementioned terms are examined in every record and recourse, and their pattern is adjusted under the retrieval of pertinent results. Several terms related to the main topic of the study are derived from the four questions being investigated, which are formulated with consideration for the Patients, Interventions, and Result frameworks [17].

RQ1: Which AI methods are available for classifying leukocytes?

RQ2: How are DL methodologies used in medical research, specifically in the classification of leukocytes?
RQ3: In the medical imaging processing area, specifically for leukocyte identification, how are DL approaches applied?
RQ4: What kind of DL is useful and effective for identifying white blood cells in images of blood smears?

3 | Evaluation Standards for Extracted Studies

The PRISMA rules are followed to apply inclusion/exclusion criteria to the papers that were initially obtained. The requirements for being included and excluded for weeding out irrelevant items are shown in Table 1. There are three phases involved in the selection of articles for research. [18] First, redundant documents are eliminated. Subsequently, the relevance of the article's heading, abstractions, and search terms is examined, and following a comprehensive examination, the additional studies were considered. Bias and conflict analysis are eliminated by the application of criteria for inclusion and exclusion. Throughout the publication's process of selection, a total of 1893 studies were gathered for assessing the existing research based on the study's emphasis. 1319 publications were retrieved after the first process of selection, which included personal sorting. The works have been reduced by employing the pertinent title. After reviewing the headline and summary, these 1319 publications have been reduced to 894 studies. Following the next step, these studies are selected based on approach and findings, yielding 633 publications. After reviewing the entire contents, the articles were filtered, leaving 286 articles. We assessed the technique, findings, impact factor of the journal, and citations to determine the worth of the remaining papers. Eighty papers are selected for the planned study after each of these parameters has been verified. The quality assessment is carried out following the conclusion of the article's scrutiny procedure (paper inclusion and exclusion). Every research article is evaluated considering the scrutinization standards. Every research paper is examined, and the quality of every research issue is evaluated. The authors carefully examine and assess each of the chosen papers.

Table 1: Qualifications for papers inclusion and exclusion.

Qualifications for inclusion	Qualifications for Exclusion
The academic publications were authored in the English language.	The academic publications were not in English.
The papers adequately addressed the central idea of the subject under investigation.	Papers that don't adequately clarify the suggested study's primary subject are disqualified.
Just appropriate academic publications with definite conclusions in addition to evidence.	The articles lacked sufficient comparison or unambiguous outcomes.
Current works published between 2016 and 2024.	The academic works are duplicates.

4 | Medical Image Processing

Medical imaging is a procedure that can give visual data about the body of a person that helps doctors and physicians diagnose and treat patients effectively [19]. Medical professionals and specialists use a variety of imaging techniques to diagnose conditions and provide treatments. These types of imaging comprise ultrasound, PET, MRI, CT, X-ray, and pictures from samples of blood [19-20, 25]. These techniques for imaging are crucial to medical image processing because they enable medical professionals to identify and diagnose a variety of chronic illnesses automatically through image analysis. To do a study, they can observe various parts of the body [21]. Over the past twenty years, there has been a significant increase in the use of medical imaging in computer-aided diagnosis. Examples of these applications include the classification of

leukocytes, the classification of tumors, and cancers of the breast identification [22, 23], images guiding treatment [24], as well imaging annotations [25, 26]. As a result, it is now an essential component of contemporary healthcare systems [27].

5 | Blood Cancer Classification Based on Data Mining (DM)

There are plenty of newly released scholarly reviews on AI methods for medical image processing in the literature. The following sections cover the most current and pertinent research on DM and DL techniques in medical imaging, namely for identifying the types of leucocytes in blood smear pictures [28]. The most recent and pertinent research is found in the proposed study by filtering the recent publications using the keywords "leucocytes detection" or "leucocytes classification." During our search, we discovered that the use of DM and DL techniques for the study of leukocytes in blood sample images is expanding exponentially. Image initial processing, segmentation, extracting features, selecting features, and classification are all interrelated processes in DM methods. [29] Image-enhancing techniques, including contrast modification, noise reduction, and sharpening, are part of the pre-processing procedure. [30] Before image segmentation, all of these procedures have been applied to an input image [31]. There are several different pre-processing methods, including Gabor, low pass, high pass, and median filters. [32] Before picture segmentation, these are typically employed for image sharpening, contrast modification, and noise reduction. [33] Several studies have focused on DM to detect and categorize leukocytes. Nevertheless, these methods have made it difficult and time-consuming to recognize nuclei accurately, segment ROI, separate borders to recover overlapping cells, extract robust features, and choose the best features [34]. Using that method, following separating ROI, the following process features extracting. Classification in classic supervised learning approaches relies on selecting the optimal features selection algorithm and a robust features descriptor [34], which are the most important elements in ensuring the technique's accuracy and efficiency. Figure 2 provides a general overview of TML. Leucocytes in microscopic blood sample images have been classified using a variety of traditional supervised learning techniques, including SVM [35], NB [36], KNN [37], and ANN [31].

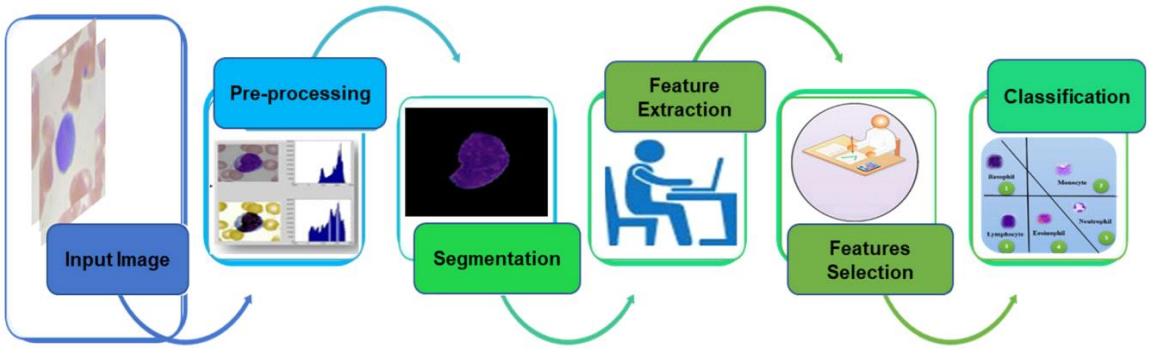

Figure 2. TML-based leucocytes segmentation and classification.

Leucocyte classification can be handled by a variety of supervised learning methods, including SVM, ANN, and Decision Trees. Hegde et al.. [38] presented a novel method wherein the authors classified WBC cells as either normal or leukemic cells by first segmenting the WBCs and then using SVM. A unique method for segmenting and classifying leukocytes in a smear of blood pictures was proposed by Zhao et al. [39]. To categorize WBCs under their five subclasses, color correlation, and morphological-based segmentation were used, followed by the extraction of texture data and classification utilizing SVM [40]. ANNs and SVMs are important components of medical image processing, but hybrids, Bayesian, Ensembles, KNN, and Tree models are also being utilized to solve problems in various sub-domains of medical imaging, including brain tumor detection, lung tumor identification, leukocyte classification, etc. [41, 42] Abdulkadir et al. [43] suggested a combination of techniques for WBC classification in blood smear images, and Sajjad et al. [12] provided a smartphone-based superior system for healthcare regarding smart cities, in which WBCs are classified using an ensemble multi-class SVM. Tantikitti et al. [44] suggested an automated technique for diagnosing dengue fever sickness. In blood smear images, leukocytes are segmented using a multi-level

threshold approach. This study uses two models of decision trees for classification. The initial model was used to determine if white blood cells were either lymphocytes or phagocytes. The second approach classifies the infection with dengue virus as either negative or positive. Work in [45] proposed a novel technique for segmenting WBC nuclei and cytoplasm using simple thresholding. Upon segmentation, some morphological procedures are carried out with elliptical curve fitting after feature extraction. The sequentially forward selection strategy is utilized to pick features, and the NB classifier is then used to categorize WBCs. Vogado et al.. [46] employed a hybrid technique to classify and segment leukocytes. In their suggested method, CNN features are employed in the given data for training the SVM classification.

6 | Blood Cancer Classification Based on Deep Learning

With the use of DL, we may create a framework where a broadly applicable learning technique is used to extract features from data rather than having to be created by individuals [47]. Deep learning produced good results in the domain of image processing for medical imaging and building a complete network with CNN was very simple [48]. To accomplish various categorization tasks, DM algorithms are trained by using physically extracted characteristics or by using other basic machine learning approaches to acquire features. [49] As a result, researchers are becoming more interested in investigating the advantages of DL approaches for WBC categorization. These days, deep learning is a potent research instrument in the fields of AI, voice analysis, the processing of human language, and imaging in medicine [50]. In the realm of medical image processing, the application of DL as a pattern detection tool is also becoming increasingly important [51]. A recent review on deep learning (DL) -oriented image processing [52] claims that numerous individuals now choose to examine healthcare data using DL methods, especially convolution networks. These techniques work especially well in fields where analyzing vast amounts of data requires intelligence comparable to that of a person. Furthermore, for obtaining valuable characteristics given many initially collected data, a solid understanding is required [53]. Nevertheless, when a large amount of data needs to be managed effectively, this work becomes difficult and takes time. As illustrated in Figure 3, DL offers complete learning and removes all unnecessary overheads associated with the choice of features as well as feature descriptor selection.

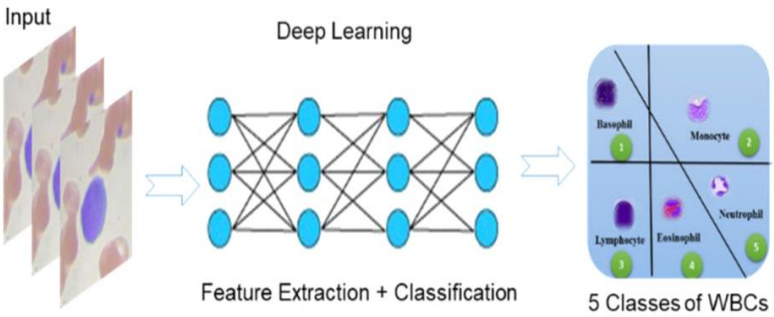

Figure 3. Deep learning for Blood Cancer classification

The key benefit of deep learning techniques is their ability to automatically identify and extract linguistically valuable characteristics from unprocessed input. That's the primary distinction between the DL versus DM approaches. [54] A sizable field of study has been drawn to employ DL-based techniques in analyzing medical images due to its unparalleled advantages. DL models fall into various categories, including CNN, DBN, LSTM, and RNN. CNN is a popular tool for imaging in medicine. Numerous convolutional, collecting, completely linked, and activation function-equipped layers make up CNN. It is trained similarly to every other regular ANN by employing the methods of gradient descent and back-propagation (as seen in Figure 4) [55]. A completely interconnected tier comes after the layers of pooling and convolution of an ordinary CNN. The results from the networks classify WBCs into their five separate groups using the Softmax method. Using the BCCD database, Banik et al. [56] developed a unique CNN model for classifying WBCs by combining the properties of the initial and final convolutional layers. The CNN structure comprising eight distinct levels was proposed by Choi et al. [57] for the classification of WBCs. An LSM-TIDC technique was presented by

Karthikeyan et al. [58] to categorize white blood cells using blood photos. Images are processed beforehand, followed by a multidirectional approach is used to extract characteristics of geometry and texture. Ultimately, DCN receives the extracted features to effectively and early detect White blood cells. The authors of [11] suggested a methodology that classified White blood cells in blood photos employing CNN. Table 2 provides a summary of a few recently published studies that used DL in blood cancer identification.

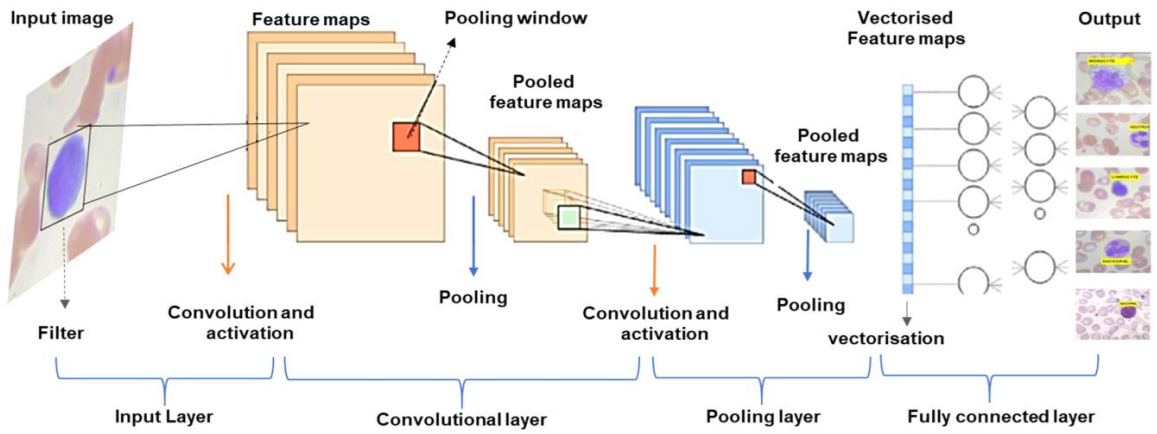

Figure 4. Building blocks of general CNN architecture for leukocyte classification.

Table 2. The major contributions to the Blood Cancer classification using DL models.

Ref.	Technique/s	key contribution
[11]	RCNN	A DL-based computer-aided method has been proposed that can quickly identify and categorize the different types of WBCs in blood sample data.
[28]	CNN& RCNN	They used CNN-based intensive machine learning for the classification of WBCs, with a precision of 94.71% for RBC & its anomalies detection and 98.68% for WBC. They also used an effective contouring aware CNN, as well as a quicker R-CNN, centered segmented strategy for dividing RBC and WBCs.
[39]	CNN & RF	With an accuracy of 92.8, an automatic technique utilizing CNN features and a random forest classifier is suggested to separate and categorize WBCs of central blood scans into the appropriate subclasses.
[44]	CNN & SVM	After incorporating a transfer learning strategy into CNN architecture to extract features, an SVM algorithm is utilized to categorize blood cells as normal and leukemic categories achieving an overall accuracy of 99.20%.
[59]	CNN	SSD and a YOLOV3 Detector Incremental Improvement Version are used for CNN-based leukocyte recognition. The mean accuracy and mean precision of the procedure were 90.09% and 93.10%, respectively.
[60]	ResNet50	Prior to network training, a leukocyte mask is first applied at the pixel level. Next, the ROI is segmented, and lastly, the trained model is applied to classify leukocytes in blood smear images with 0.98 accuracy and 0.99 precision on datasets 1 and 2.
[61]	AlexNet	White blood cell categorization achieved a standard sensitivity and accuracy of 99% through a comparative analysis of CNN-based features extraction and classical learning.
[62]	CNN	With an average accuracy in classification of 96.2%, the primary contribution of this study is an approach to classification built around a trained CNN that can identify eight distinct categories of cells in the blood floating in microscopic blood sample images.
[63]	DRNN	Using a fine-grained method based on the deep residual learning theory, leukocytes are classified from microscopic blood test images. The typical accuracy during testing was 76.84%, the top 1 accuracy was 77.80%, and the top 5 accuracy was 98.75%.
[64]	fine-tuned CNN	A refined CNN method, which is capable of classifying blood smear images into their subgroups of cells in blood with a mean accuracy of 0.97, was proposed by the authors of this paper.
[65]	RCNN & AlexNet	RBCs and other cells, including infected and WBCs, can be automatically detected and classified using a two-stage DL-based approach, which has a 72% overall accuracy rate for non-difficult infected cells.
[66]	ResNet & Inception	Using a sequentially proposed DL framework, WBCs were classified into four main types. ResNet 101 and ResNet V1 152 achieved an accuracy rate of 99.46% and 99.84%, respectively, after fine-tuning all layers over 3000 epochs.
[67]	CNN	A region of interest identification technique based on CNN and superpixels is suggested for cervical cancer cell segmentation. The experimental results showed a 94.50% accuracy rate.

[68]	CNN	CNN models that have been pre-trained are utilized as feature extractors. The extracted features from completely connected layers were then combined, and MRMR features selection and an extraordinary learning machine were added to classify WBCs with reported accuracy rates of 99% and 97.37%.
[69]	DCNN	A brand-new method for classifying ALL in colored bone marrow pictures is put forth. After segmenting the lymphocytes using a straightforward threshold method, the segment lymphocytes are subjected to CNN-based DL algorithms, which successfully categorize ALL into L1, L2, L3, and normal lymphocytes with an accuracy of 97.78%.
[70]	DCNN	To categorize WBCs into their five distinct groups, one study suggested using CNN. The validation set has an overall accuracy of 91%, whereas the training set achieves an accuracy of 99%. The reported sensitivity and specificity are 91% and 97%, respectively.
[71]	AlexNet	After pre-processing the blood smear image, characteristics are extracted using a CNN model called AlexNet. Once AlexNet has been fine-tuned, it is capable of accurately identifying blood cells as normal and pathological cells.
[72]	DCGAN & ResNet	A comprehensive detection framework built on DC-GAN and ResNet was proposed by the authors for blood cell images, and its accuracy was 91.7%. Additionally, transferable learning is used to increase the accuracy of the model.
[73]	capsule network	To classify WBCs, a novel deep-learning technique is proposed. First, data is augmented using various methods of augmentation to increase the amount of data. Next, a DL-based capsule network serves to classify WBCs into its five subclasses, reaching 96.86% accuracy.
[74]	R-CNN	To help medical professionals diagnose various hematic diseases automatically, a deep learning-based method called Faster R-CNN is proposed to detect WBCs and RBCs. The accuracy of the test is 83.25% for RBCs and 99% for Eosinophils, with the lowest testing accuracy not falling below 66%.
[75]	FPM & SOYOLO DL	This work built a hybrid method for WBC identification by combining the SOYOLO DL framework with FPM. The high-resolution WBC blood smear images are obtained using the FPM approach, and SOYOLO detects WBCs in these images with a 100% recall and 100% precision rate.
[76]	DCNN	Lightweight DL models that effectively categorize RBCs into three groups—sickle cells, normal, and other blood content—were provided in this study.
[77]	hybrid DL	A hybrid DL-based approach showing 95.17% sensitivity, 98.58% specificity, and 96.17% overall accuracy is suggested for the classification of adolescent leukemic blast and regular cells.
[78]	AlexNet	Blob detection and DL approach are used to identify WBCs in tiny blood smear images. The accuracy of this new technique is 100% for SMC-IDB and IUMS-IDB datasets, 99.7% for ALL-IDB in white cell detection, and 94.1% for leukemidical image analyzing identification.

7 | Current Challenges and Requirements

We discovered the primary demands and research problems, several important characteristics, uses, and benefits of DL approaches for image processing, especially for blood cancer detection, through analysis of the existing literature. Some common and potent machine learning algorithms have been established in the past couple of decades for healthcare image processing, including the identification of brain tumors from MRIs, the detection of blood cancer using smear visuals or the detection of lungs using CT scans [79]. However, there are still a few major obstacles that the scientific community must either accept or work towards overcoming. Specialized medical expertise, efficient DL approaches, and superior data sets that are publicly available are some of these issues.

A few of the difficulties stem from the conceptual and mathematical foundations of numerous deep learning approaches. Unsupervised and supervised approaches have to be developed to address these obstacles. To prevent these problems, the effectiveness of unsupervised or supervised image processing techniques can be undermined. Transitioning from supervised to unsupervised learning methodologies has difficulties as well, as it impacts system performance and accuracy. Applications and platforms for image processing that use deep learning techniques remain not exactly ideal, with plenty of opportunity for development. The lack of publicly available data represents the main issue facing the medical imaging industry. To solve this problem, scholars must persuade health institutions to provide healthcare data; high-quality data that is available to the general public to feed research purposes can be fascinating. Efforts that promote open datasets from various health organizations across the globe are additionally welcomed; however, certain operations (such as clinical data and limited availability of data) are needed as well. In each of these scenarios, incentive systems may be linked to monetary gains, amusement, or services rendered to these organizations in exchange for high-quality

data. Whenever large volumes of data are available, the problem gets more intriguing for research, like other domains (such as summarization of videos, Internet of Things, and controlling energy). For certain image processing applications, gathering large, accurate data sets with truth labels is essential. Such data sets can also be utilized for evaluating and organizing other tournaments. [80]

The training of predictive algorithms is the main problem in image processing and blood cancer identification and classification. To solve this issue, an optimal learning method with a more balanced capacity for generality and a computationally effective heuristics approach needs to be developed. For the development of an algorithm with impressive generality skills, an approach to learning that employs either real or arbitrary labels offers helpful instruments to deal with accessible datasets, along with effective algorithms for training. In the past few years, training utilizing DNNs achieved tremendous experimental success in a range of tasks associated with image processing. Even though this is a difficult, non-convex optimization issue, straightforward techniques like random gradient descent may uncover workable solutions that reduce the error generated by training. Even though the total quantity of parameters is far greater than the quantity of training data [49], the networks learned in this manner show a remarkable capacity to generalize. Merely reducing the training errors has become insufficient when learning a machine learning model. Selecting the incorrect global minima may also cause the estimator to behave poorly when it comes to generalization. Under such circumstances, the strategy employed to reduce the error in training essentially determines the behavior of generalization. Various global minimums with varying capacities for generality will result in numerous optimization algorithmic decisions, including beginning, updating guidelines, training rate, and terminating conditions. Without medical competence, the current DM and DL approaches are not reliable enough to be used in real-world health diagnosing platforms [32]. Developing a model for learning to assist with medical imaging processing and classifying blood cancer needs both technical and expert knowledge. We must investigate highly accurate and reliable techniques that can be applied in practical healthcare situations without the requirement for medical specialists.

8 | Future Research Directions

A significant amount of work is required from the healthcare business and research society to contribute to MAI and specifically leukocytes that are processed in the smear of blood pictures, given the considerable obstacles facing the image processing field as described in section V. We have concentrated on the most commonly brought up issue in this study, which is the lack of processed medical images and leucocyte classification datasets. To enhance image processing and WBC identification and categorization in blood smear images, a comprehensive data enhancement method, and transfer learning models are advised. To enhance the available data, a variety of data augmentation techniques are employed, such as generative adversarial networks, style transfer, and traditional picture alterations. The latest developments in deep learning models will be helpful for computer-aided diagnostic applications in the future.

These models are currently accessible on several open-source software platforms. However, due to a lack of clinical and medical competence, choosing and training a suitable machine-learning algorithm for a particular MAI problem can be difficult. Recent advancements in DL have led to improved performance in tasks involving the detection of brain tumors, the classification of leukocytes, the detection of breast cancer, and other medical image PROCESSING tasks. These include GANs, the R-CNN quicker R-CNN, and the fusion of TML and DL models. High memory requirements and computational costs are, however, their main drawbacks. Therefore, highly computational AI algorithms must be discovered for leukocyte identification in blood test data. Moreover, these lightweight models are easily implementable on devices with limited resources. DNN Models can take the place of conventional methods. Since they can classify leucocytes into five groups and can simplify the model-building process, full models are also becoming more and more popular with the recent progress of CNNs [47]. These models compete with complex DNN-based models and depend on data-driven learning techniques. Various end-to-end architectures, including CNN-based models and attention-based methods [30], are also widely used for leucocyte detection in blood images. The research community mostly uses subjective evaluation methodologies in the field of image processing. But

this work is difficult, time-consuming, and error-prone at times. Therefore, more investigation is needed to examine general assessment methods that can automatically gauge how well ML models for processing images perform from various angles.

9 | Research Contribution

The current research included a thorough analysis of artificial intelligence methods for classifying leukocytes in blood smear images. We looked at various DM and DL methods for identifying WBCs in blood sample pictures. The information was gathered from primary research projects that were released between 2014 and 2024. 87 primary research (articles published in publications, texts, seminars, and online sources) outlining artificial intelligence approaches for leucocyte categorization in blood smear photos and their applications in medical diagnostics are found in the literature reviewed for this study. Both DM and DL (as AI techniques) techniques have done similarly regarding overall contributions in the processing of medical images, according to our evaluation of the studies. The collection of all of the data collected during the present investigation will aid the scientific community by determining where to begin with subsequent studies using DM and DL algorithms for the processing of medical images. In the future, these techniques will make significant contributions to the advancement of clinical imaging, natural language processing, and speech analysis. Aside from WBCs, DM and DL approaches are utilized to identify and classify several medical image processing domains, such as MRI, CT, X-ray, and ultrasound image analysis. In this study, we studied various DM and DL algorithms used to analyze blood smear images, including SVM ANNs, Ensembles, Bayesians, neuro-fuzzy, hybrids, DL, and CNNs [12, 42].

10 | Conclusion

This work aims to identify various uses of AI in the processing of medical scans and leucocyte classifications in blood sample scans. The goal of this work is to obtain insight into complicated aspects of AI by collecting and analyzing publications to help further studies in the discipline of the processing of medical images. This study demonstrates that much more research is needed to explore the use of AI approaches for meaningful clinical scan processing as well as leucocyte categorization in blood smear images. This study also sought to explore uses for advanced DL models outside blood cancer classification. However, it has been discovered that practically all other diagnostic applications are directly or indirectly tied to AI. In the field of medical imaging, smear of blood images is an emerging topic that has received significant interest from the research community over the last three decades. This article presents the typical contributions and potential uses of deep learning and machine learning in medical image processing. Furthermore, we identified existing obstacles, future goals, and solutions for the advancement of DL models in medical image processing, including WBC categorization in blood smear pictures. In the future, we plan to broaden our survey to include diverse processed medical image areas that include MRI, CT, Ultrasound, and X-ray pictures, using the potential of DM and DL approaches.

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Author Contribution

All authors contributed equally to this work.

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Data Availability

The datasets generated during and/or analyzed during the current study are not publicly available due to the privacy-preserving nature of the data but are available from the corresponding author upon reasonable request.

Conflicts of Interest

The authors declare that there is no conflict of interest in the research.

Ethical Approval

This article does not contain any studies with human participants or animals performed by any of the authors.

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