

CLASSIFICATION AND SEGMENTATION OF LEUKEMIA USING CONVOLUTION NEURAL NETWORK

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Abstract – *Leukemia implies blood disease which is highlighted by the uncontrolled and strange creation of white platelets (leukocytes) by the bone marrow in the blood. Examining minute platelet pictures, sicknesses can be distinguished and analyzed early. Hematologist are utilizing procedure of picture handling to examine, recognize and distinguish leukemia types in patients as of late. Recognition through pictures is quick and modest technique as there is no unique need of hardware for lab testing. We have zeroed in on the progressions in the calculation of cells and measurable boundaries like mean and standard deviation what isolates white platelets from other blood parts utilizing handling apparatuses like Jupiter Journal. Image enhancement, segmentation, and feature extraction are examples of image processing steps. A CNN on the improved and preprocessed dataset. The CNN should be made to precisely recognize typical and leukemic platelets. After the CNN has been trained, it should be evaluated using a separate dataset of images of blood cells. The model's precision ought to be estimated in the assessment. The CNN can be utilized to identify leukemia cells from dataset after it has been prepared and assessed.*

Key Words: Leukaemia, CNN, Image enhancement, segmentation and feature extraction

I. INTRODUCTION

The target of a leukemia location project utilizing CNN is to make a precise and dependable framework for distinguishing and grouping leukemic cells in blood tests. The project involves training a CNN to accurately classify blood cells from images of both normal and leukemic cells. A commonplace CNN design for this task could comprise of a few convolutional and pooling layers, trailed by completely associated layers and a delicate max yield layer. To find the architecture that works best for your dataset, you can play around with it. a CNN-based method for pinpointing leukemia in blood samples. Leukemia diagnosis and treatment can be aided by this system in numerous healthcare settings. A blood malignant growth results when strange white platelets (leukemia cells) collect in the bone marrow. Leukemia cells that are unable to mature properly are replaced by healthy cells that produce functional lymphocytes as ALL progresses rapidly. The leukemia cells are conveyed in the circulation system to different organs

and tissues, including the cerebrum, liver, lymph hubs and testicles, where they proceed to develop and partition. The developing, isolating and spreading of these leukemia cells might bring about various potential side effects. Everything is commonly connected with having more B lymphatic cells than Immune system microorganisms. The body is protected from germs and infections by B and T cells, which also actively destroy infected cells. B cells especially assist with keeping microorganisms from contaminating the body while Lymphocytes annihilate the tainted cells. Numerous programmed division and leukemia discovery has been proposed over these years. The majority of the methods used local image data as their foundation. By utilizing the HSV variety model a two-stage division is utilized. Much work has been there to meet the real clinical techniques because of the perplexing idea of leukemia cells. On segmentation and detection, similar studies have been published in the literature. It exclusively relies upon the robotized division and detection of leukemia. In this paper, we propose an automated method for examining a blood smear, which can help a doctor make better diagnoses and provide better treatment. Separating the other components of the blood from the leukocytes and extracting the lymphocytes is the method that is suggested in this paper. Fractal, shape, and other texture features are extracted from that extracted lymphocyte. For cell core limit unpleasantness estimation two new highlights were proposed for leukemia recognition. In view of separated highlights, the pictures are characterized into solid and leukemia by Help vector machine (CNN).

II. OBJECTIVE

The need of the leukemia discovery by utilizing picture handling is on the grounds that the above finding tedious and exorbitant. As it is hard to separate the leukemia cells from white platelets by involving clinical techniques as it changes as for time. By this method we can undoubtedly separate the phones from white platelets significantly quicker and best way just by thinking about the infinitesimal pictures. Several modalities, such as morphology, cell phenotyping, cytochemistry, cytogenetics, and molecular genetics, are necessary for the diagnosis of ALL. In spite of mechanical advances in medication, morphology stays the bleeding edge haematological demonstrative procedure. The perception of extreme leukemic cell development an morphological oddities in cell structures during the visual

assessment of fringe bloods Mears excites the primary doubt of leukemia.

III. EXISTING SYSTEM

Leukemia is a type of cancer that affects the blood and bone marrow, the spongy tissue between the bones that makes blood cells[1]. One of adult leukemia's most common forms is acute myeloid leukemia (AML)[2]. The signs and side effects of leukemia are vague in nature and furthermore they are tantamount to the side effects of other shared messes. Manual tiny investigation of stained blood smear or bone marrow suction is the best way to a viable determination of leukemia. K means calculation is utilized for division. Classification is done with KNN, NN, and SVM[3]. The spectral features are optimized with the help of GLCM. The nearby double example is utilized for surface depiction[4]. The classifier's performance on blood microscope images was evaluated[5].

3.1 Disadvantages

- ✓ Deep learning models can be hard to decipher, which can make it trying to comprehend how the framework is making its findings.
- ✓ When medical professionals are required to explain their diagnostic results to patients or other stakeholders, this may be cause for concern.
- ✓ Deep learning models require enormous datasets to really prepare. When there are small datasets or limited data access, this can be difficult.

IV. PROPOSED SYSTEM

The first step in developing a deep learning-based system for leukemia detection is to collect a dataset of blood cell images. This dataset should include both normal and leukemic blood cells, as well as images with varying degrees of complexity and variation. Once the dataset has been collected, the images need to be pre-processed to ensure that they are all the same size and resolution. Additionally, the images may be pre-processed to enhance features or remove noise. To increase the size of the dataset, data augmentation techniques may be used to generate additional images of blood cells with different orientations, scales, and rotations.

The next step is to train a CNN on the pre-processed and augmented dataset. The CNN should be designed to accurately classify blood cells as either normal or leukemic. Once the CNN has been trained, it should be evaluated using a separate dataset of blood cell images. The evaluation should measure the accuracy, of the model. Once the CNN has been trained and evaluated, it can be used for real-time detection of leukemia cells in blood samples. This may involve analysing blood Cells dataset, and using the CNN to provide diagnostic results in real-time. Overall, a proposed

system for leukemia detection using deep learning would involve collecting and preprocessing a dataset of blood cell images, training a CNN to classify blood cells as either normal or leukemic, evaluating the performance of the CNN, and using the CNN for real-time detection of leukemia cells in blood samples.

Table 1: Comparative Analysis with Related Works

Title & Year	Algorithm	Accuracy	Loss
Automated blast cell detection for Acute Lymphoblastic Leukemia diagnosis	YOLOv4	80.0%	8.0%
Feature Extraction of White Blood Cells Using CMYK - Moment localization and Deep Learning in Acute Myeloid Leukemia Blood Smear Microscopic Image	XG Boost	82%	4%
A COMPARATIVE STUDY OF CLASSIFICATION AND SEGMENTATION OF LEUKEMIA USING CNN	CNN	94%	3%

4.1 Advantages of Proposed System

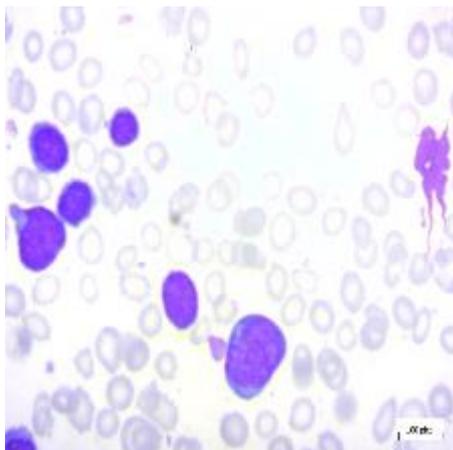
- ✓ Profound learning models, especially CNNs, are appropriate for picture based errands like leukemia recognition analyse.
- ✓ A profound learning-based framework for leukemia recognition can break down blood tests progressively, taking into consideration quicker determination and treatment.
- ✓ Via computerizing the course of leukemia location, a profound learning-based framework can decrease the expense of determination.

V. RELATED WORK

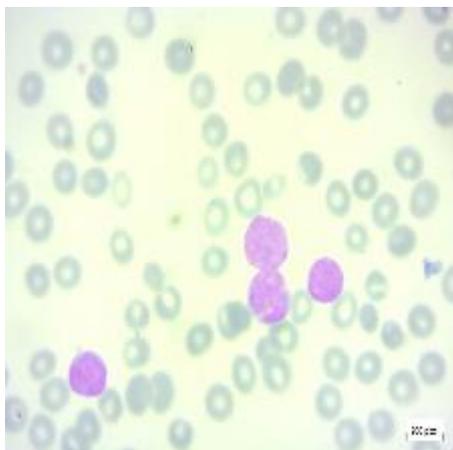
5.1 DATA PREPARATION

We have utilized the Intense Lymphoblastic Leukemia Picture dataset. Picture utilized in this venture were gotten from frontiers dataset which is a public dataset accessible

on the web. This dataset comprised of 3256 fringe blood smear pictures from patient, whose blood tests were ready and stained. These pictures had goal of 320*240. The picture documents gathered are in JPG or PNG picture design. The size of the preparation dataset will be pivotal in light of the fact that one of the attributes of profound learning is its capacity to perform well when prepared with huge informational indexes hence large number of pictures are expected to prepare CNN successfully. We have utilized the Intense Lymphoblastic Leukemia Dataset to prepare and approve our CNN model.



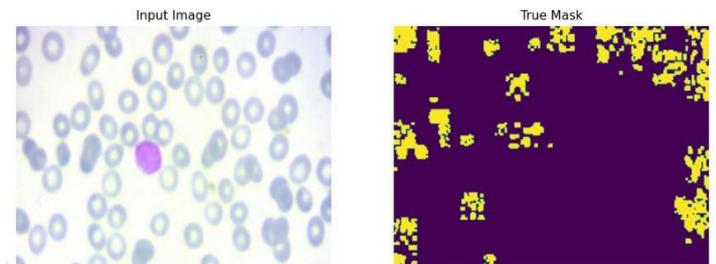
5.1.1 Image dataset1



5.1.2 Image dataset2

5.2 IMAGE PRE-PROCESSING

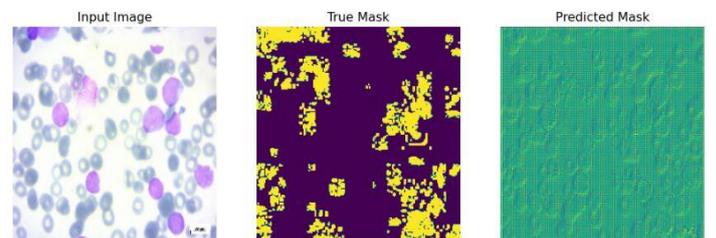
The acquired images are pre-processed to get rid of unnecessary noise and small substances that can't be measured, like blood cells. For better division consequences of the platelets, the acquired picture must be improved. A few image processing techniques, such as contrast adjustment, grey-scale detection, edge detection, and spatial smoothing filtering, are used to accomplish this.



5.2.1 IMAGE PRE-PROCESSING

5.3 IMAGE SEGMENTATION

Choosing the part of the image that interests you is the first step in this process. The region that is made up of the blood cells is outlined in this. Round Hough change is applied and not a large part of the picture division is required in light of the fact that the applied change looks just for the round objects in the picture. 5) Identifying Blood Cells: After looking for the blood cells in the image, the Circular Hough transform finds them. The capability "draw circle" draws circles around the distinguished cells. Even the circles that overlap are found. 6) Including Cells: Counting the quantity of cells drawn gives the complete number of platelets in the picture.

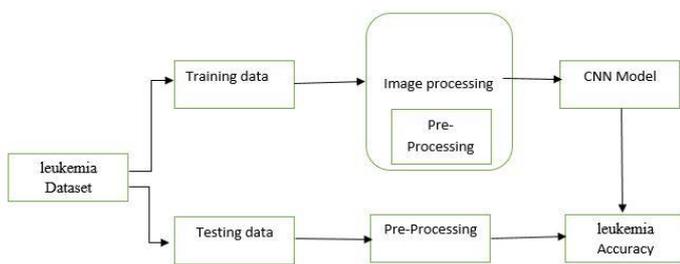


5.3.1 IMAGE SEGMENTATION

5.4 IMAGE CLASSIFICATION

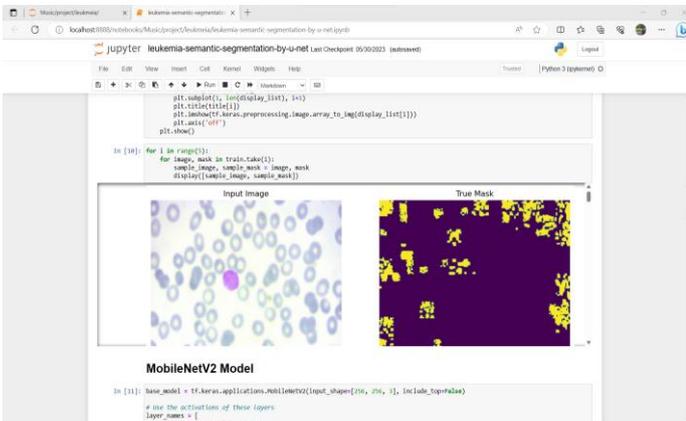
The essential aim is in this part is to recognize AML and CML utilizing geological element extraction and variety-based highlight extraction. In this last stage, the highlights extricated are utilized to give the last response. Every one of the highlights extricated are recorded into the various sections with their qualities. The proposed system calculates the feature values after receiving the classified WBC images as input. The upsides of the test picture highlights are checked with the boundaries carried out. Abnormalities are being observed using three parameters. Entire cell size (region), Core size, a fury of Core and cytoplasm

VI. SYSTEM ARCHITECTURE



6.1 System Architecture

VIII. SCREENSHOTS



8.1 image upload



8.2 Cell Detection

IX. CONCLUSION

Due to present day way of life, contamination and different elements malignant growth is turning out to be a greater amount of normal illness. Since cancer is a rapidly spreading disease, conventional methods of cancer detection take time because the transport of sample tissue (biopsy) to a cancer diagnosis facility takes time. Early treatment will likely increase a patient's chances of survival. The framework will be worked by involving highlights in minuscule pictures by

looking at changes on surface, math, colors and measurable examination as a classifier input. The system should work well, be dependable, take less time to process, make fewer mistakes, and be accurate.

REFERENCES

- [1] C.R., Valencio, M.N., Tronco, A.C.B., Domingos, C.R.B., "Knowledge Extraction Using Visualization of Hemoglobin Parameters to Identify Thalassemia", Proceedings of the 17th IEEE Symposium on Computer Based Medical Systems, 2004, pp. 1-6.
- [2] R., Adollah, M.Y., Mashor, N.F.M, Nasir, H., Rosline, H., Mahsin, H., Adilah, "Blood Cell Image Segmentation: A Review", Biomed 2008, Proceedings 21, 2008, pp. 141-144.
- [3] N., Ritter, J., Cooper, "Segmentation and Border Identification of Cells in Images of Peripheral Blood Smear Slides", 30th Australasian Computer Science Conference, Conference in Research and Practice in Information Technology, Vol. 62, 2007, pp. 161-169.
- [4] D.M.U., Sabino, L.D.F., Costa, L.D.F., E.G., Rizzatti, M.A., Zago, "A Texture Approach to Leukocyte Recognition", Real Time Imaging, Vol. 10, 2004, pp. 205-206.
- [5] M.C., Colunga, O.S., Siordia, S.J., Maybank, "Leukocyte Recognition Using EM Algorithm", MICAI 2009, LNAI 5845, Springer Verlag Berlin Heidelberg, 2009, pp. 545-555.
- [6] K.S., Srinivisan, D., Lakshmi, H., Ranganathan, N., Gunasekaran, "Non Invasive Estimation of Hemoglobin in Blood Using Color Analysis", 1st International Conference on Industrial and Information System, ICIIS 2006, Sri Lanka, 8 - 11 August 2006, pp 547-549.
- [7] W., Shitong, W., Min, "A new Detection Algorithm (NDA) Based on Fuzzy Cellular Neural Networks for White Blood Cell Detection", IEEE Transactions on Information Technology in Biomedicine, Vol. 10, No. 1, January 2006, pp. 5-10.
- [8] Pham, T., Tran, T., Phung, D., & Venkatesh, S. (2017). Predicting healthcare trajectories from medical records: A deep learning approach. Journal of Biomedical Informatics, 69, 218-229. Doi: 10.1016/j.jbi.2017.04.001.
- [9] Kourou, K., Exarchos, T. P., Exarchos, K. P., Karamouzis, M. V., & Fotiadis, D. I. (2014). Machine learning applications in cancer prognosis and prediction. Computational and structural biotechnology journal, 13, 8-17. <https://doi.org/10.1016/j.csbj.2014.11.005>.
- [10] Jiang, F., Jiang, Y., Zhi, H., Dong, Y., Li, H., Ma, S., Wang, Y., Dong, Q., Shen, H., & Wang, Y. (2017). Artificial intelligence in healthcare: past, present and future. Stroke and vascular neurology, 2(4), 230-243. <https://doi.org/10.1136/svn-2017-000101>.

[10] Sarwar, S., Dent, A., Faust, K., Richer, M., Djuric, U., Ommeren, R.V., & Diamandis, P. (2019). Physician perspectives on the integration of artificial intelligence into diagnostic pathology. *npj Digital Medicine*.

[11] Through the Microscope: Blood Cells - Life's Blood.<http://www.wadsworth.org/chemheme/heme> [3 October 2011].

[12] T., Bergen, D., Steckhan, T., Wittenberg, T., Zerfab, "Segmentation of leukocytes and erythrocytes in Blood Smear Images", 30th Annual International IEEE EMBS Conference, Vancouver, Canada, August 20 - 24, 2008, pp. 3075-3078.

[13] S. Osowski, R., Siroic, T., Markiewicz, K.Siwiek, "Application of Support Vector Machine and Genetic Algorithm for Improved Blood Cell Recognition", IEEE Transactions on Instrumentation and Measurement, Vol. 58, No. 7, July 2009, pp. 2159-2168.

[14] Y.M., Hirimitugoda, G., Wijayarathna, "Artificial Intelligence-Based Approach for Determination of Haematalogic Diseases", IEEE, 2009.

[15] S., Mohapatra, D., Patra, S., Satpathi, "Image Analysis of Blood Microscopic Images for Leukemia Detection", International Conference on Industrial Electronics, Control and Robotics, IEEE, 2010, pp. 215-219.

[16] N., H., A., Halim, M., Y., Mashor, R., Hassan, "Automatic Blasts Counting for Acute Leukemia Based on Blood Samples", International Journal of Research and Reviews in Computer Science, Vol. 2, No. 4, August 2011, pp. 971-976.