

Self Monitoring of White Blood Cells Cancer Diseases

Prof. A. A. Utikar
Department of Information
Technology
Sinhgad College of
Engineering
Pune, India

Hitesh Patil
Department of Information
Technology
Sinhgad College of
Engineering
Pune, India

Rahul Shah
Department of Information
Technology

Sinhgad College of Engineering
Pune, India

Tushar More
Department of Information
Technology

Sinhgad College of Engineering
Pune, India

Abstract— Self Monitoring analysis of white blood cell cancer disease such as Leukemia and Myeloma is a hard biomedical learn subject. Our go in the direction of present planned for the main time a novel state of the picture ask for so as help in identify the pallid blood cell disease. We divide these disease eager on two categories; every collection include comparable indication disease so as to might puzzle inside make a analysis. base on top of the doctor's diversity, single of two approach be apply. every shift toward be sensible in the way of single of the two disease category from side to side compute unlike skin. eventually, a accidental Forest classifier be obliging intended for the last choice. The prospect move in the direction of aim in the direction of early on discovery of white blood cells cancer, decrease the misdiagnosis luggage inside addition in the way of humanizing the scheme information method. also, allow the expert just in the direction of contain the last change on the result obtain as of the system.

Keywords— White blood cells disorder, categorization, bottomless knowledge, Random Forest.

I. INTRODUCTION

A. Purpose

Now a day's self-monitoring of white blood cells cancer detection it is a demanding mission in the biomedical investigate topic. There is a require to learn and make a scheme which will create it easy for end users to get the discovery without homicide their time. Maintaining the honesty of the stipulation

B. Biological Facts

In the resistant system, white blood cell is most significant part bodies. White blood cells are part of the body's secluded system. It helps the body brawl infection and other diseases. Types of white blood cells are following they are: granulocytes (neutrophils, eosinophils, and basophils), monocytes, and lymphocytes (T cells and B cells).

test the number of white blood cells in the blood is characteristically part of a complete blood cell (CBC) test. It may be used to look for circumstances such as disease, zirritation, allergies, and leukemia.

C. White Blood Cell Count in Image Processing

Image dispensation is concerning with digital imagery to extract helpful information. It is concerned in dissimilar other

topics such as layout-land use, nature credit, coin credit, Medical Imaging. Medical imaging uses the in order extracted from the digital image to improve the analytic of dissimilar diseases. The two usually used categories of white blood cell disorders divide them quantitatively into those causing extreme numbers and those causing inadequate numbers. Leukocytosis is generally fit, but it also might be dysfunctional proliferative. WBC proliferative disorders can be classed as myeloproliferative and lymph proliferative.

Some are autoimmune, but numerous are neoplastic. one more way to categorize disorders of white blood cells is qualitative. There are a lot of disorders in which the numbers of white blood cells are normal but the cells do not purpose usually. Neoplasia of WBCs can be type but is often evil. Of the diversity of tumors of the blood and lymph, cancers of WBCs can be generally classified as leukemia and lymphomas, although those categories partly cover and are often grouped as a couple.

II. RELATED WORK

Some researches were proposed to differentiate between:

A. AML and ALL disease

Acute Myeloid Leukemia is a kind of blood cancer disease. one more name for the Acute Myeloid Leukemia is Acute Myelogenous Leukemia. AML is the most usual acute leukemia disturbing adults. In the fillet marrow is present in AML is a type of cancer that begins the phase. The possibility of getting AML increases with era. But a being can get AML at any era. About 8 in 10 adults with acute leukemia have been detected an AML.

Karthikeyan and Poornima future an move toward for discovery of Leukemia in blood at previous time stages. They have second-hand an adaptive median filter for noise removal and adaptive Histogram Equalization for difference improvement in the pre-processing phase. They useful k-means and Fuzzy c-means clustering for segmentation. They computed statistical, textural and geometrical skin and practical Support Vector Machine (SVM) for categorization. Their move toward achieved 90% with Fuzzy c-means and 83% with k-means using Fuzzy Logic: cleverness, manage, and in order dataset.

one more research by Mohapatra et al. planned a Fuzzy-based Blood Image Segmentation model for the Automated Detection of Leukemia. They sensible selective middle filtering followed by unsnapping masking in pre-processing. In segmentation, they second-hand an enhanced account of fuzzy clustering method cluster followed by adjacent neighbor classification in $L^*a^*b^*$ color space like (L^* for lightness, a^* for redness greenness axis, and b^* a yellowness blueness axis).

The computed description is two novel shape features: Hausdorff Dimension and contour mark. They achieved 93% on a database of 108 blood smear images of size 512 x 512 pixels Support Vector Machine (SVM) is working for categorization and they achieved 93% on a database of 108 blood smear images of size 512 x 512 pixels.

B. AML Detection only or ALL Detection only

An come up to by Agaian et al. proposed an easy method that automatically detects and segments AML in blood smears. Segmentation was done in the CIELAB Colorspace by K-Means clustering algorithm. Hausdorff measurement kind was computed using the box-counting method and Local Binary Pattern (LBP). categorization achieved maximum correctness using the Support Vector Machine (SVM) on American Society of Hematology (ASH) for Leukemia dataset.

In this dataset comprised of 100 images a half from AML patients and a half from non-AML patients. The image size used for their categorization was 184 x 138 pixels.

an additional system proposed by Bhattacharjee and Saini for Acute Lymphoblastic Leukemia uncovering using Watershed Transformation method. They practical contrast improvement and excellence adjustment for enhancing images before segmentation. In segmentation they used a watershed algorithm, isolating the blood cell and the cell nucleus. They computed area, perimeter, circularity and form factor kind. Gaussian Mixture Models (GMM) and Binary Search Tree (BST) were applied for categorization. On one hand, GMM had achieved 93% while BSTe had achieved 86%.

C. Detection of Acute Myeloid Leukemia

Another projected approach, by Sarrafzadeh et al. focused mainly on differentiating between M2, M3 and M5 sub-types to evaluate their introduced method. The approach was practical in the formulated $L^*a^*b^*$ color space. Segmentation is performed using K-means clustering to separate leukocytes from other blood components. surface and shape features are extracted in arrange to be classified using Discriminative Dictionary Learning (DDL).

They achieved 97.53% correctness on Medical Image & Signal Processing Research Center (MISP) dataset. They used a dataset composed of 27 microscopic images of three sub-types of AML; 9 AML-M2, 10 AML-M3 and 8 AML-M5. Feature Extraction development is a procedure to obtain the description or parameters that will be used to notice ALL, AML M3 or Healthy Cells. A feature to be searched is the WBC area, nucleus ratio and granule ratio. WBC area is the area of WBC; the nucleus ratio is the ratio between the areas of the cell nucleus with the area of WBC.

III. MOTIVATION

Following had been source of motivation for this paper:

- in our time self-monitoring of white blood cells cancer detection it is a demanding task in the biomedical research topic.
- The discovery and classification of white blood is a hot issue since of its significant applications in disease diagnosis.

IV. SYSTEM ARCHITECTURE

The given figure shows the thorough flow of Self-Monitoring of White Blood Cells Cancer Diseases.

In this format user upload image as an input. Using CNN Classifier, known image file will be processed. Random forest algorithm is used to recognize the exact grouping of the mechanism to legalize the disease. A Random forest algorithm helpful for identifying the disease by analyzing the patient's medical records WBC cells. This system is used to decide and predict an automated system to correctly detect white blood cell cancer diseases for the user.

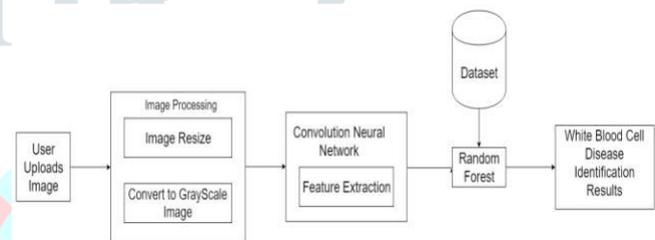


Fig 1: System Overview

In this project, CNN classifier (Convolutional Neural Network) algorithm is used to extract the features from the images and Random Forest algorithm used to identify the correct combination of the components to validate the disease.

V. PROPOSED APPROACH

This piece explains types of algorithms used for detection of diseases.

A. Convolutional Neural Network (CNN)

Convolutional Neural Networks are the type of bottomless Artificial Neural Networks that are used primarily for the purpose of classifying images (for e.g. name what they see), cluster them by the resemblance (like photo search), and perform object recognition within scenes. These are the algorithms that can recognize face details of an individual, their characteristics, street signs, tumors, platypuses and other essential aspects of visual data.

The layers involved in Convolutional Neural Network are:

1. Convolution Layer- In this layer we apply filter (3*3) on image.

2. Pooling layer- Pooling is an important part of Convolutional neural networks for object discovery.
3. Fully Connected layer- It is the fully connected layer of neurons at the end of CNN.

A CNN consists of an input layer and an output layer, as well as other manifold hidden layers. The hidden layers of a CNN typically consist of several convolutional layers, pooling layers, fully connected layers and the normalization layers. Convolutional layers be relevant a convolution operation to the input layer, passing the result to the next layer.

B. Random Forest

Random forest algorithm is a type of supervised classification algorithm.

In general, the extra number of trees in the forest, the more robust forest looks like. In the same way in the Random Forest Classifier, the advanced the number of trees in the forest, it gives the high accuracy fallout. Random forest algorithm is used to identify the correct grouping of the components to validate the disease. Random forest algorithms also helpful for identifying the disease by analyzing the patient's medical records of especially WBC cells.

Random forest algorithm achieved has successes in medical field as it is one of the most powerful algorithms which is being widely used in different applications.

VI. CONCLUSION

inside this paper, we propose the design, development and evaluation of an automated system to accurately detect white blood cells cancer diseases. It detects the types and sub-types of Leukaemia (ALL and AML) and Myeloma. It is a recognition system applied to acquired blood microscopic images then performs preprocessing, segmentation, feature extraction and classification. The proposed solution converts images to YCBCR color space and constructs the Gaussian distribution of 53 CB and CR values. Statistical, texture, size ratio and morphological features are then computed to train the classifier. Unlike existing systems, our system has the

ability to learn from misclassified tests to enhance the future accuracy of the system. Random Forest classifier is the best classifier that is able to differentiate between different types and the one which gives us the best accuracy. The system achieved 94.3 % accuracy in detecting and classifying types and sub types.

ACKNOWLEDGMENT

We would like to thank our Project Guide, Prof. A. A. Utikar for providing us with all guidance and help needed and providing great support in project activities.

REFERENCES

- [1] Partha Pratim Banik, Rappy Saha, Ki-Doo Kim "Fused convolutional neural network for white blood cell image classification", 978-1-5386-7822-0/19, 2019 IEEE.
- [2] C. Reta, L. Altamirano, J. A. Gonzalez, R. Diaz-Hernandez, H. Peregrina, I. Olmos, J. E. Alonso, R. Lobato, "Segmentation and classification of bone marrow cells images using contextual information for medical diagnosis of acute leukemias", PLOS ONE | DOI:10.1371/journal.pone.0130805, 2015.
- [3] Dr.T.Karthikeyan, N.Poornima "Microscopic image segmentation using fuzzy c means for leukemia diagnosis", ISSN: 2350-0328, Vol. 4, Issue 1 , January 2017.
- [4] Hend Mohamed, Rowan Omar, Nermeen Saeed, Ali Essam, Nada Ayman, Taraggy Mohiy and Ashraf AbdelRaouf, "Automated detection of white blood cells cancer diseases", 978-1-5386-5083-7/18, 2018 IEEE.
- [5] O. Sarrafzade, H. Rabbani, A. M. Dehnavi, A. Talebi, "Detecting different sub-types of acute myelogenous leukemia using dictionary learning and sparse representation", 978-1-4799-8339-1/15, 2015 IEEE.
- [6] S. Sameh, M. A. Azim, A. AbdelRaouf, "Narrowed coronary artery detection and classification using angiographic scans", 978-1-5386-1191-3/17, 2017 IEEE.
- [7] Athelas Blog: <https://blog.athelas.com/classifying-white-blood-cells-with-convolutional-neural-networks-2ca6da239331>
- [8] WBC Symptoms and Normal Counts : <https://www.healthline.com/health/wbc-count#results>
- [9] Kaggle Practice Data: <https://kaggle.com/paultimothymooney/blood-cell>