

ACUTE LYMPHOBLASTIC LEUKEMIA DIAGNOSIS IN BLOOD MICROSCOPIC IMAGES USING LOCAL BINARY PATTERN AND SUPERVISED CLASSIFIER

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Acute lymphoblastic leukemia (ALL) is a subtype of acute leukemia, that is well-known among adults. The average age of someone with ALL is sixty five years. The want for automation of leukemia detection arises when you consider that modern-day methods contain manual exam of the blood smear as the first step towards prognosis. that is time-ingesting, and its accuracy relies upon at the operator's capacity. on this paper, an easy method that mechanically detects and segments ALL in blood smears is provided. The proposed method differs from others in: 1) the simplicity of the advanced technique; 2) type of entire blood smear images in place of sub images; and three) use of those algorithms to section and locate nucleated cells. pc simulation concerned the following checks: comparing the effect of Hausdorff dimension on the device earlier than and after the impact of neighborhood binary pattern, evaluating the overall performance of the proposed algorithms on sub snap shots and entire pics, and comparing the outcomes of a number of the existing structures with the proposed machine. eighty microscopic blood photographs were tested, and the proposed framework controlled to gain 98% accuracy for the localization of the lymphoblast cells and to separate it from the sub pics and whole pictures.

Keywords-Image Acquisition, Color Features and Color Correlation, Nuclei Segmentation.

I.INTRODUCTION

Diagnosing leukemia is primarily based on the reality that white cell count is accelerated with immature blast cells (lymphoid or myeloid), and neutrophils and platelets are reduced. Acute lymphoblastic leukemia (ALL) is a heterogeneous clonal disorder of haemopoietic progenitor cells ("blasts"), which lose the capacity to differentiate typically and to reply to ordinary regulators of proliferation.

This loss results in deadly infection, bleeding, or organ infiltration, usually, within the absence of remedy, within a year of prognosis. ALL is showed while the marrow consists of more than 30% blasts. In this paper, most effective ALL is considered. ALL is a fast-growing cancer of the blood and bone marrow. It's far fatal if left untreated, because of its rapid unfold into the bloodstream and other vital organs. moreover ALL additionally makes up 15–20% of early life leukemia, kind of 60% of instances occur in people elderly younger than two decades. Survival in early life acute lymphoblastic leukemia is drawing close 90%, but remedy in infants (i.e., kids more youthful than 365 days) and adults desires improvement. Early prognosis of the sickness is essential for the healing of patients, specifically within the case of kids. In this paper, only acute lymphocytic leukemia is considered. Fig. 1 suggests six different pics, three depicting wholesome cells from non-ALL patients and 3 from ALL patients. This method greatly relies upon on the operator's capability and fatigue stages. Diagnostic confusion additionally takes place because of imitation of comparable signs by way of different problems. Furthermore, the identification task is generally difficult due to the form of capabilities and the often doubtful pictures purpose missing out on important indicators as to which shape of leukemia is being determined. Furthermore, due to the complicated nature of blood smear pix and variation in slide preparation strategies, plenty paintings has to be carried out to fulfill actual medical needs. consequently, these factors can lead to wrong diagnosis. in the beyond, digital picture processing techniques have helped to investigate the cells that cause more accurate, fashionable, and faraway disease prognosis systems. however, there are some complications in extracting the records from WBCs due to wide version of cells in form, size, part, and role. moreover,

seeing that illumination is imbalance, the picture contrast between cell boundaries and the history varies depending at the condition for the duration of the taking pictures technique.

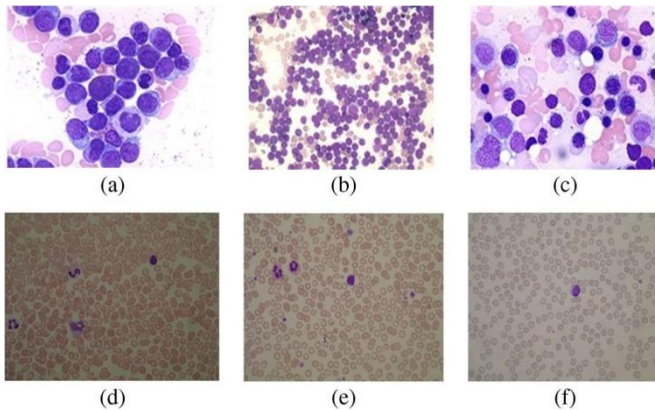


Fig 1 Images from ASH. (a)–(c) Lymphoblasts from ALL patients. (d)–(f) Healthy cells from non-ALL patients.

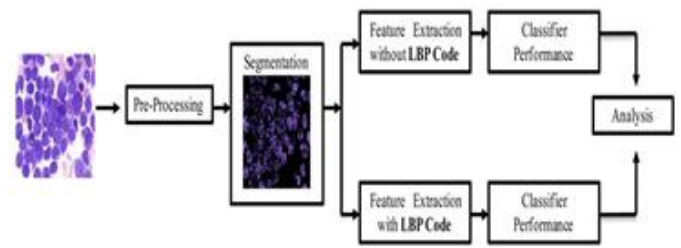
II PROCESS OVERVIEW

The gadget proposed guarantees step-by way of-step processing. Fig. 2 depicts the gadget review. The device review gives a detailed depiction of the series of steps which might be to be followed for green class of acute leukemia. step one entails preprocessing the entire pictures to conquer any heritage nonuniformity because of irregular illumination. Preprocessing also includes shade correlation wherein RGB snap shots are converted to L^*a^*b colour area images. This step ensures perceptual uniformity. This step is observed by okay-approach clustering to carry out the nucleus of every cell. Segmentation is followed by function extraction primarily based on which classification and validation are carried out.

PREPROCESSING

Photo Acquisition: For ALL, we accessed the american Society of Hematology (ASH) for their online picture bank of leukemia cells. The ASH picture bank is a web-based totally photograph library that offers complete and growing collections of snap shots regarding a wide range of hematology classes. They offer terrific images captured the use of one-of-a-kind microscopes in exclusive resolutions. Our database for ALL comprised 80 photographs—forty from ALL sufferers and forty from non-ALL patients. The decision used for our type became 184×138 pixels. The pre processing includes the image resizing, Graycolor space conversion and de-noising. Before the segmentation process, we are going to

resize the input image into fixed Row and Column size [256 256] using imresize command.



Then the input image is converted into gray scale conversion for accurate segmentation. The conversion is done by RGB2GRAY Matlab commands to reduce the effect of noise we apply the median filter for de-noising.

NUCLEI SEGMENTATION

The aim of image segmentation is to extract vital records from an enter photograph. It performs a key function since the performance of subsequent function extraction and type is predicated significantly on the precise identification of the lymphoblasts. Many algorithms for segmentation were evolved for gray-degree photographs. Segmentation on this device is executed for extracting the nuclei of the leukocytes the usage of colour-primarily based clustering. k-method remains extensively used. This speaks to the issue in designing a general-purpose clustering set of rules and the sick-posed hassle of clustering. In this paper, we chose clusters similar to nucleus (high saturation), history (high luminance and occasional saturation), and other cells (e.g. erythrocytes and leukocyte cytoplasm). here, every pixel is assigned to any such training using the properties of the cluster center.

K-MEANS CLUSTERING ALGORITHM

The k-approach set of rules calls for three consumer-distinctive parameters: the quantity of clusters okay, cluster initialization, and distance metric. Each pixel within the L^*a^*b color area is classed into any of the okay clusters through calculating the Euclidean distance among the pixel and each colour indicator. these clusters correspond to nucleus (high saturation), background (excessive luminance and low saturation), and other cells (e. g., erythrocytes and leukocyte cytoplasm) .Each pixel of the whole picture will be labeled to a selected shade depending on the minimum distance from every indicator. We take into account handiest the cluster that

carries the blue nucleus, that's required for the characteristic extraction. While performing k means segmentation of whole pix, it became discovered that, in some of the segmented pix, best the edges of the nuclei had been received as opposed to the entire pix of the nuclei. This shortcoming turned into triumph over by using employing morphological filtering..An photograph is partitioned into numerous areas relying on the features to be extracted. employing morphological filtering ensures that perceptibility

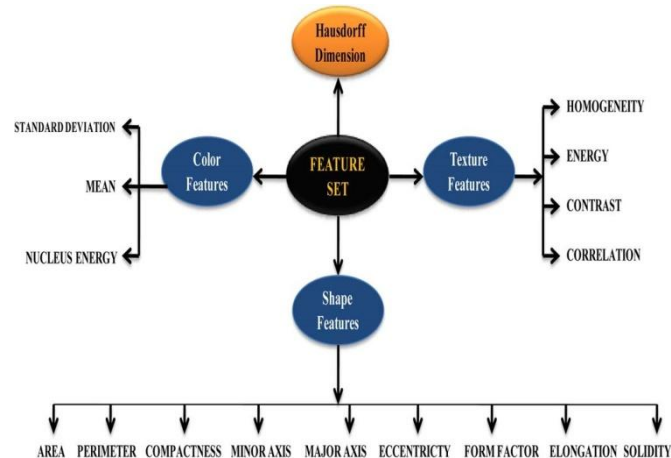


Figure 4 Feature set developed for the proposed system comprising of shape, color, texture features, and HD.

FEATURE EXTRACTION

Feature extraction in image processing is a technique of redefining a large set of redundant data into a set of features of reduced dimension. Transforming the input data into the set of features is called feature extraction. Feature selection greatly influences the classifier performance; therefore, a correct choice of features is a very crucial step. In order to construct an effective feature set, several published articles were studied, and their feature selection methodology was observed. It was noted that certain features were widely used as they gave a good classification. We implemented these features on whole images in our system. Those features were considered to boost the classifier performance. Fig. 4 gives the set of features chosen to classify the image database.

1.HD:The most important theoretical fractal dimensions are the Rényi dimension, the HD, and the packing dimension. Practically, the box-counting dimension is widely used, partly due to their ease of implementation. In a box counting algorithm, the number of boxes covering the point set is a power-law function of the box size. Fractal dimension is

estimated as the exponent of such power

2.law: All fractal dimensions are real numbers that characterize the fractalness (texture/roughness) of the objects. Lymphoblast can be differentiated using perimeter roughness of the nucleus. HD is considered an essential feature considered in our proposed system.

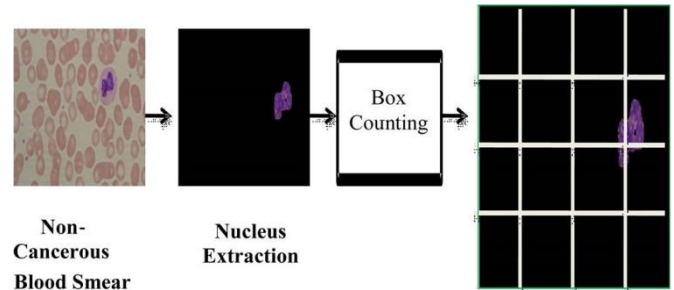


Figure 5 Superimposing of the nucleus with a grid of squares for box count measure.

The procedure for HD measurement using the box counting method is elaborated below as an algorithm:

- 1) Binary image is obtained from the gray-level image of the blood sample;
- 2) Edge detection technique is employed to trace out the nucleus boundaries;
- 3)Edges are superimposed by a grid of squares;
- 4) the HD may then be defined as follows:

$$HD = \frac{\log(R)}{\log(R(s))} \quad (1)$$

where R is the number of squares in the superimposed grid, and R(s) is the number of occupied squares or boxes (box count). Higher HD signifies higher degree of roughness.Fig. 5 illustrates the given algorithm. It shows how the nucleus from a non-cancer cell is superimposed with a grid of squares to perform suitable box counting. The finer the grid gets, the more accurate is the shape approximated.

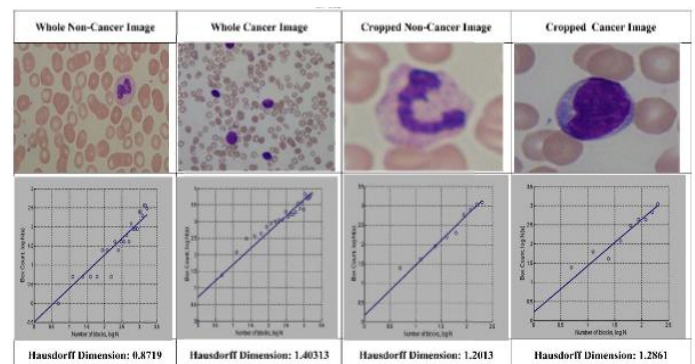


Figure 6 Results of HD


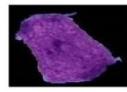
LBP: The concept of local binary pattern (LBP) was introduced for texture classification . This approach has many advantages. For example, the LBP texture features have the following characteristics:

- 1) They are robust against illumination changes;
- 2) they are very fast to compute features
- 3) they do not require many parameters to be set;
- 4) they are local grayscale transformations and scaling;
- 5) they are invariant with respect to monotonic
- 6) they have performed very well in many computer vision image retrieval applications.

The LBP method has proved to outperform many existing methods, including the linear discriminant analysis and the principal component analysis. In order to deal with textures at different scales, the LBP operator was later extended to use neighborhoods of different sizes, the result will be a binary pattern with length M. This operation is illustrated in Fig. 6. For our database of images, an (8, 1) circular neighborhood was used. The segmented images were extracted using k-means clustering, and then the LBP operator was applied on them before calculating the HD (see Fig. 7). Two sets of values were

(GLCM) and color features.

SHAPE FEATURES WITH THEIR ILLUSTRATIONS

Features	Cancerous	Normal
Images		
1.Mean	32.3699	37.1222
2.Standard Deviation	47.3662	41.0178
3.Area: The total number of none zero pixels within the image region.	6453	1985
4. Perimeter: Calculating distance between successive boundary pixels gives the perimeter.	256	807.293
5.Elongation: The nucleus bulging is measured in terms of a ratio called elongation. This is defined as the ratio between the maximum distance (<i>Rmax</i>) and the minimum distance (<i>Rmin</i>) from the center of gravity of the nucleus boundary.	1.1357	1.522
6. Eccentricity: Since lymphocytes are more circular than the myeloblasts, eccentricity is an important feature. It is a parameter that is used to measure how much the shape of a nucleus deviates from being circular.	0.4740	0.7541
7. Form-factor: A Dimensionless parameter considered which changes with surface irregularities.	1.2373	0.0383
8. Solidity: An essential feature for blast classification equals the ratio of actual area over the convex hull area	0.5679	0.2117
9.Compactness/roundedness: is the measure of a nucleus	10.1559	328.32

Shape features. One of the shape features that has proven to be a good measure for classifying ALL by their shape is compactness. The shape of the nucleus, according to haematologists, is an essential feature for discrimination of lymphoblasts. extracted: first, HD of the 80 images without applying LBP, and second, HD of the images after applying LBP. When comparing these two data sets, it was observed that the LBP operator enhanced the overall performance by a very high margin. GLCM features . Texture is defined as a function of the spatial variation in pixel intensities. The GLCM and associated texture feature calculations are image analysis techniques.

Textual characteristics. Some of these features are the following.

- 1) Energy:** Also known as uniformity (or angular second moment), it is a measure of homogeneity of image.

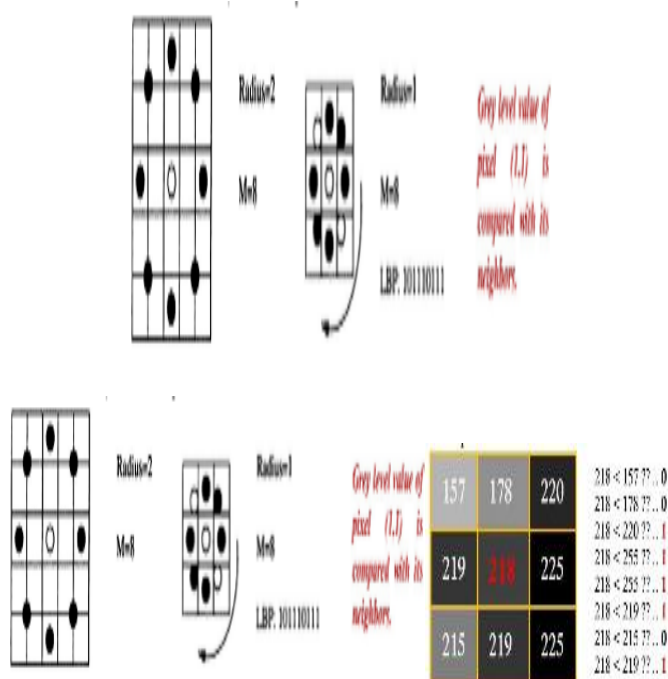


Figure 7 LBP texture classification

Additionally, the following features have been also chosen in our classification system: shape gray-level cooccurrencematrix

2) Contrast: The contrast feature is a difference moment of the regional cooccurrence matrix and is a measure of the contrast or the amount of local variations present in an image.

3) Entropy: This parameter measures the disorder of an image. When the image is not texturally uniform, entropy is very large.

4) Correlation: The correlation feature is a measure of regional-pattern linear dependence in the image.

Color features. In addition to the features aforementioned, we have used the following color-based feature.

5) Cell Energy: Also known as the measure of uniformity, it is the different Lab image components. We define feature “ δ ” to be

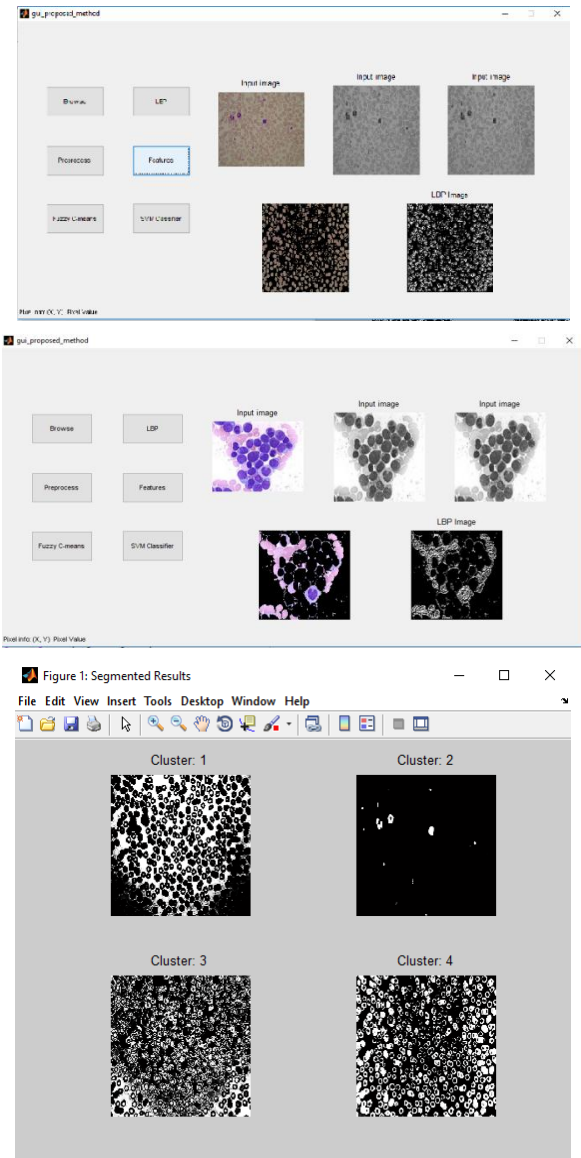
$$\delta = \sum_i \sum_j P^2(i, j) + (\sqrt{-1}) \left(\frac{\sqrt{\sum_{i=1}^n (x_i - x')^2}}{n-1} \right) \quad (2)$$

where $x' = \sum_{i=1}^n x_i / n$, $P(i, j)$ represents the normalized GLCM element for the i th row and j th column, and $\sum_i \sum_j P^2(i, j)$ represents the ASM.

III. COMPUTER SIMULATION

The selection of a classification method for type is a challenging hassle due to the fact an appropriate preference given the to be had records can drastically assist improving the accuracy in credit score scoring practice. there's a masses of statistical strategies, which intention at solving binary class duties. in this paper, we use a aid vector system (SVM) for constructing a selection surface in the function space that bisects the 2 classes, i.e., cancerous and noncancerous, and maximizes the margin of separation among training of factors. SVMs is a promising nonlinear nonparametric classification approach, which already confirmed properly outcomes within the scientific diagnostics, optical person recognition, electric load forecasting, and other fields. furthermore, the SVM is a powerful stateof-the-artwork algorithm with strong theoretical foundations based at the Vapnik–Chervonenkis theory and with strong regularization homes. Regularization refers to the generalization of the model to new statistics. plenty of the preliminary fulfillment of SVMs was attributed to the so-known as kernel trick wherein schooling records are implicitly mapped to a

excessive-dimensional function area, and a margin maximizing linear classifier is found out on this mapped area



IV DATASET ANALYSIS

For each image in the dataset, the classification/position of ALL lymphoblasts is provided by expert oncologists. Furthermore, we suggest a specific set of figure of merits to be processed in order to fairly compare different algorithms with the proposed dataset. The images of the dataset has been captured with an optical laboratory microscope coupled with a Canon PowerShot G5 camera. All images are in JPG format with 24 bit color depth, resolution 2592 x 1944. The ALL_IDB1 version 1.0 can be used both for testing segmentation capability of algorithms, as well as the classification systems and image preprocessing methods. This dataset is composed of 108 images collected during

September, 2005. It contains about 39000 blood elements, where the lymphocytes has been labeled by expert oncologists. The images are taken with different magnifications of the microscope ranging from 300 to 500.

V CONCLUSION

In this paper, we've proposed a colour-primarily based clustering for segmenting white blood cells from Acute Lymphoblastic Leukemia images. The technique efficiently combines RGB and Lab shade space based unique-threshold methods to exploit their complementary strengths. It consists of three main elements: preprocessing, segmentation, and postprocessing. historical past and red blood cells of the image are extracted through distinct thresholds in segmentation procedure. The experimental outcomes advised that an overall segmentation give high accuracy because the first step of computerized white blood cellular differential system, it indicates a good prospect in further WBC function extraction, ALL category, and diagnosis. As a result of accuracy comparsion proposed method have 98% accuracy ,that is it have 8% more than existing method.

VI FUTURESCOPE

As the extension/futurescope of the project we can do 2 mode operation in this project,through uploading the datas which is obtaining from patient. Here it will work with respect to 2 mode operation. One is patient mode and other one is doctor mode. This will detail explain on next phase

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