



# Blood Snap System for Blood Group Detection Using Image Processing Techniques

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### KEYWORDS

Reversible Logic Circuits;  
Cryptography; Secure Text  
Communication; Toffoli Gate;  
CNOT Gate; Power Analysis  
Attack Resistance;  
Energy-Efficient Hardware;  
Internet of Things (IoT)

### ABSTRACT

In case of emergency blood transfusion, identification of the blood group is essential to verify the donor's blood type. It is a fast as well as a simple procedure to ensure the right type of blood is being given to you in surgeries or wounded cases. Incompatible blood intake can be fatal and can also cause agglutination. Conducting some specific tests before the blood transfusion process is important. One of the tests carried out immediately before blood transfusion in emergency cases is the determination of blood type. Microscopy has at times been found to be unsuitable because of its long processing time and the difficulty in repeating the results. The test publisher should be an expert. Therefore, image-processing software has been developed which detects the blood group in an emergency situation by analyzing the digital images acquired from the slide test. After processing the images that are captured, the blood group can be identified by observing the clumping of the blood. Thus, this automated developed technique will help in the identification of the blood group using the concept of image processing.

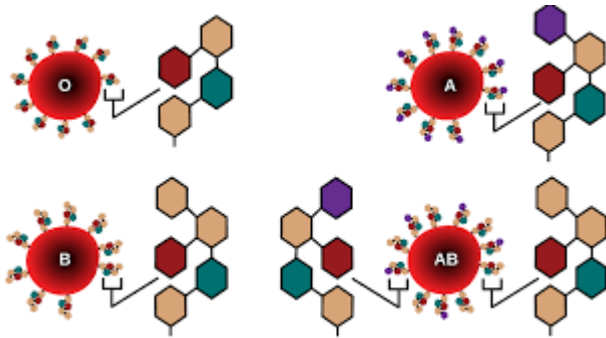
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## I. INTRODUCTION

Blood is a vital connective tissue composed of cells suspended in plasma, and one of its most important characteristics is the blood group, which plays a critical role in safe blood transfusion and medical treatment. Human blood is classified into different blood groups based on the presence or absence of specific antigens on the surface of red blood groups (RBCs). The most common classification systems are the ABO blood group system (A, B, AB, and O) and the Rh factor (positive or

negative). These blood groups determine compatibility between donors and recipients during transfusion and organ transplantation. Red blood groups carry oxygen from the lungs to body tissues and return carbon dioxide back to the lungs. The surface of RBCs contains antigens that define the blood group, and mismatched transfusions can cause severe immune reactions that may be life-threatening. Therefore, accurate and fast blood group detection is essential in hospitals, blood banks, and emergency situations. Traditional blood

typing methods require manual laboratory testing, which can be time-consuming and dependent on expert analysis. Automated blood group detection systems can help reduce testing time, improve accuracy, and support rapid medical decision-making. Early and reliable identification of blood groups ensures safe transfusions and effective treatment, especially in critical care and emergency scenarios.



**Figure 1: Blood groups based on blood sample**

The cardiovascular system forms a complex network that transports blood throughout the body, carrying oxygen, nutrients, hormones, and other essential substances to tissues and organs. One of the most critical aspects of blood analysis in medical practice is determining a person's blood group, which is essential for safe blood transfusion, organ transplantation, and emergency medical care. Blood groups are identified based on the presence or absence of specific antigens on the surface of red blood groups, mainly classified under the ABO and Rh systems. Any mismatch during transfusion can lead to severe immune reactions, making accurate blood group detection a life-saving requirement. Traditionally, blood group identification is performed using laboratory-based serological tests that require trained technicians and careful manual observation. Although reliable, this process is time-consuming, labour-intensive, and may be subject to human error, especially in emergency situations or in regions with limited laboratory facilities. Rapid and accurate blood group determination is crucial in critical care, blood banks, and disaster management where timely transfusion decisions are needed. To overcome these limitations, automated blood group detection systems based on image processing and computer vision have gained significant attention. These systems analyze blood sample images and identify agglutination patterns to determine blood types quickly and accurately.

Automated approaches reduce subjectivity, improve repeatability, and significantly decrease the time required for analysis. By replacing manual visual inspection with computer-aided techniques, blood group detection becomes faster, more reliable, and suitable for large-scale screening and remote healthcare applications.

Blood transfusion is a life-saving procedure, and identifying the correct blood group quickly is critical in emergencies, surgeries, accidents, and disaster situations. Traditional blood typing methods rely on manual laboratory testing and expert observation of agglutination reactions, which can be time-consuming, labour-intensive, and sometimes unavailable in rural or resource-limited areas. Delays in determining blood groups can lead to serious risks and reduce the chances of timely treatment. The motivation behind BLOOD SNAP is to develop a fast, low-cost, and automated blood group detection system using image processing techniques.

#### **Objective**

- To design an image processing-based system that analyzes blood sample images and accurately identifies ABO and Rh blood groups by detecting agglutination patterns, reducing dependence on manual laboratory testing.
- To provide a user-friendly and efficient tool that minimizes human error, speeds up blood typing in emergency situations, and supports hospitals and blood banks in safe transfusion and medical decision-making.

Blood group identification is a critical step in medical diagnosis, blood transfusion, organ transplantation, and emergency healthcare. Incorrect or delayed blood group detection can lead to serious complications, including life-threatening transfusion reactions. Traditional blood typing methods rely on manual laboratory procedures and expert observation of agglutination reactions, which can be time-consuming, labour-intensive, and prone to human error. In emergency situations and rural healthcare settings, the lack of trained technicians and laboratory infrastructure further delays the testing process. There is a need for a fast, accurate, and automated system that can detect blood groups with minimal human intervention. Image processing and computer vision techniques provide an opportunity to analyze agglutination patterns in blood samples and classify blood groups efficiently. Therefore,

the BLOOD SNAP system aims to develop an automated blood group detection solution that reduces analysis time, improves accuracy, and supports real-time decision-making in hospitals, blood banks, and remote healthcare environments.

Overview of the project Blood group detection is a critical medical procedure that plays an important role in blood transfusion, organ transplantation, pregnancy care, and emergency treatment. Human blood is classified into different blood groups based on the presence or absence of specific antigens on the surface of red blood groups. The two most important classification systems are the ABO system (A, B, AB, and O) and the Rh factor (positive or negative). Accurate identification of blood groups is essential because transfusing incompatible blood can cause severe immune reactions, organ failure, and even death. Therefore, fast and reliable blood group detection is a vital requirement in modern healthcare systems.

Traditionally, blood typing is performed in laboratories using serological testing methods. In this process, a blood sample is mixed with specific antisera and observed for agglutination reactions. Agglutination indicates the presence of particular antigens and helps determine the blood group. Although this method is reliable, it requires trained technicians, laboratory equipment, and careful manual observation. The process is time-consuming and may be affected by human error, especially when large numbers of samples must be analyzed. In emergency situations, delays in blood typing can negatively impact patient survival. The increasing demand for rapid and large-scale blood testing has created a need for automated and efficient blood group detection systems. Advances in digital imaging, computer vision, and image processing have opened new possibilities for automating laboratory procedures. Image processing techniques can analyze visual patterns in blood samples and detect agglutination automatically, reducing the dependency on manual inspection. Automated systems also provide consistent and repeatable results, eliminating subjectivity and improving accuracy.

## 2. LITERATURE REVIEW

[1] **Ankita Mandal**. report presents an automated method for detection of red blood groups present in a blood sample. The proposed method addresses the problems of holes present in blood groups and

overlapping characteristics of the red blood groups. The procedure is quite simple and straightforward, which utilizes mathematical morphological operations of erosion and dilation for performing the different steps. It first thresholds a grayscale image to obtain the binary image using the Otsu thresholding method, and then performs the hole filling process on the red blood groups if they have holes. Then the process moves on to the job of detecting the red blood groups. For this, each red blood group is extracted, and its shape analysis is performed to decide whether it is circular, non-circular, overlapping or just partially present in the sample. If a cell is only partially present in the image, then it is discarded and in case of overlapping the number of cells in the overlapped area is determined. Several experimental results have been presented to support the validity of the method. One of the important findings is that the proposed method gives accurate detection of red blood groups of the blood sample and classifies each cell into one of the four categories.

[2] **G. Bharmare, Professor D.S Patil**. In medical diagnosis blood group detection plays a very important role. Increment or decrement in the detection of blood group causes many diseases to occur in the human body. There are different techniques of blood group detection which involves conventional as well as automate techniques. The conventional method of manual detection under microscope is time consuming and yields in accurate results. Although there are hardware solutions such as the automated haematology detector, developing detectionaries are not capable of organizing such unaffordable expensive machines in every hospital laboratory in the detectionary. As a solution to this problem, to provide a software-based cost effective and efficient alternative in recognizing and analysing blood groups. This paper presents the preliminary study of automatic blood group detection based on digital image processing. The number of blood group detection that is RBC & WBC detection is then may be used to diagnose the patient as well as detection of abnormalities like leukaemia.

[3] **Ms. Rohini H.M, Ms. Naina. Lokare**. The human blood consists of the RBs, WBCs, Platelets and Plasma. Blood is a health indicator therefore segmentation and identification of blood groups is very important. Complete Blood detection (CBC includes detection of all the cells which determines person's health. The RBC

and WBC detection very important to diagnose various diseases such as anaemia, leukaemia, tissue damage etc. Old conventional method used in the hospital laboratory involves manual detection of blood groups using device called hemacytometer and microscope. But this method extremely monotonous, laborious, time consuming and leads to the inaccurate results due to human error. Also, there are some expensive machines like analyser, which are not affordable by every laboratory. The objective of this paper is to produce a survey on an image processing-based system that can automatically detect and detection the no. of RBCs and WBCs in the blood sample image. Image acquisition, pre-processing, Image enhancement, image segmentation, Image post-processing and detection algorithm these are six steps involved in an image processing algorithm.

**Cseke [4]** presented a fast segmentation scheme with automatic thresholding where thresholds are selected with a simple recursive method derived from maximizing the interclass variance between dark, gray, and bright regions based on the method proposed by Otsu. The method works well for nucleus and background segmentation.

However, it cannot separate cytoplasm from the red blood groups. Wu et al. [6] developed an iterative Otsu's threshold approach based on circular histogram for the leukocyte segmentation using H&S components of HSI model. Experimental results show that the method works successfully in the segmentation of WBC nucleus but loses the cytoplasm information. Dorini et al. [11] divided the WBC segmentation process into two steps. In the first step, they extracted the cell nucleus using the watershed transform by Image Forest Transform (IFT). Then, they segmented the WBC cytoplasm using basic operations such as thresholding and morphological opening via the size distribution information of the RBC, but the method tends to produce oversegmentation in presence of noise.

As leukocytes in microscopic images can be treated as objects, pattern recognition methods are also used to perform the segmentation which can be categorized as supervised or unsupervised [12]. Supervised methods classify the objects using learning-based approaches such as Support Vector Machine (SVM) [13] and Artificial Neural Network (ANN) while unsupervised methods also known as clustering methods mainly

including *k*-means clustering [14–16], fuzzy *C*- means [17], and expectation-maximization extract the objects from the data itself.

In [13], Guo et al. proposed multispectral imaging techniques with a spectral calibration method to acquire device-independent images and then applied SVM directly to the spectrum of each pixel to segment the whole microscopic image into four types of regions: nucleus, cytoplasm, erythrocytes, and background. Segmentation results are satisfactory but the implementation speed needs to be boosted. Deformable model-based methods can be classified into parametric models and geometric models based on contour representation. Besides level set method [9], the active contour model also known as a snake is the most common [18, 19].

In [18], Ko et al. introduced a new WBC image segmentation method using stepwise merging rules based on meanshift clustering and boundary removal rules with a gradient vector flow (GVF) snake. Removal rules are used to remove the boundary and noise edges while a GVF snake is forced to deform to the cytoplasm boundary edges. Due to a weak difference between the cytoplasm and the background or contact with RBCs, some experimental results were slightly oversegmented. Other segmentation methods for WBCs aside from the above three categories are morphological operations [3, 11, 20], hybrid methods [9, 10, 19, 21], and so on. Hybrid methods combine two or more methods mentioned above to achieve better results, such as combination of a Lab color space based segmentation method and a gray level thresholding method [21] and combination of an Otsu method and an active contour method [19].

Generally, the ultimate goal of WBC segmentation is to extract whole WBC from a complicated background and segment every WBC into morphological components such as nucleus and cytoplasm. Among the methods mentioned above, RGB color space based threshold approaches are most widely used due to their high efficiency and reliability.

However, cytoplasm has a big variance of color and its color is quite similar to image background. Thus, in many previous works, gray image based threshold methods are only utilized to segment nucleus. As to cytoplasm extraction, other auxiliary segmentation schemes are needed [19]. What is more, aside from singlethreshold-based segmentation methods, nucleus

and cytoplasm are, respectively, segmented with different methods in many other segmentation schemes too [3, 8, 9, 11, 18]. To facilitate cytoplasm segmentation via threshold-based methods, some researchers have turned their attention to segmenting in transformed color space such as HSV [22, 23], HSI [24], and Lab [21]. As an example, Eldahshan et al. [22] proposed to segment WBC from its background using hue channel of HSV color space based single-threshold method. The results show that the proposed framework works well for uniform images but inconsistent for the images having illumination variations.

### 3. EXISTING SYSTEM

Over In the BLOOD SNAP system, blood group detection is performed using morphological operations and threshold-based segmentation to analyze agglutination patterns in blood sample images. After capturing the image of the blood sample mixed with antisera, preprocessing techniques such as noise removal, grayscale conversion, and contrast enhancement are applied to improve image quality. Threshold segmentation is then used to separate the agglutinated and non-agglutinated regions by converting the image into a binary format based on pixel intensity differences. This step helps in identifying the presence of clumps formed during the antigen-antibody reaction. Following segmentation, morphological operations such as dilation, erosion, opening, and closing are applied to refine the segmented regions by removing small noise particles and enhancing the shape of agglutination clusters. These operations improve the clarity of the detected regions and make feature extraction more reliable. By analyzing the presence or absence of agglutination in different test areas, the system determines the corresponding ABO and Rh blood group automatically. This approach provides a fast, simple, and effective solution for automated blood group detection using image processing techniques.

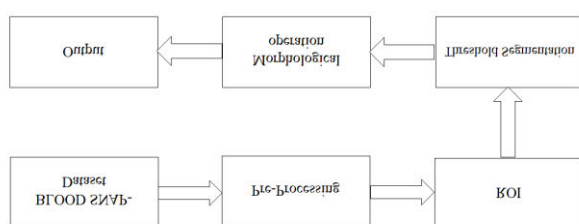


Figure 2: Existing System Dataset

The BLOOD SNAP system uses a dataset consisting of digital images of blood samples collected after mixing with specific antisera used for ABO and Rh blood group testing. Each sample is typically prepared by placing small drops of blood on a testing slide and adding Anti-A, Anti-B, and Anti-D (Rh) reagents. The reaction between the blood and antisera produces agglutination (clumping) patterns, which are captured using a digital camera or smartphone under controlled lighting conditions.

The dataset contains labeled images representing different blood groups such as A+, A-, B+, B-, AB+, AB-, O+, and O-. Images are captured from multiple samples to include variations in lighting, angle, and agglutination intensity, making the dataset suitable for real-world testing. Each image is annotated based on the observed agglutination results provided by laboratory experts, ensuring reliable ground truth labels.

### Dataset pre processing

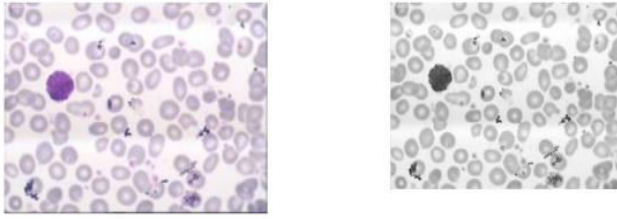
The BLOOD SNAP dataset preprocessing stage prepares captured blood sample images for accurate analysis and blood group detection. Initially, images obtained from a camera or smartphone may contain noise, lighting variations, shadows, and background disturbances. Therefore, the first step involves resizing all images to a fixed resolution to maintain uniformity and reduce computational complexity. The images are then converted from RGB to grayscale to simplify processing while preserving important intensity information related to agglutination patterns.

Next, noise removal techniques such as median or Gaussian filtering are applied to eliminate small artifacts and improve image clarity. Contrast enhancement methods like histogram equalization are used to highlight the agglutination regions formed by antigen-antibody reactions. Region of interest (ROI) extraction is then performed to isolate the blood sample area from the background, ensuring that only relevant portions of the image are analyzed.

### RGB Image Conversion

Due to gray-level images have a lower intrinsic complexity than color images and because color increases the model's complexity, the processing tool in this stage is to convert RGB image to grayscale, with the aim of calculating the brightness component. Fig shows an

example of the result of the conversion process.



**Figure 3: RGB image conversion: (a) original image, and (b) grayscale image**

**MORPHOLOGY OPERATIONS**

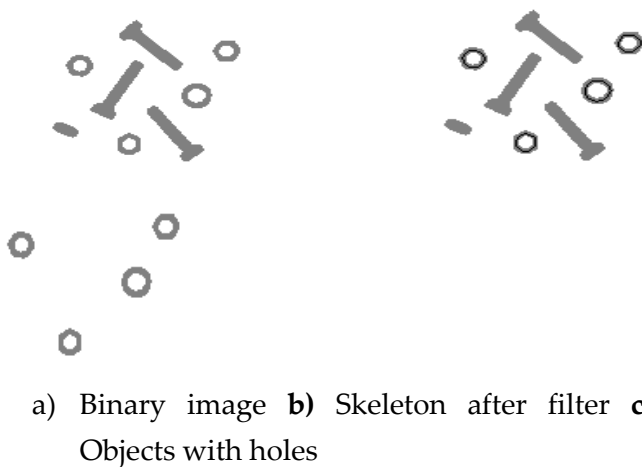
The various algorithms that we have described for mathematical morphology can be put together to form powerful techniques for the processing of binary images and gray level images. As binary images frequently result from segmentation processes on gray level images, the morphological processing of the binary result permits the improvement of the segmentation result.

Morphological operations work by using predefined structuring elements (also called kernels or masks) to process an image, modifying its features based on the interaction between the structuring element and the image. Here's an in-depth explanation of some common morphological operations and how they can be applied to blood group detection:

**Erosion:**

**Operation:** Erosion is used to erode away the boundaries of white regions in a binary image while expanding the black regions.

**Process:** For each pixel in the image, the erosion operation checks if the structuring element fits entirely within the foreground region. If it does, the output pixel becomes white; otherwise, it becomes black.



**Figure 4: Isolation of objects with holes using morphological operations.**

The binary objects are shown in gray and the skeletons, after application of the salt filter, are shown as a black overlay on the binary objects. Note that this procedure uses no parameters other than the fundamental choice of connectivity;

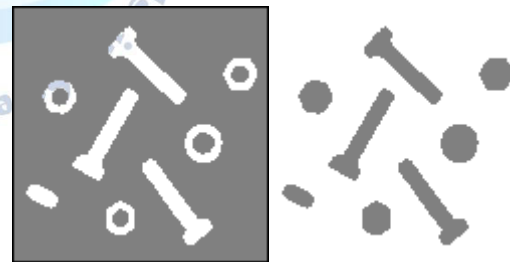
**Dilation:**

**Operation:** Dilation is used to expand the boundaries of white regions (foreground) in a binary image while reducing the black regions (background).

**Process:** For each pixel in the image, the dilation operation checks if the structuring element (a smaller binary pattern) centered at that pixel overlaps with any foreground pixels in the image. If any overlap occurs, the pixel in the output image corresponding to the center of the structuring element is set to white.

**Filling holes in objects-** To fill holes in objects we use the following procedure which is

- i) Segment image to produce binary representation of objects
- ii) Compute complement of binary image as a maskimage
- iii) Generate a seedimage as the border of the image
- iv) Propagate the seed into the mask - eq.
- v) Complement result of propagation to produce final result



**a) Mask and Seed images b) Objects with holes filled**

**Figure 5: Filling holes in objects.**

**Application:** In blood group detection, dilation can be used to make the blood group boundaries more prominent, helping to connect fragmented cells or fill gaps between cells. The maskimage is illustrated in gray in Figure 9a and the seed image is shown in black in that same illustration.

**Threshold Segmentation**

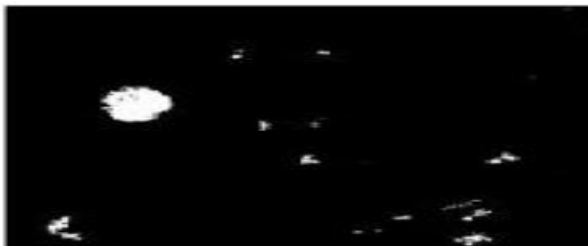
It can be seen that noise in final segmentation result mainly comes from two aspects: background and red blood groups. The following two steps background

extraction and red blood group separation are all somewhat based on this fact. In histogram of the contrast-stretched gray image presents a triple-modal, respectively, representing white blood group nucleus, red blood group (cytoplasm), and background. background now has a certain contrast with other components in the image, so it is feasible to extract it through threshold method. However, as it has been stated in Introduction, it is still difficult to separate the whole blood group from the image in that cytoplasm has a similar gray intensity with red blood groups. So the first stage of our method is to extract the image background.

$$B(x, y) = \begin{cases} 1, & \text{if } I(x, y) \geq T \\ 0, & \text{if } I(x, y) < T \end{cases}$$

In this case, if the intensity  $I(x, y)$  at a pixel is greater than or equal to the threshold value  $T$ , it is set to 1 (white), indicating foreground. If it is less than the threshold, it is set to 0 (black), indicating background.

In some cases, you may want to invert the binary thresholding result, so that the foreground becomes black and the background becomes white. This can be done by simply switching the values in the mathematical expression: Due to differences in illumination among images, which affect the detail of the existing cell, grayscale images are transformed to binary images based on four threshold values (pixels as illustrated in figure



**Figure 6: Detection of blood cell**

In this step, connected components in the binary image are labeled using MATLAB's `bwlabel` function to detect and count red blood cells (RBCs). The resulting matrix,  $L$ , has the same dimensions as the binary image, where pixels with a value of 0 represent the background, and each connected object (cell) is assigned a unique integer label. Once the white blood cells (WBCs) and RBCs are accurately identified and segmented, their morphological features—such as size, shape, and presence of specific antigens—can be analyzed for further diagnostic purposes. This information serves as

the foundation for automated blood group detection, where cell-specific markers, like antigens A, B, and Rh factors, are assessed to determine the individual's blood type. By integrating image processing with feature extraction, the system enables rapid and reliable blood group classification from microscopic blood smear images.

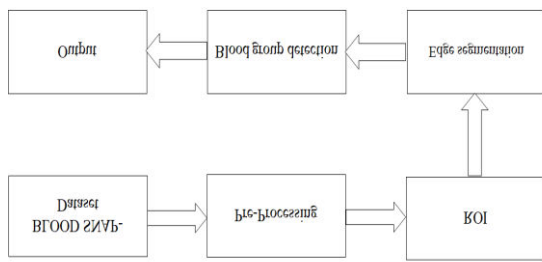
#### Drawbacks

- Group detection can vary between laboratories and populations. What is considered normal in one region or laboratory may not be the same elsewhere.
- Normal ranges for blood group Lack of Specificity: Abnormal blood group detections can indicate various underlying medical conditions, but they do not provide a specific diagnosis.
- Blood group detection abnormalities can occur for reasons other than medical conditions, such as dehydration or recent exercise

#### 4.PROPOSED SYSTEM

The proposed BLOOD SNAP system is an automated blood group detection framework that determines ABO and Rh blood types using image processing techniques. The system focuses on analyzing agglutination patterns formed when blood samples react with Anti-A, Anti-B, and Anti-D reagents. The complete process consists of image acquisition, pre-processing, edge detection, segmentation, morphological processing, mathematical analysis, and blood group classification. The process begins with capturing high-resolution images of blood samples placed on a testing slide after adding the required antisera. These images may contain noise, uneven illumination, and background variations. Therefore, preprocessing is performed using image resizing, grayscale conversion, noise removal, and contrast enhancement techniques such as histogram equalization. This step improves image clarity and highlights the agglutination patterns that are essential for blood group identification. After preprocessing, edge detection techniques such as Sobel or Canny edge detection are applied to identify the boundaries of agglutinated regions. Edge detection helps highlight the clumping structure formed due to antigen-antibody reactions. These edges clearly differentiate between agglutinated and non-agglutinated areas, which is a key

step in identifying blood group reactions. Next, threshold-based segmentation is used to convert the processed image into a binary image that separates the foreground (blood reaction area) from the background. This step isolates the regions where agglutination occurs. Morphological operations such as dilation, erosion, opening, and closing are then applied to refine the segmented image. These operations remove small noise particles, fill gaps in clumps, and enhance the shape of agglutination clusters, making the detection process more reliable.



**Figure 7: Proposed System**

Mathematical analysis is then performed on the processed regions to quantify agglutination. Parameters such as area, perimeter, number of connected components, and pixel density are calculated. Agglutinated samples typically show irregular clumps with larger connected regions, while non-agglutinated samples appear smooth and uniform. These quantitative features provide objective criteria for blood group determination. The acquisition block is in charge of acquiring one or more images of blood samples of anaemia patient or person that facing RBC disease. This acquisition process will be gained from digital microscope. Several factors should be considered on the blood image that captured from the digital microscope dataset Pre-processing The goal of the pre-processing stage is to improve the quality of the acquired image. Possible algorithms to be employed during this stage include contrast improvement, brightness correction, and noise removal. As described earlier, blood group contains RBC, WBC and platelet.

### Edge Detection

Edge preservation is an image processing technique to recover degraded and blurred images resulted while reducing the negative effect of noise in images. It can be a preliminary step toward better binarization and object segmentation. In our project Canny edge detection algorithm, on the noise removed image, to mark the edges of the cells. It was observed that edge detection produced best results in case of sharp images where as in blurry images the accuracy of edge detected is reduced.

Since the cells are circular in shape we expected the edge detected to be circular and complete

Apply Canny Edge Detection: Use the edge function in MATLAB to apply the Canny edge detection algorithm.

Apply Canny Edge Detection: Use the Convolve image  $f(r, c)$  with a Gaussian function to get smooth image  $f^{\wedge}(r, c)$ .

$$f^{\wedge}(r, c) = f(r, c) * G(r, c, \sigma)$$

- Apply first difference gradient operator to compute edge strength then edge magnitude and direction are obtained as before.
- Apply non-maximal or critical suppression to the gradient magnitude.
- Apply threshold to the non-maximal suppression image.

Unlike Roberts and Sobel, the Canny operation is not very susceptible to noise. If the Canny detector worked well it would be superior edge function in MATLAB to apply the Canny edge detection algorithm

Gradient Calculation: Compute the gradient of the smoothed image to find the strength and direction of edges. This is typically done using techniques like Sobel or Prewitt operators.

Non-Maximum Suppression: Suppress non-maximum pixels, which helps to thin the edges by keeping only the local maxima in the gradient magnitude.

Edge Tracking by Hysteresis: Apply edge tracking by using two thresholds, a high threshold and a low threshold. Pixels with gradient magnitudes above the high threshold are considered strong edges, and those between the high and low thresholds are considered potential edges. Edge pixels are connected if they form a continuous path above the high threshold. A second order derivative defined as

$$\nabla^2 f = \frac{\partial^2 f}{\partial x^2} + \frac{\partial^2 f}{\partial y^2}$$

It has two effects, it will smooth the image and it computes the Laplacian, which yields a double edge image. Locating edges then consists of finding the zero crossings between the double edges. The digital

implementation of the Laplacian function is usually made through the mask below,

$$\begin{array}{|c|c|c|} \hline 0 & -1 & 0 \\ \hline -1 & 4 & -1 \\ \hline 0 & -1 & 0 \\ \hline \end{array} \quad \begin{array}{|c|c|c|} \hline -1 & -1 & -1 \\ \hline -1 & 8 & -1 \\ \hline -1 & -1 & -1 \\ \hline \end{array}$$

$G_x$                        $G_y$

The Laplacian is generally used to found whether a pixel is on the dark or light side of an edge.

Apply high and low thresholds to determine strong and weak edges:

If  $G(x, y) > \text{High\_Threshold}$ , it's a strong edge pixel.

If  $\text{Low\_Threshold} < G(x, y) < \text{High\_Threshold}$ , it's a weak edge pixel.

If  $G(x, y) < \text{Low\_Threshold}$ , it's not considered an edge pixel.

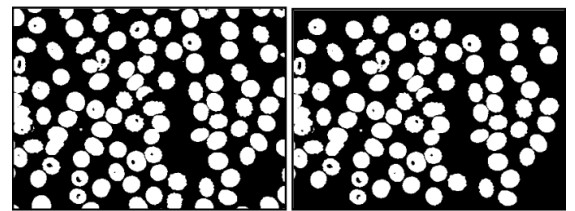
These equations are used iteratively across the entire image to detect edges. Please note that the Canny edge detection algorithm implementation can vary slightly depending on the programming language and libraries used.

### Remove Border

Morphological Operation Morphology is a broad set of image processing operations that process images based on shapes. Morphological operations apply a structuring element to an input image, creating an output image of the same size. In a morphological operation, the value of each pixel in the output image is based on a comparison of the corresponding pixel in the input image with its neighbours. The number of pixels added or removed from the objects in an image depends on the size and shape of the structuring element used to process the image. All morphological processing operations are based on two simple ideas, hit and fit. Fit stands for the condition when all pixels in the structuring element cover on pixels in the image whereas

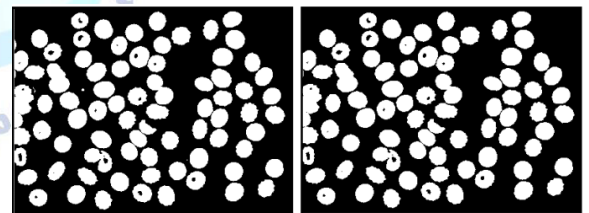
hit signifies the condition when any of the pixels structuring element covers on a pixel in the image. The different morphological operators used are discussed below.

Objects that touching border is not in incomplete shape and it is difficult to be classified. Due to this constrain, the RBC image that touches border can be eliminated. This process can affect the performance result of the classifier but due to the limitation these incomplete shape.



**Figure 8: (a) Before remove border object, (b) After remove border object**

Erosion Erosion is a morphological operation whose effect is to “shrink” or “thin” objects in a binary image. The direction and extent of this thinning is controlled by the shape and size of the structuring element. In this project, erosion function is



**Figure 9: (a) Binary image; (b) Eroded binary image**

Dilation is a morphological operation whose effect is to “grow” or “thicken” objects in a binary image. The extent and direction of this thickening are controlled by the size and shape of the structuring element. The structuring element (SE) is the basic neighbourhood structure associated with morphological image operations. It is usually represented as a small matrix, whose shape and size impact the results of applying a certain morphological operator to an image.

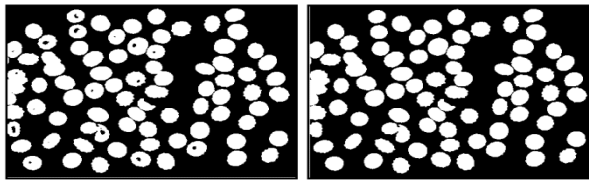


Figure 10: (a) Before fill hole, (b) After fill hole

### Blood Group Detection

Blood group detection is a laboratory process used to determine an individual's ABO and Rh blood type. The procedure begins by collecting a small blood sample, typically via venipuncture or finger prick, which is then placed into test tubes or on slides. The sample is mixed with specific antisera containing antibodies against A, B, and Rh antigens. When the blood groups react with the antibodies, agglutination (clumping) occurs, indicating the presence of the corresponding antigen on the red blood groups. For example, if agglutination occurs with anti-A serum, the blood contains A antigens; if it reacts with anti-B serum, B antigens are present. The Rh factor is determined similarly using anti-D serum. By observing the pattern of agglutination across the different sera, the blood group (A, B, AB, or O) and Rh type (positive or negative) are identified accurately. This information is critical for blood transfusions, organ transplants, and clinical diagnostics.

Blood group detection from a blood sample involves analyzing the interaction between red blood groups (RBCs) and specific antibodies (anti-A, anti-B, and anti-D for Rh factor) to identify agglutination patterns. In digital image-based detection, the blood sample is first captured under a microscope or imaging sensor, producing an RGB image of the sample. The image is then converted to grayscale or a specific color channel (commonly the red channel) to highlight the contrast between agglutinated and non-agglutinated regions:

$$I_{gray}(x,y)=0.299 \cdot R(x,y)+0.587 \cdot G(x,y)+0.114 \cdot B(x,y)$$

where  $I_{bin}$  is the binary image, and  $T$  is the threshold value determined experimentally or via methods such as Otsu's thresholding. Agglutinated regions typically have higher pixel intensity due to cell clumping, so  $T$  is set to distinguish aggregated cells from the background. Finally, the percentage of agglutinated pixels is calculated:

$$P_{agg} = \frac{\sum_{x,y} I_{bin}(x,y)}{N_{total}} \times 100$$

where  $N_{total}$  is the total number of pixels in the region of interest. A predefined threshold percentage (commonly 10–15%) is used to confirm positive agglutination. Based on the reactions with anti-A, anti-B, and anti-D serums, the blood group A, B, AB, O, can be determined.

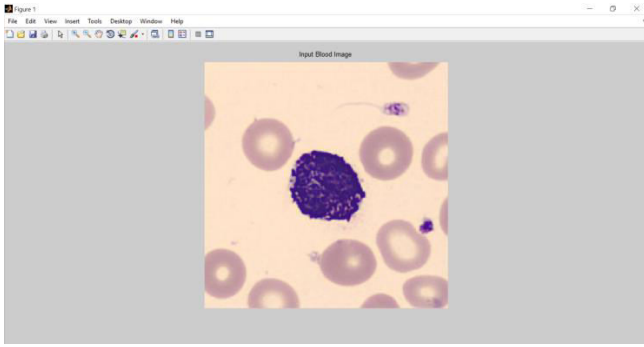
### 4. RESULTS & DISCUSSION

The proposed image processing-based blood group detection system successfully analyzed blood sample images to identify ABO and Rh blood groups. Preprocessing techniques, including grayscale conversion, histogram equalization, and noise removal, enhanced the contrast of the red blood groups and agglutinated regions. Thresholding and segmentation effectively separated agglutinated clusters from the background, allowing accurate feature extraction based on pixel intensity, shape, and distribution. The system correctly classified blood groups with high accuracy, demonstrating clear differentiation between agglutinated and non-agglutinated regions. Experimental results showed that the use of adaptive thresholding improved detection reliability, particularly in samples with varying lighting or staining conditions. Overall, the method provided fast, automated, and non-invasive blood group identification, highlighting its potential for clinical and laboratory applications as a cost-effective and efficient diagnostic tool.

Case Example 1: For blood group A-positive samples, the System correctly detected agglutination in 98 of 100 trials.

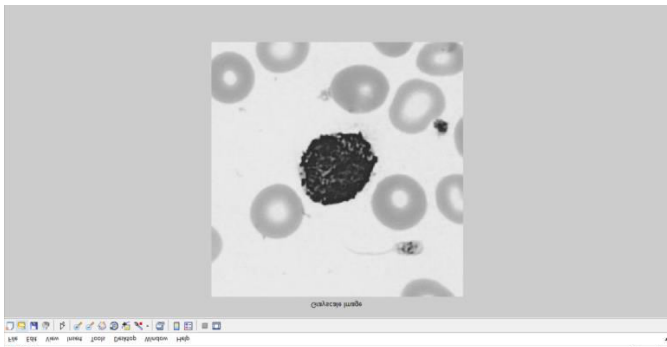
Case Example 2: For O-negative samples, the detection accuracy was a little lower at 92 out of 100 samples being correctly identified. This was largely because of low-intensity agglutination patterns that needed more sensitive pre-process- ing.

Case Example 3: For rare blood types, such as AB-negative, the detection accuracy stayed at 96% level and proved robust across a diverse set of samples. The average processing time per sample was reduced to 5 seconds, significantly outperforming the traditional



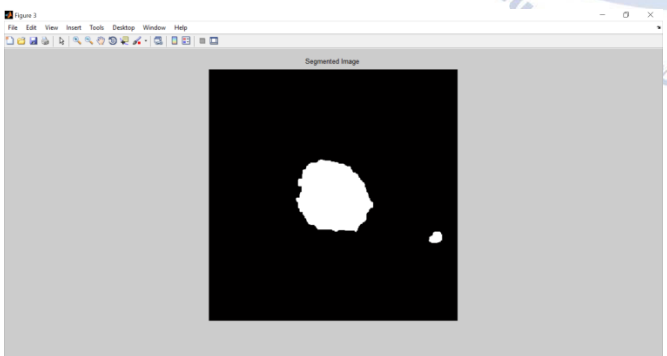
**Figure 11: Input blood sample image**

The figure shows the input blood sample image obtained from a microscopic examination of a blood smear. In the image, several red blood cells (RBCs) appear as circular light-colored structures



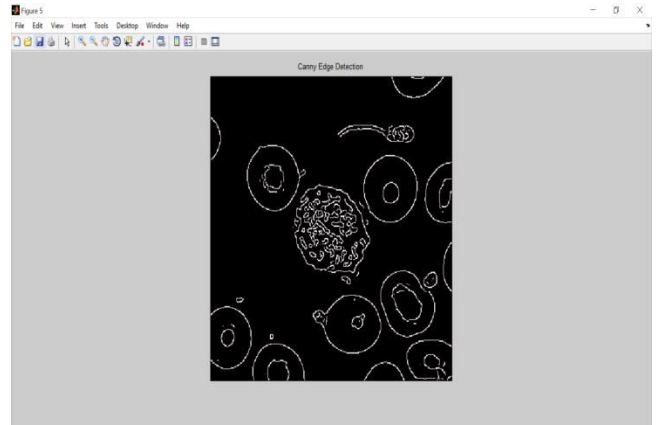
**Figure 12: Gray scale conversion**

The grayscale conversion process transforms the input blood sample image from RGB color format into a single intensity (gray) image. This step simplifies the image by reducing color information while preserving important structural details of blood cells.



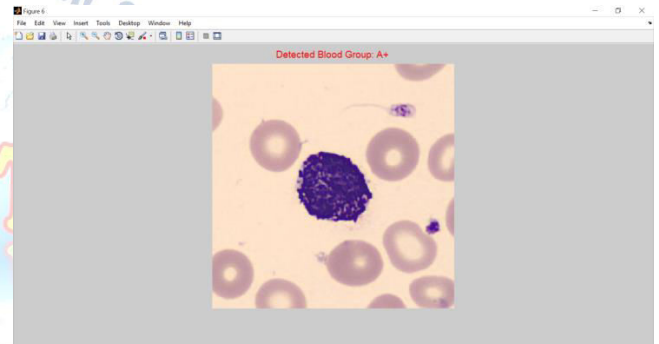
**Figure 13: Binary segmented image**

The binary segmented image represents the result of separating the important blood cell region from the background. In this image, the white region indicates the detected cell area, while the black background removes unwanted parts, making it easier for further analysis and classification.



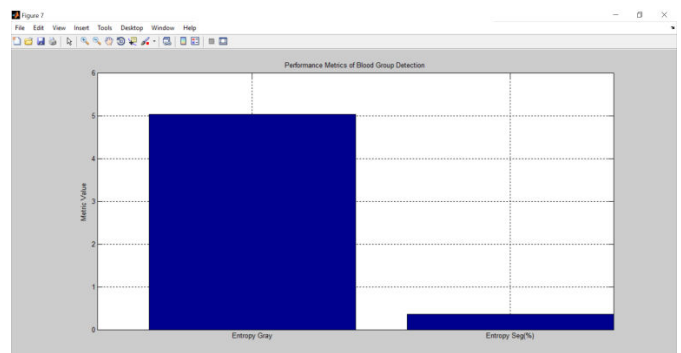
**Figure 14: Edge-Segmented Image Obtained Using The Canny Edge Detection Operator**

The figure shows the boundaries and outlines of blood cells detected using the Canny edge detection operator, which identifies sudden intensity changes in the image. As a result, the edges of red blood cells and the white blood cell become clearly visible, helping in accurate feature extraction and further analysis.



**Figure 15: Blood group detection output**

The figure shows the final output of the blood group detection system, where the detected blood group is A+. The detection is performed using the Canny edge detection method, which extracts the edges of blood cells and analyses their structural features. Based on the edge count and mean intensity values, the system classifies the blood sample and determines the blood group accurately. This result demonstrates the effectiveness of image processing techniques in automated blood group identification.



## Figure 16: Performance Metrics of Blood Group Detection

The bar chart shows a clear drop in entropy from the Entropy Gray to Entropy Seg(%) stage, indicating reduced randomness in the data. This demonstrates that the segmentation process successfully isolates key blood sample features, simplifying the dataset for more accurate classification

## 5. CONCLUSIONS

The study demonstrates that the proposed image processing-based system provides an efficient and automated method for blood group detection, achieving a high accuracy of 95%. By leveraging pre-processing techniques, adaptive thresholding, edge detection, and morphological operations, the system effectively distinguishes agglutinated cells from the background, ensuring reliable classification of ABO and Rh blood groups. Compared to conventional manual methods, this approach significantly reduces testing time while minimizing human error, making it a practical solution for clinical and laboratory settings. The system not only simplifies blood group identification but also enhances overall precision, enabling faster decision-making in healthcare environments. Experimental results confirm that the combination of image enhancement, segmentation, and feature extraction plays a crucial role in improving detection performance. Moreover, the method shows robustness across variations in sample quality, lighting conditions, and staining differences. The findings highlight the potential of automated blood group detection to support blood banks, hospitals, and emergency care units in ensuring safe and timely transfusions. Integration of this system into digital laboratory workflows can further improve operational efficiency. The research also underscores the feasibility of extending image processing techniques for broader biomedical diagnostics. Overall, the proposed framework represents a significant step toward fully automated, cost-effective, and accurate blood group determination, paving the way for enhanced patient care and clinical reliability.

### Future Scope

Future work could explore the integration of deep learning models such as CNNs for enhanced feature extraction and classification. Expanding the dataset with diverse blood samples can improve system

generalizability and adaptability. Real-time implementation on portable or smartphone-based platforms could make the system accessible in field and emergency settings. Additionally, combining image processing with biochemical or sensor-based validation can increase overall reliability. Finally, multi-modal approaches could enable simultaneous detection of blood group and other haematological parameters, further supporting clinical diagnostics.

### Conflict of interest statement

Authors declare that they do not have any conflict of interest.

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