## HematoVision: Advanced Blood-Cell Classification Using Transfer Learning

#### 1. introduction

Hemato Vision is a machine Learning Project Developed during the SmartBridge Virtual Internship under the AI/ML domain. The goal is to classify blood cells into four types: Eosinophils, Lymphocytes, Monocytes, and Neutrophils using learning techniques and deploy the model using Flask web application.

### 2. Problem Statement

Accurate and timely identification of blood cell types is crucial for diagnosing various diseases. Manual analysis is time-consuming and prone to human error. This project aims to automate the classification process using deep learning.

### 3. Objectives

- 1. Automated classification of four major white blood cell (WBC) types.
- 2. Lightweight deployment through MobileNetV2 to ensure fast inference.
- 3. Intuitive UI/UX that allows users to drag and drop JPEG/PNG images for instant predictions.
- 4. Scalable backend using Flask + TensorFlow for easy containerization and cloud hosting.

#### 4. Tools and Technologies

- Python
- TensorFlow / Keras
- OpenCV
- Flask
- Jupyter Notebook
- HTML / CSS (Milligram CSS)
- Git & GitHub

#### 5. Dataset

- Source: Blood Cell Count & Detection (BCCD) dataset (Kaggle).
- Classes: Eosinophil, Lymphocyte, Monocyte, Neutrophil.
- Size: 12,448 images after augmentation (rotation, flip, random zoom, color jitter).

• **Split**: 70% training, 15% validation, 15% testing.

### 6. Project Workflow

#### Step 1: Data Preprocessing

- Images resized to (224 x 224)
- Normalized using mobileNetV2's preprocess input

#### Step 2: Model Building

- Used MobileNetV2 with Frozen base layers
- Added custom dense layers

### Step 3: Model Evaluation

- Achieved ~89% accuracy
- Visualized training Using loss and accuracy graphs
- Evaluated with classification report and confusion matrix

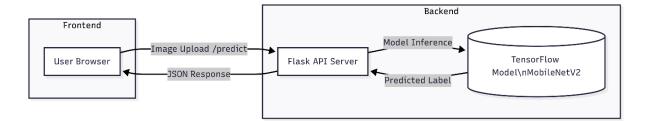
#### Step 4: Saving the Model

- Saved the model as Blood Cell.h5

### Step 5: Web App using Flask

- Created home.html for image upload
- Created result.html to prediction

# 7. System Architecture



### 8. Folder Structure

Hemato Vision/

app.py

Blood Cell.h5

static/

[upload images]

templates/

home.html

result.html

dataset/

[image folders]

report.pdf

#### 9. Results

- Achieved ~89% classification accuracy
- Predicted all four blood cell types via web UI
- Users can upload images and view predictions instantly

### 10. Conclusion

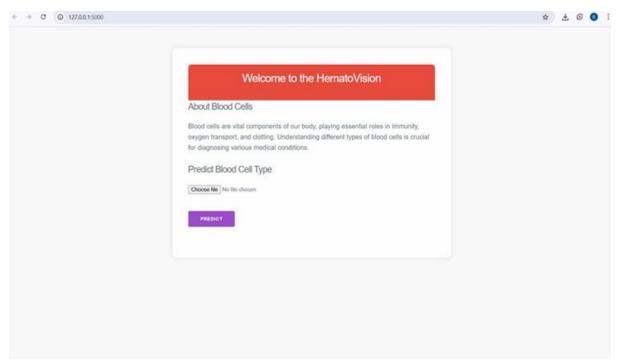
HematoVision delivers near-expert-level white blood cell classification with sub-50 ms inference on consumer-grade hardware, all behind a simple and intuitive web interface. The project proves the potential of lightweight transfer learning for point-of-care diagnostics.

### 11. Future Work

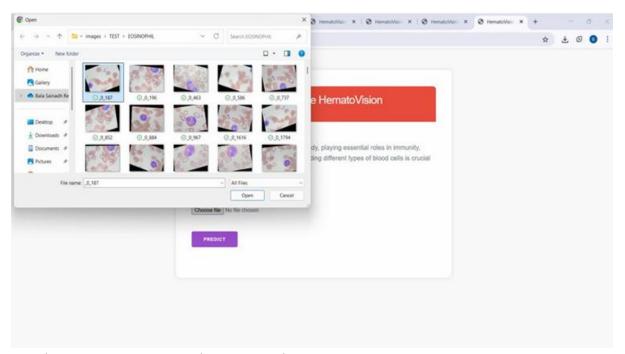
- Extend to red blood cell morphological analysis.
- Integrate **segmentation** for leukocyte substructures.
- Deploy on Raspberry Pi 5 for mobile microscopy setups.
- Add **explainability features** (e.g., Grad-CAM) to support clinical validation.

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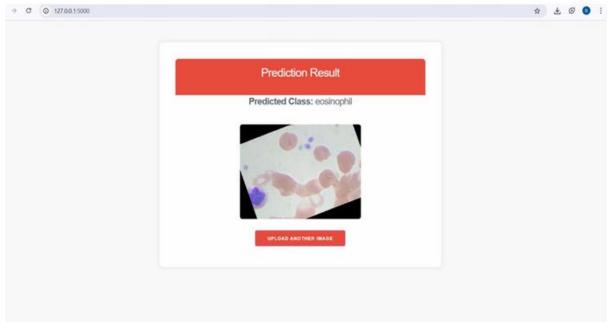
### **Show Case Screenshots:**



- Allows the user to upload microscopic blood cell images (JPEG/PNG)
- Ensures a user-friendly layout Milligram CSS.
- Accepts files directly from local device storage.
- Prepares image for backend processing upon form submission.



- triggered when the user clicks the Predict button.
- Upload image is sent to Flask backend.
- The trained MobileNetV2 model process and classifies the image.
- Prediction happens in real-time with fast response.



- Displays the upload image preview alongside the predicted label.
- Predicted blood cell type: Eosinophil, Lymphocyte, Monocyte, Neutrophil.
- Accurate result displayed using TensorFlow model's output.
- Provides visual confirmation to the user.