# IMPLEMENT DEEP LEARNING TECHNIQUES FOR MALARIA DETECTION USING IBM CLOUD

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# **Smart Bridge – Mini Project Report**

#### 1. INTRODUCTION

#### 1.1 Overview:

Malaria is a significant burden on our healthcare system and it is the major cause of death in many developing countries. It is endemic in some parts of the world which means that the disease is regularly found in the region. Therefore, early testing is necessary to detect malaria and save lives. Thus, it gives us the motivation to make malaria diagnosis more effective and faster. A specialized technology proves essential to combat this problem.

For Malaria diagnosis, the RDT and microscopic diagnosis are the most used clinical methods. RDT is an effective and faster tool. Also, it does not require the presence of a trained medical professional. But RDT has few drawbacks like susceptibility to damage by heat and humidity and higher cost compared to a light microscope. The microscopic diagnosis systems do not suffer from these shortcomings, but it requires the presence of a trained microscopist. Machine learning has recently made headlines in the medical field. The machine learning approaches have proved to be successful in the diagnosis of a disease.

#### 1.2 Purpose

The main aim of this malaria detection system is to address the challenges in the existing system by automating the process of malaria detection using Machine

#### 2. LITERATURE SURVEY

# 2.1 Existing problem

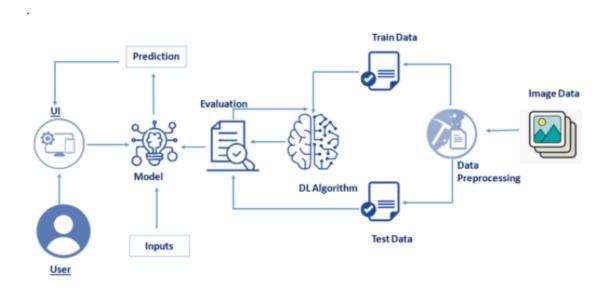
Malaria being a life threatening disease has caused deep research interests among the scientists all over the world. Earlier, malaria was mostly diagnosed in the laboratory setting requiring a great deal of human expertise. Automatic systems such as those relying on machine learning techniques were initially studied to overcome this problem. Techniques reported in this domain of study mostly considered the hand-crafted features in decision making. For example, relied on morphological factors for feature extraction and applied SVM and Principal Component Analysis (PCA) for the classification purpose. However, the accuracy achieved through these kinds of model is low compared to the more recently studied deep learning based techniques.

#### 2.2 Proposed solution

For automatic detection of malaria pathogens from the microscopic images, Convolutional Neural Network (CNN) received much attention from the researchers in recent times. Dong et al. for example evaluated the performances of three popular Convolutional Neural Network , namely LeNet-5 , AlexNet and GoogLeNet . In addition, they trained an SVM classifier for comparison purposes and concluded that CNN is advantageous over SVM in terms of the capacity of learning image features automatically. To find the optimal layer of a pretrained model to extract features from underlying malaria parasite data, Rajaraman et al. evaluated the performances of AlexNet, VGG-16 , ResNet-50 , Xception , DenseNet-121 along with their custom-built model . Liang et al. reported 97.37% accuracy in detection with the help of their 16-layered CNN model and claimed that their model is superior to transfer learning models . Hung et al. pre-trained a model on Imagenet but fine-tuned it on own data for detecting malaria parasite.

#### 3.THEORITICAL ANALYSIS:

## 3.1 Block diagram



# 3.2HARDWARE AND SOFTWARE REQUIREMENTS IN THE PROJECT:

For running a machine learning model on the system you need a system with minimum of 16 GB RAM in it and you require a good processor for high performance of the model. In the list of **software requirements** you must have:

- 1. Jupyter Notebook for programming, which can be installed by Anaconda IDE.
- 2. Python packages
- 3. A better software for running the html and css files for application building phase e.g. spyder.

## **4. EXPERIMENTAL INVESTIGATIONS:**

In the experiment of CNN-SVM and CNN-KNN, a CNN model was used to extract the features. Later these features were classified through SVM or KNN as per the objective of our experiment. The outline of

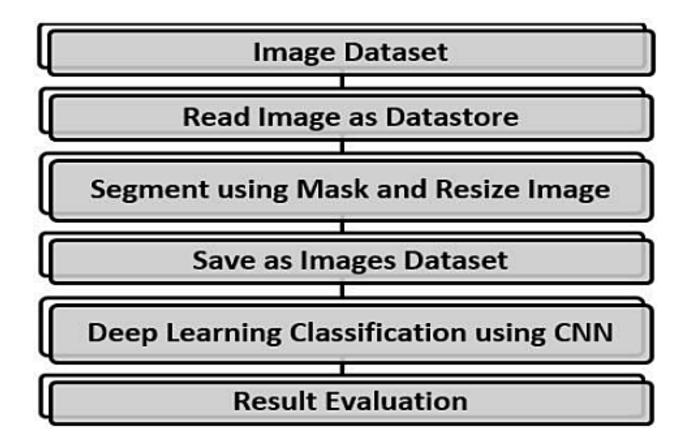
this architecture is depicted in

Support vector machine (SVM) and K-Nearest Neighbor (KNN) are the core machine learning algorithms that were used as a classifier by implicitly mapping their inputs into high-dimensional feature spaces . Earlier work reported the effectiveness of SVM or KNN classifier on the Convolutional neural network's extracted features instead of using the softmax layer. Being motivated with this work, we trained SVM and KNN models after extracting features from a CNN model.

Autoencoder Training Finally, autoencoder based architecture was explored to increase the model's performance. It is found that the autoencoder based experiments outperforms other experiments. Autoencoder is composed of two main components which are Encoder and Decoder discussed in the proposed model subsection. The main aim of an Autoencoder is dimensionality reduction, but to detect malaria parasite, it is used as a classifier. This classifier can classify infected and parasitized images from a microscopic blood smear.

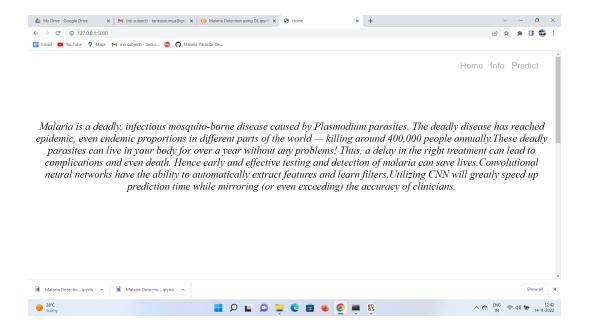
Stochastic Gradient Descent (SGD) was used as an optimizer with initial learning rate of 0.01 and momentum of 0.5 while training the student model. To update the weight, a decay of 1e-6 was applied using Keras learning rate decay function. To calculate the loss and update the weights and biases of all trainable parameters Cross Entropy Loss function was applied. Combine validation accuracy

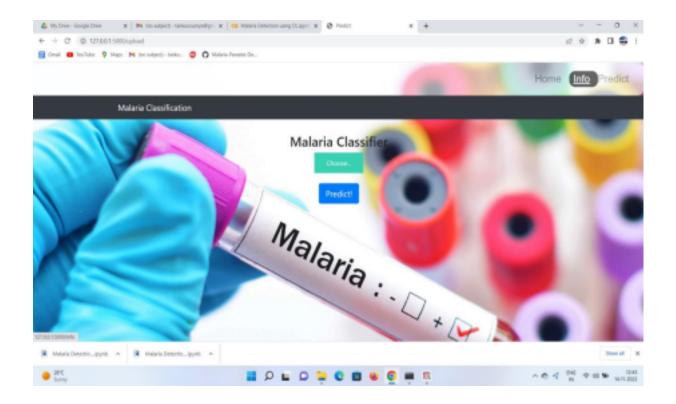
#### 5.FLOW CHART:-

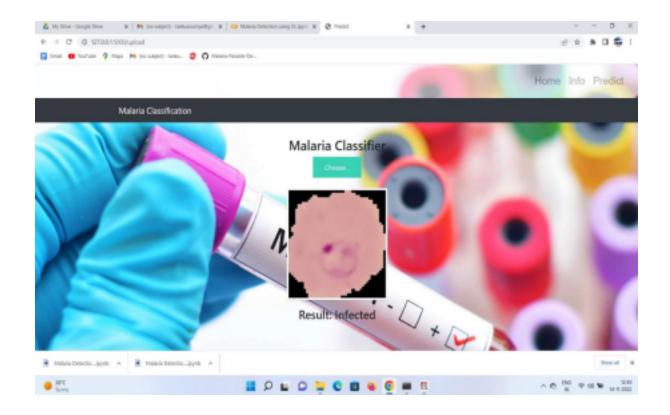


6.RESULTS:-

Final output of the project:







#### 7. ADVANTAGES & DISADVANTAGES

#### ADVANTAGES

Blood slide microscopy makes it possible to count the number of parasites and is more useful than rapid diagnostic tests for monitoring the effectiveness of malaria treatment.

**Magnification and higher resolution** – as electrons rather than light waves are used, it can be used to analyze structures which cannot otherwise be seen.

#### **DISADVANTAGES**

But RDT has few drawbacks like susceptibility to damage by heat and humidity and higher cost compared to a light microscope. The microscopic diagnosis

systems do not suffer from these shortcomings, but it requires the presence of a trained microscopist.

Machine learning has recently made headlines in the medical field.

#### 8.APPLICATIONS:-

Transfer learning can be effectively used for carrying out image recognition tasks. For example, using transfer learning, a model that is pre-trained for identifying dogs can be used to identify cats. Neural networks are trained on large datasets of images to recognise objects.

#### 9.CONCLUSION-

The correct diagnosis of malaria parasite species and correct treatment can reduce the number of malaria-infected individuals who carry the parasites between populations and may thus reduce the risk of re-introducing of malaria into other parts of Iran, outside this province, where an interruption of transmission had been previously achieved.

## **10.FUTURE SCOPE**:

Practical validation of model efficiency was performed by deploying the model in 10 different mobile phones of varying computational capacity and a server-backed web application. Data gathered from these environments show that the model can be used to perform inference under 1 s per sample in both offline (mobile only) and online (web application) mode. These inference speeds coupled with the high classification accuracy lead us to believe that this work can play a part towards building a fully automated system for malaria parasite detection which may be useful in resource-constrained areas in the foreseeable future

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[Google Scholar]

#### APPENDIX :-

# 1. Google Collab

https://drive.google.com/file/d/16rzVDqoB9OXszacIOUbaMjFQW55A Kc Ri/view?usp=sharing

#### 2. Flask Code

import os

import numpy as np #used for numerical analysis

from flask import Flask,request,render\_template# Flask-It is our framework which we are going to use to run/serve our application.

#request-for accessing file which was uploaded by the user on our application.

#render\_template- used for rendering the html pages

from tensorflow.keras.models import load\_model#to load our trained model

from tensorflow.keras.preprocessing import image

```
app=Flask(__name__)#our flask app
model=load_model(r"C:\Users\DELL\OneDrive\Desktop\iomp\MalariaDetection-DL-
main\Flask\maleria.h5")#loading the model
@app.route("/") #default route
def about():
  return render_template("about.html")#rendering html page
@app.route("/about") #route about page
def home():
  return render_template("about.html")#rendering html page
@app.route("/info") # route for info page
def information():
  return render_template("info.html")#rendering html page
@app.route("/upload") # route for uploads
def test():
  return render_template("index6.html")#rendering html page
@app.route("/predict",methods=["GET","POST"]) #route for our prediction
def upload():
  if request.method=='POST':
    f=request.files['file'] #requesting the file
    basepath=os.path.dirname('__file__')#storing the file directory
    filepath=os.path.join(basepath,"uploads",f.filename)#storing the file in uploads folder
    f.save(filepath)#saving the file
    img=image.load_img(filepath,target_size=(64,64)) #load and reshaping the image
    x=image.img_to_array(img)#converting image to array
    x=np.expand_dims(x,axis=0)#changing the dimensions of the image
```

```
pred=model.predict(x)#predicting the results
print("prediction",pred)#printing the prediction
if(pred==1):#checking the results
    result="Uninfected"
else:
    result="Infected"
    return result#resturing the result
return None

#port = int(os.getenv("PORT"))
if __name__=="__main__":
    app.run(debug=False)#running our app
#app.run(host='0.0.0.0', port=8000,debug=False)
```