BUG OFF MALARIA

# DEEP LEARNING TECHNIQUES TO DETECT MALARIA USING IBM CLOUD

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# INTRODUCTION

- Malaria is a life-threatening disease that is spread by the Plasmodium parasites. It is detected by trained microscopists who analyze microscopic blood smear images.
- Modern deep learning techniques may be used to do this analysis automatically.
- The need for the trained personnel can be greatly reduced with the development of an automatic accurate and efficient model. In our project, we have developed an automated convolutional neural network (CNN) based model which shall predict if a person has malaria using the microscopic blood smear images.
- Our deep learning-based model can detect malarial parasites from microscopic images with an accuracy of 96-97%.
- For a practical and real world application, we have made a website with which our model has been integrated with using flask.

#### PROBLEM STATEMENT

Implement Deep Learning Techniques to detect malaria using IBM Cloud

### SOLUTION



Malaria can be detected by analyzing microscopic blood smear images.

This process can be automated by using convolutional neural networks (CNN) model which shall use a training dataset of almost 28000 images to get a high accuracy in prediction as it becomes really important in the field of medicine to get diagnosed properly. We have made a website which allows the user to upload the blood smear image and our model will predict if malaria is detected or not.



#### TRAINING THE MODEL

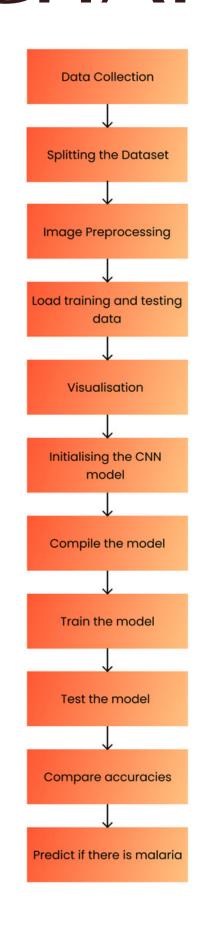
- We have used Python which is a statistical mathematical programming language.
- Import all the necessary libraries in python.
- Apply image DataGenerator functionality to train and test set
- Import required model building libraries
- Initialize the model, add convolution layers, pooling layer, flatten layer and dense layer
- Compile the model fit and save the model

#### TESTING THE MODEL

The model is to be tested with different images to know if it is predicting correctly. Import the packages that are used to load the model and get the predictions.

Pre-processing the image includes converting the image to array and resizing according to the model. Give the pre-processed image to the model to know to which class your model belongs to.

## PROCESS FLOWCHART





#### RESULT

So, after running the model through various iterations and a lot of fine-tuning, we need up with a model having an accuracy of around 96-97%, and was able to predict the uninfected and parasitized quite accurately.

The results mentioned above are presented in the next few slides:

```
In [22]: hist = malaria model.fit generator(x train, steps per epoch = 344, epochs = 30, validation data = x test, validation steps = 86, callbacks=[early stop])
  Epoch 1/30
  Epoch 2/30
  Epoch 3/30
  Epoch 4/30
  Epoch 5/30
  Epoch 6/30
  Epoch 7/30
  Epoch 8/30
  Epoch 9/30
  Epoch 10/30
  Epoch 11/30
  Epoch 12/30
  Epoch 13/30
```

```
Epoch 14/30
Epoch 15/30
Epoch 16/30
Epoch 17/30
Epoch 18/30
Epoch 19/30
Epoch 20/30
Epoch 21/30
Epoch 22/30
Epoch 23/30
Epoch 24/30
Epoch 25/30
Epoch 26/30
Epoch 27/30
Epoch 28/30
Epoch 29/30
Epoch 30/30
```

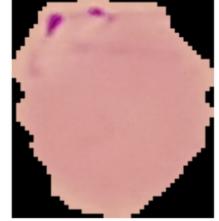
#### RESULT

#### Check whether the person is infected with malaria or not

Interactive, simple and quick way to predict malaria

#### **PREDICTION**

Upload image and check whether it is infected or not with malaria



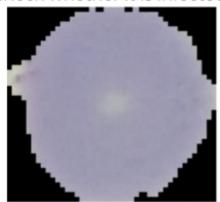
The prediction is: infected

#### Check whether the person is infected with malaria of

Interactive, simple and quick way to predict malaria

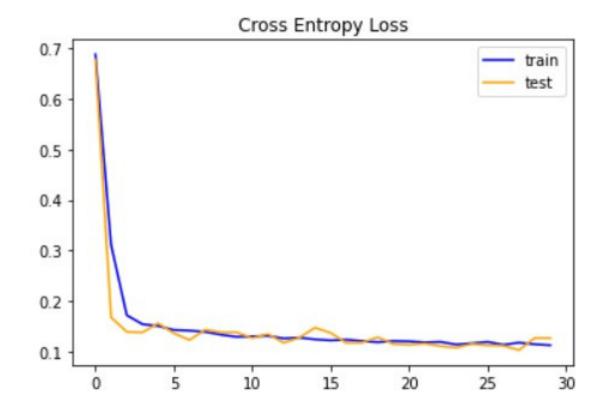
#### **PREDICTION**

Upload image and check whether it is infected or not with malaria



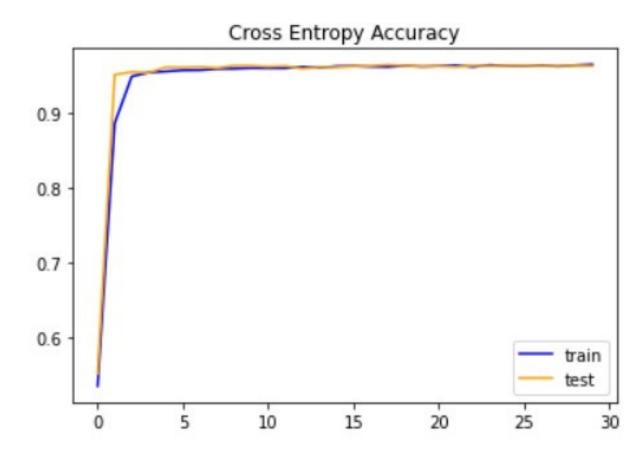
The prediction is: uninfected

Out[23]: <matplotlib.legend.Legend at 0x7f627508fe90>



```
In [25]:  # plot training and testing accuracy
plt.title('Cross Entropy Accuracy')
plt.plot(hist.history['acc'], color='blue', label='train')
plt.plot(hist.history['val_acc'], color='orange', label='test')
plt.legend()
```

Out[25]: <matplotlib.legend.Legend at 0x7f6257e8d8d0>



#### CONCLUSION



Well, malaria being one of the fatal diseases, has been a problem for a long time. But using deep learning models, like the ones that we used in our project, the detection of the disease can be done much faster and can be deal with the proper treatment accordingly.

Technologies like these neural networks have been a blessing to us and using them in the right way is gonna benefit all of us.



# THANK