

MACHINE LEARNING: AUTOMATED CELL

IMAGE ANALYSIS FOR

MALARIA INFECTION

DETECTION



GitHub <https://github.com/jnliou/project4>

- Introduction
- Solution Architecture
- Exploratory Data Analysis
- Data Science and Machine Learning
- Database
- Front-End Application
- Limitations and Future Recommendations
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- Our Team

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INTRODUCTION

What is Malaria?

Malaria is a severe and **occasionally lethal illness** caused by a parasite that commonly infects mosquitos that feeds on humans. Individuals who contract malaria typically experience severe symptoms such as high fevers, intense chills, and symptoms resembling the flu (CDC).



PREVALENCE

247 million

documented clinical cases of malaria

619,000

fatalities as of April 2023

South Africa is home to

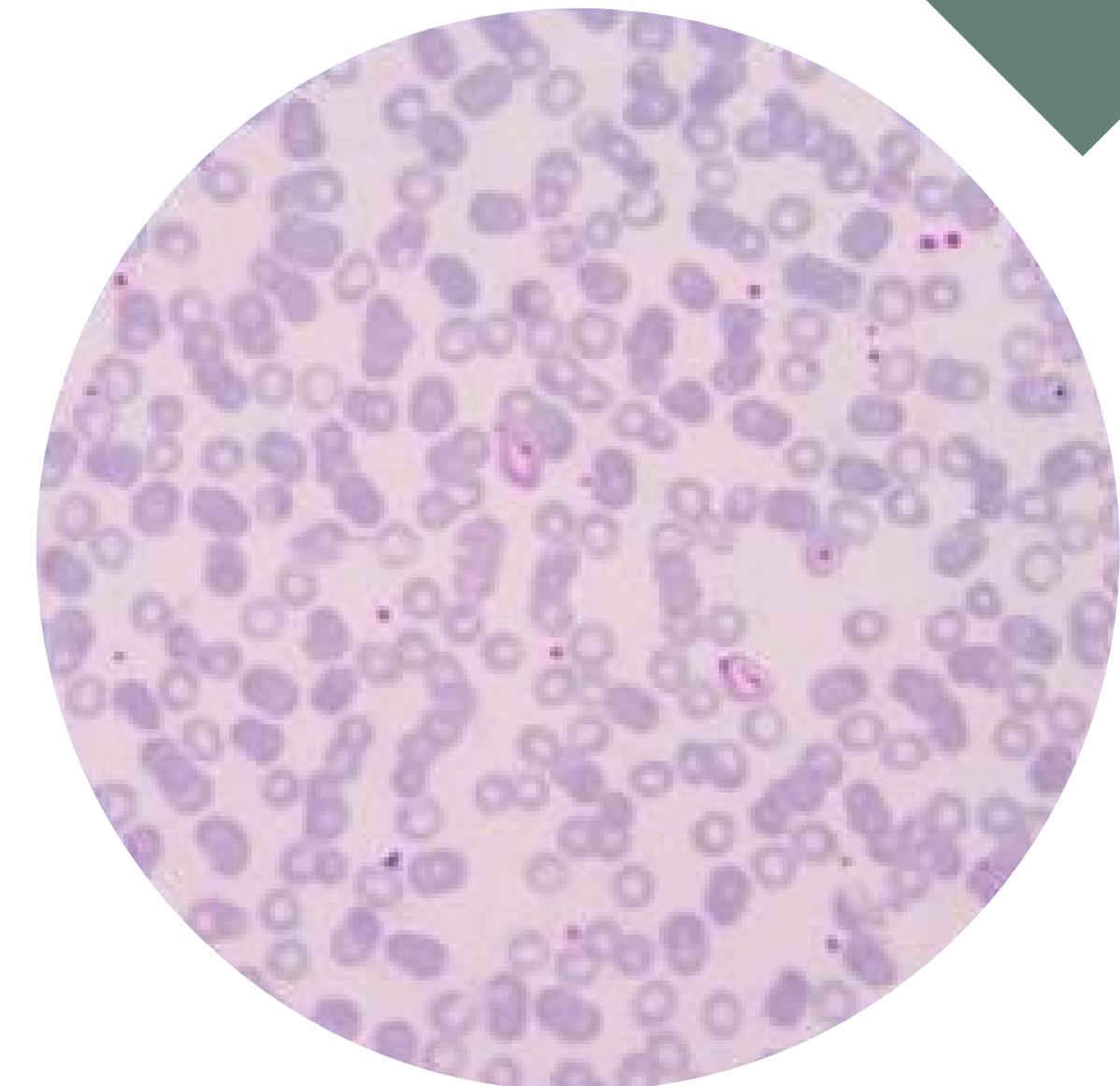
96%

victims out of which 76% are children

according to the WHO...

THE WHY?

- Leveraging Machine Learning Principles for Malaria Detection
- Develop predictive model using cell images
- Streamline healthcare diagnosis
- Empower public for timely assistance
- Facilitate learning for medical and science students

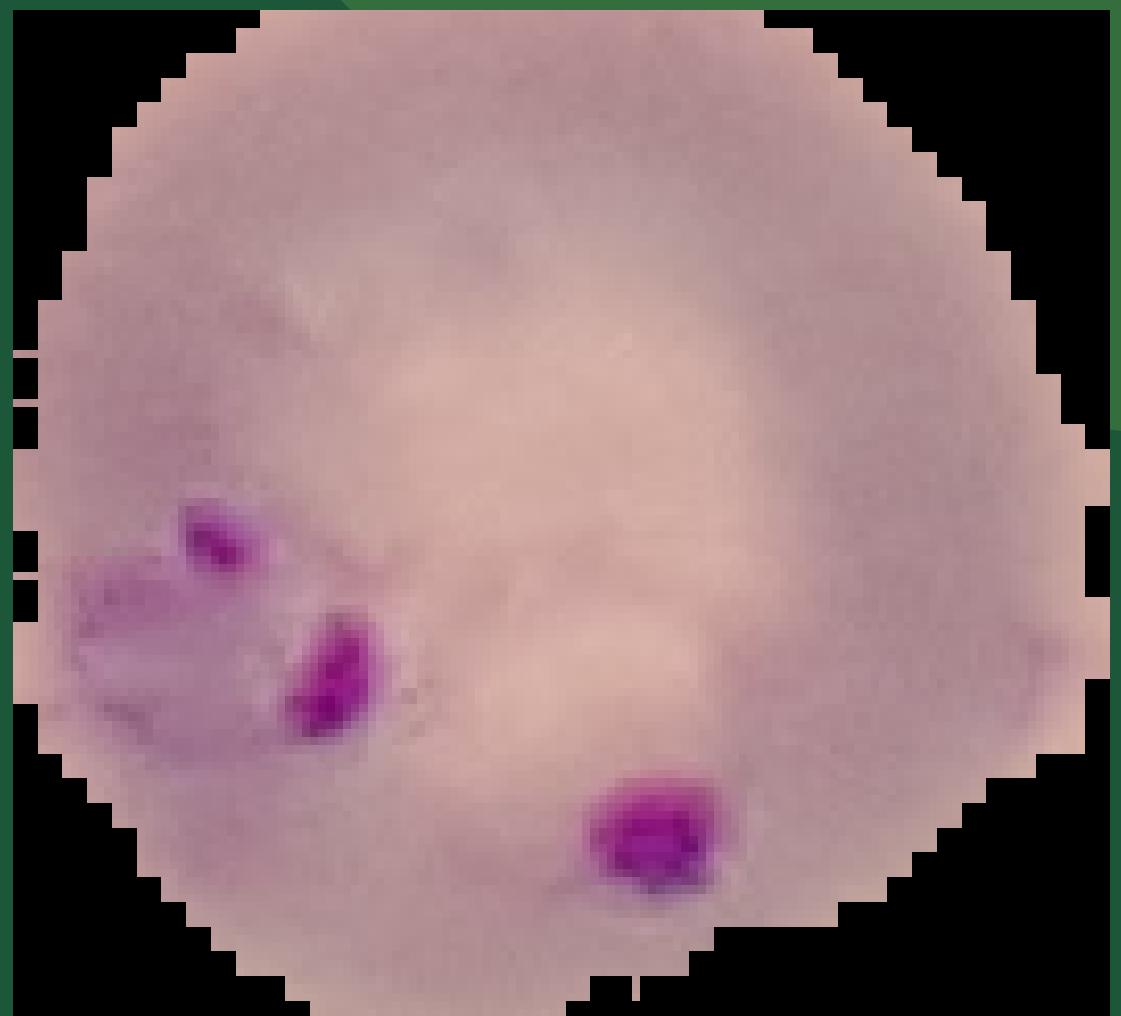


How is Malaria diagnosed?

According to the Center for Disease Control (CDC), Malaria parasites can be identified by examining **under the microscope** a drop of the patient's blood, spread out as a "**blood smear**" on a microscope slide.



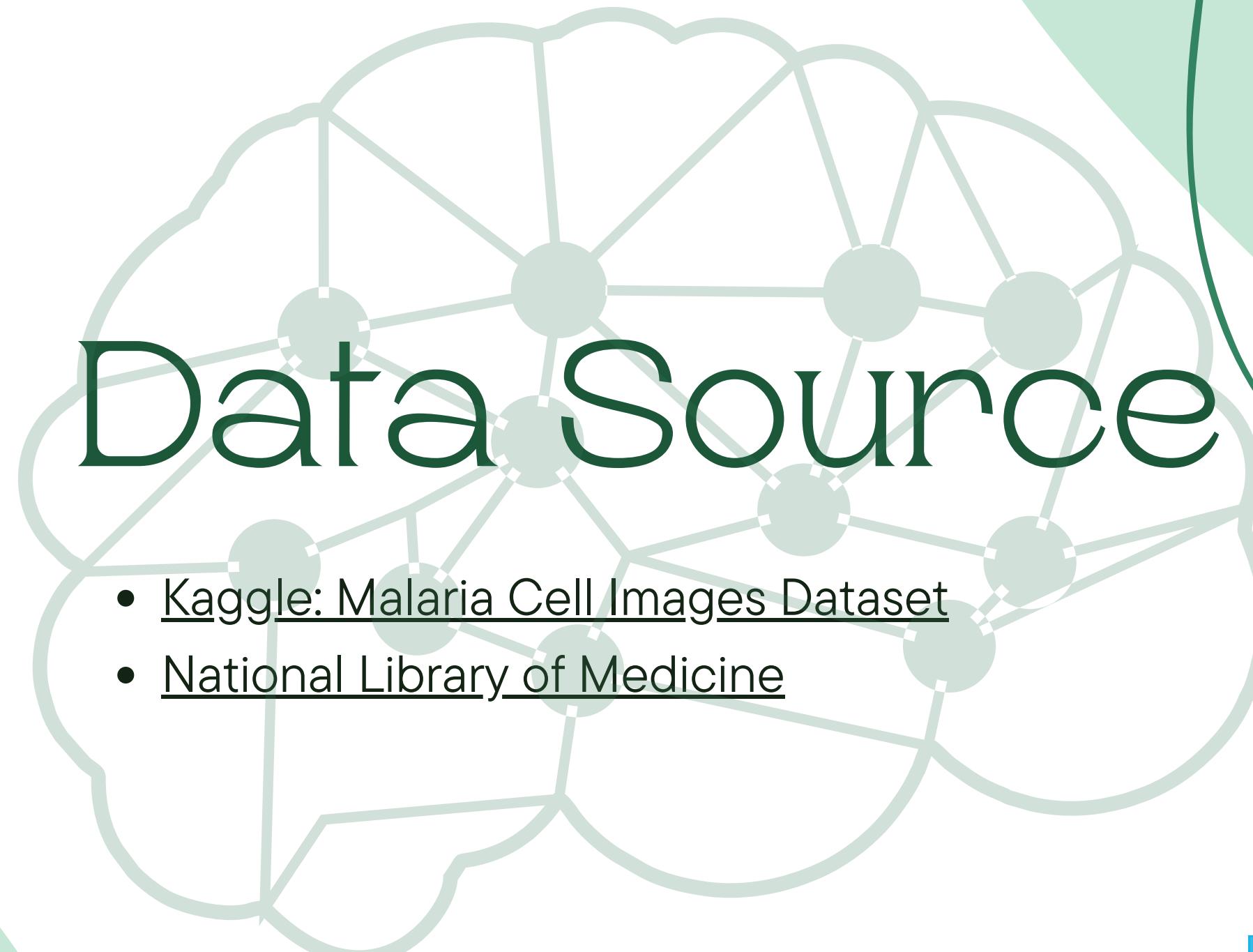
Uninfected



Infected



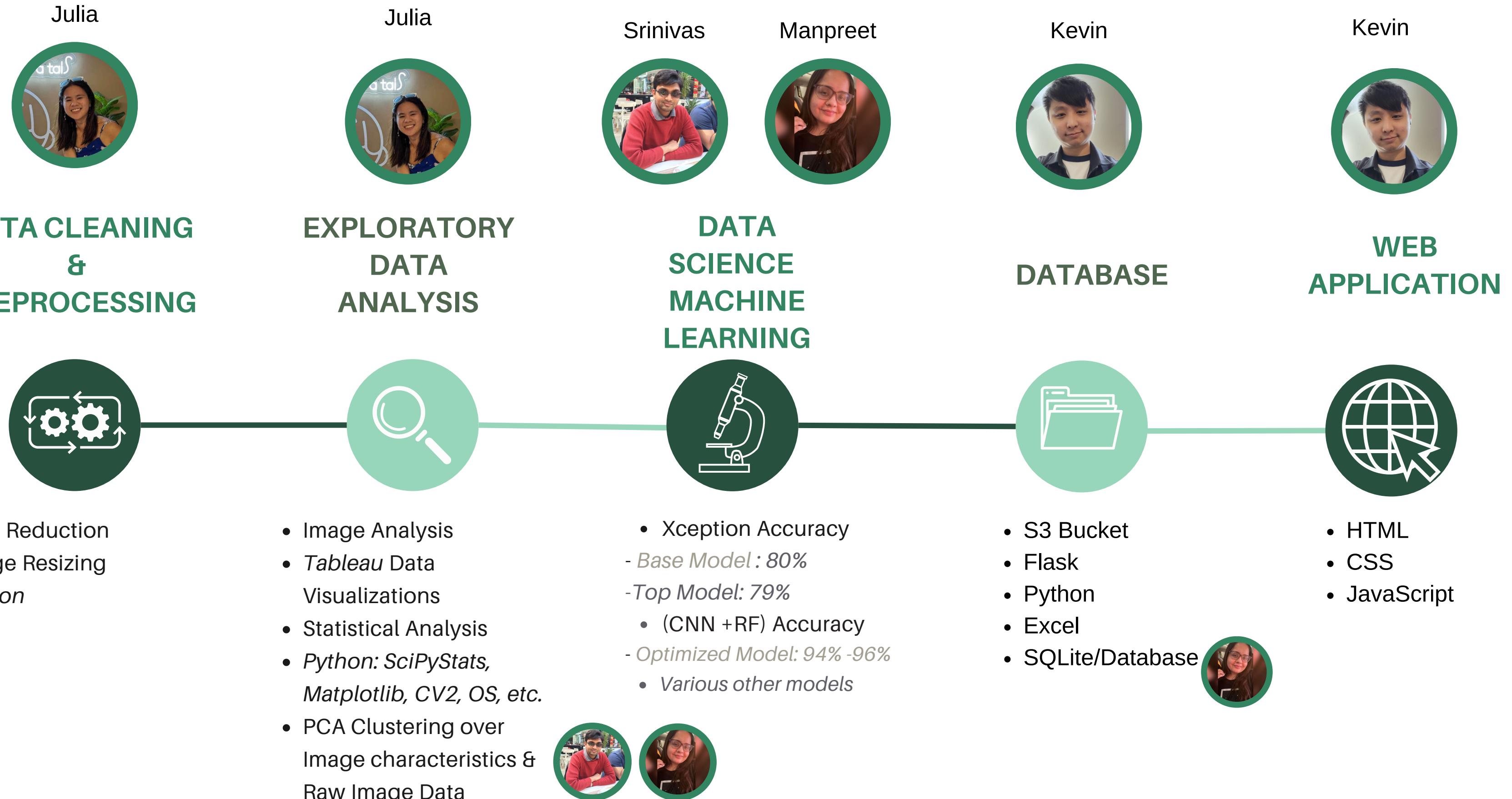
U.S. National Library
of Medicine



kaggle



Solution Architecture | Project Timeline

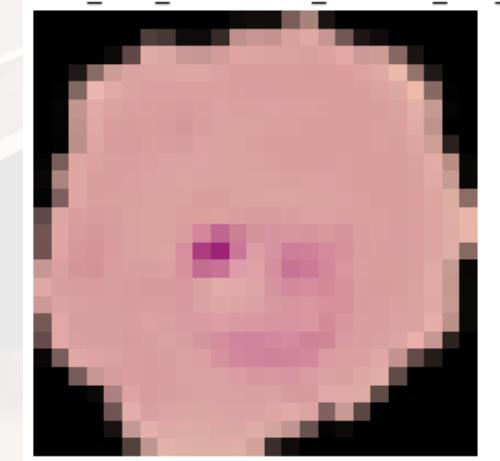
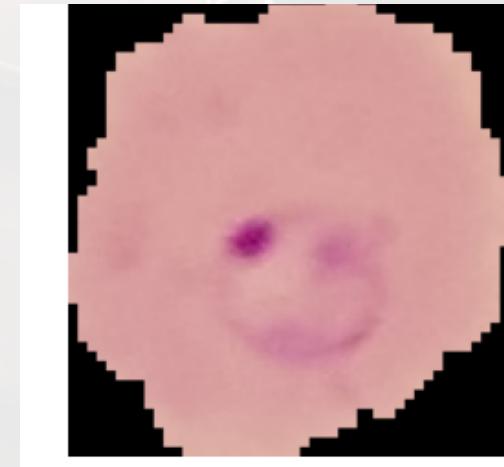


PREPROCESSING

Data Reduction

- 27600 to 5000 images
- 2500 infected 2500 uninfected
- 70% of images go into training folder 30% go into testing folder

150x150

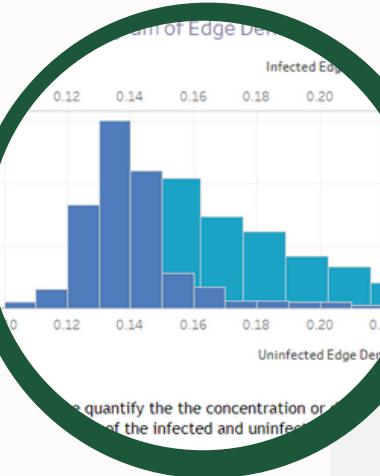


25x25

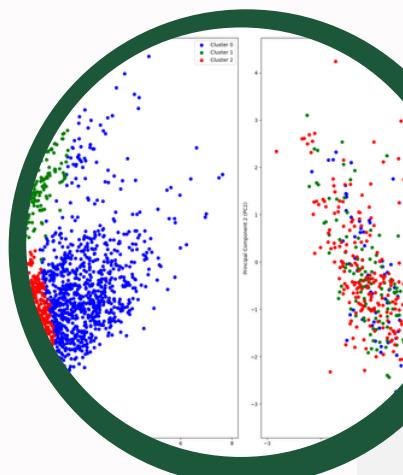
Image Resizing

- Converting image size from 150x150 to 25x25

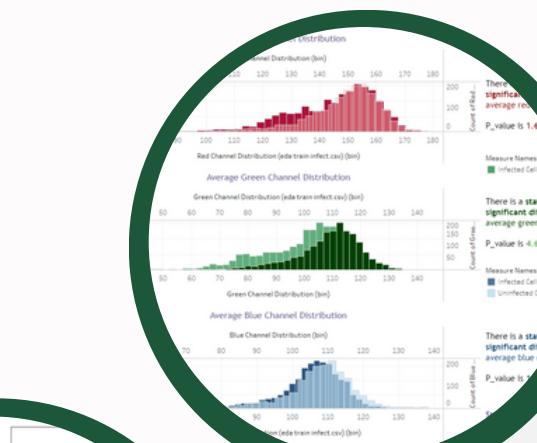
Exploratory Data Analysis



Statistical Analysis & Tableau
Mann Whitney U T-Test,
Data Visualizations via
Matplotlib and Tableau



PCA Clustering of Image Classification
Utilizing Machine Learning
Principles of PCA
Clustering for EDA &
Image Pixels



RGB Channel Distribution
Average Red, Green
& Blue Color Channel
Distrubtions



Assessing Blobs
Mean Blob Size &
Max Blob Size



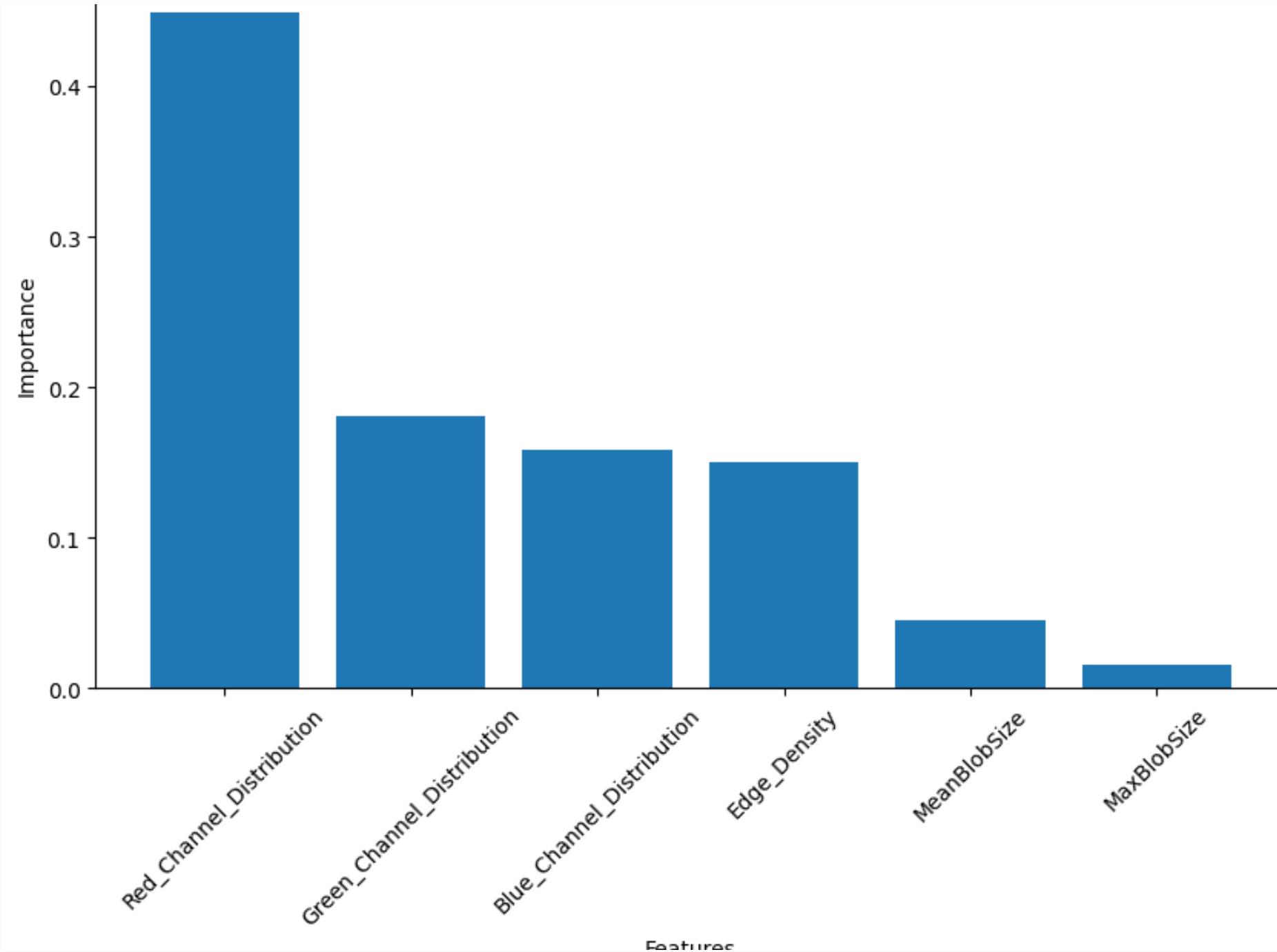
MODEL SELECTION

EDA DATASET

				EDA Features					
Types of Models →	Random Forest	Random Forest + Hyp	RF + Gradient boosting	Liner Regression Model	SVC model	SVC+Hyp	SVC + PCA	SVC+RF+NN	Decision Tree
Hypter parameter		0.10				0.10			
Parameters									
Uninfected									
Accuracy	83%	89%	83%		79%	85%	79%	81%	74%
Precision	0.81	0.88	0.80		0.78	0.83	0.72	0.77	0.72
Recall	0.83	0.90	0.86		0.85	0.83	0.88	0.84	0.70
F1-score	0.82	0.89	0.83		0.82	0.83	0.79	0.8	0.71
Infected									
Precision	0.85	0.90	0.87		0.79	0.86	0.87	0.85	0.75
Recall	0.83	0.88	0.81		0.70	0.86	0.71	0.78	0.77
F1-score	0.84	0.89	0.84		0.74	0.86	0.78	0.81	0.76
R2				0.32					
MSE				0.17					

MODEL SELECTION

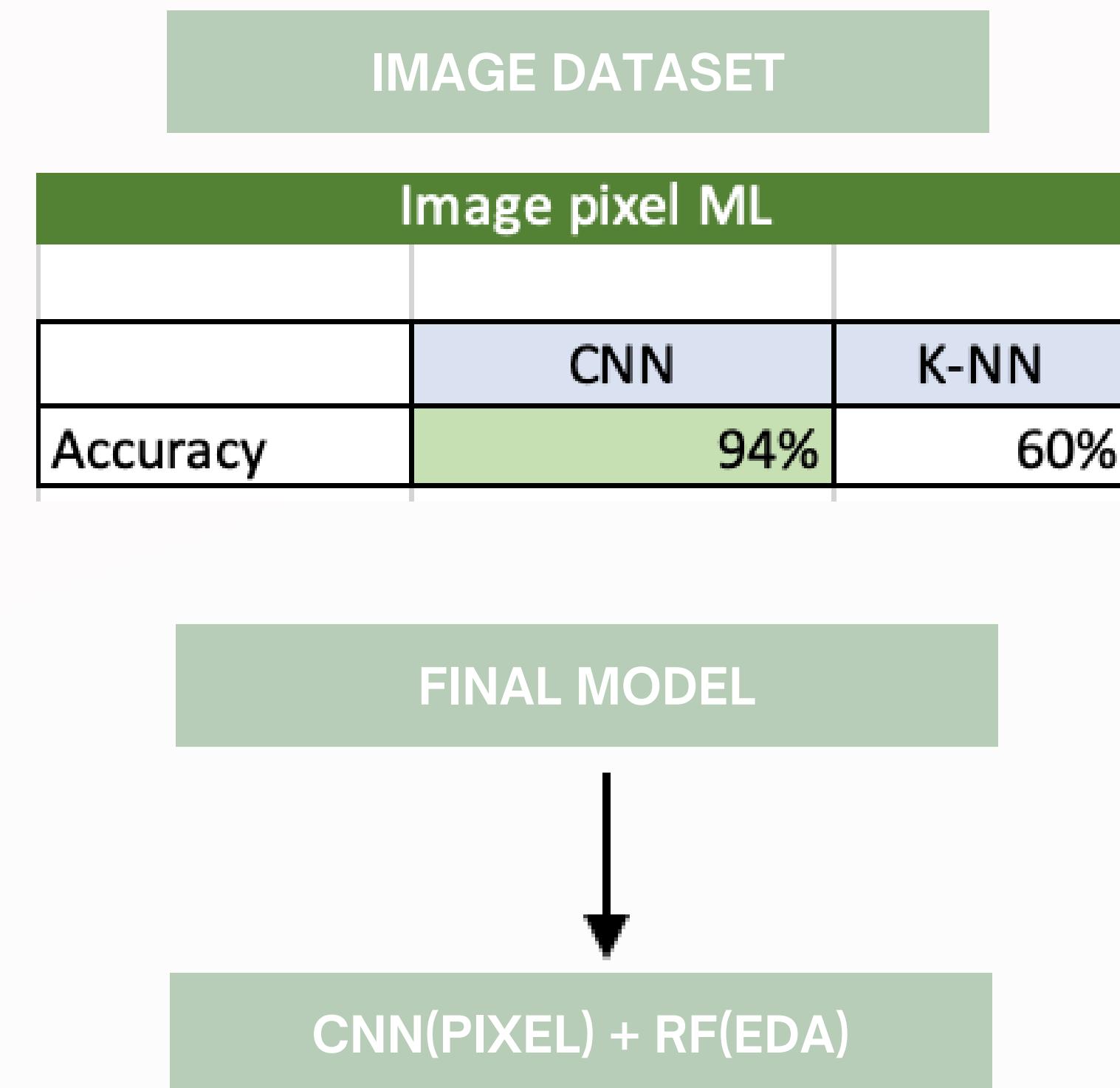
IMPORTANT FEATURES



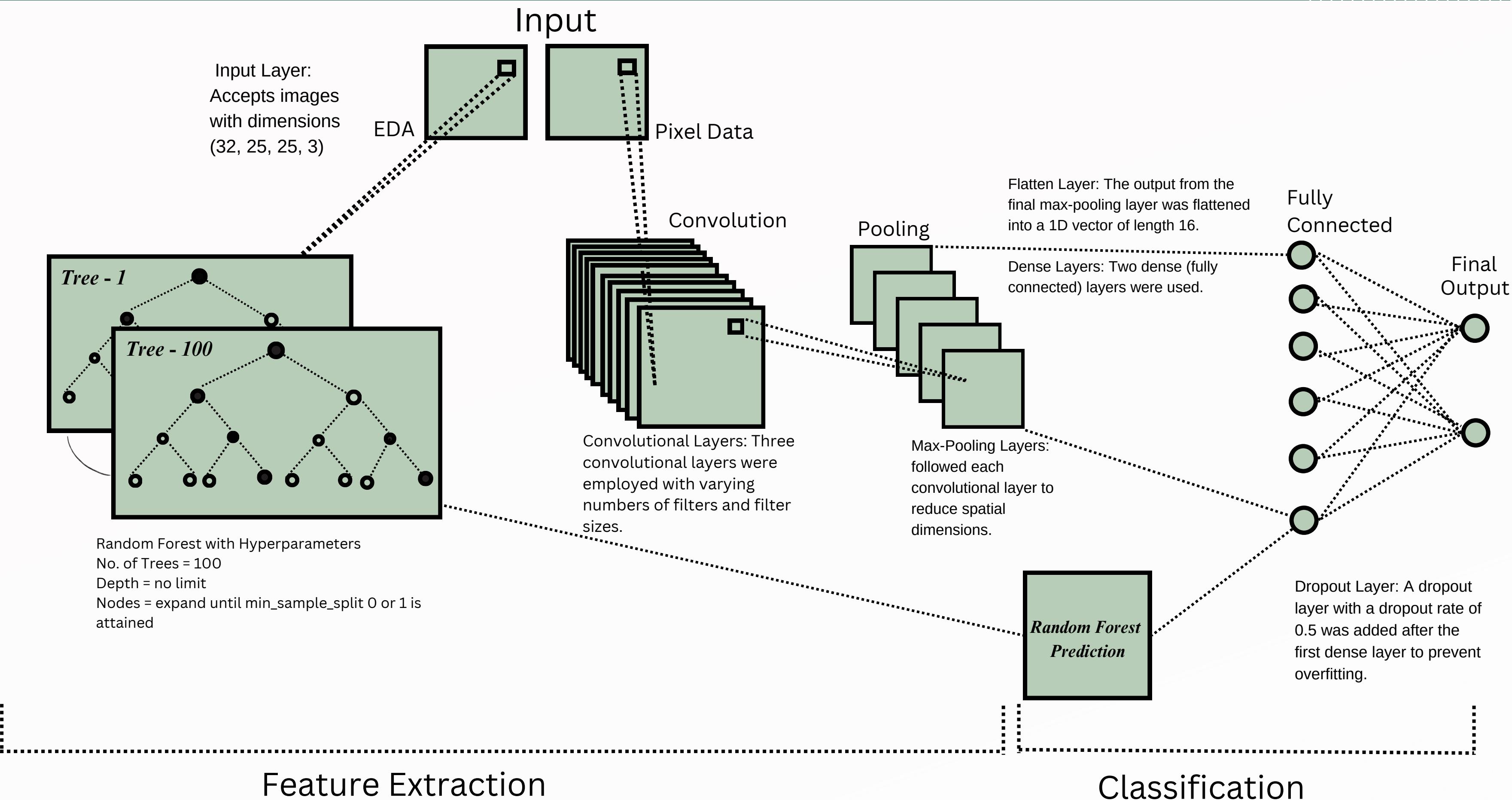
This graph shows which features of the dataset are significant in determining whether the image is infected or not infected.

Here we see that Mean and Max Blob size are less significant wrt to other features.

MODEL SELECTION

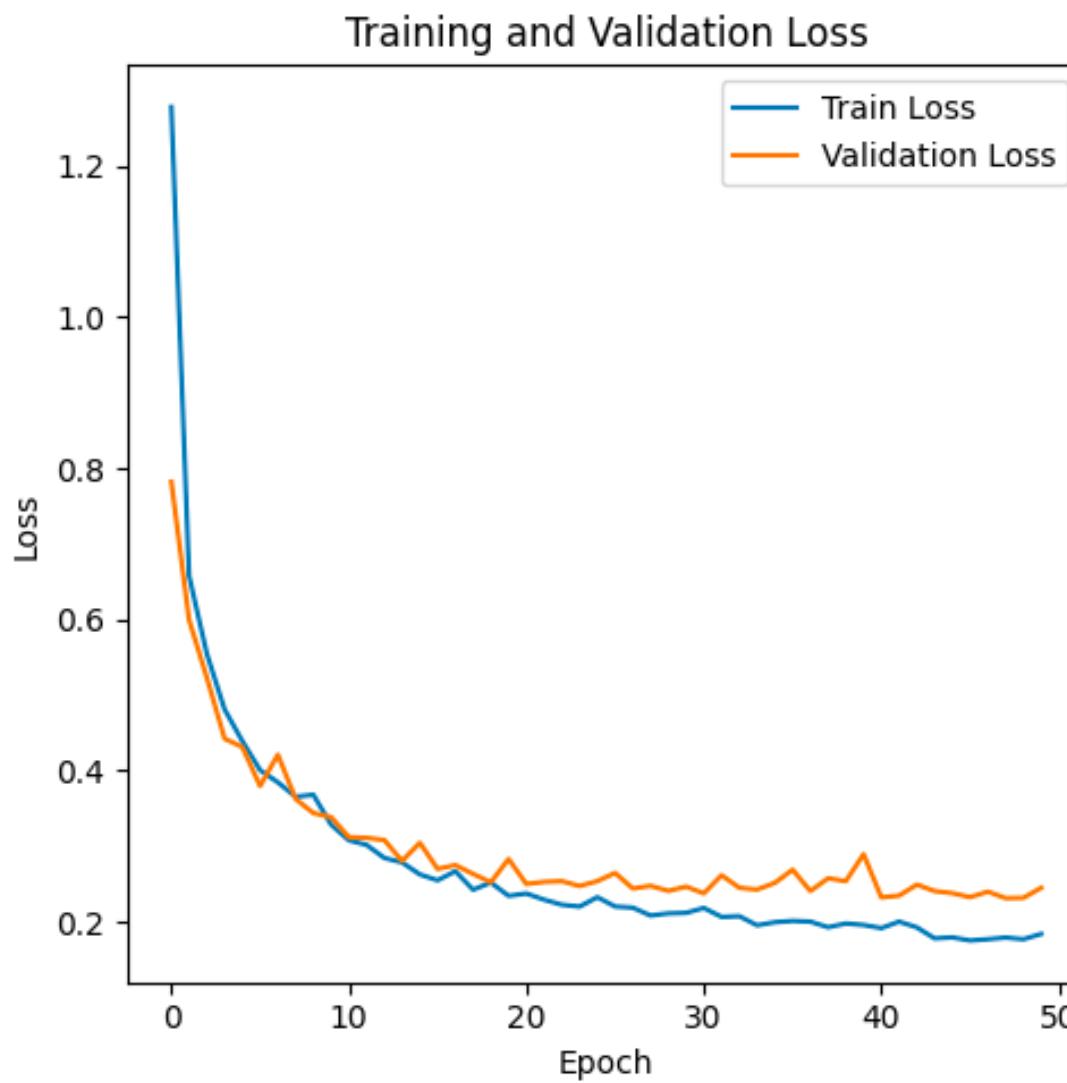
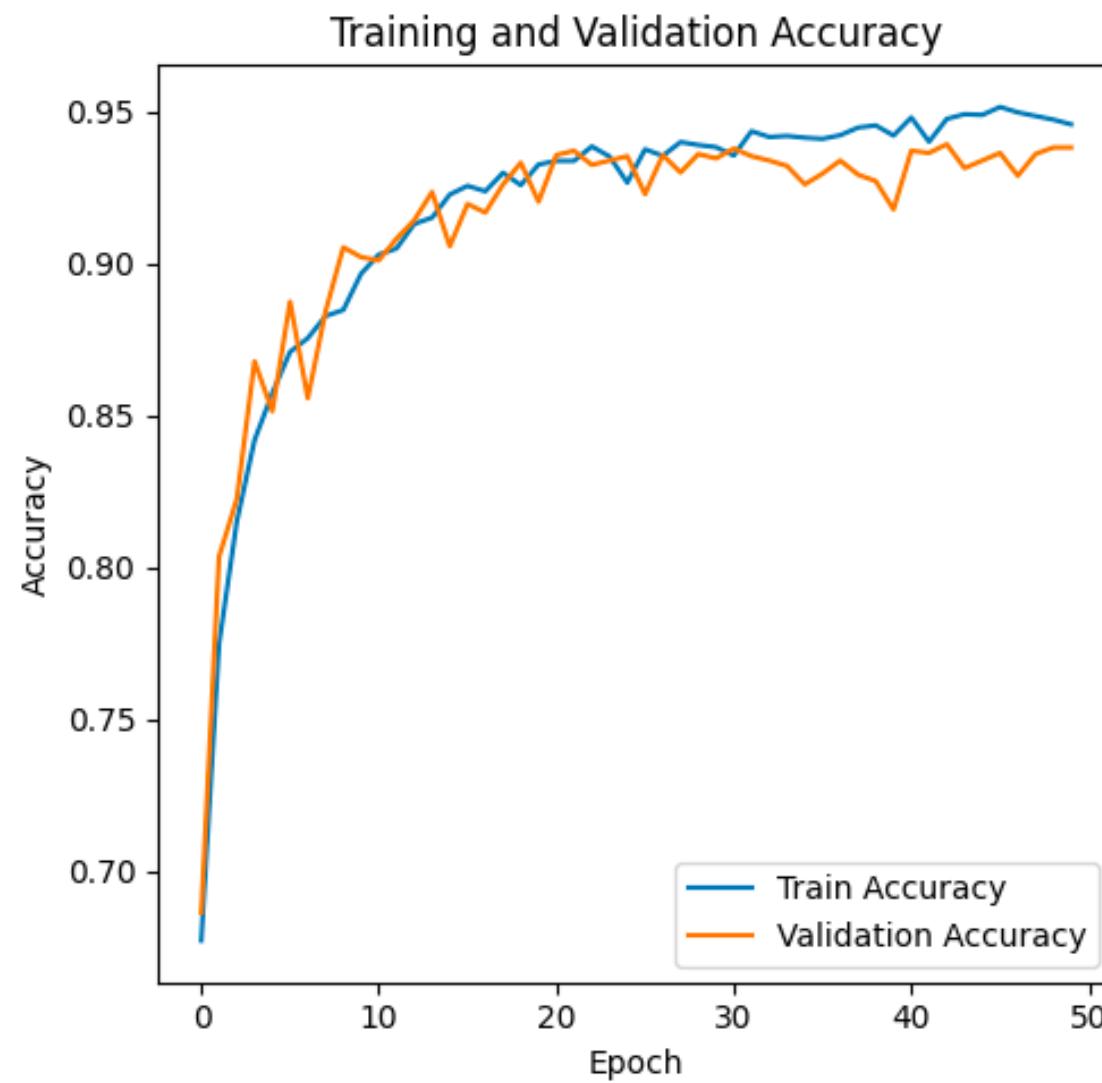


MODEL ARCHITECTURE



CNN + RF(Hyp)

ACCURACY



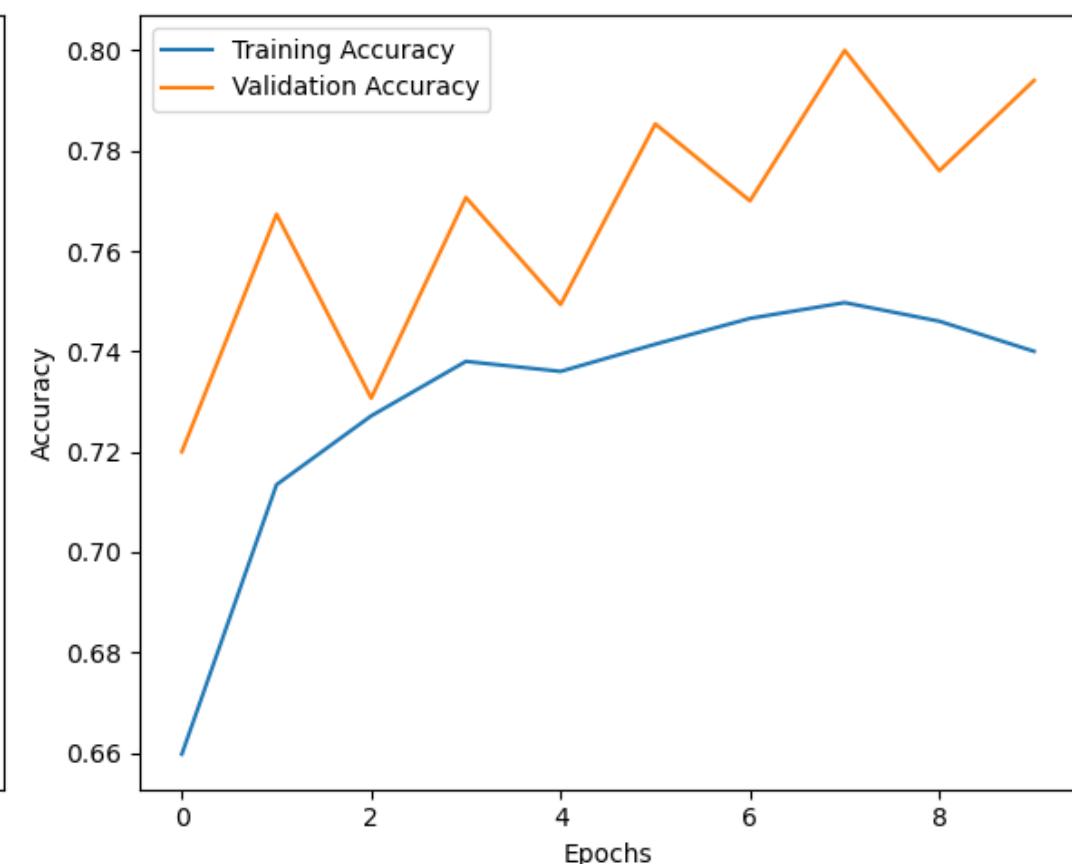
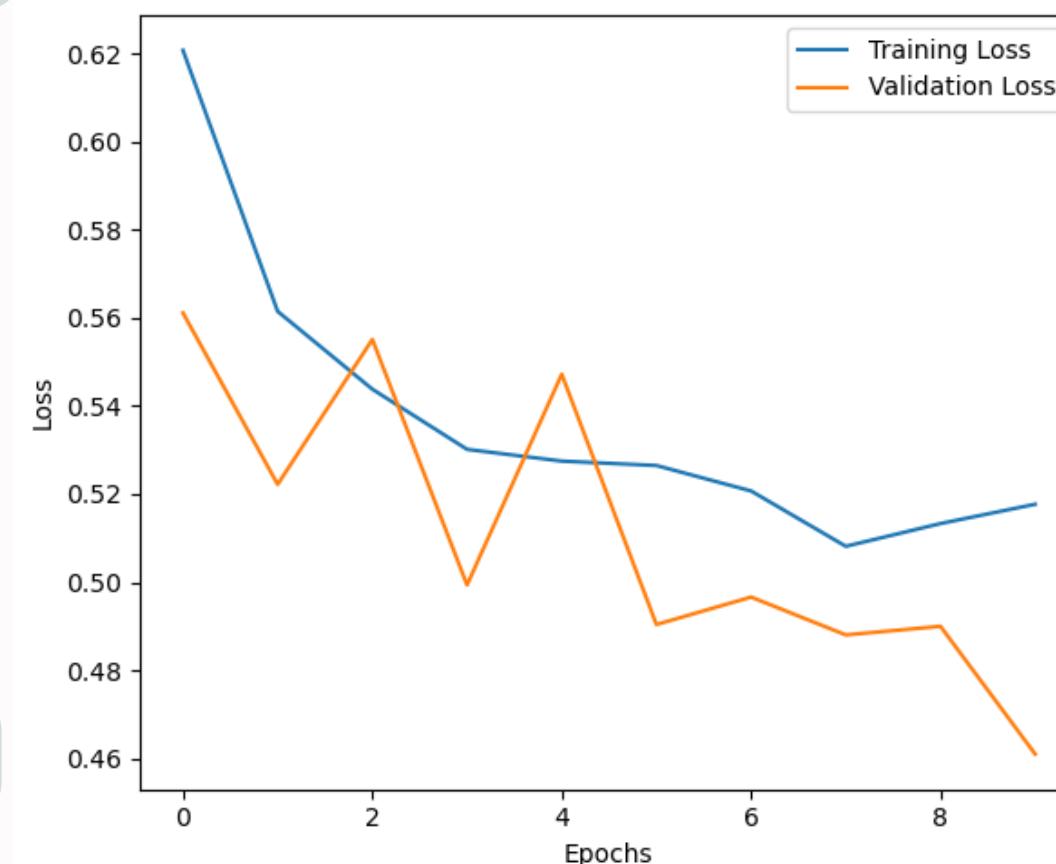
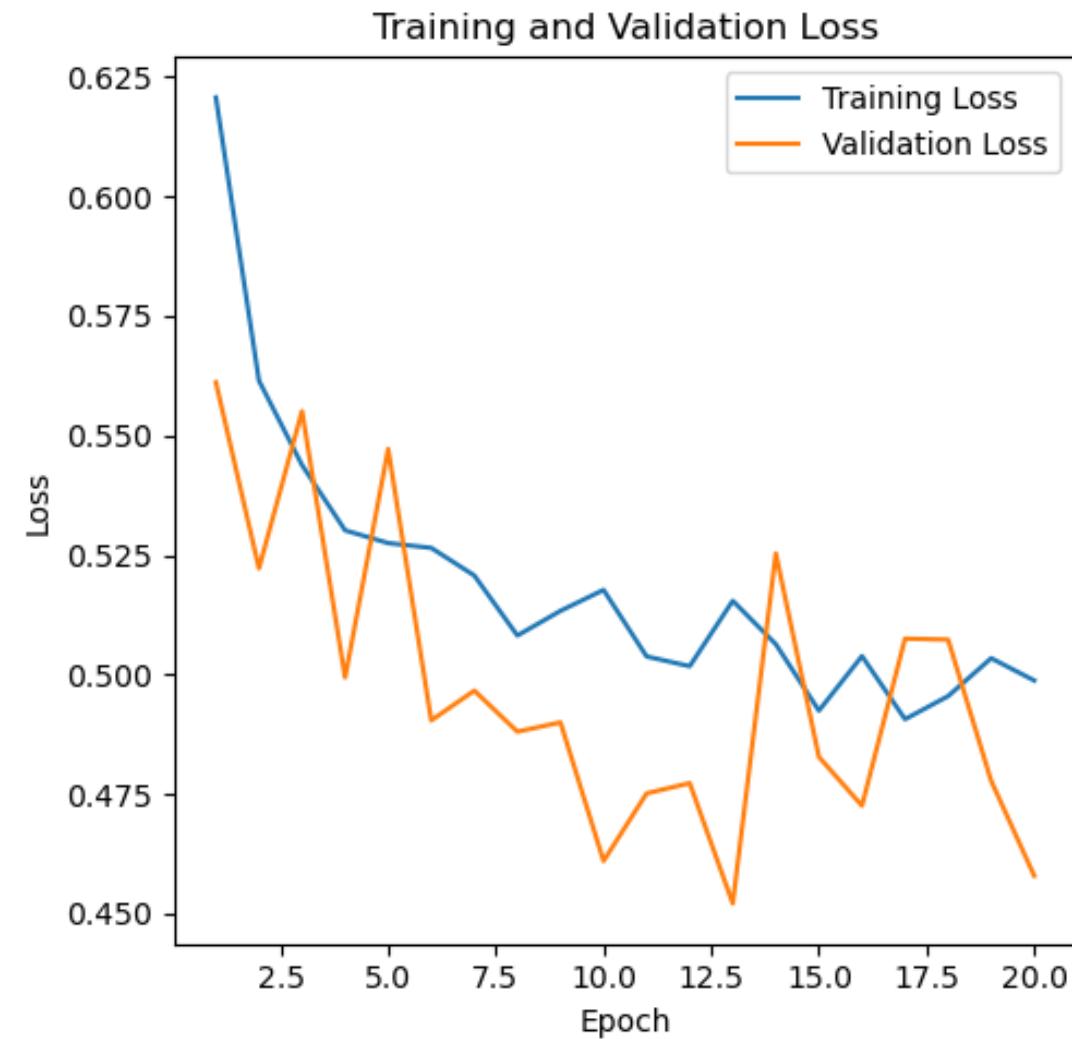
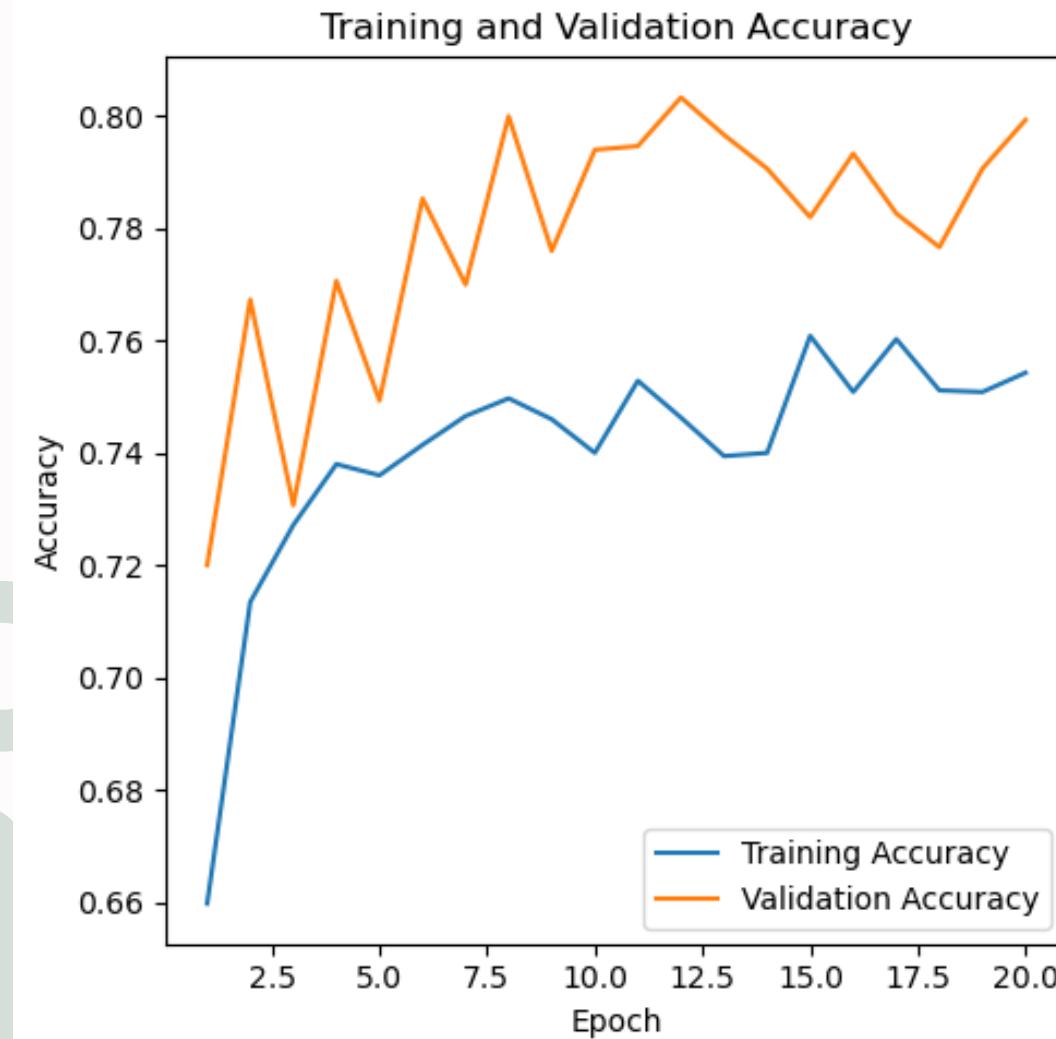
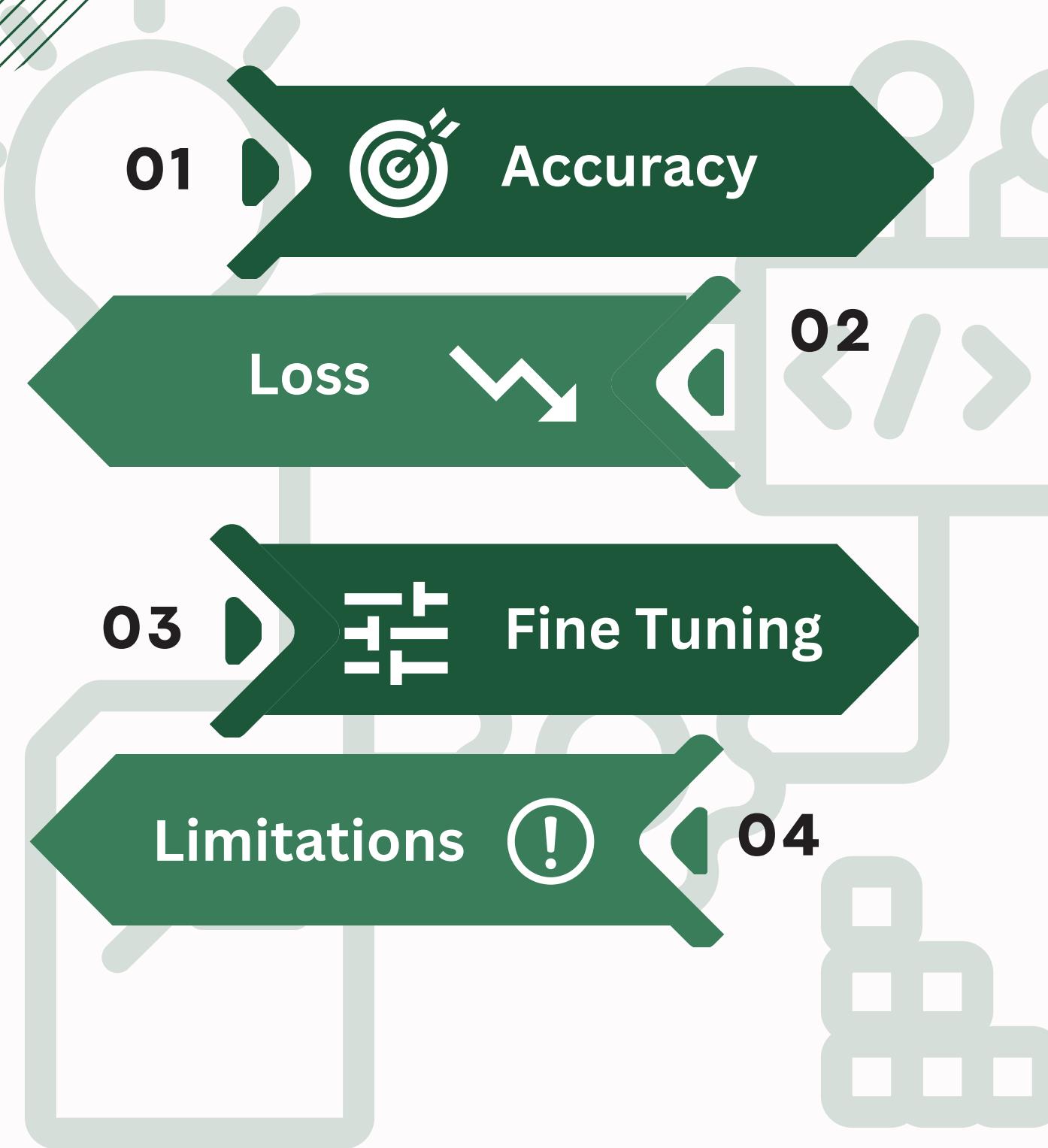
Accuracy with selected features: 0.89

	precision	recall	f1-score	support
0	0.88	0.90	0.89	750
1	0.90	0.88	0.89	750
accuracy			0.89	1500
macro avg	0.89	0.89	0.89	1500
weighted avg	0.89	0.89	0.89	1500

FINAL PREDICTIONS ->

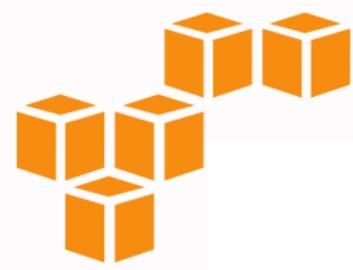
Effciency of predicting uninfected by both models is 96.13 %
Effciency of predicting infected by both models is 96.53 %

Xception



DATABASE

Connecting the ML model to our webpage!



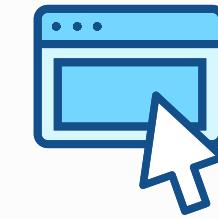
AWS & S3 Bucket

Using the cloud service from AWS and S3 storages to store our images, later on pulling these images using special API.



Python & SQLite

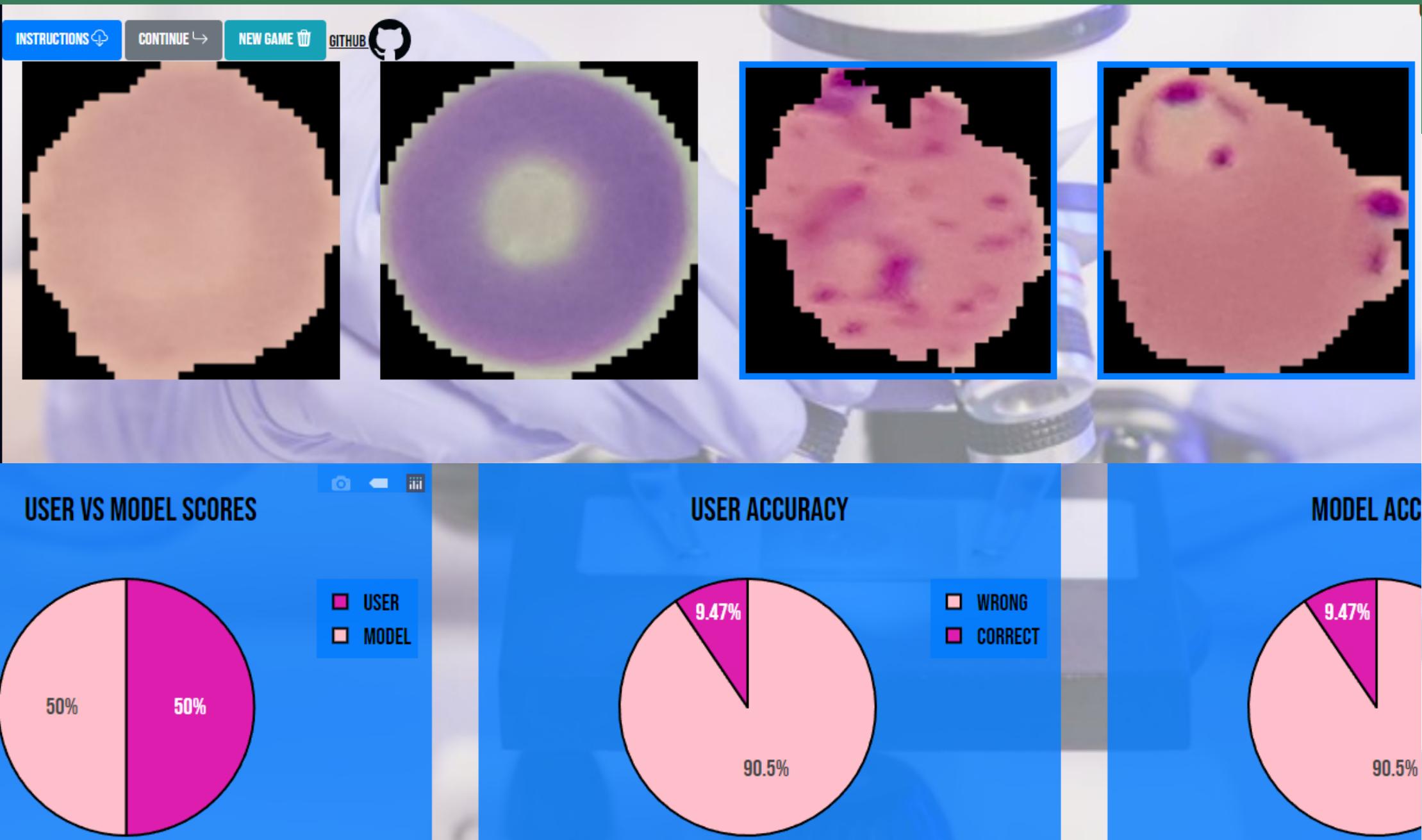
Using python with pandas to filter and create the structure for our database in SQLite, smoothing the process for our web app to read.



Python & Shutil

Reading through the data base to see which images will be needed to be stored on S3. Using shutil library to move files to folders for easier management.

Web Application



Frontend: HTML, CSS and JavaScript



Visualization: Plotly - pie charts



User Interaction

Users can **select** infected cells on the platform. Once they **submit** their selections, the data is sent to our backend for processing.



Styling & Fonts: Bootstrap and Google Fonts



AWS S3: Storage

LIMITATIONS



Limited
Diversity



Quality
Control



Varied
Image
Dimensions



Limited
Computing
Resources

CONCLUSION + REFLECTION



SIGNIFICANT
STEP
FORWARD



OPTIMIZED MODEL(CNN +RF)
ACCURACY: 94%- 96%



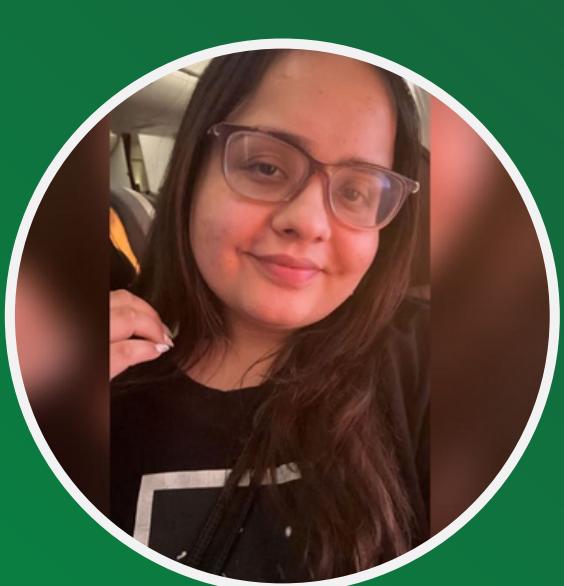
TECHNOLOGY'S IMPACT ON
REAL-WORLD HEALTH
CHALLENGES

Our Dream Team



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**Data Engineer/Analyst
Product Manager**

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[Github](#)



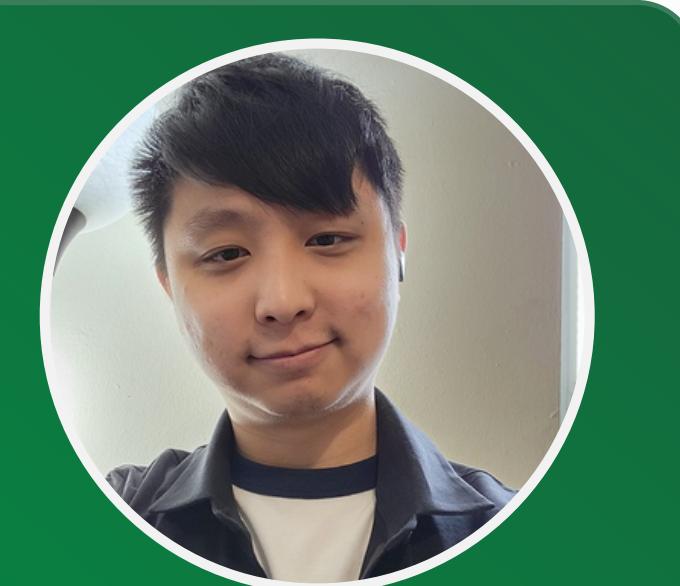
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Happy Graduation!



UNIVERSITY OF TORONTO
SCHOOL OF CONTINUING STUDIES

Thank You

Bharat Janarthanan
Anthony Inthavong
Mahesh Peiris
Nayan Saxena
Mike Gonzalez

Appendix



Data Source

<https://www.kaggle.com/datasets/iarunava/cell-images-for-detecting-malaria>

<https://ceb.nlm.nih.gov/repositories/malaria-datasets/>



Github

<https://github.com/jnliou/project4>

Tableau

<https://public.tableau.com/app/profile/julia.liou6123/viz/EDAonCellImagesofMalaria-Tableau/ExploratoryDataAnalysisonMalariaCellImages>



Project Management

<https://trello.com/b/ZZba331m/project-4>
https://miro.com/app/board/uXjVMgjTsg0/?share_link_id=288261535764

Appendix Cont'd



References

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- Team, K. (n.d.). Keras Documentation: Transfer Learning & Fine-tuning. https://keras.io/guides/transfer_learning/
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A close-up photograph of a compound light microscope. The eyepiece lens is labeled "OLYMPUS PlanFL N 4x /0.13 00/- /FN26.5". The objective lens is labeled "OLYMPUS PlanFL N 10x /1.30 Oil 0.17 /FN26.5". A small green leaf specimen is mounted on a glass slide, which is positioned on the microscope stage. The background is dark, making the metallic components of the microscope stand out.

DEMO TIME