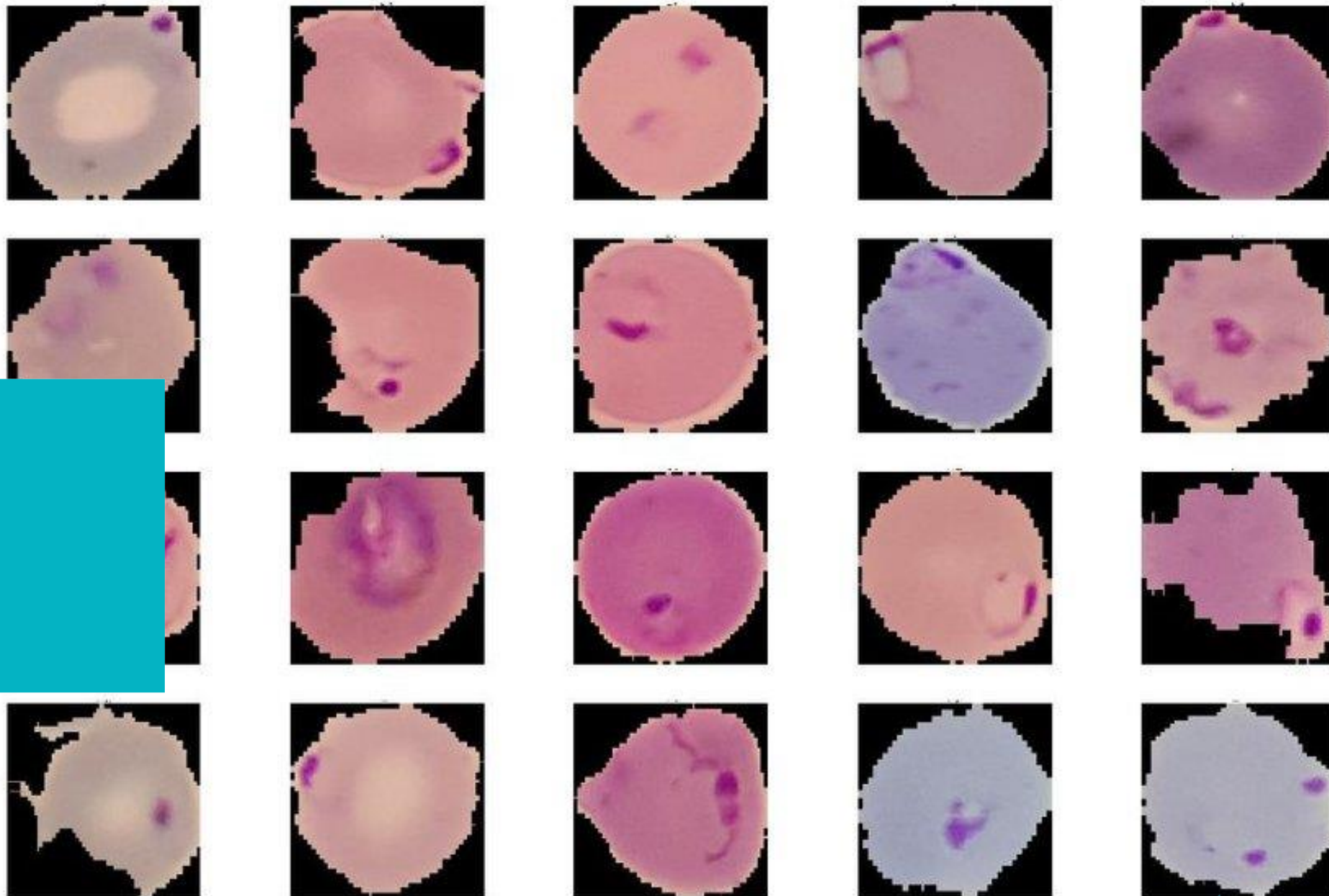
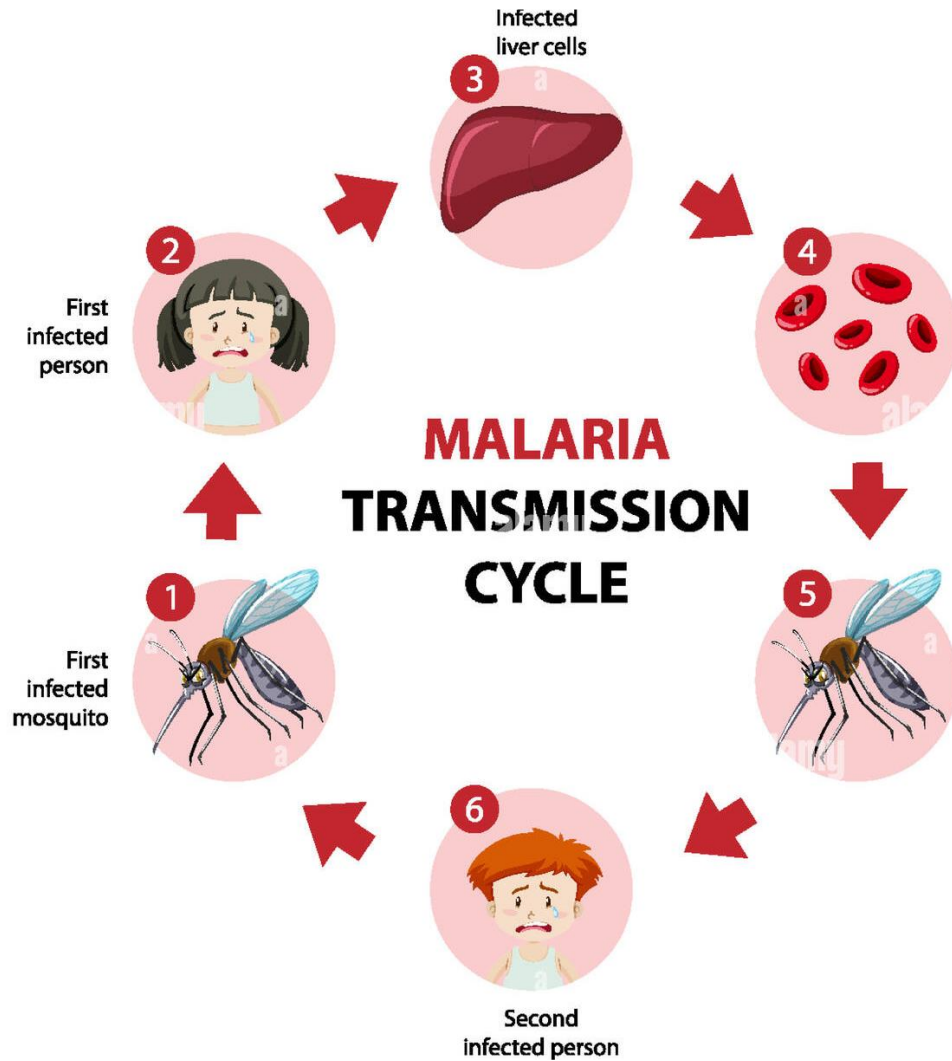


MALARIA DETECTION

Roula Krayem



INTRODUCTION



- ❑ What is Malaria?
- ❑ Malaria caused 627000 deaths in 2020
- ❑ Diagnoses methods including Blood Smear test
- ❑ Microscopic images of Red Blood Cells (RBCs)

PROBLEM DEFINITION

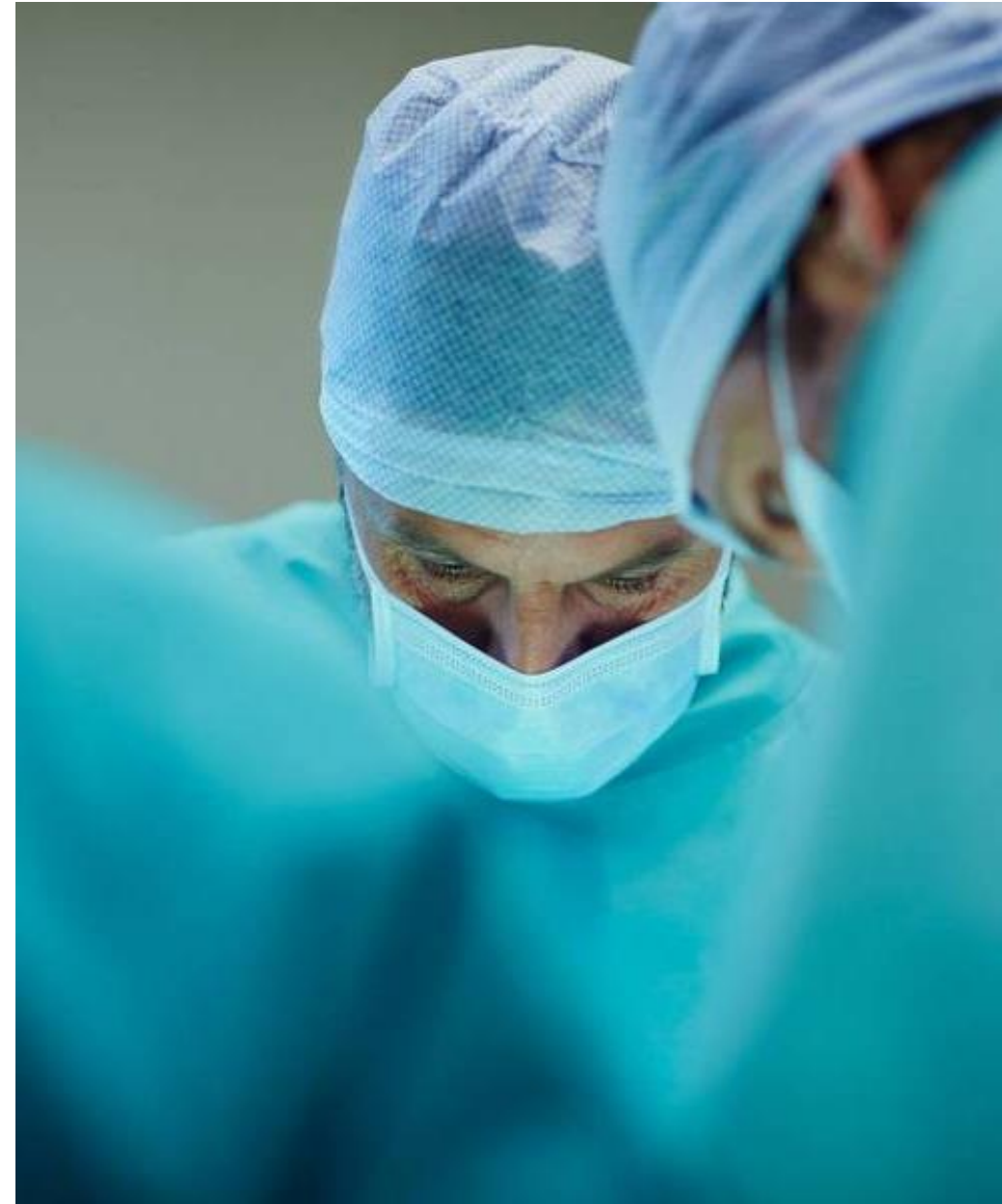
Millions of Malaria cases and hundreds of thousands of deaths annually



Providers study each RBCs sample individually under the microscope



Slow process that could delay treatment



SOLUTION

Build a Neural Network model to expedite and facilitate Malaria detection

STEP 1

Perform data
Exploration and note
key takeaways

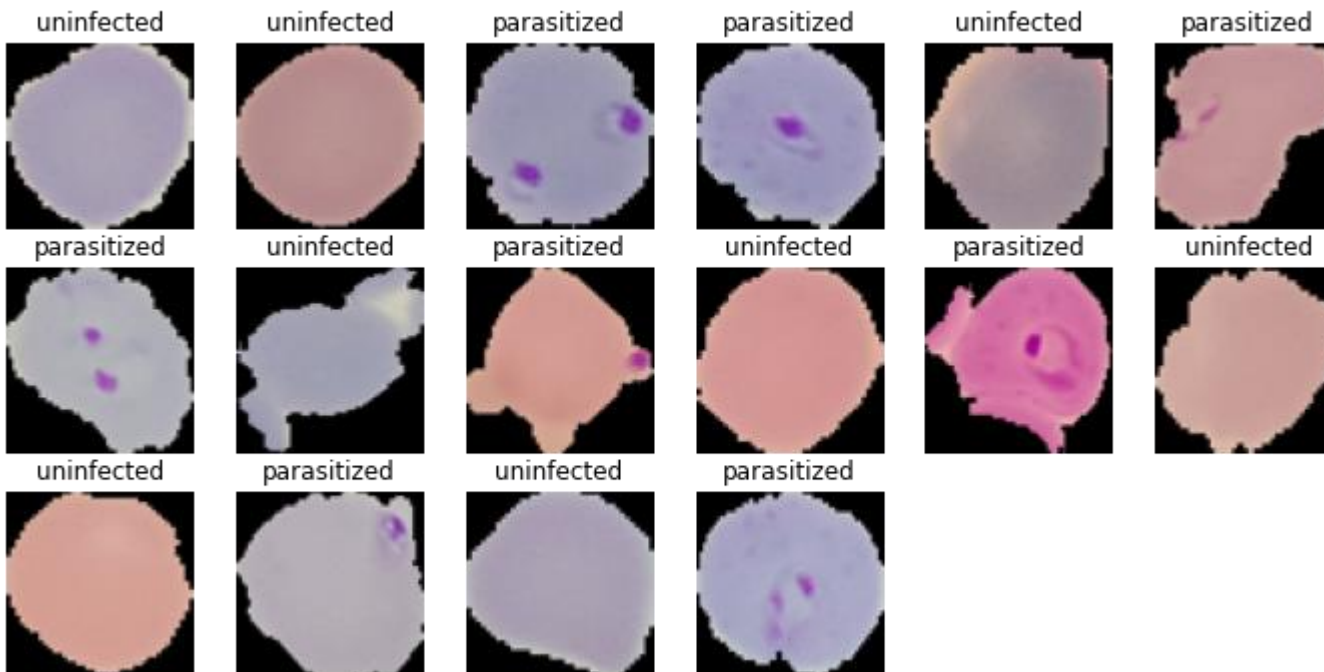
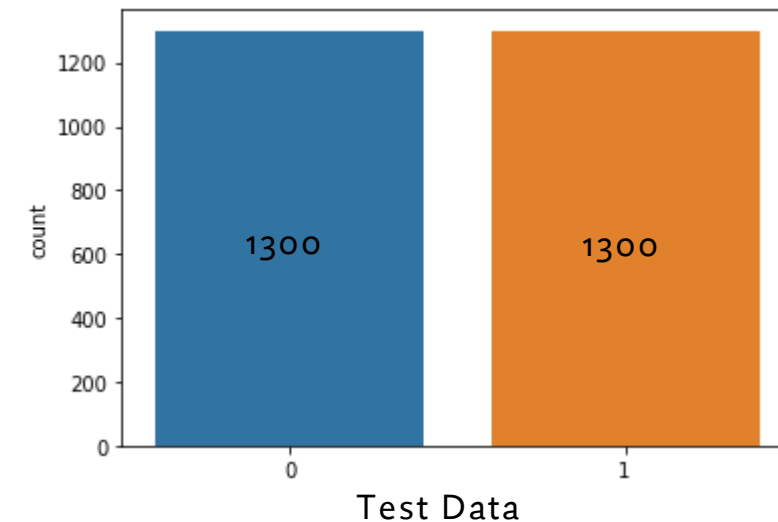
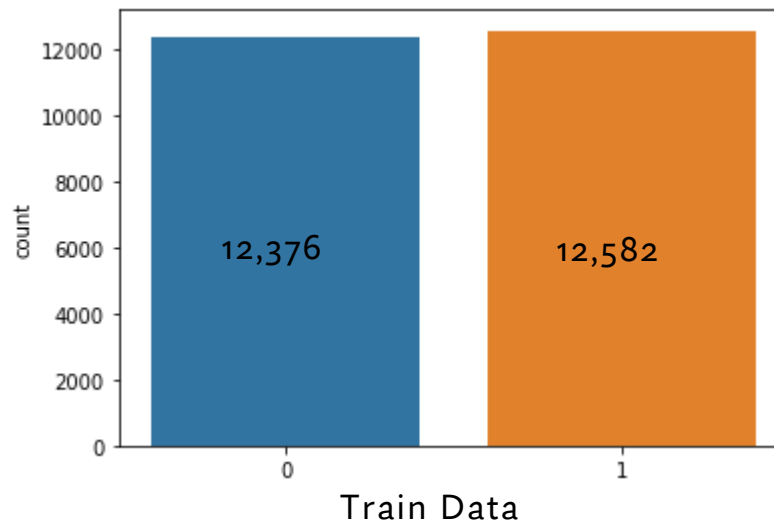
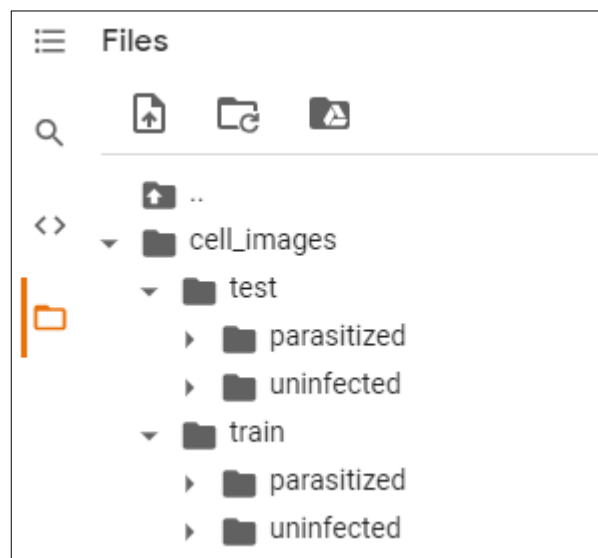
STEP 2

Build and test multiple
Neural Network
models to reach the
best performing
Model

STEP 3

Select the best
performing model and
provide
recommendations

DATA EXPLORATION



DATA PROCESSING TECHNIQUES

- ✓ Image resizing (64 x 64)
- ✓ Normalization
- ✓ One-Hot Encoding

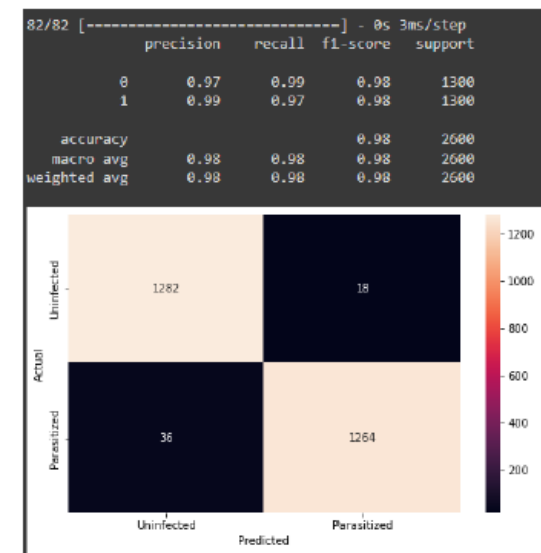
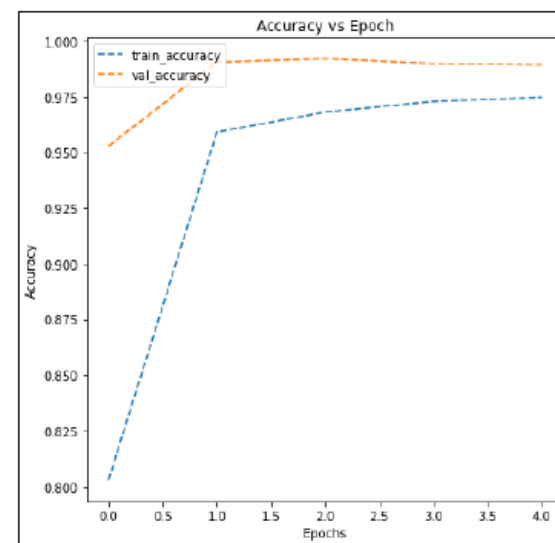


MODEL CONVERSATION

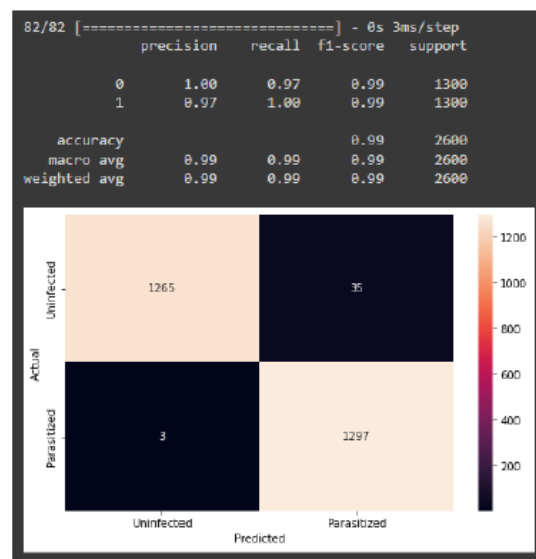
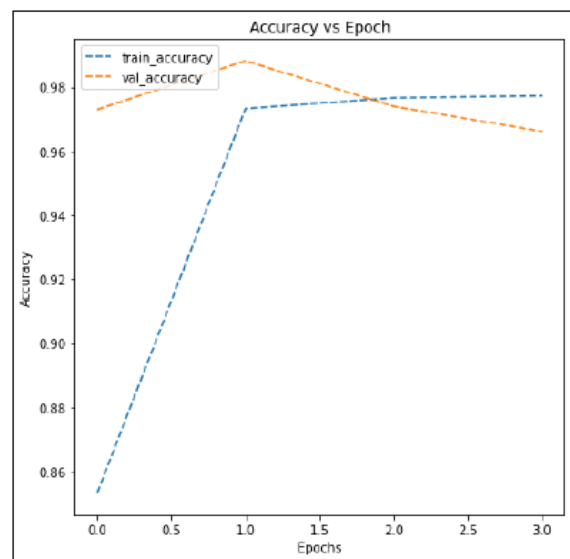


FROM SCRATCH

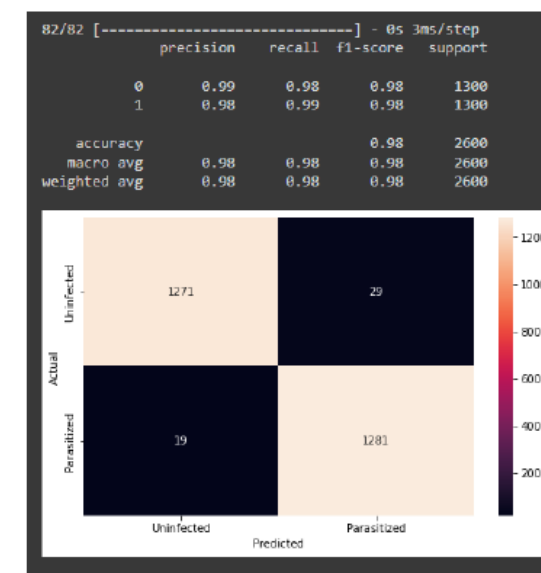
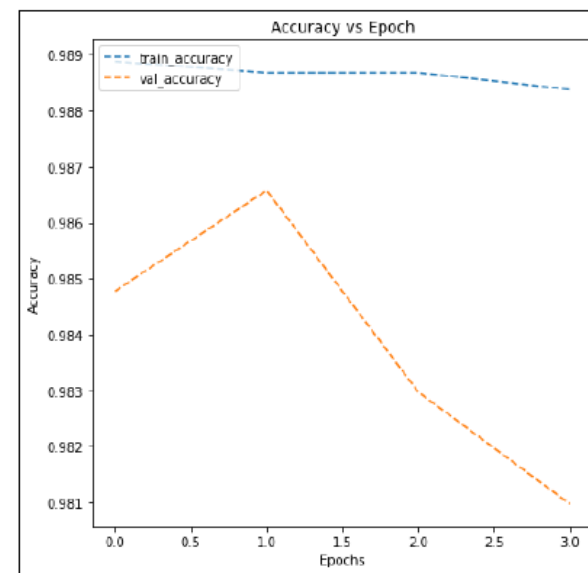
Base Model

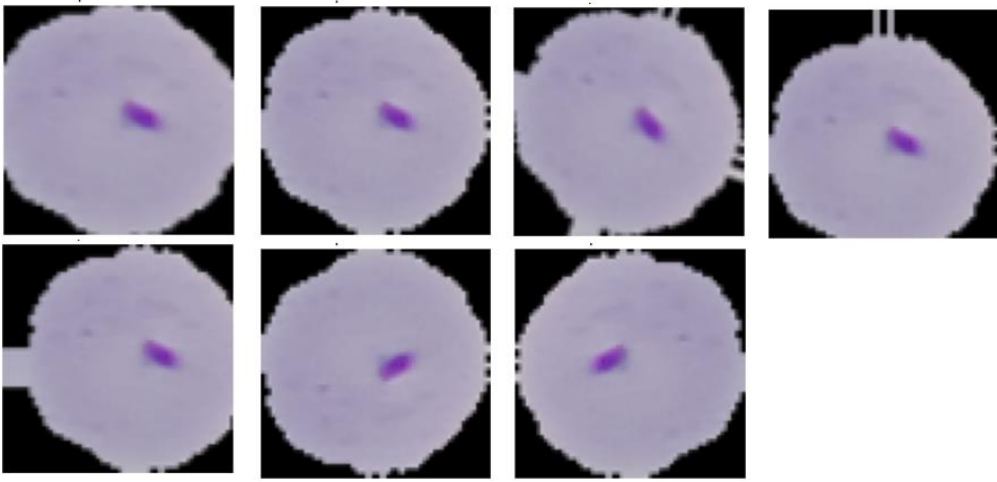


First Model



Second Model



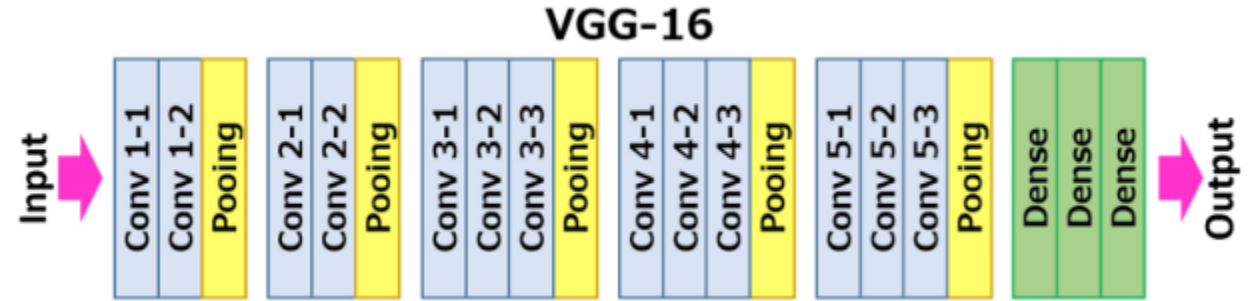


DATA AUGMENTATION

Data Image Process	Precision Unin*	Precision Para*	Recall Unin*	Recall Para*	F1 Score Unin*	F1 Score Para*	Accuracy	Misclassified Para*	Misclassified Unin*
Zoom	0.97	0.99	0.99	0.97	0.98	0.98	98%	37	17
Shear	0.98	0.99	0.99	0.98	0.98	0.98	98%	30	15
Rotation	0.98	0.98	0.98	0.98	0.98	0.98	98%	22	21
Width Shift	0.97	0.99	0.99	0.97	0.98	0.98	98%	39	11
Height Shift	0.99	0.98	0.98	0.99	0.98	0.98	98%	14	31
Vertical Flip	0.97	0.99	0.99	0.97	0.98	0.98	98%	38	15
Horizontal Flip	0.98	0.99	0.99	0.98	0.99	0.98	98%	29	10
All	0.98	0.99	0.99	0.98	0.98	0.98	98%	28	15

*Para= Parasitized and Unin = Uninfected

VGG16



Learning rate / batch size	Precision Unin*	Precision Para*	Recall Unin*	Recall Para*	F1 Score Unin*	F1 Score Para*	Accuracy	Misclassified Para*	Misclassified Unin*
0.001/32	0.98	0.84	0.82	0.98	0.89	0.91	90%	24	238
0.00001/23	0.96	0.89	0.89	0.96	0.92	0.93	92%	50	147
0.00001/65	0.95	0.91	0.91	0.95	0.93	0.93	93%	66	123
0.0000/100	0.96	0.89	0.88	0.96	0.92	0.93	92%	48	154

*Para= Parasitized and Unin = Uninfected

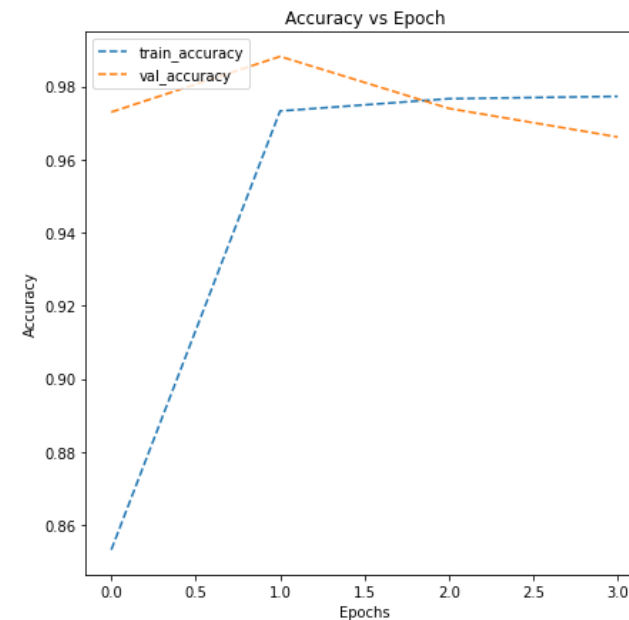
BEST MODEL

MODEL (1)

Model: "sequential"

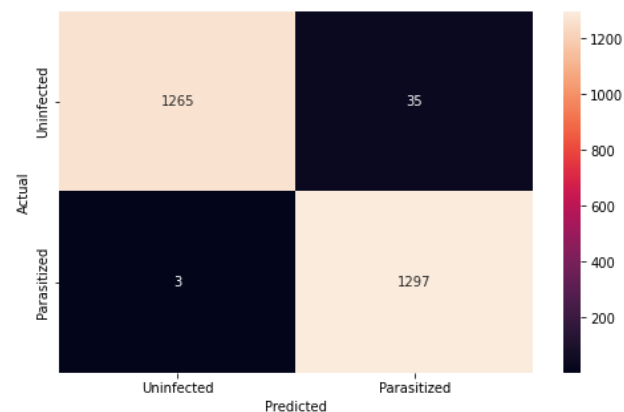
Layer (type)	Output Shape	Param #
conv2d (Conv2D)	(None, 64, 64, 32)	416
max_pooling2d (MaxPooling2D)	(None, 32, 32, 32)	0
conv2d_1 (Conv2D)	(None, 32, 32, 32)	4128
max_pooling2d_1 (MaxPooling2D)	(None, 16, 16, 32)	0
dropout (Dropout)	(None, 16, 16, 32)	0
conv2d_2 (Conv2D)	(None, 16, 16, 32)	4128
max_pooling2d_2 (MaxPooling2D)	(None, 8, 8, 32)	0
dropout_1 (Dropout)	(None, 8, 8, 32)	0
conv2d_3 (Conv2D)	(None, 8, 8, 32)	4128
max_pooling2d_3 (MaxPooling2D)	(None, 4, 4, 32)	0
conv2d_4 (Conv2D)	(None, 4, 4, 32)	4128
max_pooling2d_4 (MaxPooling2D)	(None, 2, 2, 32)	0
dropout_2 (Dropout)	(None, 2, 2, 32)	0
flatten (Flatten)	(None, 128)	0
dense (Dense)	(None, 512)	66048
dropout_3 (Dropout)	(None, 512)	0
dense_1 (Dense)	(None, 2)	1026

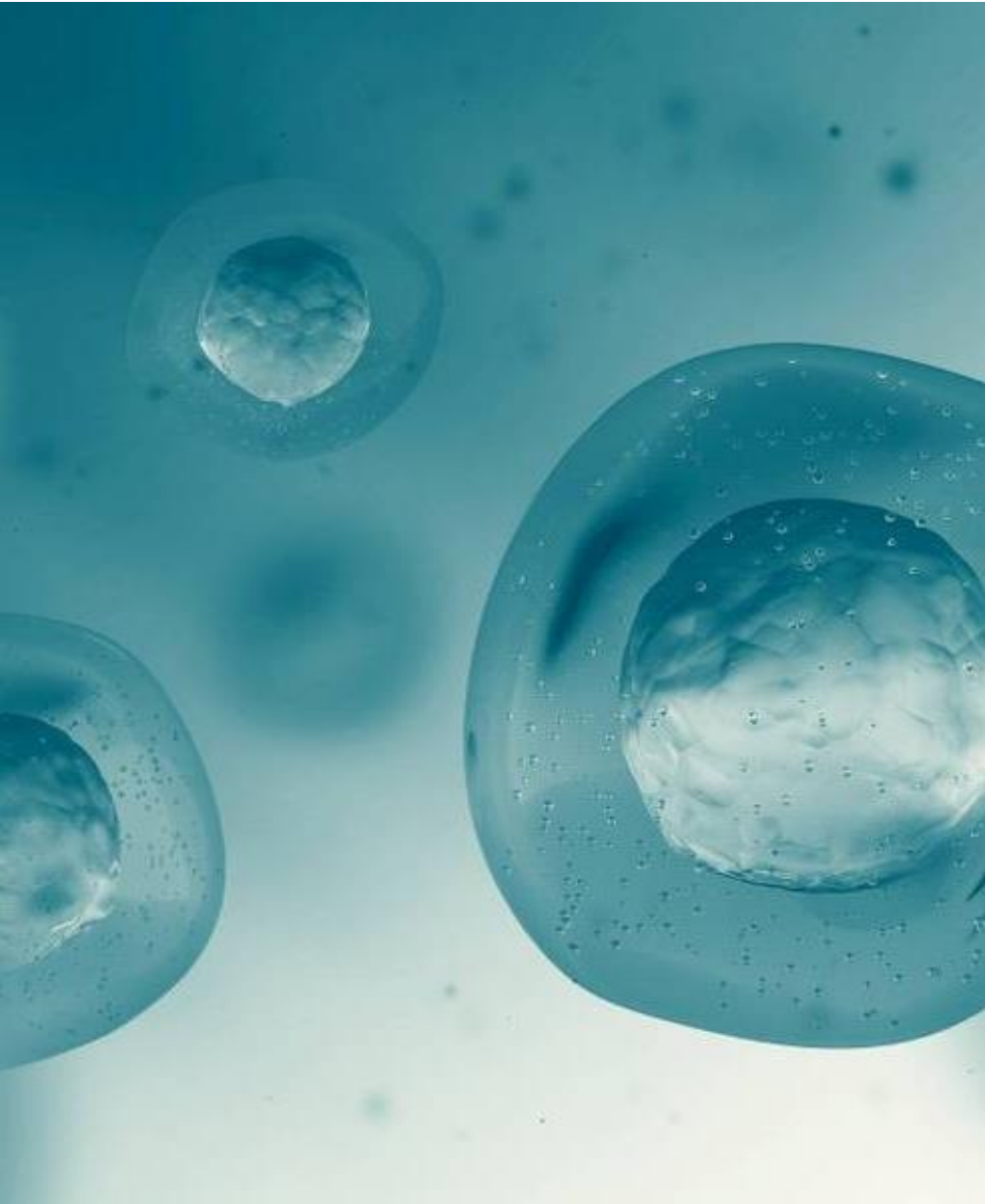
=====
 Total params: 84,002
 Trainable params: 84,002
 Non-trainable params: 0



82/82 [=====] - 0s 3ms/step

	precision	recall	f1-score	support
0	1.00	0.97	0.99	1300
1	0.97	1.00	0.99	1300
accuracy			0.99	2600
macro avg	0.99	0.99	0.99	2600
weighted avg	0.99	0.99	0.99	2600





RECOMMENDATIONS

MODEL FUTURE IMPROVEMENTS

1. Try different learning rates and batch-size when fitting the second model using the data image generator.
2. Use the Ensemble technique with multiple weak models to improve the results.

POLICYMAKERS

1. Adopting the first model to automate and accelerate Malaria detection.
2. Requiring a revision of the images classified as uninfected by a physician before making the final clinical decision.



Thank you!
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