

By: Sameeha Ramadhan

## **Table of Contents**

01 Introduction

The Goal of the Project

04 Modeling

The Models & Evaluations

02 What is Skin Cancer?

A Brief Description

05 Conclusion

Summary & Recommendations

03 The Data

Obtaining & Preprocessing







## **Business Problem:**

To reduce diagnosis timings for doctors by building an algorithm that can read images of skin lesions and determine if it is malignant (cancerous) or benign (not).



## What is Skin Cancer?



## 2 Main Layers

Epidermis (outer) & dermis (inner) skin



#### Basal

Most common



#### **Epidermis**

Is where skin cancer starts



#### Squamous

2nd, and can be cured like basal though costly



#### **UV Rays**

From the sun and tanning beds is the most common cause



#### Melanoma

3rd most common but deadliest



## The Data

1 Sourced

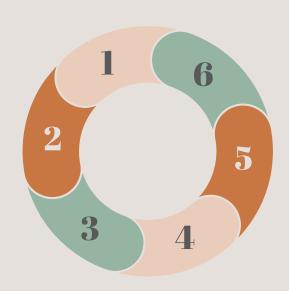
From dermascopy.org, cancer.com, & more

2 Size

Contains over 28,000 images of skin lesions

3 Structure

Contains 2 subsets "train" & "test", each with "benign" & "malignant" folders



**Distribution** 

Is balanced w/ a near even distribution

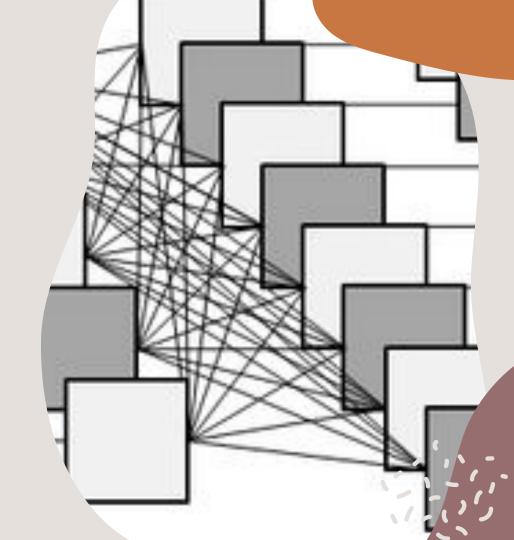
Augmentation

4 of the 5 models were fed augmented data

**Normalization** 

Images were scaled to 256 colors (0-255)

## 04 Modeling The Models & Evaluation



# Convolutional Neural Network (CNN)

#### 4 Final Models

Out of 20 tested

#### **Pre-trained**

EfficientNet-BO and ResNet50 were included. The latter performed the best.

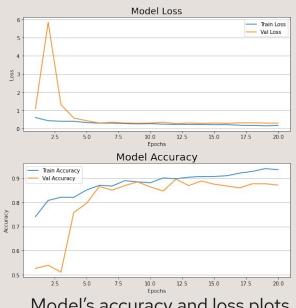
## **30 Epochs**

Each has at least 30 epochs

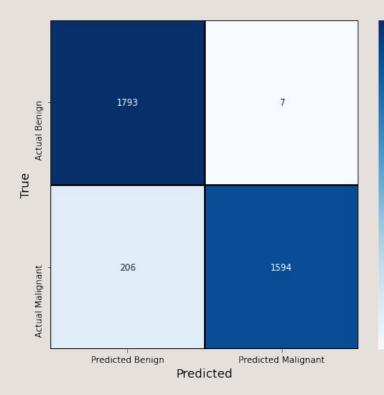
```
▶ base model= ResNet50(include top=False, weights="imagenet", input shape=(height, width,3))
model4= Sequential()
model4.add(base model)
model4.add(Conv2D(64, (3, 3), activation = 'relu'))
model4.add(Dropout(0.2))
model4.add(Conv2D(64, (3, 3), activation = 'relu'))
model4.add(MaxPooling2D(pool size = (2, 2)))
model4.add(Dropout(0.2))
model4.add(Flatten())
model4.add(Dense(512,activation='relu'))
model4.add(Dense(256,activation='relu'))
model4.add(Dropout(0.2))
model4.add(Dense(128,activation='relu'))
model4.add(Dense(64,activation='relu'))
model4.add(Dense(32,activation='relu'))
model4.add(Dense(16,activation='relu'))
model4.add(Dense(8,activation='relu'))
model4.add(Dense(2, activation='softmax'))
model4.summary()
```

ResNet50 model architecture

## **Model Evaluation**



Model's accuracy and loss plots



1600

1400

- 1200

- 1000

- 800

600

- 400

- 200

1 out of 9 misdiagnosed as healthy 1 in 257 misdiagnosed as cancer

## **Conclusion**

This project has shown how to classify benign or malignant diagnosis' from a skin lesion image. This tool can assist the medical industry and patients in providing quicker diagnosis.

## Recommendations

- Re-run the models on larger datasets.
- Re-run some of the models with a greater number of epochs (such as 100 or more on the ones with 30)
- Fine tune and test other parameters to reduce overfitting
- Build models for more complex problems, such as determining the types of cancers, skin diseases related to Agent Orange and more.
- Output the model to a user friendly application, preferably a web app.



shramadhan@gmail.com GitHub: @samtuleen linkedin.com/in/sameeha-ramadhan-3a1bba140