

# **Deep Learning Techniques for Breast Cancer Risk Prediction using IBM Cloud**

## **1. INTRODUCTION**

### **1.1 Overview**

Breast cancer is one of the main causes of cancer death worldwide. Computer-aided diagnosis systems showed the potential for improving diagnostic accuracy. But early detection and prevention can significantly reduce the chances of death. It is important to detect breast cancer as early as possible.

### **1.2 Purpose**

The goal is to classify images into two classifications of malignant and benign. As early diagnostics significantly increases the chances of correct treatment and survival. In this application, we are helping the doctors and patients to classify the Type of Tumour for the specific image given with the help of Neural Networks.

## **2. LITERATURE SURVEY**

### **2.1 Existing problem**

One of the main causes of cancer death worldwide is Breast Cancer. Computer-aided diagnosis systems showed the potential for improving diagnostic accuracy. But early detection and prevention can significantly reduce the chances of death. It is important to detect breast cancer as early as possible.

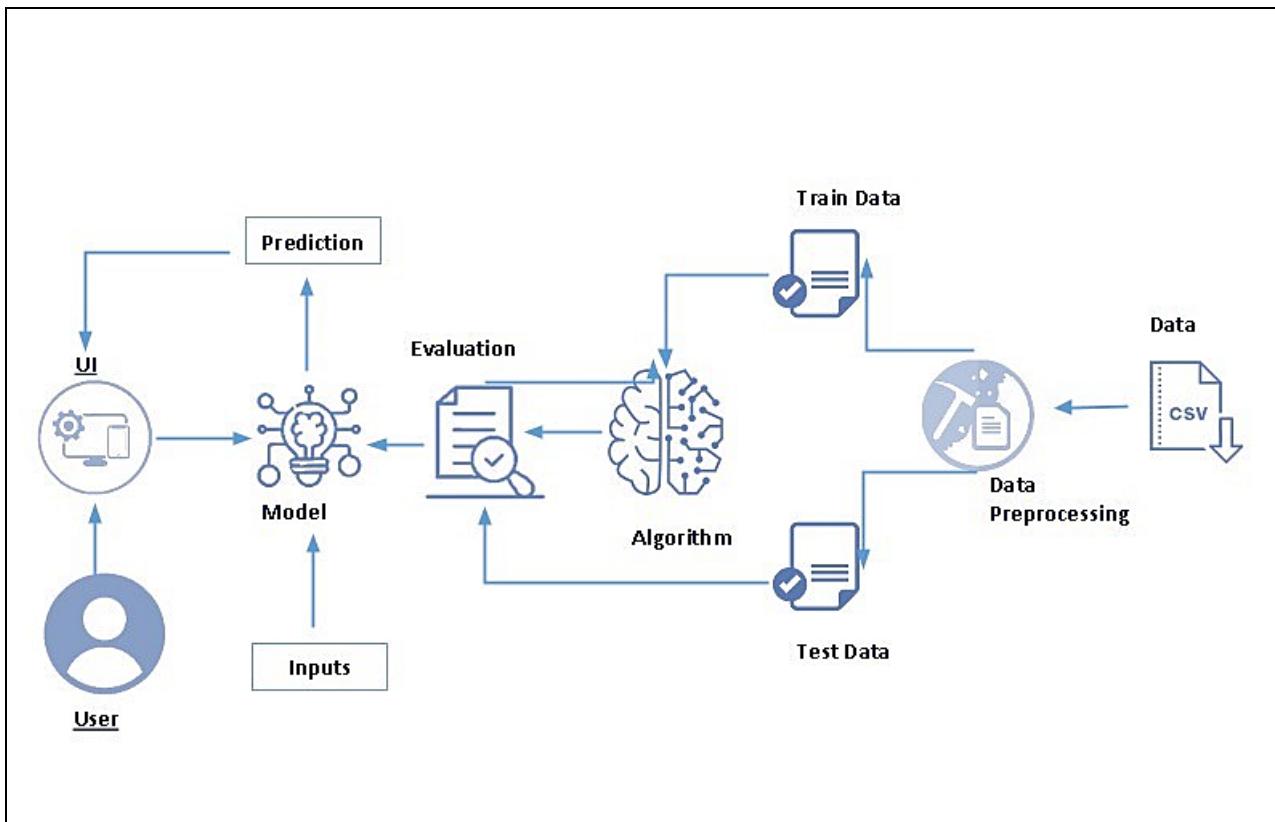
### **2.2 Proposed Solution**

This project goal is to classify images into two classifications of malignant and benign. As early diagnostics significantly increases the chances of correct treatment and

survival. In this application, we are helping the doctors and patients to classify the Type of Tumour for the specific image given with the help of Neural Networks.

### 3. THEORITICAL ANALYSIS

#### 3.1 Block Diagram



#### 3.2 Hardware / Software designing

##### Software Requirements:

- Ananconda Navigator
- Tensor flow
- Keras
- Flask

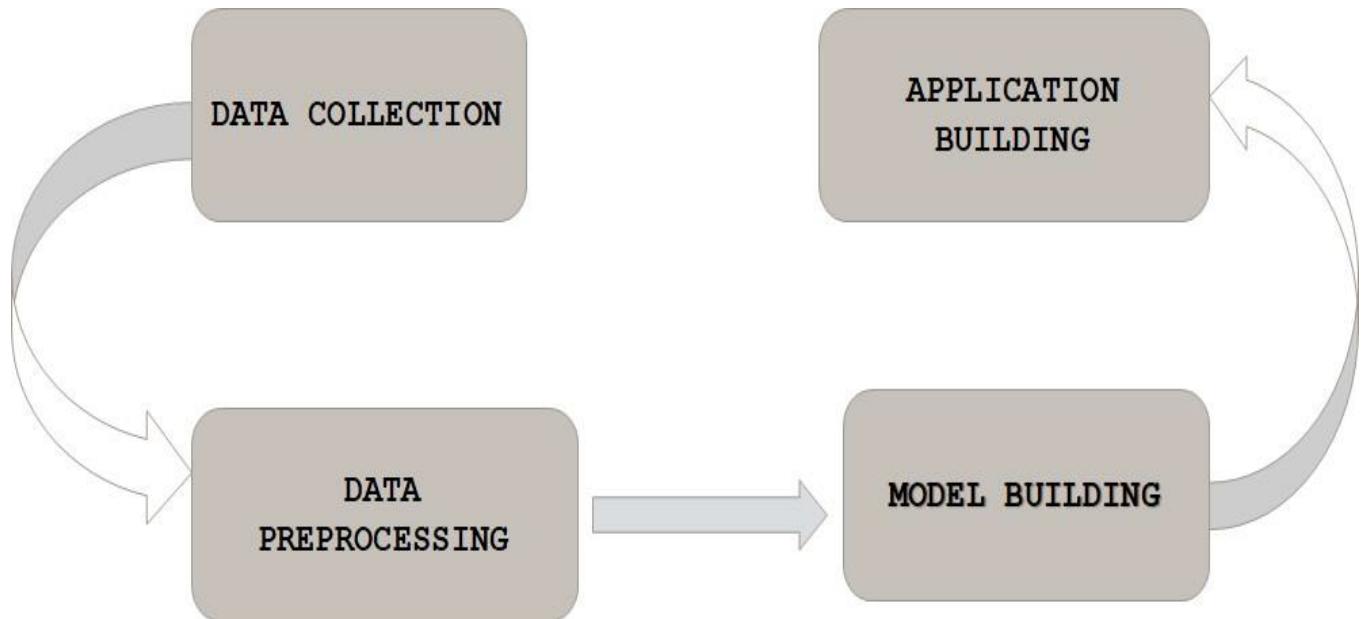
### **Hardware Requirements:**

- Processor : Intel Core i3
- Hard Disk Space : Min 100GB
- Ram : 4GB
- Display : 14.1 "Color Monitor(LCD, CRT or LED)
- Clock Speed : 1.67 GHz

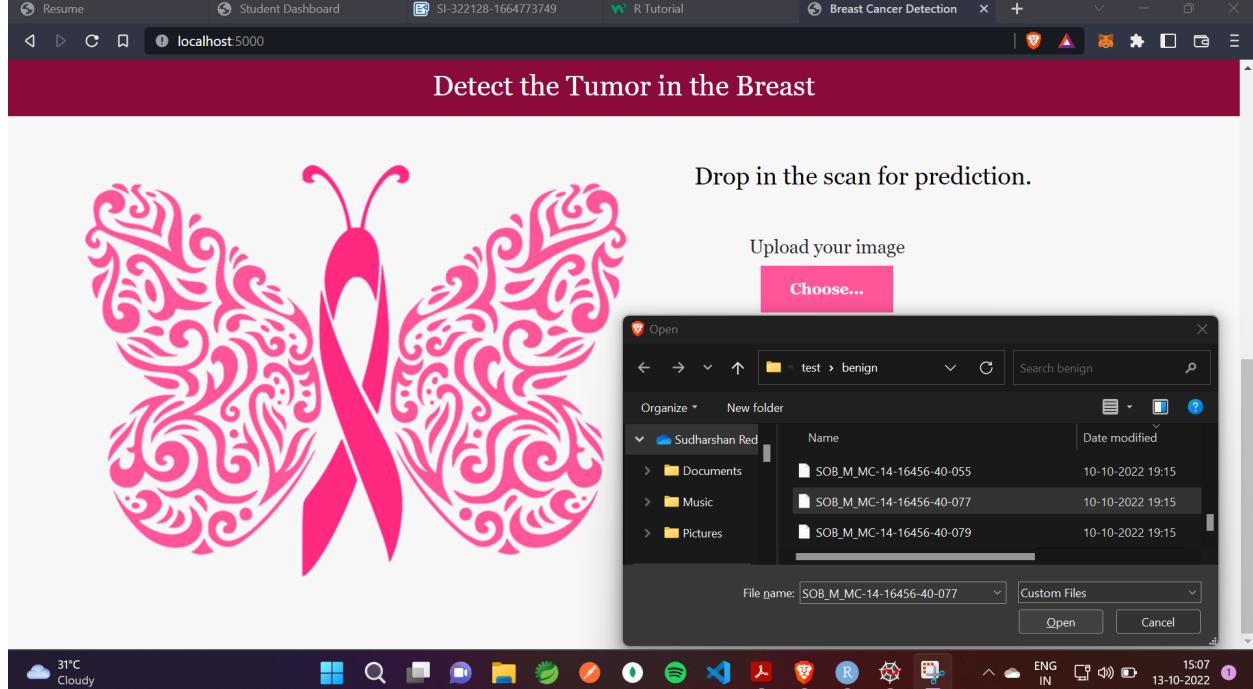
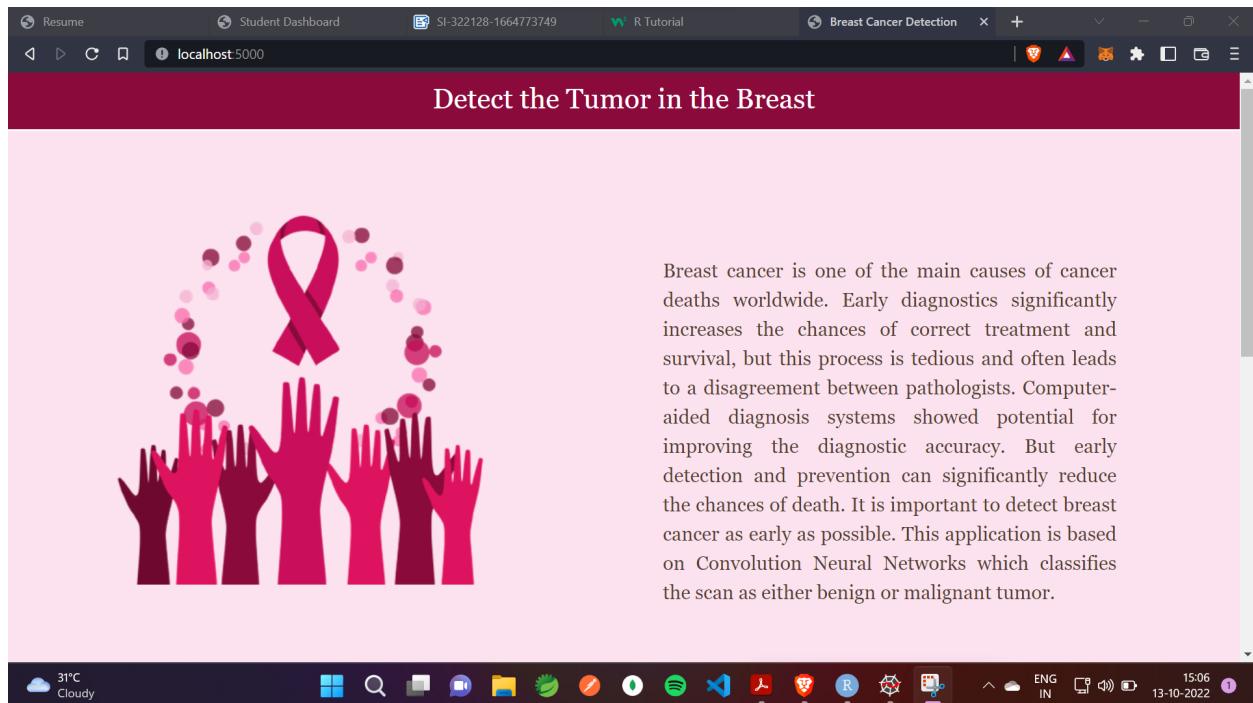
## **4. EXPERIMENTAL INVESTIGATIONS**

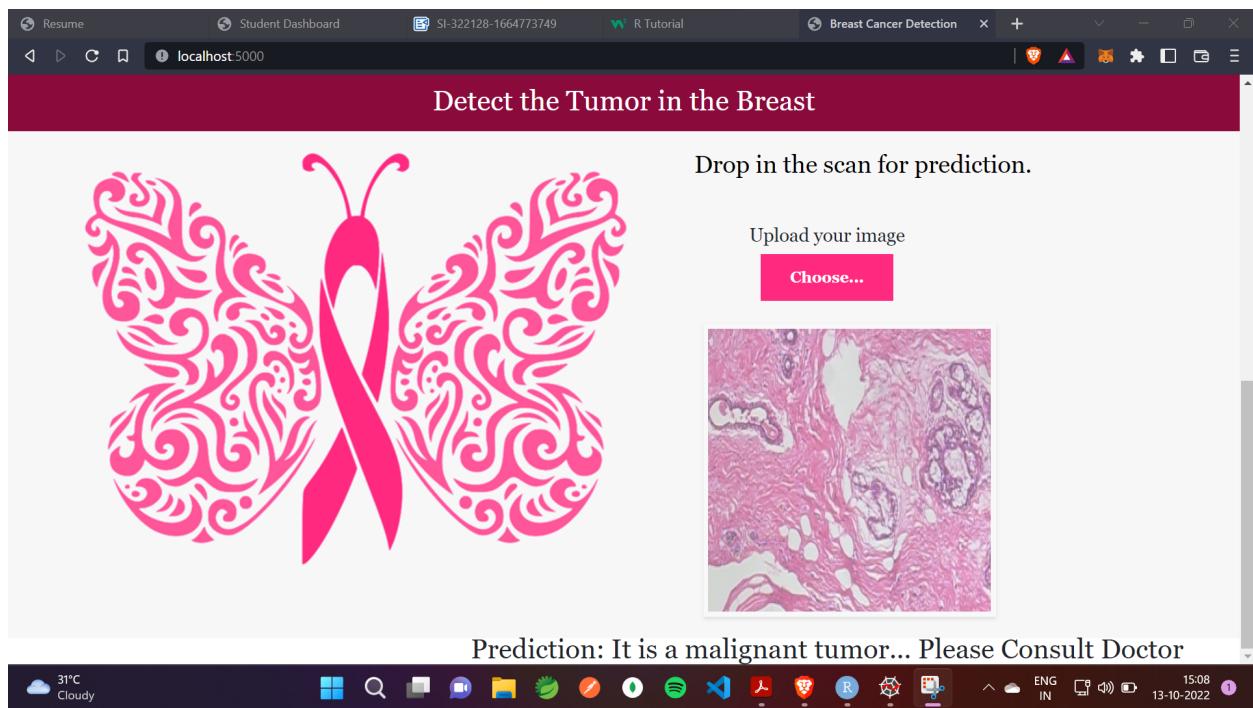
It shows that a model provided with different breast cancer scanning report images will predict the tumor and displays the output. Choose the image and click on upload, then it will predict the output.

## **5. FLOWCHART**



## **6. RESULT**





## 7. ADVANTAGES AND DISADVANTAGES

### Advantages:

- Early detection and prevention can significantly reduce the chances of death.
- Increased accuracy for Breast cancer risk prediction.
- Reduce the time complexity.

### Disadvantages:

- Requires massive datasets to train on.
- Time consuming and more resources required.

## 8. APPLICATIONS

- Deep learning and Neural networks are key technologies used in the breast cancer risk prediction .

- It presents the results obtained by processing input from uploading image.

## 9. CONCLUSION

In this project, we have established the application to predict from uploaded image based on the IBM Cloud application. Breast cancer risk prediction can only use this web app to predict the breast cancer risk.

## 10. FUTURE SCOPE

The project can be further enhanced by deploying the deep learning model obtained using a web application and larger dataset could be used for prediction to give higher accuracy and produce the better result.

## 11. BIBILOGRAPHY

- Radiology , Breast cancer risk prediction using deep learning by *Min Sun Bae,MD,PhD and Hyug-Gi Kim, PhD.*
- Deep Learning and Convolutional Neural Networks for Medical Imaging and Clinical Informatics by *Le Lu, Xiasong Wang, Gustavo Carneiro and Lin Yang*

# APPENDIX

## Source Code

- Main\_Project.ipynb

The screenshot shows a Google Colab notebook titled "Main Project.ipynb". The code cell at the top imports TensorFlow and Keras preprocessing modules, initializes data generators for training and testing datasets, and prints the number of images found. The next cell attempts to mount Google Drive, which is already mounted. The final cell imports Keras models and layers.

```
[6] import tensorflow as tf
[7] from keras.preprocessing.image import ImageDataGenerator
[8] train_datagen = ImageDataGenerator(rescale = 1./255, shear_range = 0.2, zoom_range = 0.2, horizontal_flip = True)
[9] test_datagen = ImageDataGenerator(rescale = 1./255)
[10] x_train = train_datagen.flow_from_directory(r"/content/drive/MyDrive/SmartBidge/Breast-Cancer-Risk-Prediction-main/breastcancerdataset/train", target_size=(64, 64), batch_size=32, class_mode='binary')
[11] x_test = test_datagen.flow_from_directory(r"/content/drive/MyDrive/SmartBidge/Breast-Cancer-Risk-Prediction-main/breastcancerdataset/test", target_size=(64, 64), batch_size=32, class_mode='binary')

Found 103 images belonging to 2 classes.
Found 22 images belonging to 2 classes.

[12] from google.colab import drive
[13] drive.mount('/content/drive')

Drive already mounted at /content/drive; to attempt to forcibly remount, call drive.mount("/content/drive", force_remount=True).

[14] from keras.preprocessing.image import ImageDataGenerator
[15] from keras.models import Sequential
[16] from keras.layers import Conv2D, MaxPooling2D, Dense, Dropout, Flatten
```

The screenshot shows a Google Colab notebook titled "Main Project.ipynb". The code cell contains the following Python code:

```
[10] from keras.preprocessing.image import ImageDataGenerator  
      from keras.models import Sequential  
      from keras.layers import Conv2D, MaxPooling2D, Dense, Dropout, Flatten  
      from keras.layers import MaxPooling2D  
  
[11] model=Sequential()  
  
[12] model.add(Conv2D(64,(3, 3),activation='relu', input_shape=(75, 75, 3)))  
  
[13] model.add(MaxPooling2D(pool_size = (2,2)))  
  
[14] model.add(Flatten())  
  
[15] model.add(Dense(units= 40 ,kernel_initializer='random_uniform',activation = 'relu'))  
  
[16] model.add(Dense(units= 1,activation = 'softmax',kernel_initializer= 'uniform'))  
  
[17] model.compile(loss='binary_crossentropy',optimizer='adam',metrics=['accuracy'])
```

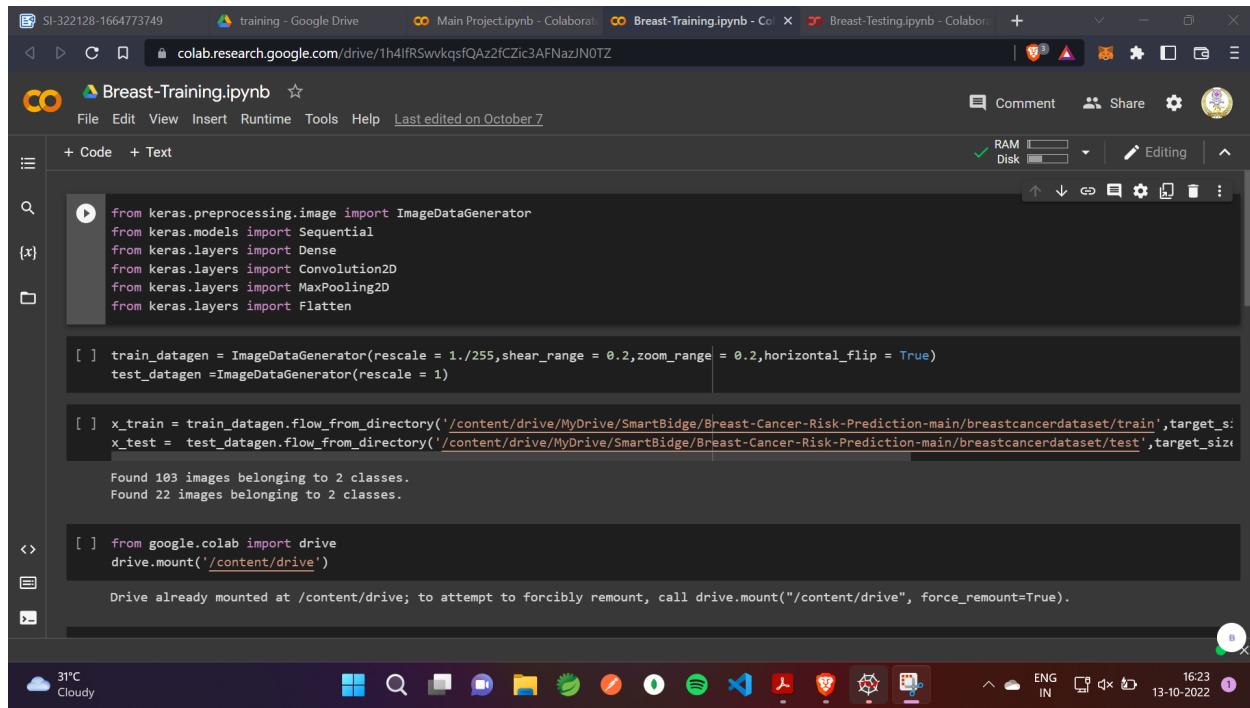
The status bar at the bottom indicates "0s completed at 4:10 PM". The taskbar at the bottom includes icons for Cloudy weather, 31°C, Microsoft Edge, File Explorer, Task View, File History, Spotify, Visual Studio Code, OneDrive, and a few others.

The screenshot shows a continuation of the Google Colab notebook "Main Project.ipynb". The code cell contains the following Python code:

```
[15] model.add(Dense(units= 40 ,kernel_initializer='random_uniform',activation = 'relu'))  
  
[16] model.add(Dense(units= 1,activation = 'softmax',kernel_initializer= 'uniform'))  
  
[17] model.compile(loss='binary_crossentropy',optimizer='adam',metrics=['accuracy'])  
  
[18] model.save('breastcancer.h5')  
  
[19] from keras.models import load_model  
      from keras.preprocessing import image  
      import numpy as np  
      from tensorflow.keras.models import load_model  
  
[20] model = load_model("breastcancer.h5")
```

The status bar at the bottom indicates "0s completed at 4:10 PM". The taskbar at the bottom includes icons for Cloudy weather, 31°C, Microsoft Edge, File Explorer, Task View, File History, Spotify, Visual Studio Code, OneDrive, and a few others.

## ● Breast-Training.ipynb



```
from keras.preprocessing.image import ImageDataGenerator
from keras.models import Sequential
from keras.layers import Dense
from keras.layers import Convolution2D
from keras.layers import MaxPooling2D
from keras.layers import Flatten

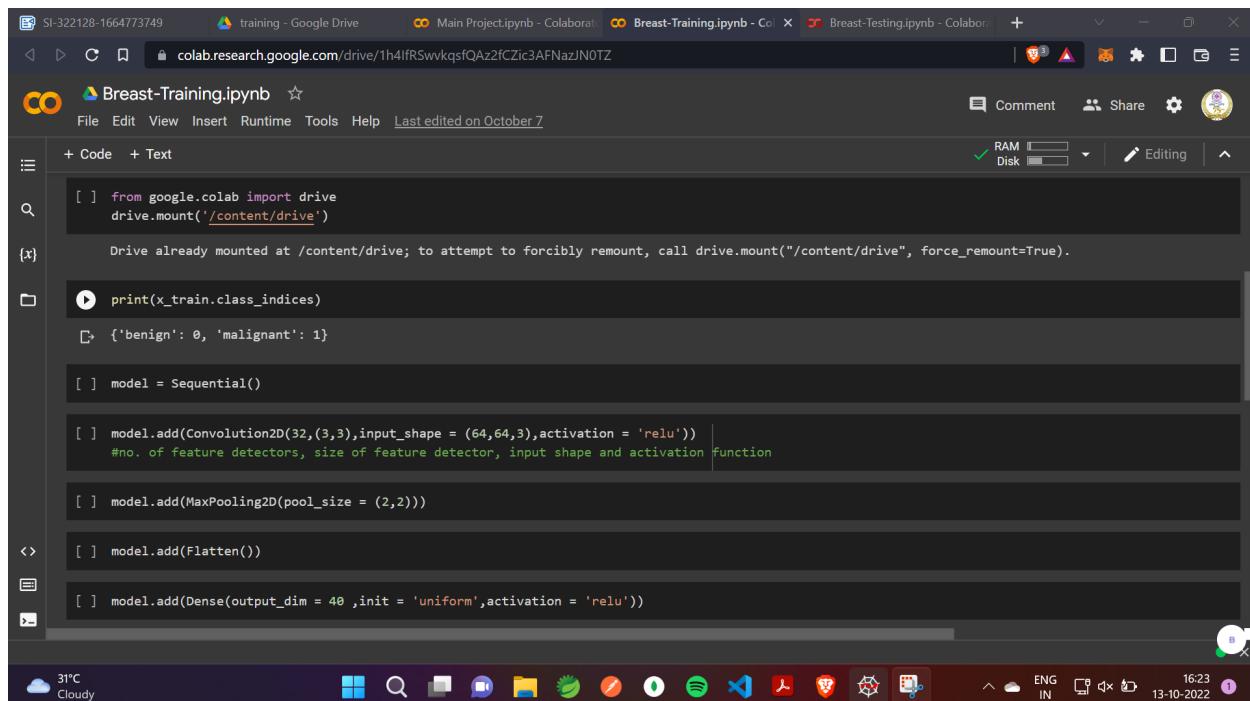
[ ] train_datagen = ImageDataGenerator(rescale = 1./255,shear_range = 0.2,zoom_range = 0.2,horizontal_flip = True)
test_datagen = ImageDataGenerator(rescale = 1)

[ ] x_train = train_datagen.flow_from_directory('/content/drive/MyDrive/SmartBidge/Breast-Cancer-Risk-Prediction-main/breastcancerdataset/train',target_size=(64,64),batch_size=32,class_mode='binary')
x_test = test_datagen.flow_from_directory('/content/drive/MyDrive/SmartBidge/Breast-Cancer-Risk-Prediction-main/breastcancerdataset/test',target_size=(64,64),batch_size=32,class_mode='binary')

Found 103 images belonging to 2 classes.
Found 22 images belonging to 2 classes.

[ ] from google.colab import drive
drive.mount('/content/drive')

Drive already mounted at /content/drive; to attempt to forcibly remount, call drive.mount("/content/drive", force_remount=True).
```



```
from google.colab import drive
drive.mount('/content/drive')

Drive already mounted at /content/drive; to attempt to forcibly remount, call drive.mount("/content/drive", force_remount=True).

[ ] print(x_train.class_indices)
{'benign': 0, 'malignant': 1}

[ ] model = Sequential()

[ ] model.add(Convolution2D(32,(3,3),input_shape = (64,64,3),activation = 'relu'))
#no. of feature detectors, size of feature detector, input shape and activation function

[ ] model.add(MaxPooling2D(pool_size = (2,2)))

[ ] model.add(Flatten())

[ ] model.add(Dense(output_dim = 40 ,init = 'uniform',activation = 'relu'))
```

The screenshot shows a Google Colab notebook titled "Breast-Training.ipynb". The code cell contains the following Python code:

```
[ ] model.add(Dense(output_dim = 1,activation = 'softmax',init ='uniform'))  
[ ] model.compile(loss = 'binary_crossentropy',optimizer = "adam",metrics = ["accuracy"])  
[ ] model.fit_generator(x_train, steps_per_epoch = 5,epochs = 10,validation_data = x_test,validation_steps = 40)  
[ ] model.save("breastcancer.h5")
```

## ● Breast-Testing.ipynb

The screenshot shows a Google Colab notebook titled "Breast-Testing.ipynb". The code cell contains the following Python code:

```
from keras.models import load_model  
from keras.preprocessing import image  
import numpy as np  
model = load_model("/content/drive/MyDrive/SmartBidge/Breast-Cancer-Risk-Prediction-main/training/ibmbreastcancer.h5")  
from google.colab import drive  
drive.mount('/content/drive')  
img = image.load_img('/content/drive/MyDrive/SmartBidge/Breast-Cancer-Risk-Prediction-main/benign.png',target_size = (64,64))
```

The screenshot shows a Jupyter Notebook interface on Google Colab. The notebook is titled "Breast-Testing.ipynb". The code cell contains the following Python code:

```
[ ] x = image.img_to_array(img)
x = np.expand_dims(x, axis = 0)

{x}
[ ] pred = np.argmax(model.predict(x), axis=-1)
#model.predict_classes(x)

[ ] pred
```