

Configuration Manual

MSc Research Project MSc Data Analytics

Yash Kutaphale Student ID: X21195960

School of Computing National College of Ireland

Supervisor: Furqan Rustam

National College of Ireland

MSc Project Submission Sheet

School of Computing

Student Name: Yash Santosh Kutaphale

Student ID: x21195960

Programme: MSc in Data Analytics **Year:** 2022-23

Module: MSc Research Project

Lecturer: Furqan Rustam

Submission Due Date: 14th August 2023

Project Title: Lung Cancer Detection Using Machine Learning And Deep Learning

Word Count: 1239 **Page Count:** 24

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Signature: Yash Kutaphale

Date: 14/08/23

PLEASE READ THE FOLLOWING INSTRUCTIONS AND CHECKLIST

Assignments that are submitted to the Programme Coordinator Office must be placed into the assignment box located outside the office.

Configuration Manual

Yash Kutaphale
Student ID: X21195960

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1. Hardware & Software

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Figure 1: Device Specifications

1.2 Windows Specifications

Figure 2: Windows Specifications

1.3 Python and Jupyter

Python Version: 3.10.1 Jupyter Version: 6.4.8

2 Data

2.1 Collection

The "IQOTHNCCD - Lung Cancer Dataset" dataset is provided from Kaggle. It provides a thorough set of information designed especially for lung cancer research.

IQ-OTH/NCCD - Lung Cancer Dataset

Includes CT scans of patients diagnosed with Lung Cancer.

Figure 4: Dataset

2.2 Model Building

2.2.1 Approch 1

Figure 5: Python Code for importing all the required libraries:

Approach 1

In this section all the necessary libraries are imported.

Figure 6: Python code for Loading Data:

Loading Data

```
# Set the path to your image dataset
dataset_directory = "C://Users//A//Desktop//CT Scan Dataset"
# Set the number of desired GLCM features to select
num features = 20
# Set the number of classes in your dataset
num\_classes = 3# Set the batch size and number of training epochs
batch_size = 32epochs = 10# Load the image collection
image_collection = imread_collection(os.path.join(dataset_directory, '*', '*.jpg'))
```
Here first the data set path is set. Then the number of GLCM features is set to 20. The number of classes is set to 3 as there are three classes present in the data set. And later the image collection is loaded.

Figure 7: Python Code for Extracting and Performing Feature Selection:

Extracting GLCM Features

The computed GLCM features are stored in a variable and then each image runs through a loop Which extracts the GLCM features and stores in an array

Performing Feature Selection

```
# Convert labels to numerical values
label_mapping = {label: i for i, label in enumerate(set(labels))}<br>numerical_labels = np.array([label_mapping[label] for label in labels])
# Perform feature selection using ANOVA F-value
selector = SelectKBest(f_classif, k=num_features)
selected_features = selector.fit_transform(glcm_features, numerical_labels)
# Get the names of the selected features
selected_feature_indices = selector.get_support(indices=True)<br>selected_feature_names = [f'Feature {idx+1}' for idx in selected_feature_indices]
# Print the names of the selected features
print('Selected Features:')
for feature_name in selected_feature_names:
     print(feature_name)
```
All the labels are converted to numerical values and feature selection is performed using ANOVA F value. The name of the selected features is fetched.

Figure 8: Python Code to Split Data into Train and Test:

Train - Test Split

```
# Split the data into train and test sets
X train, X test, y train, y test = train test split(selected features, numerical labels, test size=0.2, random state=42)
# Normalize the selected features
scalar = standardScalar()X_train_normalized = scaler.fit_transform(X_train)
X test normalized = scaler.transform(X test)
```

The data set is split into trained and test and the selecyed features are normalise using standard scaler.

Figure 9: Python code to build the CNN Model:

Developing CNN Model

```
# Create the CNN model
model = Sequential(f)Dense(128, activation='relu', input_shape=(num_features,)),
     Dense(num_classes, activation='softmax')
\left| \right\rangle# Compile the model
model.compile(optimizer='adam',
                  loss='categorical_crossentropy',
                  metrics=['accuracy'])
# Convert the Labels to one-hot encoding
y_train_encoded = to_categorical(y_train, num_classes=num_classes)
y_test_encoded = to_categorical(y_test, num_classes=num_classes)
# Train the model
model.fit(X_train_normalized, y_train_encoded,
              validation_data=(X_test_normalized, y_test_encoded),
             batch_size=batch_size,
             epochs=epochs)
# Save the trained model
model.save('lung_cancer_model.keras')
# Evaluate the model on the test dataset
_, test_accuracy = model.evaluate(X_test_normalized, y_test_encoded)
# Predict the classes for the test dataset
y_pred_probabilities = model.predict(X_test_normalized)
y_pred = np.argmax(y_pred_probabilities, axis=1)
# Compute evaluation metrics for the CNN model
accuracy = accuracy_score(y_test, y_pred)<br>precision = precision score(y_test, y_pred, average='weighted', zero_division=0)<br>precision = precision score(y_test, y_pred, average='weighted', zero_division=0)
                       precall = recall score(y_test, y_pred, average='weighted')<br>f1 = f1_score(y_test, y_pred, average='weighted')<br>f1 = f1_score(y_test, y_pred, average='weighted')
                       # Print the evaluation metrics for the CNN model
                       # Print the evaluation metrics for the CNN<br>print('CNN Test Accuracy:', test_accuracy)<br>print('CNN Accuracy:', accuracy)<br>print('CNN Precision:', precision)<br>print('CNN Recall:', recall)<br>print('CNN F1-score:', f1)
```
In this part of the code the CNN model is Built and compiled. Then the lables are converted to one hot encoding and the model is trained using the trained data set. Later the model is use for prediction and evalution matrix are computed.

Figure 10: Python Code to build the SVM Model:

Developing SVM Model

```
# Build the SVM model
svm_model = SVC(kernel='linear')
svm_model.fit(X_train_normalized, y_train)
# Predict the classes for the test dataset using the SVM model
svm_pred = svm_model.predict(X_test_normalized)
# Compute evaluation metrics for the SVM model
svm_accuracy = accuracy_score(y_test, svm_pred)<br>svm_accuracy = accuracy_score(y_test, svm_pred)<br>svm_precision = precision_score(y_test, svm_pred, average='weighted', zero_division=0)
sym_precision = precision_score(y_cest, sym_pred, average= weighted')<br>sym_recall = recall_score(y_test, sym_pred, average='weighted')<br>sym_f1 = f1_score(y_test, sym_pred, average='weighted')
# Print the evaluation metrics for the SVM model
print('SVM Accuracy:', svm_accuracy)<br>print('SVM Precision:', svm_accuracy)
print('SVM Recall:', svm_recall)<br>print('SVM F1-score:', svm_recall)
SVM Accuracy: 0.6045454545454545
```
SVM Precision: 0.5282170608028943 SVM Recall: 0.6045454545454545 SVM F1-score: 0.5493479044700212

In this part of the code the SVM model is Built. Later the model is use for prediction and evalution matrix are computed.

Figure 11: Python Code to build the Random Forest Model:

Developing Random Forest Model

```
# Build the Random Forest model
rf model = RandomForestClassifier(n estimators=100)
rf model.fit(X train normalized, \overline{v} train)
# Predict the classes for the test dataset using the Random Forest model
rf pred = rf model.predict(X test normalized)
# Compute evaluation metrics for the Random Forest model<br>rf_accuracy = accuracy_score(y_test, rf_pred)
rf_precision = precision_score(y_test, rf_pred, average='weighted', zero_division=0)
rf_recall = recall_score(y_test, rf_pred, average='weighted')
rf f1 = f1 score(y test, rf pred, average='weighted')
# Print the evaluation metrics for the Random Forest model
# Pitart Chandom Forest Accuracy:', rf_accuracy)<br>print('Random Forest Accuracy:', rf_accuracy)<br>print('Random Forest Precision:', rf_precision)<br>print('Random Forest Recall:', rf_recall)<br>print('Random Forest F1-score:', rf_f
Random Forest Accuracy: 0.5772727272727273
Random Forest Precision: 0.5731013779814579
Random Forest Recall: 0.5772727272727273
Random Forest F1-score: 0.5748873543107312
```
In this part of the code the random forest model is Built. Later the model is use for prediction and evalution matrix are computed.

Figure 12: Python Code for building the Decision Tree Model:

Developing Decesion Tree Model

```
: # Build the Decision Tree model
  dt_model = DecisionTreeClassifier()
  dt_model.fit(X_train_normalized, y_train)
  # Predict the classes for the test dataset using the Decision Tree model
  dt pred = dt model.predict(x test normalized)
  # Compute evaluation metrics for the Decision Tree model
  dt accuracy = accuracy score(y test, dt pred)
  dt_precision = precision_score(y_test, dt_pred, average='weighted', zero_division=0)
  dt_recall = recall_score(y_test, dt_pred, average='weighted')
  dt_f1 = f1_score(y_test, dt_pred, average='weighted')
  # Print the evaluation metrics for the Decision Tree model
  print ('Decision Tree Accuracy:', dt_accuracy)<br>print ('Decision Tree Precision:', dt_precision)<br>print ('Decision Tree Recall:', dt_recall)
  print('Decision Tree F1-score:', dt_f1)
  Decision Tree Accuracy: 0.5727272727272728
  Decision Tree Precision: 0.5640756556740163
  Decision Tree Recall: 0.5727272727272728
  Decision Tree F1-score: 0.5664226682408501
```
In this part of the code the Decision tree model is Built. Later the model is use for prediction and evalution matrix are computed.

Figure 13: Python Code for Developing the Gradient Boosting Machine Model:

Developing Gradient Boosting Machine Model

```
# Build the Gradient Boosting Machine (GBM) model
gbm_model = GradientBoostingClassifier(n_estimators=100)
gbm_model.fit(X_train_normalized, y_train)
# Predict the classes for the test dataset using the GBM model
gbm_pred = gbm_m model.predict(X_test-normalized)# Compute evaluation metrics for the GBM model
gbm_accuracy = accuracy_score(y_test, gbm_pred)
gbm_precision = precision_score(y_test, gbm_pred, average='weighted', zero_division=0)
gbm_recall = recall_score(y_test, gbm_pred, average='weighted')
gbm_f1 = f1_score(y_test, gbm_pred, average='weighted')# Print the evaluation metrics for the GBM model
# Pitt Circle evolution metrics for the born model<br>print('Gradient Boosting Machine Accuracy:', gbm_accuracy)<br>print('Gradient Boosting Machine Precision:', gbm_precision)<br>print('Gradient Boosting Machine Recall:', gbm_reca
Gradient Boosting Machine Accuracy: 0.6409090909090909
```

```
Gradient Boosting Machine Precision: 0.6317883302319494
Gradient Boosting Machine Recall: 0.6409090909090909
Gradient Boosting Machine F1-score: 0.6227518225735617
```
In this part of the code the Gradient boosting machine model is Built. Later the model is use for prediction and evalution matrix are computed.

Figure 14: Python Code for developing the KNN Model:

Developing KNN Model

In this part of the code the KNN model is Built. Later the model is use for prediction and evalution matrix are computed.

Figure 15: Python Code for building Logistic Regression Model:

Developing Logistic Regression Model

```
: # Build the Loaistic Rearession model
  from sklearn.linear model import LogisticRegression
  logreg model = LogisticRegression(max iter=1000)
  logreg_model.fit(X_train_normalized, y_train)
  # Predict the classes for the test dataset using the Logistic Regression model<br>logreg_pred = logreg_model.predict(X_test_normalized)
  # Compute evaluation metrics for the Logistic Regression model
  logreg_accuracy = accuracy_score(y_test, logreg_pred)
  logreg_precision = precision_score(y_test, logreg_pred, average='weighted', zero_division=0)
  logreg_recall = recall_score(y_test, logreg_pred, average='weighted')
  logreg_f1 = f1_score(y_test, logreg_pred, average='weighted')
  # Print the evaluation metrics for the Logistic Regression model
  print('Logistic Regression Accuracy:', logreg_accuracy)<br>print('Logistic Regression Accuracy:', logreg_accuracy)<br>print('Logistic Regression Precision:', logreg_precision)
  print('Logistic Regression Recall:', logreg_recall)<br>print('Logistic Regression Recall:', logreg_recall)<br>print('Logistic Regression F1-score:', logreg_f1)
  Logistic Regression Accuracy: 0.6136363636363636
  Logistic Regression Precision: 0.531367490956532
```
Logistic Regression Recall: 0.6136363636363636 Logistic Regression F1-score: 0.5638501191348974

In this part of the code the Logistic regression model is Built. Later the model is use for prediction and evalution matrix are computed.

Figure 16: Python Code to develop the ResNet Model:

Developing ResNet Model

```
: # Create a fully connected neural network
  fc_model = sequential([Dense(128, activation='relu', input_shape=(num_features,)),
       Dense(64, activation='relu'),
       Dense(num_classes, activation='softmax')
  \overline{1}# Compile the model
  for Compile the model<br>form is done in the compile (optimizer = 'adam',<br>loss='categorical_crossentropy',
                      metrics=['accuracy'])
  # Train the modelfc_model.fit(X_train_normalized, y_train_encoded,
                   validation_data=(X_test_normalized, y_test_encoded),
                  batch_size=batch_size,
                   epochs=epochs)
  # Evaluate the model on the test dataset
  _, fc_test_accuracy = fc_model.evaluate(X_test_normalized, y_test_encoded)
  # Predict the classes for the test dataset using the fully connected model
  fc y pred probabilities = fc model.predict(x test normalized)
  f \circ f pred = np.argmax(f \circ f pred_probabilities, axis=1)
  # Compute evaluation metrics for the fully connected model
  fc_accuracy = accuracy_score(y_test, fc_y_pred)<br>fc_accuracy = accuracy_score(y_test, fc_y_pred)<br>fc_precision = precision_score(y_test, fc_y_pred, average='weighted', zero_division=0)
  recall = recall score(y_test, fc_y_pred, average='weighted')<br>fc_f1 = f1_score(y_test, fc_y_pred, average='weighted')<br>fc_f1 = f1_score(y_test, fc_y_pred, average='weighted')
  # Print the evaluation metrics for the fully connected model
  print('Fully Connected Test Accuracy:', fc_test_accuracy)<br>print('Fully Connected Accuracy:', fc_accuracy)<br>print('Fully Connected Precision:', fc_precision)
  print('Fully Connected Recall:', fc_recall)
  print('Fully Connected F1-score:', fc_f1)
  Fully Connected Test Accuracy: 0.6409090757369995
   Fully Connected Accuracy: 0.6409090909090909
  Fully Connected Precision: 0.5548832897290247
  Eully Connected Recall: 0.6409090909090909
  Fully Connected F1-score: 0.594112181524769
```
In this part of the code the ResNet model is Built and compiled. The model is then trained and Later the model is use for prediction and evalution matrix are computed.

__

Figure 17: Python Code to build Efficient Model:

Developing EfficientNet Model

```
# After Normalizing the selected features:<br>scaler = StandardScaler()
    scaler = StandardScaler()<br>X_train_normalized = scaler.fit_transform(X_train)<br>X_test_normalized = scaler.transform(X_test)
     # Function to pad the features to 32x32<br>def pad_features(features, target_size=(32, 32)):
           pad_reatures(reatures, target_size=(32, 32)):<br>padded_features = np.zeros((features.shape[0], target_size[0], target_size[1]))<br>offset_x = (target_size[0] - features.shape[1]) // 2<br>offset_y = (target_size[1] - features.shape
    # Reshape the data to (samples, 20, 20)<br>X_train_reshaped = X_train_normalized.reshape(-1, num_features, 1)
     X_test_reshaped = X_test_normalized.reshape(-1, num_features, 1)
     # Pad the reshaped data
     X_train_padded = pad_features(X_train_reshaped)<br>X_test_padded = pad_features(X_test_reshaped)
    # Add an extra dimension for the channel (since EfficientNet expects it)<br>X_train_padded = X_train_padded[..., np.newaxis]<br>X_test_padded = X_test_padded[..., np.newaxis]
     # Now use the padded data for EfficientNetB0:
    whow use the punker boxes of the extended of the contract of efficient phase = Efficient base of the extended EU and the state of t
    x = usualneragerousingzo()(x)<br>x = Dense(128, activation='relu')(x)<br>predictions = Dense(num_classes, activation='softmax')(x)<br>efficientnet_model = Model(inputs=efficientnet_base.input, outputs=predictions)
efficientnet_model.compile(optimizer='adam',<br>loss='categorical_crossentropy',
                                                    metrics=['accuracy'])
{\it efficient net\_model.fit(x\_train\_padded, \textit{ y\_train\_encoded, \textit{ y\_test\_encoded, \textit{ y\_test\_encoded, \textit{ x\_test\_padded, y\_test\_encoded, \textit{ x\_test\_padded, \textit{ y\_test\_encoded, \textit{ x\_test\_padded, \textit{ y\_test\_pmodel}}}, \textit{ x\_test\_pmodel, \textit{ y\_test\_pmodel, \textit{ y\_test\_pmodel, \textit{ x\_test\_pmodel, \textit{ y\_test\_pmodel, \textit{ x\_test\_pmodel, \textit{ x\_test\_pmodel, \textit{ x\_test\_pmodel, \textit{ x\_test\_pmodel, \textit{ x\_test\_pmodel,batch_size=batch_size,
                                               epochs=epochs)
_, efficientnet_test_accuracy = efficientnet_model.evaluate(X_test_padded, y_test_encoded)<br>efficientnet_y_pred_probabilities = efficientnet_model.predict(X_test_padded)<br>efficientnet_y_pred = np.argmax(efficientnet_y_pred_p
efficientnet_accuracy = accuracy_score(y_test, efficientnet_y_pred)
efficientnet_acturaty = acturaty_soure(y_test, efficientnet_y_pred, average='weighted', zero_division=0)<br>efficientnet_precision = precision_score(y_test, efficientnet_y_pred, average='weighted', zero_division=0)<br>efficientn
nrint('EfficientNet Test Accuracy:', efficientnet test accuracy)
print('EfficientNet Test Accuracy:', efficientnet_test_a<br>print('EfficientNet Accuracy:', efficientnet_accuracy)<br>print('EfficientNet Precision:', efficientnet_precision)<br>print('EfficientNet Recall:', efficientnet_recall)<br>pr
EfficientNet Test Accuracy: 0.4954545497894287
EfficientNet Accuracy: 0.4954545454545455
EfficientNet Precision: 0.24547520661157027
EfficientNet Recall: 0.4954545454545455
EfficientNet F1-score: 0.32829510914617294
```
After normalising the selected features using standard scaler tye features are reshaped into 32x32. The data is reshaped to 20x20 and padded. After that extra dimension is added as the model used is EfficientNet Then the padded data is used to built and compiled model. The evaluation metrices are printed in the end.

Figure 18: Python Code to generate comparison of results:

Comparison of the Results

```
# Compare the evaluation metrics of the models
print('Evaluation Metrics:')
print(print('CNN Accuracy:', accuracy)<br>print('SVM Accuracy:', svm_accuracy)
princ("Swit Accuracy;", "swin_accuracy;"<br>print("Random Forest Accuracy:", rf_accuracy)<br>print("Decision Tree Accuracy:", dt_accuracy)
print('Gradient Boosting Machine Accuracy:', gbm_accuracy)
print ("k-Nearest Neighbors Accuracy;", som_accuracy<br>print ("k-Nearest Neighbors Accuracy;", knn_accuracy)<br>print ("Logistic Regression Accuracy;", logreg_accuracy)<br>print ("ResNet Accuracy;", ef_accuracy)<br>print ("FfficientN
print('CNN Precision:', precision)<br>print('SVM Precision:', svm_precision)
print('Random Forest Precision:', rf_precision)<br>print('Random Forest Precision:', rf_precision)<br>print('Decision Tree Precision:', dt_precision)<br>print('Gradient Boosting Machine Precision:', gbm_precision)
print('k-Nearest Neighbors Precision:', knn_precision)<br>print('k-Nearest Neighbors Precision:', knn_precision)<br>print('Logistic Regression Precision:', logreg_precisi
                                                                             , logreg_precision)
print('ResNet Precision:', fc_precision)
print('EfficientNet Precision:', efficientnet_precision)<br>print('EfficientNet Precision:', efficientnet_precision)<br>print('----------------------')
print('CNN Recall:', recall)<br>print('SVM Recall:', svm_recall)
print('Random Forest Recall:', rf_recall)<br>print('Random Forest Recall:', dt_recall)<br>print('Gradient Boosting Machine Recall:', gbm_recall)
print('k-Nearest Neighbors Recall:', knn_recall)
print('Logistic Regression Recall:', logreg_recall)
print('ResNet Recall:', fc_recall)
print('EfficientNet Recall:', efficientnet_recall)
print('---1)
```

```
print('-------------------------')
princ('CNN F1-score:', f1)<br>print('SVM F1-score:', svm_f1)
print('Random Forest F1-score:', rf_f1)<br>print('Decision Tree F1-score:', dt_f1)
print('Gradient Boosting Machine F1-score:', gbm_f1)<br>print('k-Nearest Neighbors F1-score:', knn_f1)<br>print('Logistic Regression F1-score:', logreg_f1)
print('ResNet F1-score:', fc_f1)
print('EfficientNet F1-score:', efficientnet f1)
```
Evaluation Metrics:

CNN Accuracy: 0.6272727272727273 SVM Accuracy: 0.6045454545454545 Random Forest Accuracy: 0.5772727272727273 Decision Tree Accuracy: 0.5727272727272728 Gradient Boosting Machine Accuracy: 0.6409090909090909 k-Nearest Neighbors Accuracy: 0.6545454545454545 Logistic Regression Accuracy: 0.6136363636363636 ResNet Accuracy: 0.6409090909090909 EfficientNet Accuracy: 0.4954545454545455 CNN Precision: 0.5621333417713508 SVM Precision: 0.5282170608028943 Random Forest Precision: 0.5731013779814579 Decision Tree Precision: 0.5640756556740163 Gradient Boosting Machine Precision: 0.6317883302319494 k-Nearest Neighbors Precision: 0.6508510638297873 Logistic Regression Precision: 0.531367490956532 ResNet Precision: 0.5548832897290247 EfficientNet Precision: 0.24547520661157027

```
CNN Recall: 0.6272727272727273
SVM Recall: 0.6045454545454545
Random Forest Recall: 0.5772727272727273
Decision Tree Recall: 0.5727272727272728
Gradient Boosting Machine Recall: 0.6409090909090909
k-Nearest Neighbors Recall: 0.654545454545454545<br>Logistic Regression Recall: 0.613636363636363636
ResNet Recall: 0.6409090909090909
EfficientNet Recall: 0.4954545454545455
CNN F1-score: 0.5653481316109108
SVM F1-score: 0.5493479044700212
```

```
Pandom Forest F1-500re: 0.5748873543107312<br>Decision Tree F1-500re: 0.5748873543107312
Gradient Boosting Machine F1-score: 0.6227518225735617
k-Nearest Neighbors F1-score: 0.6323060991988145
Logistic Regression F1-score: 0.5638501191348974
ResNet F1-score: 0.594112181524769
EfficientNet F1-score: 0.32829510914617294
```
In the end the results of all the models are compared.

Figure 19: Python Code to Show visualization of the Results:

Visual Representation of Results

Here the evaluation metrices of all the models is compared visually using bar graph.

Figure 20: Python Code to plot Confusion Matrix for CNN Model:

The Confusion matrix of CNN model is plotted here in this section.

2.2.2 Approch 2

Figure 21: Python Code to import all the required libraries in Approach 2

Approach 2

import numpy as np import pandas as nd import matplotlib.pyplot as plt import matplotlib. image as mpimg from PIL import Image import seaborn as sns import cv2 import random import os import imageio import plotly.graph_objects as go import plotly.express as px
import plotly.express as px
import plotly.figure_factory as ff
from plotly.subplots import make_subplots from collections import Counter from sklearn.preprocessing import StandardScaler from sklearn model selection import train test split from sklearn.neighbors import LocalOutlierFactor from sklearn.metrics import accuracy_score, recall_score, precision_score, classification_report, confusion_matrix, plot_confusio from sklearn.model_selection import RandomizedSearchCV, cross_val_score, RepeatedStratifiedKFold from imblearn.over_sampling import SMOTE import tensorflow as tf import tensorflow_addons as tfa import keras from keras.models import Sequential from keras.layers import Dense, Dropout, Activation, Flatten from keras.layers import Conv2D, MaxPooling2D, GlobalAveragePooling2D, BatchNormalization from keras.applications import resnet from tensorflow.keras.applications import EfficientNetB0, EfficientNetB1, EfficientNetB2, EfficientNetB3, EfficientNetB4, Efficie from keras.applications.resnet import ResNet50 from keras.preprocessing.image import ImageDataGenerator, load_img, img_to_array, array_to_img

All the necessary libraries required for the second approach are imported in this section.

__

Figure 22: Python Code to Load the dataset for approach 2:

Loading Data for Approach 2

```
# Set the directory and categories
directory = r"C:\Users\A\Desktop\CT Scan Dataset"
categories = ['Bengin cases', 'Malignant cases', 'Normal cases']
# Analyzing image size variations<br>size_data = {}
for i in categories:
    path = os.path.join(directory, i)
    class_{num} = categories.index(i)temp\_dict = \{\}for file in os.listdir(path):
         filepath = os.path.join(path, file)
        height, width, channels = imageio.imread(filepath).shape<br>if str(height) + ' x ' + str(width) in temp_dict:
             temp_dict[str(height) + ' x ' + str(width)] += 1
         else:
             temp_dict[str(height) + ' x ' + str(width)] = 1
    size\_data[i] = temp\_dictsize_data
{'Bengin cases': {'512 x 512': 120},
  'Malignant cases': {'512 x 512': 501,
   '404 \times 511 : 1.
  '512 x 801': 28,
```
'512 x 623': 31}, 'Normal cases': $\{$ '512 x 512': 415, '331 x 506': 1}}

In this part of the code data set directory and categories are set.

Figure 23: Python Code for preprocessing and visualization of data:

Preprocessing and visualizing images

```
# Preprocessing and visualization of images
img\_size = 256for i in categories:
    r and comples = 0, 3<br>
fig, ax = plt.subplots(samples, 3, figsize=(15, 15))<br>
fig.suptitle(i)
     path = os.path.join(directory, i)
     class_num = categories.index(i)<br>for curr_cnt, file in enumerate(os.listdir(path)):<br>filepath = os.path.join(path, file)
          img = cv2.imread(filename, 0)img0 = cv2.resize(img, (img_size, img_size))
          img1 = cv2.GaussianBlur(img0, (5, 5), 0)ax[cnt, 0].imshow(img)ax[cnt, 1].imshow(img@)ax[cnt, 2].imshow(img1)\cot + 1if cnt == samples:
               break
plt.show()
```
The preprocessing step is carried out here. The images are resize and GaussianBlur effect is used. And the results are visualized.

Bengin cases

Malignant cases

Normal cases

Preparing Data for Model Building

```
: # Preparing data<br>data = []
    img size = 256for i in categories:<br>
path = os.path.join(directory, i)<br>
class_num = categories.index(i)<br>
for file in os.listdir(path):
                filepath = os.path.join(path, file)img = cv2.imread(filepath, 0)
                # preprocess here<br>img = cv2.resize(img, (img_size, img_size))<br>data.append([img, class_num])
    random.shuffle(data)
   x, y = [], []<br>for feature, label in data:<br>x.append(feature)
          y.append(label)
   print('X length:', len(X))<br>print('y counts:', Counter(y))
    # normalize
   x = np.array(X) \cdot reshape(-1, img_size, img_size, 1)<br>x = x / 255.0y = np.array(y)X length: 1097
    y counts: Counter({1: 561, 2: 416, 0: 120})
```
Here the data is prepared and normalised for building the model.

Figure 26: Python Code to split into Train and Test and reshape Data using SMOTE

```
# Splitting the data into training and validation sets<br># Using stratification to maintain the distribution of the target variable in both training and validation sets<br>X_train, X_valid, y_train, y_valid = train_test_split(X
# Displaying the distribution of classes in the training and validation sets
print(Counter(y_train), Counter(y_valid))
# Checking the shape of the training data
print(len(X_train), X_train.shape)
# Reshaping the training data to a 2D array in preparation for SMOTE oversampling X_train = X_train.reshape(X_train.shape[0], img_size*img_size*1)
print(len(X_train), X_train.shape)
# Using SMOTE (Synthetic Minority Over-sampling Technique) to address class imbalance<br># This will create synthetic samples for the minority classes
print('Before SMOTE:', Counter(y_train))<br>smote = SMOTE()
X_train_sampled, y_train_sampled = smote.fit_resample(X_train, y_train) print('After SMOTE:', Counter(y_train_sampled))
# Reshaping the data back to its original shape after SMOTE oversampling
x_{\text{train}} = \bar{x}_{\text{train}}.reshape(x_{\text{train}} \cdot \bar{x}_{\text{map}}) img size, img size, 1)<br>x_{\text{train\_sampled}} = x_{\text{train\_sampled}.\text{reshape}(x_{\text{train\_sampled}.\text{shape[0]},\text{img\_size},\text{img\_size}, 1)# Checking the shape of the training data after SMOTE oversampling print(len(X_train), X_train.shape)
print(len(X_train_sampled), X_train_sampled.shape)
Counter({1: 420, 2: 312, 0: 90}) Counter({1: 141, 2: 104, 0: 30})<br>822 (822, 256, 256, 1)<br>822 (822, 65536)
Before SMOTE: Counter({1: 420, 2: 312, 0: 90})
After SMOTE: Counter({2: 420, 1: 420, 0: 420})<br>822 (822, 256, 256, 1)
1260 (1260, 256, 256, 1)
```
The dataset is split to train and test and synthetic samples are created for minority classes using SMOTE.

Figure 28: Python Code to Build CNN Model:

Building CNN Model

```
: # Initialize the CNN model using Keras Sequential API
  cnn_model = Sequential()
  # Add the first convolutional layer with 64 filters of size 3x3
  # Using the ReLU activation function
  cnn_model.add(Conv2D(64, (3, 3), input_shape=X_train.shape[1:]))<br>cnn_model.add(Conv2D(64, (3, 3), input_shape=X_train.shape[1:]))
  # Add a max-pooling layer to down-sample the feature maps
  cnn_model.add(MaxPooling2D(pool_size=(2, 2)))
  # Add the second convolutional layer with 64 filters of size 3x3
  # Using the ReLU activation function directly within the Conv2D layer
  cnn model.add(Conv2D(64, (3, 3), activation='relu'))
  # Add another max-pooling layer
  cnn_model.add(MaxPooling2D(pool_size=(2, 2)))
  # Flatten the feature maps into a one-dimensional vector to prepare for the fully connected layers
  cnn model.add(Flatten())
  # Add a dense (fully connected) layer with 16 neurons
  cnn_model.add(Dense(16))
  # Add the output layer with 3 neurons (one for each class)
  # Using the softmax activation function to produce a probability distribution
  cnn_model.add(Dense(3, activation='softmax'))
  # Display a summary of the model's architecture
  cnn_model.summary()
  # Compile the CNN model specifying the loss function, optimizer, and evaluation metric
  cnn_model.compile(loss='sparse_categorical_crossentropy', optimizer='adam', metrics=['accuracy'])
```
The CNN model is built in this part of the code. First convolutional layer with 64 filters is added using the ReLU activation function. Later a max-pooling layer is added to down sample the feature maps.

Again a convolutional layer is added followed by a max-pooling layer. In the end dense layersare added.

The model is trained and the classes are predicted. Classification report of the model is fetched.

Confusion Matrix for the Model

Confusion matrix is generated for the CNN model.

Figure 30: Python Code to Visualize the Results:

Visualization of the Results

```
# Training and validation accuracy and loss curves
plt.figure(figsize=(15,5))
# Accuracy
plt.subplot(1, 2, 1)
plt.plot(history.history['accuracy'], label='Training Accuracy')<br>plt.plot(history.history['val_accuracy'], label='Validation Accuracy')
plt.title('Model Accuracy')
plt.xlabel('Epoch')<br>plt.ylabel('Epoch')
plt.legend()
# Loss# Loss<br>plt.subplot(1, 2, 2)<br>plt.plot(history.history['loss'], label='Training Loss')<br>plt.plot(history.history['val_loss'], label='Validation Loss')<br>http://wodol_loss')
pit.piot(mistory.mistory<br>plt.title('Model Loss')<br>plt.xlabel('Epoch')<br>plt.ylabel('Loss')
plt.legend()
plt.tight_layout()
plt.show()
```
Line graph is printed for the model accuracy and loss is plotted using Matplotlib.

