```
In [1]: import os
        import cv2
        import numpy as np
        import librosa
        import random
        import pywt
        from skimage.measure import block reduce
        from sklearn.model selection import train test split
        from sklearn.decomposition import PCA
        import tensorflow as tf
        from tensorflow.keras import layers, models
        from sklearn.metrics import confusion matrix, classification_report, ConfusionMatrixDisplay
        import matplotlib.pyplot as plt
        from tensorflow import keras
        from keras.models import Model
        import matplotlib.pyplot as plt
        from tensorflow.keras.models import Sequential
        from tensorflow.keras.layers import Dense,Dropout
        from tensorflow.keras.optimizers import Adam
        from tensorflow.keras.utils import to categorical
        from tensorflow.keras import layers
        from tensorflow.keras.layers import Conv2D, MaxPooling2D, Flatten, Dense, AveragePooling2D
        from tensorflow.keras import layers, optimizers
        from sklearn.ensemble import RandomForestClassifier
        from sklearn.metrics import accuracy score
```

C:\Users\nehit\anaconda3\Lib\site-packages\paramiko\transport.py:219: CryptographyDeprecationWarning: Blowfish has be en deprecated

"class": algorithms.Blowfish,

WARNING:tensorflow:From C:\Users\nehit\anaconda3\Lib\site-packages\keras\src\losses.py:2976: The name tf.losses.spars e\_softmax\_cross\_entropy is deprecated. Please use tf.compat.v1.losses.sparse\_softmax\_cross\_entropy instead.

```
In [94]: normal_path = 'binaryclassification/normal sound/264_1309356143724_D.wav'
abnormal_path = "binaryclassification/abnormal sound/201108222245.wav"
```

```
In [95]: # Directory paths for normal and abnormal heartbeats
    normal_dir = 'binaryclassification/normal sound'
    abnormal_dir = 'binaryclassification/abnormal sound'

In [97]: audio_data, sampling_rate = librosa.load('binaryclassification/normal sound/264_1309356143724_D.wav', sr=None, mono=Tr
    print(f"The sampling rate given: {sampling_rate} Hz")
```

The sampling rate given: 4000 Hz

## down sample

```
In [98]: #function for plotting Amplitude vs time

def plot_audio_waveform(audio_data, sample_rate):
    # Calculate the duration of the audio signal
    duration = len(audio_data) / sample_rate

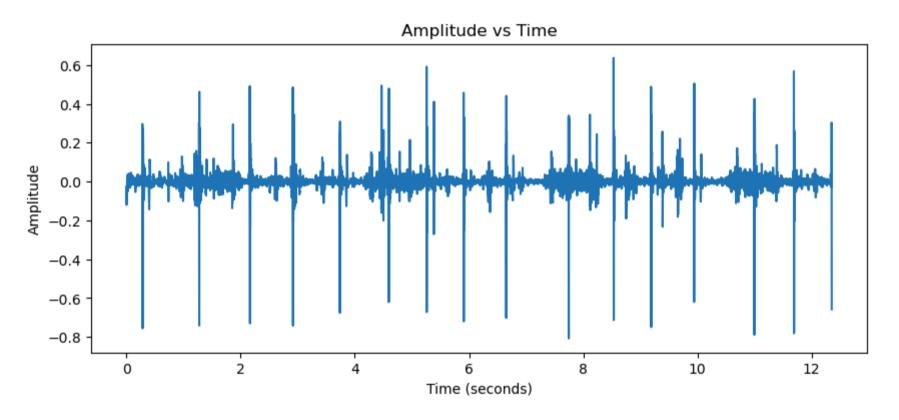
# Create a time array for plotting
    time = np.linspace(0, duration, len(audio_data))

# Plot the amplitude versus time
    plt.figure(figsize=(10, 4))
    plt.plot(time, audio_data)
    plt.title('Amplitude vs Time')
    plt.xlabel('Time (seconds)')
    plt.ylabel('Amplitude')

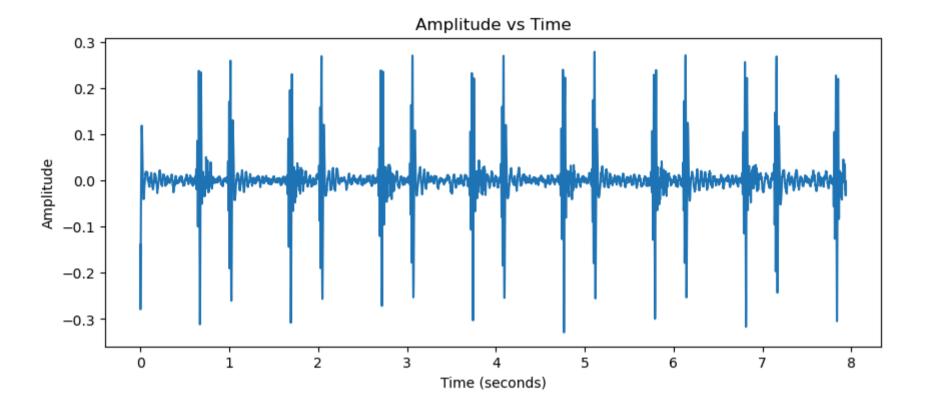
plt.show()
```

```
In [99]: load_normal = load_audio(normal_path)
    load_abnormal = load_audio(abnormal_path)
    print("NORMAL")
    plot_audio_waveform(load_normal,1000)
    print("ABNORMAL")
    plot_audio_waveform(load_abnormal,1000)
```

NORMAL



ABNORMAL



```
In [102]: #LOAD THE DATA WUTH THE REQUIRED SAMPLE RATE -1000
          def load(path: str):
              sample_rate, data = wav.read(path)
              new sample rate = 1000
              resampled_data = resample(data, int(len(data) * (new_sample_rate / sample_rate)))
              time = np.linspace(
                  0, # start
                  len(resampled_data) / new_sample_rate,
                  num = len(resampled data)
              return [time, resampled_data]
          #segmentation of data
          def segment(data):
              seg\_size = 3000 \# note sampled at 1000, and for t=3s, s = 1000*3s
              segmented_data = [data[1][i:i+seg_size] for i in range(0,len(data[1]),seg_size)]
              if len(segmented_data[len(segmented_data)-1]) < seg_size:</pre>
                  segmented_data.pop()
              return segmented_data
```

```
In [57]: #Loading the data :
         folder normal = path normal
         folder abnormal = path abnormal
         files_normal_path = os.listdir(folder_normal)
         files abnormal path = os.listdir(folder abnormal)
         files normal = []
         files_abnormal = []
         for file in files normal path:
             if file.endswith(".wav"):
                 files normal.append(os.path.join(folder_normal, file))
         for file in files_abnormal_path:
             if file.endswith(".wav"):
                 files abnormal.append(os.path.join(folder abnormal, file))
         data normal = []
         data_abnormal = []
         for i in files_normal:
             p1 = load(i)
             p2 = segment(p1)
             for j in p2:
                 data_normal.append(j)
                 dataset.append(j)
         for i in files abnormal:
             p1 = load(i)
             p2 = segment(p1)
             for j in p2:
                 data_abnormal.append(j)
                 dataset.append(j)
```

```
In [58]: print(f"The size of dataset we have: {len(dataset)}")
```

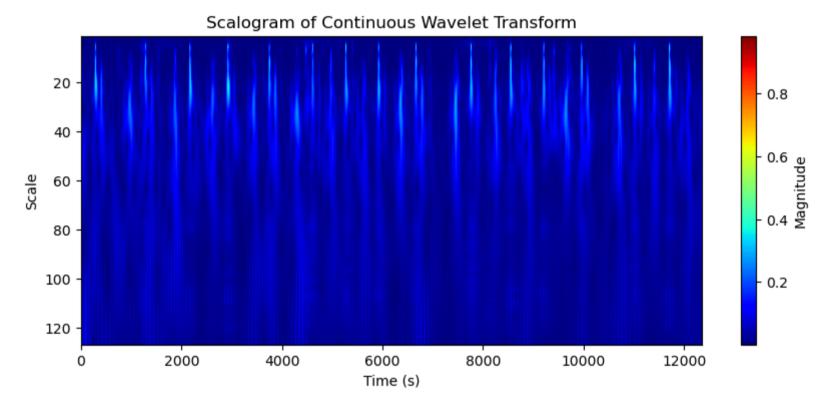
The size of dataset we have: 1672

```
In [59]: | def extract cwt features(audio data, scales=np.arange(1, 128)):
             # Perform Continuous Wavelet Transform (CWT) using pywt
             cwt data, = pywt.cwt(audio data, scales, 'morl')
             # Take absolute values of CWT coefficients as features
             cwt features = np.abs(cwt_data)
             return cwt features
         temp = extract_cwt_features(dataset[0][0])
         temp.shape
Out[59]: (127, 3000)
In [60]: def plot cwt scalogram(audio data, sampling rate, scales=np.arange(1,128)):
             # scaler = StandardScaler()
             # Calculate the CWT coefficients
             coeffs, _ = pywt.cwt(audio_data, scales, 'morl')
             # coeffs = scaler.fit transform(coeffs)
             print(f"The shape of cwt extract: {coeffs.shape}")
             # Calculate time axis
             time = np.arange(0, len(audio data)) / sampling rate
             # Plot scalogram
             plt.figure(figsize=(10, 4))
             plt.imshow(np.abs(coeffs), extent=[0, len(audio_data), scales[-1], scales[0]],
                        aspect='auto', cmap='jet')
             plt.colorbar(label='Magnitude')
             plt.xlabel('Time (s)')
             plt.ylabel('Scale')
             plt.title('Scalogram of Continuous Wavelet Transform')
             plt.show()
```

```
In [61]: print("NORMAL")
    plot_cwt_scalogram(load_normal,1000)
    print("ABNORMAL")
    plot_cwt_scalogram(load_abnormal,1000)
```

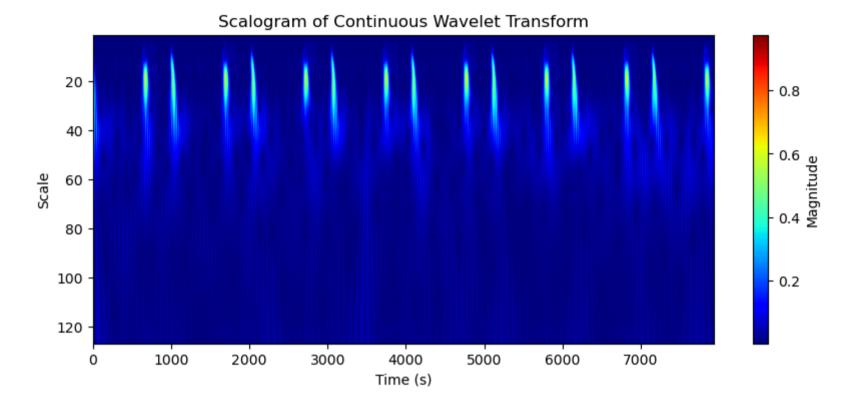
## NORMAL

The shape of cwt extract: (127, 12351)



## ABNORMAL

The shape of cwt extract: (127, 7936)



```
In [62]: #spillting the dataset

# Split the dataset into features (audio data) and labels
X = [entry[0] for entry in dataset] # Features (audio data)
y = [entry[1] for entry in dataset] # Labels

# Split the dataset into training and testing sets
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.2, random_state=42)
```

```
In [63]: def extract_cwt_features_from_audio_dataset(audio_dataset, sample_rate, scales=np.arange(1, 128)):
    cwt_features_dataset = []

for audio_data in audio_dataset:
    # Compute CWT features for each audio sample
    cwt_data, _ = pywt.cwt(audio_data, scales, 'morl')

# Take absolute values of CWT coefficients as features
    cwt_features = np.abs(cwt_data)

pca = PCA(n_components=100) # Assuming you want to reduce to 100 dimensions

# Fit and transform the data matrix using PCA
    reduced_data = pca.fit_transform(cwt_features)

# Append CWT features to the dataset
    cwt_features_dataset.append(reduced_data)

return cwt_features_dataset
```

```
In [64]: X train extracted = extract cwt features from audio dataset(X train,1000)
         X test extracted = extract cwt features from audio dataset(X test,1000)
         C:\Users\nehit\anaconda3\Lib\site-packages\sklearn\decomposition\ pca.py:640: RuntimeWarning: invalid value encoun
         tered in divide
           self.explained variance ratio = self.explained variance / total var
         C:\Users\nehit\anaconda3\Lib\site-packages\sklearn\decomposition\ pca.py:640: RuntimeWarning: invalid value encoun
         tered in divide
           self.explained variance ratio = self.explained variance / total var
         C:\Users\nehit\anaconda3\Lib\site-packages\sklearn\decomposition\ pca.py:640: RuntimeWarning: invalid value encoun
         tered in divide
           self.explained variance ratio = self.explained variance / total var
         C:\Users\nehit\anaconda3\Lib\site-packages\sklearn\decomposition\ pca.py:640: RuntimeWarning: invalid value encoun
         tered in divide
           self.explained variance ratio = self.explained variance / total var
         C:\Users\nehit\anaconda3\Lib\site-packages\sklearn\decomposition\ pca.py:640: RuntimeWarning: invalid value encoun
         tered in divide
           self.explained variance ratio = self.explained variance / total var
         C:\Users\nehit\anaconda3\Lib\site-packages\sklearn\decomposition\ pca.py:640: RuntimeWarning: invalid value encoun
         tered in divide
           self.explained variance ratio = self.explained variance / total var
         C:\Users\nehit\anaconda3\Lib\site-packages\sklearn\decomposition\ pca.py:640: RuntimeWarning: invalid value encoun ▼
In [65]: Xtrain = np.array(X_train_extracted)
         Xtest = np.array(X test extracted)
         ytrain = np.array(y train)
         ytest = np.array(y test)
         Xtrain.shape, ytrain.shape, Xtest.shape, ytest.shape
Out[65]: ((1337, 127, 100), (1337,), (335, 127, 100), (335,))
```

```
In [66]: import seaborn as sns

def plot_confusion_matrix(y_true, y_pred):
    # Compute confusion matrix
    cm = confusion_matrix(y_true, y_pred)

# Plotting the confusion matrix with default labels '0' and '1'
    plt.figure(figsize=(8, 6))
    sns.heatmap(cm, annot=True, fmt="d", cmap="Blues", xticklabels=['0', '1'], yticklabels=['0', '1'])
    plt.xlabel("Predicted labels")
    plt.ylabel("True labels")
    plt.title("Confusion Matrix")
    plt.show()
```

#MODEL 1

```
In [67]: model1 = Sequential()
         # Add convolutional layers and pooling layers
         model1.add(Conv2D(64, kernel size=(3, 3), activation='relu', input shape=(127, 100, 1)))
         model1.add(AveragePooling2D(pool size=(2, 2), strides=(2, 2)))
         model1.add(Dropout(0.5))
         model1.add(Conv2D(32, kernel size=(3, 3), activation='relu'))
         model1.add(AveragePooling2D(pool size=(2, 2), strides=(2, 2)))
         model1.add(Dropout(0.5))
         model1.add(Conv2D(16, kernel size=(3, 3), activation='relu'))
         model1.add(AveragePooling2D(pool size=(2, 2), strides=(2, 2)))
         model1.add(Conv2D(8, kernel size=(3, 3), activation='relu'))
         model1.add(AveragePooling2D(pool size=(2, 2)))
         model1.add(Flatten())
         model1.add(Dropout(0.5))
         # Add dense Layers
         model1.add(Dense(8, activation='relu'))
         model1.add(Dense(2, activation='softmax'))
         adam = Adam(learning rate=0.001)
         # Compile the model with sparse categorical crossentropy
         model1.compile(optimizer='adam', loss='sparse categorical crossentropy', metrics=['accuracy'])
         model1.summary()
```

Model: "sequential\_2"

Layer (type)	Output Shape	Param #
conv2d_7 (Conv2D)	(None, 125, 98, 64)	640
<pre>average_pooling2d_3 (Avera gePooling2D)</pre>	(None, 62, 49, 64)	0
dropout_3 (Dropout)	(None, 62, 49, 64)	0
conv2d_8 (Conv2D)	(None, 60, 47, 32)	18464
<pre>average_pooling2d_4 (Avera gePooling2D)</pre>	(None, 30, 23, 32)	0
dropout_4 (Dropout)	(None, 30, 23, 32)	0
conv2d_9 (Conv2D)	(None, 28, 21, 16)	4624
<pre>average_pooling2d_5 (Avera gePooling2D)</pre>	(None, 14, 10, 16)	0
conv2d_10 (Conv2D)	(None, 12, 8, 8)	1160
<pre>average_pooling2d_6 (Avera gePooling2D)</pre>	(None, 6, 4, 8)	0
flatten_3 (Flatten)	(None, 192)	0
dropout_5 (Dropout)	(None, 192)	0
dense_6 (Dense)	(None, 8)	1544
dense_7 (Dense)	(None, 2)	18
	=======================================	

Total params: 26450 (103.32 KB) Trainable params: 26450 (103.32 KB) Non-trainable params: 0 (0.00 Byte)

```
In [68]: model1_accuracies = []
for i in range(3):
    print("step: ",i+1)
    model1.fit(Xtrain, ytrain, epochs=15, batch_size=32)
    test_loss_m1, test_accuracy_m1 = model1.evaluate(Xtest, ytest, verbose=2)
    model1_accuracies.append(test_accuracy_m1)

mean_test_accuracy_m1 = np.mean(model1_accuracies)
print("Mean test accuracy Model 1:", mean_test_accuracy_m1)
```

```
step: 1
Epoch 1/15
42/42 [============= ] - 13s 230ms/step - loss: 0.6911 - accuracy: 0.5206
Epoch 2/15
42/42 [============== ] - 10s 229ms/step - loss: 0.6878 - accuracy: 0.5460
Epoch 3/15
Epoch 4/15
42/42 [============= ] - 10s 229ms/step - loss: 0.6788 - accuracy: 0.5445
Epoch 5/15
Epoch 6/15
42/42 [============= ] - 10s 231ms/step - loss: 0.6755 - accuracy: 0.5497
Epoch 7/15
42/42 [============= ] - 10s 234ms/step - loss: 0.6766 - accuracy: 0.5527
Epoch 8/15
Epoch 9/15
42/42 [============= ] - 9s 219ms/step - loss: 0.6743 - accuracy: 0.5535
Epoch 10/15
Epoch 11/15
42/42 [============== ] - 9s 224ms/step - loss: 0.6727 - accuracy: 0.5460
Epoch 12/15
42/42 [============= ] - 9s 223ms/step - loss: 0.6715 - accuracy: 0.5535
Epoch 13/15
Epoch 14/15
42/42 [============= ] - 9s 223ms/step - loss: 0.6721 - accuracy: 0.5527
Epoch 15/15
11/11 - 1s - loss: 0.6769 - accuracy: 0.5403 - 887ms/epoch - 81ms/step
step: 2
Epoch 1/15
42/42 [============== ] - 10s 231ms/step - loss: 0.6725 - accuracy: 0.5580
Epoch 2/15
Epoch 3/15
42/42 [============== ] - 10s 231ms/step - loss: 0.6726 - accuracy: 0.5542
Epoch 4/15
```

```
Epoch 5/15
Epoch 6/15
42/42 [============= ] - 10s 229ms/step - loss: 0.6652 - accuracy: 0.5512
Epoch 7/15
42/42 [============= ] - 10s 243ms/step - loss: 0.6674 - accuracy: 0.5535
Epoch 8/15
42/42 [============== ] - 10s 231ms/step - loss: 0.6692 - accuracy: 0.5497
Epoch 9/15
Epoch 10/15
42/42 [============ ] - 9s 220ms/step - loss: 0.6626 - accuracy: 0.5415
Epoch 11/15
42/42 [============== ] - 9s 221ms/step - loss: 0.6687 - accuracy: 0.5497
Epoch 12/15
Epoch 13/15
42/42 [============ ] - 9s 222ms/step - loss: 0.6609 - accuracy: 0.5527
Epoch 14/15
42/42 [============== ] - 9s 221ms/step - loss: 0.6649 - accuracy: 0.5625
Epoch 15/15
42/42 [============= ] - 9s 224ms/step - loss: 0.6590 - accuracy: 0.5527
11/11 - 1s - loss: 0.6636 - accuracy: 0.5881 - 512ms/epoch - 47ms/step
step: 3
Epoch 1/15
Epoch 2/15
42/42 [============== ] - 9s 225ms/step - loss: 0.6604 - accuracy: 0.5797
Epoch 3/15
Epoch 4/15
Epoch 5/15
Epoch 6/15
Epoch 7/15
Epoch 8/15
Epoch 9/15
42/42 [=================== ] - 11s 256ms/step - loss: 0.6518 - accuracy: 0.5789
```

```
curacy: 0.6537
Epoch 2/20
42/42 [============= ] - ETA: 0s - loss: 0.6438 - accuracy: 0.5894
Epoch 2: val accuracy did not improve from 0.65373
curacy: 0.5343
Epoch 3/20
42/42 [============== ] - ETA: 0s - loss: 0.6411 - accuracy: 0.5871
Epoch 3: val accuracy did not improve from 0.65373
curacy: 0.6358
Epoch 4/20
42/42 [============= ] - ETA: 0s - loss: 0.6385 - accuracy: 0.6096
Epoch 4: val accuracy did not improve from 0.65373
curacy: 0.6179
Epoch 5/20
42/42 [============== ] - ETA: 0s - loss: 0.6390 - accuracy: 0.5916
Epoch 5: val accuracy did not improve from 0.65373
curacy: 0.6507
Epoch 6/20
42/42 [============= ] - ETA: 0s - loss: 0.6408 - accuracy: 0.6148
Epoch 6: val accuracy did not improve from 0.65373
42/42 [============== ] - 10s 241ms/step - loss: 0.6408 - accuracy: 0.6148 - val_loss: 0.6514 - val_ac
curacy: 0.6239
Epoch 7/20
42/42 [============= ] - ETA: 0s - loss: 0.6350 - accuracy: 0.5991
Epoch 7: val accuracy did not improve from 0.65373
curacy: 0.6328
Epoch 8/20
Epoch 8: val accuracy did not improve from 0.65373
curacy: 0.6448
Epoch 9/20
42/42 [============== ] - ETA: 0s - loss: 0.6373 - accuracy: 0.6043
Epoch 9: val accuracy did not improve from 0.65373
```

```
curacy: 0.6269
Epoch 10/20
Epoch 10: val accuracy did not improve from 0.65373
42/42 [============== ] - 10s 239ms/step - loss: 0.6402 - accuracy: 0.5991 - val loss: 0.6441 - val ac
curacy: 0.6149
Epoch 11/20
Epoch 11: val accuracy did not improve from 0.65373
curacy: 0.6209
Epoch 12/20
Epoch 12: val accuracy did not improve from 0.65373
curacy: 0.6090
Epoch 13/20
42/42 [============== ] - ETA: 0s - loss: 0.6324 - accuracy: 0.6193
Epoch 13: val_accuracy did not improve from 0.65373
curacy: 0.6269
Test Loss: 0.6548299193382263, Test Accuracy: 0.6537313461303711
```

```
In [70]: model1.load_weights('best_model_weights.h5')
```

## In [71]: # Evaluate the model on the test data from sklearn.metrics import accuracy\_score, precision\_score, recall\_score, f1\_score, roc\_auc\_score y\_pred = model1.predict(Xtest) y\_pred\_classes = np.argmax(y\_pred, axis=1) # Convert probabilities to class labels # Calculate evaluation metrics accuracy = accuracy\_score(ytest, y\_pred\_classes) precision = precision\_score(ytest, y\_pred\_classes, average='weighted') recall = recall\_score(ytest, y\_pred\_classes, average='weighted') f1 = f1\_score(ytest, y\_pred\_classes, average='weighted') # Print the evaluation metrics print(f"Accuracy: {accuracy}") print(f"Precision: {precision}") print(f"Recall: {recall}") print(f"F1-score: {f1}")

11/11 [======] - 1s 44ms/step

Accuracy: 0.6537313432835821 Precision: 0.652074790880761 Recall: 0.6537313432835821 F1-score: 0.6523173605655931

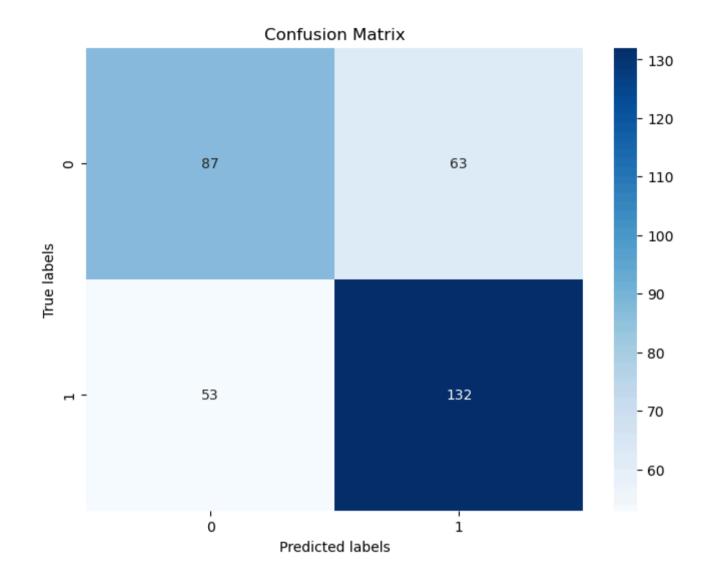
```
In [72]: from sklearn.metrics import confusion matrix
         def calculate sensitivity specificity(y true, y pred):
             # Calculate confusion matrix
             cm = confusion matrix(y true, y pred)
             # Extract TP, TN, FP, FN from confusion matrix
             TP = cm[1, 1] # True Positives
             TN = cm[0, 0] # True Negatives
             FP = cm[0, 1] # False Positives
             FN = cm[1, 0] # False Negatives
             # Calculate Sensitivity (True Positive Rate, TPR)
             sensitivity = TP / (TP + FN)
             # Calculate Specificity (True Negative Rate, TNR)
             specificity = TN / (TN + FP)
             return sensitivity, specificity
         # Assuming y true contains the true binary labels (0 and 1) and y pred contains the predicted labels
         sensitivity, specificity = calculate sensitivity specificity(ytest, y pred classes)
         print(f"Sensitivity (True Positive Rate, TPR): {sensitivity}")
         print(f"Specificity (True Negative Rate, TNR): {specificity}")
```

Sensitivity (True Positive Rate, TPR): 0.7135135135135136

Specificity (True Negative Rate, TNR): 0.58

```
In [82]: from sklearn.metrics import confusion matrix
         def display confusion matrix(y true, y pred, labels=None):
             cm = confusion_matrix(y_true, y_pred, labels=labels)
             if labels is None:
                 labels = sorted(set(y_true).union(y_pred))
             header = "|{:^10}|".format("True \\ Predicted")
             header += "|".join(["{:^10}]".format(label) for label in labels]) + "|"
             print(header)
             print("-" * len(header))
             for i, row_label in enumerate(labels):
                 row = "|{:^10}|".format(row label)
                 row += "|".join(["{:^10}".format(cm[i, j]) for j in range(len(labels))]) + "|"
                 print(row)
             print("-" * len(header))
         display_confusion_matrix(ytest, y_pred_classes)
         plot_confusion_matrix(ytest,y_pred_classes)
```

True \ Pre	dicted	0	1	1	1
0   1	87   53		63 132		



#MODEL 2

```
In [77]: X train flattened = np.array([feature.flatten() for feature in X train extracted])
         X test flattened = np.array([feature.flatten() for feature in X test extracted])
         X train flattened.shape, X test flattened.shape
Out[77]: ((1337, 12700), (335, 12700))
In [78]: random forest = RandomForestClassifier(n estimators=100, random state=42)
         # Train the classifier
         random forest.fit(X train flattened, ytrain)
         # Make predictions on the test set
         y pred 1 = random forest.predict(X test flattened)
         # Calculate accuracy
         accuracy = accuracy score(ytest, y pred 1)
         precision = precision_score(ytest, y_pred_1, average='weighted')
         recall = recall_score(ytest, y_pred_1, average='weighted')
         f1 = f1 score(ytest, y pred 1, average='weighted')
         print(f"Accuracy: {accuracy}")
         print(f"Precision: {precision}")
         print(f"Recall: {recall}")
         print(f"F1-score: {f1}")
         Accuracy: 0.573134328358209
         Precision: 0.5655104947471202
         Recall: 0.573134328358209
         F1-score: 0.5596455967400165
In [83]: display confusion matrix(ytest, y pred 1)
         |True \ Predicted|
                         57
                                    93
                         50
                                   135
```

```
In [84]: # Assuming y_true contains the true binary labels (0 and 1) and y_pred contains the predicted labels
sensitivity_1, specificity_1 = calculate_sensitivity_specificity(ytest, y_pred_1)

print(f"Sensitivity (True Positive Rate, TPR): {sensitivity_1}")
print(f"Specificity (True Negative Rate, TNR): {specificity_1}")

Sensitivity (True Positive Rate, TPR): 0.7297297297297297
Specificity (True Negative Rate, TNR): 0.38

#MODEL 3
```

```
In [87]: from tensorflow.keras.layers import Input, Flatten, Dense
         from tensorflow.keras.models import Model
         from tensorflow.keras.applications import ResNet50
         from tensorflow.keras.optimizers import Adam
         # Load pre-trained ResNet50 model
         resnet base = ResNet50(weights='imagenet', include top=False, input shape=(127, 100, 3))
         # Freeze ResNet Layers so they won't be trained
         for layer in resnet base.layers:
             layer.trainable = False
         # Add custom layers for classification on top of ResNet50
         x = Flatten()(resnet base.output)
         x = Dense(256, activation='relu')(x)
         predictions = Dense(2, activation='softmax')(x) # 'num classes' is the number of output classes
         # Create the model
         model 4 = Model(inputs=resnet base.input, outputs=predictions)
         # Compile the model
         model 4.compile(optimizer=Adam(), loss='sparse categorical crossentropy', metrics=['accuracy'])
         # Now you can train or evaluate the model using your data
```

```
In [89]: from sklearn.metrics import accuracy_score, precision_score, recall_score, f1_score
# Assuming you have X_train, y_train, X_test, y_test after data preprocessing and splitting
# Train your modified ResNet-50 V2 model
history = model_4.fit(X_train_rgb, ytrain, epochs=13, batch_size=32, validation_data=(X_test_rgb, ytest))
```

```
Epoch 1/13
42/42 [============== ] - 42s 822ms/step - loss: 2.9671 - accuracy: 0.5183 - val loss: 0.9451 - val ac
curacy: 0.5522
Epoch 2/13
curacy: 0.5821
Epoch 3/13
curacy: 0.6000
Epoch 4/13
curacy: 0.5493
Epoch 5/13
curacy: 0.5194
Epoch 6/13
curacy: 0.5134
Epoch 7/13
curacy: 0.6149
Epoch 8/13
curacy: 0.5194
Epoch 9/13
curacy: 0.5164
Epoch 10/13
curacy: 0.6179
Epoch 11/13
42/42 [=============== ] - 32s 775ms/step - loss: 0.6245 - accuracy: 0.6350 - val loss: 0.6728 - val ac
curacy: 0.6388
Epoch 12/13
curacy: 0.6269
Epoch 13/13
curacy: 0.5284
```

```
In [34]: y pred 2 = model 4.predict(X test rgb)
         y pred classes 2 = np.argmax(y pred 2, axis=1) # Convert probabilities to class labels
         # Calculate evaluation metrics
         accuracy = accuracy score(ytest, y pred classes 2)
         precision = precision score(ytest, y pred classes 2, average='weighted')
         recall = recall score(ytest, y pred classes 2, average='weighted')
         f1 = f1 score(ytest, y pred classes 2, average='weighted')
         # Print the evaluation metrics
         print(f"Accuracy: {accuracy}")
         print(f"Precision: {precision}")
         print(f"Recall: {recall}")
         print(f"F1-score: {f1}")
         11/11 [======== ] - 9s 547ms/step
         Accuracy: 0.6029850746268657
         Precision: 0.6295732474937124
         Recall: 0.6029850746268657
         F1-score: 0.5291781909116685
In [35]: display confusion matrix(ytest, y pred classes 2)
         |True \ Predicted|
                         25
                                   121
                         12
                                   177
In [36]: # Assuming y true contains the true binary labels (0 and 1) and y pred contains the predicted labels
         sensitivity 2, specificity 2 = calculate sensitivity specificity(ytest, y pred classes 2)
         print(f"Sensitivity (True Positive Rate, TPR): {sensitivity 2}")
         print(f"Specificity (True Negative Rate, TNR): {specificity 2}")
         Sensitivity (True Positive Rate, TPR): 0.9365079365079365
         Specificity (True Negative Rate, TNR): 0.17123287671232876
         #MODEL 4(Model 1 features extracted and a random classifier)
```

```
In [44]: from sklearn.ensemble import RandomForestClassifier
    from sklearn.metrics import accuracy_score

# Extract features from the second last fully connected layer
    feature_extractor_m1 = tf.keras.Model(inputs=model1.inputs, outputs=model1.layers[-2].output)

# Predict the features for train and test data
    train_features_m1 = feature_extractor_m1.predict(Xtrain)
    test_features_m1 = feature_extractor_m1.predict(Xtest)

# Model features using Random Forest classifier
    rf_classifier_m1 = RandomForestClassifier(n_estimators=100, random_state=42)
    rf_classifier_m1.fit(train_features_m1, ytrain)

# Predict using the Random Forest classifier
    predictions_m1 = rf_classifier_m1.predict(test_features_m1)

# Calculate accuracy
    accuracy_m1 = accuracy_score(ytest, predictions_m1)
    print("Accuracy of Random Forest classifier on extracted features model 1:", accuracy_m1)
```

Accuracy of Random Forest classifier on extracted features model 1: 0.5791044776119403