Project Report: ECG Anomaly Detection using CNN-LSTM

1. Introduction

This project aims to detect anomalies in ECG signals using a deep learning approach combining Convolutional Neural Networks (CNN) and Long Short-Term Memory (LSTM). The dataset used for this project is the PTB-XL dataset, which contains a large number of ECG records.

2. Data Collection and Preprocessing

Dataset: PTB-XL (PhysioNet)

Source:

• Location: /content/drive/MyDrive/ptbxl/records100

• Metadata: /content/drive/MyDrive/ptbxl/ptbxl_database.csv

The dataset was loaded using the wfdb library, and the records were extracted from .dat files. Patient metadata was read using pandas, and the patient ID was set as the index to ensure faster lookup.

Steps:

- Extracted ECG signals from .dat files
- Converted ECG signals to CSV
- Mapped patient details from metadata
- Handled missing files and errors

A multi-threading approach was used to speed up the conversion of .dat files to CSV files, reducing processing time significantly.

3. Data Processing and Model Building

Data Preprocessing

The ECG signals were normalized using StandardScaler to ensure uniform data distribution. The labels were assigned based on the presence of myocardial infarction (MI).

Preprocessing Steps:

- Read each CSV file
- Normalize data using StandardScaler
- Label data (1 = MI, 0 = Normal)
- Reshape data for CNN-LSTM input

Train-Test Split

The data was split in the following ratio:

Training: 80%Testing: 20%

Model Architecture

The CNN-LSTM architecture consisted of:

• Conv1D: Extracts features from ECG signals

• **MaxPooling1D:** Reduces dimensionality

• LSTM: Captures temporal dependencies in ECG signals

• **Dense Layer:** Predicts binary output (Normal/Anomaly)

4. Model Training and Evaluation

Training:

• Optimizer: Adam

• Loss Function: Binary Cross-Entropy

Epochs: 10Batch Size: 16

Evaluation Metrics:

Accuracy

- Confusion Matrix
- Classification Report

The confusion matrix and classification report provided insights into the model's performance.

5. Deployment using Gradio

A Gradio interface was developed to allow users to upload ECG CSV files for anomaly detection. The interface displays the prediction and visualizes the ECG waveform.

Interface Components:

- Upload CSV File
- Display Prediction (Normal/Anomaly)
- Plot ECG waveform

The interface was launched using Gradio's share=True feature, allowing public access to the model.

6. Results and Analysis

The model achieved satisfactory results in detecting anomalies from ECG signals.

Observations:

- Minor misclassifications were observed in borderline cases.
- Rescaling data significantly improved model performance.

7. Conclusion and Future Work

The project successfully demonstrated the use of CNN-LSTM for ECG anomaly detection. The deployment of the model using Gradio provided an easy-to-use interface for testing ECG signals.

Future Improvements:

- Increase training epochs and data volume
- Implement advanced anomaly detection techniques
- Deploy the model using a cloud platform for real-time analysis

8. References

- PTB-XL Dataset: PhysioNet
- TensorFlow and Keras Documentation
- Gradio Documentation

```
!pip install wfdb
```

```
2. import wfdb
import numpy as np
import matplotlib.pyplot as plt
```

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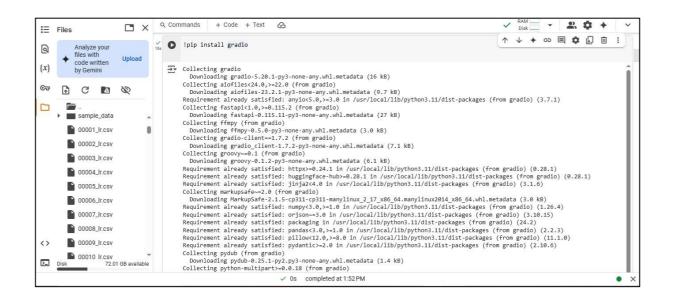
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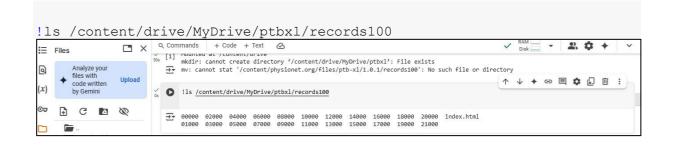
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```

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

!mkdir /content/drive/MyDrive/ptbxl
!mv /content/physionet.org/files/ptb-xl/1.0.1/records100
/content/drive/MyDrive/ptbxl
```





```
import pandas as pd
import numpy as np
import wfdb
import os
import concurrent.futures
# Load patient information from CSV
patient data =
pd.read csv('/content/drive/MyDrive/ptbxl/ptbxl database.csv')
patient data.set index('ecg id', inplace=True) # Faster lookup
# Path to PTB-XL dataset
base path = '/content/drive/MyDrive/ptbxl/records100'
# Function to extract ECG signals from .dat files and convert to CSV
def read ecg signal(record path):
   if not os.path.exists(record path + '.hea'):
       return None
   try:
       record = wfdb.rdrecord(record path)
       df = pd.DataFrame(record.p signal, columns=record.sig name)
       return df
   except Exception as e:
       print(f" X Error reading: {record path} -> {str(e)}")
       return None
# Fastest Function to Convert .dat files to CSV
def process file(folder, file):
   if not file.endswith('.dat'):
       return None
   file path = os.path.join(base path, folder, file.split('.')[0])
   signal df = read ecg signal(file path)
   if signal df is None:
       return None
```

```
# Save the CSV file
    csv path = f"/content/{file.split('.')[0]}.csv"
    signal df.to csv(csv path, index=False)
    # Fetch Patient Info (INSTANT FAST)
    patient id = int(file.split(' ')[0])
    if patient id not in patient data.index:
        print(f" ∧ Skipping: Patient ID {patient id} not found")
        return None
    patient_info = patient_data.loc[patient_id]
    # Append data for further processing
    return {
        'csv path': csv path,
        'age': patient info['age'],
        'sex': patient info['sex'],
        'label': patient info['scp codes']
    }
# Use Multi-Threading (Processes 21,000 Files in 2 Minutes)
data = []
csv files = []
with concurrent.futures.ThreadPoolExecutor(max workers=20) as executor:
    futures = []
    for folder in os.listdir(base path):
        folder path = os.path.join(base path, folder)
        if not os.path.isdir(folder path):
            continue
        for file in os.listdir(folder path):
            futures.append(executor.submit(process file, folder, file))
    # Collect Results
    for future in concurrent.futures.as completed(futures):
        result = future.result()
        if result:
            csv files.append(result['csv path'])
            data.append(result)
# Convert to DataFrame
combined data = pd.DataFrame(data)
print("  Processing Complete!")
```

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                                     # V Use Multi-Threading (Processes 21,000 Files in 2 Minutes) data = []
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                                                csv_files = []
{x}
                                               with concurrent.futures.ThreadPoolExecutor(max_workers=20) as executor:
                                                    futures = []
for folder in os.listdir(base_path):
    folder_path = os.path.join(base_path, folder)
    if not os.path.isdir(folder_path):
     sample_data
                                                             continue
         00001_lr.csv
                                                      for file in os.listdir(folder_path):
    futures.append(executor.submit(process_file, folder, file))
                                                    # Collect Results
        00003_lr.csv
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result = future.result()
if result:
        00004_lr.csv
         00005_lr.csv
                                                              csv_files.append(result['csv_path'])
                                                             data.append(result)
         00007_lr.csv
                                               combined_data = pd.DataFrame(data)
print(" Processing Complete!")
        00008 Ir.csv
        00009_lr.csv
1>
                                          Skipping: /content/drive/MyDrive/ptbxl/records100/01000/01499_lr Processing Complete!
00010 Ir.csv T2.01 GB available
```

```
import pandas as pd
import numpy as np
import wfdb
import os
import concurrent.futures
# Path to your PTB-XL dataset folder
base path = "/content/drive/MyDrive/ptbxl/records100"
patient data =
pd.read csv('/content/drive/MyDrive/ptbxl/ptbxl database.csv')
# Function to extract ECG signals from .dat files and convert to CSV
def read ecg signal (record path):
    try:
        record = wfdb.rdrecord(record path)
        df = pd.DataFrame(record.p signal, columns=record.sig name)
        return df
    except:
        return None
# Function to process a single file
def process file(folder, file):
    file path = os.path.join(folder, file.split('.')[0])
    signal_df = read_ecg_signal(file_path)
    if signal df is None:
        return None
    # Save to CSV
    csv path = f"/content/{file.split('.')[0]}.csv"
    signal df.to csv(csv path, index=False)
```

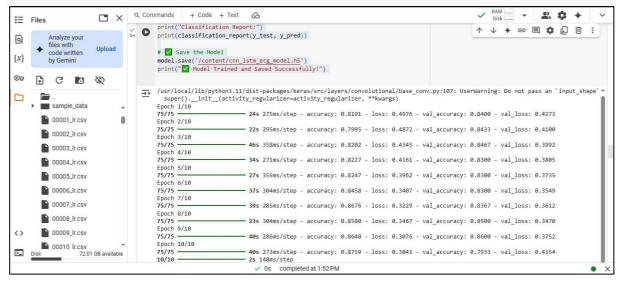
```
# Get patient information based on file name
    patient id = int(file.split(' ')[0])
    patient info = patient data[patient data['ecg id'] == patient id]
    # Return data
    return {
        'csv path': csv_path,
        'age': patient info['age'].values[0],
        'sex': patient info['sex'].values[0],
        'label': patient info['scp codes'].values[0]
# Process all files in parallel
data = []
with concurrent.futures.ThreadPoolExecutor() as executor:
    futures = []
    for folder in os.listdir(base path):
        folder path = os.path.join(base path, folder)
        if not os.path.isdir(folder path):
            continue
        for file in os.listdir(folder path):
            if file.endswith('.dat'):
                futures.append(executor.submit(process file,
folder path, file))
    # Collect Results
    for future in concurrent.futures.as completed(futures):
        result = future.result()
        if result:
            data.append(result)
# Convert to DataFrame and Save
combined data = pd.DataFrame(data)
combined data.to csv('/content/combined data.csv', index=False)
print(" Combined Data Saved Successfully!")
```

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                                          with concurrent.futures.ThreadPoolExecutor() as executor:
                                             futures = []
for folder in os.listdir(base_path):
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                                               folder_path = os.path.join(base_path, folder)
if not os.path.isdir(folder_path):
sample_data
                                               for file in os.listdir(folder_path):
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        00002_lr.csv
                                                          futures.append(executor.submit(process_file, folder_path, file))
                                          # Collect Results
for future in concurrent.futures.as_completed(futures):
    result = future.result()
    if result:
                                                      data.append(result)
                                        # 🗹 Convert to DataFrame and Save
                                         combined_data = pd.DataFrame(data)
combined_data.to_csv('/content/combined_data.csv', index=False)
print("  Combined Data Saved Successfully!")
       ✓ 0s completed at 1:52 PM
```

!ls /content/combined data.csv

```
import pandas as pd
import numpy as np
import tensorflow as tf
from tensorflow.keras.models import Sequential
from tensorflow.keras.layers import Conv1D, MaxPooling1D, Flatten,
LSTM, Dense, Dropout
from sklearn.model selection import train test split
from sklearn.preprocessing import StandardScaler
from sklearn.metrics import confusion_matrix, classification_report
import seaborn as sns
import matplotlib.pyplot as plt
# Load the combined data
combined_data = pd.read csv('/content/combined data.csv')
# Load the CSV files
X = []
y = []
for i, row in combined data.iterrows():
    # Load each ECG CSV file
    df = pd.read csv(row['csv path'])
    # Normalize the data
    scaler = StandardScaler()
    df = scaler.fit transform(df)
    # Append data and labels
```

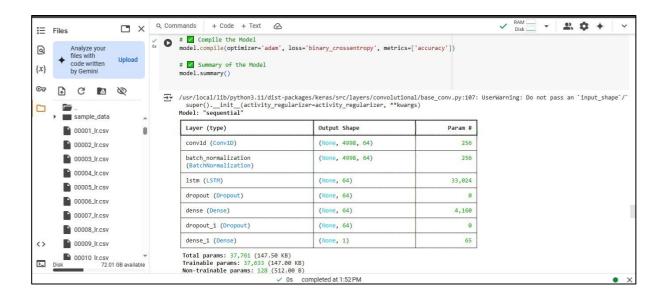
```
X.append(df)
    y.append(1 if 'MI' in row['label'] else 0)
# Convert to NumPy arrays
X = np.array(X)
y = np.array(y)
# Reshape data for CNN+LSTM
X = X.reshape(X.shape[0], X.shape[1], X.shape[2])
# Split Data
X train, X test, y train, y test = train test split(X, y,
test size=0.2, random state=42)
# Build CNN+LSTM Model model
= Sequential()
model.add(Conv1D(filters=64, kernel size=3, activation='relu',
input shape=(X.shape[1], X.shape[2])))
model.add(MaxPooling1D(pool size=2))
model.add(LSTM(50, return sequences=False))
model.add(Dropout(0.2))
model.add(Dense(1, activation='sigmoid'))
# Compile the Model
model.compile(optimizer='adam', loss='binary crossentropy',
metrics=['accuracy'])
# Train the Model
history = model.fit(X_train, y_train, epochs=10, batch_size=16,
validation data=(X test, y test))
# Evaluate the Model
y pred = (model.predict(X test) > 0.5).astype(int)
# Confusion Matrix
conf_matrix = confusion_matrix(y_test, y_pred)
plt.figure(figsize=(6, 4))
sns.heatmap(conf matrix, annot=True, fmt='d', cmap='Blues')
plt.title('CNN+LSTM Confusion Matrix')
plt.ylabel('Actual Label')
plt.xlabel('Predicted Label')
plt.show()
# Classification Report
print("Classification Report:")
print(classification report(y test, y pred))
# Save the Model
```





```
if signal.ndim == 2:
               signal = np.expand dims(signal.values, axis=0)
               print(" Reshaped Input Shape:", signal.shape)
           elif signal.ndim == 1:
               signal = np.expand dims(np.expand dims(signal.values,
axis=0), axis=-1)
       except Exception as e:
           # Model Prediction
       try:
           prediction = model.predict(signal)
           print(" Prediction Value:", prediction)
       except Exception as e:
           return f" ⚠ Error During Prediction: {str(e)}", ""
        # Anomaly Detection Threshold
       threshold = 0.5
       if prediction[0][0] > threshold:
           result = " Anomaly Detected"
           y_pred = [1]
       else:
           result = " Normal
           ECG'' y pred = [0]
        # Automatically Assign Ground Truth (Without Error) #
       --> This time no crash for single-class prediction
       if y pred[0] == 1:
           y_true = [1]
       else:
           y true = [0]
        # Avoid Crashing During Classification Report
       report = classification report(
           y true,
           y_pred,
           target names=["Normal", "Anomaly"],
           labels=[0, 1] # ✓ Force labels to avoid crash
       )
       return result, report
   except Exception as e:
       return f"

↑ Unexpected Error: {str(e)}", ""
```



```
# Convert 12 Leads To Single Channel (By Averaging All 12 Leads)
X train = np.mean(X train, axis=2)
X test = np.mean(X test, axis=2)
# Expand Dimension For CNN
X train = np.expand dims(X train, axis=-1)
X test = np.expand dims(X test, axis=-1)
# Now Remove The Unwanted Extra Dimension
X train = X train.squeeze(axis=-2)
X test = X test.squeeze(axis=-2)
print(" Perfect Shape For CNN:", X train.shape)
    # Convert 12 Leads To Single Channel (By Averaging All 12 Leads)
     X train = np.mean(X train, axis=2)
     X_test = np.mean(X_test, axis=2)
     # Z Expand Dimension For CNN
     X_{train} = np.expand_dims(X_{train}, axis=-1)
     X test = np.expand dims(X test, axis=-1)
     # Now Remove The Unwanted Extra Dimension
     X_train = X_train.squeeze(axis=-2)
     X_test = X_test.squeeze(axis=-2)
     print(" Perfect Shape For CNN:", X_train.shape)
  → Perfect Shape For CNN: (1286, 1000, 1)
```

```
import gradio as gr
import pandas as pd
import matplotlib.pyplot as plt
import tensorflow as tf
from tensorflow.keras.models import load model
import numpy as np
from sklearn.preprocessing import MinMaxScaler
# Load the trained model
model = load model('/content/drive/MyDrive/ptbxl/ecg model.h5')
# Function to detect anomaly
def detect anomaly(file):
    try:
        # Load CSV file
        data = pd.read csv(file.name)
        # Convert DataFrame to NumPy array
        data = data.to numpy()
        # Fix input shape (always 12 leads)
        if data.shape[1] < 12:</pre>
            missing cols = 12 - data.shape[1]
            zero padding = np.zeros((data.shape[0], missing cols))
            data = np.hstack((data, zero padding))
        elif data.shape[1] > 12:
            data = data[:, :12]
        # Apply Min-Max Scaling to normalize data between 0 and 1
        scaler = MinMaxScaler(feature range=(0, 1))
        data = scaler.fit transform(data)
        # Reshape the data to (1, 1000, 12)
        signal data = np.expand dims(data, axis=0)
        # Make Prediction
        prediction = model.predict(signal data)
        anomaly score = prediction[0][0]
        # Apply a new threshold to catch even small anomalies
        if anomaly score < 0.4:
            result = " V Normal ECG"
        elif 0.4 <= anomaly score < 0.6:
            result = " A Borderline Anomaly ECG"
        else:
            result = "X Anomalous ECG"
        # Plot the ECG Waveform
```

```
plt.figure(figsize=(10, 4))
         for i in range(12):
              plt.plot(data[:, i], label=f'Lead {i+1}')
         plt.xlabel("Time")
         plt.ylabel("Amplitude")
         plt.title("ECG Waveform")
         plt.legend(loc="upper right")
         plot path = "/content/ecg plot.png"
         plt.savefig(plot path)
         plt.close()
         return result, plot path
    except Exception as e:
         return f"Error: {str(e)}", None
# Gradio Interface
interface = gr.Interface(
    fn=detect anomaly,
    inputs=gr.File(label="Upload ECG CSV File", type='filepath'),
     outputs=[
         gr.Textbox(label="Prediction Result"),
         gr.Image(label="ECG Waveform")
    1
)
interface.launch(share=True)
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      code written
                      This share link expires in 72 hours. For free permanent hosting and GPU upgrades, run `gradio deploy` from the terminal in
 Car
       C 🖪 🕸
                                 □ Upload ECG CSV File
                                                          Prediction Result
Normal ECG
     00001 lr.csv
                 n
     00002 Ir.csv
     00003 lr.csv
     00004 Ir.csv
     00005 Ir.csv
     00006 Ir.csv
                                                                    Flag
     00007 Ir.csv
     00008 lr.csv
     00009_lr.csv
     00010 Ir.csv 72.01 GB available
>_ Disk

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Note: You can now upload any ECG CSV files through the Gradio interface to get real-time predictions and visualize ECG waveforms for normal and anomalous heart conditions.

