ECG Findings (Demo)

Setting Up xLSTMTime: A Comprehensive Guide for Beginners

XLSTM -Time for PTBXL ECG Dataset

1. Download the zip file **xLSTMTime-main.zip** from Google Drive and extract it to the desired location

Google Drive link -

It includes original code from the author of the xLSTM-Time model along with modification required for the PTBXL dataset

The original Github repository for xLSTM-Time can be found here:

https://github.com/muslehal/xLSTMTime

If you want to create a virtual environment then follow step 2 otherwise skip step 2

- 2. Create Virtual Environment & Installing required packages
 - > Open Terminal (you can use vscode terminal or your machine terminal)
 - > Locate to **xLSTMTime-main** using the **cd** command
 - > Run python -m venv .venv
 - > Run **source venv/bin/activate** to activate newly created virtual environment
 - > Inside the virtual environment Run pip install -r requirements.txt
- 3. Installing required packages without a virtual environment
 - > Run **pip install -r requirements.txt** to have the required packages in the global python environment
- 4. Try to run main.py

Understanding Code Structure and Process Flow

all_six_datasets - This folder has datasets which were used by the author. Datasets are in .csv format. To keep it consistent new datasets are all added here inside folders with names defined in *DSETS* in datautils.py. Make sure there is a .csv dataset file inside dedicated folders.

data_preprocessing.py - This file reads ECG signal files from *the PTB-XL-Dataset* folder and creates a .csv file that can be fed to the model.

Understanding the format in which we can feed the data is extremely crucial. First of all, in the data_preprocessing.ipynp file, we are processing the PTBXL dataset, which is also present in the main folder.

The format of the data is called 'expanded form' which was used in the book (Modern Time Series Forecasting with Python) we followed throughout the course. Expanded form for PTBXL:

- * Each ecg_id has 0 999 timesteps because we are using 10s long ECG signal and 100Hz frequency
- * we have total 21799 ecg_ids, all the ecg_ids are stacked up with their 999 timesteps
- * we have a total of 21799 * 1000 = 21799000 rows and 12 columns for 12 leads in our CSV file

main.py - it has all the crucial parameters to set, which is passed to 'args' to model

1. Parameters/hyperparameters

- --dset: dataset name from the list 'DSETS' defined in datautils.py
- --context_points: Sequence length (window size) of time-series data fed into the model. If you expect long-range dependencies, larger values might be better. If patterns are shorter, smaller values might be sufficient and faster to train. A good starting point would be to test values equal to your data's primary periodicity.
- --target_points: Forecast horizon the number of time steps the model predicts into the future. In your Dataset_PTBXL, this corresponds to pred_len. Integer, default: 96. Experimentation: This is determined by your forecasting goal. However, smaller forecast horizons are generally easier to predict.
- --scaler: Scaling method for input data. String, default: 'standard'.
 Options could include 'standard' (StandardScaler), 'minmax'
 (MinMaxScaler), or None (no scaling).
- --features: Specifies which features to use from the dataset 'M' (Multivariate) or 'S' (Single/Univariate).
- --batch_size: Number of samples processed in each iteration of training. Larger batch sizes can lead to faster training but might require more memory. Smaller batch sizes can be more stable but might take longer.
- -num_workers: Number of subprocesses to use for data loading. Helps speed up data loading. Integer, default: 1. Experimentation: Set to the number of CPU cores.
- --patch_len: Length of each patch. Integer, default: 12.
- --stride: Stride between patches. Integer, default: 12. If stride == patch_len, the patches are non-overlapping. If stride < patch_len, the patches overlap.
 - Patching can help the model focus on local patterns and can reduce the computational costs for very long sequences. If you have strong long-range dependencies, patching might hurt performance. If the data is locally smooth, overlapping patches might help.
- --revin: Flag to use RevIN for normalization. Integer, default: 1 (True).
 RevIN is a technique that can help with the instability of training deep
 time series models. If your model trains unstably, RevIN can be very
 helpful.
- --n_epochs: Number of training epochs.
- --is_train: Flag to indicate whether to train the model (1) or run in testing mode (0).
- -n_layers: More layers can capture more complex relationships but can also lead to overfitting and longer training times.
- --d_model: Dimension of the transformer layers (embedding dimension).
- --dropout: Higher dropout can help with overfitting but might slow down training.
- --head_dropout: Head dropout rate. Used to prevent overfitting on heads.

- **2. find_Ir()** This function is defined in main.py and sets up the necessary components for training your xLSTM-Time model (data, model, loss function, callbacks) and then uses a learning rate finder to determine a suitable learning rate before starting the full training process, returns suggested learning rate
- **3. train_func()** This function takes the suggested learning rate as an argument and trains the model by setting a few configurations in the function *get_dls(args)* defined in *datautils.py* and a class defined in *src/data/pred_dataset.py*
- **4. test_func()** This functions saves models in saved_models/-dset (as defined above) and stores predicted, target values in list named *out*, returns *out*
- **5. plot_features_actual_vs_predicted()** Based on number of feature range defined, it plots actual vs predicted values for different features.

datautils.py - it has a list DSETS with names of datasets

It prepares the dataset for training a deep learning model using PyTorch

1. get_dls() - it has a condition for each dataset defined in *DSETS* for checking the current dataset defined in main.py

Each condition block forwards parameters to the Dataset loading class defined in

pred_dataset.py . The get_dls function loads the dataset and validates it by checking the shape of a sample.

Key Components

 Imports: Uses PyTorch for deep learning, NumPy and Pandas for data handling, and custom modules (DataLoaders, Dataset_PTBXL) for dataset processing.

2. Parameters:

- context_points: Past time steps used for prediction.
- target_points: Future time steps to predict.
- batch_size and num_workers: For data batch processing and parallelism.
- o features: Specifies which dataset features to use.

3. Dataset Loading:

 The dataset should be in the path specified in root_path and is loaded from patient.csv.

4. Usage:

- The Params class sets the configuration values.
- The get_dls function returns a DataLoaders object containing the training data.
- After loading, it prints the shape of the first training sample.

5. Customization:

- Modify root_path and data_path for your dataset location.
- Adjust parameters for batch size, features, and time points.

src/data/pred_dataset.py - This file contains classes for datasets to handle the training, testing, validation split, defining features in the dataset, and size/shape of the dataset.

Dataset_PTBXL and Dataset_longterm classes are added to handle these datasets. **Dataset_PTBXL class:**

This class Dataset_PTBXL is a custom implementation of a dataset used for time-series forecasting with ECG data. It inherits from torch.utils.data.Dataset, and is designed to work with PyTorch for training deep learning models. Here's an explanation of its key components:

Constructor (__init__ method)

The __init__ method initializes the dataset, sets various configuration options, and prepares the data for loading. The key arguments are:

- root_path: The directory where the dataset is stored.
- split: The dataset split type (train, val, or test).
- **size**: Defines the sequence length, label length, and prediction length. Default values are set based on 4-hour intervals.
- features: Specifies the features to use ('M' for multiple features, 'S' for single feature).
- target: The target column (usually the column you're trying to predict).
- **scale**: Whether to normalize (scale) the data.
- **timeenc**: Defines how time encoding works (not implemented in this code snippet).
- freq: Frequency of time steps (e.g., seconds, minutes).
- time_col_name: The column name containing timestamps (e.g., 'date').
- use_time_features: Whether to use time-related features for the model.
- train_split & test_split: Ratios for splitting the data into training, validation, and test sets.

Reading and Processing the Data (__read_data__ method)

- Data Loading: The dataset is read from a CSV file located at data_path within the root_path. The data is then processed and split into training, validation, and test sets based on the specified ratios.
- Scaling: If scale=True, the dataset is normalized using StandardScaler. The scaler is fit on the training data and then applied to the entire dataset.

- **Feature Selection**: The features argument determines which columns of the data to use. If 'M', it takes all columns starting from the 5th column (features). If 'S', it only takes the target column.
- **Timestamps**: The time_col_name (e.g., 'date') is converted to a datetime format. Although the time features are not fully implemented in this code, there's an option (timeenc) to add them if needed.

Data Slicing

The dataset is split into windows of data:

- **seq_len**: The length of the input sequence (how many past time steps are used for prediction).
- **label_len**: The length of the label sequence (the length of the true values used for training).
- **pred_len**: The length of the prediction sequence (how many future time steps the model predicts).
- Based on the current split (train, val, test), the data is sliced accordingly.

Indexing (__getitem__ method)

- **Data Retrieval**: For each sample, the __getitem__ method retrieves the input sequence (seq_x) and the output sequence (seq_y), using the slicing window defined earlier.
- If use_time_features is enabled, time features (such as date/time related data) could be added, but they are currently commented out.
- The method returns the data as a PyTorch tensor via the _torch function.

Length of Dataset (__len__ method)

The __len__ method returns the number of possible samples in the dataset (i.e., the number of windows that can be created from the data). It ensures that the sequence length and prediction length are respected.

Inverse Transformation (inverse_transform method)

The inverse_transform method reverses the scaling operation, allowing you to convert the scaled data back to its original form using the fitted scaler.

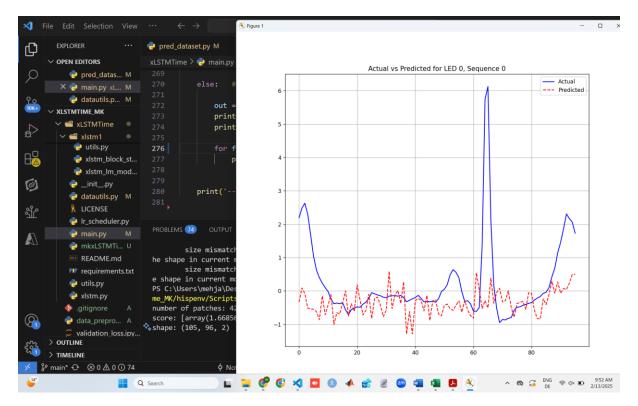
xlstm1/slstm/cell.py - sLSTM needs GPU for faster processing, in this file it can be defined. There are 2 options CUDA or Vanilla. There is no mention of MacBook's MPS framework.

saved_models - Inside this folder, you can find trained models which can be tested by setting *-is_train* parameter in main.py 0.

Model # 2 (xLSTMTime_cw512_tw96_patch12_stride12_epochs10_model2)

ECG data for one patient with 1 Lead and 100 Hz in each second and 10 seconds for one patient therefore 1000 time steps. (no well tuning of parameters)

| Context points | 512 |
|----------------|------|
| Target points | 96 |
| Batch size | 64 |
| Ir | 1e-3 |
| n_layers | 3 |
| d_model | 256 |
| dropout | 0.2 |
| Patch_len | 12 |
| Stride | 12 |



Model # 3 (xLSTMTime_cw100_tw50_patch12_stride12_epochs50_model3)

ECG data for one patient with 1 Lead and 100 Hz in each second and 10 seconds for one patient therefore 1000 time steps. (Hyperparameters Tuned)

Configuration in main.py

```
args = {
    'context_points': 100,
                               # Sequence Length
    'target_points': 50,
                                 # Forecast Horizon
    'batch size': 32,
                                  # Batch Size
    'lr': 1e-3,
                                  # Learning Rate
    'n_layers': 2,
                                  # Number of Layers
    'd model': 128,
                                  # Hidden Size
                                  # Dropout Rate
    'dropout': 0.2,
    'patch_len': 12,
                                  # Patch Length
    'stride': 12,
                                  # Stride Between Patches
}
```

Configuration in pred_dataset.py

```
def init (self, root path, split='test', size=None
```

Why Do We Include 12 Extra Time Steps (label_len)?

Instead of predicting **directly from 100 to 150**, we start seq_y from **time step 88**, not 100.

Reason:

The model needs a transition period to "warm up" before making predictions.

- The first 12 time steps (88 to 99) help the model adjust.
- The next **50 time steps (100 to 149)** are the actual predictions

```
Time steps: 0 1 2 ... 88 89 90 ... 99 100 101 102 ... 149

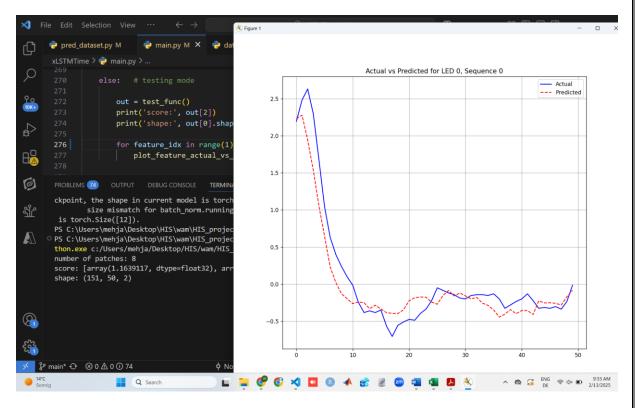
Sequence X: |------|

Sequence Y: |------|
Label part: |-----|

Prediction part: |------|
```

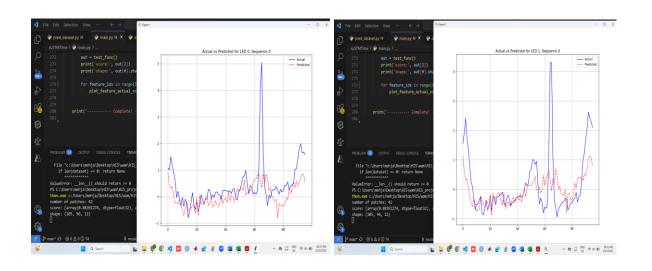
How This Works for All 1000 Rows

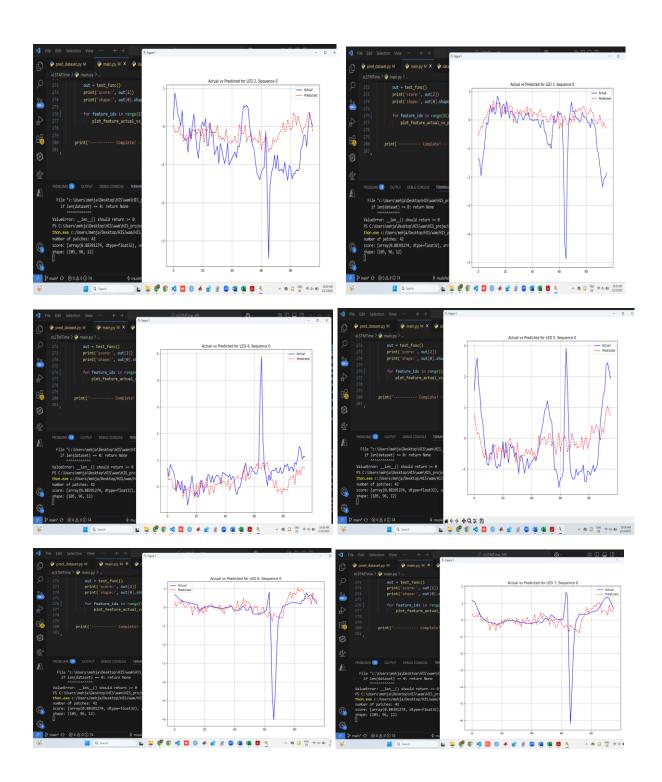
- If you slide the window forward (index=1), your next training set starts at time step 1 instead of 0.
- The model keeps moving forward until it reaches the end of the 1000 rows.
- The last possible starting index would be around 850, since seq_x needs 100 rows.

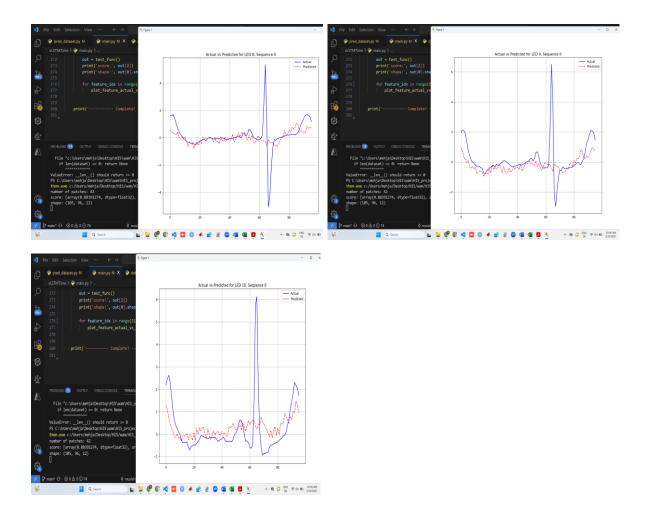


Model # 1 (xLSTMTime_cw512_tw96_patch12_stride12_epochs50_model1)

| Context points | 512 |
|----------------|------|
| Target points | 96 |
| Batch size | 32 |
| Ir | 1e-3 |
| n_layers | 3 |
| d_model | 128 |
| dropout | 0.2 |
| Patch_len | 12 |
| Stride | 12 |
| | |







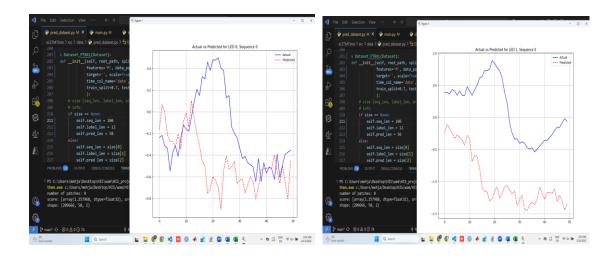
Model # 3 (xLSTMTime_cw100_tw50_patch12_stride12_epochs20_model3)

ECG data for one patient with 12 Leads and 100 Hz in each second and 10 seconds for one patient therefore 1000 time steps.

```
args = {
    'context_points': 100,
    'target_points': 50,
    'batch_size': 32,
    'features' : 'M',
    'lr': 1e-3,
    'n_layers': 2,
    'd_model': 128,
    'dropout': 0.2,
    'patch_len': 12,
    'stride': 12
}
```

def __init__(self, root_path, split='train', size=None,

```
if self.features == 'M' or self.features == 'MS':
      cols_data = df_raw.columns[5:] #takes features from 5th column becuase
LED column starts
      df_data = df_raw[cols_data]
```

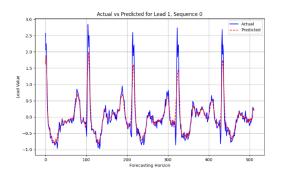


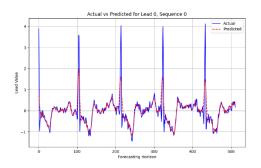
```
Model # 7
  (xLSTMTime_cw512_tw512_patch12_stride12_epochs3_model1)
context_points - 512
target_points -512
batch_size - 64
scaler - 'standard'
n_layers - 3
drop_out - 0.2
```

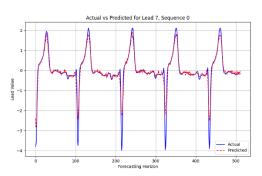
```
target='lead_0', scale=True, timeenc=0, freq='t',
                 use time features=False):
        # size [seq len, label len, pred len]
        if size is None:
            self.seq len = 512
            self.label len = 256
            self.pred_len = 512
        else:
            self.seq len = size[0]
            self.label_len = size[1]
            self.pred_len = size[2]
        assert split in ['train', 'test', 'val']
        type_map = {'train': 0, 'val': 1, 'test': 2}
        self.set type = type map[split]
        self.features = features
        self.target = target
        self.scale = scale
        self.timeenc = timeenc
        self.freq = freq
        self.use_time_features = use_time_features
       self.root_path = root_path
       self.data_path = data_path
        self._read_data_()
    def _read_data (self):
        self.scaler = StandardScaler()
        df_raw = pd.read_csv(os.path.join(self.root_path, self.data_path))
       # Assuming 80% train, 10% val, 10% test split
        #data leak consider
       total_ecgs = df_raw['ecg_id'].nunique()
       train_split = int(0.8 * total_ecgs)
       val_split = int(0.9 * total_ecgs)
        if self.set_type == 0: # train
            df_data = df_raw[df_raw['ecg_id'] < train_split]</pre>
        elif self.set_type == 1: # val
            df_data = df_raw[(df_raw['ecg_id'] >= train_split) &
(df_raw['ecg_id'] < val_split)]</pre>
        else: # test
            df_data = df_raw[df_raw['ecg_id'] >= val_split]
        if self.features == 'M' or self.features == 'MS':
            cols_data = df_data.columns[5:] #from 0-4 cols has static data
from col 5 all the lead values are taken for prediction.
           self.data = df data[cols data].values
```

```
elif self.features == 'S':
        self.data = df data[[self.target]].values
    if self.scale:
        self.scaler.fit(self.data)
        self.data = self.scaler.transform(self.data)
    # Static features
    self.static_data = df_data[['patient_id', 'age', 'sex']].values
def _getitem_(self, index):
   s begin = index * self.seq len
    s_end = s_begin + self.seq_len
    seq_x = self.data[s_begin:s_end]
    seq_y = self.data[s_begin:s_end] # Same as seq_x for autoencoder-like
    static_features = self.static_data[index]
    return _torch(seq_x, seq_y, static_features)
def _len_(self):
    return len(self.data) // self.seq_len
def inverse_transform(self, data):
    return self.scaler.inverse_transform(data)
```

```
number of patches: 42
Mean Squared Error: 0.23169969022274017
Mean Absolute Error: 0.18223978579044342
score: [array(0.23169969, dtype=float32), array(0.18223979, dtype=float32)]
shape: (4333, 512, 12)
```



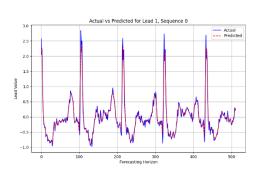


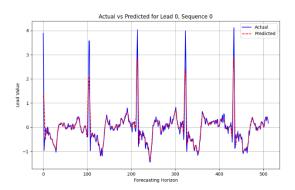


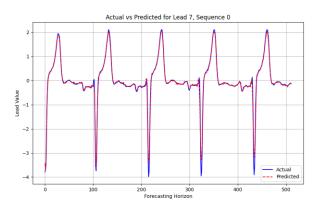
> Model # 8 (xLSTMTime_cw512_tw512_patch12_stride12_epochs10_model1)

| <pre>suggested lr: 0.00014174741629268049 number of patches: 42</pre> |
|--|
| epoch train_loss valid_loss valid_mse valid_mae time Better model found at epoch 0 with valid_loss value: 0.37884962379918646. |
| |
| |
| 0 0.461706 0.378850 0.636115 0.378850 03:59 |
| Better model found at epoch 1 with valid_loss value: 0.2536268659037474. |
| 1 0.303038 0.253627 0.342350 0.253627 03:53 |
| Better model found at epoch 2 with valid_loss value: 0.20673561052172282. |
| 2 0.215587 0.206736 0.246705 0.206736 04:04 |
| Better model found at epoch 3 with valid_loss value: 0.1759277597300153. |
| 3 0.180243 0.175928 0.189917 0.175928 03:59 |
| Better model found at epoch 4 with valid_loss value: 0.1567956681012722. |
| 4 0.158784 0.156796 0.159887 0.156796 04:05 |
| Better model found at epoch 5 with valid_loss value: 0.14053334161972797. 5 0.144197 0.140533 0.139423 0.140533 04:14 |
| Better model found at epoch 6 with valid_loss value: 0.12856590117809544. |
| 6 0.134027 0.128566 0.126795 0.128566 04:08 |
| Better model found at epoch 7 with valid_loss value: 0.12243929840306225. |
| 7 0.127413 0.122439 0.121297 0.122439 04:05 |
| Better model found at epoch 8 with valid_loss value: 0.1199722272431413. |
| 8 0.123773 0.119972 0.119005 0.119972 03:57 |
| Better model found at epoch 9 with valid_loss value: 0.11947969198502048. |
| 9 0.122437 0.119480 0.118637 0.119480 04:09 |
| Complete! |

number of patches: 42
Mean Squared Error : 0.1186368465423584
Mean Absolute Error : 0.11947967112064362
score: [array(0.11863685, dtype=float32), array(0.11947967, dtype=float32)]
shape: (4333, 512, 12)







Model # 5 (xLSTMTime_cw900_tw100_patch24_stride24_epochs20_model5)

ECG data for all patients with all Leads and 100 Hz in each second and 10 seconds for one patient therefore 1000 time steps.

```
args = {
    'context_points': 900,
                                  # Sequence Length (you may adjust based on data)
    'target_points': 100,
                                   # Forecast Horizon
                                 # Batch Size, adjust based on memory
    'batch_size': 32,
    'lr': 1e-3,
                                    # Learning Rate (lowered for deeper model)
                                  # Increase the number of layers to capture more complexity
    'n_layers': 4,
                                 # Hidden Size (increase for more capacity)

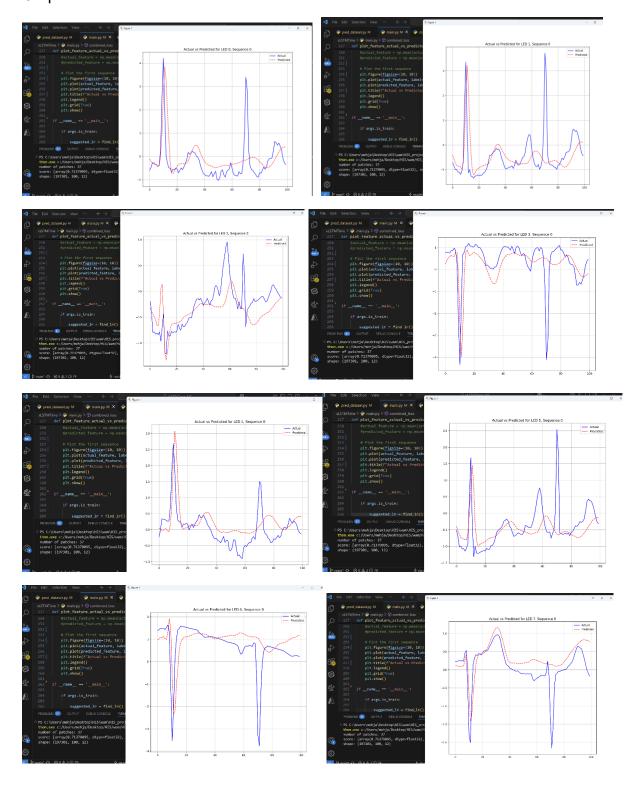
# Dropout Rate (increased for better regularization)

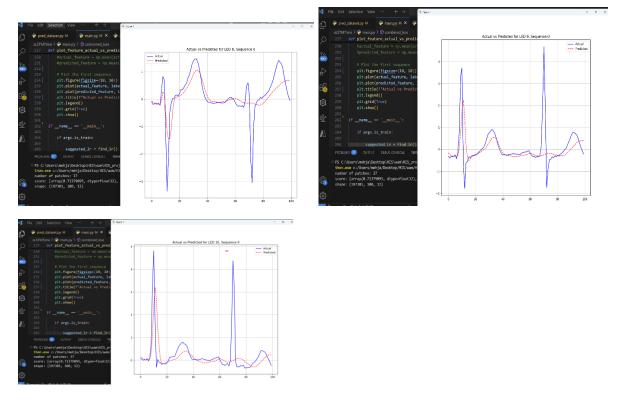
# Patch Length (adjusted for larger data)
    'd_model': 256,
    'dropout': 0.2,
    'patch len': 24,
    'stride': 24,
                                   # Stride Between Patches (adjusted for patch length)
    'epochs': 20,
                                   # number of epochs
```

```
# size [seq len, label len, pred len]
# info
if size == None:
   self.seq len = 100
   self.label len = 12
   self.pred_len = 50
else:
        self.seq len = size[0]
        self.label_len = size[1]
        self.pred_len = size[2]
    # init
    assert split in ['train', 'test', 'val']
    type_map = {'train': 0, 'val': 1, 'test': 2}
    self.set_type = type_map[split]
    self.features = features
    self.target = target
    self.scale = scale
    self.timeenc = timeenc
    self.freq = freq
    self.time_col_name = time_col_name
    self.use_time_features = use_time_features
    # train test ratio
    self.train_split, self.test_split = train_split, test_split
    self.root_path = root_path
    self.data_path = data_path
    self.__read_data__()
def __read_data__(self):
    self.scaler = StandardScaler()
    df_raw = pd.read_csv(os.path.join(self.root_path,
                                       self.data_path))
    df_raw.columns: [time_col_name, ...(other features), target feature]
    cols = list(df_raw.columns)
    #cols.remove(self.target) if self.target
    #cols.remove(self.time col name)
    #df_raw = df_raw[[self.time_col_name] + cols + [self.target]]
    num_train = int(len(df_raw) * self.train_split)
    num_test = int(len(df_raw) * self.test_split)
    num_vali = len(df_raw) - num_train - num_test
```

```
border1s = [0, num_train - self.seq_len, len(df_raw) - num_test -
self.seq len]
        border2s = [num train, num train + num vali, len(df raw)]
        border1 = border1s[self.set type]
        border2 = border2s[self.set type]
        if self.features == 'M' or self.features == 'MS':
            cols_data = df_raw.columns[5:] #takes features from 5th column
becuase LED column starts
            df_data = df_raw[cols_data]
        elif self.features == 'S':
            df_data = df_raw[[self.target]]
        if self.scale:
            train data = df data[border1s[0]:border2s[0]]
            self.scaler.fit(train data.values)
            data = self.scaler.transform(df_data.values)
            data = df data.values
        df_stamp = df_raw[[self.time_col_name]][border1:border2]
        df_stamp[self.time_col_name] =
pd.to_datetime(df_stamp[self.time_col_name])
        self.data_x = data[border1:border2]
        self.data_y = data[border1:border2]
        #self.data_stamp = data_stamp
    def getitem_(self, index):
        s_begin = index
        s_end = s_begin + self.seq_len
        r_begin = s_end - self.label_len
        r_end = r_begin + self.label_len + self.pred_len
        seq_x = self.data_x[s_begin:s_end]
        seq_y = self.data_y[r_begin:r_end]
        #seq_x_mark = self.data_stamp[s_begin:s_end]
        #seq_y_mark = self.data_stamp[r_begin:r_end]
        if self.use_time_features: return _torch(seq_x, seq_y) #removed
seq_x_mark and seq_y_mark
        else: return _torch(seq_x, seq_y)
    def __len__(self):
        return len(self.data_x) - self.seq_len - self.pred_len + 1
    def inverse_transform(self, data):
        return self.scaler.inverse transform(data)
```

Output



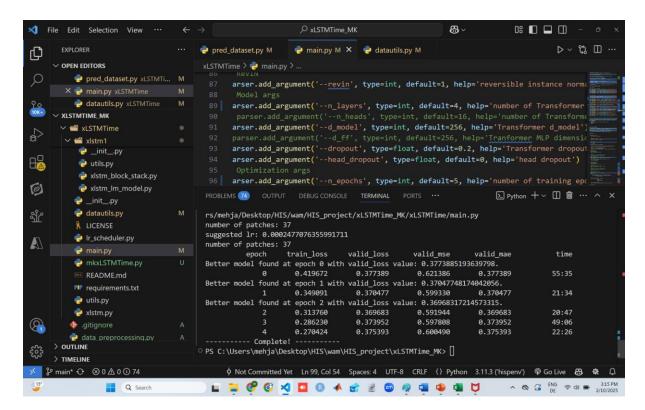


Model # 6:

Dataset: MIT-BIH Long-Term ECG Database.

The MIT-BIH Long-Term ECG Database is a collection of 7 long-duration electrocardiogram (ECG) recordings (14 to 22 hours each), with manually reviewed beat annotations.

> All the parameters are exactly the same as for above model.



Output:

