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Meta-Learning for Enhanced Disease Prediction from EHR Data

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Abstract-The healthcare sector is becoming more dependent on electronic health records (EH for disease forecasting, risk evaluation, and mortality analysis. Although AI-driven models have en ced disease prediction, they frequently focus on common diseases and face difficulties with new or Furthermore, these models require large datasets for better accuracy, posing challenges in dive se or lin ed data scenarios. To solve these issues, this research proposes a novel Long Short-Terr Men from EHRs. The Attention network-based meta-learning framework for prediction tasks using ti framework is designed to address challenges such as limited sample sizes, balance labels, nd the ability to predict unseen diseases. The proposed model is capable of handling mu asks related to irregular 516 patterns and anomalies in time-series signals. The meta-learning approach ena s the system to leverage knowledge from previous tasks, enhancing its ability to predict new and p ious unseen diseases from ECG data. The proposed LSTM-Attention model is evaluated against endonal models like Support Vector Machine (SVM), Random Forest (RF), and XGBoost. Experimer al ult demonstrate that the proposed (in) model outperforms these models, achieving superior performance edicting HRV, arrhythmia, and chieves the highest accuracy (0.92), precision abnormalities from ECG signals. The LSTM-Atten de (0.90), recall (0.91), F1 score (0.91), and ROC-AU the prediction time for the proposed model 0.93). N breov is 95 seconds, significantly faster than other mod

Keywords—*Electronic Health Record; Meta-Learn*, *Electrocardiogram; LSTM with Attention; Disease* Prediction; Accuracy.

I. INTRODUCTION

hth Records (EHR) have received a lot of attention in healthcare analytics Continuous patient Electronic recently [1]. Many studies aducted to create strategies for predicting clinical risks such as death, en k hospital readmission, the nset of a thronic disease, and deterioration of an existing condition. The reasons for this include: 1 the yzing patient EHRs due to factors such as noise, sparsity, and inconsistency, and 2) the ne or relia e health risk prediction models to assist clinical decision-makers in identifying potential on so that patients can receive better care. In response to these challenges, a variety of eat ave been developed, ranging from more standard methods to Deep Learning (DL) compute hms alg models [2

A straiffunt difference between healthcare challenges and applications in domains such as robot vision, such evaluation, and machine translation is the scarcity of available sample datasets, as well as the high cost or inability to collect additional samples [3]. Each data sample is paired with a specific patient in the context of personalized patient risk prediction, which aims to forecast a specific clinical risk for each individual. With a narrower focus on a certain condition, the already small world population of 7.5 billion people decreases even further. Unfortunately, there is only a limited quantity of patient data available in a certain EHR corpus for training a risk prediction model. Furthermore, the clinical dangers being addressed are extremely detailed. A lack of comprehensive understanding of the biochemical pathways underlying the majority of deadly diseases

complicates the design of effective treatments. To develop models that can reliably predict clinical risk, it is essential to maximize the use of limited patient samples alongside existing knowledge about clinical risk and predictive models. This study specifically focuses on predicting illness using ECG. ECGs provide a wealth of critical physiological information for diagnosing a wide range of cardiac conditions. Their intricate patterns carry vital insights, yet understanding these signals often requires highly trained specialists; even experienced doctors may struggle to identify subtle trends or apply their knowledge to new or challenging patients.

A. Current Challenges and Proposed Solution

Recognizing the complexity of ECG analysis underscores the need for advanced computational tech ques to streamline and enhance diagnostic accuracy. The traditional Machine Learning (ML) pipeline for diagnosis involves two main steps: feature extraction and model development. Initial meth ds p focused on using a single characteristic for diagnosis. However, these standard appropriate nave p ven insufficient due to the high level of manual involvement and the requirement for speized To address these limitations, DL, particularly convolutional neural networks ome powerful alternatives for disease prediction. These networks automate feature extractihize penormance, thus and op minimizing manual intervention in parameter selection. However, DL mo equire large datasets for accurate predictions and are often applicable only for identifying specific disease They are not suitable for predicting rare diseases or conditions with minimal samples.

To address these challenges, a meta-learning approach is propored in this work. The LSTM-Attention model is chosen as the base model for meta-learning. The proport 1 system approach set yzes ECG data and is capable of predicting HRV, arrhythmia, and other abnormalities in CG and ings. This model is beneficial for predicting rare diseases from ECG signals, thereby improving patient health to facilitating appropriate treatment.

B. Problem Statement and Research Contribution

Identifying diseases from ECG requires more data and is affected by data quality issues such as noise, sparsity, and inconsistency. Traditional nethods fail to identify rare diseases due to limited datasets. This paper proposes a solution that overcomes the scallences by using meta-learning combined with an LSTM-Attention model. The key contribution of the searches

- The meta-learning an area of utilized to predict HRV, arrhythmia, and other abnormalities using a single model from hinimal CG data.
- An LSTM-Attention QL model is proposed to capture temporal features from ECG data, with attention mechanism incomponent to identify the most significant features for enhanced prediction accuracy.
- The propered LSTM. Attention model is compared with traditional models such as SVM, XGBoost, and RF rateg quark metrics and execution time.
- The EHR late is sourced from MIMIC-III, which is highly unstructured and underwent extensive preproceeding to enhance the accuracy of DL-based meta-learning prediction.

The research paper is organized as follows: Section I details the importance of disease prediction from EHR, talking es in current methods, and the need for meta-learning. Section II discusses recent research work on EHR data. Section III covers the theoretical concept of meta-learning and the LSTM-Attention model. Section IV discusses the experimental setup, data used, and outcomes of the proposed model in disease prediction. Section V concludes the research with future work.

II. RELATED WORK

Various risk prediction models have been developed using ML, DL, and EHR data. Some of the recent works in this area are highlighted in this section. Using feature extraction, the study [4] proposed an EHR risk prediction model. By initially extracting structured text from EHR data using Natural Language Processing, the researchers [5] recommend a system that employs ML techniques to categorize the text as an indicator of "good" or "bad" quality and then use it for prediction. Logistic regression and SVM were utilized as ML models. The research [6] proposed a system consisting of two models. The initial model included the development of interoperable EHR systems with a uniform database structure. Module two covered tasks including clearing and retrieving data from the EHR system, as well as evaluating and forecasting data. Using the proposed decision support system, the proposed system's accuracy in forecasting diabetes disease and the EHR system's interoperability were assessed.

The study [7] developed a multi-task learning (MTL) model to make clinical prediction timeries EHR data. To demonstrate that MTL systems can overcome task imbalance and interdy [8] nce, tr compares their performance to that of traditional single-task models using pren al È dataset. With uniform input durations and variables, these models may be applied to all parents in EHR system. Transfer learning was employed in one study [9] to address the diminishing data proble personalized models. The study [10] aimed to increase disease prediction accuracy by utilizing EHR temporal ta through a new hybrid DL architecture. The architecture incorporates both CNN and LSTM networks indings support the notion that predictive model development should shift toward including com ex n ral network topologies, potentially leading to more personalized models. The paper [11] proposes hat integrates LSTM and Graph Neural Networks (GNNs) to forecast opioid overdose risks, util illness development and patient ng ten interaction graphs. The model's dependence on ΕĤ ata may raise privacy issues, and although ensi ties for actors in real-time scenarios. The research interpretability has improved, it may still prov e diffiç [12] examines DL techniques for clinical decision port, utilizing electronic health record data to predict illness stages, detect genetic markers, and anticipate hepitalization requirements. Although DL provides great accuracy, it remains constrained by the availability and quality of labeled data, as well as the complexity of integrating various electronic health reord for nats.

The study [13] employs LSTM t ctured EHR data for the automated prediction of surgical site te str infections, surpassing convention model h as random forests in both precision and area under the ROC curve. Although LSTM mod hib xcellent accuracy, they necessitate extensive, pristine, and meticulously annotated datasets, which can be fficult to acquire consistently across various healthcare environments. The lel that integrates LSTM and machine learning techniques to predict newresearch [14] proposes a h rid m onset deliriu patient data, surpassing the performance of conventional models such as logistic ed 🐧 regressio tGBM. he efficacy is significantly reliant on the quality of EHR data and feature selection, and and the m may ncounter difficulties with infrequent or unrecognized conditions. Paper [15] suggests extractin cting lung cancer from EHR datasets using a DL architecture combined with Natural and essing (NLP). The text mining model can automatically forecast occurrences from input datasets, La guage l otimal cancer predictions. The model was tested in a new context to see how well it performs and allow for nt DL with NLP is to different datasets. The evaluation demonstrated that the prediction accuracy z restr as honer than that of existing methods. However, there are some limitations in the DL models. The scarcity of labeled datasets and datasets with long-tailed class distributions complicates a wide range of medical tasks. Because general practitioners may be unfamiliar with rare diseases and struggle to differentiate between them, an AI-based decision-making model could help improve diagnostic accuracy. In recent research, many algorithms have been proposed for handling EHR data and predicting diseases. However, there remains a gap between research and real-time implementation. Proposed models often struggle with unseen data, rare

diseases, and new conditions. Most studies use data collected from the internet, which is typically sourced from specific groups of people within certain locations or age ranges. This limitation makes such data insufficient for deploying models in clinical settings. The research aims to address these problems using an LSTM-Attention-based meta-learning approach.

III. THEORETICAL BACKGROUND

This section details the theoretical foundation of two important concepts in this research: meta-learning and LSTM-Attention networks. It also explains how to integrate the LSTM-Attention model into the meta-learning framework. The nomenclature used in the meta-learning and LSTM-Attention networks is provided in Appendix 1 for reference.

A. Meta-Learning

In this case, the model is designed to learn from a limited number of samples. The key co learning is to build on prior knowledge rather than starting from scratch to achiev []. This approach draws upon foundational concepts from supervised machine learning. t accepts an e is a odeľ observation x and assigns it a label y. D represents the training data, while ϕ s the model parameters. Training is an optimization challenge in supervised learning, to maximize the elihood of the parameters based on the D : $\arg \max_{\phi} \log p(\phi|D)$. The issue is reframed as the determined t of the limit of the parameter's marginal probability: $\arg \max_{\phi} \log p(\phi)$ and the data's m morobability given the parameters: $\arg \max_{\phi} \log p(D|\phi)$. A regularizer, such as $\log p(\phi)$ (e.g., weigh dec an be used, and optimization is performed over the dataset with $\sum_{i} \log p(y_i | x_i, \phi)$. Equations, pervised learning for optimizing ribe model parameters based on task-specific data.

$$\arg \max_{\phi} \log p(\phi|D) = \arg \max_{\phi} \log \frac{p(\phi)}{n(D)}$$
(1)

$$q \max_{\phi} \log p(\phi|D) \tag{2}$$

$$= \arg \max_{\phi} \log p(D|\phi) + \log p(\phi)$$
(3)

$$= \arg \max_{\phi} \sum_{i} \log p(y_i | x_i, \phi) + \log p(\phi)$$
(4)

Given $D = \{(x_1, y_1), \dots, (x_k, y_k)\}$ where the enotes the input and y_i denotes the corresponding labels, metalearning integrates prior expressionce with limited new data inputs. To merge existing knowledge with new data points, the problem despress d in Equation (5)

$$\mathcal{L}_{\phi} \log p(\phi|D, D_{meta-train}) \tag{5}$$

sents prior or meta-training data. Maintaining D_{meta-train} is difficult due to Wher emands. Meta-learning characterizes the meta-training dataset through meta-parameters significant mory obtaine by θ $\arg \max_{\theta} p(\theta | D_{meta-train})$. The meta-parameters θ , derived from $D_{meta-train}$, encapsulate prior infor ion essential for the rapid execution of new tasks. This task becomes a maximum likelihood issue: a log_ $(D, D_{meta-train})$. Using prior meta-training data, the goal is to optimize the likelihood of the arg oncerning the new data. The likelihood process is regarded as an integration of the θ . To estimate irame gration, a point estimate θ^* for the θ is used. Meta-training $p(\theta^*|D_{meta-train})$ involves acquiring metaparameters from existing meta-training data, while adaptation $p(\phi|D, \theta^*)$ focuses on deriving parameters for a novel task using both new data and established meta-parameters. Equations (6-10) describe meta-learning for optimizing model parameters using prior knowledge from meta-training data to enable fast adaptation to new tasks.

$$\log p(\phi|D, D_{meta-train}) = \log \frac{p(\phi, D, D_{meta-train})}{p(D, D_{meta-train})}$$
(6)

$$= \log \int \frac{p(\phi, D)p(\theta, D_{meta-train})}{p(D, D_{meta-train})} d\theta$$
(7)

$$= \log \int \frac{p(\phi, D, \theta) p(\theta, D_{meta-train})}{p(D, \theta) p(D_{meta-train})} d\theta$$
(8)

$$= \log \int p(\phi|D,\theta) \, p(\theta|D_{meta-train}) \, d\theta$$

$$\approx \log p(\phi|D, \theta^*) p(\theta^*|D_{meta-train})$$

Where $\theta^* = \arg \max_{\theta} p(\theta | D_{meta-train})$. To estimate the new task $\arg \max_{\phi} \log p(\phi | D, D_{meta-train})$, as $\arg \max_{\phi} \log p(\phi | D, \theta^*)$, where ϕ represents the task-specific parameters and θ^* denotes the initial information distributed across all tasks.

In summary, acquiring a new skill involves two stages. The first stage is gaining proficince with the detalearning parameter $\theta^* = \arg \max_{\theta} p(\theta | D_{meta-train})$, followed by refining it using a limited number or examples $\phi^* = \arg \max_{\phi} p(\phi | D, \theta^*)$.



Fig. 1. Meta-Lorning for new task prediction.

Figure 1 shows the working of verta-learning in new task prediction and it illustrates the involvement of meta-parameters in new tasks an above neta-parameters are updated based on task-specific parameters using a limited number of new input/output pairs. Ultimately, the updated ϕ^* is utilized for prediction. The algorithm 1 gives the working of petal participation model.

Algorithm 1: Neta-Learning	
Input: Mean training data $D_{meta-train}$, New task data D_{new} , Parameters $ heta, \phi$	
Output Adapted model parameters $oldsymbol{\phi}^*$, performance.	
Step 1: Initelize Meta-parameters $\boldsymbol{\theta}$.	
For each tast T_i in $D_{meta-train}$:	
a. Extract the training data D_i for task T_i .	
Train the model on task T_i using the current θ .	
c. Update meta-parameters $\boldsymbol{\theta}$.	
$\theta^* = \arg \max_{\theta} p(\theta D_{meta-train})$	
Step 2: Adapt at New Task	
a. Extract New Task Data	

b. Initialize Task-Specific Parameters ${oldsymbol{\phi}}$

c. Update task-specific parameters ϕ using new task data **D** and the prior θ^* .

 $\phi^* = \arg \max_{\phi} p(\phi | D, \theta^*)$

Step 3: Evaluate the Adapted Model

Step 4: Repeat Steps 2 and 3 to adapt the model to different tasks, using the prior θ^* .

B. LSTM-Attention

This section details the importance of combining LSTM and the Attention mechanism. First, the architecture and working of LSTM and the Attention mechanism are detailed. Next, the proposed model is explained

LSTM: When handling large volumes of sequencing data, LSTM is recommended as a training articled [2]. The gated state enables LSTM to exercise more control over the transmission state compared to the standard RNN. After a long state $c_{(t-1)}$, the forget gate $f_{(t)}$ retains or discards the input. The input gate $c_{(t)}$ determines whether to allow the input data to the current state of the memory cell $c_{(t)}$. The output rate $o_{(t)}$ functions similarly to the input gate by deciding to send the signal from the current to the next over. Equations (11-15) illustrate the calculation principles for each gate structure [24].

$$f_{(t)} = \sigma (W_f x_{(t)} + V_f h_{(t-1)} + b_f)$$
(11)

$$i_{(t)} = \sigma (W_i x_{(t)} + V_i h_{(t-1)} + b_i)$$
(12)

$$o_{(t)} = \sigma(W_o x_{(t)} + V_o h_{(t-1)} + b)$$
(13)

$$c_{(t)} = f_t \otimes c_{(t-1)} + b_t \otimes ta \left(u_{t} + V_c h_{(t-1)} + b_c \right)$$
(14)

$$h_{(t)} = o_t Stanher_{(t)}$$
 (15)

The input sequence $x = (x_1, x_2 \dots x_n)$ and the opposed unce $h = (h_1, h_2 \dots h_n)$ are defined as follows. The weight matrix W and vector V re-established, with h representing the threshold. The sigmoid activation function $\sigma(x) = \frac{1}{1 + e^{-x}}$ is defined, and the dot products indicated by \otimes .

Attention Mechanism: The attention n chanism (AM) addresses the problem of information overload when those resources to more critical tasks. A larger number of computational resources are limit cati parameters allows the model t tore m hformation and effectively represent features within neural information overload. Applying an AM helps the network prioritize networks, however, this co nd relevant input data, filter g out i essential information and focusing only on what is necessary for the task at hand. The attention va mined through two processes: (1) defining attention distributions for each is det input datasé distributions to compute the weighted average of the input data. Equations (16) to (putational concepts of the AM. the co

$$h_t = RNN(x_t, h_{t-1}) \tag{16}$$

$$c_i = \sum_{j=1}^n \alpha_{ij} h_j \tag{17}$$

$$\alpha_{ij} = \frac{exp(s(h_t, \overline{h_s}))}{\sum_{k=1}^{n} exp(s(h_t, \overline{h_s}))}$$
(18)

$$s(h_t, \overline{h_s}) = h_t, \overline{h_s} \tag{19}$$

The hidden state variable h_t is generated by merging the input x with the previous hidden state h_{t-1} . The c_i represents the weighted average of all hidden layers as well as the following hidden layer unit. The weight ratio of the hidden layer units is represented by α . Consequently, the target value is $\overline{h_s}$, where s represents the weight calculation method.

Proposed Model: Combining the AM into the LSTM network enables the model to achieve high prediction accuracy. Regardless of the input signal sequence length, conventional LSTM may transform it into a fixed length, which limits the model's ability to learn from extensive sequences. To enhance prediction accuracy, the AM fortifies connections between hidden layers and highlights significant information by assigning weight components. The network encoder utilizes the AM to retain intermediate outputs. The proposed model is trained to detect correlations between input and output sequences by selectively learning meaningful representations from the input sequence. Figure 5 illustrates how the AM computes the score for each variable by utilizing intermediate variables obtained from the hidden layers of the LSTM. The weight assigned to enable reflects its relative importance. Ultimately, the vector layer captures important features from the input, integrates it, and assigns additional weight.



Fig. 2. LSTM ane. ion a hitecture.

After integrating LSTM with the AM, Equations ()-() present the conditions for updating each parameter.

$$\begin{pmatrix} f_{t}^{n} \\ p \\ \sigma \\ \sigma \\ \tau^{n} \\ \sigma^{n} \\ \tau^{n} \\ \tau^$$

(20)

$$S_i^t = W_s tanh (W_h h_{(t-1)}^n + W_x x_i + b_s)$$
 (21)

$$\beta_i^t = \frac{exp(s_k^t)}{\sum_{k=1}^n exp(s_k^t)} \tag{22}$$

$$\sum_{i}^{N} \beta_{i}^{t} = 1 \tag{23}$$

where Z_{L+M}^n represents the n-th layer parameters, σ denotes the sigmoid function. The embedding matrix is indicated a Ed_{t-1}^n with L referring to the LSTM dimension. The context vector v_t represents the relevant input vector at the term t, with M denoting the dimension of v_t . The relevance score s_i^t determines the attention weights 2^t one networks for the attributes $X = \{x_1, x_2, ..., x_N\}$ can obtain attention weights after acquiring their r scores. The weight matrices applied in the proposed network at the time t' are detailed in Equations (24-26).

$$w_{t',ih}' = w_{t'}, ih\beta_{t'}$$
⁽²⁴⁾

$$w_{t',oh}' = w_{t'}, oh\beta_{t'}$$
⁽²⁵⁾

$$w_{t',fh}' = w_{t'}, fh\beta_{t'}$$
⁽²⁶⁾

Enhancing the weights allows the network to focus on capturing significant data. The LSTM-Attention mechanism improves prediction accuracy by identifying the most important temporal features from the limited ECG data. The algorithm 2 gives the working of LSTM-Attention model.

Algorithm 2: LSTM-Attention Model

Input: Sequence of input data X. LSTM hidden states h , Attention weights α	
Output: Accuracy.	•
Step 1: LSTM Update	
a. Forget gate: $f_t = \sigma(W_f, x_t + V_f, h_{t-1} + b_f)$	
b. Input gate: $i_t = \sigma(W_i, x_t + V_i, h_{t-1} + b_i)$	
c. Output gate: $\boldsymbol{o}_t = \boldsymbol{\sigma}(\boldsymbol{W}_o, \boldsymbol{x}_t + \boldsymbol{V}_o, \boldsymbol{h}_{t-1} + \boldsymbol{b}_o)$	
d. Memory cell update: $c_t = f_t \odot c_{t-1} + i_t \odot tanh(W_c, x_t + V_c, h_{t-1} + b_c)$	
e. Hidden state update: $h_t = o_t \odot tanh(c_t)$	
Step 2: Attention Mechanism	
a. Calculate hidden state at time t : $h_t = RNN(x_t, h_{t-1})$	
b. Compute the attention score for each input: $s_i^t = W_s \cdot tanh(W_h, h_t, y, W_x, x, y, b_s)$	
c. Calculate the attention weight $\beta_i^t = \frac{exp(s_i^t)}{\sum_{k=1}^n exp(s_k^t)}$	
d. Ensure that the attention weights sum to $1 \sum_{i}^{N} \beta_{i}^{t} = 1$	
Step 3: Update the LSTM parameters using Attention a. $w'_{t',ih} = w_{t'}, ih\beta_{t'}$	
D. $W_{t'ab} = W_{t'} O D D_{t'}$	

c. $w'_{t',fh} = w_{t'}, fh\beta_{t'}$

Step 4: Evaluate the LSTM-Attention Model using accuracy Step 5: Repeat Steps 2 and 3 if evaluation outcommunitatisfied

C. LSTM-Attention-based Meta-Learning

The LSTM-Attention model capture and most important temporal features from ECG, which helps identify abnormalities in the ECG. The integration of 250 A-Attention in the meta-learning concept can handle a single model for a variety of task predictions using ECG data. This helps the model to learn from past data and adapt quickly and accurately to new tasks

a meta-learning is detailed as follows: The ECG input is given to the The process of LSTM₁ ttentic bas LSTM model, which capt nporal features that help understand the health condition of an individual. es the te The AM refi dentifies the most important ones from the LSTM output by adjusting the importance to all features, but not all features contribute equally to the prediction weight m TM giv etrics n is trained on each task, such as HRV prediction, arrhythmia prediction, and task. LST Atten. task, the model uses specific features from the ECG related to that task. Meta-learning abnorm ities. g training to optimize the meta-parameters for quick adaptation to various tasks. If new ECG data is used du ase comes, the proposed framework can identify the new or unseen disease using the metawif new d earned during training. Therefore, the framework predicts the new task using previous training arame knowledge with minimal data.

• RESULT AND DISCUSSION

The section details the experimental setup, data used in the research, and the evaluation of the proposed model in comparison with existing models.

A. Experimental Set-up

The experiment was conducted on the Google Colaboratory platform. The data are stored in Google Drive for easy access. The Graphics Processing Unit (GPU) available in Colab was used to run the code. The code is

written in Python. The scikit-learn library is used for data processing and model evaluation. The model is imported from TensorFlow. The proposed method block diagram is given in Figure 3. The MIMIC-III data is collected and the required preprocessing is done. Then, the three different tasks are taken, and the LSTM-attention model is designed and tuned for each task individually using the meta-learning concept. If new data comes in, the proposed meta-learning-based LSTM-attention model can predict the disease accurately. The outcome of the model is evaluated using standard metrics.



Fig. 3. Block diagram of the posed method for disease prediction from EHR data.

B. Data Collection and Processin

The MIMIC III dataset ichticludes medical information for intensive care units, was used in this hat contains the medical records of more than 60,000 individuals who were pilot study. MIMIC-III is database hospitalized in the nits of Beth Israel Deaconess Medical Center between 2001 and 2012. This database hold ensive ta. The study dataset primarily focuses on ECG data stored in EDF (European Data Format) f The **DF** format is widely used for storing time-series data from various biological signals. The ts of a EDF file include the metadata within the header section and the signal data in the main main compon. ile. h context of the dataset, the EDF files likely contain multiple channels of ECG data collected body of th ECG signal, representing the heart's electrical activity, is crucial for diagnosing and monitoring time. Th cardia tions. QRS points refer to annotations on the QRS complex in an ECG signal. Accurate recognition S complex is essential for tasks such as HRV analysis, arrhythmia detection, and other abnormalities. The first pre-processing step is to identify the missing data and fill in the missing values using the imputation method. Next, the data using the Synthetic Minority Over-sampling Technique (SMOTE) method [19] to ensure the data is balanced, with each label in the dataset having equal samples. For feature extraction, the HRV distribution is calculated using standard deviation and rolling windows. Figure 4 shows the outcome of the HRV analysis and rolling window.



Finally, segmentation and normalization are performed on the data. In segmentation, by continuous signal is divided into meaningful intervals. For normalization, the standardization termique happle is to convert the values of the data into a range from 0 to 1. This helps to reduce the memory isage and complexity of model training. Figure 5 provides the signal plot before and after normalization.



Fig. 5. Signal plot before a fiter tormalization. *C. Result Comparison*

After processing the provided to the meta-learning framework, which includes the LSTMate the effectiveness of the proposed LSTM-Attention model, it is compared with Attention mo o e as SVN Random Forest (RF), and XGBoost. All the models are implemented within the other mo ork. The three tasks-HRV prediction, arrhythmia prediction, and abnormality meta-learn fram rmed using the same dataset. The processed data is split into training, validation, and test prediction are ratio. The LSTM-Attention model is trained on each task individually, and metrics such as in a 7 set loss are used during the training and validation processes. The results of the LSTM-Attention accui and lel for each task are shown in Figures 6-8.



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Fig. 8. Performance plot of propsoed model for Abnormality prediction.

After training and validation, the model is tested to evaluate it nce. The evaluation metrics such berf iddit as accuracy, precision, recall, F1 score, and ROC-AUC are com n to processing time. The highest metrics of accuracy, precision, recall, F1 score, and RO by the proposed model, with values re atta AU of 0.92, 0.90, 0.91, 0.91, and 0.93, respectively. Nex es for accuracy (0.87), precision (0.83), recall .he hi est s (0.85), F1 score (0.84), and ROC-AUC (0.88) are RF. XGboost shows better results, and finally, SVM tained/ performs the worst. The metric values for all model shown in Table 1. The table also provides the formulas to calculate the metrics, where TP, TN, FP, FN represe the True Positive, True Negative, False Positive, and False Negative for the model's prediction

	Tuble IV Terrormance companion of provided 201101 Internation intolder what conventional models					
	Methods	Fongula	SVM [20]	Random	XGBoost	Proposed
				Forest [21]	[22]	LSTM-
						Attention
		TP/TN	0.85	0.87	0.86	0.92
	Accuracy	TP + TN + FP + FN				
		TP	0.81	0.83	0.82	0.90
	Precision	TP + FP				
	Reca.	TP	0.84	0.85	0.83	0.91
		TP + FN				
	F. Scor	2 * Precision * Recall	0.82	0.84	0.83	0.91
		Precision + Recall				
V	ROC_AUC	$\begin{pmatrix} 1 & TP \end{pmatrix} (FP)$	0.86	0.88	0.87	0.93
		$\int_{0} \left(\frac{TP + FN}{TP + FN} \right)^{d} \left(\frac{FP + TN}{FP + TN} \right)$				
	Processing	-	120	115	110	95
	Time					

 Table 1. Performance comparison of provided LSTM-Attention model with conventional models

Finally, processing time is important when deploying the model in real-time applications. The proposed model takes just 95 seconds to predict the disease from ECG data, while the other models—SVM, Random Forest, and XGBoost—take 120 seconds, 115 seconds, and 110 seconds, respectively. The quality metrics show the effectiveness of the proposed model in disease prediction, and the processing time indicates that the model can be deployed in real-time applications. Figures 9–14 show the comparison plots for the quality metrics and processing time attained by the models. The visual graphs help in a better understanding of the model evaluation.



Fig. 9. Accuracy comparison of proposed LSTN actention-based Meta-learning model with conventional *mol. Vs.*



Fig. 10. Precision comparison of proposed LSTM-Attention-based Meta-learning model with conventional models.



Fig. 11. Recall comparison of proposed LSTM-Attention-based Materian model with conventional







V. CONCLUTION

The research aims to propose an efficient model for predicting all types of diseases from ECG data. While many reactive models are available, they are often generalized and fail to account for unique patient data, are carly when dealing with rare diseases. To address this issue, the research proposes a novel LSTM-Attention-based meta-learning model for predicting diseases from ECG signals in EHR data. The ECG data in EHR is often limited, but by using the proposed model, it is capable of predicting three different tasks: HRV, arrhythmia, and abnormality prediction. The integration of AI models and meta-learning can solve many problems in the healthcare field by providing accurate predictions with limited data, unseen data, etc. The proposed model is compared with SVM, RF, and XGBoost, and it achieves the highest accuracy of 0.92, outperforming RF with an accuracy of 0.87. The proposed model improves prediction accuracy by 5%. Additionally, compared to other models, the proposed model requires minimal time (95 seconds) to predict diseases from ECG data. The quality and processing time of the model ensure that it can be implemented in clinical applications. The proposed framework will help healthcare professionals predict and treat diseases as early as possible.

In the future, to enhance the accuracy of the proposed model, multi-modal data will be used. Other data such as medical imaging, will be incorporated with ECG data to provide a comprehensive health profile. T deploy the proposed model in real-time, it will be further refined for integration into a wearable device for continuous health monitoring and immediate disease prediction.

Declaration

Ethics approval and consent to participate Not applicable.

Clinical trial number Not applicable.

Consent for publication Not applicable.

Competing interests The authors declare no competing interests.

Data Availability The dataset generated and/or analysed during the current <u>https://mimic.mit.edu/</u>.

Authors Contribution

C.G., S.M.B., S.T.A., S.K.M. participated in designing the methodology, concept and implemented the code and performed experiments, and wrote the manuscript. S.M. adated, supervised, reviewed and edited the manuscript.

on MIMIC III dataset [27].

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Appendix 1. Nomenclature

<u> </u>	Training data set Model parameters
$p(\phi)$	Prior probability of the model parameters
D _{meta-train}	Meta-training data set
	Meta-parameters learned from the meta-
θ	training data
<u> </u>	Point estimate of the meta-parameters
f_t	Forget gate in LSTM
l _t	Input gate in LSTM
h_t	Hiddon store in LSTM
$\sigma(x)$	Sigmoid activation
h_t	Hidden state vector generate by RNN at time
$lpha_{i,j}$	Attention weight the <i>i</i> -th element and <i>j</i> -th
α_{ii}	for azed, tention weight
$\frac{l_j}{s(h_t, h_s)}$	Sim. vity function between hidden states
v_t	Context vector at time <i>t</i> in AM
β_i^t	A cention weight at time t for the i -th input
W(t'),ih	Weight matrix for input-hidden layer connections at time t'
W(t'),oh	Weight matrix for output-hidden layer connections at time <i>t</i> '
Z ⁿ _(L+M)	<i>n</i> -th layer parameters of the LSTM-Attention network
W _s	Weight matrix for the similarity function in AM
W	Weight matrix for hidden state in AM
V r	Weight matrix for input in AM
<i>b</i> .	Bias term for the similarity function in AM