

HOSPITAL READMISSION

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Hospital Readmission Prediction of ICU Patients Using Deep Learning Algorithms

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Abstract—Deep learning healthcare applications have evolved over the last few years. This advancement leads to new applications and possibilities in the field of health care. Due to different variants of deep learning algorithms like convolutional and recurrent neural network these advancements in healthcare application made possible.

Index Terms—Deep learning, LSTM, Healthcare, Recurrent Neural Network, GRU.

I. INTRODUCTION

A. Hospital Readmission

Hospital Readmissions are distinguished as a sign of the low quality of consideration, for example, inadequate release arranging and care coordination. Emergency clinic readmission[1] happens when a patient release from a medical clinic is readmitted inside a predetermined interim. Admittance of a patient can happen due to any decided or undecided reasons. These factors drove Health care associations to consider managing readmission rates; an expansion in readmission rates prompts an increment in punishments. There has been an expanded use of Readmission [2] as a result rule in wellbeing administrations investigation and as a quality criterion for well being frameworks. For Medicare patients, hospitalizations are exceptionally distressing, and it is considerably more when they occur in frequent readmissions. A ton of research concentrates demonstrated that emergency clinics could take part in a few exercises like explaining patient release directions, planning with post-intense consideration suppliers, lively cleaning system to lessen the rate of readmissions of patients. Be that as it may, these individual subsequent meet-ups can be costly.

B. Machine Learning For Health Care

Machine learning leads to provide [34]elligence by providing new knowledge. Since the onset of machine learning, it has been widely used in achieving paramount goals as the healthcare sector is evolving. The healthcare sector is producing an enormous mass of data, and it is beyond human capability to manually analyze this vast data and produce inferences based on that. [24]chine learning has proven to be a benefit at this level. The machine learning algorithms automatically find patterns in the data, which enable to get inferences from the

data easily. Machine learning can assist healthcare providers in a variety of tasks like hospital readmission of patients suffering from chronic illness. This can not only lower the lifetime risk of the patient but also can save millions of dollars if the algorithm can predict efficiently and correctly. Other applications of machine learning in the healthcare domain includes drugs combination which should be avoided taking together, classifying images for different diseases like skin cancer. Keeping this in the picture, it can be said that machine learning can revolutionize the healthcare sector.

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C. Deep Learning

Deep learning is one of the extended section of Machine learning(ML), which works on a specific ML algorithm, that is the artificial neural network. It's leveraging the recent advancements in computing power and thus using specific purpose neural network to learn from a plethora of data and make predictions based on the patterns detected. The structure consists of several hidden layers which are meant to extract the information by exploiting the structures present in the data. Deep neural networks are specialized in solving problems, including data, which is greatly structured. It also pledge to substitute hand-engineered attributes with dimensionality reduction methods that are hierarchical, unsupervised, or semi-supervised. Multiple hidden [46]ts are being used between input and output in case of Deep Neural Network (DNN). Similar to the artificial neural network, the complex non-linear correlation can be modeled using DNN[2].

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D. Recurrent Neural Network

An artificial neural network has a specific kind called recurrent neural network(RNN), and it has directed cycles between connection. The internal state of the network exhibits dynamic temporal behavior. Unlike ANN, random series can be managed by using the internal state of RNN. This is the fundamental structure introduced in the 1980s: neuron-like units in a network, a directed connection between every neuron. There is a time-dependent continuous valued activation in every unit. There are input, output, and hidden layers. One input vec[30] is supplied at every time steps. At every time step, the non-linear function of the weighted sum of all the

activations of connected units is calculated by each non-input unit[3].

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E. Long Short term Memory

(LSTM) network, a deep learning RNN, is used by numerous researches published by Schmidhuber & Hochreiter in 1997[4]. This deep learning system solved the vanishing gradient issue of standard RNN. The recurrent gates called forget gate are introduced in LSTM. There are an unlimited number of virtual layers in LSTM-RNNs, through which the error can flow backward. LSTM can learn from events that occurred thousands of time steps ago so it can be used for very deep learning tasks. LSTM has a unique architecture that consists of a memory cell, which can keep the state which it is currently in, over duration. Non-Linear gating units are used in LSTM to control the course of data in and out of the memory cell. Over time, many improvements have been suggested and made in the standard architecture of LSTM to make it more efficient. Today, LSTMs are being used in a variety of learning problems which differ in scale and nature significantly when compared to the problems which were initially solved by LSTM.

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II. LITERATURE REVIEW

During the last decade several research work is carried out in the field of healthcare. Hospital readmission prediction is important and crucial application of healthcare for the improved quality of life of an individual.

After stunning successes in cognitive domains, deep learning is expected to transform healthcare [21]. Most remarkable results thus far in health have been in diagnostic imaging [7], [9], which is a natural step given recordbreaking results in computer vision. However, diagnostic imaging is only a small part of the story. A full intelligent medical system should be able to reason about the past (historical illness), present (diagnosis) and future (prognosis). Here we adopt the notion of reasoning as algebraically manipulating previously acquired knowledge in order to answer a new question [1]. For that we learn to embed discrete medical objects into continuous vectors, which lend themselves to a wide host of powerful algebraic and statistical operators.

Author [5] developed an approach that helps clinicians determine the optimal initial dose of a drug to safely and quickly reach a therapeutic aPTT window. In [6] author uses a super learner algorithm is used for predicting hospital mortality in patients. In [7] author used a novel called Interpretable Mimic Learning is used, to learn interpretable phenotype features for making robust prediction while mimicking the performance of deep learning models metaheuristics. In [8] Combination of convolution and recurrent neural network model is used for object detection. In [9] authors objective was to reduce overfitting using batch normalization. In [10] author introduced interpretable mimic learning that uses gradient boosting trees to learn interpretable models and at the same time achieves strong prediction performance as deep learning models. In [11] author found the proportional split of data is

useful for deep learning applications when dealing with large volumes of data. In [12] author leveraged longitudinal EHR data for early detection of heart failure using Recurrent Neural Network. Researchers have recently begun attempting to apply neural network based methods (or deep learning) to EHR to utilize its ability to learn complex patterns from data. Previous studies such as phenotype learning or representation learning, however, have not fully addressed the sequential nature of EHR is especially related to our work in that both studies use RNN for sequence prediction. However, while uses regular times series of real-valued variables collected from ICU patients to predict diagnosis codes, we use discrete medical codes (e.g. diagnosis, medication, procedure) extracted from longitudinal patient visit records. Also, in each visit we make a prediction about predict diagnosis, medication order in the next visit and the time to next visit. In [13] author uses Long short term memory (LSTM) RNN to predict disordered proteins, such as SPOT-disorder. In [14] State-of-the-art deep and recurrent neural network models were applied to achieve best results on classifying opioid users. In [15] author used Gated Recurrent Unit (GRU) to solve the vanishing gradient problem of a standard RNN. In [16] author used LSTM to overcome the vanishing gradient problem of traditional RNN. In [17] author Combines the use of time frequency heat map representations with a deep convolutional neural network (CNN) to classify heart sounds. In [18] author used Statistical Analysis is used to find the stability and readmission rates. In [19] author implemented a 13-layer deep learning CNN algorithm for the automated EEG analysis.

III. METHODOLOGY

A. Data description: MIMIC-III

Medical Information Mart for Intensive Care (MIMIC-III)[20] consists of data about patients admitted to various critical care units in a large hospital. MIMIC-III is generally viewed as a large and single-center database. A large number of different parameters are present in MIMIC III database. These parameters include information such as vital signs, medications, laboratory measurements, observations, fluid balance, procedure codes, diagnostic codes, imaging reports, hospital length of stay, survival data, and others. The database consists of information of around 58,576 distinct patients who were admitted to various critical care units of the hospital between 2001 and 2012. The data comprises of patients aged 16 years or above only. Descriptive statistics were performed on this dataset, and it was found that the median age is 65.8 years for adult patients(Q1-Q3: 52:877:8). Out of total patients, there were 55.9% male patients and only 11.5% of the cases had in-hospital mortality. It was calculated from the data that on an average a patient stayed for 2.1 days in ICU and had an average of 6.9 Days of hospital stay. MIMIC-III database have several idiosyncratic properties about it, and these are mentioned below:

- The dataset is a huge database accumulated throughout more than a decade and consists of very detailed information of each patient under care.

- MIMIC-III database requires a user to oblige by a user data agreement and once it is accepted the analysis is boundless.
- It contains discharge summaries as well as reports of ECG, imaging studies and information about various codes like International Classification of Disease, 9th Edition (ICD-9) codes, Diagnosis Related Group (DRG) codes, and Current Procedural Terminology (CPT) codes.
- It is a time series data. Clinical variables are recorded concerning time for each patient. Figure 1 shows the chart of timing clinical variables recorded in the dataset.

The details of dataset used in this research is given in table I below.

Data	Statistics
Total Patients	58,576
Patients Selected For Study	6,587
Number of classes	2
Number of patients readmitted in 30 days	2,461
Number of patients readmitted after 30 days	4,126
Number of female patients	2,916
Number of male patients	3,671
Mean age of all patients	90.68

TABLE I
DATASET SUMMARY

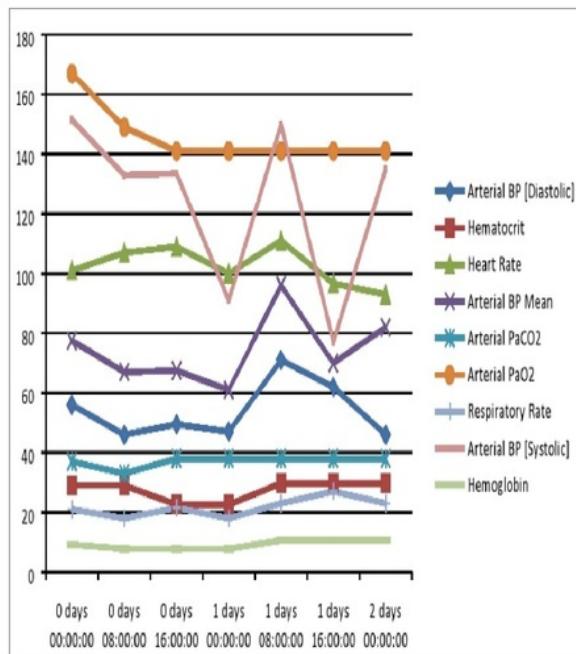


Fig. 1. Some of the Clinical recordings associated with each patient[20]

B. Data Preprocessing

1) *Data Extraction:* MIMIC-III dataset consisted of around 58,576 patients who were diagnosed as suffering from different types of disease and hence mortality due to those diseases. These diseases include Pulmonary disease, Circulatory disease, Trauma, a disease of the digestive system and many more. Since the dataset is very large, we only consider data of those patients who were readmitted again which gives the details of 6,587 patients. The data set is divided into two classes:-

- Patients readmitted in 30 days.
- Patients readmitted after 30 days.

2) *Normalization:* Normalization is used to remove the bias among the attributes. It brings the attributes on a standard scale. It standardizes the range of independent features or variables of data, called feature scaling. We use min-max normalization[22] here.

- Min-Max Normalization Let a matrix "A":-

$$A = \begin{bmatrix} a_{11} & a_{12} \\ a_{21} & a_{22} \end{bmatrix} \quad (1)$$

So for L2 normalization we calculate

$$a = \min(a_{11}, a_{21}), \quad (2)$$

$$b = \max(a_{11}, a_{21}) \quad (3)$$

$$c = \min(a_{12}, a_{22}), \quad (4)$$

$$d = \max(a_{12}, a_{22}) \quad (5)$$

And "A" becomes:

$$A = \begin{bmatrix} (a_{11} - a)/(b - a) & (a_{12} - c)/(d - c) \\ (a_{21} - a)/(b - a) & (a_{22} - c)/(d - c) \end{bmatrix} \quad (6)$$

C. Feature Selection

Features are selected through Nature Inspired Algorithms:

1) *Bat Algorithm:* Bat Algorithm is an optimization algorithm that mimics the echolocation behavior of microbats[22]. Bats uses echolocation to distinguish between food/prey and they also sense distance and other background barriers. Bats fly at random with a speed u_i at point x_i with a constant frequency μ_{min} , changing wavelength and loudness L_0 to look for the prey. They can spontaneously change the frequency of their released pulses and regulate the rate of pulse ejection r in the range of $[0, 1]$, based on how close is their prey. Even though the loudness can change in different ways, we suppose that the loudness changes from a high (positive) L_0 to a least fixed value L_{min} . Rules to update position x_i and speed u_i of bats at time step t is given by:

$$\mu_i = \mu_{min} + (\mu_{max} - \mu_{min})\beta \quad (7)$$

$$u_i^t = u_i^{t-1} + (x_i^{t-1} - x_b)\mu_i \quad (8)$$

$$x_i^t = x_i^{t-1} + u_i^t \quad (9)$$

where $\beta \in [0, 1]$ is a arbitrary vector extract from a rectangular distribution and after comparing all the solution of n bats we

found x_b the global best solution. For the local search part a local random walk is used to find a new solution for each bat once a solution is selected from the current best solution:

$$x_{new} = x_{old} + \epsilon L^t \quad (10)$$

where $\epsilon \in [-1,1]$ is an arbitrary number. Average loudness of all the n bats at that time is L^t .

Variation of Loudness and Pulse Emission:

$$L_i^{t+1} = \alpha L_i^t \quad (11)$$

$$r_i^{t+1} = r_i^0 [1 - \exp(-\gamma t)] \quad (12)$$

⁴³ α & γ are constant.

2) Grey Wolf Optimizer: Gre ⁴ Wolf Optimizer(GWO) Algorithm imitate the nature of grey wolves[23]. The GWO algorithm imitates the leadership pecking order and coursing procedure of grey wolves in nature. Four types of grey wolves ²² selected for replicate the leadership hierarchy such as alpha, beta, delta, and omega. Also, the three main steps of coursing, hunting for prey, encircling prey, and attacking victim, are implemented.

Alpha wolf is a leader male or female. They make decisions like the sleeping place, hunting, etc. Other wolves acknowledge alpha wolf by their tail down. Beta wolf help alpha wolf in making decisions. They are an advisor for alpha and discipliner for the pack. They also ensure all subordinate obey the order of alpha and give feedback to alpha. Delta wolves also called subordinate. They dominate omega wolves.

Categories of Delta wolves:-

- Scouts - Watch boundaries.
- Sentinels - Protect pack.
- Elders - Alpha or beta wolf sometimes.
- Hunters - Help alpha and beta wolf in hunting.

⁴² Care Taker - Care ill weak and wounded wolves.

Omega wolves are like the scapegoat in the pack. They are having weak fitness and are allowed last to eat.

Search Process in GWO Algorithm:-

- Searching
- Encircling
- Attacking the prey

a) *Searching*:

- Search according to aplha, beta and delta wolves. They separate to look for prey and come together to ambush on prey.
- Modeled by using w random variable greater than 1 or less than -1.
- When $|w| > 1$ wolves are required to separate from prey to find better solution.

b) *Encircling Prey*: It is the process in which wolves encircle the prey for attack. It is modeled as:

$$\vec{d} = |\vec{c} \cdot \vec{x}_p(t) - \vec{x}(t)| \quad (13)$$

$$\vec{x}(t+1) = \vec{x}_p(t) - \vec{w} \vec{d} \quad (14)$$

Where t is the current iteration, \vec{w} , \vec{c} are coefficient vectors and \vec{x}_p is position of prey and \vec{x} is the position of grey wolf.

Updation of Coefficients

$$\vec{w} = 2\vec{a}\vec{r}1 - \vec{a} \quad (15)$$

$$\vec{c} = 2\vec{r}2 \quad (16)$$

Where \vec{a} linearly decrease from two to zero over the course of iteration. $\vec{r}1$ and $\vec{r}2$ are arbitrary vectors ⁴³ [23,1].

c) *Attacking*: The attack is mostly led by alpha. The beta ³⁴ delta may also join in hunting. We first save three best solutions i.e. alpha, beta and delta which are the best answers and update the locations of other wolves based on these three. It is modeled as:

$$\vec{d}_\alpha = |\vec{c}_1 \vec{x}_\alpha - \vec{x}| \quad (17)$$

$$\vec{d}_\beta = |\vec{c}_2 \vec{x}_\beta - \vec{x}| \quad (18)$$

$$\vec{d}_\delta = |\vec{c}_3 \vec{x}_\delta - \vec{x}| \quad (19)$$

$$\vec{x}_1 = \vec{x}_\alpha - \vec{w}_1 \vec{d}_\alpha \quad (20)$$

$$\vec{x}_2 = \vec{x}_\beta - \vec{w}_2 \vec{d}_\beta \quad (21)$$

$$\vec{x}_3 = \vec{x}_\delta - \vec{w}_3 \vec{d}_\delta \quad (22)$$

$$\vec{x}(t+1) = (\vec{x}_1 + \vec{x}_2 + \vec{x}_3)/3 \quad (23)$$

Grey wolf ¹⁶ack it's target when it is immobile. In GWO vector w is a arbitrary value within an range [-2a,2a] a decrease from two to zero throughout the iteration. When $|w| < 1$ wolves ambush on victim.

D. Feature Extraction

For feature extraction Convolution Neural Network(CNN) is ¹²d.

1) *Convolutional Neural Network*: A convolution neural network(CNN)[24] is an efficient machine learning approach from deep learning and is close to standard multi layered networks. It consists ¹³ of three layers:

- Convolution Layer
- Pooling Layer
- Fully Connected Layer

Convolutional layers: This layer is the most important layer among all in CNN. It contains rectangular grids or cubic blocks called filters. The filter is moved at every block of input neuron to produce output. These filters are trainable. Output is determined by equation:

$$WI^x + b \quad (24)$$

There are three hyperparameters in convolutionlayer that decides the output:

- Depth
- Stride
- Zero Padding

The number of filters used in convolution layer with certain stride is the depth of the output neuron. For Example say, the RGB image, with a depth of three. Given an image of size (5x5x3) and given ten filters of size (3x3x3). Here, when the

value of stride is given 1, the filters hovers a single pixel over an image then the output neuron is of size (3x3x10). Whilst when the value of stride is given 2, the filters will skip two pixels, and output neuron is of size (2x2x1) [37].

Zero padding is the process of filling zeros around the edge of the input neuron. Zero padding is commonly used to adjust the size of input neuron during the filter process when we need to main the output neuron based on input neuron size.

$$\text{Output} = (W - F + 2P)/S + 1 \quad (25)$$

where the input neurons size is W , filter's (kernel) size is F , paddings size is P , and stride is S . Linear algebra are also used in the convolutional neural network [36]. Now consider that given a matrix of dimension $A \times B$ (A is the number of rows and B is the number of columns). Now two dimensional convolution operation with two matrices (A_M, B_M) is a dimension of matrix M , and (A_N, B_N) is a dimension of matrix N . Convolution of these matrix is given by below equation:

$$P(i, j) = \sum_{a=0}^{A_M-1} \sum_{b=0}^{B_M-1} M(a, b) * N(i - a, j - b) \quad (26)$$

Max-pooling layer: Pooling layer executes the next operation after each convolution layer. These layers are used to reduce the size of the neurons. They are small rectangular grids that acquires small portion of convolutional layer and filters it to give a result from that block. The most commonly used layer is max pooling that fetch that maximum pixel from the block. A expression for a max-pooling layer is expressed in a equation:

$$h_i^j(a, b) = \max \{N(a), b \in N(b) \} h_j^{l-1}(\bar{x}, \bar{b}) \quad (27)$$

Fully Connected layers: The final layer of a convolutional neural network(CNN) is a fully connected layer that is formed from the attachment of all preceding neurons. It reduces the spatial information as it is fully connected like in artificial neural network. It contains neurons beginning at input neurons to the output neurons.

Details of each layer and their details are presented in Table II.

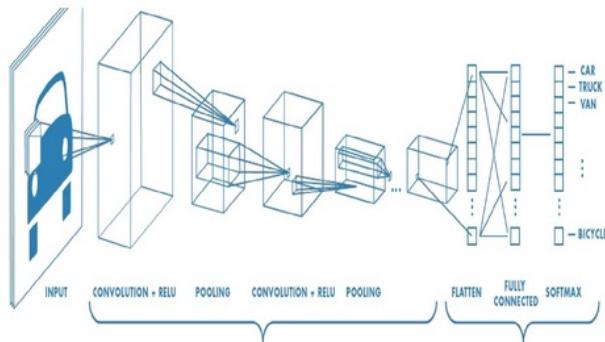


Fig. 2. Convolution Neural Network Architecture

Layers	Kernel Size	Filters Used	Input Shape	Output Shape
Conv1D	5	32	(98,30)	(98,32)
Conv1D	5	32	(98,32)	(98,32)
Conv1D	5	64	(98,32)	(98,64)
MaxPooling1D	2	1	(98,64)	(98,32)

TABLE II
CONVOLUTION NEURAL NETWORK SUMMARY

Algorithms	Number of Features Selected
Bat Algorithm	21
Grey Wolf Optimizer	17
Convolution Neural Network	32

TABLE III
FEATURES SELECTED BY DIFFERENT ALGORITHMS USED

E. Algorithms

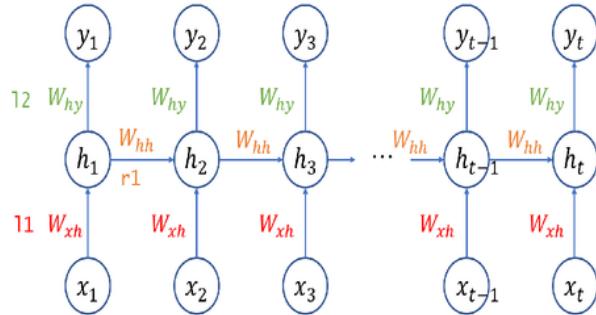
Recurrent neural networks (RNN) are utilized to represent the time dependency in the time series data[25]. As most of the healthcare data is a series of temporal recordings, hence RNNs have been widely adopted in the healthcare domain. We are usually provided with a series of observations $x_1 \dots x_T$ and we train a classifier to generate hypotheses \hat{y} . The recurrent connections are added in feed-forward neural networks to make it RNN. The output of a neuron in a typical NN is as follows:

$$y_i^t = \sigma(W_i x^t + b_i) \quad (28)$$

Where the weight matrix is represented as W_i , the bias represented as b_i and σ represents the sigmoid function. While in the case of RNN, a neuron is fed with the output of the neuron at time $t - 1$. The following equation shows the new activation function:

$$y_i^t = (W_i x^t + V_i x^{t-1} + b_i) \quad (29)$$

As RNN uses the previous outputs as recurrent connection, their current output depends upon the previous states. This property of RNN makes it very useful in sequence labeling tasks. The backpropagation through time [56] can be used to train RNNs. It was demonstrated by that learning long-term dependencies is difficult using gradient descent. This is mainly because the backpropagating error can vanish which makes the network inefficient in learning long-range dependencies, or frequently explode which makes convergence impossible. LSTM models were designed to tackle the issue of vanishing gradients and were developed to model long-range dependencies efficiently. LSTMs can fix this by maintaining an internal state that represents the memory cell of the LSTM neuron. This internal state can only be written and read via gates which decides what information will pass through gates. The following diagram shows the recurrent neural network.



$$h_t = \tanh(l1(x_t) + r1(h_{t-1}))$$

$$y_t = l2(h_t)$$

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Fig. 3. An recurrent neural network

1) **Long Short Term Memory (LSTM) Networks:** Long Short Term Memory networks are commonly called LSTMs. They [32] a particular kind of RNN, that are good in understanding long-term dependencies. They were introduced by [26] and were polished and familiarized by many people. They work very effectively on a large diversity of problems and are now used by many researchers. To solve long-term dependency problem, LSTMs are precisely formulated. Their default behaviour is to remembering something for a long duration of time. All recurrent neural networks contains a series of replicating chunks of the neural network. In traditional RNNs, this replicating chunk have a straightforward architecture, such as a single tank layer. LSTMs also have this series-like architecture, but the recurrent chunk has a disimilar architecture. Rather than having a single layer, there are four layers which are connecting extraordinarily. The intermediate information is stored in a single hidden layer 1 and its state changes over time (t_{t-1} , t_t , t_{t+1}). On the final hidden state vector l_T , we used a fully connected layer followed by sigmoid function. For loss function, we used log loss. The following equations can be used for calculation of the current hidden layer l_t .

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$$f_t = \sigma(W_f X_t + R_f l_{t-1} + b_f) \quad (30)$$

$$u_t = \sigma(W_u X_t + R_u l_{t-1} + b_u) \quad (31)$$

$$o_t = \sigma(W_o X_t + R_o l_{t-1} + b_o) \quad (32)$$

$$\tilde{C}_t = \Phi(W_C X_t + R_C l_{t-1} + b_c) \quad (33)$$

$$C_t = f_t C_{t-1} + i_t \tilde{C}_t \quad (34)$$

$$l_t = o_t \phi(C_t) \quad (35)$$

u_t , o_t and f_t are input gate, output gate and forget gate respectively. Forget gate decides which information to discard, input gates are used to update cell and output gates are used to

decide the output of the cell state [20]. Cell states are completely overridden in classical RNN, but LSTM has the potential to eliminate or append data to the cell state. The input weight of each gate, recurrent weight, and the bias are expressed as W^* , R^* , b^* respectively where * can be f, u, o and c. Here σ , Φ stands for an element wise operation of the sigmoid (logistic) and tanh function respectively. For matrix multiplication. The candidate values are computed in equation(3.33), and equation(3.34) old state is multiplied by f_t , and this helps in neglecting the things we decided to neglect. Then we add $24 * \delta C_t$ in it. This is the new candidate values, increased by how much we planned to update each state value. The final output of an LSTM unit is given by equation(3.35). Here * represents the Hadamard (element-wise) multiplication process. Figure 3 shows the LSTM network.

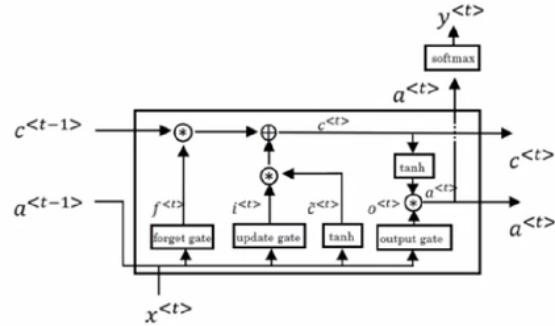


Fig. 4. A LSTM Network

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2) **Bidirectional Recurrent Neural Networks (BRNN):** Bidirectional Recurrent Neural Networks also known as BRNN is just like RNN but it trains simultaneously on both directions of the time dependent data[27]. This algorithm gives the better result in both regression and classification problem. BRNN computes both forward ($\vec{1}$) and backward ($\overleftarrow{1}$) hidden sequence.

$$\vec{1}_t = \mathcal{O}(W_{x\vec{1}} x_t + W_{\vec{1}\vec{1}} \vec{1}_{t-1} + b_{\vec{1}}) \quad (36)$$

$$\overleftarrow{1}_t = \mathcal{O}(W_{x\overleftarrow{1}} x_t + W_{\overleftarrow{1}\overleftarrow{1}} \overleftarrow{1}_{t+1} + b_{\overleftarrow{1}}) \quad (37)$$

$$y_t = W_{\vec{1}y} \vec{1}_t + W_{\overleftarrow{1}y} \overleftarrow{1}_t + b_o \quad (38)$$

The long range context can be accessed in both directions by combining Bidirectional RNNs with LSTM which gives bidirectional LSTM[28].

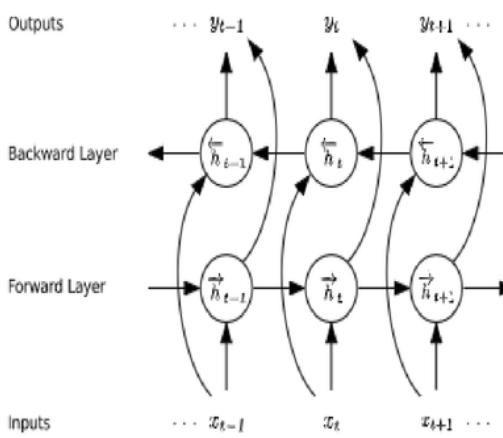


Fig. 5. Bidirectional Recurrent Neural Networks

F. Flow Diagram

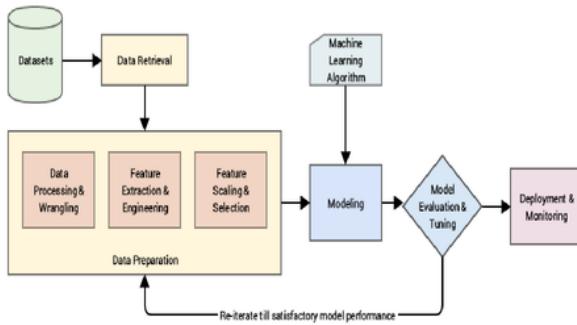


Fig. 6. Flow Diagram

According to the research objectives, the report will describe the work flow as below:

- Step 1:** Collect a vast readmission dataset with several features.
- Step 2:** Do data-preprocessing to deal with missing values.
- Step 3:** Convolution layers can be used for feature extraction, i.e., to retrieve the meaningful data out of the dataset.
- Step 4:** Do Feature scaling to normalize the dataset to reduce biases.
- Step 5:** Feature selection can also be done using nature-inspired algorithms to fetch important features from the dataset.
- Step 6:** Feeding the retrieved dataset into the mode [49]
- Step 7:** Calculate the Accuracy, FScore, AUC, etc. based on the actual and predicted output.
- Step 8:** If the results are satisfied, then stop else goto step 2 and redo the procedure with different hyperparameters, algorithms, etc.

IV. RESULTS

A. Implementation Details

The entire dataset was split using stratified sampling and 10% of the dataset was used as a validation set and 48% random search was used for hyperparameter optimization. The rest 8% of the dataset was used for training purpose and was tested on 10% of the dataset. The LSTM model was trained with 30 epochs using Adam optimizer. LSTM layer uses 100 memory cell with no dropout and 25 memory cell with 20% dropout and these architectures are found after validation performance. Also train this model with Bat Algorithm. Another model train on this dataset was convolution LSTM (CLSTM). It is a hybrid model in which convolution layers are used for feature extraction. The CLSTM model was trained with 30 epochs and the batch size of 1000. First, the convolution layer was used with 45 filters of size 5×5 . Then the LSTM layer that uses 100 memory cell with 20% dropout. Also train this model with Grey Wolf Optimizer Algorithm. In the training phase, the model was saved and also compute these performance metrics (AUC, F1, precision, and recall). Several built-in modules of python will be utilized, namely:

- Numpy
- Pandas
- Scikit-Learn
- Keras
- Matplotlib
- NiaPy

The computations were performed on Ubuntu 16.04 LTS with following hardware specifications:

- Processor Intel i7 6th Gen. broadwell processor.
- RAM 24 GB DDR4
- Secondary Memory 250 GB SSD
- Graphics Card Nvidia 1080Ti 11GB

B. Results Obtained

This segment shows the readmission prediction results acquired from various predictive models. The prediction accuracy will be differentiate on the basis of two variables final accuracy and AUC under ROC (Receiver Operating Characteristic) Curve[29]. Both final accuracy and area under ROC curve relies on the classifier's capability to rank examples for positive class, however in the case of final accuracy, it also relies on the capability to compute the threshold in the ordering to distinguish the positive class. Applying various models on the data, with default attributes, we got results as referenced in Table IV. Also we compare deep learning results with baseline models in Table V to show how deep learning helps in improving the accuracy.

- 1) Curves showing Receiver Operating Curve (ROC) and Loss Curve of different models:

Algorithm type	Algorithm	AUC	Accuracy	F1 Score Micro	F1 Score Macro
Deep Learning without Nature Inspired Algorithms	Long Short Term Memory	0.88	78.84%	0.80	0.76
	Long Short Term Memory Auto-Encoders	0.73	70.00%	0.70	0.62
	Convolution Long Short Term Memory	0.75	70.98%	0.71	0.66
	Convolution Long Short Term Memory Auto-Encoders	0.87	80.32%	0.80	0.78
	Bi-Directional Long Short Term Memory Auto-Encoders	0.85	77.99%	0.78	0.75
Deep Learning with Nature Inspired Algorithms	Long Short Term Memory with Bat Algorithm	0.75	70.42%	0.70	0.63
	Long Short Term Memory with Grey Wolf optimizer Algorithm	0.85	81.57%	0.79	0.76

TABLE IV
Comparison among different Models implemented

Methods	Area Under ROC Curve	Accuracy
Logistic Regression	0.54	58.09%
Support Vector Machines	0.54	59.37%
Decision Tree	0.52	55.68%
Random Forest	0.53	60.49%

TABLE V
Comparison with Baseline Models

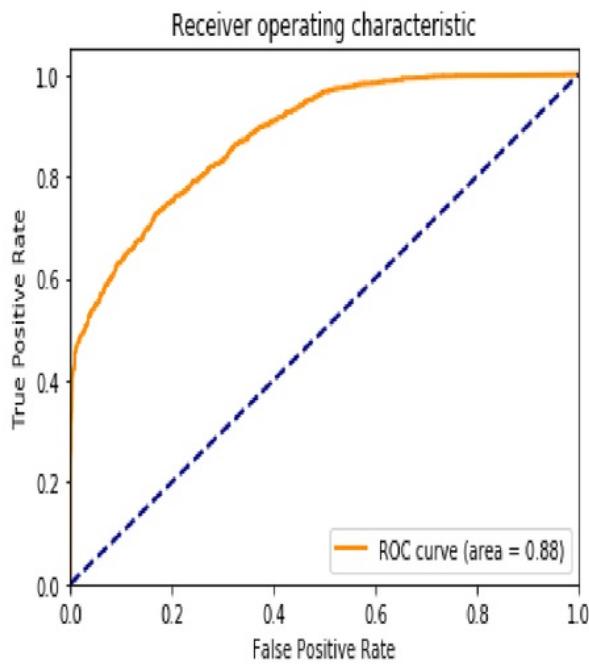


Fig. 7. Receiver Operating Characteristic Curve: Long Short Term Memory (LSTM).

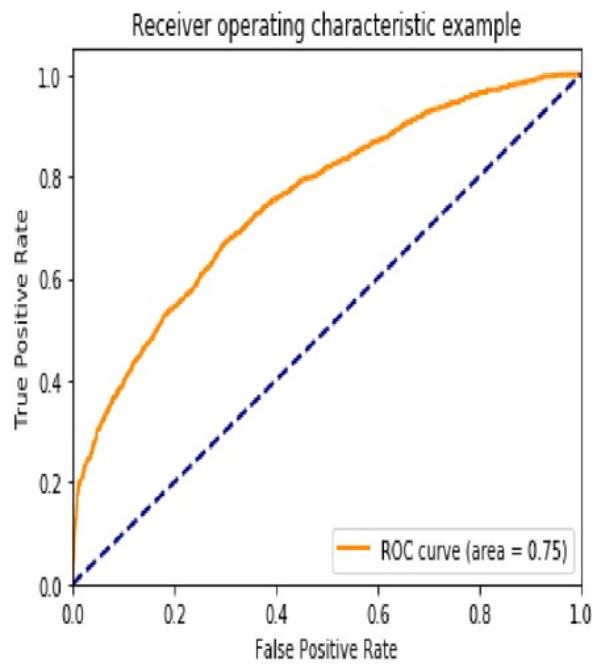


Fig. 9. Receiver Operating Characteristic Curve: Convolution Long Short Term Memory.

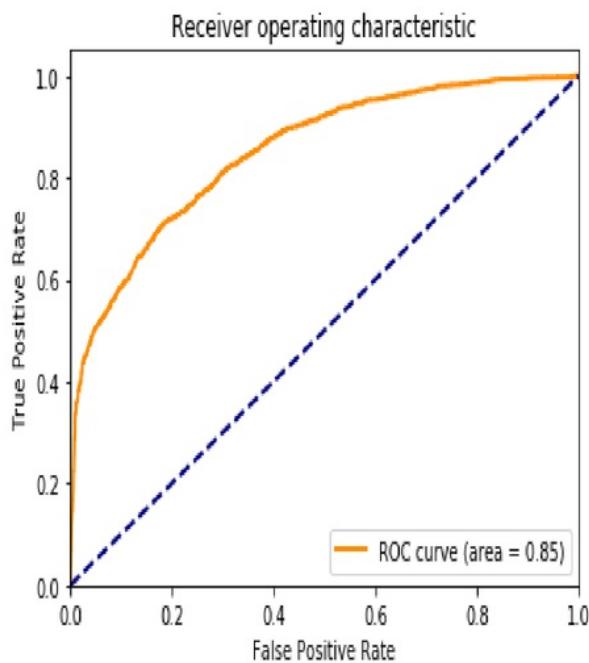


Fig. 8. Receiver Operating Characteristic Curve Bi-Directional Long Short Term Memory.

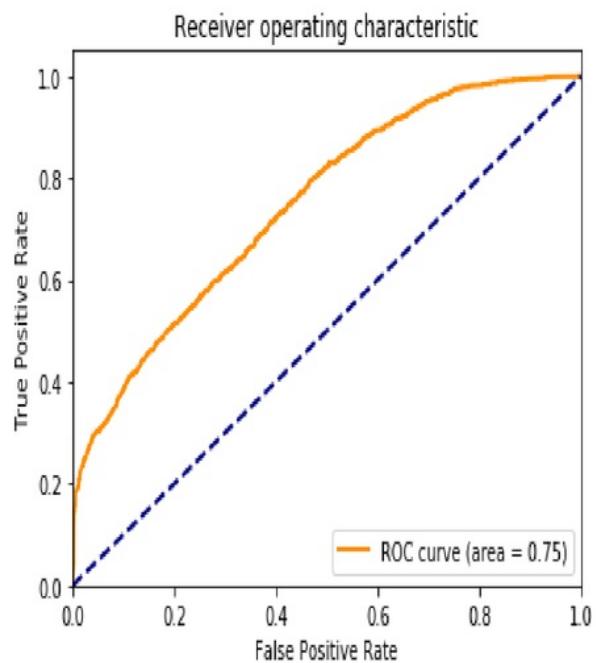
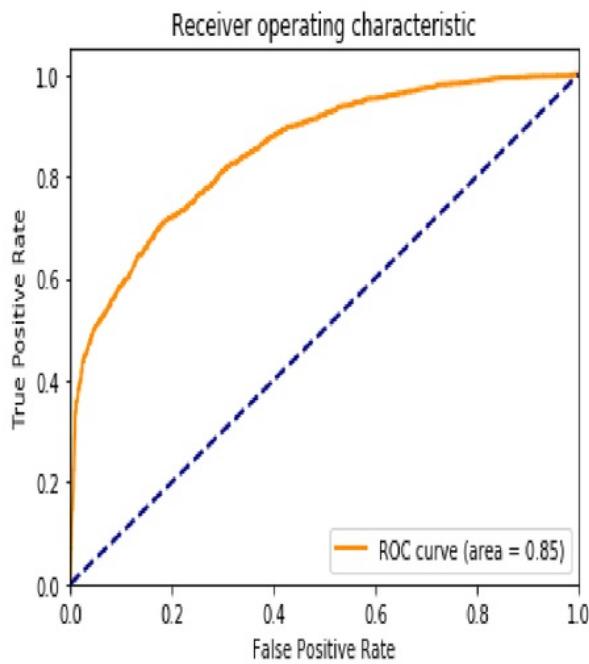
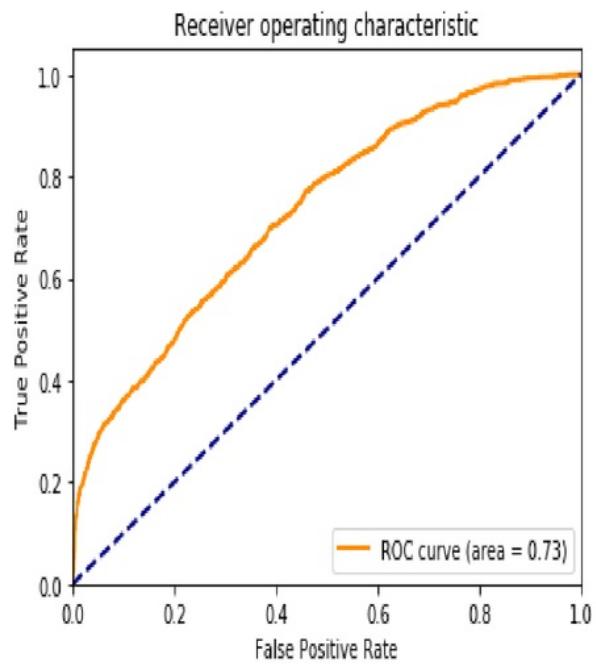


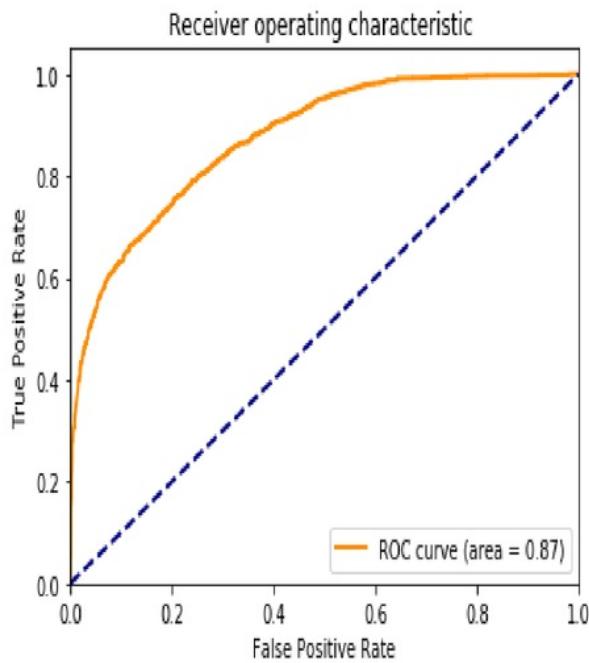
Fig. 10. Receiver Operating Characteristic Curve: Long Short Term Memory with Bat Algorithm.



6
Fig. 11. Receiver Operating Characteristic Curve: Long Short Term Memory with Grey Wolf Optimizer Algorithm.



6
Fig. 13. Receiver Operating Characteristic Curve: Long Short Term Memory Auto-Encoders.



6
Fig. 12. Receiver Operating Characteristic Curve Convolution Long Short Term Memory Auto Encoders.

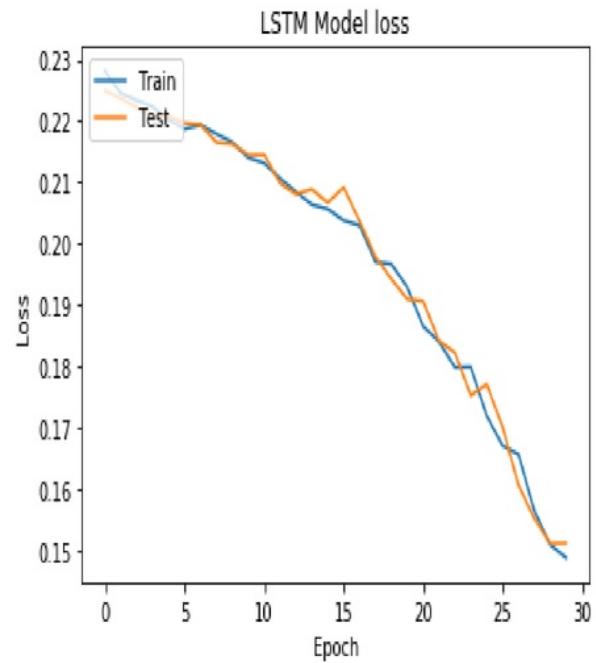


Fig. 14. Loss Curve: Long Short Term Memory.

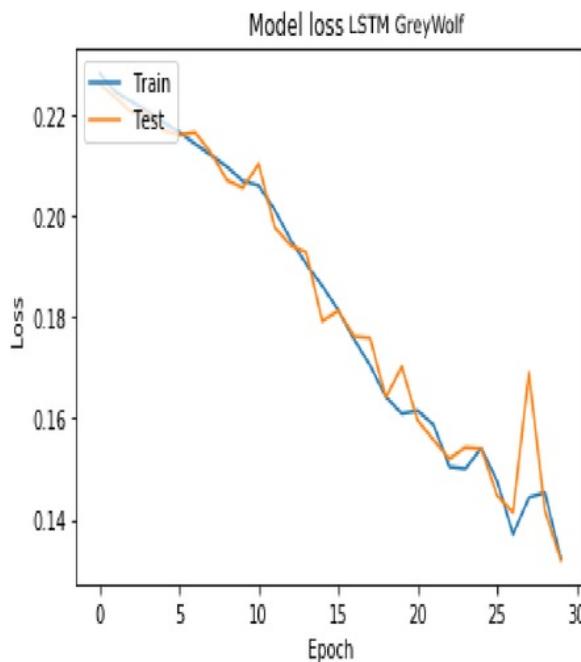


Fig. 15. Loss Curve: Long Short Term Memory Grey Wolf.

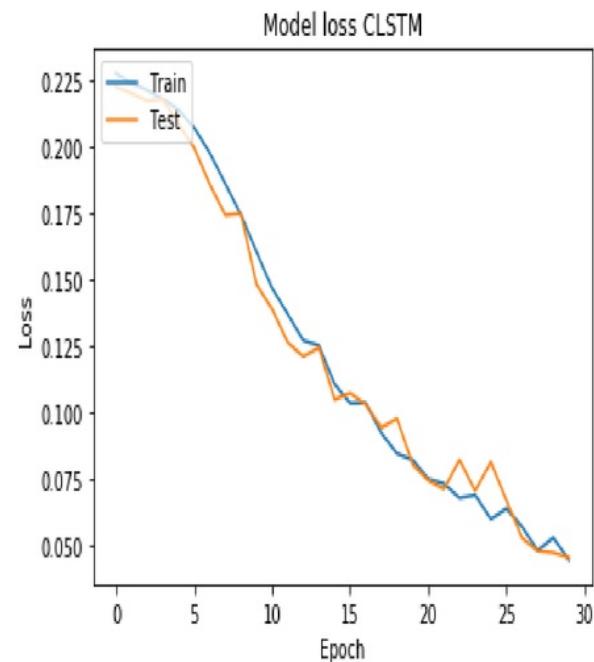


Fig. 17. Loss Curve: Convolution Long Short Term Memory.

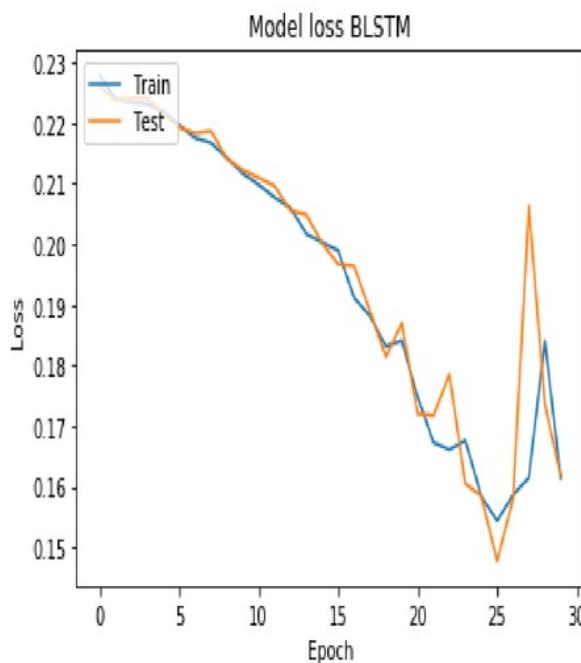


Fig. 16. Loss Curve: Bidirectional Long Short Term Memory.

V. CONCLUSION

Halth Care Readmissions is extremely difficult task for both sufferer at a health center. In our thesis, we design a robust algorithm to determine the amount of patients admitted again to a health care. We measured the performance of different deep learning algorithms with and without Nature-inspired algorithms to predict readmission probability and concluded that Long Short Term Memory with grey wolf optimizer performed better than the remaining ML algorithms in the prediction value. We also establish that the result of the combination of LSTM and Convolution layer was remarkable on this dataset. This architecture can be used in current's health system to aim at high possibility patients, decrease the degree of readmission, and provide excellent health care.

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