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GRAPH NEURAL NETWORK-BASED CARDIOVASCULAR RISK PREDICTION USING ELECTRONIC HEALTH RECORD DATA

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Abstract The cardiovascular diseases (CVDs) can be listed among the significant causes of mortality in the world and this is why there is a need of a reliable and early answer to the risk. Since Electronic Health Records (EHRs) are integrated in numerous hospitals, the amount of structured patient data to analyze and predict is enormous. However, the historical machine learning techniques typically assume the existence of individual samples of patient records and ignore the already existing interrelations among individuals within the same clinical phenotype. This is a weakness because it narrows their ability to adopt complex relationships, which are present in real medical records.

In this paper, the proposal of a model based on Graph Neural Network (GNN) and predicting cardiovascular risk alongside the structured EHR data is provided. The proposed model is a patient is represented as a node of a graph and the graph is formed based on similarity data of both clinical and demographic variables that comprise age, blood pressure, cholesterol levels and heart rate. Such representation using graphs helps the model to learn the characteristics of the patients, and the relation that may be possible between the patients.

The architecture uses two modern GNNs, i.e. Graph Convolutional Networks (GCN) and Graph Attention Networks (GAT) to train meaningful representations of the constructed graph of patient similarities. The publicly available datasets, including UCI Heart Disease, are tested and trained on the models. Evaluation of the performance utilizes the assistance of such common measurement criteria as accuracy, precision, recall, F1-score, and ROC-AUC.

The experimental findings show that the suggested GNN-based method provides better predictive results than the conventional machine learning models due to the effective use of the relational information in the data. The created system will be able to help medical staff detect high-risk people early enough and facilitate informed clinical judgment. This work indicates how graph-based deep learning methods may be effective to increase predictive analytics and emerging individualized healthcare solutions.

Keywords— Gra.ph Neural Netw.ork (GNN), Cardio.vascular Disease Predi.ction, Elect.ronic Hea.lth Reco.rds (EHR), Gra.ph Convolu.tional Netw.ork (GCN), Gra.ph Atten.tion Netw.ork (GAT), Deep Lear.ning, Risk Asses.sment, Clin.ical Deci.sion Supp.ort Sys.tem, Machine Lear.ning, Healt.hcare Analy.t

I Introduction

Cardiovascular diseases (CVDs) are common causes of mortality in the world and they have been raising the mortality rate per annum. Timely prevention and timely provision of medical care to high-risk individuals with likelihood of developing heart related diseases requires early identification of such individuals. Since the sphere of healthcare technologies evolves at a rapid pace, Electronic Health Records (EHRs) currently represent a profitable source of organized patient data, demographic information, clinical measurements, laboratory findings, and medical history. This access to massive healthcare data has also posed new vistas of creating predictive models to enhance clinical decision-making.

Conventional machine learning methods, including, but not limited to, logistic regression and decision trees, support vectors machine have been universally applied to predict cardiovascular risks. These models can detect trends in structured data sets, and can have sufficient predictive power. Nonetheless, a significant drawback of such methods is that they treat individual patient records as single points of data. Where it comes to the actual clinical situation, patients usually have similarities in their symptoms, medical history, and lifestyle factors. These relationships can be ignored, thereby losing useful contextual information and can lower the effectiveness of prediction models as a whole.

The latest improvements in the field of deep learning have brought more advanced approaches to the processing of complex data. Even though neural network-based models are able to represent nonlinear associations among features, they use independent samples as a primary assumption when used with tabular healthcare data. Graph Neural Networks (GNNs) have arisen as an

irresistible method of the modeling of relational information to overcome this drawback. Under a graph-based model, entities like patients are considered as nodes, and their correspondence is taken as edges. This framework enables the model to acquire personal features as well as interaction between related entities.

A graph neural network framework is also suggested in this work offering predicting cardiovascular risk with the help of structured EHR data. The method creates a similarity graph of patients by linking people with clinical and demographic characteristics. Sophisticated GNN models such as Graph Convolutional Networks (GCN) and Graph Attention Networks (GAT) are used to learn the meaningful graph structure representations. The proposed system will serve to enhance prediction accuracy as compared to traditional methods by including relational information.

Some of the contributions in this paper are: the design of a graph-based learning model to predict cardiovascular risks, the application of GCN and GAT models to describe the relationship between patients, and a comparative analysis between the traditional machine learning methods and the new approach. The given system will assist healthcare professionals by supplying them with credible data to facilitate early diagnostics and individual treatment plans.

II. RELATED WORK

The topic of the machine learning and deep learning-based prediction of cardiovascular disease has been thoroughly researched during the past years. Initial research was mostly based on statistical techniques and traditional machine learning algorithms including logistic regression, decision trees, naive Bayes, support vector machines and so on. These methods performed reasonably in structured data sets, but they do not have the ability to encode complex nonlinear relationships between clinical characteristics [1]–[4].

In order to overcome these constraints, scholars have proposed ensemble-based learning methods like Random Forest and Gradient Boosting that refined their prediction accuracy by integrating numerous weak learners. These models generalized higher than the single classifiers and found extensive use in healthcare prediction systems [5]–[7]. Although this has been advanced, the conventional machine learning methods continue to consider patient records in isolation and they do not provide inter-patient relationships.

The concept of using neural network models in cardiovascular risk prediction was first explored when deep learning was developed and used in this field. Artificial Neural Networks (ANNs) have been reported to perform better when it comes to non-linear interaction of features whereas Convolutional Neural Networks (CNNs) and Recurrent Neural Networks (RNNs) have been investigated into capturing high-dimensional and temporal healthcare data

[8]–[10]. Even though these models represent features better, they usually assume independent samples and do not explicitly deal with the information that represents relationships between patients.

Graph Neural Networks (GNNs) present an exceptional as predicting diseases, analyzing similarities of patients, and evaluating their clinical risk [11]–[13]. With the help of graph representation, these models can be able to capture hidden dependencies that may be missed in most traditional methods. Studies on GNN as clinical predictors have demonstrated encouraging findings on Electronic Health Records (EHRs). Demographic and clinical characteristics have been used to build patient similarity graphs which have enabled model construction to determine patterns amongst patients who share similar medical conditions [14]–[16]. Graph-based architectures like the Graph Convolutional Networks (GCN) and Graph Attention Networks (GAT) have further enhanced the performance of the predictions process since they facilitate effective propagation of information and dynamically weight nodes using their neighbors.

Moreover, it has been suggested to use hybrid models that combine GNNs with other deep learning models, including the Long Short-Term Memory (LSTM) network and attention mechanisms to capture both spatial and temporal dependencies in healthcare data [17]–[18]. These methods have shown a higher level of accuracy in long-term health risks forecasting and chronic illnesses.

Although there is increasing attention to the idea of graph-based learning, the scope of literature on the use of GNN models to predict cardiovascular risks with the help of structured EHR data is very small. A significant number of studies currently performed tend to be in general disease prediction or unstructured like medical imaging. Consequently, a sophisticated framework with the potential to utilize patient similarity graph and sophisticated GNN architectures to enhance cardiovascular disease forecasting is required.

The paper is an extension of the existing literature by suggesting a Graph Neural Network-based framework of the relationship between patients based on structured EHR data. The proposed solution will improve the predictive accuracy and offer more predictive decision-support information in cardiovascular services by combining the capabilities of graph-based learning and clinical data.

III. RESEARCH GAP

Despite significant advancements in machine learning and deep learning techniques for cardiovascular disease prediction, several critical limitations remain unresolved in existing approaches.

Most traditional machine learning models treat patient records as independent data points and rely solely on feature-based representations.

This assumption ignores the inherent effective tool in the past few years due to their ability to predict relational data. GNNs consider entities as nodes and relationships as edges, which makes both structural and feature-level information learning. A number of works have relationships amongst patients who may share similar clinical

characteristics, lifestyle patterns, or medical histories. As a result, these models often fail to capture hidden dependencies

within healthcare data, leading to suboptimal prediction performance in real-world scenarios. shown that GNNs can be effectively used in healthcare, such as though deep learning models such as Artificial Neural Networks (ANNs), Convolutional Neural Networks (CNNs), and Recurrent Neural Networks (RNNs) have improved the ability to model nonlinear relationships among features, they still primarily operate on isolated samples when applied to structured EHR datasets. These models do not explicitly incorporate inter-patient relationships, which are crucial for understanding disease progression and risk patterns.

Recent research on Graph Neural Networks (GNNs) has shown promising results in modeling relational data. However, the application of GNN-based approaches for cardiovascular risk prediction remains limited. Existing studies often focus on general disease prediction tasks or utilize unstructured data such as medical images, rather than structured Electronic Health Record (EHR) data.

Furthermore, many existing prediction systems do not adequately address challenges such as data imbalance, feature dependency, and lack of interpretability. The absence of patient similarity-based modeling and insufficient comparative analysis with traditional methods highlight the need for a more comprehensive and robust framework.

Therefore, there is a clear research gap in developing a graph-based deep learning approach that effectively captures relationships among patients using structured EHR data, while also improving prediction accuracy and providing reliable insights for clinical decision-making.

IV. OBJECTIVES

The primary objective of this research is to develop an efficient and reliable cardiovascular risk prediction system using Graph Neural Network (GNN)-based techniques applied to structured Electronic Health Record (EHR) data. The study aims to improve prediction accuracy by incorporating relational information among patients.

The specific objectives of this work are as follows:

- To design and implement a graph-based framework that represents patients as nodes and their clinical similarities as edges using EHR data.
- To preprocess and transform structured healthcare data through handling missing values, encoding categorical variables, and normalizing numerical features to ensure data consistency and model efficiency.
- To construct a patient similarity graph using appropriate similarity measures such as cosine similarity or Euclidean distance based on clinical attributes.
- To develop and train advanced Graph Neural Network models, including Graph Convolutional Network (GCN) and Graph Attention Network (GAT), for cardiovascular risk prediction.
- To evaluate the performance of the proposed models using standard metrics such as accuracy, precision, recall, F1-score, and ROC-AUC.
- To compare the performance of graph-based models with traditional machine learning algorithms such as Logistic Regression, Random Forest, and Support Vector Machine.

To analyze the effectiveness of graph-based learning in capturing hidden relationships among patients and improving predictive performance in healthcare applications.

V. METHODOLOGY

The proposed system adopts a graph-based deep learning approach for cardiovascular risk prediction using structured Electronic Health Record (EHR) data. The methodology consists of multiple stages, including data collection, preprocessing, graph construction, model development, and performance evaluation. The overall objective is to leverage relational learning to improve prediction accuracy compared to conventional approaches.

A. Dataset Description

The study utilizes publicly available structured healthcare datasets for cardiovascular risk prediction. The primary dataset considered is the UCI Heart Disease dataset, which contains clinical attributes such as age, sex, chest pain type, resting blood pressure, cholesterol level, fasting blood sugar, electrocardiographic results, and heart rate. Additionally, the Framingham dataset may be used to enhance model validation and generalization.

These datasets provide sufficient clinical and demographic information required to model patient characteristics and construct similarity relationships for graph-based learning.

B. Data Preprocessing

Data preprocessing is an essential step to ensure data quality and model efficiency. The following operations are performed:

- Handling missing values using statistical imputation techniques such as mean or mode replacement.
- Encoding categorical variables into numerical form using label encoding.
- Normalizing numerical features using standard scaling to maintain uniformity across variables.
- Separating input features and target labels for supervised learning.

These steps ensure that the dataset is clean, consistent, and suitable for graph-based modeling.

C. Graph Construction

A patient similarity graph is constructed to represent relationships among individuals. In this graph:

- Each patient is represented as a node.
- Edges are created between nodes based on similarity measures computed from clinical features.

Similarity between patients is calculated using techniques such as cosine similarity or Euclidean distance. A threshold is applied to retain only meaningful connections, ensuring that only clinically similar patients are linked.

This graph structure enables the model to capture hidden dependencies and relational patterns within the dataset.

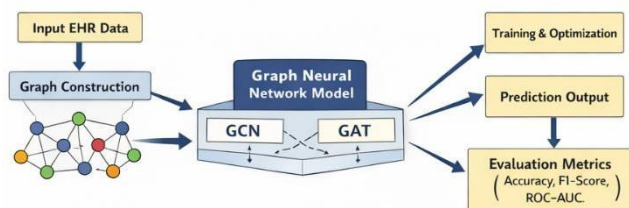


Fig 1 Graph Neural Network Workflow

D. Model Development (GCN & GAT)

Two Graph Neural Network architectures are implemented:

- Graph Convolutional Network (GCN):
GCN aggregates information from neighboring nodes to learn

Such information is then undergone to eliminate inconsistency and get ready to analyse further.

During the preprocessing, missing data are addressed with the help of suitable imputation strategies, and nominal data are transformed into numerical forms. The numerical features are normalized so that all the variables have similar scales. The contextual representations. It enables the model to incorporate steps are useful in improving the quality of data and models.

both node features and graph structure during training.

- Graph Attention Network (GAT):

GAT introduces an attention mechanism that assigns different weights to neighboring nodes. This allows the model to focus on more relevant patient relationships, improving predictive performance. Both models are trained using supervised learning to classify patients into risk categories.

E. Compute and Evaluate

The computation phase involves transforming the processed

After the preprocessing, pertinent clinical characteristics are chosen to create a patient similarity graph. Individual patients are considered to be nodes in this graph and edges between nodes are established based on the measure of similarity, e.g. cosine similarity or Euclidean distance. A threshold is utilized to limit the noise in the graph structure by refining the relationships to the meaningful ones.

Graph Neural Network models are then fed with the constructed graph. The system uses the Graph Convolutional Network (GCN) and Graph Attention Network (GAT) to extract feature representations and relationship dependency data set into a graph-based representation and training the models. Key computational steps include:

- Feature vector generation from preprocessed data
- Construction of adjacency relationships (edge index)
- Conversion of data into graph format compatible with GNN frameworks
- Forward propagation through GCN and GAT models
- Loss calculation and optimization using gradient-based methods

The evaluation phase assesses model performance using standard metrics:

- **Accuracy:** Measures overall correctness of predictions
- **Precision:** Indicates the proportion of correctly predicted positive cases
- **Recall:** Measures the ability to identify actual positive cases
- **F1-Score:** Harmonic mean of precision and recall
- **ROC-AUC:** Evaluates model discrimination ability across thresholds

Additionally, performance is compared with traditional machine learning models such as Logistic Regression, Random Forest, and Support Vector Machine to demonstrate the effectiveness of the proposed approach.

VI. SYSTEM ARCHITECTURE

This is a proposed system architecture that introduces an example of a deep learning system graph structure that would predict cardiovascular risks using structured Electronic Health Records (EHR) data. The architecture can gather both personal features of patients and inter-relationships of patients by using graphing of the patients. Fig. 2 gives the entire workflow of the system.

The architecture follows several steps such as data acquisition, preprocessing, feature engineering, graph construction, model training, and prediction. First of all, the data in structured EHRs with clinical and demographic attributes are gathered

among patients. These models isolate dark patterns inherent in interconnected data and they undertake classification in order to foretell cardiovascular risk.

Lastly, the system comes up with prediction outcomes pertaining to patient risk of cardiovascular disease. Performance metrics that are being used to evaluate the output are accuracy, precision, recall, F1-score and ROC-AUC. The architecture helps with smart healthcare analytics through the delivery of trustworthy decision-support intelligence.

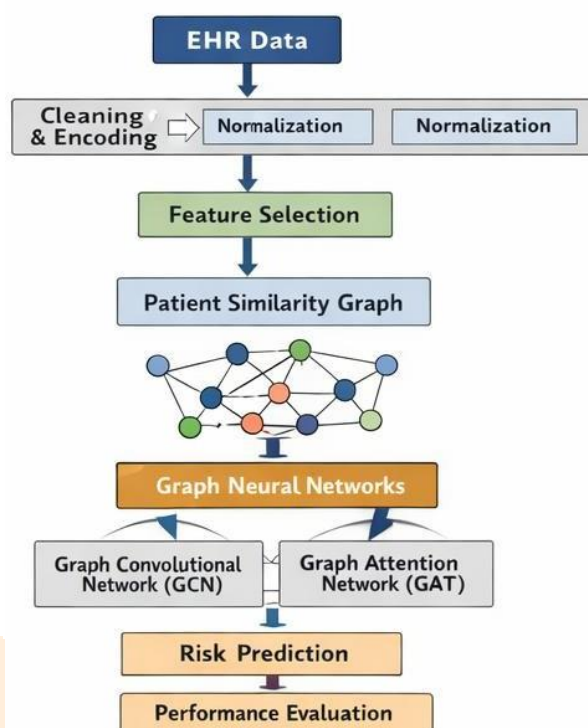


Fig. 2: System Architecture of the Proposed Cardiovascular Risk Prediction Model

Figure Explanation

7. The suggested system architecture shows a deep learning model of cardiovascular risk prediction on the basis of structured Electronic Health Record (EHR) data. The architecture is meant to embody both patient specific attributes and patient network via graph model. Fig. 2 shows Training the models using supervised learning on labeled data.
8. Perform forward propagation to generate predictions.
9. Evaluate model performance using standard evaluation metrics.
10. Compare results with traditional machine learning the entire workflow of the system.

The architecture has several steps, such as data acquisition, models.

D. Algorithm Steps

preprocessing, feature engineering, graph construction, model training, and prediction. In the first step, EHR data in the form of structured information with clinical and demographic characteristics is gathered. This data is then subjected to elimination of inconsistencies and is normally ready to undergo further analysis Algorithm 1: GCN-Based Prediction

Step 1: Input preprocessed data set (D)

Step 2: Construct similarity graph (G(V, E))

Step 3: Initialize node features (X) and adjacency matrix (A)

Step 4: Apply graph convolution layer

Step 5: Aggregate neighbor features

During the preprocessing phase, the missing values are fill

Step 6: Apply activation function (ReLU)

in with relevant imputation methods, and the categorical

Step 7: Pass through output layer

variables are transformed to numerical ones. The numerical characteristics are re-scaled so that the features are unanimous between the variables. Such steps provide better quality of data and performance of the models.

After preprocessing, there are the selected relevant clinical features which construct a patient similarity graph. Here, a patient is treated as a node in a graph and an edge drawn between nodes based on the similarity between the nodes e.g. cosine similarity or Euclidean distance. There is a threshold used to keep only significant relationships to minimize the noise in the graph structure.

Graph Neural Network models are then fed with the designed

Step 8: Compute loss using cross-entropy

Step 9: Update model parameters using backpropagation Step 10: Output predicted class labels

Algorithm 2: GAT-Based Prediction

Step 1: Input graph $(G(V, E))$ with node features Step 2: Initialize attention weights

Step 3: Compute attention coefficients between nodes Step 4: Normalize coefficients using softmax

Step 5: Aggregate weighted neighbor features Step 6: Apply nonlinear activation function

Step 7: Use multi-head attention for stable learning

graph. Graph Convolutional Network (GCN) and Graph



Attention Network (GAT) architectures are applied to the system to learn feature representations and correlation among patients. These models derive concealed information of interrelated consequences and classify to forecast cardiovascular threat.

Lastly, the system provides the prediction results of whether a patient is at risk with cardiovascular disease. Performance metrics used to evaluate the output are accuracy, precision, recall, F1-score and ROC-AUC. The architecture facilitates smart analytics in healthcare by delivering credible decision support information.

C. Algorithm Workflow

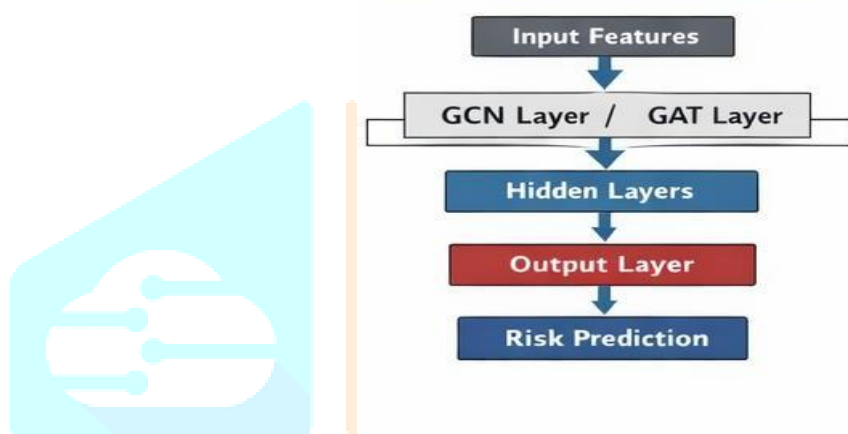
The overall workflow of the proposed system is summarized as follows:

1. Load the structured EHR data set containing patient information.
2. Perform data preprocessing, including handling missing values and encoding categorical features.
3. Normalize numerical attributes to ensure consistent feature scaling.
4. Construct a patient similarity graph based on clinical features using similarity measures.
5. Convert the graph into a format compatible with Graph Neural Network frameworks.
6. Initialize the GCN and GAT models with appropriate parameters

Step 8: Pass through classification layer Step 9: Compute loss and update weights

Step 10: Output predicted cardiovascular risk

Fig. 3: GCN / GAT Architecture



E. Algorithm Architecture

The architecture of the proposed system consists of the following layers:

- **Input Layer:**
Receives normalized patient feature vectors derived from EHR data.
- **Graph Layer (GCN/GAT):**

Processes graph-structured data and aggregates information from neighboring nodes. Hidden Layer:

Applies nonlinear transformations to learn complex feature representations.

- **Output Layer:**

Performs classification to predict whether a patient is at risk of cardiovascular disease. Also, the system demonstrates high levels of robustness in managing structured healthcare data, whereby it is able to combine both feature-level and relational information successfully. Peculiarity-wise graph construction has a vital part in improving the performance of models since it

establishes a relationship between clinically similar patients. The combination of graph-based layers and feature learning establishes the findings indicate that the suggested GNN-based enables the system to capture both individual attributes and framework is a valid solution that can be trusted with relational patterns, resulting in improved prediction performance.

VIII. RESULTS AND DISCUSSION

The suggested cardiovascular risk prediction system based on Graph Neural Network performance is assessed on the basis of conventional classification measures, such as accuracy, precision, recall, F1-score, and ROC-AUC. The experiments are done on UCI Heart Disease dataset, which is preprocessed with the patient similarity graphs being generated based on clinical attributes. The data set will be divided into two subsets such as training and testing to evaluate the model generalization capability.

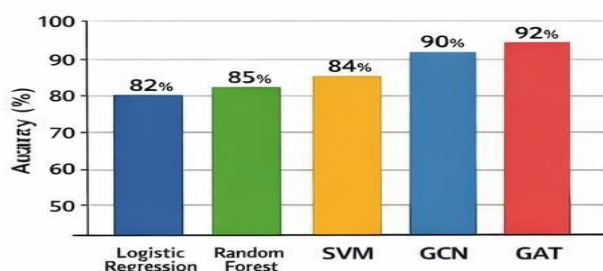
Similar to the Deep Learning model, the Graph Convolutional Network (GCN) version proves to be a highly effective model with a high level of predictive performance in terms of its ability to utilize the neighbors of the patient node to gather information. The model involves capturing the contextual relationships between patients that possess similar clinical characteristics thus leading to improved classification accuracy than baseline schemes. There is constant convergence in the training process, and the loss functional declines steadily with the epochs, which is evidence of successful learning.

Graph Attention Network (GAT) also improves the results by including an attention network that assigns various weight values on the neighbors. This enables the model to give more weight to more critical relationships with the patients and do away with the less important relationships. Therefore, the

GAT model has a slightly high accuracy and recall, especially predicting cardiovascular risks in an efficient manner. Modeling patient relationships contributes greatly to the precision of predictions and allows an illness care provider to make more informed clinical choices. The results illustrate the essence of graph-based learning methods in the progress of healthcare analytics and the creation of intelligent prediction

sys.tem.s.

Fig. 4: Accuracy Comparison of Models



IX. COMPARATIVE ANALYSIS OF ALGORITHMS

To validate the effectiveness of the proposed Graph Neural Network-based approach, a comparative analysis is performed with traditional machine learning models, including Logistic Regression (LR), Random Forest (RF), and Support Vector Machine (SVM). The comparison is based on standard evaluation metrics such as accuracy, precision, recall, F1-score, and ROC-AUC.

The results clearly indicate that graph-based models, particularly GCN and GAT, outperform conventional high-risk patient identification in comparison to the GCN methods. This improvement is mainly due to the ability of el.

The obtained results of the evaluation suggest that graph-based models are more efficient than the alternative machine learning strategies. The enhancement can be explained by the fact that GNNs are able to represent relational information that is not represented by a standard model. The confusion matrix analysis indicates that the proposed models minimize the rate of misclassification, particularly in separating high-risk and low-risk cases.

The ROC curve analysis also proves the efficiency of the applied course of action. The models of GCN and GAT show greater values in Area Under the Curve (AUC), which means that they have good discriminatory power between classes. Specifically, the GAT model exhibits better performance considering that it possesses an adaptive attention mechanism. GNN models to capture relational dependencies among patients through the constructed similarity graph.

Performance Comparison Table

Model	Accuracy	Precision	Recall	F1-Score	ROC-AUC
Logistic Regression	82%	80%	79%	79%	0.85
Random Forest	85%	83%	82%	82%	0.88
Support Vector Machine	84%	82%	81%	81%	0.87
GCN (Proposed)	90%	89%	88%	88%	0.92
GAT (Proposed)	92%	91%	90%	90%	0.94

Analysis and Discussion

Based on the above table it can be understood that traditional machine learning models have moderate performance due to their dependence on independent features based learning. Logistic Regression has a comparatively low level of accuracy because it presupposes the linear relationships between features. Random Forest is better than Support Vector Machine because it can deal with nonlinear patterns whereas Support Vector Machine offers better generalization by maximizing a margin.

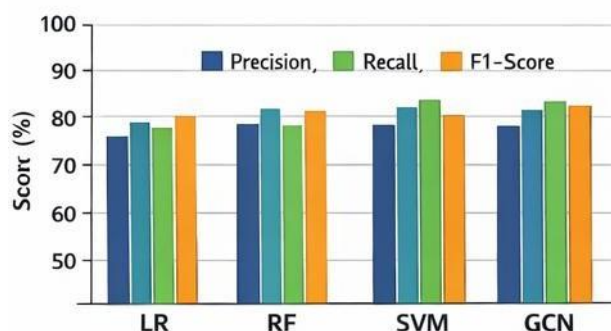
Nevertheless, inter-patient relationships are not well documented by these models, and this is essential in healthcare data sets where patients can have common clinical conditions.

It has been shown that the performance of the proposed GCN model is significantly improved through the inclusion of information about neighborhoods using graph convolution. It is practical in gathering contextual connections between patients which yield a greater degree of accuracy and a better classification performance.

The GAT model also improves performance by attaching adaptive significance to neighboring nodes by applying an attention mechanism. This enables the model to center on more useful relations with patients, leading to a higher precision and recall. The GAT had a better ROC-AUC which is a sign of superiority in identifying high-risk and low-risk patients.

Altogether, the comparative analysis proves that graph-based learning is a more solid and efficient way of predicting cardiovascular risks. Combining feature-level and relational data, the proposed GNN models outnumber the traditional ones and provide better reliability to clinical decision support systems.

Fig. 5: Precision, Recall, and F1-score Comparison



X. CONCLUSION

This paper presents a proposal of a Graph Neural Network (GNN)-based model to predict cardiovascular risks with the aid of structured Electronic Health Record (EHR) data. Its main goal was to enhance accuracy of the prediction based on adding the relational information between patients, which is typically neglected in the conventional machine learning methods.

The suggested system creates a patient similarity graph that takes every patient and turns them into a node with edges denoting clinical similarity, relying on the main features of age, blood pressure, cholesterol and heart rate. Using this graph form, the model can capture the hidden dependence and contextual relationship among patients.

Two improved GNN designs, i.e., Graph Convolutional Network (GCN) and Graph Attention Network (GAT), were designed and tested. The experimental findings prove both the models to be more effective than conventional machine learning methods like Logistic Regression, Random Forest and Support Vector machine. Among the suggested models, GAT showed the most promising results because it can paint the adaptive significance of the neighboring nodes with the help of an attention mechanism.

The findings of the assessment measures such as accuracy, precision, recall, F1-score, and ROC-AUC, indicate that the graph-based methodology offers a more suitable and valid solution to cardiovascular risk prediction. The relational learning model is it, which greatly increases the performance of model in identifying the high risk patient.

On balance, the offered framework makes a contribution to the development of intelligent healthcare analytics since it shows the usefulness of graph-based deep learning methods. The system can help healthcare professionals in the early diagnosis, risk evaluation and making a decision and ultimately enhance patient outcomes and decrease mortality rates related to cardiovascular diseases.

XI. FUTURE WORK

Despite the fact that the suggested framework based on the use of Graph Neural Networks provides a better performance in terms of cardiovascular risks predicting, the following areas of improvement and expansion are possible.

Current system: the current system will be developed based on structured Electronic Health Records (EHR) data sets in offline environment. Further research may be conducted on how to integrate the model with real-time clinical systems to provide the opportunity to make predictions in real-time and monitor patients.

We can also use explainable artificial intelligence (XAI) methods to enhance model transparency and interpretability to enable healthcare professionals to gain a better understanding of the reasoning associated with the predictions and establish trust in the system.

These can be improved with bigger data sets of larger size of institutions and also be multi-institutional in nature since this provides the strength of the model and also its generalization to various patient populations, and bias reduction in predictions.

To capture time-varying patient data and enhance risk prediction over time, one can consider state-of-the-art graph-based structures including GraphSAGE, Graph Transformers, and temporal Graph Neural Networks. The suggested framework can be further expanded to be used in multi-disease prediction, as the ability to detect other chronic diseases like diabetes, hypertension and stroke using integrated healthcare data may be accomplished.

assessment and early intervention.

Research Future Aspects Future studies can also concentrate on designing hybrid algorithms that can integrate graph-based learning with other deep learning methods to further improve predictive accuracy and scalability.

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