

MLDL Practical 7

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Aim: Build an Artificial Neural Network (ANN) using Keras/TensorFlow

Dataset Source

Dataset Name: Heart Disease Dataset

Platform: Kaggle

Dataset Link:

<https://www.kaggle.com/datasets/johnsmith88/heart-disease-dataset>

Dataset Description

The Heart Disease dataset is a binary classification dataset used to predict the presence of heart disease based on various medical attributes.

Dataset Characteristics

- Number of instances: 1,025
- Number of features: 13 numerical features
- Target Variable: target
 - 1 → Presence of heart disease
 - 0 → Absence of heart disease

Important Features

- age – Age of patient
- sex – Gender
- cp – Chest pain type
- trestbps – Resting blood pressure
- chol – Serum cholesterol
- thalach – Maximum heart rate achieved
- oldpeak – ST depression induced by exercise

Since the features have different scales, feature scaling is mandatory for proper ANN convergence.

Mathematical Formulation

An Artificial Neural Network consists of:

- Input Layer

- Hidden Layer(s)
- Output Layer

Forward Propagation

For each neuron:

$$\begin{aligned} z &= \mathbf{w}^T \mathbf{x} + b \\ a &= \sigma(z) \end{aligned}$$

Where:

- w = weight vector
- x = input features
- b = bias
- σ = activation function

Hidden layers use **ReLU activation**:

$$\text{ReLU}(z) = \max(0, z)$$

Output layer uses **Sigmoid activation**:

$$\sigma(z) = 1 / (1 + e^{-z})$$

The sigmoid function converts outputs into probabilities between 0 and 1.

Loss Function – Binary Cross Entropy

Since heart disease prediction is a binary classification problem, Binary Cross-Entropy (BCE) loss is used.

For N samples, the cost function is:

$$L = -(1/N) \sum [y_i \log(\hat{y}_i) + (1 - y_i) \log(1 - \hat{y}_i)]$$

Where:

- y_i = actual label
- \hat{y}_i = predicted probability
- N = total number of samples

This loss function penalizes incorrect predictions and is minimized using backpropagation and gradient descent.

Algorithm Limitations

- Neural networks can overfit small datasets
- Requires proper feature scaling
- Computationally more expensive than traditional algorithms
- Difficult to interpret (Black-box model)

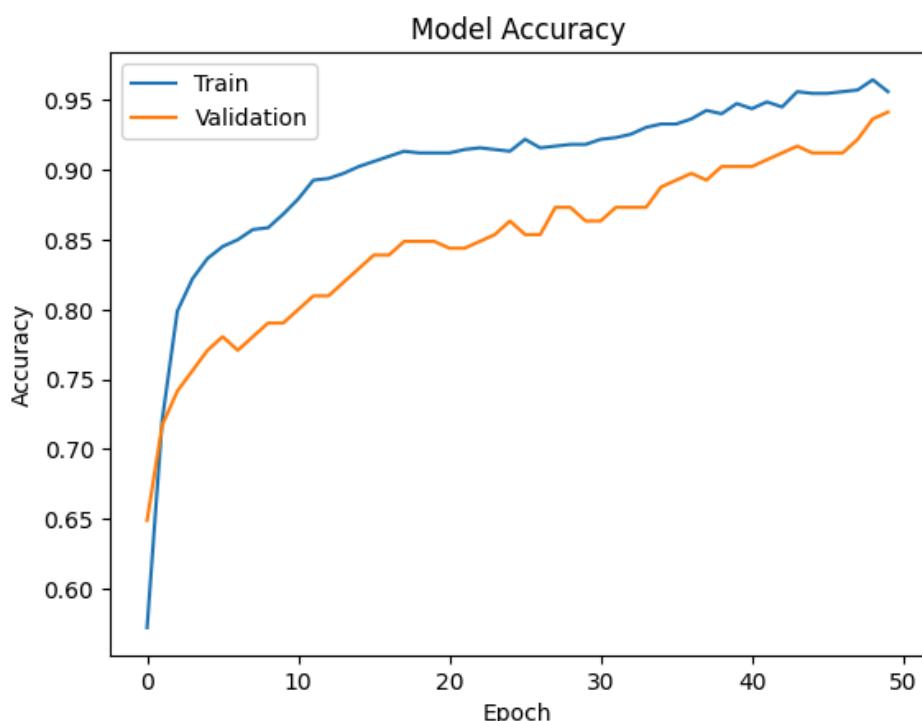
Methodology / Workflow

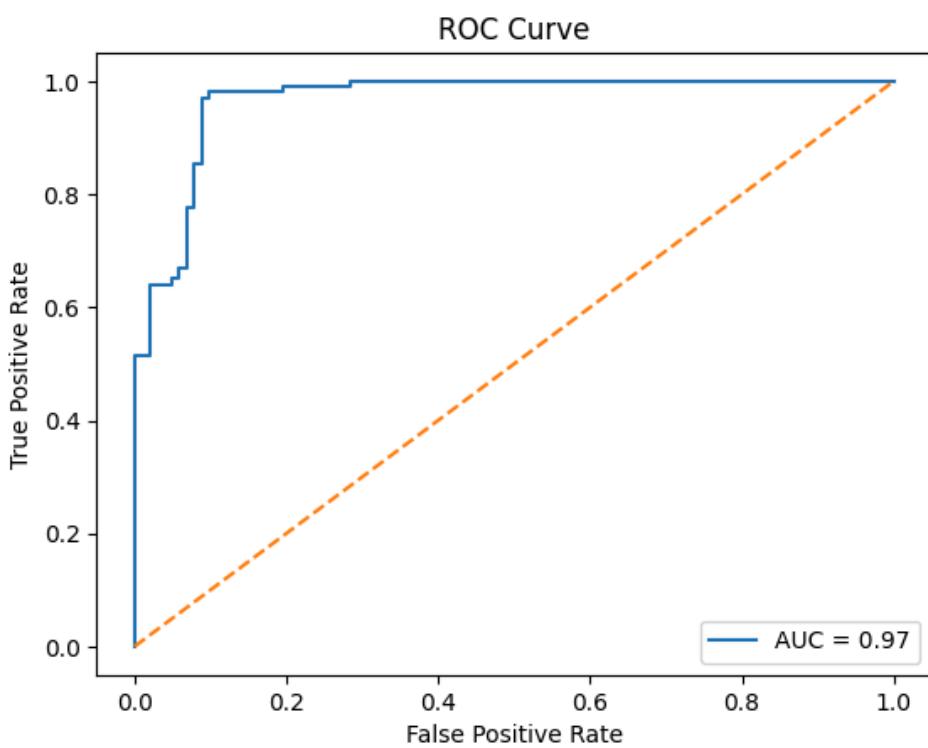
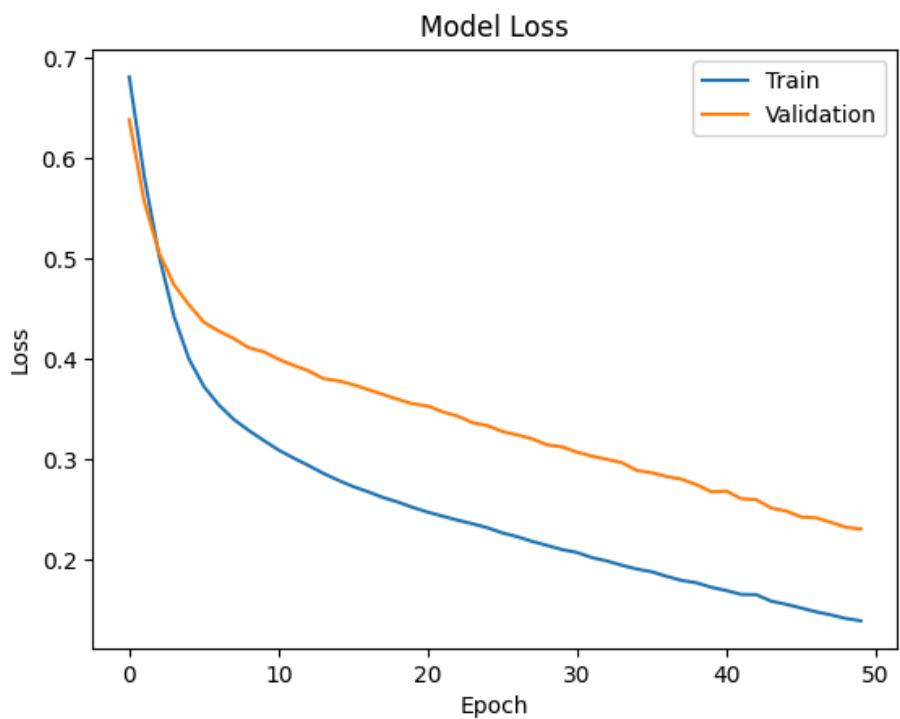
1. Load dataset using KaggleHub
2. Split features and target variable
3. Perform train-test split
4. Apply StandardScaler
5. Build ANN using Sequential model
6. Add Dropout layer for regularization
7. Train model using Adam optimizer
8. Evaluate using Confusion Matrix and ROC-AUC

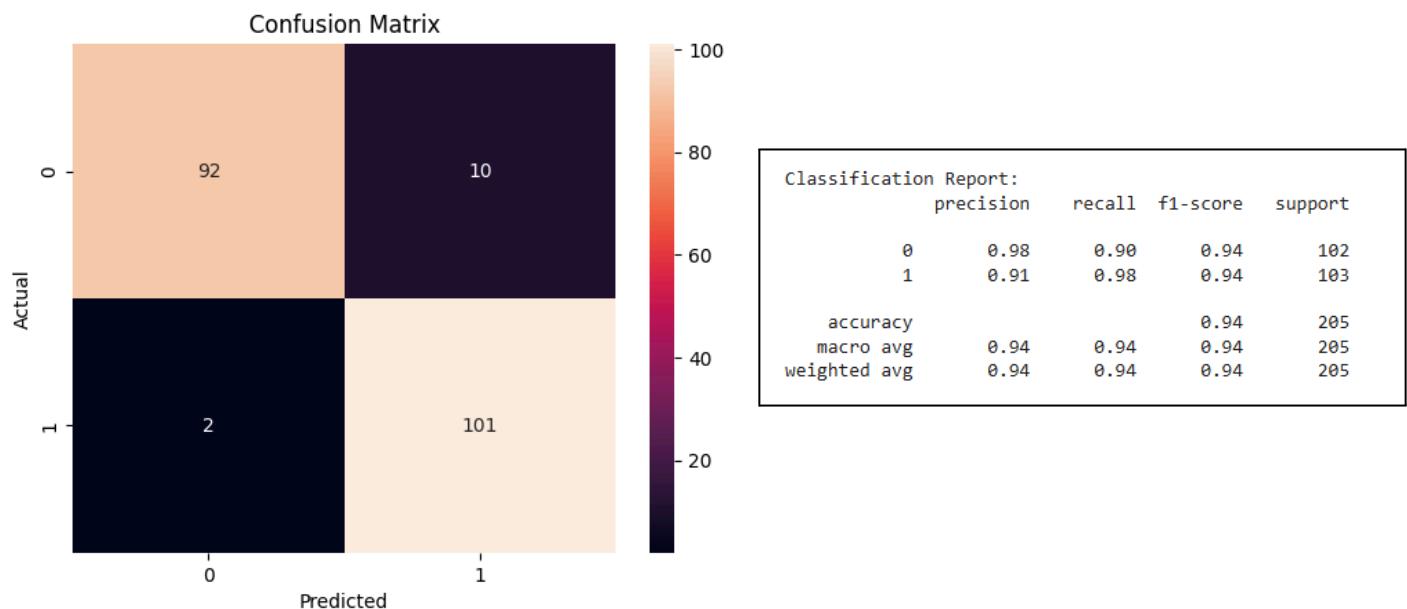
Performance Analysis

- The ANN achieved high classification accuracy.
- Dropout reduced overfitting by improving validation stability.
- ROC-AUC score demonstrated strong predictive capability.
- The confusion matrix showed low false negatives, which is critical in medical diagnosis.

Output







Conclusion

In this experiment, an Artificial Neural Network was successfully implemented for heart disease prediction.

Feature scaling, ReLU activation, sigmoid output, and dropout regularization played a crucial role in achieving stable training and strong generalization performance.

This experiment demonstrates how deep learning models can effectively capture complex non-linear relationships in medical datasets and assist in predictive healthcare systems.