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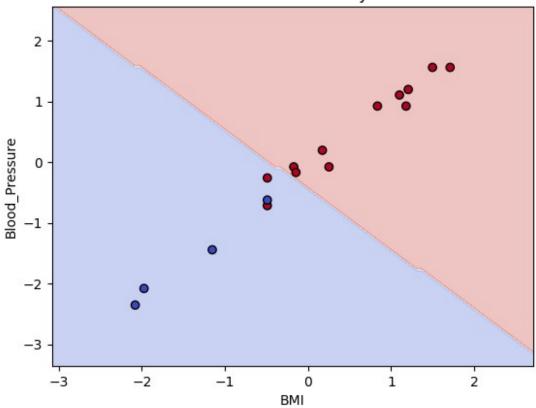
Batch: T(4)

# **Logistic Regression from Scratch**

```
import numpy as np
import pandas as pd
import matplotlib.pyplot as plt
from sklearn.model selection import train test split
from sklearn.preprocessing import StandardScaler
class LogisticRegressionScratch:
    def init (self, learning rate=0.01, iterations=1000):
        self.learning rate = learning rate
        self.iterations = iterations
        self.weights = None
        self.bias = None
    def sigmoid(self, z):
        return 1 / (1 + np.exp(-z))
    def fit(self, X, y):
        m, n = X.shape
        self.weights = np.zeros(n)
        self.bias = 0
        for in range(self.iterations):
            linear model = np.dot(X, self.weights) + self.bias
            y pred = self.sigmoid(linear model)
            dw = (1/m) * np.dot(X.T, (y_pred - y))
            db = (1/m) * np.sum(y pred - y)
            self.weights -= self.learning rate * dw
            self.bias -= self.learning rate * db
    def predict(self, X):
        linear model = np.dot(X, self.weights) + self.bias
        y pred = self.sigmoid(linear model)
        return (y pred >= 0.5).astype(int)
# Load dataset
file path = "/content/Records-Patient.csv"
df = pd.read csv(file path)
```

```
# Selecting two features for visualization
target col = 'Risk Level (Target)'
features = ['BMI', 'Blood Pressure'] # Change these as needed
X = df[features]
y = df[target col]
# Splitting dataset
X_train, X_test, y_train, y_test = train_test_split(X, y,
test size=0.2, random state=42)
# Standardizing features
scaler = StandardScaler()
X train = scaler.fit transform(X train)
X test = scaler.transform(X test)
# Training the model
model = LogisticRegressionScratch(learning_rate=0.1, iterations=1000)
model.fit(X train, y train)
# Predictions and accuracy
y pred = model.predict(X test)
accuracy = np.mean(y_pred == y_test)
print(f"Accuracy: {accuracy * 100:.2f}%")
# Plotting decision boundary
def plot decision boundary(X, y, model):
    x_{min}, x_{max} = X[:, 0].min() - 1, X[:, 0].max() + 1
    y_{min}, y_{max} = X[:, 1].min() - 1, X[:, 1].max() + 1
    xx, yy = np.meshgrid(np.linspace(x min, x max, 100),
                         np.linspace(y min, y max, 100))
    grid = np.c [xx.ravel(), yy.ravel()]
    Z = model.predict(grid)
    Z = Z.reshape(xx.shape)
    plt.contourf(xx, yy, Z, alpha=0.3, cmap='coolwarm')
    plt.scatter(X[:, 0], X[:, 1], c=y, edgecolors='k',
cmap='coolwarm')
    plt.xlabel(features[0])
    plt.vlabel(features[1])
    plt.title('Decision Boundary')
    plt.show()
plot decision boundary(X test, y test.values, model)
Accuracy: 87.50%
```

#### **Decision Boundary**



```
import numpy as np
import pandas as pd
import matplotlib.pyplot as plt
# Logistic function (sigmoid)
def logistic(x):
    return 1 / (1 + np.exp(-x))
# Log loss function
def log_loss(y, y_dash):
    return - (y * np.log(y_dash)) - ((1 - y) * np.log(1 - y_dash))
# Cost function using vectorization
def cost_func_vec(y, y_dash):
    m = \overline{len(y)}
    loss_vec = np.array([log_loss(y[i], y_dash[i]) for i in range(m)])
    cost = np.dot(loss_vec, np.ones(m)) / m
    return cost
# Cost function in terms of model parameters (using vectorization)
def cost logreg vec(X, y, w, b):
    m, n = X.shape
    z = np.matmul(X, w) + b
```

```
v dash = logistic(z)
    return cost func vec(y, y dash)
# Gradient computation
def compute gradients(X, y, w, b):
    m = len(y)
    z = np.matmul(X, w) + b
    y dash = logistic(z)
    dw = np.dot(X.T, (y dash - y)) / m
    db = np.sum(y dash - y) / m
    return dw, db
# Gradient Descent for logistic regression
def gradient descent(X, y, w, b, learning rate, iterations):
    cost history = []
    for i in range(iterations):
        dw, db = compute gradients(X, y, w, b)
        # Update weights and bias
        w -= learning rate * dw
        b -= learning rate * db
        # Compute cost after updating
        cost = cost logreg vec(X, y, w, b)
        cost history.append(cost)
    return w, b, cost history
# Full logistic regression model
def logistic regression(X, y, learning rate=0.01, iterations=1000):
    m, n = X.shape
    w = np.zeros(n) # Initialize weights as zero
                     # Initialize bias as zero
    w, b, cost history = gradient descent(X, y, w, b, learning rate,
iterations)
    return w, b, cost history
# Load dataset
file path = "/content/Records-Patient.csv"
df = pd.read csv(file path)
# Selecting two features for visualization and the target
features = ['BMI', 'Blood Pressure'] # Select two features for
visualization
target col = 'Risk Level (Target)'
```

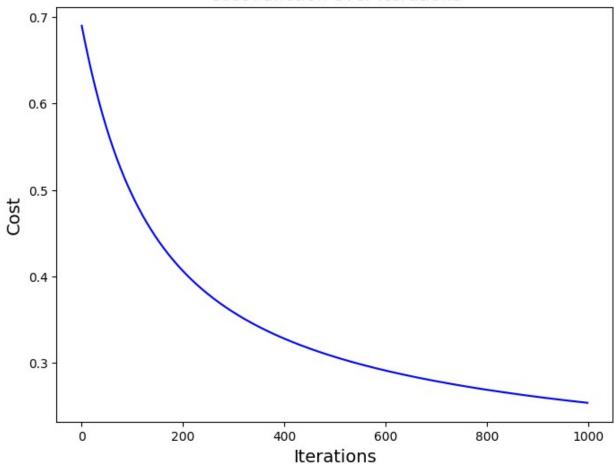
```
X = df[features].values
y = df[target col].values
# Normalize the features
X mean = np.mean(X, axis=0)
X \text{ std} = \text{np.std}(X, axis=0)
X = (X - X_mean) / X_std
# Train the logistic regression model
learning rate = 0.01
iterations = 1000
w, b, cost history = logistic regression(X, y, learning rate,
iterations)
# Plotting the cost function history
plt.figure(figsize=(8, 6))
plt.plot(range(len(cost_history)), cost_history, color='blue')
plt.xlabel("Iterations", fontsize=14)
plt.ylabel("Cost", fontsize=14)
plt.title("Cost Function over Iterations", fontsize=14)
plt.show()
# Predict function
def predict(X, w, b):
    z = np.matmul(X, w) + b
    return logistic(z)
# Example prediction
predictions = predict(X, w, b)
predictions = (predictions >= 0.5).astype(int)
print("Predictions:", predictions)
# Plotting the decision boundary
def plot_decision_boundary(X, y, w, b):
    plt.figure(figsize=(8, 6))
    # Create grid points to plot decision boundary
    x_{min}, x_{max} = X[:, 0].min() - 1, X[:, 0].max() + 1
    y_{min}, y_{max} = X[:, 1].min() - 1, X[:, 1].max() + 1
    xx, yy = np.meshgrid(np.linspace(x min, x max, 100),
                         np.linspace(y min, y max, 100))
    Z = predict(np.c [xx.ravel(), yy.ravel()], w, b)
    Z = Z.reshape(xx.shape)
    # Plot decision boundary
    plt.contourf(xx, yy, Z, alpha=0.8, cmap='coolwarm')
    # Plot data points
    plt.scatter(X[:, 0], X[:, 1], c=y, edgecolors='k', marker='o',
```

```
cmap='coolwarm')

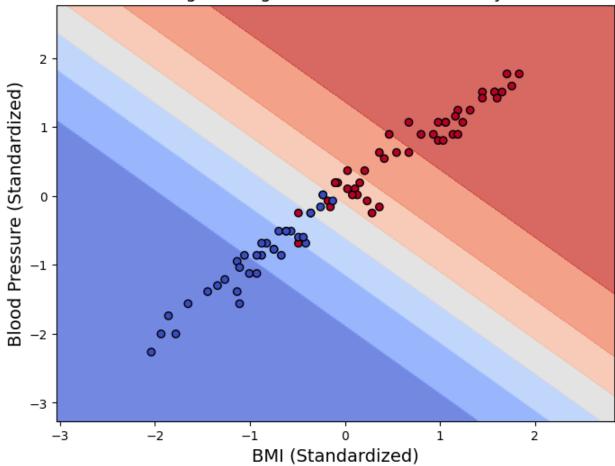
plt.xlabel('BMI (Standardized)', fontsize=14)
 plt.ylabel('Blood Pressure (Standardized)', fontsize=14)
 plt.title("Logistic Regression Decision Boundary", fontsize=14)
 plt.show()

# Plot the decision boundary
plot_decision_boundary(X, y, w, b)
```

#### Cost Function over Iterations







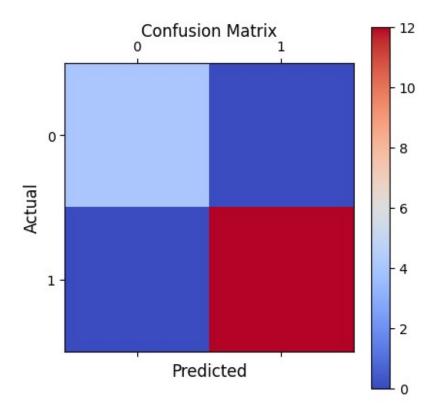
# **Logistic Regression with Library**

```
import numpy as np
import pandas as pd
import matplotlib.pyplot as plt
from sklearn.model_selection import train_test_split
from sklearn.preprocessing import StandardScaler
from sklearn.linear_model import LogisticRegression
from sklearn.metrics import accuracy_score, classification_report,
confusion_matrix

# Load the patient records dataset
file_path = "/content/Records-Patient.csv" # Update the file path if
necessary
df = pd.read_csv(file_path)

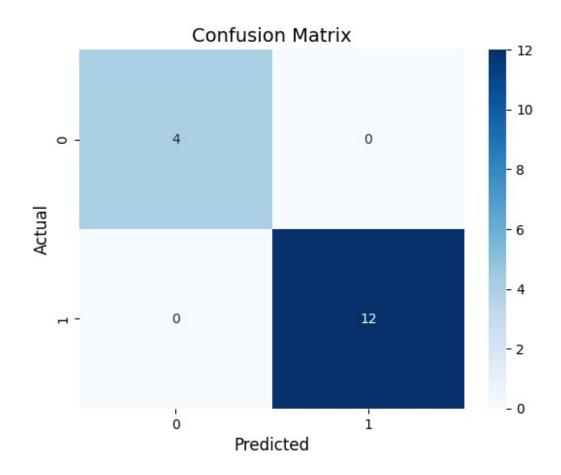
# Features and target selection
X = df[['Age', 'BMI', 'Blood_Pressure', 'Cholesterol', 'Smoking',
```

```
'Family History'll.values
y = df['Risk Level (Target)'].values # Target column
# Split into training and testing datasets
X train, X test, y train, y test = train test split(X, y,
test size=0.2, random state=42)
# Standardize the features
scaler = StandardScaler()
X train = scaler.fit transform(X train)
X test = scaler.transform(X test)
# Initialize and train the Logistic Regression model
model = LogisticRegression()
model.fit(X_train, y_train)
# Predictions
y_pred = model.predict(X_test)
# Evaluate the model
accuracy = accuracy score(y test, y pred)
conf matrix = confusion matrix(y test, y pred)
class report = classification report(y test, y pred)
# Display results
print(f'Accuracy: {accuracy:.2f}')
print('Confusion Matrix:\n', conf matrix)
print('Classification Report:\n', class report)
# Plot confusion matrix
plt.matshow(conf matrix, cmap='coolwarm')
plt.title('Confusion Matrix', pad=20)
plt.colorbar()
plt.xlabel('Predicted', fontsize=12)
plt.ylabel('Actual', fontsize=12)
plt.show()
Accuracy: 1.00
Confusion Matrix:
 [[4 0]
 [ 0 12]]
Classification Report:
                            recall f1-score
                                               support
               precision
                                                     4
           0
                   1.00
                             1.00
                                       1.00
           1
                   1.00
                             1.00
                                       1.00
                                                    12
                                       1.00
                                                    16
    accuracy
                   1.00
                             1.00
                                       1.00
                                                    16
   macro avg
```



```
# Import necessary libraries
import numpy as np
import pandas as pd
import matplotlib.pyplot as plt
import seaborn as sns
from sklearn.model selection import train test split
from sklearn.preprocessing import StandardScaler
from sklearn.linear model import LogisticRegression
from sklearn.metrics import accuracy score, confusion matrix,
classification report
# Load the dataset
file path = "/content/Records-Patient.csv" # Update with your dataset
path
df = pd.read_csv(file_path)
# Features and target
X = df[['Age', 'BMI', 'Blood_Pressure', 'Cholesterol', 'Smoking',
'Family History']].values
y = df['Risk Level (Target)'].values
# Split the dataset into training and testing sets
```

```
X train, X test, y train, y test = train test split(X, y,
test size=0.2, random state=42)
# Feature Scaling
scaler = StandardScaler()
X train = scaler.fit transform(X train)
X_test = scaler.transform(X_test)
# Initialize and train the Logistic Regression model
model = LogisticRegression()
model.fit(X train, y train)
# Make predictions
y pred = model.predict(X test)
# Evaluate the model
accuracy = accuracy_score(y_test, y_pred)
conf matrix = confusion_matrix(y_test, y_pred)
report = classification report(y test, y pred)
# Print results
print(f"Accuracy: {accuracy:.4f}")
print("Confusion Matrix:\n", conf matrix)
print("Classification Report:\n", report)
# Visualization of the confusion matrix
sns.heatmap(conf_matrix, annot=True, fmt="d", cmap="Blues")
plt.xlabel("Predicted", fontsize=12)
plt.ylabel("Actual", fontsize=12)
plt.title("Confusion Matrix", fontsize=14)
plt.show()
Accuracy: 1.0000
Confusion Matrix:
 [[ 4 0]
 [ 0 12]]
Classification Report:
               precision
                            recall f1-score
                                                support
           0
                                        1.00
                                                     4
                   1.00
                             1.00
           1
                   1.00
                             1.00
                                        1.00
                                                    12
                                        1.00
                                                    16
    accuracy
                             1.00
                                        1.00
                                                    16
   macro avg
                   1.00
weighted avg
                   1.00
                             1.00
                                        1.00
                                                    16
```



### **Activation Functions**

Sigmoid Function

```
import math

def sigmoid(x):
    return 1 / (1 + math.exp(-x))

sigmoid(100)

1.0

sigmoid(-2)

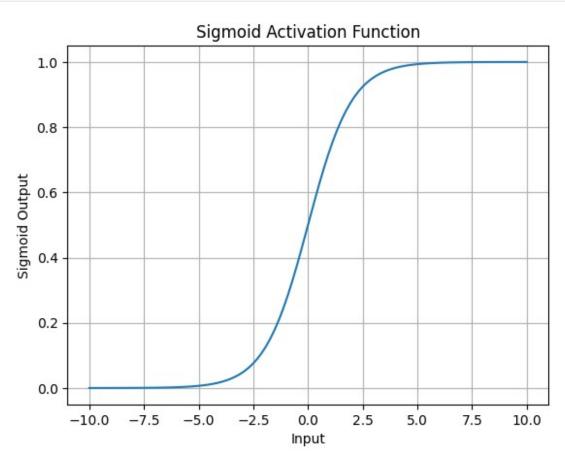
0.11920292202211755

import numpy as np
import matplotlib.pyplot as plt

def plot_sigmoid():
```

```
x = np.linspace(-10, 10, 100)
y = 1 / (1 + np.exp(-x))

plt.plot(x, y)
plt.xlabel('Input')
plt.ylabel('Sigmoid Output')
plt.title('Sigmoid Activation Function')
plt.grid(True)
plt.show()
```



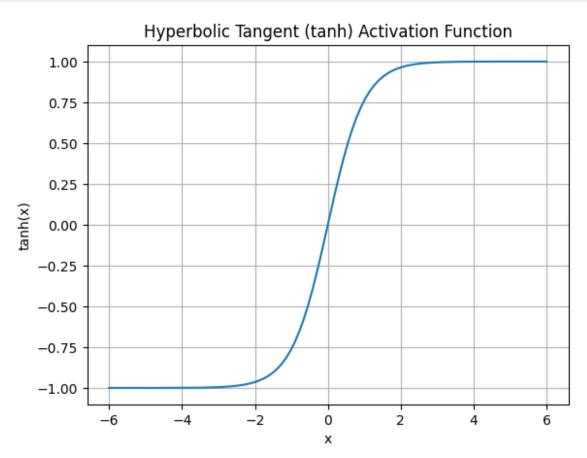
### **TANH Function**

```
def tanh(x):
    return (math.exp(x)-math.exp(-x))/(math.exp(x)+math.exp(-x))
tanh(-2)
-0.964027580075817
tanh(12)
```

```
0.9999999999244972
import numpy as np
import matplotlib.pyplot as plt

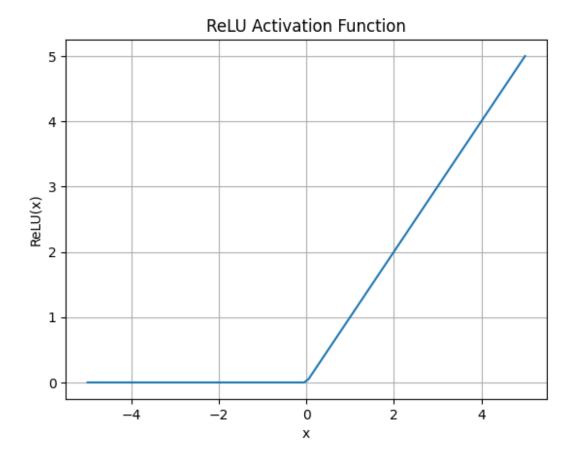
def plot_tanh():
    x = np.linspace(-6, 6, 100)
    tanh = np.tanh(x)

    plt.plot(x, tanh)
    plt.title("Hyperbolic Tangent (tanh) Activation Function")
    plt.xlabel("x")
    plt.ylabel("tanh(x)")
    plt.grid(True)
    plt.show()
```



### **RELU**

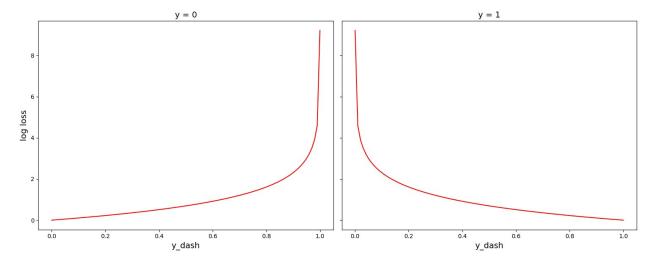
```
def relu(x):
  return max(0,x)
relu(-70)
0
relu(9)
9
import numpy as np
import matplotlib.pyplot as plt
def plot_relu():
    x = np.linspace(-5, 5, 100)
    relu = np.maximum(0, x)
    plt.plot(x, relu)
    plt.title("ReLU Activation Function")
    plt.xlabel("x")
plt.ylabel("ReLU(x)")
    plt.grid(True)
    plt.show()
    plot relu()
```



## **Log Loss Function**

```
# Log loss
def log loss(y, y_dash):
    """Computes log loss for inputs true value (0 or 1) and predicted
value (between 0 and 1)
   Args:
             (scalar): true value (0 or 1)
     y_dash (scalar): predicted value (probability of y being 1)
    Returns:
      loss (float): nonnegative loss corresponding to y and y dash
    loss = - (y * np.log(y_dash)) - ((1 - y) * np.log(1 - y_dash))
    return loss
y, y_dash = 1, 0.9
print(f"log_loss({y}, {y_dash}) = {log_loss(y, y_dash)}")
y, y dash = 0, 0.3
print(f"log_loss({y}, {y_dash}) = {log_loss(y, y_dash)}")
y, y_{dash} = 1, 0.7
print(f"log_loss({y}, {y_dash}) = {log_loss(y, y_dash)}")
```

```
y, y dash = 0, 0.2
print(f"log_loss({y}, {y_dash}) = {log_loss(y, y_dash)}")
log loss(1, 0.9) = 0.10536051565782628
\log \log (0, 0.3) = 0.35667494393873245
log loss(1, 0.7) = 0.35667494393873245
log loss(0, 0.2) = 0.2231435513142097
\# Log loss for y = 0 and y = 1
fig, ax = plt.subplots(1, 2, figsize = (15, 6), sharex = True, sharey
= True)
y_dash = np.linspace(0.0001, 0.9999, 100)
ax[0].plot(y_dash, log_loss(0, y_dash), color = 'red')
ax[0].set title("y = 0", fontsize = 14)
ax[0].set_xlabel("y_dash", fontsize = 14)
ax[0].set ylabel("log loss", fontsize = 14)
ax[1].plot(y dash, log loss(1, y dash), color = 'red')
ax[1].set_title("y = 1", fontsize = 14)
ax[1].set_xlabel("y_dash", fontsize = 14)
plt.tight layout()
plt.show()
```

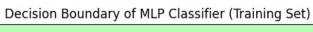


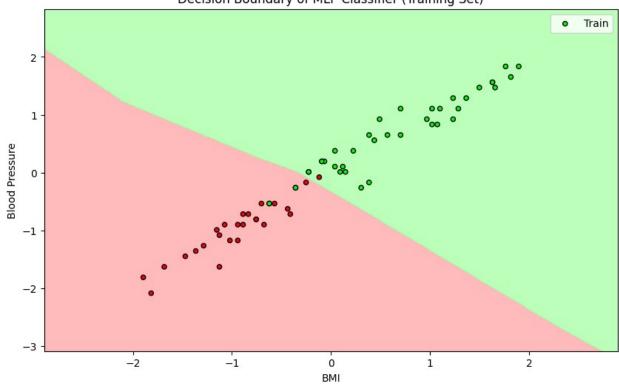
# Sklearn Implementation of MultiLayer Perceptron(MLP)

```
from sklearn.model_selection import train_test_split
from sklearn.preprocessing import StandardScaler
from sklearn.neural_network import MLPClassifier
from sklearn.metrics import accuracy_score, classification_report
import numpy as np
import pandas as pd
import matplotlib.pyplot as plt
```

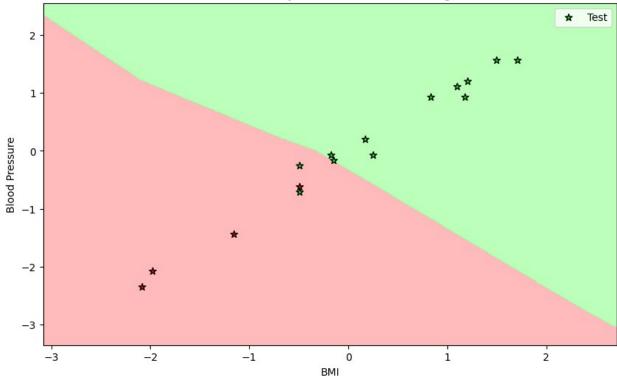
```
from matplotlib.colors import ListedColormap
# Load the Patient Records dataset
data = pd.read csv("/content/Records-Patient.csv")
X = data.iloc[:, 1:-1].values # Use all features except Patient ID
and Label
y = data.iloc[:, -1].values # Target labels
# Split the dataset into training and testing sets
X train, X test, y train, y test = train test split(X, y,
test size=0.2, random state=42)
# Standardize the features
scaler = StandardScaler()
X train = scaler.fit transform(X train)
X test = scaler.transform(X test)
# Define the main MLP classifier (trained on all features)
mlp = MLPClassifier(hidden layer sizes=(10, 10), max iter=1000,
random state=42)
mlp.fit(X train, y train)
# Make predictions
y pred = mlp.predict(X test)
# Evaluate the model
accuracy = accuracy score(y test, y pred)
print("Accuracy:", accuracy)
print("\nClassification Report:\n", classification report(y test,
y pred))
# Train a separate classifier for visualization (only on BMI & Blood
Pressure)
# ============
X vis = data.iloc[:, [2, 3]].values # Selecting BMI and Blood
Pressure
y_vis = data.iloc[:, -1].values # Target labels
# Split and scale the visualization dataset
X vis train, X vis test, y vis train, y vis test =
train test split(X vis, y vis, test size=0.2, random state=42)
scaler vis = StandardScaler()
X vis train = scaler vis.fit transform(X vis train)
X vis test = scaler vis.transform(X vis test)
# Define a new MLP model for visualization
mlp_vis = MLPClassifier(hidden_layer_sizes=(10, 10), max iter=1000,
random state=42)
mlp vis.fit(X vis train, y vis train)
```

```
# Plot decision boundary for training set
x \min, x \max = X \text{ vis train}[:, 0].\min() - 1, X \text{ vis train}[:, 0].\max() +
y_{min}, y_{max} = X_{vis}train[:, 1].min() - 1, <math>X_{vis}train[:, 1].max() + 1
xx, yy = np.meshgrid(np.arange(x_min, x_max, 0.01),
                     np.arange(y_min, y_max, 0.01))
Z train = mlp vis.predict(np.c [xx.ravel(), yy.ravel()])
Z_train = Z_train.reshape(xx.shape)
# Define color maps
cmap_light = ListedColormap(['#FFAAAA', '#AAFFAA'])
cmap bold = ListedColormap(['#FF0000', '#00FF00'])
plt.figure(figsize=(10, 6))
plt.contourf(xx, yy, Z_train, alpha=0.8, cmap=cmap_light)
plt.scatter(X vis train[:, 0], X vis train[:, 1], c=y vis train,
cmap=cmap bold, edgecolor='k', s=20, label='Train')
plt.title("Decision Boundary of MLP Classifier (Training Set)")
plt.xlabel("BMI")
plt.ylabel("Blood Pressure")
plt.legend()
plt.show()
# ================
# Plot decision boundary for testing set
x_{min}, x_{max} = X_{vis}test[:, 0].min() - 1, <math>X_{vis}test[:, 0].max() + 1
y \min, y \max = X \text{ vis test}[:, 1].\min() - 1, X \text{ vis test}[:, 1].\max() + 1
xx, yy = np.meshgrid(np.arange(x min, x max, 0.01),
                     np.arange(y min, y max, 0.01))
Z test = mlp vis.predict(np.c [xx.ravel(), yy.ravel()])
Z test = Z test.reshape(xx.shape)
plt.figure(figsize=(10, 6))
plt.contourf(xx, yy, Z test, alpha=0.8, cmap=cmap light)
plt.scatter(X_vis_test[:, 0], X_vis_test[:, 1], c=y_vis_test,
cmap=cmap_bold, edgecolor='k', s=50, label='Test', marker='*')
plt.title("Decision Boundary of MLP Classifier (Testing Set)")
plt.xlabel("BMI")
plt.ylabel("Blood Pressure")
plt.legend()
plt.show()
```









# Keras Implementation of MultiLayer Perceptron(MLP)

```
# Import necessary libraries
import numpy as np
import matplotlib.pyplot as plt
from keras.models import Sequential
from keras.layers import Dense
from sklearn.datasets import make_classification
from sklearn.model selection import train test split
from sklearn.preprocessing import StandardScaler
from sklearn.metrics import classification report,
ConfusionMatrixDisplay
# Step 1: Prepare a synthetic dataset
X, y = make classification(
    n samples=1000, n features=20, n informative=15, n redundant=5,
random state=42
# Split the dataset into training (80%) and testing (20%) sets
X_train, X_test, y_train, y_test = train_test_split(X, y,
test size=0.2, random state=42)
```

```
# Standardize the data (helps with convergence and performance)
scaler = StandardScaler()
X train = scaler.fit transform(X train) # Fit the scaler on training
data and transform it
X_test = scaler.transform(X_test) # Transform the testing data using
the same scaler
# Step 2: Build the ANN model
model = Sequential([
   Dense(32, activation='relu', input dim=X train.shape[1]), #
Hidden layer with 32 neurons and ReLU activation
   Dense(16, activation='relu'), # Another hidden layer with 16
neurons and ReLU activation
   Dense(1, activation='sigmoid') # Output layer with 1 neuron and
sigmoid activation for binary classification
# Step 3: Compile the model
model.compile(optimizer='adam', loss='binary crossentropy',
metrics=['accuracy'])
# Step 4: Train the model
history = model.fit(X train, y train, epochs=20, batch size=32,
validation split=0.2, verbose=1)
# Step 5: Evaluate the model
loss, accuracy = model.evaluate(X test, y test)
print(f"\nTest Loss: {loss:.4f}")
print(f"Test Accuracy: {accuracy:.4f}")
# Step 6: Generate predictions
y pred = (model.predict(X test) > 0.5).astype(int)
# Step 7: Classification Report
print("\nClassification Report:")
print(classification report(y test, y pred))
# Step 8: Visualize confusion matrix
ConfusionMatrixDisplay.from predictions(y test, y pred)
plt.show()
Epoch 1/20
/usr/local/lib/python3.11/dist-packages/keras/src/layers/core/
dense.py:87: UserWarning: Do not pass an `input shape`/`input dim`
argument to a layer. When using Sequential models, prefer using an
`Input(shape)` object as the first layer in the model instead.
  super(). init (activity regularizer=activity regularizer,
**kwarqs)
```

```
20/20 ______ 2s 16ms/step - accuracy: 0.5341 - loss:
1.0082 - val accuracy: 0.4938 - val loss: 0.8741
Epoch 2/20
                ———— 0s 5ms/step - accuracy: 0.5357 - loss:
20/20 ——
0.7633 - val accuracy: 0.5375 - val loss: 0.7021
Epoch 3/20 _____ 0s 7ms/step - accuracy: 0.6023 - loss:
0.6426 - val accuracy: 0.6938 - val loss: 0.6158
0.5541 - val accuracy: 0.7188 - val loss: 0.5557
Epoch 5/20 ______ 0s 7ms/step - accuracy: 0.7891 - loss:
0.5064 - val accuracy: 0.7625 - val loss: 0.5119
Epoch 6/20 ______ 0s 5ms/step - accuracy: 0.8585 - loss:
0.4480 - val accuracy: 0.8125 - val loss: 0.4713
Epoch 7/20
                ———— 0s 5ms/step - accuracy: 0.8883 - loss:
0.3956 - val accuracy: 0.8313 - val loss: 0.4364
Epoch 8/20
                _____ 0s 5ms/step - accuracy: 0.8906 - loss:
20/20 —
0.3623 - val accuracy: 0.8250 - val loss: 0.4074
Epoch 9/20
20/20 ————— Os 5ms/step - accuracy: 0.9176 - loss:
0.3330 - val accuracy: 0.8500 - val loss: 0.3845
0.3112 - val accuracy: 0.8500 - val loss: 0.3647
Epoch 11/20 ______ 0s 5ms/step - accuracy: 0.9110 - loss:
0.2744 - val accuracy: 0.8562 - val loss: 0.3479
Epoch 12/20
20/20 ———— Os 7ms/step - accuracy: 0.9133 - loss:
0.2567 - val accuracy: 0.8500 - val loss: 0.3302
Epoch 13/20
                Os 5ms/step - accuracy: 0.9112 - loss:
20/20 ——
0.2367 - val accuracy: 0.8500 - val loss: 0.3161
Epoch 14/20 Os 7ms/step - accuracy: 0.9092 - loss:
0.2279 - val accuracy: 0.8562 - val loss: 0.3067
Epoch 15/20

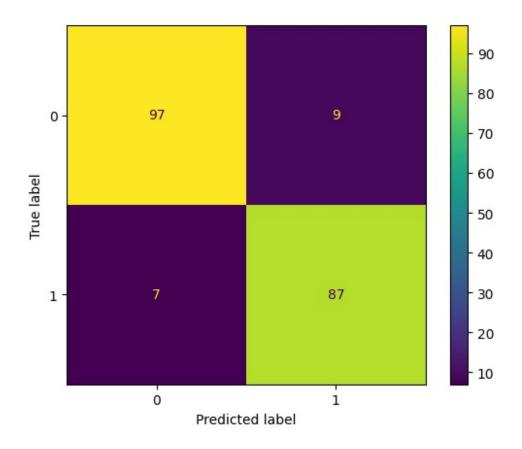
Os 5ms/step - accuracy: 0.9466 - loss:
0.1986 - val accuracy: 0.8562 - val loss: 0.2936
Epoch 16/20
20/20 ———— Os 7ms/step - accuracy: 0.9436 - loss:
0.1851 - val accuracy: 0.8625 - val_loss: 0.2910
Epoch 17/20 ______ 0s 6ms/step - accuracy: 0.9365 - loss:
0.1860 - val accuracy: 0.8750 - val_loss: 0.2795
```

Test Loss: 0.1964 Test Accuracy: 0.9200

7/7 — 0s 11ms/step

#### Classification Report:

CCGSSTITCGCT	on report.			
	precision	recall	f1-score	support
0	0.93	0.92	0.92	106
1	0.91	0.93	0.92	94
accuracy			0.92	200
macro avg	0.92	0.92	0.92	200
weighted avg	0.92	0.92	0.92	200

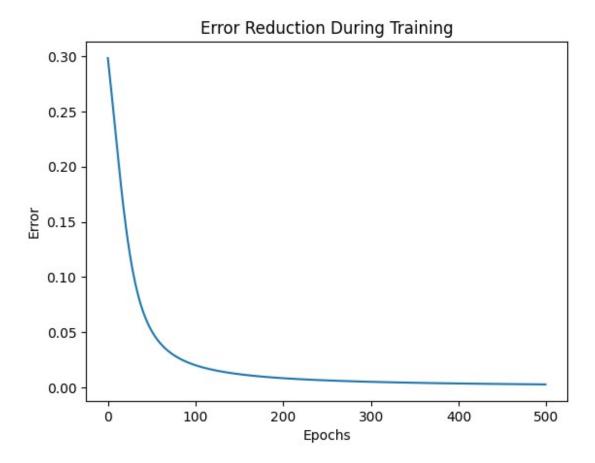


# **Backward Propogation from Sratch**

```
import math
import numpy as np
import matplotlib.pyplot as plt
# Sigmoid Activation Function
def sigmoid(x):
    return 1 / (1 + np.exp(-x))
# Derivative of Sigmoid Function
def sigmoid derivative(x):
    return x * (1 - x)
# Feedforward function
def feed forward(b1, b2, w1, w2, x):
    hidden = []
    output = []
    # Hidden Layer Activation
    hidden.append(sigmoid(b1 + np.dot(w1[:2], x))) # First neuron
    hidden.append(sigmoid(b1 + np.dot(w1[2:], x))) # Second neuron
```

```
# Output Layer Activation
   output.append(sigmoid(b2 + np.dot(w2[:2], hidden))) # First
output neuron
   output.append(sigmoid(b2 + np.dot(w2[2:], hidden))) # Second
output neuron
    return hidden, output
# Error Calculation
def find error(output, desired):
    return sum((np.array(output) - np.array(desired))**2) / 2
# Backpropagation
def back propagate(w1, w2, hidden, output, desired, x, alpha):
   # Compute error terms for output layer
   delta output = [(output[i] - desired[i]) *
sigmoid derivative(output[i]) for i in range(len(output))]
   # Compute error terms for hidden layer
   delta hidden = []
   for i in range(len(hidden)):
        temp = sum(delta output[j] * w2[i + j * len(hidden)] for j in
range(len(output)))
        delta hidden.append(temp * sigmoid_derivative(hidden[i]))
   # Update weights for hidden-to-output layer
   for i in range(len(w2)):
        w2[i] -= alpha * delta_output[i // len(hidden)] * hidden[i %
len(hidden)]
   # Update weights for input-to-hidden layer
   for i in range(len(w1)):
        w1[i] -= alpha * delta hidden[i // len(x)] * x[i % len(x)]
# Initialization
w1 = [0.15, 0.20, 0.25, 0.30] # Weights for input to hidden layer
w2 = [0.40, 0.45, 0.50, 0.55] # Weights for hidden to output layer
x = [0.05, 0.10] # Input values
b1 = 0.35 # Bias for hidden layer
b2 = 0.60 # Bias for output layer
desired = [0.01, 0.99] # Desired output
epochs = 500 # Training iterations
alpha = 0.5 # Learning rate
error = []
# Training Loop
for in range(epochs):
   hidden, output = feed_forward(b1, b2, w1, w2, x)
    error.append(find error(output, desired))
   back propagate(w1, w2, hidden, output, desired, x, alpha)
```

```
# Plot Error Reduction Over Time
plt.plot(error)
plt.xlabel("Epochs")
plt.ylabel("Error")
plt.title("Error Reduction During Training")
plt.show()
# Final Output after Training
print("Final Output:", output)
```



Final Output: [0.06389456919363433, 0.9404153154587165]

### Link of ipynb file:

https://colab.research.google.com/drive/1N6h82U250j5U4lCFPQ0aLbzefTKoO-JM?usp=sharing