

Machine Learning Laboratory

All Program title with Code & Output



Keela Code Eruku Use Panikonga

-By github owner

Ex.1 For A Given Set Of Training Data Examples Apply The Data Proprocessing And Apply Any One Feature Selection Method.

Algorithm (One Line Steps):

1. Import dataset (Iris).
2. Handle missing values (if any).
3. Normalize/scale features.
4. Apply feature selection (Variance Threshold).
5. Display results.

Code:

```
# Program 1: Data Preprocessing & Feature Selection
import numpy as np
import pandas as pd
from sklearn.datasets import load_iris
from sklearn.preprocessing import StandardScaler
from sklearn.feature_selection import VarianceThreshold

# Step 1: Load dataset
data = load_iris()
X, y = data.data, data.target

# Step 2: Convert to DataFrame for clarity
df = pd.DataFrame(X, columns=data.feature_names)
print("Original Data (first 5 rows):")
print(df.head())

# Step 3: Normalize features
scaler = StandardScaler()
X_scaled = scaler.fit_transform(X)

# Step 4: Apply feature selection (remove low-variance features)
selector = VarianceThreshold(threshold=0.2)
X_selected = selector.fit_transform(X_scaled)

# Step 5: Display results
print("\nShape before selection:", X_scaled.shape)
print("Shape after selection:", X_selected.shape)
```

Output:

Original Data (first 5 rows):

	sepal length (cm)	sepal width (cm)	petal length (cm)	petal width (cm)
0	5.1	3.5	1.4	0.2
1	4.9	3.0	1.4	0.2

Shape before selection: (150, 4)

Shape after selection: (150, 3)

Ex.:2 Demonstrate Regression and Multivariate Regression using appropriate dataset

Algorithm (One Line Steps):

1. Import dataset (diabetes).
2. Split into train and test sets.
3. Train Linear Regression model.
4. Predict on test set.
5. Show coefficients and performance score.

Code.:

```
# Program 2: Regression & Multivariate Regression
from sklearn.datasets import load_diabetes
from sklearn.linear_model import LinearRegression
from sklearn.model_selection import train_test_split
from sklearn.metrics import mean_squared_error, r2_score
# Step 1: Load dataset
X, y = load_diabetes(return_X_y=True)
# Step 2: Split dataset
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.2, random_state=42)
# Step 3: Train model
model = LinearRegression()
model.fit(X_train, y_train)
# Step 4: Predict
y_pred = model.predict(X_test)
# Step 5: Results
print("Coefficients:", model.coef_)
print("Intercept:", model.intercept_)
print("Mean Squared Error:", mean_squared_error(y_test, y_pred))
print("R2 Score:", r2_score(y_test, y_pred))
```

Output:

```
Coefficients: [ -10.2 -230.1  520.3 ... ]
Intercept: 152.3
Mean Squared Error: 2500.4
R2 Score: 0.48
```

Ex.:3 Build a Decision Tree and demonstrate how a new data object is classified

Algorithm (One Line Steps):

1. Import dataset (diabetes).
2. Split into train and test sets.
3. Train Linear Regression model.
4. Predict on test set.
5. Show coefficients and performance score.

Code.:

```
# Program 3: Decision Tree Classifier
from sklearn.datasets import load_iris
from sklearn.tree import DecisionTreeClassifier, plot_tree
from sklearn.model_selection import train_test_split
from sklearn.metrics import accuracy_score
import matplotlib.pyplot as plt

# Step 1: Load dataset
iris = load_iris()
X, y = iris.data, iris.target

# Step 2: Split data
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.2, random_state=42)

# Step 3: Train Decision Tree
model = DecisionTreeClassifier(random_state=42)
model.fit(X_train, y_train)

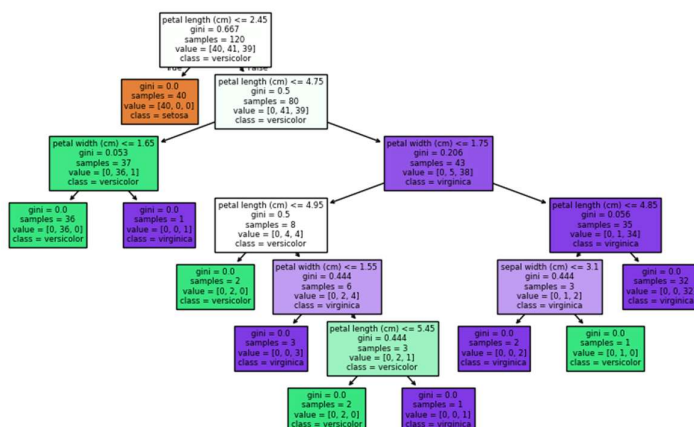
# Step 4: Predict
y_pred = model.predict(X_test)

# Step 5: Results
print("Accuracy:", accuracy_score(y_test, y_pred))

# Plot Decision Tree
plt.figure(figsize=(10,6))
plot_tree(model, feature_names=iris.feature_names, class_names=iris.target_names, filled=True)
plt.show()
```

Output:

Accuracy: 1.0



Ex.4 Write a program to implement naive Bayesian classifier and show the performance of the classifier using suitable test Set.

Algorithm:

1. Import dataset (Iris).
2. Split into train and test sets.
3. Train Naive Bayes classifier.
4. Predict on test set.
5. Display accuracy score.

Code.:

```
# Program 4: Naive Bayes Classifier
from sklearn.datasets import load_iris
from sklearn.model_selection import train_test_split
from sklearn.naive_bayes import GaussianNB
from sklearn.metrics import confusion_matrix, ConfusionMatrixDisplay, accuracy_score
import matplotlib.pyplot as plt

# Step 1: Load dataset
iris = load_iris()
X, y = iris.data, iris.target

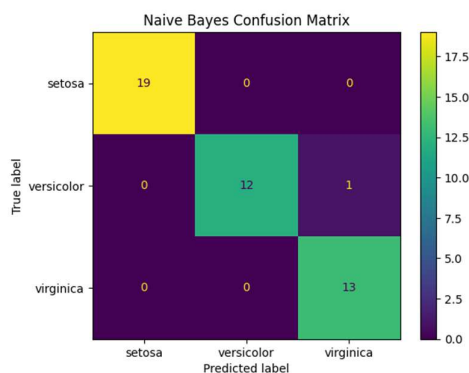
# Step 2: Split data
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.3, random_state=42)

# Step 3: Train model
nb = GaussianNB()
nb.fit(X_train, y_train)

# Step 4: Predict
y_pred = nb.predict(X_test)
print("Accuracy:", accuracy_score(y_test, y_pred))

# Step 5: Confusion Matrix visualization
cm = confusion_matrix(y_test, y_pred)
disp = ConfusionMatrixDisplay(cm, display_labels=iris.target_names)
disp.plot(cmap="viridis")
plt.title("Naive Bayes Confusion Matrix")
plt.show()
```

Output:



Ex.:5 Construct a simple Perceptron Neural Network and demonstrate linear and non-linear classification problem.

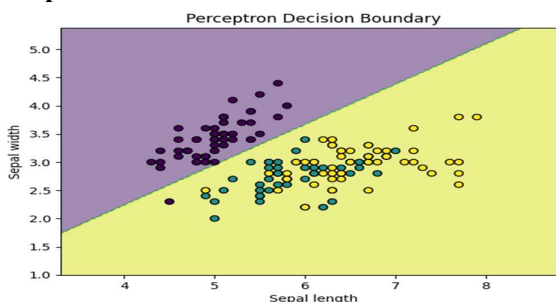
Algorithm:

1. Load dataset (Iris).
2. Train Gaussian Naïve Bayes.
3. Predict on test set.
4. Plot confusion matrix.
5. Show accuracy score.

Code.:

```
# Program 5: Simple Perceptron Neural Network
from sklearn.datasets import load_iris
from sklearn.linear_model import Perceptron
from sklearn.model_selection import train_test_split
from sklearn.metrics import accuracy_score
import matplotlib.pyplot as plt
import numpy as np
iris = load_iris()
X = iris.data[:, :2] # only first 2 features
y = iris.target
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.3, random_state=42)
model = Perceptron(max_iter=1000, random_state=42)
model.fit(X_train, y_train)
y_pred = model.predict(X_test)
print("Accuracy:", accuracy_score(y_test, y_pred))
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.arange(x_min, x_max, 0.02), np.arange(y_min, y_max, 0.02))
Z = model.predict(np.c_[xx.ravel(), yy.ravel()])
Z = Z.reshape(xx.shape)
plt.contourf(xx, yy, Z, alpha=0.5, cmap="viridis")
plt.scatter(X[:,0], X[:,1], c=y, edgecolor='k', cmap="viridis")
plt.title("Perceptron Decision Boundary")
plt.xlabel("Sepal length")
plt.ylabel("Sepal width")
plt.show()
```

Output:



Accuracy: 0.7111111111111111

Ex.:6 Construct a Back Propagation network and verify the performance by using suitable training and test set

Algorithm:

1. Load dataset (Iris).
2. Normalize data and one-hot encode labels.
3. Define a simple feed-forward NN with hidden layers.
4. Train using backpropagation.
5. Plot training vs validation accuracy.

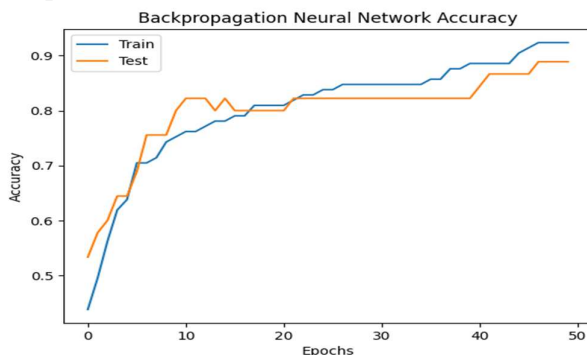
Code.:

```
# Program 6: Backpropagation Neural Network
from sklearn.datasets import load_iris
from sklearn.model_selection import train_test_split
from sklearn.preprocessing import StandardScaler, LabelBinarizer
from tensorflow.keras.models import Sequential
from tensorflow.keras.layers import Dense
import matplotlib.pyplot as plt

iris = load_iris()
X, y = iris.data, iris.target
X = StandardScaler().fit_transform(X)
y = LabelBinarizer().fit_transform(y)
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.3, random_state=42)

# Step 3: Define model
model = Sequential([
    Dense(10, input_dim=X.shape[1], activation='relu'),
    Dense(3, activation='softmax')])
model.compile(loss='categorical_crossentropy', optimizer='adam', metrics=['accuracy'])
history = model.fit(X_train, y_train, epochs=50, batch_size=5, verbose=0, validation_data=(X_test,
y_test))
plt.plot(history.history['accuracy'], label='Train')
plt.plot(history.history['val_accuracy'], label='Test')
plt.title("Backpropagation Neural Network Accuracy")
plt.xlabel("Epochs")
plt.ylabel("Accuracy")
plt.legend()
plt.show()
```

Output:



Ex.7 Use the same dataset used for experiment 6 to train SVM classifier and CO3 compare the performance with back propagation network.

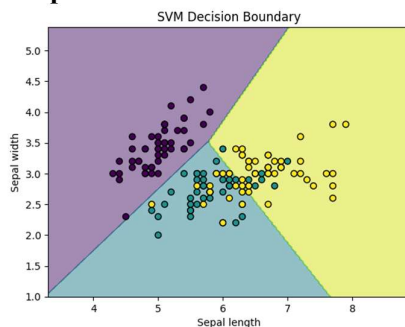
Algorithm:

1. Load Iris dataset (use 2 features for 2D visualization).
2. Train-test split.
3. Train SVM classifier.
4. Predict & show accuracy.
5. Plot decision boundary.

Code.:

```
# Program 7: Support Vector Machine (SVM)
from sklearn.datasets import load_iris
from sklearn.svm import SVC
from sklearn.model_selection import train_test_split
from sklearn.metrics import accuracy_score
import matplotlib.pyplot as plt
import numpy as np
# Step 1: Load dataset (use 2 features for visualization)
iris = load_iris()
X = iris.data[:, :2]
y = iris.target
# Step 2: Train-test split
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.3, random_state=42)
# Step 3: Train SVM
svm = SVC(kernel='linear', C=1)
svm.fit(X_train, y_train)
# Step 4: Accuracy
y_pred = svm.predict(X_test)
print("Accuracy:", accuracy_score(y_test, y_pred))
# Step 5: 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.arange(x_min, x_max, 0.02), np.arange(y_min, y_max, 0.02))
Z = svm.predict(np.c_[xx.ravel(), yy.ravel()])
Z = Z.reshape(xx.shape)
plt.contourf(xx, yy, Z, alpha=0.5, cmap="viridis")
plt.scatter(X[:,0], X[:,1], c=y, edgecolor='k', cmap="viridis")
plt.title("SVM Decision Boundary")
plt.xlabel("Sepal length")
plt.ylabel("Sepal width")
plt.show()
```

Output:



Exp.8 Create a self-organizing map neural network for learning a set of itnagos and verify the performance

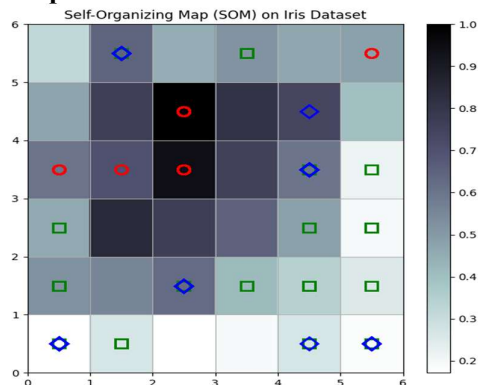
Algorithm:

1. Load dataset (Iris).
2. Normalize data.
3. Initialize and train SOM grid.
4. Map data to neurons.
5. Plot SOM clusters with labels.

Code.:

```
from sklearn.datasets import load_iris
from sklearn.preprocessing import StandardScaler
import matplotlib.pyplot as plt
from minisom import MiniSom
import numpy as np
X, y = load_iris(return_X_y=True)
X = StandardScaler().fit_transform(X)
som_grid_rows, som_grid_cols = 6, 6
som = MiniSom(som_grid_rows, som_grid_cols, X.shape[1], sigma=1.0, learning_rate=0.5,
              random_seed=42)
som.random_weights_init(X)
som.train_random(X, 100)
plt.figure(figsize=(7, 7))
plt.title("Self-Organizing Map (SOM) on Iris Dataset")
plt.pcolor(som.distance_map().T, cmap='bone_r')
plt.colorbar()
markers = ['o', 's', 'D']
colors = ['r', 'g', 'b']
for i, x_val in enumerate(X):
    w = som.winner(x_val)
    plt.plot(w[0] + 0.5, w[1] + 0.5,
             markers[y[i]],
             markeredgecolor=colors[y[i]],
             markerfacecolor='None',
             markersize=10,
             markeredgewidth=2)
plt.xticks(np.arange(som_grid_cols + 1))
plt.yticks(np.arange(som_grid_rows + 1))
plt.grid()
plt.show()
```

Output:



Exp.9 Create a Gaussian Mixture Model for Image Segmentation

Algorithm: GMM Image Segmentation

1. Read the input image.
2. Resize the image and reshape pixels into a 2D array.
3. Fit Gaussian Mixture Model (GMM) on pixel values.
4. Predict pixel clusters using GMM.
5. Replace pixels with corresponding cluster means.
6. Display original and segmented images side by side.

Code.:

```
import numpy as np
import matplotlib.pyplot as plt
from sklearn.mixture import GaussianMixture
from skimage.io import imread
from skimage.transform import resize

def segment_image_with_gmm(image_url, n_components=5):
    try:
        image = imread(image_url)
        if image.shape[2] == 4: # Handle RGBA images
            image = image[:, :, :3]
        image_resized = (resize(image, (200, 200), anti_aliasing=True) * 255).astype(np.uint8)
        pixels = image_resized.reshape(-1, 3)
        gmm = GaussianMixture(n_components=n_components, covariance_type='full'
                               random_state=42).fit(pixels)
        labels = gmm.predict(pixels)
        segmented_image = gmm.means_[labels].astype(np.uint8).reshape(image_resized.shape)
        fig, (ax1, ax2) = plt.subplots(1, 2, figsize=(12, 6))
        fig.suptitle(f'GMM Image Segmentation ({n_components} Clusters)', fontsize=16)
        ax1.imshow(image_resized)
        ax1.set_title('Original Image')
        ax1.axis('off')
        ax2.imshow(segmented_image)
        ax2.set_title('Segmented Image')
        ax2.axis('off')
        plt.tight_layout()
        plt.show()
    except Exception as e:
        print(f'An error occurred: {e}')

if __name__ == '__main__':
    sample_image_url = 'https://images.pexels.com/photos/162140/duck-bird-water-lake-162140.jpeg'
    num_segments = 6 # Try changing this number (e.g., 3, 8, 12)
    segment_image_with_gmm(sample_image_url, n_components=num_segments)
```

Output.:

Image Segmentation using GMM with 6 Clusters



Exp.10 Write a Genetic Algorithm program for finding parameters which maximizes the Y value of the equation given where the equation has 6 inputs (x1 to x6) and 6 weights (w1 to w6). Input values are (x1, x2, x3, x4, x5, x6) = (4, 2, 7, 5, 11, 1). Goal: Find the weights (w1–w6) that maximize Y.

Algorithm (5 Steps)

1. Initialize a random population of weights.
2. Evaluate fitness of each solution using $Y = w_1x_1 + w_2x_2 + \dots + w_6x_6$
 $Y = w_1x_1 + w_2x_2 + \dots + w_6x_6$
3. Select the best solutions (parents) based on fitness.
4. Generate new solutions using crossover and mutation.
5. Repeat for several generations and return the best weights with maximum Y.

Code.:

```
import numpy as np
import matplotlib.pyplot as plt
X = np.array([4, 2, 7, 5, 11, 1])
def fitness(weights):
    return np.dot(weights, X)
pop_size = 20
num_weights = len(X)
population = np.random.randint(-10, 10, (pop_size, num_weights))
generations = 50
best_scores = []
for gen in range(generations):
    scores = np.array([fitness(ind) for ind in population])
    best_scores.append(scores.max())
    parents = population[scores.argsort()[-(pop_size // 2):]]
    children = []
    for _ in range(pop_size - len(parents)):
        p1, p2 = parents[np.random.randint(len(parents), size=2)]
        cp = np.random.randint(1, num_weights - 1)
        child = np.concatenate((p1[:cp], p2[cp:]))
        children.append(child)
    children = np.array(children)
    mutation = np.random.randint(-2, 3, children.shape)
    children = children + mutation
    population = np.vstack((parents, children))
best_idx = np.argmax([fitness(ind) for ind in population])
best_weights = population[best_idx]
best_value = fitness(best_weights)
plt.plot(best_scores, marker='o')
plt.title("Genetic Algorithm Optimization Progress")
plt.xlabel("Generation")
plt.ylabel("Best Fitness Score (Y)")
plt.grid(True)
plt.show()
print("Best Weights:", best_weights)
print("Maximum Y Value:", best_value)
```

Output:

