REGRESSION

```
import numpy as np
import matplotlib.pyplot as plt
from sklearn.datasets import make regression
from sklearn.linear model import LinearRegression
from sklearn.model_selection import train_test_split
from sklearn.metrics import mean_squared_error
# Generate sample data
X, y = make regression(n samples=100, n features=1, noise=10, random state=42)
# Split the data 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)
# Initialize and fit the linear regression model
model = LinearRegression()
model.fit(X train, y train)
# Make predictions
y pred train = model.predict(X train)
y_pred_test = model.predict(X_test)
# Calculate mean squared error
mse train = mean squared error(y train, y pred train)
mse_test = mean_squared_error(y_test, y_pred_test)
# Plot the data and the regression line
plt.scatter(X train, y train, color='blue', label='Training data')
plt.scatter(X_test, y_test, color='red', label='Testing data')
plt.plot(X_train, y_pred_train, color='green', label='Regression line')
plt.title('Linear Regression')
plt.xlabel('X')
plt.ylabel('y')
plt.legend()
plt.show()
# Print mean squared error
print("Mean Squared Error (Training):", mse_train)
print("Mean Squared Error (Testing):", mse_test)
```

PCA

```
from sklearn.datasets import load_iris
from sklearn.decomposition import PCA
import matplotlib.pyplot as plt
# Load the Iris dataset
iris = load_iris()
X = iris.data
y = iris.target
# Perform PCA
pca = PCA(n_components=2) # Reduce to 2 principal components
X_pca = pca.fit_transform(X)
# Plot the PCA-transformed data
plt.figure(figsize=(8, 6))
for i in range(3):
  plt.scatter(X_pca[y == i, 0], X_pca[y == i, 1], label=iris.target_names[i])
plt.title('PCA of Iris Dataset')
plt.xlabel('Principal Component 1')
plt.ylabel('Principal Component 2')
plt.legend()
plt.show()
```

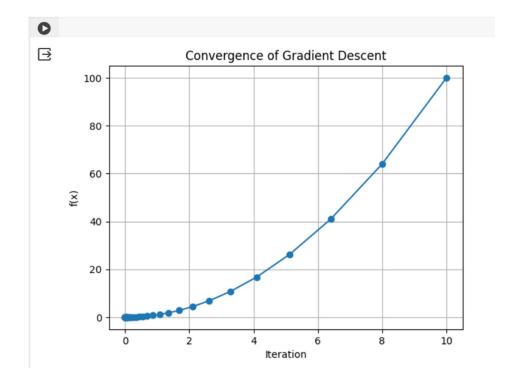
SVM -

```
import numpy as np
import matplotlib.pyplot as plt
from sklearn.datasets import make_blobs
from sklearn import sym
# Generate synthetic dataset with blobs
X, y = make_blobs(n_samples=100, centers=2, random_state=42)
# Print the dataset
print("Dataset:")
print("Number of samples: ", len(X))
# Train SVM model
clf = svm.SVC(kernel='linear')
clf.fit(X, y)
# Plot data points
plt.figure(figsize=(10, 6))
plt.scatter(X[:, 0], X[:, 1], c=y, cmap=plt.cm.coolwarm, s=30, edgecolors='k')
plt.xlabel('Feature 1')
plt.ylabel('Feature 2')
# Plot decision boundary
ax = plt.gca()
xlim = ax.get xlim()
ylim = ax.get_ylim()
# Create grid to evaluate model
xx = np.linspace(xlim[0], xlim[1], 30)
yy = np.linspace(ylim[0], ylim[1], 30)
YY, XX = np.meshgrid(yy, xx)
xy = np.vstack([XX.ravel(), YY.ravel()]).T # Mesh grid for the first two features
Z = clf.decision_function(xy).reshape(XX.shape)
# Plot decision boundary and margins
ax.contour(XX, YY, Z, colors='k', levels=[-1, 0, 1], alpha=0.5,
      linestyles=['--', '-', '--'])
ax.scatter(clf.support vectors [:, 0], clf.support vectors [:, 1], s=100,
      linewidth=1, facecolors='none', edgecolors='k')
plt.title('SVM Decision Boundary')
plt.show()
```

LAB 5: IMPLEMENTATION OF GRADIENT DESCENT ALGORITHM

Gradient descent: An optimization algorithm that iteratively adjusts parameters to minimize a given function by moving in the direction of the steepest descent. It's a core method in machine learning for training models and finding optimal solutions.

```
import numpy as np
import matplotlib.pyplot as plt
# Define the function
def f(x):
  return x**2
# Define the gradient of the function
def grad f(x):
  return 2*x
# Gradient Descent Algorithm
def gradient descent(learning rate, iterations):
  x = 10  # Initial guess
  history = [x]
  for i in range(iterations):
     gradient = grad f(x)
     x = x - learning rate * gradient
     history.append(x)
  return history
# Setting the hyperparameters
learning rate = 0.1
iterations = 50
# Running the gradient descent
history = gradient descent(learning rate, iterations)
# Visualizing the convergence
plt.plot(history, f(np.array(history)), '-o')
plt.xlabel('Iteration')
plt.ylabel('f(x)')
plt.title('Convergence of Gradient Descent')
plt.grid(True)
plt.show()
```



LAB 6: IMPLEMENTATION OF NAÏVE BAYESIAN CLASSIFICATION

Naive Bayesian classification is a probabilistic machine learning technique based on Bayes' theorem, assuming independence among features, commonly used for classification tasks where each feature contributes independently to the prediction of the class label.

Code-

```
from sklearn.datasets import load iris
from sklearn.model selection import train test split
from sklearn.naive bayes import GaussianNB
from sklearn.metrics import accuracy score
import numpy as np
iris = load iris()
X = iris.data
y = iris.target
# Splitting 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)
# Initializing Gaussian Naive Bayes classifier
clf = GaussianNB()
# Training the classifier
clf.fit(X_train, y_train)
# Calculating class priors
class counts = np.bincount(y train)
class priors = class counts / len(y train)
# Making predictions using Bayesian Decision Theory
posterior probs = clf.predict proba(X test)
y pred = np.argmax(posterior probs * class priors, axis=1)
print("Predicted Labels:", y pred)
# Calculate accuracy
accuracy = accuracy score(y test, y pred)
print("Accuracy:", accuracy)
```

Output-

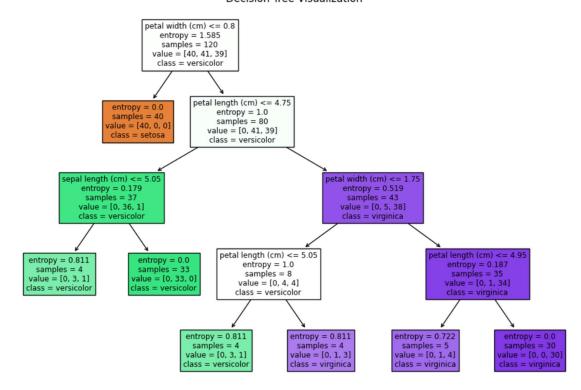
LAB 7: IMPLEMENTATION OF DECISION TREE CLASSIFIER

A decision tree classifier is a supervised machine learning algorithm that recursively partitions the feature space into regions, forming a tree-like structure, where each internal node represents a decision based on a feature, and each leaf node represents a class label.

```
from sklearn.datasets import load iris
from sklearn.model selection import train test split, GridSearchCV
from sklearn.tree import DecisionTreeClassifier, plot tree
import matplotlib.pyplot as plt
iris = load iris()
X = iris.data
y = iris.target
# Splitting 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)
# Initializing Decision Tree Classifier
dt classifier = DecisionTreeClassifier()
# Defining hyperparameters grid for tuning
param grid = {
  'criterion': ['gini', 'entropy'],
  'max depth': [3, 4, 5, 6, 7, 8, 9, 10],
  'min samples split': [2, 5, 10],
  'min samples leaf': [1, 2, 4]
# Performing grid search cross-validation to find the best hyperparameters
grid search = GridSearchCV(dt classifier, param grid, cv=5)
grid search.fit(X train, y train)
# Getting the best hyperparameters
best params = grid search.best params
print("Best Hyperparameters:", best params)
# Initializing Decision Tree Classifier with best hyperparameters
best dt classifier = DecisionTreeClassifier(**best params)
# Training the classifier with the best hyperparameters
best dt classifier.fit(X train, y train)
# Visualizing the constructed decision tree
```

Best Hyperparameters: {'criterion': 'entropy', 'max_depth': 5, 'min_samples_leaf': 4, 'min_samples_split': 2}

Decision Tree Visualization



LAB 8: IMPLEMENTATION OF BASIC ENSEMBLE METHOD AND BOOSTING METHOD

Ensemble methods combine multiple models to improve prediction accuracy, such as Random Forest, which aggregates the predictions of multiple decision trees. Boosting is a sequential ensemble technique where each subsequent model focuses on correcting the errors made by the previous ones, as seen in AdaBoost and XGBoost, leading to improved overall performance.

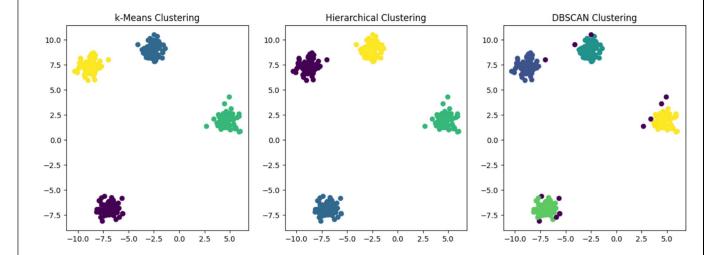
```
from sklearn.datasets import load iris
from sklearn.model selection import train test split
from sklearn.ensemble import RandomForestClassifier, AdaBoostClassifier
from xgboost import XGBClassifier
from sklearn.metrics import accuracy score
iris = load iris()
X = iris.data
y = iris.target
# Splitting 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)
# Initializing individual models
random forest = RandomForestClassifier(random state=42)
adaboost = AdaBoostClassifier(random state=42)
xgboost = XGBClassifier(random state=42)
# Training individual models
random forest.fit(X train, y train)
adaboost.fit(X train, y train)
xgboost.fit(X train, y train)
# Making predictions
rf predictions = random forest.predict(X test)
adaboost predictions = adaboost.predict(X test)
xgboost predictions = xgboost.predict(X test)
# Calculating accuracies
rf accuracy = accuracy score(y test, rf predictions)
adaboost accuracy = accuracy score(y test, adaboost predictions)
xgboost accuracy = accuracy score(y test, xgboost predictions)
print("Random Forest Accuracy:", rf accuracy)
print("AdaBoost Accuracy:", adaboost accuracy)
print("XGBoost Accuracy:", xgboost accuracy)
```

Output-
Random Forest Accuracy: 1.0 AdaBoost Accuracy: 1.0 XGBoost Accuracy: 1.0

LAB 9: IMPLEMENTATION OF CLUSTERING METHODS

Clustering algorithms such as KMeans, Hierarchical, and DBSCAN are fundamental techniques in unsupervised machine learning, used to partition datasets into distinct groups based on similarity measures, with KMeans clustering assigning data points to clusters by minimizing the within-cluster variance, Hierarchical clustering forming nested clusters by iteratively merging or splitting clusters, and DBSCAN identifying dense regions of data points, effectively handling noise and arbitrary cluster shapes.

```
import numpy as np
import matplotlib.pyplot as plt
from sklearn.datasets import make blobs
from sklearn.cluster import KMeans, AgglomerativeClustering, DBSCAN
# Generating synthetic data
X, = make blobs(n samples=300, centers=4, cluster std=0.60, random state=42)
# Implementing k-Means clustering
kmeans = KMeans(n clusters=4)
kmeans labels = kmeans.fit predict(X)
# Implementing Hierarchical clustering (Agglomerative)
hierarchical = AgglomerativeClustering(n clusters=4)
hierarchical labels = hierarchical.fit predict(X)
# Implementing DBSCAN clustering
dbscan = DBSCAN(eps=0.5, min samples=5)
dbscan\ labels = dbscan.fit\ predict(X)
# Visualizing clustering results
plt.figure(figsize=(15, 5))
plt.subplot(131)
plt.scatter(X[:, 0], X[:, 1], c=kmeans labels, cmap='viridis')
plt.title("k-Means Clustering")
plt.subplot(132)
plt.scatter(X[:, 0], X[:, 1], c=hierarchical labels, cmap='viridis')
plt.title("Hierarchical Clustering")
plt.subplot(133)
plt.scatter(X[:, 0], X[:, 1], c=dbscan labels, cmap='viridis')
plt.title("DBSCAN Clustering")
plt.show()
```



LAB 10: IMPLEMENTATION OF K-NEAREST NEIGHBORS (KNN)

K-Nearest Neighbors (KNN) is a simple yet powerful non-parametric classification algorithm that predicts the class of a data point based on the majority class of its nearest neighbors in the feature space, making it effective for both classification and regression tasks.

```
from sklearn.datasets import make classification
from sklearn.model selection import train test split
from sklearn.neighbors import KNeighborsClassifier
from sklearn.tree import DecisionTreeClassifier
from sklearn.metrics import accuracy score
# Generate synthetic data
X, y = make classification(n samples=1000, n features=20, n classes=2, random state=42)
# 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)
# Implement KNN classifier
def knn classifier(k):
  knn = KNeighborsClassifier(n neighbors=k)
  knn.fit(X train, y train)
  y pred = knn.predict(X test)
  return accuracy score(y test, y pred)
# Implement Decision Tree classifier
def decision tree classifier():
  dt = DecisionTreeClassifier(random state=42)
  dt.fit(X train, y train)
  y pred = dt.predict(X test)
  return accuracy score(y test, y pred)
# Evaluate KNN classifier with different values of K
k values = [1, 3, 5, 7, 9]
knn accuracies = []
for k in k values:
  accuracy = knn classifier(k)
  knn accuracies.append(accuracy)
# Evaluate Decision Tree classifier
dt accuracy = decision tree classifier()
# Print accuracies
print("Decision Tree Accuracy:", dt accuracy)
print("KNN Accuracies for Different Values of K:")
```

```
for k, accuracy in zip(k_values, knn_accuracies): print("K =", k, "Accuracy:", accuracy)
```

Decision Tree Accuracy: 0.875 KNN Accuracies for Different Values of K:

K = 1 Accuracy: 0.78
K = 3 Accuracy: 0.81
K = 5 Accuracy: 0.81
K = 7 Accuracy: 0.81
K = 9 Accuracy: 0.8

LAB 11: IMPLEMENTATION OF LOGISTIC REGRESSION

Logistic regression is a statistical model used for binary classification, which predicts the probability of an instance belonging to a particular class based on input features, utilizing the logistic (sigmoid) function to map input values to probabilities.

```
from sklearn.datasets import load iris
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
iris = load iris()
# Extracting features (X) and target variable (y)
X = iris.data
y = iris.target
# Splitting 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)
# Standardizing the features
scaler = StandardScaler()
X train = scaler.fit transform(X train)
X \text{ test} = \text{scaler.transform}(X \text{ test})
# Initializing and fitting the logistic regression model
log reg = LogisticRegression(max iter=1000)
log reg.fit(X train, y train)
# Printing some values
print("Coefficients:", log reg.coef )
print("Intercept:", log reg.intercept )
# Predicting on the test set
y_pred = log_reg.predict(X_test)
# Printing predicted values
print("Predicted values:", y_pred)
# Calculating accuracy
accuracy = accuracy_score(y_test, y_pred)
print("Accuracy:", accuracy)
```