

## Logistic

class LogisticRegression:

```
def __init__(self, learning_rate=0.01, num_itrtns=10000):
```

```
    self.lr = learning_rate
```

```
    self.num_itrtns = num_itrtns
```

```
    self.wghts = None
```

```
    self.bias = None
```

```
def sigmoid(self, z):
```

```
    return 1 / (1 + np.exp(-z))
```

```
def fit(self, X, y):
```

```
    num_smpls, num_fts = X.shape
```

```
    self.wghts = np.zeros(num_fts)
```

```
    self.bias = 0
```

```
    # Gradient descent
```

```
    for _ in range(self.num_itrtns):
```

```
        model = np.dot(X, self.wghts) + self.bias
```

```
        predictions = self.sigmoid(model)
```

```
        # Gradient calculation
```

```
        dw = (1 / num_smpls) * np.dot(X.T, (predictions - y))
```

```
        db = (1 / num_smpls) * np.sum(predictions - y)
```

```
        # Update weights and bias
```

```
        self.wghts -= self.lr * dw
```

```
        self.bias -= self.lr * db
```

```
def predict(self, X):
```

```
    model = np.dot(X, self.wghts) + self.bias
```

```

predictions = self.sigmoid(model)
predictions_cls = [1 if i > 0.5 else 0 for i in predictions]
return predictions_cls

```

## KNN

```

def euclidean_distance(p1, p2):
    return np.sqrt(np.sum((p1 - p2)**2))

def vote(neighbours):
    class_counter = Counter(neighbours)
    return class_counter.most_common(1)[0][0]

def knn(x_train, y_train, test_point, k):
    distances = []
    for i in range(len(x_train)):
        distance = euclidean_distance(test_point, x_train.iloc[i])
        distances.append((distance, y_train.iloc[i]))
    distances = sorted(distances)
    neighbours = np.asarray(distances[:k])
    label = vote(neighbours[:, 1])
    return label

```

## SVM

```

class SVM:
    def __init__(self, learning_rate=0.001, n_iters=1000):
        self.lr = learning_rate
        self.n_iters = n_iters
        self.w = None
        self.b = None

```

```

def fit(self, X, y):
    n_samples, n_features = X.shape
    self.w = np.zeros(n_features)
    self.b = 0

    for _ in range(self.n_iters):
        for idx, x_i in enumerate(X.values):
            condition = y.iloc[idx] * (np.dot(x_i, self.w) + self.b) >= 1
            if condition:
                self.w -= self.lr * (2 * self.w)
            else:
                self.w -= self.lr * (2 * self.w - np.dot(x_i, y.iloc[idx]))
                self.b -= self.lr * y.iloc[idx]

def predict(self, X):
    prediction = np.dot(X, self.w) + self.b
    return np.sign(prediction).astype(int)

```

Linear

```

def fit(X_train, y_train):
    m=0
    b=0
    num=0
    den=0
    for i in range(X_train.shape[0]):
        num = num + ((X_train[i] - X_train.mean()) * (y_train[i] - y_train.mean()))
        den = den + ((X_train[i] - X_train.mean()) * (X_train[i] - X_train.mean()))
    m = num/den
    b = y_train.mean() - (m * X_train.mean())
    print(m)
    print(b)

```

```
return m,b
```

```
def predict(m,b,X_test):
```

```
    print(X_test)
```

```
    y = m*X_test + b
```

```
    return y
```

## **Decision tree**

```
def entropy(y):
```

```
    class_counts = {}
```

```
    for label in y:
```

```
        if label in class_counts:
```

```
            class_counts[label] += 1
```

```
        else:
```

```
            class_counts[label] = 1
```

```
    entropy = 0
```

```
    total_samples = len(y)
```

```
    for count in class_counts.values():
```

```
        probability = count / total_samples
```

```
        entropy -= probability * math.log2(probability)
```

```
    return entropy
```

```
def information_gain(y, feature):
```

```
    total_entropy = entropy(y)
```

```
    unique_values = feature.unique()
```

```
    weighted_entropies = 0
```

```
    for value in unique_values:
```

```
        subset_y = y[feature == value]
```

```
        weighted_entropies += len(subset_y) / len(y) * entropy(subset_y)
```

```

information_gain= total_entropy - weighted_entropies

return information_gain

class Node:

    def __init__(self, feature=None, value=None, entropy=None, information_gain=None, left=None,
right=None):

        self.feature = feature

        self.value = value

        self.entropy = entropy

        self.information_gain = information_gain

        self.left = left

        self.right = right

    def build_decision_tree(X, y):

        if entropy(y) == 0:

            # If all instances have the same class, create a leaf node

            return Node(value=y.iloc[0])

        if X.empty:

            # If no features left, create a leaf node with the majority class

            return Node(value=y.value_counts().idxmax())

        # Find the best feature to split on

        best_feature = None

        max_info_gain = -1

        for feature_name in X.columns:

            current_info_gain = information_gain(y, X[feature_name])

            if current_info_gain > max_info_gain:

                max_info_gain = current_info_gain

                best_feature = feature_name

        # Create a node with the best feature

        node = Node(feature=best_feature, entropy=entropy(y), information_gain=max_info_gain, value={})

        # Recursively build the left and right subtrees

        unique_values = X[best_feature].unique()

        for value in unique_values:

```

```

subset_X = X[X[best_feature] == value].drop(columns=[best_feature])
subset_y = y[X[best_feature] == value]
child_node = build_decision_tree(subset_X, subset_y)
if node.value is None:
    node.value = {value: child_node}
else:
    node.value[value] = child_node
return node

def predict(node, instance):
    if node.feature is None:
        return node.value
    else:
        value = instance[node.feature]
        if value in node.value:
            return predict(node.value[value], instance)
        else:
            return node.value

Predictions = [predict(tree, instance) for _, instance in X_test.iterrows()]

```

## Naïve Bayes

```

from sklearn.preprocessing import LabelEncoder

label_encoder = LabelEncoder()

for column in df.select_dtypes(include=['object']).columns:
    df[column] = label_encoder.fit_transform(df[column])

def naive_bayes(X, y):
    class_probs = {}
    feature_probs = {}
    num_samples, num_features = X.shape
    unique_classes = np.unique(y)
    for c in unique_classes:

```

```

# Calculate class probabilities
class_probs[c] = np.sum(y == c) / num_samples

# Calculate feature probabilities for each class
features_given_class = X[y == c]
feature_probs[c] = np.sum(features_given_class, axis=0) / np.sum(y == c)

return class_probs, feature_probs

def predict(X, class_probs, feature_probs):
    predictions = []
    for sample in X:
        class_scores = {}
        for c, class_prob in class_probs.items():
            # Calculate the probability of the sample belonging to each class
            feature_probs_given_class = feature_probs[c]
            log_prob = np.sum(np.log(sample * feature_probs_given_class + (1 - sample) * (1 -
            feature_probs_given_class)))
            class_scores[c] = np.log(class_prob) + log_prob
        # Predict the class with the highest probability
        predicted_class = max(class_scores, key=class_scores.get)
        predictions.append(predicted_class)
    return predictions

```

## PCA

```

def calculate_mean(data):
    return np.mean(data, axis=0)

def calculate_covariance_matrix(data, mean):
    n_samples = data.shape[0]
    covariance_matrix = np.dot((data - mean).T, (data - mean)) / (n_samples - 1)
    return covariance_matrix

def perform_eigenvalue_decomposition(cov_matrix):
    eigenvalues, eigenvectors = np.linalg.eig(cov_matrix)

```

```

return eigenvalues, eigenvectors

def sort_eigenvectors(eigenvalues, eigenvectors):
    sorted_indices = np.argsort(eigenvalues)[::-1]
    sorted_eigenvalues = eigenvalues[sorted_indices]
    sorted_eigenvectors = eigenvectors[:, sorted_indices]
    return sorted_eigenvalues, sorted_eigenvectors

def select_principal_components(eigenvectors, num_components):
    selected_components = eigenvectors[:, :num_components]
    return selected_components

def transform_data(data, selected_components):
    transformed_data = np.dot(data, selected_components)
    return transformed_data

def pca(data, num_components):
    mean = calculate_mean(data)
    covariance_matrix = calculate_covariance_matrix(data, mean)
    eigenvalues, eigenvectors = perform_eigenvalue_decomposition(covariance_matrix)
    sorted_eigenvalues, sorted_eigenvectors = sort_eigenvectors(eigenvalues, eigenvectors)

    selected_components = select_principal_components(sorted_eigenvectors, num_components)

    transformed_data = transform_data(data, selected_components)
    return transformed_data

```

## **RANDOM FOREST**

```

class RandomForest:
    def __init__(self, n_estimators=100, max_depth=None):
        self.n_estimators = n_estimators
        self.max_depth = max_depth
        self.trees = []

    def fit(self, X, y):

```



```

for _ in range(self.n_estimators):
    tree = DecisionTreeClassifier(max_depth=self.max_depth)
    indices = np.random.choice(len(X), len(X), replace=False)
    tree.fit(X[indices], y[indices])
    self.trees.append(tree)

def predict(self, X):
    predictions = np.array([tree.predict(X) for tree in self.trees])
    return np.apply_along_axis(lambda x: np.bincount(x).argmax(), axis=0, arr=predictions)

```

## K-Means

```

import pandas as pd
import matplotlib.pyplot as plt
import numpy as np

def euclidean_distance(x1, x2):
    return np.sqrt(np.sum((x1 - x2) ** 2))

def initialize_centroids(X, k):
    centroids = X[np.random.choice(X.shape[0], k, replace=False)]
    return centroids

def assign_clusters(X, centroids):
    distances = np.array([np.linalg.norm(X - centroid, axis=1) for centroid in centroids])
    clusters = np.argmin(distances, axis=0)
    return clusters

def update_centroids(X, clusters, k):
    centroids = np.array([X[clusters == i].mean(axis=0) for i in range(k)])
    return centroids

```

```

def kmeans(X, k, max_iters=100):
    centroids = initialize_centroids(X, k)
    for _ in range(max_iters):
        prev_centroids = centroids.copy()
        clusters = assign_clusters(X, centroids)
        centroids = update_centroids(X, clusters, k)
        if np.all(prev_centroids == centroids):
            break
    return clusters, centroids

# Load the dataset
df = pd.read_csv("student_clustering.csv")
X = df[['cgpa', 'iq']].values

# Choose the optimal number of clusters based on the elbow method
optimal_k = 4

# Run K-means with the optimal K
clusters, centroids = kmeans(X, optimal_k)

# Visualize the clustering results
plt.figure(figsize=(10, 6))
for i in range(optimal_k):
    plt.scatter(X[clusters == i][:, 0], X[clusters == i][:, 1], label=f'Cluster {i}', alpha=0.6, s=100)
plt.scatter(centroids[:, 0], centroids[:, 1], s=300, c='red', label='Centroids')
plt.xlabel('CGPA')
plt.ylabel('IQ')
plt.title('K-means Clustering')
plt.legend()
plt.show()

```

**ANN\_And**

**Perceptron:**

**import numpy as np**

**weight = np.array([0, 0])**

**bias = 0**

**learning\_rate = 1**

**def activation(yin):**

**if yin > 0:**

**return 1**

**elif yin == 0:**

**return 0**

**else:**

**return -1**

**epochs = 100**

**input\_data = np.array([[1, 1], [1, -1], [-1, 1], [-1, -1]])**

**targets = np.array([1, -1, -1, -1])**

**for epoch in range(epochs):**

**for i in range(len(input\_data)):**

**yin = bias + np.dot(input\_data[i], weight)**

**y = activation(yin)**

**if y != targets[i]:**

**weight[0] += learning\_rate \* targets[i] \* input\_data[i][0]**

**weight[1] += learning\_rate \* targets[i] \* input\_data[i][1]**

**bias += learning\_rate \* targets[i]**

```
print("Final weight:", weight)
```

```
print("Final bias:", bias)
```

```
def predict(x1, x2):
```

```
    yin = bias + np.dot([x1, x2], weight)
```

```
    return activation(yin)
```

```
# Test the predict function
```

```
print("Input (1, 1): Predicted Output:", predict(1, 1))
```

```
print("Input (1, -1): Predicted Output:", predict(1, -1))
```

```
print("Input (-1, 1): Predicted Output:", predict(-1, 1))
```

```
print("Input (-1, -1): Predicted Output:", predict(-1, -1))
```