

1) scatter plot and hill climbing

```
x1 = np.random.rand(25)
y1 = np.random.rand(25)

x2 = np.random.rand(25)
y2 = np.random.rand(25)

plt.scatter(x1, y1, color='blue', label='Group 1')
plt.scatter(x2, y2, color='red', label='Group 2')

plt.title("Scatter plot with two colors")
plt.xlabel("X_axis")
plt.ylabel("Y_axis")
plt.legend()
plt.show()
```

Hill climbing:

```
def hill_climb(start, step_size, max_iterations):
    def fitness(x):
        return -x**2 + 4 # The function we want to maximize

    current = start
    for _ in range(max_iterations):
        neighbor = current + step_size
        if fitness(neighbor) > fitness(current):
            current = neighbor
        else:
            break # No better neighbor found
    return current, fitness(current)

solution, value = hill_climb(start=2, step_size=-0.1, max_iterations=100)
print("Best solution:", solution)
print("Best value:", value)
```

2) 3d plot and BFS

```
import numpy as np
import matplotlib.pyplot as plt
from mpl_toolkits.mplot3d import Axes3D # Needed for 3D plotting

# Creating meshgrid for X and Y
x = np.linspace(-5, 5, 50)
y = np.linspace(-5, 5, 50)
X, Y = np.meshgrid(x, y)

# Define Z as a function of X and Y
Z = np.sin(np.sqrt(X**2 + Y**2)) # Example surface

# Plotting
fig = plt.figure()
ax = fig.add_subplot(111, projection='3d')
ax.plot_surface(X, Y, Z, cmap='viridis') # You can change color map

ax.set_title("3D Surface Plot")
ax.set_xlabel("X axis")
ax.set_ylabel("Y axis")
ax.set_zlabel("Z axis")

plt.show()
```

BFS

```
from collections import deque

# BFS function
def bfs(graph, start):
    visited = set()
    queue = deque([start])

    while queue:
        node = queue.popleft()
        if node not in visited:
            print(node, end=' ')
            visited.add(node)
            queue.extend(neighbor for neighbor in graph[node] if neighbor not in visited)

# Example graph (adjacency list)
graph = {
    'A': ['B', 'C'],
```

```

'B': ['D', 'E'],
'C': ['F'],
'D': [],
'E': ['F'],
'F': []
}

```

```

# Call BFS
bfs(graph, 'A')

```

3) contour plot and A*

contour plot :

```

import numpy as np
import matplotlib.pyplot as plt

```

```

x = np.linspace(-5, 5, 100)
y = np.linspace(-5, 5, 100)
X, Y = np.meshgrid(x, y)
Z = np.sin(np.sqrt(X**2 + Y**2)) # function of X, Y

```

```

plt.contourf(X, Y, Z, cmap='plasma') # filled contour
plt.colorbar()
plt.title("Contour Plot")
plt.xlabel("X axis")
plt.ylabel("Y axis")
plt.show()

```

A*

```

import heapq

```

```

def heuristic(a, b):
    # Manhattan distance
    return abs(a[0] - b[0]) + abs(a[1] - b[1])

```

```

def astar(grid, start, goal):
    rows, cols = len(grid), len(grid[0])
    open_list = []
    heapq.heappush(open_list, (0 + heuristic(start, goal), 0, start, [start]))
    visited = set()

```

```

    while open_list:
        f, cost, current, path = heapq.heappop(open_list)

```

```

        if current == goal:
            return path

```

```

    if current in visited:
        continue
    visited.add(current)

    for dx, dy in [(-1,0),(1,0),(0,-1),(0,1)]: # 4 directions
        nx, ny = current[0] + dx, current[1] + dy
        if 0 <= nx < rows and 0 <= ny < cols and grid[nx][ny] == 0:
            next_node = (nx, ny)
            heapq.heappush(open_list, (cost + 1 + heuristic(next_node, goal), cost + 1, next_node, path
+ [next_node]))

    return None # No path found

# Example grid (0 = open, 1 = wall)
grid = [
    [0, 0, 0, 0],
    [1, 1, 0, 1],
    [0, 0, 0, 0],
    [0, 1, 1, 0],
    [0, 0, 0, 0]
]

start = (0, 0)
goal = (4, 3)

path = astar(grid, start, goal)
print("Path:", path)

```

4) Heat map and Min-max algorithm

```

import seaborn as sns
import numpy as np
import matplotlib.pyplot as plt

data = np.random.rand(10, 5) # 2D data: 10 rows × 5 columns
sns.heatmap(data, cmap='viridis')

plt.title("Heat Map")
plt.show()

```

min –max

Minimax Algorithm Implementation for a Game Tree

```

def minimax(depth, node_index, is_maximizing_player, scores, height):

```

```

# Base case: leaf node
if depth == height:
    return scores[node_index]

if is_maximizing_player:
    return max(
        minimax(depth + 1, node_index * 2, False, scores, height),
        minimax(depth + 1, node_index * 2 + 1, False, scores, height)
    )
else:
    return min(
        minimax(depth + 1, node_index * 2, True, scores, height),
        minimax(depth + 1, node_index * 2 + 1, True, scores, height)
    )

# Example leaf node scores
scores = [3, 5, 6, 9, 1, 2, 0, -1]

# Height of the tree = log2(len(scores)) = 3
tree_height = 3

# Call minimax starting from root (depth=0), root index=0, maximizing player's turn
best_value = minimax(0, 0, True, scores, tree_height)
print("The optimal value is:", best_value)

```

5) Box plot and Alpha beta pruning

```

import seaborn as sns
import numpy as np
import matplotlib.pyplot as plt

data = np.random.rand(100, 101) # 100 samples, 101 features
sns.boxplot(data=data)

plt.title("Box Plot")
plt.xlabel("Features")
plt.ylabel("Values")
plt.show()

```

Alpha beta

```

def alphabeta(depth, node_index, is_max, scores, alpha, beta, height):
    if depth == height:
        return scores[node_index]

```

```

if is_max:
    best = float('-inf')
    # Explore left and right child
    for i in range(2):
        val = alphabeta(depth + 1, node_index * 2 + i, False, scores, alpha, beta, height)
        best = max(best, val)
        alpha = max(alpha, best)
        if beta <= alpha:
            break #  $\beta$  cut-off
    return best
else:
    best = float('inf')
    # Explore left and right child
    for i in range(2):
        val = alphabeta(depth + 1, node_index * 2 + i, True, scores, alpha, beta, height)
        best = min(best, val)
        beta = min(beta, best)
        if beta <= alpha:
            break #  $\alpha$  cut-off
    return best

# Example leaf node values
scores = [3, 5, 6, 9, 1, 2, 0, -1]

# Tree height = 3 (log2 of 8 leaf nodes)
tree_height = 3

# Start at root: depth=0, node_index=0, maximizing=True
optimal = alphabeta(0, 0, True, scores, float('-inf'), float('inf'), tree_height)
print("The optimal value is:", optimal)

```

6) Navy base classifier

```

import pandas as pd
from sklearn.model_selection import train_test_split
from sklearn.preprocessing import LabelEncoder
from sklearn.naive_bayes import GaussianNB
from sklearn.metrics import accuracy_score

# Load Titanic dataset from CSV
# Replace with your actual path if needed
data = pd.read_csv('titanic.csv')

# Select features and target (you can customize as needed)
features = ['Pclass', 'Sex', 'Age', 'SibSp', 'Parch', 'Fare']

```

```

target = 'Survived'

# Handle missing values (simple strategy)
data = data[features + [target]].dropna()

# Encode categorical columns
le = LabelEncoder()
data['Sex'] = le.fit_transform(data['Sex']) # male=1, female=0

# Split into input (X) and output (y)
X = data[features]
y = data[target]

# Split into training and testing (70-30)
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.3, random_state=0)

# Train Naive Bayes classifier
model = GaussianNB()
model.fit(X_train, y_train)

# Predict and evaluate
y_pred = model.predict(X_test)
print("Accuracy of Naive Bayes classifier:", accuracy_score(y_test, y_pred))

```

7) KNN

```

import pandas as pd
from sklearn.model_selection import train_test_split
from sklearn.preprocessing import StandardScaler
from sklearn.neighbors import KNeighborsClassifier
from sklearn.metrics import accuracy_score

# Load the glass dataset from local CSV file
# Replace 'glass.csv' with the correct filename or path if needed
data = pd.read_csv('glass.csv')

# Separate features (X) and target (y)
X = data.iloc[:, :-1] # All columns except the last (features)

```

```

y = data.iloc[:, -1] # Last column (target class)

# Split dataset into 70% training and 30% testing
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.3, random_state=0)

# Standardize the feature values
scaler = StandardScaler()
X_train = scaler.fit_transform(X_train)
X_test = scaler.transform(X_test)

# KNN with Euclidean distance (default)
knn_euclidean = KNeighborsClassifier(n_neighbors=3, metric='euclidean')
knn_euclidean.fit(X_train, y_train)
y_pred_euclidean = knn_euclidean.predict(X_test)
print("Accuracy with Euclidean distance:", accuracy_score(y_test, y_pred_euclidean))

# KNN with Manhattan distance
knn_manhattan = KNeighborsClassifier(n_neighbors=3, metric='manhattan')
knn_manhattan.fit(X_train, y_train)
y_pred_manhattan = knn_manhattan.predict(X_test)
print("Accuracy with Manhattan distance:", accuracy_score(y_test, y_pred_manhattan))

```

8) K means

```

import matplotlib.pyplot as plt
from sklearn.datasets import load_iris
from sklearn.cluster import KMeans
from sklearn.preprocessing import StandardScaler

# Load and scale data
iris = load_iris()
X = StandardScaler().fit_transform(iris.data)

# Apply K-Means (3 clusters for 3 Iris species)
kmeans = KMeans(n_clusters=3, random_state=0)
labels = kmeans.fit_predict(X)

```



```
# Scatter plot using first two features
plt.scatter(X[:, 0], X[:, 1], c=labels, cmap='viridis')
plt.title("K-Means Clustering (Iris Dataset)")
plt.xlabel("Feature 1")
plt.ylabel("Feature 2")
plt.show()
```

9) single and complete linkage

```
import matplotlib.pyplot as plt
from sklearn.datasets import load_iris
from sklearn.preprocessing import StandardScaler
from scipy.cluster.hierarchy import dendrogram, linkage

# Load and standardize the Iris dataset
iris = load_iris()
X = StandardScaler().fit_transform(iris.data)

# Function to plot dendrogram
def plot_dendrogram(X, method):
    Z = linkage(X, method=method)
```

```

plt.figure(figsize=(6, 4))
plt.title(f'Dendrogram - {method} linkage')
dendrogram(Z)
plt.xlabel("Samples")
plt.ylabel("Distance")
plt.tight_layout()
plt.show()

# Plot dendrograms for both linkage methods
for method in ['single', 'complete']:
    plot_dendrogram(X, method)

```

11)PCA and LDA

```

from sklearn.datasets import load_iris
from sklearn.decomposition import PCA
from sklearn.discriminant_analysis import LinearDiscriminantAnalysis as LDA
import matplotlib.pyplot as plt

# Load the Iris dataset
iris = load_iris()
X = iris.data    # Features (4 columns)
y = iris.target  # Labels (0 = setosa, 1 = versicolor, 2 = virginica)

print("Shape of original data:", X.shape)
# -----

```

```

# PCA (Principal Component Analysis)
# -----
print("\n--- PCA ---")
pca = PCA(n_components=2)    # Reduce to 2 dimensions
X_pca = pca.fit_transform(X)

print("Shape after PCA:", X_pca.shape)
print("Explained Variance Ratio:", pca.explained_variance_ratio_)

# Plot PCA result
plt.figure(figsize=(6, 4))
plt.title("PCA - Iris Dataset")
plt.scatter(X_pca[:, 0], X_pca[:, 1], c=y, cmap='viridis')
plt.xlabel("PCA 1")
plt.ylabel("PCA 2")
plt.show()

# -----
# LDA (Linear Discriminant Analysis)
# -----
print("\n--- LDA ---")
lda = LDA(n_components=2)
X_lda = lda.fit_transform(X, y)

print("Shape after LDA:", X_lda.shape)

# Plot LDA result
plt.figure(figsize=(6, 4))
plt.title("LDA - Iris Dataset")
plt.scatter(X_lda[:, 0], X_lda[:, 1], c=y, cmap='plasma')
plt.xlabel("LDA 1")
plt.ylabel("LDA 2")
plt.show()

```

12) Write a Program to develop simple single layer perceptron to implement AND, OR Boolean functions.

```

# Step function
def step(x):
    return 1 if x >= 0 else 0

# Training function
def train(X, y):
    w1, w2 = 0, 0 # weights
    b = 0          # bias
    lr = 1         # learning rate

```

```

for _ in range(10): # 10 iterations
    for i in range(4): # for all 4 input examples
        x1, x2 = X[i]
        z = w1 * x1 + w2 * x2 + b
        y_pred = step(z)
        error = y[i] - y_pred
        # update weights and bias
        w1 += lr * error * x1
        w2 += lr * error * x2
        b += lr * error

    return w1, w2, b

# Prediction
def predict(X, w1, w2, b):
    results = []
    for x1, x2 in X:
        z = w1 * x1 + w2 * x2 + b
        results.append(step(z))
    return results

# Input and Output for AND
X = [[0,0], [0,1], [1,0], [1,1]]
y = [0, 0, 0, 1]

# Train and predict
w1, w2, b = train(X, y)
print("AND Gate Output:", predict(X, w1, w2, b))

```

output:
AND Predictions: [0, 0, 0, 1]
OR Predictions: [0, 1, 1, 1]