**Objective:** Solve the Tic-Tac-Toe problem using the Depth First Search technique.

```
# Set up the game board as a list
board = ["-", "-", "-",
# Define a function to print the game board
def print board():
  print(board[0] + " | " + board[1] + " | " + board[2])
  print(board[3] + " | " + board[4] + " | " + board[5])
  print(board[6] + " | " + board[7] + " | " + board[8])
# Define a function to handle a player's turn
def take turn(player):
  print(player + "'s turn.")
  position = input("Choose a position from 1-9: ")
  while position not in ["1", "2", "3", "4", "5", "6", "7", "8", "9"]:
     position = input("Invalid input. Choose a position from 1-9: ")
  position = int(position) - 1
  while board[position] != "-":
        position = int(input("Position already taken. Choose a different
position: ")) - 1
  board[position] = player
  print board()
```

```
# Define a function to check if the game is over
def check game over():
  # Check for a win
  if (board[0] == board[1] == board[2] != "-") or \
    (board[3] == board[4] == board[5] != "-") or \
    (board[6] == board[7] == board[8] != "-") or \
    (board[0] == board[3] == board[6] != "-") or \
    (board[1] == board[4] == board[7] != "-") or \
    (board[2] == board[5] == board[8] != "-") or \
    (board[0] == board[4] == board[8] != "-") or \
    (board[2] == board[4] == board[6] != "-"):
    return "win"
  # Check for a tie
  elif "-" not in board:
    return "tie"
  # Game is not over
  else:
    return "play"
# Define the main game loop
def play game():
  print board()
  current player = "X"
  game over = False
  while not game over:
     take turn(current player)
     game result = check game over()
     if game result == "win":
       print(current player + " wins!")
```

```
game_over = True
elif game_result == "tie":
    print("It's a tie!")
    game_over = True
else:
    # Switch to the other player
    current_player = "O" if current_player == "X" else "X"
```

# Start the game play\_game()

**Objective:** Show that the 8-puzzle states are divided into two disjoint sets, such that any state is reachable from any other state in the same set, while no state is reachable from any state in the other set.

```
# Python3 program to print the path from root
# node to destination node for N*N-1 puzzle
# algorithm using Branch and Bound
# The solution assumes that instance of
# puzzle is solvable
# Importing copy for deepcopy function
import copy
# Importing the heap functions from python
# library for Priority Queue
from heapq import heappush, heappop
# This variable can be changed to change
# the program from 8 puzzle(n=3) to 15
# puzzle(n=4) to 24 puzzle(n=5)...
n = 3
# bottom, left, top, right
row = [1, 0, -1, 0]
col = [0, -1, 0, 1]
```

```
# A class for Priority Queue
class priorityQueue:
  # Constructor to initialize a
  # Priority Queue
  def __init__(self):
     self.heap = []
  # Inserts a new key 'k'
  def push(self, k):
    heappush(self.heap, k)
  # Method to remove minimum element
  # from Priority Queue
  def pop(self):
    return heappop(self.heap)
  # Method to know if the Queue is empty
  def empty(self):
    if not self.heap:
       return True
    else:
       return False
# Node structure
class node:
  def __init__(self, parent, mat, empty_tile_pos,
          cost, level):
```

```
# Stores the parent node of the
     # current node helps in tracing
     # path when the answer is found
     self.parent = parent
     # Stores the matrix
     self.mat = mat
     # Stores the position at which the
     # empty space tile exists in the matrix
     self.empty tile pos = empty tile pos
     # Stores the number of misplaced tiles
     self.cost = cost
     # Stores the number of moves so far
     self.level = level
  # This method is defined so that the
  # priority queue is formed based on
  # the cost variable of the objects
  def lt (self, nxt):
     return self.cost < nxt.cost
# Function to calculate the number of
```

# Function to calculate the number of
# misplaced tiles ie. number of non-blank
# tiles not in their goal position
def calculateCost(mat, final) -> int:

```
count = 0
  for i in range(n):
    for j in range(n):
       if ((mat[i][j]) and
         (mat[i][j] != final[i][j])):
         count += 1
  return count
def newNode(mat, empty_tile_pos, new_empty_tile_pos,
       level, parent, final) -> node:
  # Copy data from parent matrix to current matrix
  new mat = copy.deepcopy(mat)
  # Move tile by 1 position
  x1 = \text{empty tile pos}[0]
  y1 = empty tile pos[1]
  x2 = new empty tile pos[0]
  y2 = new empty tile pos[1]
  new_mat[x1][y1], new_mat[x2][y2] = new_mat[x2][y2],
new mat[x1][y1]
  # Set number of misplaced tiles
  cost = calculateCost(new mat, final)
  new node = node(parent, new mat, new empty tile pos,
            cost, level)
  return new node
```

```
# Function to print the N x N matrix
def printMatrix(mat):
  for i in range(n):
     for j in range(n):
       print("%d " % (mat[i][j]), end = " ")
     print()
# Function to check if (x, y) is a valid
# matrix coordinate
def isSafe(x, y):
  return x \ge 0 and x < n and y \ge 0 and y < n
# Print path from root node to destination node
def printPath(root):
  if root == None:
     return
  printPath(root.parent)
  printMatrix(root.mat)
  print()
# Function to solve N*N - 1 puzzle algorithm
# using Branch and Bound. empty tile pos is
# the blank tile position in the initial state.
def solve(initial, empty_tile_pos, final):
```

```
# Create a priority queue to store live
# nodes of search tree
pq = priorityQueue()
# Create the root node
cost = calculateCost(initial, final)
root = node(None, initial,
       empty tile pos, cost, 0)
# Add root to list of live nodes
pq.push(root)
# Finds a live node with least cost,
# add its children to list of live
# nodes and finally deletes it from
# the list.
while not pq.empty():
  # Find a live node with least estimated
  # cost and delete it from the list of
  # live nodes
  minimum = pq.pop()
  # If minimum is the answer node
  if minimum.cost == 0:
     # Print the path from root to
     # destination;
     printPath(minimum)
     return
```

```
# Generate all possible children
     for i in range(4):
       new tile pos = [
         minimum.empty_tile_pos[0] + row[i],
         minimum.empty_tile_pos[1] + col[i], ]
       if isSafe(new tile pos[0], new tile pos[1]):
         # Create a child node
         child = newNode(minimum.mat,
                   minimum.empty tile pos,
                   new tile pos,
                   minimum.level + 1,
                   minimum, final,)
         # Add child to list of live nodes
         pq.push(child)
# Driver Code
# Initial configuration
# Value 0 is used for empty space
initial = [[1, 2, 3],
       [5, 6, 0],
       [7, 8, 4]]
# Solvable Final configuration
# Value 0 is used for empty space
final = [[1, 2, 3],
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```

# Blank tile coordinates in # initial configuration empty\_tile\_pos = [ 1, 2 ]

# Function call to solve the puzzle solve(initial, empty\_tile\_pos, final)

## **Result:**

1 2 3

5 6 0

7 8 4

1 2 3

5 0 6

7 8 4

1 2 3

5 8 6

7 0 4

1 2 3

5 8 6

0 7 4

**Objective:** To represent and evaluate different scenarios using predicate logic and knowledge rules.

```
from sympy import symbols
from sympy.logic.boolalg import Implies, And, Or, Not
from sympy.logic.inference import satisfiable
# Define predicates (functions returning boolean values)
Professor = symbols('Professor')
Teaches = symbols('Teaches')
Researches = symbols('Researches')
Student = symbols('Student')
Subject = symbols('Subject')
# Define individuals
Alice = symbols('Alice')
Bob = symbols('Bob')
Math = symbols('Math')
AI = symbols('AI')
# Define knowledge rules
knowledge base = [
  Implies(Professor, Teaches), # Professors teach
  Implies(Teaches, Subject), # If someone teaches, there is a subject
```

```
Implies(Researches, Professor), # If someone researches, they are a
professor
   Implies(Student, Not(Teaches)) # Students do not teach
]

# Scenario: Alice is a professor and researches AI
facts = [Professor.subs(Professor, Alice), Researches.subs(Researches,
Alice)]

# Check if Alice teaches
query = Teaches.subs(Teaches, Alice)
result = satisfiable(And(*knowledge_base, *facts, query))
print(f'Does Alice teach? {'Yes' if result else 'No'}")
```

Does Alice teach? Yes

**Objective:** To apply the Find-S and Candidate Elimination algorithms to a concept learning task and compare their inductive biases and outputs.

```
import numpy as np
# Sample training data: (Sky, AirTemp, Humidity, Wind, Water,
Forecast, EnjoySport)
data = np.array([
  ['Sunny', 'Warm', 'Normal', 'Strong', 'Warm', 'Same', 'Yes'],
  ['Sunny', 'Warm', 'High', 'Strong', 'Warm', 'Same', 'Yes'],
  ['Rainy', 'Cold', 'High', 'Strong', 'Warm', 'Change', 'No'],
  ['Sunny', 'Warm', 'High', 'Strong', 'Cool', 'Change', 'Yes']
1)
# Extract features (X) and labels (Y)
X = data[:, :-1] \# All columns except the last one
Y = data[:, -1] # Last column (labels)
# ----- FIND-S Algorithm -----
def find s algorithm(X, Y):
  specific hypothesis = X[0].copy() # Start with the first positive
example
  for i, row in enumerate(X[1:]):
     if Y[i+1] == 'Yes': # Consider only positive examples
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```

```
for j in range(len(specific hypothesis)):
         if row[j] != specific hypothesis[j]:
            specific hypothesis[j] = '?' # Generalize when necessary
  return specific hypothesis
# ----- CANDIDATE ELIMINATION Algorithm -----
def candidate elimination algorithm(X, Y):
  num attributes = X.shape[1]
  specific hypothesis = X[0].copy() # Start with first positive example
  general hypothesis = [['?'] * num attributes] # Most general
hypothesis
  for i, row in enumerate(X):
     if Y[i] == 'Yes': # Positive example \rightarrow specialize S
       for j in range(num attributes):
         if specific hypothesis[j] != row[j]:
            specific hypothesis[j] = '?'
       # Remove inconsistent general hypotheses
       general_hypothesis = [gh for gh in general hypothesis if
any(gh[k]!= specific hypothesis[k] and gh[k]!= '?' for k in
range(num_attributes))]
    elif Y[i] == 'No': # Negative example \rightarrow generalize G
       new general hypotheses = []
       for gh in general hypothesis:
         for j in range(num attributes):
            if gh[j] == '?': # Try specializing general hypothesis
              if specific_hypothesis[j] != '?':
                 new hypothesis = gh.copy()
                 new hypothesis[j] = specific hypothesis[j]
```

```
new_general_hypotheses.append(new_hypothesis)
general_hypothesis += new_general_hypotheses

# Remove overly specific hypotheses
general_hypothesis = [gh for gh in general_hypothesis if any(h != '?'
for h in gh)]
return specific_hypothesis, general_hypothesis

# Run both algorithms
find_s_result = find_s_algorithm(X, Y)
specific_h, general_h = candidate_elimination_algorithm(X, Y)

# Display results
print("\nFind-S Hypothesis:", find_s_result)
print("\nCandidate Elimination Algorithm:")
print("Most Specific Hypothesis:", specific_h)
print("Most General Hypotheses:", general_h)
```

```
Find-S Hypothesis: ['Sunny' 'Warm' '?' 'Strong' '?' '?']

Candidate Elimination Algorithm:

Most Specific Hypothesis: ['Sunny' 'Warm' '?' 'Strong' '?' '?']

Most General Hypotheses: []
```

**Objective:** To construct a decision tree using the ID3 algorithm on a simple classification dataset.

```
import numpy as np
import pandas as pd
from collections import Counter
import math
```

```
# Sample dataset: (Outlook, Temperature, Humidity, Wind, PlayTennis)
data = pd.DataFrame([
  ['Sunny', 'Hot', 'High', 'Weak', 'No'],
  ['Sunny', 'Hot', 'High', 'Strong', 'No'],
  ['Overcast', 'Hot', 'High', 'Weak', 'Yes'],
  ['Rain', 'Mild', 'High', 'Weak', 'Yes'],
  ['Rain', 'Cool', 'Normal', 'Weak', 'Yes'],
  ['Rain', 'Cool', 'Normal', 'Strong', 'No'],
  ['Overcast', 'Cool', 'Normal', 'Strong', 'Yes'],
  ['Sunny', 'Mild', 'High', 'Weak', 'No'],
  ['Sunny', 'Cool', 'Normal', 'Weak', 'Yes'],
  ['Rain', 'Mild', 'Normal', 'Weak', 'Yes'],
  ['Sunny', 'Mild', 'Normal', 'Strong', 'Yes'],
  ['Overcast', 'Mild', 'High', 'Strong', 'Yes'],
  ['Overcast', 'Hot', 'Normal', 'Weak', 'Yes'],
  ['Rain', 'Mild', 'High', 'Strong', 'No']
], columns=['Outlook', 'Temperature', 'Humidity', 'Wind', 'PlayTennis'])
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```

```
# Function to calculate entropy
def entropy(target col):
  counts = Counter(target col)
  total = len(target col)
  return -sum((count / total) * math.log2(count / total) for count in
counts.values())
# Function to calculate Information Gain
def info gain(data, split attribute, target name):
  total entropy = entropy(data[target name]) # Parent entropy
  values = data[split attribute].unique() # Unique values of the
attribute
  weighted entropy = sum(
     (len(subset := data[data[split attribute] == val]) / len(data)) *
entropy(subset[target name])
     for val in values
  return total entropy - weighted entropy # Information gain
# Function to build the decision tree
def id3(data, features, target):
  # Base cases
  if len(set(data[target])) == 1: # Pure class
     return data[target].iloc[0]
  if not features: # No more features to split on
     return data[target].mode()[0]
  # Select the best feature
  best feature = max(features, key=lambda f: info gain(data, f, target))
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```

```
# Create a subtree
  tree = {best feature: {}}
  features = [f for f in features if f != best feature] # Remove used
feature
  for value in data[best_feature].unique(): # For each unique value of
the best feature
     subset = data[data[best feature] == value]
     tree[best feature][value] = id3(subset, features, target) # Recursive
call
  return tree
# Build decision tree
features = list(data.columns[:-1])
decision tree = id3(data, features, 'PlayTennis')
# Function to classify a new example
def classify(tree, sample):
  if not isinstance(tree, dict):
     return tree # Leaf node reached
  root = next(iter(tree))
  return classify(tree[root][sample[root]], sample) if sample[root] in
tree[root] else 'Unknown'
# Display the decision tree
import pprint
print("\nDecision Tree:")
pprint.pprint(decision tree)
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```

```
# Test new example
sample = {'Outlook': 'Sunny', 'Temperature': 'Cool', 'Humidity': 'High',
'Wind': 'Strong'}
print("\nNew Sample Classification:", classify(decision_tree, sample))
```

**Objective:** To assess how the ID3 algorithm performs on datasets with varying characteristics and complexity, examining overfitting, underfitting, and decision tree depth.

```
import numpy as np
import pandas as pd
import math
from collections import Counter
from sklearn.model selection import train test split
from sklearn.metrics import accuracy_score
import pprint
# Function to compute entropy
def entropy(target col):
  counts = Counter(target col)
  total = len(target col)
  return -sum((count / total) * math.log2(count / total) for count in
counts.values())
# Function to compute information gain
def info gain(data, split attribute, target name):
  total entropy = entropy(data[target name]) # Parent entropy
  values = data[split attribute].unique() # Unique values of the
attribute
  weighted entropy = sum(
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```

```
(len(subset := data[data[split attribute] == val]) / len(data)) *
entropy(subset[target name])
     for val in values
  return total entropy - weighted entropy # Information gain
# Function to build the decision tree
def id3(data, features, target, depth=0, max_depth=None):
  # Base cases
  if len(set(data[target])) == 1: # Pure class
     return data[target].iloc[0]
  if not features or (max depth is not None and depth >= max depth):
# No more features to split or max depth reached
     return data[target].mode()[0]
  # Select best feature
  best feature = max(features, key=lambda f: info gain(data, f, target))
  # Create subtree
  tree = {best feature: {}}
  features = [f for f in features if f!= best_feature] # Remove used
feature
  for value in data[best_feature].unique(): # Split on each unique value
     subset = data[data[best feature] == value]
     tree[best feature][value] = id3(subset, features, target, depth + 1,
max depth)
  return tree
```

```
# Function to classify new examples
def classify(tree, sample):
  if not isinstance(tree, dict):
     return tree # Leaf node reached
  root = next(iter(tree))
  return classify(tree[root][sample[root]], sample) if sample[root] in
tree[root] else 'Unknown'
# Function to predict multiple samples
def predict(tree, X):
  return [classify(tree, x) for _, x in X.iterrows()]
# Function to assess performance on different datasets
def evaluate_id3(data, max depth=None):
  X = data.iloc[:, :-1]
  y = data.iloc[:, -1]
  # Split into training and test sets
  X train, X test, y train, y test = train test split(X, y, test size=0.3,
random state=42)
  # Train the decision tree
  features = list(X.columns)
  tree = id3(pd.concat([X train, y train], axis=1), features,
y train.name, max depth=max depth)
  # Make predictions
  y train pred = predict(tree, X train)
  y test pred = predict(tree, X_test)
```

```
# Compute accuracy
  train acc = accuracy score(y train, y train pred)
  test acc = accuracy score(y test, y test pred)
  # Print results
  print(f"\nDecision Tree (Max Depth = {max depth if max depth else
'Full'}):")
  pprint.pprint(tree)
  print(f"\nTraining Accuracy: {train acc:.4f}")
  print(f"Test Accuracy: {test acc:.4f}")
  print("-" * 40)
# Sample dataset: (Simple, Medium, and Complex)
data simple = pd.DataFrame([
  ['Sunny', 'Hot', 'High', 'Weak', 'No'],
  ['Sunny', 'Hot', 'High', 'Strong', 'No'],
  ['Overcast', 'Hot', 'High', 'Weak', 'Yes'],
  ['Rain', 'Mild', 'High', 'Weak', 'Yes'],
  ['Rain', 'Cool', 'Normal', 'Weak', 'Yes'],
  ['Rain', 'Cool', 'Normal', 'Strong', 'No'],
  ['Overcast', 'Cool', 'Normal', 'Strong', 'Yes'],
  ['Sunny', 'Mild', 'High', 'Weak', 'No'],
  ['Sunny', 'Cool', 'Normal', 'Weak', 'Yes'],
  ['Rain', 'Mild', 'Normal', 'Weak', 'Yes'],
  ['Sunny', 'Mild', 'Normal', 'Strong', 'Yes'],
  ['Overcast', 'Mild', 'High', 'Strong', 'Yes'],
  ['Overcast', 'Hot', 'Normal', 'Weak', 'Yes'],
  ['Rain', 'Mild', 'High', 'Strong', 'No']
], columns=['Outlook', 'Temperature', 'Humidity', 'Wind', 'PlayTennis'])
```

# Run evaluations evaluate\_id3(data\_simple, max\_depth=1) # Underfitting scenario evaluate\_id3(data\_simple, max\_depth=3) # Balanced tree evaluate\_id3(data\_simple, max\_depth=None) # Full tree (potential overfitting)

```
Decision Tree (Max Depth = 1):
{'Humidity': {'High': 'No', 'Normal': 'Yes'}}
Training Accuracy: 0.7778
Test Accuracy: 0.6000
Decision Tree (Max Depth = 3):
{'Humidity': {'High': {'Outlook': {'Overcast': 'Yes',
                                    'Rain': {'Wind': {'Strong': 'No',
                                                     'Weak': 'Yes'}},
                                    'Sunny': 'No'}},
              'Normal': 'Yes'}}
Training Accuracy: 1.0000
Test Accuracy: 0.8000
Decision Tree (Max Depth = Full):
{'Humidity': {'High': {'Outlook': {'Overcast': 'Yes',
                                   'Rain': {'Wind': {'Strong': 'No',
                                                     'Weak': 'Yes'}},
                                    'Sunny': 'No'}},
              'Normal': 'Yes'}}
Training Accuracy: 1.0000
Test Accuracy: 0.8000
```

**Objective:** To examine different types of machine learning approaches (Supervised, Unsupervised, Semi-supervised, and Reinforcement Learning) by setting up a basic classification problem and exploring how each type applies differently.

#### **Source Code:**

```
import numpy as np
import pandas as pd
import matplotlib.pyplot as plt
from sklearn.model_selection import train_test_split
from sklearn.tree import DecisionTreeClassifier
from sklearn.svm import SVC
from sklearn.metrics import accuracy_score
from sklearn.cluster import KMeans
from sklearn.semi_supervised import LabelPropagation
import random
```

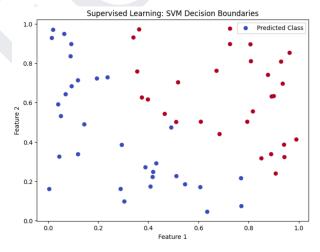
```
# Generate a synthetic dataset (features: X1, X2, class: Y) np.random.seed(42)  X = \text{np.random.rand}(200, 2) \# 200 \text{ samples}, 2 \text{ features}   Y = \text{np.where}(X[:, 0] + X[:, 1] > 1, 1, 0) \# \text{ Simple linear classification}  boundary
```

# Split into training and test sets

```
X train, X test, Y train, Y test = train test split(X, Y, test size=0.3,
random state=42)
# ------ 1. SUPERVISED LEARNING ------
print("\n--- Supervised Learning: Decision Tree ---")
dt model = DecisionTreeClassifier()
dt model.fit(X train, Y train)
Y pred = dt model.predict(X test)
print("Decision Tree Accuracy:", accuracy score(Y test, Y pred))
print("\n--- Supervised Learning: Support Vector Machine (SVM) ---")
svm model = SVC()
svm model.fit(X train, Y train)
Y pred svm = svm model.predict(X test)
print("SVM Accuracy:", accuracy_score(Y_test, Y_pred_svm))
# ------ 2. UNSUPERVISED LEARNING ------
print("\n--- Unsupervised Learning: K-Means Clustering ---")
kmeans = KMeans(n clusters=2, random state=42)
clusters = kmeans.fit predict(X test) # No true labels used
print("Cluster Assignments:", clusters[:10]) # Show first 10 cluster
labels
# ----- 3. SEMI-SUPERVISED LEARNING
print("\n--- Semi-Supervised Learning: Label Propagation ---")
labels = np.copy(Y train)
random unlabeled = np.random.choice(len(Y train), size=int(0.8 *
len(Y_train)), replace=False)
labels[random unlabeled] = -1 # Hide 80% of labels
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```

```
lp model = LabelPropagation()
lp model.fit(X train, labels)
Y pred lp = lp model.predict(X test)
print("Semi-Supervised Accuracy:", accuracy score(Y test,
Y pred lp))
# ----- 4. REINFORCEMENT LEARNING
print("\n--- Reinforcement Learning: Q-Learning for Classification ---")
Q table = np.zeros((2, 2)) # 2 states (classes), 2 actions (predict 0 or 1)
alpha = 0.1 \# Learning rate
gamma = 0.9 # Discount factor
# Q-learning process
for _ in range(1000):
  state = random.choice([0, 1]) # Randomly choose a class state
  action = np.argmax(Q table[state]) if np.random.rand() > 0.2 else
np.random.choice([0, 1]) # \varepsilon-greedy policy
  reward = 1 if action == state else -1 # Reward is 1 if prediction is
correct
  Q table[state, action] = (1 - alpha) * Q table[state, action] + alpha *
(reward + gamma * np.max(Q_table[action]))
# Testing Q-learning-based classification
Y pred q = [np.argmax(Q table[y]) for y in Y test] # Predict based on
learned Q-table
print("Q-Learning Accuracy:", accuracy score(Y test, Y pred q))
# ------ Plot the dataset -----
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```

```
plt.figure(figsize=(8, 6))
plt.scatter(X_test[:, 0], X_test[:, 1], c=Y_pred_svm, cmap='coolwarm',
label='Predicted Class')
plt.title('Supervised Learning: SVM Decision Boundaries')
plt.xlabel('Feature 1')
plt.ylabel('Feature 2')
plt.legend()
plt.show()
```



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To understand how Find-S and Candidate **Objective:** Elimination algorithms search through the hypothesis space in concept learning tasks, and to observe the role of inductive bias in shaping the learned concept.

```
import numpy as np
import pandas as pd
# Sample dataset for concept learning
data = np.array([
  ['Sunny', 'Warm', 'Normal', 'Strong', 'Warm', 'Same', 'Yes'],
  ['Sunny', 'Warm', 'High', 'Strong', 'Warm', 'Same', 'Yes'],
  ['Rainy', 'Cold', 'High', 'Strong', 'Warm', 'Change', 'No'],
  ['Sunny', 'Warm', 'High', 'Strong', 'Cool', 'Change', 'Yes']
])
# Convert to Pandas DataFrame
columns = ['Sky', 'Temperature', 'Humidity', 'Wind', 'Water', 'Forecast',
'EnjoySport']
df = pd.DataFrame(data, columns=columns)
# Separate features and target
X = df.iloc[:, :-1].values
Y = df.iloc[:, -1].values
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```

```
# ------ FIND-S ALGORITHM ------
def find s(X, Y):
  hypothesis = np.array(X[0]) # Start with the first positive example
  for i, sample in enumerate(X):
    if Y[i] == 'Yes': # Only consider positive examples
       for j in range(len(sample)):
         if sample[i] != hypothesis[i]:
            hypothesis[i] = '?' # Generalize differing attributes
  return hypothesis
# ----- CANDIDATE ELIMINATION ALGORITHM
def candidate elimination(X, Y):
  num attributes = X.shape[1]
  # Initialize the most specific and most general hypotheses
  S = ['\emptyset'] * num attributes # Most specific hypothesis
  G = ['?'] * num attributes # Most general hypothesis
  version_space = [] # Store hypotheses during training
  for i, sample in enumerate(X):
    if Y[i] == 'Yes': # Positive example
       for j in range(num attributes):
         if S[i] == '\emptyset':
            S[i] = \text{sample}[i] \# \text{Assign first positive sample}
         elif S[i] != sample[i]:
            S[j] = '?' \# Generalize
       version_space.append(tuple(S)) # Track the version space
```

```
elif Y[i] == 'No': # Negative example
       for j in range(num attributes):
         if S[j] != '?' and S[j] != sample[j]:
            G[j] = S[j] # Specialize
       version space.append(tuple(G))
  return S, G, version space
# Run Find-S Algorithm
find s hypothesis = find s(X, Y)
print("\nFinal Hypothesis (Find-S):", find_s_hypothesis)
# Run Candidate Elimination Algorithm
S_{final}, G_{final}, version_space = candidate elimination(X, Y)
print("\nFinal Specific Hypothesis (S):", S final)
print("Final General Hypothesis (G):", G_final)
print("Version Space:", version space)
Result:
```

```
Final Hypothesis (Find-S): ['Sunny' 'Warm' '?' 'Strong' '?' '?']
```

```
Final Specific Hypothesis (S): ['Sunny' 'Warm' '?' 'Strong' '?' '?']
Final General Hypothesis (G): ['?' '?' '?' '?' '?']
Version Space: [('Sunny', 'Warm', '?', 'Strong', '?', '?'), ('?', '?', '?',
'?', '?', '?')]
```

**Objective:** To go through all stages of a real-life machine learning project, from data collection to model fine-tuning, using a regression dataset like the "California Housing Prices."

#### **Source Code:**

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, GridSearchCV
from sklearn.preprocessing import StandardScaler
from sklearn.ensemble import RandomForestRegressor
from sklearn.metrics import mean\_absolute\_error, mean\_squared\_error,
r2\_score
from sklearn.linear\_model import LinearRegression
from sklearn.pipeline import Pipeline
from sklearn.impute import SimpleImputer
from sklearn.compose import ColumnTransformer
from sklearn.preprocessing import OneHotEncoder

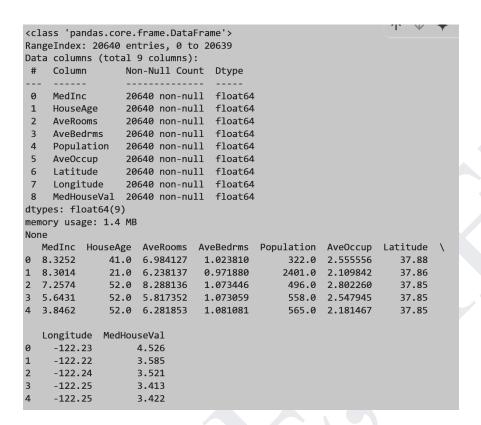
# Load California Housing dataset from sklearn.datasets import fetch\_california\_housing california = fetch\_california\_housing(as\_frame=True) df = california.frame

```
# Display basic dataset information
print(df.info())
print(df.head())
# Check for missing values
print("\nMissing Values:\n", df.isnull().sum())
# Summary statistics
print("\nDataset Statistics:\n", df.describe())
# Visualizing correlations
plt.figure(figsize=(10, 6))
sns.heatmap(df.corr(), annot=True, cmap="coolwarm", linewidths=0.5)
plt.title("Feature Correlation Heatmap")
plt.show()
# Plot target variable distribution
sns.histplot(df['MedHouseVal'], bins=30, kde=True)
plt.title("Median House Value Distribution")
plt.show()
# Define features and target
X = df.drop(columns=["MedHouseVal"])
y = df["MedHouseVal"]
# Identify numerical and categorical columns
num features = X.select dtypes(include=["float64",
"int64"]).columns.tolist()
cat_features = X.select_dtypes(include=["object"]).columns.tolist() # In
this dataset, no categorical features exist.
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```

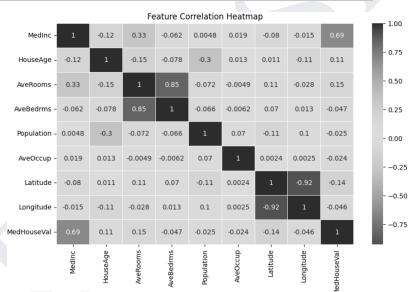
```
# Data Preprocessing Pipeline
num transformer = Pipeline(steps=[
  ("imputer", SimpleImputer(strategy="median")),
  ("scaler", StandardScaler())
1)
cat transformer = Pipeline(steps=[
  ("encoder", OneHotEncoder(handle unknown="ignore"))
])
preprocessor = ColumnTransformer(transformers=[
  ("num", num transformer, num features),
  ("cat", cat transformer, cat features) # No categorical features in this
dataset, but kept for extensibility
])
# Split data into training and test sets
X train, X test, y train, y test = train test split(X, y, test size=0.2,
random state=42)
# Apply transformations
X_train = preprocessor.fit_transform(X_train)
X_{test} = preprocessor.transform(X_{test})
# Train a Linear Regression Model
lr model = LinearRegression()
lr_model.fit(X_train, y_train)
# Predictions
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```

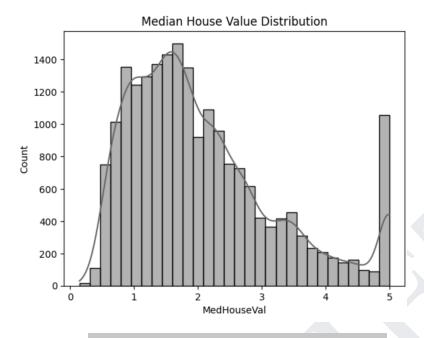
```
y pred lr = lr model.predict(X test)
# Evaluate Performance
print("\n--- Linear Regression Performance ---")
print(f"R<sup>2</sup> Score: {r2 score(y test, y pred lr):.4f}")
print(f"MAE: {mean absolute error(y test, y pred lr):.4f}")
print(f"RMSE: {np.sqrt(mean squared error(y test, y pred lr)):.4f}")
# Train a Random Forest Model
rf_model = RandomForestRegressor(n estimators=100,
random state=42)
rf model.fit(X train, y train)
# Predictions
y_pred_rf = rf_model.predict(X_test)
# Evaluate Performance
print("\n--- Random Forest Performance ---")
print(f"R<sup>2</sup> Score: {r2 score(y test, y pred rf):.4f}")
print(f"MAE: {mean absolute error(y test, y pred rf):.4f}")
print(f"RMSE: {np.sqrt(mean squared error(y test, y pred rf)):.4f}")
# Define hyperparameter grid
param grid = {
  "n estimators": [50, 100, 200],
  "max depth": [10, 20, None],
  "min_samples_split": [2, 5, 10]
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```

```
# Grid Search with Cross-Validation
grid search =
GridSearchCV(RandomForestRegressor(random state=42), param grid,
cv=3, scoring="r2", n jobs=-1)
grid_search.fit(X_train, y_train)
# Best model evaluation
best rf = grid search.best estimator
y_pred_best = best_rf.predict(X_test)
print("\n--- Best Random Forest Performance ---")
print(f"Best Parameters: {grid search.best params }")
print(f"R² Score: {r2_score(y_test, y_pred_best):.4f}")
print(f"MAE: {mean absolute error(y test, y pred best):.4f}")
print(f"RMSE: {np.sqrt(mean_ squared error(y test,
y_pred_best)):.4f}")
# Extract feature importances
importances = best rf.feature importances
feature names = num features # No categorical features in this dataset
# Plot Feature Importance
plt.figure(figsize=(10, 5))
sns.barplot(x=importances, y=feature names, palette="viridis")
plt.title("Feature Importance in Random Forest Model")
plt.xlabel("Importance Score")
plt.ylabel("Feature")
plt.show()
```



Missin	ng Values:					
MedIn	ic 0					
HouseA	ige 0					
AveRoo	oms 0					
AveBed	Irms 0					
Popula	ition 0					
Ave0cc	up 0					
Latitu	ide 0					
Longit	ude 0					
MedHou	ıseVal 0					
dtype:	int64					
Datase	et Statistics:					
	MedInc	HouseAge	AveRooms	AveBedrms	Population	\
count	20640.000000	20640.000000	20640.000000	20640.000000	20640.000000	
mean	3.870671	28.639486	5.429000	1.096675	1425.476744	
std	1.899822	12.585558	2.474173	0.473911	1132.462122	
min	0.499900	1.000000	0.846154	0.333333	3.000000	
25%	2.563400	18.000000	4.440716	1.006079	787.000000	
50%	3.534800	29.000000	5.229129	1.048780	1166.000000	
75%	4.743250	37.000000	6.052381	1.099526	1725.000000	
max	15.000100	52.000000	141.909091	34.066667	35682.000000	
	Ave0ccup	Latitude	Longitude	MedHouseVal		
count	20640.000000	20640.000000	20640.000000	20640.000000		
mean	3.070655	35.631861	-119.569704	2.068558		
std	10.386050	2.135952	2.003532	1.153956		
min	0.692308	32.540000	-124.350000	0.149990		
25%	2.429741	33.930000	-121.800000	1.196000		
50%	2.818116	34.260000	-118.490000	1.797000		
75%	3.282261	37.710000	-118.010000	2.647250		
	3.202201	37.710000	-119.010000	2.04/200		





--- Linear Regression Performance ---

R<sup>2</sup> Score: 0.5758 MAE: 0.5332 RMSE: 0.7456

--- Random Forest Performance ---

R<sup>2</sup> Score: 0.8053 MAE: 0.3274 RMSE: 0.5051

# **Experiment-10**

**Objective:** To perform binary and multiclass classification on the MNIST dataset, analyze performance metrics, and perform error analysis.

## **Source Code:**

import numpy as np Prof.Ranjitha R, ECE, SIET

```
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, classification report,
confusion matrix
from tensorflow.keras.datasets import mnist
# Load MNIST dataset
(X_train, y_train), (X_test, y_test) = mnist.load_data()
# Normalize images (scale pixel values between 0 and 1)
X train = X train / 255.0
X \text{ test} = X \text{ test} / 255.0
# Flatten images from 28x28 to 1D (784 features)
X train flat = X train.reshape(X train.shape[0], -1)
X test flat = X test.reshape(X test.shape[0], -1)
print("Dataset Shape:", X train.shape, X test.shape)
# Convert to a binary classification problem (0 vs. not 0)
y train binary = (y train == 0).astype(int)
y test binary = (y \text{ test} == 0).astype(int)
# Train a logistic regression model
binary clf = LogisticRegression(max iter=1000)
binary clf.fit(X train flat, y train binary)
```

```
# Predictions
y pred binary = binary clf.predict(X test flat)
# Evaluate Performance
print("\n--- Binary Classification (0 vs. Others) ---")
print("Accuracy:", accuracy score(y test binary, y pred binary))
print(classification report(y test binary, y pred binary))
# Train a logistic regression model for multiclass classification
multi clf = LogisticRegression(max iter=1000,
multi class="multinomial", solver="lbfgs")
multi clf.fit(X train flat, y train)
# Predictions
y pred multi = multi clf.predict(X test flat)
# Evaluate Performance
print("\n--- Multiclass Classification (All Digits) ---")
print("Accuracy:", accuracy score(y test, y pred multi))
print(classification_report(y_test, y_pred_multi))
# Compute Confusion Matrix
conf matrix = confusion matrix(y test, y pred multi)
# Plot Confusion Matrix
plt.figure(figsize=(10, 7))
sns.heatmap(conf matrix, annot=True, fmt="d", cmap="coolwarm",
xticklabels=range(10), yticklabels=range(10))
plt.xlabel("Predicted Label")
plt.ylabel("True Label")
plt.title("Confusion Matrix (Multiclass)")
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```

```
# Identify misclassified examples
misclassified_idxs = np.where(y_test != y_pred_multi)[0]

# Display some misclassified images
plt.figure(figsize=(12, 6))
for i, idx in enumerate(misclassified_idxs[:10]):
    plt.subplot(2, 5, i + 1)
    plt.imshow(X_test[idx], cmap="gray")
    plt.title(f"True: {y_test[idx]}, Pred: {y_pred_multi[idx]}")
    plt.axis("off")
plt.tight_layout()
plt.show()
```

```
Downloading data from <a href="https://storage.googleapis.com/tensorflow/tf-keras-datasets/mnist.npz">https://storage.googleapis.com/tensorflow/tf-keras-datasets/mnist.npz</a>
11490434/11490434
                                              - 0s Ous/step
Dataset Shape: (60000, 28, 28) (10000, 28, 28)
--- Binary Classification (0 vs. Others) ---
Accuracy: 0.9917
                 precision recall f1-score
                                                        support
                                   0.99
                       1.00
                                               1.00
                                                            9020
                       0.95
                                   0.96
                                               0.96
                                                             980
                                                           10000
                                               0.99
    accuracy
                       0.97
                                   0.98
                                               0.98
                                                           10000
   macro avg
weighted avg
                       0.99
                                   0.99
                                               0.99
                                                           10000
```

Multiclass Classification (All Digits) Accuracy: 0.9258									
•	precision	recall	f1-score	support					
0	0.95	0.98	0.96	980					
1	0.96	0.98	0.97	1135					
2	0.93	0.90	0.91	1032					
3	0.90	0.91	0.91	1010					
4	0.94	0.94	0.94	982					
5	0.90	0.87	0.88	892					
6	0.94	0.95	0.95	958					
7	0.93	0.92	0.93	1028					
8	0.88	0.88	0.88	974					
9	0.91	0.92	0.91	1009					
accuracy			0.93	10000					
macro avg	0.92	0.92	0.92	10000					
weighted avg	0.93	0.93	0.93	10000					

