BMS COLLEGE OF ENGINEERING

(An Autonomous Institution Affiliated to VTU, Belagavi)

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DEPARTMENT OF MACHINE LEARNING

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ADVANCED MACHINE LEARNING(22AM6PCAML)

Lab 3- AML Laboratory Programs

Submitted by

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Valuation Report (to be filled by the faculty)

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Comments:	Faculty Signature: with date	

1. Construct a Bayesian network considering medical data. Use this model to demonstrate the diagnosis of heart patients using standard Heart Disease Data Set.

```
In [ ]: !pip install pgmpy
     In [ ]: import pandas as pd
                 from pgmpy.models import BayesianNetwork
from pgmpy.estimators import MaximumLikelihoodEstimator
                 from pgmpy.inference import VariableElimination
     In [ ]: # Load the Heart Disease Data Set (replace with your dataset)
                 data = pd.read_csv('heart dis.csv')
    In [43]: print(data)
                 Age Gender
0 30-39 Male
                                                     ChestPain BloodPressure Cholesterol ECGResult \
                                  Male
                                             Typical Angina
                                                                                              High
                                                                                High
                 ypical Angina
Atypical Angina
2 50-59 Male Non-Anginal Pain
3 60-69 Female Asymptoma+*-
4 40-49 For-*
                                                                                               High Abnormal
                                                                              Normal
                                                                    Normaı
High
Normal
                                                                                                Normal Abnormal
                 4 40-49 Female Atypical Angina
                                                                                                              Normal
                                                                                                  High Abnormal
                 5 50-59 Male Typical Angina
6 60-69 Male Non-Anginal Pain
                                                                                                Normal
                                                                                                              Norma]
                     ExerciseAngina STDepression Diagnosis
                                                         2.0
                                  False
                                                         0.5
                                                                            0
                                    True
                                                          3.0
                                    True
                                  False
In [46]: # Define the variables and their dependencies
model = BayesianNetwork([
                   el = BayesianNetwork([
('Age', 'ChestPain'),
('Gender', 'ChestPain'),
('Age', 'BloodPressure'),
('Gender', 'BloodPressure'),
('Gender', 'Cholesterol'),
('Gender', 'Cholesterol'),
('Age', 'EGGResult'),
('ChestPain', 'ECGResult'),
('ChestPain', 'ExerciseAngina'),
('ECGResult', 'ExerciseAngina'),
('ECGResult', 'ExerciseAngina'),
('ESCRESULT', 'ExerciseAngina'),
('STDepression', 'Diagnosis')
             ])
             # Estimate the parameters (probabilities) using Maximum Likelihood Estimation
              model.fit(data, estimator=MaximumLikelihoodEstimator)
             # Perform inference on the model
infer = VariableElimination(model)
              # Query the model for the posterior probabilities
              query = infer.query(variables=['Diagnosis'], evidence={
   'Age': '50-59',
   'Gender': 'Male',
   'character'
                    'Gender': 'Male',
'ChestPain': 'Typical Angina',
'BloodPressure': 'High',
'Cholesterol': 'High',
'ECGResult': 'Abnormal',
'ExerciseAngina': True,
                    'STDepression': 2.5
              # Print the posterior probabilities
             print(query)
                 | Diagnosis | phi(Diagnosis) |
                 | Diagnosis(0) | 0.0000 |
                 +-----
                 | Diagnosis(1) | 1.0000 |
```

Application:

 Medical diagnosis: Bayesian networks are widely used in medical diagnosis systems to model the dependencies between symptoms, diseases, and test results.

- Risk assessment: Bayesian networks can be used to model and assess risks in various domains, such as finance, insurance, and engineering.
- Natural language processing: Bayesian networks have been applied in tasks such as text categorization, sentiment analysis, and information retrieval.

2. Build an Artificial Neural Network by implementing the Backpropagation algorithm and test the same using appropriate data sets.

```
In [1]: # Importing libraries
             import pandas as pd
import numpy as np
             import matplotlib.pyplot as plt
             data = pd.read_csv('Breast cancer.csv')
del data['Unnamed: 32']
   In [2]: X = data.iloc[:, 2:].values
y = data.iloc[:, 1].values
              # Encoding categorical data
             from sklearn.preprocessing import LabelEncoder
labelencoder_X_1 = LabelEncoder()
y = labelencoder_X_1.fit_transform(y)
             # Splitting the dataset into the Training set and Test set from sklearn.model_selection import train_test_split
             X_train, X_test, y_train, y_test = train_test_split(X, y, test_size = 0.1, random_state = 0)
              #Feature Scaling
             from sklearn.preprocessing import StandardScaler
             sc = StandardScaler()
X_train = sc.fit_transform(X_train)
             X_test = sc.transform(X_test)
   In [4]: !pip install keras
             Collecting keras
                Downloading keras-2.12.0-py2.py3-none-any.whl (1.7 MB)
                                                                      -- 1.7/1.7 MB 5.8 MB/s eta 0:00:00
              Installing collected packages: keras
              Successfully installed keras-2.12.0
 In [5]: import keras
            from keras.models import Sequential
from keras.layers import Dense, Dropout
 In [6]: # Initialising the ANN
            classifier = Sequential()
            classifier.add(Dense(units=16, kernel initializer='uniform', activation='relu', input dim=30))
            # Adding dropout to prevent overfitting classifier.add(Dropout(rate=0.1))
  In [8]: # Adding the second hidden layer
            classifier.add(Dense(units=16, kernel_initializer='uniform', activation='relu'))
            # Adding dropout to prevent overfitting
           classifier.add(Dropout(rate=0.1))
            classifier.add(Dense(units=1, kernel initializer='uniform', activation='sigmoid'))
In [10]: # Compiling the ANN
            classifier.compile(optimizer='adam', loss='binary_crossentropy', metrics=['accuracy'])
In [11]: # Fitting the ANN to the Training set
    classifier.fit(X train, y_train, batch_size=100, epochs=50)
    # Long scroll ahead but worth
    # The batch size and number of epochs have been set using trial and error. Still looking for more efficient ways. Open to suggest
```

```
Epoch 1/50
                  ======== ] - 2s 5ms/step - loss: 0.6926 - accuracy: 0.6094
6/6 [=====
Epoch 2/50
6/6 [=====
                    ======== ] - 0s 3ms/step - loss: 0.6902 - accuracy: 0.6406
Epoch 3/50
6/6 [====
                                      0s 3ms/step - loss: 0.6869 - accuracy: 0.6816
Epoch 4/50
                                       0s 3ms/step - loss: 0.6817 - accuracy: 0.7637
Epoch 5/50
6/6 [=====
                                       0s 4ms/step - loss: 0.6730 - accuracy: 0.8457
Epoch 6/50
                                     - 0s 4ms/step - loss: 0.6602 - accuracy: 0.9043
6/6 [=====
Epoch 7/50
6/6 [=====
                                     - 0s 4ms/step - loss: 0.6409 - accuracy: 0.9199
Epoch 8/50
6/6 [=====
Epoch 9/50
                                      0s 4ms/step - loss: 0.6156 - accuracy: 0.9375
6/6 [===
                                      0s 4ms/step - loss: 0.5830 - accuracy: 0.9414
Epoch 10/50
                                       0s 4ms/step - loss: 0.5418 - accuracy: 0.9414
Epoch 11/50
                                       0s 4ms/step - loss: 0.4900 - accuracy: 0.9414
Epoch 12/50
6/6 [======
                                     - 0s 4ms/step - loss: 0.4393 - accuracy: 0.9395
Epoch 13/50
                                      Os 3ms/step - loss: 0.3886 - accuracy: 0.9336
6/6 [=======
Epoch 14/50
6/6 [===
                                     - 0s 3ms/step - loss: 0.3348 - accuracy: 0.9395
Epoch 15/50
6/6 [===
                                       0s 3ms/step - loss: 0.2933 - accuracy: 0.9434
Epoch 16/50
                                       0s 4ms/step - loss: 0.2570 - accuracy: 0.9453
Epoch 17/50
6/6 [====
                                       0s 3ms/step - loss: 0.2302 - accuracy: 0.9512
Epoch 18/50
                      ======== ] - 0s 3ms/step - loss: 0.2012 - accuracy: 0.9531
6/6 [====
Epoch 19/50
                   ======== ] - 0s 3ms/step - loss: 0.1856 - accuracy: 0.9609
6/6 [=====
Epoch 20/50
Epoch 21/50
                    =======] - 0s 3ms/step - loss: 0.1550 - accuracy: 0.9707
Epoch 22/50
                                      0s 3ms/step - loss: 0.1427 - accuracy: 0.9688
Fnoch 23/50
                                      0s 3ms/step - loss: 0.1312 - accuracy: 0.9648
6/6 [=====
Epoch 24/50
6/6 [=====
                                      0s 4ms/step - loss: 0.1272 - accuracy: 0.9668
Epoch 25/50
                                       0s 3ms/step - loss: 0.1243 - accuracy: 0.9707
6/6 [======
Epoch 26/50
6/6 [====
                                       0s 3ms/step - loss: 0.1194 - accuracy: 0.9668
Epoch 27/50
6/6 [=====
Epoch 28/50
                                       0s 4ms/step - loss: 0.1103 - accuracy: 0.9727
                                       0s 3ms/step - loss: 0.1024 - accuracy: 0.9746
Epoch 29/50
                                       0s 3ms/step - loss: 0.1007 - accuracy: 0.9766
Epoch 30/50
                                       0s 3ms/step - loss: 0.0965 - accuracy: 0.9766
6/6 [=====
Epoch 31/50
6/6 [=====
                                       0s 4ms/step - loss: 0.0945 - accuracy: 0.9785
Epoch 32/50
6/6 [=====
Epoch 33/50
                                       0s 3ms/step - loss: 0.0914 - accuracy: 0.9785
6/6 [=====
                   ======== ] - 0s 3ms/step - loss: 0.0878 - accuracy: 0.9805
Epoch 34/50
6/6 [=======]
Epoch 35/50
                                       0s 3ms/step - loss: 0.0894 - accuracy: 0.9727
6/6 [=====
                                       0s 3ms/step - loss: 0.0835 - accuracy: 0.9785
Epoch 36/50
                                       0s 3ms/step - loss: 0.0802 - accuracy: 0.9785
Epoch 37/50
6/6 [===
                                       0s 4ms/step - loss: 0.0780 - accuracy: 0.9844
Epoch 38/50
6/6 [=====
                         =======] - 0s 4ms/step - loss: 0.0788 - accuracy: 0.9844
Epoch 39/50
                  ======== 1 - 0s 3ms/step - loss: 0.0790 - accuracy: 0.9785
6/6 [=====
Epoch 40/50
```

```
Epoch 41/50
                                       ====] - 0s 4ms/step - loss: 0.0733 - accuracy: 0.9805
         6/6 [====
         Epoch 42/50
                                6/6 [====
         Epoch 43/50
         6/6 [=====
                                             - 0s 4ms/step - loss: 0.0775 - accuracy: 0.9805
         Epoch 44/50
                                             - 0s 4ms/step - loss: 0.0705 - accuracy: 0.9824
         6/6 [==:
         Epoch 45/50
         6/6 [====
                                    ======] - 0s 4ms/step - loss: 0.0684 - accuracy: 0.9824
         Epoch 46/50
                                            - 0s 4ms/step - loss: 0.0722 - accuracy: 0.9805
         6/6 [=====
Epoch 47/50
                                            - 0s 4ms/step - loss: 0.0691 - accuracy: 0.9824
         6/6 [====
         Epoch 48/50
                                            - 0s 3ms/step - loss: 0.0669 - accuracy: 0.9824
         6/6 [===
         Epoch 49/50
                                 =======] - 0s 4ms/step - loss: 0.0650 - accuracy: 0.9805
         6/6 [===
         Epoch 50/50
                              6/6 [======
Out[11]: <keras.callbacks.History at 0x120f3816590>
In [12]: # Predicting the Test set results
y_pred = classifier.predict(X_test)
y_pred = (y_pred > 0.5)
         2/2 [======] - 0s 22ms/step
In [13]: # Making the Confusion Matrix
from sklearn.metrics import confusion_matrix
         cm = confusion_matrix(y_test, y_pred)
In [14]: print("Our accuracy is {}%".format(((cm[0][0] + cm[1][1])/57)*100))
         Our accuracy is 96.49122807017544%
In [15]: sns.heatmap(cm,annot=True)
         plt.savefig('h.png')
                                                                                  - 30
                                                                                   - 25
                          33
                                                          2
        0 -
                                                                                   20
                                                                                  - 15
                                                                                   10
                           0
                                                                                   5
                                                          1
```

Application:

- Neural networks: Backpropagation is the primary learning algorithm used to train artificial neural networks for various tasks, including pattern recognition, image classification, and speech recognition.
- Function approximation: Backpropagation can be used to approximate complex functions by training neural networks with appropriate architectures and activation functions.
- Time series prediction: Backpropagation-based neural networks can be applied to predict future values in time series data, such as stock market prices or weather forecasting.

3. Demonstrate the working of EM algorithm to cluster a set of data stored in a .CSV file.

```
In [3]: import numpy as np
             import pandas as pd
from scipy.stats import multivariate_normal
In [9]: class GMM:
    def __init__(self, n_clusters, max_iterations=100):
        self.n_clusters = n_clusters
        self.max_iterations = max_iterations
                         self.weights = None
                         self.means = None
self.covariances = None
                   def fit(self, X):
                         n_samples, n_features = X.shape
                         # Initialize parameters randomly
                         self.weights = np.ones(self.n_clusters) / self.n_clusters
self.means = np.random.rand(self.n_clusters, n_features)
self.covariances = np.array([np.eye(n_features)] * self.n_clusters)
                         # EM algorithm
                         for _ in range(self.max_iterations):
    # E-step: Calculate responsibilities
                               responsibilities = self._expectation(X)
                               # M-step: Update parameters
self._maximization(X, responsibilities)
                   def predict(self, X):
    # E-step: Calculate responsibilities
                         responsibilities = self._expectation(X)
                          # Assign samples to clusters based on responsibilities
                         labels = np.argmax(responsibilities, axis=1)
                         return labels
                 def _expectation(self, X):
                      n_samples = X.shape[0]
                      # Calculate probabilities using the current parameters
probabilities = np.zeros((n_samples, self.n_clusters))
for k in range(self.n_clusters):
    probabilities[:, k] = self.weights[k] * multivariate_normal.pdf(X, self.means[k], self.covariances[k])
                      # Calculate responsibilities using Bayes' rule
responsibilities = probabilities / np.sum(probabilities, axis=1, keepdims=True)
                       return responsibilities
                def _maximization(self, X, responsibilities):
                      n_samples = X.shape[0]
                      # Update weights
self.weights = np.mean(responsibilities, axis=0)
                       # Update means
                      # Optate meural
for k in range(self.n_clusters):
    weight_k = self.weights[k]
    mean_k = np.sum(responsibilities[:, k].reshape(-1, 1) * X, axis=0) / (n_samples * weight_k)
    self.means[k] = mean_k
                      weight_k = self.m_clusters):
    weight_k = self.weights[k]
    diff = X - self.means[k]
    covariance_k = np.dot((responsibilities[:, k].reshape(-1, 1) * diff).T, diff) / (n_samples * weight_k)
                             self.covariances[k] = covariance_k
          # Load the dataset from CSV file
data = pd.read_csv('Em algo.csv')
          X = data.values
          # Create and fit the GMM model
          gmm = GMM(n_clusters)
```

gmm.fit(X)

```
from sklearn.preprocessing import LabelEncoder
# label encoder = LabelEncoder()
# X_encoded = label_encoder.fit_transform(X_categorical)
# Make predictions
labels = gmm.predict(X)
print("Cluster labels:", labels)
\begin{smallmatrix}1&1&1&1&2&2&2&1&2\\1&1&1&1&1&2&2&2&1&2\\\end{smallmatrix}
1 2 2 2 2 2 1 2 2 2 2 2 2 1 2 2 2 1 1 1 2 1 2 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1
1\; 1\; 1\; 1\; 1\; 1\; 1\; 1\; 1\; 1\; 1\; 1\; 1\; 2\; 2\; 0\; 2\; 0\; 2\; 2\; 2\; 2\; 2\; 2\; 2\; 2\; 1\; 2\; 1\; 1\; 1\; 1\; 2\; 2\; 1\; 1\; 1\; 1\; 1\; 2\; 2
1 2 1 2 1 2 1 1 1 1 1 1 1 1 1 1 1 1 2 2 1 2 2 2 2 2 2 2 2 2 2 1 2 1 2 2 2 1 1
2 1 1 1 1 1 1 1 1 1 2 2 1 1 1 1 1 2 2 2 2 2 2 1 1 1 2 2 2 2 2 1 1 1 2 2 1 2 2 1 1 1 2 2 1 2 2 1
```

Application:

- Clustering: The EM algorithm is used in clustering algorithms like Gaussian Mixture Models (GMM) to estimate the parameters of each cluster and assign data points to the most likely cluster.
- Image segmentation: EM algorithm can be used to segment images by modeling the distribution of pixel intensities in different regions.
- Missing data imputation: EM algorithm can be applied to impute missing values in datasets by estimating the missing values based on observed data.
- 4. Demonstrate Pre processing (Data Cleaning, Integration and Transformation) activity on suitable data: For example: Identify and Delete Rows that Contain Duplicate Data by considering an appropriate dataset. Identify and Delete Columns That Contain a Single Value by considering an appropriate dataset.

```
In [1]: import pandas as pd
         # Load the dataset into a pandas DataFrame
        data1= pd.read_csv('Emp info.csv')
print(data1)
                                          Email
                                                         Phone Country
                             john@example.com 123-456-7890
             John
                                                                    USA
                            emily@example.com 987-654-3210
            John
                    25
                             iohn@example.com 123-456-7890
                                                                    USA
                            sarah@example.com 555-555-555
            Sarah
                            peter@example.com 111-222-3333
emily@example.com 987-654-3210
            Peter
                    28
                                                                    USA
                   26 emily@example.com 98/-03-25

25 john.doe@example.com 123-456-7890

25 john.doe@example.com 555-555-5555
            Emily
             John
                                                                    USA
           Sarah
                            sarah@example.com 555-555-5555 Canada
In [2]: # Identify and delete duplicate rows based on all columns
        data = data1.drop_duplicates()
         # Print the updated dataset after removing duplicates
        print(data)
             Name Age
                                         Email
                                                         Phone Country
                            john@example.com 123-456-7890
emily@example.com 987-654-3210
           Emily 30
                                                                     UK
                   35
                            sarah@example.com 555-555-5555
            Peter
                    28
                            peter@example.com 111-222-3333
                                                                    USA
                   25 john.doe@example.com 123-456-7890
In [3]: # # Identify and delete columns that contain a single value
         # Assuming your dataset is stored in a DataFrame called 'data
         data.drop(data.columns[-1], axis=1, inplace=True)
         print(data)
  In [4]: data2 = pd.read_csv('Employee Salary.csv')
           print(data2)
               Name Employee_id
           1 Emily
              Sarah
              Emily
           6 John
7 Sarah
 In [12]: #merging the datasets
           merged_data = pd.merge(data1 ,data2, on = 'Employee_id')
           print(merged_data)
               Employee_id Experience_Years Age Gender Salary
1 5 28 Female 30000
                                                 21
                                                        Male
                                                                60000
                                             2 22
                                                        Male
                                                                10000
                                                 62
                                                        Male
                                                                40000
                                                 54 Female
                                                 21
                                                                40000
                                                      Female
                                                      Female
Female
                                                                30000
           10
                         11
                                                 26
                                                      Female
                                                                20000
           12
                         13
                                            14
                                                        Male
                                                                10000
```

Applications:

- Data cleaning: Pre-processing techniques like handling missing values, removing outliers, and correcting inconsistent data are used to improve the quality of data before further analysis.
- Feature scaling and normalization: Pre-processing can involve scaling or normalizing features to ensure that they have similar ranges or distributions, which can improve the performance of certain algorithms like SVM or K-means.
- Dimensionality reduction: Techniques like Principal Component Analysis (PCA) or Feature Selection can be used to reduce the number of features while preserving relevant information and reducing computational complexity.

5. For a given set of training data examples stored in a .CSV file, implement and demonstrate the Candidate-Elimination algorithm. Output a description of the set of all hypotheses consistent with the training examples.

```
In [1]:
                                                 import numpy as np
import pandas as pd
                                                  for dirname, _, filenames in os.walk('trainingdata.csv'):
    for filename in filenames:
                                                                            print(os.path.join(dirname, filename))
                     In [2]: import random
                                                  import csv
                    def s_0(n):
                                                               return ('0',)*n
                     In [4]: def more_general(h1, h2):
                                                                more_general_parts = []
for x, y in zip(h1, h2):
    mg = x == "?" or (x != "0" and (x == y or y == "0"))
    more_general_parts.append(mg)
                                                                return all(more_general_parts)
                                                 11 = [1, 2, 3]
12 = [3, 4, 5]
                                                  list(zip(l1, l2))
                    Out[4]: [(1, 3), (2, 4), (3, 5)]
In [5]: # min_generalizations
def fulfills(example, hypothesis):
                                             ### the implementation is the same as for hypotheses:
                                              return more_general(hypothesis, example)
                               def min_generalizations(h, x):
                                            mall_general factors(n, x).
h_new = list(h)
for i in range(len(h)):
    if not fulfills(x[i:i+1], h[i:i+1]):
        h_new[i] = '?' if h[i] != '0' else x[i]
return [tuple(h_new)]
In [6]:  \begin{aligned} &\text{min\_generalizations(h=('0', '0' , 'sunny'),} \\ && & & & & \\ && & & & \\ && & & & \\ && & & & \\ && & & & \\ && & & & \\ && & & & \\ && & & & \\ && & & & \\ && & & & \\ && & & \\ && & & & \\ && & & \\ && & & \\ && & & \\ && & & \\ && & & \\ && & & \\ && & & \\ && & & \\ && & & \\ && & & \\ && & & \\ && & & \\ && & & \\ && & & \\ && & & \\ && & & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && & \\ && \\ && & \\ && \\ && \\ && \\ && \\ && \\ && \\ && \\ && \\ && \\ && \\ && \\ &
Out[6]: [('rainy', 'windy', '?')]
 In [7]: def min_specializations(h, domains, x):
                                            if x[i]!= val:

h_new = h[:i] + (val,) + h[i+1:]

results.append(h_new)

elif h[i]!= "0":

h_new = h[:i] + ('0',) + h[i+1:]

results.append(h_new)
                                              return results
Out[8]: [('a', 'x'), ('c', 'x'), ('?', '0')]
```

```
In [12]: def get_domains(examples):
    d = [set() for i in examples[0]]
                    for x in examples:
    for i, xi in enumerate(x):
        d[i].add(xi)
return [list(sorted(x)) for x in d]
              get_domains(examples)
G = set([g_0(len(domains))])
                    S = set([s_0(len(domains))])
i=0
                    print("\n G[{0}]:".format(i),G)
print("\n S[{0}]:".format(i),S)
                     for xcx in examples:
                         if cx=='Y': # x is positive example
                         ar xx= Y: #x is positive example
   G = {g for g in G if fulfills(x, g)}
   S = generalize_S(x, G, S)
else: #x is negative example
   S = {s for s in S if not fulfills(x, s)}
                         G = specialize G(x, domains, G, S)
print("\n G[{0}]:".format(i),G)
print("\n S[{0}]:".format(i),S)
                   return
In [14]: def generalize_S(x, G, S):
    S_prev = list(S)
                   for s in S_prev:
   if s not in S:
                         continue
if not fulfills(x, s):
                               S.remove(s)
                               S.remove(s)
Splus = min generalizations(s, x)
## keep only generalizations that have a counterpart in G
S.update([h for h in Splus if any([more_general(g,h) for g in G])])
## remove hypotheses less specific than any other in S
                               S.difference_update([h for h in S if any([more_general(h, h1) for h1 in S if h != h1])])
                   return S
for g in G_prev:
if g not in G:
                               continue
                          if fulfills(x, g):
                              Tulriis(x, g):
    G.remove(g)
    Gminus = min_specializations(g, domains, x)
    ## keep only specializations that have a conuterpart in S
    G.update([h for h in Gminus if any([more_general(h, s) for s in S])])
                               for s in s])])
## remove hypotheses less general than any other in G
```

Applications:

- Concept learning: The Candidate-Elimination algorithm is used to learn concepts from labeled data and update the hypothesis space iteratively.
- Inductive logic programming: The algorithm can be used to learn logical rules or hypotheses from examples and background knowledge.
- 6. Implement and demonstrate the FIND-S algorithm for finding the most specific hypothesis based on a given set of training data samples. Read the training data from a .CSV file and show the output for test cases. Develop an interactive program by Comparing the result by implementing LIST THEN ELIMINATE algorithm.

```
In [1]: import numpy as np
import pandas as pd

In [41]: import csv
num_attributes = 6
    a = []
    print("\n The Given Training Data Set \n")
    file = 'ws.csv'
    with open(file, 'r') as csvfile:
        reader = csv.reader(csvfile)
        for row in reader:
            a.append (row)
            print(row)

        type(reader)

The Given Training Data Set

        ['sunny', 'Warm', 'Normal', 'Strong', 'Warm', 'Same', 'Yes']
        ['sunny', 'Warm', 'High', 'Strong', 'Warm', 'Same', 'Yes']
        ['sunny', 'Warm', 'High', 'Strong', 'Warm', 'Change', 'No']
        ['sunny', 'Warm', 'High', 'Strong', 'Warm', 'Change', 'Yes']

Out[41]: _csv.reader

In [42]: a[0][-1]

Out[42]: ['sunny', 'Warm', 'Normal', 'Strong', 'Warm', 'Same']

In [43]: print("\n The initial value of hypothesis: ")
        hypothesis = ['0'] * num_attributes
        print(hypothesis)
```

```
for j in range(0,num_attributes):
    hypothesis[j] = a[0][j]
            hypothesis
            The initial value of hypothesis: ['0', '0', '0', '0', '0', '0']
  Out[43]: ['Sunny', 'Warm', 'Normal', 'Strong', 'Warm', 'Same']
  In [44]: hypothesis == a[0][:-1]
  Out[44]: True
  In [45]: print("\n Find S: Finding a Maximally Specific Hypothesis\n")
            hypothesis[j]='?
else :
                hypothesis[j]= a[i][j]
print("\n\nFor Training instance No:{} the hypothesis is ".format(i), hypothesis)
            \label{print("$n$ The Maximally Specific Hypothesis for a given Training Examples :$\n"$)} \\
            print(hypothesis)
            Find S: Finding a Maximally Specific Hypothesis
           Sunny Warm Normal Strong Warm Same
           For Training instance No:0 the hypothesis is ['Sunny', 'Warm', 'Normal', 'Strong', 'Warm', 'Same']
           Sunny Warm High Strong Warm Same
           For Training instance No:1 the hypothesis is ['Sunny', 'Warm', '?', 'Strong', 'Warm', 'Same']
           For Training instance No:2 the hypothesis is ['Sunny', 'Warm', '?', 'Strong', 'Warm', 'Same']
           Sunny Warm High Strong Cool Change
           For Training instance No:3 the hypothesis is \ ['Sunny', 'Warm', '?', 'Strong', '?', '?']
            The Maximally Specific Hypothesis for a given Training Examples :
           ['Sunny', 'Warm', '?', 'Strong', '?', '?']
In [18]: import pandas as pd
          def list_then_eliminate(training_data):
              positive_examples = training_data[training_data['Play'] == 'No']
              if positive_examples.empty:
    return None
              hypothesis = positive examples.iloc[0, :-1]
              for _, example in training_data.iterrows():
                  if example['Play'] == 'No':
    for attr in hypothesis.keys():
                           if example[attr] != hypothesis[attr]:
    hypothesis[attr] = '?'
             return hypothesis
          # Load the training data
          training_data = pd.read_csv('dataset 1.csv')
print(training_data)
              Outlook Temperature Humidity Windy Play
                                        High False No
High False Yes
          1 Overcast
                               Hot
                 Rain
                               Mild
                                        High False Yes
                 Rain
                               Cool
                                      Normal False Yes
                Sunny
                              Mild
                                      Normal
```

```
In [14]: # Run the LIST THEN ELIMINATE algorithm
hypothesis_space = list_then_eliminate(training_data)

# Output the hypothesis space
print("Hypothesis space:")
print(hypothesis_space)

Hypothesis space:
Outlook Sunny
Temperature Hot
Humidity High
Windy False
Name: 0, dtype: object
```

Application:

- Concept learning: The FIND-S algorithm is used to learn the most specific hypothesis from labeled examples in the form of attribute-value pairs.
- Rule induction: FIND-S can be applied to induce classification rules from data, where each rule represents a specific condition that leads to a particular class label.
