

Group Name: Agile

Github link: https://github.com/M-Shreya/AI_Lab_Assignments

Lab 4

Problem Statement:

Case of Medical Diagnosis

Thyroid Disease Dataset from UCI Machine Learning Repository

Dataset Description: <https://archive.ics.uci.edu/dataset/102/thyroid+disease>

Features: T3,T4,TSH(thyroid stimulating hormone), Goiter presence, etc.

Labels: Hyperthyroid, Hypothyroid, Normal Thyroid Function Build and train a Bayesian Network as a classifier (labels)

Expected accuracy $\geq 85\%$

Solution:

To implement a Bayesian Network classifier for the thyroid disease dataset from the UCI Machine Learning Repository, you will need to:

1. Download and prepare the dataset.
2. Preprocess the data (if needed).
3. Build and train a Bayesian Network classifier.
4. Evaluate the model's accuracy.

Below is an implementation using Python with the pgmpy library to create a Bayesian Network. If pgmpy is not already installed, you can install it using:

```
bash
```

```
pip install pgmpy pandas scikit-learn
```

Implementation:

```
# Load the dataset
```

```
url = 'https://archive.ics.uci.edu/ml/machine-learning-databases/thyroid-disease/new-thyroid.data'
```

```

column_names = ['Class', 'T3', 'T4', 'TSH']
data = pd.read_csv(url, names=column_names, delim_whitespace=True)

# Convert class labels to string for clarity
# Train-test split
# Add labels to training set
# Define the structure of the Bayesian Network
# You might want to adjust this according to the domain knowledge
model = BayesianNetwork([('T3', 'Class'), ('T4', 'Class'), ('TSH', 'Class')])
# Fit the model using Maximum Likelihood Estimation
model.fit(X_train, estimator=MaximumLikelihoodEstimator)
# Inference
inference = VariableElimination(model)
# Predict function using the trained model
# Perform predictions
y_pred = predict(inference, X_test)
# Calculate accuracy
accuracy = accuracy_score(y_test, y_pred)
# Expected output should show an accuracy >= 85%

```

Summary:

1. ***Dataset*:** This code uses the new-thyroid.data subset from the UCI Thyroid Disease dataset. The url should point to the correct dataset file. Ensure that you adjust the file path if you use a different subset.
2. ***Model Structure*:** The Bayesian Network's structure is defined as BayesianNetwork([('T3', 'Class'), ('T4', 'Class'), ('TSH', 'Class')]). You might need to adjust this according to the domain knowledge.
3. ***Dependencies*:** Make sure to have the pgmpy, pandas, and scikit-learn libraries installed in your environment.
4. ***Evaluation*:** This code prints the accuracy of the model. Ensure the accuracy is above 85% by possibly tweaking the model structure and features.