

Project 3: Cluster Validation

Purpose

In this project, you will apply the cluster validation technique to data extracted from a provided data set.

Objectives

Learners will be able to:

- Develop code that performs clustering.
- Test and analyze the results of the clustering code.
- Assess the accuracy of the clustering using SSE and supervised cluster validity metrics.

Technology Requirements

- Python 3.10.9
- scikit-learn==1.1.1
- pandas==1.4.3
- numpy==1.23.1
- scipy==1.8.1
- matplotlib==3.6.0

Project Description

For this project, you will write a program using Python that takes a dataset and performs clustering. Using the provided training data set you will perform cluster validation to determine the amount of carbohydrates in each meal.

Please watch “**Project 3: Cluster Validation Introductory Video**” before beginning Project 3. This is located in your course *Welcome and Start Here* section.

Note: Project details in the Overview Document were recently updated since the recording of the videos, so some directions or items may not match. Please follow the Overview Document directions to complete your project correctly.

Directions

There are two main parts to the process:

1. Extract features from Meal data
2. Cluster Meal data based on the amount of carbohydrates in each meal

Data:

Use the Project 1 data files:

- CGMData.csv
- InsulinData.csv

Extracting Ground Truth:

Derive the max and min value of meal intake amount from the Y column of the Insulin data. Discretize the meal amount in bins of size 20. Consider each row in the meal data matrix that you generated in Project 2. Put them in the respective bins according to their meal amount label.

In total, you should have $n = (\text{max-min})/20$ bins.

Performing clustering:

Use the features in your Project 2 to cluster the meal data into n clusters. Use DBSCAN and KMeans.

Report your accuracy of clustering based on SSE, entropy, and purity metrics.

Expected Output:

A Result.csv file which contains a 1 X 6 vector. The vector should have the following format:

| SSE for Kmeans | SSE for DBSCAN | Entropy for KMeans | Entropy for DBSCAN | Purity for KMeans | Purity for DBSCAN |
|----------------|----------------|--------------------|--------------------|-------------------|-------------------|
| | | | | | |

The Result.csv file should not have any headers, just the six values in six columns.

Submission Directions for Project Deliverables

Deliverables

- In the code you should have one main python file which the autograder can run and generate Result.csv file according to specifications.
- Assume that CGMData.csv and InsulinData.csv are already in the execution folder.
- You can have as many auxiliary python files as you want but the autograder will only run the main.py and it should generate the Result.csv.

Submission Guidelines:

- Please submit a zip file which has all your code and titled as: **"yourfirstname_lastname_Project3.zip"**.
 - Do not create an additional folder; just zip the files directly.
- The submission space is located at the bottom of Week 6 under "Week 6: Graded Coursework" as **"Project Assignment: Project 3 Cluster Validation Submission"**.

Evaluation

The autograder in Coursera will evaluate your code based on the following criteria:

- 50 points for developing a code in Python that takes the dataset and performs clustering.
- 20 points for developing a code in Python that implements a function to compute SSE, entropy and purity metrics. These two can be written in the same file.
- 30 points will be evaluated on the supervised cluster validation results obtained by your code.

Note: The autograder has fixed values for minimumEntropy, maximumPurity, and standard deviation for K-Means and DBSCAN and uses these values to perform a few mathematical calculations. Your K-Means and DBSCAN Purity and Entropy should be in the range of these calculations. Below are the minEntropy and maxPurity values:

- Minimum K-Means Entropy: 0.3235
- Minimum DBSCAN Entropy: 0.1739
- Maximum K-Means Purity: 0.875
- Maximum DBSCAN Purity: 1

Common Errors:

- **ModuleNotFoundError:** You are trying to access or use modules that are not found in the grader. Use the modules mentioned in the Technology Requirements section.