

CSE 347/447 Data Mining: Project 1 – Clustering Algorithms

Due on 11:59 PM, March 4, 2020

Submission Instructions: Please submit your Code and Report as .zip file to CourseSite. Schedule a meeting with TA (zhk219@lehigh.edu) after submission.

Note that copying code/report from another student or source is not allowed and may result in an F in the grade. Partial credit will be given for partial solutions.

Remember that you can be at most 3 days late; for every late date you lose 10% of your grade.

Datasets: Cho and Iyer are gene sequences datasets which need to be clustered according to the similar gene patterns. You can download them at: https://drive.google.com/open?id=1I00QuY1lu0ahnEy-hkEN_MTwATm2uXiZ and https://drive.google.com/open?id=1KHTaMp_9yfID6Q1wg1YsLnfmwu2g2C7A.

A short description of the datasets can be found at <https://drive.google.com/open?id=1Nm71bYSUsIp8cFGq7UQmgJUhCcY2yuqT>.

Complete the following tasks:

- Implement k-means and spectral clustering algorithms to find clusters of genes which exhibit similar expression profiles. Compare the two methods and discuss their pros and cons.

You are required to validate your clustering results using the following methods:

- Choose an external index and compare the clustering results from different clustering algorithms with an external index (the ground truth clusters are provided in the data sets).
- Choose an internal index and compare the clustering results.

Your final submission should include the following:

- Code: Two clustering algorithms implemented with two functions, respectively. Your code should allow the users to choose either of the clustering algorithm. Just an example:
cin 'the method you choose:'
cout 'clustering result is:'
- Report: Describe the flow of all the implemented algorithms. Compare the performance of these approaches using external and internal index on the two given datasets. State the pros and cons of each algorithm and any findings you get from the experiments.

If you choose MATLAB (every Lehigh student has free access), you could use the GUI interface with menu bars. This is not required but you will find it useful. Enjoy yourself, and if you have any question, feel free to ask.