
Introduction to Machine Learning

Work 1 Clustering exercise

Course 2021-2022

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1 Description of the work

The aim of the exercise is to analyze different clustering algorithms using several data sets from the UCI repository. To this end, first of all you will implement several clustering algorithms using Python 3.7 and PyCharm IDE.

1.1 Methodology of the analysis

You will analyze the behavior of different clustering algorithms in well-known data sets from the UCI repository. These data sets are defined in .arff format. So, you will be able to analyze them with the Weka environment, too. A guide can be found at:

<https://www.cs.auckland.ac.nz/courses/compsci367s1c/tutorials/IntroductionToWeka.pdf>

<http://www.cs.utexas.edu/users/ml/tutorials/Weka-tut/>

The Weka is used to analyze if your code in Python is correct or not.

This work is divided in six tasks:

1. Implement your code for reading the arff file in Python and store the information in memory. Be careful, some of the data sets contain numerical and categorical data and may also contain missing values. For this exercise it is not necessary to store the class of the data set. However, for future assignments you will be requested to use the class. In the following address:

<https://docs.scipy.org/doc/scipy/reference/generated/scipy.io.arff.loadarff.html>

You will find a code for reading and writing arff files in Python. You can implement your own parser (do not use any other library) or you can analyze the code based on `scipy.io` library in the abovementioned link, execute it, and modify it accordingly to your needs. If you need it, for the numerical datasets, you can transform the continuous variables/attributes into categorical variables using function `pandas.qcut`.

2. Implement in a Python file the code that uses the **OPTICS** (*Ordering Points To Identify the Clustering Structure*) from the `sklearn` library and apply it to the data of the file. Test 3 different distance metrics, such as *Euclidean*, *Cosine*, *l1* or *l2* (among others), and evaluate what happen when you set up the algorithm with two different approximations to the nearest neighbors: *auto*, *ball_tree*, *kd_tree* or *brute*.
3. Implement **your own K-Means** (KM) algorithm and apply it to the data of the file. **Note that you are not allowed to use** `sklearn` library or any other library.
4. Choose one of the following algorithms and **implement your own code**: **K-Modes**, **K-Medoids**, or **K-Prototypes**. **Note that you are not allowed to use** `sklearn` or any other library that implements these algorithms. Next, apply it to different data sets.

5. Implement **your own code** in one of the fuzzy clustering algorithms: Fuzzy C-Means (FCM) (Bezdek, 1981) or Possibilistic C-Means (PCM) (Krishnapuram - Keller, 1993).
6. Analyze the algorithms in three data sets (see Section 2) considering the following restrictions:
 - a. Two should contain all the attributes categorical or mixed attributes, and one with all the attributes numerical attributes.
 - b. At least two of them should be large enough to be able to extract conclusions.

1.2 Work to deliver

In this work, you will use and analyze OPTICS algorithm and you will implement and analyze K-Means, another algorithm of your choice (K-Modes, K-Medoids, or K-Prototypes), and a Fuzzy Clustering algorithm. You may select 3 data sets for your analysis. At the end, you will find a list of the data sets available (see Section 2). The data sets are in a zip file in Racó at UPC and in campus virtual at UB (campusvirtual.ub.edu).

The idea is that you implement **your own code in Python 3.7** and you will use it to produce the results of the analysis. The development will be in the **PyCharm IDE** and you will deliver the project with your code in campus virtual at UB.

Once you have obtained the results, you will show them in several ways:

1. **Compare using tables and/or graphs the clustering algorithms using some clustering validation metrics.** Some examples are: **Adjusted Rand Index, Purity, Davies–Bouldin index, F-measure.** You can use these ones or other ones from the literature that best suit your evaluation. **For the evaluation metrics, you can use the ones defined in sklearn library.**
2. The results obtained can be compared to the true values. **To show the results, you can use a confusion matrix, for example.**

From the tables and graphs, you will reason and extract conclusions about the results obtained. For example, some questions that may help you to comment your results:

- Which information can be obtained for each data set using each algorithm? Is it the same or not?
- Which clustering algorithm do you consider is the best one for datasets with categorical data, with numerical data and with mixed data?
- Did you find differences among algorithms? According to the data sets chosen, which algorithm gives you more advice for knowing the underlying information in the data set?
- Can you explain the setup that you have used for each algorithm?
- In the case of the K-Means and the other algorithms where you have to choose the K, which has been the best K value?

- Have you implemented any improvement on the basic algorithms? For example, you can introduce/use a performance measure, like Silhouette, to decide which the best K value is. Another example, you can use K-Means++ for choosing the initial values (or "seeds") for the K-Means algorithm. Alternatively, you can use G-Means that is an algorithm for determining automatically the best K value.
- In the case of Fuzzy Clustering algorithm, you can optimize the C value. Have you done the optimization? Which are the results? In case that you have not included the optimization, how many C- values have you tested for each data set? And which value do you consider it is the best one?

You should deliver a report as well as the code in Python 3.7 and PyCharm project in Campus virtual in a zip file by **October 31, 2021**. Please, the name of the zip file should contain the name and surname of each member of the group, and the number of your work. For example, considering that this assignment is Work 1 (we will use the acronym w1) and the students are Bart and Lisa Simpson, the name of the file will be: *BartSimpsonLisaSimpson_w1.zip*

The maximum extension allowed for the report is 10 pages in a two-column report or 20 pages in a one-column report. Please take a look at the report guidelines at the end of this document.

The report will contain:

- Details about the implementation of your algorithms, including the decisions made during the implementation and the setup of the different parameters.
- The evaluation of the algorithms, including tables and/or graphs that show your results with comments about them.
- Justify your results and, in addition, reason each one of the questions defined above in your evaluation. Moreover, add any comment or observation that you consider important from your results.
- **It is extremely important that you explain how to execute your code. Moreover, call files from the relative path to the project, not the global paths of your computer.**

According to the Merriam-Webster online dictionary, to "plagiarize" means:

- to steal and pass off (the ideas or words of another) as one's own
- to use (another's production) without crediting the source
- to commit literary theft
- to present as new and original an idea or product derived from an existing source

In other words, **Plagiarism** is an act of fraud. It involves both stealing someone else's work and lying about it afterwards. **A copy of the practical implies a mark of 0 for both the group that makes the copy and the group who passed the code (if it exists).**

2 Data sets

Below, you will find a table that shows in detail the data sets that you can use in this work. All these data sets are obtained from the UCI machine learning repository. First column describes the name of the domain or data set. Next columns show #Cases = Number of cases or instances in the data set, #Num. = Number of numeric attributes, #Nom. = Number of nominal attributes, #Cla. = Number of classes, Dev.Cla. = Deviation of class distribution, Maj.Cla. = Percentage of instances belonging to the majority class, Min.Cla. = Percentage of instances belonging to the minority class, MV = Percentage of values with missing values (it means the percentage of unknown values in the data set). When the columns contain a '-', it means a 0. For example, the Glass data set contains 0 nominal attributes and it is complete as it does not contain missing values.

Domain	#Cases	#Num.	#Nom.	#Cla.	Dev.Cla.	Maj.Cla.	Min.Cla.	MV
<i>Adult</i>	48,842	6	8	2	26.07%	76.07%	23.93%	0.95%
<i>Audiology</i>	226	-	69	24	6.43%	25.22%	0.44%	2.00%
<i>Autos</i>	205	15	10	6	10.25%	32.68%	1.46%	1.15%
* <i>Balance scale</i>	625	4	-	3	18.03%	46.08%	7.84%	-
* <i>Breast cancer Wisconsin</i>	699	9	-	2	20.28%	70.28%	29.72%	0.25%
* <i>Bupa</i>	345	6	-	2	7.97%	57.97%	42.03%	-
* <i>cmc</i>	1,473	2	7	3	8.26%	42.70%	22.61%	-
<i>Horse-Colic</i>	368	7	15	2	13.04%	63.04%	36.96%	23.80%
* <i>Connect-4</i>	67,557	-	42	3	23.79%	65.83%	9.55%	-
<i>Credit-A</i>	690	6	9	2	5.51%	55.51%	44.49%	0.65%
* <i>Glass</i>	214	9	-	2	12.69%	35.51%	4.21%	-
* <i>TAO-Grid</i>	1,888	2	-	2	0.00%	50.00%	50.00%	-
<i>Heart-C</i>	303	6	7	5	4.46%	54.46%	45.54%	0.17%
<i>Heart-H</i>	294	6	7	5	13.95%	63.95%	36.05%	20.46%
* <i>Heart-Statlog</i>	270	13	-	2	5.56%	55.56%	44.44%	-
<i>Hepatitis</i>	155	6	13	2	29.35%	79.35%	20.65%	6.01%
<i>Hypothyroid</i>	3,772	7	22	4	38.89%	92.29%	0.05%	5.54%
* <i>Ionosphere</i>	351	34	-	2	14.10%	64.10%	35.90%	-
* <i>Iris</i>	150	4	-	3	-	33.33%	33.33%	-
* <i>Kropt</i>	28,056	-	6	18	5.21%	16.23%	0.10%	-
* <i>Kr-vs-kp</i>	3,196	-	36	2	2.22%	52.22%	47.78%	-
<i>Labor</i>	57	8	8	2	14.91%	64.91%	35.09%	55.48%
* <i>Lymph</i>	148	3	15	4	23.47%	54.73%	1.35%	-
<i>Mushroom</i>	8,124	-	22	2	1.80%	51.80%	48.20%	1.38%
* <i>Mx</i>	2,048	-	11	2	0.00%	50.00%	50.00%	-
* <i>Nursery</i>	12,960	-	8	5	15.33%	33.33%	0.02%	-
* <i>Pen-based</i>	10,992	16	-	10	0.40%	10.41%	9.60%	-
* <i>Pima-Diabetes</i>	768	8	-	2	15.10%	65.10%	34.90%	-
* <i>SatImage</i>	6,435	36	-	6	6.19%	23.82%	9.73%	-
* <i>Segment</i>	2,310	19	-	7	0.00%	14.29%	14.29%	-
<i>Sick</i>	3,772	7	22	2	43.88%	93.88%	6.12%	5.54%
* <i>Sonar</i>	208	60	-	2	3.37%	53.37%	46.63%	-
<i>Soybean</i>	683	-	35	19	4.31%	13.47%	1.17%	9.78%
* <i>Splice</i>	3,190	-	60	3	13.12%	51.88%	24.04%	-
* <i>Vehicle</i>	946	18	-	4	0.89%	25.77%	23.52%	-
<i>Vote</i>	435	-	16	2	11.38%	61.38%	38.62%	5.63%
* <i>Vowel</i>	990	10	3	11	0.00%	9.09%	9.09%	-
* <i>Waveform</i>	5,000	40	-	3	0.36%	33.84%	33.06%	-
* <i>Wine</i>	178	13	-	3	5.28%	39.89%	26.97%	-
* <i>Zoo</i>	101	1	16	7	11.82%	40.59%	3.96%	-

3 Report guidelines

I believe that it will be of great help for you some general **guidelines** for writing the report. It is not a complete list, it contains some comments and suggestions that you may consider in the assignments.

First of all, include a **front cover** with the name and surnames of the group members and a title of the work (this front cover is not part of the length of the report). The font size should be 11 or 12, not smaller and recall that figures should be also large enough for easier readability.

Analyze the different parameters and use tables or plots to show the results of your evaluation, and justify your decision of the final parameters. Not just saying we have tested several parameters and the best one is X or Y.

Additionally, it is important to **justify your findings**, not to just plot the graph with no comments on it about your observations and your judgement of the behavior of the algorithm in your data. Recall that when you add a comment on a plot, you should also link the comment to the figure/table/plot you are talking about. For example, as shown in Figure X or this result indicates that ... (see Figure X).

For the reader of a report, it is difficult to compare graphs if they contain different ranges in their axes and if they are placed in different pages (i.e., more than one page in the middle). It is supposed that **your findings are based on the experiments and the results obtained**. You cannot extract a conclusion if your results do not support it.

Presenting the results in isolation for every one of the datasets helps you to **fix the parameters** and extract conclusions considering the properties of a particular dataset. However, an **overall evaluation** of the methods/algorithms tested over the different datasets is also important to denote the applicability of the method/algorithm to different domains.

If you have read and implemented improvements on the basic algorithms, please add the comment and the references that justify your decision too. The report should contain a section with **references** or bibliography. Remember to add references at the end of the report. (references are not part of the length of the report)

Apart from showing your results and describing them, you **should also answer the questions that are requested in the description of the work**.

Conclusions in a report are important, please, remember to add them in your reports. Not only the general conclusions about what you have done, the conclusions of your findings and your observations of the results.