Introduction to Machine Learning Work 1 Clustering exercise

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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 the clustering algorithms using **Python 2.7**.

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 four tasks:

Make or use a parser to read the .arff file in Python and save the information in a matrix. In the following address:

https://docs.scipy.org/doc/scipy-0.19.1/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 <code>scipy.io</code> library in the abovementioned link, execute it, and modify it accordingly to your needs. For the numerical datasets, you can transform the continuous variables/attributes into categorical variables using function <code>pandas.qcut</code>.

- 1. 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.
- 2. Implement one of the following clustering algorithms: Bisecting K-Means (BKM) algorithm or a K-Harmonic Means (KHM) algorithm, and apply it to the data of the file.
- 3. Implement one of the following fuzzy clustering algorithms: Fuzzy C-Means (FCM) Clustering (Bezdek, 1981), Possibililistic C-Means (PCM) Clustering (Krishnapuram Keller, 1993), or Fuzzy Possibililistic C-Means (FPCM) (N. Pal K. Pal Bezdek, 1997).
- 4. Analyze the algorithms in three data sets (see Section 2). 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 implement and analyze K-Means, Bisecting K-Means or K-Harmonic Means 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 folder in Racó.

The idea is that you implement **your own code in Python 2.7** and you will use it to produce the results of the analysis.

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?
- 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, Bisecting K-Means or K-Harmonic Means, which has been the best K value? Have you implemented any improvement on the basic algorithms? For example, you can introduce a performance measure to decide which the best K value is. Another example, in K-Means++ it is defined an algorithm for choosing the initial values (or "seeds") for the k-means clustering algorithm.
- 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 2.7 in Racó in a zip file by November, 5th 2017.

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.
- Reason each one of the questions defined above in your evaluation and add any comment or observation that you consider important from your results.
- Additionally, you should explain how to execute your code.

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