Instructions

Please submit your solution by **Tuesday**, **February 18 (11:59 PM PT)**. Submissions should be made on **Gradescope**. Please complete homework **individually**. Please include the code of your solutions in the submission with a write-up describing how to run the code.

You are allowed to use any third-party libraries.

You will need the following files for this Homework: iris.csv (available at http://tinyurl.com/dsc148-w24)

1. DBSCAN Algorithm (10 points):

Consider the following figure:

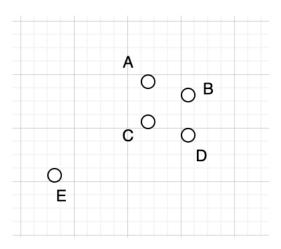


Figure 1: Points

There are 5 points: A, B, C, D, E. The distance matrix between these points is as follows:

	A	В	С	D	Е
A	0	2	2	5	6
В	2	0	2	3	7
\mathbf{C}	2	2	0	3	6
D	5	3	3	0	7
Ε	6	7	6	7	0

Table 1: Distance matrix

If we cluster the above points using the DBSCAN algorithm with $\epsilon = 4$ and minimum points = 3,

- (a) (5 points) How many clusters are formed? Draw the outline of your clusters. Explain your reasoning.
- (b) (5 points) With respect to the above figure, state one advantage of DBSCAN over k-means algorithm.

2. Association Rule Mining (20 points): Consider the following table:

$\overline{\mathbf{Id}}$	Movies watched		
1	Titanic, A star is born, Crazy Rich Asians		
2	Titanic, Inception, Crazy Rich Asians		
3	Titanic, Crazy Rich Asians, Avatar, Iron Man		
4	A star is born, Inception, Crazy Rich Asians, Avengers		
5	A star is born, Inception, Crazy Rich Asians, Avatar, Avengers		

Table 2: Movies

- (a) (10 points) Find all frequent patterns (i.e., movie combinations) whose support ≥ 0.5 . **Hint:** A single movie is also considered a combination.
- (b) (10 points) Find all the rules (X -> Y) (s, c) where s represents support and c represents confidence such that $s \ge 0.5$, $c \ge 0.6$.
- 3. **PCA (20 points):** Consider the following two plots. These are plots of training data points X in \mathbb{R}^2 belonging to 2 classes.

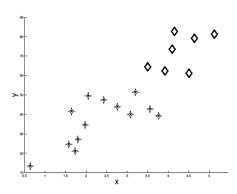


Figure 2: Dataset-1

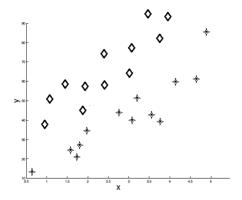


Figure 3: Dataset-2

Answer the following questions for each dataset.

- (a) (10 points) Draw all two principal components in the picture (you can take a screenshot). You are expected to draw the rough directions of the principal components instead of accurate computations. **Hint**: There are 2 principal components for each dataset.
- (b) (10 points) After projecting all the points onto one of the principal components, is it possible to correctly classify all the points by just a threshold function? If yes, which principal component should we project onto and why? If no, please explain your reasoning.

Hint: Classifying with respect to a threshold function means you just need a threshold to classify them. For example,

```
\begin{array}{l} \textbf{if} \ x < 0.64 \ \textbf{then} \\ \text{predict label-1} \\ \textbf{else} \\ \text{predict label-0} \\ \textbf{end if} \end{array}
```

Here 0.64 is considered as the threshold.

4. PCA, k-Means and GMM clustering (50 points):

In this question, we will using the Iris dataset to predict Species of the iris plant. ('iris.csv')

- (a) (5 points) In the data preparation step do the following:
 - (i) Split data into features (X) and label (y). Our label is the column Species and features include all the other columns except Species and Id.
 - (ii) Standardize the features (X_standardized) by removing the mean (i.e mean=0) and scaling to unit variance. (Hint: use sklearn.preprocessing.StandardScaler())
- (b) (15 points) Project the 4-dimensional standardized data (X_standardized) onto 2 dimensions using PCA (sklearn.decomposition.PCA()). Visualize the scatterplot of the first two principal components of the data. In the scatterplot assign each data point a color based on its species with the following dictionary:

```
{'Iris-setosa': 'r', 'Iris-versicolor': 'g', 'Iris-virginica': 'b'}.
```

- (c) (10 points) Cluster the 4-dimensional standardized data (X_standardized) into 3 clusters using GMM clustering. Score the clustering accuracy with sklearn.metrics.cluster.adjusted_rand_score(). (Hint: use sklearn.mixture.GaussianMixture())
- (d) (10 points) Cluster the 4-dimensional standardized data (X_standardized) into 3 clusters using K-means clustering. Score the clustering accuracy with sklearn.metrics.cluster.adjusted_rand_score(). (Hint: use sklearn.cluster.KMeans())
- (e) (10 points) Briefly compare the result from part (c) and part (d). Explain why Gaussian Mixture algorithm performs better than k-Means algorithm.

Suggested reading: http://scikit-learn.org/stable/modules/generated/sklearn.metrics.adjusted_rand_score.html