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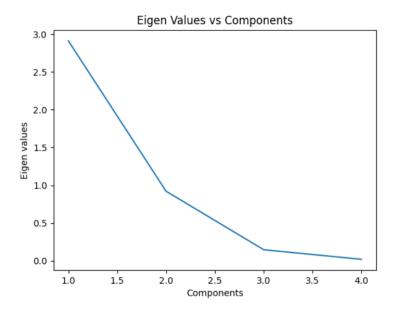


Figure 1 Eigenvalue vs. components

- 1. The eigenvalue decrease corresponding to each component increasing successively.
- 2. Because attributes are more dependent on first eigen vector so it has more spread around 1st eigen vector.



2 a.

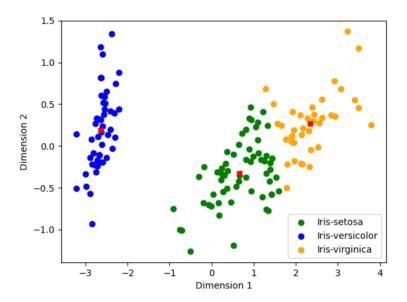


Figure 2 K-means (K=3) clustering on Iris flower dataset

- 1. The clustering looks quite accurate forming good clusters as K-Means is an unsupervised algorithm.
- 2. K-means algorithm assumes cluster boundaries to be circular in 2D. From the output, the boundary doesn't seem to be clearly circular.
- **b.** The value for distortion measure is 63.8738
- c. The purity score after examples are assigned to the clusters is 0.887



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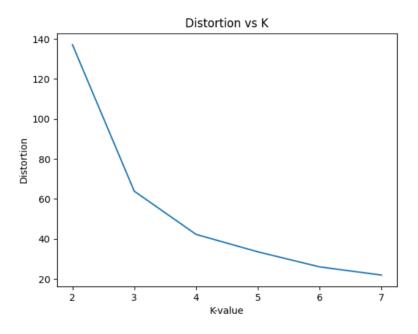


Figure 3 Number of clusters(K) vs. distortion measure

- 1. The distortion measure decreases with an increase in K.
- 2. Distortion measure decreases drastically for k=2 to k=3 then very gradually because our data has 3 species which also indicate optimal K value.
- 3. Number of species in data set is 3, So intuitively k=3 should be the number of optimum clusters. The elbow and distortion measure plot closely follow the intuition.

Table 1 Purity score for K value = 2,3,4,5,6 & 7

K value	Purity score
2	0.667
3	0.887
4	0.687
5	0.673
6	0.513
7	0.513



Inferences:

- 1. The highest purity score is obtained with K = 3.
- 2. Purity score increases from k=2 to k=3 and then decreases for further K values.
- 3. Because the number of species in our data is 3 so purity score is highest for k=3.
- 4. Except k=3, distortion measures decreases with increase in purity score.

4 a.

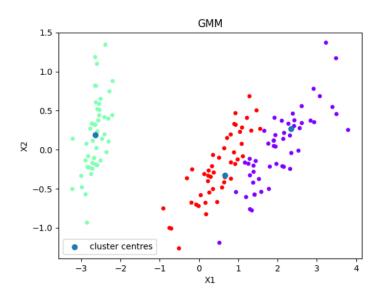


Figure 4 GMM (K=3) clustering on Iris flower dataset

- 1. Predicted results are very close to actual ones that's why GMM looks quite accurate.
- 2. GMM algorithm assumes cluster boundaries to be elliptical in 2D. From the output, the boundary seems little bit elliptical.
- 3. Yes, from the graphs we can see that the boundaries in K-means were circular while in GMM they are elliptical.
- b. The value for distortion measure is -280.87
- c. The purity score after examples are assigned to the clusters is 0.98



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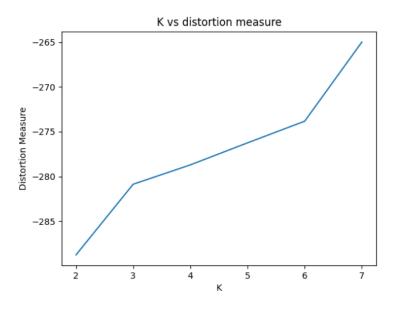


Figure 5 Number of clusters(K) vs. distortion measure

- 1. The distortion measure increase with an increase in K.
- 2. Because there are three species, the distortion measure has a steeper slope between k=2 and k=3, then progressively grows until k=6, then abruptly increases beyond that.
- 3. From the number of species in the given dataset, intuitively k=3 should be the optimal value of k. The elbow method also follow the intuition.

Table 2 Purity score for K value = 2,3,4,5,6 & 7

K value	Purity score
2	0.667
3	0.98
4	0.833
5	0.767
6	0.64
7	0.627



Inferences:

- 1. The highest purity score is obtained with K = 3.
- 2. Purity score increases for k=2 to k=3 then decreases for k=2 to k=7.
- 3. Because the number of species in our data is 3 so purity score is highest for k=3.
- 4. Except k=3, distortion measure decreases with increase in purity score.
- 5. By inferences, we can say that GMM is more accurate than K-means.

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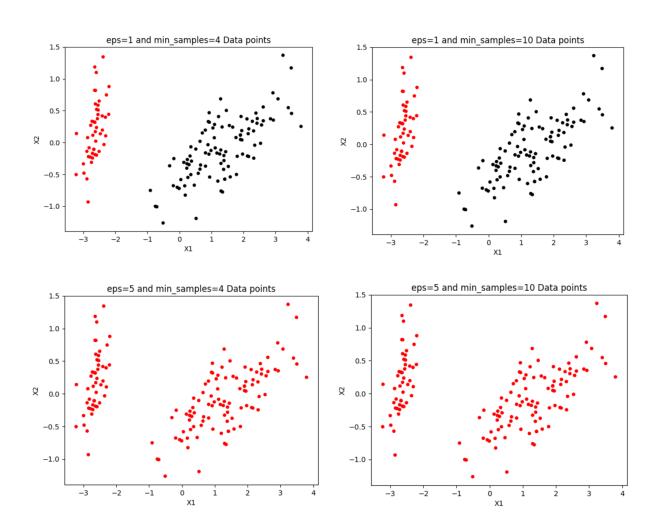




Figure 6 DBSCAN clustering on Iris flower dataset

Inferences:

- 1. Due to our eps values, accuracy is not that good.
- 2. The number of clusters are less than both K-means and GMM and boundaries are neither circular nor elliptical in DBSCAN.

b.

Eps	Min_samples	Purity Score
1	4	0.667
	10	0.667
5	4	0.333
	10	0.333

Inferences: 0.333

- 1. For the same eps value, increasing min_samples doesn't change purity score.
- 2. For the same min_samples, increasing eps value decrease purity score.