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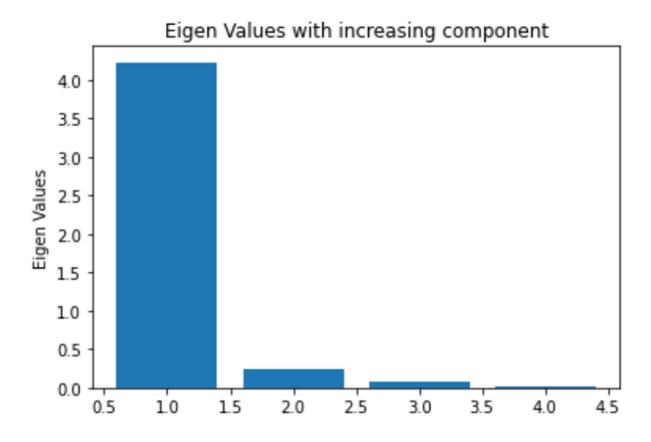


Figure 1 Eigenvalue vs. components

- 1. With each component increase, the eigenvalue decreases. Until component = 2, there is a significant drop. Following that, the reduction is less.
- 2. This is to be expected, given eigenvalues are a measure of variance along the principal components, and PCA is done in such a way that data variance diminishes as principal components are added.



2 a.

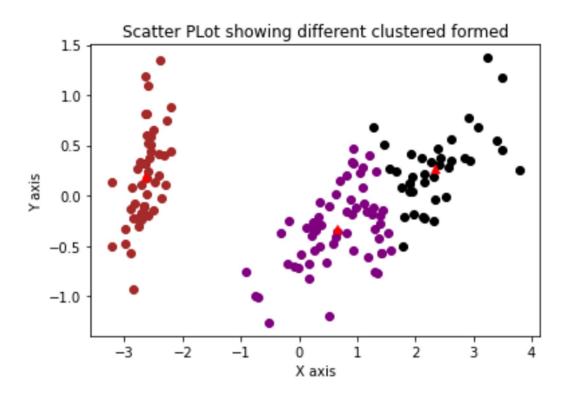


Figure 2 K-means (K=3) clustering on Iris flower dataset

- 1. The clusters appear to be well defined and circular in shape, with significant densities of datapoints in each cluster centre.
- 2. Yes, the boundaries seem to be circular.
- **b.** The value for distortion measure is 63.874
- c. The purity score after examples are assigned to the clusters is 88.7%



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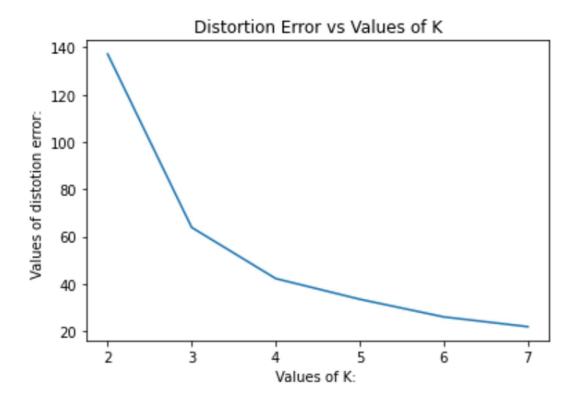


Figure 3 Number of clusters(K) vs. distortion measure

- 1. The distortion measure decreases with an increase in K.
- 2. More centers are developed when more clusters are formed. Because the distortion measure is the total of the distance between cluster points and their centers, it decreases as K increases.
- 3. We can see from the graph that K=3 is the best value for best results when using the elbow method. And the outcome is exactly what we expected, with K=3 having the greatest purity score of all the Ks.

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

K value	Purity score
2	66.667
3	88.667
4	68.667
5	67.333
6	50.667
7	50.667



Inferences:

- 1. The highest purity score is obtained with K =3
- 2. Purity initially increases as the k value rises. Then it starts to go downhill. At the ideal k value, purity is at its greatest.
- 3. This is because as we get closer to the actual number of clusters in the dataset ie.3, more datapoints are correctly classified, and purity score rises; however, once we reach the maximum value at K=3, we start making more clusters than are actually present in the dataset, and thus begin incorrectly classifying datapoints, lowering our purity score.
- 4. They have a similar tendency once the purity score's k value reaches its maximum k; both begin to decline after that. In addition, the elbow point of distortion measurement yields the k value, which corresponds to the highest purity score.

4 a.

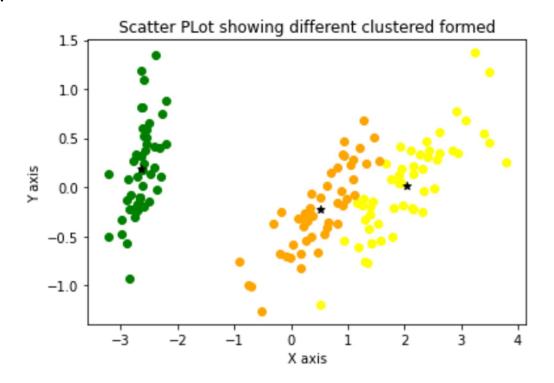


Figure 4 GMM (K=3) clustering on Iris flower dataset



Inferences:

- 1. The clusters formed in the above plot are elliptical in shape.
- 2. GMM algorithm assumes cluster boundaries to be elliptical in 2D. From the output, the boundary seems to be elliptical.
- 3. The GMM has elliptical cluster boundaries while the K means has circular.
- 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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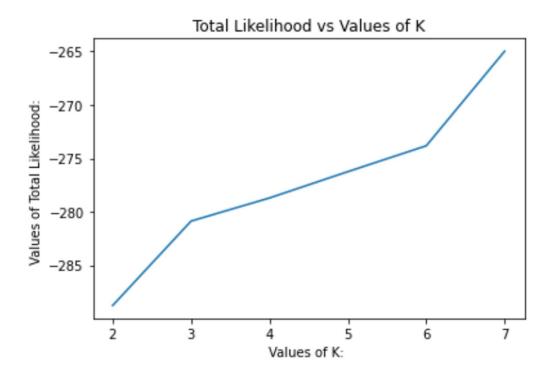


Figure 5 Number of clusters(K) vs. distortion measure

Inferences:

1. The distortion measure increase with an increase in K. Justify the observed trend.



- 2. As the number of species in the given dataset is, we can intuitively say that optimum clusters is
- 3. Yes, the elbow method allows us to follow intuition.

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

K value	Purity score
2	66.67
3	98
4	83.33
5	76.67
6	64.0
7	62.27

- 1. The highest purity score is obtained with K = 3
- 2. Purity initially increases as the k value rises. Then it starts to go downhill. At the ideal k value, purity is at its greatest.
- 3. This is because, as we get closer to the actual number of clusters in the dataset, a greater number of datapoints are correctly classified, and the purity score rises; however, once we reach the maximum value at K=3, we begin to make more clusters than are actually present in the dataset, and thus begin incorrectly classifying datapoints, lowering our purity score.
- 4. They have a similar tendency once the purity score's k value reaches its maximum k; both begin to decline after that. In addition, the elbow point of distortion measurement yields the k value, which corresponds to the highest purity score.
- 5. The GMM model appears to better suit the data than the K-Means model. This is to be expected, given that GMM is a probabilistic soft classifier.



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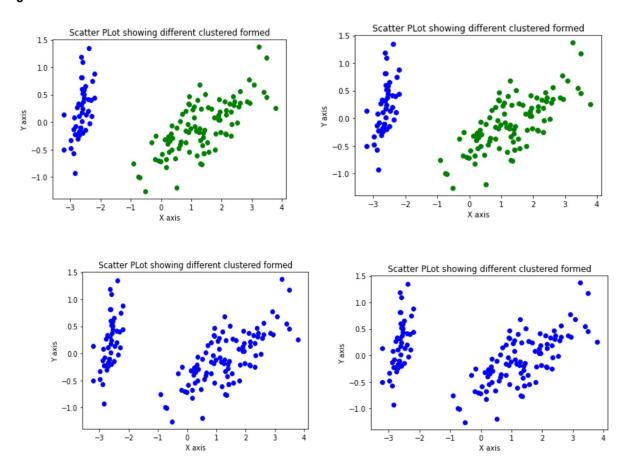


Figure 6 DBSCAN clustering on Iris flower dataset

Inferences:

- 1. We are not getting a good accuracy the reason maybe that we are not taking an appropriate value of radius and Min_sample.
- 2. The DBCSAN is not able to separate the two clusters in the right as they don't seem to have a common point where their frequency differ. GMM has elliptical boundary, K means has circular and DBSCAN's boundary can take any shape.

b.



Eps	Min_samples	Purity Score
1	4	66.67
	10	66.67
4	4	33.33
	10	33.33

Inferences:

- 1. Min_samples doesn't affect purity scores value.
- 2. For the same min_samples, increasing eps value decreased the purity score

THANK YOU