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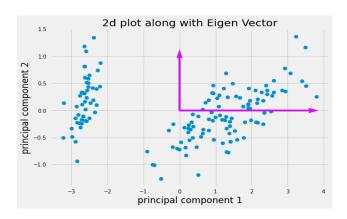


Figure 1 Eigenvalue vs. components

### Inference

1. Eigen values are decreasing corresponding to each component increase or decrease successively. Since the eigen values contains the characteristic of whole data then therefore eigen value(principle comp 1) > eigen value(principle comp 2) > eigen value(principle comp 3) >......> eigen value(principle comp n)

## 2 a.

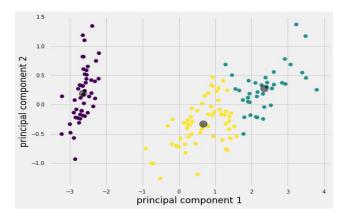


Figure 2 K-means (K=3) clustering on Iris flower dataset



#### Inferences:

- 1. From the plot we can see that any point is at minimum distance with respect to allotted centre than the other two centers.
- 2. No here it does not seem to be. Because it only searches at a perticluar distance over all the direction from the centers. So it may not possible always such that all point lie on circumference.
- **b.** The value for distortion measure is 63.87.
- c. The purity score after examples are assigned to the clusters is 0.88.

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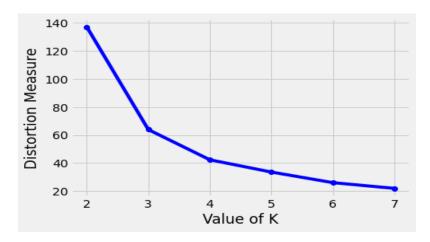


Figure 3 Number of clusters(K) vs. distortion measure

- 1. Distortion measure decreases with an increase in K.
- 2. If the number of cluster increased than it the model will have more number of choices to get fit into.
- 3. From the plot we can take as K = 3 or 4 for more precise model with help of elbow method.

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

K value	Purity score
2	0.66
3	0.88
4	0.84
5	0.90
6	0.89
7	0.96



#### Inferences:

- 1. The highest purity score is obtained with K = 7.
- 2. Increasing the value of K it increases the purity score.
- 3. If the number of cluster increased than it the model will have more number of choices to get fit into.
- 4. Yes as the distortion measure decreases then the purity score increases.

#### 4 a.

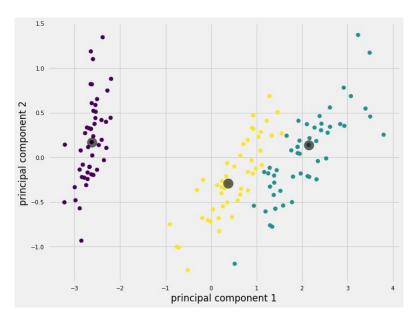


Figure 4 GMM (K=3) clustering on Iris flower dataset

- 1. From the plot we can see that any point is at minimum distance with respect to allotted centre than the other two centers
- 2. No here this is not elliptical.GMM only uses elliptical area only to find the number the point lying in a cluster.
- 3. GMM is much better in prediction than K-Means as purity score of GMM is higher.
- **b.** The value for log likelihood is -0.87.
- c. The purity score after examples are assigned to the clusters is 0.98



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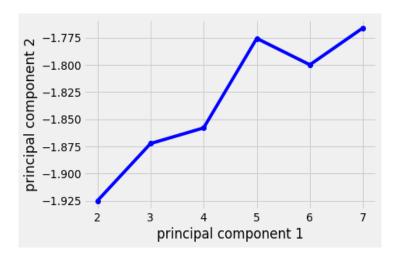


Figure 5 Number of clusters(K) vs. distortion measure

#### Inferences:

- 1. The distortion measure decreases with an increase in K.
- 2. As the more number of cluster is given then it become easier to divide the data.
- 3. The number of optimum clusters is 3 or 4.Yes since between 3 to 4 it converges as difference become negligible.

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

K value	Purity score
2	0.66
3	0.98
4	0.98
5	0.94
6	0.90
7	0.96

- 1. The highest purity score is obtained with K = .3 and 4
- 2. The increament is random on increasing the number of K.
- 3. As the distortion measure decreases, the purity score increases.
- 4. Since GMM is soft clustering techniques. Hence GMM has better purity score than K Means.



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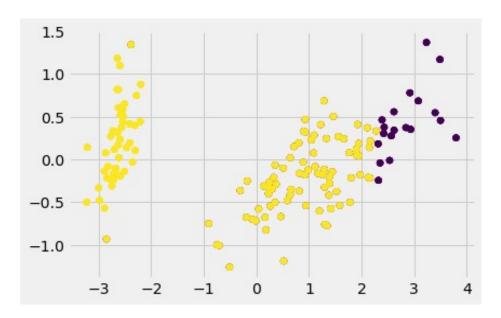


Figure 6 DBSCAN clustering on Iris flower dataset

### Inferences:

- 1. DBSCAN only take two cluster to divide the whole set of data.
- 2. Yes in GMM, we will have to give the prior information about number of clusters while that is not in the case of DBSCAN.

b.

Eps	Min_samples	Purity Score
1	4	0.66
	10	0.66
5	4	0.33
	10	0.33

- 1. For the same eps value, does increasing min\_samples does not really increases purity score.
- 2. For the same min\_samples, increasing eps value the purity score decreases.