

# KMEANS and Kernel KMEANS

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## Abstract

Kernel principal component analysis (kernel-PCA) is a prominent non-linear extension of one of the most used classical dimensionality reduction algorithms (PCA-Principal Component Analysis). In this paper, we present an inhaustive comparison between the classical PCA and the kernel-PCA. We use the moon dataset

## Keywords

Dimensionality reduction, PCA, kernel-PCA

## 1 INTRODUCTION

K-Means clustering is a fast, robust, and simple algorithm that gives reliable results when data sets are distinct or well separated from each other in a linear fashion. It is best used when the number of cluster centers, is specified due to a well-defined list of types shown in the data. However, it is important to keep in mind that K-Means clustering may not perform well if it contains heavily overlapping data, if the Euclidean distance does not measure the underlying factors well, or if the data is noisy or full of outliers

## 2 PARTITION CLUSTER-ING

### 2.1 KMEANS

### 2.2 Kernel KMEANS

## 3 EXPERIMENTAL RE-SULT

### 3.1 Dataset

### 3.2 Performance analysis

## 4 CONCLUSION

## References

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