

# Data mining



# Finding patterns in data: Dimensionality reduction

*Many algorithms*

PCA

Factor analysis

Independent component analysis (ICA)

Singular value decomposition (SVD – closely related to PCA)

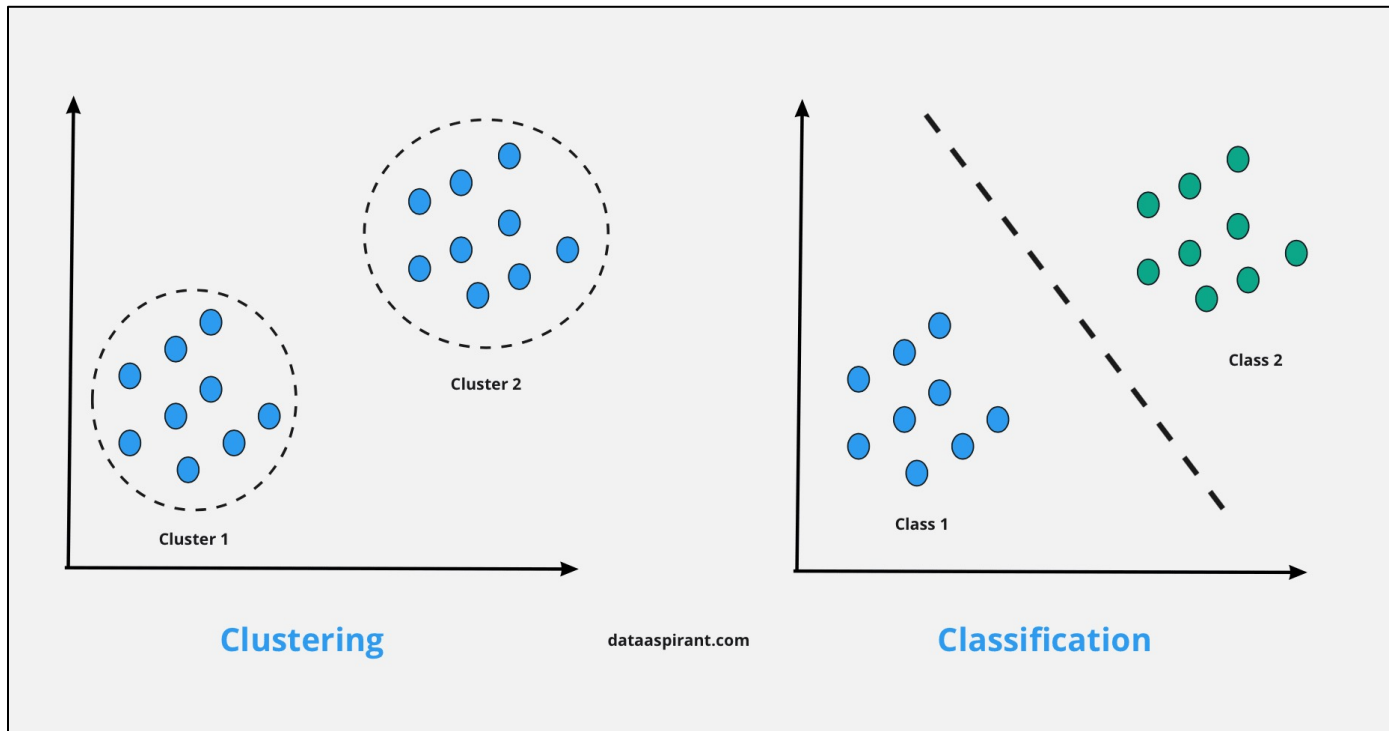
Non-negative matrix factorization (NMF)

demixed PCA (dPCA)

Linear discriminant analysis (LDA)

Bespoke statespace analyses

# Finding patterns in data: Clustering



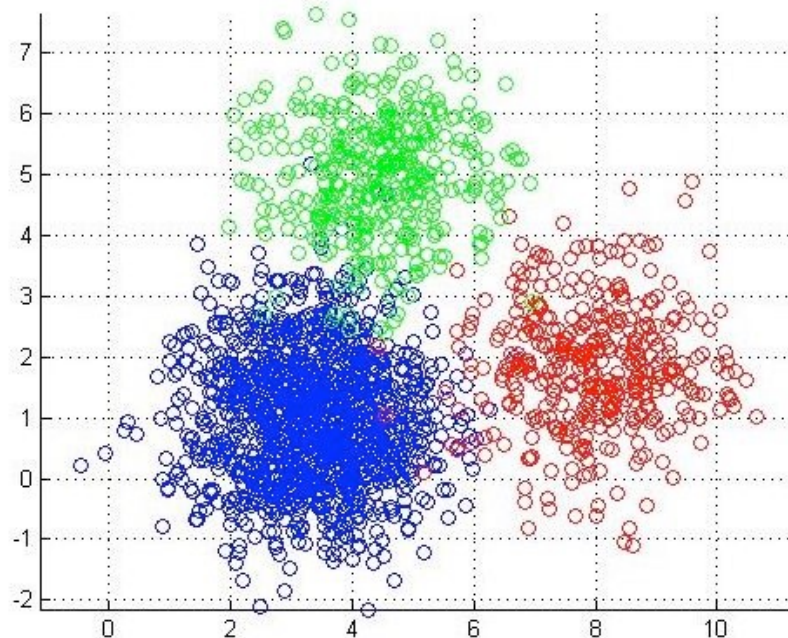
# Finding patterns in data: Clustering

*Even more algorithms!*

K-means

# K-means

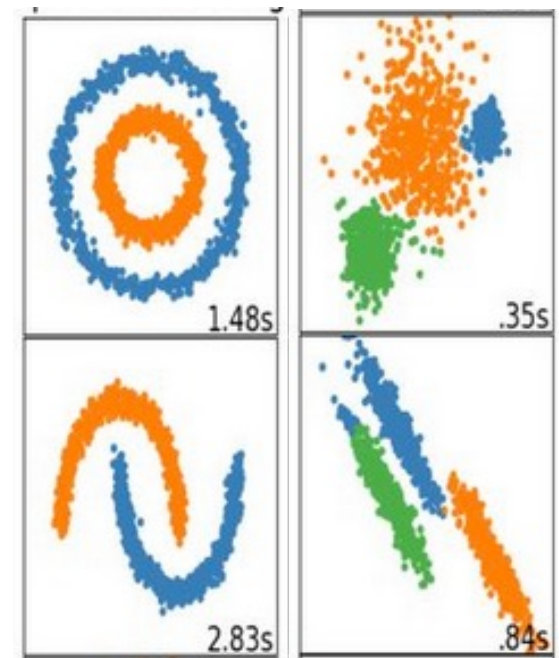
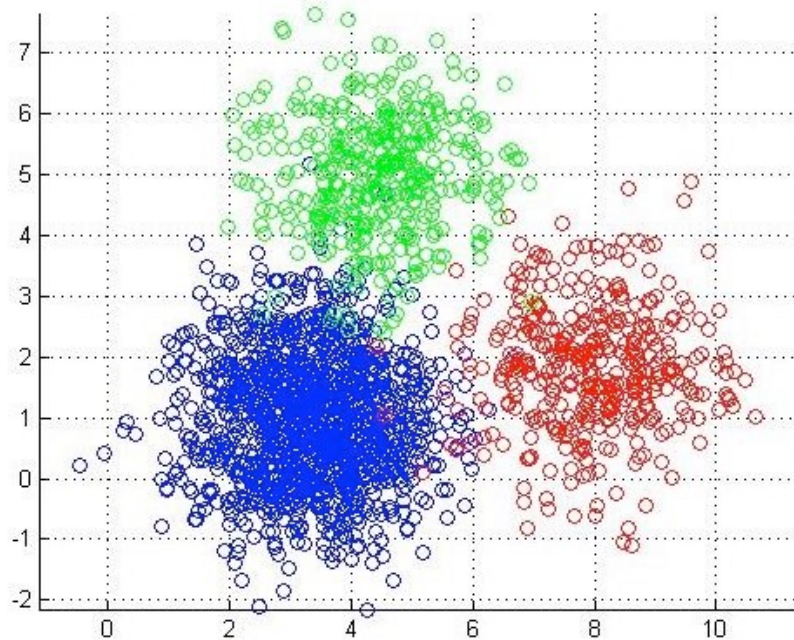
- $K$  = the number of clusters (set by the experimenter)
- Minimizes the total sum of squared distances from each point to its respective cluster center (in  $n$ -dimensional space)



# Finding patterns in data: Clustering

*Even more algorithms!*

K-means



# Finding patterns in data: Clustering

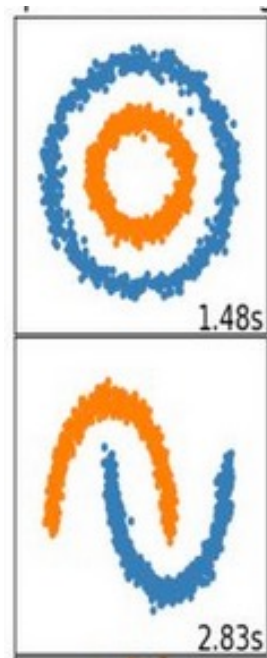
*Even more algorithms!*

K-means

Spectral Clustering – *group based on graph distances*

Density-Based Spatial Clustering of Applications with Noise (DBSCAN)

- *group based on distances between nearest points*



# Finding patterns in data: Clustering

*Even more algorithms!*

K-means

Spectral Clustering

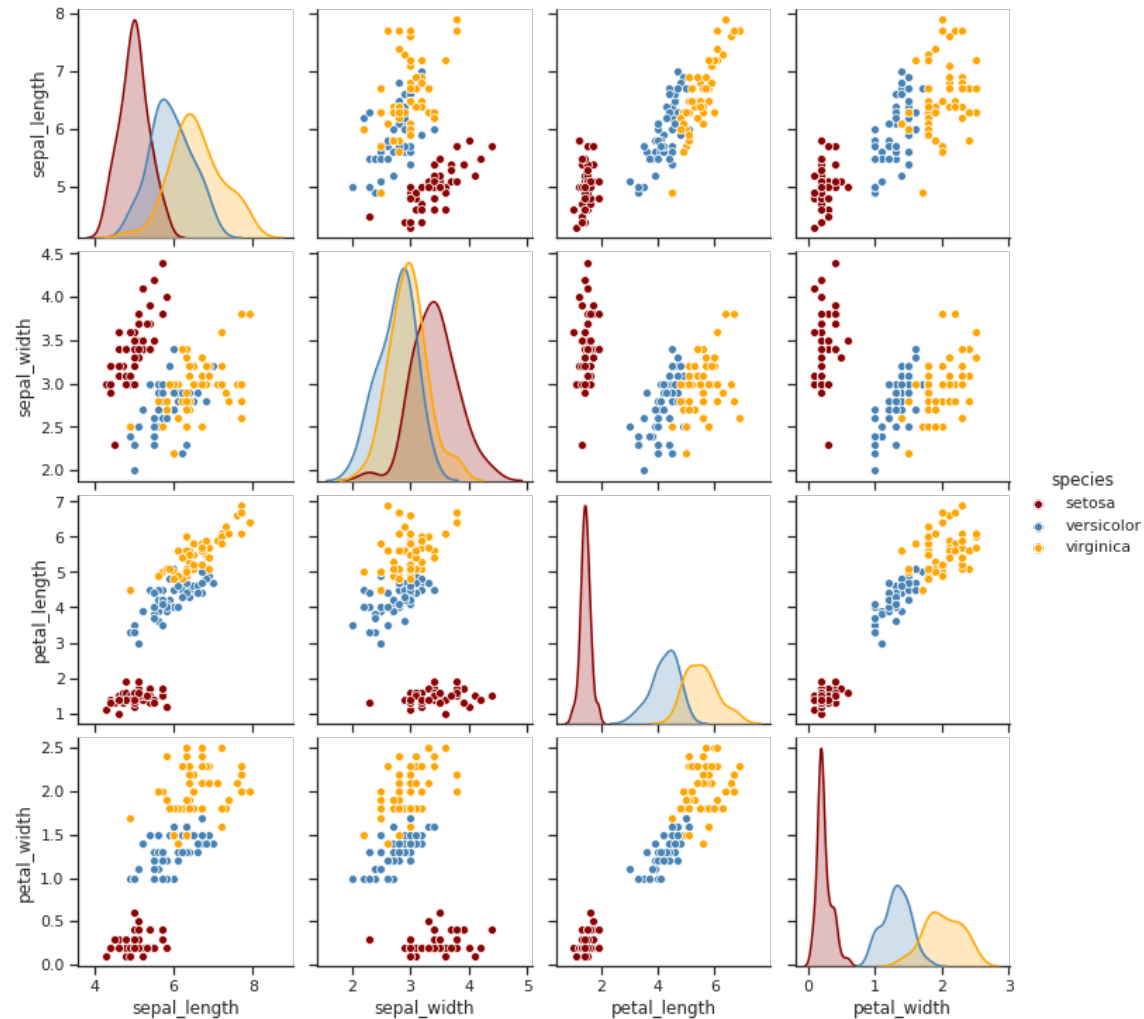
Density-Based Spatial Clustering of Applications with Noise (DBSCAN)

Gaussian Mixture Models (GMM) using Expectation Maximization (EM)



# Gaussian Mixture Models (GMM) using Expectation Maximization (EM)

- “soft” clustering (=assigns probabilities)
- tries to assign data to different Gaussian distributions



# Finding patterns in data: Clustering

*Even more algorithms!*

K-means

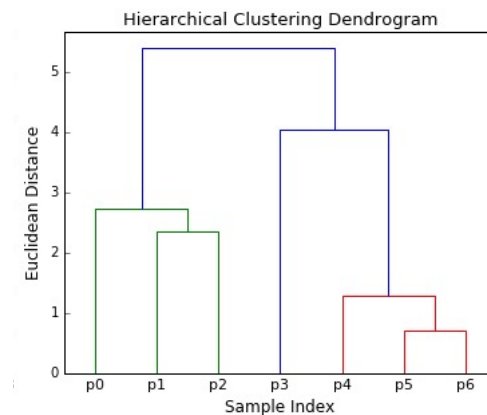
Spectral Clustering

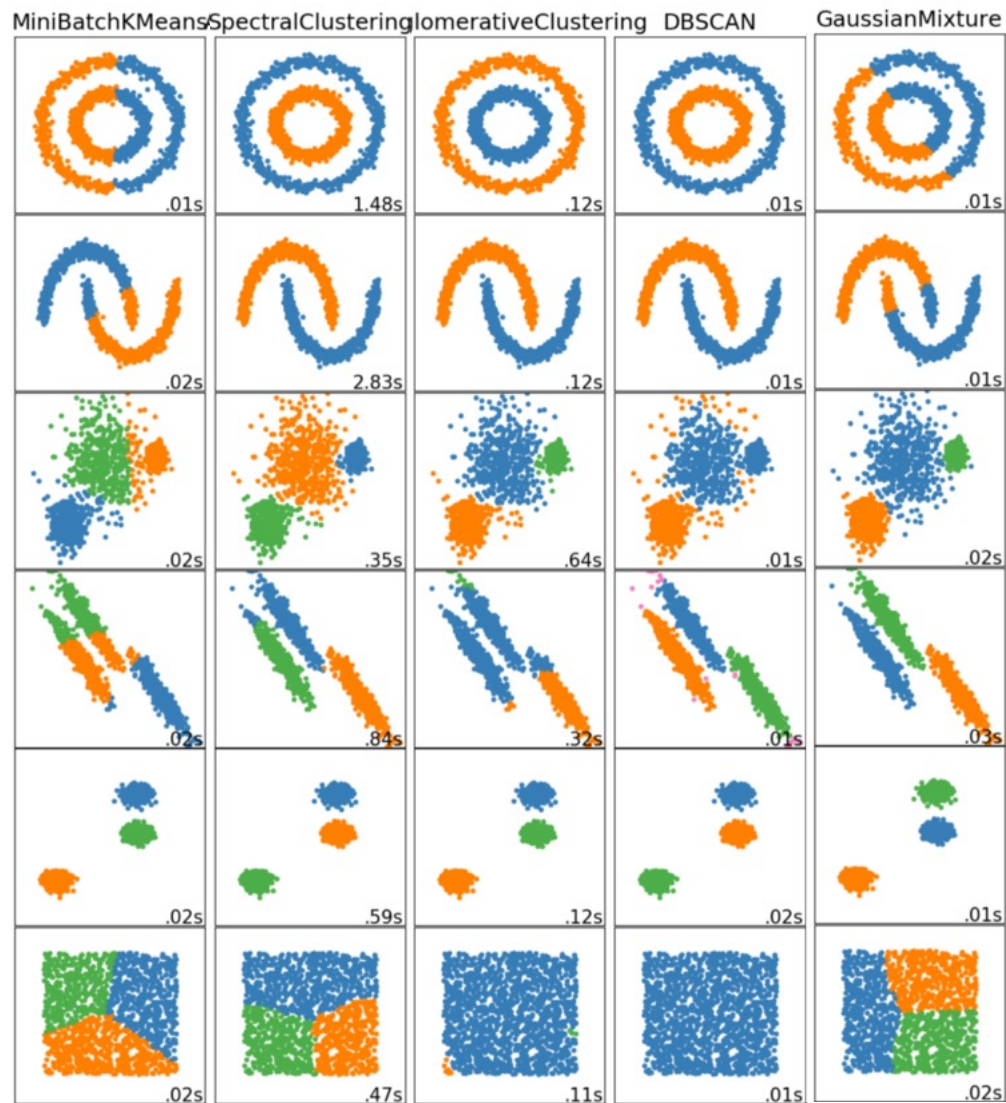
Density-Based Spatial Clustering of Applications with Noise (DBSCAN)

Gaussian Mixture Models (GMM) using Expectation Maximization (EM)

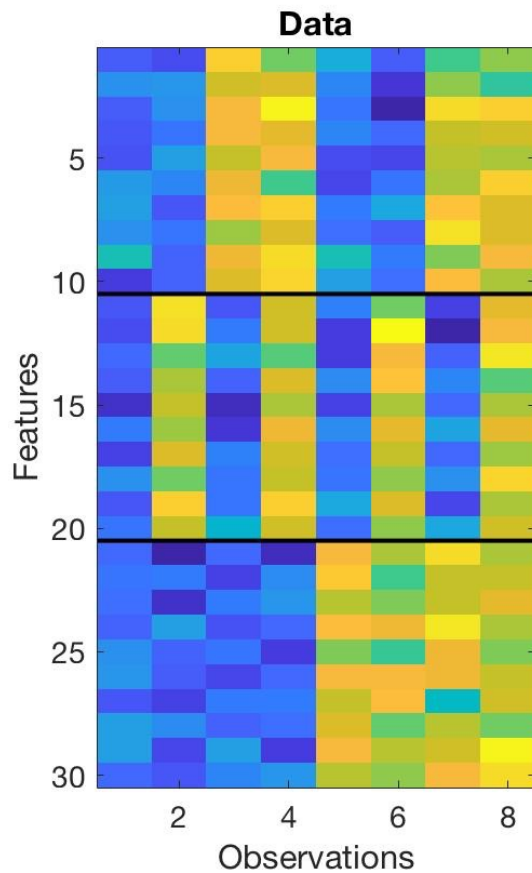
Hierarchical clustering algorithms

- “bottom up” (assign each point to a cluster -> successively merge)
- “top down” (start with one cluster -> split)





# K-means - demo



```
% Let's do it again with a more complex data set
% Additional patterns
pattern2 = [5 10 5 10 5 10 5 10];
pattern3 = [5 5 5 5 10 10 10 10];

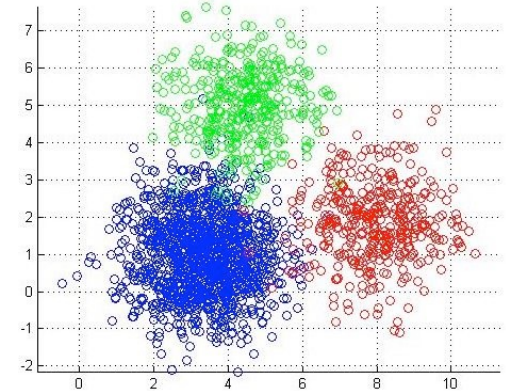
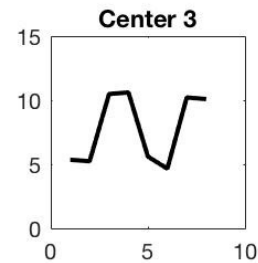
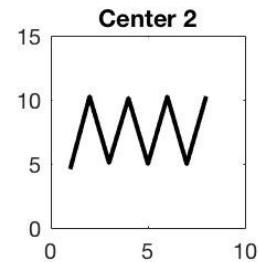
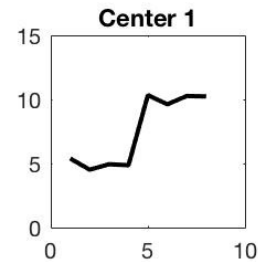
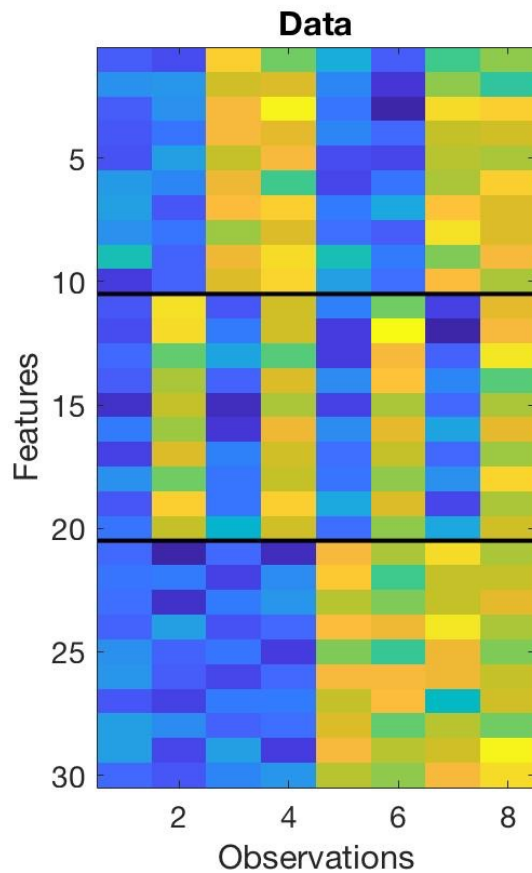
% and create two more populations that follow different patterns
pop2 = []; pop3 = [];
for k = 1:10
    for j = 1:8
        noise = normrnd(0,var); % noise should be independent for this simulation
        pop2(k,j) = pattern2(j)+noise;
        noise = normrnd(0,var);
        pop3(k,j) = pattern3(j) + noise;
    end
end
pop = [pop1;pop2;pop3]; % Our full feature matrix is all of these subpopulations together
```

# K-means - demo

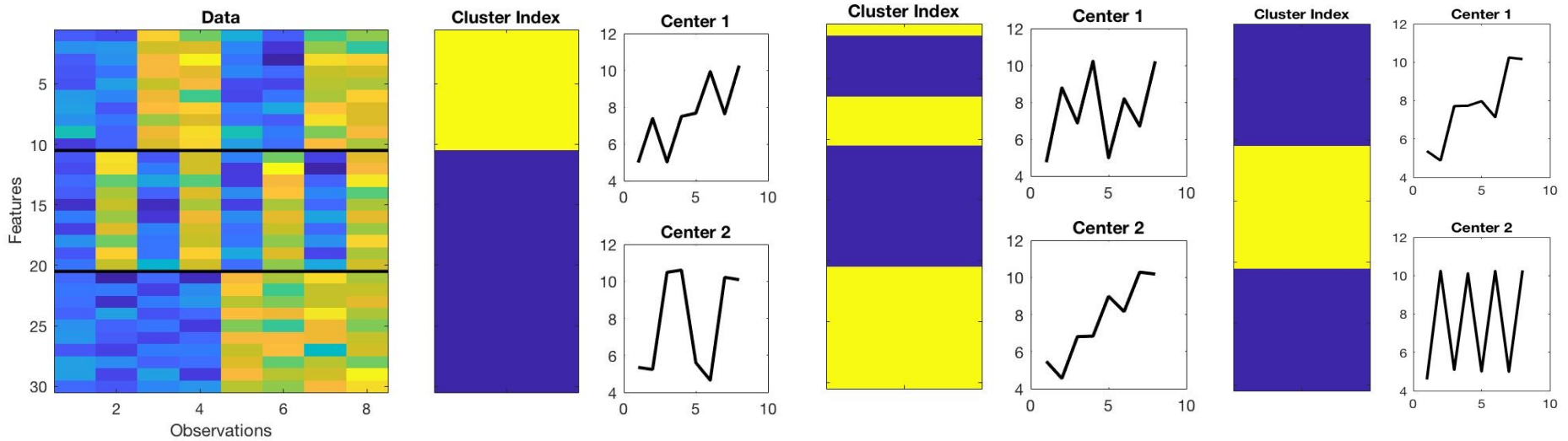
30 observations in 8-dimensions

Matlab: `[idx,centers] = kmeans(pop,3); %first run kmeans with 3 clusters`

R: `pop_cluster <- pop %>% kmeans(centers = 3)`



# K-means - demo



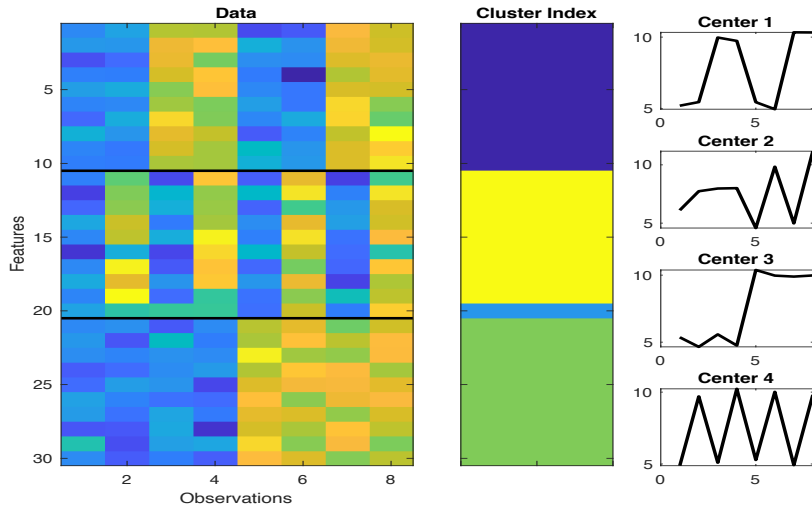
## Considerations:

K-means optimization can be inconsistent

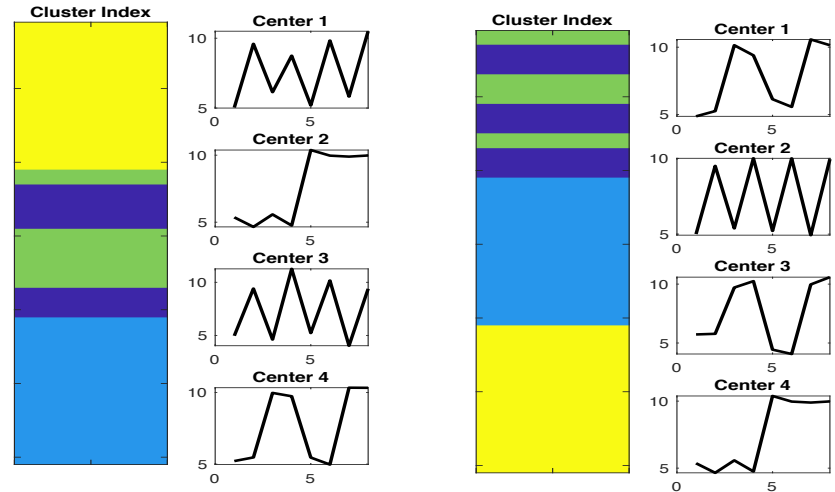
-> run with a random seed multiple times, find the global optimal solution

# K-means - demo

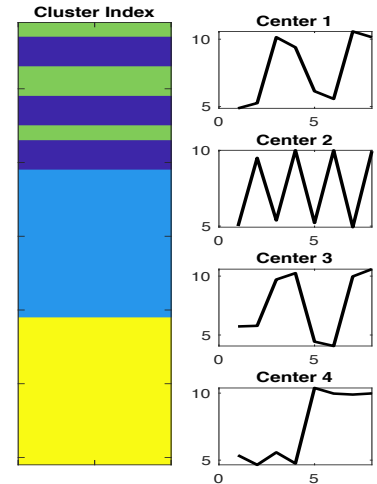
Run 1



Run 2



Run 3



## Considerations:

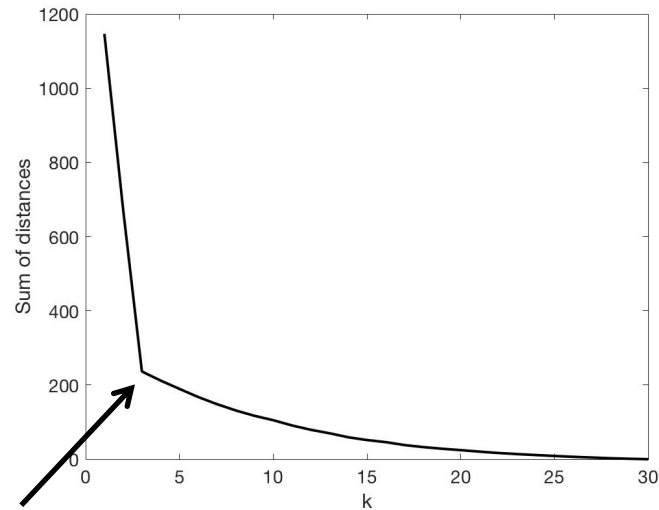
K-means optimization can be inconsistent

-> run with a random seed multiple times, find the global optimal solution

K is unknown

-> elbow method

Find K with the elbow method:

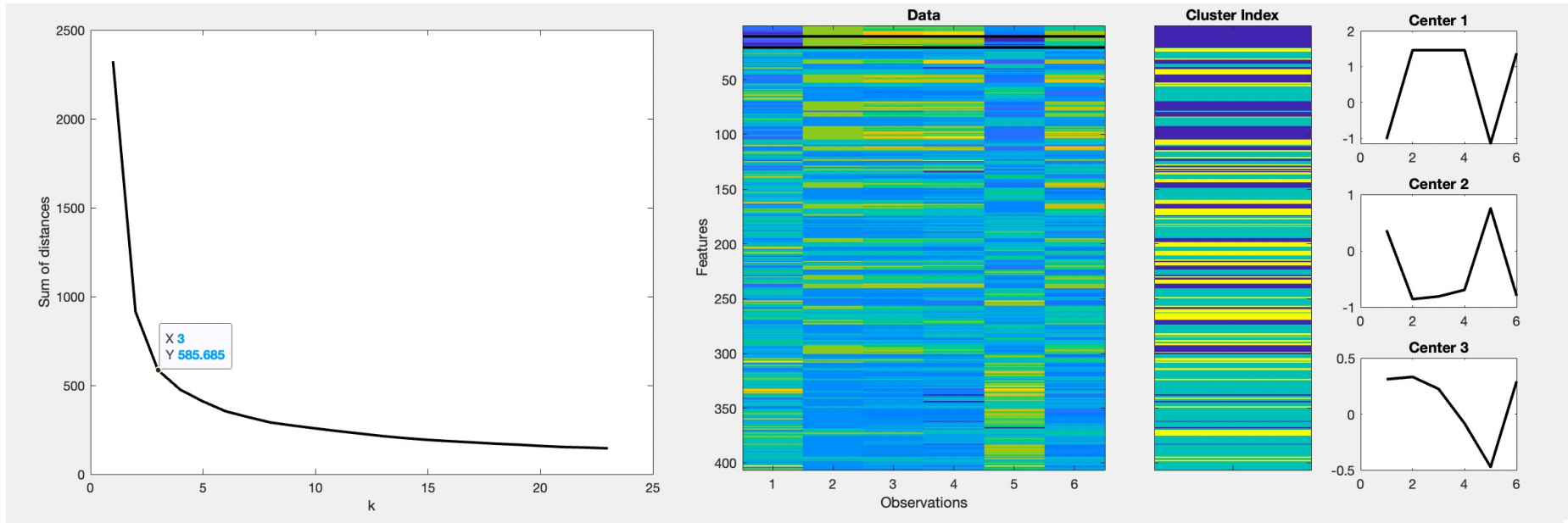


Elbow  
at k=3

```
numiter = 1000; %to find the optimal solution for each value of k, we'll need to rerun the algo
numk = length(pop(:,1)); %we'll test each value of k until we have 1 per feature (e.g. neuron)
distances = [];
for k = 1:numk
    d = [];
    for iter = 1:numiter
        [~,~,sumd] = kmeans(pop,k); %sumd is the sum of distances to each cluster center
        d(iter) = sum(sumd); %the total distance is the sum of sumd
    end
    distances(k) = min(d); %the minimum distance is the optimal solution for that k
end
```



# K-means



## Considerations:

K-means optimization can be inconsistent

-> run with a random seed multiple times, find the global optimal solution

K is unknown

-> elbow method

K-means struggles with irregular data (e.g. unequal numbers or variance)

# A note on model comparison

## The situation:

You have a lot of data, and want to find explanatory variables

## The problem:

Adding more variables will always add explanatory power

- But –
1. it may be a small improvement
  2. it may be *overfitting*

## What to do:

Formal model comparisons estimate model fit with penalties for increasing number of parameters:

*Akaike Information Criterion (AIC) & Bayesian Information Criterion (BIC)*

Also remember to hold out data when data mining!

# Homework #9

(second part)

## HW9: Data mining

You have recorded pupil responses in a subject viewing different images.

The data are saved in *data.txt*, which includes 500 trials where each trial is 1200ms long

1. Plot the mean pupil response over all trials
2. Do PCA across trials (Hint: each PC should be 1200 elements long, and there should be 500 of them).  
How much variance does the first PC account for? How many components account for  $\geq 90\%$  of the variance?
3. Plot the first principal component.
4. Run k-means 10 times with  $k=2$ . For each run, plot the cluster centers you obtain.