

Time Series Clustering with R

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Clustering Analysis (Data Segmentation)

• What it is ?

- It is an **unsupervised** machine learning technique (unlabeled data).
- Grouping data into different **clusters**.
- Minimizing dissimilarity **within** clusters or Maximizing dissimilarity **between** clusters.

• Applications

- Market/ Customers/ Product Segmentation.
- Recommendation Systems.
Ex: "The similar products you may also like"
"RECOMMENDED FOR YOU"
- Grouping Search Result.
Ex: [Yippy \(formerly Clusty\)](#)
[Topical Clustering of Search Results](#)

Yippy



clustering analysis

Search

Results 1-20 of about 60,800,077 | Details

Sources Sites Time Topics

Top 122 Results

- + Statistics (23)
- + Data Mining (12)
- + Software (13)
- + University (10)
- + Research (12)
- + Search (10)
- + Discriminant, Model (6)
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- Tutorial (5)
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- + Free (5)
- Exploration (4)
- Clustering Algorithms (3)
- Expression, Gene (4)
- Database (3)
- Failover (3)
- 2015 Internet Security Threat Report (2)
- GIS (2)
- Detection (2)
- Windows Clustering (2)
- County (2)
- Other Topics (28)

[Cluster analysis - Wikipedia](#) [new window](#) [preview](#)

Cluster **analysis** or **clustering** is the task of grouping a set of objects in such a way that objects in the same group (called a cluster) are more similar ...

https://en.wikipedia.org/wiki/Cluster_analysis - - Yippy Index V

[Cluster Analysis: Basic Concepts and Algorithms](#) [new window](#) [preview](#)

8 Cluster **Analysis**: Basic Concepts and Algorithms Cluster analysis divides data into groups (clusters) that are meaningful, useful, or both. If meaningful groups are the goal ...

<https://www-users.cs.umn.edu/~kumar/dmbook/ch8.pdf> - - Yippy Index V

[Conduct and Interpret a Cluster Analysis - Statistics.com](#) [new window](#) [preview](#)

The Cluster **Analysis** is an explorative **analysis** that tries to identify structures within the data. Cluster **analysis** is also called segmentation **analysis**.

www.statisticssolutions.com/cluster-analysis-2 - - Yippy Index V

[Quick-R: Cluster Analysis - statmethods.net](#) [new window](#) [preview](#)

Learn R functions for cluster **analysis**. This section describes three of the many approaches: hierarchical agglomerative, partitioning, and model based.

www.statmethods.net/advstats/cluster.html - - Yippy Index V

[Statistics.com - Cluster Analysis](#) [new window](#) [preview](#)

This course covers how to use various cluster **analysis** methods, such as hierarchical **clustering**, to identify possible clusters in multivariate data.

<https://www.statistics.com/cluster-analysis> - - Yippy Index V

[k-means clustering - Wikipedia](#) [new window](#) [preview](#)

k-means **clustering** is a method of vector quantization, originally from signal processing, that is popular for cluster **analysis** in data mining. k-means **clustering** aims ...

https://en.wikipedia.org/wiki/K-means_clustering - - Yippy Index V

[Cluster Analysis - norusis.com](#) [new window](#) [preview](#)

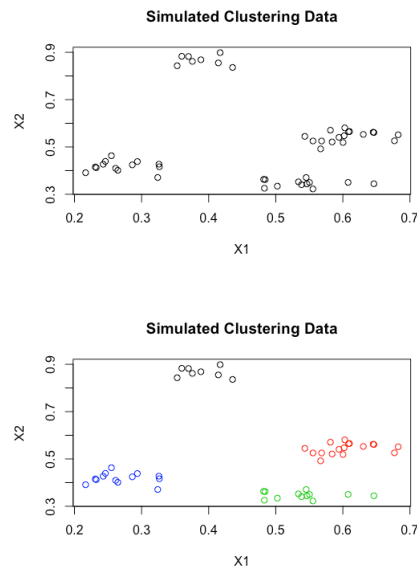
363 Cluster **Analysis** depends on, among other things, the size of the data file. Methods commonly used for small data sets are impractical for data files with ...

Example Data

```
#### SIMULATED DATA ####
library(MixSim)

set.seed(3)
Q <- MixSim(BarOmega = 0.00002, MaxOmega = NULL, K = 4, p = 2, hom = TRUE)
A = simdataset(n = 50, Pi = Q$Pi, Mu = Q$Mu, S = Q$S)
sim_d = data.frame(A$X)
sim_d$true_cluster = A$id
plot(sim_d[,1:2], main = "Simulated Clustering Data")
plot(sim_d[,1:2], col = sim_d$true_cluster, main = "Simulated Clustering Data")
```

Example Data



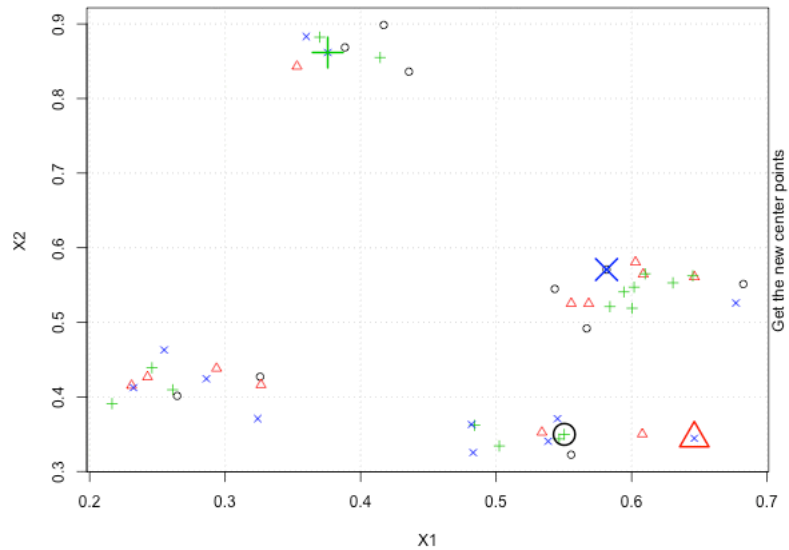
	X1	X2	true_cluster
1	0.4356460	0.8359740	1
2	0.3530190	0.8431142	1
3	0.3757343	0.8616696	1
4	0.3699114	0.8822238	1
5	0.3598940	0.8830917	1
6	0.4144158	0.8550112	1
7	0.3884225	0.8685076	1
8	0.4170783	0.8985235	1
9	0.5942671	0.5407596	2
10	0.6085898	0.5644635	2
11	0.5814696	0.5707407	2
12	0.5433074	0.5447630	2
13	0.6018648	0.5471925	2
14	0.6824566	0.5513103	2
15	0.5668619	0.4918365	2
16	0.6099432	0.5650591	2
17	0.6028487	0.5806603	2
18	0.6769366	0.5259041	2
19	0.5839923	0.5210776	2
20	0.6463144	0.5608912	2
21	0.5683214	0.5251682	2
22	0.6451486	0.5622831	2
23	0.5554161	0.5253096	2
24	0.6003407	0.5188467	2
25	0.6304850	0.5529242	2

Well-Known Methods

• K-means Clustering

1. Specify the number of clusters (K).
2. Define what is **similar** or **dissimilar**
→ **Distance!**
3. Algorithm
 1. Random assign all the points to a unique cluster (k), where $k = 1, 2, 3, \dots, K$.
 2. Calculate the mean center point for each cluster (m_1, m_2, \dots, m_K).
 3. Reassign the point to the cluster that it has closest distance to those mean center points.
 4. Repeat 2. ~ 3., until the assignment stop changing.

K-means Animation!



Definition for (Dis)similarity

- Suppose we have a measured point, x_{ij} , where
Points: $i = 1, 2, 3, \dots, N$
Attributes: $j = 1, 2, 3, \dots, p$

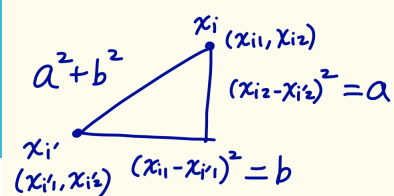
- Dissimilarity between point i and i'

$$D(x_i, x_{i'}) = \sum_{j=1}^p d_j(x_{ij}, x_{i'j})$$

- Euclidean Distance

$$d_j(x_{ij}, x_{i'j}) = (x_{ij} - x_{i'j})^2.$$

$$d(x_i, x_{i'}) = \sum_{j=1}^p (x_{ij} - x_{i'j})^2 = \|x_i - x_{i'}\|^2$$



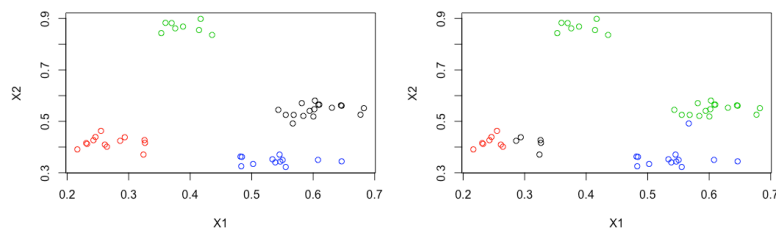
K-means Clustering

• Things Need To Be Noticed

1. You won't always get the same results. So, do it more times!

```
#### Things need to be noticed ####
set.seed(2223) # good case
k = kmeans(sim_d[,1:2], centers=4, nstart = 1)
plot(sim_d[,1:2], col = k$cluster, main = "Good Case")

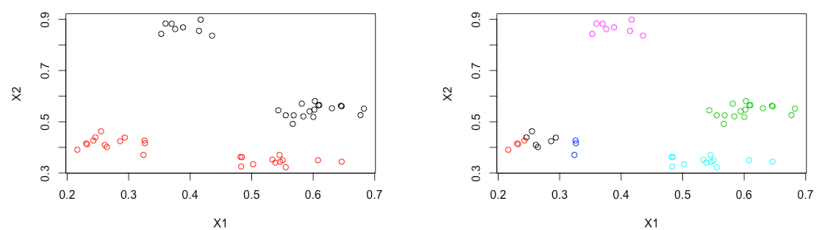
set.seed(14) # bad case
k = kmeans(sim_d[,1:2], centers=4, nstart = 1)
plot(sim_d[,1:2], col = k$cluster, main = "Bad Case")
```



K-means Clustering

• Things Need To Be Noticed

2. How to choose the correct number of clusters ?



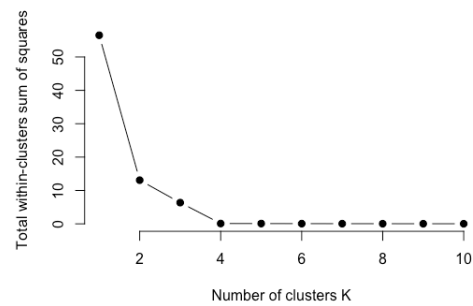
$$\sum (x_i - \bar{x})^2 = \sum (x_i - \bar{x}_k)^2 + \sum (\bar{x}_k - \bar{x})^2$$

TOTAL Sum of Squares = **WITHIN** Groups SS + **BETWEEN** Groups SS
 (TOTAL variance = **WITHIN** variance + **BETWEEN** variance)

K-means Clustering

- An Intuitive Method for finding the best k

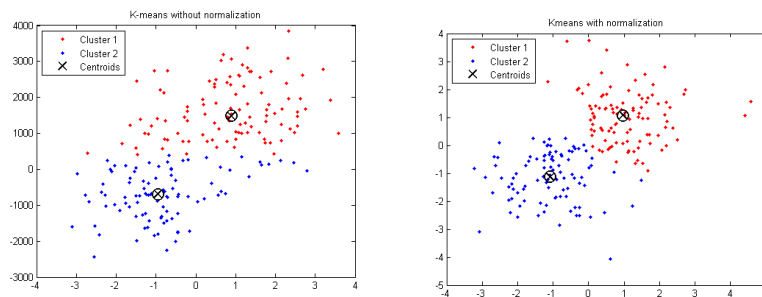
```
# find the best number of cluster
k.max <- 10
wss <- sapply(1:k.max, function(k){
  kmeans(sim_d, k, nstart=30, iter.max = 15 )$tot.withinss
})
plot(1:k.max, wss, type="b", pch = 19, frame = FALSE, xlab="Number of clusters K",
     ylab="Total within-clusters sum of squares")
```



K-means Clustering

- Things Need To Be Noticed

3. Standardize the data or not?



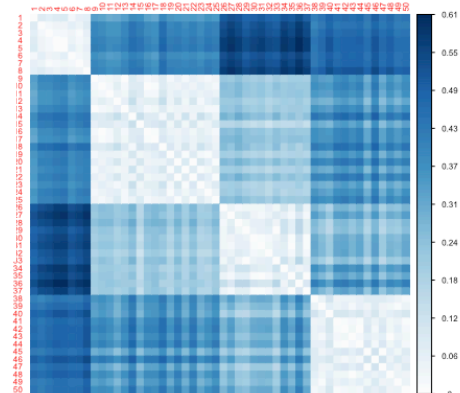
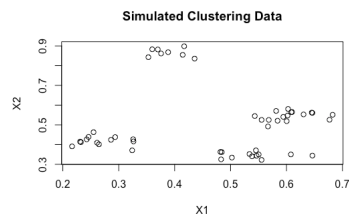
The source of pics : [here](#)

Hierarchical Clustering

• Dissimilarity Matrix

```
#### Dissimilarity Matrix ####  
library("corrplot")
```

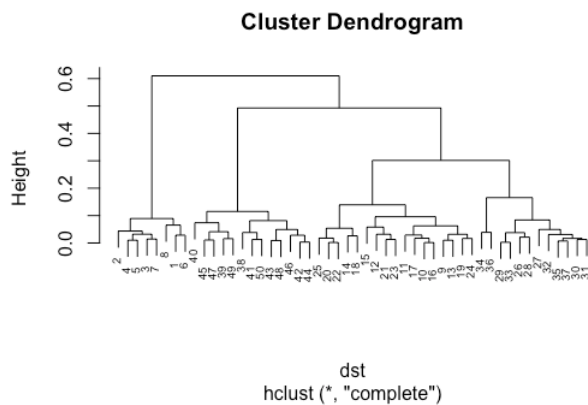
```
dst = dist(sim_d[,1:2], method = "euclidean")  
corrplot(as.matrix(dst), is.corr = FALSE, method = "color",  
         tl.cex = 0.5, cl.cex = 0.5, mar = c(0.5, 0.5, 0.5, 0.5))
```



Hierarchical Clustering

• Algorithm

```
# Hierarchical clustering  
dst = dist(sim_d[,1:2], method = "euclidean")  
hc = hclust(dst)  
plot(hc, cex = 0.6)  
rect.hclust(hc, k = 4, border = 2:5)
```



Hierarchical Clustering

• Find the Best k

```
#### Find the best cluster ####
library(dtw)
dst = dist(sim_d[,1:2], method = "euclidean")
hc = hclust(dst)

wss <- function(d) {
  sum(scale(d, scale = FALSE)^2)
}

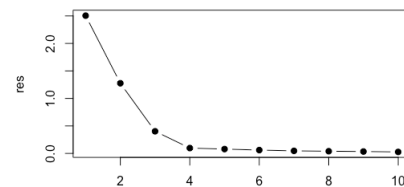
wrap = function(i, hc, x) {
  cl = cutree(hc, i)
  spl = split(x, cl)
  wss = sum(sapply(spl, wss))
  wss
}

res = sapply(seq.int(1, 10), wrap, h = hc, x = sim_d[,1:2])
plot(seq_along(res), res, type = "b", pch = 19)

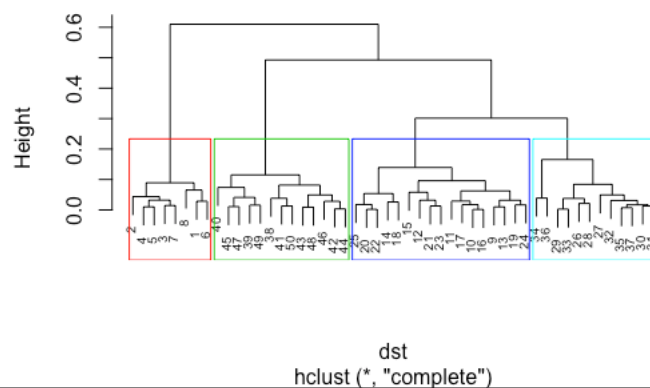
k = 4
plot(hc, cex = 0.6)
rect.hclust(hc, k, border = 2:5)
cutree(hc, k)
```

Hierarchical Clustering

• Find the Best k



Cluster Dendrogram



Hierarchical Clustering

• Things Need To Know

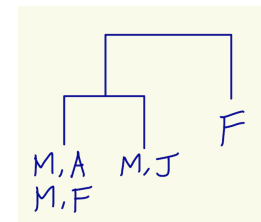
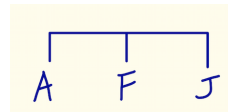
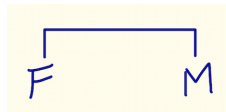
1. No need to specify the number of clusters. One Dendrogram can show many clusters.
2. It assumes that all the clusters are nested with a hierarchical structure, but it may not be the case.

Ex:

$x_1 = \{ \text{Male, Female} \}$, $x_2 = \{ \text{Americans, Japanese, French} \}$

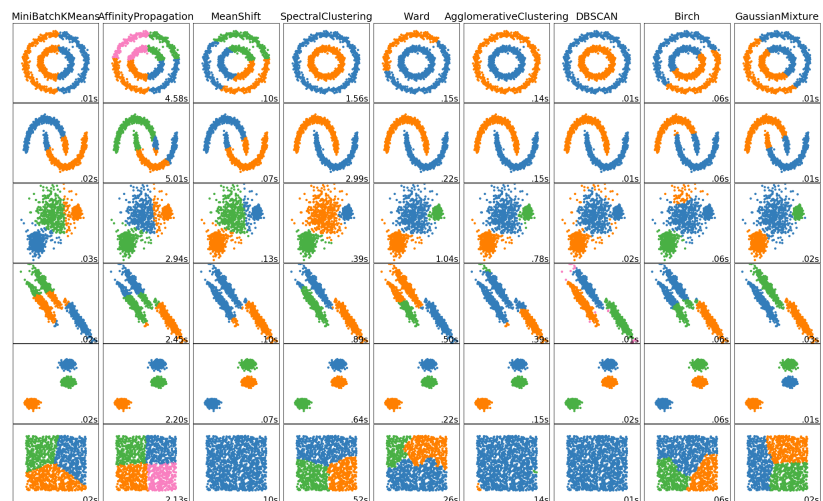
Best 2 clusters : by Gender

Best 3 clusters : by Nationality



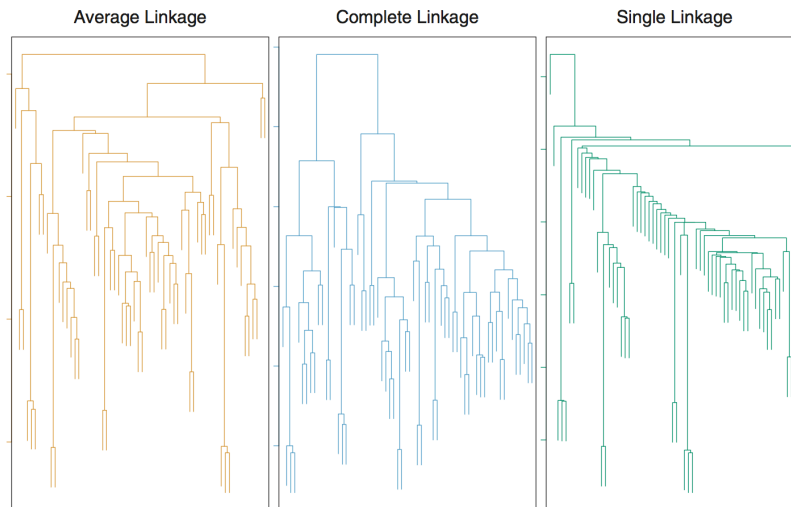
Further Topics

The source of the graph : [here](#)



Further Topics

The source of the graph : [here](#)



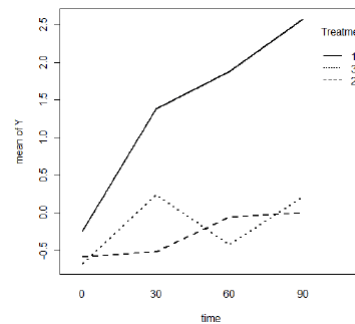
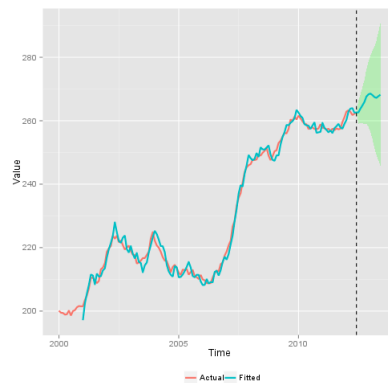
Time Series Data

• General Definition

The data is taken by time and believe that time will be an influential factor to the data.

Ex : [Sales data](#) ,

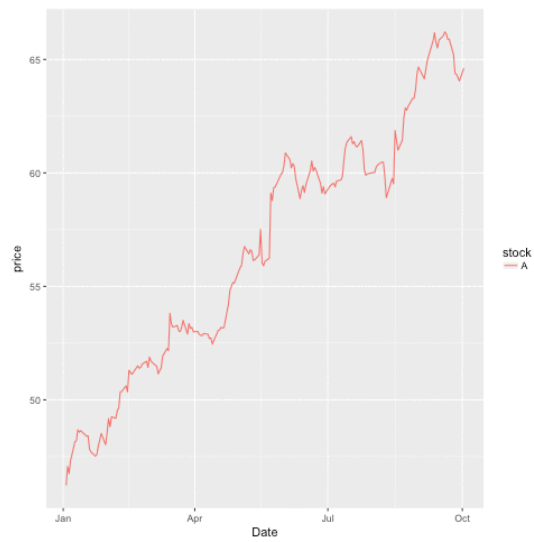
[Difference in temperature under different treatments](#)



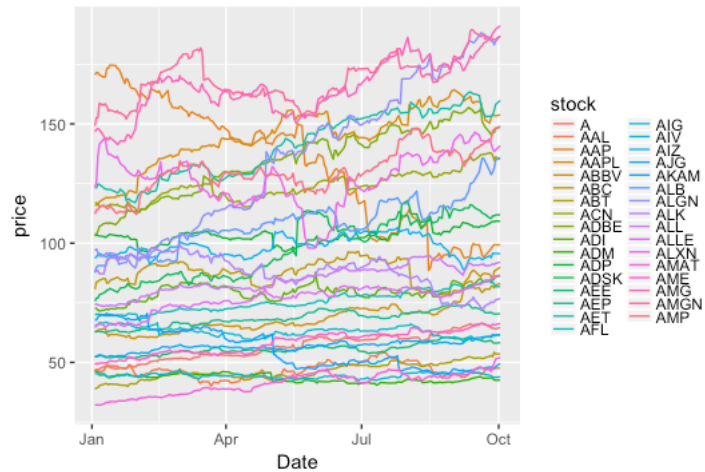
Example Data



Example Data



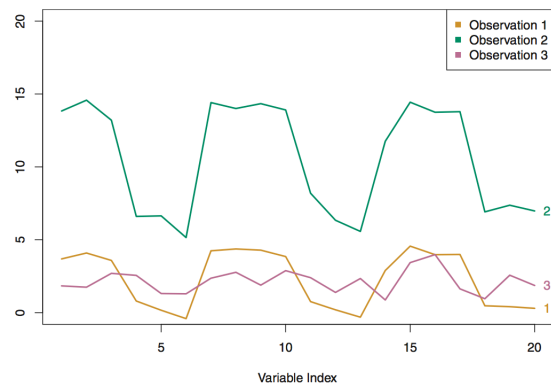
Example Data



Distance between Time Series

- Euclidean Distance may not be what we want.

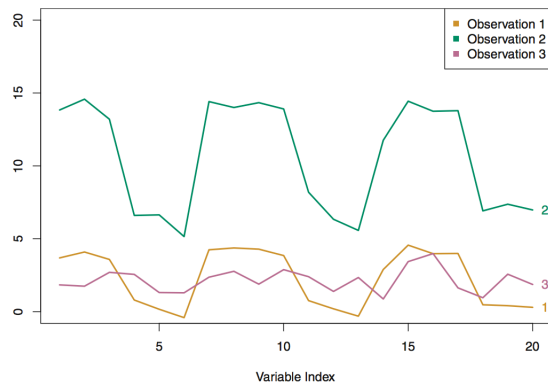
(The source of this pic : [here](#))



Distance between Time Series

• Think about Correlation

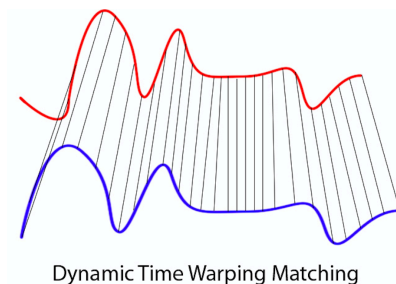
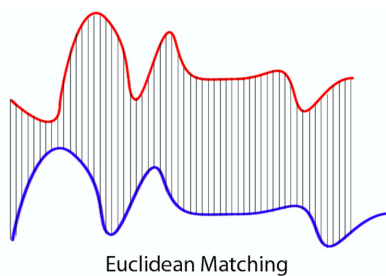
$$\rho(x_i, x_{i'}) = \frac{\sum_j (x_{ij} - \bar{x}_i)(x_{i'j} - \bar{x}_{i'})}{\sqrt{\sum_j (x_{ij} - \bar{x}_i)^2 \sum_j (x_{i'j} - \bar{x}_{i'})^2}}$$



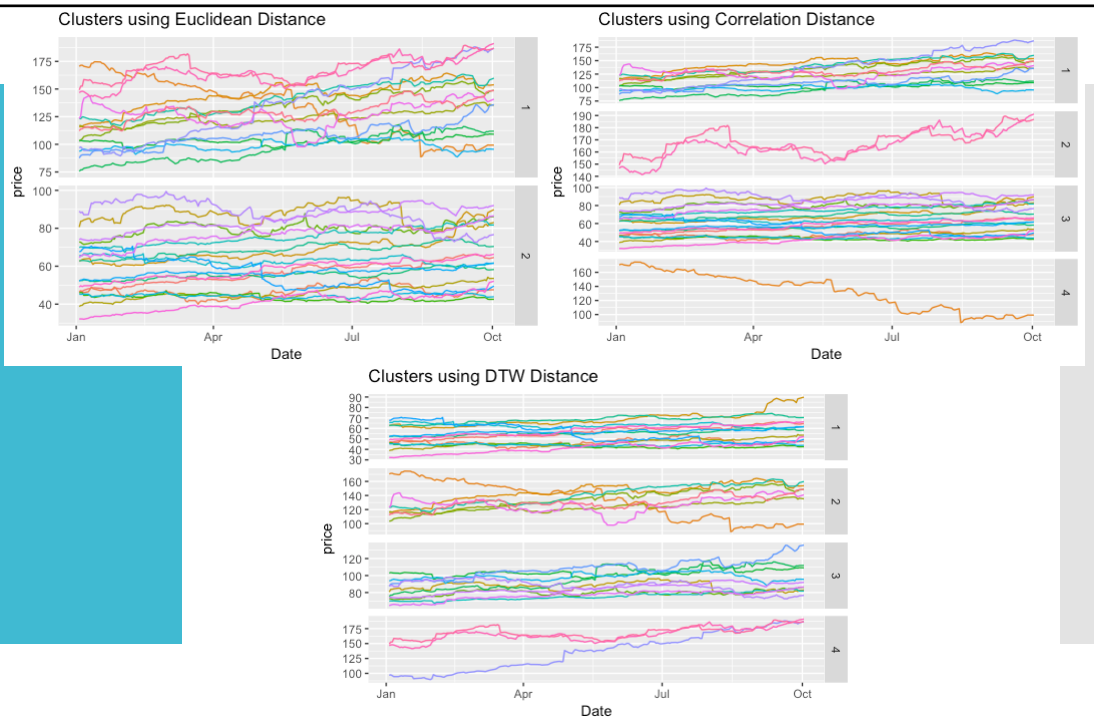
Distance between Time Series

• Think about Dynamic Time Warping (DTW) Distance

(The source of this pic : [here](#))



Results



References and Resources

• Books

- [An Introduction to Statistical Learning with Applications in R](#)
- [The Elements of Statistical Learning: Data Mining, Inference and Prediction](#)



• R Packages

- [MixSim](#) – Generate clustering data
- [corrplot](#) – Plot the dissimilarity matrix
- [tidyverse](#) – Data Manipulation
- [ggplot2](#) – Data Visualization
- [amap](#) – K-means with Correlation Distance

Q and A

- My personal website and Github
 - <http://yintingchou.com/>
 - https://github.com/choux130/EU_TSClustering_101117

Demo

- Rstudio!