Data Analytics Course at Indian Institute of Foreign Trade

Day 19: UNSUPERVISED LEARNING Using R

by

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CONTENTS

SL No.	Topics (Dimensionality Reduction)	SL No.	Topics
1	Principal Component Analysis	3	Cluster Analysis
2	Factor Analysis	4	Association Rule Mining

- A dimensionality reduction technique
- Reduces the dimensionality of multivariate data without compromising much on the variation in the original data set.
- Achieved by transforming the original varaible into a new set of variables namely prinipal components (PCAs)
- PCAs are uncorrelated and ordered
- Hence the first few of them account for most of the variation in the orignal variables

- Desribes the variation in a set of correlated variables $x = (x_1, x_2, ---, x_q)$ by a set of uncorrelated variables $y = (y_1, y_2, ---, y_q)$
- Each principal component is a linear combination of the x variables.
- The new variables are derived in decreasing order of importance.
- Hence y₁ account for maximum possible variation in x among all linear combinations of x
- y₂ account for maximum possible of the remaining variation subject to being uncorrelated to y1. and so on.

- A dimensionality reduction technique
- Large number of correlated variables can be reduced to a manageable number of uncorrelated or independent factors.
- The emphasis is on the identification of underlying factors that might explain the dimensions associated with large data sets

$$y_i = a_{i1}x_1 + a_{i2}x_2 + a_{i3}x_3 + ---+ a_{iq}x_q$$

Where y_i: estimate of ith principal component, a_i: weight or score coefficient, x_i: ith variable and k: number of variables

The coefficients are selected such that

- the first principal component explains largest portion of the total variation
- •the second first principal component accounts for the most of the residual variance, etc.

- Helps to understand the variability in large data sets with inter correlated variables using a smaller number of uncorrelated factors.
- Explaining variability of a set of n variables using m factors where m < n
- The emphasis is on the identification of underlying factors that might explain the dimensions associated with large data

Objectives

- Reduces the complexity of a large set of variables by summarizing them in a smaller set of components or factors
- Tries to improve the interpretation of complex data through logical factors

Computation of sample Principal Components

- The first principal component is that linear combination of original variables whose sample variance is greatest amongst all possible such linear combinations
- The second principal component is the linear combination of original variables that account for maximum proportion of the remaining variance subject to being uncorrelated with the first principal component and so on
- The first principal component is

$$y_1 = a_{11}x_1 + a_{12}x_2 + a_{13}x_3 + --- + a_{1q}x_q$$

The variance of y_1 be increased by increasing $a_1 = (a_{11}, ---, a_{1q})$, a restriction must be placed on these coefficients.

The sensible restriction or constraint is to ensure that sum of squares of the coefficients should be equal to one

$$a_1'a_1 = 1$$

Computation of sample Principal Components

- To choose the elements of a_1 which maximizes the variance of y_1 subject to the constraint of a_1 ' $a_1 = 1$
- Since y_1 is a linear combination of x, the sample variance of y_1 is given by

$$Var(y_1) = a_1'Sa_1$$

where S is the sample covariance matrix of x

The coefficients a₁ of first principal component y₁ is computed by solving

Maximize

$$z = a_1'Sa_1$$
Subject to
$$a_1'a_1 = 1$$

The solution to the above problem (using Lagrange multiplier method) is the Eigen vector of S corresponding to the largest Eigen value of S denoted by λ_1

Computation of sample Principal Components

In general, the coefficients a_i of first principal component y_i is computed by solving

Maximize

$$z = a_i$$
'Sa_i
Subject to
 a_i 'a_i = 1

The solution to the above problem (using Lagrange multiplier method) is the Eigen vector of S corresponding to the ith largest Eigen value of S denoted by λ_i

Since $a_i'a_i = 1$, the variance of ith principal component y_1 will be λ_i

Steps

- Prepare correlation matrix
- Extract a set of principal components using correlation matrix
- Determine the number of principal components
- Interpret results

Example: Suppose a researcher wants to determine the underlying benefits consumers seek from the purchase of a toothpaste. A sample of 30 respondents was interviewed. The respondents were asked to indicate their degree of agreement with the following statements using a 7 point scale (1: strongly disagree, 7: strongly agree)

- 1. It is important to buy a toothpaste that prevents cavities
- 2. I like a toothpaste that gives shiny teeth
- 3. A toothpaste should strengthen your gums
- 4. I prefer toothpaste that freshens breath
- 5. Prevention of tooth decay is not an important benefit offered by a toothpaste
- 6. The most important consideration in buying a toothpaste is attractive teeth

Dataset: **PCA_Factor_Analysis_Example.csv**

Step 1: Normalize the data

z transform:

Transformed data = (Data – Mean) / SD

Reading ghe file to R

>mydata = mydata[,2:7]

Transforming the variables

>myzdata = scale(mydata)

Step 2: Check for Correlation

- Variables must be correlated for data reduction.
- > cor(myzdata)

Correlation Matrix

		x 1	x2	х3	x4	x5	x6
	x1	1.000	053	.873	086	858	.004
	x2	053	1.000	155	.572	.020	.640
	x 3	.873	155	1.000	248	778	018
Correlation	x4	086	.572	248	1.000	007	.640
	x5	858	.020	778	007	1.000	136
	x6	.004	.640	018	.640	136	1.000

High correlation between x₁, x₃ & x₅

Good correlation between x₂, x₄ & x₆

Step 4: Method used: Principle Component Analysis

> mymodel = princomp(myzdata)

>summary(mymodel)

Used to identify minimum number of components accounting for maximum variance in the data; Eigen Values: Amount of variance attributed to a component.

Total Variance = 6 (Sum of all Eigen values)

Prop. variance for PC1= Eigen value of PC1 / Total Variance (2.731/6 = 0.455)

Component	SD	Variance	Proportion of Variance	Cumulative Proportion of Variance
PC 1	1.653	2.732	0.455	0.455
PC 2	1.489	2.217	0.369	0.825
PC 3	0.665	0.442	0.074	0.899
PC 4	0.584	0.341	0.057	0.955
PC 5	0.427	0.182	0.030	0.986
PC 6	0.292	0.085	0.014	1.000
Total		6.000		

Step 4: Determine the number of Components

- 1.Based on Eigen Values: Only components with Eigen value > 1.0 or Eigen value > 0.7 are selected.
- 2. Based on cumulative % variance: Factors extracted should account for at least 65 % of variance

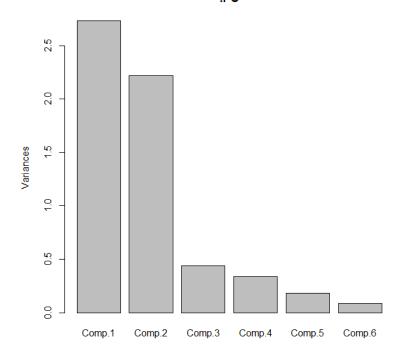
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PC 6	0.292	0.085	0.014	1.000
Total		6.000		

Number of factors selected: 2

Step 4: Determine the number of Factors

>plot(mymodel)

3. Based on Scree plot: Plot of the Eigen values against the number of factors in order of extraction. The number of components is identified based on slope change of scree plot



Step 5: Calculate Component Scores – Eigen Vectors

>loadings(mymodel)

$$y_i = a_{i1}x_1 + a_{i2}x_2 + a_{i3}x_3 + ---+ a_{ik}x_k$$

	Component			
	y ₁ y ₂			
X ₁	0.562	-0.170		
X ₂	-0.182	-0.534		
X ₃	0.566 -0.088			
X ₄	-0.207 -0.530			
X ₅	-0.526	0.236		
X ₆	-0.107	-0.585		

Step 5: Interpret Components – Eigen Vectors

	Component				
	y ₁ y ₂				
X ₁	0.562	-0.170			
X ₂	-0.182 -0.534				
X ₃	0.566 -0.088				
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X ₅	-0.526 0.236				
X ₆	-0.107 -0.585				

Component 1 is correlated with x₁, x₃ & x₅

Component 2 is correlated with x₂, x₄ & x₆

Step 5: Interpret Components

	Component			
	У ₁	y ₂		
Prevention of Cavities	0.562	-0.170		
x ₂	-0.182	-0.534		
Strong Gum	0.566	-0.088		
X ₄	-0.207	-0.530		
Non Prevention of Tooth Decay	-0.526	0.236		
X ₆	-0.107	-0.585		

Interpretation

Component 1 (y₁) represents the health related benefits

Step 5: Interpret Components

	Component			
	У ₁	y ₂		
Prevention of Cavities	0.562	-0.170		
Shiny Teeth	-0.182	-0.534		
Strong Gum	0.566	-0.088		
Fresh Breath	-0.207	-0.530		
Non Prevention of Tooth Decay	-0.526	0.236		
Attractive Teeth	-0.107	-0.585		

Interpretation

Component 2 (y₂) represents the social related benefits

Step 6: Reduced Data Set

>pc = mymodel\$scores

>cbind(pc[,1], pc[,2])

Respondent	PC1	PC2	Respondent	PC1	PC2
1	1.953	-0.071	16	1.412	0.1352
2	-1.6763	0.9852	17	1.261	0.6098
3	2.4298	0.6577	18	2.5041	-0.2372
4	-0.0908	-1.6975	19	-1.2981	1.3974
5	-1.5154	2.7238	20	-1.2777	-1.7423
6	1.6696	0.0148	21	-1.449	1.7912
7	1.0622	1.1536	22	0.9783	-0.2455
8	2.0882	-0.5402	23	-1.4107	0.8217
9	-1.29	1.3543	24	-0.9281	-2.6799
10	-2.7958	-1.6321	25	1.4305	-0.0294
11	2.0398	0.3893	26	-1.0791	-2.2053
12	-1.6682	0.9421	27	1.4698	0.106
13	2.4379	0.6146	28	-1.5875	-1.2162
14	-0.4251	-1.9974	29	-0.8027	-3.2699
15	-1.6509	1.8801	30	-1.7904	1.987

EXPLORATORY FACTOR ANALYSIS

- Many times it may not be possible to measure some the concepts directly
- Such cases the concepts are examined indirectly by collecting information on variables which can be directly measurable and assumed to be indicators of the concepts of interest.

Example

It is difficult to conclude a student is interested in science or arts directly. The students scores on science subjects or arts subjects can be an indicator for the students interest in science or arts.

- Concepts which cannot be measured directly are called latent variables or factors
- The variables which can be directly measured and related to latent variables are called manifest variables

The method of analysis to uncover the relationship between latent variables and manifest variables is factor analysis

The method is based on multiple regression, except in factor analysis manifest variables is regressed on unobservable latent variables

Types of Factor Analysis: Exploratory and Confirmatory

Exploratory Factor Analysis

Used to investigate the relationship between factors and manifest variables without making any assumption about which manifest variables is related to which factors

Confirmatory Factor Analysis

Used to test whether a specific factor model postulated a priori on the relationship between factors and manifest variables is correct or not

Factor analysis model

A regression model linking the manifest variables to a set of unobserved (or unobservable) latent variables

Assumes that the observed relationships between the manifest variables are the result of relationship between manifest variables and latent variables

The relationship between the manifest variables is measured using covariance matrix or correlation matrix.

Factor analysis model

Let a set of observed or manifest variables $x = (x_1, x_2, - - -, x_q)$ be linked to k unobserved latent variables or common factors f_1 , f_2 , - - -, f_k , where k < q by the regression model given by

$$x_{1} = \lambda_{11}f_{1} + \lambda_{12}f_{2} + - - - + \lambda_{1k}f_{k} + \mu_{1}$$

$$x_{2} = \lambda_{21}f_{1} + \lambda_{22}f_{2} + - - - + \lambda_{2k}f_{k} + \mu_{2}$$

$$\vdots$$

$$x_{q} = \lambda_{q1}f_{1} + \lambda_{q2}f_{2} + - - - + \lambda_{qk}f_{k} + \mu_{q}$$

Where

 $\lambda_{j} \, \text{are regression coefficients of the } x \, \text{variables on the common factors known as factor loadings}$

Shows how each observed variable x_i , depends on the common factors

Factor analysis model

The regression model is

$$x = \Lambda f + \mu$$

Assumptions

The random disturbance terms μ_1 , μ_2 , - - -, μ_q are uncorrelated with each other and with the factors f_1 , f_2 , - - -, f_k .

Hence correlation between the observed variables arise from their relationship with the common factors.

The factors f_1 , f_2 , - - -, f_k also uncorrelated and occur in the standardized form with mean zero and standard deviation one

Principal Component Method of factor analysis model

Very similar to principal component analysis but not operating directly on S or R but on the reduced covariance matrix S*

$$S^* = S - \psi$$

Steps

- Prepare correlation matrix
- Extract a set of factors using correlation matrix
- Determine the number of factors
- Rotate factors to increase interpretability
- Interpret results

Example: Suppose a researcher wants to determine the underlying benefits consumers seek from the purchase of a toothpaste. A sample of 30 respondents was interviewed. The respondents were asked to indicate their degree of agreement with the following statements using a 7 point scale (1: strongly disagree, 7: strongly agree)

- 1. It is important to buy a toothpaste that prevents cavities
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Transforming the variables

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Step 2: Check for Correlation

Variables must be correlated for data reduction.

> cor(myzdata)

Correlation Matrix

		x 1	x2	х3	x4	x5	x6
	x1	1.000	053	.873	086	858	.004
	x2	053	1.000	155	.572	.020	.640
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Correlation	x4	086	.572	248	1.000	007	.640
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High correlation between x1, x3 & x5

Good correlation between x2, x4 & x6

Step 3: Check for Sampling (factor) adequacy

- >library(psych)
- >KMO(myzdata)

Statistics	Value	Criteria		
Kaiser, Meyer, Olkin (KMO)	0.66	> 0.5		

Step 4: Identifying the number of factors

Compute eigen values

Choose the factors with eigen values > 1

>s = cov(myzdata)

>s_eigen = eigen(s)

> variance = s_eigen\$values

Factor	Variance	% Variance	Cum % Variance
F1	2.731188	45.52	45.52
F2	2.218119	36.97	82.49
F3	0.441598	7.36	89.85
F4	0.341258	5.69	95.54
F5	0.182628	3.04	98.58
F6	0.085209	1.42	100.00
Total	6		

Step 4: Determine the number of Factors

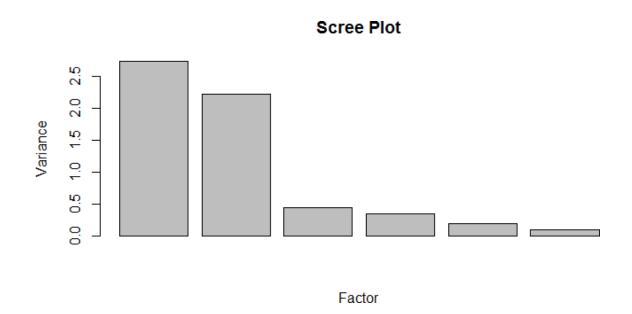
- 1. Based on Eigen Values: Only factors with Eigen value > 1.0 are selected
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Factor	Variance	% Variance	Cum % Variance
F1	2.731188	45.52	45.52
F2	2.218119	36.97	82.49
F3	0.441598	7.36	89.85
F4	0.341258	5.69	95.54
F5	0.182628	3.04	98.58
F6	0.085209	1.42	100.00
Total	6		

Number of factors selected: 2

Step 4: Determine the number of Factors

- >barplot(variance, xlab = "Factor", ylab = 'Variance', main = "Scree Plot")
- 3. Based on Scree plot: Plot of the eigen values against the number of factors in order of extraction. The number of factors is identified based on slope change of scree plot



Number of factors selected: 2

Step 5: Calculate Factor Scores

>mymodel = factanal(myzdata, 2)

	Component		
	1 2		
x1	0.968	0.000	
x2	0.000	0.749	
х3	0.898	-0.140	
x4	0.000	0.784	
x5	-0.887	0.236	
x6	0.000	0.830	

Interpretation is difficult when the variables are evenly loaded on many factors Solution: Rotation

Step 5: Calculate Factor Scores: Rotation

A process by which a solution is made more interpretable without changing its underlying mathematical properties.

Types of rotations

- 1. Orthogonal rotation
- 2. Oblique rotation

Orthogonal rotation

Restricts the rotated factors to being uncorrelated

Oblique rotation

allows the rotated factors to be correlated

Step 5: Calculate Factor Scores: Rotation

A process by which a solution is made more interpretable without changing its underlying mathematical properties.

Commonly used rotation: Orthogonal rotation

Commonly used orthogonal rotation: varimax rotation

Try to achieve factors with a few large loadings and as many near – zero loadings as possible

Step 5: Calculate Factor Scores: Rotation

- >myrotatedmodel = factanal(myzdata, 2, rotation = "varimax", scores = "regression")
- > myrotatedmodel

	Component		
	1 2		
x1	0.968	0.000	
x2	0.000	0.749	
х3	0.898	-0.140	
x4	0.000	0.784	
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Step 5: Interpret Components – Eigen Vectors

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Component 1 is correlated with x1, x3 & x5

Component 2 is correlated with x2, x4 & x6

Step 5: Interpret Components

	Component		
	1	2	
Prevention of Cavities	0.968	0.000	
x2	0.000	0.749	
Strong Gum	0.898	-0.140	
x4	0.000	0.784	
Non Prevention of Tooth Decay	-0.887	0.236	
х6	0.000	0.830	

Interpretation

Component 1 represents the health related benefits

Step 5: Interpret Components

	Component	
	1	2
Prevention of Cavities	0.968	0.000
Shiny Teeth	0.000	0.749
Strong Gum	0.898	-0.140
Fresh Breath	0.000	0.784
Non Prevention of Tooth Decay	-0.887	0.236
Attractive Teeth	0.000	0.830

Interpretation

Component 2 represents the social related benefits

Step 6: Reduced Data Set

>output = myrotatedmodel\$scores

>output

· μι	·				
Respondent	Factor1	Factor2	Respondent	Factor1	Factor2
1	1.3046	-0.2413	16	0.8934	-0.342
2	-1.2952	-0.2556	17	0.5714	-0.5502
3	1.1629	-0.7569	18	1.501	-0.24
4	0.1747	1.0108	19	-0.9845	-0.6938
5	-1.428	-1.3608	20	-0.4187	1.259
6	0.9864	-0.2511	21	-1.3132	-0.8042
7	0.4605	-0.9084	22	0.5706	-0.0143
8	1.1867	-0.0515	23	-0.9855	-0.1067
9	-0.7678	-0.6358	24	0.0209	1.7277
10	-1.1191	1.3473	25	0.8821	-0.1881
11	1.0738	-0.6495	26	-0.3011	1.5195
12	-1.0785	-0.1976	27	0.4675	-0.2914
13	1.3796	-0.6989	28	-0.5606	0.7776
14	0.0978	1.2286	29	0.1111	2.1146
15	-1.394	-0.7847	30	-1.1988	-0.9623

A technique used to classify objects or cases into relatively homogeneous groups called clusters

Cluster

A collection of data objects similar to one another within the same cluster and dissimilar to the objects in other clusters

Cluster analysis

A procedure for grouping a set of data objects into clusters

•A technique used to classify objects or cases into relatively homogeneous groups called clusters

Example: A survey was done to study the consumers attitude towards shopping. The consumers need to be clustered based on their attitude towards shopping. The respondents were asked to express their degree of agreement with the following statements on a 7 point scale (1: strongly disagree, 7: strongly agree).

x1: Shopping is fun

x2: Shopping is bad for your budget

x3: I combine shopping with eating out

x4: I try to get the best buys when shopping

x5: I don't care about shopping

x6: You can save a lot os money by comparing prices

Dataset: Cluster_Analysis_Example.csv

Step 1: Choose Type of clustering - Agglomerative Clustering

- Hierarchical Clustering characterized by development of a hierarchy or tree like structure
- Starts with each object or record as separate clusters
- Clusters are formed by grouping objects in to bigger and bigger clusters until all objects are in one cluster.
- The objects grouped based on linkage measure
- Commonly used linkage measure is Euclidean distance d,
- Euclidean distance between two records i and j, d_{ij} is defined as

$$d_{ij} = \sqrt{\sum_{k=1}^{q} (x_{ik} - x_{jk})^2}$$

Step 1: Choose Type of clustering - Agglomerative Clustering

- The data are not partitioned into a particular number of classes or groups at a single step
- Consists of a series of partitions that may run from a single cluster containing all individuals to n clusters, each contain a single individual
- Produce partitions by a series of successive fusions of the n individuals into groups
- Fusion once made are irreversible, when the algorithm has placed two individuals in the same group they cannot subsequently appear in different groups

Types of Linkage

1. Single Linkage:

Based on minimum distance

The first two objects clustered are those having minimum distance between them

$$d_{\mathbf{A}B} = \min_{\substack{i \in A \\ i \in B}} (d_{ij})$$

2. Complete Linkage:

Based on maximum distance

The distance between two clusters is calculated as the distance between two furthest points $\frac{1}{2} = \frac{1}{2} \frac{1}{2$

$$d_{AB} = \max_{\substack{i \in A \\ i \in B}} (d_{ij})$$

Where d_{AB} is the distance between two clusters A and B and d_{ij} is the distance between individuals i and j found from the initial inter – individual distance matrix

Types of Linkage

3. Average Linkage:

Based on average distance

The distance between two clusters is defined as the average of the distance between all pairs of points

Preferred method

$$d_{AB} = \frac{1}{n_A n_B} \sum_{i \in A} \sum_{j \in B} d_{ij}$$

Where n_A and n_B are the numbers of individuals in clusters A and B

Step2: Choose Method

Variance method:

- Generates clusters with minimum within cluster variance
- Uses Ward's Procedure

Ward's Procedure:

- For each cluster means for all the variables are computed
- For each object or record, the Euclidean distance to the cluster mean is computed

R Code

Read data to mydata and compute distance

Generate Clusters

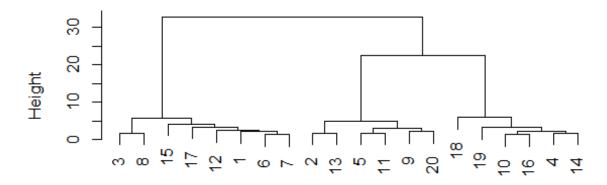
```
>mymodel = hclust(distance, method =
"ward.D")
```

Plot Dendogram

>plot(mymodel)

Decide on number of clusters: Dendrogram

Cluster Dendrogram



distance hclust (*, "ward.D")

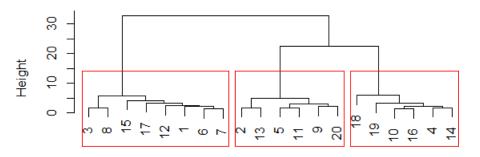
Decide on number of clusters: Dendrogram

Stages is given in x axis and distance in y axis

When one move from 3 cluster to 2 cluster the distance increases drastically. So 3 cluster may be appropriate

- >groups = cutree(mymodel, k = 3)
- > rect.hclust(mymodel, k = 3, border = "red")

Cluster Dendrogram



distance hclust (*, "ward.D")

Identification of cluster membership for each record

>output = cbind(mydata, groups >print(output)

Respondent id	x1	x2	х3	x4	x5	х6	groups
1	6	4	7	3	2	3	1
2	2	3	1	4	5	4	2
3	7	2	6	4	1	3	1
4	4	6	4	5	3	6	3
5	1	3	2	2	6	4	2
6	6	4	6	3	3	4	1
7	5	3	6	3	3	4	1
8	7	3	7	4	1	4	1
9	2	4	3	3	6	3	2
10	3	5	3	6	4	6	3
11	1	3	2	3	5	3	2
12	5	4	5	4	2	4	1
13	2	2	1	5	4	4	2
14	4	6	4	6	4	7	3
15	6	5	4	2	1	4	1
16	3	5	4	6	4	7	3
17	4	4	7	2	2	5	1
18	3	7	2	6	4	3	3
19	4	6	3	7	2	7	3
20	2	3	2	4	7	2	2

Cluster Profile

> aggregate(mydata, by = list(groups), FUN = mean)

Variables		Cluster Mean			
		2	3		
x1 (shopping is fun)	4.750	2	3.875		
x2 (shopping upsets my budget)	3.5	3.500	5		
x3 (I combine shopping with eating out)	4.875	2.25	3.875		
x4 (I try to get best buys when shopping)	3. 5	4.25	4.625		
x5 (I don't care about shopping)	3	4.75	3.25		
x6 (save a lot by comparing prices	4	4	4.875		

Cluster 1: High on x1 & x3 but low on x5 Fun loving and concerned

Cluster 2: Low on x1 & x3 but High on x5
Careless & no fun in shopping (apathetic)

Cluster 3: High on x2 x4 & x6

Concerned about spending money (Economical)

k mean clustering

Partitions n individuals in a set of multivariate data into k groups or clusters (G_1 , G_2 , - - -, G_k)

k is given or a possible range is specified

Common approach is to identify the k groups which minimizes the within – group sum of squares (WGSS)

$$WGSS = \sum_{j=1}^{q} \sum_{l=1}^{k} \sum_{i \in G_i} (x_{ij} - \overline{X}_j^{(1)})^2$$

Where $\bar{x}_{j}^{(l)} = \frac{1}{n_{i}} \sum_{i \in G_{i}} x_{ij}$ is the mean of the individuals in group G_{l} on variable j

Computing WGSS for each value of k and choose that of value of k which minimize WGSS is almost impossible

One option is to plot WGSS for different values of k and choose the optimum k at which the slope of the curve changes

k mean clustering

Computing WGSS for each value of k and choose that of value of k which minimize WGSS is almost impossible

Moreover as number of cluster increases BGSS decreases or BGSS / Total SS will increase

One option is to plot BGSS/ Total SS for different values of k and choose the optimum k at which the curve flattens or slope changes

Example: Cluster the data given in cluster_Analysis_example.csv using k mean method

```
>mynewmodel = kmeans(mydata,3)
> mynewmodel
>cluster = mynewmodel$cluster
>output = cbind(mydata, cluster)
> write.csv(output, "E:/output.csv")

To find optimum k, compute BGSS / Total SS for different values of k
> kmeans(mydata,k, k = 1,2, - - -)
```

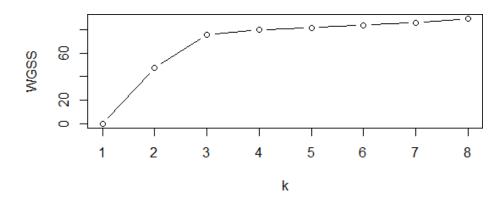
Example: Cluster the data given in cluster_Analysis_example.csv using k mean method

optimum k, > kmeans(mydata,k, k =1,2, - - -)

k	WGSS/Total SS
1	0.0
2	47.5
3	75.8
4	79.6
5	81.4
6	83.7
7	85.8
8	89.2
•	

Example: Cluster the data given in cluster_Analysis_example.csv using k mean method

optimum k, > plot(k, WGSS, type = "b")



The curve flattens after k = 3, hence optimum k is 3

A modeling technique based upon the logic that if a customer buy a certain group of items, he is more (or less) likely to buy another group of items

Example:

Those who buy cigarettes are more likely to buy match box also.

Association Rule Mining:

Developing rules that predict the occurrence of of an item based on the occurrence of other items in the transaction

Example

ld	Items
1	Milk, Bread
2	Bread, Biscuits, Toys, Eggs
3	Milk, Biscuits, Toys, Fruits
4	Bread, Milk, Toys, Biscuits
5	Milk, Bread, Biscuits, Fruits

{Milk, Bread} → {Biscuits} with probability = 2 / 3

Itemset:

A collection of one or more items

k – itemset

An itemset consisting of k items

ld	Items
1	Milk, Bread
2	Bread, Biscuits, Toys, Eggs
3	Milk, Biscuits, Toys, Fruits
4	Bread, Milk, Toys, Biscuits
5	Milk, Bread, Biscuits, Fruits

Support count:

Frequency of occurrence of an itemset

Example

{Milk, Bread, Biscuits} = 2

ld	Items
1	Milk, Bread
2	Bread, Biscuits, Toys, Eggs
3	Milk, Biscuits, Toys, Fruits
4	Bread, Milk, Toys, Biscuits
5	Milk, Bread, Biscuits, Fruits

Support:

Proportion or fraction of transaction that contain an itemset

Example

{Milk, Bread, Biscuits} = 2 / 5

ld	Items
1	Milk, Bread
2	Bread, Biscuits, Toys, Eggs
3	Milk, Biscuits, Toys, Fruits
4	Bread, Milk, Toys, Biscuits
5	Milk, Bread, Biscuits, Fruits

Frequent Itemset

An itemset whose support is greater than or equal to minimum support

Confidence

Conditional probability that an item will appear in transactions that contain another items

Example

Confidence that Toys will appear in transaction containing Milk & Biscuits

= $\{Milk, Biscuits, Toys\} / \{Milk, Biscuits\} = 2 / 3 = 0.67$

Id	Items
1	Milk, Bread
2	Bread, Biscuits, Toys, Eggs
3	Milk, Biscuits, Toys, Fruits
4	Bread, Milk, Toys, Biscuits
5	Milk, Bread, Biscuits, Fruits

Association Rule Mining

1. Frequent Itemset Generation

Fix minimum support value

Generate all itemsets whose support ≥ minimum support

2. Rule Generation

Fix minimum confidence value

Generate high confidence rules from each frequent itemset

Frequent Itemset Generation: Apriori Algorithm

- a. Fix minimum support count
- b. Generate all itemsets of length = 1
- c. Calculate the support for each itemset
- d. Eliminate all itemsets with support count < minimum support count
- e. Repeat steps c & d for itemsets of length = 2, 3, ---

Frequent Itemset Generation: Apriori Algorithm

Example:

Minimum Support count = 2

ld	Items
1	A,C,D
2	B,C,E
3	A,B,C,E
4	B,E
5	A,E
6	A,C,E

Frequent Itemset Generation: Apriori Algorithm

Example:

Minimum Support count = 2

Step 1:

Generate itemsets of length = 1 & calculate support

Item	Support count
А	4
В	3
С	4
D	1
Е	5

Frequent Itemset Generation: Apriori Algorithm

Example:

Minimum Support count = 2

Step 2:

Item	Support count
А	4
В	3
С	4
D	1
E	5

Frequent Itemset Generation: Apriori Algorithm

Example:

Minimum Support count = 2

Step 2:

Item	Support count
А	4
В	3
С	4
E	5

Frequent Itemset Generation: Apriori Algorithm

Example:

Minimum Support count = 2

Step 3:

generate itemsets of length = 2

Item	Support count
A, B	1
A, C	3
A,E	3
B, C	2
B, E	3
C,E	3

Frequent Itemset Generation: Apriori Algorithm

Example:

Minimum Support count = 2

Step 4:

Item	Support count
A, B	1
A, C	3
A,E	3
B, C	2
B, E	3
C,E	3

Frequent Itemset Generation: Apriori Algorithm

Example:

Minimum Support count = 2

Step 4:

Item	Support count
A, C	3
A,E	3
B, C	2
B, E	3
C,E	3

Frequent Itemset Generation: Apriori Algorithm

Example:

Minimum Support count = 2

Step 5:

generate itemsets of length = 3

Item	Support count
A, C, E	2
B, C, E	2

Step 6:

generate itemsets of length = 4

Itemset	Support Count
A, B, C, E	1

Frequent Itemset Generation: Apriori Algorithm

Example:

Minimum Support count = 2

Result:

Item	Support count	Support
A, C, E	2	0.33
B, C, E	2	0.33
A,C	3	0.50
A , E	3	0.50
B,C	2	0.33
B,E	3	0.50
C,E	3	0.50

Association Rule Mining: Apriori Algorithm

Example:

Minimum Support = 0.50

Minimum Confidence = 0.5

Item	Support count	Support
A, C, E	2	0.33
B, C, E	2	0.33
A, C	3	0.50
A,E	3	0.50
B,C	2	0.33
B,E	3	0.50
C,E	3	0.50

Association Rule Mining: Apriori Algorithm

Example:

Minimum Support = 0.50

Minimum Confidence = 0.5

Item	Support	Confidence
A→C	0.50	0.75
A → E	0.50	0.75
$B \rightarrow E$	0.50	1.00
$C \rightarrow E$	0.50	0.75
C→A	0.50	0.75
E→A	0.50	0.60
E→B	0.50	0.60
E→C	0.50	0.60

Association Rule Mining: Other Measures

Lift

Lift $(A \longrightarrow C) = Confidence (A \longrightarrow C) / Support (C)$

Example

Item	Confidence	Support	Lift
$A \longrightarrow C$	0.75	C = 0.67	1.12
A →E	0.75	E = 0.83	0.93

Criteria : Lift ≥ 1

Lift (A, C) = 1.12 > Lift (A, E) indicates that A has a greater impact on the frequency of C than it has on the frequency of E

R Code

```
Reading the file and variables
>target = mydata$items
>ident = mydata$ld
Making transactions
>library(arules)
>transactions = as(split(target, ident),"transactions")
Generating rules
>library(arules)
>myrules = apriori(transactions, parameter = list(support = 0.5, confidence = 0.25, minlen = 2))
Displaying rules
>myrules
>inspect(myrules)
```