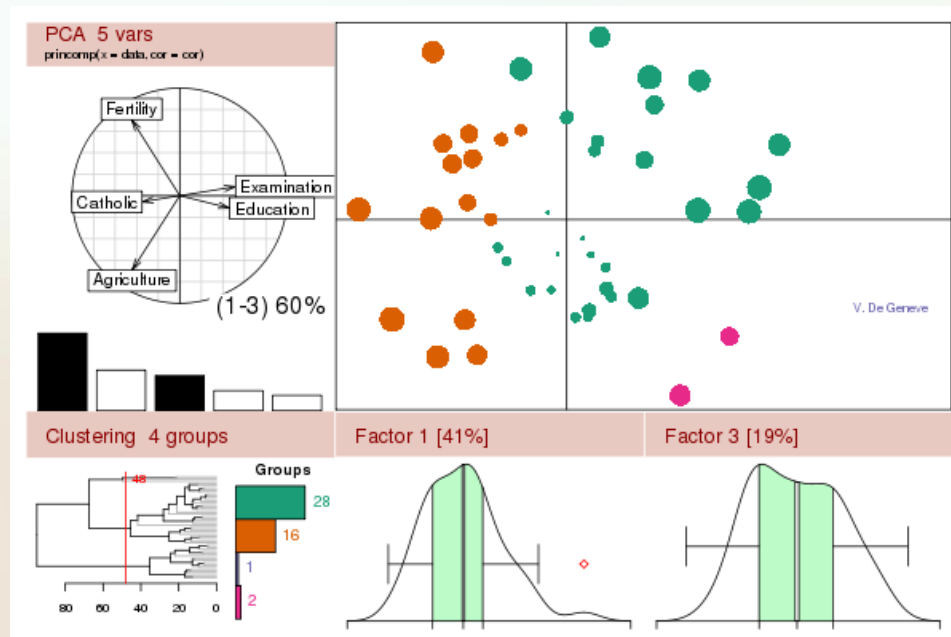


# R Language

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# R Introduction

- GNU Project Developed by John Chambers @ Bell Lab
- Free software environment for [statistical computing](#) and [graphics](#)
- Functional programming language written primarily in [C](#), [Fortran](#)



# R Technical Introduction

- R is functional programming language
- R is an interpreted language
- R is object oriented-language
- R works in an environment level

# R Downloadable links

- Two sources to get R-environment
  - [R-Project](#)
  - [R-Studio](#) (Preferred)

# Why we R?

- I. For Statistical Analysis
- II. For Data Visualization
- III. For Mathematical Functions and modeling

# Getting Started

- We will work as we go
- Declaring a variable
- `> x = 11`
- `> print(x)`
- `> y <- 11`
- `> X // Error`
- You can use `'='`, `'<-'` or `'->'` to assign a variable

# Check your variables

- You can see it in workspace section
- Or use the following command  
> `ls()`
- To remove a variable from Workspace memory
- > `rm(x)`

# Variable name rule

- Object name can use characters, numbers or period
- But number may not occur first, you can use period as first character, but then you can expect it to be skipped when you call '>ls' command
- a
- .a
- a.1
- 1a



# String

- > string = “notice double quote”
- > string <- ‘it works with single quote too’

# Numeric Operations

- `> a + a`
- `> x - y`
- `> m / n`
- `> x^2`
- `> log(2)`
- `> sqrt(y)`
- `> exp(z)`
- `> log2(1024)`
- `> abs(-10)`

# Vectors and operation

- > v\_number = c(1,2,3,4,5)
- > v\_gender = c('male' , 'female')
- > seq ( from = 1, to = 10, by = 1)
- > rep ( 1, times =10)
- > rep ( 1:3, times = 2)

# Vectors Operation & Extraction

- `> x = seq(from =1 , to=10, by=2)`
- `> y = seq ( from = 2, to 10, by=2)`
- `> x + y`
- `> x[1]` # To extract first element
- `> x[-1]` # To extract all except first element
- `> x[1:3]` # To extract 1<sup>st</sup> three elements
- `> x[c(1,3)]` # To extract 1<sup>st</sup> and 3<sup>rd</sup> element
- `> y[ y < 6 ]` # To extract element less than 6

# Matrix

- `> m_seq1 = matrix(1:9 , nrow=3, byrow =TRUE)`
- `> m_seq2 = matrix(1:9 , nrow=3, byrow =FALSE)`

# Reading Data

- > read.csv (file=~ /Dataset/titanic.csv" , header=TRUE, sep=',')
- > read.csv2 (file=~ /Dataset/titanic.csv" , header=T)
- > read.csv2 (file.choose() , header=TRUE)
- > read.table (file.choose() , header=TRUE, sep=',')

# Testing Data

- > `dim(data1)`      # To check dimension of data
- > `head(data1)`      # To check first 6 entries of data
- > `tail(data1)`      # To check last 6 entries of data
- > `data1[c(23,5,7,55), ]`      # To extract specific data
- > `name(data1)`      # To check field names
- > `attach(cap)`      # To make properties recognizable
- > `mean(Age)`      # To find Mean
- > `median (Age)`      # To find Median
- > `detach (cap)`      # To find Median

# Getting Summary

- You can ask for data Summary using
  - > `summary(data)`
- You can also ask what values are available using levels.
  - > `class(data$value)`
  - > `levels(data$value)`
  - > `x = c(1,0,1,0,0,0,1,1,0)`
  - > `x<- as.factor(x)`
  - > `class(x)`
  - > `summary(x)`

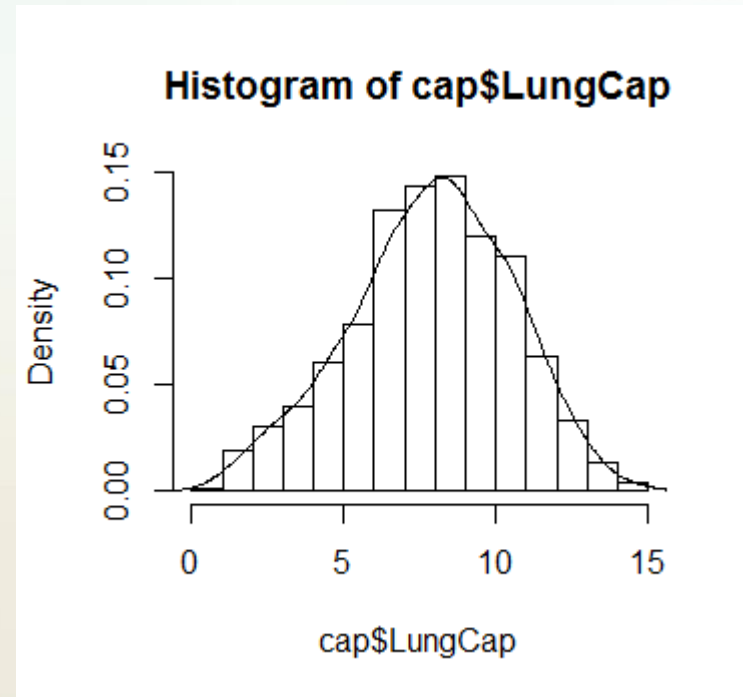


# If & blocks

- `> attach(titanic_data)`
- `> summary(age[sex=="women"])`
- `> # Sub setting data`
- `> childData <- titanic_data[age=="child" & sex=="male", ]`
- `> detach(titanic_data)`
- `> areWomanandchild <- titanic_data$sex == "women" & titanic_data$age == "child"`
- `> titanic_data_with_classification <- cbind(titanic_data, areWomanandchild) #you can also use 'rbind' for row wise binding`

# Histogram

- `> hist(cap$lungCap, prob = T)`
- `> lines(density(cap$lungCap))`

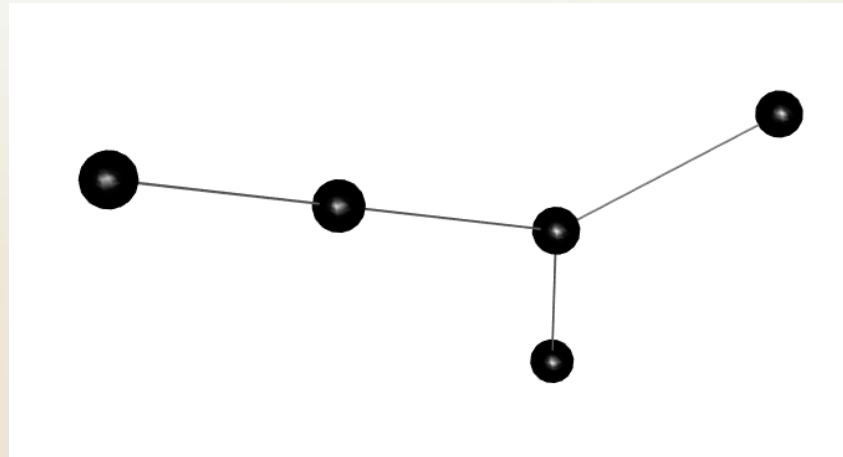
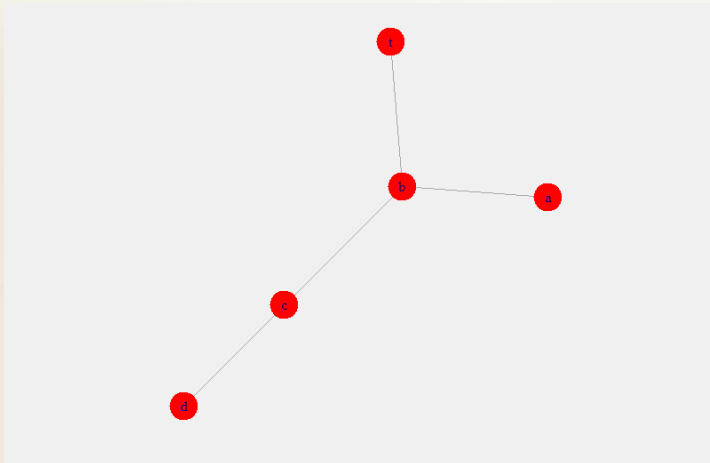


# Installing and loading packages

- Packages are simple modules that provides common functionalities.
- Packages are open sourced as well so you can create your own and contribute
- > `install.packages('igraph')`
- > `library(igraph)`
- > `help(package = igraph)`

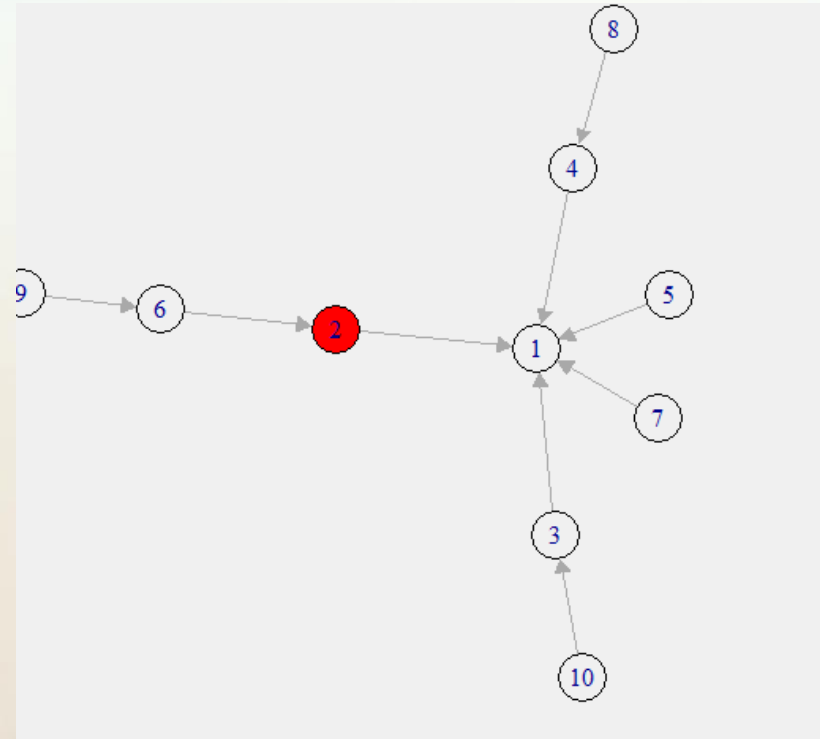
# Creating A simple graph

- `> g.manual <- graph.formula(a—b, b—c, c—d, t—b)`
- `> tkplot(g.manual, vertex.color="white", vertex.size=15)`
- `> rglplot(g.manual)`
- `> plot(g.manual)`



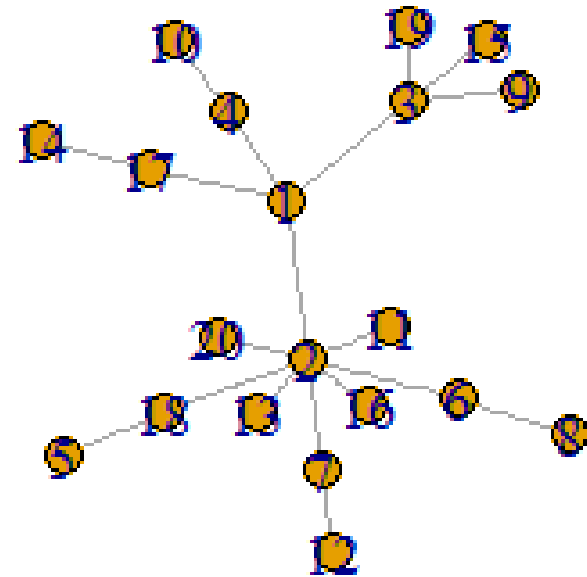
# Taking out a node

- > g.barabasi = barabasi.game(n=10, p = 0.5)
- > V(g.barabasi)[which.max(betweenness(g.barabasi, v = V(g.barabasi)))]\$color="red"
- > tkplot(v.barabasi)



# Playing more

- > `g.random = erdos.renyi.game(10, 0.3)`
- > `g.mst = minimum.spanning.tree(g.random)`



# You can find

- > closeness(g)
- > betweenness(g)
- > degree(g)
- Other important commands are
- > max(v)
- > min(v)
- > which.max(v)
- > which.min(v)

## You can Also try

Function	Description
<code>aging.prefatt.game</code>	Evolving graph, based on preferential attachment and aging
<code>bipartite.random.game</code>	Generate Bi-partite graph using random model
<code>degree.sequence.game</code>	Generate random graph with given degree sequence
<code>forest.fire.game</code>	Evolve a graph based on fire spreading phenomena
<code>graph.adjacency</code>	Create a graph from adjacency matrix
<code>graph.bipartite</code>	Creates a bi-partite graph
<code>graph.complementer</code>	Creates a complementary graph for a given graphs
<code>graph.empty</code>	Creates an empty graph