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COURSE: MSc CS

**SUBJECT: SOCIAL NETWORK
ANALYSIS**

PRACTICAL: 1-8

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1		Write a program to compute the following for a given a network: (i) number of edges, (ii) number of nodes; (iii) degree of node; (iv) node with lowest degree; (v) the adjacency list; (vi) matrix of the graph.	
2		Perform following tasks: (i) View data collection forms and/or import onemode/two-mode datasets; (ii) Basic Networks matrices transformations	
3		Compute the following node level measures: (i) Density; (ii) Degree; (iii) Reciprocity; (iv) Transitivity; (v) Centralization; (vi) Clustering.	
4		For a given network find the following: (i) Length of the shortest path from a given node to another node; (ii) the density of the graph	
5		Write a program to distinguish between a network as a matrix, a network as an edge list, and a network as a sociogram (or “network graph”) using 3 distinct networks representatives of each.	
6		Write a program to exhibit structural equivalence, automorphic equivalence, and regular equivalence from a network.	
7		Create sociograms for the persons-by-persons network and the committee-bycommittee network for a given relevant problem. Create one-mode network and two-node network for the same	
8		Perform SVD analysis of a network.	

Practical 1: Write a program to compute the following for a given a network: (i) number of edges, (ii) number of nodes; (iii) degree of node; (iv) node with lowest degree; (v) the adjacency list; (vi) matrix of the graph.

Code:

```
# Install and load the igraph package
install.packages("igraph")
library(igraph)

# Create a graph object 'g' using graph.formula function with edges 1-2, 1-3,
2-3, 2-4, 3-5, 4-5, 4-6, 4-7, 5-6, 6-7
g <- graph.formula(1-2, 1-3, 2-3, 2-4, 3-5, 4-5, 4-6, 4-7, 5-6, 6-7)

# Plot the graph object 'g'
plot(g)

# Count the number of edges in 'g'
ecount(g)

# Count the number of vertices in 'g'
vcount(g)

# Calculate the degree of each vertex in 'g'
degree(g)

# Create another graph object 'dg' using graph.formula function with edges 1-
>2, 1->3, 2<-3
dg <- graph.formula(1->2, 1->3, 2<+3)

# Plot the graph object 'dg'
plot(dg)

# Calculate the in-degree of each vertex in 'dg'
degree(dg, mode="in")

# Calculate the out-degree of each vertex in 'dg'
degree(dg, mode="out")

# Print the name of the vertex with the minimum degree in 'dg'
V(dg)$name[degree(dg)==min(degree(dg))]

# Print the name of the vertex with the maximum degree in 'dg'
V(dg)$name[degree(dg)==max(degree(dg))]
```

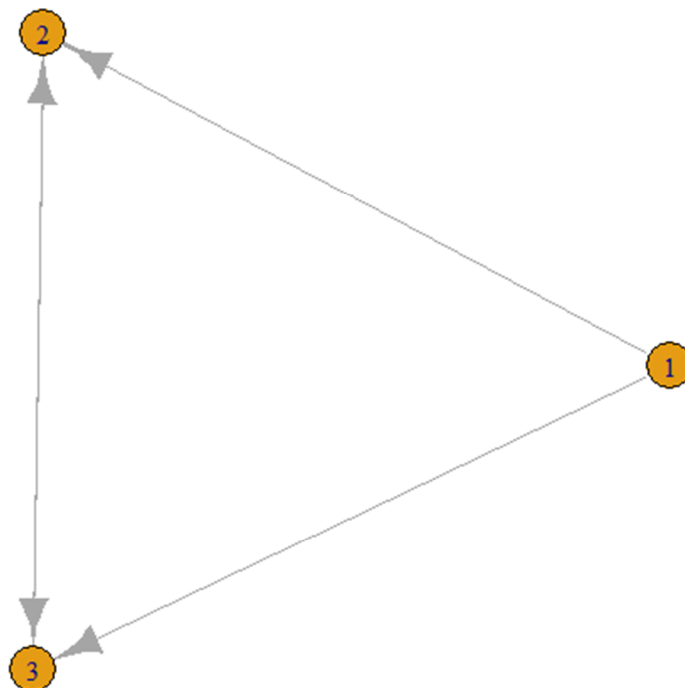
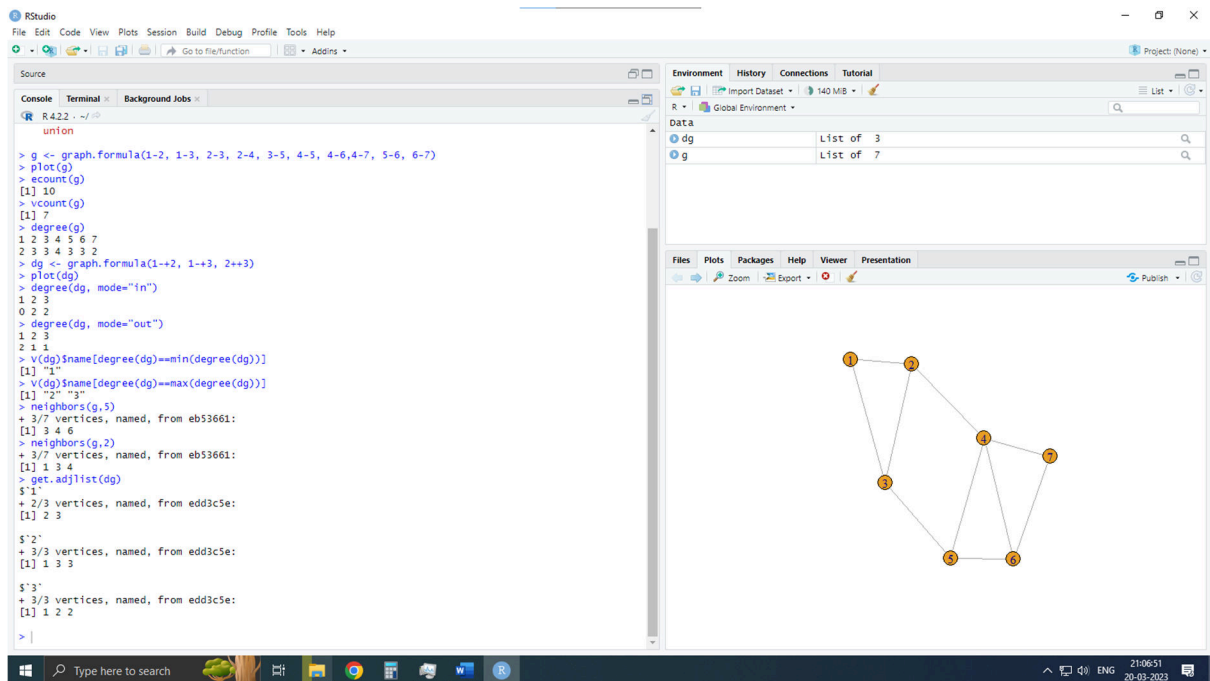
```
# Find the neighbors of vertex 5 in 'g'
neighbors(g,5)

# Find the neighbors of vertex 2 in 'g'
neighbors(g,2)

# Get the adjacency list of 'dg'
get.adjlist(dg)

# Get the adjacency matrix of 'g'
get.adjacency(g)
```

OUTPUT



Practical 2: Perform following tasks: (i) View data collection forms and/or import onemode/two-mode datasets; (ii) Basic Networks matrices transformations

Code:

```
# Get the current working directory
getwd()

# Set the working directory to "d:/SNA_pract"
setwd("d:/SNA_pract")

# Read the nodes.csv file into a data frame 'nodes'
nodes <- read.csv("nodes.csv", header = T, , as.is = T)

# Print the first few rows of 'nodes'
head(nodes)

# Read the edges.csv file into a data frame 'links'
links <- read.csv("edges.csv", header = T, as.is = T)

# Print the first few rows of 'links'
head(links)

# Create a graph object 'net' from the data frames 'nodes' and 'links'
net <- graph.data.frame(d = links,
                        vertices = nodes,
                        directed = T)

# Convert the graph object 'net' to an adjacency matrix 'm'
m = as.matrix(net)

# Print the adjacency matrix of 'net'
get.adjacency(net)

# Plot the graph object 'net'
plot(net)
```

OUTPUT

The screenshot shows the RStudio interface with the following components:

- Source Editor:** Contains R code for reading a CSV file, creating a graph object, and generating an adjacency matrix.
- Console:** Displays the output of the R code, including the structure of the 'links' data frame and the resulting 'dgcmatrix' object.
- Environment:** Lists the objects in the global environment: 'links' (52 obs. of 4 variables), 'm' (Formal class dgcmatrix), 'net' (List of 17), and 'nodes' (17 obs. of 5 variables).
- Files:** Shows the file explorer with a file named 'edges.csv'.

R Code in Source Editor:

```
R 4.2.2. ~ /
File Edit Code View Plots Session Build Debug Profile Tools Help
Go to file/function Addins
Source
Console Terminal Background Jobs
> links <- read.csv("edges.csv", header = T, as.is = T)
> head(links)
  from to weight type
1 s01 s02    10 hyperlink
2 s01 s02    12 hyperlink
3 s01 s03    22 hyperlink
4 s01 s04    21 hyperlink
5 s04 s11    22 mention
6 s05 s13    21 mention
> #basic Networks matrices transformations
> net <- graph.data.frame(d = links,
+                         vertices = nodes,
+                         directed = T)
> m = as.matrix(net)
> get.adjacency(net)
17 x 17 sparse matrix of class "dgcmatrix"
[[ suppressing 17 column names 's01', 's02', 's03' ... ]]

s01 . 2 1 1 . . . . . 1 . .
s02 1 . 1 . . . . . 1 1 . . .
s03 1 . . 1 1 . . 1 1 1 . . .
s04 . 1 . . 1 . . . 1 1 . . . 1
s05 1 1 . . . . 1 . . . . 1 .
s06 . . . . 1 . . . . . 1 1 .
s07 . 1 . . . . 1 1 . . 1 . .
s08 . 1 . . . . 1 2 . . . . .
s09 . . . . . 1 . . . . . .
s10 . 1 . . . . . . . . . . .
s11 . . . . . . . . . . . .
s12 . . . . . 1 . . . . 1 1 .
s13 . . . . . . . . . 1 . 1 .
s14 . . . . . . . . . 1 . 1 .
s15 2 . . 1 1 . . . . . . .
s16 . . . . 1 . . . . . . 1 .
s17 . . . 1 . . . . . . . . .
> |
```

Environment Data:

Object	Details
links	52 obs. of 4 variables
m	Formal class dgcmatrix
net	List of 17
nodes	17 obs. of 5 variables

Files:

- edges.csv

System Information: 21:05:17, 20-03-2023

Practical 3: Compute the following node level measures: (i) Density; (ii) Degree; (iii) Reciprocity; (iv) Transitivity; (v) Centralization; (vi) Clustering.

Code:

```
# Load the igraph library
library(igraph)

# Create a graph object 'g'
g <- graph.formula(1-2, 1-3, 2-3, 2-4, 3-5, 4-5, 4-6,4-7, 5-6, 6-7)

# Density
# Number of vertices
vcount(g)

# Number of edges
ecount(g)

# Density of the graph
ecount(g) / (vcount(g) * (vcount(g) - 1) / 2)

# Degree
degree(g)

# Reciprocity:
# Create a directed graph 'dg'
dg <- graph.formula(1->2, 1->3, 2->3)

# Plot the directed graph 'dg'
plot(dg)

# Reciprocity of the directed graph 'dg'
reciprocity(dg)

# Formula for reciprocity
(2 * dyad.census(dg)$mut / ecount(dg))

# Transitivity
# Create a famous graph 'kite'
kite <- graph.famous("Krackhardt_Kite")

# Find the adjacent triangles in the 'kite' graph
atri <- adjacent.triangles(kite)

# Plot the 'kite' graph with vertex labels as adjacent triangles
plot(kite, vertex.label = atri)
```



```
# Local transitivity of the directed graph 'dg'
transitivity(dg, type = "local")

# Proportion of adjacent triangles to all possible triangles in the 'kite'
graph
adjacent.triangles(kite) / (degree(kite) * (degree(kite) - 1) / 2)

# Centralization
# Degree of centrality
centralization.degree(g, mode = "in", normalized = T)

# Closeness Centralization
closeness(g)
centralization.closeness(g, mode = "all", normalized = TRUE)

# Betweenness Centrality
betweenness(g, directed = T, weights = NA)
edge.betweenness(g, directed = T, weights = NA)
centralization.betweenness(g, directed = T, normalized = T)

# Eigenvector centrality
centralization.evcent(g, directed = T, normalized = T)

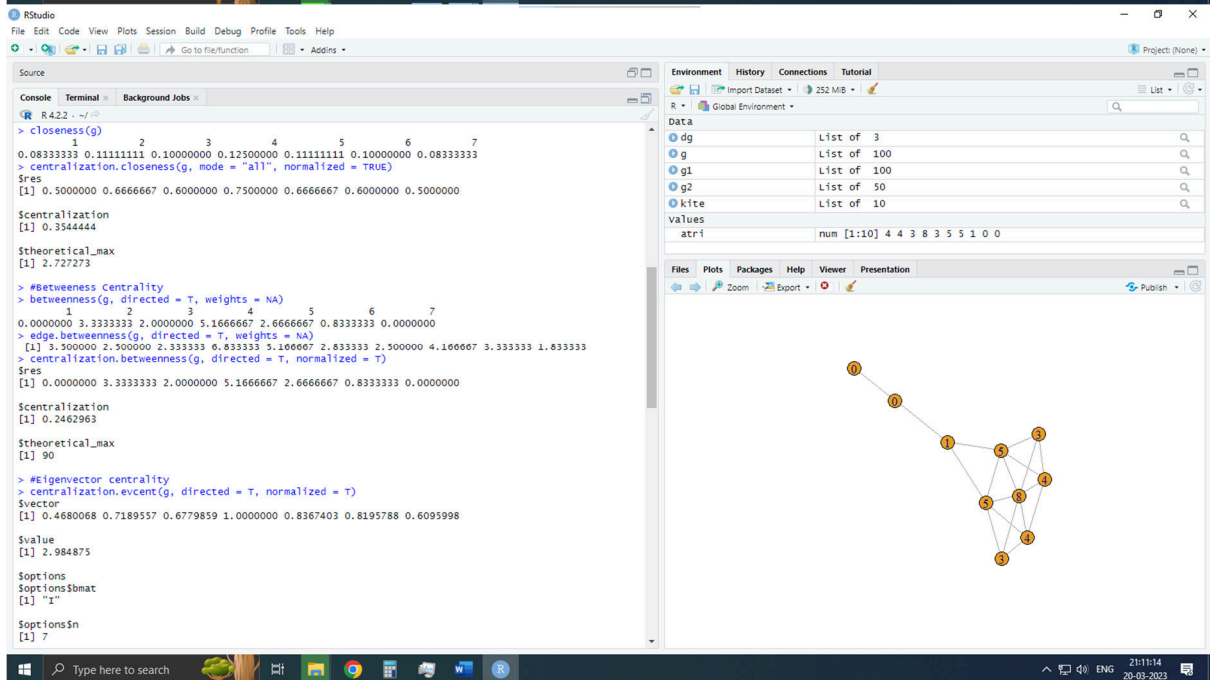
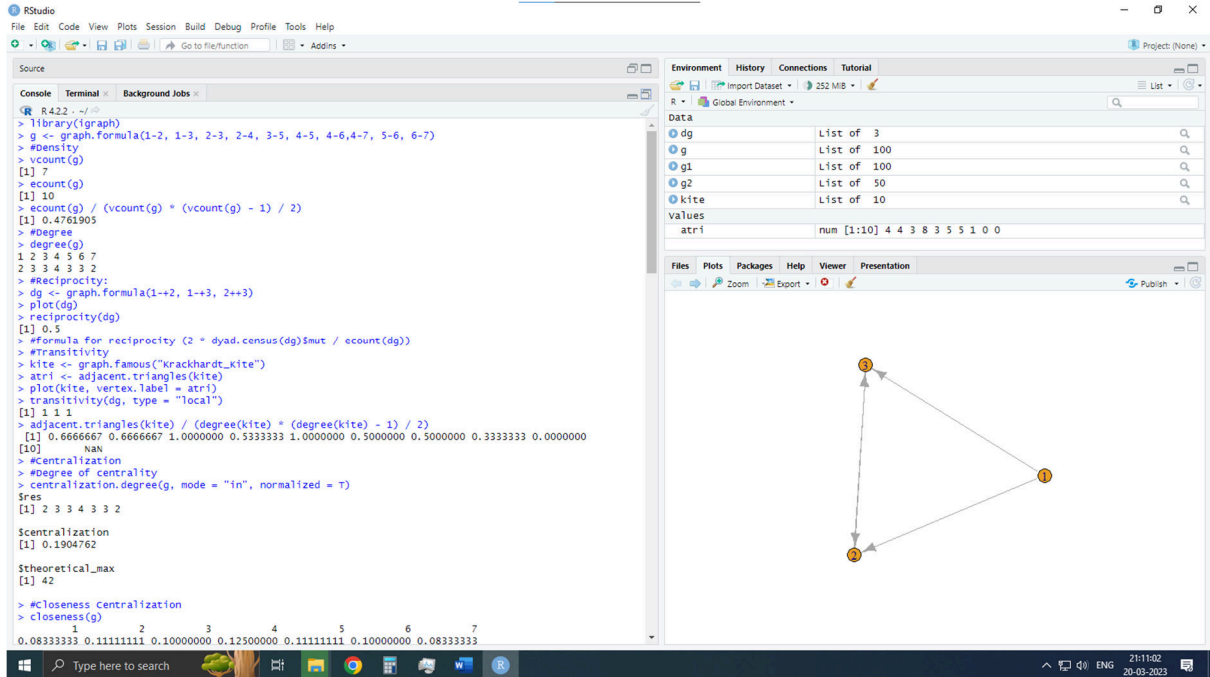
# Clustering
# Create two graphs 'g1' and 'g2'
g2 <- barabasi.game(50, p = 2, directed = F)
g1 <- watts.strogatz.game(1, size = 100, nei = 5, p = 0.05)

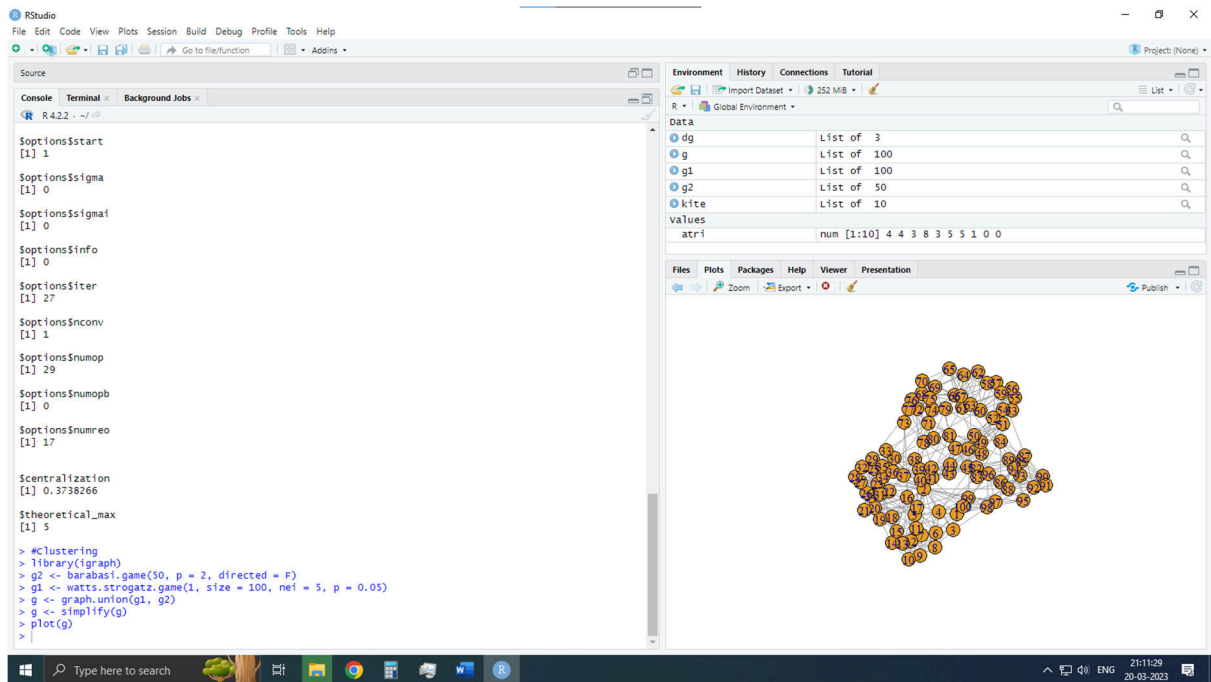
# Combine the two graphs 'g1' and 'g2'
g <- graph.union(g1, g2)

# Simplify the combined graph 'g'
g <- simplify(g)

# Plot the simplified graph 'g'
plot(g)
```

OUTPUT





Practical 4: For a given network find the following: (i) Length of the shortest path from a given node to another node; (ii) the density of the graph

Code:

```
library(igraph)

# creating a matrix from a table
matt <- as.matrix(read.table(text=
      "node  R  S  T  U
      R  7  5  0  0
      S  7  0  0  2
      T  0  6  0  0
      U  4  0  1  0", header=T))

# storing the row names in nms and removing the first column
nms <- matt[,1]
matt <- matt[, -1]

# setting the column and row names to be the same
colnames(matt) <- rownames(matt) <- nms

# replacing NA values with 0
matt[is.na(matt)] <- 0

# creating a weighted graph from the matrix
g <- graph.adjacency(matt, weighted=TRUE)

# plotting the graph
plot(g)

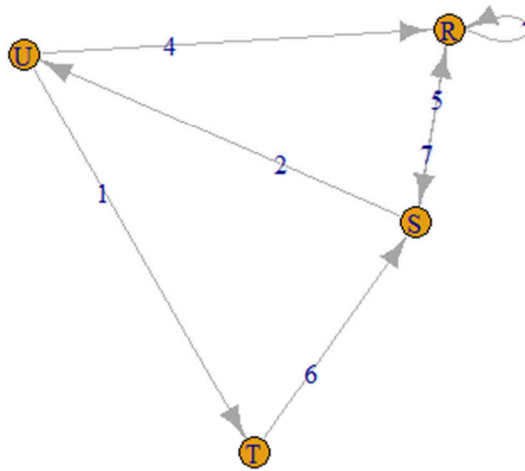
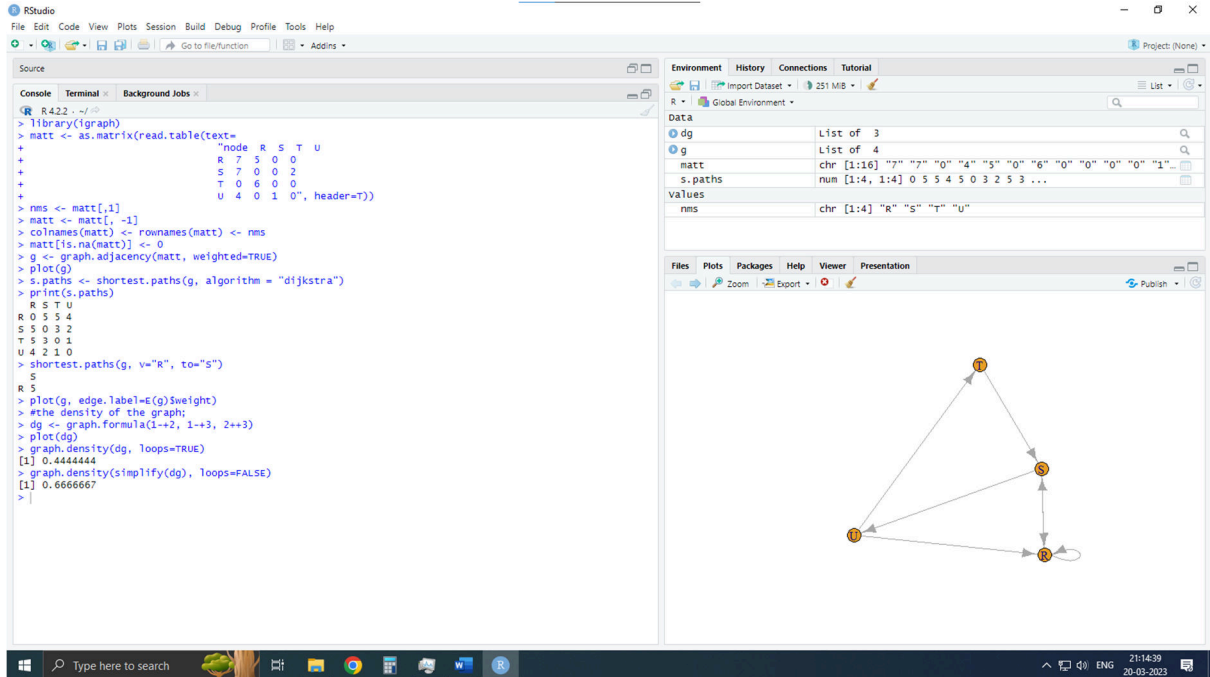
# calculating the shortest paths between all pairs of nodes
s.paths <- shortest.paths(g, algorithm = "dijkstra")
print(s.paths)

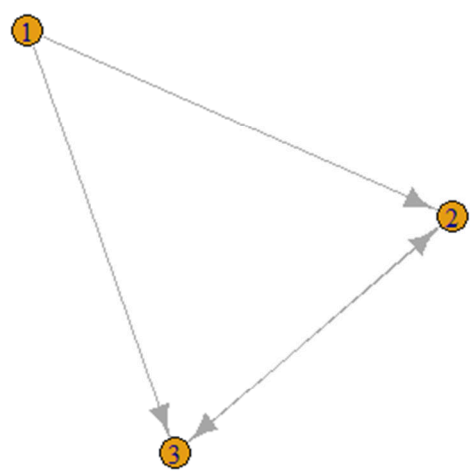
# calculating the shortest path between R and S
shortest.paths(g, v="R", to="S")

# plotting the graph with edge weights as labels
plot(g, edge.label=E(g)$weight)

# calculating the density of the graph
dg <- graph.formula(1-+2, 1-+3, 2++3)
plot(dg)
graph.density(dg, loops=TRUE)
graph.density(simplify(dg), loops=FALSE)
```

OUTPUT





Practical 5: Write a program to distinguish between a network as a matrix, a network as an edge list, and a network as a sociogram (or “network graph”) using 3 distinct networks representatives of each.

Code:

```
# Load igraph package
library(igraph)

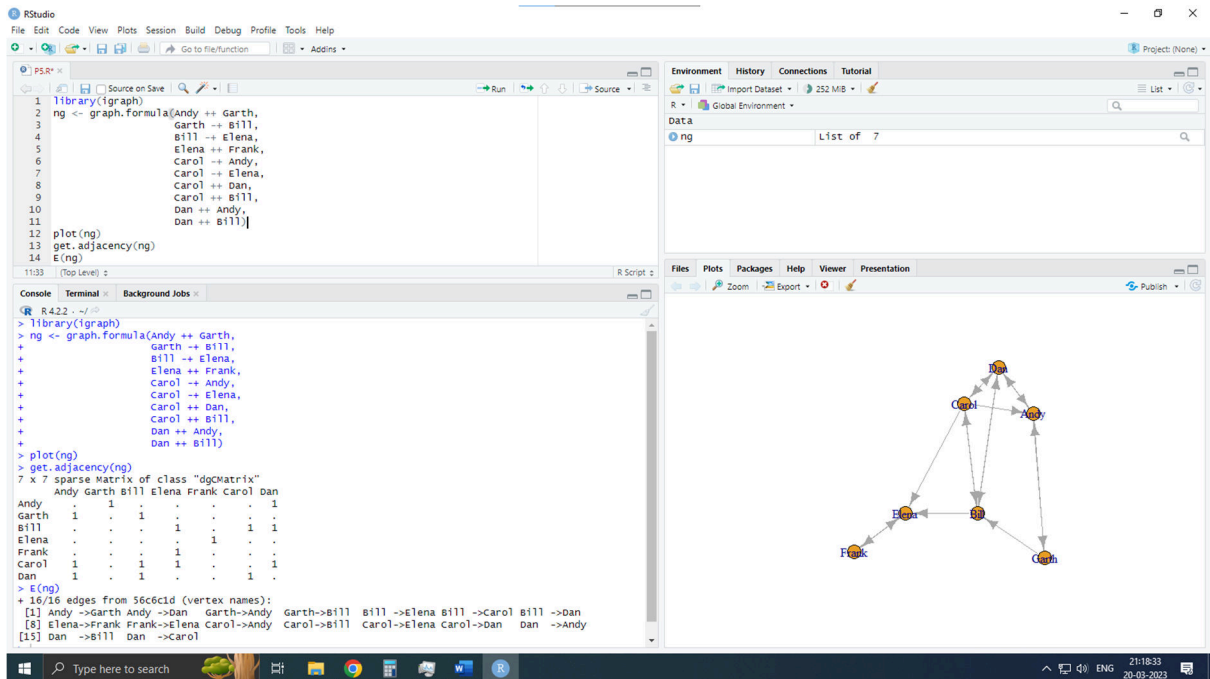
# Define network using graph formula notation
ng <- graph.formula(Andy ++ Garth,
                   Garth -+ Bill,
                   Bill -+ Elena,
                   Elena ++ Frank,
                   Carol -+ Andy,
                   Carol -+ Elena,
                   Carol ++ Dan,
                   Carol ++ Bill,
                   Dan ++ Andy,
                   Dan ++ Bill)

# Plot the network
plot(ng)

# Display adjacency matrix of the network
get.adjacency(ng)

# Display edges of the network
E(ng)
```

OUTPUT



Practical 6: Write a program to exhibit structural equivalence, automorphic equivalence, and regular equivalence from a network.

Codes:

```
# Install and load necessary packages
install.packages("sna")
library(sna)
library(igraph)

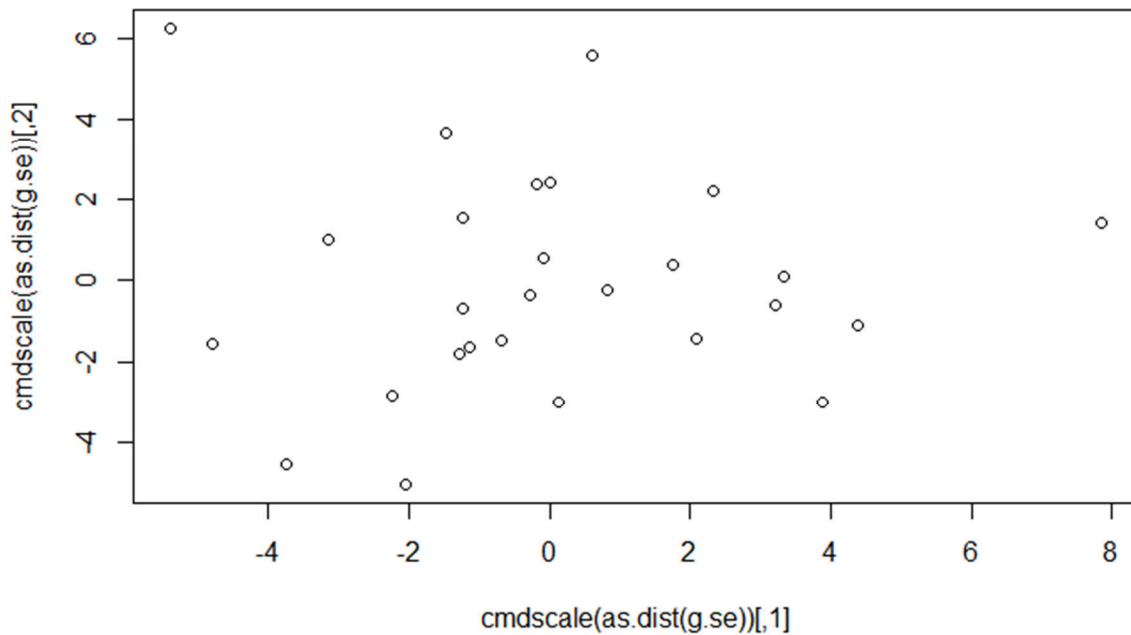
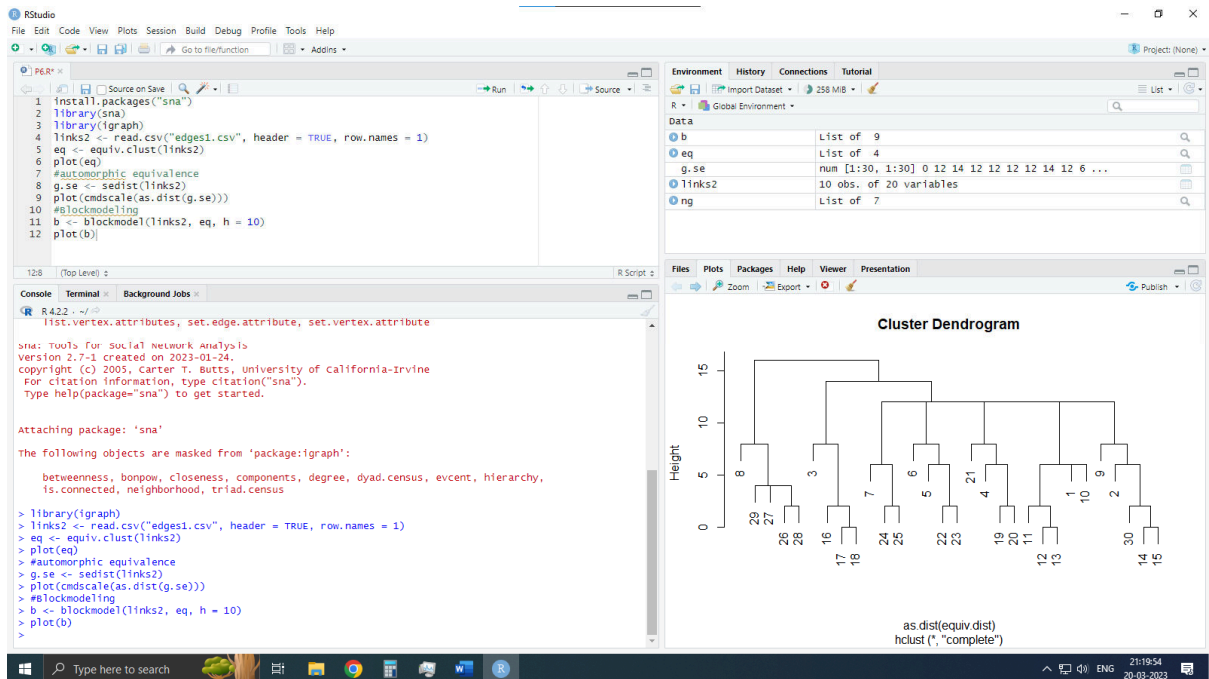
# Read data from file
links2 <- read.csv("edges1.csv", header = TRUE, row.names = 1)

# Equivalence clustering
eq <- equiv.clust(links2)
plot(eq)

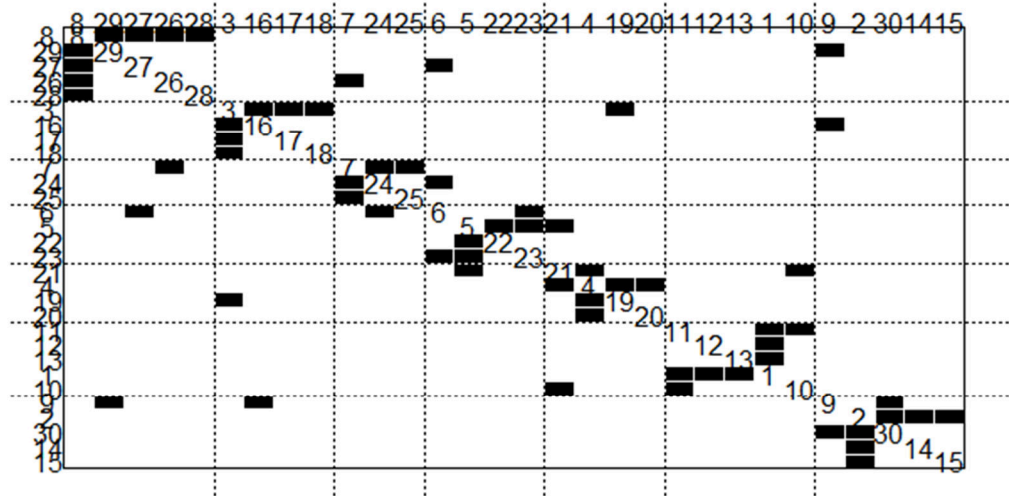
# Automorphic equivalence
g.se <- sedist(links2)
plot(cmdscale(as.dist(g.se)))

# Blockmodeling
b <- blockmodel(links2, eq, h = 10)
plot(b)
```

OUTPUT



Relation - 1



Practical 7: Perform SVD analysis on network

Code:

```
# Load the igraph library
library(igraph)

# Create a 9x4 matrix with specific values
a <- matrix(c(1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 0, 0, 0, 0, 0,
              0, 0, 0, 0, 1, 1, 1, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 1, 1, 1), 9, 4)

# Print the matrix to the console
print(a)

# Perform singular value decomposition on the matrix
svd(a)
```

OUTPUT

```
RStudio
File Edit Code View Plots Session Build Debug Profile Tools Help
Go to file/function Addins

Source
Console Terminal Background Jobs
R 4.2.2 ~ /
> a <- matrix(c(1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 0, 0, 0, 0, 0,
+              0, 0, 0, 0, 1, 1, 1, 0, 0, 0, 0, 0, 0, 0, 0, 1, 1, 1), 9, 4)
> print(a)
      [,1] [,2] [,3] [,4]
[1,]    1    1    0    0
[2,]    1    1    0    0
[3,]    1    1    0    0
[4,]    1    0    1    0
[5,]    1    0    1    0
[6,]    1    0    1    0
[7,]    1    0    0    1
[8,]    1    0    0    1
[9,]    1    0    0    1
> svd(a)
$d
[1] 3.464102e+00 1.732051e+00 1.732051e+00
[4] 1.922963e-16

$u
      [,1]      [,2]      [,3]
[1,] -0.3333333  0.4714045 -1.741269e-16
[2,] -0.3333333  0.4714045 -3.692621e-16
[3,] -0.3333333  0.4714045 -5.301858e-17
[4,] -0.3333333 -0.2357023 -4.082483e-01
[5,] -0.3333333 -0.2357023 -4.082483e-01
[6,] -0.3333333 -0.2357023 -4.082483e-01
[7,] -0.3333333 -0.2357023  4.082483e-01
[8,] -0.3333333 -0.2357023  4.082483e-01
[9,] -0.3333333 -0.2357023  4.082483e-01
      [,4]
[1,] 7.760882e-01
[2,] -1.683504e-01
[3,] -6.077378e-01
[4,] 6.774193e-17
[5,] 6.774193e-17
[6,] 6.774193e-17
[7,] 5.194768e-17
[8,] 5.194768e-17
[9,] 5.194768e-17

$V
      [,1]      [,2]      [,3] [,4]
[1,] -0.8660254  0.0000000 -4.378026e-17  0.5
[2,]  0.2886751  0.8164966  2.500507e-16  0.5
```