

Social network analysis with R sna package

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Different kinds of networks

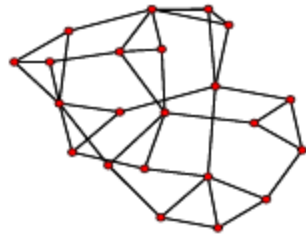
- Random graphs
 - a graph that is generated by some random process
- Scale free network
 - whose degree distribution follows a power law
- Small world
 - most nodes are not neighbors of one another, but most nodes can be reached from every other by a small number of hops or steps

Differ by Graph index

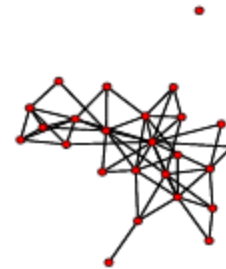
- Degree distribution
- average node-to-node distance
 - average shortest path length
- clustering coefficient
 - Global, local

network examples

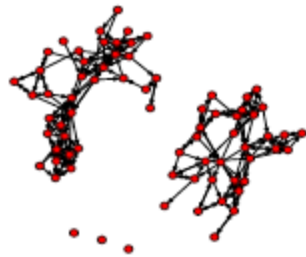
Taro Exchange



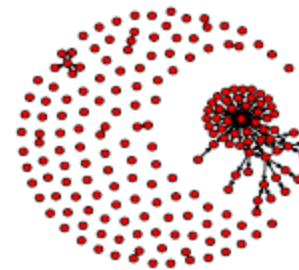
Texas SAR EMON



Coleman Friendship Network



Year 2000 MIDs



GLI-Graph level index

- Array statistic

- Mean
- Variance
- Standard deviation
- Skr
- ...

- Graph statistic

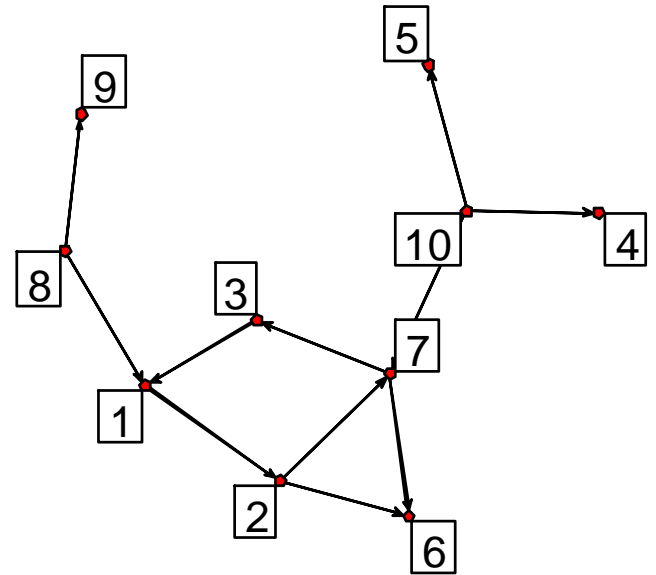
- Degree
- Density
- Reciprocity
- Centralization
- ...

Simple graph measurements

- Degree
 - Number of links to a vertex (indegree, outdegree...)
- Density
 - sum of tie values divided by the number of possible ties
- Reciprocity
 - the proportion of dyads which are symmetric
- Mutuality
 - the number of complete dyads
- Transitivity
 - the total number of transitive triads is computed

Example

- Degree
 - $\text{sum}(g) = 11$
- Density
 - $\text{gden}(g) = 11/90 = 0.1222$
- Reciprocity
 - $\text{grecip}(g, \text{measure}="dyadic") = 0.7556$
 - $\text{grecip}(g, \text{measure}="edgewise") = 0$
- Mutuality
 - $\text{mutuality}(g) = 0$
- Transtivity
 - $\text{gtrans}(g) = 0.1111$



Path and Cycle statistics

- `kpath.census`
- `kcycle.census`
 - `dyad.census`
 - `Triad.census`

Multi graph measurements

- Graph mean

- In dichotomous case. graph mean corresponds to graph's density

$$\bar{\delta}_H = \frac{1}{|V_U|^2} \sum_{x=1}^{|V_U|} \sum_{y=1}^{|V_U|} \delta_H(x, y)$$

- Graph covariance

- gcov/gscov

$$Cov(H_i, H_j) = \frac{1}{|V_U|^2} \sum_{x=1}^{|V_U|} \sum_{y=1}^{|V_U|} ((\delta_i(x, y) - \bar{\delta}_{H_i}) (\delta_j(x, y) - \bar{\delta}_{H_j}))$$

- Graph correlation

- gcor/gscor

$$\rho(H_i, H_j) = \frac{Cov(H_i, H_j)}{\sqrt{Var(H_i) Var(H_j)}}$$

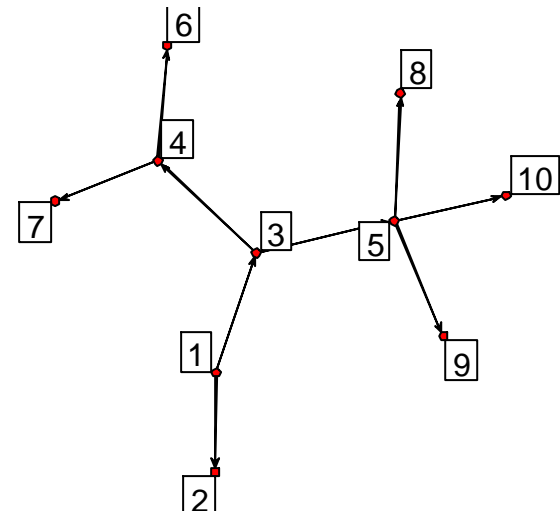
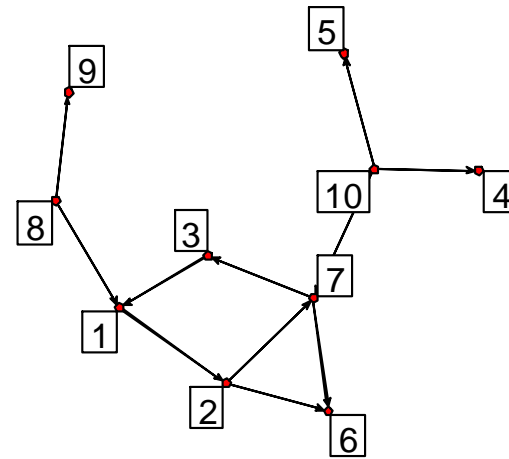
- Structural covariance

- unlabeled graph

$$Cov_S(G_i, G_j | \mathcal{P}_i, \mathcal{P}_j) = \max_{L_a \in \mathcal{P}_i, L_b \in \mathcal{P}_j} Cov(L_a(G_i), L_b(G_j))$$

Example

- $gcov(g1,g2) = -0.001123596$
- $gscov(g1,g2,exchange.list=1:10) = -0.001123596$
- $gscov(g1,g2)=0.04382022$
 - unlabeled graph
- $gcor(g1,g2) = -0.01130756$
- $gscor(g1,g2,exchange.list=1:10) = -0.01130756$
- $gscor(g1,g2) = 0.4409948$
 - unlabeled graph



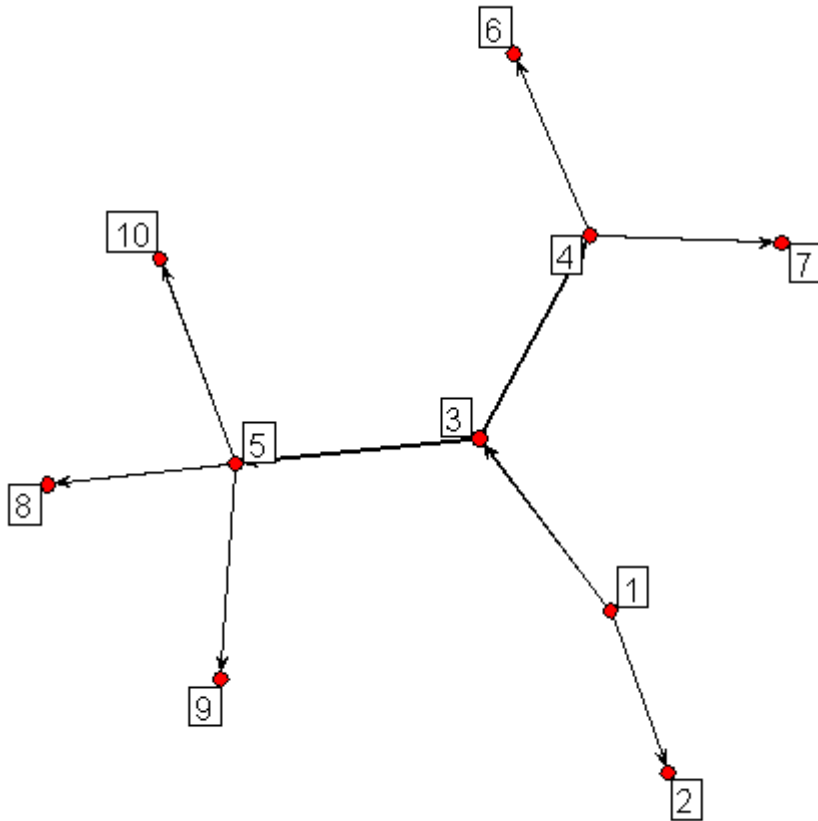
gCOV

```
> v=10
> G=rgraph(v)
> H=rgraph(v)
> g1=G-sum(G)/(v*(v-1))
> g2=H-sum(H)/(v*(v-1))
> diag(g1)=0
> diag(g2)=0
> sum(g1*g2)/(v*(v-1)-1)
[1] -0.01947566
> gcov(G,H)
[1] -0.01947566
> |
```

Measure of structure

- Connectedness $1 - \left(\frac{V}{N(N-1)/2} \right)$
 - '0' for no edges
 - '1' for $\binom{N}{2}$ edges
- Hierarchy $1 - \left(\frac{V}{MaxV} \right)$
 - '0' for all two-way links
 - '1' for all one-way links
- Efficiency $1 - \left(\frac{V}{MaxV} \right)$
 - '0' for $\binom{N}{2}$ edges
 - '1' for N-1 edges
- Least Upper Boundedness (lubness)
 - '0' for all vertex link into one
 - '1' for all outtree

Example



- Outtree
 - Connectedness=1
 - Hierarchy=1
 - Efficiency=1
 - Lubness=1

Graph centrality

- Degree
 - Number of links to a vertex(indegree, outdegree...)
- Betweenness
 - Number of shortest paths pass it
- Closeness
 - Length to all other vertices
- Centralization by 3 ways above
 - '0' for all vertices has equal position(central score)
 - '1' for 1 vertex be the center of the graph
- See also
 - evcent, bonpow, graphcent, infocent, prestige

Example

```
> centralization(g,degree,mode="graph")
```

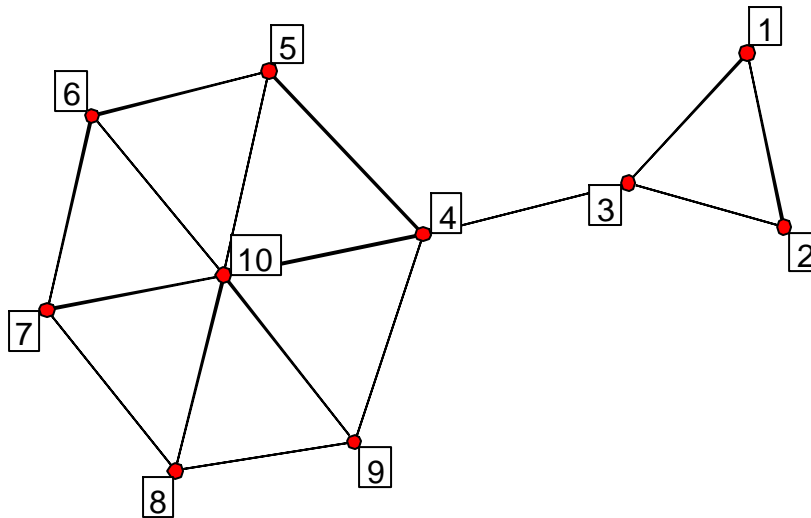
```
[1] 0.1944444
```

```
> centralization(g,betweenness,mode="graph")
```

```
[1] 0.1026235
```

```
> centralization(g,closeness,mode="graph")
```

```
[1] 0
```



Mode="graph" means only consider indegree

Example

```
> centralization(g,degree,mode="graph")
```

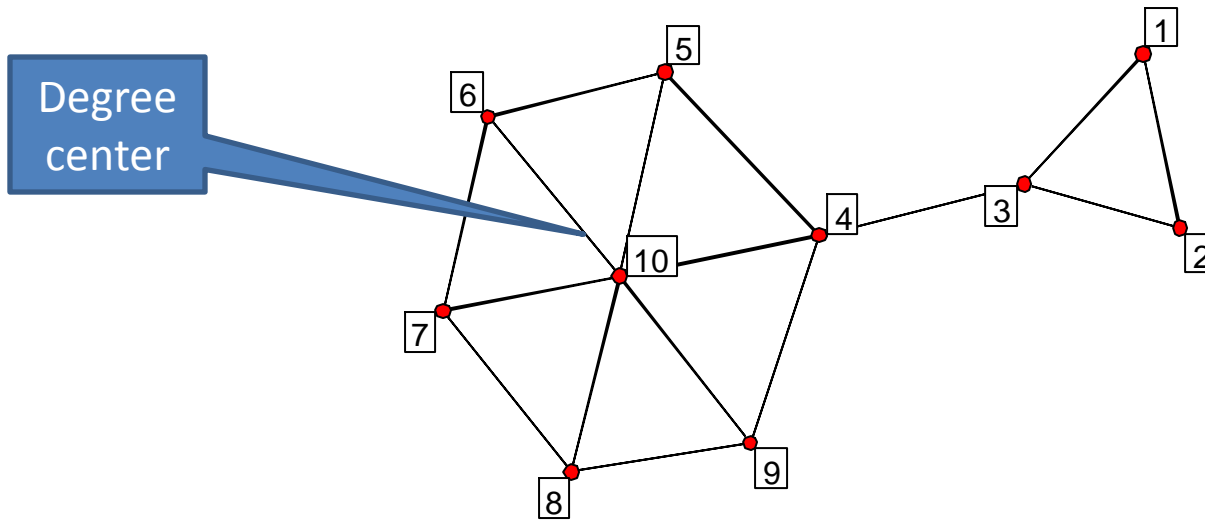
```
[1] 0.1944444
```

```
> centralization(g,betweenness,mode="graph")
```

```
[1] 0.1026235
```

```
> centralization(g,closeness,mode="graph")
```

```
[1] 0
```



Mode="graph" means only consider indegree

Example

```
> centralization(g,degree,mode="graph")
```

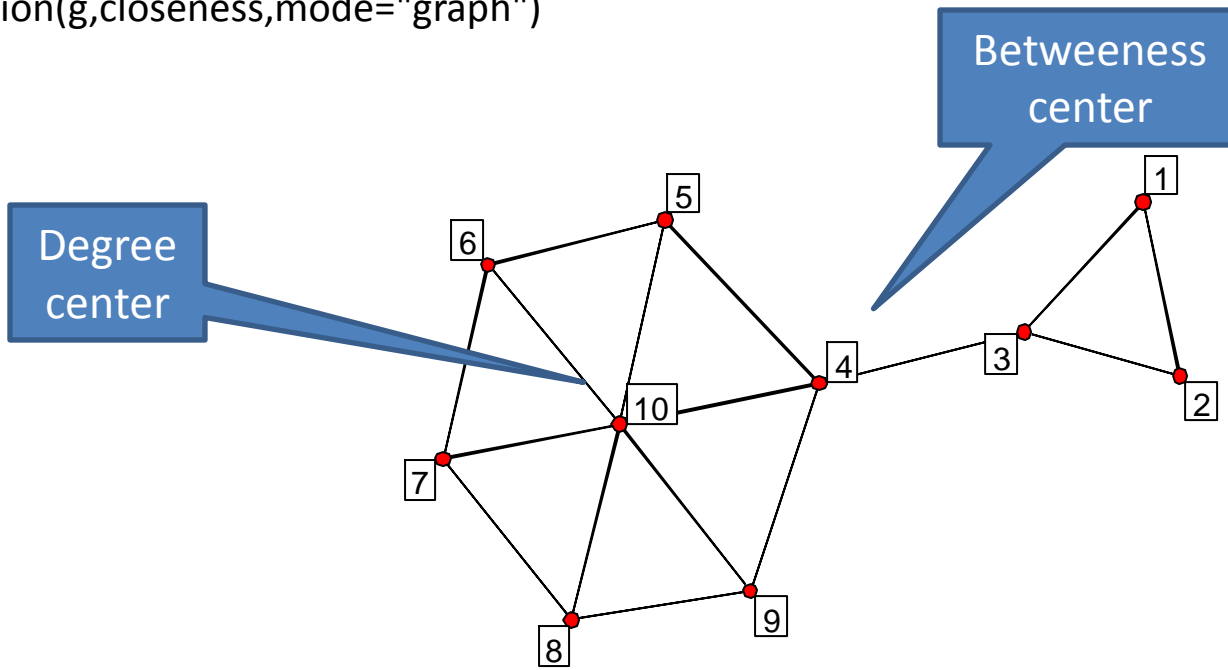
```
[1] 0.1944444
```

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> centralization(g,betweenness,mode="graph")
```

```
[1] 0.1026235
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> centralization(g,closeness,mode="graph")
```

```
[1] 0
```



Mode="graph" means only consider indegree

Example

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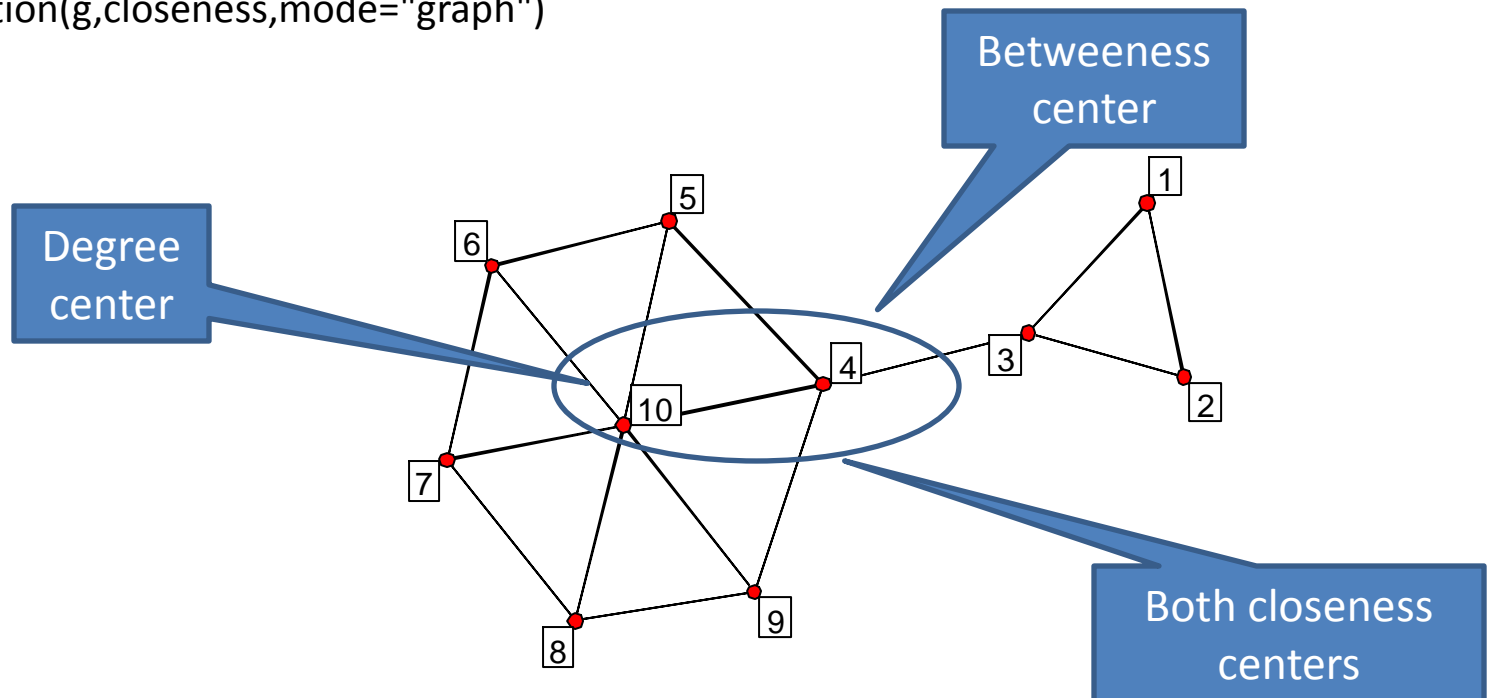
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[1] 0.1944444
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> centralization(g,betweenness,mode="graph")
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[1] 0.1026235
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```

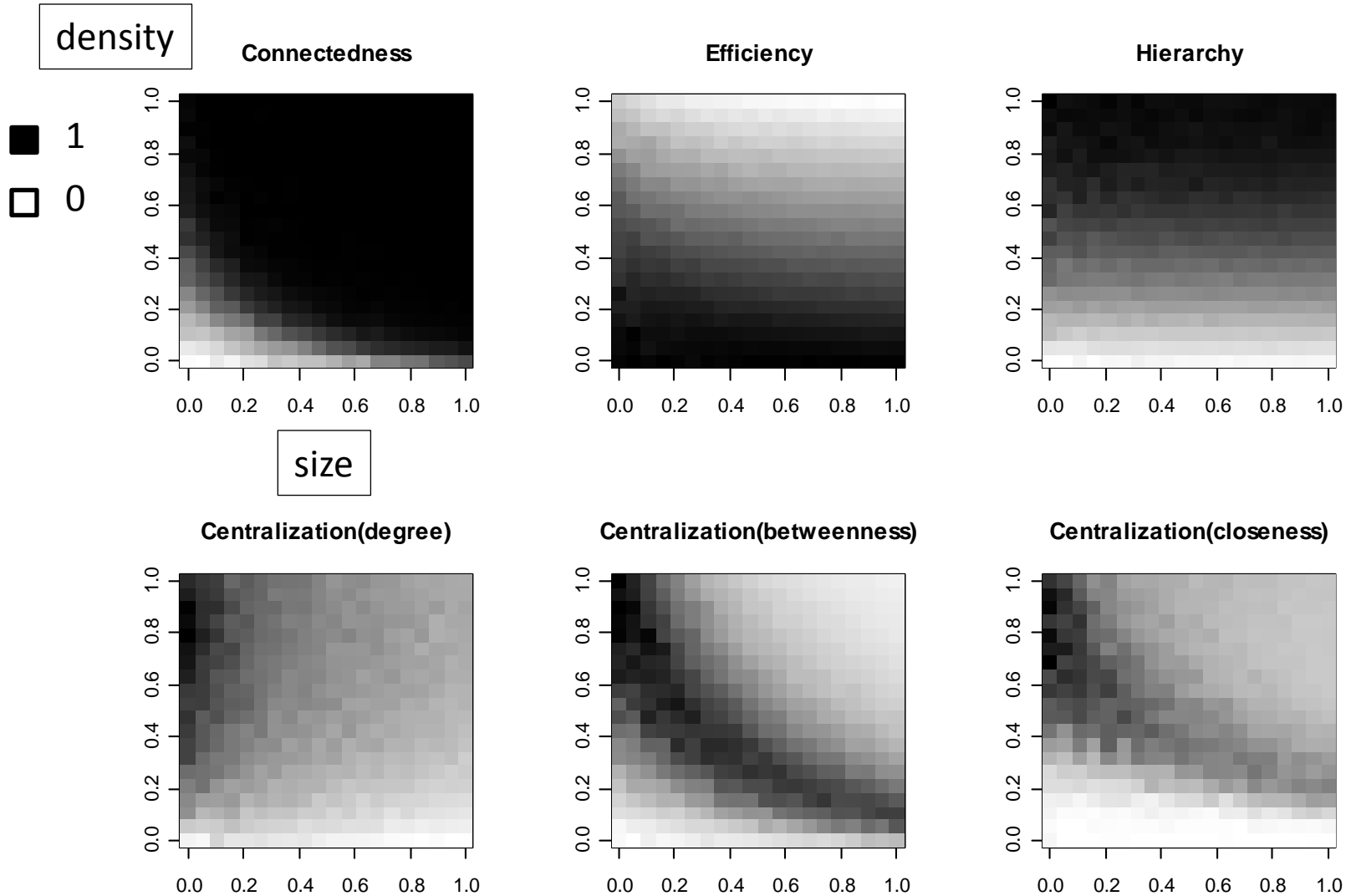
```
[1] 0
```



Mode="graph" means only consider indegree

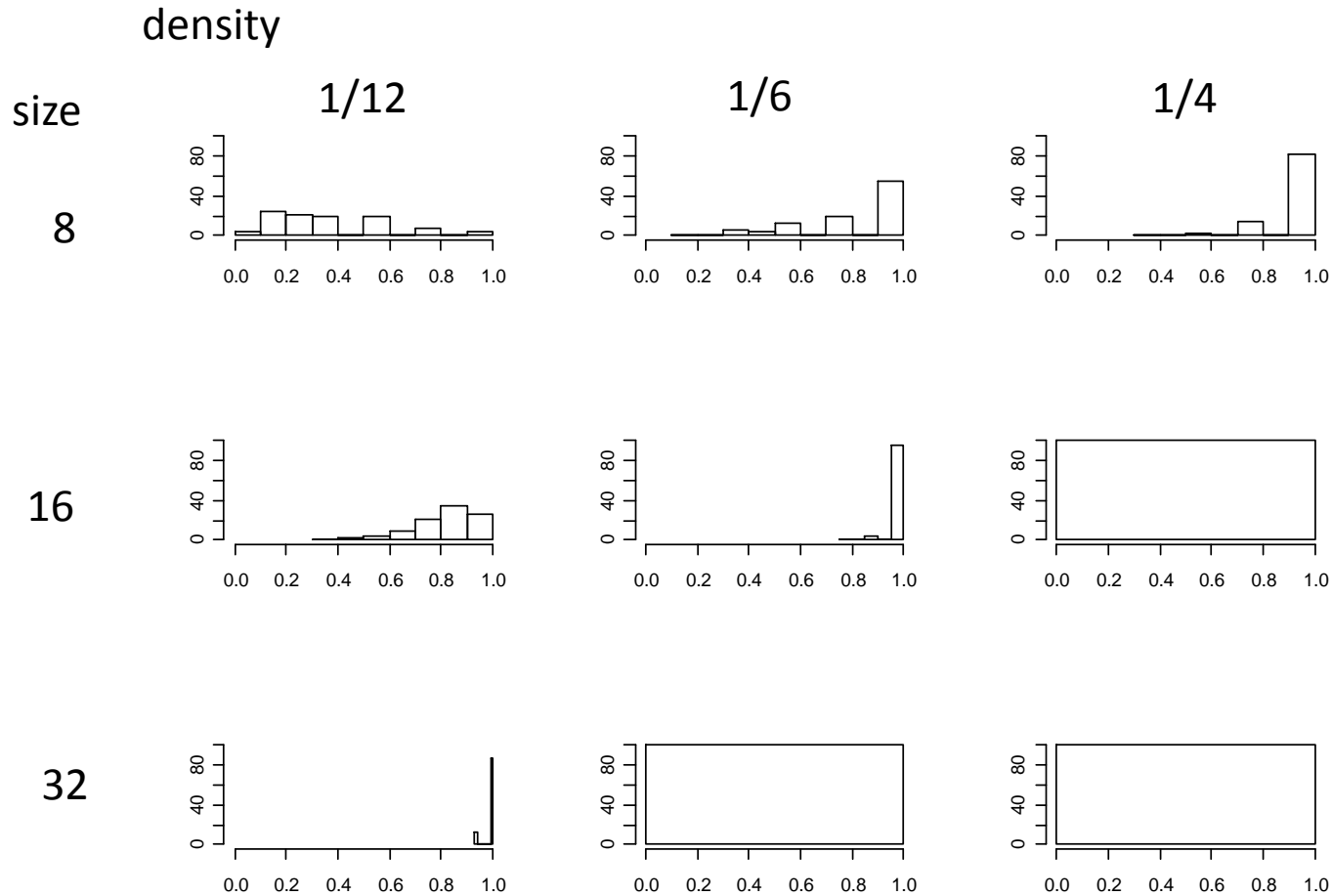
GLI relation

GLI map

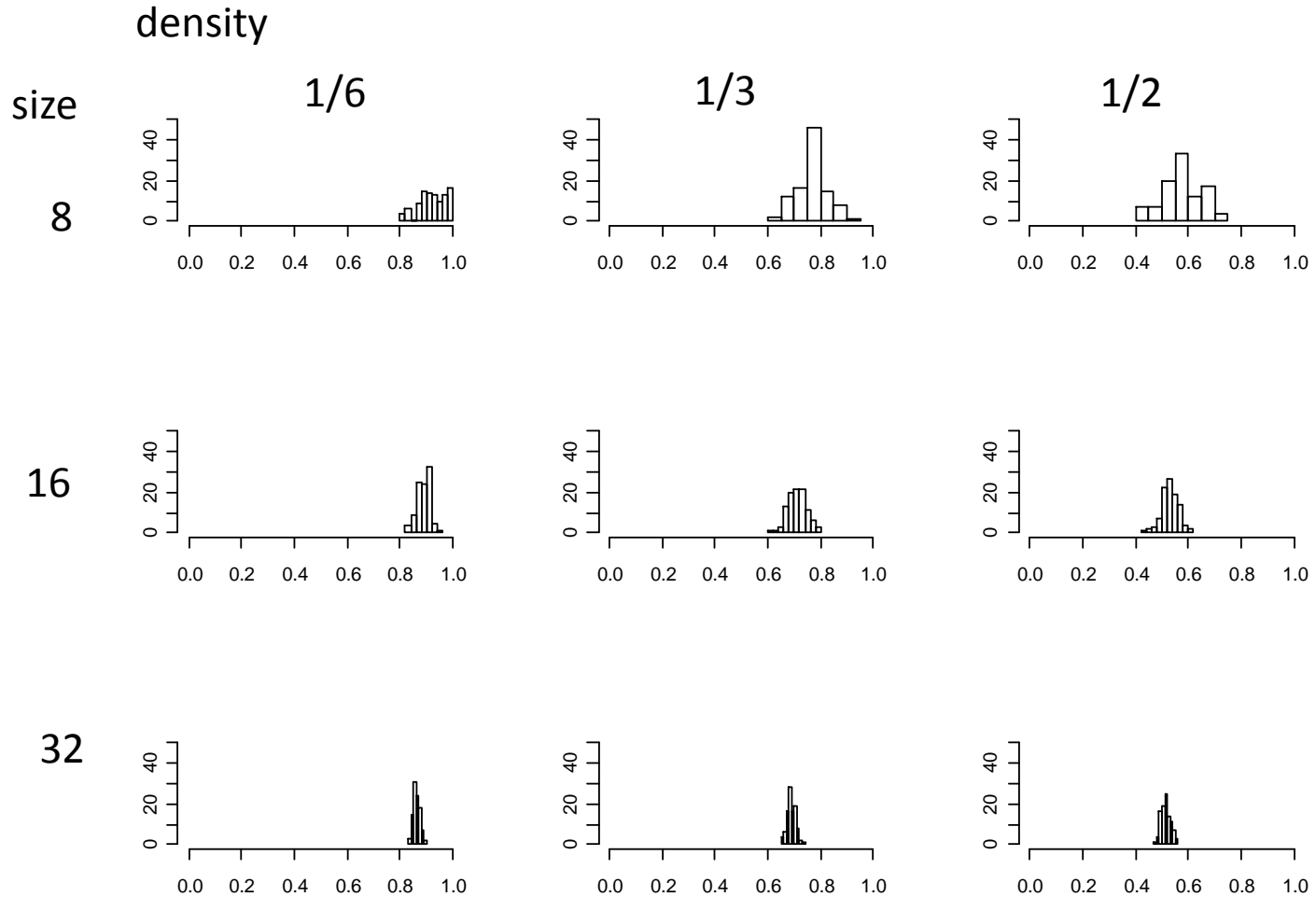


Anderson, B.S.; Butts, C. T.; and Carley, K. M. (1999). "The Interaction of Size and Density with Graph-Level Indices."

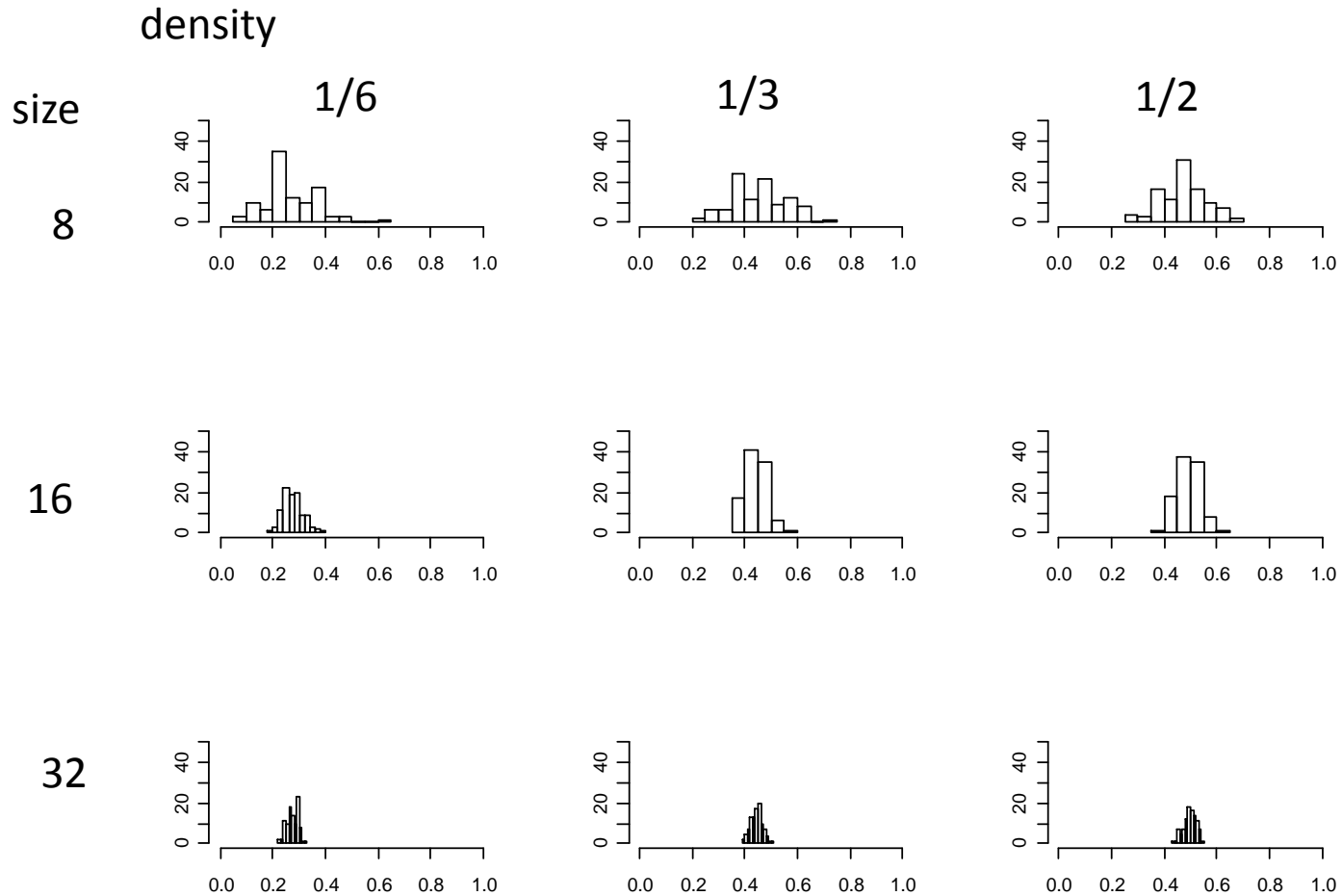
connectedness distribution by graph size and density



efficiency distribution by graph size and density



hierarchy distribution by graph size and density



GLI map R code

```
compare<-function(size,den)
{
  g=rgraph(n=size,m=100,tprob=den)
  gli1=apply(g,1,connectedness)
  gli2=apply(g,1,efficiency)
  gli3=apply(g,1,hierarchy)
  gli4=apply(g,1,function(x) centralization(x,degree))
  gli5=apply(g,1,function(x) centralization(x,betweenness))
  gli6=apply(g,1,function(x) centralization(x,closeness))
  x1=mean(gli1,na.rm=T)
  x2=mean(gli2,na.rm=T)
  x3=mean(gli3,na.rm=T)
  x4=mean(gli4,na.rm=T)
  x5=mean(gli5,na.rm=T)
  x6=mean(gli6,na.rm=T)
  return(c(x1,x2,x3,x4,x5,x6))
}
```

```
nx=20
ny=20
res=array(0,c(nx,ny,6))
size=5:26
den=seq(0.05,0.5,length.out=20)
for(i in 1:nx)
  for(j in 1:ny)
    res[i,j]=compare(size[i],den[j])

#image(res,col=gray(1000:1/1000))

par(mfrow=c(2,3))
image(res[,1],col=gray(1000:1/1000),main="Connectedness")
image(res[,2],col=gray(1000:1/1000),main="Efficiency")
image(res[,3],col=gray(1000:1/1000),main="Hierarchy")
image(res[,4],col=gray(1000:1/1000),main="Centralization(degree)")
image(res[,5],col=gray(1000:1/1000),main="Centralization(betweenness)")
image(res[,6],col=gray(1000:1/1000),main="Centralization(closeness)")
```

GLI distribution R code

```
par(mfrow=c(3,3))
for(i in 1:3)
  for(j in 1:3)
    hist(centralization(rgraph(4*2^i,100,tprob=j/4),betweenness),main="",xlab="",ylab="",xlim=range(0:1),ylim=range(0:50))
    hist(centralization(rgraph(4*2^i,100,tprob=j/4),degree),main="",xlab="",ylab="",xlim=range(0:1),ylim=range(0:50))
    hist(hierarchy(rgraph(4*2^i,100,tprob=j/6)),main="",xlab="",ylab="",xlim=range(0:1),ylim=range(0:50))
    hist(efficiency(rgraph(4*2^i,100,tprob=j/6)),main="",xlab="",ylab="",xlim=range(0:1),ylim=range(0:50))
    hist(connectedness(rgraph(4*2^i,100,tprob=j/12)),main="",xlab="",ylab="",xlim=range(0:1),ylim=range(0:100))
```

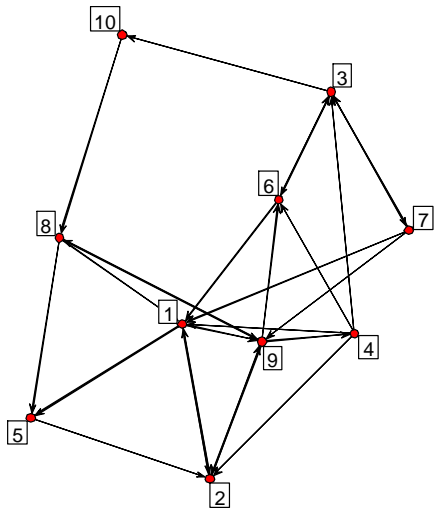
Graph distance

Clustering, MDS

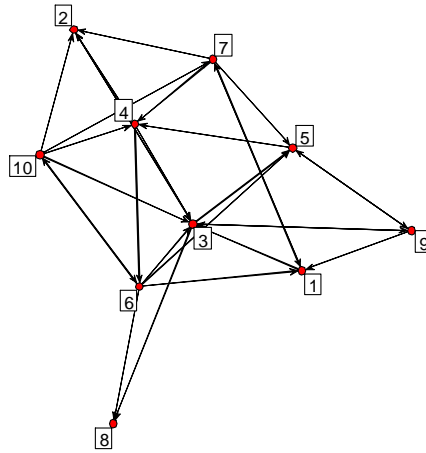
Distance between graphs

- Hamming(labeling) distance
 - $|\{e : (e \in E(G_1), e \notin E(G_2)) \wedge (e \notin E(G_1), e \in E(G_2))\}|$
number of addition/deletion operations required to turn the edge set of G1 into that of G2
 - ‘hdist’ for typical hamming distance matrix
- Structure distance
 - $d_S(G, H | L_G, L_H) = \min_{L_G, L_H} d(\ell(G), \ell(H))$
 - ‘structdist’ & ‘sdmat’ for structure distance with exchange.list of vertices

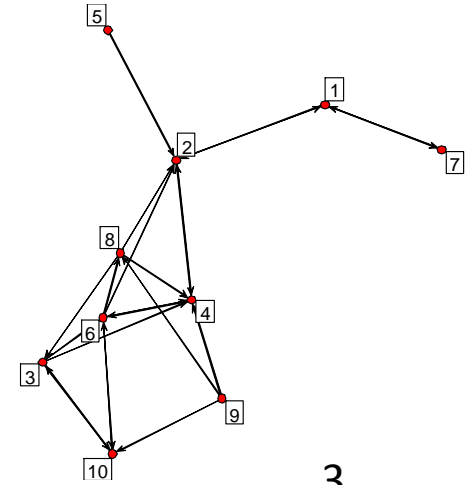
Example



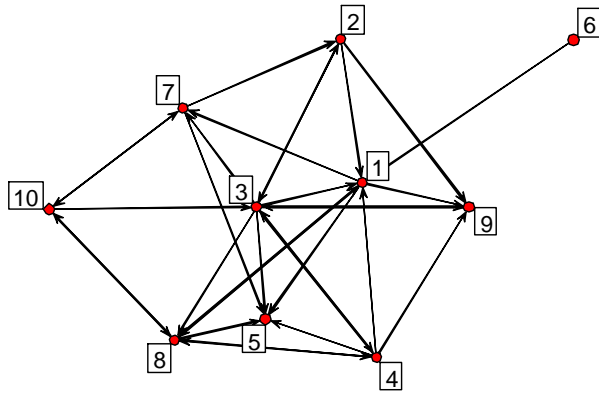
1



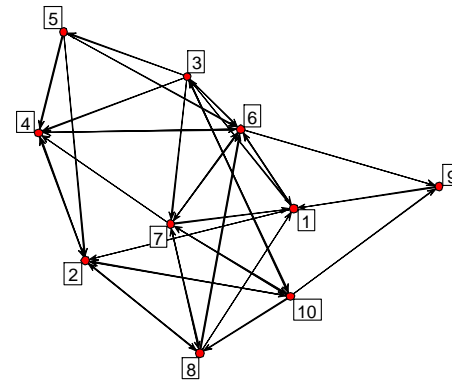
2



3



4



5

Example

hdist(g)

	1	2	3	4	5
1	0	44	29	35	39
2	44	0	35	35	39
3	29	35	0	44	34
4	35	35	44	0	48
5	39	39	34	48	0

sdmatrix(g)

	[, 1]	[, 2]	[, 3]	[, 4]	[, 5]
[1,]	0	24	23	25	27
[2,]	24	0	25	27	29
[3,]	23	25	0	26	28
[4,]	25	27	26	0	28
[5,]	27	29	28	28	0

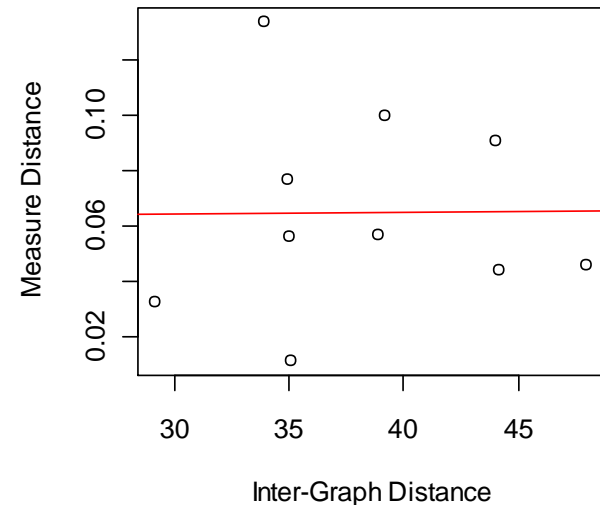
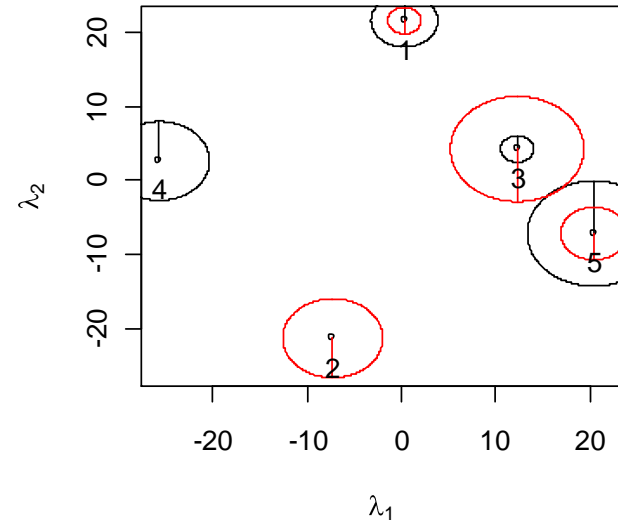
structdist(g)

	1	2	3	4	5
1	0	22	21	23	25
2	22	0	21	21	23
3	21	21	0	20	24
4	23	23	20	0	20
5	25	23	22	20	0

Inter-Graph MDS

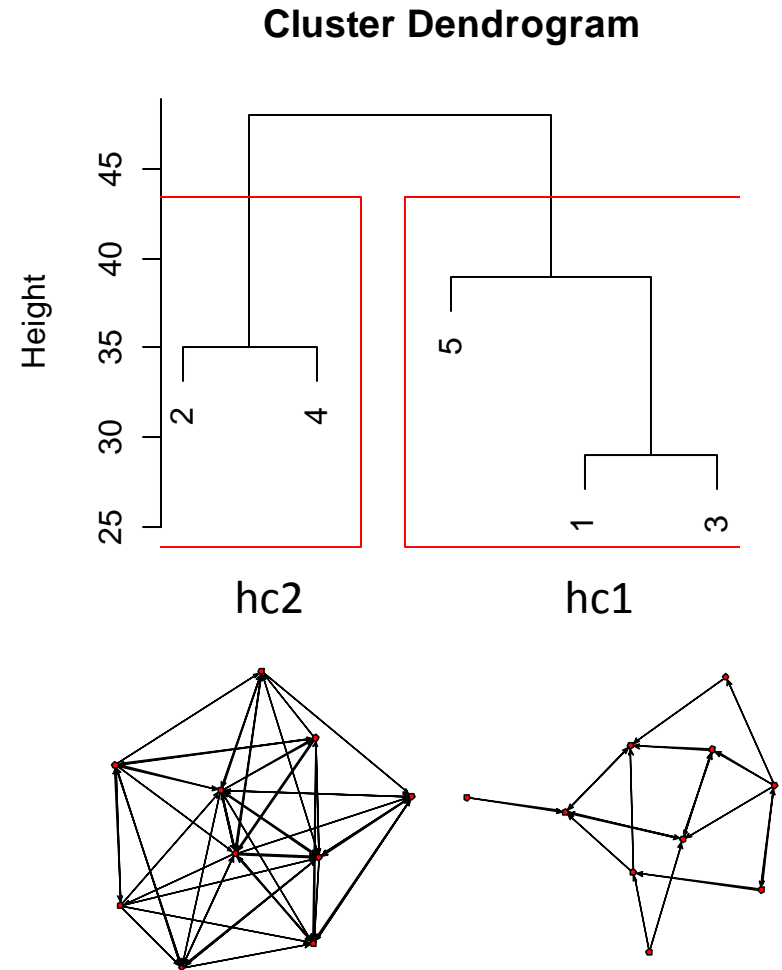
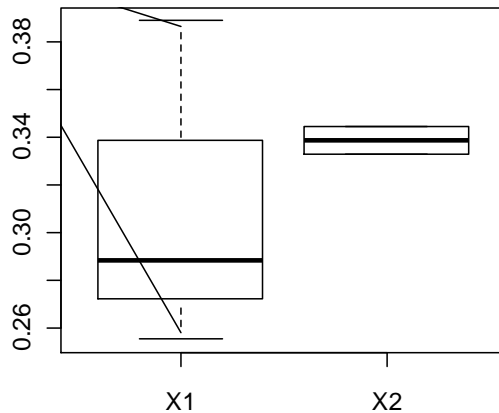
- ‘gdist.plotstats’
 - Plot by distances between graphs
 - Add graph level index as third or fourth dimension

```
> g.h<-hdist(g) #sample graph used before  
> gdist.plotdiff(g.h,gden(g),lm.line=TRUE)  
> gdist.plotstats(g.h,cbind(gden(g),grecip(g)))
```



Graph clustering

- Use hamming distance
 - `g.h=hdist(g)`
 - `g.c<-hclust(as.dist(g.h))`
 - `rect.hclust(g.c,2)`
 - `g.cg<-gclust.centralgraph(g.c,2,g)`
 - `gplot(g.cg[1,,])`
 - `gplot(g.cg[2,,])`
 - `gclust.boxstats(g.c,2,gden(g))`



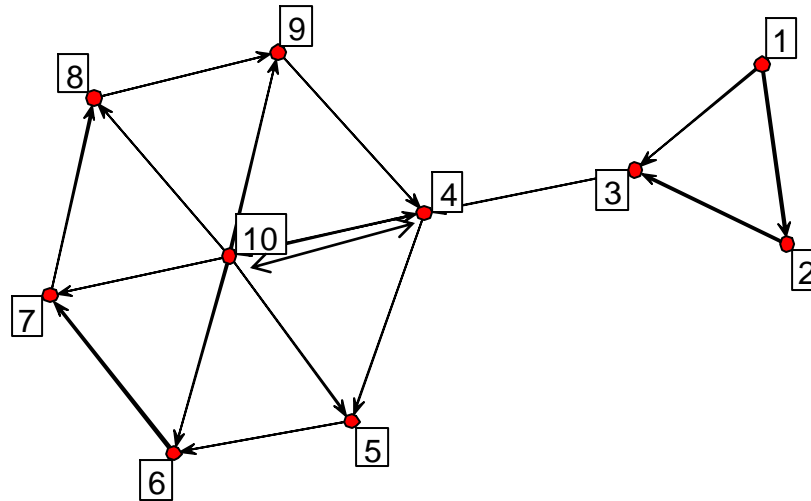
Distance between vertices

- Structural equivalence
 - ‘sedist’ with 4 methods:
 1. correlation: the product-moment correlation
 2. euclidean: the euclidean distance
 3. hamming: the Hamming distance
 4. gamma: the gamma correlation
- Path distance
 - ‘geodist’ with shortest path distance and the number of shortest pathes

Breiger, R.L.; Boorman, S.A.; and Arabie, P. (1975). “An Algorithm for Clustering Relational Data with Applications to Social Network Analysis and Comparison with Multidimensional Scaling.”

Brandes, U. (2000). “Faster Evaluation of Shortest-Path Based Centrality Indices.”

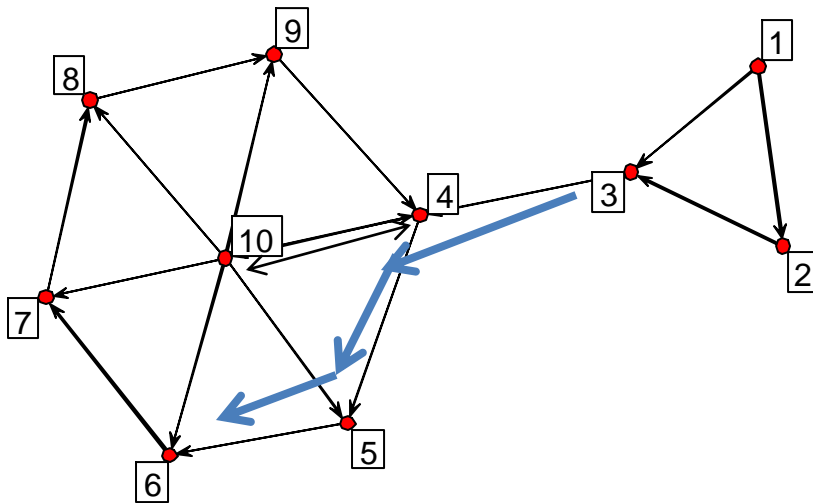
'sedist' Example



sedist(g) = sedist(g, mode="graph")

	[, 1]	[, 2]	[, 3]	[, 4]	[, 5]	[, 6]	[, 7]	[, 8]	[, 9]	[, 10]
[1,]	0	0	3	7	5	5	5	5	5	9
[2,]	0	0	1	7	5	5	5	5	5	9
[3,]	3	1	0	6	6	6	6	6	4	8
[4,]	7	7	6	0	4	6	6	6	4	6
[5,]	5	5	6	4	0	2	4	4	4	4
[6,]	5	5	6	6	2	0	2	4	4	6
[7,]	5	5	6	6	4	2	0	2	4	6
[8,]	5	5	6	6	4	4	2	0	2	6
[9,]	5	5	4	4	4	4	4	2	0	6
[10,]	9	9	8	6	4	6	6	6	6	0

'geodist' Example



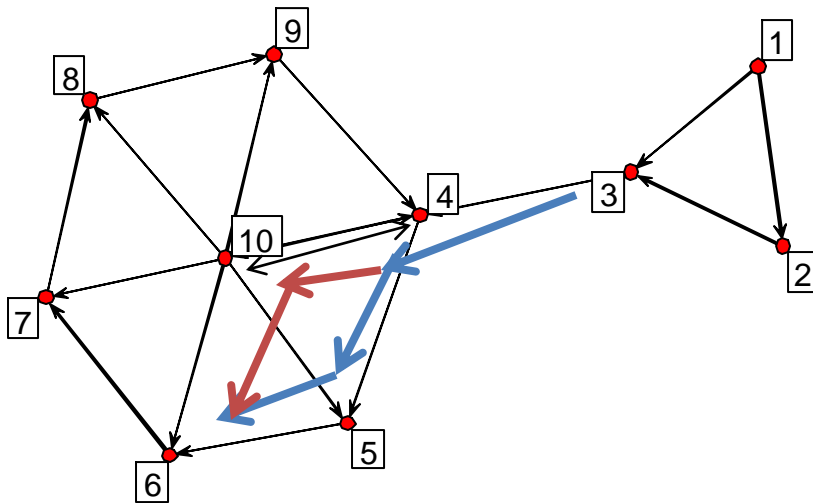
geodist(g)
\$counts

	[, 1]	[, 2]	[, 3]	[, 4]	[, 5]	[, 6]	[, 7]	[, 8]	[, 9]	[, 10]
[1,]	1	1	1	1	1	2	1	1	1	1
[2,]	0	1	1	1	1	2	1	1	1	1
[3,]	0	0	1	1	1	2	1	1	1	1
[4,]	0	0	0	1	1	2	1	1	1	1
[5,]	0	0	0	1	1	1	1	1	1	1
[6,]	0	0	0	1	1	1	1	1	1	1
[7,]	0	0	0	1	1	2	1	1	1	1
[8,]	0	0	0	1	1	2	1	1	1	1
[9,]	0	0	0	1	1	2	1	1	1	1
[10,]	0	0	0	1	1	1	1	1	1	1

\$gdist

	[, 1]	[, 2]	[, 3]	[, 4]	[, 5]	[, 6]	[, 7]	[, 8]	[, 9]	[, 10]
[1,]	0	1	1	2	3	4	4	4	4	3
[2,]	Inf	0	1	2	3	4	4	4	4	3
[3,]	Inf	Inf	0	1	2	3	3	3	3	2
[4,]	Inf	Inf	Inf	0	1	2	2	2	2	1
[5,]	Inf	Inf	Inf	5	0	1	2	3	4	6
[6,]	Inf	Inf	Inf	4	5	0	1	2	3	5
[7,]	Inf	Inf	Inf	3	4	5	0	1	2	4
[8,]	Inf	Inf	Inf	2	3	4	4	0	1	3
[9,]	Inf	Inf	Inf	1	2	3	3	3	0	2
[10,]	Inf	Inf	Inf	1	1	1	1	1	1	0

'geodist' Example



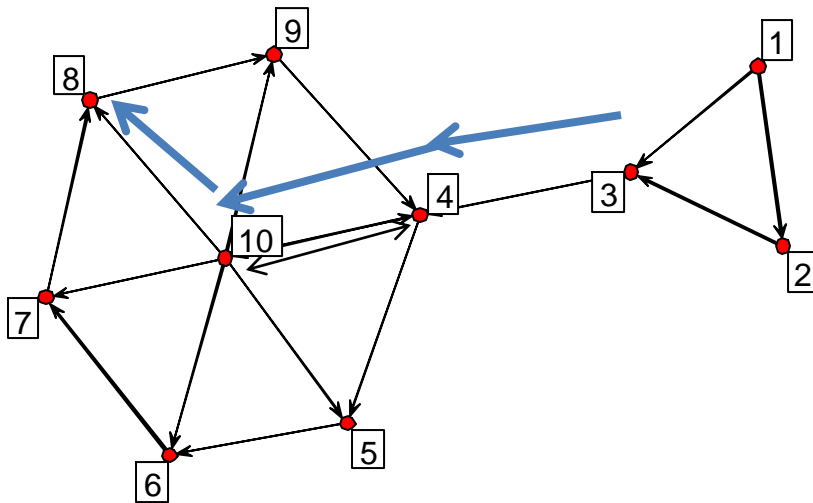
geodist(g)
\$counts

	[, 1]	[, 2]	[, 3]	[, 4]	[, 5]	[, 6]	[, 7]	[, 8]	[, 9]	[, 10]
[1,]	1	1	1	1	1	2	1	1	1	1
[2,]	0	1	1	1	1	2	1	1	1	1
[3,]	0	0	1	1	1	2	1	1	1	1
[4,]	0	0	0	1	1	2	1	1	1	1
[5,]	0	0	0	1	1	1	1	1	1	1
[6,]	0	0	0	1	1	1	1	1	1	1
[7,]	0	0	0	1	1	2	1	1	1	1
[8,]	0	0	0	1	1	2	1	1	1	1
[9,]	0	0	0	1	1	2	1	1	1	1
[10,]	0	0	0	1	1	1	1	1	1	1

\$gdist

	[, 1]	[, 2]	[, 3]	[, 4]	[, 5]	[, 6]	[, 7]	[, 8]	[, 9]	[, 10]
[1,]	0	1	1	2	3	4	4	4	4	3
[2,]	Inf	0	1	2	3	4	4	4	4	3
[3,]	Inf	Inf	0	1	2	3	3	3	3	2
[4,]	Inf	Inf	Inf	0	1	2	2	2	2	1
[5,]	Inf	Inf	Inf	5	0	1	2	3	4	6
[6,]	Inf	Inf	Inf	4	5	0	1	2	3	5
[7,]	Inf	Inf	Inf	3	4	5	0	1	2	4
[8,]	Inf	Inf	Inf	2	3	4	4	0	1	3
[9,]	Inf	Inf	Inf	1	2	3	3	3	0	2
[10,]	Inf	Inf	Inf	1	1	1	1	1	1	0

'geodist' Example



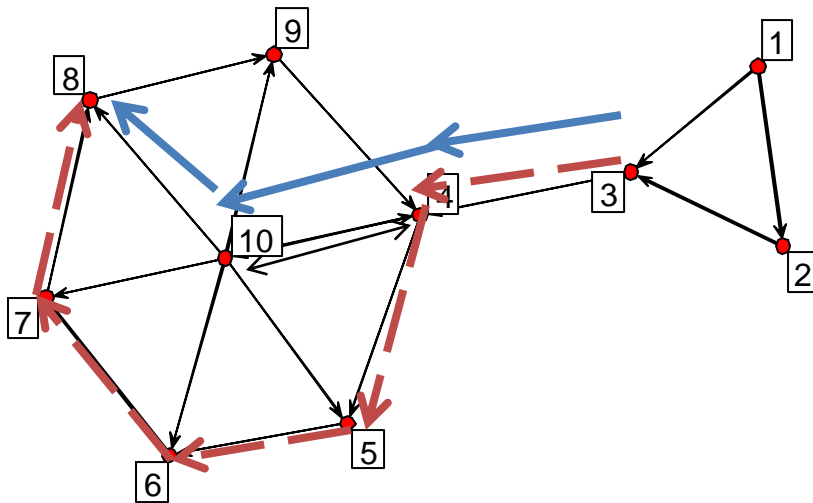
geodist(g)
\$counts

	[, 1]	[, 2]	[, 3]	[, 4]	[, 5]	[, 6]	[, 7]	[, 8]	[, 9]	[, 10]
[1,]	1	1	1	1	1	2	1	1	1	1
[2,]	0	1	1	1	1	2	1	1	1	1
[3,]	0	0	1	1	1	2	1	1	1	1
[4,]	0	0	0	1	1	2	1	1	1	1
[5,]	0	0	0	1	1	1	1	1	1	1
[6,]	0	0	0	1	1	1	1	1	1	1
[7,]	0	0	0	1	1	2	1	1	1	1
[8,]	0	0	0	1	1	2	1	1	1	1
[9,]	0	0	0	1	1	2	1	1	1	1
[10,]	0	0	0	1	1	1	1	1	1	1

\$gdist

	[, 1]	[, 2]	[, 3]	[, 4]	[, 5]	[, 6]	[, 7]	[, 8]	[, 9]	[, 10]
[1,]	0	1	1	2	3	4	4	4	4	3
[2,]	Inf	0	1	2	3	4	4	4	4	3
[3,]	Inf	Inf	0	1	2	3	3	3	3	2
[4,]	Inf	Inf	Inf	0	1	2	2	2	2	1
[5,]	Inf	Inf	Inf	5	0	1	2	3	4	6
[6,]	Inf	Inf	Inf	4	5	0	1	2	3	5
[7,]	Inf	Inf	Inf	3	4	5	0	1	2	4
[8,]	Inf	Inf	Inf	2	3	4	4	0	1	3
[9,]	Inf	Inf	Inf	1	2	3	3	3	0	2
[10,]	Inf	Inf	Inf	1	1	1	1	1	1	0

'geodist' Example



geodist(g)
\$counts

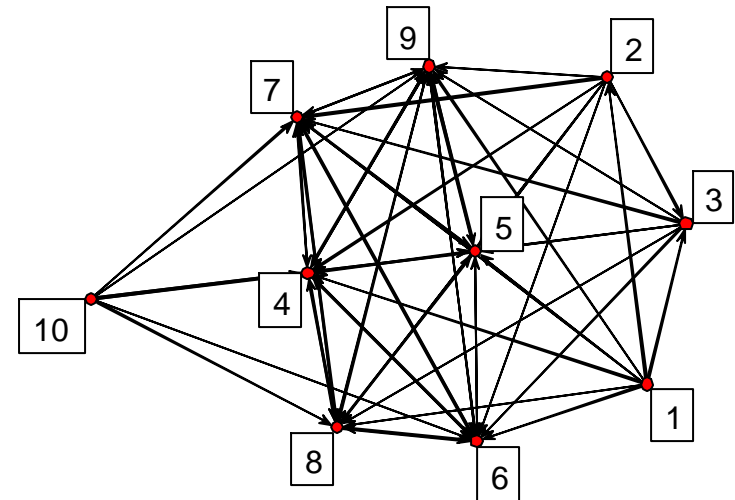
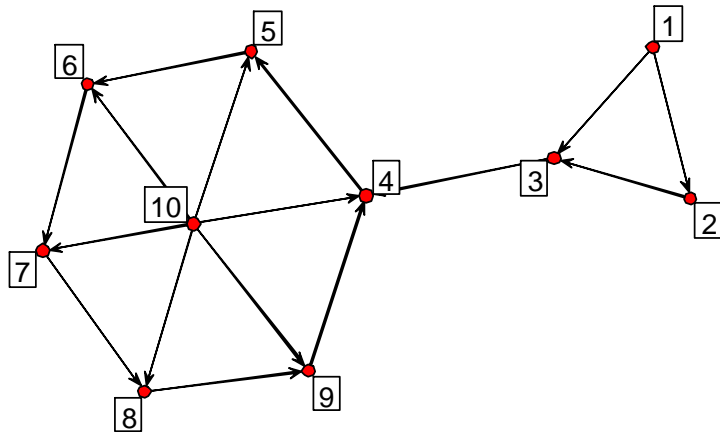
	[, 1]	[, 2]	[, 3]	[, 4]	[, 5]	[, 6]	[, 7]	[, 8]	[, 9]	[, 10]
[1,]	1	1	1	1	1	2	1	1	1	1
[2,]	0	1	1	1	1	2	1	1	1	1
[3,]	0	0	1	1	1	2	1	1	1	1
[4,]	0	0	0	1	1	2	1	1	1	1
[5,]	0	0	0	1	1	1	1	1	1	1
[6,]	0	0	0	1	1	1	1	1	1	1
[7,]	0	0	0	1	1	2	1	1	1	1
[8,]	0	0	0	1	1	2	1	1	1	1
[9,]	0	0	0	1	1	2	1	1	1	1
[10,]	0	0	0	1	1	1	1	1	1	1

\$gdist

	[, 1]	[, 2]	[, 3]	[, 4]	[, 5]	[, 6]	[, 7]	[, 8]	[, 9]	[, 10]
[1,]	0	1	1	2	3	4	4	4	4	3
[2,]	Inf	0	1	2	3	4	4	4	4	3
[3,]	Inf	Inf	0	1	2	3	3	3	3	2
[4,]	Inf	Inf	Inf	0	1	2	2	2	2	1
[5,]	Inf	Inf	Inf	5	0	1	2	3	4	6
[6,]	Inf	Inf	Inf	4	5	0	1	2	3	5
[7,]	Inf	Inf	Inf	3	4	5	0	1	2	4
[8,]	Inf	Inf	Inf	2	3	4	4	0	1	3
[9,]	Inf	Inf	Inf	1	2	3	3	3	0	2
[10,]	Inf	Inf	Inf	1	1	1	1	1	1	0

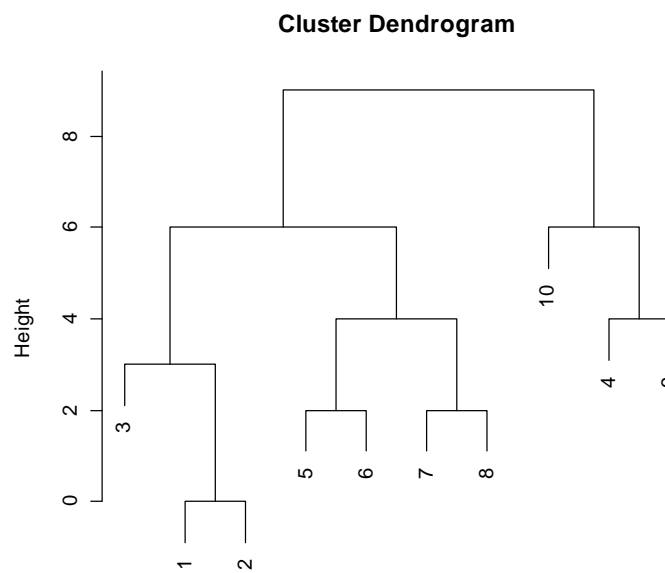
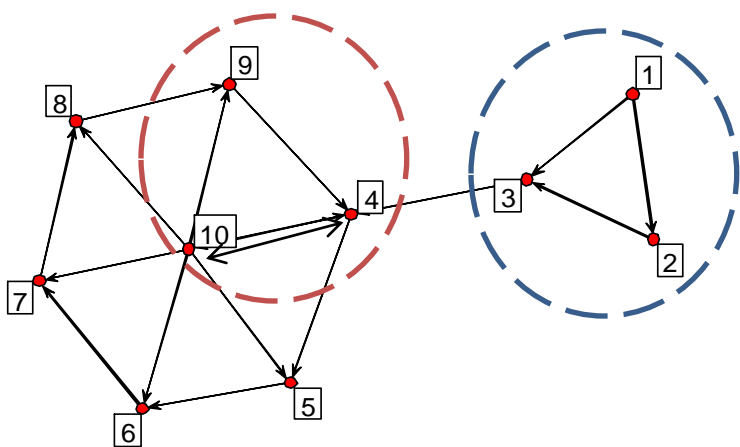
'geodist' reachability

- `gplot(reachability(g),label=1:10)`



Graph vertices clustering by 'sedist'

- General clustering methods
- 'equiv.clust' for vertices clustering by Structural equivalence('sedist')



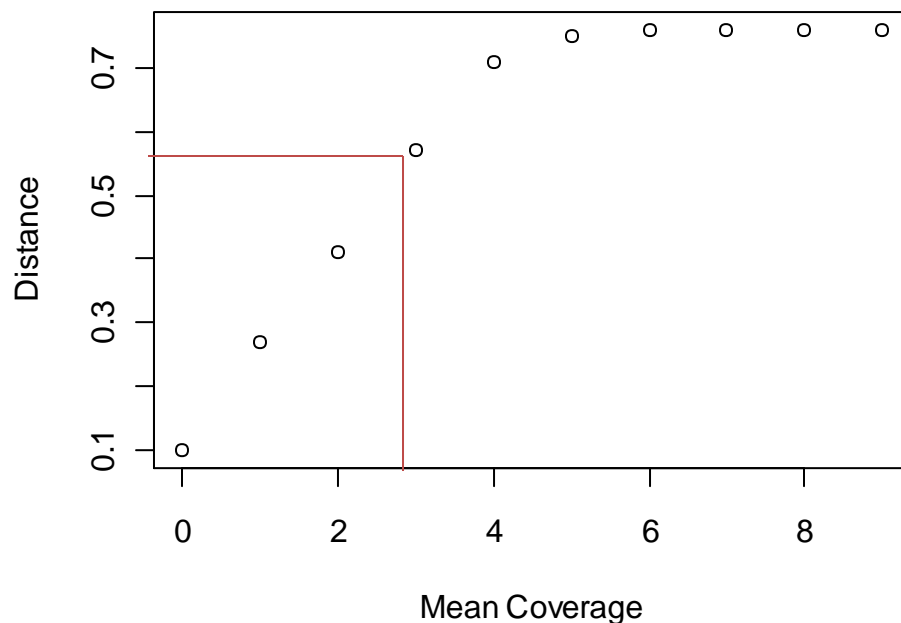
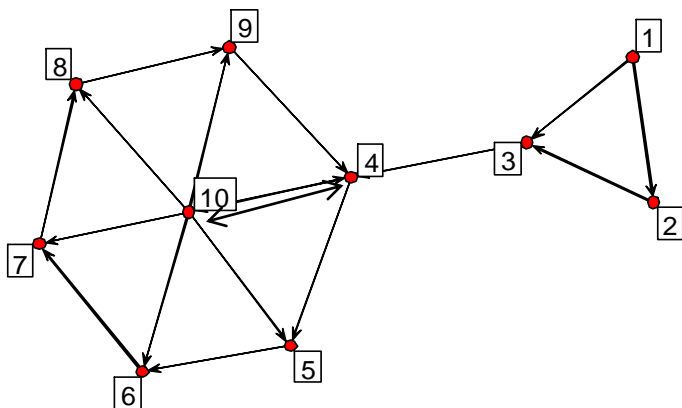
```
as.dist(equiv.dist)
hclust(*, "complete")
```

Graph structure by 'geodist'

- `structure.statistics`

```
> ss<-structure.statistics(g)
```

```
> plot(0:9,ss,xlab="Mean Coverage",ylab="Distance")
```



Graph cov based function

Regression, principal component,
canonical correlation

Multi graph measurements

- Graph mean

- In dichotomous case. graph mean corresponds to graph's density

$$\bar{\delta}_H = \frac{1}{|V_U|^2} \sum_{x=1}^{|V_U|} \sum_{y=1}^{|V_U|} \delta_H(x, y)$$

- Graph covariance

- gcov/gscov

$$Cov(H_i, H_j) = \frac{1}{|V_U|^2} \sum_{x=1}^{|V_U|} \sum_{y=1}^{|V_U|} ((\delta_i(x, y) - \bar{\delta}_{H_i}) (\delta_j(x, y) - \bar{\delta}_{H_j}))$$

- Graph correlation

- gcor/gscor

$$\rho(H_i, H_j) = \frac{Cov(H_i, H_j)}{\sqrt{Var(H_i) Var(H_j)}}$$

- Structural covariance

- unlabeled graph

$$Cov_S(G_i, G_j | \mathcal{P}_i, \mathcal{P}_j) = \max_{L_a \in \mathcal{P}_i, L_b \in \mathcal{P}_j} Cov(L_a(G_i), L_b(G_j))$$

Correlation statistic model

- Canonical correlation
 - netcancor
- Linear regression
 - netlm
- Logistic regression
 - netlogit
- Linear autocorrelation model
 - lnam
 - nacf

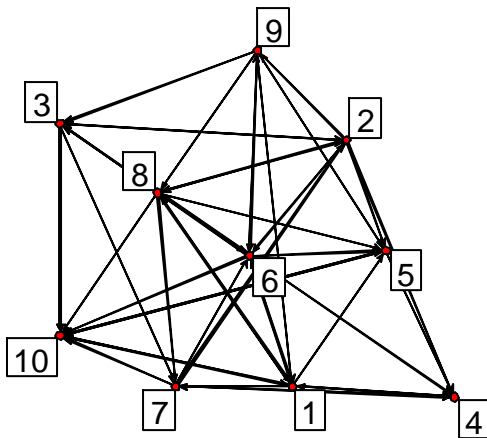
Random graph models

Graph evolution

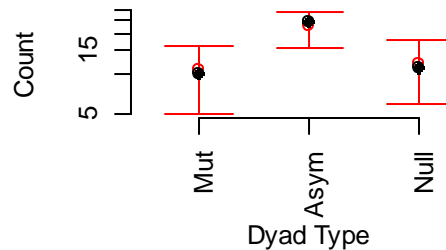
- Random
- Biased
- 4 Phases

Biased net model

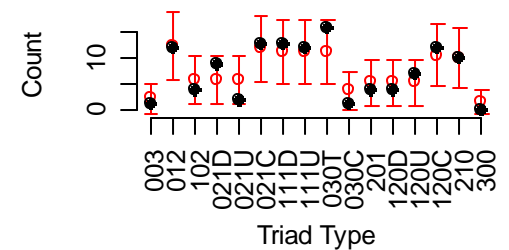
- graph generate: rgnb
- graph prediction: bn



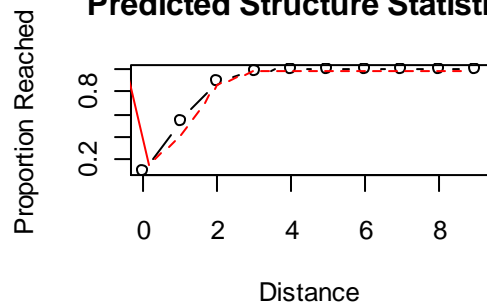
Predicted Dyad Census



Predicted Triad Census



Predicted Structure Statistics



Graph statistic test

- `cugtest`
- `qaptest`

Thanks

