

---

# **igraph** – a package for network analysis

---

Gábor Csárdi  
[Gabor.Csardi@unil.ch](mailto:Gabor.Csardi@unil.ch)

Department of Medical Genetics,  
University of Lausanne, Lausanne, Switzerland

# Outline

---

1. Why another graph package?
2. igraph architecture, data model and data representation
3. Manipulating graphs
4. Features and their time complexity

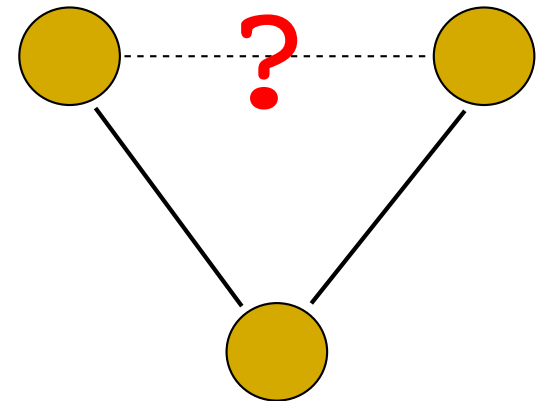
# Why another graph package?

---

- graph is slow. RBGL is slow, too.

---

```
1 > ba2 # graph & RBGL
2 A graphNEL graph with undirected edges
3 Number of Nodes = 100000
4 Number of Edges = 199801
```



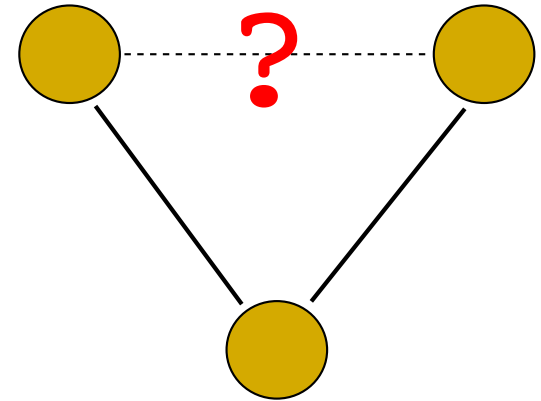
# Why another graph package?

---

- graph is slow. RBGL is slow, too.

---

```
1 > ba2 # graph & RBGL
2 A graphNEL graph with undirected edges
3 Number of Nodes = 100000
4 Number of Edges = 199801
5 > system.time(RBGL::transitivity(ba2))
6 user system elapsed
7 7.517 0.000 7.567
```

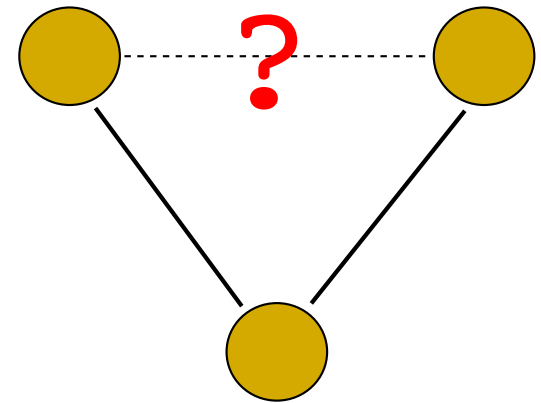


# Why another graph package?

---

- graph is slow. RBGL is slow, too.

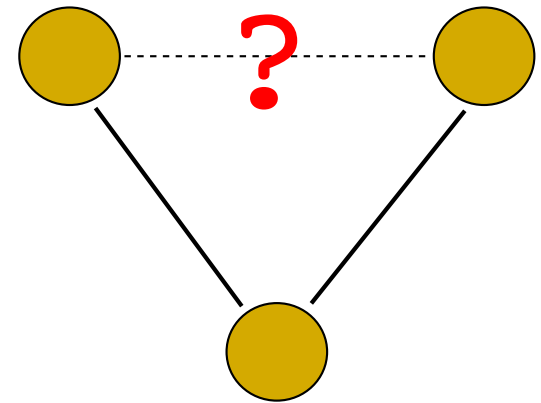
```
1 > ba2 # graph & RBGL
2 A graphNEL graph with undirected edges
3 Number of Nodes = 100000
4 Number of Edges = 199801
5 > system.time(RBGL::transitivity(ba2))
6 user system elapsed
7 7.517 0.000 7.567
8 > summary(ba) # igraph
9 Vertices: 1e+05
10 Edges: 199801
11 Directed: FALSE
12 No graph attributes.
13 No vertex attributes.
14 No edge attributes.
```



# Why another graph package?

- graph is slow. RBGL is slow, too.

```
1 > ba2 # graph & RBGL
2 A graphNEL graph with undirected edges
3 Number of Nodes = 100000
4 Number of Edges = 199801
5 > system.time(RBGL::transitivity(ba2))
6 user system elapsed
7 7.517 0.000 7.567
8 > summary(ba) # igraph
9 Vertices: 1e+05
10 Edges: 199801
11 Directed: FALSE
12 No graph attributes.
13 No vertex attributes.
14 No edge attributes.
15 > system.time(igraph::transitivity(ba))
16 user system elapsed
17 0.328 0.000 0.335
```



# Why another graph package?

---

- sna is slow. network is slow, too.

---

```
1 > net2                                     # SNA & network
2   Network attributes:
3     vertices = 1e+05
4     directed = TRUE
5     hyper = FALSE
6     loops = FALSE
7     multiple = FALSE
8     bipartite = FALSE
9     total edges= 199801
10    missing edges= 0
11    non-missing edges= 199801
12    ...
```

# Why another graph package?

---

- sna is slow. network is slow, too.

---

```
1 > net2                                # SNA & network
2   Network attributes:
3     vertices = 1e+05
4     directed = TRUE
5     hyper = FALSE
6     loops = FALSE
7     multiple = FALSE
8     bipartite = FALSE
9     total edges= 199801
10    missing edges= 0
11    non-missing edges= 199801
12    ...
13 > gtrans(net2)
14 Error in matrix(0, nr = network.size(x), nc = network.size(x)) :
15   too many elements specified
```

---



# Why another graph package?

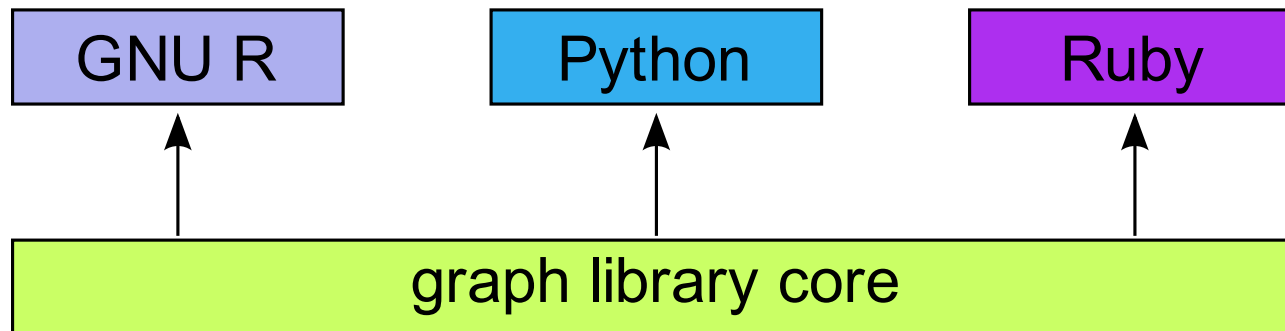
---

- graph is slow. RBGL is slow, too.
- sna is slow. network is slow, too.
- A generic solution was needed, i.e. a common C layer, that can be interfaced from C/C++, R, Python, etc.

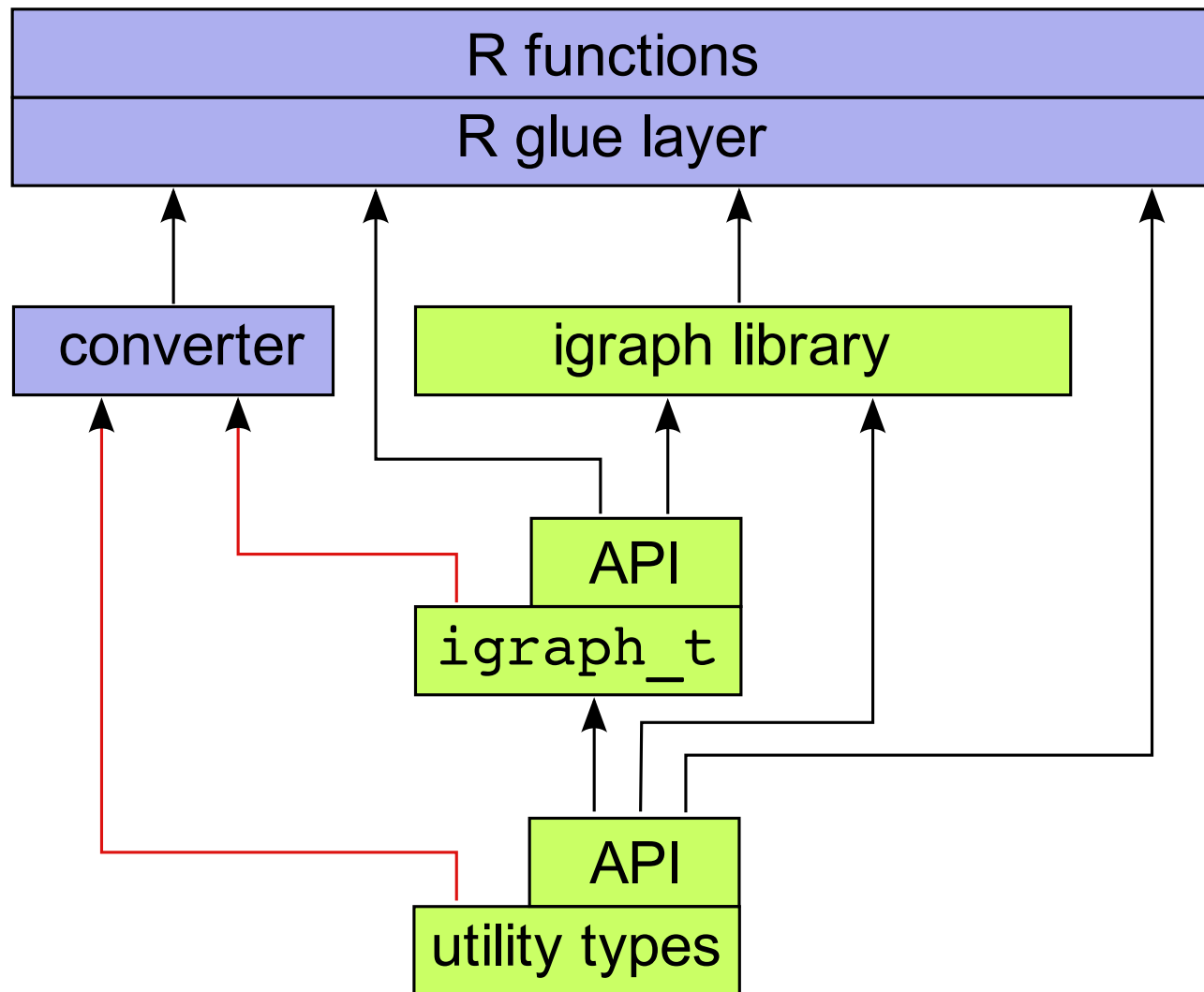
# Why another graph package?

---

- graph is slow. RBGL is slow, too.
- sna is slow. network is slow, too.
- A generic solution was needed, i.e. a common C layer, that can be interfaced from C/C++, R, Python, etc.



# The igraph architecture



# Dependencies

---

- Standard C/C++ libraries.



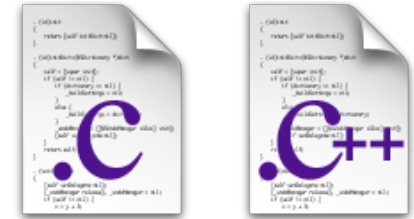
# Dependencies

- Standard C/C++ libraries.
- stats package, this is part of base.



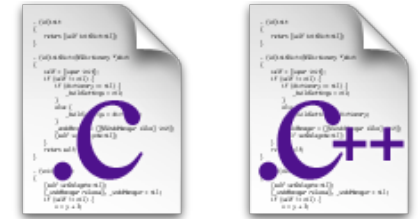
# Dependencies

- Standard C/C++ libraries.
- stats package, this is part of base.
- Optional: libxml2 library, for reading GraphML files (included in Windows builds).



# Dependencies

- Standard C/C++ libraries.
- stats package, this is part of base.
- Optional: libxml2 library, for reading GraphML files (included in Windows builds).
- Optional: GMP library, graph automorphisms (not included in Windows builds).



# Dependencies

- Standard C/C++ libraries.
- stats package, this is part of base.
- Optional: libxml2 library, for reading GraphML files (included in Windows builds).
- Optional: GMP library, graph automorphisms (not included in Windows builds).
- Suggested packages: stats4, rgl, tcltk, RSQLite, digest, graph, Matrix.

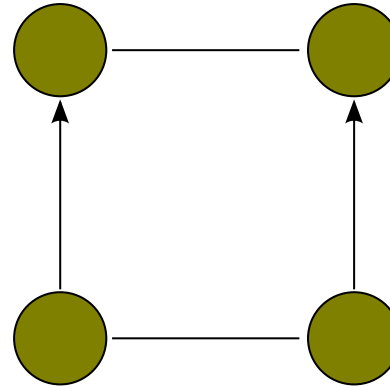




# The igraph data model, what cannot be represented

---

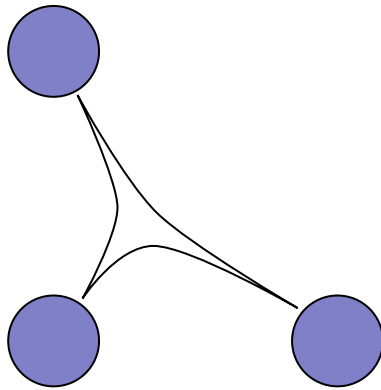
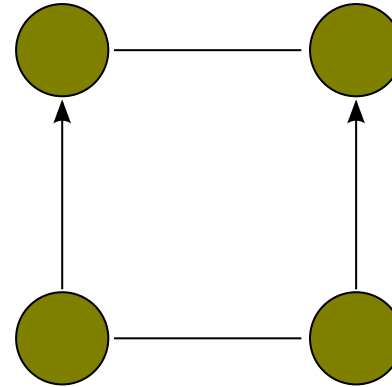
“Mixed” graphs, with undirected and directed edges.  
You can “emulate” them via graph attributes.



# The igraph data model, what cannot be represented

---

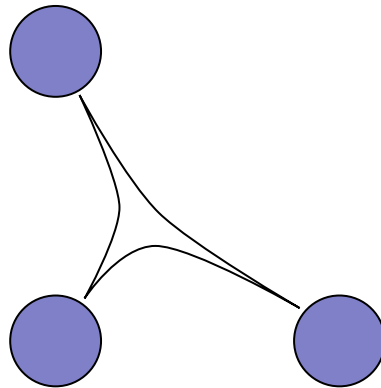
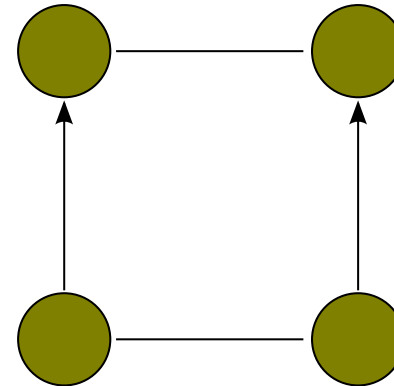
“Mixed” graphs, with undirected and directed edges.  
You can “emulate” them via graph attributes.



Hypergraphs. Perhaps see the `hypergraph` package.

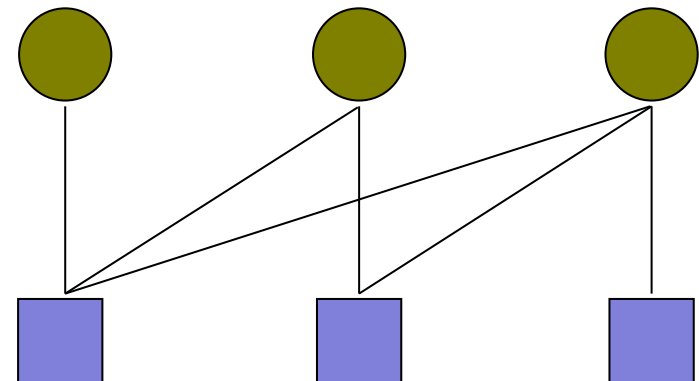
# The igraph data model, what cannot be represented

“Mixed” graphs, with undirected and directed edges.  
You can “emulate” them via graph attributes.



Hypergraphs. Perhaps see the hypergraph package.

No direct support for bipartite (two-mode) graphs.  
It is possible to handle them via graph attributes.



# Graph representation, sparse graphs

---

Flat data structures, indexed edge lists. Easy to handle, good for many kind of questions.

Alice	Bob
Bob	Diana
Cecil	Diana
Alice	Esmeralda
Diana	Esmeralda
Cecil	Fabien
Esmeralda	Fabien
Bob	Gigi
Cecil	Gigi
Diana	Gigi
Alice	Helen
Bob	Helen
Diana	Helen
Esmeralda	Helen
Fabien	Helen
Gigi	Helen
Diana	Iannis
Esmeralda	Iannis
Alice	Jennifer
Helen	Jennifer

# Graph representation, sparse graphs

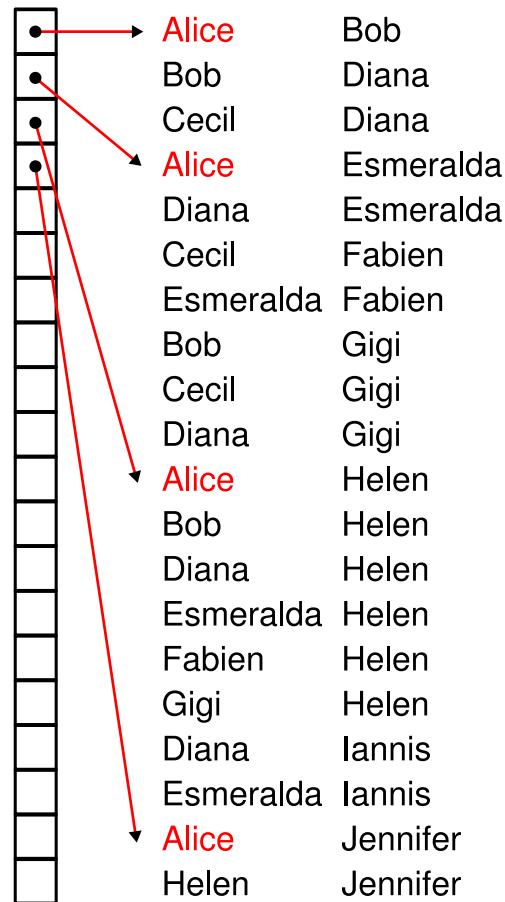
---

Flat data structures, indexed edge lists. Easy to handle, good for many kind of questions.

Alice	Bob
Bob	Diana
Cecil	Diana
Alice	Esmeralda
Diana	Esmeralda
Cecil	Fabien
Esmeralda	Fabien
Bob	Gigi
Cecil	Gigi
Diana	Gigi
Alice	Helen
Bob	Helen
Diana	Helen
Esmeralda	Helen
Fabien	Helen
Gigi	Helen
Diana	Iannis
Esmeralda	Iannis
Alice	Jennifer
Helen	Jennifer

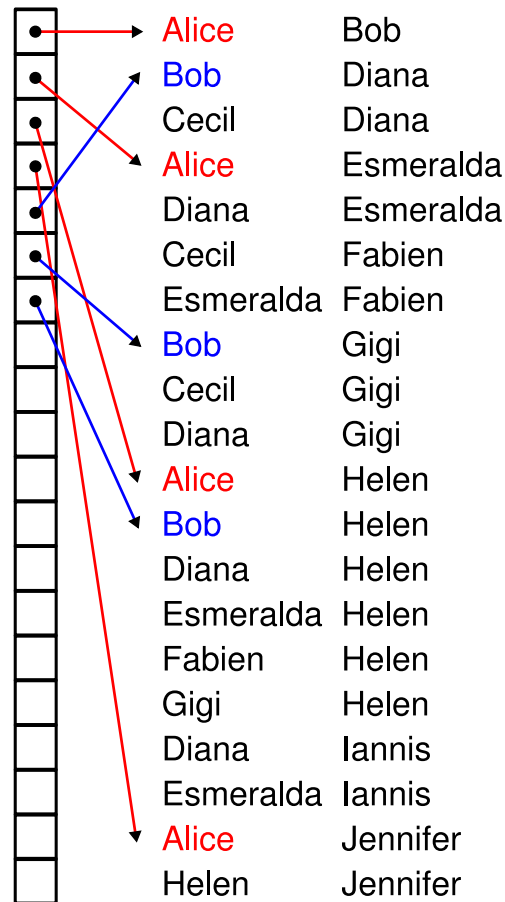
# Graph representation, sparse graphs

Flat data structures, indexed edge lists. Easy to handle, good for many kind of questions.



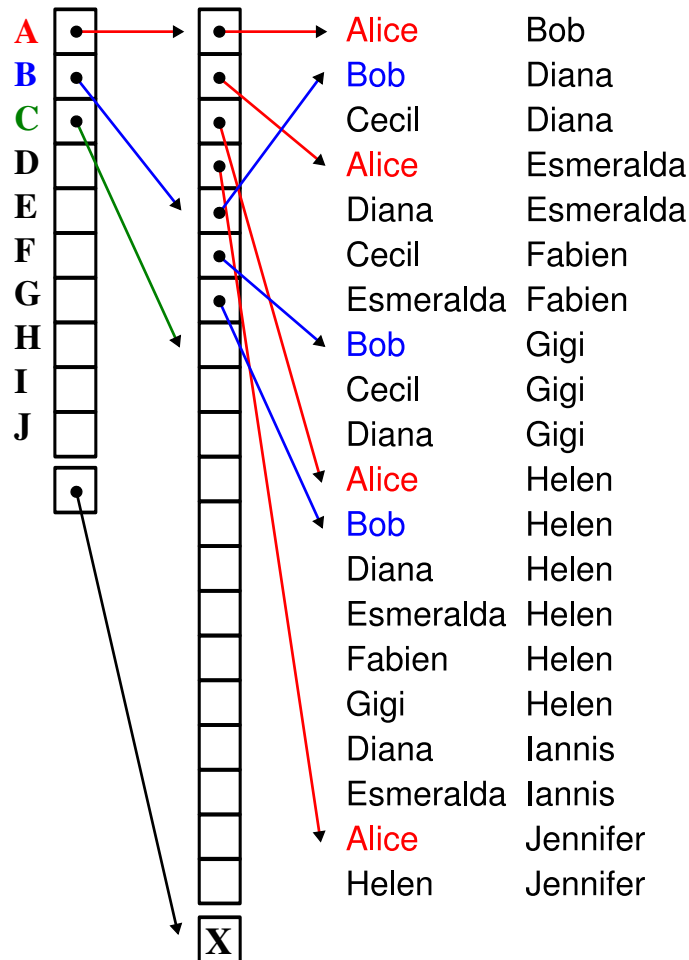
# Graph representation, sparse graphs

Flat data structures, indexed edge lists. Easy to handle, good for many kind of questions.



# Graph representation, sparse graphs

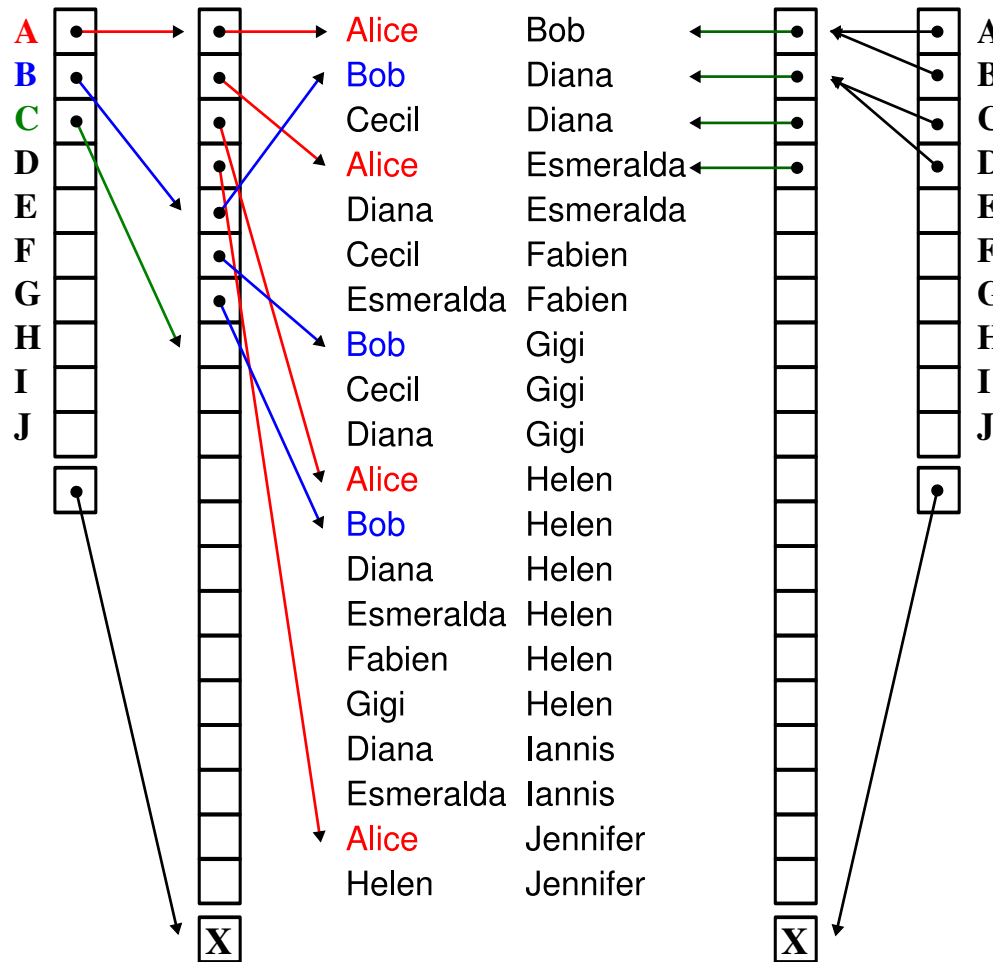
Flat data structures, indexed edge lists. Easy to handle, good for many kind of questions.





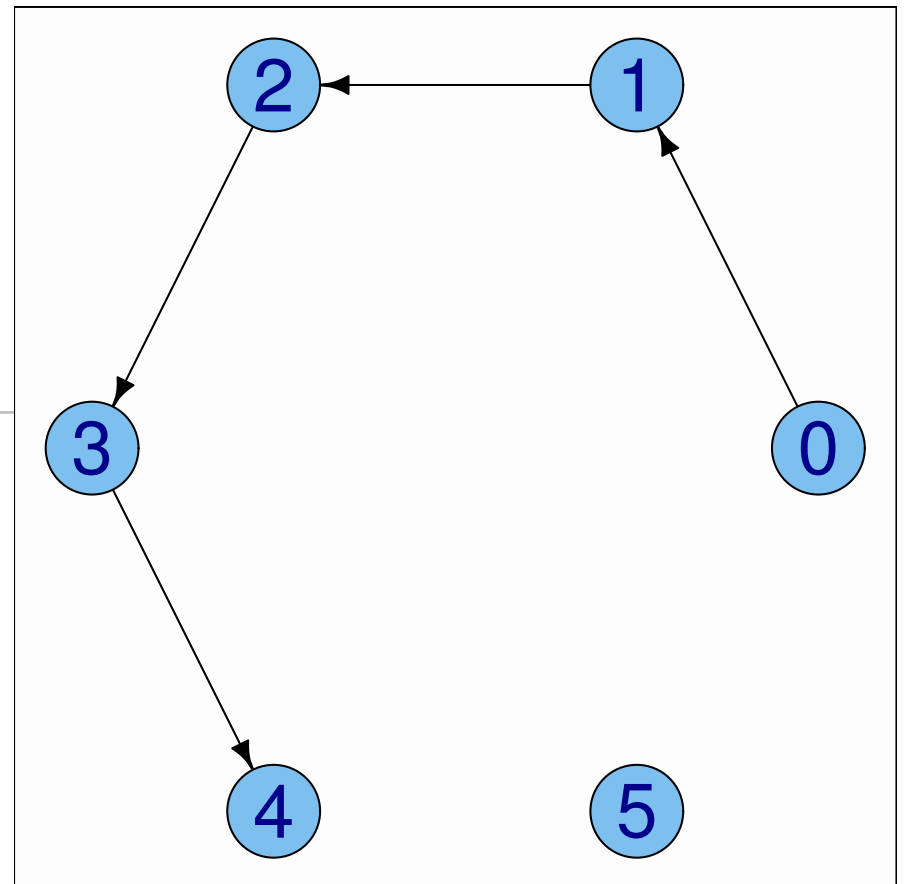
# Graph representation, sparse graphs

Flat data structures, indexed edge lists. Easy to handle, good for many kind of questions.



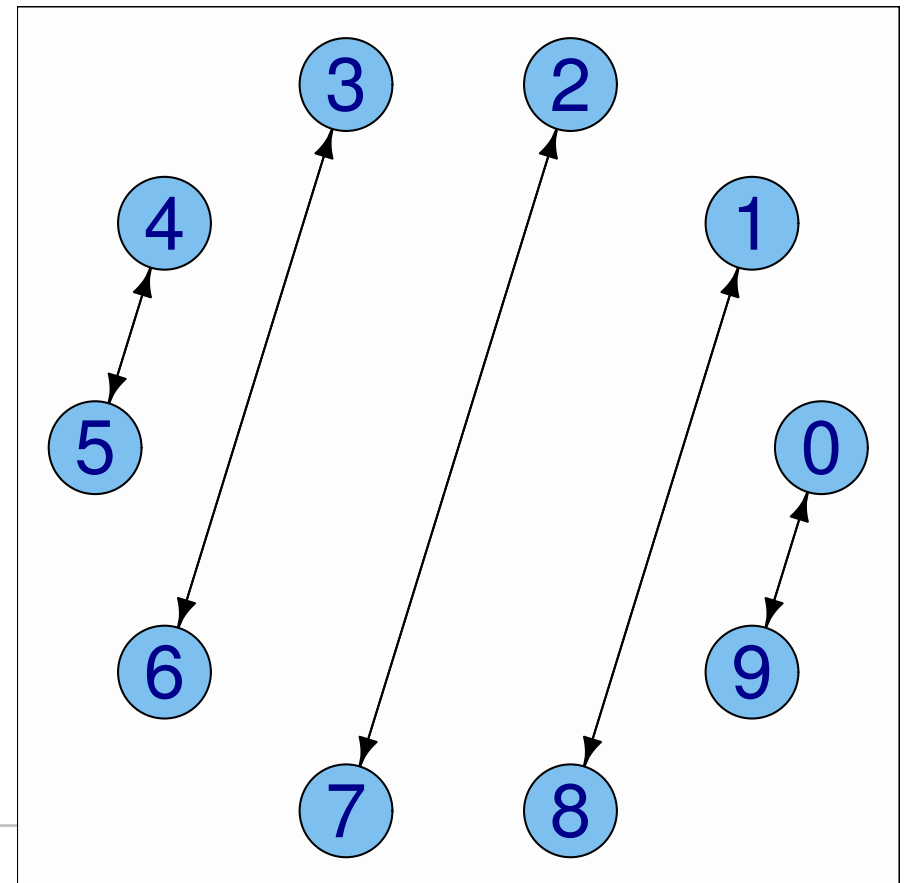
# Creating graphs, via vertex ids

```
1 > g <- graph( c(0,1, 1,2, 2,3, 3,4), n=6, directed=TRUE )
2 > g
3 Vertices: 6
4 Edges: 4
5 Directed: TRUE
6 Edges:
7
8 [0] 0 -> 1
9 [1] 1 -> 2
10 [2] 2 -> 3
11 [3] 3 -> 4
```



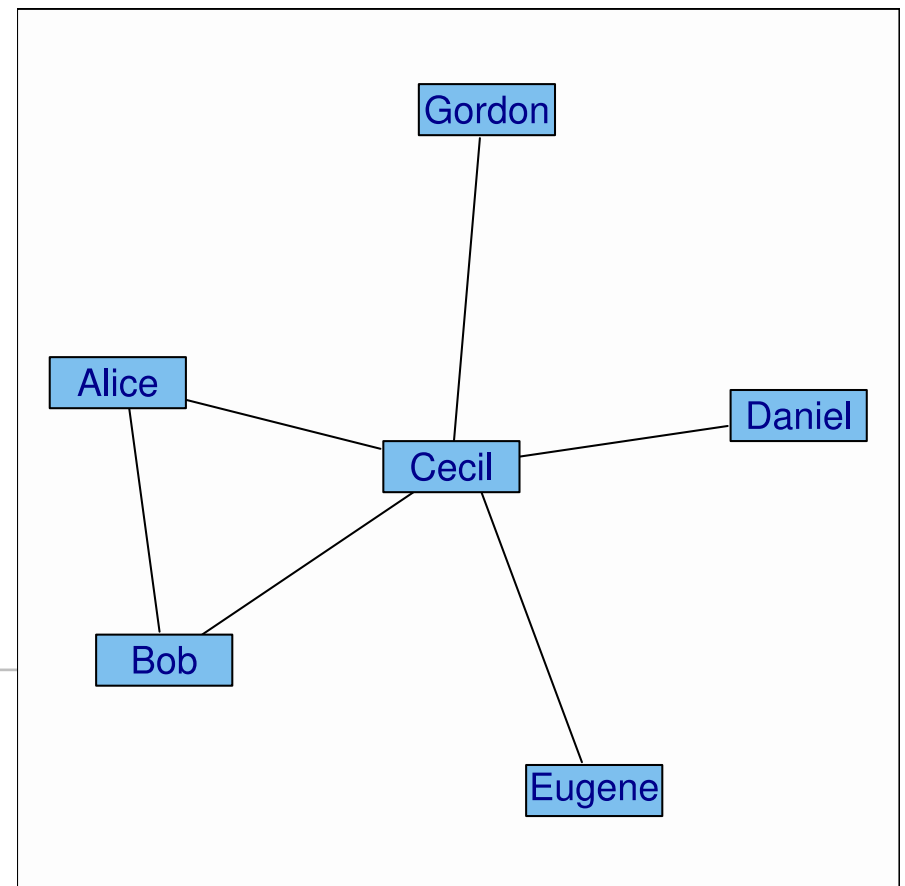
# Creating graphs, via vertex ids

```
1 > el <- cbind(0:9, 9:0)
2 > g <- graph( t(el), directed=TRUE)
3 > g
4 Vertices: 10
5 Edges: 10
6 Directed: TRUE
7 Edges:
8
9 [0] 0 -> 9
10 [1] 1 -> 8
11 [2] 2 -> 7
12 [3] 3 -> 6
13 [4] 4 -> 5
14 [5] 5 -> 4
15 [6] 6 -> 3
16 [7] 7 -> 2
17 [8] 8 -> 1
18 [9] 9 -> 0
```



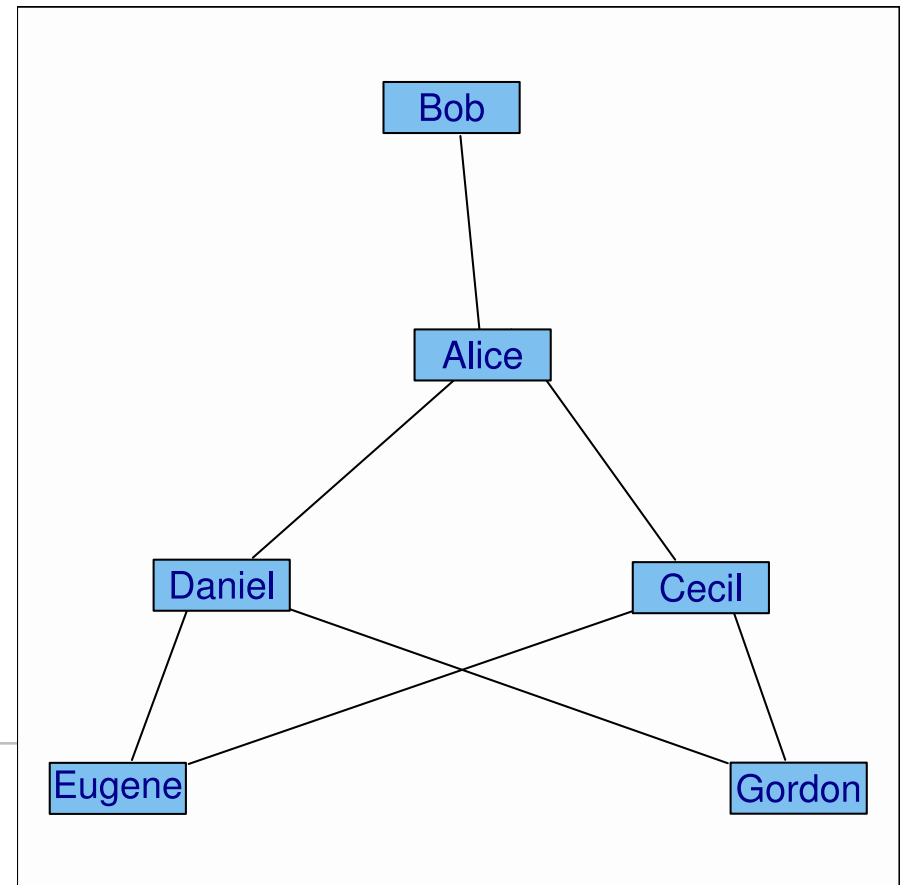
# Creating graphs, graph.formula

```
1 # A simple undirected graph
2 > g <- graph.formula( Alice-Bob-Cecil-Alice,
3                       Daniel-Cecil-Eugene, Cecil-Gordon )
4 > g
5 Vertices: 6
6 Edges: 6
7 Directed: FALSE
8 Edges:
9
10 [0] Alice -- Bob
11 [1] Bob   -- Cecil
12 [2] Alice -- Cecil
13 [3] Cecil -- Daniel
14 [4] Cecil -- Eugene
15 [5] Cecil -- Gordon
```



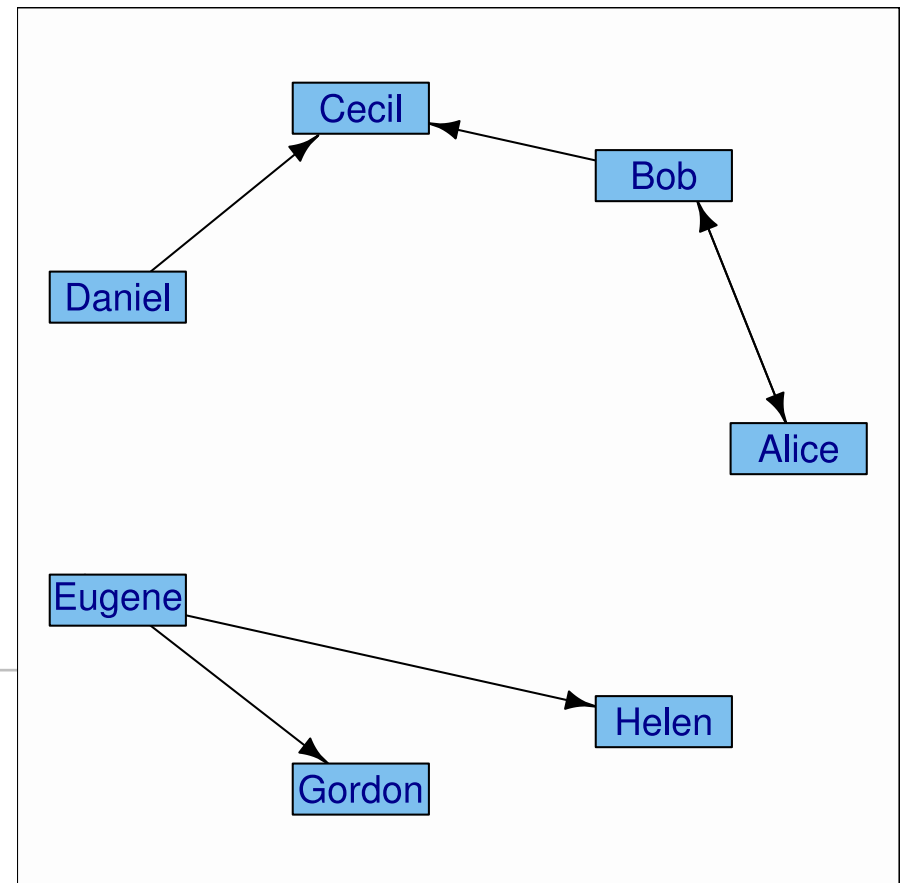
# Creating graphs, graph.formula

```
1 # Another undirected graph, ":" notation
2 > g2 <- graph.formula( Alice-Bob:Cecil:Daniel,
3                       Cecil:Daniel-Eugene:Gordon )
4 > g2
5 Vertices: 6
6 Edges: 7
7 Directed: FALSE
8 Edges:
9
10 [0] Alice -- Bob
11 [1] Alice -- Cecil
12 [2] Alice -- Daniel
13 [3] Cecil -- Eugene
14 [4] Cecil -- Gordon
15 [5] Daniel -- Eugene
16 [6] Daniel -- Gordon
```



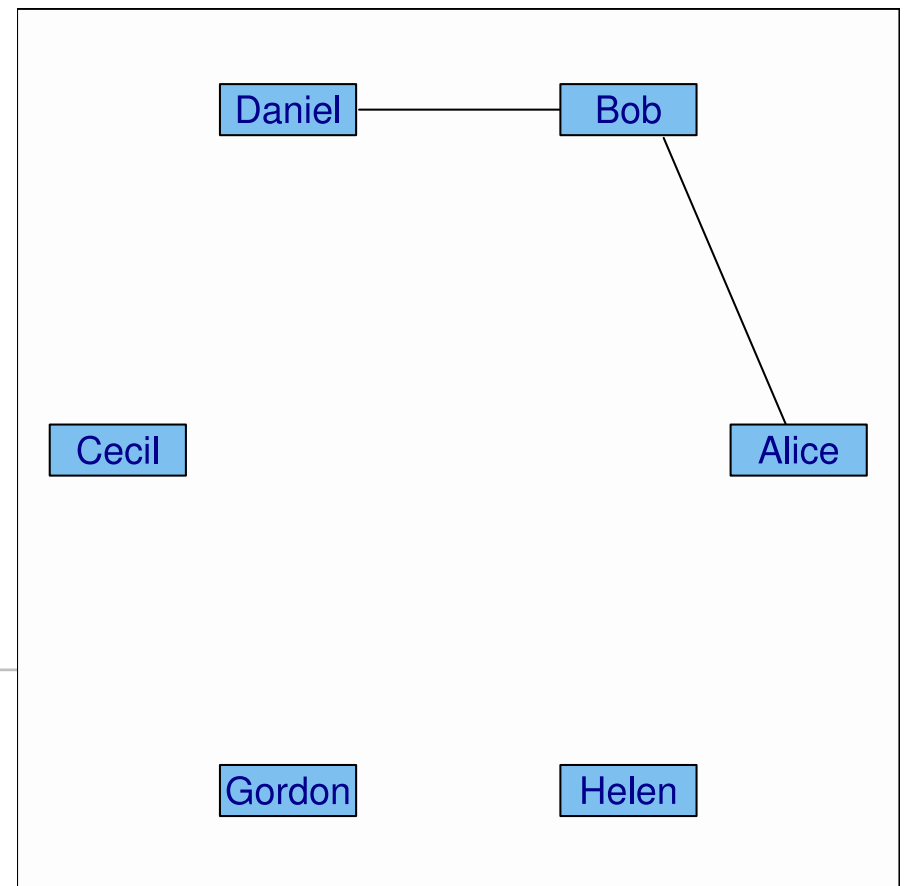
# Creating graphs, graph.formula

```
1 # A directed graph
2 > g3 <- graph.formula( Alice +--+ Bob ---+ Cecil
3     +--+ Daniel, Eugene ---+ Gordon:Helen )
4 > g3
5 Vertices: 7
6 Edges: 6
7 Directed: TRUE
8 Edges:
9
10 [0] Bob    -> Alice
11 [1] Alice  -> Bob
12 [2] Bob    -> Cecil
13 [3] Daniel -> Cecil
14 [4] Eugene -> Gordon
15 [5] Eugene -> Helen
```



# Creating graphs, graph.formula

```
1 # A graph with isolate vertices
2 > g4 <- graph.formula( Alice -- Bob -- Daniel,
3                       Cecil:Gordon, Helen )
4 > g4
5 Vertices: 6
6 Edges: 2
7 Directed: FALSE
8 Edges:
9
10 [0] Alice  -- Bob
11 [1] Bob    -- Daniel
12 > V(g4)
13 Vertex sequence:
14 [1] "Alice" "Bob" "Daniel"
15 [4] "Cecil" "Gordon" "Helen"
```

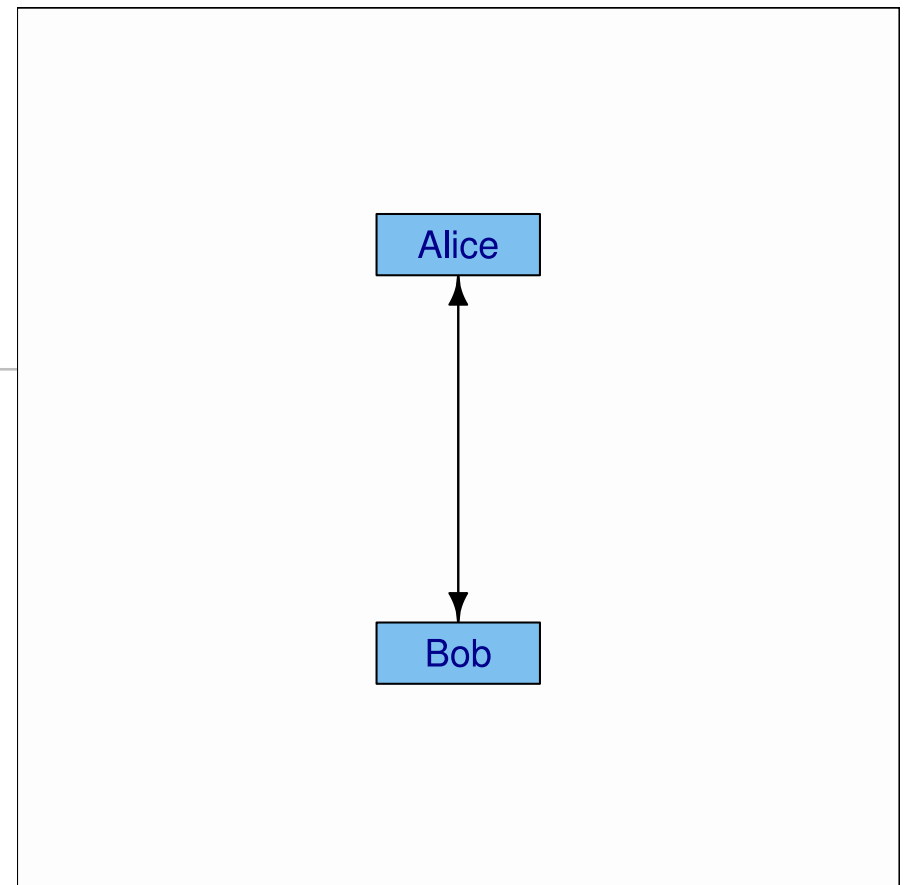


# Creating graphs, graph.formula

---

```
1 # "Arrows" can be arbitrarily long
2 > g5 <- graph.formula( Alice +-----+ Bob )
3 > g5
4 Vertices: 2
5 Edges: 2
6 Directed: TRUE
7 Edges:
8
9 [0] Bob -> Alice
10 [1] Alice -> Bob
```

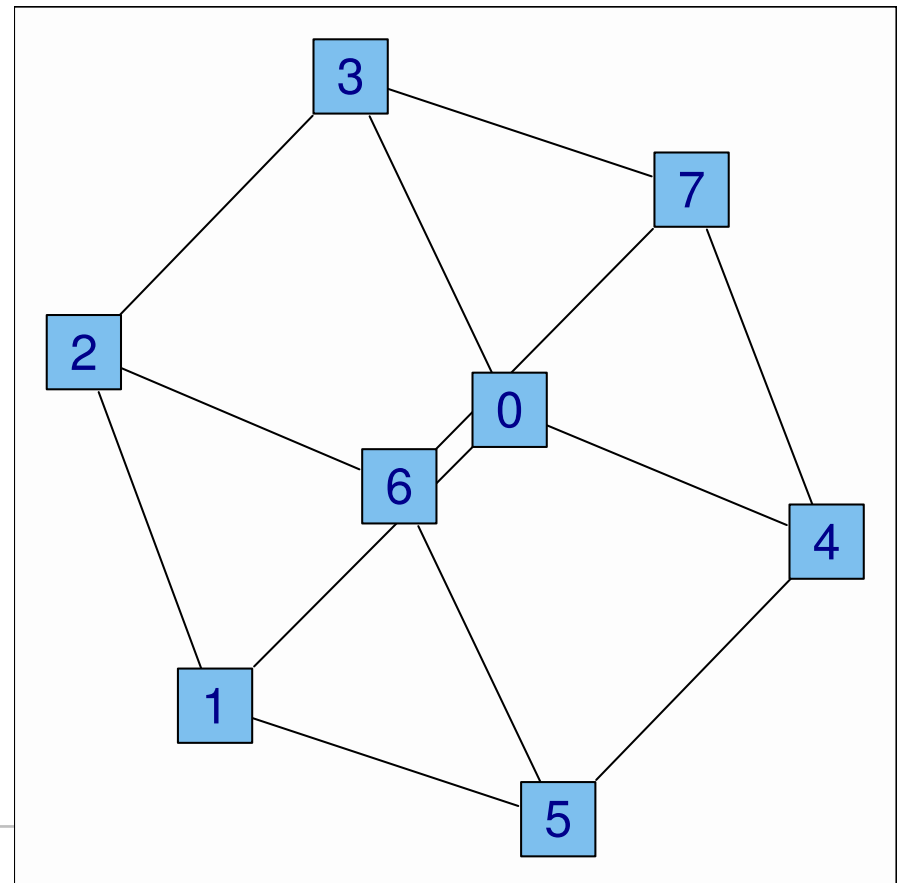
---





# Creating graphs, graph.famous

```
1 > graph.famous("Cubical")
2 Vertices: 8
3 Edges: 12
4 Directed: FALSE
5 Edges:
6
7 [0] 0 -- 1
8 [1] 1 -- 2
9 [2] 2 -- 3
10 [3] 0 -- 3
11 [4] 4 -- 5
12 [5] 5 -- 6
13 [6] 6 -- 7
14 [7] 4 -- 7
15 [8] 0 -- 4
16 [9] 1 -- 5
17 [10] 2 -- 6
18 [11] 3 -- 7
```



# Creating graphs, `graph.data.frame`

---

```
1 > traits <- read.csv("traits.csv", head=F)
2 > traits
3           V1 V2 V3
4 1 Alice Anderson 48 F
5 2 Bob Bradford 33 M
6 3 Cecil Connor 45 F
7 4 David Daugher 34 M
8 5 Esmeralda Escobar 21 F
9 6 Frank Finley 36 M
10 7 Gabi Garbo 44 F
11 8 Helen Hunt 40 F
12 9 Iris Irving 25 F
13 10 James Jones 47 M
14 > colnames(traits) <- c("name", "age", "gender")
15 > traits[,1] <- sapply(strsplit(as.character(traits[,1]), " "), "[", 1)
```

---

# Creating graphs, graph.data.frame

---

```
1 > relations <- read.csv("relations.csv", head=F)
2 > relations
3           V1           V2 V3 V4 V5
4  1      Bob      Alice  N  4  4
5  2    Cecil      Bob   N  5  5
6  3    Cecil      Alice  Y  5  5
7  4    David      Alice  N  3  4
8  5    David      Bob   N  4  2
9  6 Esmeralda      Alice  Y  4  3
10 7     Frank      Alice  N  3  2
11 8     Frank Esmeralda  N  4  4
12 9     Gabi      Bob   Y  5  5
13 10    Gabi      Alice  N  3  0
14 11    Helen      Alice  N  4  1
15 12    Iris      Cecil  N  0  1
16 ...
17 > colnames(relations) <- c("from", "to", "same.room",
18     "friendship", "advice")
```

---

# Creating graphs, `graph.data.frame`

---

```
1 > orgnet <- graph.data.frame(relations, vertices=traits)
2 > summary(orgnet)
3 Vertices: 10
4 Edges: 34
5 Directed: TRUE
6 No graph attributes.
7 Vertex attributes: name, age, gender.
8 Edge attributes: same.room, friendship, advice.
```

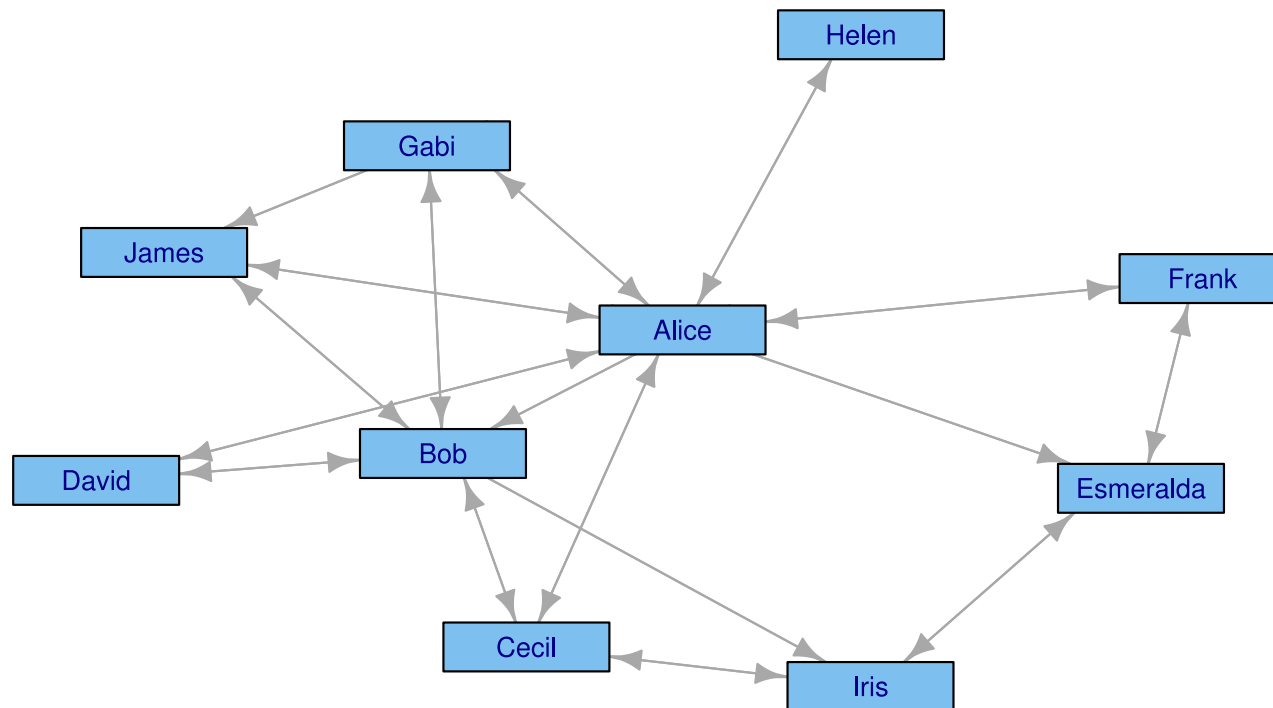
---

# Creating graphs, graph.data.frame

---

```
1 > plot(orgnet, layout=layout.kamada.kawai, vertex.label=V(orgnet)$name,  
2     vertex.shape="rectangle", vertex.size=20, asp=FALSE)
```

---

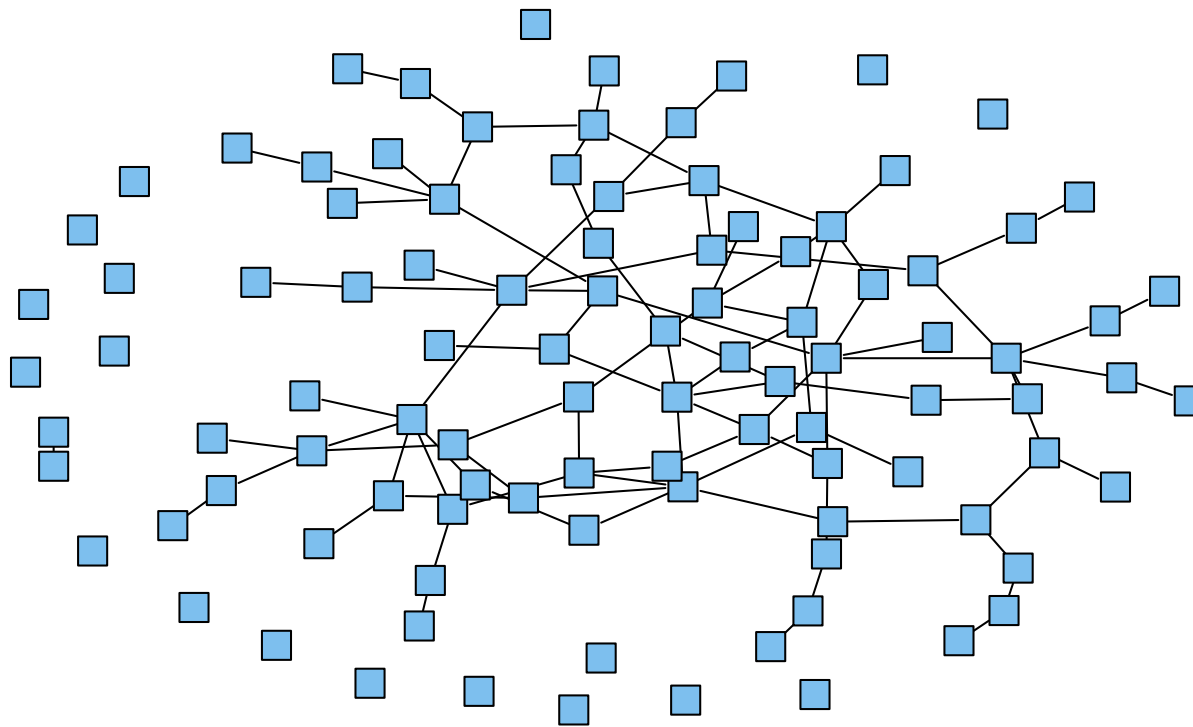


# Creating graphs, random graphs

---

```
1 > er <- erdos.renyi.game(100, 100, type="gnm")  
2 > plot(er, vertex.size=5, vertex.label=NA, asp=FALSE, vertex.shape="square",  
3     layout=layout.fruchterman.reingold, edge.color="black")
```

---

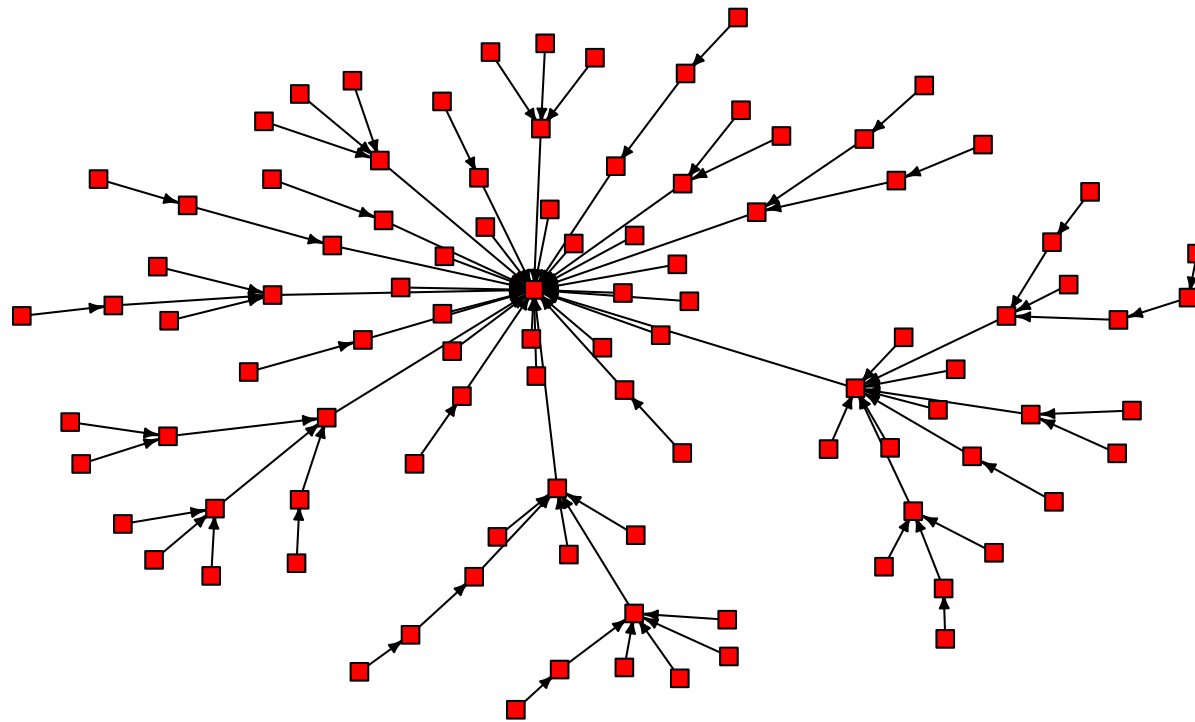


# Creating graphs, random graphs

---

```
1 > ba <- ba.game(100, power=1, m=1)
2 > plot(ba, vertex.size=3, vertex.label=NA, asp=FALSE, vertex.shape="square",
3       layout=layout.fruchterman.reingold, edge.color="black",
4       edge.arrow.size=0.5)
```

---



## Meta data: graph/vertex/edge attributes

---

- Assigning attributes: `set/get.graph/vertex/edge.attribute`.



## Meta data: graph/vertex/edge attributes

---

- Assigning attributes: `set/get.graph/vertex/edge.attribute`.
- $V(g)$  and  $E(g)$ .

## Meta data: graph/vertex/edge attributes

---

- Assigning attributes: `set/get.graph/vertex/edge.attribute`.
- `V(g)` and `E(g)`.
- Easy access of attributes:

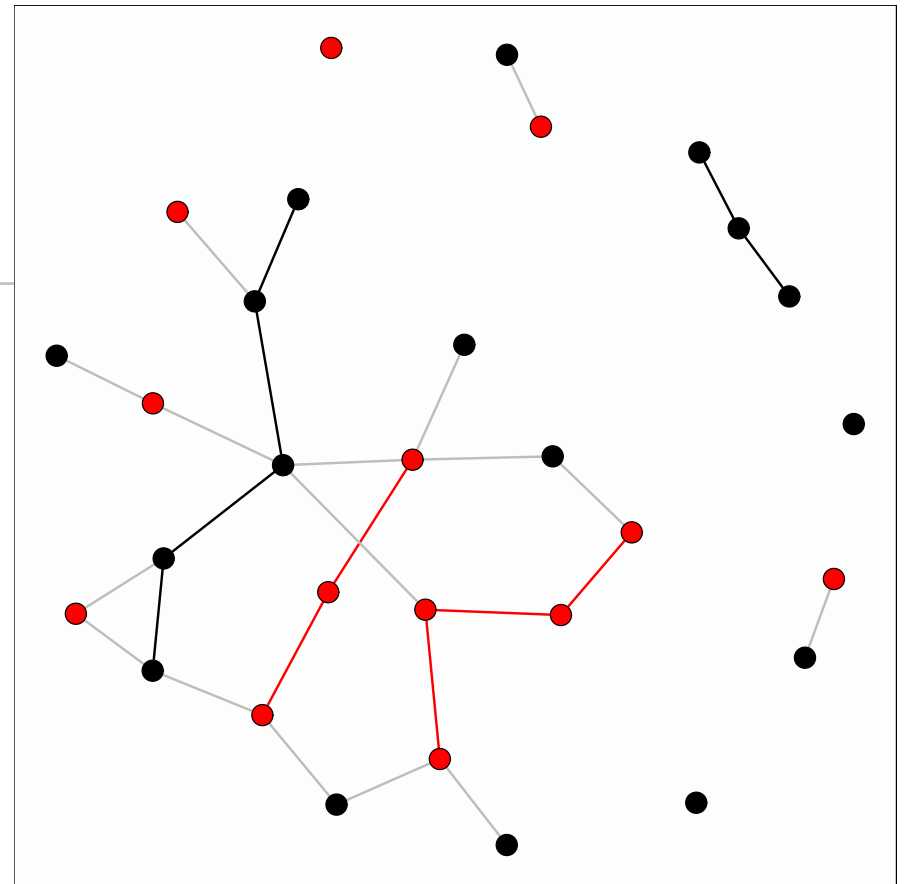
---

```
1 > g <- erdos.renyi.game(30, 2/30)
2 > V(g)$color <- sample( c("red", "black"),
3                         vcount(g), rep=TRUE)
4 > V(g)$color
5 [1] "red" "black" "red" "black" "black" "black" "red" "red" "red"
6 [10] "black" "black" "black" "red" "red" "black" "red" "black" "black"
7 [19] "red" "red" "black" "black" "red" "black" "black" "red" "black"
8 [28] "black" "black" "red"
9 > E(g)$color <- "grey"
```

---

# Vertex/edge selection with attributes

```
1 > red <- V(g)[ color == "red" ]  
2 > bl <- V(g)[ color == "black" ]  
3 > E(g)[ red %--% red ]$color <- "red"  
4 > E(g)[ bl %--% bl ]$color <- "black"  
5 > plot(g, vertex.size=5,  
6     layout=layout.fruchterman.reingold,  
7     vertex.label=NA)
```



# Visualizing graphs

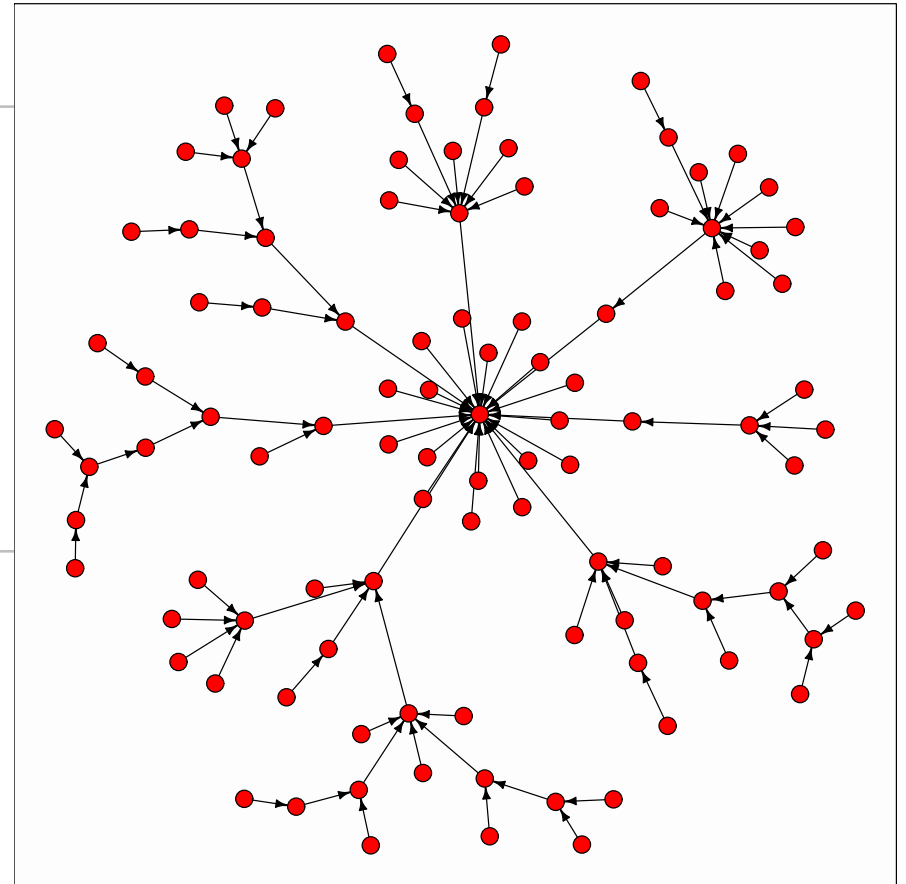
---

- Three functions with (almost) identical interfaces.

# Visualizing graphs

- Three functions with (almost) identical interfaces.
- `plot` Uses traditional R graphics, non-interactive, 2d. Publication quality plots in all formats R supports.

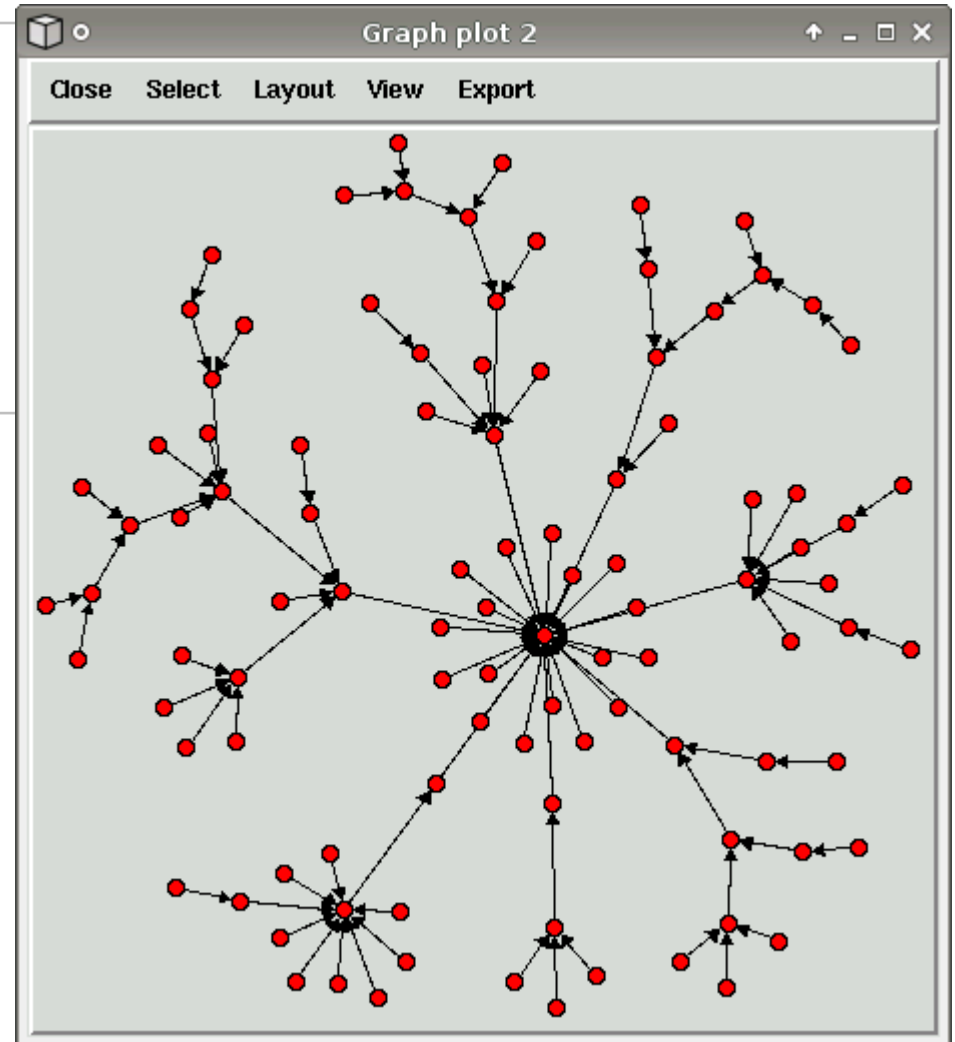
```
1 > g <- barabasi.game(100, m=1)
2 > igraph.par("plot.layout",
3           layout.fruchterman.reingold)
4 > plot(g, vertex.size=4, vertex.label=NA,
5       edge.arrow.size=0.7,
6       edge.color="black",
7       vertex.color="red", frame=TRUE)
```



# Visualizing graphs

tkplot Uses Tcl/Tk via the tcltk package, interactive, 2d.

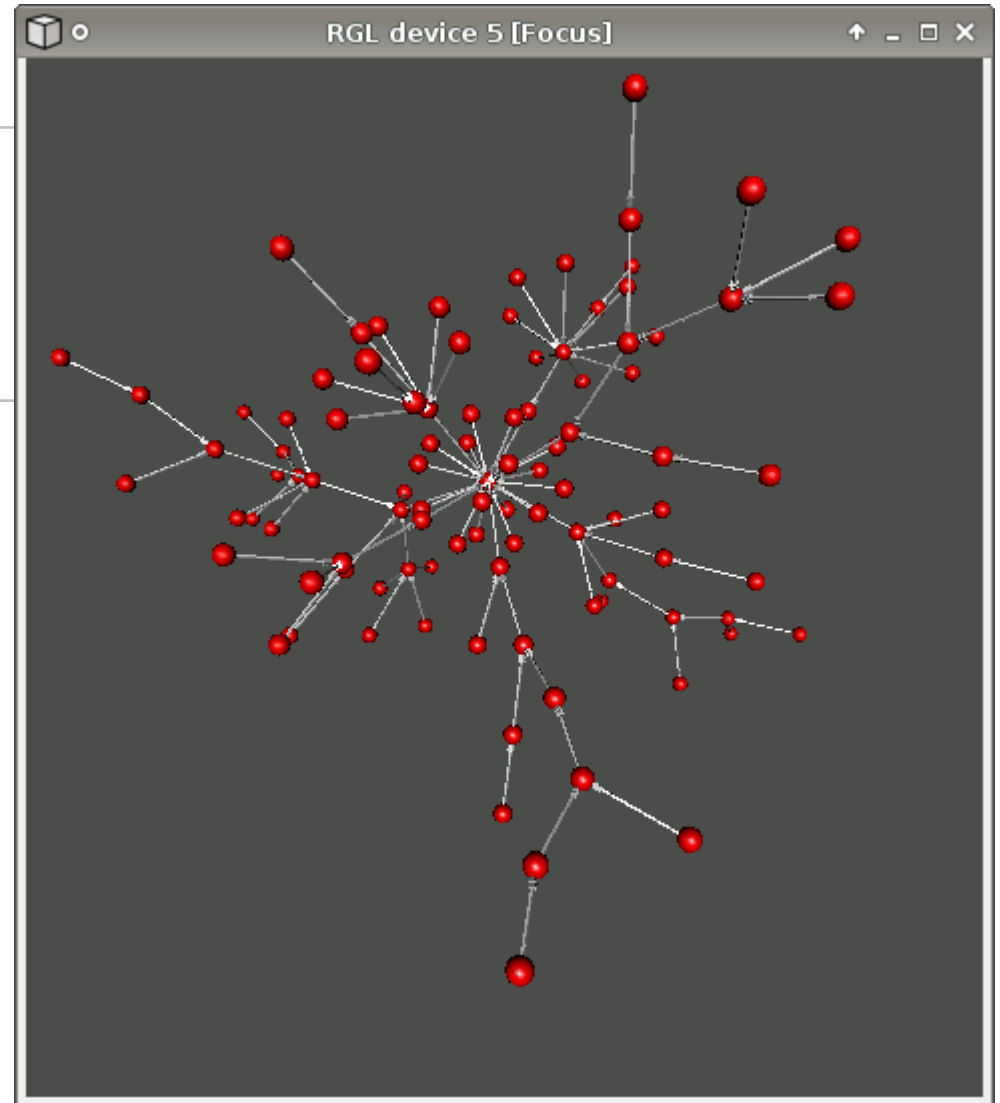
```
1 > id <- tkplot(g, vertex.size=4,  
2   vertex.label=NA,  
3   edge.color="black",  
4   edge.arrow.size=0.7,  
5   vertex.color="red")  
6 > coords <- tkplot.getcoords(id)
```



# Visualizing graphs

rglplot Needs the rgl package.

```
1 > co <- layout.kamada.kawai(g, dim=3)
2 > rglplot(g, vertex.size=5,
3       vertex.label=NA,
4       layout=co)
```



## Working with a somewhat bigger graph

---

```
1 > vertices <- read.csv("http://cneurocv.s.rmki.kfki.hu/igraph/judicial.csv")
2 > edges <- read.table("http://cneurocv.s.rmki.kfki.hu/igraph/allcites.txt")
3 > jg <- graph.data.frame(edges, vertices=vertices, dir=TRUE)
4 > summary(jg)
5 Vertices: 30288
6 Edges: 216738
7 Directed: TRUE
8 No graph attributes.
9 Vertex attributes: name, usid, parties, year, overruled, overruling,
10   oxford, liihc, indeg, outdeg, hub, hubrank, auth, authrank, between, incent.
11 No edge attributes.
```

---



# Working with a somewhat bigger graph

---

```
1 > is.connected(jg)                # Is it connected?
2 [1] FALSE
```

# Working with a somewhat bigger graph

---

```
1 > is.connected(jg)           # Is it connected?
2   [1] FALSE
3
4 > no.clusters(jg)           # How many components?
5   [1] 4881
```

# Working with a somewhat bigger graph

---

```
1 > is.connected(jg)           # Is it connected?
2 [1] FALSE
3
4 > no.clusters(jg)           # How many components?
5 [1] 4881
6
7 > table(clusters(jg)$csize)  # How big are these?
8
9      1      3      4 25389
10 4871      8      1      1
```

# Working with a somewhat bigger graph

---

```
1 > is.connected(jg)           # Is it connected?
2 [1] FALSE
3
4 > no.clusters(jg)           # How many components?
5 [1] 4881
6
7 > table(clusters(jg)$csize)  # How big are these?
8
9      1      3      4 25389
10 4871      8      1      1
11
12 > max(degree(jg, mode="in")) # Vertex degree
13 [1] 248
14 > max(degree(jg, mode="out"))
15 [1] 195
16 > max(degree(jg, mode="all"))
17 [1] 313
```

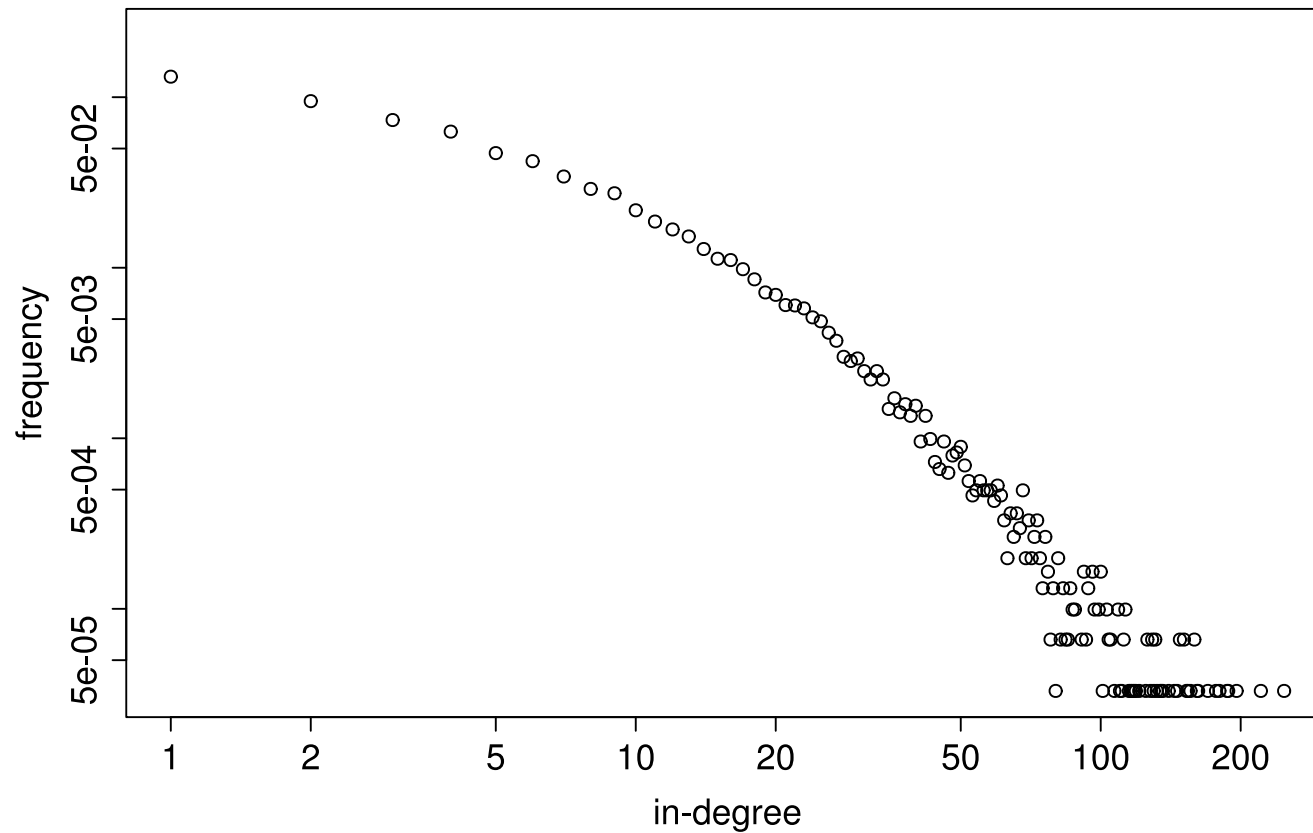
---

# Working with a somewhat bigger graph

---

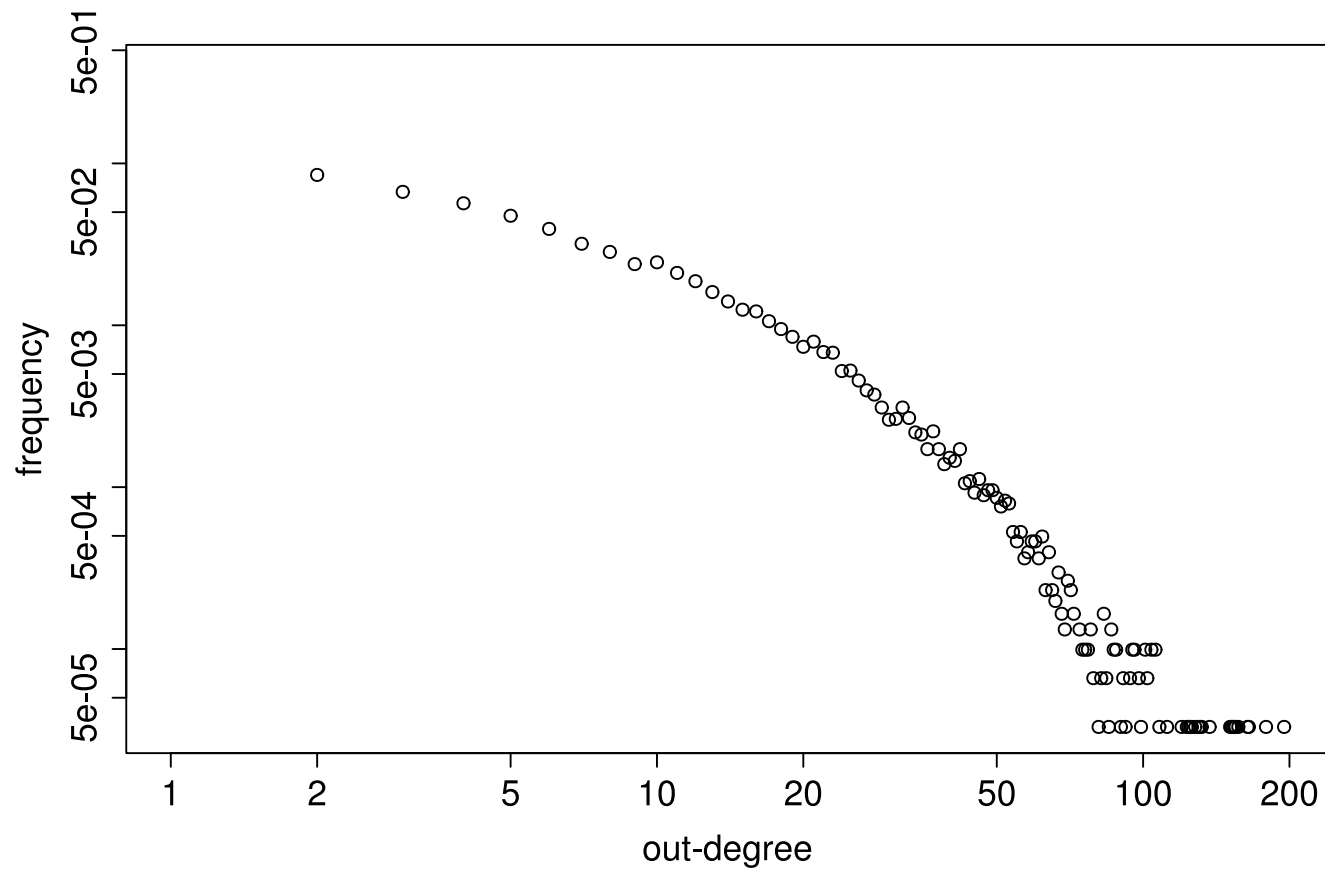
```
1 # In-degree distribution  
2 > plot(degree.distribution(jg, mode="in"), log="xy")
```

---



# Working with a somewhat bigger graph

```
1 # Out-degree distribution  
2 plot(degree.distribution(jg, mode="out"), log="xy")
```



# Working with a somewhat bigger graph

---

```
1 # Taking the largest component
2 > cl <- clusters(jg)
3 > jg2 <- subgraph(jg, which(cl$membership == which.max(cl$csizes)-1)-1)
4 > summary(jg2)
5 Vertices: 25389
6 Edges: 216718
7 Directed: TRUE
8 No graph attributes.
9 Vertex attributes: name, usid, parties, year, overruled, overruling,
10   oxford, liihc, indeg, outdeg, hub, hubrank, auth, authrank,
11   between, incent.
12 No edge attributes.
```

---

# Working with a somewhat bigger graph

---

```
1 > graph.density(jg2) # Density
2 [1] 0.0003362180
```



# Working with a somewhat bigger graph

---

```
1 > graph.density(jg2) # Density
2 [1] 0.0003362180
3
4 > transitivity(jg2) # Transitivity
5 [1] 0.1260031
```

# Working with a somewhat bigger graph

---

```
1 > graph.density(jg2) # Density
2 [1] 0.0003362180
3
4 > transitivity(jg2) # Transitivity
5 [1] 0.1260031
6
7 # Transitivity of a random graph of the same size
8 > g <- erdos.renyi.game(vcount(jg2), ecount(jg2), type="gnm")
9 > transitivity(g)
10 [1] 0.00064649
```

# Working with a somewhat bigger graph

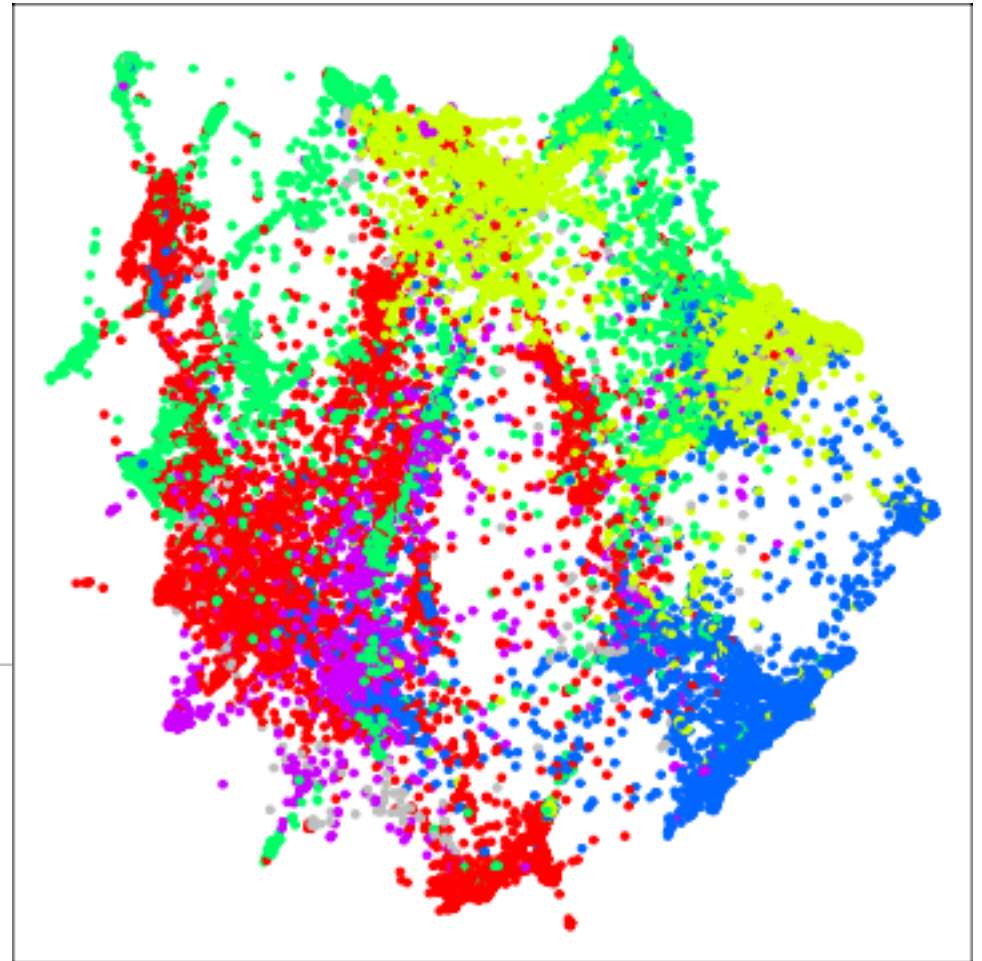
---

```
1 > graph.density(jg2) # Density
2 [1] 0.0003362180
3
4 > transitivity(jg2) # Transitivity
5 [1] 0.1260031
6
7 # Transitivity of a random graph of the same size
8 > g <- erdos.renyi.game(vcount(jg2), ecount(jg2), type="gnm")
9 > transitivity(g)
10 [1] 0.00064649
11
12 # Transitivity of a random graph with the same degrees
13 > g2 <- degree.sequence.game(degree(jg2,mode="all"), method="v1")
14 > transitivity(g2)
15 [1] 0.004107072
```

---

# Community structure detection

```
1 > fc <- fastgreedy.community(simplify(as.undirected(jg2)))
2 > memb <- community.to.membership(jg2,
3     fc$merges,
4     which.max(fc$modularity))
5 > lay <- layout.drl(jg2)
6 > jg3 <- graph.empty(n=vcount(jg2))
7 > colbar <- rainbow(5)
8 > col <- colbar[memb$membership+1]
9 > col[is.na(col)] <- "grey"
10 > plot(jg3, layout=lay, vertex.size=1,
11     vertex.label=NA, asp=FALSE,
12     vertex.color=col,
13     vertex.frame.color=col)
```



# Functionality, what can be calculated?

Fast (millions)	creating graphs (most of the time) • structural modification (add/delete edges/vertices) • subgraph • simplify • graph.decompose • degree • clusters • graph.density • is.simple, is.loop, is.multiple • articulation points and biconnected components • ARPACK stuff: page.rank, hub.score, authority.score, eigenvector centrality • transitivity • Burt's constraint • dyad & triad census, graph motifs • $k$ -cores • MST • reciprocity • modularity • closeness and (edge) betweenness <i>estimation</i> • <i>shortest paths from one source</i> • <i>generating <math>G_{n,p}</math> and <math>G_{n,m}</math> graphs</i> • <i>generating PA graphs with various PA exponents</i> • <i>topological sort</i>
Slow (10000)	closeness • diameter • betweenness • all-pairs shortest paths, average path length • most layout generators •
Very slow (100)	cliques • cohesive blocks • edge/vertex connectivity • maximum flows and minimum cuts • power centrality • alpha centrality • (sub)graph isomorphism

## Connection to other network/graph software

---

- graph package: `igraph.to.graphNEL`, `igraph.from.graphNEL`.

## Connection to other network/graph software

---

- graph package: `igraph.to.graphNEL`, `igraph.from.graphNEL`.
- Sparse matrices (Matrix package), `get.adjacency` and `graph.adjacency` supports them.

## Connection to other network/graph software

---

- graph package: `igraph.to.graphNEL`, `igraph.from.graphNEL`.
- Sparse matrices (Matrix package), `get.adjacency` and `graph.adjacency` supports them.
- `sna` and `network` R packages. Currently through adjacency matrices. Use namespaces!



## Connection to other network/graph software

---

- graph package: `igraph.to.graphNEL`, `igraph.from.graphNEL`.
- Sparse matrices (Matrix package), `get.adjacency` and `graph.adjacency` supports them.
- `sna` and `network` R packages. Currently through adjacency matrices. Use namespaces!
- Pajek. `.net` file format is supported.

## Connection to other network/graph software

---

- graph package: `igraph.to.graphNEL`, `igraph.from.graphNEL`.
- Sparse matrices (Matrix package), `get.adjacency` and `graph.adjacency` supports them.
- `sna` and `network` R packages. Currently through adjacency matrices. Use namespaces!
- Pajek. `.net` file format is supported.
- Visone. Use GraphML format.

## Connection to other network/graph software

---

- graph package: `igraph.to.graphNEL`, `igraph.from.graphNEL`.
- Sparse matrices (Matrix package), `get.adjacency` and `graph.adjacency` supports them.
- `sna` and `network` R packages. Currently through adjacency matrices. Use namespaces!
- Pajek. `.net` file format is supported.
- Visone. Use GraphML format.
- Cytoscape. Use GML format.

## Connection to other network/graph software

---

- graph package: `igraph.to.graphNEL`, `igraph.from.graphNEL`.
- Sparse matrices (Matrix package), `get.adjacency` and `graph.adjacency` supports them.
- `sna` and `network` R packages. Currently through adjacency matrices. Use namespaces!
- Pajek. `.net` file format is supported.
- Visone. Use GraphML format.
- Cytoscape. Use GML format.
- GraphViz. `igraph` can write `.dot` files.

# Connection to other network/graph software

---

- graph package: `igraph.to.graphNEL`, `igraph.from.graphNEL`.
- Sparse matrices (Matrix package), `get.adjacency` and `graph.adjacency` supports them.
- `sna` and `network` R packages. Currently through adjacency matrices. Use namespaces!
- Pajek. `.net` file format is supported.
- Visone. Use GraphML format.
- Cytoscape. Use GML format.
- GraphViz. `igraph` can write `.dot` files.
- In general. The GraphML and GML file formats are fully supported, many programs can read/write these.

# Acknowledgements

---

Tamás Nepusz

Peter McMahan, the BLISS, Walktrap, Spinglass, DrL projects

All the people who contributed code, sent bug reports, suggestions

The R project