

R software for network analysis

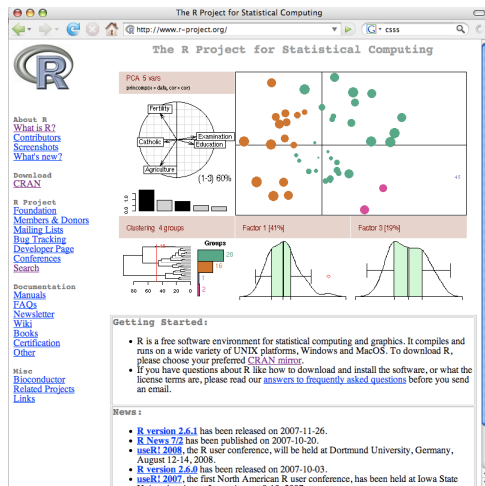
Dave Hunter

Penn State Dept. of Statistics
Joint with Mark and Carter and many others

MURI networks grant meeting, November 18, 2008

The R project and statnet

www.r-project.org



R: A statistical environment

- Open-source
- Extendible; ~1500 user-created packages exist
- Mostly written in C
- Well-maintained (if not always well-documented) by a core group of the world's top computational statisticians

www.statnet.org

Thanks Carter!

statnet

Introduction
Installation
Online Users Guide
Resources
Papers and Preprints
statnet Users Group
About Us
Citing statnet
License and source code attribution requirements

statnet

Software tools for the analysis, simulation and visualization of network data.

Welcome to statnet!

Version 2.1 released April 27, 2008

Special Issue of the Journal of Statistical Software

with 9 papers on statnet came out May 8, 2008

This website provides information on, background material for and access to the statnet suite of packages for network analysis. Directions for downloading statnet can be found under installation on the navigation bar to the left. The packages are written for the R statistical computing environment, so it runs on any computing platform that supports R. If you do not already have R installed, you will need to install it via the main Rweb resource-site, www.r-project.org. Instructions for installing R can also be found under [Installation](#).

statnet: A suite of R packages for network analysis

Includes packages:

- sna
- network
- ergm
- latentnet
- degreenet
- ...

Special issue (v 24) of *J. Stat. Soft.* devoted to statnet

A simple R package exploiting the R/C interface

```
% ls -RF1 pakij/
```

```
DESCRIPTION*
```

```
R/
```

```
src/
```

```
pakij//R:
```

```
my_runif.R
```

```
zzz.R*
```

```
pakij//src:
```

```
my_runif.c*
```

```
% R CMD INSTALL pakij
```

A simple R package exploiting the R/C interface

```
% ls -RF1 pakij/  
DESCRIPTION*  
R/  
src/
```

```
pakij//R:  
my_runif.R  
zzz.R*
```

```
pakij//src:  
my_runif.c*
```

```
% R CMD INSTALL pakij
```

pakij/R/my_runif.R

```
my_runif <- function(n) {  
  .C("hey_diddle_diddle",  
     n=as.integer(n),  
     foo=double(n)  
  )$foo  
}
```

A simple R package exploiting the R/C interface

```
% ls -RF1 pakij/  
DESCRIPTION*  
R/  
src/
```

```
pakij//R:  
my_runif.R  
zzz.R*
```

```
pakij//src:  
my_runif.c*
```

```
% R CMD INSTALL pakij
```

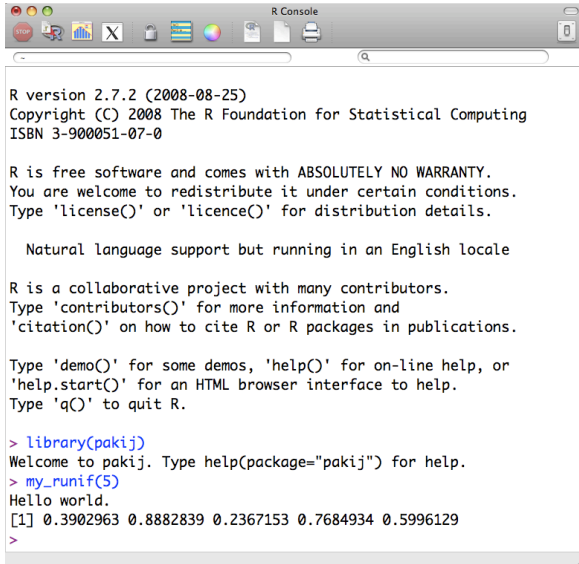
pakij/R/my_runif.R

```
my_runif <- function(n) {  
  .C("hey_diddle_diddle",  
    n=as.integer(n),  
    foo=double(n)  
  )$foo  
}
```

pakij/src/my_runif.c

```
#include <R.h>  
  
void hey_diddle_diddle (int *n, double *answer) {  
  Rprintf("Hello world.\n");  
  GetRNGstate(); /* R: enable uniform RNG */  
  for (int i=0; i<*n; i++)  
    answer[i] = unif_rand();  
  PutRNGstate(); /* Disable RNG before return */  
}
```

Using the pakij package



```
R Console  
R version 2.7.2 (2008-08-25)  
Copyright (C) 2008 The R Foundation for Statistical Computing  
ISBN 3-900051-07-0  
  
R is free software and comes with ABSOLUTELY NO WARRANTY.  
You are welcome to redistribute it under certain conditions.  
Type 'license()' or 'licence()' for distribution details.  
  
Natural language support but running in an English locale  
  
R is a collaborative project with many contributors.  
Type 'contributors()' for more information and  
'citation()' on how to cite R or R packages in publications.  
  
Type 'demo()' for some demos, 'help()' for on-line help, or  
'help.start()' for an HTML browser interface to help.  
Type 'q()' to quit R.  
  
> library(pakij)  
Welcome to pakij. Type help(package="pakij") for help.  
> my_runif(5)  
Hello world.  
[1] 0.3902963 0.8882839 0.2367153 0.7684934 0.5996129  
>
```

The R API

- Entry points in the R executable callable from C code
- Example: Carter exploits this in the `network` package.

In network/R:

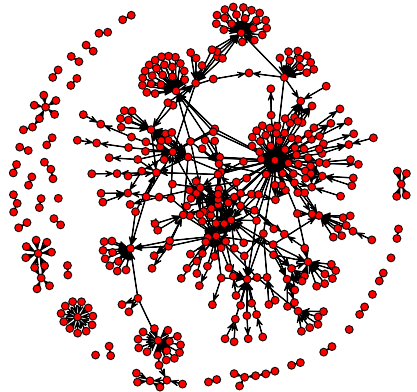
```
add.edge<-function(x, tail, head, names.eval=NULL,
                  vals.eval=NULL, edge.check=FALSE, ...){
  #Check to be sure we were called with a network
  if(!is.network(x))
    stop("add.edge requires an argument of class network.")
  #Do the deed
  invisible(.Call("addEdge_R", x, tail, head, names.eval, vals.eval,
                 edge.check, PACKAGE="network"))
}
```

In network/src:

```
SEXP addEdge_R(SEXP x, SEXP tail, SEXP head, SEXP namesEval,
              SEXP valsEval, SEXP edgeCheck) {
  ...
}
```

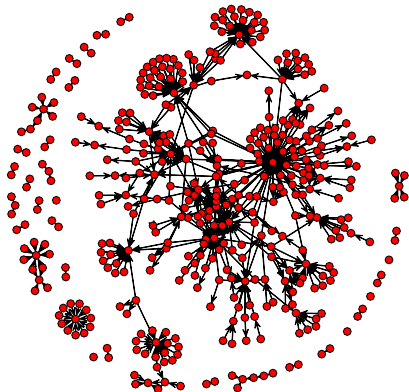

An E. Coli network (Salgado et al., 2001) in R

```
> plot(ecoli1)
```



An E. Coli network (Salgado et al., 2001) in R

```
> plot(ecoli1)  
> class(ecoli1)  
[1] "network"
```

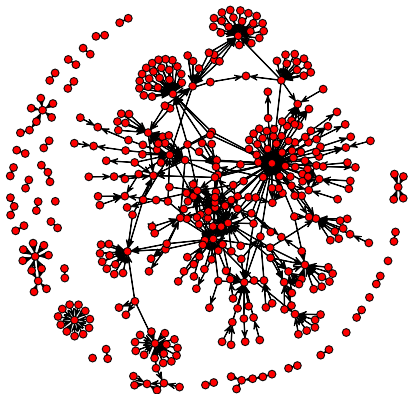


An E. Coli network (Salgado et al., 2001) in R

```
> plot(ecoli1)
> class(ecoli1)
[1] "network"
> summary(ecoli1)
Network attributes:
vertices: 423
  directed : TRUE
  hyper : FALSE
  loops : FALSE
  multiple : FALSE
  bipartite : FALSE
total edges= 519
density = 0.002907465
```

Vertex attributes:

```
self:
FALSE TRUE
 364   59
```



An E. Coli network (Salgado et al., 2001) in R

```
> plot(ecoli1)
```

```
> class(ecoli1)
```

```
[1] "network"
```

```
> summary(ecoli1)
```

```
Network attributes:
```

```
vertices: 423
```

```
directed : TRUE
```

```
hyper : FALSE
```

```
loops : FALSE
```

```
multiple : FALSE
```

```
bipartite : FALSE
```

```
total edges= 519
```

```
density = 0.002907465
```

```
Vertex attributes:
```

```
self:
```

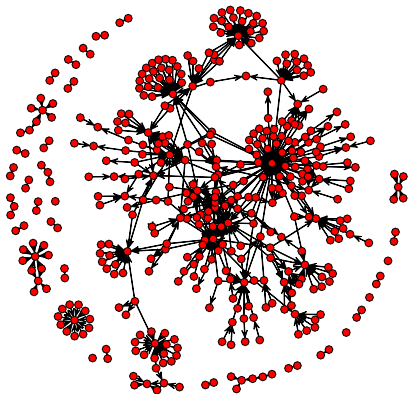
```
FALSE TRUE
```

```
364 59
```

```
> summary(ecoli1 ~ edges + odegree(0:5))
```

```
edges odegree0 odegree1 odegree2 odegree3 odegree4 odegree5
```

```
519 81 229 71 25 13 3
```



A simple ERGM using the `ergm` package in R

The `ecoli1` network has 519 edges, 423 nodes (thus 423×422 possible directed edges). Take $g(y) = \#$ edges in y :

$$\begin{aligned} P(Y = y) &\propto \exp\{\theta \times \# \text{ edges in } y\} \\ &= \left(\frac{p}{1-p}\right)^{(\# \text{ edges in } y)} \end{aligned}$$

A simple ERGM using the `ergm` package in R

The `ecoli1` network has 519 edges, 423 nodes (thus 423×422 possible directed edges). Take $g(y) = \#$ edges in y :

$$\begin{aligned} P(Y = y) &\propto \exp\{\theta \times \# \text{ edges in } y\} \\ &= \left(\frac{p}{1-p}\right)^{(\# \text{ edges in } y)} \end{aligned}$$

```
> modell <- ergm(ecoli1 ~ edges)
> modell$coef
  edges
-5.837562
```

This means the MLE is $\hat{\theta} = -5.84$. It is exact because $g(y)$ in this example ensures independent Y_{ij} , so MPLE=MLE!

A simple ERGM using the ergm package in R

The `ecoli1` network has 519 edges, 423 nodes (thus 423×422 possible directed edges). Take $g(y) = \#$ edges in y :

$$\begin{aligned} P(Y = y) &\propto \exp\{\theta \times \# \text{ edges in } y\} \\ &= \left(\frac{p}{1-p}\right)^{(\# \text{ edges in } y)} \end{aligned}$$

```
> modell <- ergm(ecoli1 ~ edges)
> modell$coef
  edges
-5.837562
```

This means the MLE is $\hat{\theta} = -5.84$. It is exact because $g(y)$ in this example ensures independent Y_{ij} , so MPLE=MLE!

Observe:

```
> log(519 / (423 * 422 - 519))
[1] -5.837562
```

Another simple ERGM

Let's consider an edge to have one of four categories, depending on whether or not each of its endpoints is self-regulating:

```
> model2 <- ergm(ecoli1 ~ nodemix("self"))
> summary(model2)
```

Pseudolikelihood Results:

	Estimate	Std. Error	MCMC s.e.	p-value
<code>mix.self.FALSE.FALSE</code>	-6.61974	0.07543	NA	<1e-04
<code>mix.self.TRUE.FALSE</code>	-7.48923	0.28874	NA	<1e-04
<code>mix.self.FALSE.TRUE</code>	-4.22686	0.05730	NA	<1e-04
<code>mix.self.TRUE.TRUE</code>	-5.04049	0.21389	NA	<1e-04

This choice of $g(y)$ also ensures independent Y_{ij} , so the MLE and the MPLE are the same. Thus, the MLE is found analytically and there is no MCMC standard error.