

GIBBS Function

#Gibbs function receives four parameters: "roe" for correlation, "N" for number of realizations, "lag" for determining how many realizations to skip between saves, and "burnin" for determining how many realizations to skip before starting to save. Gibbs generates N independent random normal values.

```
gibbs <- function(roe,N,lag,burnin) {  
  
  #Set N to be Nations*lag+burnin  
  N <- N*lag + burnin  
  #Initialize x and y  
  x <- 80  
  y <- 80  
  #Initialize vectors  
  x.s <- NULL  
  y.s <- NULL  
  
  for(i in 1:N) {  
  
    #generate a "y"  
    y <- rnorm(1, roe*x, sqrt(1-roe^2))  
    #generate an "x"  
    x <- rnorm(1, roe*y, sqrt(1-roe^2))  
  
    #if i is greater than burnin and if i is a multiple of the lag,  
    store x and y  
    if(i > burnin) {  
      if(i %% lag == 0) {  
        x.s <- c(x.s,x)  
        y.s <- c(y.s,y)  
      }  
    }  
  }  
  vectors <- list("x" = x.s, "y" = y.s)  
  return(vectors)  
}
```

##1##

```
#Run Gibbs with N=500, lag=1, and burnin=0
```

```
roe <- 0.75
```

```
N <- 500
```

```
lag <- 1
```

```
burnin <- 0
```

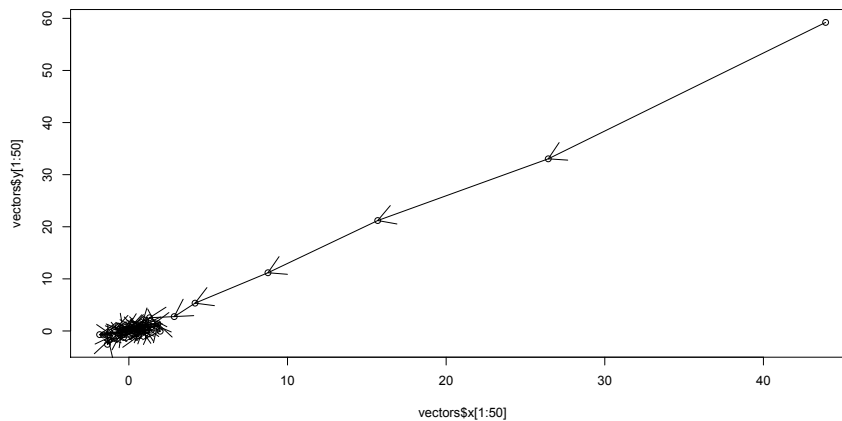
```
vectors <- gibbs(roe,N,lag,burnin)
```

#a

```
#Plot first 50 values and connect the points in order generated  
using arrows command
```

```
plot(vectors$x[1:50],vectors$y[1:50])
```

```
arrows(vectors$x[1:49],vectors$y[1:49],vectors$x[2:50],vectors$y[  
2:50],size=0.1,open=T)
```



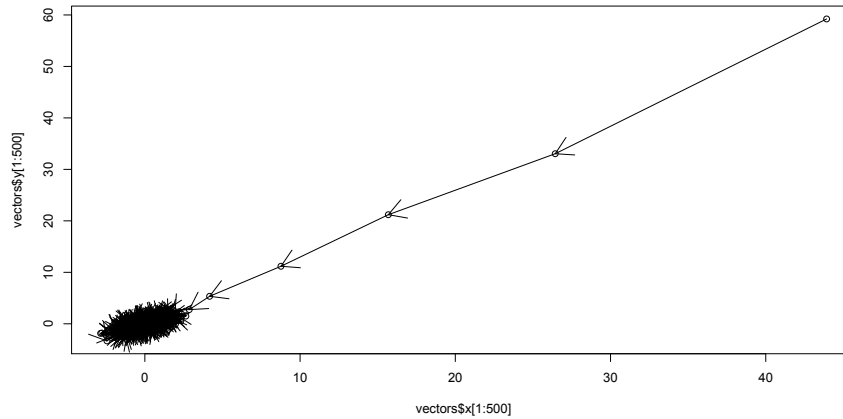
#b

```
#Plot all 500 values with arrows to illustrate possible values  
for lag and burnin
```

```
plot(vectors$x[1:500],vectors$y[1:500])
```

```
arrows(vectors$x[1:499],vectors$y[1:499],vectors$x[2:500],vectors  
$y[2:500],size=0.1,open=T)
```

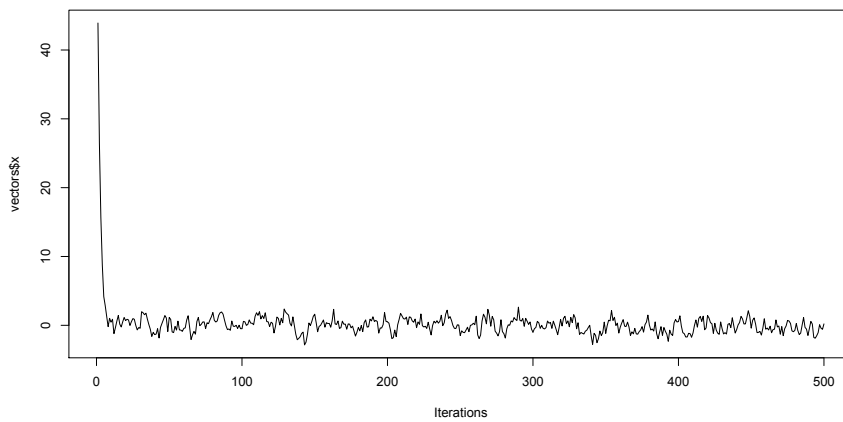
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#c

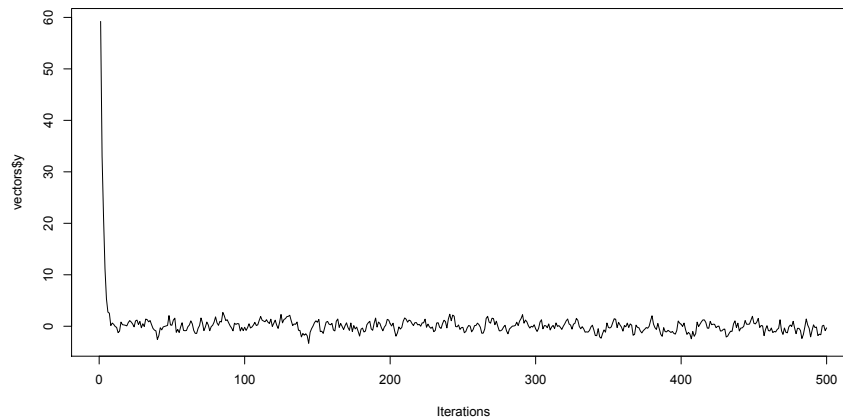
#Use x.s and y.s from initial run to determine values for burnin
based on drop in ts.plot - take larger drop to set burnin.

```
ts.plot(vectors$x,xlab="Iterations")
```



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```
ts.plot(vectors$y,xlab="Iterations")
```

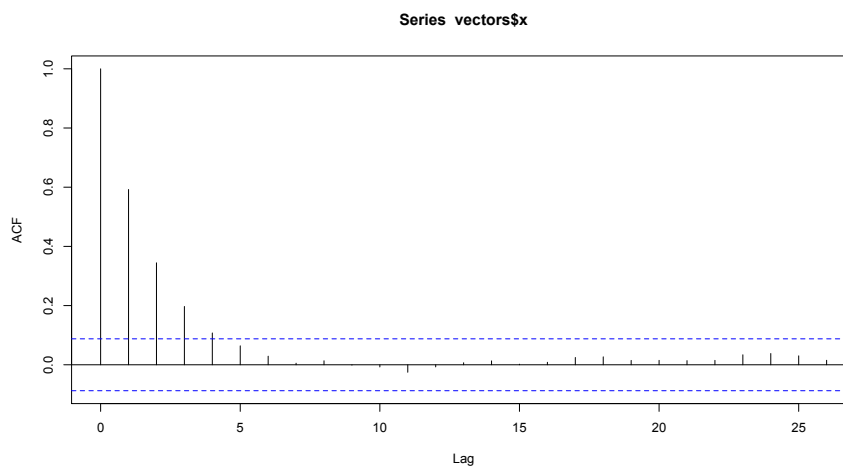


```
#Set burnin to 20 (10 may work but choose 20 to be conservative)
```

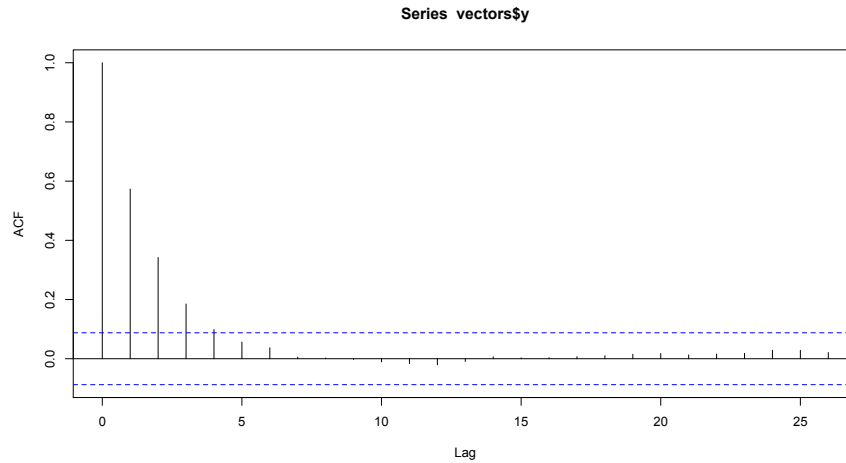
#d

```
#Use x.s and y.s from initial run to determine values for lag  
based on largest acf - autocorrelation - for both vectors, x.s  
and y.s (conservatively select bar that falls below the line).
```

```
acf(vectors$x)
```



`acf(vectors$y)`



```
#Set lag to 5
```

#e

```
#Run Gibbs with N=10000, lag=5, and burnin=20 and verify  
normality
```

```
roe <- 0.75
```

```
N <- 10000
```

```
lag <- 5
```

```
burnin <- 20
```

```
vectors <- gibbs(roe,N,lag,burnin)
```

```
mean(vectors$x)
```

```
#[1] -0.003423303
```

```
var(vectors$x)
```

```
#[1] 1.014041
```

```
quantile(vectors$x)
```

```
#          0%          25%          50%          75%         100%  
#-3.792598217 -0.683614847  0.002497831  0.655396187  3.577334894
```

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```
mean(vectors$y)
#[1] -0.000683986
```

```
var(vectors$y)
#[1] 1.006094
```

```
quantile(vectors$y)
#          0%          25%          50%          75%         100%
#-3.61015779 -0.68406521 -0.01410746  0.66701307  3.92869627
```

##2##

```
roe <- -0.9
N <- 15000
lag <- 5
burnin <- 20
```

```
r.s <- NULL
p.q1 <- NULL
p.q2 <- NULL
p.q1n4 <- NULL
p <- 0
```

```
while(roe <= .9) {
```

```
  #create roe vector
  r.s <- c(r.s,roe)
```

```
  #Run gibbs to generate two independent normal random variables
  vectors <- gibbs(roe,N,lag,burnin)
```

```
  #initialize xy.s - vector to store scenarios
```

```
  xy.s <- NULL
  #restrict x and y to quadrant 1
  xy.s <- (vectors$x > 0 & vectors$y > 0)
  count <- 0
  count <- length(xy.s[xy.s=="TRUE"])
```

```
  p <- count/N
  #accumulate probabilities for quadrant 1
  p.q1 <- c(p.q1,p)
```

```
  xy.s <- NULL
```

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```
#restrict x and y to quadrant 2
xy.s <- (vectors$x < 0 & vectors$y > 0)
count <- 0
  count <- length(xy.s[xy.s=="TRUE"])
p <- count/N
#accumulate probabilities for quadrant 1
p.q2 <- c(p.q2,p)

xy.s <- NULL
#restrict x and y to quadrants 1 and 4
xy.s <- (vectors$x > 0)
count <- 0
  count <- length(xy.s[xy.s=="TRUE"])
p <- count/N
#accumulate probabilities for quadrants 1 and 4
p.q1n4 <- c(p.q1n4,p)

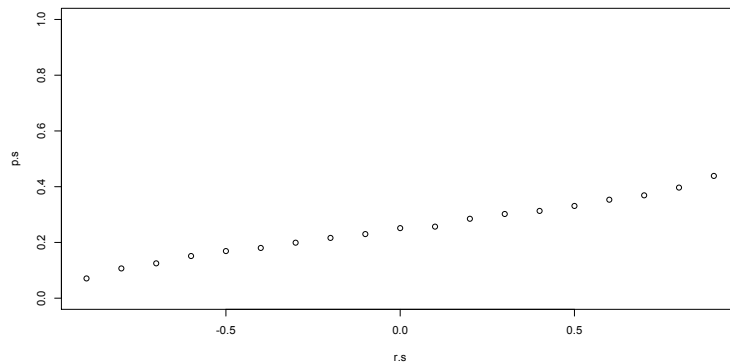
#increment roe by 0.1
roe <- roe + 0.1
}

#Plot probability vs. roe with vertical axis limits (0,1) for
each scenario
```

#a

```
plot(r.s,p.q1,ylim=c(0,1))
```

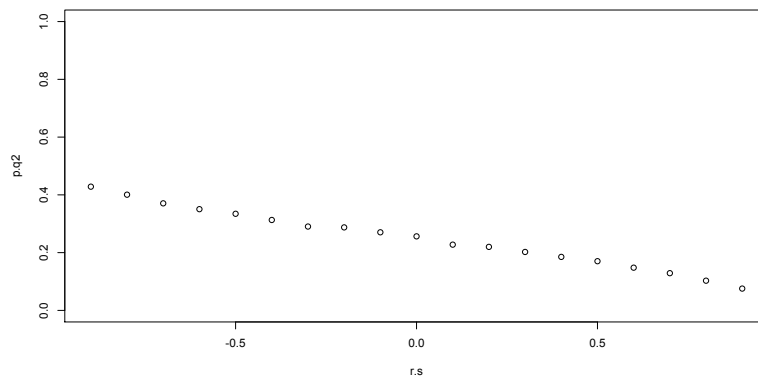
$P(X > 0 \text{ and } Y > 0)$ approaches 0 as ρ approaches -1, is 0.25 when $\rho=0$, and approaches 0.5 as ρ approaches 1.



#b

```
plot(r.s,p.q2,ylim=c(0,1))
```

$P(X < 0 \text{ and } Y > 0)$ approaches 0.5 as ρ approaches -1, is 0.25 when $\rho=0$, and approaches 0 as ρ approaches 1.



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#c

```
plot(r.s,p.q1n4,ylim=c(0,1))
```

$P(X > 0)$ is 0.5 as roe moves from -0.9 to 0.9.

