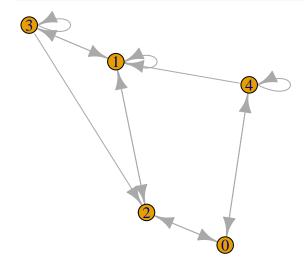
# Ma2

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```
Various packages in use:
```

```
set.seed(98765432)
library(igraph)
##
## Attaching package: 'igraph'
## The following objects are masked from 'package:stats':
##
       decompose, spectrum
##
## The following object is masked from 'package:base':
##
##
       union
library(expm)
## Loading required package: Matrix
##
## Attaching package: 'expm'
## The following object is masked from 'package:Matrix':
##
##
       expm
w -> wide t -> tilde h -> hat m -> matrix
```



b)

```
irreducibility <- function(P, M){
    P.new <- P
    for(i in 2:M){
        Pk <- P % % i
            P.new <- P.new + Pk
}

P.bar <- (1/M)*P.new

if(all(P.bar > 0)){
        print("Transition Matrix is Irreducible")

}
else{
        print("YTransition Matrix is Reducible")
}

irreducibility(P,5)
```

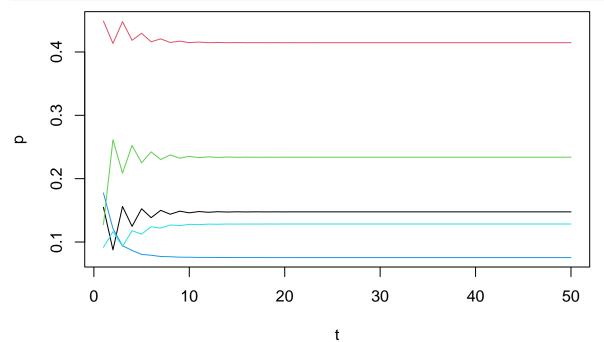
## [1] "Transition Matrix is Irreducible"

```
c)
K = 50
M = 5
p0 <- as.vector(c(0.013, 0.214, 0.197, 0.375, 0.201))

K <- 50
pk <- matrix(nrow = K, ncol = M)
for (i in 1:K) {
    pk[i, ] <- p0 %*% (P%*%i)
}

plot(x = 1:K, y = pk[, 1], type = "l", col = 1, xlab = "t", ylab = "p",
ylim = c(min(pk), max(pk)))

lines(x = 1:K, y = pk[, 2], type = "l", col = 2)
lines(x = 1:K, y = pk[, 3], type = "l", col = 3)
lines(x = 1:K, y = pk[, 4], type = "l", col = 4)
lines(x = 1:K, y = pk[, 5], type = "l", col = 5)</pre>
```



d) Stationary distribution by computing eigenvalues of P

```
pi <- eigen(t(P))$vectors[,1]/sum(eigen(t(P))$vectors[,1])

pi_confirm <- pi %*% P

print(pi_confirm - pi)

## 0 1 2 3 4

## [1,] 5.551115e-17 -1.665335e-16 1.110223e-16 4.163336e-17 -5.551115e-17

e) Stationary distribution by iterting 50 times: π = π · P<sup>50</sup>

pi_converged <- p0 %*% (P%^%50)</pre>
```

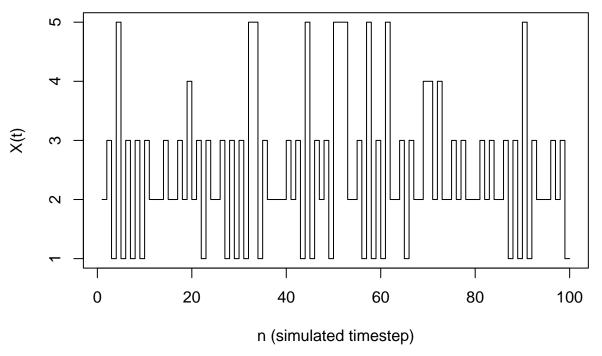
As we see the difference is small

```
print(pi-pi_converged)
```

```
## 0 1 2 3 4
## [1,] 4.76654e-11 2.764483e-11 -5.232206e-11 -4.16657e-12 -1.882194e-11
```

```
problem 2.2
a) function for simulating large samples
simulMarkov<-function(p0,mP,n){</pre>
M<-dim(mP)[1]
X<-rep(0,n)</pre>
X[1] <- sample(M,1,prob=p0)</pre>
for(t in 2:n) X[t] <- sample(M,1,prob=mP[X[t-1],])</pre>
return(X)
}
n<-100
X<-rep(0,n)</pre>
X[1] <-sample(M,1,prob=p0)</pre>
for(t in 2:n) X[t] <- sample(M,1,prob=P[X[t-1],])</pre>
problem 2.2
b)
plot(x = 1:n, y = X, main = "Simulation of markov Chain based on (P, p0)",
     xlab = "n (simulated timestep)", ylab = "X(t)", type = "S")
```

## Simulation of markov Chain based on (P, p0)

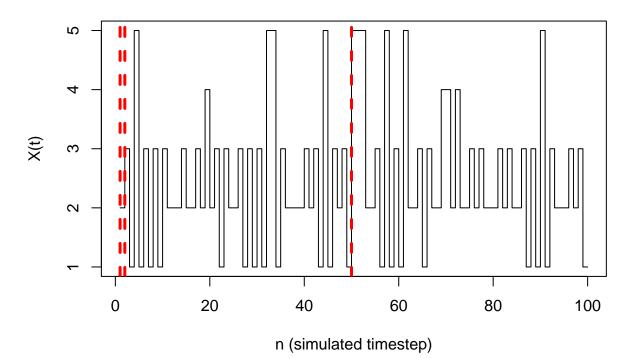


```
##problem 2.2
c)
iz<-4
S_all<-which(X==iz)
Tn<-length(S_all)
Tn
```

## [1] 4

# 

## Simulation of markov Chain based on (P, p0)



```
\frac{\text{problem } 2.3}{\text{a})}
```

```
#set.seed(1234)
X.10000 <- simulMarkov(p0=p0, mP=P,n=10000)

iz.10000 = 4 #X.10000[1]
S_all<-which(X.10000 == iz.10000)

Tn.10000<-length(S_all)

Tn.10000
## [1] 834
b)
tau.10000<-S_all[2:Tn.10000]-S_all[1:(Tn.10000-1)]

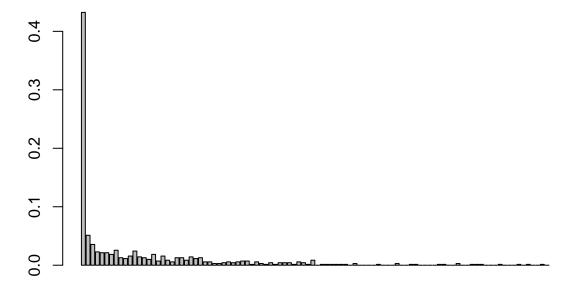
f_i0<-c(1:100)
for(j in 1:100){
f_i0[j]<-(1/10000)*length(which(tau.10000==j))</pre>
```

 $f_i0$ 

```
c)
set.seed(123)
i0 <- 4
Tn.1 <- length(which(simulMarkov(p0 = p0, mP=P, n=10000) == i0))
tau.1 <- S_all[2:Tn.1] - S_all[1:(Tn.1 - 1)]
K <- 100

f_hat <- numeric(K)

for (s in 1:K) {
    f_hat[s] <- length(which(tau.1 == s))
}
f_hat <- f_hat / (Tn.1 - 1)</pre>
barplot(f_hat, xlab = "f_hat")
```



### f\_hat

```
i)
mu_hat_i0 \leftarrow (1/(Tn.1-1))*sum(tau.1[-1]) #AKA tou_bar
mu_hat_i0
## [1] 12.21479
  ii)
mu\_tilde\_i0 \leftarrow sum((1:K) * f\_hat)
mu_tilde_i0
## [1] 11.5633
 iii)
pi_i0 <- pi_converged[i0]</pre>
pi_i0
## [1] 0.07545784
rbind("1 / pi_i0" = 1 / pi_i0,
"mu_hat_i0" = mu_hat_i0,
"mu_tilde_i0" = mu_tilde_i0)
##
                    [,1]
## 1 / pi_i0 13.25243
## mu_hat_i0 12.21479
## mu_tilde_i0 11.56330
  d)
n = 10000
m = n - 1
X = simulMarkov(p0 = p0, mP=P, n=n)
```

```
wtpi <- whpi <- numeric(5)</pre>
    for(i in 1:5){
          wtpi[i] <-m^(-1)*length((which(X==i)))
    }
wtpi
## [1] 0.13981398 0.42854285 0.23202320 0.07760776 0.12211221
sum(wtpi)
## [1] 1.0001
  e)
Emprircal Transition matrix:
P_{\text{hat}} \leftarrow \text{matrix}(\text{nrow} = 5, \text{ncol} = 5, 0)
wtpi <-c()
whpi <-c()
for (t in 1:(length(X) - 1)) P_hat[X[t],
              X[t + 1] <- P_hat[X[t], X[t + 1]] + 1
 for (i in 1:5) P_hat[i, ] <- P_hat[i, ] / sum(P_hat[i, ])</pre>
P_hat
##
              [,1]
                        [,2]
                                   [,3]
                                              [,4]
## [1,] 0.0000000 0.0000000 0.4907010 0.00000000 0.5092990
## [2,] 0.0000000 0.5376896 0.3633606 0.09894982 0.0000000
## [3,] 0.4312204 0.5687796 0.0000000 0.00000000 0.0000000
## [4,] 0.0000000 0.4484536 0.0992268 0.45231959 0.0000000
## [5,] 0.3259623 0.2571663 0.0000000 0.00000000 0.4168714
problem 2.4
we apply the amse matrix as in 2.3 (e)
  a)
P_hat
             [,1]
                        [,2]
                                   [,3]
                                              [,4]
## [1,] 0.0000000 0.0000000 0.4907010 0.00000000 0.5092990
## [2,] 0.0000000 0.5376896 0.3633606 0.09894982 0.0000000
## [3,] 0.4312204 0.5687796 0.0000000 0.00000000 0.0000000
## [4,] 0.0000000 0.4484536 0.0992268 0.45231959 0.0000000
## [5,] 0.3259623 0.2571663 0.0000000 0.00000000 0.4168714
comparing by subtracting P from \hat{P} b)
P-P_hat
##
               0
## 0 0.00000000 0.00000000 0.007298999
                                             0.00000000 -0.007298999
## 1 0.00000000 -0.010689615 0.002639440
                                             0.008050175 0.000000000
## 2 0.019779646 -0.019779646 0.000000000
                                             0.00000000 0.00000000
## 3 0.000000000 0.025546392 0.014773196 -0.040319588 0.000000000
## 4 0.002037674 -0.008166257 0.000000000 0.000000000 0.006128583
```

as we see the difference is about two to thre decimals