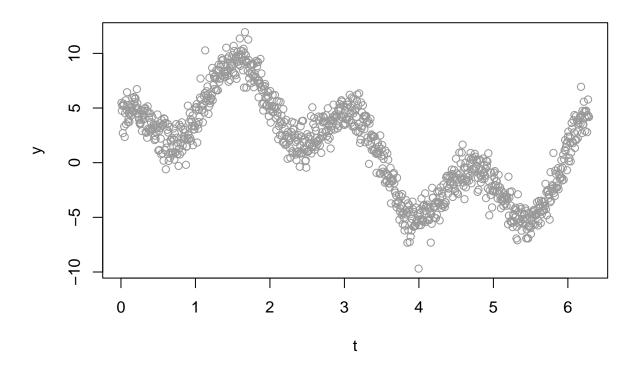
R입문 기말고사

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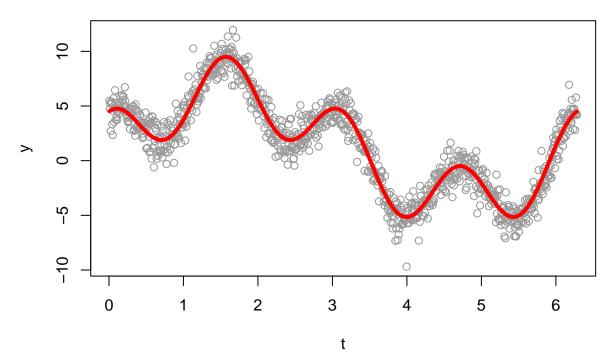
library(tidyverse)

1번

```
### 1)
e=c()
for(i in 1:1000){
   e[i] = rnorm(1)
    }
head(e)
## [1] 0.9497786 0.1874779 0.6281326 -1.9025805 0.7670813 0.5950860
### 2)
t=c()
for(i in 1:1000){
   t[i]=(2*pi/1000)*i
x1=c()
for(i in 1:1000){
   x1[i]=sin(t[i])
x2=c()
for(i in 1:1000){
   x2[i]=cos(t[i]*4)
    }
head(x1)
## [1] 0.006283144 0.012566040 0.018848440 0.025130095 0.031410759 0.037690183
## [1] 0.9996842 0.9987370 0.9971589 0.9949510 0.9921147 0.9886517
### 3)
y=c()
for(i in 1:1000){
    y[i] = 1.5 + 5*x1[i] + 3*x2[i] + e[i]
plot(t,y,col="gray60")
```



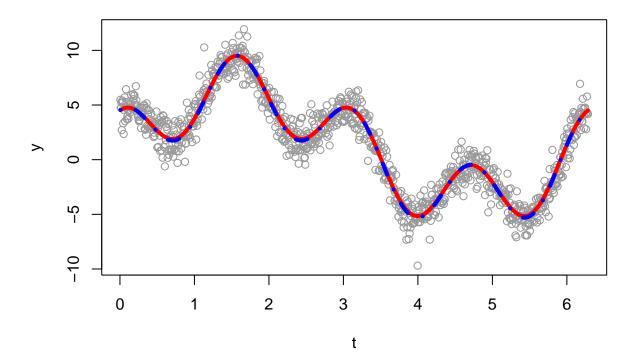
```
### 4)
X=cbind(rep(1,1000),x1,x2)
head(X)
                   x1
## [1,] 1 0.006283144 0.9996842
## [2,] 1 0.012566040 0.9987370
## [3,] 1 0.018848440 0.9971589
## [4,] 1 0.025130095 0.9949510
## [5,] 1 0.031410759 0.9921147
## [6,] 1 0.037690183 0.9886517
beta = rbind(1.5,5,3)
beta
##
        [,1]
## [1,] 1.5
## [2,] 5.0
## [3,] 3.0
Xbeta=array(X %*% beta)
plot(t,y,col="gray60")
lines(t,Xbeta,col="red",lwd=4)
```



```
### 6)
beta_hat=solve(t(X) %*% X) %*% t(X) %*% y
beta_hat

## [,1]
## 1.439189
## x1 4.997604
## x2 3.089615
β̂은 β와 거의 유사하다.

### 7)
Xbeta_hat=array(X %*% beta_hat)
plot(t,y,col="gray60")
lines(t,Xbeta,col="red",lwd=4)
lines(t,Xbeta_hat,col="blue",lty=4,lwd=4)
```



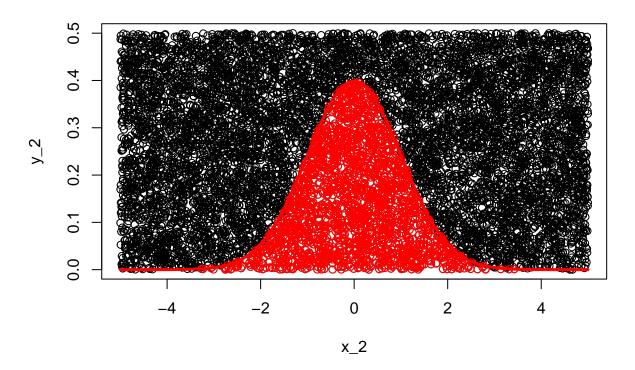
```
### 1)

x_2=runif(10000,-5,5)
y_2=runif(10000)*0.5

x_21=seq(from=-5, to=5, by=0.01)
y_21=(1/sqrt(2*pi))*exp((-1/2)*(x_21**2))
check=function(x,y){
    y<(1/sqrt(2*pi))*exp((-1/2)*(x**2))
    }

tst = c()
for (i in 1:10000) tst[i] = check(x_2[i],y_2[i])
sum(tst)

## [1] 2013
plot(x_2,y_2)
lines(x_21,y_21,col='red',lwd=3)
points(x_2[tst],y_2[tst],col='red')
```



빨간 점들 중에서 -1.96~1.96 사이의 점의 개수 비율 sum(x_2[tst]>=-1.96 & x_2[tst]<=1.96) / sum(tst)

```
## [1] 0.9483358
```

```
### 2)
x_norm=c()
count=0
for(i in 1:1000){
    x_norm[i]=rnorm(1)
    if(x_norm[i]>=-1.96 & x_norm[i]<=1.96){
        count = count +1
        }
    }
count</pre>
```

[1] 957

3번

```
### type A 생존확률
nor1=c()
spec=c()
nor8=c()
count=c()
for(i in 1:10000){
    nor1[i]=sum(cumprod(rbinom(20,size=1,0.5))) # 일반인 1이 해결한 수
```

```
spec[i]=sum(cumprod(rbinom(20-nor1[i]-1,size=1,0.95))) # 장인이 해결한 수
                            if(spec[i]>=(20-nor1[i]-1-1)){
                                          count[i] = 1
                           }else{
                                          nor8[i]=sum(cumprod(rbinom(20-nor1[i]-1-spec[i]-1,size=1,0.5)))
                                                       if(nor8[i] == (20-nor1[i]-1-spec[i]-1)){
                                                                     count[i] = 1
                                                       }else{
                                                                     count[i] = 0
                                                                     }
                                          }
sum(count)/10000
## [1] 0.4404
### type b 생존 확률
type_b=rbinom(10000, 20, 0.5)
sum(type_b <= 7)/10000
## [1] 0.1296
type A가 많이 유리하다
4번
df=read_csv('https://raw.githubusercontent.com/guebin/2021IR/master/_notebooks/covid19.csv') head(df)
### 1) df %>% group_by(year) %>% summarise(sum=sum(cases))
2)
df \%>\% filter(year==2020 \& month==2 \& day<=15) \%>\% group\_by(prov) \%>\% summarise(prov\_cases=sum(cases))
%>% arrange(desc(prov_cases)) #### 경기
3)
\label{eq:cases} $$ df \%>\% filter(year==2020 \& month==2 \& day>15) \%>\% group\_by(prov) \%>\% summarise(prov\_cases=sum(cases)) $$ (a) $$ (a) $$ (b) $$ (b) $$ (b) $$ (c) $$ (
%>% arrange(desc(prov cases))
```

대구