# HW2\_Yi\_Xiong

### Yi Xiong

## 2021/2/14

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its mean $\overline{x}$ , and variance $\hat{\sigma}^2$ set.seed(6666)  x<-rnorm(100)  # mean  (x_mean<-mean(x))	pute

a) Repeat this experiment 100 times and obtain 100 estimates of  $\overline{x}$ , and  $\widehat{\sigma}^2$ 

```
xt \leftarrow data.frame(T = 1:100)
for (i in 1:100) {
  xt <- cbind(xt, rnorm(100))</pre>
colnames(xt)[-1] <- paste0("S", 1:100)</pre>
# mean
head(xt_mean <- colMeans(xt[, -1]))</pre>
                           S2
## -0.13158945 0.09087463 0.08560193 -0.04104514 0.14779573 0.12226931
# variance
head(xt_var <- apply(xt[, -1], 2, var))</pre>
## 0.9685090 0.7678836 1.0244348 0.9493001 1.0950379 1.0968314
b) Compute the mean, variance of \overline{x}_i, for 1 \le i \le 100
(xt_mean_mean <- mean(xt_mean))</pre>
## [1] 0.008002791
# variance
(xt_mean_var <- var(xt_mean))</pre>
## [1] 0.00922678
c) Compute the mean, variance of \widehat{\sigma}_i^2, for 1 \le i \le 100
# mean
(xt_var_mean <- mean(xt_var))</pre>
## [1] 0.9880721
# variance
(xt_var_var <- var(xt_var))</pre>
## [1] 0.02034064
2) Generate \varepsilon_t \sim \text{iid N}(0,1) for 1 \le t \le 500
```

,

```
t<-500
set.seed(6666)
x<-rnorm(500)
```

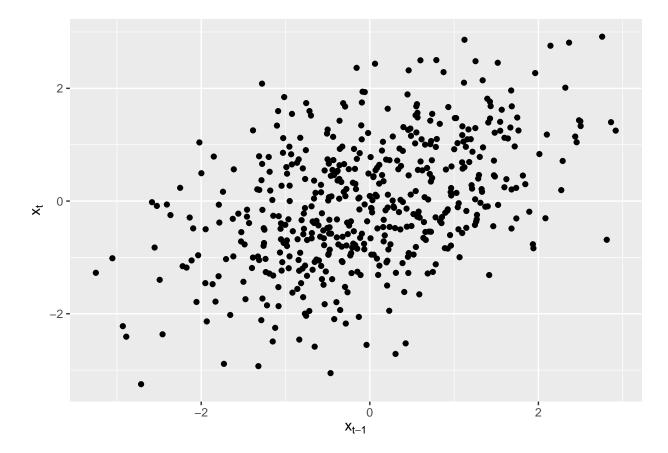
a) Define  $x_t = \varepsilon_t + 0.5\varepsilon_{t-1}$  called MA(1) process

```
xt<-c(x[1])
for (t in 2:500){
  xt[t]<-x[t]+0.5*x[t-1]
}</pre>
```

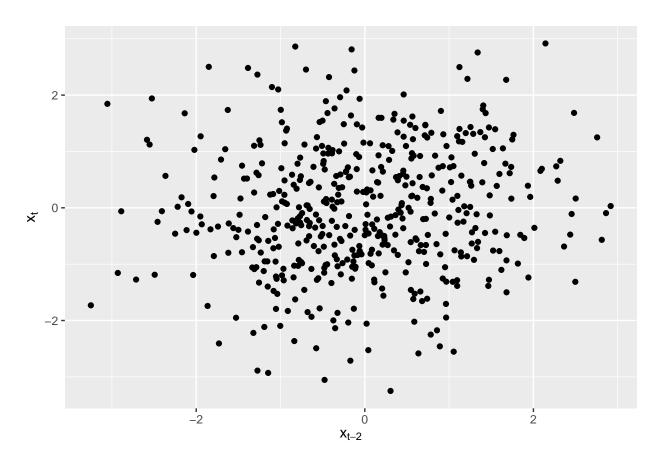
b) Plot  $x_t$  (vs)  $x_{t\text{-}1}$  ,  $x_t$  (vs)  $x_{t\text{-}2}$  ,  $x_t$  (vs)  $_t$ 

```
library(ggplot2)
# t vs t-1

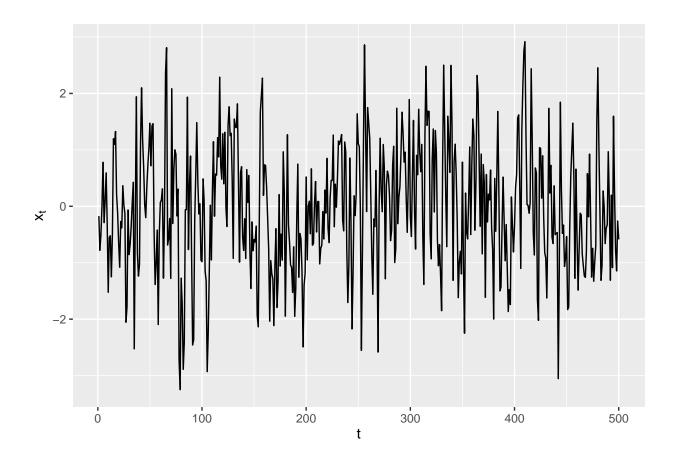
df<-data.frame(x=xt[1:(t-1)],y=xt[2:t])
p<-ggplot(df,aes(x=x,y=y))+
   geom_point()+
   xlab(expression(x[t-1]))+
   ylab(expression(x[t]))
p</pre>
```



```
# t vs t-2
df<-data.frame(x=xt[1:(t-2)],y=xt[3:t])
p<-ggplot(df,aes(x=x,y=y))+
   geom_point()+
   xlab(expression(x[t-2]))+
   ylab(expression(x[t]))
p</pre>
```



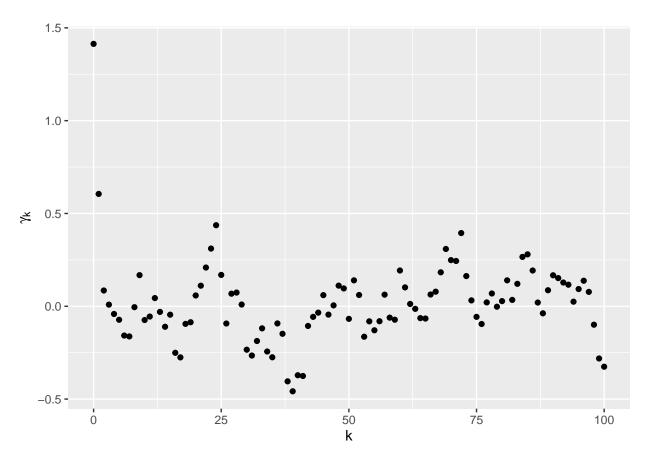
```
# xt vs t
df<-data.frame(x=1:t,y=xt)
p<-ggplot(df,aes(x=x,y=y))+
  geom_line()+
  xlab(expression(t))+
  ylab(expression(x[t]))
p</pre>
```



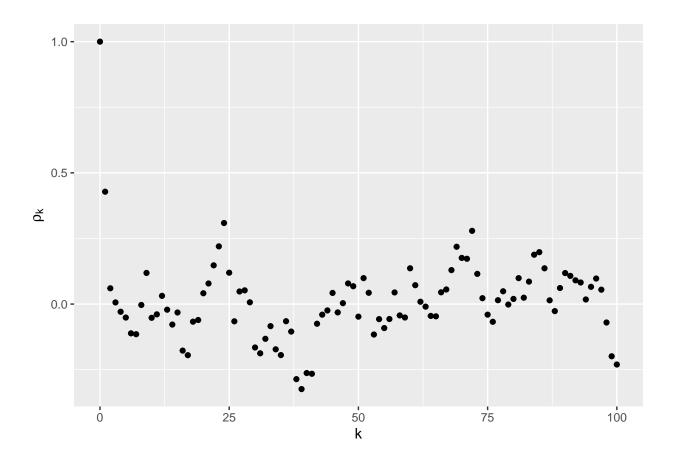
### c) Compute the mean, variance and autocorrelation of $\mathbf{x}_t$ and plot

```
# mean
(xt_mean<-mean(xt))</pre>
## [1] -0.03393109
# var
(xt_var<-var(xt))</pre>
## [1] 1.252277
# autocorrelation
getCov<-function(k){</pre>
  x1<-xt[1:100]
  x2 < -xt[(k+1):(100+k)]
  cov(x1,x2)
}
k<-0:100
cov1<-sapply(k,getCov)</pre>
dfCov<-data.frame(k=k,cov=cov1)</pre>
p<-ggplot(dfCov,aes(x=k,y=cov))+</pre>
```

```
geom_point()+
xlab(expression(k))+
ylab(expression(gamma[k]))
p
```



```
p<-cov1/cov1[1]
dfP<-data.frame(k=k,p=p)
p<-ggplot(dfP,aes(x=k,y=p))+
   geom_point()+
   xlab(expression(k))+
   ylab(expression(rho[k]))
p</pre>
```

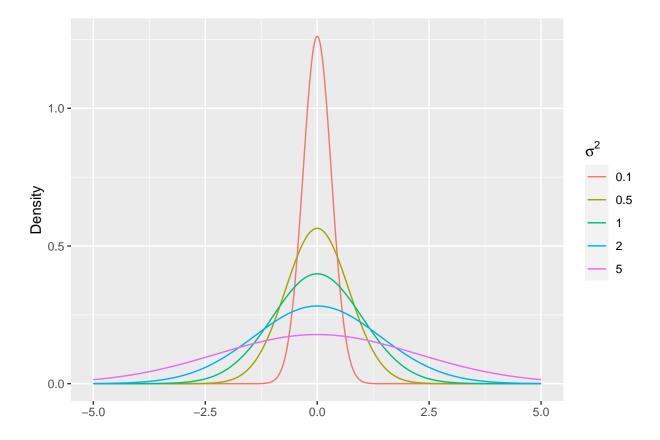


3) Plot  $N\left(0,\sigma^2\right)$  for  $\sigma^2=$  0.1, 0.5, 1.0, 2.0, 5.0 on the same plot and comment

```
# ggplot() + xlim(-5, 5) + geom_function(
   fun = function(x)
#
     dnorm(x, sd = sqrt(0.1)),
  colour = "red"
# ) + geom_function(
  fun = function(x)
     dnorm(x, sd = sqrt(0.5)),
  colour = "blue"
# ) + geom_function(
   fun = function(x)
     dnorm(x, sd = sqrt(1)),
   colour = "green"
# ) + geom_function(
   fun = function(x)
     dnorm(x, sd = sqrt(2)),
  colour = "yellow"
# ) + geom_function(
# fun = function(x)
     dnorm(x, sd = sqrt(5)),
# colour = "orange"
```

```
# ) +
# ylab("Density")+
# scale_colour_manual(values=cols)

x1<-seq(-5,5,length.out=1001)
d<-sapply(c(0.1,0.5,1,2,5), function(x)dnorm(x1,sd = sqrt(x)))
d<-data.frame(x=rep(x1,times=5),density=c(d),sigma=rep(c(0.1,0.5,1,2,5),each=1001))
d$sigma<-as.factor(d$sigma)
p<-ggplot(d,aes(x=x,y=density,color=sigma))+
    geom_line()+
    xlab("")+
    ylab("Density")+
    scale_color_discrete(name = expression(sigma^2))
p</pre>
```



As  $\sigma^2$  decreases, the tail gets thiner, the overall spread decreases, and the peak at  $\mu = 0$  increases,