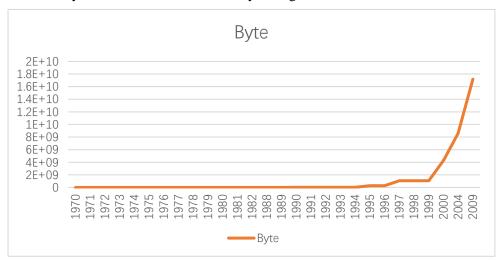
## HW Unit 1

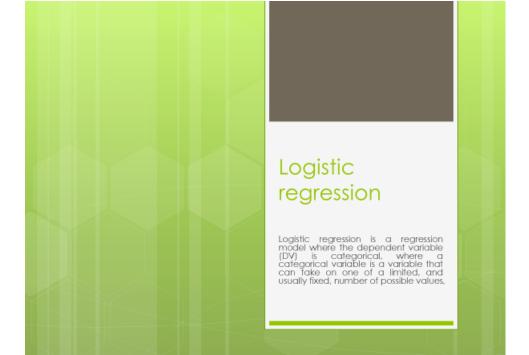
Q1. Calculate the increase of memory of PCs over the last 30 years and check whether the

year	Byte
1970	262144
1971	262144
1972	262144
1973	262144
1974	262144
1975	262144
1976	262144
1977	262144
1978	262144
1979	262144
1980	262144
1981	262144
1982	262144
1988	2097152
1989	2097152
1990	2097152
1991	16777216
1992	16777216
1993	16777216
1994	16777216
1995	16777216
1996	2.68E+08
1997	2.68E+08
1998	1.07E+09
1999	1.07E+09
2000	1.07E+09
2004	4.29E+09
2009	8.59E+09
2014	1.72E+10

FMRI analysis could have been done 20 years ago.



2. prepare 2-5 slides explaining logistic regression



# Logistic regression VS multiple linear regression

- There are many similarities between logistic regression and multiple linear regression, the biggest difference is that their dependent variables are different, and the others are almost the same.
- If the dependent variable is continuous, that is, multiple linear regression, if it is the two distribution, that is the logistic regression.

# **Application**

- To explore the risk factors of a disease and predict the probability of a disease according to the risk factors.
- If we have established the logistic regression model, we can predict the probability of a disease or a certain situation under different independent variables according to the model.
- 3. I have done that and my account is Bingren Sun026.

#### HW Unit 2

dpois(x,lambda)

```
1. \quad memory.df = read.csv("byte.csv",header = TRUE) \\ plot(memory.df\$Byte \sim memory.df\$year,type = "o",main = "The development of internal memory")
```

```
memory")

2. splines.reg.11 = smooth.spline(x = memory.df$year, y = memory.df$Byte, spar = 0.2)

splines.reg.12 = smooth.spline(x = memory.df$year, y = memory.df$Byte, spar = 1)

splines.reg.13 = smooth.spline(x = memory.df$year, y = memory.df$Byte, spar = 2)

lines(splines.reg.11, col = "green", lwd = 2)

lines(splines.reg.12, col = "pink", lwd = 2)

lines(splines.reg.13, col = "blue", lwd = 2)

3. lambda=4

x=6

dpois(x,lambda)

lambda=5

x=0
```

#### HW Unit 3

1. #install.packages("digest",repos='http://cran.us.r-project.org')

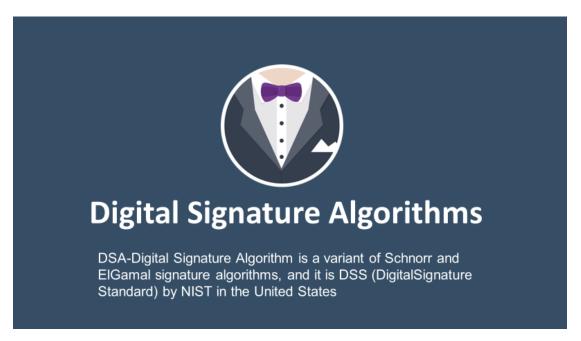
library(digest)

digest("I learn a lot from this class when I am proper listening to the professor", "sha256")

digest("I do not learn a lot from this class when I am absent and playing on my

Iphone", "sha256")

2. Make 3-5 slides (in PPTX) on the DSA (Digital Signature Algorithms)



## **Digital Signature**



Digital signature is a common physical signature similar to that written on paper, but it is implemented in the field of public key cryptography, and is used to identify digital information.



Digital signature is a digital string which can not be forged by others only. It is also an effective proof of the authenticity of information sent by the sender of the information.



A set of digital signatures usually defines two complementary operations, one for signature and the other for verification.



## **Digital Signature Algorithms**









- ✓ DSA is based on RSA algorithm.
- ✓ The private key X and the public key y are called a pair of keys (x, y). The private key can only be held solely by the signer himself, and the public key can be published publicly. Key pairs can be used continuously over a period of time.
- ✓ It follows the principle of "to signature with the private key and to verify with the public key"



#### 3. library(rjson)

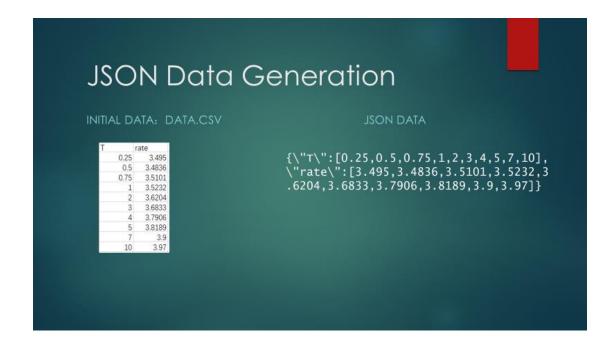
library(readr)

data <- read\_csv("~/R/Home-Work-for-BDIF/data.csv")

json\_ratedata <- toJSON(data,method = "C")</pre>

data

json\_ratedata



# JSON Data Generation R CODE: |ibrary(rjson) |ibrary(readr) | data <- read\_csv("~/R/Home-Work-for-BDIF/data.csv") | json\_ratedata <- toJSON(data,method = "C") | data | json\_ratedata

4. #install.packages("rjson", repos="http://cran.us.r-project.org")
 library(rjson)
 json\_file = "http://crix.hu-berlin.de/data/crix.json"
 json\_data = fromJSON(file=json\_file)
 crix <- Reduce(rbind,json\_data)
 crix\_data\_frame <- as.data.frame(crix)
 lst <- lapply(json\_data,function(x){
 df<-data.frame(date=x\$date,price=x\$price)</pre>

```
return(df)

})

crix_data_frame <- Reduce(rbind,lst)

plot(crix_data_frame$date,crix_data_frame$price)

#library(forecast)

#library(tseries)

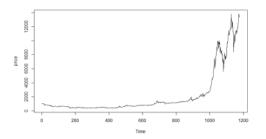
plot(crix_data_frame)

HW Unit 4
```



```
| #figure5 | hist(ret, col = "grey", breaks = 40, freq = FALSE) | lines(density(ret), lwd = 2) | par(mfrow = c(1, 2)) | # histogram of returns | hist(ret, col = "grey", breaks = 20, freq = FALSE, ylim = c(1, 2) | # histogram of returns | hist(ret, col = "grey", breaks = 20, freq = FALSE, ylim = c(0, 25), xlab = NA) | lines(density(ret), lwd = 2) | mu = mean(ret) | sigma = sd(ret) | x = seq(1,n|2,2) | x = seq(2,n|2,2) | x = seq(4, 4, length = 100) | curve(dnorm(x, mean = mean(ret), sd = sd(ret)), add = TRUE, col = "red", lwd = 2) | # qq-plot | qqnorm(ret) | qqline(ret, col = "blue", lwd = 3) | # figure5 | library(foreast) | library(f
```

# Q1-Figures



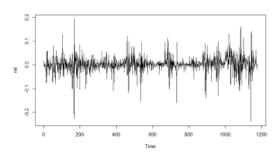
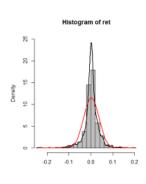


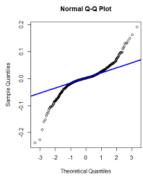
Figure 3

Figure 4



# Q1-Figures





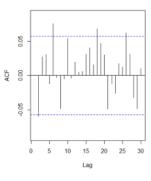


Figure 5

Figure 6



```
Q2
# install and load packages
libraries = c("zoo", "tseries")
lapply(libraries, function(x) if (!(x %in%
installed.packages())) {
  install.packages(x)
lapply(libraries, library, quietly = TRUE, character.only = TRUE)
```

# Q2

```
tsdiag(fit1)
Box.test(fit1$residuals, lag = 1)
                                                                                      fitr4 = aritina(ret_order = c(2, 1, 3))
tsdiag(fitr4)
```

```
# to conclude, 202 is better than 213
fit202 = arima(ret, order = c(2, 0, 2))
tsdiag(fit202)
tsdiag(fit4)
tsdiag(fit4)

AIC(fit202, k = log(length(ret)))
AIC(fit4, k = log(length(ret)))
AIC(fit74, k = log(length(ret)))
fit202$\frac{1}{3}$inse(crpre$\frac{1}{3}$pred, col = "red", lty = 3, lwd = 3]
lines(crpre$\frac{1}{3}$pred = 2. Crpre$\frac{1}{3}$pred = 3; lwd = 3]

# arima202 predict
fit202 = arima(ret, order = c(2, 0, 2))
crpre = predict(fit202, n.ahead = 30)

# arima202 predict
fit202 = arima(ret, order = c(2, 0, 2))
crpre = predict(fit202, n.ahead = 30)

# arima202 predict
fit202 = arima(ret, order = c(2, 0, 2))
crpre = predict(fit202, n.ahead = 30)

# arima202 predict
fit202 = arima(ret, order = c(2, 0, 2))
crpre = predict(fit202, n.ahead = 30)

# arima202 predict
fit202 = arima(ret, order = c(2, 0, 2))
crpre = predict(fit202, n.ahead = 30)

# arima202 predict
fit202 = arima(ret, order = c(2, 0, 2))
crpre = predict(fit202, n.ahead = 30)

# arima202 predict
fit202 = arima(ret, order = c(2, 0, 2))
crpre = predict(fit202, n.ahead = 30)

# arima202 predict
fit202 = arima(ret, order = c(2, 0, 2))
crpre = predict(fit202, n.ahead = 30)

# arima202 predict
fit202 = arima(ret, order = c(2, 0, 2))
crpre = predict(fit202, n.ahead = 30)

# arima202 predict
fit202 = arima(ret, order = c(2, 0, 2))
crpre = predict(fit202, n.ahead = 30)

# arima202 predict
fit202 = arima(ret, order = c(2, 0, 2))
crpre = predict(fit202, n.ahead = 30)

# arima202 predict
fit202 = arima(ret, order = c(2, 0, 2))
crpre = predict(fit202, n.ahead = 30)

# arima202 predict
fit202 = arima(ret, order = c(2, 0, 2))
crpre = predict(fit202, n.ahead = 30)

# arima202 predict
fit202 = arima(ret, order = c(2, 0, 2))
crpre = predict(fit202, n.ahead = 30)

# arima202 predict
fit202 = arima(ret, order = c(2, 0, 2))
crpre = predict(fit202, n.ahead = 30)

# arima202 predict
fit202 = arima(ret, order = c(2, 0, 2))
crpre = predict(fit202, n.ahead = 30)

# arima202 predict
fit202 = arima(ret, order = c(2, 0, 2))
crpre = predict(fit202, n.ahead = 30)

# arima202
```

## Q2-figure

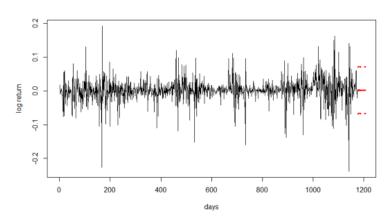


Figure 7



### Question1

```
library(rjson)
json_file = "http://crix.hu-berlin.de/data/crix.json"
json_data = fromJSON(file=json_file)
crix_data_frame <- as.data.frame(json_data)</pre>
w=crix_data_frame
dim(w)
n=dim(w)
a = seq(1,n[2],2)
b = seq(2,n[2],2)
data=t(w[1,a])
price=t(w[1,b])
#figure3
ts.plot(price)
#figure4
ret=diff(log(price))
ts.plot(ret)
#figure5
hist(ret, col = "grey", breaks = 40, freq = FALSE)
```

```
lines(density(ret), lwd = 2)
par(mfrow = c(1, 2))
# histogram of returns
hist(ret, col = "grey", breaks = 20, freq = FALSE, ylim = c(0, 25), xlab = NA)
lines(density(ret), lwd = 2)
mu = mean(ret)
sigma = sd(ret)
x = seq(-4, 4, length = 100)
curve(dnorm(x, mean = mean(ret), sd = sd(ret)), add = TRUE, col = "red",
     lwd = 2)
    # qq-plot
    qqnorm(ret)
    qqline(ret, col = "blue", lwd = 3)
    #figure6
    library(forecast)
    library(tseries)
    Acf(ret)
    Question2
    \#rm(list = ls(all = TRUE))
    #graphics.off()
    # install and load packages
    libraries = c("zoo", "tseries")
    lapply(libraries, function(x) if (!(x %in% installed.packages())) {
       install.packages(x)
    })
    lapply(libraries, library, quietly = TRUE, character.only = TRUE)
    # d order
    Box.test(ret, type = "Ljung-Box", lag = 20)
    # stationary test
```

```
adf.test(ret, alternative = "stationary")
kpss.test(ret, null = "Trend")
par(mfrow = c(1, 2))
# acf plot
autocorr = acf(ret, lag.max = 20, ylab = "Sample Autocorrelation", main = NA, lwd = 2,
ylim = c(-0.3, 1)
# LB test of linear dependence
print(cbind(autocorr$lag, autocorr$acf))
Box.test(ret, type = "Ljung-Box", lag = 1, fitdf = 0)
Box.test(autocorr$acf, type = "Ljung-Box")
# plot of pacf
autopcorr = pacf(ret, lag.max = 20, ylab = "Sample Partial Autocorrelation", main = NA,
ylim = c(-0.3, 0.3), lwd = 2)
print(cbind(autopcorr$lag, autopcorr$acf))
# arima model
par(mfrow = c(1, 1))
auto.arima(ret)
fit1 = arima(ret, order = c(1, 0, 1))
tsdiag(fit1)
Box.test(fit1\$residuals, lag = 1)
# aic
aic = matrix(NA, 6, 6)
for (p in 0:4) {
  for (q in 0:3) {
  a.p.q = arima(ret, order = c(p, 0, q))
  aic.p.q = a.p.q$aic
   aic[p + 1, q + 1] = aic.p.q
  }
}
aic
```

```
# bic
bic = matrix(NA, 6, 6)
for (p in 0:4) {
  for (q in 0:3) {
     b.p.q = arima(ret, order = c(p, 0, q))
     bic.p.q = AIC(b.p.q, k = log(length(ret)))
     bic[p + 1, q + 1] = bic.p.q
  }
}
bic
# select p and q order of ARIMA model
fit4 = arima(ret, order = c(2, 0, 3))
tsdiag(fit4)
Box.test(fit4\$residuals, lag = 1)
fitr4 = arima(ret, order = c(2, 1, 3))
tsdiag(fitr4)
Box.test(fitr4\$residuals, lag = 1)
# to conclude, 202 is better than 213
fit202 = arima(ret, order = c(2, 0, 2))
tsdiag(fit202)
tsdiag(fit4)
tsdiag(fitr4)
AIC(fit202, k = log(length(ret)))
AIC(fit4, k = log(length(ret)))
AIC(fitr4, k = log(length(ret)))
fit202$aic
fit4$aic
fitr4$aic
```



```
#install.packages("bitops")
#install.packages("stringr")
library(RCurl)
#install.packages("XML")
library(RCurl)
library(XML)
url1 =
"http://shakespeare.mit.edu/romeo_juliet/full.ht
ml"
url2 =
"http://shakespeare.mit.edu/julius_caesar/full.ht
ml"
url3 =
"http://shakespeare.mit.edu/hamlet/full.html"
html1 = readLines(url1, encoding = "UTF-8")
html2 = readLines(url2, encoding = "UTF-8")
html1 = htmlParse(html2, encoding = "UTF-8")
html2 = htmlParse(html2, encoding = "UTF-8")
html3 = htmlParse(html2, encoding = "UTF-8")
html3 = htmlParse(html3, encoding = "UTF-8")
html3 = htmlParse(html3, encoding = "UTF-8")
html3 = htmlParse(html3, encoding = "UTF-8")
}

#install.packages("bitops")
#install.packages("bitops")
#install.packages("bitops")
#install.packages("bitops")
#install.packages("bitops")
| library(bitops)
| library(stringr)
| abs1 = lapply(url1, FUN = function(x) htmlParse(x, encoding = "Latin-1"))
| abs2 = lapply(url2, FUN = function(x) htmlParse(x, encoding = "Latin-1"))
| abs3 = lapply(url3, FUN = function(x) htmlParse(x, encoding = "Latin-1"))
| clean_txt = function(x) htmlParse(x, encoding = "Latin-1")
| clean_txt = function(x) htmlParse(x, encoding = "Latin-1")
| clean_txt = function(x) htmlParse(x, encoding = "UTF-8")
| clean_t
```

```
Q1
```

```
cleantxt1 = lapply(abs1,clean_txt)
cleantxt2 = lapply(abs2,clean_txt)
cleantxt3 = lapply(abs3,clean_txt)
vec_abs1 = unlist(cleantxt1)
vec_abs2 = unlist(cleantxt2)
vec_abs3 = unlist(cleantxt3)

###Text Mining
install.packages("tm")
install.packages("SnowballC")
library(tm)
library(SnowballC)
abs1 = Corpus(VectorSource(vec_abs1))
abs2 = Corpus(VectorSource(vec_abs2))
abs3 = Corpus(VectorSource(vec_abs3))
```

```
abs_dtm1 = DocumentTermMatrix(abs1, control = list(
stemming = TRUE, stopwords = TRUE, minWordLength
= 3,
    removeNumbers = TRUE, removePunctuation = TRUE))
abs_dtm2 = DocumentTermMatrix(abs2, control = list(
    stemming = TRUE, stopwords = TRUE, minWordLength
= 3,
    removeNumbers = TRUE, removePunctuation = TRUE))
abs_dtm3 = DocumentTermMatrix(abs3, control = list(
    stemming = TRUE, stopwords = TRUE, minWordLength
= 3,
    removeNumbers = TRUE, removePunctuation = TRUE))
##WordCloud
instal.packages("ggplot2")
install.packages("ggplot2")
install.packages("wordcloud")
library(wordcloud)
freq1 = colSums(as.matrix(abs_dtm1))
freq2 = colSums(as.matrix(abs_dtm2))
freq3 = colSums(as.matrix(abs_dtm3))
wf1 = data.frame(word=names(freq1), freq=freq1)
wf2 = data.frame(word=names(freq2), freq=freq2)
wf3 = data.frame(word=names(freq3), freq=freq3)
```

```
#Romeo and Juliet
plot1 = ggplot(subset(wf1, freq>15), aes(word, freq1))
plot1 = plot1 + geom_bar(stat="identity")
plot1 = plot1 + geom_bar(stat="identity")
plot1 = plot1 + theme(axis.text.x=element_text(angle=45, hjust=1))
plot1
freq1 = colSums(as.matrix(abs_dtm1))
dark2_1 = brewer.pal(6, "Dark2")
wordcloud(names(freq2), freq2, max.words=100, rot.per=0.2, colors=dark2_1)
#Hamlet
plot3 = ggplot(subset(wf3, freq>15), aes(word, freq3))
plot3 = plot3 + geom_bar(stat="identity")
plot3
freq1 = colSums(as.matrix(abs_dtm2))
dark2_1 = brewer.pal(6, "Dark2")
wordcloud(names(freq1), freq1, max.words=100, rot.per=0.2, colors=dark2_1)
#Julius Caeser
plot2 = ggplot(subset(wf2, freq>15), aes(word, freq2))
plot2 = plot2 + geom_bar(stat="identity")
plot2 = plot2 + geom_bar(stat="identity")
plot2 = plot2 + geom_bar(stat="identity")
plot2
```

## Q1-Figures



g quidensem qui de l'entre l'e

Figure 3

Figure 1 Figure 2



# Q2

# Q2-Figures

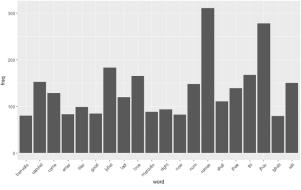


Figure 4

# Q2-figure

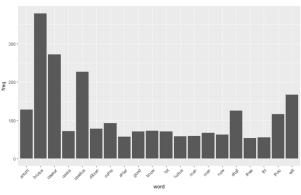


Figure 5

# Q2-figure

