# Computational Statistics

Lab 5

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### Question 1

0 -

#### 1.1

```
## Warning: package 'ggplot2' was built under R version 3.3.2
lottery <- read.csv2(".../data/lottery.csv")
q11 <- ggplot(lottery, aes(x = Day_of_year, y = Draft_No)) + geom_point()
plot(q11)</pre>
```

data <- data.frame(x=lottery\$Day\_of\_year, y=lottery\$Draft\_No)</pre>

100

The data looks random although there might be some sort of skewness in the right side of the graph were there are a lacking some observations and therefore having a lower probability of getting selected.

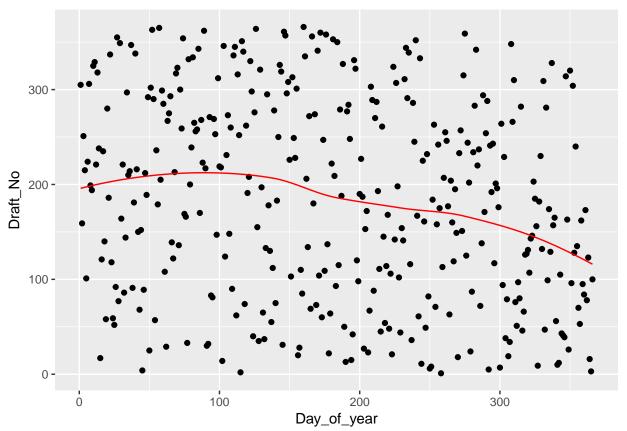
200

Day\_of\_year

300

```
loessfit <- loess(y ~ x, data=data)
data$pred <- predict(loessfit, data$x)

q12 <- q11 + geom_path(data = data, aes(x=x, y=pred), col = "red")
plot(q12)</pre>
```



The fit (line) doesn't seem straight and seems to have a decreasing trend which would support previous statements of people born on a days later on in a year has a lower probability of beeing selected.

```
teststat <- function(model) {
  function(data) {
    xa <- data$x[which.min(data$y)]
    xb <- data$x[which.max(data$y)]

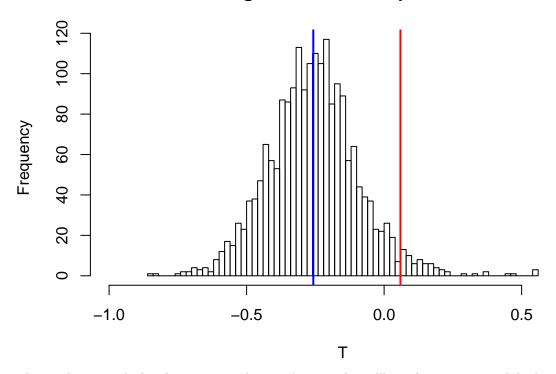
  fit <- model(y ~ x, data)

    ya <- predict(fit, xa)
    yb <- predict(fit, xb)

    (yb - ya) / (xb - xa)
}</pre>
```

```
}
teststat_boot <- function(data, idx, stat) {</pre>
    data <- data[idx,]</pre>
    stat(data)
}
library(boot)
B <- 2000
set.seed(123456)
npboot <- boot(data=data, statistic=teststat_boot, R=B, stat=teststat(model=loess))</pre>
pvalue <- sum(npboot$t > 0) / B
pvalue
## [1] 0.0595
hist(npboot\$t, xlim = c(-1,0.7), breaks = 50,
     main = "Histogram for bootstrap t-values", xlab ="T")
abline(v=pvalue, col = "red", lwd = 2)
abline(v=mean(npboot$t), col="blue", lwd=2)
```

### Histogram for bootstrap t-values



The p-value was calculated to 0.0595 and we can't reject the nullhypothesis at a 0.05 alpha-level. We conclude Ho and that the lottery is random.

```
teststat_permutation<- function(data, B, stat) {
    n <- nrow(data)
    statistics <- rep(0, B)
    newdata <- data.frame(x=data$x, y=sample(data$y, n))

for (b in 1:B) {
    statistics[b] <- stat(newdata)
    newdata$y <- sample(data$y, n)
  }

sum(abs(statistics) >= abs(stat(data))) / B
}

set.seed(123456)
pvalue2 <- teststat_permutation(data, B, teststat(loess))
pvalue2</pre>
```

#### ## [1] 0.0925

We conclude Ho in this test as well, the lottery is random.

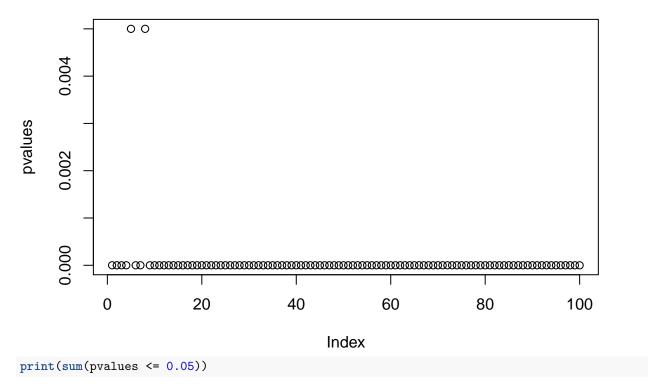
```
genranddata <- function(x, alpha) {
    data.frame(x=x, y=pmax(0, pmin(alpha * x + rnorm(length(x), mean=183, sd=10), 366)))
}

alphas <- seq(0.1, 10, by=0.1)
pvalues <- rep(0, length(alphas))

set.seed(123456)

for (i in 1:length(alphas)) {
    newdata <- genranddata(data$x, alphas[i])
    pvalues[i] <- teststat_permutation(newdata, 200, teststat(loess))
}

plot(pvalues)</pre>
```



### ## [1] 100

Since we reject Ho in every test the power will be 1 becaus we can't have any Type-2 errors when we reject everything. This isn't strange since the test-data is obviously non-random and gives us an indication that the test-statistic is a good one.

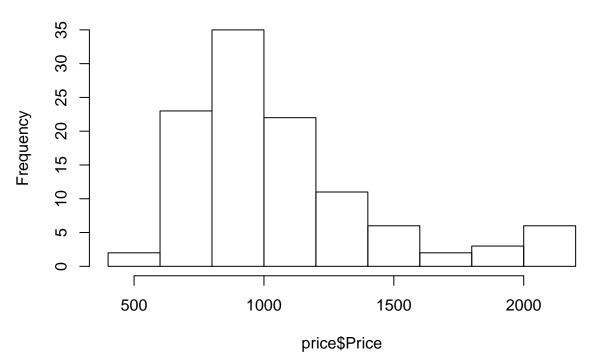
### Question 2

### 2.1

```
price <- read.csv("../data/prices1.csv", sep=";")
mean(price$Price)

## [1] 1080.473
hist(price$Price)</pre>
```

## Histogram of price\$Price



Looks like a Gamma distribution.

### 2.2

## [1] 1080.828

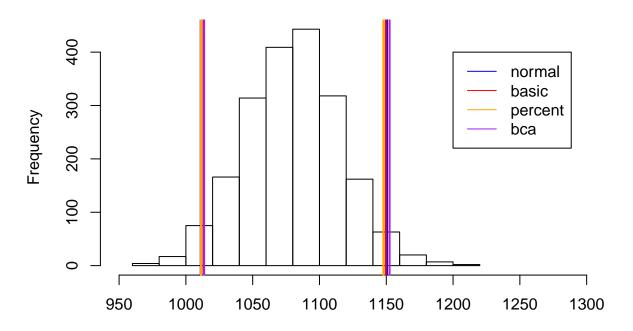
```
bootmean <- function(data,ind){
  data <- data[ind]
  mean(data)
}

B <- 2000
estmean <- boot(data = price$Price,bootmean, R = B)

2*estmean$t0 - mean(estmean$t)</pre>
```

```
mean(estmean$t- estmean$t0)
## [1] -0.355
sum((estmean$t-mean(estmean$t))^2)/(B-1)
## [1] 1264.299
cibo <- boot.ci(estmean)</pre>
## Warning in boot.ci(estmean): bootstrap variances needed for studentized
## intervals
print(cibo)
## BOOTSTRAP CONFIDENCE INTERVAL CALCULATIONS
## Based on 2000 bootstrap replicates
##
## CALL :
## boot.ci(boot.out = estmean)
## Intervals :
## Level
             Normal
       (1011, 1151) (1013, 1150)
## 95%
             Percentile
## Level
                                   BCa
         (1011, 1148)
                        (1014, 1152)
## 95%
## Calculations and Intervals on Original Scale
hist(estmean$t, xlim = c(950,1300), main = "Bootstrap estimation of mean", xlab = "")
abline(v= cibo$normal[2:3], col = "blue", lwd = 2)
abline(v= cibo$basic[4:5], col = "red", lwd = 2)
abline(v= cibo$percent[4:5], col = "orange", lwd = 2)
abline(v= cibo$bca[4:5], col = "purple", lwd = 2)
legend(y=400,x =1200,c("normal","basic","percent","bca"),lty = c(1,1),
       col = c("blue","red","orange","purple"))
```

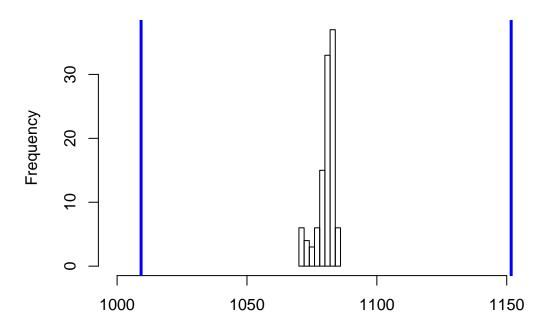
### **Bootstrap estimation of mean**



```
jackknife <- function(data,B,tstat){</pre>
  stopifnot( B <= length(data))</pre>
  est \leftarrow rep(1,times = B)
  for(i in 1:B){
    est[i] <- tstat(data[-i])</pre>
  }
  return(est)
jackest<-jackknife(price$Price, B = length(price$Price), tstat = mean)</pre>
mean(jackest)
## [1] 1080.473
n <- length(price$Price)</pre>
tstar<- n*mean(price$Price)-(n-1)*jackest</pre>
JT<-mean(tstar)</pre>
jackvar <-(1/(n*(n-1)))*sum((tstar- JT)^2)
lowerci <- JT - 1.96*sqrt(jackvar)</pre>
upperci <- JT + 1.96*sqrt(jackvar)</pre>
\# mean(tstar) + qt(0.975, length(price\$Price)-1)*sqrt(var(tstar)/length(price\$Price))
```

```
# mean(tstar) - qt(0.975,length(price$Price)-1)*sqrt(var(tstar)/length(price$Price))
hist(jackest,breaks = 10, xlim = c(1000,1180), xlab = "" ,main = "Jackknife estimate")
abline(v=lowerci, col ="blue", lwd = 3)
abline(v=upperci, col ="blue", lwd = 3)
```

### Jackknife estimate



```
library(knitr)
```

```
## Warning: package 'knitr' was built under R version 3.3.2

cimatrix<-rbind(
    c("Normal",cibo$normal[2],mean(estmean$t),cibo$normal[2]),
    c("Percent",cibo$percent[4],mean(estmean$t),cibo$percent[5]),
    c("Basic",cibo$basic[4],mean(estmean$t),cibo$basic[5]),
    c("Bca",cibo$bca[4],mean(estmean$t),cibo$bca[5]),
    c("Jackknife",lowerci,mean(jackest),upperci)
)

colnames(cimatrix) <- c("Name","Lower","Mean","Upper")

kable(cimatrix, digits = 3)</pre>
```

Name	Lower	Mean	Upper
Normal	1011.13731332721	1080.11772727273	1011.13731332721
Percent	1011.30801927524	1080.11772727273	1147.81474498594
Basic	1013.13070955952	1080.11772727273	1149.63743527021
Bca	1013.62920802339	1080.11772727273	1152.45553911654
Jackknife	1009.23782103407	1080.47272727273	1151.70763351138