# Statistics CH4

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# 4.2.0.1 Excercise

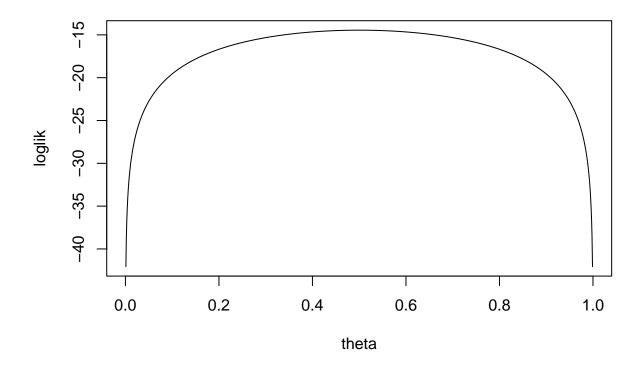
a.

```
X = c(3, 0, 2, 1, 3, 2, 1, 0, 2, 1)
theta_hat_mom = (7/3 - mean(X))/2
theta_hat_mom
```

## [1] 0.4166667

b.

```
theta = seq(0, 1, 0.001)
loglik = 2*log(2/3*theta)+3*log(1/3*theta)+3*log(2/3*(1-theta))+2*log(1/3*(1-theta))
plot(theta, loglik, type='l')
```



## c.

```
theta_hat_mle = theta[which.max(loglik)]
theta_hat_mle
```

## [1] 0.5

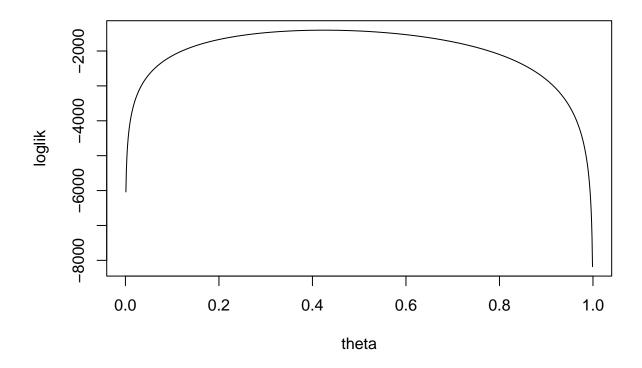
```
mean_hat_mle = 7/3 - 2*theta_hat_mle
mean_hat_mle
```

## [1] 1.333333

# **4.2.0.2** Exercise

a.

```
theta = seq(0, 1, 0.001)
loglik = 1184*log(1-theta)+874*log(theta)
plot(theta, loglik, type='l')
```



## b.

```
theta_hat_mle = theta[which.max(loglik)]
theta_hat_mle
```

## [1] 0.425

# **4.2.0.4** Exercise

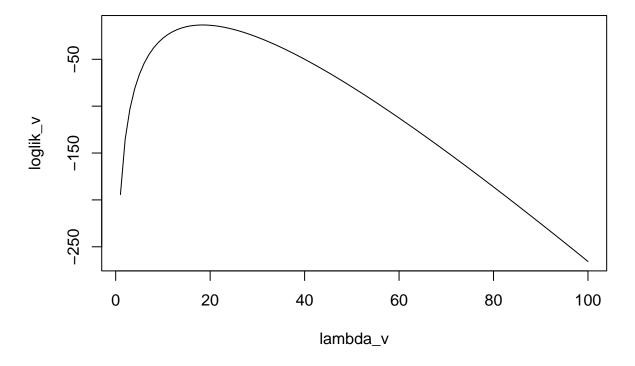
b.

```
X = c(17, 25, 17, 18, 15)
e_hat_mom = mean(X)
e_hat_mom
```

## [1] 18.4

 $\mathbf{c}.$ 

```
loglik = function(lambda) {
    X = c(17, 25, 17, 18, 15)
    result = sum(sapply(X, dpois, lambda=lambda, log = TRUE))
    return(result)
}
lambda_v = seq(1, 100, 1)
loglik_v = sapply(lambda_v, loglik)
plot(lambda_v, loglik_v, type='l')
```



## d.

```
lambda_hat_mle = lambda_v[which.max(loglik_v)]
lambda_hat_mle
```

## [1] 18

e.

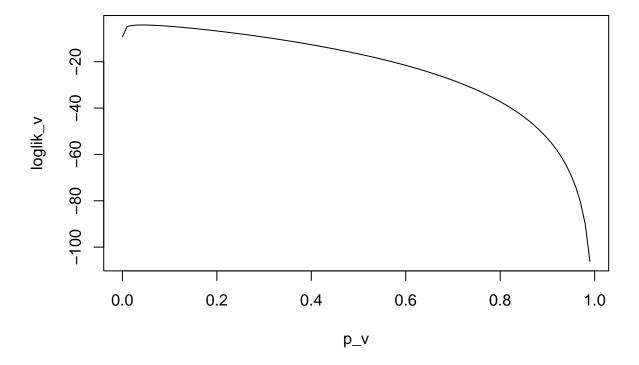
```
opt = optimize(loglik, c(1, 100), maximum = TRUE)
lambda_hat_mle2 = opt$maximum
lambda_hat_mle2
```

## [1] 18.40002

### **4.2.0.6** Exercise

c.

```
loglik = function(p) {
    X = 23
    result = dgeom(X, prob = p, log = TRUE)
    return(result)
}
p_v = seq(0.0001,1,0.01)
loglik_v = loglik(p_v)
plot(p_v, loglik_v, type='l')
```



## d.

```
p_hat_mle = p_v[which.max(loglik_v)]
p_hat_mle
```

## [1] 0.0401

e.

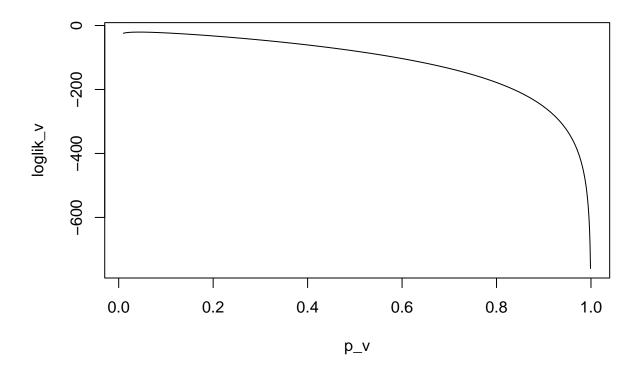
```
opt = optimize(loglik, c(0,1), maximum=TRUE)
p_hat_mle1 = opt$maximum
p_hat_mle1
```

## [1] 0.04166196

### **4.2.0.7** Exercise

a.

```
loglik = function(p) {
    X = 20:24
    result = sum(dgeom(X, prob = p, log = TRUE))
    return(result)
}
p_v = seq(0.01,1,0.001)
loglik_v = sapply(p_v, loglik)
plot(p_v, loglik_v, type='l')
```



## b.

```
opt = optimize(loglik, c(0,1), maximum=TRUE)
p_hat_mle = opt$maximum
p_hat_mle
```

## [1] 0.04347581

#### 4.2.0.8 Exercise

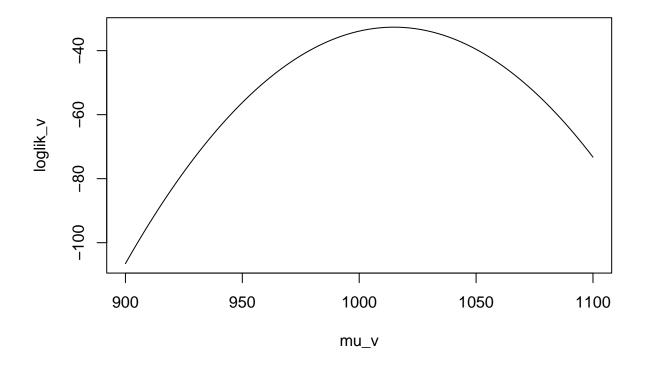
b.

```
X = c(983, 1041, 1026, 1025, 1033, 968, 1028)
mu_hat_mom = mean(X)
mu_hat_mom
```

## [1] 1014.857

c.

```
loglik = function(mu) {
   X = c(983, 1041, 1026, 1025, 1033, 968, 1028)
   result = sum(dnorm(X, mean = mu, sd = 25, log = TRUE))
   return(result)
}
mu_v = seq(900, 1100, 1)
loglik_v = sapply(mu_v, loglik)
plot(mu_v, loglik_v, type = 'l')
```



```
mu_hat_mle = mu_v[which.max(loglik_v)]
mu_hat_mle
```

## [1] 1015

d.

```
opt = optimize(loglik, c(900, 1100), maximum = TRUE)
mu_hat_mle2 = opt$maximum
mu_hat_mle2
```

## [1] 1014.857

### **4.2.0.9** Exercise

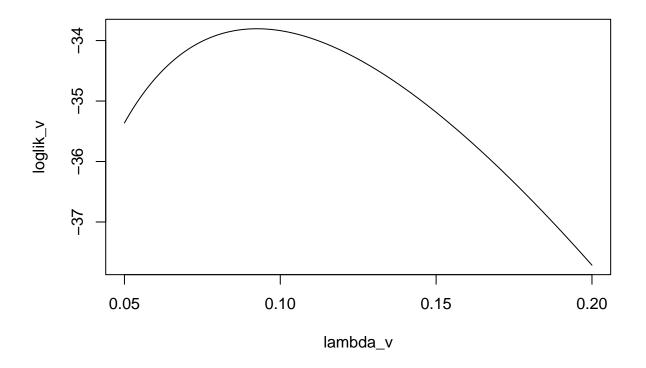
b.

```
X = c(8.5, 2.3, 11.0, 22.5, 13.6, 5.8, 27.3, 13.1, 0.9, 3.1)
mean(X)
```

## [1] 10.81

c.

```
loglik = function(lambda){
    X = c(8.5, 2.3, 11.0, 22.5, 13.6, 5.8, 27.3, 13.1, 0.9, 3.1)
    result = sum(dexp(X, rate = lambda, log = TRUE))
    return(result)
}
expecta = seq(5, 20, 0.01)
lambda_v = 1/expecta
loglik_v = sapply(lambda_v, loglik)
plot(lambda_v, loglik_v, type = 'l')
```



```
lambda_hat_mle = lambda_v[which.max(loglik_v)]
1/lambda_hat_mle
```

## [1] 10.81

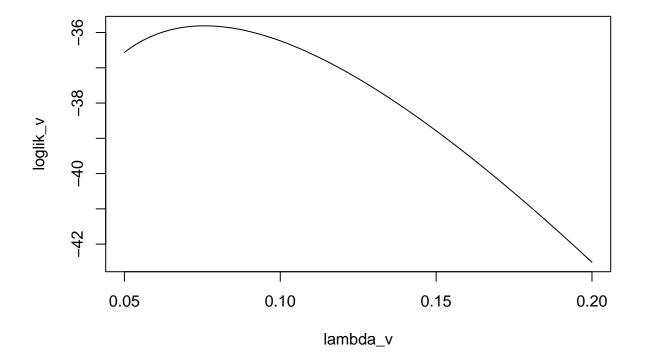
d.

```
opt = optimize(loglik, c(0, 1), maximum = TRUE)
lambda_hat_mle1 = opt$maximum
1/lambda_hat_mle1
```

#### 4.2.0.10 Exercise

a.

```
loglik = function(lambda){
    X = c(8.5, 2.3, 11.0, 22.5, 13.6, 5.8, 27.3, 13.1, 0.9, 3.1)
    result = sum(dexp(X, rate = lambda, log = TRUE)) + pexp(24, rate = lambda, lower.tail = FALSE, log.p = return(result)
}
expecta = seq(5, 20, 0.01)
lambda_v = 1/expecta
loglik_v = sapply(lambda_v, loglik)
plot(lambda_v, loglik_v, type = 'l')
```



```
lambda_hat_mle = lambda_v[which.max(loglik_v)]
1/lambda_hat_mle
```

## [1] 13.21

b.

```
opt = optimize(loglik, c(0, 1), maximum = TRUE)
lambda_hat_mle1 = opt$maximum
1/lambda_hat_mle1
```

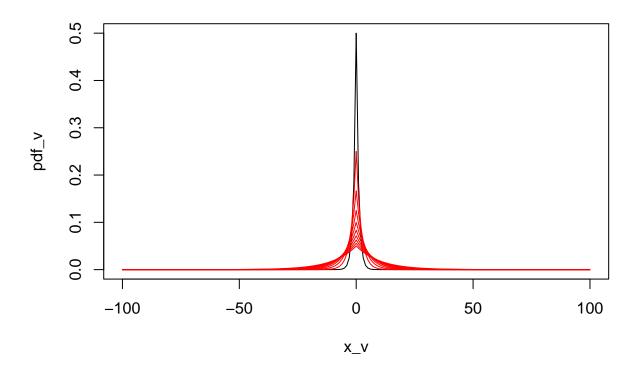
## [1] 13.21264

### **4.2.0.11** Exercise

a.

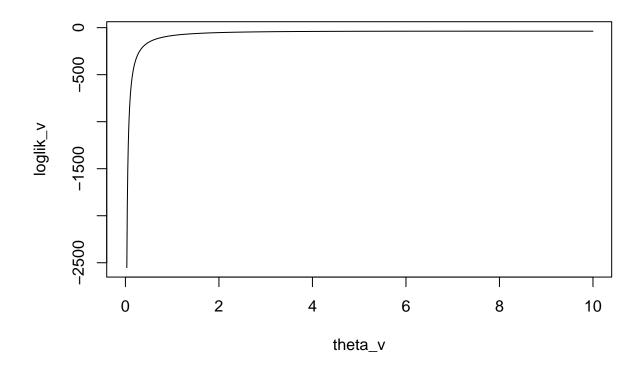
```
pdf = function(x, theta){
    result = 1/(2*theta)*exp(-1*abs(x)/theta)
    return(result)
}

x_v = seq(-100, 100, 1)
pdf_v = pdf(x_v, 1)
plot(x_v, pdf_v, type = 'l')
for (i in 2:10) {
    pdf_v = pdf(x_v, i)
    lines(x_v, pdf_v, type = 'l', col = 'red')
}
```



##b.

```
loglik = function(theta){
    X = c(4.77,5.52,0.38,1.13,21.86,13.79,12.61,1.01,13.96,2.37)
    result = sum(log(pdf(X, theta)))
    return(result)
}
theta_v = seq(0, 10, 0.01)
loglik_v = sapply(theta_v, loglik)
plot(theta_v, loglik_v, type = 'l')
```



```
theta_hat_mle = theta_v[which.max(loglik_v)]
theta_hat_mle
```

## [1] 7.74

c.

```
opt = optimize(loglik, c(0, 10), maximum = TRUE)
theta_hat_mle1 = opt$maximum
theta_hat_mle1
```

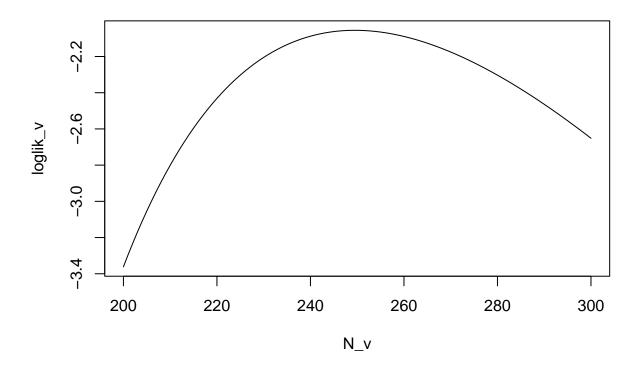
## [1] 7.739999

# **4.2.0.12** Exercise

b.

```
loglik = function(N){
    X = 20
    result = dhyper(X, m = 100, n = N-100, k = 50, log = TRUE)
```

```
return(result)
}
N_v = seq(200, 300, 1)
loglik_v = loglik(N_v)
plot(N_v, loglik_v, type = 'l')
```



```
N_hat_mle = N_v[which.max(loglik_v)]
N_hat_mle
```

## [1] 249

c.

```
opt = optimize(loglik, c(200, 300), maximum = TRUE)

## Warning in dhyper(X, m = 100, n = N - 100, k = 50, log = TRUE): NaNs produced

## Warning in optimize(loglik, c(200, 300), maximum = TRUE): NA/Inf replaced by

## maximum positive value

## Warning in dhyper(X, m = 100, n = N - 100, k = 50, log = TRUE): NaNs produced
```

- ## Warning in optimize(loglik, c(200, 300), maximum = TRUE): NA/Inf replaced by
  ## maximum positive value
- ## Warning in dhyper(X, m = 100, n = N 100, k = 50, log = TRUE): NaNs produced
- ## Warning in optimize(loglik, c(200, 300), maximum = TRUE): NA/Inf replaced by
  ## maximum positive value
- ## Warning in dhyper(X, m = 100, n = N 100, k = 50, log = TRUE): NaNs produced
- ## Warning in optimize(loglik, c(200, 300), maximum = TRUE): NA/Inf replaced by ## maximum positive value
- ## Warning in dhyper(X, m = 100, n = N 100, k = 50, log = TRUE): NaNs produced
- ## Warning in optimize(loglik, c(200, 300), maximum = TRUE): NA/Inf replaced by ## maximum positive value
- ## Warning in dhyper(X, m = 100, n = N 100, k = 50, log = TRUE): NaNs produced
- ## Warning in optimize(loglik, c(200, 300), maximum = TRUE): NA/Inf replaced by
  ## maximum positive value
- ## Warning in dhyper(X, m = 100, n = N 100, k = 50, log = TRUE): NaNs produced
- ## Warning in optimize(loglik, c(200, 300), maximum = TRUE): NA/Inf replaced by ## maximum positive value
- ## Warning in dhyper(X, m = 100, n = N 100, k = 50, log = TRUE): NaNs produced
- ## Warning in optimize(loglik, c(200, 300), maximum = TRUE): NA/Inf replaced by ## maximum positive value
- ## Warning in dhyper(X, m = 100, n = N 100, k = 50, log = TRUE): NaNs produced
- ## Warning in optimize(loglik, c(200, 300), maximum = TRUE): NA/Inf replaced by ## maximum positive value
- ## Warning in dhyper(X, m = 100, n = N 100, k = 50, log = TRUE): NaNs produced
- ## Warning in optimize(loglik, c(200, 300), maximum = TRUE): NA/Inf replaced by ## maximum positive value
- ## Warning in dhyper(X, m = 100, n = N 100, k = 50, log = TRUE): NaNs produced
- ## Warning in optimize(loglik, c(200, 300), maximum = TRUE): NA/Inf replaced by ## maximum positive value
- ## Warning in dhyper(X, m = 100, n = N 100, k = 50, log = TRUE): NaNs produced

- ## Warning in optimize(loglik, c(200, 300), maximum = TRUE): NA/Inf replaced by
  ## maximum positive value
- ## Warning in dhyper(X, m = 100, n = N 100, k = 50, log = TRUE): NaNs produced
- ## Warning in optimize(loglik, c(200, 300), maximum = TRUE): NA/Inf replaced by
  ## maximum positive value
- ## Warning in dhyper(X, m = 100, n = N 100, k = 50, log = TRUE): NaNs produced
- ## Warning in optimize(loglik, c(200, 300), maximum = TRUE): NA/Inf replaced by ## maximum positive value
- ## Warning in dhyper(X, m = 100, n = N 100, k = 50, log = TRUE): NaNs produced
- ## Warning in optimize(loglik, c(200, 300), maximum = TRUE): NA/Inf replaced by ## maximum positive value
- ## Warning in dhyper(X, m = 100, n = N 100, k = 50, log = TRUE): NaNs produced
- ## Warning in optimize(loglik, c(200, 300), maximum = TRUE): NA/Inf replaced by
  ## maximum positive value
- ## Warning in dhyper(X, m = 100, n = N 100, k = 50, log = TRUE): NaNs produced
- ## Warning in optimize(loglik, c(200, 300), maximum = TRUE): NA/Inf replaced by ## maximum positive value
- ## Warning in dhyper(X, m = 100, n = N 100, k = 50, log = TRUE): NaNs produced
- ## Warning in optimize(loglik, c(200, 300), maximum = TRUE): NA/Inf replaced by ## maximum positive value
- ## Warning in dhyper(X, m = 100, n = N 100, k = 50, log = TRUE): NaNs produced
- ## Warning in optimize(loglik, c(200, 300), maximum = TRUE): NA/Inf replaced by ## maximum positive value
- ## Warning in dhyper(X, m = 100, n = N 100, k = 50, log = TRUE): NaNs produced
- ## Warning in optimize(loglik, c(200, 300), maximum = TRUE): NA/Inf replaced by ## maximum positive value
- ## Warning in dhyper(X, m = 100, n = N 100, k = 50, log = TRUE): NaNs produced
- ## Warning in optimize(loglik, c(200, 300), maximum = TRUE): NA/Inf replaced by ## maximum positive value
- ## Warning in dhyper(X, m = 100, n = N 100, k = 50, log = TRUE): NaNs produced

```
## Warning in optimize(loglik, c(200, 300), maximum = TRUE): NA/Inf replaced by
## maximum positive value
## Warning in dhyper(X, m = 100, n = N - 100, k = 50, log = TRUE): NaNs produced
## Warning in optimize(loglik, c(200, 300), maximum = TRUE): NA/Inf replaced by
## maximum positive value
## Warning in dhyper(X, m = 100, n = N - 100, k = 50, log = TRUE): NaNs produced
## Warning in optimize(loglik, c(200, 300), maximum = TRUE): NA/Inf replaced by
## maximum positive value
## Warning in dhyper(X, m = 100, n = N - 100, k = 50, log = TRUE): NaNs produced
## Warning in optimize(loglik, c(200, 300), maximum = TRUE): NA/Inf replaced by
## maximum positive value
## Warning in dhyper(X, m = 100, n = N - 100, k = 50, log = TRUE): NaNs produced
## Warning in optimize(loglik, c(200, 300), maximum = TRUE): NA/Inf replaced by
## maximum positive value
## Warning in dhyper(X, m = 100, n = N - 100, k = 50, log = TRUE): NaNs produced
## Warning in optimize(loglik, c(200, 300), maximum = TRUE): NA/Inf replaced by
## maximum positive value
## Warning in dhyper(X, m = 100, n = N - 100, k = 50, log = TRUE): NaNs produced
## Warning in optimize(loglik, c(200, 300), maximum = TRUE): NA/Inf replaced by
## maximum positive value
## Warning in dhyper(X, m = 100, n = N - 100, k = 50, log = TRUE): NaNs produced
## Warning in optimize(loglik, c(200, 300), maximum = TRUE): NA/Inf replaced by
## maximum positive value
## Warning in dhyper(X, m = 100, n = N - 100, k = 50, log = TRUE): NaNs produced
N_hat_mle1 = opt$maximum
N_hat_mle1
```

## [1] 299.9999

#### 4.2.0.14 Exercise

```
x=c(983,1041,1026,1025,1033,968,1028)
loglik=-Inf
for (m in seq(900,1100,0.1)){
   for (s in seq(10,40,0.1)){
      tmp=sum(log(dnorm(x,m,s)))  # compute the loglikelihood by summing
      if (tmp>loglik){
        loglik=tmp
            mu=m
            sigma=s
      }
   }
}
mu
## [1] 1014.9
```

#### 4.2.0.15 Exercise

## [1] 25.7

```
x=c(983,1041,1026,1025,1033,968,1028)
11 = function(theta,x) {
                           # We define an R function that computes the log likelihood
  mu=theta[1]
   sigma=theta[2]
   -sum(log(dnorm(x,mu,sigma))) # compute loglikelihood by summing
                                 # add a minus sign, because optim *minimizes*.
optim(c(1000,25),11,x=x,method="BFGS")
## $par
## [1] 1014.86313 25.69854
## $value
## [1] 32.65757
##
## $counts
## function gradient
##
         21
                 18
## $convergence
## [1] 0
##
## $message
## NULL
```