

Inferential Statistics Assignment

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Introduction

I went through the first and the third assignment, here reported in this order. This report has been created using r markdown and latex.

1. First Assignment

First of all I've defined some constants and a function to make my code clearer and understandable. I've also included some libraries that I've used in this part of the assignment.

```
library(plotrix)
library(rmarkdown)
set.seed(29)

#tau function
tau <- function(theta){
  out <- theta/(1-theta)
  out2 <- log(out)
  return(out2)
}

theta0 = 2/3 ##Neg bin success probability
r = 10 ##Neg bin size param
n <- c(10,50) ## Sample sizes
alpha = 0.05 ## Wald c.i. .95
```

In this segment I've created a 2*N matrix with N = 10⁵ used to collect some results in order to simulate the distribution of $\hat{\tau}$. Every element is obtained evaluating $\hat{\theta}$ with the function *tau*. $\hat{\theta}$ is defined as

$$\hat{\theta} = \frac{r * n}{n * (r + \bar{y})}$$

where r is the *index* parameter of the negative binomial, n is the size of the r. ve. and \bar{y} is the sample mean. Those results were obtained finding the maximum of the likelihood function $l(\hat{\theta})$.

```

N = 1e5
sim.N.tau <-matrix(NA, nrow = N, ncol = 2)
for(i in 1:N){
  xnegbin10 <- rnbinom(n=10,size=10,prob=theta0)
  xnegbin50 <- rnbinom(n=50,size=10,prob=theta0)
  sim.N.tau[i,1] <- tau(r*n[1]/(r*n[1]+sum(xnegbin10)))
  sim.N.tau[i,2] <- tau(r*n[2]/(r*n[2]+sum(xnegbin50)))
}

```

In the next cell I've managed to calculate what would have been the true value of mean and variance of the distribution of $\hat{\tau}$. In order to do so I've calculated the variance of $\hat{\theta}$ with Cramér-Rao's theorem, knowing that the Maximum Likelihood Estimator is asymptotically efficient. In formulas:

$$Var(\hat{\theta}) = I_n(\theta)^{-1}$$

where $I_n(\theta)$ is the Fisher information

$$I_n(\theta) = n * I(\theta)$$

knowing that our r. ve. is built from Y_1, \dots, Y_n i.i.d. variables and

$$I(\theta) = \frac{\theta^2 * (1 - \theta)}{r * n}$$

which is obtained by the definition of Fisher information. After that I've used the Delta Method to obtain the variance and the mean of the gaussian curve that represent our ideal density (and distribution), and I've got the following results:

$\hat{\tau}$'s variance:

$$var(\hat{\tau}) = \left(\frac{d\tau(\theta)}{d\theta}\right)^2 * Var(\hat{\theta})$$

$\hat{\tau}$'s mean:

$$E[\hat{\tau}] = \tau(\theta)$$

Evaluating variance and mean in θ_0 bring us to the real p.d.f. of $\hat{\tau}$

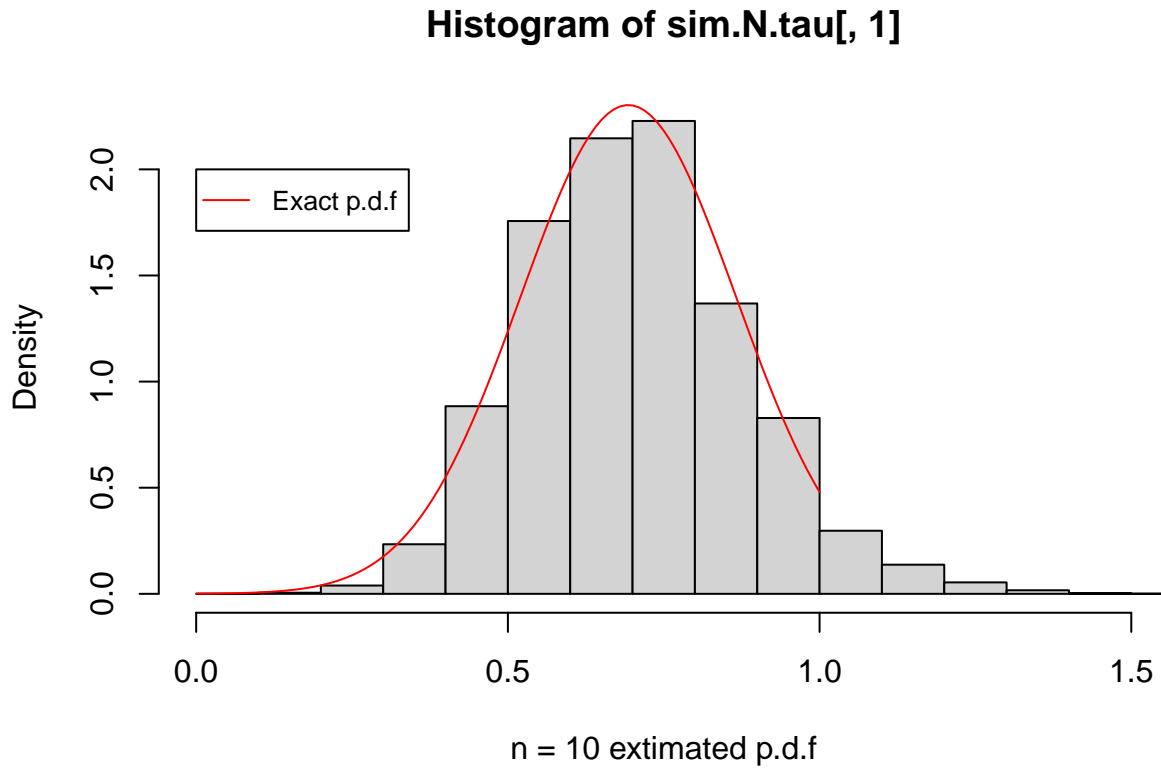
```

true_mean = (tau(theta0))
true_var10_o <- (theta0^2*(1-theta0))/(r*n[1])
true_var10 <- true_var10_o*(1/(theta0*(1-theta0)))^2
true_var50_o <- (theta0^2*(1-theta0))/(r*n[2])
true_var50 <- true_var50_o*(1/(theta0*(1-theta0)))^2

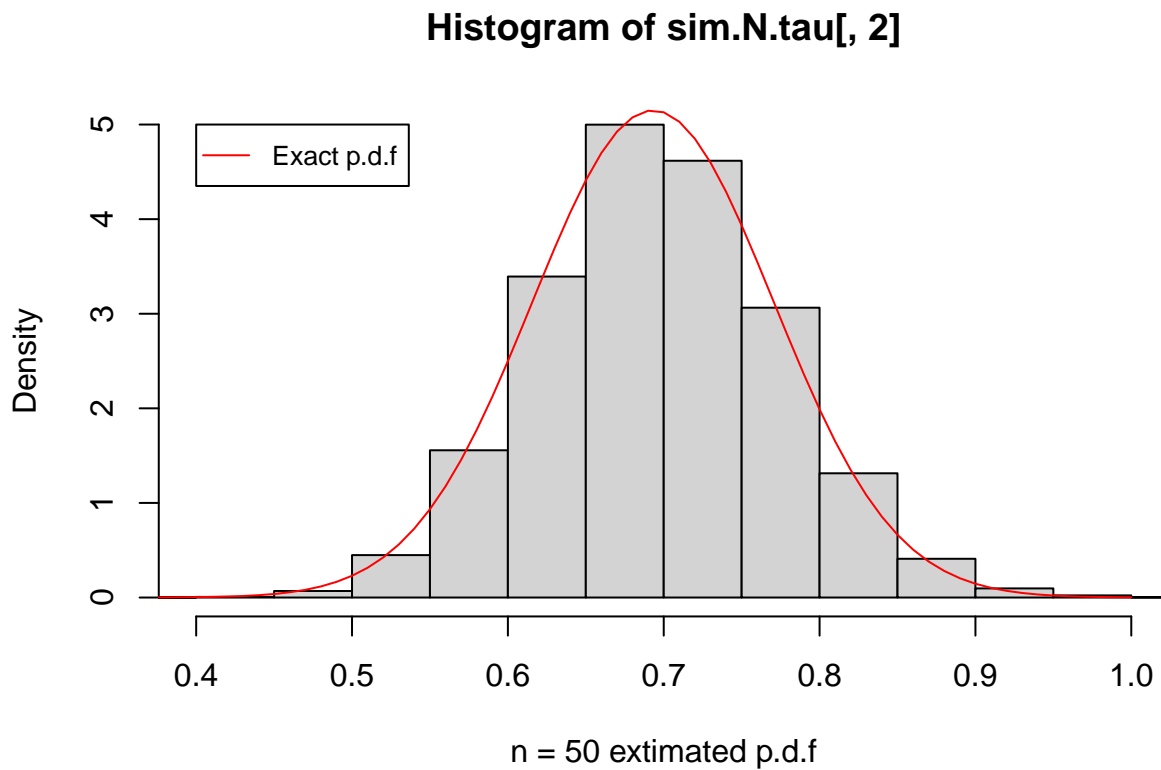
```

I've made a couple of plot to see if our theoretical results are supported by our simulations.

Plot for $n = 10$:



Plot for $n = 50$:



We can observe that the estimated p.d.f. fits perfectly the exact p.d.f.. I've runned the simulation with just 10^5 trials but to obtain something that fits even better under the curve I could have runned with more samples with the perks of having to wait a large amount of time to execute the code.

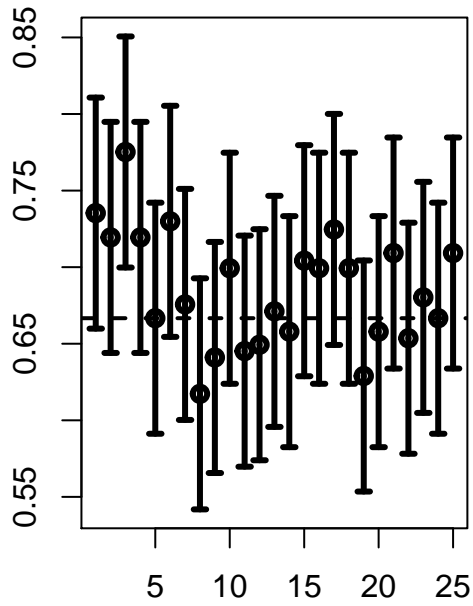
Coverage Probability, Wald C.I.

In the following lines I've made a simulation to study the coverage probability of the .95 Wald confidence interval for τ and θ

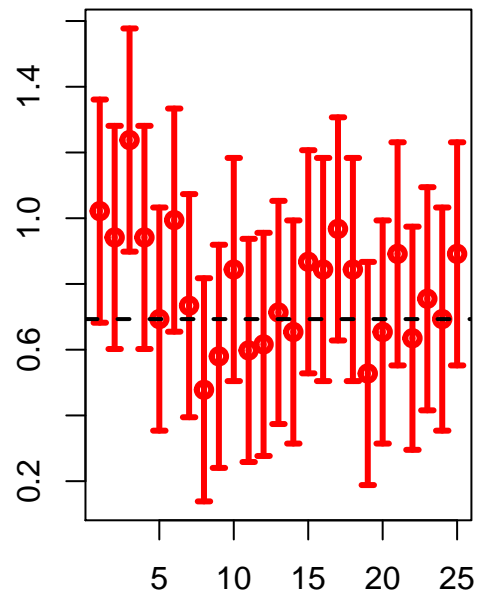
```
sim.N.CI_the <- matrix(NA, nrow = N, ncol = 4)
sim.N.CI_tau <- matrix(NA, nrow = N, ncol = 4)
for(i in 1:N){
  xnegbin10 <- rnbino(n=n[1],size=r,prob=theta0)
  xnegbin50 <- rnbino(n=n[2],size=r,prob=theta0)
  sim.N.CI_the[i,1:2] <- r*n[1]/(r*n[1]+sum(xnegbin10))
  sim.N.CI_the[i,1:2] <- sim.N.CI_the[i,1:2] + c(-1,1)*qnorm(p = alpha/2,
    lower.tail = FALSE)*sqrt(true_var10_o)
  sim.N.CI_the[i,3:4] <- r*n[2]/(r*n[2]+sum(xnegbin50))
  sim.N.CI_the[i,3:4] <- sim.N.CI_the[i,3:4] + c(-1,1)*qnorm(p = alpha/2,
    lower.tail = FALSE)*sqrt(true_var50_o)
  sim.N.CI_tau[i,1:2] <- tau(r*n[1]/(r*n[1]+sum(xnegbin10)))
  sim.N.CI_tau[i,1:2] <- sim.N.CI_tau[i,1:2] + c(-1,1)*qnorm(p = alpha/2,
    lower.tail = FALSE)*sqrt(true_var10)
  sim.N.CI_tau[i,3:4] <- tau(r*n[2]/(r*n[2]+sum(xnegbin50)))
  sim.N.CI_tau[i,3:4] <- sim.N.CI_tau[i,3:4] + c(-1,1)*qnorm(p = alpha/2,
    lower.tail = FALSE)*sqrt(true_var50)
}
theta10.inside <- apply(sim.N.CI_the[,1:2], MARGIN = 1,
  function(x)ifelse(theta0>=x[1]&theta0<=x[2],1,0))
theta50.inside <- apply(sim.N.CI_the[,3:4], MARGIN = 1,
  function(x)ifelse(theta0>=x[1]&theta0<=x[2],1,0))
tau10.inside <- apply(sim.N.CI_tau[,1:2], MARGIN = 1,
  function(x)ifelse(true_mean>=x[1]&true_mean<=x[2],1,0))
tau50.inside <- apply(sim.N.CI_tau[,3:4], MARGIN = 1,
  function(x)ifelse(true_mean>=x[1]&true_mean<=x[2],1,0))
```

In the plots below I've reported the first 25 confidence intervals for θ and τ

Confidence interval plot for θ and τ with $n = 10$:

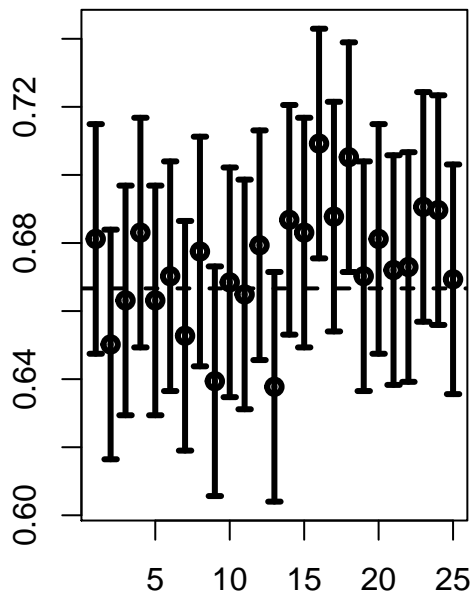


Observed C.I. for theta (n=10)

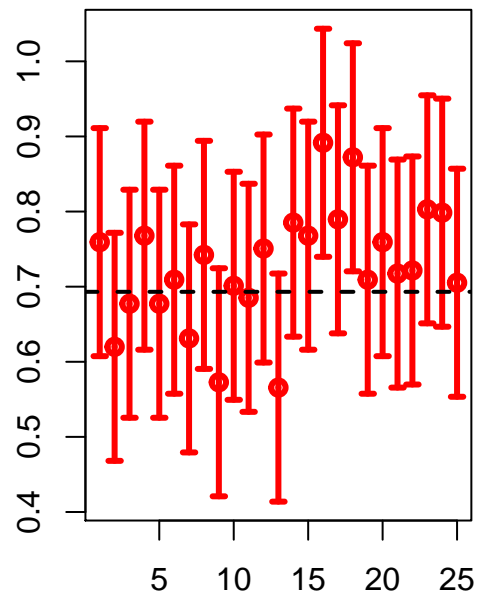


Observed C.I. for tau (n=10)

Confidence interval plot for θ and τ with $n = 50$:



Observed C.I. for theta (n=50)



Observed C.I. for tau (n=50)

Here's some results:

```
mean(theta10.inside) #Confidence probability for theta n=10
```

```
## [1] 0.95415
```

```
mean(theta50.inside) #Confidence probability for theta n=50
```

```
## [1] 0.94976
```

```
mean(tau10.inside) #Confidence probability for tau n=10
```

```
## [1] 0.94768
```

```
mean(tau50.inside) #Confidence probability for tau n=50
```

```
## [1] 0.94857
```

We can notice that in every scenario the confidence probability is close to .95 (even better some times). It's easy to state that our confidence intervals for $\hat{\theta}$ and $\hat{\tau}$ have the same behaviour.

2. Third Assignment

I've defined some recurrent values and settings below:

```
set.seed(29)
n = c(10,15,10,15) #Number of elements for every col
var = c(1,1,1,1) #Variance for every col
mu = c(0,0,0,0) #Mean for every col
k=4 #Number of cols
N = 1e6 #Number of iterations for simulation
```

In order to estimate the distribution of the F-Statistics for ANOVA, we need to calculate SSR and SSE for every trial, where SSR and SSE are defined as follows:

$$SSR := \sum_{j=1}^k \left(\sum_{i=1}^{n_j} (\bar{Y}_{*j} - \bar{Y})^2 \right)$$

$$SSE := \sum_{j=1}^k \left(\sum_{i=1}^{n_j} (Y_{ij} - \bar{Y}_{*j})^2 \right)$$

where \bar{Y}_{*j} is the observed mean for treatment j , \bar{Y} is the overall mean and Y_{ij} is the observed sample ij .

For every iteration l , after computing SSR_l and SSE_l , I'm going to store the value

$$F_{\text{obs}}[l] := \frac{\frac{SSR_l}{k-1}}{\frac{SSE_l}{n-k}}$$

in the array called *sim.N.F*.

```
sim.N.F <- rep(0,N) #initialized the array
for(l in 1:N){
  #defined the observed sample as follows
  #used var[1] as variance as every sample comes from N(0,1)
  y <- rnorm(sum(n), mean = 0, sd = sqrt(var[1]))
  #Treatment j mean
  estMuj <- c(sum(y[1:10])/n[1],
              sum(y[11:25])/n[2],
              sum(y[26:35])/n[3],
              sum(y[36:50])/n[4])
  #Overall mean
  estMu <- sum(y)/sum(n)
  SSR <- 0 #init SSR
  SSE <- 0 #init SSE
  for (i in 1:k){
    offset <- sum(n[1:(i-1)])
    for(j in 1:n[i]){
```

```

        SSR = SSR + (estMuj[i]-estMu)^2
        SSE = SSE + (y[offset+j]-estMuj[i])^2
    }
}
sim.N.F[1] <- (SSR/(k-1))/(SSE/(sum(n)-k)) #value of F observed (Fobs)
}

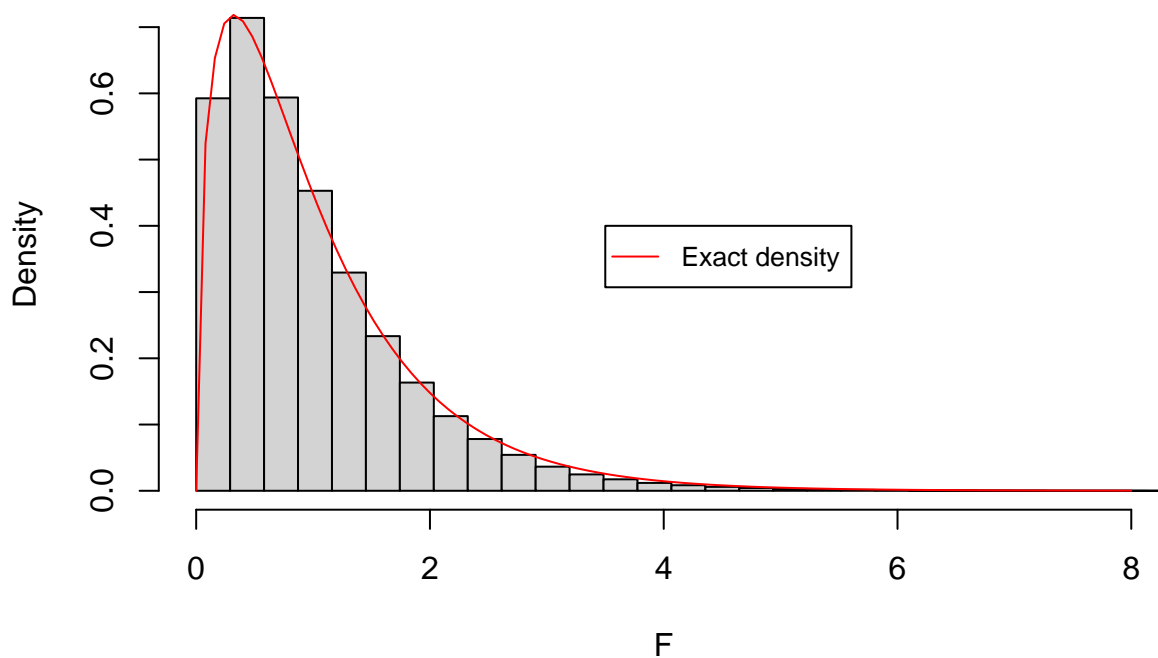
```

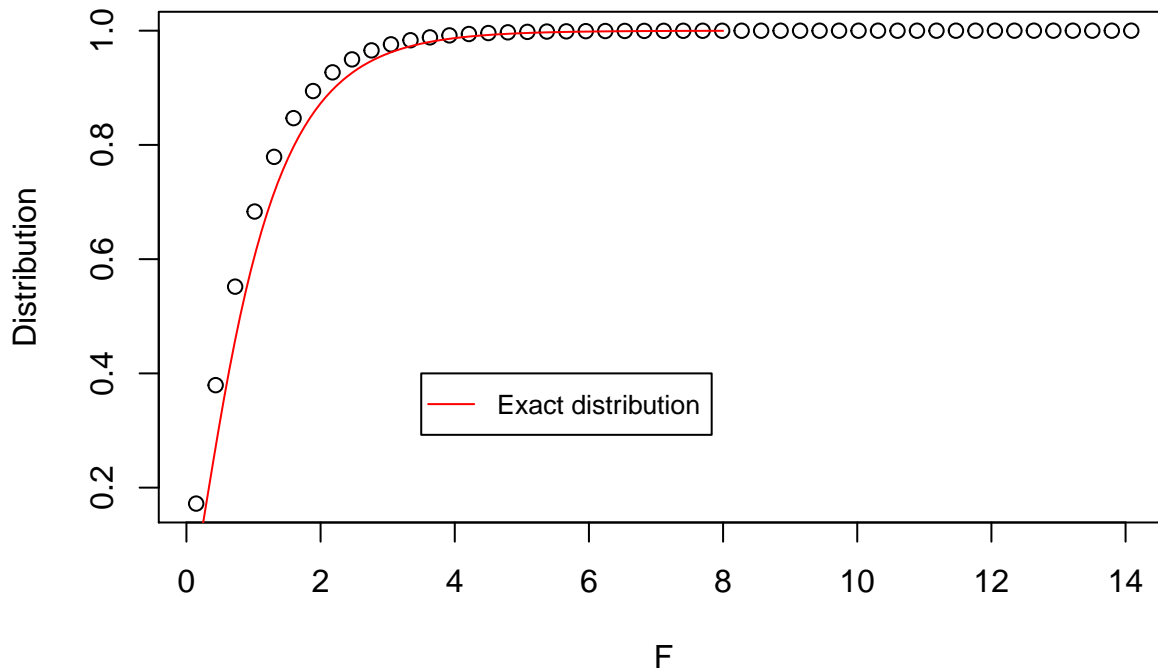
From theory we know that

$$\frac{\frac{SSR}{k-1}}{\frac{SSE}{n-k}} \sim \frac{\chi_{k-1}^2}{\chi_{n-k}^2}$$

This new r.v. should follow a F distribution with parameters k-1 and n-k. In the plots below there's a comparison between our simulated sample density/distribution and the real density/distribution of F.

Histogram of sim.N.F





Now we are asked to change the value of the variance as follows:

```
var <- c(1/2,2,1/2,2) #Vector of variances
```

From theory we know that the ANOVA test works with *r.v.'s* with the same variance, so I expect to see that our estimated p.d.f. won't fit the real p.d.f. as good as the previous simulation. In the next code lines I've re-executed the analysis with the changes I just made.

```
sim.N.F_n <- rep(0, N)
for(l in 1:N){
  offset <- 1
  y_n <- rep(0, 50)
  for (i in 1:k){
    y_n[offset:sum(n[1:i])] <- rnorm(n[i],
      mean = mu[i], sd = sqrt(var[i]))
    offset <- offset + n[i]
  }
  #Treatment j mean
  estMuj <- c(sum(y_n[1:10])/n[1],
    sum(y_n[11:25])/n[2],
    sum(y_n[26:35])/n[3],
    sum(y_n[36:50])/n[4])
  #Overall mean
  estMu <- sum(y_n)/sum(n)
  SSR <- 0
  SSE <- 0
  for (i in 1:k){
    offset <- sum(n[1:(i-1)])
```

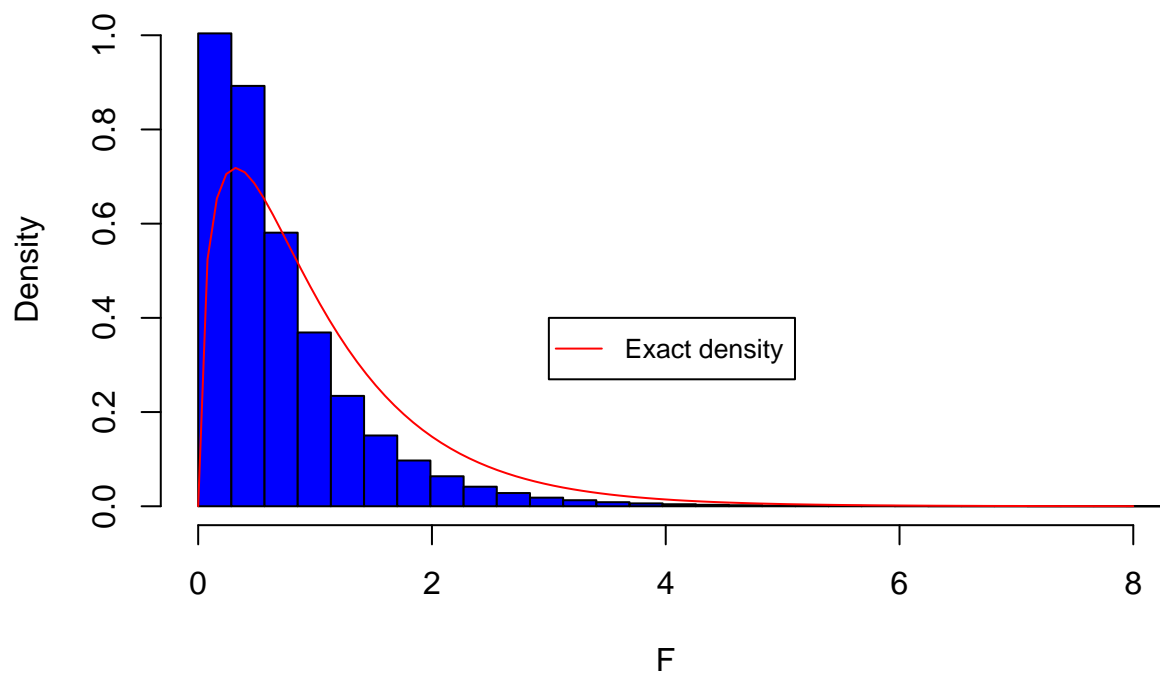
```

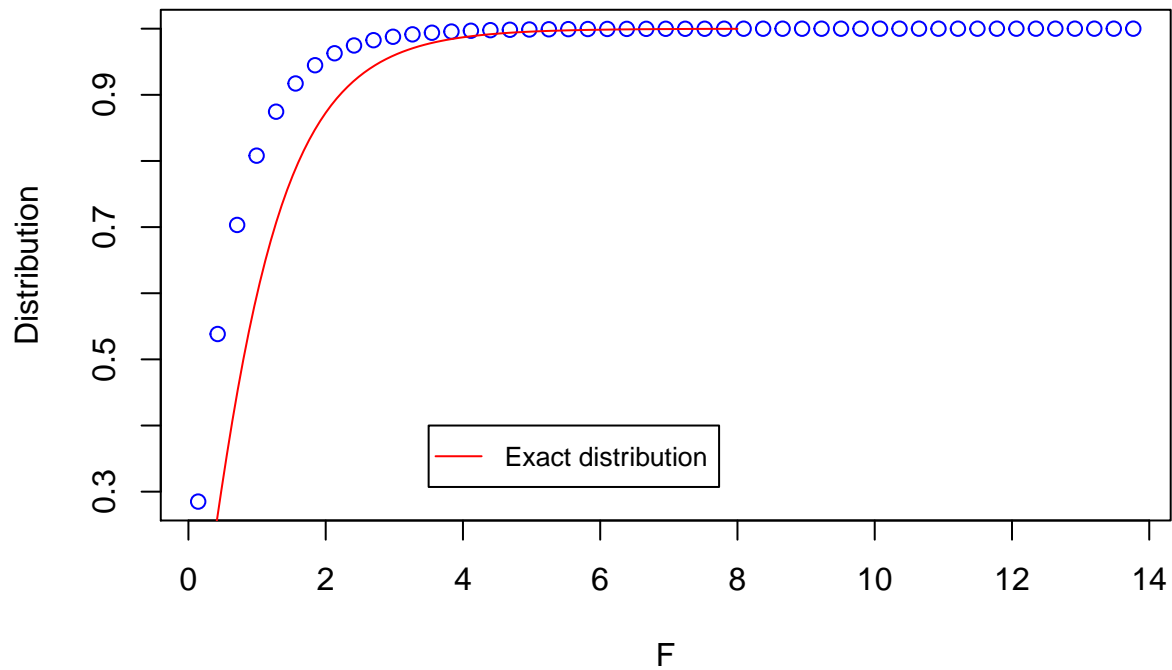
    for(j in 1:n[i]){
      SSR = SSR + (estMuj[i]-estMu)^2
      SSE = SSE + (y_n[offset+j]-estMuj[i])^2
    }
  }
  #value of new F observed (Fobs)
  sim.N.F_n[l] <- (SSR/(k-1))/(SSE/(sum(n)-k))
}

```

Those are the plots obtained from the latest analysis:

Histogram of sim.N.F_n





We can observe that, with this changes in the variance array, our simulated p.d.f. for f-statistics does not follow the exact p.d.f.. As I've stated before, It's probably due to the fact that ANOVA works for r.v.'s with the same variance and this is not the case.