1)

Suppose a random variable X has pdf as $f(x)=2e^{-2(x-1)}$, x>1. Which of the following represents P(0 < X < 4)? (Note: you do not need to solve for exact number).

- 1. $\int_0^4 2e^{-2(x-1)} dx$; 2. $\int_1^4 2e^{-2(x-1)} dx$; 3. $\int_0^4 x2e^{-2(x-1)} dx$; 4. $\sum_{x=0}^4 2e^{-2(x-1)}$; 5. $\int_0^\infty x2e^{-2(x-1)} dx$;

A random variable X has pdf

$$f(x) = \frac{2^x}{x!}e^{-2}, x = 0,1,2...$$

Find P(X = 1).

Rscript:

((2)/factorial(1))*exp(-2)

Answer:

0.2707

Then find $P(-2 \le X \le 4)$.

Rscript:

 $sum(((2^{(0:3)})/factorial(0:3))*exp(-2))$

or

x_range<-c(0:3)
f.x<-function(x) ((2^x)/factorial(x))*exp(-2)
f_x<-function(x) f.x(x)*(x %in% x_range)
sum(f_x(x_range))</pre>

Note: I don't know which way you prefer I don't want my marks to be cut because I use a longer code instead of one line code. So please let me know in future which one should I use and Please don't cut marks for writing both the scripts here as I don't know what exactly you want here.

Answer:

0.8571

Give your answers to at least four decimal places.

3)

If two carriers of the gene for albinism marry and have children, then each of their children has a probability of 1/4 of being albino. Let the random variable Y denote the number of their albino children out of all 3 of their children. Then Y follows a binomial(n, p) distribution. Find the values for n and p.

Answer:

n=3

p=1/4=.25

For Y following a binomial (n = 3, p = 0.25) distribution, compute the following:

$$P(Y \le 2) =$$

$$E(Y) =$$

$$Var(Y) =$$

Give your answers to at least four decimal places.

Rscript:

sum(dbinom((0:2),size=3,p=0.25))

Answer:

 $P(Y \le 2) = 0.9844$

Rscript:

y < -c(0:3)

sum(y*dbinom(y,size=3,p=0.25))

Answer:

E(Y) = 0.7500

Rscript:

y < -c(0:3)

EY<-sum(y*dbinom(y,size=3,p=0.25)) sum((y-EY)^2*dbinom(y,size=3,p=0.25))

Answer:

Var(Y) = 0.5625

For X following a Chi-square distribution with degree of freedom m = 3, compute the following:

$$P(1 < X < 4) =$$

$$E(X) =$$

$$Var(X) =$$

Give your answers to at least four decimal places.

Also, use a Monte Carlo simulation with sample size n=100,000 to estimate P(1 < X < 4). What is your Monte Carlo estimate? Does it agrees with the answer above?

Rscript:

x<-function c(0:Inf)
integrate(function(x) dchisq(x,df=3), lower=1,upper=4)</pre>

Answer:

$$P(1 < X < 4) = .5398$$

Rscript:

x<-function c(0:Inf) integrate(function(x) x*dchisq(x,df=3), lower=0,upper=Inf)

Answer:

E(X) = 3 (mean is same as degree of freedom for chi-square distribution)

Rscript:

x<-function c(0:Inf)

EX<-integrate(function(x) x*dchisq(x,df=3), lower=0,upper=Inf)\$value integrate(function(x) $((x - EX)^2)*dchisq(x,df=3)$, lower=0,upper=Inf)

Answer:

Var(x) = 6 (Variance is twice the degree of freedom for chi-square distribution)

Rscript:

X<-rchisq(n=100000, df=3) mean((1<x) & (x<4))

Answer:

.5378 Yes the monte carlo estimate agrees with above.

Suppose X follows a Chi-square distribution with degree of freedom m = 5 so that E(X) = 5 and Var(X) = 10. Also, let Y = 4X - 10. Find E(Y) and Var(Y). Does Y follow a Chi-square distribution with degree of freedom m=10?

$$E(Y) =$$

$$Var(Y) =$$

Does Y follow a Chi-square distribution with degree of freedom m = 10?

Answer:

Y = 4X-10
Using the formula
$$E(aX+b)=aE(X)+b$$

 $E(Y) = 4*E(X) - 10$
 $E(Y) = 20-10 = 10$

Using the formula
$$Var(aX+b) = a^2Var(X)$$

 $Var(Y) = 4^2Var(X)$
 $Var(Y) = 16*10=160$

For Y to follow chi-square distribution with degree of freedom m=10 then it should follow 2 conditions namely;

- 1. E(Y) = 10 as mean for chi square distribution is same as degree of freedom
- 2. Var(Y) = 20 as variance for chi square distribution is twice the degree of freedom.

E(Y) is 10 but Var(Y) is 160 so Y doesn't follow a chi square distribution.

The Zyxin gene expression values are distributed according to $N(\mu = 1.6, \sigma = 0.4)$

(a) What is the probability that a randomly chosen patient have the Zyxin gene expression values between 1 and 1.6? Rscript:

x<-c(-Inf,Inf)

integrate(function(x) dnorm(x,mean=1.6,sd=0.4), lower=1,upper=1.6)\$value

Answer:

0.4332

(b) Use a Monte Carlo simulation of sample size n=500,000 to estimate the probability in part (a). Give your R code, and show the value of your estimate.

Rscript:

x<-rnorm(n=500000, mean=1.6, sd=.4) mean((1<x) & (x<1.6))

Answer:

0.4334

(c) What is the probability that exactly 2 out of 5 patients have the Zyxin gene expression values between 1 and 1.6? Please show your work on how to arrive at the answer. Give your answer to at least four decimal places.

Rscript:

dbinom(2,size =5, prob=.4332)

Answer:

Now this normal distribution become binomial distribution with set =5 and probability =0.4332.

We have to find P(X=2) which we can calculate by dbinom(2,size=5, prob=.4332)

Which gives us the answer - .3417

.3417 is the probability of exactly 2 patients out of 5 having the zyxin gen expression values between 1 and 1.6.

(a) Hand in a R script that calculates the mean and variance of two random variables $X\sim F(m=2,n=5)$ and $Y\sim F(m=10,n=5)$ from their density functions.

Rscript:

Mean of X:

x < -c(0, Inf)

integrate(function(x) x*df(x,df1=2,df2=5), lower=0,upper=Inf)\$value

Answer:

1.6667

Variance of X:

x < -c(0, Inf)

 $EX < -integrate(function(x) \ x*df(x,df1=2,df2=5), \ lower=0, upper=Inf) \\ value integrate(function(x) \ (x-EX)^2*df(x,df1=2,df2=5), \ lower=0, upper=Inf) \\$

Answer:

13.8889

Mean of Y:

y < -c(0, Inf)

integrate(function(y) y*df(y,df1=10,df2=5), lower=0,upper=Inf)\$value

Answer:

1.6667

Variance of Y:

y < -c(0, Inf)

EY<-integrate(function(y) y*df(y,df1=10,df2=5), lower=0,upper=Inf)\$value integrate(function(y) (y-EY)^2*df(y,df1=10,df2=5), lower=0,upper=Inf)

Answer:

7.2222

(b) Use the formula in Table 3.4.1 to calculate the means and variances directly.

Mean of X:

$$\frac{n}{n-2} = 5/3 = 1.6667$$

Variance of X:

$$\frac{2n^2(m+n-2)}{m(n-2)^2(n-4)} = 50*5/18 = 13.8889$$

Mean of Y:

$$\frac{n}{n-2} = 5/3 = 1.6667$$

Variance of Y:

$$\frac{2n^2(m+n-2)}{m(n-2)^2(n-4)} = 50*13/90 = 7.2222$$

(c) Run your script in (a), and check that your answers agree with those from part (b).

Yes, my scripts agrees with the answers from part (b)