

# R LANGUAGE

## DATA LOADING(TYPE 2)

```
a1=c(64,78,48,11,47,50,47,06,63,34)
a2=c (22,43,77,76,66,39,44,34,84,85)
a3=c (24,66,18,20,10,45,62,96,09,44)
a=cbind (a1, a2, a3)
summary (a)
boxplot (a)
```

## DATA LOADING(TYPE 3)

### MATRIAL STATUS DATA

```
status=matrix(c(180,210,70,40), nrow=1,ncol=4, byrow=T)
rownames (status)=c("Marital Status")
colnames (status)=c("Never married", "Married", "Divorced", "Widowed")
status
barplot (status, beside=T,main="Graphical Representation",
legend.text=rownames (status),col='green')
```

## FREQUENCY COUNT

```
a=c(1,1,1,3,4,3,5,5,6,6,6,2,1,3,7,7,7,4,3,5,7,1,2,3,4,8)
b=table (a)
b
cbind (b)
plot(b,col='red',xlab='Numbers',ylab='Frequency',lwd=3)
```

## HISTOGRAM

### TYPE 1

```
a=rep(c(149.55,154.55,159.55,164.55,169.55,174.55,179.55,184.55),c(1,3,24,58,60,27,2,2))
H=hist (a, breaks=c(144.55,149.55,154.55,159.55,164.55,169.55,174.55,179.55,184.55))
H$counts= H$counts/(5*sum (H$counts))
plot (H, xlab="Height",ylab="Relative Frequency Density",main="Hisogram",ylim=c(0,0.1))
# With normal approximation
curve (dnorm (x,mean(a), sd(a)),col="blue", add=TRUE)
```

## TYPE 2

```
a=rep(c(149.55,154.55,159.55,164.55,169.55,174.55,179.55, 184.55),c(1,3,24,58,60,27,2,2))
H=hist (a, breaks=c(144.55,149.55,154.55,159.55,164.55,169.55,174.55,179.55, 184.55))
H$counts= H$counts/5
plot (H,xlab="Height",ylab="Frequency Density", main="Hisogram",ylim=c(0,15))
# With normal approximation
b=length(a)
curve (b*dnorm(x,mean(a), sd(a)),col="blue", add=TRUE)
```

## TYPE 3

```
a=rep(c(149.55,154.55,159.55,164.55,169.55,174.55,179.55,184.55),c(1,3,24,58,60,27,2,2))
H=hist(a,breaks=c(144.55,149.55,154.55,159.55,164.55,169.55,174.55,179.55,184.55))
plot(H,xlab="Height",ylab="Frequency",main="Hisogram",ylim=c(0,70))
b=length(a)
b1=min(diff (H$breaks))
curve(b*b1*dnorm(x,mean(a),sd(a)),col="blue", add=TRUE)
```

## BOX PLOT

### TYPE 1

```
a=c(31,71,69,59,70,74,41,87,37,46,56,34,62,15,53,81,76,24,21,8)
summary (a)
boxplot (a)
boxplot (a, col = "red")
```

### TYPE 2

```
a=c(31,71,69,59,70,74,41,87,37,46,56,34,62,15,53,81,76,24,21,8)
b=c(10,23,55,60,62,68,53,78,60,87,78,93,94,90,65,88,76,96,86,66)
dat=cbind (a,b)
colnames(dat) =c("Marks A","Marks B")
boxplot (dat)
boxplot(dat,col =c("red","blue"))
```

## STEM- LEAF PLOT

```
a=c(11,13,18,16,21,29,24,30,39,34,33)
```

```
stem (a)
```

```
b=c(11,13,18,16,21,29,24,30,39,34,33,48, 47,42,50,55,53,52,61,69,65,62)
```

```
stem (b)
```

```
stem (b, scale=2)
```

## FREQUENCY POLYGON

```
a=c(0,1,2,3,4,5,6,7,8)
```

```
b=c (0,4,33,76,50,26,8,1,0)
```

```
plot (a,b,type="o",xlab="Number of Peas",ylab="Frequency")
```

## PIE CHART

### TYPE 2

```
states=c("Punjab","Haryana", "Tamil Nadu","Andhra Pradesh", "Uttar Pradesh", "Others")
```

```
procurement=c(5486,1248,589,3987,1296, 1654)
```

```
rice.proc=data.frame(states,procurement)
```

```
rice.proc=transform(rice.proc,percentage=(rice.proc$procurement/14260)*100)
```

```
rice.proc
```

```
attach(rice.proc)
```

```
pie(procurement,labels=c("Punjab(38.47%)", "Haryana (8.75%)", "Tamil Nadu(4.13%)",
```

```
"Andhra Pradesh (27.96%)", "Uttar Pradesh(9.09%)","Others(11.60%)"),
```

```
main="Procurement of Rice('000 tonnes)\nduring oct. 1993 to Sept. 1994",
```

```
col=c("royalblue4","orchid","chocolate4","olivedrab3","orange3","purple 1","violetred4"))
```

```
detach(rice.proc)
```

## PLOT OGIVES

```
Height=c(149.55,154.55,159.55,164.55,169.55,174.55,179.55,184.55)
```

```
a = c(1,3,24,58,60,27,2,2)
```

```
b=cumsum (a)
```

```
plot (Height,b, type="o",ylab="Cumulative Frequency", col="blue",xlim = c(144,185))
```

```
a1=rev(a)
```

```
b1=cumsum (a1)
```

```
b2=rev (b1)
```

```
lines (Height-5,b2, type="o",col="red")
```

## DESCRIPTIVE STATISTICS

### TYPE1

```
x=c(70,48,84,21,84,33,64,36,61,50,27,87,41,65,33)
```

```
max (x)-min(x) # Range
```

```
var (x) # Variance
```

```
sd(x) # Standard Deviation
```

```
sum (abs (x-40))/length(x) # Mean deviation about 50
```

```
sum (abs (x-mean (x)))/length(x) # Mean deviation about mean
```

```
quantile (x) # Quartiles
```

```
IQR=67.5-64.5 # Using Quartile Data
```

```
(sd(x)/mean (x))*100 # Coefficient of Variation
```

### TYPE2

```
x=c (60,39,75,33,69,47,69,32,59,52,49,63)
```

```
r=3 # r=1,2,3,4
```

```
sum (x^r)/length (x) # Raw Moments
```

```
sum ( (x-mean (x))^r)/length(x) # Central Moments
```

## CORRELATION

```
x=c(126,157,153,156,152,135,145,132,153,143,165,132,180,161,170,176,165,163,157,180)
y=c(292,367,361,366,343,326,331,307,343,322,363,296,412,377,394,384,370,365,349,403)
z=data.frame(x,y)
colnames (z)=c("Marks in School", "Marks in College")
plot (z,col="blue") # Scatter Plot
cor (x,y) # Correlation Coefficient
cor (z) # Correlation Matrix
```

## REGRESSION

### TYPE1

```
x=c(126,157,153,156, 152,135,145,132, 153,143,165,132,180,161,170,176,165,163,157,180)
y=c(292,367,361,366,343,326,331,307,343,322,363,296,412,377,394,384,370,365,349,403)
fit=lm(y~x)
summary(fit)
plot (x,y,xlab = 'Marks in School', ylab='Marks in College',col='blue',cex=1.2,
pch=16,main='Regression Fit')
abline(fit,lwd=2,col='green')
x1 = data.frame (x=seq(min(x),max(x),0.5))
y1=predict (fit, interval="conf",newdata=x1)
y2=predict (fit, interval="pred",newdata=x1)
matlines(x1$x,y1,lty=c(1,2,2),col=c("green", "red", "red"),lwd=c(2,1,1))
matlines(x1$x,y2,lty=c(1,3,3),col=c("green", "black", "black"),lwd=c (2,1,1))
```

## TYPE2

```
x=c(107,110,121,124,129,132,131,153,142,151,158,152,164,152,162,156,175,178,172,177,174,183,  
172,176)
```

```
y=c(99,82,102,81,89,87,59,112,74,91,116,91,96,85,131,136, 162,121,137,82,165,159,91,193)
```

```
fit=lm(y~x + I(x^2)) # Polynomial Fit
```

```
summary (fit)
```

```
plot (x,y,xlab='X',ylab='Y',ylim=c(20,200), col='blue',main='Polynomial Fit',
```

```
cex=1.2,pch=16)
```

```
x1 = data.frame(x=seq(min(x), max (x),0.5))
```

```
lines(x1$x,predict(fit,newdata=x1),col="green",lwd=2)
```

```
y1=predict (fit,interval="conf",newdata=x1)
```

```
y2=predict (fit,interval="pred",newdata=x1)
```

```
matlines(x1$x,y1,lty=c(1,2,2),col=c("green", "red", "red"),lwd=c(2,1,1))
```

```
matlines(x1$x,y2,lty=c(1,3,3),col=c("green", "black", "black"),lwd=c(2,1,1))
```

## ESTIMATE

```
x1=rnorm(100, 0, 1) # Generate 100 Random Samples from N(0,1)
```

```
xbar1=mean (x1)
```

```
sd1=sd(x1) # var (sd) is sample variance(sd) in R with denominator n-1
```

```
x2=rnorm(500, 0, 1) # Generate 500 Random Samples from N(0,1)
```

```
xbar2=mean (x2)
```

```
sd2=sd(x2)
```

```
x3=rnorm(5000, 0, 1) # Generate 5000 Random Samples from N(0,1)
```

```
xbar3=mean (x3)
```

```
sd3=sd(x3)
```

```
hist(x3,col='green',main="Histogram") # Histogram for sample size 5000
```

## Exponential curve fitting

```
x=c(0,.01,.03,.05,.07,.09,.11,.13,.15,.17,.19,.21)
y=c(1,1.03,1.06,1.38,2.09,3.54,6.41,12.6,22.1,1.39,65.32,99.78)
y1=log(y)
fit=lm(y1~x)
a=exp(fit$coefficients[1])
b=fit$coefficients[2]
summary(fit)
plot(x,y,main = "Exp Fit")
x1=seq(min(x),max(x)+0.1,0.01)
y1=a*exp(b*x1)
lines(x1,y1,lty=1,col="blue",lwd=2)
```

## NORMAL DISTRIBUTION GRAPH

```
x=seq(-5,5,0.1)
f=dnorm(x,mean = 0.4,sd=1.2)
plot(x,f,type="l",xlim = c(min(x),max(x)),ylim = c(0,1),lwd=2,col="blue",ylab="f(x)" )
x=seq(-5,5,0.1)
f=dnorm(x,mean = 0.4,sd=1.2)
f1=dnorm(x,mean = 0.4,sd=0.6)
f2=dnorm(x,mean = 0.4,sd=1.8)
plot(x,f,type="l",xlim = c(min(x),max(x)),ylim = c(0,1),lwd=2,col="blue",ylab="f(x)" )
lines(x,f1,type="l",xlim = c(min(x),max(x)),ylim = c(0,1),lwd=2,col="red")
lines(x,f2,type="l",xlim = c(min(x),max(x)),ylim = c(0,1),lwd=2,col="green")
legend(3,0.8,legend =
c("N(0.4,1.2)", "N(0.4,0.6)", "N(0.4,1.8)",col=c("blue", "red", "green"),lty=c(1,1,1)))
```

## BINOMIAL CURVE FITTING

```
x=c(0,1,2,3,4,5,6,7,8,9)
f=c(7,33,54,38,35,15,7,4,1,1)
n=sum(f)
dat=rep(x,f)
hist(dat,col="blue",xlab="x",main="Histogram") #Idea about the distribution
lam=mean(dat) #estimate of lambda=bar(x)
f1=f*0 #Initialize expected frequency
f1[1]=exp(-lam) #Initial value of e^(-lam)
for(i in 1:8)
{
  f1[i+1]=(lam/i)*f1[i]
}
f1[10]=1-sum(f1) #adjusting total probability
Ef=n*f1 #expected frequency
out=data.frame(x,Ef,f)
colnames(out) = c("|x|","|Expected Frequency|","|Observed Frequency|")
format(out,scientific=F,digit=3)
```

## BINOMIAL PLOT DISTRIBUTION

```
n=10
p=0.2
x=0:n
f=dbinom(x, n, p, log = FALSE)
plot (x,f,type="h",xlim =
c(min(x),max(x)),ylim =
c(0,1),lwd=2, col="blue",ylab = "p(X=x)")
points(x,f,pch=16,cex=1,col="dark red")
```



## NORMAL CURVE FITTING

```
x=seq (146.5, 181.5,5) # Midvalues
f=c(1,3,24,58,60,27,2,2)
n=sum (f)
dat=rep(x,f)
hist(dat,xlab="Height",main="Histogram",ylim=c(0,max(f)+3)) # ylim needed for curve
xbar=mean (dat) # Estimate of mu = mean (x)
s=sd(dat)
curve (n*5*dnorm(x,xbar,s),col="blue", add=T) # Add curve
xi=seq(144,184,5) # Intervalvalues
x1= (xi-xbar)/s # Normalize
y1=dnorm(x1) # Standard normal value
F1=pnorm(x1) # Df values
F2=c (0,F1,1) # Extended DF, including -Inf, Inf
F3= diff (F2) # Actual Probability values
Ef=n+F3 # Expected frequency
x1=c(-Inf,x, Inf) # Extended x, including -Inf, Inf
f1=c (0,f,0) # Extended f, including -Inf, Inf
Out=data.frame (x1,Ef,f1)
colnames (Out) =c("Ix(Mid) |", "| Expected Frequency ", "I Observed Frequency ")
format (Out, scientific = F,digit =3)
```

## CIRCULAR SYSTEMATIC SAMPLING

```
N=90
n=17
r=sample(c(1:90),size=1) #One sample using SRS
r
k=N/n
k=5 #Check the above k and write the nearest integer
samp=seq(r,r+k*n,k) #Generate systematic sample
samp=samp%%90*1 #Use nodulo function i.e remainder
```