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Course :- MCA

Semester :- 1st

Paper Name :- Scripting language and R lab

Paper Code :- PMC 103

Type of Paper :- Regular

Ans 3 - # Dplyr library function.

```
library(dplyr)
setwd("G:/MCA")
mydata <- read.csv("most runs.csv")
mydata
```

Descriptive statistics

```
summary(mydata)
dim(mydata)
str(mydata)
names(mydata)
```

select function

```
my_subdata <- select(mydata, batsman, overage)
my_subdata
```

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filter & arrange funⁿ

mysubdata 1 ← filter (mydata, average > 50)

mysubdata 1

my subdata 2 ← arrange (mydata, desc (Average))

my subdata 3 ← arrange (mydata, desc (strike rate))

Top and Bottom 5 average Batsman

head (mysubdata 2)

tail (mysubdata 2)

Different Plot of Dataset

Histogram

hist (mydata \$ average, col = c ("blue", "green", "red"),
x lab = "Average", y lab = "Players", break = 50)

Scattered Plot

plot (mydata \$ strike rate, col = c ("blue", "green", "red"),
x lab = "players", y lab = "strike rate")

Barplot

barplot (mydata \$ average, col = c ("blue", "green", "red"),
x lab = "players", y lab = "Average")

Box plot

```
boxplot(mydata$average, col=c('blue', 'green', 'red')  
xlab="players", ylab="Average")
```

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Ans 4. # descriptive statistics
 summary(mydata)
 dim(mydata)
 str(mydata)
 names(mydata)

Inferential statistics

Chi-squared test

model <- chisq.test(mydata)

model

#Output p-value = 0.446283 > 0.05

Thus 'mydata' is highly correlated and we

accept the NULL Hypothesis

Correlation coefficient

cor(mydata\$Batsman, mydata\$runs)

#Output 0.99324 > 0.8

Thus Batsman & runs is strongly correlated

to each other

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Anova Test

```
mysubdata4 <- anov(mydata $ runs ~ mydata $ average)
```

```
mysubdata4
```

Output $P(>F)$ is 0.0013 as this value is less than
0.05 then we reject NULL hypothesis and accept
the alternative hypothesis.

T-test

```
t.test(mydata, mu=100)
```

Here p-value is $0.446283 > 0.05$

so we accept the NULL hypothesis.

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