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Course :- MCA 1sem
Paper name & code :- Scripting Language & R Lab :

As 3 : # Dplyr library function

```
library (dplyr)
Setwd ("G:/MCA")
mydata <- read.csv ("mostruns.csv")
mydata
# Descriptive Statistics
summary (mydata)
dim (mydata)
str (mydata)
names (mydata)
# select function.
mysubdata <- select (mydata, batsmans, average)
mysubdata
# filter and arrange function.
mysubdata1 <- filter (mydata, average > 50)
mysubdata1
```

```
mysubdata 2 <- arrange (mydata, desc (
  average))
mysubdata 2
```

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```
mySubdata3 <- arrange (my data, desc (strikerate))
```

```
# Top 4 Bottom 5 average batsman.
```

```
head (mySubdata2)
```

```
tail (mySubdata2)
```

```
# mutate function (to add a column to data set
```

```
my data <- mutate (mydata, Performance = runs - balls)
```

```
# Different Plot of Data set
```

```
# Histogram
```

```
hist (mydata $ average, col = c ('blue', 'green', 'yellow'))
```

```
xlab = "Average", ylab = "Players", break = 50)
```

```
# Scattered Plot
```

```
plot (mydata $ strikerate, col = c ('blue', 'green', 'red'),
```

```
xlab = "Players", ylab = "strikerate")
```

```
# Bar Plot
```

```
barplot (mydata $ average, col = c ('blue', 'green', 'red'),
```

```
xlab = "Players", ylab = "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 \$ Batman , mydata \$ runs)

output 0.99324 > 0.8

Thus Bat man & runs is strongly correlated to

each other

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A move test

~~mydata~~.

my subdata 4 \leftarrow aov (mydata\$run ~ mydata\$ average

my subdata 4

output $Pv(>F)$ is 0.0013 as this values is less than

0.05 then we reject NULL Hypothesis and accept

the alternative Hypothesis .

T-Test

This gives us the T- Score for the data set

t.test (mydata , mu = 100)

Here p-value is 0.446283 > 0.05

So we accept the NULL Hypothesis .

Frathans