

VERSION 1.0

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THE TITAN INSURANCE COMPANY

CASE STUDY

PRESENTED BY: SMDM GROUP 13

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THE TITAN INSURANCE COMPANY

A. DESCRIBE THE FIVE PER CENT SIGNIFICANCE TEST YOU WOULD APPLY TO THESE DATA TO DETERMINE WHETHER NEW SCHEME HAS SIGNIFICANTLY RAISED OUTPUTS?

We shall use Paired t-test for this case as we are interested in evaluating the effectiveness of new scheme that has been Implemented in Titan. We are to infer whether the average output has increased or not. Hence this is a case of two population means where you have two samples in which observations in one sample can be paired with observations in the other sample i.e. Before-and-after execution of the scheme.

B. WHAT CONCLUSION DOES THE TEST LEAD TO?

We have used R as the tool to arrive at a decision;

Since we have to find whether the new scheme has significantly raised output we can formulate the hypotheses as follows:

Null Hypothesis-> $H_0 : \mu_1 = \mu_2$

Alternate Hypothesis-> $H_1 : \mu_1 < \mu_2$

Following is the R code for the same:

Calculating the paired t-test for the 2 samples

<<R-Code>>

```
paired.t.test=t.test(Old.Scheme, New.Scheme, paired = T, alternative = "less")
```

<<R-Output>>

Paired t-test

data: Old.Scheme and New.Scheme

t = -1.5559, df = 29, p-value = 0.06529

alternative hypothesis: true difference in means is less than 0

95 percent confidence interval:

-Inf 368.1762

sample estimates:

mean of the differences -4000

Inference:

The p-value 0.06529 is greater than 0.05 significant level of the test, hence we accept the Null Hypothesis with 95% confidence.

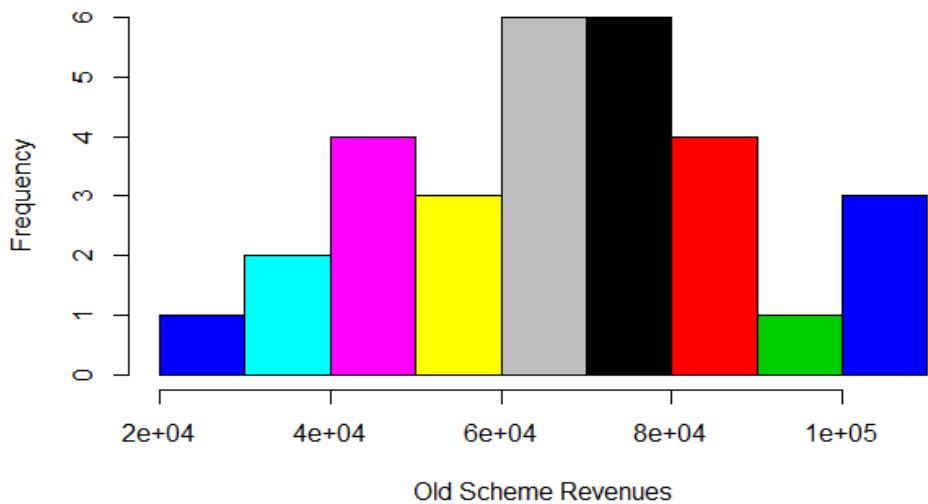
This implies that there is no significant increase in the average output after implementation of the new scheme. Hence the Titan Insurance company can stick to the older scheme which would be less expensive than the new scheme.

C. WHAT RESERVATIONS HAVE YOU ABOUT THIS RESULT?

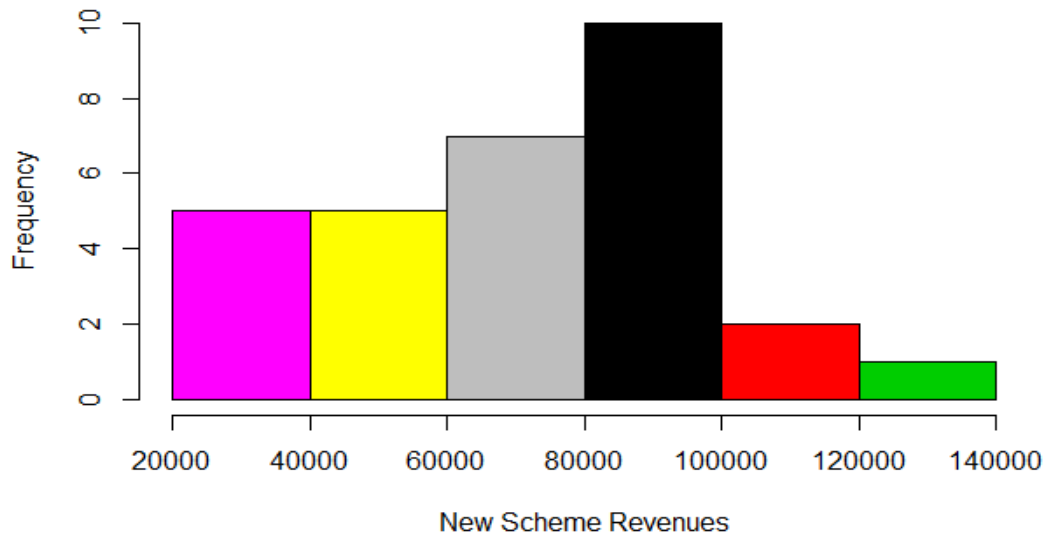
The following reservations are there about the result while performing the paired t-test

- The p-value of 0.0625 implies that there is a weak evidence of rejecting the alternate hypothesis.
- The dependent variable must be continuous (interval/ratio).
- The observations are independent of one another.
- The dependent variable should be approximately normally distributed.

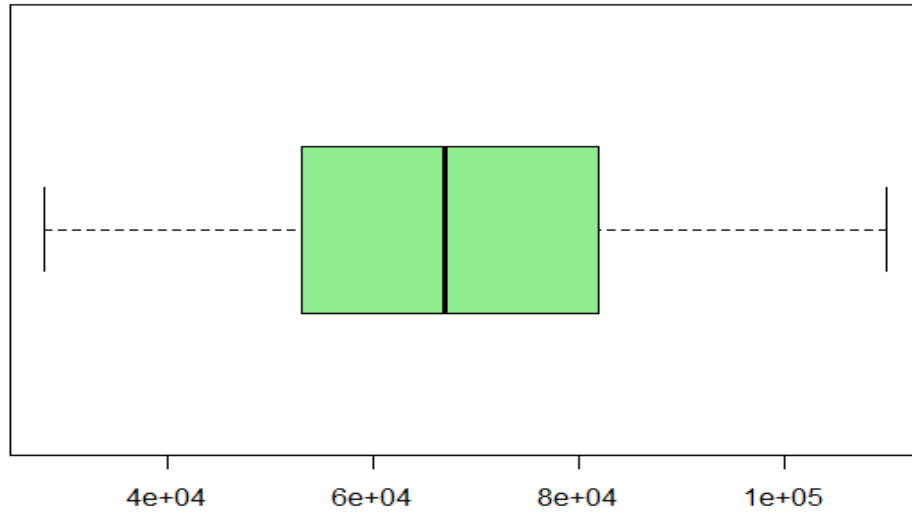
Old Scheme follows a Normal Distribution Approximately



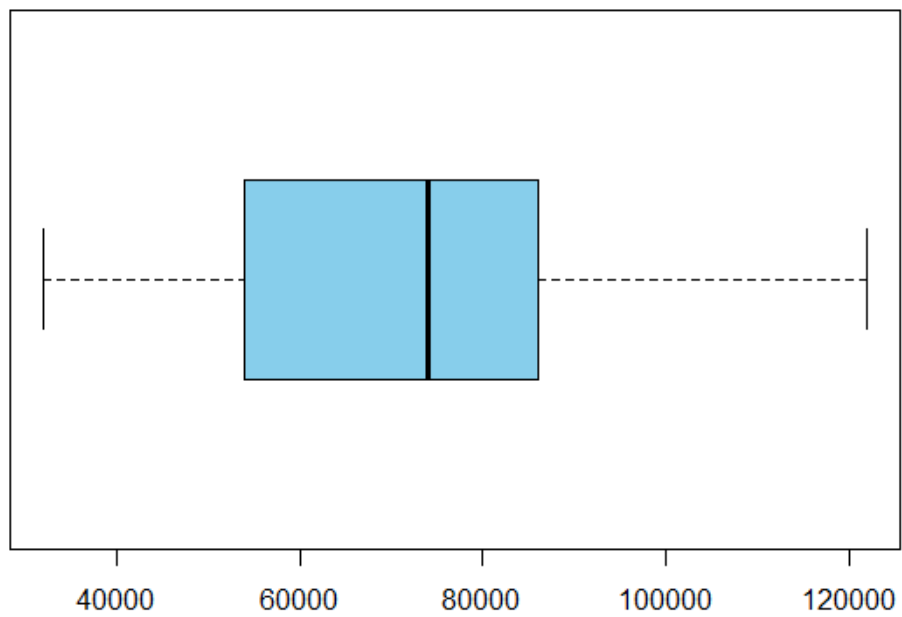
New Scheme follows a Normal Distribution Approximately



Old Scheme has no outliers



New Scheme has no outliers



D. SUPPOSE IT HAS BEEN CALCULATED THAT IN ORDER FOR TITAN TO BREAK EVEN, THE AVERAGE OUTPUT MUST INCREASE BY £5000. IF THIS FIGURE IS ALTERNATIVE HYPOTHESIS, WHAT IS:

(I) THE PROBABILITY OF A TYPE 1 ERROR?

As we have taken a significance level of 5%, hence the probability of Type 1 error is 5%

(II) THE POWER OF THE TEST?

The power of the test is 0.4592 i.e. 45.92%

<<R CODE>>

finding out Std deviation for the samples

SD1=sd(Old.Scheme)

SD2=sd(New.Scheme)

Calculating pooled SD

pooledSD=sqrt((SD1^2+SD2^2)/(2))

Calculating the power of the test

As the question states that the avg mean has to increase by 5000 in the new scheme to achieve breakeven, Hence 5000 needs to be added to 4000(mean diff of existing sample)

power.t.test(n=30,delta=9000,sd=pooledSD,power = NULL,sig.level = 0.05, type

"two.sample", alternative = "one.sided")

<<R OUTPUT>>

n = 30

delta = 9000

sd = 22332.11

sig.level = 0.05

power = 0.4592797

alternative = one.sided

(III) THE PROBABILITY OF TYPE 2 ERROR?

The Probability of type 2 error is (1-power) of the test

Power=1-0.4592

= 0.5408

=54.08%

E. WHAT SAMPLE SIZE WOULD MAKE THE PROBABILITIES OF TYPE 1 AND TYPE 2 ERRORS EQUAL?

The number of sample when the Type 1 is equal to Type 2 error is 133.948~134 Nos.

<<R CODE>>

calculating the no. of sample when the Type 1 and Type 2 error is equal

```
power.t.test(n=NULL,delta = 9000,sd = pooledSD,power = 0.95,sig.level = 0.05, type = "two.sample",alternative = "one.sided")
```

<<R OUTPUT>>

n = 133.948

delta = 9000

sd = 22332.11

sig.level = 0.05

power = 0.95

alternative = one.sided

PS: Refer Annexure I for entire program of R

ANNEXURE I

R-CODE

```
# Assignment on Titan Insurance company
#set working directory
setwd("F:/1.00 Home/4.00 PERSONAL/4.00 GreatLakes/1.00 1st Residency 18-22/R
Classes")
#Read the data from the source
mydata=read.csv("Titan.csv",header = T)
attach(mydata) # attaching mydata to R DB
mydata
# Calculating the paired t-test for the 2 samples
Paired.t.test=t.test(Old.Scheme, New.Scheme, paired = T, alternative = "less")
# Displaying the result
Paired.t.test
# finding out Std deviation for the samples
SD1=sd(Old.Scheme)
SD1
SD2=sd(New.Scheme)
SD2
# Calculating pooled SD
pooledSD=sqrt((SD1^2+SD2^2)/(2))
pooledSD
# Calculating the power of the test
# As the question states that the avg mean has to increase by 5000 in the new scheme to
achieve breakeven, Hence 5000 needs to be added to 4000(mean diff of existing sample)
power.t.test(n=30,delta=9000,sd=pooledSD,power = NULL,sig.level = 0.05, type =
"two.sample",alternative = "one.sided")
```

```

# Calculating the probability of Type 2 error (1-power)
1-0.459

# Probability of type 2 error = 0.5408

# calculating the no. of sample when the Type 1 and Type 2 error is equal
power.t.test(n=NULL,delta = 9000,sd = pooledSD,power = 0.95,sig.level = 0.05, type =
"two.sample",alternative = "one.sided")

# Histogram to check out the normality of old scheme
hist(Old.Scheme, main = "Old Scheme follows a Normal Distribution Approximately", col =
c(20:50), ylab = "Frequency", xlab = "Old Scheme")

# Histogram to check out the normality of New Scheme
hist(New.Scheme, main = "New Scheme follows a Normal Distribution Approximately", col
= c(30:60), ylab = "Frequency", xlab = "Old Scheme")

#Boxplot to check out the presence of outliers in Old scheme
boxplot(Old.Scheme,main="Old Scheme has No Outliers",col="light green")

#Boxplot to check out the presence of outliers in New Scheme
boxplot(New.Scheme,main="New Scheme has No Outliers",col="Sky Blue")

```


R-OUTPUT

```
> mydata
  Old.Scheme New.Scheme
1      57000      62000
2     103000     122000
3      59000      54000
4      75000      82000
5      84000      84000
6      73000      86000
7      35000      32000
8     110000     104000
9      44000      38000
10     82000     107000
11     67000      84000
12     64000      85000
13     78000      99000
14     53000      39000
15     41000      34000
16     39000      58000
17     80000      73000
18     87000      53000
19     73000      66000
20     65000      78000
21     28000      41000
22     62000      71000
23     49000      38000
24     84000      95000
25     63000      81000
26     77000      58000
27     67000      75000
28     101000     94000
29     91000     100000
30     50000      68000

> Paired.t.test=t.test(Old.Scheme, New.Scheme, paired = T, alternative = "les
s")
> Paired.t.test

      Paired t-test

data:  Old.Scheme and New.Scheme
t = -1.5559, df = 29, p-value = 0.06529
alternative hypothesis: true difference in means is less than 0
95 percent confidence interval:
      -Inf 368.1762
sample estimates:
mean of the differences
      -4000

> SD1=sd(Old.Scheme)
> SD1
[1] 20455.98
> SD2=sd(New.Scheme)
> SD2
[1] 24062.39
> pooledSD=sqrt((SD1^2+SD2^2)/(2))
```

```
> pooledSD
[1] 22332.11
> power.t.test(n=30,delta=9000,sd=pooledSD,power = NULL,sig.level = 0.05, type = "two.sample",alternative = "one.sided")
```

Two-sample t test power calculation

```
      n = 30
  delta = 9000
    sd = 22332.11
sig.level = 0.05
  power = 0.4592797
alternative = one.sided
```

NOTE: n is number in *each* group

```
> 1-0.459
[1] 0.541
> power.t.test(n=NULL,delta = 9000,sd = pooledSD,power = 0.95,sig.level = 0.05, type = "two.sample",alternative = "one.sided")
```

Two-sample t test power calculation

```
      n = 133.948
  delta = 9000
    sd = 22332.11
sig.level = 0.05
  power = 0.95
alternative = one.sided
```

NOTE: n is number in *each* group

```
> hist(Old.Scheme,
+      main = "Old Scheme follows a Normal Distribution Approximately",
+      col = c(20:50), ylab = "Frequency", xlab = "Old Scheme")
> hist(New.Scheme,
+      main = "New Scheme follows a Normal Distribution Approximately",
+      col = c(30:60), ylab = "Frequency", xlab = "Old Scheme")
> boxplot(Old.Scheme,main="Old Scheme has No Outliers",col="light green")
> boxplot(New.Scheme,main="New Scheme has No Outliers",col="Sky Blue")
```