Using Randomized Complete Block Design to find if there is significant difference in the various weight loss methods

Srikar

2021-09-08

Introduction:

<u>ANOVA</u> is a statistical test used to determine whether or not there is a significant d ifference between the means of treatments. The analysis of variance is the systemat ic algebraic procedure of decomposing (i.e., Partitioning) overall variation. If we find there is significant difference, we conduct a post-hoc test to check where the difference occurs.

Randomized Complete Block Design (RCBD) is the most common design of exper iments in many disciplines, including agriculture, engineering, medical, etc. In addition to the experimental error reducing ability, the design widens the generalization of the study findings. The repeated application of the treatments under investigation is known as <u>replication</u>. If the treatment is applied only once we have no means of knowing about the variations in the results of a treatments. Only when we repeat the application of the treatment several times, we can estimate the experimental error.

Objective:

To check whether the different weight loss methods result in the same amount of weight lost

Data Description:

The data set contains information on 20 people who undertook one of four diets (re ferred to as diet A, B, C and D). The aim of the study was to see which diet was be st for losing weight. The variable 'Method' describes the various dieting methods a nd WeightL shows how much weight the participants had lost in pounds. Replicati ons shows the repeated application of the methods.

Data Summary:

```
library(readx1)
data<- read_excel("C:/Users/Srikar/Desktop/SS/R/Sem 5/Design of Exp/Practical
6/dataset.xlsx")
head(data,6)</pre>
```

Method	Replication	WeightL
<chr></chr>	<dbl></dbl>	<dbl></dbl>
Α	1	21
Α	2	47
Α	3	27
Α	4	75
Α	5	30
В	1	88

summary(data)

Method	Replications	WeightL	
Length:20	Min. :1	Min. :17.00	
Class :characte	1st Qu.:2	1st Qu.:30.00	
Mode :charac	Median :3	Median :44.50	
	Mean :3	Mean :50.95	
	3rd Qu.:4	3rd Qu.:70.50	
	Max. :5	Max. :95.00	

We observe that the range of values are from 17 kgs to 95 kgs which are the minimum and maximu m values.

dim(data)

[1] 20 3

There are 20 observations with 3 variables: Method, replications and weight names(data)

[1] "Method" "Replications" "WeightL"

Hypothesis Statement:

Null Hypothesis (H 0): There is no significant difference in the different die meth ods ($\mu 1 = \mu 2 = \mu 3 = \mu 4$) where ($\mu 1, \mu 2, \mu 3, \mu 4$) are mean weight loss by different methods f ollowed

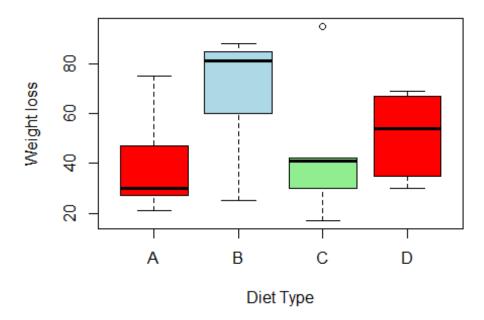
Alternative Hypothesis (H 1): At least two diet methods have significant differe nce. ($\mu 1 \neq \mu 2 \neq \mu 3 \neq \mu 4$)

#We keep the significance level as 5% or 0.05

Procedure

#1)Plotting boxplot

Types of weightloss methods and the weight lost



```
Table.A <- A.box$stats
colnames(Table.A)<-A.box$names
rownames(Table.A)<-c('min','lower quartile','median','upper quartile','max')
Table.A</pre>
```

		Box plot table		
	А	В	С	D
min	21	25	17	30
lower quartile	27	60	30	35
median	30	81	41	54
upper quartile	47	85	42	67
max	75	88	42	69

The above boxplot represents the above table. The minimum weight lost using any of the methods was found in method C. There is only one outlier in found in method C. The max imum weight loss was found in Method B. The median weight loss was highest in method B.

#2) Constructing the anova model

model=aov(data\$WeightL~data\$Method+data\$Replications, data = data)
summary(model)

		ANOVA Table for RCBD design			
	Df	Sum Sq	Mean Sq	F value	P-value
data\$Method	3	2196	732.1	1.162	0.357
data\$Replicatio	1	46	46.2	0.073	0.79
Residuals	15	9449	629.9		

We observe that the weight loss methods and the replications are not significant as their p -value is more than 0.05. Hence, we accept the null hypothesis. There is no significant difference in the different weight loss methods.

Conclusion:

All the weight loss methods lead to same amount of weight loss.