



# STATISTICAL ANALYSIS PROCEDURE OF RANDOMISED BLOCK DESIGN (RBD) FOR HORTICULTURAL DATA ON POTATO PLANT

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## Abstract

This paper was attempted to find out best treatment among 10 treatments by using Randomised Block Design analysis. RBD is most suitable for Agricultural field experiments when fertility gradient is in one direction. In this design, the treatments are assigned to the experimental plots in a random manner in blocks. This article considers the application, advantages and disadvantages and statistical analysis of randomized block designs.

**Key words :** RBD, ANOVA, CD.

## Introduction

In RBD, the entire experimental material is divided into number of homogeneous blocks. Each and every treatment should appear only once in each block.;

RBD follows all the basic principles of Experimental Designs

1. Replication
2. Randomisation
3. Local Control

RBD follows ANOVA-II way classification

### Assumptions of ANOVA

1. Independence of cases – this is an assumption of the model that simplifies the statistical analysis.
2. Normality – the distributions of the residuals are normal.
3. Equality (or homogeneity) of variances, called homoscedasticity.

### Advantages of RBD

1. This design has complete flexibility, *i.e.*, any

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number of replications can be included in this design

2. The amount of information got in RBD is more as compared CRD.
3. The statistical analysis is simple and easy. Even some observations are missing for certain treatments, the data can be analysed by the missing plot technique.

### Disadvantages of RBD

1. When the number of treatments is increased, the block size will increase.
2. If the block size is large maintaining homogeneity is difficult and hence when more number of treatments is present this design may not be suitable.

## Materials and Method

A varietal trial is conducted on Potato (*Solanum tuberosum*) with 10 varieties and each variety is replicated 3 times at College of Horticulture, Venkataramannagudem. Yield per plant in gms are as follows. Analyse the data and draw your conclusion

**Null Hypothesis for Treatments**

$H_0$  : There is no significant difference among mean

Treatments	R-I	R-II	R-III
T <sub>1</sub>	138.20	153.40	170.40
T <sub>2</sub>	173.80	187.00	208.20
T <sub>3</sub>	218.40	216.20	247.20
T <sub>4</sub>	250.20	273.40	247.00
T <sub>5</sub>	163.40	206.60	157.20
T <sub>6</sub>	207.80	173.00	237.40
T <sub>7</sub>	268.20	180.60	226.40
T <sub>8</sub>	193.20	178.20	126.60
T <sub>9</sub>	236.80	182.60	166.20
T <sub>10</sub>	266.80	184.60	185.40

Sol:

Treatments	Replications			Treatment Totals	Treatment Means
	R <sub>1</sub>	R <sub>2</sub>	R <sub>3</sub>		
T <sub>1</sub>	138.20	153.40	170.40	T <sub>1</sub> =462	$\alpha_1 = \frac{T_1}{r} = \frac{462}{3} = 154$
T <sub>2</sub>	173.80	187.00	208.20	T <sub>2</sub> =569	$\alpha_2 = \frac{T_2}{r} = \frac{569}{3} = 189.67$
T <sub>3</sub>	218.40	216.20	247.20	T <sub>3</sub> =681.80	$\alpha_3 = \frac{T_3}{r} = \frac{681.8}{3} = 227.27$
T <sub>4</sub>	250.20	273.40	247.00	T <sub>4</sub> =770.60	$\alpha_4 = \frac{T_4}{r} = \frac{770.60}{3} = 256.87$
T <sub>5</sub>	163.40	206.60	157.20	T <sub>5</sub> =527.20	$\alpha_5 = \frac{T_5}{r} = \frac{527.20}{3} = 175.73$
T <sub>6</sub>	207.80	173.00	237.40	T <sub>6</sub> =618.20	$\alpha_6 = \frac{T_6}{r} = \frac{618.20}{3} = 206.07$
T <sub>7</sub>	268.20	180.60	226.40	T <sub>7</sub> =675.20	$\alpha_7 = \frac{T_7}{r} = \frac{675.20}{3} = 225.07$
T <sub>8</sub>	193.20	178.20	126.60	T <sub>8</sub> =498	$\alpha_8 = \frac{T_8}{r} = \frac{498}{3} = 166$
T <sub>9</sub>	236.80	182.60	166.20	T <sub>9</sub> =585.60	$\alpha_9 = \frac{T_9}{r} = \frac{585.60}{3} = 195.20$
T <sub>10</sub>	266.80	184.60	185.40	T <sub>10</sub> =636.80	$\alpha_{10} = \frac{T_{10}}{r} = \frac{636.80}{3} = 212.27$
Replication Totals	R <sub>1</sub> =2116.80	R <sub>2</sub> =1935.60	R <sub>3</sub> =1972.00	GT=6024.40	
Replication	$\beta_1 = \frac{R_1}{k} =$	$\beta_2 = \frac{R_2}{k} =$	$\beta_3 = \frac{R_3}{k} =$		
Means	$\frac{2116.8}{10} = 211$	$\frac{1935.60}{10} = 193.56$	$\frac{1972}{10} = 197.21$		

The mathematica model for RBD is  $x_{ij} = \mu + \alpha_i + \beta_j + \epsilon_{ij}$ .

of varieties

$$H_0 : \alpha_1 = \alpha_2 = \dots = \alpha_{10}$$

**Alternative Hypothesis for Treatments**

$H_1$  : There is significant difference among mean of varieties

$$H_1 : \alpha_1 \neq \alpha_2 \neq \dots \neq \alpha_{10}$$

**Null Hypothesis for Blocks**

$H_0$  : There is no significant difference among mean of Replications / Blocks

$$H_0 : \beta_1 = \beta_2 = \beta_3$$

**Alternative Hypothesis for Treatments**

$H_1$  : There is significant difference among mean of Replications / Blocks

$$H_1 : \beta_1 \neq \beta_2 \neq \beta_3$$

k is no. Of varieties = 10  
r is no. Of replications = 3

$$GM = \frac{GT}{kr} = \frac{6024.40}{10 * 3} = \frac{6024.40}{30} = 200.8133$$

$$CF = \frac{GT^2}{kr} = \frac{6024.40^2}{10 * 3} = \frac{6024.40 * 6024.40}{30}$$

$$= \frac{36293395}{30} = 1209780$$

$$Total SS = \sum_{i=1}^k \sum_{j=1}^r x_{ij}^2 - CF = (x_{1,1}^2 + x_{1,2}^2 + \dots + x_{10,3}^2) - CF$$

$$= (138.2^2 + 153.4^2 + \dots + 185.4^2) - 1209780$$

$$= 1255244 - 1209780 = 45464.39$$

$$Trt SS = \frac{T_1^2 + T_2^2 + \dots + T_k^2}{r} - CF$$

$$= \frac{462^2 + 569^2 + \dots + 636.8^2}{3} - 1209780$$

$$= \frac{3708332}{3} - 1209780 = 1236111 - 1209780 = 26330.93$$

$$Rep/Block SS = \frac{R_1^2 + R_2^2 + \dots + R_r^2}{k} - CF$$

Prepare the following ANOVA Table

Sources of Variation	Degrees of freedom	Sum of Squares	Mean Sum of Squares	F-Cal	F-tab
Treatments	k-1=10-1=9	Trt SS=26330.93	$Trt MSS = \frac{Trt.SS}{k-1}$ $= \frac{26330.93}{9} = 2925.659$	$\frac{Trt.MSS}{Error MSS}$ $\frac{2925.659}{960.8862} = 3.0448*$	$F_{k-1,(k-1)(r-1)}$ $F_{9,18} = 2.4563 \text{ at } 5\%$ $F_{9,18} = 3.5971 \text{ at } 1\%$
Replications	r-1=3-1=2	Rep SS=1837.5147	$Rep SS MSS = \frac{Rep.SS}{r-1}$ $= \frac{1837.5147}{2} = 918.7573$	$\frac{Rept.MSS}{Error MSS}$ $\frac{918.7573}{960.8862} = 0.9562$	$F_{r-1,(k-1)(r-1)}$ $F_{2,18} = 3.5546 \text{ at } 5\%$ $F_{2,18} = 6.0129 \text{ at } 1\%$
Error	(k-1)(r-1)=9*2=18	Error SS= 17295.95	$Error MSS = \frac{Error.SS}{(k-1)(r-1)}$ $= \frac{17295.95}{18} = 960.8862$	-----	-----
Total	kr-1=10*3-1=29	Total SS=45464.39	-----	-----	-----

$$= \frac{2116.80^2 + 1935.6^2 + 1972^2}{10} - 1209780$$

$$= \frac{12116174}{10} - 1209780$$

$$= 1211617.4 - 1209780 = 1837.5147$$

Error SS = Total SS – Trt SS – Rep SS

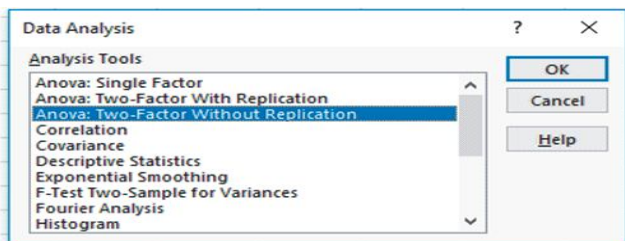
$$= 45464.39 - 26330.93 - 1837.5147 = 17295.95$$

**Analysis procedure of RBD in MS-Excel**

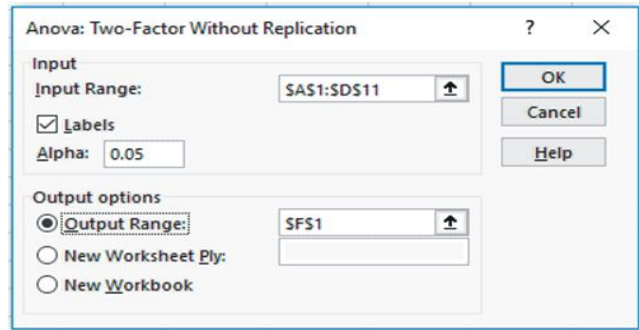
In MS-Excel sheet, enter data as below picture

	A	B	C	D	E
1	Treatments	R-I	R-II	R-III	
2	T1	138.2	153.4	170.4	
3	T2	173.8	187	208.2	
4	T3	218.4	216.2	247.2	
5	T4	250.2	273.4	247	
6	T5	163.4	206.6	157.2	
7	T6	207.8	173	237.4	
8	T7	268.2	180.6	226.4	
9	T8	193.2	178.2	126.6	
10	T9	236.8	182.6	166.2	
11	T10	266.8	184.6	185.4	

Select **Data** menu in Excel → choose **Data analysis**



choose **Anova: Two Factor Without Replication**  
→ press **OK**



Give Input Range: **\$A\$1:\$D\$11** → Put tick mark on **Labels** → Give **Alpha: 0.05** for 5% level of significance → Give **output Range \$F\$1** under Output options (i.e. Output will display from F1 cell) → press **OK**

	F	G	H	I	J	K	L
Anova: Two-Factor Without Replication							
	<b>SUMMARY</b>						
	Count	Sum	Average	Variance			
T1	3	462	154	259.48			
T2	3	569	189.6667	301.1733			
T3	3	681.8	227.2667	299.2133			
T4	3	770.6	256.8667	207.5733			
T5	3	527.2	175.7333	724.1733			
T6	3	618.2	206.0667	1039.093			
T7	3	675.2	225.0667	1919.773			
T8	3	498	166	1220.52			
T9	3	585.6	195.2	1365.16			
T10	3	636.8	212.2667	2230.573			
R-I	10	2116.8	211.68	1990.491			
R-II	10	1935.6	193.56	1082.212			
R-III	10	1972	197.2	1774.729			
<b>ANOVA</b>							
	Source of Variation	SS	df	MS	F	P-value	F crit
	Rows	26330.93	9	2925.659	3.04475	0.02122	2.456281
	Columns	1837.515	2	918.7573	0.956156	0.403048	3.554557
	Error	17295.95	18	960.8862			
	<b>Total</b>	<b>45464.39</b>	<b>29</b>				

In the ANOVA table of above picture, Rows means Treatments or Varieties and Columns means Blocks or replications.

**Results and Discussions**

Here F-cal (3.0488) value is > F-tab at 5% level of significance with  $F_{k-1,(k-1)(r-1)}^{F_{9,18}} = 2.4563$

So, we reject Null Hypothesis for treatments at 5% level of significance. *i.e.*, there is slightly significant difference among variety means.

**Standard Errors**

$$SEM = \sqrt{\frac{\text{Error } MSS}{r}} = \sqrt{\frac{960.8862}{3}} = 17.8968$$

$$SED = \sqrt{2} * SEM = \sqrt{2} * 17.8968 = 1.41 * 17.8968 = 25.3099$$

$$CD = SED * t_{\text{tab value at error } d.f} = 25.3099 * 2.1009 = 53.1741$$

**Bar notation**

T <sub>4</sub>	T <sub>3</sub>	T <sub>7</sub>	T <sub>10</sub>	T <sub>6</sub>	T <sub>9</sub>	T <sub>2</sub>	T <sub>5</sub>	T <sub>8</sub>	T <sub>1</sub>
256.87	227.27	225.07	212.27	206.07	195.20	189.67	175.73	166.00	154.00

We arrange the treatment means into decreasing order

1. Those pairs underscored are non-significant
2. Those pairs not scored are significant

Among 10 treatments, T<sub>4</sub> is best treatment

T<sub>4</sub> treatment is significantly different from T<sub>9</sub>, T<sub>2</sub>, T<sub>5</sub>, T<sub>8</sub>, T<sub>1</sub>

$$CV\% = \sqrt{\frac{EMSS}{GM}} * 100 = \sqrt{\frac{960.8862}{200.81}} * 100 =$$

$$\frac{30.9982}{200.81} * 100 = 0.1544 * 100 = 15.44\%$$

Less CV% indicates more consistency of data.

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