

The **augmentedRCBD** package: A brief introduction

J. Aravind¹ , Mukesh Sankar S.²

Division of Germplasm Conservation, ICAR-National Bureau of Plant Genetic Resources, New Delhi
Division of Genetics, ICAR-Indian Agricultural Research Institute, New Delhi

2018-06-18

Contents

Overview	1
Installation	1
Testing	1
Citing augmentedRCBD	5
Session Info	5
References	6

Overview

Installation

The package can be installed using the following functions:

```
# Install from CRAN
install.packages('augmentedRCBD', dependencies=TRUE)

# Install development version from Github
devtools::install_github("aravind-j/augmentedRCBD")
```

Then the package can be loaded using the function

```
library(augmentedRCBD)
```

Testing

```
# Example data
blk <- c(rep(1,7),rep(2,6),rep(3,7))
trt <- c(1, 2, 3, 4, 7, 11, 12, 1, 2, 3, 4, 5, 9, 1, 2, 3, 4, 8, 6, 10)
y1 <- c(92, 79, 87, 81, 96, 89, 82, 79, 81, 81, 91, 79, 78, 83, 77, 78, 78,
70, 75, 74)
y2 <- c(258, 224, 238, 278, 347, 300, 289, 260, 220, 237, 227, 281, 311, 250,
240, 268, 287, 226, 395, 450)
data <- data.frame(blk, trt, y1, y2)
# Convert block and treatment to factors
data$blk <- as.factor(data$blk)
data$trt <- as.factor(data$trt)
# Results for variable y1
out1 <- augmentedRCBD(data$blk, data$trt, data$y1, method = "lsd",
alpha = 0.05, group = TRUE, console = TRUE)
```

Augmented design details

=====

```

Number of blocks      "3"
Number of treatments  "12"
Number of check treatments "4"
Number of test treatments "8"
Check treatments      "1, 2, 3, 4"

```

ANOVA, Treatment Adjusted

=====

	Df	Sum Sq	Mean Sq	F value	Pr(>F)
Block (ignoring Treatments)	2	360.1	180.04	6.675	0.0298 *
Treatment (eliminating Blocks)	11	285.1	25.92	0.961	0.5499
Treatment: Check	3	52.9	17.64	0.654	0.6092
Treatment: Test and Test vs. Check	8	232.2	29.02	1.076	0.4779
Residuals	6	161.8	26.97		

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

ANOVA, Block Adjusted

=====

	Df	Sum Sq	Mean Sq	F value	Pr(>F)
Treatment (ignoring Blocks)	11	575.7	52.33	1.940	0.215
Treatment: Check	3	52.9	17.64	0.654	0.609
Treatment: Test	7	505.9	72.27	2.679	0.125
Treatment: Test vs. Check	1	16.9	16.87	0.626	0.459
Block (eliminating Treatments)	2	69.5	34.75	1.288	0.342
Residuals	6	161.8	26.97		

Treatment means

=====

	Treatment	Block	Means	SE	r	Min	Max	Adjusted Means
1	1	1	84.66667	3.844188	3	79	92	84.66667
2	10	3	74.00000	NA	1	74	74	77.25000
3	11	1	89.00000	NA	1	89	89	86.50000
4	12	1	82.00000	NA	1	82	82	79.50000
5	2	2	79.00000	1.154701	3	77	81	79.00000
6	3	3	82.00000	2.645751	3	78	87	82.00000
7	4	4	83.33333	3.929942	3	78	91	83.33333
8	5	2	79.00000	NA	1	79	79	78.25000
9	6	3	75.00000	NA	1	75	75	78.25000
10	7	1	96.00000	NA	1	96	96	93.50000
11	8	3	70.00000	NA	1	70	70	73.25000
12	9	2	78.00000	NA	1	78	78	77.25000

Coefficient of variation

=====

6.372367

Overall adjusted mean

=====

81.0625

Standard errors

=====

	Std. Error of Diff.	CD (5%)
Control Treatment Means	4.240458	10.37603
Two Test Treatments (Same Block)	7.344688	17.97180
Two Test Treatments (Different Blocks)	8.211611	20.09309
A Test Treatment and a Control Treatment	6.360687	15.56404

Treatment groups

=====

Method : lsd

	Treatment	Adjusted Means	SE	df	lower.CL	upper.CL	Group
8	8	73.25000	5.609598	6	59.52381	86.97619	1
9	9	77.25000	5.609598	6	63.52381	90.97619	12
10	10	77.25000	5.609598	6	63.52381	90.97619	12
5	5	78.25000	5.609598	6	64.52381	91.97619	12
6	6	78.25000	5.609598	6	64.52381	91.97619	12
2	2	79.00000	2.998456	6	71.66304	86.33696	12
12	12	79.50000	5.609598	6	65.77381	93.22619	12
3	3	82.00000	2.998456	6	74.66304	89.33696	12
4	4	83.33333	2.998456	6	75.99637	90.67029	12
1	1	84.66667	2.998456	6	77.32971	92.00363	12
11	11	86.50000	5.609598	6	72.77381	100.22619	12
7	7	93.50000	5.609598	6	79.77381	107.22619	2

Results for variable y2

```
out2 <- augmentedRCBD(data$blk, data$trt, data$y1, method = "lsd",
alpha = 0.05, group = TRUE, console = TRUE)
```

Augmented design details

=====

Number of blocks	"3"
Number of treatments	"12"
Number of check treatments	"4"
Number of test treatments	"8"
Check treatments	"1, 2, 3, 4"

ANOVA, Treatment Adjusted

=====

	Df	Sum Sq	Mean Sq	F value	Pr(>F)
Block (ignoring Treatments)	2	360.1	180.04	6.675	0.0298 *
Treatment (eliminating Blocks)	11	285.1	25.92	0.961	0.5499
Treatment: Check	3	52.9	17.64	0.654	0.6092
Treatment: Test and Test vs. Check	8	232.2	29.02	1.076	0.4779
Residuals	6	161.8	26.97		

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

ANOVA, Block Adjusted

=====

	Df	Sum Sq	Mean Sq	F value	Pr(>F)
Treatment (ignoring Blocks)	11	575.7	52.33	1.940	0.215
Treatment: Check	3	52.9	17.64	0.654	0.609
Treatment: Test	7	505.9	72.27	2.679	0.125
Treatment: Test vs. Check	1	16.9	16.87	0.626	0.459
Block (eliminating Treatments)	2	69.5	34.75	1.288	0.342
Residuals	6	161.8	26.97		

Treatment means

=====

	Treatment	Block	Means	SE	r	Min	Max	Adjusted Means
1	1		84.66667	3.844188	3	79	92	84.66667
2	10	3	74.00000	NA	1	74	74	77.25000
3	11	1	89.00000	NA	1	89	89	86.50000
4	12	1	82.00000	NA	1	82	82	79.50000
5	2		79.00000	1.154701	3	77	81	79.00000
6	3		82.00000	2.645751	3	78	87	82.00000
7	4		83.33333	3.929942	3	78	91	83.33333
8	5	2	79.00000	NA	1	79	79	78.25000
9	6	3	75.00000	NA	1	75	75	78.25000
10	7	1	96.00000	NA	1	96	96	93.50000
11	8	3	70.00000	NA	1	70	70	73.25000
12	9	2	78.00000	NA	1	78	78	77.25000

Coefficient of variation

=====

6.372367

Overall adjusted mean

=====

81.0625

Standard errors

=====

	Std. Error of Diff.	CD (5%)
Control Treatment Means	4.240458	10.37603
Two Test Treatments (Same Block)	7.344688	17.97180
Two Test Treatments (Different Blocks)	8.211611	20.09309
A Test Treatment and a Control Treatment	6.360687	15.56404

Treatment groups

=====

Method : lsd

	Treatment	Adjusted Means	SE	df	lower.CL	upper.CL	Group
8	8	73.25000	5.609598	6	59.52381	86.97619	1
9	9	77.25000	5.609598	6	63.52381	90.97619	12
10	10	77.25000	5.609598	6	63.52381	90.97619	12
5	5	78.25000	5.609598	6	64.52381	91.97619	12
6	6	78.25000	5.609598	6	64.52381	91.97619	12
2	2	79.00000	2.998456	6	71.66304	86.33696	12
12	12	79.50000	5.609598	6	65.77381	93.22619	12
3	3	82.00000	2.998456	6	74.66304	89.33696	12

4	4	83.33333	2.998456	6	75.99637	90.67029	12
1	1	84.66667	2.998456	6	77.32971	92.00363	12
11	11	86.50000	5.609598	6	72.77381	100.22619	12
7	7	93.50000	5.609598	6	79.77381	107.22619	2

Citing *augmentedRCBD*

Warning in citation("augmentedRCBD"): no date field in DESCRIPTION file of package 'augmentedRCBD'

Warning in citation("augmentedRCBD"): could not determine year for 'augmentedRCBD' from package DESCRIPTION file

To cite package 'augmentedRCBD' in publications use:

J. Aravind, Mukesh Sankar S., Dhammaprakash Pandhari Wankhede and Vikender Kaur (NA). *augmentedRCBD: Analysis of Augmented Randomised Complete Block Design*. R package version 0.0.0.9000. <https://github.com/aravind-j/augmentedRCBD>

A BibTeX entry for LaTeX users is

```
@Manual{,
  title = {augmentedRCBD: Analysis of Augmented Randomised Complete Block Design},
  author = {{J. Aravind} and {Mukesh Sankar S.} and {Dhammaprakash Pandhari Wankhede} and {Vikender Kaur}},
  note = {R package version 0.0.0.9000},
  url = {https://github.com/aravind-j/augmentedRCBD},
  year = {2018},
}
```

Session Info

```
sessionInfo()
```

```
R version 3.5.0 (2018-04-23)
Platform: x86_64-w64-mingw32/x64 (64-bit)
Running under: Windows >= 8 x64 (build 9200)
```

```
Matrix products: default
```

```
locale:
```

```
[1] LC_COLLATE=English_India.1252 LC_CTYPE=English_India.1252
[3] LC_MONETARY=English_India.1252 LC_NUMERIC=C
[5] LC_TIME=English_India.1252
```

```
attached base packages:
```

```
[1] stats      graphics  grDevices  utils      datasets  methods    base
```

```
other attached packages:
```

```
[1] augmentedRCBD_0.0.0.9000
```

```
loaded via a namespace (and not attached):
```

```
[1] zoo_1.8-1      reshape2_1.4.3  splines_3.5.0
```

[4] lattice_0.20-35	colorspace_1.3-2	htmltools_0.3.6
[7] yaml_2.1.18	base64enc_0.1-3	survival_2.42-3
[10] rlang_0.2.0	R.oo_1.22.0	pillar_1.2.2
[13] glue_1.2.0	withr_2.1.2	R.utils_2.6.0
[16] gdtools_0.1.7	bindrcpp_0.2.2	uuid_0.1-2
[19] emmeans_1.1.3	multcomp_1.4-8	bindr_0.1.1
[22] plyr_1.8.4	multcompView_0.1-7	stringr_1.3.0
[25] munsell_0.4.3	commonmark_1.5	gtable_0.2.0
[28] R.methodsS3_1.7.1	moments_0.14	zip_1.0.0
[31] devtools_1.13.5	mvtnorm_1.0-7	codetools_0.2-15
[34] coda_0.19-1	memoise_1.1.0	evaluate_0.10.1
[37] knitr_1.20	gbRd_0.4-11	TH.data_1.0-8
[40] Rcpp_0.12.16	xtable_1.8-2	scales_0.5.0
[43] backports_1.1.2	ggplot2_2.2.1	digest_0.6.15
[46] stringi_1.1.7	dplyr_0.7.4	grid_3.5.0
[49] rprojroot_1.3-2	bibtex_0.4.2	Rdpack_0.7-0
[52] tools_3.5.0	sandwich_2.4-0	magrittr_1.5
[55] lazyeval_0.2.1	tibble_1.4.2	pkgconfig_2.0.1
[58] MASS_7.3-49	Matrix_1.2-14	xml2_1.2.0
[61] estimability_1.3	assertthat_0.2.0	rmarkdown_1.9
[64] roxygen2_6.0.1	officer_0.3.0	flextable_0.4.4
[67] R6_2.2.2	nlme_3.1-137	compiler_3.5.0

References