

# Package ‘midrangeMCP’

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**Type** Package

**Title** Multiples Comparisons Procedures Based on Studentized Midrange and Range Distributions

**Version** 3.1.1

**Description** Apply tests of multiple comparisons based on studentized 'midrange' and 'range' distributions. The tests are: Tukey Midrange ('TM' test), Student-Newman-Keuls Midrange ('SNKM' test), Means Grouping Midrange ('MGM' test) and Means Grouping Range ('MGR' test). The first two tests were published by Batista and Ferreira (2020) <[doi:10.1590/1413-7054202044008020](https://doi.org/10.1590/1413-7054202044008020)>. The last two are being published.

**Imports** stats, utils, graphics, grDevices, SMR, writexl, xtable, tcltk

**Suggests** tkrplot, testthat, knitr, rmarkdown, covr

**License** GPL (>= 2)

**URL** <https://bendeivide.github.io/midrangeMCP/>,  
<https://github.com/bendeivide/midrangeMCP>

**BugReports** <https://github.com/bendeivide/midrangeMCP/issues>

**LazyData** TRUE

**RoxygenNote** 7.1.1

**NeedsCompilation** no

**Author** Ben Deivide [aut, cre] (<<https://orcid.org/0000-0001-7019-8794>>),  
Daniel Furtado [aut] (<<https://orcid.org/0000-0002-4371-5239>>),  
Diego Arthur [ctb]

**Maintainer** Ben Deivide <[ben.deivide@ufsj.edu.br](mailto:ben.deivide@ufsj.edu.br)>

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guimidrangeMCP	<i>Graphical User Interface for midrangeMCP function</i>
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### Description

guimidrangeMCP A Graphical User Interface (GUI) for function that returns the MGM, MGR, SNKM and TM tests

### Usage

```
guimidrangeMCP(gui = TRUE)
```

### Arguments

gui Logical argument, TRUE or FALSE. The default is TRUE

### Value

guimidrangeMCP presents a GUI for the results of the four multiple comparison procedures MGM, MGR, SNKM and TM tests. In addition, the GUI returns a graph of the results, as well as the export of these results to three types of file extension and latex code.

### Examples

```
# Loading package
library(midrangeMCP)
if (interactive()) {
  guimidrangeMCP(gui = FALSE)
}
```

MRbarplot

*Plotting the result of the multiple comparison procedures***Description**

MRbarplot creates a bar plot with vertical or horizontal bars to compare the mean treatments by the tests: means grouping based on midrange, means grouping based on range, Student-Newman-Keuls and Tukey based on midrange.

**Usage**

```
MRbarplot(x, MCP = "all", col = grDevices::heat.colors(10), horiz = FALSE, ...)
```

**Arguments**

x	An object of the MRtest function
MCP	Allows choosing the multiple comparison test. The <i>default</i> is "all". This option will perform all tests available in the MRtest object.
col	A specification for the plotting color. The <i>default</i> is <code>heat.colors(10)</code> .
horiz	a logical value. If FALSE, the bars are drawn vertically with the first bar to the left. If TRUE, the bars are drawn horizontally with the first at the bottom.
...	Parameters of the <a href="#">barplot</a> function

**Details**

The MCP argument allows choosing several tests of multiple comparisons from the MRtest object. For plots in papers, use `col = gray.colors(10)`. For details, see [colors](#) function.

**Value**

MRbarplot return the bar plot of the tests chosen ("MGM", "MGR", "SNKM" and "TM") to evaluate the treatment means.

**Examples**

```
# Simulated data (completely randomized design)

rv <- c(100.08, 105.66, 97.64, 100.11, 102.60, 121.29, 100.80,
        99.11, 104.43, 122.18, 119.49, 124.37, 123.19, 134.16,
        125.67, 128.88, 148.07, 134.27, 151.53, 127.31)

# Treatments
treat <- factor(rep(LETTERS[1:5], each = 4))

# Anova
res <- aov(rv~treat)
```

```

# Loading the midrangeMCP package
library(midrangeMCP)

# Choosing tests
results <- MRtest(y = res, trt = "treat", alpha = 0.05,
                 main = "Multiple Comparison Procedures",
                 MCP = c("MGM", "TM"))

MRbarplot(results, MCP = "all") # It will be shown two
                                # graphs. First, for the
                                # results of \code{'MGM'}
                                # and the second for the
                                # results of \code{'TM'}.

MRbarplot(results, MCP = "MGM") # It will be shown
                                # only the graph
                                # for the result of
                                # \code{'MGM'}

# Plot for papers
MRbarplot(results, MCP = "all", col = gray.colors(10))

```

---

MRtest

*Multiple comparison procedures to the means of a factor using the studentized range and midrange distributions.*

---

## Description

MRtest applies the Means grouping based on midrange, Means Grouping based on Range, Student-Newman-Keuls based on midrange and Tukey based on midrange tests. These are new tests for multiple comparisons proposed by the authors (BATISTA, 2016), that are being published.

## Usage

```

MRtest(
  y,
  trt = NULL,
  dferror = NULL,
  mserror = NULL,
  replication = NULL,
  alpha = 0.05,
  main = NULL,
  MCP = "all",
  ismean = FALSE
)

```

**Arguments**

y	Model (aov or lm), numeric vector containing the response variable or the mean of the treatments.
trt	Constant (y = model) or a vector containing the treatments.
dferror	Degrees of freedom of the Mean Square Error.
mserror	Mean Square Error.
replication	Number de repetitions of the treatments in the experiment. For unbalanced data should be informed the harmonic mean of repetitions. This argument should be informed only if ismean = TRUE.
alpha	Significant level. The default is $\alpha = 0.05$ .
main	Title of the analysis.
MCP	Allows choosing the multiple comparison test; the <i>default</i> is "all". This option will go perform all tests. However, the options are: the Means grouping based on midrange test ("MGM"), Means Grouping based on Range test ("MGR"), the Student-Newman-Keuls based on midrange test ("SNKM") and the Tukey based on midrange test ("TM").
ismean	Logic. If FALSE (default), the y argument represents a model (aov or lm) or a numeric vector containing the response variable. If TRUE the y argument represents the mean of treatments.

**Details**

The MCP argument allows you to choose various tests of multiple comparisons at once. For example, `MCP = c("MGM", "MGR")`, and so on.

**Value**

MRtest returns the print of a list of results. First, the summary of y. Second, the statistics of the test chosen. And finally, the mean group results for each test. If MRtest function is stored in an object, the results will be printed and also stored in the object.

**Examples**

```
# Simulated data (completely randomized design)

# Response variable
rv <- c(100.08, 105.66, 97.64, 100.11, 102.60, 121.29, 100.80,
        99.11, 104.43, 122.18, 119.49, 124.37, 123.19, 134.16,
        125.67, 128.88, 148.07, 134.27, 151.53, 127.31)

# Treatments
treat <- factor(rep(LETTERS[1:5], each = 4))

# Anova
res <- anova(aov(rv~treat))
DFerror <- res$Df[2]
MSerror <- res$`Mean Sq`[2]
```

```

# Loading the midrangeMCP package
library(midrangeMCP)

# applying the tests
results <- MRtest(y = rv,
                 trt = treat,
                 dferror = DFerror,
                 mserror = MSerror,
                 alpha = 0.05,
                 main = "Multiple Comparison Procedure: MGM test",
                 MCP = c("MGM"))

# Other option for the MCP argument is "all". All tests are used.

results$Groups      # Results of the tests
results$Statistics  # Main arguments of the tests
results$Summary     # Summary of the response variable

# Using the y argument as aov or lm model
res <- aov(rv~treat)

MRtest(y = res, trt = "treat", alpha = 0.05,
       main = "Multiple Comparison Procedure: MGM test",
       MCP = c("MGM"))

# For unbalanced data: It will be used the harmonic mean of
#                       the number of experiment replicates

# Using the previous example
rv <- rv[-1]
treat <- treat[-1]

res <- lm(rv~treat) # Linear model

# Multiple comparison procedure: MGR test
MRtest(y = res, trt = "treat", alpha = 0.05,
       main = "Multiple Comparison Procedure: MGR test",
       MCP = c("MGR"))

# Assuming that the available data are the averages
# of the treatments and the analysis of variance

# Analysis of Variance Table

# Response: rv
#           Df Sum Sq Mean Sq F value    Pr(>F)
# treat      4 4135.2 1033.80  14.669 4.562e-05 ***
# Residuals 15 1057.1   70.47

mean.treat <- c(100.87, 105.95, 117.62, 127.97, 140.30)
treat <- factor(LETTERS[1:5])
DFerror <- 15

```

```

MSerror <- 70.47488
replic <- 4

MRtest(y = mean.treat,
       trt = treat,
       dferror = DFerror,
       mserror = MSerror,
       replication = replic,
       alpha = 0.05,
       main = "Multiple Comparison Procedure: MGM test",
       MCP = c("MGM"),
       ismean = TRUE)

```

---

MRwrite

*Export the results of the MRtest function.*


---

## Description

The x object from a MRtest is written to file arguments.

## Usage

```
MRwrite(x, MCP = "all", extension = "csv", dataMR = "all")
```

## Arguments

x	object from the MRtest functions.
MCP	Allows choosing the multiple comparison test. The <i>default</i> is "all". This option will go perform all tests from the MRtest object.
extension	Type of format of the file. Four options "csv", "txt" "xlsx" and "latex". The <i>default</i> is "csv".
dataMR	Allows to choose the results to be written. Three options are available: "groups", "summary" or "all". The option "groups" writes the treatment mean groups evaluated by the chosen test in the MCP argument. The "summary" writes the descriptive statistics of the response variable. The options "all" should be chosen for both results.

## Details

Note that the choice of the tests in the MRwrite function must be in accordance with the tests chosen in the x argument.

## Value

MRwrite writes the most important results for the chosen tests in the MCP argument.

**Examples**

```
# Simulated data (completely randomized design)

rv <- c(100.08, 105.66, 97.64, 100.11, 102.60, 121.29, 100.80,
        99.11, 104.43, 122.18, 119.49, 124.37, 123.19, 134.16,
        125.67, 128.88, 148.07, 134.27, 151.53, 127.31)

# Treatments
treat <- factor(rep(LETTERS[1:5], each = 4))

# Anova
res <- aov(rv~treat)

# Loading the midrangeMCP package
library(midrangeMCP)

# Choosing any tests
results <- MRtest(y = res, trt = "treat", alpha = 0.05,
                  main = "Multiple Comparison Procedures",
                  MCP = c("MGM", "TM"))

#Export file in latex (Output in Console)
MRwrite(results, MCP = "all", extension = "latex", dataMR = "all")

#Observation: The MRwrite function export
#              only one extension at a time
```



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