

# Rcmdr Plug-In Package for the EZR (Easy R especially for medical statistics)

## Description

This package provides an R Commander plug-in EZR (Easy R), which adds a variety of statistical functions, including survival analyses, ROC analyses, metaanalyses, sample size calculation, and so on, to the R commander. EZR enables point-and-click easy access to a variety of statistical functions as shown below, especially for medical statistics. A report that introduced the investigation of EZR was published in *Bone Marrow Transplantation* (Nature Publishing Group) as an Open article. This report can be used as a simple manual. It can be freely downloaded from the journal website (URL:

<http://www.nature.com/bmt/journal/vaop/ncurrent/pdf/bmt2012244a.pdf>). A complete manual of EZR is available only in Japanese (Chugai Igakusha, ISBN: 978-4-498-10901-8, URL: <http://www.chugaiigaku.jp/item/detail.php?id=1660> or Nankodo, ISBN: 978-4-524-26158-1, URL: <http://www.nankodo.co.jp/g/g9784524261581/>)

## Details

Package: RcmdrPlugin.EZR

Type: Package

Version: 1.31

Date: 2015-12-01

License: GPL (>= 2)

## Installation of EZR

See <http://www.jichi.ac.jp/saitama-sct/SaitamaHP.files/statmedEN.html> for detailed information. Briefly, EZR is platform-independent. RcmdrPlugin.EZR should be installed with R commander (Rcmdr) and the other packages that Rcmdr or EZR depends on. Packages required by EZR can be easily installed at the same time with the installation of EZR by copying and pasting the following command to the R console window after ">".

```
install.packages(pkgs="RcmdrPlugin.EZR", dependencies=TRUE)
```

After installation, R commander can be started by the command, `library(Rcmdr)` from the R console. Packages required by Rcmdr are installed at the first start-up of R commander. EZR can be loaded by selecting RcmdrPlugin.EZR from the "Tools" > "Load Rcmdr plug-in(s)" menu. Answer "Yes" to "Restart now?".

On Windows, EZR plugin package will be loaded when R commander is started, if the following sentence is added to the Rprofile.site file in etc folder in the R folder (C:\Program Files\R\R-X.XX.X\etc).

```
options(Rcmdr=list(plugins="RcmdrPlugin.EZR"))
```

In addition, if the following phrase is added to the command in "Target:" column on the

"Shortcut" tab of the "Property" of "R" shortcut on the desktop (which can be opened by right-clicking on the shortcut), R commander will start at the same time with launching R.

```
R_DEFAULT_PACKAGES="Rcmdr"
```

Therefore, if these two options were added, EZR can be started just by double-clicking on the "R" shortcut on the desktop.

In OS X, these can be done by opening the Terminal.app located in the Utilities folder, followed by copying and pasting the following command.

```
echo "options(Rcmdr=list(plugins='RcmdrPlugin.EZR'))" >> ~/.Rprofile
echo "library(Rcmdr)" >> ~/.Rprofile
echo "local({" >> ~/.Rprofile
echo "old <- getOption('defaultPackages')" >> ~/.Rprofile
echo "options(defaultPackages = c(old, 'Rcmdr'))" >> ~/.Rprofile
echo "})" >> ~/.Rprofile
```

The default data folder of Windows EZR can be changed by right-clicking on this "R" shortcut on the desktop, selecting "Properties", and replacing the folder name in the "Start in:" column on the "Shortcut" tab.

## EZR statistical functions

EZR includes following statistical functions.

### *For discrete variables*

- Frequency distributions/cr Confidence interval for a proportion
- One sample proportion test
- Confidence interval for a difference between two proportions
- Confidence interval for a ratio of two proportions
- Compare two proportions (Fisher's exact test and Chi-square test)
- Compare proportions of two paired samples (McNemar test)
- Compare proportions of more than two paired samples (Cochran Q test)
- Cochran-Armitage test for trend in proportions
- Logistic regression

### *For continuous variables*

- Numerical summaries
- Smirnov-Grubbs test for outliers
- Kolmogorov-Smirnov test for normal distribution
- Confidence interval for a mean
- Single-sample t-test
- Two-variances F-test
- Two-sample t-test
- Paired t-test
- Bartlett's test
- One-way ANOVA
- Repeated-measures ANOVA

Multi-way ANOVA  
ANCOVA  
Test for Pearson's correlation  
Linear regression

*For nonparametric tests for continuous variables*

Mann-Whitney U test  
Wilcoxon's signed rank test  
Kruskal-Wallis test  
Friedman test  
Jonckheere-Terpstra test  
Spearman's rank correlation test

*For survival analysis*

Kaplan-Meier survival curve and logrank test  
Logrank trend test  
Cox proportional hazard regression  
Cox proportional hazard regression with time-dependent covariate  
Cumulative incidence of competing events and Gray test  
Fine-Gray proportional hazard regression for competing events

*For diagnostic test analysis*

Accuracy of qualitative test  
Kappa statistics for agreement of two tests  
Compute positive and negative predictive values  
ROC curve analysis for quantitative test  
Compare two ROC curves  
Cronbach's alpha coefficient for reliability

*For matched-pair analysis*

Extract matched controls (This function relies on optmatch package and is limited to academic use.)  
Mantel-Haenszel test for matched proportions  
Conditional logistic regression for matched-pair analysis  
Stratified Cox proportional hazard regression for matched-pair analysis

*For meta-analysis and meta-regression test*

Meta-analysis and meta-regression test for proportions  
Meta-analysis and meta-regression test for means  
Meta-analysis and meta-regression test for hazard ratios

*For sample size and power calculation*

Calculate sample size from control and desired response rates  
Calculate sample size from proportion and confidence interval  
Calculate sample size or power for comparison with specified proportion  
Calculate sample size or power for comparison between two proportions  
Calculate sample size for non-inferiority trial of two proportions

Calculate sample size from standard deviation and confidence interval  
Calculate sample size or power for comparison between two means  
Calculate sample size for non-inferiority trial of two means  
Calculate sample size or power for comparison between two paired means  
Calculate sample size or power for comparison between two survival curves  
Calculate sample size for non-inferiority trial of two survival curves

*For drawing graphs*

Bar graph(Frequencies)  
Pie chart(Frequencies)  
Stem-and-leaf display  
Histogram  
QQ plot  
Bar graph(Means)  
Line graph(Means)  
Line graph(Repeated measures)  
Boxplot  
Dot chart  
Ordered chart  
Scatterplot  
Scatterplot matrix  
Adjusted survival curve  
Stacked cumulative incidences

*Statistical functions from original R commander*

Principal-components analysis  
Factor analysis  
k-means cluster analysis  
Hierarchical cluster analysis  
Summarize hierarchical clustering  
Add hierarchical clustering to data set  
Linear hypothesis  
Variance-inflation factor  
Breusch-Pagan test for heteroscedasticity  
Durbin-Watson test for autocorrelation  
RESET test for nonlinearity  
Bonferroni outlier test  
Basic diagnostic plots  
Residual quantile-comparison plot  
Component+residual plots  
Added-variable plots  
Influence plot  
Effect plots

## **Basic operations in EZR**

These EZR functions can be started by point-and-click access using the items on the menu bar.  
See `EZRdialogs` for details. R commander automatically creates and executes corresponding R commands that appear in the "Script window". Results are shown in the "Output window". If any

errors or warnings are noted, messages will appear in the "Message window". The created commands can be saved by selecting "File" > "Save script" on the menu bar. The output can be saved by selecting "File" > "Save output". By saving the commands, users can reproduce the analyses and can also share the procedure with the other investigators.

The following EZR functions can be executed by typing the commands in the "Script window" and clicking on the "Submit" button.

Following functions are built to create a formatted table for presentation.

`w.twoway(table, filename)` function copies the results of two-way table analyses to the clipboard or text file.

`w.ttest(table, filename)` function copies the results of t-test to the clipboard or text file.

`w.survival(table, filename)` function copies the results of survival analyses to the clipboard or text file.

`w.ci(table, filename)` function copies the results of cumulative incidence analyses to the clipboard or text file.

`w.multi(table, filename)` function copies the results of multivariate regression analyses to the clipboard or text file.

"table" can be omitted except for logistic regression analysis and Fine & Gray proportional hazard regression analysis, in which "odds" and "crr.table" should be specified for "table"(default is "cox.table" to copy the results of Cox proportional hazard regression analysis).

If "filename" is omitted, the formatted table will be copied to the clipboard, which can be pasted into a spreadsheet.

`Mantel.Byar()` function is for Mantel-Byar test, which should be performed after executing "Cox proportional hazard modeling with time-dependent covariate".

## Translations

EZR comes with translations from English into Japanese.

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## References

Its complete manual is available only in Japanese (Chugai Igakusha, ISBN: 978-4-498-10901-8, URL: <http://www.chugaiigaku.jp/item/detail.php?id=1660> or Nankodo, ISBN: 978-4-524-26158-1, URL: <http://www.nankodo.co.jp/g/g9784524261581/>), but a report that introduced the investigation of EZR was published in *Bone Marrow Transplantation* (Nature Publishing Group) as an Open article. This report can be used as a simple manual. It can be freely downloaded from the journal website as shown below.

Yoshinobu Kanda (2013). Investigation of the freely available easy-to-use software EZR for medical statistics. *Bone Marrow Transplantation* (Open article, URL: <http://www.nature.com/bmt/journal/vaop/ncurrent/pdf/bmt2012244a.pdf>).

EZR web site: Division of Hematology, Saitama Medical Center, Jichi Medical University. URL: <http://www.jichi.ac.jp/saitama-sct/SaitamaHP.files/statmedEN.html>