

# Package ‘olsrr’

October 14, 2022

**Type** Package

**Title** Tools for Building OLS Regression Models

**Version** 0.5.3

**Description** Tools designed to make it easier for users, particularly beginner/intermediate R users to build ordinary least squares regression models. Includes comprehensive regression output, heteroskedasticity tests, collinearity diagnostics, residual diagnostics, measures of influence, model fit assessment and variable selection procedures.

**Depends** R(>= 3.3)

**Imports** car, data.table, ggplot2, goftest, graphics, gridExtra,  
nortest, Rcpp, stats, utils

**Suggests** covr, descriptr, knitr, rmarkdown, testthat, vdiff, xplorerr

**License** MIT + file LICENSE

**URL** <https://olsrr.rsquaredacademy.com/>,  
<https://github.com/rsquaredacademy/olsrr>

**BugReports** <https://github.com/rsquaredacademy/olsrr/issues>

**Encoding** UTF-8

**LazyData** true

**VignetteBuilder** knitr

**RoxygenNote** 6.1.1

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auto *Test Data Set*

**Description**

Test Data Set

**Usage**

auto

**Format**

An object of class `tbl_df` (inherits from `tbl`, `data.frame`) with 74 rows and 11 columns.

---

cement	<i>Test Data Set</i>
--------	----------------------

---

**Description**

Test Data Set

**Usage**

cement

**Format**

An object of class `data.frame` with 13 rows and 6 columns.

---

fitness	<i>Test Data Set</i>
---------	----------------------

---

**Description**

Test Data Set

**Usage**

fitness

**Format**

An object of class `data.frame` with 31 rows and 7 columns.

---

hsb	<i>Test Data Set</i>
-----	----------------------

---

**Description**

Test Data Set

**Usage**

hsb

**Format**

An object of class `data.frame` with 200 rows and 15 columns.

---

olsrr	<i>olsrr package</i>
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---

**Description**

Tools for teaching and learning OLS regression

**Details**

See the README on [GitHub](#)

---

ols_aic	<i>Akaike information criterion</i>
---------	-------------------------------------

---

**Description**

Akaike information criterion for model selection.

**Usage**

```
ols_aic(model, method = c("R", "STATA", "SAS"))
```

**Arguments**

model	An object of class <code>lm</code> .
method	A character vector; specify the method to compute AIC. Valid options include R, STATA and SAS.

**Details**

AIC provides a means for model selection. Given a collection of models for the data, AIC estimates the quality of each model, relative to each of the other models. R and STATA use loglikelihood to compute AIC. SAS uses residual sum of squares. Below is the formula in each case:

*R & STATA*

$$AIC = -2(\text{loglikelihood}) + 2p$$

*SAS*

$$AIC = n * \ln(SSE/n) + 2p$$

where  $n$  is the sample size and  $p$  is the number of model parameters including intercept.

**Value**

Akaike information criterion of the model.

## References

- Akaike, H. (1969). “Fitting Autoregressive Models for Prediction.” *Annals of the Institute of Statistical Mathematics* 21:243–247.
- Judge, G. G., Griffiths, W. E., Hill, R. C., and Lee, T.-C. (1980). *The Theory and Practice of Econometrics*. New York: John Wiley & Sons.

## See Also

Other model selection criteria: [ols\\_apc](#), [ols\\_fpe](#), [ols\\_hsp](#), [ols\\_mallows\\_cp](#), [ols\\_msep](#), [ols\\_sbc](#), [ols\\_sbic](#)

## Examples

```
# using R computation method
model <- lm(mpg ~ disp + hp + wt + qsec, data = mtcars)
ols_aic(model)

# using STATA computation method
model <- lm(mpg ~ disp + hp + wt + qsec, data = mtcars)
ols_aic(model, method = 'STATA')

# using SAS computation method
model <- lm(mpg ~ disp + hp + wt + qsec, data = mtcars)
ols_aic(model, method = 'SAS')
```

---

ols\_apc

*Amemiya's prediction criterion*

---

## Description

Amemiya's prediction error.

## Usage

```
ols_apc(model)
```

## Arguments

model            An object of class `lm`.

## Details

Amemiya's Prediction Criterion penalizes R-squared more heavily than does adjusted R-squared for each addition degree of freedom used on the right-hand-side of the equation. The higher the better for this criterion.

$$((n + p)/(n - p))(1 - (R^2))$$

where  $n$  is the sample size,  $p$  is the number of predictors including the intercept and  $R^2$  is the coefficient of determination.

### Value

Amemiya's prediction error of the model.

### References

- Amemiya, T. (1976). Selection of Regressors. Technical Report 225, Stanford University, Stanford, CA.
- Judge, G. G., Griffiths, W. E., Hill, R. C., and Lee, T.-C. (1980). The Theory and Practice of Econometrics. New York: John Wiley & Sons.

### See Also

Other model selection criteria: [ols\\_aic](#), [ols\\_fpe](#), [ols\\_hsp](#), [ols\\_mallows\\_cp](#), [ols\\_msep](#), [ols\\_sbc](#), [ols\\_sbic](#)

### Examples

```
model <- lm(mpg ~ disp + hp + wt + qsec, data = mtcars)
ols_apc(model)
```

---

ols\_coll\_diag

*Collinearity diagnostics*

---

### Description

Variance inflation factor, tolerance, eigenvalues and condition indices.

### Usage

```
ols_coll_diag(model)

ols_vif_tol(model)

ols_eigen_cindex(model)
```

### Arguments

model            An object of class `lm`.

## Details

Collinearity implies two variables are near perfect linear combinations of one another. Multicollinearity involves more than two variables. In the presence of multicollinearity, regression estimates are unstable and have high standard errors.

### *Tolerance*

Percent of variance in the predictor that cannot be accounted for by other predictors.

Steps to calculate tolerance:

- Regress the  $k$ th predictor on rest of the predictors in the model.
- Compute  $R^2$  - the coefficient of determination from the regression in the above step.
- $Tolerance = 1 - R^2$

### *Variance Inflation Factor*

Variance inflation factors measure the inflation in the variances of the parameter estimates due to collinearities that exist among the predictors. It is a measure of how much the variance of the estimated regression coefficient  $\beta_k$  is inflated by the existence of correlation among the predictor variables in the model. A VIF of 1 means that there is no correlation among the  $k$ th predictor and the remaining predictor variables, and hence the variance of  $\beta_k$  is not inflated at all. The general rule of thumb is that VIFs exceeding 4 warrant further investigation, while VIFs exceeding 10 are signs of serious multicollinearity requiring correction.

Steps to calculate VIF:

- Regress the  $k$ th predictor on rest of the predictors in the model.
- Compute  $R^2$  - the coefficient of determination from the regression in the above step.
- $Tolerance = 1/1 - R^2 = 1/Tolerance$

### *Condition Index*

Most multivariate statistical approaches involve decomposing a correlation matrix into linear combinations of variables. The linear combinations are chosen so that the first combination has the largest possible variance (subject to some restrictions), the second combination has the next largest variance, subject to being uncorrelated with the first, the third has the largest possible variance, subject to being uncorrelated with the first and second, and so forth. The variance of each of these linear combinations is called an eigenvalue. Collinearity is spotted by finding 2 or more variables that have large proportions of variance (.50 or more) that correspond to large condition indices. A rule of thumb is to label as large those condition indices in the range of 30 or larger.

## Value

ols\_coll\_diag returns an object of class "ols\_coll\_diag". An object of class "ols\_coll\_diag" is a list containing the following components:

vif_t	tolerance and variance inflation factors
eig_cindex	eigen values and condition index



## References

Belsley, D. A., Kuh, E., and Welsch, R. E. (1980). *Regression Diagnostics: Identifying Influential Data and Sources of Collinearity*. New York: John Wiley & Sons.

## Examples

```
# model
model <- lm(mpg ~ disp + hp + wt + drat, data = mtcars)

# vif and tolerance
ols_vif_tol(model)

# eigenvalues and condition indices
ols_eigen_cindex(model)

# collinearity diagnostics
ols_coll_diag(model)
```

---

ols_correlations	<i>Part and partial correlations</i>
------------------	--------------------------------------

---

## Description

Zero-order, part and partial correlations.

## Usage

```
ols_correlations(model)
```

## Arguments

model            An object of class `lm`.

## Details

`ols_correlations()` returns the relative importance of independent variables in determining response variable. How much each variable uniquely contributes to `rsquare` over and above that which can be accounted for by the other predictors? Zero order correlation is the Pearson correlation coefficient between the dependent variable and the independent variables. Part correlations indicates how much `rsquare` will decrease if that variable is removed from the model and partial correlations indicates amount of variance in response variable, which is not estimated by the other independent variables in the model, but is estimated by the specific variable.

**Value**

ols\_correlations returns an object of class "ols\_correlations". An object of class "ols\_correlations" is a data frame containing the following components:

Zero-order	zero order correlations
Partial	partial correlations
Part	part correlations

**References**

Morrison, D. F. 1976. Multivariate statistical methods. New York: McGraw-Hill.

**Examples**

```
model <- lm(mpg ~ disp + hp + wt + qsec, data = mtcars)
ols_correlations(model)
```

---

ols\_fpe

*Final prediction error*


---

**Description**

Estimated mean square error of prediction.

**Usage**

```
ols_fpe(model)
```

**Arguments**

model            An object of class lm.

**Details**

Computes the estimated mean square error of prediction for each model selected assuming that the values of the regressors are fixed and that the model is correct.

$$MSE((n + p)/n)$$

where  $MSE = SSE/(n - p)$ ,  $n$  is the sample size and  $p$  is the number of predictors including the intercept

**Value**

Final prediction error of the model.

## References

Akaike, H. (1969). "Fitting Autoregressive Models for Prediction." *Annals of the Institute of Statistical Mathematics* 21:243–247.

Judge, G. G., Griffiths, W. E., Hill, R. C., and Lee, T.-C. (1980). *The Theory and Practice of Econometrics*. New York: John Wiley & Sons.

## See Also

Other model selection criteria: [ols\\_aic](#), [ols\\_apc](#), [ols\\_hsp](#), [ols\\_mallows\\_cp](#), [ols\\_msep](#), [ols\\_sbc](#), [ols\\_sbic](#)

## Examples

```
model <- lm(mpg ~ disp + hp + wt + qsec, data = mtcars)
ols_fpe(model)
```

---

ols\_hadi

*Hadi's influence measure*

---

## Description

Measure of influence based on the fact that influential observations in either the response variable or in the predictors or both.

## Usage

```
ols_hadi(model)
```

## Arguments

model            An object of class `lm`.

## Value

Hadi's measure of the model.

## References

Chatterjee, Samprit and Hadi, Ali. *Regression Analysis by Example*. 5th ed. N.p.: John Wiley & Sons, 2012. Print.

## See Also

Other influence measures: [ols\\_leverage](#), [ols\\_pred\\_rsqr](#), [ols\\_press](#)

**Examples**

```
model <- lm(mpg ~ disp + hp + wt, data = mtcars)
ols_hadi(model)
```

---

ols\_hsp

*Hocking's Sp*


---

**Description**

Average prediction mean squared error.

**Usage**

```
ols_hsp(model)
```

**Arguments**

model            An object of class `lm`.

**Details**

Hocking's  $Sp$  criterion is an adjustment of the residual sum of Squares. Minimize this criterion.

$$MSE/(n - p - 1)$$

where  $MSE = SSE/(n - p)$ ,  $n$  is the sample size and  $p$  is the number of predictors including the intercept

**Value**

Hocking's  $Sp$  of the model.

**References**

Hocking, R. R. (1976). "The Analysis and Selection of Variables in a Linear Regression." *Biometrics* 32:1-50.

**See Also**

Other model selection criteria: [ols\\_aic](#), [ols\\_apc](#), [ols\\_fpe](#), [ols\\_mallows\\_cp](#), [ols\\_msep](#), [ols\\_sbc](#), [ols\\_sbic](#)

**Examples**

```
model <- lm(mpg ~ disp + hp + wt + qsec, data = mtcars)
ols_hsp(model)
```

---

ols_launch_app	<i>Launch shiny app</i>
----------------	-------------------------

---

**Description**

Launches shiny app for interactive model building.

**Usage**

```
ols_launch_app()
```

**Examples**

```
## Not run:  
ols_launch_app()  
  
## End(Not run)
```

---

ols_leverage	<i>Leverage</i>
--------------	-----------------

---

**Description**

The leverage of an observation is based on how much the observation's value on the predictor variable differs from the mean of the predictor variable. The greater an observation's leverage, the more potential it has to be an influential observation.

**Usage**

```
ols_leverage(model)
```

**Arguments**

model            An object of class `lm`.

**Value**

Leverage of the model.

**References**

Kutner, MH, Nachtsheim CJ, Neter J and Li W., 2004, Applied Linear Statistical Models (5th edition). Chicago, IL., McGraw Hill/Irwin.

**See Also**

Other influence measures: [ols\\_hadi](#), [ols\\_pred\\_rsqr](#), [ols\\_press](#)

## Examples

```
model <- lm(mpg ~ disp + hp + wt + qsec, data = mtcars)
ols_leverage(model)
```

---

ols_mallows_cp	<i>Mallow's Cp</i>
----------------	--------------------

---

## Description

Mallow's Cp.

## Usage

```
ols_mallows_cp(model, fullmodel)
```

## Arguments

model	An object of class lm.
fullmodel	An object of class lm.

## Details

Mallows' Cp statistic estimates the size of the bias that is introduced into the predicted responses by having an underspecified model. Use Mallows' Cp to choose between multiple regression models. Look for models where Mallows' Cp is small and close to the number of predictors in the model plus the constant (p).

## Value

Mallow's Cp of the model.

## References

Hocking, R. R. (1976). "The Analysis and Selection of Variables in a Linear Regression." *Biometrics* 32:1–50.

Mallows, C. L. (1973). "Some Comments on Cp." *Technometrics* 15:661–675.

## See Also

Other model selection criteria: [ols\\_aic](#), [ols\\_apc](#), [ols\\_fpe](#), [ols\\_hsp](#), [ols\\_msep](#), [ols\\_sbc](#), [ols\\_sbic](#)

## Examples

```
full_model <- lm(mpg ~ ., data = mtcars)
model <- lm(mpg ~ disp + hp + wt + qsec, data = mtcars)
ols_mallows_cp(model, full_model)
```

---

ols_msep	<i>MSEP</i>
----------	-------------

---

**Description**

Estimated error of prediction, assuming multivariate normality.

**Usage**

```
ols_msep(model)
```

**Arguments**

model            An object of class `lm`.

**Details**

Computes the estimated mean square error of prediction assuming that both independent and dependent variables are multivariate normal.

$$MSE(n+1)(n-2)/n(n-p-1)$$

where  $MSE = SSE/(n-p)$ ,  $n$  is the sample size and  $p$  is the number of predictors including the intercept

**Value**

Estimated error of prediction of the model.

**References**

Stein, C. (1960). "Multiple Regression." In Contributions to Probability and Statistics: Essays in Honor of Harold Hotelling, edited by I. Olkin, S. G. Ghurye, W. Hoeffding, W. G. Madow, and H. B. Mann, 264–305. Stanford, CA: Stanford University Press.

Darlington, R. B. (1968). "Multiple Regression in Psychological Research and Practice." Psychological Bulletin 69:161–182.

**See Also**

Other model selection criteria: [ols\\_aic](#), [ols\\_apc](#), [ols\\_fpe](#), [ols\\_hsp](#), [ols\\_mallows\\_cp](#), [ols\\_sbc](#), [ols\\_sbic](#)

**Examples**

```
model <- lm(mpg ~ disp + hp + wt + qsec, data = mtcars)
ols_msep(model)
```

---

`ols_plot_added_variable`*Added variable plots*

---

### Description

Added variable plot provides information about the marginal importance of a predictor variable, given the other predictor variables already in the model. It shows the marginal importance of the variable in reducing the residual variability.

### Usage

```
ols_plot_added_variable(model, print_plot = TRUE)
```

### Arguments

<code>model</code>	An object of class <code>lm</code> .
<code>print_plot</code>	logical; if TRUE, prints the plot else returns a plot object.

### Details

The added variable plot was introduced by Mosteller and Tukey (1977). It enables us to visualize the regression coefficient of a new variable being considered to be included in a model. The plot can be constructed for each predictor variable.

Let us assume we want to test the effect of adding/removing variable  $X$  from a model. Let the response variable of the model be  $Y$

Steps to construct an added variable plot:

- Regress  $Y$  on all variables other than  $X$  and store the residuals ( $Y$  residuals).
- Regress  $X$  on all the other variables included in the model ( $X$  residuals).
- Construct a scatter plot of  $Y$  residuals and  $X$  residuals.

What do the  $Y$  and  $X$  residuals represent? The  $Y$  residuals represent the part of  $Y$  not explained by all the variables other than  $X$ . The  $X$  residuals represent the part of  $X$  not explained by other variables. The slope of the line fitted to the points in the added variable plot is equal to the regression coefficient when  $Y$  is regressed on all variables including  $X$ .

A strong linear relationship in the added variable plot indicates the increased importance of the contribution of  $X$  to the model already containing the other predictors.

### Deprecated Function

`ols_avplots()` has been deprecated. Instead use `ols_plot_added_variable()`.



## References

Chatterjee, Samprit and Hadi, Ali. Regression Analysis by Example. 5th ed. N.p.: John Wiley & Sons, 2012. Print.

Kutner, MH, Nachtsheim CJ, Neter J and Li W., 2004, Applied Linear Statistical Models (5th edition). Chicago, IL., McGraw Hill/Irwin.

## See Also

[ols\_plot\_resid\_regressor()], [ols\_plot\_comp\_plus\_resid()]

## Examples

```
model <- lm(mpg ~ disp + hp + wt, data = mtcars)
ols_plot_added_variable(model)
```

---

ols\_plot\_comp\_plus\_resid  
*Residual plus component plot*

---

## Description

The residual plus component plot indicates whether any non-linearity is present in the relationship between response and predictor variables and can suggest possible transformations for linearizing the data.

## Usage

```
ols_plot_comp_plus_resid(model, print_plot = TRUE)
```

## Arguments

model	An object of class lm.
print_plot	logical; if TRUE, prints the plot else returns a plot object.

## Deprecated Function

ols\_rpc\_plot() has been deprecated. Instead use ols\_plot\_comp\_plus\_resid().

## References

Chatterjee, Samprit and Hadi, Ali. Regression Analysis by Example. 5th ed. N.p.: John Wiley & Sons, 2012. Print.

Kutner, MH, Nachtsheim CJ, Neter J and Li W., 2004, Applied Linear Statistical Models (5th edition). Chicago, IL., McGraw Hill/Irwin.

**See Also**

[ols\_plot\_added\_variable()], [ols\_plot\_resid\_regressor()]

**Examples**

```
model <- lm(mpg ~ disp + hp + wt + qsec, data = mtcars)
ols_plot_comp_plus_resid(model)
```

---

ols\_plot\_cooksd\_bar     *Cooks' D bar plot*

---

**Description**

Bar Plot of cook's distance to detect observations that strongly influence fitted values of the model.

**Usage**

```
ols_plot_cooksd_bar(model, print_plot = TRUE)
```

**Arguments**

model	An object of class lm.
print_plot	logical; if TRUE, prints the plot else returns a plot object.

**Details**

Cook's distance was introduced by American statistician R Dennis Cook in 1977. It is used to identify influential data points. It depends on both the residual and leverage i.e it takes it account both the  $x$  value and  $y$  value of the observation.

Steps to compute Cook's distance:

- Delete observations one at a time.
- Refit the regression model on remaining  $n - 1$  observations
- examine how much all of the fitted values change when the  $i$ th observation is deleted.

A data point having a large cook's d indicates that the data point strongly influences the fitted values.

**Value**

ols\_plot\_cooksd\_bar returns a list containing the following components:

outliers	a data.frame with observation number and cooks distance that exceed threshold
threshold	threshold for classifying an observation as an outlier

**Deprecated Function**

ols\_cooksd\_barplot() has been deprecated. Instead use ols\_plot\_cooksd\_bar().

**See Also**

[ols\_plot\_cooksd\_chart()]

**Examples**

```
model <- lm(mpg ~ disp + hp + wt, data = mtcars)
ols_plot_cooksd_bar(model)
```

---

ols\_plot\_cooksd\_chart *Cooks' D chart*

---

**Description**

Chart of cook's distance to detect observations that strongly influence fitted values of the model.

**Usage**

```
ols_plot_cooksd_chart(model, print_plot = TRUE)
```

**Arguments**

model	An object of class lm.
print_plot	logical; if TRUE, prints the plot else returns a plot object.

**Details**

Cook's distance was introduced by American statistician R Dennis Cook in 1977. It is used to identify influential data points. It depends on both the residual and leverage i.e it takes it account both the  $x$  value and  $y$  value of the observation.

Steps to compute Cook's distance:

- Delete observations one at a time.
- Refit the regression model on remaining  $n - 1$  observations
- examine how much all of the fitted values change when the  $i$ th observation is deleted.

A data point having a large cook's  $d$  indicates that the data point strongly influences the fitted values.

**Value**

ols\_plot\_cooksd\_chart returns a list containing the following components:

outliers	a data.frame with observation number and cooks distance that exceed threshold
threshold	threshold for classifying an observation as an outlier

**Deprecated Function**

`ols_cooksd_chart()` has been deprecated. Instead use `ols_plot_cooksd_chart()`.

**See Also**

[`ols_plot_cooksd_bar()`]

**Examples**

```
model <- lm(mpg ~ disp + hp + wt, data = mtcars)
ols_plot_cooksd_chart(model)
```

---

`ols_plot_dfbetas`      *DFBETAs panel*

---

**Description**

Panel of plots to detect influential observations using DFBETAs.

**Usage**

```
ols_plot_dfbetas(model, print_plot = TRUE)
```

**Arguments**

`model`            An object of class `lm`.  
`print_plot`       logical; if `TRUE`, prints the plot else returns a plot object.

**Details**

DFBETA measures the difference in each parameter estimate with and without the influential point. There is a DFBETA for each data point i.e if there are  $n$  observations and  $k$  variables, there will be  $n * k$  DFBETAs. In general, large values of DFBETAS indicate observations that are influential in estimating a given parameter. Belsley, Kuh, and Welsch recommend 2 as a general cutoff value to indicate influential observations and  $2/\sqrt{(n)}$  as a size-adjusted cutoff.

**Value**

list; `ols_plot_dfbetas` returns a list of `data.frame` (for intercept and each predictor) with the observation number and DFBETA of observations that exceed the threshold for classifying an observation as an outlier/influential observation.

**Deprecated Function**

`ols_dfbetas_panel()` has been deprecated. Instead use `ols_plot_dfbetas()`.

**References**

Belsley, David A.; Kuh, Edwin; Welsh, Roy E. (1980). Regression Diagnostics: Identifying Influential Data and Sources of Collinearity.

Wiley Series in Probability and Mathematical Statistics. New York: John Wiley & Sons. pp. ISBN 0-471-05856-4.

**See Also**

[ols\_plot\_dffits()]

**Examples**

```
model <- lm(mpg ~ disp + hp + wt + qsec, data = mtcars)
ols_plot_dfbetas(model)
```

---

ols_plot_dffits	<i>DFFITs plot</i>
-----------------	--------------------

---

**Description**

Plot for detecting influential observations using DFFITs.

**Usage**

```
ols_plot_dffits(model, print_plot = TRUE)
```

**Arguments**

model	An object of class lm.
print_plot	logical; if TRUE, prints the plot else returns a plot object.

**Details**

DFFIT - difference in fits, is used to identify influential data points. It quantifies the number of standard deviations that the fitted value changes when the *i*th data point is omitted.

Steps to compute DFFITs:

- Delete observations one at a time.
- Refit the regression model on remaining  $n - 1$  observations
- examine how much all of the fitted values change when the *i*th observation is deleted.

An observation is deemed influential if the absolute value of its DFFITs value is greater than:

$$2\sqrt{(p + 1)/(n - p - 1)}$$

where *n* is the number of observations and *p* is the number of predictors including intercept.

**Value**

ols\_plot\_dffits returns a list containing the following components:

outliers	a data.frame with observation number and DFFITs that exceed threshold
threshold	threshold for classifying an observation as an outlier

**Deprecated Function**

ols\_dffits\_plot() has been deprecated. Instead use ols\_plot\_dffits().

**References**

Belsley, David A.; Kuh, Edwin; Welsh, Roy E. (1980). Regression Diagnostics: Identifying Influential Data and Sources of Collinearity.

Wiley Series in Probability and Mathematical Statistics. New York: John Wiley & Sons. ISBN 0-471-05856-4.

**See Also**

[ols\_plot\_dfbetas()]

**Examples**

```
model <- lm(mpg ~ disp + hp + wt + qsec, data = mtcars)
ols_plot_dffits(model)
```

---

ols\_plot\_diagnostics *Diagnostics panel*

---

**Description**

Panel of plots for regression diagnostics.

**Usage**

```
ols_plot_diagnostics(model, print_plot = TRUE)
```

**Arguments**

model	An object of class lm.
print_plot	logical; if TRUE, prints the plot else returns a plot object. #’ @section Deprecated Function: ols_diagnostic_panel() has been deprecated. Instead use ols_plot_diagnostics().

**Examples**

```
model <- lm(mpg ~ disp + hp + wt + qsec, data = mtcars)
ols_plot_diagnostics(model)
```

---

ols_plot_hadi	<i>Hadi plot</i>
---------------	------------------

---

**Description**

Hadi's measure of influence based on the fact that influential observations can be present in either the response variable or in the predictors or both. The plot is used to detect influential observations based on Hadi's measure.

**Usage**

```
ols_plot_hadi(model, print_plot = TRUE)
```

**Arguments**

model	An object of class <code>lm</code> .
print_plot	logical; if <code>TRUE</code> , prints the plot else returns a plot object.

**Deprecated Function**

`ols_hadi_plot()` has been deprecated. Instead use `ols_plot_hadi()`.

**References**

Chatterjee, Samprit and Hadi, Ali. Regression Analysis by Example. 5th ed. N.p.: John Wiley & Sons, 2012. Print.

**See Also**

[`ols_plot_resid_pot()`]

**Examples**

```
model <- lm(mpg ~ disp + hp + wt, data = mtcars)
ols_plot_hadi(model)
```

---

ols\_plot\_obs\_fit      *Observed vs fitted values plot*

---

**Description**

Plot of observed vs fitted values to assess the fit of the model.

**Usage**

```
ols_plot_obs_fit(model, print_plot = TRUE)
```

**Arguments**

model            An object of class lm.  
print\_plot      logical; if TRUE, prints the plot else returns a plot object.

**Details**

Ideally, all your points should be close to a regressed diagonal line. Draw such a diagonal line within your graph and check out where the points lie. If your model had a high R Square, all the points would be close to this diagonal line. The lower the R Square, the weaker the Goodness of fit of your model, the more foggy or dispersed your points are from this diagonal line.

**Deprecated Function**

ols\_ovsp\_plot() has been deprecated. Instead use ols\_plot\_obs\_fit().

**Examples**

```
model <- lm(mpg ~ disp + hp + wt, data = mtcars)  
ols_plot_obs_fit(model)
```

---

ols\_plot\_reg\_line      *Simple linear regression line*

---

**Description**

Plot to demonstrate that the regression line always passes through mean of the response and predictor variables.

**Usage**

```
ols_plot_reg_line(response, predictor, print_plot = TRUE)
```



**Arguments**

response	Response variable.
predictor	Predictor variable.
print_plot	logical; if TRUE, prints the plot else returns a plot object.

**Deprecated Function**

ols\_reg\_line() has been deprecated. Instead use ols\_plot\_reg\_line().

**Examples**

```
ols_plot_reg_line(mtcars$mpg, mtcars$disp)
```

---

ols_plot_resid_box	<i>Residual box plot</i>
--------------------	--------------------------

---

**Description**

Box plot of residuals to examine if residuals are normally distributed.

**Usage**

```
ols_plot_resid_box(model, print_plot = TRUE)
```

**Arguments**

model	An object of class lm.
print_plot	logical; if TRUE, prints the plot else returns a plot object.

**Deprecated Function**

ols\_rsd\_boxplot() has been deprecated. Instead use ols\_plot\_resid\_box().

**See Also**

Other residual diagnostics: [ols\\_plot\\_resid\\_fit](#), [ols\\_plot\\_resid\\_hist](#), [ols\\_plot\\_resid\\_qq](#), [ols\\_test\\_correlation](#), [ols\\_test\\_normality](#)

**Examples**

```
model <- lm(mpg ~ disp + hp + wt, data = mtcars)
ols_plot_resid_box(model)
```

---

ols\_plot\_resid\_fit      *Residual vs fitted plot*

---

### Description

Scatter plot of residuals on the y axis and fitted values on the x axis to detect non-linearity, unequal error variances, and outliers.

### Usage

```
ols_plot_resid_fit(model, print_plot = TRUE)
```

### Arguments

model	An object of class lm.
print_plot	logical; if TRUE, prints the plot else returns a plot object.

### Details

Characteristics of a well behaved residual vs fitted plot:

- The residuals spread randomly around the 0 line indicating that the relationship is linear.
- The residuals form an approximate horizontal band around the 0 line indicating homogeneity of error variance.
- No one residual is visibly away from the random pattern of the residuals indicating that there are no outliers.

### Deprecated Function

ols\_rvsp\_plot() has been deprecated. Instead use ols\_plot\_resid\_fit().

### See Also

Other residual diagnostics: [ols\\_plot\\_resid\\_box](#), [ols\\_plot\\_resid\\_hist](#), [ols\\_plot\\_resid\\_qq](#), [ols\\_test\\_correlation](#), [ols\\_test\\_normality](#)

### Examples

```
model <- lm(mpg ~ disp + hp + wt, data = mtcars)
ols_plot_resid_fit(model)
```

---

ols\_plot\_resid\_fit\_spread  
*Residual fit spread plot*

---

### Description

Plot to detect non-linearity, influential observations and outliers.

### Usage

```
ols_plot_resid_fit_spread(model, print_plot = TRUE)
```

```
ols_plot_fm(model, print_plot = TRUE)
```

```
ols_plot_resid_spread(model, print_plot = TRUE)
```

### Arguments

model	An object of class lm.
print_plot	logical; if TRUE, prints the plot else returns a plot object.

### Details

Consists of side-by-side quantile plots of the centered fit and the residuals. It shows how much variation in the data is explained by the fit and how much remains in the residuals. For inappropriate models, the spread of the residuals in such a plot is often greater than the spread of the centered fit.

### Deprecated Function

ols\_rfs\_plot(), ols\_fm\_plot() and ols\_rsd\_plot() has been deprecated. Instead use ols\_plot\_resid\_fit\_spread(), ols\_plot\_fm() and ols\_plot\_resid\_spread().

### References

Cleveland, W. S. (1993). Visualizing Data. Summit, NJ: Hobart Press.

### Examples

```
# model
model <- lm(mpg ~ disp + hp + wt, data = mtcars)

# residual fit spread plot
ols_plot_resid_fit_spread(model)

# fit mean plot
ols_plot_fm(model)

# residual spread plot
```

```
ols_plot_resid_spread(model)
```

---

```
ols_plot_resid_hist
```

*Residual histogram*

---

**Description**

Histogram of residuals for detecting violation of normality assumption.

**Usage**

```
ols_plot_resid_hist(model, print_plot = TRUE)
```

**Arguments**

`model`            An object of class `lm`.  
`print_plot`       logical; if `TRUE`, prints the plot else returns a plot object.

**Deprecated Function**

`ols_rsd_hist()` has been deprecated. Instead use `ols_plot_resid_hist()`.

**See Also**

Other residual diagnostics: [ols\\_plot\\_resid\\_box](#), [ols\\_plot\\_resid\\_fit](#), [ols\\_plot\\_resid\\_qq](#), [ols\\_test\\_correlation](#), [ols\\_test\\_normality](#)

**Examples**

```
model <- lm(mpg ~ disp + hp + wt, data = mtcars)
ols_plot_resid_hist(model)
```

---

```
ols_plot_resid_lev
```

*Studentized residuals vs leverage plot*

---

**Description**

Graph for detecting outliers and/or observations with high leverage.

**Usage**

```
ols_plot_resid_lev(model, print_plot = TRUE)
```

**Arguments**

model            An object of class lm.  
print\_plot       logical; if TRUE, prints the plot else returns a plot object.

**Deprecated Function**

ols\_rsdlev\_plot() has been deprecated. Instead use ols\_plot\_resid\_lev().

**See Also**

[ols\_plot\_resid\_stud\_fit()], [ols\_plot\_resid\_lev()]

**Examples**

```
model <- lm(read ~ write + math + science, data = hsb)
ols_plot_resid_lev(model)
```

---

ols\_plot\_resid\_pot       *Potential residual plot*

---

**Description**

Plot to aid in classifying unusual observations as high-leverage points, outliers, or a combination of both.

**Usage**

```
ols_plot_resid_pot(model, print_plot = TRUE)
```

**Arguments**

model            An object of class lm.  
print\_plot       logical; if TRUE, prints the plot else returns a plot object.

**Deprecated Function**

ols\_potrsd\_plot() has been deprecated. Instead use ols\_plot\_resid\_pot().

**References**

Chatterjee, Samprit and Hadi, Ali. Regression Analysis by Example. 5th ed. N.p.: John Wiley & Sons, 2012. Print.

**See Also**

[ols\_plot\_hadi()]

## Examples

```
model <- lm(mpg ~ disp + hp + wt, data = mtcars)
ols_plot_resid_pot(model)
```

---

ols\_plot\_resid\_qq      *Residual QQ plot*

---

## Description

Graph for detecting violation of normality assumption.

## Usage

```
ols_plot_resid_qq(model, print_plot = TRUE)
```

## Arguments

model	An object of class lm.
print_plot	logical; if TRUE, prints the plot else returns a plot object.

## Deprecated Function

ols\_rsd\_qqplot() has been deprecated. Instead use ols\_plot\_resid\_qq().

## See Also

Other residual diagnostics: [ols\\_plot\\_resid\\_box](#), [ols\\_plot\\_resid\\_fit](#), [ols\\_plot\\_resid\\_hist](#), [ols\\_test\\_correlation](#), [ols\\_test\\_normality](#)

## Examples

```
model <- lm(mpg ~ disp + hp + wt, data = mtcars)
ols_plot_resid_qq(model)
```

---

`ols_plot_resid_regressor`*Residual vs regressor plot*

---

**Description**

Graph to determine whether we should add a new predictor to the model already containing other predictors. The residuals from the model is regressed on the new predictor and if the plot shows non random pattern, you should consider adding the new predictor to the model.

**Usage**

```
ols_plot_resid_regressor(model, variable, print_plot = TRUE)
```

**Arguments**

<code>model</code>	An object of class <code>lm</code> .
<code>variable</code>	New predictor to be added to the model.
<code>print_plot</code>	logical; if TRUE, prints the plot else returns a plot object.

**Deprecated Function**

`ols_rvsr_plot()` has been deprecated. Instead use `ols_plot_resid_regressor()`.

**See Also**

[`ols_plot_added_variable()`], [`ols_plot_comp_plus_resid()`]

**Examples**

```
model <- lm(mpg ~ disp + hp + wt, data = mtcars)
ols_plot_resid_regressor(model, 'drat')
```

---

`ols_plot_resid_stand` *Standardized residual chart*

---

**Description**

Chart for identifying outliers.

**Usage**

```
ols_plot_resid_stand(model, print_plot = TRUE)
```

**Arguments**

model            An object of class `lm`.  
 print\_plot      logical; if TRUE, prints the plot else returns a plot object.

**Details**

Standardized residual (internally studentized) is the residual divided by estimated standard deviation.

**Value**

`ols_plot_resid_stand` returns a list containing the following components:

outliers        a data.frame with observation number and standardized residuals that exceed threshold

for classifying an observation as an outlier

threshold       threshold for classifying an observation as an outlier

**Deprecated Function**

`ols_srsd_chart()` has been deprecated. Instead use `ols_plot_resid_stand()`.

**See Also**

[`ols_plot_resid_stud()`]

**Examples**

```
model <- lm(mpg ~ disp + hp + wt, data = mtcars)
ols_plot_resid_stand(model)
```

---

`ols_plot_resid_stud`    *Studentized residual plot*

---

**Description**

Graph for identifying outliers.

**Usage**

```
ols_plot_resid_stud(model, print_plot = TRUE)
```

**Arguments**

model            An object of class `lm`.  
 print\_plot      logical; if TRUE, prints the plot else returns a plot object.



**Details**

Studentized deleted residuals (or externally studentized residuals) is the deleted residual divided by its estimated standard deviation. Studentized residuals are going to be more effective for detecting outlying Y observations than standardized residuals. If an observation has an externally studentized residual that is larger than 3 (in absolute value) we can call it an outlier.

**Value**

ols\_plot\_resid\_stud returns a list containing the following components:

outliers            a data.frame with observation number and studentized residuals that exceed threshold

for classifying an observation as an outlier

threshold            threshold for classifying an observation as an outlier

**Deprecated Function**

ols\_srsd\_plot() has been deprecated. Instead use ols\_plot\_resid\_stud().

**See Also**

[ols\_plot\_resid\_stand()]

**Examples**

```
model <- lm(mpg ~ disp + hp + wt, data = mtcars)
ols_plot_resid_stud(model)
```

---

ols\_plot\_resid\_stud\_fit

*Deleted studentized residual vs fitted values plot*

---

**Description**

Plot for detecting violation of assumptions about residuals such as non-linearity, constant variances and outliers. It can also be used to examine model fit.

**Usage**

```
ols_plot_resid_stud_fit(model, print_plot = TRUE)
```

**Arguments**

model                An object of class lm.

print\_plot           logical; if TRUE, prints the plot else returns a plot object.

**Details**

Studentized deleted residuals (or externally studentized residuals) is the deleted residual divided by its estimated standard deviation. Studentized residuals are going to be more effective for detecting outlying Y observations than standardized residuals. If an observation has an externally studentized residual that is larger than 2 (in absolute value) we can call it an outlier.

**Value**

ols\_plot\_resid\_stud\_fit returns a list containing the following components:

outliers	a data.frame with observation number, fitted values and deleted studentized residuals that exceed the threshold for classifying observations as outliers/influential observations
threshold	threshold for classifying an observation as an outlier/influential observation

**Deprecated Function**

ols\_dsrvsp\_plot() has been deprecated. Instead use ols\_plot\_resid\_stud\_fit().

**See Also**

[ols\_plot\_resid\_lev()], [ols\_plot\_resid\_stand()], [ols\_plot\_resid\_stud()]

**Examples**

```
model <- lm(mpg ~ disp + hp + wt + qsec, data = mtcars)
ols_plot_resid_stud_fit(model)
```

---

ols\_plot\_response      *Response variable profile*

---

**Description**

Panel of plots to explore and visualize the response variable.

**Usage**

```
ols_plot_response(model, print_plot = TRUE)
```

**Arguments**

model	An object of class lm.
print_plot	logical; if TRUE, prints the plot else returns a plot object.

**Deprecated Function**

ols\_resp\_viz() has been deprecated. Instead use ols\_plot\_response().

**Examples**

```
model <- lm(mpg ~ disp + hp + wt, data = mtcars)
ols_plot_response(model)
```

---

ols_pred_rsq	<i>Predicted rsquare</i>
--------------	--------------------------

---

**Description**

Use predicted rsquared to determine how well the model predicts responses for new observations. Larger values of predicted R2 indicate models of greater predictive ability.

**Usage**

```
ols_pred_rsq(model)
```

**Arguments**

model            An object of class lm.

**Value**

Predicted rsquare of the model.

**See Also**

Other influence measures: [ols\\_hadi](#), [ols\\_leverage](#), [ols\\_press](#)

**Examples**

```
model <- lm(mpg ~ disp + hp + wt + qsec, data = mtcars)
ols_pred_rsq(model)
```

---

ols_prep_avplot_data	<i>Added variable plot data</i>
----------------------	---------------------------------

---

**Description**

Data for generating the added variable plots.

**Usage**

```
ols_prep_avplot_data(model)
```

**Arguments**

model            An object of class lm.

**Examples**

```
model <- lm(mpg ~ disp + hp + wt, data = mtcars)
ols_prep_avplot_data(model)
```

---

ols\_prep\_cdplot\_data    *Cooks' D plot data*

---

**Description**

Prepare data for cook's d bar plot.

**Usage**

```
ols_prep_cdplot_data(model)
```

**Arguments**

model            An object of class lm.

**Examples**

```
model <- lm(mpg ~ disp + hp + wt, data = mtcars)
ols_prep_cdplot_data(model)
```

---

ols\_prep\_cdplot\_outliers  
                          *Cooks' d outlier data*

---

**Description**

Outlier data for cook's d bar plot.

**Usage**

```
ols_prep_cdplot_outliers(k)
```

**Arguments**

k                Cooks' d bar plot data.

**Examples**

```
model <- lm(mpg ~ disp + hp + wt, data = mtcars)
k <- ols_prep_cdplot_data(model)
ols_prep_cdplot_outliers(k)
```

---

ols\_prep\_dfbeta\_data *DFBETAs plot data*

---

**Description**

Prepares the data for dfbetas plot.

**Usage**

```
ols_prep_dfbeta_data(d, threshold)
```

**Arguments**

d	A tibble or data.frame with dfbetas.
threshold	The threshold for outliers.

**Examples**

```
model <- lm(mpg ~ disp + hp + wt + qsec, data = mtcars)
dfb <- dfbetas(model)
n <- nrow(dfb)
threshold <- 2 / sqrt(n)
dbetas <- dfb[, 1]
df_data <- data.frame(obs = seq_len(n), dbetas = dbetas)
ols_prep_dfbeta_data(df_data, threshold)
```

---

ols\_prep\_dfbeta\_outliers  
*DFBETAs plot outliers*

---

**Description**

Data for identifying outliers in dfbetas plot.

**Usage**

```
ols_prep_dfbeta_outliers(d)
```

**Arguments**

d                    A tibble or data.frame.

**Examples**

```
model <- lm(mpg ~ disp + hp + wt + qsec, data = mtcars)
dfb <- dfbetas(model)
n <- nrow(dfb)
threshold <- 2 / sqrt(n)
dbetas <- dfb[, 1]
df_data <- data.frame(obs = seq_len(n), dbetas = dbetas)
d <- ols_prep_dfbeta_data(df_data, threshold)
ols_prep_dfbeta_outliers(d)
```

---

ols\_prep\_dsrvf\_data    *Deleted studentized residual plot data*

---

**Description**

Generates data for deleted studentized residual vs fitted plot.

**Usage**

```
ols_prep_dsrvf_data(model)
```

**Arguments**

model                An object of class lm.

**Examples**

```
model <- lm(mpg ~ disp + hp + wt + qsec, data = mtcars)
ols_prep_dsrvf_data(model)
```

---

ols\_prep\_outlier\_obs    *Cooks' D outlier observations*

---

**Description**

Identify outliers in cook's d plot.

**Usage**

```
ols_prep_outlier_obs(k)
```

**Arguments**

k                      Cooks' d bar plot data.

**Examples**

```
model <- lm(mpg ~ disp + hp + wt, data = mtcars)
k <- ols_prep_cdplot_data(model)
ols_prep_outlier_obs(k)
```

---

ols\_prep\_regress\_x      *Regress predictor on other predictors*

---

**Description**

Regress a predictor in the model on all the other predictors.

**Usage**

```
ols_prep_regress_x(data, i)
```

**Arguments**

data                    A data.frame.  
i                        A numeric vector (indicates the predictor in the model).

**Examples**

```
model <- lm(mpg ~ disp + hp + wt, data = mtcars)
data <- ols_prep_avplot_data(model)
ols_prep_regress_x(data, 1)
```

---

ols\_prep\_regress\_y      *Regress y on other predictors*

---

**Description**

Regress y on all the predictors except the ith predictor.

**Usage**

```
ols_prep_regress_y(data, i)
```

**Arguments**

`data` A data.frame.  
`i` A numeric vector (indicates the predictor in the model).

**Examples**

```
model <- lm(mpg ~ disp + hp + wt, data = mtcars)
data <- ols_prep_avplot_data(model)
ols_prep_regress_y(data, 1)
```

---

```
ols_prep_rfplot_fmdata
      Residual fit spread plot data
```

---

**Description**

Data for generating residual fit spread plot.

**Usage**

```
ols_prep_rfplot_fmdata(model)
ols_prep_rfplot_rsdata(model)
```

**Arguments**

`model` An object of class `lm`.

**Examples**

```
model <- lm(mpg ~ disp + hp + wt + qsec, data = mtcars)
ols_prep_rfplot_fmdata(model)
ols_prep_rfplot_rsdata(model)
```



---

`ols_prep_rstudlev_data`*Studentized residual vs leverage plot data*

---

**Description**

Generates data for studentized residual vs leverage plot.

**Usage**

```
ols_prep_rstudlev_data(model)
```

**Arguments**

`model` An object of class `lm`.

**Examples**

```
model <- lm(read ~ write + math + science, data = hsb)
ols_prep_rstudlev_data(model)
```

---

`ols_prep_rvsrplot_data`*Residual vs regressor plot data*

---

**Description**

Data for generating residual vs regressor plot.

**Usage**

```
ols_prep_rvsrplot_data(model)
```

**Arguments**

`model` An object of class `lm`.

**Examples**

```
model <- lm(mpg ~ disp + hp + wt + qsec, data = mtcars)
ols_prep_rvsrplot_data(model)
```

---

ols\_prep\_srchart\_data *Standardized residual chart data*

---

**Description**

Generates data for standardized residual chart.

**Usage**

```
ols_prep_srchart_data(model)
```

**Arguments**

model            An object of class lm.

**Examples**

```
model <- lm(read ~ write + math + science, data = hsb)
ols_prep_srchart_data(model)
```

---

ols\_prep\_srplot\_data *Studentized residual plot data*

---

**Description**

Generates data for studentized residual plot.

**Usage**

```
ols_prep_srplot_data(model)
```

**Arguments**

model            An object of class lm.

**Examples**

```
model <- lm(read ~ write + math + science, data = hsb)
ols_prep_srplot_data(model)
```

---

ols_press	<i>PRESS</i>
-----------	--------------

---

### Description

PRESS (prediction sum of squares) tells you how well the model will predict new data.

### Usage

```
ols_press(model)
```

### Arguments

model            An object of class `lm`.

### Details

The prediction sum of squares (PRESS) is the sum of squares of the prediction error. Each fitted to obtain the predicted value for the *i*th observation. Use PRESS to assess your model's predictive ability. Usually, the smaller the PRESS value, the better the model's predictive ability.

### Value

Predicted sum of squares of the model.

### References

Kutner, MH, Nachtsheim CJ, Neter J and Li W., 2004, Applied Linear Statistical Models (5th edition). Chicago, IL., McGraw Hill/Irwin.

### See Also

Other influence measures: [ols\\_hadi](#), [ols\\_leverage](#), [ols\\_pred\\_rsq](#)

### Examples

```
model <- lm(mpg ~ disp + hp + wt + qsec, data = mtcars)
ols_press(model)
```

---

ols\_pure\_error\_anova *Lack of fit F test*

---

### Description

Assess how much of the error in prediction is due to lack of model fit.

### Usage

```
ols_pure_error_anova(model, ...)
```

### Arguments

model	An object of class lm.
...	Other parameters.

### Details

The residual sum of squares resulting from a regression can be decomposed into 2 components:

- Due to lack of fit
- Due to random variation

If most of the error is due to lack of fit and not just random error, the model should be discarded and a new model must be built.

### Value

ols\_pure\_error\_anova returns an object of class "ols\_pure\_error\_anova". An object of class "ols\_pure\_error\_anova" is a list containing the following components:

lackoffit	lack of fit sum of squares
pure_error	pure error sum of squares
rss	regression sum of squares
ess	error sum of squares
total	total sum of squares
rms	regression mean square
ems	error mean square
lms	lack of fit mean square
pms	pure error mean square
rf	f statistic
lf	lack of fit f statistic
pr	p-value of f statistic
pl	p-value pf lack of fit f statistic

mpred	data.frame containing data for the response and predictor of the model
df_rss	regression sum of squares degrees of freedom
df_ess	error sum of squares degrees of freedom
df_lof	lack of fit degrees of freedom
df_error	pure error degrees of freedom
final	data.frame; contains computed values used for the lack of fit f test
resp	character vector; name of response variable
preds	character vector; name of predictor variable

**Note**

The lack of fit F test works only with simple linear regression. Moreover, it is important that the data contains repeat observations i.e. replicates for at least one of the values of the predictor x. This test generally only applies to datasets with plenty of replicates.

**References**

Kutner, MH, Nachtsheim CJ, Neter J and Li W., 2004, Applied Linear Statistical Models (5th edition). Chicago, IL., McGraw Hill/Irwin.

**Examples**

```
model <- lm(mpg ~ disp, data = mtcars)
ols_pure_error_anova(model)
```

---

ols\_regress                      *Ordinary least squares regression*

---

**Description**

Ordinary least squares regression.

**Usage**

```
ols_regress(object, ...)
```

```
## S3 method for class 'lm'
```

```
ols_regress(object, ...)
```

**Arguments**

object	An object of class "formula" (or one that can be coerced to that class): a symbolic description of the model to be fitted or class <code>lm</code> .
...	Other inputs.

**Value**

ols\_regress returns an object of class "ols\_regress". An object of class "ols\_regress" is a list containing the following components:

r	square root of rsquare, correlation between observed and predicted values of dependent variable
rsq	coefficient of determination or r-square
adjr	adjusted rsquare
sigma	root mean squared error
cv	coefficient of variation
mse	mean squared error
mae	mean absolute error
aic	akaike information criteria
sbc	bayesian information criteria
sbic	sawa bayesian information criteria
prsq	predicted rsquare
error_df	residual degrees of freedom
model_df	regression degrees of freedom
total_df	total degrees of freedom
ess	error sum of squares
rss	regression sum of squares
tss	total sum of squares
rms	regression mean square
ems	error mean square
f	f statistis
p	p-value for f
n	number of predictors including intercept
betas	betas; estimated coefficients
sbetas	standardized betas
std_errors	standard errors
tvalues	t values
pvalues	p-value of tvalues
df	degrees of freedom of betas
conf_lm	confidence intervals for coefficients
title	title for the model
dependent	character vector; name of the dependent variable
predictors	character vector; name of the predictor variables
mvars	character vector; name of the predictor variables including intercept
model	input model for ols_regress

## Interaction Terms

If the model includes interaction terms, the standardized betas are computed after scaling and centering the predictors.

## References

<https://www.ssc.wisc.edu/~hemken/Stataworkshops/stdBeta/Getting>

## Examples

```
ols_regress(mpg ~ disp + hp + wt, data = mtcars)

# if model includes interaction terms set iterm to TRUE
ols_regress(mpg ~ disp * wt, data = mtcars, iterm = TRUE)
```

---

ols_sbc	<i>Bayesian information criterion</i>
---------	---------------------------------------

---

## Description

Bayesian information criterion for model selection.

## Usage

```
ols_sbc(model, method = c("R", "STATA", "SAS"))
```

## Arguments

model	An object of class <code>lm</code> .
method	A character vector; specify the method to compute BIC. Valid options include R, STATA and SAS.

## Details

SBC provides a means for model selection. Given a collection of models for the data, SBC estimates the quality of each model, relative to each of the other models. R and STATA use loglikelihood to compute SBC. SAS uses residual sum of squares. Below is the formula in each case:

*R & STATA*

$$AIC = -2(\text{loglikelihood}) + \ln(n) * 2p$$

*SAS*

$$AIC = n * \ln(SSE/n) + p * \ln(n)$$

where  $n$  is the sample size and  $p$  is the number of model parameters including intercept.

**Value**

The bayesian information criterion of the model.

**References**

Schwarz, G. (1978). "Estimating the Dimension of a Model." *Annals of Statistics* 6:461–464.

Judge, G. G., Griffiths, W. E., Hill, R. C., and Lee, T.-C. (1980). *The Theory and Practice of Econometrics*. New York: John Wiley & Sons.

**See Also**

Other model selection criteria: [ols\\_aic](#), [ols\\_apc](#), [ols\\_fpe](#), [ols\\_hsp](#), [ols\\_mallows\\_cp](#), [ols\\_msep](#), [ols\\_sbic](#)

**Examples**

```
# using R computation method
model <- lm(mpg ~ disp + hp + wt + qsec, data = mtcars)
ols_sbc(model)

# using STATA computation method
model <- lm(mpg ~ disp + hp + wt + qsec, data = mtcars)
ols_sbc(model, method = 'STATA')

# using SAS computation method
model <- lm(mpg ~ disp + hp + wt + qsec, data = mtcars)
ols_sbc(model, method = 'SAS')
```

---

ols\_sbic

*Sawa's bayesian information criterion*

---

**Description**

Sawa's bayesian information criterion for model selection.

**Usage**

```
ols_sbc(model, full_model)
```

**Arguments**

model            An object of class lm.  
full\_model       An object of class lm.



**Details**

Sawa (1978) developed a model selection criterion that was derived from a Bayesian modification of the AIC criterion. Sawa's Bayesian Information Criterion (BIC) is a function of the number of observations  $n$ , the SSE, the pure error variance fitting the full model, and the number of independent variables including the intercept.

$$SBIC = n * \ln(SSE/n) + 2(p + 2)q - 2(q^2)$$

where  $q = n(\sigma^2)/SSE$ ,  $n$  is the sample size,  $p$  is the number of model parameters including intercept  $SSE$  is the residual sum of squares.

**Value**

Sawa's Bayesian Information Criterion

**References**

Sawa, T. (1978). "Information Criteria for Discriminating among Alternative Regression Models." *Econometrica* 46:1273–1282.

Judge, G. G., Griffiths, W. E., Hill, R. C., and Lee, T.-C. (1980). *The Theory and Practice of Econometrics*. New York: John Wiley & Sons.

**See Also**

Other model selection criteria: [ols\\_aic](#), [ols\\_apc](#), [ols\\_fpe](#), [ols\\_hsp](#), [ols\\_mallows\\_cp](#), [ols\\_msep](#), [ols\\_sbc](#)

**Examples**

```
full_model <- lm(mpg ~ ., data = mtcars)
model <- lm(mpg ~ disp + hp + wt + qsec, data = mtcars)
ols_sbic(model, full_model)
```

---

ols\_step\_all\_possible *All possible regression*

---

**Description**

Fits all regressions involving one regressor, two regressors, three regressors, and so on. It tests all possible subsets of the set of potential independent variables.

**Usage**

```
ols_step_all_possible(model, ...)

## S3 method for class 'ols_step_all_possible'
plot(x, model = NA, print_plot = TRUE,
     ...)
```

**Arguments**

model	An object of class lm.
...	Other arguments.
x	An object of class ols_best_subset.
print_plot	logical; if TRUE, prints the plot else returns a plot object.

**Value**

ols\_step\_all\_possible returns an object of class "ols\_step\_all\_possible". An object of class "ols\_step\_all\_possible" is a data frame containing the following components:

n	model number
predictors	predictors in the model
rsquare	rsquare of the model
adjr	adjusted rsquare of the model
predrsq	predicted rsquare of the model
cp	mallow's Cp
aic	akaike information criteria
sbic	sawa bayesian information criteria
sbc	schwarz bayes information criteria
gmsep	estimated MSE of prediction, assuming multivariate normality
jp	final prediction error
pc	amemiya prediction criteria
sp	hocking's Sp

**Deprecated Function**

ols\_all\_subset() has been deprecated. Instead use ols\_step\_all\_possible().

**References**

Mendenhall William and Sinsich Terry, 2012, A Second Course in Statistics Regression Analysis (7th edition). Prentice Hall

**See Also**

Other variable selection procedures: [ols\\_step\\_backward\\_aic](#), [ols\\_step\\_backward\\_p](#), [ols\\_step\\_best\\_subset](#), [ols\\_step\\_both\\_aic](#), [ols\\_step\\_forward\\_aic](#), [ols\\_step\\_forward\\_p](#)

**Examples**

```
model <- lm(mpg ~ disp + hp, data = mtcars)
k <- ols_step_all_possible(model)
k

# plot
plot(k)
```

---

ols\_step\_all\_possible\_betas

*All possible regression variable coefficients*

---

**Description**

Returns the coefficients for each variable from each model.

**Usage**

```
ols_step_all_possible_betas(object, ...)
```

**Arguments**

object	An object of class lm.
...	Other arguments.

**Value**

ols\_step\_all\_possible\_betas returns a data.frame containing:

model_index	model number
predictor	predictor
beta_coef	coefficient for the predictor

**Examples**

```
## Not run:
model <- lm(mpg ~ disp + hp + wt, data = mtcars)
ols_step_all_possible_betas(model)

## End(Not run)
```

---

ols\_step\_backward\_aic *Stepwise AIC backward regression*

---

### Description

Build regression model from a set of candidate predictor variables by removing predictors based on akaike information criterion, in a stepwise manner until there is no variable left to remove any more.

### Usage

```
ols_step_backward_aic(model, ...)

## Default S3 method:
ols_step_backward_aic(model, progress = FALSE,
  details = FALSE, ...)

## S3 method for class 'ols_step_backward_aic'
plot(x, print_plot = TRUE, ...)
```

### Arguments

model	An object of class <code>lm</code> ; the model should include all candidate predictor variables.
...	Other arguments.
progress	Logical; if TRUE, will display variable selection progress.
details	Logical; if TRUE, will print the regression result at each step.
x	An object of class <code>ols_step_backward_aic</code> .
print_plot	logical; if TRUE, prints the plot else returns a plot object.

### Value

`ols_step_backward_aic` returns an object of class `"ols_step_backward_aic"`. An object of class `"ols_step_backward_aic"` is a list containing the following components:

model	model with the least AIC; an object of class <code>lm</code>
steps	total number of steps
predictors	variables removed from the model
aics	akaike information criteria
ess	error sum of squares
rss	regression sum of squares
rsq	rsquare
arsq	adjusted rsquare

**Deprecated Function**

ols\_stepaic\_backward() has been deprecated. Instead use ols\_step\_backward\_aic().

**References**

Venables, W. N. and Ripley, B. D. (2002) Modern Applied Statistics with S. Fourth edition. Springer.

**See Also**

Other variable selection procedures: [ols\\_step\\_all\\_possible](#), [ols\\_step\\_backward\\_p](#), [ols\\_step\\_best\\_subset](#), [ols\\_step\\_both\\_aic](#), [ols\\_step\\_forward\\_aic](#), [ols\\_step\\_forward\\_p](#)

**Examples**

```
# stepwise backward regression
model <- lm(y ~ ., data = surgical)
ols_step_backward_aic(model)

# stepwise backward regression plot
model <- lm(y ~ ., data = surgical)
k <- ols_step_backward_aic(model)
plot(k)

# final model
k$model
```

---

ols\_step\_backward\_p    *Stepwise backward regression*

---

**Description**

Build regression model from a set of candidate predictor variables by removing predictors based on p values, in a stepwise manner until there is no variable left to remove any more.

**Usage**

```
ols_step_backward_p(model, ...)
```

## Default S3 method:

```
ols_step_backward_p(model, prem = 0.3,
  progress = FALSE, details = FALSE, ...)
```

## S3 method for class 'ols\_step\_backward\_p'

```
plot(x, model = NA, print_plot = TRUE,
  ...)
```

**Arguments**

<code>model</code>	An object of class <code>lm</code> ; the model should include all candidate predictor variables.
<code>...</code>	Other inputs.
<code>prem</code>	p value; variables with p more than <code>prem</code> will be removed from the model.
<code>progress</code>	Logical; if TRUE, will display variable selection progress.
<code>details</code>	Logical; if TRUE, will print the regression result at each step.
<code>x</code>	An object of class <code>ols_step_backward_p</code> .
<code>print_plot</code>	logical; if TRUE, prints the plot else returns a plot object.

**Value**

`ols_step_backward_p` returns an object of class `"ols_step_backward_p"`. An object of class `"ols_step_backward_p"` is a list containing the following components:

<code>model</code>	final model; an object of class <code>lm</code>
<code>steps</code>	total number of steps
<code>removed</code>	variables removed from the model
<code>rsquare</code>	coefficient of determination
<code>aic</code>	akaike information criteria
<code>sbc</code>	bayesian information criteria
<code>sbic</code>	sawa's bayesian information criteria
<code>adjr</code>	adjusted r-square
<code>rmse</code>	root mean square error
<code>mallows_cp</code>	mallow's Cp
<code>indvar</code>	predictors

**Deprecated Function**

`ols_step_backward()` has been deprecated. Instead use `ols_step_backward_p()`.

**References**

Chatterjee, Samprit and Hadi, Ali. Regression Analysis by Example. 5th ed. N.p.: John Wiley & Sons, 2012. Print.

**See Also**

Other variable selection procedures: [ols\\_step\\_all\\_possible](#), [ols\\_step\\_backward\\_aic](#), [ols\\_step\\_best\\_subset](#), [ols\\_step\\_both\\_aic](#), [ols\\_step\\_forward\\_aic](#), [ols\\_step\\_forward\\_p](#)

**Examples**

```
# stepwise backward regression
model <- lm(y ~ ., data = surgical)
ols_step_backward_p(model)

# stepwise backward regression plot
model <- lm(y ~ ., data = surgical)
k <- ols_step_backward_p(model)
plot(k)

# final model
k$model
```

---

ols\_step\_best\_subset *Best subsets regression*

---

**Description**

Select the subset of predictors that do the best at meeting some well-defined objective criterion, such as having the largest R<sup>2</sup> value or the smallest MSE, Mallows's Cp or AIC.

**Usage**

```
ols_step_best_subset(model, ...)
```

```
## S3 method for class 'ols_step_best_subset'
plot(x, model = NA, print_plot = TRUE,
     ...)
```

**Arguments**

model	An object of class lm.
...	Other inputs.
x	An object of class ols_step_best_subset.
print_plot	logical; if TRUE, prints the plot else returns a plot object.

**Value**

ols\_step\_best\_subset returns an object of class "ols\_step\_best\_subset". An object of class "ols\_step\_best\_subset" is a data frame containing the following components:

n	model number
predictors	predictors in the model
rsquare	rsquare of the model
adjr	adjusted rsquare of the model

predrsq	predicted rsquare of the model
cp	mallow's Cp
aic	akaike information criteria
sbic	sawa bayesian information criteria
sbc	schwarz bayes information criteria
gmsep	estimated MSE of prediction, assuming multivariate normality
jp	final prediction error
pc	amemiya prediction criteria
sp	hocking's Sp

### Deprecated Function

`ols_best_subset()` has been deprecated. Instead use `ols_step_best_subset()`.

### References

Kutner, MH, Nachtsheim CJ, Neter J and Li W., 2004, Applied Linear Statistical Models (5th edition). Chicago, IL., McGraw Hill/Irwin.

### See Also

Other variable selection procedures: [ols\\_step\\_all\\_possible](#), [ols\\_step\\_backward\\_aic](#), [ols\\_step\\_backward\\_p](#), [ols\\_step\\_both\\_aic](#), [ols\\_step\\_forward\\_aic](#), [ols\\_step\\_forward\\_p](#)

### Examples

```
model <- lm(mpg ~ disp + hp + wt + qsec, data = mtcars)
ols_step_best_subset(model)

# plot
model <- lm(mpg ~ disp + hp + wt + qsec, data = mtcars)
k <- ols_step_best_subset(model)
plot(k)
```

---

ols\_step\_both\_aic      *Stepwise AIC regression*

---

### Description

Build regression model from a set of candidate predictor variables by entering and removing predictors based on akaike information criteria, in a stepwise manner until there is no variable left to enter or remove any more.



**Usage**

```
ols_step_both_aic(model, progress = FALSE, details = FALSE)

## S3 method for class 'ols_step_both_aic'
plot(x, print_plot = TRUE, ...)
```

**Arguments**

model	An object of class <code>lm</code> .
progress	Logical; if TRUE, will display variable selection progress.
details	Logical; if TRUE, details of variable selection will be printed on screen.
x	An object of class <code>ols_step_both_aic</code> .
print_plot	logical; if TRUE, prints the plot else returns a plot object.
...	Other arguments.

**Value**

`ols_step_both_aic` returns an object of class `"ols_step_both_aic"`. An object of class `"ols_step_both_aic"` is a list containing the following components:

model	model with the least AIC; an object of class <code>lm</code>
predictors	variables added/removed from the model
method	addition/deletion
aics	akaike information criteria
ess	error sum of squares
rss	regression sum of squares
rsq	rsquare
arsq	adjusted rsquare
steps	total number of steps

**Deprecated Function**

`ols_stepaic_both()` has been deprecated. Instead use `ols_step_both_aic()`.

**References**

Venables, W. N. and Ripley, B. D. (2002) *Modern Applied Statistics with S*. Fourth edition. Springer.

**See Also**

Other variable selection procedures: [ols\\_step\\_all\\_possible](#), [ols\\_step\\_backward\\_aic](#), [ols\\_step\\_backward\\_p](#), [ols\\_step\\_best\\_subset](#), [ols\\_step\\_forward\\_aic](#), [ols\\_step\\_forward\\_p](#)

**Examples**

```
## Not run:
# stepwise regression
model <- lm(y ~ ., data = stepdata)
ols_step_both_aic(model)

# stepwise regression plot
model <- lm(y ~ ., data = stepdata)
k <- ols_step_both_aic(model)
plot(k)

# final model
k$model

## End(Not run)
```

---

ols_step_both_p	<i>Stepwise regression</i>
-----------------	----------------------------

---

**Description**

Build regression model from a set of candidate predictor variables by entering and removing predictors based on p values, in a stepwise manner until there is no variable left to enter or remove any more.

**Usage**

```
ols_step_both_p(model, ...)

## Default S3 method:
ols_step_both_p(model, pent = 0.1, prem = 0.3,
  progress = FALSE, details = FALSE, ...)

## S3 method for class 'ols_step_both_p'
plot(x, model = NA, print_plot = TRUE, ...)
```

**Arguments**

model	An object of class <code>lm</code> ; the model should include all candidate predictor variables.
...	Other arguments.
pent	p value; variables with p value less than pent will enter into the model.
prem	p value; variables with p more than prem will be removed from the model.
progress	Logical; if TRUE, will display variable selection progress.
details	Logical; if TRUE, will print the regression result at each step.
x	An object of class <code>ols_step_both_p</code> .
print_plot	logical; if TRUE, prints the plot else returns a plot object.

**Value**

ols\_step\_both\_p returns an object of class "ols\_step\_both\_p". An object of class "ols\_step\_both\_p" is a list containing the following components:

model	final model; an object of class lm
orders	candidate predictor variables according to the order by which they were added or removed from the model
method	addition/deletion
steps	total number of steps
predictors	variables retained in the model (after addition)
rsquare	coefficient of determination
aic	akaike information criteria
sbc	bayesian information criteria
sbic	sawa's bayesian information criteria
adjr	adjusted r-square
rmse	root mean square error
mallows_cp	mallow's Cp
indvar	predictors

**Deprecated Function**

ols\_stepwise() has been deprecated. Instead use ols\_step\_both\_p().

**References**

Chatterjee, Samprit and Hadi, Ali. Regression Analysis by Example. 5th ed. N.p.: John Wiley & Sons, 2012. Print.

**Examples**

```
# stepwise regression
model <- lm(y ~ ., data = surgical)
ols_step_both_p(model)

# stepwise regression plot
model <- lm(y ~ ., data = surgical)
k <- ols_step_both_p(model)
plot(k)

# final model
k$model
```

---

ols\_step\_forward\_aic *Stepwise AIC forward regression*

---

### Description

Build regression model from a set of candidate predictor variables by entering predictors based on akaike information criterion, in a stepwise manner until there is no variable left to enter any more.

### Usage

```
ols_step_forward_aic(model, ...)

## Default S3 method:
ols_step_forward_aic(model, progress = FALSE,
  details = FALSE, ...)

## S3 method for class 'ols_step_forward_aic'
plot(x, print_plot = TRUE, ...)
```

### Arguments

model	An object of class <code>lm</code> .
...	Other arguments.
progress	Logical; if TRUE, will display variable selection progress.
details	Logical; if TRUE, will print the regression result at each step.
x	An object of class <code>ols_step_forward_aic</code> .
print_plot	logical; if TRUE, prints the plot else returns a plot object.

### Value

`ols_step_forward_aic` returns an object of class `"ols_step_forward_aic"`. An object of class `"ols_step_forward_aic"` is a list containing the following components:

model	model with the least AIC; an object of class <code>lm</code>
steps	total number of steps
predictors	variables added to the model
aics	akaike information criteria
ess	error sum of squares
rss	regression sum of squares
rsq	rsquare
arsq	adjusted rsquare

### Deprecated Function

`ols_stepaic_forward()` has been deprecated. Instead use `ols_step_forward_aic()`.

## References

Venables, W. N. and Ripley, B. D. (2002) Modern Applied Statistics with S. Fourth edition. Springer.

## See Also

Other variable selection procedures: [ols\\_step\\_all\\_possible](#), [ols\\_step\\_backward\\_aic](#), [ols\\_step\\_backward\\_p](#), [ols\\_step\\_best\\_subset](#), [ols\\_step\\_both\\_aic](#), [ols\\_step\\_forward\\_p](#)

## Examples

```
# stepwise forward regression
model <- lm(y ~ ., data = surgical)
ols_step_forward_aic(model)

# stepwise forward regression plot
model <- lm(y ~ ., data = surgical)
k <- ols_step_forward_aic(model)
plot(k)

# final model
k$model
```

---

ols\_step\_forward\_p      *Stepwise forward regression*

---

## Description

Build regression model from a set of candidate predictor variables by entering predictors based on  $p$  values, in a stepwise manner until there is no variable left to enter any more.

## Usage

```
ols_step_forward_p(model, ...)
```

## Default S3 method:

```
ols_step_forward_p(model, penter = 0.3,
  progress = FALSE, details = FALSE, ...)
```

## S3 method for class 'ols\_step\_forward\_p'

```
plot(x, model = NA, print_plot = TRUE,
  ...)
```

**Arguments**

model	An object of class <code>lm</code> ; the model should include all candidate predictor variables.
...	Other arguments.
penter	p value; variables with p value less than penter will enter into the model
progress	Logical; if TRUE, will display variable selection progress.
details	Logical; if TRUE, will print the regression result at each step.
x	An object of class <code>ols_step_forward_p</code> .
print_plot	logical; if TRUE, prints the plot else returns a plot object.

**Value**

`ols_step_forward_p` returns an object of class "`ols_step_forward_p`". An object of class "`ols_step_forward_p`" is a list containing the following components:

model	final model; an object of class <code>lm</code>
steps	number of steps
predictors	variables added to the model
rsquare	coefficient of determination
aic	akaike information criteria
sbc	bayesian information criteria
sbic	sawa's bayesian information criteria
adjr	adjusted r-square
rmse	root mean square error
mallows_cp	mallow's Cp
indvar	predictors

**Deprecated Function**

`ols_step_forward()` has been deprecated. Instead use `ols_step_forward_p()`.

**References**

Chatterjee, Samprit and Hadi, Ali. Regression Analysis by Example. 5th ed. N.p.: John Wiley & Sons, 2012. Print.

Kutner, MH, Nachtsheim CJ, Neter J and Li W., 2004, Applied Linear Statistical Models (5th edition). Chicago, IL., McGraw Hill/Irwin.

**See Also**

Other variable selection procedures: [ols\\_step\\_all\\_possible](#), [ols\\_step\\_backward\\_aic](#), [ols\\_step\\_backward\\_p](#), [ols\\_step\\_best\\_subset](#), [ols\\_step\\_both\\_aic](#), [ols\\_step\\_forward\\_aic](#)

**Examples**

```
# stepwise forward regression
model <- lm(y ~ ., data = surgical)
ols_step_forward_p(model)

# stepwise forward regression plot
model <- lm(y ~ ., data = surgical)
k <- ols_step_forward_p(model)
plot(k)

# final model
k$model
```

---

ols_test_bartlett	<i>Bartlett test</i>
-------------------	----------------------

---

**Description**

Test if k samples are from populations with equal variances.

**Usage**

```
ols_test_bartlett(data, ...)

## Default S3 method:
ols_test_bartlett(data, ..., group_var = NULL)
```

**Arguments**

data	A data.frame or tibble.
...	Columns in data.
group_var	Grouping variable.

**Details**

Bartlett's test is used to test if variances across samples is equal. It is sensitive to departures from normality. The Levene test is an alternative test that is less sensitive to departures from normality.

**Value**

ols\_test\_bartlett returns an object of class "ols\_test\_bartlett". An object of class "ols\_test\_bartlett" is a list containing the following components:

fstat	f statistic
pval	p-value of fstat
df	degrees of freedom

**Deprecated Function**

ols\_bartlett\_test() has been deprecated. Instead use ols\_test\_bartlett().

**References**

Snedecor, George W. and Cochran, William G. (1989), Statistical Methods, Eighth Edition, Iowa State University Press.

**See Also**

Other heteroskedasticity tests: [ols\\_test\\_breusch\\_pagan](#), [ols\\_test\\_f](#), [ols\\_test\\_score](#)

**Examples**

```
# using grouping variable
library(descriptor)
ols_test_bartlett(mtcars, 'mpg', group_var = 'cyl')

# using variables
ols_test_bartlett(hsb, 'read', 'write')
```

---

```
ols_test_breusch_pagan
```

```
Breusch pagan test
```

---

**Description**

Test for constant variance. It assumes that the error terms are normally distributed.

**Usage**

```
ols_test_breusch_pagan(model, fitted.values = TRUE, rhs = FALSE,
  multiple = FALSE, p.adj = c("none", "bonferroni", "sidak", "holm"),
  vars = NA)
```

**Arguments**

model	An object of class lm.
fitted.values	Logical; if TRUE, use fitted values of regression model.
rhs	Logical; if TRUE, specifies that tests for heteroskedasticity be performed for the right-hand-side (explanatory) variables of the fitted regression model.
multiple	Logical; if TRUE, specifies that multiple testing be performed.
p.adj	Adjustment for p value, the following options are available: bonferroni, holm, sidak and none.
vars	Variables to be used for heteroskedasticity test.



**Details**

Breusch Pagan Test was introduced by Trevor Breusch and Adrian Pagan in 1979. It is used to test for heteroskedasticity in a linear regression model. It test whether variance of errors from a regression is dependent on the values of a independent variable.

- Null Hypothesis: Equal/constant variances
- Alternative Hypothesis: Unequal/non-constant variances

**Computation**

- Fit a regression model
- Regress the squared residuals from the above model on the independent variables
- Compute  $nR^2$ . It follows a chi square distribution with  $p - 1$  degrees of freedom, where  $p$  is the number of independent variables,  $n$  is the sample size and  $R^2$  is the coefficient of determination from the regression in step 2.

**Value**

ols\_test\_breusch\_pagan returns an object of class "ols\_test\_breusch\_pagan". An object of class "ols\_test\_breusch\_pagan" is a list containing the following components:

bp	breusch pagan statistic
p	p-value of bp
fv	fitted values of the regression model
rhs	names of explanatory variables of fitted regression model
multiple	logical value indicating if multiple tests should be performed
padj	adjusted p values
vars	variables to be used for heteroskedasticity test
resp	response variable
preds	predictors

**Deprecated Function**

ols\_bp\_test() has been deprecated. Instead use ols\_test\_breusch\_pagan().

**References**

T.S. Breusch & A.R. Pagan (1979), A Simple Test for Heteroscedasticity and Random Coefficient Variation. *Econometrica* 47, 1287–1294

Cook, R. D.; Weisberg, S. (1983). "Diagnostics for Heteroskedasticity in Regression". *Biometrika*. 70 (1): 1–10.

**See Also**

Other heteroskedasticity tests: [ols\\_test\\_bartlett](#), [ols\\_test\\_f](#), [ols\\_test\\_score](#)

## Examples

```
# model
model <- lm(mpg ~ disp + hp + wt + drat, data = mtcars)

# use fitted values of the model
ols_test_breusch_pagan(model)

# use independent variables of the model
ols_test_breusch_pagan(model, rhs = TRUE)

# use independent variables of the model and perform multiple tests
ols_test_breusch_pagan(model, rhs = TRUE, multiple = TRUE)

# bonferroni p value adjustment
ols_test_breusch_pagan(model, rhs = TRUE, multiple = TRUE, p.adj = 'bonferroni')

# sidak p value adjustment
ols_test_breusch_pagan(model, rhs = TRUE, multiple = TRUE, p.adj = 'sidak')

# holm's p value adjustment
ols_test_breusch_pagan(model, rhs = TRUE, multiple = TRUE, p.adj = 'holm')
```

---

ols\_test\_correlation *Correlation test for normality*

---

## Description

Correlation between observed residuals and expected residuals under normality.

## Usage

```
ols_test_correlation(model)
```

## Arguments

model            An object of class lm.

## Value

Correlation between fitted regression model residuals and expected values of residuals.

## Deprecated Function

ols\_corr\_test() has been deprecated. Instead use ols\_test\_correlation().

## See Also

Other residual diagnostics: [ols\\_plot\\_resid\\_box](#), [ols\\_plot\\_resid\\_fit](#), [ols\\_plot\\_resid\\_hist](#), [ols\\_plot\\_resid\\_qq](#), [ols\\_test\\_normality](#)

**Examples**

```
model <- lm(mpg ~ disp + hp + wt + qsec, data = mtcars)
ols_test_correlation(model)
```

---

ols_test_f	<i>F test</i>
------------	---------------

---

**Description**

Test for heteroskedasticity under the assumption that the errors are independent and identically distributed (i.i.d.).

**Usage**

```
ols_test_f(model, fitted_values = TRUE, rhs = FALSE, vars = NULL,
  ...)
```

**Arguments**

model	An object of class <code>lm</code> .
fitted_values	Logical; if TRUE, use fitted values of regression model.
rhs	Logical; if TRUE, specifies that tests for heteroskedasticity be performed for the right-hand-side (explanatory) variables of the fitted regression model.
vars	Variables to be used for for heteroskedasticity test.
...	Other arguments.

**Value**

`ols_test_f` returns an object of class `"ols_test_f"`. An object of class `"ols_test_f"` is a list containing the following components:

f	f statistic
p	p-value of f
fv	fitted values of the regression model
rhs	names of explanatory variables of fitted regression model
numdf	numerator degrees of freedom
dendf	denominator degrees of freedom
vars	variables to be used for heteroskedasticity test
resp	response variable
preds	predictors

**Deprecated Function**

`ols_f_test()` has been deprecated. Instead use `ols_test_f()`.

## References

Wooldridge, J. M. 2013. *Introductory Econometrics: A Modern Approach*. 5th ed. Mason, OH: South-Western.

## See Also

Other heteroskedasticity tests: [ols\\_test\\_bartlett](#), [ols\\_test\\_breusch\\_pagan](#), [ols\\_test\\_score](#)

## Examples

```
# model
model <- lm(mpg ~ disp + hp + wt + qsec, data = mtcars)

# using fitted values
ols_test_f(model)

# using all predictors of the model
ols_test_f(model, rhs = TRUE)

# using fitted values
ols_test_f(model, vars = c('disp', 'hp'))
```

---

ols\_test\_normality      *Test for normality*

---

## Description

Test for detecting violation of normality assumption.

## Usage

```
ols_test_normality(y, ...)
```

## S3 method for class 'lm'

```
ols_test_normality(y, ...)
```

## Arguments

`y`                    A numeric vector or an object of class `lm`.

`...`                Other arguments.

**Value**

ols\_test\_normality returns an object of class "ols\_test\_normality". An object of class "ols\_test\_normality" is a list containing the following components:

kolmogorv	kolmogorv smirnov statistic
shapiro	shapiro wilk statistic
cramer	cramer von mises statistic
anderson	anderson darling statistic

**Deprecated Function**

ols\_norm\_test() has been deprecated. Instead use ols\_test\_normality().

**See Also**

Other residual diagnostics: [ols\\_plot\\_resid\\_box](#), [ols\\_plot\\_resid\\_fit](#), [ols\\_plot\\_resid\\_hist](#), [ols\\_plot\\_resid\\_qq](#), [ols\\_test\\_correlation](#)

**Examples**

```
model <- lm(mpg ~ disp + hp + wt + qsec, data = mtcars)
ols_test_normality(model)
```

---

ols\_test\_outlier      *Bonferroni Outlier Test*

---

**Description**

Detect outliers using Bonferroni p values.

**Usage**

```
ols_test_outlier(model, cut_off = 0.05, n_max = 10, ...)
```

**Arguments**

model	An object of class lm.
cut_off	Bonferroni p-values cut off for reporting observations.
n_max	Maximum number of observations to report, default is 10.
...	Other arguments.

**Examples**

```
# model
model <- lm(y ~ ., data = surgical)
ols_test_outlier(model)
```

---

ols_test_score	<i>Score test</i>
----------------	-------------------

---

### Description

Test for heteroskedasticity under the assumption that the errors are independent and identically distributed (i.i.d.).

### Usage

```
ols_test_score(model, fitted_values = TRUE, rhs = FALSE, vars = NULL)
```

### Arguments

model	An object of class <code>lm</code> .
fitted_values	Logical; if TRUE, use fitted values of regression model.
rhs	Logical; if TRUE, specifies that tests for heteroskedasticity be performed for the right-hand-side (explanatory) variables of the fitted regression model.
vars	Variables to be used for for heteroskedasticity test.

### Value

`ols_test_score` returns an object of class `"ols_test_score"`. An object of class `"ols_test_score"` is a list containing the following components:

score	f statistic
p	p value of score
df	degrees of freedom
fv	fitted values of the regression model
rhs	names of explanatory variables of fitted regression model
resp	response variable
preds	predictors

### Deprecated Function

`ols_score_test()` has been deprecated. Instead use `ols_test_score()`.

### References

Breusch, T. S. and Pagan, A. R. (1979) A simple test for heteroscedasticity and random coefficient variation. *Econometrica* 47, 1287–1294.

Cook, R. D. and Weisberg, S. (1983) Diagnostics for heteroscedasticity in regression. *Biometrika* 70, 1–10.

Koenker, R. 1981. A note on studentizing a test for heteroskedasticity. *Journal of Econometrics* 17: 107–112.

**See Also**

Other heteroskedasticity tests: [ols\\_test\\_bartlett](#), [ols\\_test\\_breusch\\_pagan](#), [ols\\_test\\_f](#)

**Examples**

```
# model
model <- lm(mpg ~ disp + hp + wt, data = mtcars)

# using fitted values of the model
ols_test_score(model)

# using predictors from the model
ols_test_score(model, rhs = TRUE)

# specify predictors from the model
ols_test_score(model, vars = c('disp', 'wt'))
```

---

rivers

*Test Data Set*

---

**Description**

Test Data Set

**Usage**

rivers

**Format**

An object of class `data.frame` with 20 rows and 6 columns.

---

rvsr\_plot\_shiny

*Residual vs regressors plot for shiny app*

---

**Description**

Graph to determine whether we should add a new predictor to the model already containing other predictors. The residuals from the model is regressed on the new predictor and if the plot shows non random pattern, you should consider adding the new predictor to the model.

**Usage**

```
rvsr_plot_shiny(model, data, variable, print_plot = TRUE)
```

**Arguments**

model	An object of class <code>lm</code> .
data	A <code>data.frame</code> or <code>tibble</code> .
variable	Character; new predictor to be added to the model.
print_plot	logical; if TRUE, prints the plot else returns a plot object.

**Examples**

```
model <- lm(mpg ~ disp + hp + wt, data = mtcars)
rvsr_plot_shiny(model, mtcars, 'drat')
```

---

stepdata	<i>Test Data Set</i>
----------	----------------------

---

**Description**

Test Data Set

**Usage**

```
stepdata
```

**Format**

An object of class `data.frame` with 20000 rows and 7 columns.

---

surgical	<i>Surgical Unit Data Set</i>
----------	-------------------------------

---

**Description**

A dataset containing data about survival of patients undergoing liver operation.

**Usage**

```
surgical
```



**Format**

A data frame with 54 rows and 9 variables:

**bcs** blood clotting score

**pindex** prognostic index

**enzyme\_test** enzyme function test score

**liver\_test** liver function test score

**age** age, in years

**gender** indicator variable for gender (0 = male, 1 = female)

**alc\_mod** indicator variable for history of alcohol use (0 = None, 1 = Moderate)

**alc\_heavy** indicator variable for history of alcohol use (0 = None, 1 = Heavy)

**y** Survival Time

**Source**

Kutner, MH, Nachtsheim CJ, Neter J and Li W., 2004, Applied Linear Statistical Models (5th edition). Chicago, IL., McGraw Hill/Irwin.

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