

# Package ‘plotor’

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**Type** Package

**Title** Odds Ratio Tools for Logistic Regression

**Version** 0.8.0

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**Description** Produces odds ratio analyses with comprehensive reporting tools. Generates plots, summary tables, and diagnostic checks for logistic regression models fitted with 'glm()' using binomial family. Provides visualisation methods, formatted reporting tables via 'gt', and tools to assess logistic regression model assumptions.

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**Encoding** UTF-8

**Imports** broom, car, cli, detectseparation, dplyr, forcats, ggplot2, glue, gt, janitor, purrr, rlang, scales, stats, stringr, tibble, tidyr, tidyselect

**RoxygenNote** 7.3.3

**Suggests** datasets, knitr, labelled, magrittr, MASS, medicaldata (>= 0.2.0.9001), NHANES, R4HCR, rmarkdown, svglite, testthat (>= 3.0.0), vdiff

**VignetteBuilder** knitr

**URL** <https://github.com/craig-parylo/plotor>,  
<https://craig-parylo.github.io/plotor/>

**BugReports** <https://github.com/craig-parylo/plotor/issues>

**Config/testthat/edition** 3

**Depends** R (>= 4.1.0)

**Config/Remotes** higgil3425/medicaldata

**NeedsCompilation** no

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**Repository** CRAN

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check_or	<i>Check OR</i>
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Description

Performs a series of tests to ensure that assumptions for logistic regression are met, with optional detailed feedback if any tests fail.

Usage

```
check_or(glm_model_results, confint_fast_estimate = FALSE, details = TRUE)
```

Arguments

- glm\_model\_results  
Results from a binomial Generalised Linear Model (GLM), as produced by `stats::glm()`.
- confint\_fast\_estimate  
Boolean (default = FALSE) Use a faster estimate of the confidence interval?  
Note: this assumes normally distributed data, which may not be suitable for your data.
- details  
Boolean (default = TRUE) Show detailed feedback for any failed tests?

Value

Logical, TRUE if all assumption tests pass, FALSE if one or more tests fail

Examples

```
# Load the Titanic dataset
df <- datasets::Titanic |>
  dplyr::as_tibble() |>
  # convert aggregated counts to individual observations
  dplyr::filter(n > 0) |>
  tidyr::uncount(weights = n) |>
  # convert character variables to factors
  dplyr::mutate(dplyr::across(dplyr::where(is.character), as.factor))

# Perform logistic regression using `glm`
lr <- stats::glm(
  data = df,
```

```

    family = binomial,
    formula = Survived ~ Class + Sex + Age
  )

  # Check the model for logistic regression assumption violations
  check_or(lr)

```

plot\_or

*Plot OR*

## Description

Produces an Odds Ratio plot to visualise the results of a logistic regression analysis.

## Usage

```

plot_or(
  glm_model_results,
  conf_level = 0.95,
  confint_fast_estimate = FALSE,
  assumption_checks = TRUE
)

```

## Arguments

glm_model_results	Results from a binomial Generalised Linear Model (GLM), as produced by <code>stats::glm()</code> .
conf_level	Numeric value between 0.001 and 0.999 (default = 0.95) specifying the confidence level for the confidence interval.
confint_fast_estimate	Boolean (default = FALSE) indicating whether to use a faster estimate of the confidence interval. Note: this assumes normally distributed data, which may not be suitable for your data.
assumption_checks	Boolean (default = TRUE) indicating whether to conduct checks to ensure that the assumptions of logistic regression are met.

## Value

The function returns an object of class `gg` and `ggplot`, which can be customised and extended using various `ggplot2` functions.

## See Also

- See vignette('using\_plotlor', package = 'plotlor') for more details on usage.
- More details and examples can be found on the website: <https://craig-parylo.github.io/plotlor/index.html>

## Examples

```
# Load required libraries
library(plotor)
library(datasets)
library(dplyr)
library(ggplot2)
library(stats)
library(forcats)
library(tidyr)

# Load the Titanic dataset
df <- datasets::Titanic |>
  as_tibble() |>
  # convert aggregated counts to individual observations
  filter(n > 0) |>
  uncount(weights = n) |>
  # convert character variables to factors
  mutate(across(where(is.character), as.factor))

# Perform logistic regression using `glm`
lr <- glm(
  data = df,
  family = 'binomial',
  formula = Survived ~ Class + Sex + Age
)

# Produce the Odds Ratio plot
plot_or(lr)
```

---

table\_or

---

*Table OR*


---

## Description

Produces a formatted table displaying the outputs from the Odds Ratio analysis, including details on covariate characteristics and model results.

## Usage

```
table_or(
  glm_model_results,
  conf_level = 0.95,
  output = c("tibble", "gt"),
  output_type = c("multivariable", "uni_and_multi"),
  confint_fast_estimate = FALSE,
  assumption_checks = TRUE,
  anonymise_counts = FALSE,
  use_model_data_only = TRUE
)
```

## Arguments

<code>glm_model_results</code>	Results from a binomial Generalised Linear Model (GLM), as produced by <code>stats::glm()</code> .
<code>conf_level</code>	Numeric value between 0.001 and 0.999 (default = 0.95) specifying the confidence level for the confidence interval.
<code>output</code>	String describing the output type (default = "tibble"). Options include "tibble" and "gt".
<code>output_type</code>	String description of the output type (default = "multivariable"). Options include "multivariable" and "uni_and_multi". Selecting "multivariable" will produce a summary table of the supplied multivariable model. Selecting "uni_and_multi" will produce a summary table showing estimates of the Odds Ratio, Confidence Intervals and p-values produced using a univariable logistic regression model for each predictor along with the adjusted Odds Ratio, Confidence Intervals and p-values from the supplied multivariable model.
<code>confint_fast_estimate</code>	Boolean (default = FALSE) indicating whether to use a faster estimate of the confidence interval. Note: this assumes normally distributed data, which may not be suitable for your data.
<code>assumption_checks</code>	Boolean (default = TRUE) indicating whether to conduct checks to ensure that the assumptions of logistic regression are met.
<code>anonymise_counts</code>	Boolean (default = FALSE) indicating whether to anonymise counts in the output table. If TRUE, counts less than 10 are suppressed and otherwise rounded to the nearest multiple of 5.
<code>use_model_data_only</code>	Boolean (default = FALSE) indicating whether to use only the subset of data that was used as part of the multivariable model, or set to TRUE to use the full set of data provided to the multivariable model. Note, any records containing missing values for any of the outcome or predictor variables are automatically excluded from the multivariable model by <code>stats::glm</code> , so the overall number of records used in multivariable models can be much lower than the total number of records supplied to the function. Set to TRUE to increase comparability between the univariable and multivariable models, set to FALSE to gain a more holistic view of the individual relationships between predictors and outcome.

## Details

The table includes the following information:

- **Covariate Characteristics:**
  - Number of observations for each characteristic
  - Number of observations resulting in the outcome of interest
  - Conversion rate of the outcome based on the number of observations
- **Model Results:**

- Estimated Odds Ratio, standard error, and p-value
- Calculated confidence interval for the specified confidence level

A visualisation of the Odds Ratio plot is also provided for an at-a-glance view of the model results.

If `anonymise_counts` is set to `TRUE`, counts below 10 are suppressed as `<10`, and other counts are rounded to the nearest multiple of 5. This feature is helpful when working with sensitive data.

## Value

The returned object depends on the output parameter:

- If `output = 'tibble'`, the function returns an object of class `"tbl_df"`, `"tbl"`, and `"data.frame"`.
- If `output = 'gt'`, the function returns an object of class `"gt_tbl"` and `"list"`.

## Examples

```
# Load the Titanic dataset
df <- datasets::Titanic |>
  dplyr::as_tibble() |>
  # convert aggregated counts to individual observations
  dplyr::filter(n > 0) |>
  tidyr::uncount(weights = n) |>
  # convert character variables to factors
  dplyr::mutate(dplyr::across(dplyr::where(is.character), as.factor))

# Perform logistic regression using `glm`
lr <- stats::glm(
  data = df,
  family = 'binomial',
  formula = Survived ~ Class + Sex + Age
)

# Produce the Odds Ratio table as a tibble
table_or(lr)

# Produce the Odds Ratio table as a gt object
table_or(lr, output = 'gt')
```

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