

# Package ‘CIplot’

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**Type** Package

**Title** Functions to Plot Confidence Interval

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**Description** Plot confidence interval from the objects of statistical tests such as `t.test()`, `var.test()`, `cor.test()`, `prop.test()` and `fisher.test()` ('htest' class), Tukey test [`TukeyHSD()`], Dunnett test [`glht()` in 'multcomp' package], logistic regression [`glm()`], and Tukey or Games-Howell test [`posthocTGH()` in 'userfriendlyscience' package].  
Users are able to set the styles of lines and points.  
This package contains the function to calculate odds ratios and their confidence intervals from the result of logistic regression.

**Imports** MASS, multcomp

**Suggests** BSDA, fmsb, userfriendlyscience

**License** GPL (>= 2)

**URL** <https://github.com/toshi-ara/CIplot>

**BugReports** <https://github.com/toshi-ara/CIplot/issues/>

**RoxygenNote** 6.0.1

**Collate** 'CIplot.R' 'CIplot.default.R' 'CIplot.htest.R'  
'CIplot.TukeyHSD.R' 'CIplot.glht.R' 'CIplot.glm.R'  
'CIplot.ORci.R' 'CIplot.posthocTGH.R' 'ORci.R' 'print.ORci.R'

**NeedsCompilation** no

**Repository** CRAN

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CIplot

*Plot Confidential Interval***Description**

A function to plot confidential interval for such as `htest`, `TukeyHSD`, `glht` (**multcomp**), `glm` (logistic regression only!) and `posthocTGH` (**userfriendlyscience**) objects.

**Usage**

```
CIplot(x, ...)
```

```
## Default S3 method:
```

```
CIplot(x, xlog = FALSE, xlim = NULL, xlab = NULL,
       yname = TRUE, las = 0, pch = 21, pcol = 1, pcolbg = "white",
       pcex = 1, conf.level = 0.95, cilty = 1, cilwd = 1, cicol = 1, v,
       vltty = 2, vlwd = 1, vcol = 1, main = NULL, ...)
```

```
## S3 method for class 'htest'
```

```
CIplot(x, xlog = FALSE, xlim = NULL, xlab = NULL,
       yname = FALSE, v = NULL, ...)
```

```
## S3 method for class 'TukeyHSD'
```

```
CIplot(x, xlab = "Differences in mean", v = 0, ...)
```

```
## S3 method for class 'glht'
```

```
CIplot(x, xlab = "Differences in mean", v = 0, ...)
```

```
## S3 method for class 'glm'
```

```
CIplot(x, conf.level = 0.95, xlog = TRUE,
       xlab = "Odds Ratio", v = 1, ...)
```

```
## S3 method for class 'ORci'
```

```
CIplot(x, xlog = TRUE, xlab = "Odds Ratio", v = 1, ...)
```

```
## S3 method for class 'posthocTGH'
```

```
CIplot(x, xlab = "Differences in mean", v = 0, ...)
```

**Arguments**

<code>x</code>	default: matrix or data.frame class with 3 columns ('any name', <code>lwr</code> , <code>upr</code> ), or an object: <code>htest</code> , <code>TukeyHSD</code> , <code>glht</code> ( <b>multcomp</b> ), <code>glm</code> (logistic regression only!) or <code>posthocTGH</code> ( <b>userfriendlyscience</b> ).
<code>...</code>	other options for x-axis.
<code>xlog</code>	(logical) if log is TRUE, the x axis is drawn logarithmically. Default is FALSE.

xlim	the x limits (x1, x2) of the plot.
xlab	a title for the plot.
yname	If yname is TRUE, the name of comparison between groups are shown.
las	numeric in 0,1,2,3; the style of axis labels. Default is 0. see also par.
pch	plotting 'character', i.e., symbol to use.
pcol	color code or name of the points.
pcolbg	background (fill) color for the open plot symbols given by 'pch = 21:25'.
pcex	character (or symbol) expansion of points.
conf.level	default and glm object only. the confidence interval. Default is 0.95. see also <a href="#">ORci</a> .
cilty	line types of conficence intervals.
cilwd	line width of conficence intervals.
cicol	color code or name of conficence intervals.
v	the x-value(s) for vertical line.
vlty	line types of vertical line.
vlwd	line width of vertical line.
vcol	color code or name of vertical line.
main	a main title for the plot.

**Note**

CIplot was made based on plot.TukeyHSD.

```
# File src/library/stats/R/TukeyHSD.R
# Part of the R package, https://www.R-project.org
#
# Copyright (C) 2000-2001 Douglas M. Bates
# Copyright (C) 2002-2015 The R Core Team
```

**See Also**

plot, axis, points, par.

**Examples**

```
##### default (matrix or data.frame)
require(graphics)
x <- matrix(c(3, 1, 5,
             4, 2, 6), 2, 3, byrow = TRUE)
colnames(x) <- c("esti", "lwr", "upr")
rownames(x) <- c("A", "B")
CIplot(x, xlab = "difference", v = 2, las = 1)

##### 'htest' objects
require(graphics)
```

```
## t test
set.seed(1234)
a <- rnorm(10, 10, 2); b <- rnorm(10, 8, 2)
x <- t.test(a, b)
CIplot(x)

## binomial test
x <- binom.test(5, 20)
CIplot(x, xlim = c(0, 1))

## Fisher's exact test
x <- matrix(c(10, 7, 8, 9), 2, 2, byrow = TRUE)
res <- fisher.test(x)
CIplot(res, xlog = TRUE)

##### 'TukeyHSD' objects
require(graphics)

## Tukey test
aov1 <- aov(breaks ~ tension + wool, data = warpbreaks)
x <- TukeyHSD(aov1)

oldpar <- par(no.readonly = TRUE)
par(mfrow = c(1, 2))
CIplot(x, las = 1)
par(oldpar)

## example of line type and color
aov1 <- aov(breaks ~ tension, data = warpbreaks)
x <- TukeyHSD(aov1)
CIplot(x, las = 1,
       pcol = 2:4, pcolbg = 2:4, cicol = 2:4,
       vltty = 1, vcol = "gray")

##### 'glht' objects
require(graphics)

## Tukey test
require(multcomp)
aov1 <- aov(breaks ~ tension, data = warpbreaks)
x <- glht(aov1, linfct = mcp(tension = "Tukey"))
CIplot(x, las = 1)

## Dunnett test
x <- glht(aov1, linfct = mcp(tension = "Dunnett"))
CIplot(x, las = 1)

##### 'glm' object: logistic regression only!
## odds ratio
require(graphics)
```

```

require(MASS)
data(birthwt)
x <- glm(low ~ age + lwt + smoke + ptl + ht + ui, data = birthwt,
         family = binomial)
CIplot(x, las = 1)

##### 'posthocTGH' object
## Tukey or Games-Howell method
require(graphics)
if (require(userfriendlyscience)) {
  x <- posthocTGH(warpbreaks$breaks, warpbreaks$tension)
  CIplot(x, las = 1)
}

```

---

ORci

*Calculate odds ratios and their confidence intervals from glm object*


---

### Description

Calculate odds ratios and their confidence intervals from glm object

### Usage

```
ORci(x, conf.level = 0.95)
```

### Arguments

**x** glm object (logistic regression only!).  
**conf.level** the confidence interval. Default is 0.95.

### Value

an object ORci and matrix classes with four columns.

**OR** odds ratio

**lwr** lower confidence interval

**upr** upper confidence interval

**p.value** P value by logistic regression

### Examples

```

require(graphics)
require(MASS)
data(birthwt)
x <- glm(low ~ age + lwt + smoke + ptl + ht + ui, data = birthwt,
         family = binomial)
OR1 <- ORci(x)
CIplot(OR1, las = 1)

```

---

print.ORci	<i>Print Methods for Odds Ratios and their Confidence Intervals of ORci object</i>
------------	--

---

### Description

Print odds ratios and their confidence intervals of ORci object.

### Usage

```
## S3 method for class 'ORci'  
print(x, ...)
```

### Arguments

x	ORci object.see also <a href="#">ORci</a> .
...	other options for print such as digits.

### See Also

glm, [ORci](#).

### Examples

```
require(MASS)  
data(birthwt)  
x <- glm(low ~ age + lwt + smoke + ptl + ht + ui, data = birthwt,  
         family = binomial)  
OR1 <- ORci(x)  
print(OR1, digits = 3)
```

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