

# Package ‘adjROC’

October 12, 2022

**Type** Package

**Title** Computing Sensitivity at a Fix Value of Specificity and Vice  
Versa

**Version** 0.2.0

**Author** E. F. Haghish

**Maintainer** E. F. Haghish <haghish@uio.no>

**Description** For a binary classification the adjusted sensitivity and specificity are measured for a given fixed threshold. If the threshold for either sensitivity or specificity is not given, the crossing point between the sensitivity and specificity curves are returned.

**License** MIT + file LICENSE

**Imports** ROCit, ggplot2

**Encoding** UTF-8

**RoxygenNote** 7.1.2

**URL** <https://github.com/haghish/adjROC>,  
<https://www.sv.uio.no/psi/english/people/aca/haghish/>

**BugReports** <https://github.com/haghish/adjROC/issues>

**NeedsCompilation** no

**Repository** CRAN

**Date/Publication** 2022-03-23 13:30:06 UTC

## R topics documented:

adjroc . . . . .	2
<b>Index</b>	<b>4</b>

---

adjroc

*adjroc*


---

### Description

computes adjusted sensitivity, adjusted specificity, or the crossing point between sensitivity and specificity for different thresholds

### Usage

```
adjroc(
  score,
  class,
  method = "emp",
  sensitivity = NULL,
  specificity = NULL,
  plot = FALSE
)
```

### Arguments

score	A numeric array of diagnostic score i.e. the estimated probability of each diagnosis
class	A numeric array of equal length of "score", including the actual class of the observations
method	Specifies the method for estimating the ROC curve. Three methods are supported, which are "empirical", "binormal", and "nonparametric"
sensitivity	numeric. Specify the threshold of sensitivity
specificity	numeric. Specify the threshold of specificity
plot	logical. if TRUE, the sensitivity and specificity will be plotted

### Value

data.frame including cutoff point, and adjusted sensitivity and specificity based on the specified threshold

### Examples

```
# random classification and probability score
score <- runif(10000, min=0, max=1)
class <- sample(x = c(1,0), 10000, replace=TRUE)

# calculate adjusted sensitivity, when specificity threshold is 0.90:
adjroc(score = score, class = class, specificity = 0.9, plot = TRUE)

# calculate adjusted specificity, when sensitivity threshold equals 0.9
```

```
adjroc(score = score, class = class, sensitivity = 0.9, plot = TRUE)

# calculate the meeting point between sensitivity and specificity
adjroc(score = score, class = class, plot = TRUE)
```

# Index

adjroc, [2](#)