

# Package ‘phiDelta’

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**Type** Package

**Title** Tool for Phi Delta Analysis of Features

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**Author** Nikolas Rothe and Ursula Neumann

**Maintainer** Ursula Neumann <ursula.neumann@uni-marburg.de>

**Description** Analysis of features by phi delta diagrams. In particular, functions for reading data and calculating phi and delta as well as the functionality to plot it. Moreover it is possible to do further analysis on the data by generating rankings. For more information on phi delta diagrams, see also Giuliano Armano (2015) <[doi:10.1016/j.ins.2015.07.028](https://doi.org/10.1016/j.ins.2015.07.028)>.

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<b>borders</b>	<i>borders of the phi delta space</i>
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**Description**

calculates the corners of the phi delta space

**Usage**

```
borders(ratio)
```

**Arguments**

**ratio**      is the ratio of positive and negative of the data. The default is 1

**Value**

a matrix. Each row represents a corner in the following order: top, right, bottom, left

**Author(s)**

rothe

**Examples**

```
borders(1.0)
borders(0.5)
borders(2)
```

---

calculate_delta	<i>calculate delta</i>
-----------------	------------------------

---

**Description**

calculates delta out of specificity and sensitivity depending on the ratio

**Usage**

```
calculate_delta(spec, sens, ratio = 1)
```

**Arguments**

spec	is the specificity, the true negative rate
sens	is the sensitivity, the true positive rate
ratio	is the ratio of positive and negative of the data. The default is 1

**Value**

delta

**Author(s)**

rothe

**Examples**

```
calculate_delta(1,0)  
calculate_delta(0.5,0.3)
```

---

calculate_entropy	<i>calculate entropy</i>
-------------------	--------------------------

---

**Description**

calculates the entropy of a specificity and sensitivity tuple considering the ratio

**Usage**

```
calculate_entropy(spec, sens, ratio = 1)
```

**Arguments**

spec	numeric, is the specificity, the true negative rate
sens	numeric, is the sensitivity, the true positive rate
ratio	numeric, is the ratio of positive and negative of the data

**Value**

entropy of the tuple

**Author(s)**

rothe

**Examples**

```
calculate_entropy(1,0)
calculate_entropy(0.5,0.6,0.7)
```

---

calculate\_phi            *calculate phi*

---

**Description**

calculates phi out of specificity and sensitivity depending on the ratio

**Usage**

```
calculate_phi(spec, sens, ratio = 1)
```

**Arguments**

- |       |   |
|-------|---|
| spec  | is the specificity, the true negative rate                          |
| sens  | is the sensitivity, the true positive rate                          |
| ratio | is the ratio of positive and negative of the data. The default is 1 |

**Value**

phi

**Author(s)**

rothe

**Examples**

```
calculate_phi(1,0)
calculate_phi(0.5,0.3)
```

---

calculate_ratio	<i>calculate ratio</i>
-----------------	------------------------

---

**Description**

calculates the ratio between positive and negative samples

**Usage**

```
calculate_ratio(stats)
```

**Arguments**

stats	c_statistics
-------	--------------

**Value**

ratio

**Author(s)**

rothe

**Examples**

```
x <- c_statistics(climate_data)
ratio <- calculate_ratio(x)
```

---

climate_data	<i>Meteorological data for feature selection analysis</i>
--------------	---

---

**Description**

A dataset with meteorological data from a weather station in Frankfurt (Oder), Germany from february 2016

**Usage**

```
climate_data
```

## Format

a data frame with 29 entries and following 7 variables

`RainBool` classification variable: if it has not rained: 0, if it has rained: 1

`date` index variable from 1 to 29

`Tmin` temperature minimum of the day

`Tmax` temperature maximum of the day

`SunAvg` sunshine duration of the day

`RelHumAvg` average relative humidity of the day

`WindForceAvg` average wind force of the day

## References

modified data from <http://wetterstationen.meteomedia.de/>

`crossings`

*Diagram crossings*

## Description

adds crossings to the plot depending on the ratio

## Usage

```
crossings(ratio, col = "darkblue", ...)
```

## Arguments

- `ratio` is the ratio of positive and negative of the data
- `col` the color of the lines. Default is darkblue
- `...` further graphical parameters, see [par](#)

## Author(s)

Neumann

## Examples

```
x <- c_statistics(climate_data)
ratio <- calculate_ratio(x)
phiDelta_plot_from_data(x, crossing = FALSE)
crossings(ratio, col = "green")
```

---

**c\_matrices***confusion matrices*

---

**Description**

calculates the confusion matrices from the c\_statistics

**Usage**

```
c_matrices(stats)
```

**Arguments**

stats	c_statistics
-------	--------------

**Value**

a matrix. Each column represents a feature. Each row describes in this order: true negative, FALSE negative, true positive, FALSE negative

**Author(s)**

rothe

**Examples**

```
x <- c_statistics(climate_data)
cmat <- c_matrices(x)
```

---

**c\_statistics***Raw Confusion Statistics*

---

**Description**

reformats the raw file data to c\_statistics data so it can be used for most of the functions in this package. it can be used directly after loading data from a file like .csv

**Usage**

```
c_statistics(file)
```

**Arguments**

file	raw data from a file, for example the output of <a href="#">read.csv</a> . the file must be formated as follows: The first column contains the output of the classifier. It should only be 1 or 0. The other columns represent the features. The names of the columns 2.. are considered as the names of the features
------	---

**Value**

dataframe, first column are the labels, 0 is a negative sample, 1 a positive the other columns contain the

**Author(s)**

rothe

**Examples**

```
data("climate_data")
x <- c_statistics(climate_data)
```

<code>dist_to_middle</code>	<i>distance to the middle of the space</i>
-----------------------------	--

**Description**

calculates the euclidic distance of a phi delta tuple to the middle of the phi delta space. This could be used for a rating of the features

**Usage**

```
dist_to_middle(phi, delta, ratio)
```

**Arguments**

<code>phi</code>	numeric value or vector of phi
<code>delta</code>	numeric value or vector of delta
<code>ratio</code>	is the ratio of positive and negative of the data. The default is 1

**Value**

the euclidic distance of the tuple to the middle

**Author(s)**

rothe

**Examples**

```
dist_to_middle(1,0,1)
dist_to_middle(0.5,0.3,1)
```

---

<code>dist_to_top</code>	<i>distance to top or bottom</i>
--------------------------	----------------------------------

---

**Description**

calculates the distance of the tuple to the closer corner of top and bottom of the phi delta space with ratio 1. This can be used for a ranking of the features

**Usage**

```
dist_to_top(phi, delta)
```

**Arguments**

<code>phi</code>	numeric value or vector of phi
<code>delta</code>	numeric value or vector of delta

**Value**

distance to the top or the bottom corner

**Author(s)**

rothe

**Examples**

```
dist_to_top(1,0)  
dist_to_top(0.5,0.3)
```

---

<code>iso_accuracy</code>	<i>isometric accuracy lines</i>
---------------------------	---------------------------------

---

**Description**

adds isometric lines for the accuracy to the plot depending on the ratio

**Usage**

```
iso_accuracy(ratio = 1, granularity = 0.25, lty = "longdash",  
            col = "blue", ...)
```

**Arguments**

<code>ratio</code>	numeric value for the ratio of positive and negative of the data
<code>granularity</code>	numeric value between 0 and 1 for the granularity of the lines. It is a value for the distance between 2 lines
<code>lty</code>	the type of line, see <a href="#">par</a>
<code>col</code>	the color of the lines
<code>...</code>	further graphical parameters, see <a href="#">par</a>

**Author(s)**

rothe

**Examples**

```
x <- c_statistics(climate_data)
ratio <- calculate_ratio(x)
phiDelta_plot_from_data(x)
iso_accuracy(ratio, col = "green")
```

<code>iso_entropy_curve</code>	<i>isometric entropy</i>
--------------------------------	--------------------------

**Description**

draws isometric curves for the entropy by calculating the entropy for all points in a grid and connecting those within a epsilon environment of the value

**Usage**

```
iso_entropy_curve(x, ratio = 1, eps = 0.001, grid_granularity = 0.01)
```

**Arguments**

<code>x</code>	numeric, is the offset for the points
<code>ratio</code>	numeric, is the ratio
<code>eps</code>	numeric, the epsilon for entropies to be selected
<code>grid_granularity</code>	numeric between 0 and 1, defines the granularity of the grid

**Author(s)**

Neumann

---

iso\_negative\_predictive\_value  
*isometric negative predictive value lines*

---

**Description**

adds isometric lines for the negative predictive value to the plot depending on the ratio

**Usage**

```
iso_negative_predictive_value(ratio = 1, granularity = 0.25,
                             lty = "longdash", col = "blue", ...)
```

**Arguments**

ratio	numeric value for the ratio of positive and negative of the data
granularity	numeric value between 0 and 1 for the granularity of the lines. It is a value for the distance between 2 lines
lty	the type of line, see <a href="#">par</a>
col	the color of the lines
...	further graphical parameters, see <a href="#">par</a>

**Author(s)**

rothe

**Examples**

```
x <- c_statistics(climate_data)
ratio <- calculate_ratio(x)
phiDelta_plot_from_data(x)
iso_negative_predictive_value(ratio, col = "green")
```

---

iso\_precision      *isometric precision lines*

---

**Description**

adds isometric lines for the precision to the plot depending on the ratio

**Usage**

```
iso_precision(ratio = 1, granularity = 0.25, lty = "longdash",
              col = "blue", ...)
```

**Arguments**

<code>ratio</code>	numeric value for the ratio of positive and negative of the data
<code>granularity</code>	numeric value between 0 and 1 for the granularity of the lines. It is a value for the distance between 2 lines
<code>lty</code>	the type of line, see <a href="#">par</a>
<code>col</code>	the color of the lines
<code>...</code>	further graphical parameters, see <a href="#">par</a>

**Author(s)**

rothe

**Examples**

```
x <- c_statistics(climate_data)
ratio <- calculate_ratio(x)
phiDelta_plot_from_data(x)
iso_precision(ratio, col = "green")
```

*iso\_sensitivity*      *isometric sensitivity lines*

**Description**

adds isometric lines for the sensitivity to the plot depending on the ratio

**Usage**

```
iso_sensitivity(ratio = 1, granularity = 0.25, col = "blue",
                lty = "longdash", ...)
```

**Arguments**

<code>ratio</code>	numeric value for the ratio of positive and negative of the data
<code>granularity</code>	numeric value between 0 and 1 for the granularity of the lines. It is a value for the distance between 2 lines
<code>col</code>	the color of the lines
<code>lty</code>	the type of line, see <a href="#">par</a>
<code>...</code>	further graphical parameters, see <a href="#">par</a>

**Author(s)**

Neumann

## Examples

```
x <- c_statistics(climate_data)
ratio <- calculate_ratio(x)
phiDelta_plot_from_data(x)
iso_specificity(ratio, col = "green")
```

---

iso\_specificity      *isometric specificity lines*

---

## Description

adds isometric lines for the specificity to the plot depending on the ratio

## Usage

```
iso_specificity(ratio = 1, granularity = 0.25, col = "blue",
lty = "longdash", ...)
```

## Arguments

ratio	numeric value for the ratio of positive and negative of the data
granularity	numeric value between 0 and 1 for the granularity of the lines. It is a value for the distance between 2 lines
col	the color of the lines
lty	the type of line, see <a href="#">par</a>
...	further graphical parameters, see <a href="#">par</a>

## Author(s)

rothe

## Examples

```
x <- c_statistics(climate_data)
ratio <- calculate_ratio(x)
phiDelta_plot_from_data(x)
iso_specificity(ratio, col = "green")
```

<code>n_matrices</code>	<i>normalized confusion matrices</i>
-------------------------	--------------------------------------

### Description

normalizes the confusion matrices

### Usage

```
n_matrices(c_matrices)
```

### Arguments

<code>c_matrices</code>	confusion matrices
-------------------------	--------------------

### Value

a matrix. Each column represents a feature. Each row describes in this order: true negative rate, FALSE negative rate, true positive rate, FALSE negative rate

### Author(s)

rothe

### Examples

```
x <- c_statistics(climate_data)
cmat <- c_matrices(x)
nmat <- n_matrices(cmat)
```

<code>phiDelta.convert</code>	<i>Conversion of specificity and sensitivity to phi and delta</i>
-------------------------------	---

### Description

converts specificity and sensitivity to phi and delta depending on the ratio

### Usage

```
phiDelta.convert(spec, sens, ratio = 1)
```

### Arguments

<code>spec</code>	is the specificity, the true negative rate
<code>sens</code>	is the sensitivity, the true positive rate
<code>ratio</code>	is the ratio of positive and negative of the data. The default is 1

**Value**

List with phi and delta vectors

**Author(s)**

neumann

**Examples**

```
phiDelta.convert(1,0)
phiDelta.convert(0.5,0.3, ratio = 0.8)
```

---

phiDelta.plot

*Plot of phi delta diagram*

---

**Description**

Plots delta against phi within the phi delta diagram shape

**Usage**

```
phiDelta.plot(phi, delta, ratio = 1, names = NULL, border = "red",
              filling = "grey", crossing = TRUE, iso_specificity = FALSE,
              iso_sensitivity = FALSE, iso_neg_predictive_value = FALSE,
              iso_precision = FALSE, iso_accuracy = FALSE, highlighted = NULL)
```

**Arguments**

phi	numeric value or vector of phi
delta	numeric value or vector of delta
ratio	numeric, is the ratio of positive and negative of the data
names	string with feature names
border	the color of the border of the shape. NA for no border
filling	the color to fill the shape with
crossing	logical, if the crossing should be drawn
iso_specificity	logical, if isometric lines of the specificity should be drawn
iso_sensitivity	logical, if isometric lines of the sensitivity should be drawn
iso_neg_predictive_value	logical, if isometric lines of the negative predictive value should be drawn
iso_precision	logical, if isometric lines of the precision should be drawn
iso_accuracy	logical, if isometric lines of the accuracy should be drawn
highlighted	numeric vector, indices of the points to highlight highlighted points will be orange

**Author(s)**

rothe

**Examples**

```
x <- climate_data
phiDelta <- phiDelta.stats(x[,-1],x[,1])
phiDelta.plot(phiDelta$phi, phiDelta$delta)
phiDelta.plot(phiDelta$phi, phiDelta$delta,
ratio = phiDelta$ratio,
border = "green",
iso_neg_predictive_value = TRUE,
crossing = FALSE)
```

**phiDelta.stats**

*Phi delta statistics from dataframe*

**Description**

calculates phi, delta and the ratio directly from the dataframe with provided information and generates a list with the names of the features, their phi and delta value and the ratio

**Usage**

```
phiDelta.stats(data, labels, ratio_corrected = TRUE)
```

**Arguments**

<code>data</code>	dataframe without labels
<code>labels</code>	vector of labels
<code>ratio_corrected</code>	logical, if true phi and delta will be calculated in respect to the ratio of positive and negative samples

**Value**

dataframe, first column are the names of the features second column the phi values third column the delta values

**Author(s)**

rothe

**Examples**

```
x <- climate_data
phiDelta <- phiDelta.stats(x[,-1],x[,1], ratio_corrected = FALSE)
with_ratio <- phiDelta.stats(x[,-1],x[,1])
```

`phiDelta_from_data`      *phi delta matrix*

### Description

calculates phi and delta directly from the stats and generates a matrix with the names of the features, their phi and their delta value

### Usage

```
phiDelta_from_data(stats, ratio_corrected = TRUE)
```

### Arguments

stats	c_statistics
ratio_corrected	logical, if true phi and delta will be calculated in respect to the ratio of positive and negative samples

### Value

dataframe, first column are the names of the features second column the phi values third column the delta values

### Author(s)

rothe

### Examples

```
x <- c_statistics(climate_data)
phiDelta <- phiDelta_from_data(x, ratio_corrected = FALSE)
with_ratio <- phiDelta_from_data(x)
```

`phiDelta_plot_from_data`      *phi delta plot of raw statistic data*

### Description

this will create a basic plot directly out of the statistic data (c\_statistics)

### Usage

```
phiDelta_plot_from_data(stats, names = NULL, ratio_corrected = TRUE, ...)
```

**Arguments**

<code>stats</code>	matrix of the statistic data of the features and the classifier
<code>names</code>	vector with feature names
<code>ratio_corrected</code>	logical, if true the plot will consider the ratio of the positive and negative data samples
<code>...</code>	further parameters for the diagram see <a href="#">phiDelta.plot</a>

**Author(s)**

rothe

**Examples**

```
x <- c_statistics(climate_data)
phiDelta_plot_from_data(x)
phiDelta_plot_from_data(x, ratio_corrected = FALSE, iso_spec = TRUE, iso_sens = TRUE)
```

<code>rank_stats</code>	<i>ranking of the features</i>
-------------------------	--------------------------------

**Description**

this function puts together a number of rankings of the features

**Usage**

```
rank_stats(stats, ratio_corrected = FALSE, delta_dist = 1)
```

**Arguments**

<code>stats</code>	c_statistics, the data input
<code>ratio_corrected</code>	logical, true if ratio should be considered
<code>delta_dist,</code>	numeric, the delta value of the anchor for the geometrical ranking see <a href="#">symmetric_distance</a>

**Author(s)**

rothe

---

symmetric\_distance      *X symmetric distance of a point*

---

### Description

calculates the Distance from the positive anchor and the negative anchor to the point and returns the smaller one. That means, if y is positive the distance to the positive anchor will be return, if it is negative, the negative anchor distance will be calculated

### Usage

```
symmetric_distance(x, y, anchor)
```

### Arguments

x, y	numerical, in this case phi and delta but in general the input coordinates
anchor	vector (x,y) the anchor for the calculation of the distance

### Value

the smaller distance of (x,y) to either the positive or negative anchor

### Examples

```
symmetric_distance(0.5,0.5,c(0,0))
```

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