

Package ‘MetricsWeighted’

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Type Package

Title Weighted Metrics, Scoring Functions and Performance Measures for Machine Learning

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Description Provides weighted versions of several metrics, scoring functions and performance measures used in machine learning, including average unit deviances of the Bernoulli, Tweedie, Poisson, and Gamma distributions, see Jorgensen B. (1997, ISBN: 978-0412997112). The package also contains a weighted version of generalized R-squared, see e.g. Cohen, J. et al. (2002, ISBN: 978-0805822236). Furthermore, 'dplyr' chains are supported.

License GPL (>= 2)

Depends R (>= 3.1.0)

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URL <https://github.com/mayer79/MetricsWeighted>

BugReports <https://github.com/mayer79/MetricsWeighted/issues>

Imports stats, graphics

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accuracy	<i>Accuracy</i>
----------	-----------------

Description

Calculates weighted accuracy, i.e. the weighted proportion of elements in predicted that are equal to those in actual. The higher, the better.

Usage

```
accuracy(actual, predicted, w = NULL, ...)
```

Arguments

actual	Observed values.
predicted	Predicted values.
w	Optional case weights.
...	Further arguments passed to <code>weighted_mean</code> .

Value

A numeric vector of length one.

See Also

[classification_error](#).

Examples

```
accuracy(c(0, 0, 1, 1), c(0, 0, 1, 1))
accuracy(c(1, 0, 0, 1), c(0, 0, 1, 1), w = 1:4)
```

AUC

Area under the ROC

Description

Function modified from `glmnet` package (modified to ensure deterministic results). Calculates weighted AUC, i.e. the area under the receiver operating curve. The larger, the better.

Usage

```
AUC(actual, predicted, w = NULL, ...)
```

Arguments

actual	Observed values (0 or 1).
predicted	Predicted values of any value (not necessarily between 0 and 1).
w	Optional case weights.
...	Further arguments passed by other methods.

Details

The unweighted version can be different from the weighted one with unit weights due to ties in predicted.

Value

A numeric vector of length one.

See Also

[gini_coefficient](#).

Examples

```
AUC(c(0, 0, 1, 1), c(0.1, 0.1, 0.9, 0.8))
AUC(c(1, 0, 0, 1), c(0.1, 0.1, 0.9, 0.8))
# different from last due to ties 'in predicted':
AUC(c(1, 0, 0, 1), c(0.1, 0.1, 0.9, 0.8), w = rep(1, 4))
```

classification_error *Classification Error*

Description

Calculates weighted classification error, i.e. the weighted proportion of elements in predicted that are unequal to those in observed. Equals 1 - accuracy, thus lower values are better.

Usage

```
classification_error(actual, predicted, w = NULL, ...)
```

Arguments

actual	Observed values.
predicted	Predicted values.
w	Optional case weights.
...	Further arguments passed to accuracy.

Value

A numeric vector of length one.

See Also

[accuracy](#).

Examples

```
classification_error(c(1, 0, 0, 1), c(0, 0, 1, 1))
classification_error(c(1, 0, 0, 1), c(0, 0, 1, 1), w = 1:4)
```

deviance_bernoulli *Bernoulli Deviance*

Description

Calculates weighted average of unit Bernoulli deviance. Defined as twice logLoss. The smaller the deviance, the better.

Usage

```
deviance_bernoulli(actual, predicted, w = NULL, ...)
```

Arguments

actual	Observed values (0 or 1).
predicted	Predicted values strictly between 0 and 1.
w	Optional case weights.
...	Further arguments passed to logLoss.

Value

A numeric vector of length one.

See Also

[logLoss](#).

Examples

```
deviance_bernoulli(c(0, 0, 1, 1), c(0.1, 0.1, 0.9, 0.8))  
deviance_bernoulli(c(0, 0, 1, 1), c(0.1, 0.1, 0.9, 0.8), w = 1:4)
```

deviance_gamma *Gamma Deviance*

Description

Weighted average of (unscaled) unit Gamma deviance, see e.g. the reference below. Special case of Tweedie deviance with Tweedie parameter 2. The smaller the deviance, the better.

Usage

```
deviance_gamma(actual, predicted, w = NULL, ...)
```

Arguments

actual	Strictly positive observed values.
predicted	Strictly positive predicted values.
w	Optional case weights.
...	Further arguments passed to <code>weighted_mean</code> .

Value

A numeric vector of length one.

References

Jorgensen, B. (1997). *The Theory of Dispersion Models*. Chapman & Hall/CRC. ISBN 978-0412997112.

See Also

[deviance_tweedie](#).

Examples

```
deviance_gamma(1:10, c(1:9, 12))
deviance_gamma(1:10, c(1:9, 12), w = 1:10)
```

deviance_normal	<i>Normal Deviance</i>
-----------------	------------------------

Description

Weighted average of (unscaled) unit normal deviance. This equals the weighted mean-squared error, see e.g. the reference below. The smaller the deviance, the better.

Usage

```
deviance_normal(actual, predicted, w = NULL, ...)
```

Arguments

actual	Observed values.
predicted	Predicted values.
w	Optional case weights.
...	Further arguments passed to <code>mse</code> .

Value

A numeric vector of length one.

References

Jorgensen, B. (1997). The Theory of Dispersion Models. Chapman & Hall/CRC. ISBN 978-0412997112.

See Also

[deviance_tweedie](#), [mse](#).

Examples

```
deviance_normal(1:10, c(1:9, 12))
deviance_normal(1:10, c(1:9, 12), w = 1:10)
```

deviance_poisson	<i>Poisson Deviance</i>
------------------	-------------------------

Description

Weighted average of unit Poisson deviance, see reference below. Special case of Tweedie deviance with Tweedie parameter 1.

Usage

```
deviance_poisson(actual, predicted, w = NULL, ...)
```

Arguments

actual	Observed non-negative values.
predicted	Strictly positive predicted values.
w	Optional case weights.
...	Further arguments passed to <code>weighted_mean</code> .

Value

A numeric vector of length one.

References

Jorgensen, B. (1997). The Theory of Dispersion Models. Chapman & Hall/CRC. ISBN 978-0412997112.

See Also

[deviance_tweedie](#).

Examples

```
deviance_poisson(0:2, c(0.1, 1, 3))
deviance_poisson(0:2, c(0.1, 1, 3), w = c(1, 1, 1))
deviance_tweedie(0:2, c(0.1, 1, 3), tweedie_p = 1)
deviance_tweedie(0:2, c(0.1, 1, 3), tweedie_p = 1.01)
deviance_poisson(0:2, c(0.1, 1, 3), w = 1:3)
```

deviance_tweedie	<i>Tweedie Deviance</i>
------------------	-------------------------

Description

Weighted average of (unscaled) unit Tweedie deviance with parameter p . This includes the normal deviance ($p = 0$), the Poisson deviance ($p = 1$), as well as the Gamma deviance ($p = 2$), see reference below and https://en.wikipedia.org/wiki/Tweedie_distribution for the specific deviance formula. For $0 < p < 1$, the distribution is not defined. The smaller the deviance, the better.

Usage

```
deviance_tweedie(actual, predicted, w = NULL, tweedie_p = 0, ...)
```

Arguments

actual	Observed values.
predicted	Predicted values.
w	Optional case weights.
tweedie_p	Tweedie power.
...	Further arguments passed to <code>weighted_mean</code> .

Value

A numeric vector of length one.

References

Jorgensen, B. (1997). The Theory of Dispersion Models. Chapman & Hall/CRC. ISBN 978-0412997112.

See Also

[deviance_normal](#), [deviance_poisson](#), [deviance_gamma](#).

Examples

```
deviance_tweedie(1:10, c(1:9, 12), tweedie_p = 0)
deviance_tweedie(1:10, c(1:9, 12), tweedie_p = 1)
deviance_tweedie(1:10, c(1:9, 12), tweedie_p = 2)
deviance_tweedie(1:10, c(1:9, 12), tweedie_p = 1.5)
deviance_tweedie(1:10, c(1:9, 12), tweedie_p = 1.5, w = 1:10)
```

elementary_score *Elementary Scoring Function for Expectiles and Quantiles*

Description

Weighted average of the elementary scoring function for expectiles resp. quantiles at level α with parameter θ , see reference below. Every choice of θ gives a scoring function consistent for the expectile resp. quantile at level α . Note that the expectile at level $\alpha = 0.5$ is the expectation (mean). The smaller the score, the better.

Usage

```
elementary_score_expectile(  
  actual,  
  predicted,  
  w = NULL,  
  alpha = 0.5,  
  theta = 0,  
  ...  
)
```

```
elementary_score_quantile(  
  actual,  
  predicted,  
  w = NULL,  
  alpha = 0.5,  
  theta = 0,  
  ...  
)
```

Arguments

actual	Observed values.
predicted	Predicted values.
w	Optional case weights.
alpha	Optional level of expectile resp. quantile.
theta	Optional parameter.
...	Further arguments passed to <code>weighted_mean</code> .

Value

A numeric vector of length one.

References

Ehm, W., Gneiting, T., Jordan, A. and Krüger, F. (2016), Of quantiles and expectiles: consistent scoring functions, Choquet representations and forecast rankings. *J. R. Stat. Soc. B*, 78: 505-562, <doi.org/10.1111/rssb.12154>.

See Also

[murphy_diagram](#).

Examples

```
elementary_score_expectile(1:10, c(1:9, 12), alpha = 0.5, theta = 11)
elementary_score_quantile(1:10, c(1:9, 12), alpha = 0.5, theta = 11)
```

f1_score

F1 Score

Description

Calculates weighted F1 score or F measure defined as the harmonic mean of precision and recall, see https://en.wikipedia.org/wiki/Precision_and_recall for some background. The higher, the better.

Usage

```
f1_score(actual, predicted, w = NULL, ...)
```

Arguments

actual	Observed values (0 or 1).
predicted	Predicted values (0 or 1).
w	Optional case weights.
...	Further arguments passed to precision and recall.

Value

A numeric vector of length one.

See Also

[precision](#), [recall](#).

Examples

```
f1_score(c(0, 0, 1, 1), c(0, 0, 1, 1))
f1_score(c(1, 0, 0, 1), c(0, 0, 1, 1), w = 1:4)
```

gini_coefficient	<i>Gini Coefficient</i>
------------------	-------------------------

Description

Calculates weighted Gini coefficient, obtained as $2 * \text{AUC} - 1$. Up to ties in predicted equivalent to Somer's D. The larger the Gini coefficient, the better.

Usage

```
gini_coefficient(actual, predicted, w = NULL, ...)
```

Arguments

actual	Observed values (0 or 1).
predicted	Predicted values of any value (not necessarily between 0 and 1).
w	Optional case weights.
...	Further arguments passed to AUC.

Value

A numeric vector of length one.

See Also

[AUC](#).

Examples

```
gini_coefficient(c(0, 0, 1, 1), 2 * c(0.1, 0.1, 0.9, 0.8))
gini_coefficient(c(0, 0, 1, 1), c(0.1, 0.6, 0.9, 0.5))
gini_coefficient(c(0, 0, 1, 1), c(0.1, 0.6, 0.9, 0.5), w = 1:4)
```

logLoss	<i>Log Loss/Binary Cross Entropy</i>
---------	--------------------------------------

Description

Calculates weighted logloss resp. cross entropy. Equals half of the unit Bernoulli deviance. The smaller, the better.

Usage

```
logLoss(actual, predicted, w = NULL, ...)
```

Arguments

actual	Observed values (0 or 1).
predicted	Predicted values strictly larger than 0 and smaller than 1.
w	Optional case weights.
...	Further arguments passed to <code>weighted_mean</code> .

Value

A numeric vector of length one.

See Also

[deviance_bernoulli](#).

Examples

```
logLoss(c(0, 0, 1, 1), c(0.1, 0.1, 0.9, 0.8))
logLoss(c(0, 0, 1, 1), c(0.1, 0.1, 0.9, 0.8), w = 1:4)
```

mae

Mean Absolute Error

Description

Calculates weighted mean absolute error of predicted values. The smaller the value, the better.

Usage

```
mae(actual, predicted, w = NULL, ...)
```

Arguments

actual	Observed values.
predicted	Predicted values.
w	Optional case weights.
...	Further arguments passed to <code>weighted_mean</code> .

Value

A numeric vector of length one.

Examples

```
mae(1:10, c(1:9, 12))
mae(1:10, c(1:9, 12), w = 1:10)
```

mape	<i>Mean Absolute Percentage Error</i>
------	---------------------------------------

Description

Calculates weighted mean absolute percentage error of predicted values. The smaller, the better.

Usage

```
mape(actual, predicted, w = NULL, ...)
```

Arguments

actual	Strictly positive observed values.
predicted	Predicted values.
w	Optional case weights.
...	Further arguments passed to <code>weighted_mean</code> .

Value

A numeric vector of length one.

Examples

```
mape(1:10, c(1:9, 12))  
mape(1:10, c(1:9, 12), w = 1:10)
```

medae	<i>Median Absolute Error</i>
-------	------------------------------

Description

Calculates weighted median absolute error of predicted values. The smaller the value, the better.

Usage

```
medae(actual, predicted, w = NULL, ...)
```

Arguments

actual	Observed values.
predicted	Predicted values.
w	Optional case weights.
...	Further arguments passed to <code>weighted_mean</code> .

Value

A numeric vector of length one.

Examples

```
medae(1:10, c(2:10, 100))
medae(1:10, c(2:10, 100), w = 1:10)
```

mse	<i>Mean-Squared Error</i>
-----	---------------------------

Description

Calculates weighted mean-squared error of prediction. Equals mean unit normal deviance. The smaller, the better.

Usage

```
mse(actual, predicted, w = NULL, ...)
```

Arguments

actual	Observed values.
predicted	Predicted values.
w	Optional case weights.
...	Further arguments passed to <code>weighted_mean</code> .

Value

A numeric vector of length one.

See Also

[rmse](#), [deviance_normal](#).

Examples

```
mse(1:10, c(1:9, 12))
mse(1:10, c(1:9, 12), w = 1:10)
```

multi_metric	<i>Multiple Metrics</i>
--------------	-------------------------

Description

Provides a way to create a list of metrics/scoring functions/performance measures from a parametrized function like the Tweedie deviance or the elementary scoring functions for expectiles.

Usage

```
multi_metric(fun, ...)
```

Arguments

fun	A metric/scoring function/performance measure with additional parameter to be varied.
...	Further arguments passed to fun, including one varying parameter (specified by a vector).

Value

A named list of functions.

See Also

[performance](#).

Examples

```
data <- data.frame(act = 1:10, pred = c(1:9, 12))
multi <- multi_metric(fun = deviance_tweedie,
  tweedie_p = c(0, seq(1, 3, by = 0.1)))
performance(data, actual = "act", predicted = "pred", metrics = multi)
multi <- multi_metric(fun = r_squared, deviance_function = deviance_tweedie,
  tweedie_p = c(0, seq(1, 3, by = 0.1)))
performance(data, actual = "act", predicted = "pred", metrics = multi)
multi <- multi_metric(fun = elementary_score_expectile,
  theta = 1:11, alpha = 0.1)
performance(data, actual = "act", predicted = "pred",
  metrics = multi, key = "theta")
multi <- multi_metric(fun = elementary_score_expectile,
  theta = 1:11, alpha = 0.5)
performance(data, actual = "act", predicted = "pred",
  metrics = multi, key = "theta")
```

murphy_diagram	<i>Murphy diagram</i>
----------------	-----------------------

Description

Murphy diagram of the elementary scoring function for expectiles resp. quantiles at level α for different values of θ . Can be used to study and compare performance of one or multiple models. If the plot needs to be customized, set `plot = FALSE` to get the resulting data instead of the plot.

Usage

```
murphy_diagram(
  actual,
  predicted,
  w = NULL,
  alpha = 0.5,
  theta = seq(-2, 2, length.out = 100),
  functional = c("expectile", "quantile"),
  plot = TRUE,
  ...
)
```

Arguments

<code>actual</code>	Observed values.
<code>predicted</code>	Predicted values.
<code>w</code>	Optional case weights.
<code>alpha</code>	Level of expectile resp. quantile. The default $\alpha = 0.5$ corresponds to the expectation resp. median.
<code>theta</code>	Vector of evaluation points.
<code>functional</code>	Either "expectile" or "quantile".
<code>plot</code>	Should plot (TRUE) be returned or the data to be plotted?
<code>...</code>	Further arguments passed to <code>plot</code> .

Value

A named list of functions.

References

Ehm, W., Gneiting, T., Jordan, A. and Krüger, F. (2016), Of quantiles and expectiles: consistent scoring functions, Choquet representations and forecast rankings. *J. R. Stat. Soc. B*, 78: 505-562, <doi.org/10.1111/rssb.12154>.

See Also

[elementary_score](#).

Examples

```
y <- 1:10
predicted <- 1.1 * y
murphy_diagram(y, predicted, theta = seq(0.9, 1.2, by = 0.01))
two_models <- cbind(m1 = predicted, m2 = 1.2 * y)
murphy_diagram(y, two_models, theta = seq(0.9, 1.3, by = 0.01))
```

performance

Performance

Description

Applies one or more metrics to a `data.frame` containing columns with actual and predicted values as well as an optional column with case weights. The results are returned as a `data.frame` and can be used in a `dplyr` chain.

Usage

```
performance(
  data,
  actual,
  predicted,
  w = NULL,
  metrics = rmse,
  key = "metric",
  value = "value",
  ...
)
```

Arguments

<code>data</code>	A <code>data.frame</code> containing actual, predicted and possibly <code>w</code> .
<code>actual</code>	The column name in <code>data</code> referring to actual values.
<code>predicted</code>	The column name in <code>data</code> referring to predicted values.
<code>w</code>	The optional column name in <code>data</code> referring to case weights.
<code>metrics</code>	Either a function or a named list of functions. Each function represents a metric and has four arguments: <code>observed</code> , <code>predicted</code> , <code>case weights</code> and <code>...</code> . If not a named list but a single function, the name of the function is guessed by <code>deparse(substitute(...))</code> , which would not provide the actual name of the function if called within <code>lapply</code> etc. In such cases, you can pass a named list with one element, e.g. <code>list(rmse = rmse)</code> .

key	Name of the resulting column containing the name of the metric. Defaults to "metric".
value	Name of the resulting column with the value of the metric. Defaults to "value".
...	Further arguments passed to the metric functions, e.g. if the metric is "r_squared", you could pass the relevant deviance function as additional argument (see examples).

Value

Data frame with one row per metric and two columns: key and value.

Examples

```

ir <- iris
fit_num <- lm(Sepal.Length ~ ., data = ir)
ir$fitted <- fit_num$fitted
performance(ir, "Sepal.Length", "fitted")
performance(ir, "Sepal.Length", "fitted", metrics = r_squared)
performance(ir, "Sepal.Length", "fitted",
             metrics = c(`R-squared` = r_squared, rmse = rmse))
performance(ir, "Sepal.Length", "fitted", metrics = r_squared,
             deviance_function = deviance_gamma)
performance(ir, "Sepal.Length", "fitted", metrics = r_squared,
             deviance_function = deviance_tweedie, tweedie_p = 2)

## Not run:
library(dplyr)

iris %>%
  mutate(pred = predict(fit_num, data = .)) %>%
  performance("Sepal.Length", "pred")

# Same
iris %>%
  mutate(pred = predict(fit_num, data = .)) %>%
  performance("Sepal.Length", "pred", metrics = rmse)

# Grouped by Species
iris %>%
  mutate(pred = predict(fit_num, data = .)) %>%
  group_by(Species) %>%
  do(performance(., "Sepal.Length", "pred"))

# Multiple measures
iris %>%
  mutate(pred = predict(fit_num, data = .)) %>%
  performance("Sepal.Length", "pred",
             metrics = list(rmse = rmse, mae = mae, `R-squared` = r_squared))

# Grouped by Species
iris %>%
  mutate(pred = predict(fit_num, data = .)) %>%
  group_by(Species) %>%

```

```
do(performance(. , "Sepal.Length", "pred",
              metrics = list(rmse = rmse, mae = mae,
                            `R-squared` = r_squared)))

## End(Not run)
```

precision	<i>Precision</i>
-----------	------------------

Description

Calculates weighted precision, see https://en.wikipedia.org/wiki/Precision_and_recall for the (unweighted) version. The higher, the better.

Usage

```
precision(actual, predicted, w = NULL, ...)
```

Arguments

actual	Observed values (0 or 1).
predicted	Predicted values (0 or 1).
w	Optional case weights.
...	Further arguments passed to <code>weighted_mean</code> .

Value

A numeric vector of length one.

See Also

[recall](#), [f1_score](#).

Examples

```
precision(c(0, 0, 1, 1), c(0, 0, 1, 1))
precision(c(1, 0, 0, 1), c(0, 0, 1, 1), w = 1:4)
```

prop_within	<i>Proportion Within</i>
-------------	--------------------------

Description

Calculates weighted proportion of predictions that are within a given tolerance around the actual values. The larger the value, the better.

Usage

```
prop_within(actual, predicted, w = NULL, tol = 1, ...)
```

Arguments

actual	Observed values.
predicted	Predicted values.
w	Optional case weights.
tol	Predictions in the closed interval from actual - tol to actual + tol count as within.
...	Further arguments passed to <code>weighted_mean</code> .

Value

A numeric vector of length one.

Examples

```
prop_within(1:10, c(1:9, 12))
prop_within(1:10, c(1:9, 12), w = 1:10)
data <- data.frame(act = 1:10, pred = c(1:9, 12), w = 1:10)
multi <- multi_metric(fun = prop_within, tol = 0:3)
performance(data, actual = "act", predicted = "pred", w = "w",
  metrics = multi, key = "Proportion within")
```

recall	<i>Recall</i>
--------	---------------

Description

Calculates weighted recall, see https://en.wikipedia.org/wiki/Precision_and_recall for the (unweighted) definition. The higher, the better.

Usage

```
recall(actual, predicted, w = NULL, ...)
```

Arguments

actual	Observed values (0 or 1).
predicted	Predicted values (0 or 1).
w	Optional case weights.
...	Further arguments passed to <code>weighted_mean</code> .

Value

A numeric vector of length one.

See Also

[precision](#), [f1_score](#).

Examples

```
recall(c(0, 0, 1, 1), c(0, 0, 1, 1))
recall(c(1, 0, 0, 1), c(0, 0, 1, 1), w = 1:4)
```

 rmse

Root-Mean-Squared Error

Description

Weighted root-mean-squared error of predicted values. Equals the square root of mean-squared error. Smaller values are better.

Usage

```
rmse(actual, predicted, w = NULL, ...)
```

Arguments

actual	Observed values.
predicted	Predicted values.
w	Optional case weights.
...	Further arguments passed to <code>mse</code> .

Value

A numeric vector of length one.

See Also

[mse](#).

Examples

```
rmse(1:10, c(1:9, 12))
rmse(1:10, c(1:9, 12), w = 1:10)
```

r_squared	<i>Generalized R-Squared</i>
-----------	------------------------------

Description

Returns (weighted) proportion of deviance explained, see reference below. For the mean-squared error as deviance, this equals the usual (weighted) R-squared. The higher, the better.

Usage

```
r_squared(
  actual,
  predicted,
  w = NULL,
  deviance_function = mse,
  reference_mean = NULL,
  ...
)
```

Arguments

actual	Observed values.
predicted	Predicted values.
w	Optional case weights.
deviance_function	A positive (deviance) function taking four arguments: "actual", "predicted", "w" and "...".
reference_mean	An optional reference mean used to derive the null deviance. Recommended in out-of-sample applications.
...	Further arguments passed to weighted_mean and deviance_function.

Details

The deviance gain is calculated regarding the null model derived from the actual values. While fine for in-sample considerations, this is only an approximation for out-of-sample considerations. There, it is recommended to calculate null deviance regarding the in-sample (weighted) mean. This value can be passed by the argument reference_mean.

Value

A numeric vector of length one.

References

Cohen, Jacob. et al. (2002). Applied Multiple Regression/Correlation Analysis for the Behavioral Sciences (3rd ed.). Routledge. ISBN 978-0805822236.

See Also

[deviance_normal](#), [mse](#).

Examples

```
r_squared(1:10, c(1, 1:9))
r_squared(1:10, c(1, 1:9), w = 1:10)
r_squared(0:2, c(0.1, 1, 2), deviance_function = deviance_poisson)
r_squared(0:2, c(0.1, 1, 2), w = rep(1, 3),
          deviance_function = deviance_poisson)
r_squared(0:2, c(0.1, 1, 2), w = rep(1, 3),
          deviance_function = deviance_tweedie, tweedie_p = 1)

# respect to 'own' deviance formula
myTweedie <- function(actual, predicted, w = NULL, ...) {
  deviance_tweedie(actual, predicted, w, tweedie_p = 1.5, ...)
}
r_squared(1:10, c(1, 1:9), deviance_function = myTweedie)
```

r_squared_bernoulli *Pseudo R-Squared regarding Bernoulli deviance*

Description

Wrapper to `r_squared` with `deviance_function = deviance_bernoulli`.

Usage

```
r_squared_bernoulli(actual, predicted, w = NULL, reference_mean = NULL, ...)
```

Arguments

actual	Observed values.
predicted	Predicted values.
w	Optional case weights.
reference_mean	An optional reference mean used to derive the null deviance. Recommended in out-of-sample applications.
...	Further arguments passed to <code>r_squared</code> .

Details

The deviance gain is calculated regarding the null model derived from the actual values. While fine for in-sample considerations, this is only an approximation for out-of-sample considerations. There, it is recommended to calculate null deviance regarding the in-sample (weighted) mean. This value can be passed by the argument `reference_mean`.

Value

A numeric vector of length one.

See Also

[r_squared](#).

Examples

```
r_squared(c(0, 0, 1, 1), c(0.1, 0.1, 0.9, 0.8), w = 1:4,
  deviance_function = deviance_bernoulli)
r_squared_bernoulli(c(0, 0, 1, 1), c(0.1, 0.1, 0.9, 0.8), w = 1:4)
```

r_squared_gamma

Pseudo R-Squared regarding Gamma deviance

Description

Wrapper to `r_squared` with `deviance_function = deviance_gamma`.

Usage

```
r_squared_gamma(actual, predicted, w = NULL, reference_mean = NULL, ...)
```

Arguments

<code>actual</code>	Observed values.
<code>predicted</code>	Predicted values.
<code>w</code>	Optional case weights.
<code>reference_mean</code>	An optional reference mean used to derive the null deviance. Recommended in out-of-sample applications.
<code>...</code>	Further arguments passed to <code>r_squared</code> .

Details

The deviance gain is calculated regarding the null model derived from the actual values. While fine for in-sample considerations, this is only an approximation for out-of-sample considerations. There, it is recommended to calculate null deviance regarding the in-sample (weighted) mean. This value can be passed by the argument `reference_mean`.

Value

A numeric vector of length one.

See Also

[r_squared](#).

Examples

```
r_squared(1:10, c(1:9, 12), w = 1:10, deviance_function = deviance_gamma)
r_squared_gamma(1:10, c(1:9, 12), w = 1:10)
```

r_squared_poisson	<i>Pseudo R-Squared regarding Poisson deviance</i>
-------------------	--

Description

Wrapper to `r_squared` with `deviance_function = deviance_poisson`.

Usage

```
r_squared_poisson(actual, predicted, w = NULL, reference_mean = NULL, ...)
```

Arguments

actual	Observed values.
predicted	Predicted values.
w	Optional case weights.
reference_mean	An optional reference mean used to derive the null deviance. Recommended in out-of-sample applications.
...	Further arguments passed to <code>r_squared</code> .

Details

The deviance gain is calculated regarding the null model derived from the actual values. While fine for in-sample considerations, this is only an approximation for out-of-sample considerations. There, it is recommended to calculate null deviance regarding the in-sample (weighted) mean. This value can be passed by the argument `reference_mean`.

Value

A numeric vector of length one.

See Also

[r_squared](#).

Examples

```
r_squared(0:2, c(0.1, 1, 2), w = rep(1, 3),
          deviance_function = deviance_poisson)
r_squared_poisson(0:2, c(0.1, 1, 2), w = rep(1, 3))
```

`weighted_cor`*Weighted Pearson Correlation*

Description

Calculates weighted Pearson correlation coefficient between observed and predicted values by the help of `stats::cov.wt`.

Usage

```
weighted_cor(actual, predicted, w = NULL, na.rm = FALSE, ...)
```

Arguments

<code>actual</code>	Observed values.
<code>predicted</code>	Predicted values.
<code>w</code>	Optional case weights.
<code>na.rm</code>	Should missing values in observed or predicted be removed? Default is FALSE.
<code>...</code>	Further arguments passed to <code>stats::cov.wt</code> .

Value

A length-one numeric vector.

See Also

[weighted_mean](#).

Examples

```
weighted_cor(1:10, c(1, 1:9))
weighted_cor(1:10, c(1, 1:9), w = 1:10)
```

weighted_mean	<i>Weighted Mean</i>
---------------	----------------------

Description

Returns weighted mean of a numeric vector. In contrast to `stats::weighted.mean`, `w` does not need to be specified.

Usage

```
weighted_mean(x, w = NULL, ...)
```

Arguments

<code>x</code>	Numeric vector.
<code>w</code>	Optional non-negative, non-missing case weights.
<code>...</code>	Further arguments passed to <code>mean</code> or <code>weighted.mean</code> .

Value

A length-one numeric vector.

See Also

[weighted_quantile](#).

Examples

```
weighted_mean(1:10)
weighted_mean(1:10, w = NULL)
weighted_mean(1:10, w = 1:10)
```

weighted_median	<i>Weighted Median</i>
-----------------	------------------------

Description

Calculates weighted median. For odd sample sizes consistent with unweighted quantiles.

Usage

```
weighted_median(x, w = NULL, ...)
```

Arguments

x Numeric vector.
w Optional non-negative case weights.
... Further arguments passed to weighted_quantile.

See Also

[weighted_quantile](#).

Examples

```
n <- 21
x <- seq_len(n)
quantile(x, probs = 0.5)
weighted_median(x, w = rep(1, n))
weighted_median(x, w = x)
quantile(rep(x, x), probs = 0.5)
```

weighted_quantile *Weighted Quantiles*

Description

Calculates weighted quantiles based on the generalized inverse of the weighted ECDF. If no weights are passed, uses stats::quantile.

Usage

```
weighted_quantile(
  x,
  w = NULL,
  probs = seq(0, 1, 0.25),
  na.rm = TRUE,
  names = TRUE,
  ...
)
```

Arguments

x Numeric vector.
w Optional non-negative case weights.
probs Vector of probabilities.
na.rm Ignore missing data?
names Return names?
... Further arguments passed to stats::quantile in the unweighted case. Not used in the weighted case.

See Also

[weighted_median](#).

Examples

```
n <- 10
x <- seq_len(n)
quantile(x)
weighted_quantile(x)
weighted_quantile(x, w = rep(1, n))
quantile(x, type = 1)
weighted_quantile(x, w = x) # same as Hmisc::wtd.quantile
weighted_quantile(x, w = x, names = FALSE)
weighted_quantile(x, w = x, probs = 0.5, names = FALSE)

# Example with integer weights
x <- c(1, 1:11, 11, 11)
w <- seq_along(x)
weighted_quantile(x, w)
quantile(rep(x, w)) # same
```

weighted_var

Weighted Variance

Description

Calculates weighted variance, see `stats::cov.wt` or https://en.wikipedia.org/wiki/Sample_mean_and_covariance#Weighted_samples for details.

Usage

```
weighted_var(x, w = NULL, method = c("unbiased", "ML"), na.rm = FALSE, ...)
```

Arguments

<code>x</code>	Numeric vector.
<code>w</code>	Optional non-negative, non-missing case weights.
<code>method</code>	Specifies how the result is scaled. If "unbiased", the denominator is reduced by -1, unlike "ML". See <code>stats::cov.wt</code> for details.
<code>na.rm</code>	Should missing values in <code>x</code> be removed? Default is FALSE.
<code>...</code>	Further arguments passed to <code>stats::cov.wt</code> .

Value

A length-one numeric vector.

See Also

[weighted_mean](#).

Examples

```
weighted_var(1:10)
weighted_var(1:10, w = NULL)
weighted_var(1:10, w = rep(1, 10))
weighted_var(1:10, w = 1:10)
weighted_var(1:10, w = 1:10, method = "ML")
```

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