

Package ‘mlf’

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Type Package

Title Machine Learning Foundations

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Description Offers a gentle introduction to machine learning concepts for practitioners with a statistical pedigree: decomposition of model error (bias-variance trade-off), nonlinear correlations, information theory and functional permutation/bootstrap simulations. Székely GJ, Rizzo ML, Bakirov NK. (2007). <[doi:10.1214/009053607000000505](https://doi.org/10.1214/009053607000000505)>. Reshef DN, Reshef YA, Finucane HK, Grossman SR, McVean G, Turnbaugh PJ, Lander ES, Mitzenmacher M, Sabatti PC. (2011). <[doi:10.1126/science.1205438](https://doi.org/10.1126/science.1205438)>.

Imports stats, utils

URL <http://mlf-project.us/>

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boot

*Bootstrap Confidence Intervals via Resampling***Description**

Provides nonparametric confidence intervals via percentile-based resampling for given `mlf` function.

Usage

```
boot(x, y, func, reps, conf.int)
```

Arguments

x, y	numeric vectors of data values
func	specify <code>mlf</code> function
reps	(optional) number of resamples. Defaults to 500
conf.int	(optional) numeric value indicating level of confidence. Defaults to 0.90.

Examples

```
# Sample data
a <- rnorm(25, 80, 35)
b <- rnorm(25, 100, 50)

mlf::mic(a, b)
mlf::boot(a, b, mic)
```

bvto	<i>Bias-Variance Trade-Off</i>
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Description

Provides estimated error decomposition from model predictions (mse, bias, variance).

Usage

```
bvto(truth, estimate)
```

Arguments

truth	test data vector or baseline accuracctruth to test against.
estimate	predicted vector

Examples

```
# Sample data
test <- rnorm(25, 80, 35)
predicted <- rnorm(25, 80, 50)

mlf::bvto(test, predicted)
```

distcorr	<i>Distance Correlation</i>
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Description

Provides pairwise correlation via distance covariance normalized by distance standard deviation.
Allows for non-linear dependencies.

Usage

```
distcorr(x, y)
```

Arguments

x, y	numeric vectors of data values
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References

Székely GJ, Rizzo ML, Bakirov NK. Measuring and testing dependence by correlation of distances. Ann Stat. 2007. 35(6):2769-2794.

Examples

```
# Sample data
a <- rnorm(25, 80, 35)
b <- rnorm(25, 100, 50)

mlf::distcorr(a, b)
```

entropy

Entropy

Description

Estimates uncertainty in univariate probability distribution.

Usage

```
entropy(x, bins)
```

Arguments

<code>x</code>	numeric or discrete data vector
<code>bins</code>	specify number of bins if numeric or integer data class.

Examples

```
# Sample numeric vector
a <- rnorm(25, 80, 35)
mlf::entropy(a, bins = 2)

# Sample discrete vector
b <- as.factor(c(1,1,1,2))
mlf::entropy(b)
```

get_bias

Bias

Description

Estimates squared bias by decomposing model prediction error.

Usage

```
get_bias(truth, estimate)
```

Arguments

- | | |
|----------|--|
| truth | test data vector or baseline accuracy to test against. |
| estimate | predicted vector |

Examples

```
# Sample data
test <- rnorm(25, 80, 35)
predicted <- rnorm(25, 80, 50)

mlf::get_bias(test, predicted)
```

get_mse*Mean Squared Error*

Description

Estimates mean squared error from model predictions.

Usage

```
get_mse(truth, estimate)
```

Arguments

- | | |
|----------|--|
| truth | test data vector or baseline accuracy to test against. |
| estimate | predicted vector |

Examples

```
# Sample data
test <- rnorm(25, 80, 35)
predicted <- rnorm(25, 80, 50)

mlf::get_mse(test, predicted)
```

<code>get_var</code>	<i>Variance</i>
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Description

Estimates squared variance by decomposing model prediction error.

Usage

```
get_var(estimate)
```

Arguments

<code>estimate</code>	predicted vector
-----------------------	------------------

Examples

```
# Sample data
test <- rnorm(25, 80, 35)
predicted <- rnorm(25, 80, 50)

mlf::get_var(predicted)
```

<code>jointentropy</code>	<i>Joint Entropy</i>
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Description

Estimated difference between two probability distributions.

Usage

```
jointentropy(x, y, bins)
```

Arguments

<code>x, y</code>	numeric or discrete data vectors
<code>bins</code>	specify number of bins

Examples

```
# Sample numeric vector  
a <- rnorm(25, 80, 35)  
b <- rnorm(25, 90, 35)  
mlf::jointentropy(a, b, bins = 2)  
  
# Sample discrete vector  
a <- as.factor(c(1,1,2,2))  
b <- as.factor(c(1,1,1,2))  
mlf::jointentropy(a, b)
```

kld

Kullback-Leibler Divergence

Description

Provides estimated difference between individual entropy and cross-entropy of two probability distributions.

Usage

```
kld(x, y, bins)
```

Arguments

x, y	numeric or discrete data vectors
bins	specify number of bins

Examples

```
# Sample numeric vector  
a <- rnorm(25, 80, 35)  
b <- rnorm(25, 90, 35)  
mlf::kld(a, b, bins = 2)  
  
# Sample discrete vector  
a <- as.factor(c(1,1,2,2))  
b <- as.factor(c(1,1,1,2))  
mlf::kld(a, b)
```

mi*Mutual Information***Description**

Estimates Kullback-Leibler divergence of joint distribution and the product of two respective marginal distributions. Roughly speaking, the amount of information one variable provides about another.

Usage

```
mi(x, y)
```

Arguments

<code>x, y</code>	numeric or discrete data vectors
-------------------	----------------------------------

Examples

```
# Sample data
a <- rnorm(25, 80, 35)
b <- rnorm(25, 100, 50)

mlf::mi(a, b)
```

mic*Maximal Information Criterion***Description**

Information-theoretic approach for detecting non-linear pairwise dependencies. Employs heuristic discretization to achieve highest normalized mutual information.

Usage

```
mic(x, y)
```

Arguments

<code>x, y</code>	numeric or discrete data vectors
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References

Reshef DN, Reshef YA, Finucane HK, Grossman SR, McVean G, Turnbaugh PJ, Lander ES, Mitzenmacher M, Sabeti PC. Detecting novel associations in large data sets. *Science*. 2011. 334(6062):1518-1524.

Examples

```
# Sample data  
a <- rnorm(25, 80, 35)  
b <- rnorm(25, 100, 50)  
  
mlf::mic(a, b)
```

perm

Permutation Test

Description

Provides nonparametric statistical significance via sample randomization.

Usage

```
perm(x, y, func, reps)
```

Arguments

x, y	numeric vectors of data values
func	specify mlf function: (distcorr or mic).
reps	(optional) number of resamples. Defaults to 500.

Examples

```
# Sample data  
a <- rnorm(25, 80, 35)  
b <- rnorm(25, 100, 50)  
  
mlf::mic(a, b)  
mlf::perm(a, b, mic)
```

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