

Package ‘Tariff’

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Type Package

Title Replicate Tariff Method for Verbal Autopsy

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Description Implement the Tariff algorithm for coding cause-of-death from verbal autopsies. The Tariff method was originally proposed in James et al (2011) <DOI:10.1186/1478-7954-9-31> and later refined as Tariff 2.0 in Serina, et al. (2015) <DOI:10.1186/s12916-015-0527-9>. Note that this package was not developed by authors affiliated with the Institute for Health Metrics and Evaluation and thus unintentional discrepancies may exist between the this implementation and the implementation available from IHME.

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RoxygenNote 6.1.0

NeedsCompilation no

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plot.tariff	<i>Plot CSMF of the results obtained from Tariff algorithm</i>
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Description

This function plots the CSMF of the fitted results.

Usage

```
## S3 method for class 'tariff'
plot(x, top = NULL, min.prob = 0, ...)
```

Arguments

x	fitted object from <code>tariff</code>
top	maximum causes to plot
min.prob	minimum fraction for the causes plotted
...	Arguments to be passed to/from graphic function

Examples

```
data("RandomVA3")
test <- RandomVA3[1:200, ]
train <- RandomVA3[201:400, ]
allcauses <- unique(train$cause)
fit <- tariff(causes.train = "cause", symps.train = train,
             symps.test = test, causes.table = allcauses)
plot(fit, top = 10, main = "Top 5 population COD distribution")
plot(fit, min.prob = 0.05, main = "Ppulation COD distribution (at least 5%)")
```

print.tariff_summary	<i>Print method for the summary of the results obtained from Tariff algorithm</i>
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Description

This function prints the summary message of the fitted results.

Usage

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

Arguments

x	summary object for Tariff fit
...	not used

RandomVA3	<i>400 records of Sample Input</i>
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Description

This is a dataset consisting of 400 arbitrary sample input deaths randomly sampled from cleaned PHMRC data.

Format

400 arbitrary input records.

Examples

```
data(RandomVA3)
head(RandomVA3$train)
head(RandomVA3$test)
```

SampleCategory3	<i>Grouping of causes in RandomVA3</i>
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Description

This is a matrix specifying a default grouping of the causes used in RandomVA3.

Format

17 by 2 matrix

Examples

```
data(SampleCategory3)
SampleCategory3
```

summary.tariff	<i>Summary of the results obtained from Tariff algorithm</i>
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Description

This function prints the summary message of the fitted results.

Usage

```
## S3 method for class 'tariff'
summary(object, top = 5, id = NULL, ...)
```

Arguments

object	fitted object from <code>tariff</code>
top	number of top CSMF to show
id	the ID of a specific death to show
...	not used

Examples

```
data("RandomVA3")
test <- RandomVA3[1:200, ]
train <- RandomVA3[201:400, ]
allcauses <- unique(train$cause)
fit <- tariff(causes.train = "cause", symps.train = train,
             symps.test = test, causes.table = allcauses)
correct <- which(fit$causes.test[,2] == test$cause)
accuracy <- length(correct) / dim(test)[1]
summary(fit)
summary(fit, top = 10)
summary(fit, id = "p849", top = 3)
```

tariff	<i>Replicate Tariff methods</i>
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Description

This function implements Tariff method.

Usage

```
tariff(causes.train, symps.train, symps.test, causes.table = NULL,
       use.rank = TRUE, nboot.rank = 1, use.sig = TRUE, nboot.sig = 500,
       use.top = FALSE, ntop = 40, ...)
```

Arguments

<code>causes.train</code>	character vector of causes, or the column name of cause in the training data
<code>symps.train</code>	N.train by S matrix
<code>symps.test</code>	N.test by S matrix
<code>causes.table</code>	list of causes in the data
<code>use.rank</code>	logical indicator for whether using ranks instead of scores
<code>nboot.rank</code>	number of re-sampling for baseline rank comparison. Default to 1, which re-samples training data to have a uniform cause distribution of the same size. Set this to 0 removes bootstrapping the training dataset.
<code>use.sig</code>	logical indicator for whether using significant Tariff only
<code>nboot.sig</code>	number of re-sampling for testing significance.
<code>use.top</code>	logical indicator for whether the tariff matrix should be cleaned to have only top symptoms
<code>ntop</code>	number of top tariff kept for each cause
<code>...</code>	not used

Value

<code>score</code>	matrix of score for each cause within each death
<code>causes.train</code>	vector of most likely causes in training data
<code>causes.test</code>	vector of most likely causes in testing data
<code>csmf</code>	vector of CSMF
<code>causes.table</code>	cause list used for output, i.e., list of existing causes in the training data
<code>use.rank</code>	logical indicator for whether using ranks instead of scores

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References

James, S. L., Flaxman, A. D., Murray, C. J., & Population Health Metrics Research Consortium. (2011). *Performance of the Tariff Method: validation of a simple additive algorithm for analysis of verbal autopsies*. *Population Health Metrics*, 9(1), 1-16.

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Tyler H. McCormick, Zehang R. Li, Clara Calvert, Amelia C. Crampin, Kathleen Kahn and Samuel J. Clark(2016) *Probabilistic cause-of-death assignment using verbal autopsies*, <http://arxiv.org/abs/1411.3042> To appear, *Journal of the American Statistical Association*

Examples

```
data("RandomVA3")
test <- RandomVA3[1:200, ]
train <- RandomVA3[201:400, ]
allcauses <- unique(train$cause)
fit <- tariff(causes.train = "cause", symps.train = train,
symps.test = test, causes.table = allcauses)
correct <- which(fit$causes.test[,2] == test$cause)
accuracy <- length(correct) / dim(test)[1]
```

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