Package 'synthesizer'

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Type Package

Title Synthesize Data Based on Empirical Quantile Functions and Rank Order Matching	
Version 0.4.0	
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Description Data is synthesized using a combination of inverse transform sampling using the empirical quantile functions for each variable, and then copying the rank order structure from the original dataset. The syntesizer method has a tunable parameter allowing to gradually move from realistic and possibly unsafe synthetic data to decorrelated data of less utility. License EUPL	
URL https://github.com/markvanderloo/synthesizer	
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make_synthesizer

Create a function that generates synthetic data

Description

Create a function that accepts a non-negative integer n, and that returns synthetic data sampled from the emperical (multivariate) distribution of x.

Usage

```
make_synthesizer(x, ...)
## S3 method for class 'numeric'
make_synthesizer(x, ...)
## S3 method for class 'integer'
make_synthesizer(x, ...)
## S3 method for class 'logical'
make_synthesizer(x, ...)
## S3 method for class 'factor'
make_synthesizer(x, ...)
## S3 method for class 'character'
make_synthesizer(x, ...)
## S3 method for class 'ts'
make_synthesizer(x, ...)
## S3 method for class 'data.frame'
make_synthesizer(x, rankcor = 1, ...)
```

Arguments

x [vector|data.frame] Template data to be synthesized.
... arguments passed to other methods

rankcor [numeric] in (0,1] The correlations between the ranks of the real data and synthetic data. Either a single number or a vector of the form c("variable1"=x1,...).

Only used if x is a data frame.

Value

A function accepting a single integer argument: the number of synthesized values or records to return.

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See Also

```
Other synthesis: synthesize()
```

Examples

```
synth <- make_synthesizer(cars$speed)
synth(10)

synth <- make_synthesizer(iris)
synth(6)
synth(150)
synth(250)</pre>
```

pmse

Compute the pMSE metric between synthetic and real data

Description

The propensity mean squared error is defined as $\frac{1}{N} \sum_{i=1}^{N} (p_i - c)^2$, where c is the number of synthetic records, divided by the sum of the number of synthetic and real records.

Usage

```
pmse(synth, real, model = c("lr", "rf"), nrep = NULL)
```

Arguments

synth [data.frame] Synthesized data.

real [real] Data to compare with the synthesized data.

model [character] Model used to compute propensity scores. Options are "1r": lo-

gistic regression, and "rf": random forest.

nrep [integer] Number of model repetitions to average the pMSE value over. Ig-

nored for 1r.

Value

[numeric] scalar.

Examples

```
scars <- synthesize(cars)
pmse(scars, cars)</pre>
```

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synthesize

Create synthetic version of a dataset

Description

Create n values or records based on the emperical (multivariate) distribution of y. For data frames it is possible to decorrelate synthetic from the original variables by lowering the value for the rankcor parameter.

Usage

```
synthesize(x, n = NROW(x), rankcor = 1)
```

Arguments

x [vector|data.frame] data to synthesize.

n [integer] Number of values or records to synthesize.

rankcor [numeric] in [0,1]. Either a single rank correlation value that is applied to all

variables, or a vector of the form c(variable1=ut1lity1,...). Variables not

explicitly mentioned will have rankcor=1. See also the note below.

Value

A data object of the same type and structure as x.

Note

The utility of a synthetic variable is lowered by decorelating the rank correlation between the real and synthetic data. If rankcor=1, the synthetic data will ordered such that it has the same rank order as the original data. If rankcor=0, no such reordering will take place. For values between 0 and 1, blocks of data are randomly selected and randomly permuted iteratively until the rank correlation between original and synthetic data drops below the parameter.

See Also

Other synthesis: make_synthesizer()

Examples

```
synthesize(cars$speed,10)
synthesize(cars)
synthesize(cars,25)

s1 <- synthesize(iris, rankcor=1)
s2 <- synthesize(iris, rankcor=0.5)
s3 <- synthesize(iris, rankcor=c("Species"=0.5))
oldpar <- par(mfrow=c(2,2), pch=16, las=1)</pre>
```

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```
plot(Sepal.Length ~ Sepal.Width, data=iris, col=iris$Species, main="Iris")
plot(Sepal.Length ~ Sepal.Width, data=s1, col=s1$Species, main="Synthetic Iris")
plot(Sepal.Length ~ Sepal.Width, data=s2, col=s2$Species, main="Low utility Iris")
plot(Sepal.Length ~ Sepal.Width, data=s3, col=s3$Species, main="Low utility Species")
par(oldpar)
```

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