# Package 'SKIT'

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Title SKIT

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**Description** Conduct the Semiparametric Kernel Independence Test between two vectors when there are excess zeros.

Imports stats, utils

License GPL-2 Needs

**Compilation** yes

## **R** topics documented:

SKIT-package Semiparametric Kernel Independence Test (SKIT) with excess zeros

#### Description

Conduct the test of independence between two vectors when there are excess zeros.

### Details

The function SKIT() compute the test statistics and compute their *p*-values based on bootstrap samples.

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#### skit

#### Description

Performs the test of independence between two vectors when there are excess zeros.

#### Usage

skit(x, y, bandwidth=NULL, nboot=1000, print=0)

#### Arguments

Х	Numeric vector possibly containing zeros.
У	Numeric vector possibly containing zeros with same length as x.
bandwidth	Bandwith parameter (see details). The default is NULL so that a default value will be computed from the data.
nboot	Number of bootstrap samples to estimate p-values. If p-values are not desired, then set to 0. The default is 1000.
print	0 or 1 to print information. The default is 0.

#### Details

Non-numeric values in x or y will be removed from both vectors.

If bandwidth = NULL, then the bandwidth parameter will be computed for the observed data, and re-computed for each bootstrap sample. It is computed as  $sigma*n^{(-0.2)}$ , where n = length (x), sigma = max(c(sd(x[x != 0]), sd(y[y != 0]))).

#### Value

A list containing the observed test statistics, bootstrap-estimated p-values, and the bandwith parameter for the observed data.

The overall test statistic T is computed as  $T_1 + T_2 + T_3 + T_4$  and for each component, p-value is estimated via bootstrap.

#### Examples

```
set.seed(123)
n <- 500
b00 <- 0.09; b10 <- 0.21; b01 <- 0.21; b11 <- 0.49
id <- sample(1:4, n, replace = TRUE, prob = c(b00, b10, b01, b11))
subn <- table(factor(id, levels = 1:4))
n2 <- subn[names(subn) == "2"]
n3 <- subn[names(subn) == "3"]
n4 <- subn[names(subn) == "4"]
Biv <- matrix(0, nrow = n, ncol = 2)
if(n2 != 0) Biv[id == 2,1] <- rnorm(n2, 0.4, 0.1)
if(n3 != 0) Biv[id == 3,2] <- rnorm(n3, 0.4, 0.1)
if(n4 != 0) Biv[id == 4,] <- cbind(rnorm(n4, 0.4, 0.1), rnorm(n4, 0.4, 0.1))
x <- Biv[,1]</pre>
```

y <- Biv[,2]
skit(x, y, nboot=0)</pre>