

Package ‘tEDM’

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Title Temporal Empirical Dynamic Modeling

Version 1.0

Description Inferring causation from time series data through empirical dynamic modeling (EDM), with methods such as convergent cross mapping from Sugihara et al. (2012) <[doi:10.1126/science.1227079](https://doi.org/10.1126/science.1227079)>, partial cross mapping as outlined in Leng et al. (2020) <[doi:10.1038/s41467-020-16238-0](https://doi.org/10.1038/s41467-020-16238-0)>, and cross mapping cardinality as described in Tao et al. (2023) <[doi:10.1016/j.fmre.2023.01.007](https://doi.org/10.1016/j.fmre.2023.01.007)>.

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Encoding UTF-8

RoxygenNote 7.3.2

URL <https://stsc1.github.io/tEDM/>, <https://github.com/stsc1/tEDM>

BugReports <https://github.com/stsc1/tEDM/issues>

Depends R (>= 4.1.0)

LinkingTo Rcpp, RcppThread, RcppArmadillo

Imports dplyr, ggplot2, methods, Rcpp

Suggests RcppThread, RcppArmadillo, readr, plot3D, spEDM, knitr, rmarkdown, purrr, tidyr, cowplot

VignetteBuilder knitr

NeedsCompilation yes

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ccm	<i>convergent cross mapping</i>
-----	---------------------------------

Description

convergent cross mapping

Usage

```
## S4 method for signature 'data.frame'
ccm(
  data,
  cause,
  effect,
  libsizes = NULL,
  E = 3,
  tau = 0,
  k = E + 1,
  theta = 1,
  algorithm = "simplex",
  lib = NULL,
  pred = NULL,
  threads = length(libsizes),
  parallel.level = "low",
  bidirectional = TRUE,
  progressbar = TRUE
)
```

Arguments

- data observation data.
- cause name of causal variable.
- effect name of effect variable.
- libsizes (optional) number of time points used.
- E (optional) embedding dimensions.
- tau (optional) step of time lags.

k (optional) number of nearest neighbors.
 theta (optional) weighting parameter for distances, useful when algorithm is smap.
 algorithm (optional) prediction algorithm.
 lib (optional) libraries indices.
 pred (optional) predictions indices.
 threads (optional) number of threads to use.
 parallel.level (optional) level of parallelism, low or high.
 bidirectional (optional) whether to examine bidirectional causality.
 progressbar (optional) whether to show the progress bar.

Value

A list
 xmap cross mapping results
 varname names of causal and effect variable
 bidirectional whether to examine bidirectional causality

References

Sugihara, G., May, R., Ye, H., Hsieh, C., Deyle, E., Fogarty, M., Munch, S., 2012. Detecting Causality in Complex Ecosystems. Science 338, 496–500.

Examples

```

sim = logistic_map(x = 0.4,y = 0.4,step = 45,beta_xy = 0.5,beta_yx = 0)
ccm(sim,"x","y",libsizes = seq(5,35,5),E = 8,k = 7,threads = 1)

```

cmc	<i>cross mapping cardinality</i>
-----	----------------------------------

Description

cross mapping cardinality

Usage

```

## S4 method for signature 'data.frame'
cmc(
  data,
  cause,
  effect,
  libsizes = NULL,
  E = 3,

```

```

    tau = 0,
    k = pmin(E^2),
    lib = NULL,
    pred = NULL,
    threads = length(libsizes),
    parallel.level = "low",
    bidirectional = TRUE,
    progressbar = TRUE
  )

```

Arguments

<code>data</code>	observation data.
<code>cause</code>	name of causal variable.
<code>effect</code>	name of effect variable.
<code>libsizes</code>	(optional) number of time points used.
<code>E</code>	(optional) embedding dimensions.
<code>tau</code>	(optional) step of time lags.
<code>k</code>	(optional) number of nearest neighbors.
<code>lib</code>	(optional) libraries indices.
<code>pred</code>	(optional) predictions indices.
<code>threads</code>	(optional) number of threads to use.
<code>parallel.level</code>	(optional) level of parallelism, low or high.
<code>bidirectional</code>	(optional) whether to examine bidirectional causality.
<code>progressbar</code>	(optional) whether to show the progress bar.

Value

A list

- `xmap` cross mapping results
- `cs` causal strength
- `varname` names of causal and effect variable
- `bidirectional` whether to examine bidirectional causality

References

Tao, P., Wang, Q., Shi, J., Hao, X., Liu, X., Min, B., Zhang, Y., Li, C., Cui, H., Chen, L., 2023. Detecting dynamical causality by intersection cardinal concavity. *Fundamental Research*.

Examples

```

sim = logistic_map(x = 0.4, y = 0.4, step = 45, beta_xy = 0.5, beta_yx = 0)
cmc(sim, "x", "y", E = 4, k = 15, threads = 1)

```

embedded	<i>embedding time series data</i>
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Description

embedding time series data

Usage

```
## S4 method for signature 'data.frame'
embedded(data, target, E = 3, tau = 0)
```

Arguments

data	observation data.
target	name of target variable.
E	(optional) embedding dimensions.
tau	(optional) step of time lags.

Value

A matrix

Examples

```
embedded(data.frame(t = 1:5), "t", 3)
```

fnn	<i>false nearest neighbours</i>
-----	---------------------------------

Description

false nearest neighbours

Usage

```
## S4 method for signature 'data.frame'
fnn(
  data,
  target,
  lib = NULL,
  pred = NULL,
  E = 2:10,
  tau = 0,
```

```
    rt = 10,  
    eps = 2,  
    threads = length(E)  
  )
```

Arguments

data	observation data.
target	name of target variable.
lib	(optional) libraries indices.
pred	(optional) predictions indices.
E	(optional) embedding dimensions.
tau	(optional) step of time lags.
rt	(optional) escape factor.
eps	(optional) neighborhood diameter.
threads	(optional) number of threads to use.

Value

A vector

References

Kennel M. B., Brown R. and Abarbanel H. D. I., Determining embedding dimension for phase-space reconstruction using a geometrical construction, Phys. Rev. A, Volume 45, 3403 (1992).

Examples

```
sim = logistic_map(x = 0.4,y = 0.4,step = 45,beta_xy = 0.5,beta_yx = 0)  
fnn(sim,"x",threads = 1)
```

ic	<i>intersection cardinality</i>
----	---------------------------------

Description

intersection cardinality

Usage

```
## S4 method for signature 'data.frame'
ic(
  data,
  column,
  target,
  lib = NULL,
  pred = NULL,
  E = 2:10,
  tau = 0,
  k = E + 2,
  threads = length(pred),
  parallel.level = "low"
)
```

Arguments

data	observation data.
column	name of library variable.
target	name of target variable.
lib	(optional) libraries indices.
pred	(optional) predictions indices.
E	(optional) embedding dimensions.
tau	(optional) step of time lags.
k	(optional) number of nearest neighbors used in prediction.
threads	(optional) number of threads to use.
parallel.level	(optional) level of parallelism, low or high.

Value

A list

- xmap cross mapping performance
- varname name of target variable
- method method of cross mapping
- tau step of time lag

References

Tao, P., Wang, Q., Shi, J., Hao, X., Liu, X., Min, B., Zhang, Y., Li, C., Cui, H., Chen, L., 2023. Detecting dynamical causality by intersection cardinal concavity. Fundamental Research.

Examples

```
sim = logistic_map(x = 0.4, y = 0.4, step = 45, beta_xy = 0.5, beta_yx = 0)
ic(sim, "x", "y", E = 4, k = 15:30, threads = 1)
```

logistic_map	<i>logistic map</i>
--------------	---------------------

Description

logistic map

Usage

```
logistic_map(
  x,
  y = NULL,
  z = NULL,
  step = 15,
  alpha_x = 3.6,
  alpha_y = 3.72,
  alpha_z = 3.68,
  beta_xy = 0.05,
  beta_xz = 0.05,
  beta_yx = 0.2,
  beta_yz = 0.2,
  beta_zx = 0.35,
  beta_zy = 0.35,
  threshold = Inf,
  transient = 1
)
```

Arguments

x	value x.
y	(optional) value y.
z	(optional) value z.
step	(optional) number of simulation time steps.
alpha_x	(optional) growth parameter for x.
alpha_y	(optional) growth parameter for y.
alpha_z	(optional) growth parameter for y.
beta_xy	(optional) cross-inhibition from x to y.
beta_xz	(optional) cross-inhibition from x to z.
beta_yx	(optional) cross-inhibition from y to x.
beta_yz	(optional) cross-inhibition from y to z.
beta_zx	(optional) cross-inhibition from z to x.
beta_zy	(optional) cross-inhibition from z to y.
threshold	(optional) set to NaN if the absolute value exceeds this threshold.
transient	(optional) transients to be excluded from the results.

Value

A data.frame

Examples

```
logistic_map(x = 0.2)
```

multispatialccm

multispatial convergent cross mapping

Description

multispatial convergent cross mapping

Usage

```
## S4 method for signature 'list'
multispatialccm(
  data,
  cause,
  effect,
  libsizes,
  E = 3,
  tau = 0,
  k = E + 1,
  boot = 99,
  seed = 42,
  threads = length(libsizes),
  parallel.level = "low",
  bidirectional = TRUE,
  progressbar = TRUE
)
```

Arguments

data	observation data.
cause	name of causal variable.
effect	name of effect variable.
libsizes	number of time points used in prediction.
E	(optional) embedding dimensions.
tau	(optional) step of time lags.
k	(optional) number of nearest neighbors used in prediction.
boot	(optional) number of bootstraps to perform.

seed (optional) random seed.
 threads (optional) number of threads to use.
 parallel.level (optional) level of parallelism, low or high.
 bidirectional (optional) whether to examine bidirectional causality.
 progressbar (optional) whether to show the progress bar.

Value

A list
 xmap cross mapping results
 varname names of causal and effect variable
 bidirectional whether to examine bidirectional causality

References

Clark, A.T., Ye, H., Isbell, F., Deyle, E.R., Cowles, J., Tilman, G.D., Sugihara, G., 2015. Spatial convergent cross mapping to detect causal relationships from short time series. *Ecology* 96, 1174–1181.

Examples

```

set.seed(42)
obs = runif(15,0,0.1)
sim = vector("list",15)
for (i in seq_along(obs)){
  sim[[i]] = logistic_map(x = obs[i],y = obs[i],step = 15,beta_xy = 0.5,beta_yx = 0)
}
lst = list(x = do.call(cbind, lapply(sim, function(df) df$x)),
          y = do.call(cbind, lapply(sim, function(df) df$y)))
multispatialccm(lst,"x","y",libsizes = 5:15,E = c(7,2),k = 6,threads = 1)

```

pcm

partial cross mapping

Description

partial cross mapping

Usage

```
## S4 method for signature 'data.frame'
pcm(
  data,
  cause,
  effect,
  conds,
  libsizes = NULL,
  E = 3,
  tau = 0,
  k = E + 1,
  theta = 1,
  algorithm = "simplex",
  lib = NULL,
  pred = NULL,
  threads = length(libsizes),
  parallel.level = "low",
  bidirectional = TRUE,
  cumulate = FALSE,
  progressbar = TRUE
)
```

Arguments

<code>data</code>	observation data.
<code>cause</code>	name of causal variable.
<code>effect</code>	name of effect variable.
<code>conds</code>	name of conditioning variables.
<code>libsizes</code>	(optional) number of time points used.
<code>E</code>	(optional) embedding dimensions.
<code>tau</code>	(optional) step of time lags.
<code>k</code>	(optional) number of nearest neighbors.
<code>theta</code>	(optional) weighting parameter for distances, useful when <code>algorithm</code> is <code>smap</code> .
<code>algorithm</code>	(optional) prediction algorithm.
<code>lib</code>	(optional) libraries indices.
<code>pred</code>	(optional) predictions indices.
<code>threads</code>	(optional) number of threads to use.
<code>parallel.level</code>	(optional) level of parallelism, low or high.
<code>bidirectional</code>	(optional) whether to examine bidirectional causality.
<code>cumulate</code>	(optional) serial or cumulative computation of partial cross mapping.
<code>progressbar</code>	(optional) whether to show the progress bar.

Value

A list

pxmap partial cross mapping results

xmap cross mapping results

varname names of causal and effect variable

bidirectional whether to examine bidirectional causality

References

Leng, S., Ma, H., Kurths, J. et al. Partial cross mapping eliminates indirect causal influences. Nat Commun 11, 2632 (2020).

Examples

```
sim = logistic_map(x = 0.4,y = 0.4,z = 0.4,step = 45,
                  beta_xy = 0.5, beta_xz = 0,
                  beta_yx = 0, beta_yz = 0.5,
                  beta_zx = 0, beta_zy = 0)
pcm(sim,"x","z","y",libsizes = seq(5,35,5),E = 10,k = 7,threads = 1)
```

simplex

simplex forecast

Description

simplex forecast

Usage

```
## S4 method for signature 'data.frame'
simplex(
  data,
  column,
  target,
  lib = NULL,
  pred = NULL,
  E = 2:10,
  tau = 0,
  k = E + 1,
  threads = length(E)
)

## S4 method for signature 'list'
simplex(
  data,
```

```

    column,
    target,
    lib = NULL,
    pred = NULL,
    E = 2:10,
    tau = 0,
    k = E + 1,
    threads = length(E)
  )

```

Arguments

data	observation data.
column	name of library variable.
target	name of target variable.
lib	(optional) libraries indices.
pred	(optional) predictions indices.
E	(optional) embedding dimensions.
tau	(optional) step of time lags.
k	(optional) number of nearest neighbors used in prediction.
threads	(optional) number of threads to use.

Value

A list

- xmap forecast performance
- varname name of target variable
- method method of cross mapping
- tau step of time lag

References

Sugihara G. and May R. 1990. Nonlinear forecasting as a way of distinguishing chaos from measurement error in time series. *Nature*, 344:734-741.

Examples

```

sim = logistic_map(x = 0.4, y = 0.4, step = 45, beta_xy = 0.5, beta_yx = 0)
simplex(sim, "x", "y", k = 7, threads = 1)

```

smap

smap forecast

Description

smap forecast

Usage

```
## S4 method for signature 'data.frame'
smap(
  data,
  column,
  target,
  lib = NULL,
  pred = NULL,
  E = 3,
  tau = 0,
  k = E + 1,
  theta = c(0, 1e-04, 3e-04, 0.001, 0.003, 0.01, 0.03, 0.1, 0.3, 0.5, 0.75, 1, 1.5, 2, 3,
    4, 6, 8),
  threads = length(theta)
)
```

Arguments

data	observation data.
column	name of library variable.
target	name of target variable.
lib	(optional) libraries indices.
pred	(optional) predictions indices.
E	(optional) embedding dimensions.
tau	(optional) step of time lags.
k	(optional) number of nearest neighbors used in prediction.
theta	(optional) weighting parameter for distances.
threads	(optional) number of threads to use.

Value

A list

xmap forecast performance

varname name of target variable

method method of cross mapping

References

Sugihara G. 1994. Nonlinear forecasting for the classification of natural time series. *Philosophical Transactions: Physical Sciences and Engineering*, 348 (1688):477-495.

Examples

```
sim = logistic_map(x = 0.4,y = 0.4,step = 45,beta_xy = 0.5,beta_yx = 0)
smap(sim,"x","y",E = 8,k = 7,threads = 1)
```

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