Package 'DSWE'

July 18, 2022

```
Title Data Science for Wind Energy
Version 1.6.1
Description Data science methods used in wind energy applications.
     Current functionalities include creating a multi-dimensional power curve model,
     performing power curve function comparison, covariate matching, and energy decomposition.
     Relevant works for the developed functions are:
     funGP() - Prakash et al. (2022) <doi:10.1080/00401706.2021.1905073>,
     AMK() - Lee et al. (2015) <doi:10.1080/01621459.2014.977385>,
     tempGP() - Prakash et al. (2022) <doi:10.1080/00401706.2022.2069158>,
     ComparePCurve() - Ding et al. (2021) <doi:10.1016/j.renene.2021.02.136>,
     deltaEnergy() - Latiffianti et al. (2022) <doi:10.1002/we.2722>,
     syncSize() - Latiffianti et al. (2022) <doi:10.1002/we.2722>,
     imptPower() - Latiffianti et al. (2022) <doi:10.1002/we.2722>,
     All other functions - Ding (2019, ISBN:9780429956508).
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AMK	Additive Multiplicative Kernel Regression		

Description

An additive multiplicative kernel regression based on Lee et al. (2015).

Usage

```
AMK(
   trainX,
   trainY,
   testX,
   bw = "dpi_gap",
   nMultiCov = 3,
```

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```
fixedCov = c(1, 2),
cirCov = NA
)
```

Arguments

trainX a matrix or dataframe of input variable values in the training dataset.

trainY a numeric vector for response values in the training dataset.

testX a matrix or dataframe of test input variable values to compute predictions.

bw a numeric vector or a character input for bandwidth. If character, bandwidth

computed internally; the input should be either 'dpi' or 'dpi_gap'. Default is

'dpi_gap'. See details for more information.

nMultiCov an integer or a character input specifying the number of multiplicative covariates

in each additive term. Default is 3 (same as Lee et al., 2015). The character inputs can be: 'all' for a completely multiplicative model, or 'none' for a

completely additive model. Ignored if the number of covariates is 1.

fixedCov an integer vector specifying the fixed covariates column number(s), default value

is c(1,2). Ignored if nMultiCov is set to 'all' or 'none' or if the number of

covariates is less than 3.

cirCov an integer vector specifying the circular covariates column number(s) in trainX,

default value is NA.

Details

This function is based on Lee et al. (2015). Main features are:

- Flexible number of multiplicative covariates in each additive term, which can be set using nMultiCov.
- Flexible number and columns for fixed covariates, which can be set using fixedCov. The default option c(1,2) sets the first two columns as fixed covariates in each additive term.
- Handling the data with gaps when the direct plug-in estimator used in Lee et al. fails to return a finite bandwidth. This is set using the option bw = 'dpi_gap' for bandwidth estimation.

Value

a numeric vector for predictions at the data points in testX.

References

Lee, Ding, Genton, and Xie, 2015, "Power curve estimation with multivariate environmental factors for inland and offshore wind farms," Journal of the American Statistical Association, Vol. 110, pp. 56-67, doi: 10.1080/01621459.2014.977385.

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Examples

```
data = data1
trainX = as.matrix(data[c(1:100),2])
trainY = data[c(1:100),7]
testX = as.matrix(data[c(101:110),2])
AMK_prediction = AMK(trainX, trainY, testX, bw = 'dpi_gap', cirCov = NA)
```

BayesTreePCFit

Tree based power curve estimate

Description

Tree based power curve estimate

Usage

```
BayesTreePCFit(trainX, trainY, testX, nTree = 50)
```

Arguments

trainX a matrix or dataframe to be used in modelling

trainY a numeric or vector as a target

testX a matrix or dataframe, to be used in computing the predictions

nTree a numeric value specifying number of trees to be constructed in model

Value

a vector or numeric predictions on user provided test data

```
data = data1
trainX = as.matrix(data[c(1:100),2])
trainY = data[c(1:100),7]
testX = as.matrix(data[c(100:110),2])

Bart_prediction = BayesTreePCFit(trainX, trainY, testX)
```

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ComparePCurve	Power curve	comparison
p		

Description

Power curve comparison

Usage

```
ComparePCurve(
  data,
  xCol,
  xCol.circ = NULL,
  yCol,
  testCol,
  testSet = NULL,
  thrs = 0.2,
  conflevel = 0.95,
  gridSize = c(50, 50),
  powerbins = 15,
  baseline = 1,
  limitMemory = TRUE,
  opt_method = "nlminb",
  sampleSize = list(optimSize = 500, bandSize = 5000),
  rngSeed = 1
)
```

Arguments

data	A list of data sets to be compared, the difference in the mean function is always computed as $(f(data2) - f(data1))$
xCol	A numeric or vector stating column number of covariates
xCol.circ	A numeric or vector stating column number of circular covariates
yCol	A numeric value stating the column number of the response
testCol	A numeric/vector stating column number of covariates to used in generating test set. Maximum of two columns to be used.
testSet	A matrix or dataframe consisting of test points, default value NULL, if NULL computes test points internally using testCol variables. If not NULL, total number of test points must be less than or equal to 2500.
thrs	A numeric or vector representing threshold for each covariates
conflevel	A numeric between $(0,1)$ representing the statistical significance level for constructing the band
gridSize	A numeric / vector to be used in constructing test set, should be provided when testSet is NuLL, else it is ignored. Default is c(50,50) for 2-dim input which

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is converted internally to a default of c(1000) for 1-dim input. Total number of test points (product of gridSize vector components) must be less than or equal to 2500. powerbins A numeric stating the number of power bins for computing the scaled difference, default is 15. baseline An integer between 0 to 2, where 1 indicates to use power curve of first dataset as the base for metric calculation, 2 indicates to use the power curve of second dataset as the base, and 0 indicates to use the average of both power curves as the base. Default is set to 1. A boolean (True/False) indicating whether to limit the memory use or not. DelimitMemory fault is true. If set to true, 5000 datapoints are randomly sampled from each dataset under comparison for inference opt_method A string specifying the optimization method to be used for hyperparameter estimation. Current options are: 'L-BFGS-B', 'BFGS', and 'nlminb'. Default is set to 'nlminb'. sampleSize A named list of two integer items: optimSize and bandSize, denoting the sample size for each dataset for hyperparameter optimization and confidence band computation, respectively, when limitMemory = TRUE. Default value is list(optimSize = 500, bandSize = 5000).

Value

a list containing:

rngSeed

• weightedDiff - a numeric, % difference between the functions weighted using the density of the covariates

Random seed for sampling data when limitMemory = TRUE. Default is 1.

- weightedStatDiff a numeric, % statistically significant difference between the functions weighted using the density of the covariates
- scaledDiff a numeric, % difference between the functions scaled to the orginal data
- scaledStatDiff a numeric, % statistically significant difference between the functions scaled to the original data
- unweightedDiff a numeric, % difference between the functions unweighted
- unweightedStatDiff a numeric, % statistically significant difference between the functions unweighted
- reductionRatio a list consisting of shrinkage ratio of features used in testSet
- mu1 a vector of prediction on testset using the first data set
- mu2 a vector of prediction on testset using the second data set
- muDiff a vector of the difference in prediction (mu2 mu1) for each test point
- band a vector for the confidence band at all the testpoints for the two functions to be the same at a given cofidence level.
- · confLevel a numeric representing the statistical significance level for constructing the band
- testSet a vector/matrix of the test points either provided by user, or generated internally
- estimatedParams a list of estimated hyperaparameters for the Gaussian process model
- matchedData a list of two matched datasets as generated by covariate matching

References

For details, see Ding et al. (2021) available doi: 10.1016/j.renene.2021.02.136.

Examples

```
data1 = data1[1:100, ]
data2 = data2[1:100, ]
data = list(data1, data2)
xCol = 2
xCol.circ = NULL
yCol = 7
testCol = 2
testSet = NULL
thrs = 0.2
confLevel = 0.95
gridSize = 20
function_comparison = ComparePCurve(data, xCol, xCol.circ, yCol, testCol, testSet, thrs, confLevel, gridSize)
```

ComputeWeightedDifference

Percentage weighted difference between power curves

Description

Computes percentage weighted difference between power curves based on user provided weights instead of the weights computed from the data. Please see details for more information.

Usage

```
ComputeWeightedDifference(
  muDiff,
  weights,
  base,
  statDiff = FALSE,
  confBand = NULL
)
```

Arguments

muDiff a vector of pointwise difference between two power curves on a testset as ob-

tained from ComparePCurve() or funGP() function.

weights a vector of user specified weights for each element of muDiff. It can be based

on any probability distribution of user's choice. The weights must sum to 1.

base	a vector of predictions from a power curve; to be used as the denominator in computing the percentage difference. It can be either mu1 or mu2 as obtained from ComparePCurve() or funGP() function.
statDiff	a boolean specifying whether to compute the statistical significant difference or not. Default is set to FALSE, i.e. statistical significant difference is not computed.

If set to TRUE, confBand must be provided.

confBand a vector of pointwise confidence band for all the points in the testset as obtained

from ComparePCurve() or funGP() function, named as band. Should only be

provided when statDiff is set to TRUE. Default value is NULL.

Details

The function is a modification to the percentage weighted difference defined in Ding et al. (2021). It computes a weighted difference between power curves on a testset, where the weights have to be provided by the user based on any probability distribution of their choice rather than the weights being computed from the data. The weights must sum to 1 to be valid.

Value

a numeric percentage weighted difference or statistical significant percetage weighted difference based on whether statDiff is set to FALSE or TRUE.

References

For details, see Ding et al. (2021) available at doi: 10.1016/j.renene.2021.02.136.

```
ws_test = as.matrix(seq(4.5,8.5,length.out = 10))

userweights = dweibull(ws_test, shape = 2.25, scale = 6.5)
userweights = userweights/sum(userweights)
data1 = data1[1:100, ]
data2 = data2[1:100, ]
datalist = list(data1, data2)
xCol = 2
xCol.circ = NULL
yCol = 7
testCol = 2
output = ComparePCurve(data = datalist, xCol = xCol, yCol = yCol,
testCol = testCol, testSet = ws_test)
weightedDiff = ComputeWeightedDifference(output$muDiff, userweights, output$mu1)
weightedStatDiff = ComputeWeightedDifference(output$muDiff, userweights, output$mu1,
statDiff = TRUE, confBand = output$band)
```

CovMatch 9

|--|

Description

The function aims to take list of two data sets and returns the after matched data sets using user specified covariates and threshold

Usage

```
CovMatch(data, xCol, xCol.circ, thrs, priority)
```

Arguments

data	a list, consisting of data sets to match, also each of the individual data set can be dataframe or a matrix
xCol	a vector stating the column position of covariates used
xCol.circ	a vector stating the column position of circular variables
thrs	a numerical or a vector of threshold values for each covariates, against which matching happens It should be a single value or a vector of values representing threshold for each of the covariate
priority	a boolean, default value False, otherwise computes the sequence of matching

Value

a list containing:

- originalData The data sets provided for matching
- matchedData The data sets after matching
- MinMaxOriginal The minimum and maximum value in original data for each covariate used in matching
- MinMaxMatched The minimum and maximum value in matched data for each covariates used in matching

References

Ding, Y. (2019). Data Science for Wind Energy. Chapman & Hall, Boca Raton, FL.

```
data1 = data1[1:100, ]
data2 = data2[1:100, ]
data = list(data1, data2)
xCol = 2
xCol.circ = NULL
```

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```
thrs = 0.1
priority = FALSE
matched_data = CovMatch(data, xCol, xCol.circ, thrs, priority)
```

data1

Wind Energy data set containing 47,542 data points

Description

A dataset containing the power produced and other attributes of almost 47,542 records.

Usage

```
data(data1)
```

Format

A data frame with 47,542 rows and 7 variables

Details

- Data.point sequence of integers displaying each record
- V wind speed
- D wind direction
- · air.density air density
- I turbulence intensity
- S_b wind shear
- Y wind power

data2

Wind Energy data set containing 48,068 data points

Description

A dataset containing the power produced and other attributes of almost 48,068 records.

Usage

```
data(data2)
```

Format

A data frame with 48,068 rows and 7 variables

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Details

- Data.point sequence of integers displaying each record
- V wind speed
- D wind direction
- air.density air density
- I turbulence intensity
- S_b wind shear
- Y wind power

deltaEnergy

Energy decomposition for wind turbine performance comparison

Description

Energy decomposition compares energy production from two datasets and separates it into turbine effects (deltaE.turb) and weather/environment effects (deltaE.weather).

Usage

```
deltaEnergy(
  data,
  powercol,
  timecol = 0,
  xcol,
  sync.method = "minimum power",
  imput = TRUE,
  vcol = NULL,
  vrange = NULL,
  rated.power = NULL,
  sample = TRUE,
  size = 2500,
  timestamp.min = 10
)
```

Arguments

data	A list of two data sets to be compared. A difference is always computed as (data2 - data1).	
powercol	A numeric stating the column number of power production.	
timecol	A numeric stating the column number of data time stamp. Default value is zero. A value other than zero should be provided when sync.method = 'time'.	
xcol	A numeric or vector stating the column number(s) of power curve input covariates/features (environmental or weather variables are recommended)	

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sync.method A string specifying data synchronization method. Default value 'minimum power';

other options include 'time' and 'random'.

imput A boolean (TRUE/FALSE) indicating whether power imputation should be per-

formed before calculating energy decomposition. The recommended and default value is TRUE. Change to FALSE when data have been preprocessed or imputed before.#' @param vcol A numeric stating the column number of wind speed. It

is required when imput = TRUE.

vcol A numeric stating the column number of wind speed.

vrange A vector of cut-in, rated, and cut-out wind speed. Values should be provided

when imput = TRUE.

rated.power A numerical value stating the wind turbine rated power.

sample A boolean (TRUE/FALSE) indicating whether to use sample or the whole data

sets to train the power curve to be used for power imputation. Default value is

TRUE. It is only used when imput = TRUE.

size A numeric stating the size of sample when sample = TRUE. Default value is

2500. It is only used when imput = TRUE and sample = TRUE.

timestamp.min A numerical value stating the resolution of the datasets in minutes. It is the

difference between two consecutive time stamps at which data were recorded.

Default value is 10.

Value

a list containing:

- deltaE.turb A numeric,
- deltaE.weather A numeric.
- deltaE.hat A numeric,
- deltaE.obs A numeric,
- estimated energy A numeric vector of the total energy calculated from each of f1(x2), f1(x1), f2(x2), f1(x2). If power is in kW, these values will be in kWh.
- data A list of two datasets used to calculate energy decomposition, i.e. synchronized. When imput = TRUE, the power column is the result from imputation.

References

Latiffianti, E, Ding, Y, Sheng, S, Williams, L, Morshedizadeh, M, Rodgers, M (2022). "Analysis of leading edge protection application on wind turbine performance through energy and power decomposition approaches". Wind Energy. 2022; 1-19. doi: 10.1002/we.2722.

```
data = list(data1[1:100,], data2[1:120,])
powercol = 7
timecol = 1
xcol = c(2:6)
```

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```
sync.method = 'time'
imput = TRUE
vcol = 2
vrange = c(5,12,25)
rated.power = 100
sample = FALSE
Decomposition = deltaEnergy(data, powercol, timecol, xcol, sync.method, imput, vcol, vrange, rated.power, sample)
```

funGP

Function comparison using Gaussian Process and Hypothesis testing

Description

Function comparison using Gaussian Process and Hypothesis testing

Usage

```
funGP(
  datalist,
  xCol,
  yCol,
  confLevel = 0.95,
  testset,
  limitMemory = TRUE,
  opt_method = "nlminb",
  sampleSize = list(optimSize = 500, bandSize = 5000),
  rngSeed = 1
)
```

Arguments

datalist	A list of data sets to compute a function for each of them
xCol	A numeric or vector stating the column number of covariates
yCol	A numeric value stating the column number of target
confLevel	A single value representing the statistical significance level for constructing the band
testset	Test points at which the functions will be compared
limitMemory	A boolean (True/False) indicating whether to limit the memory use or not. Default is true. If set to true, 5000 datapoints are randomly sampled from each dataset under comparison for inference.
opt_method	A string specifying the optimization method to be used for hyperparameter estimation. Current options are: 'L-BFGS-B', 'BFGS', and 'nlminb'. Default is set to 'nlminb'.

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sampleSize A named list of two integer items: optimSize and bandSize, denoting the sample size for each dataset for hyperparameter optimization and confidence band computation, respectively, when limitMemory = TRUE. Default value is list(optimSize = 500, bandSize = 5000).

rngSeed Random seed for sampling data when limitMemory = TRUE. Default is 1.

Value

a list containing:

- muDiff A vector of pointwise difference between the predictions from the two datasets (mu2-mu1)
- mu1 A vector of test prediction for first data set
- mu2 A vector of test prediction for second data set
- band A vector of the allowed statistical difference between functions at testpoints in testset
- confLevel A numeric representing the statistical significance level for constructing the band
- testset A matrix of test points to compare the functions
- estimatedParams A list of estimated hyperparameters for GP

References

Prakash, A., Tuo, R., & Ding, Y. (2022). "Gaussian process aided function comparison using noisy scattered data," Technometrics, Vol. 64, No. 1, pp. 92-102, doi: 10.1080/00401706.2021.1905073.

Examples

```
datalist = list(data1[1:100,], data2[1:100, ])
xCol = 2
yCol = 7
confLevel = 0.95
testset = seq(4,10,length.out = 20)
function_diff = funGP(datalist, xCol, yCol, confLevel, testset)
```

imptPower

Power imputation

Description

Good power curve modeling requires valid power values in the region between cut-in and cut-out wind speed. However, when turbine is not operating, the power production will be recorded as zero or negative. This function replaces those values with predicted values obtained from the estimated tempGP power curve model using one input variable - the wind speed.

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Usage

```
imptPower(
  data,
  powercol,
  vcol,
  vrange,
  rated.power = NULL,
  sample = TRUE,
  size = 2500
)
```

Arguments

data A list of two data sets that require imputation.

powercol A numeric stating the column number of power production.

vcol A numeric stating the column number of wind speed.
vrange A vector of cut-in, rated, and cut-out wind speed.

rated.power A numerical value stating the wind turbine rated power.

sample A boolean (TRUE/FALSE) indicating whether to use sample or the whole data

sets to train the power curve.

size A numeric stating the size of sample when sample = TRUE. Default value is

2500. It is only used when sample = TRUE.

Value

a list containing datasets with the imputed power.

References

Latiffianti, E, Ding, Y, Sheng, S, Williams, L, Morshedizadeh, M, Rodgers, M (2022). "Analysis of leading edge protection application on wind turbine performance through energy and power decomposition approaches". Wind Energy. 2022; 1-19. doi: 10.1002/we.2722.

```
data = list(data1[1:100,], data2[1:120, ])
powercol = 7
vcol = 2
vrange = c(5,12,25)
rated.power = 100
sample = FALSE
imputed.dat = imptPower(data, powercol, vcol, vrange, rated.power, sample)
```

16 KnnPCFit

KnnPCFit KNN: Fit

Description

The function models the powercurve using KNN, against supplied arguments

Usage

```
KnnPCFit(data, xCol, yCol, subsetSelection = FALSE)
```

Arguments

data a dataframe or a matrix, to be used in modelling

xCol a vector or numeric values stating the column number of features

yCol a numerical or a vector value stating the column number of target

subsetSelection

a boolean, default value is FALSE, if TRUE returns the best feature column

a boolean, default value is FALSE, if TRUE returns the best feature column number as xCol

Value

a list containing:

- data The data set provided by user
- xCol The column number of features provided by user or the best subset column number
- yCol The column number of target provided by user
- bestK The best k nearest neighbor calculated using the function
- RMSE The RMSE calculated using the function for provided data using user defined features and best obtained K
- MAE The MAE calculated using the function for provided data using user defined features and best obtained K

```
data = data1[c(1:100),]
xCol = 2
yCol = 7
subsetSelection = FALSE
knn_model = KnnPCFit(data, xCol, yCol, subsetSelection)
```

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KnnPredict KNN: Predict

Description

The function can be used to make prediction on test data using trained model

Usage

```
KnnPredict(knnMdl, testData)
```

Arguments

knnMdl

a list containing:

- knnMdl\$data The data set provided by user
- knnMdl\$xCol The column number of features provided by user or the best subset column number
- knnMdl\$yCol The column number of target provided by user
- knn\$bestK The best k nearest neighbor calculated using the function KnnFit

testData

a data frame or matrix, to compute the predictions

Value

a numeric / vector with prediction on test data using model generated by KnnFit

```
data = data1[c(1:100),]
xCol = 2
yCol = 7
subsetSelection = FALSE
knn_model = KnnPCFit(data, xCol, yCol, subsetSelection)
testData = data1[c(101:110), ]
prediction = KnnPredict(knn_model, testData)
```

18 KnnUpdate

KnnUpdate

KNN: Update

Description

The function can be used to update KNN model when new data is provided

Usage

```
KnnUpdate(knnMdl, newData)
```

Arguments

knnMd1

a list containing:

- knnMdl\$data The data set provided by user
- knnMdl\$xCol The column number of features provided by user or the best subset column number
- knnMdl\$yCol The column number of target provided by user
- knn\$bestK The best k nearest neighbor calculated using the function KnnFit

newData

a dataframe or a matrix, to be used for updating the model

Value

a list containing:

- data The updated data using old data set and new data
- xCol The column number of features provided by user or the best subset column number
- yCol The column number of target provided by user
- bestK The best k nearest neighbor calculated for the new data using user specified features and target

```
data = data1[c(1:100),]
xCol = 2
yCol = 7
subsetSelection = FALSE
knn_model = KnnPCFit(data, xCol, yCol, subsetSelection)
newData = data1[c(101:110), ]
knn_newmodel = KnnUpdate(knn_model, newData)
```

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|--|

Description

predict function for tempGP objects. This function computes the prediction f(x) or f(x) + g(t) depending on the temporal distance between training and test points and whether the time indices for the test points are provided.

Usage

```
## S3 method for class 'tempGP'
predict(object, testX, testT = NULL, trainT = NULL, ...)
```

Arguments

object	An object of class tempGP.
testX	A matrix with each column corresponding to one input variable.
testT	A vector of time indices of the test points. When NULL, only function $f(x)$ is used for prediction. Default is NULL.
trainT	Optional argument to override the existing trainT indices of the tempGP object.
	additional arguments for future development

Value

A vector of predictions at the testpoints in testX.

```
data = DSWE::data1
trainindex = 1:100 #using the first 100 data points to train the model
traindata = data[trainindex,]
xCol = 2 #input variable columns
yCol = 7 #response column
trainX = as.matrix(traindata[,xCol])
trainY = as.numeric(traindata[,yCol])
tempGPObject = tempGP(trainX, trainY)
testdata = DSWE::data1[101:110,] # defining test data
testX = as.matrix(testdata[,xCol, drop = FALSE])
predF = predict(tempGPObject, testX)
```

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SplinePCFit	Smoothing spline Anova method
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Description

Smoothing spline Anova method

Usage

```
SplinePCFit(data, xCol, yCol, testX, modelFormula = NULL)
```

Arguments

data	a matrix or dataframe to be used in modelling
xCol	a numeric or vector stating the column number of feature covariates
yCol	a numeric value stating the column number of target
testX	a matrix or dataframe, to be used in computing the predictions
modelFormula	default is NULL else a model formula specifying target and features. Please refer 'gss' package documentation for more details

Value

a vector or numeric predictions on user provided test data

Examples

```
data = data1[c(1:100),]
xCol = 2
yCol = 7
testX = data1[c(101:110), ]
Spline_prediction = SplinePCFit(data, xCol, yCol, testX)
```

SvmPCFit

SVM based power curve modelling

Description

SVM based power curve modelling

Usage

```
SvmPCFit(trainX, trainY, testX, kernel = "radial")
```

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Arguments

trainX a matrix or dataframe to be used in modelling

trainY a numeric or vector as a target

testX a matrix or dataframe, to be used in computing the predictions kernel default is 'radial' else can be 'linear', 'polynomial' and 'sigmoid'

Value

a vector or numeric predictions on user provided test data

Examples

```
data = data1
trainX = as.matrix(data[c(1:100),2])
trainY = data[c(1:100),7]
testX = as.matrix(data[c(101:110),2])

Svm_prediction = SvmPCFit(trainX, trainY, testX)
```

syncSize Data synchronization

Description

Data synchronization is meant to make a pair of data to have the same size. It is performed by removing some data points from the larger dataset. This step is important when comparing energy production between two data sets because energy production is time-based.

Usage

```
syncSize(data, powercol, timecol = 0, xcol, method = "minimum power")
```

Arguments

data	A list of two data sets to be synchronized.
powercol	A numeric stating the column number of power production.
timecol	A numeric stating the column number of data time stamp. Default value is zero. A value other than zero should be provided when method = 'time'.
xcol	A numeric or vector stating the column number(s) of power curve input covariates/features (to be used for energy decomposition).
method	A string specifying data synchronization method. Default value 'minimum power other options include 'time' and 'random'.

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Value

a list containing the synchronized datasets.

References

Latiffianti, E, Ding, Y, Sheng, S, Williams, L, Morshedizadeh, M, Rodgers, M (2022). "Analysis of leading edge protection application on wind turbine performance through energy and power decomposition approaches". Wind Energy. 2022; 1-19. doi: 10.1002/we.2722.

Examples

```
data = list(data1[1:200,], data2[1:180, ])
powercol = 7
timecol = 1
xcol = c(2:6)
method = 'random'
sync.dat = syncSize(data, powercol, timecol, xcol, method)

data = list(data1[500:700,], data2[600:750, ])
powercol = 7
timecol = 1
xcol = c(2:6)
method = 'time'
sync.dat = syncSize(data, powercol, timecol, xcol, method)
```

tempGP

temporal Gaussian process

Description

A Gaussian process based power curve model which explicitly models the temporal aspect of the power curve. The model consists of two parts: f(x) and g(t).

Usage

```
tempGP(
   trainX,
   trainY,
   trainT = NULL,
   fast_computation = TRUE,
   limit_memory = 5000L,
   optim_control = list(batch_size = 100L, learn_rate = 0.05, max_iter = 5000L, tol = 1e-06, beta1 = 0.9, beta2 = 0.999, epsilon = 1e-08, logfile = NULL)
)
```

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Arguments

trainX A matrix with each column corresponding to one input variable.

trainY A vector with each element corresponding to the output at the corresponding

row of trainX.

trainT A vector for time indices of the data points. By default, the function assigns

natural numbers starting from 1 as the time indices.

fast_computation

A Boolean that specifies whether to do exact inference or fast approximation.

Default is TRUE.

limit_memory An integer or NULL. The integer is used sample training points during prediction

to limit the total memory requirement. Setting the value to NULL would result in no sampling, that is, full training data is used for prediction. Default value is

5000.

optim_control A list parameters passed to the Adam optimizer when fast_computation is set to TRUE. The default values have been tested rigorously and tend to strike a

balance between accuracy and speed.

• batch_size: Number of training points sampled at each iteration of Adam.

• learn_rate: The step size for the Adam optimizer.

• max_iter: The maximum number of iterations to be performed by Adam.

• tol: Gradient tolerance.

• beta1: Decay rate for the first moment of the gradient.

• beta2: Decay rate for the second moment of the gradient.

• epsilon: A small number to avoid division by zero.

 logfile: A string specifying a file name to store hyperparameters value for each iteration.

Value

An object of class tempGP with the following attributes:

- trainX same as the input matrix trainX.
- trainY same as the input vector trainY.
- thinningNumber the thinning number computed by the algorithm.
- model F A list containing the details of the model for predicting function f(x):
 - X The input variable matrix for computing the cross-covariance for predictions, same as trainX unless the model is updated. See updateData.tempGP method for details on updating the model.
 - y The response vector, again same as trainY unless the model is updated.
 - weightedY The weighted response, that is, the response left multiplied by the inverse of the covariance matrix.
- modelG A list containing the details of the model for predicting function g(t):
 - residuals The residuals after subtracting function f(x) from the response. Used to predict g(t). See updateData.tempGP method for updating the residuals.

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- time_index The time indices of the residuals, same as trainT.
- estimated Params Estimated hyperparameters for function f(x).
- llval log-likelihood value of the hyperparameter optimization for f(x).
- gradval gradient vector at the optimal log-likelihood value.

References

Prakash, A., Tuo, R., & Ding, Y. (2022). "The temporal overfitting problem with applications in wind power curve modeling." Technometrics. doi: 10.1080/00401706.2022.2069158.

See Also

predict.tempGP for computing predictions and updateData.tempGP for updating data in a tempGP object.

Examples

```
data = DSWE::data1
trainindex = 1:100 #using the first 100 data points to train the model
traindata = data[trainindex,]
xCol = 2 #input variable columns
yCol = 7 #response column
trainX = as.matrix(traindata[,xCol])
trainY = as.numeric(traindata[,yCol])
tempGPObject = tempGP(trainX, trainY)
```

updateData

Updating data in a model

Description

updateData is a generic function to update data in a model.

Usage

```
updateData(object, ...)
```

Arguments

object A model object

... additional arguments for passing to specific methods

Value

The returned value would depend on the class of its argument object.

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See Also

```
updateData.tempGP
```

updateData.tempGP

Update the data in a tempGP object

Description

This function updates trainX, trainY, and trainT in a tempGP object. By default, if the new data has m data points, the function removes top m data points from the tempGP object and appends the new data at the bottom, thus keeping the total number of data points the same. This can be overwritten by setting replace = FALSE to keep all the data points (old and new). The method also updates modelG by computing and updating residuals at the new data points. modelF can be also be updated by setting the argument updateModelF to TRUE, though not required generally (see comments in the Arguments.)

Usage

```
## $3 method for class 'tempGP'
updateData(
  object,
  newX,
  newY,
  newT = NULL,
  replace = TRUE,
  updateModelF = FALSE,
   ...
)
```

Arguments

object	An object of class tempGP.

newX A matrix with each column corresponding to one input variable.

newY A vector with each element corresponding to the output at the corresponding

row of newX.

newT A vector with time indices of the new datapoints. If NULL, the function assigns

natural numbers starting with one larger than the existing time indices in trainT.

replace A boolean to specify whether to replace the old data with the new one, or to

add the new data while still keeping all the old data. Default is TRUE, which replaces the top m rows from the old data, where m is the number of data points

in the new data.

updateModelF A boolean to specify whether to update modelF as well. If the original tempGP

model is trained on a sufficiently large dataset (say one year), updating modelF regularly may not result in any significant improvement, but can be computa-

tionally expensive.

. . . additional arguments for future development

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Value

An updated object of class tempGP.

```
data = DSWE::data1
trainindex = 1:100 #using the first 100 data points to train the model
traindata = data[trainindex,]
xCol = 2 #input variable columns
yCol = 7 #response column
trainX = as.matrix(traindata[,xCol])
trainY = as.numeric(traindata[,yCol])
tempGPObject = tempGP(trainX, trainY)
newdata = DSWE::data1[101:110,] # defining new data
newX = as.matrix(newdata[,xCol, drop = FALSE])
newY = as.numeric(newdata[,yCol])
tempGPupdated = updateData(tempGPObject, newX, newY)
```

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