

# Package ‘CommonSplines’

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**Title** Regression Spline and Smoothing Spline

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**URL** <https://github.com/YuchenKid/CommonSplines>

**Description** This is an R package that covers commonly seen nonparametric regression using spline-based methods. For regression spline, commonly seen basis functions are provided such as truncated power basis, natural cubic spline basis, and B-spline basis. For regularization, penalties on squared second-order derivative are provided, i.e., cubic smoothing spline. This package mainly refer to “Friedman, J., Hastie, T., & Tibshirani, R. (2001). The elements of statistical learning (Vol. 1, pp. 337-387). New York: Springer series in statistics,” Chapter 5.

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**LazyData** true

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**Suggests** knitr,  
rmarkdown

**VignetteBuilder** knitr

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---

bs_basis	<i>Generate an evaluated basis matrix for B-splines</i>
----------	---

---

## Description

#' This function generates B-spline basis. The B-splines are defined following the recursive formulas due to de Boor. Only univariate input can be used.

## Usage

```
bs_basis(x, order, knots)
```

## Arguments

x	Predictor variable vector.
order	The order of basis functions. order=degree+1
knots	The knots used to construct the B-splines, including innerknots, boundary knots and phantom knots. It can be generated by bs_knots.

## Value

Basis matrix evaluated at each x value.

## Examples

```
x<-seq(0, 1, 0.001)
knots <- seq(0, 1, 0.1)
order<-4
knots<-bs_knots(x,real_knots=knots,order=order)

basis<-bs_basis(x,order,knots)
plot(x,rep(0,length(x)),type="l",ylim=c(0,1))
for (i in 1: (length(knots)-order)){
  lines(x,basis[,i])
}
```

---

bs_knots	<i>Add phantom knots for B-splines</i>
----------	--

---

**Description**

Add phantom knots for B-splines

**Usage**

```
bs_knots(x, df = NULL, real_knots = NULL, q = FALSE, order)
```

**Arguments**

x	Predictor variable vector.
df	Degrees of freedom. One can supply df rather than knots.
real_knots	The innerknots and boundary knots that define the spline. The knots can all be innerknots. The knots provided can be quantiles of x or real values. More explanation of knots, df, q can be seen in generate_knots.
q	A boolean variable define whether knots provided are quantiles or real values. When q=TRUE, real_knots are quantiles of x. When q=FALSE, real_knots are real values of x. Default is FALSE.
order	The order of basis functions. order=degree+1

**Value**

The knots used to construct the B-splines, including innerknots, boundary knots and phantom knots.

---

bs_predict	<i>Prediction using regression spline with B-spline basis</i>
------------	---

---

**Description**

This function provides prediction at value of interest using regression spline with B-spline basis. The B-splines are generated by bs\_basis and trained by the bs\_train. The return value of bs\_train can be used as an argument of bs\_predict

**Usage**

```
bs_predict(x_test, order = NULL, knots = NULL, beta = NULL,
           basis = NULL)
```

**Arguments**

x_test	The input values at which evaluations are required.
order	The order of basis functions. order=degree+1
knots	Breakpoints that define the spline. knots should be in terms of real-values of x and contain inner, boundary and phantom knots. It can be the return value of bs_knots.
beta	The coefficients of nonparametric regression.
basis	The return value of function bs_train. Instead of specify knots, order and beta, One can supply basis directly.

**Value**

The evaluated output at `x_test`.

**See Also**

`bs_basis`, `bs_train`, `bs_knots`.

**Examples**

```
x<-seq(0, 1, 0.001)
y <- x^3 * 3 - x^2 * 2 + x + exp(1)+rnorm(length(x),0,0.1)
plot(x,y)
knots <- seq(0.1, 0.9, 0.01)
order<-4
basis<-bs_train(x,y,order,knots)

x_test<-seq(0, 1, 0.01)
fit<-bs_predict(x_test,basis=basis)
plot(x_test,fit)
lines(x_test,x_test^3 * 3 - x_test^2 * 2 + x_test + exp(1),col="red")
```

---

<code>bs_train</code>	<i>Train regression coefficients for B-splines.</i>
-----------------------	---

---

**Description**

Train regression coefficients for B-splines.

**Usage**

```
bs_train(x, y, order, real_knots = NULL, df = NULL, q = FALSE)
```

**Arguments**

<code>x</code>	The input vector of training dataset.
<code>y</code>	The output vector of training dataset.
<code>order</code>	The order of B-spline functions. The default is <code>order=4</code> for cubic B-splines.
<code>real_knots</code>	The innerknots and boundary knots that define the spline. Phantom knots should not be included. Phantom knots will be generated by <code>bs_knots</code> . The knots provided can be quantiles of <code>x</code> or real values. More explanation of knots, <code>df</code> , <code>q</code> can be seen in <code>generate_knots</code> .
<code>df</code>	Degrees of freedom. One can supply <code>df</code> rather than knots.
<code>q</code>	A boolean variable define whether knots provided are quantiles or real values. When <code>q=TRUE</code> , <code>real_knots</code> are quantiles of <code>x</code> . When <code>q=FALSE</code> , <code>real_knots</code> are real values of <code>x</code> . Default is <code>FALSE</code> .

**Value**

A list with the following components:

beta	The coefficients of nonparametric regression.
basis	The B-spline basis matrix of dimension $c(\text{length}(x), \text{df})$ . $\text{df} = \text{length}(\text{innerknots}) + \text{order}$ .
knots	The knots used to construct the B-splines, including innerknots, boundary knots and phantom knots
order	The order of basis functions. $\text{order} = \text{degree} + 1$

**See Also**

generate\_knots, bs\_knots.

**Examples**

```
x<-seq(0, 1, 0.001)
y <- x^3 * 3 - x^2 * 2 + x + exp(1)+rnorm(length(x),0,0.1)
plot(x,y)
knots <- seq(0, 1, 0.1)
order<-4

basis<-bs_train(x,y,order,knots)
plot(x,rep(0,length(x)),type="l",ylim=c(0,1))
for (i in 1: (length(knots)+order)){
  lines(x,basis$basismatrix[,i])
}
```

---

cal\_loo\_cv\_error

---

*Calculate leave-one-out CV error*


---

**Description**

Calculate leave-one-out CV error

Calculate leave-one-out CV error

**Usage**

```
cal_loo_cv_error(y, f_hat, S)
```

```
cal_loo_cv_error(y, f_hat, S)
```

**Arguments**

y	response variable values
f_hat	fitted response variable values
S	smoother matrix
y	response variable values
f_hat	fitted response variable values
S	smoother matrix

**Value**

leave-one-out cross-validation error

leave-one-out cross-validation error

---

css_predict	<i>Prediction using smoothing spline with squared 2nd derivative penalty</i>
-------------	--

---

**Description**

This function takes the coefficients trained by CubicSmoothingSpline.Train and evaluate the output at x\_test

**Usage**

```
css_predict(x_test, knots = NULL, beta = NULL, basis = NULL)
```

**Arguments**

x_test	The input values at which evaluations are required.
knots	Breakpoints that define the spline. knots should be in terms of real-values of x It can be the return value of generate_knots.
beta	The coefficients of nonparametric regression.
basis	The return value of function css_train. Instead of specify knots and beta,One can supply basis directly.

**Value**

The evaluated output at x\_test.

**Examples**

```
x<-seq(0, 1, 0.0015)
y <- x^3 * 3 - x^2 * 2 + x + exp(1)+rnorm(length(x),0,0.1)
plot(x,y)
lambda<-0.001
basis<-css_train(x,y,lambda)

x_test<-seq(0, 1, 0.1)
fit<-css_predict(x_test=x_test,basis=basis)

plot(x_test,fit)
lines(x_test,x_test^3 * 3 - x_test^2 * 2 + x_test + exp(1),col="red")
```

---

css_train	<i>Train a smoothing spline with squared 2nd derivative penalty using natural cubic spline</i>
-----------	--

---

## Description

This function trains a smoothing spline with squared 2nd derivative penalty. It has an explicit, finite-dimensional, unique minimizer which is a natural cubic spline.

## Usage

```
css_train(x, y, lambda)
```

## Arguments

x	The input vector of training dataset.
y	The output vector of training dataset.
lambda	A fixed smoothing parameter.

## Value

A list with the following components:

beta	The coefficients of natural splines.
S	The smoother matrix.
knots	The knots used to construct the B-splines, including innerknots, boundary knots and phantom knots

## References

"Friedman, J., Hastie, T., & Tibshirani, R. (2001). The elements of statistical learning (Vol. 1, pp. 337-387). New York: Springer series in statistics," Chapter 5.4.

## Examples

```
x<-seq(0, 1, 0.001)
y <- x^3 * 3 - x^2 * 2 + x + exp(1)+rnorm(length(x),0,0.1)
plot(x,y)
lambda<-0.001

basis<-css_train(x,y,lambda)
cat("the knots chosen are: ",basis$knots)
```

---

generate_knots	<i>Generate knots when real value is not specified.</i>
----------------	---

---

### Description

Generate knots when real value is not specified.

### Usage

```
generate_knots(x_train, df = NULL, knots = NULL, q = FALSE)
```

### Arguments

x_train	The input vector of training dataset.
df	Degrees of freedom. One can supply df rather than knots; generate_knots then chooses (df + 1) knots at uniform quantiles of x. The default, df = 4, sets 5 knots with 3 inner knots at uniform quantiles of x.
knots	Breakpoints that define the spline, in terms of quantiles or real values of x. The default is five knots at uniform quantiles c(0, .25, .5, .75, 1). Typical values are the mean or median for one knot, quantiles for more knots.
q	A boolean variable define whether knots provided are quantiles or real values. When q=TRUE, knots provided are quantiles of x. When q=FALSE, knots provided are real values of x.

### Value

A vector of knots in terms of real values of x.

---

ncs_basis	<i>Generate an evaluated basis matrix for natural cubic splines</i>
-----------	---

---

### Description

Generate an evaluated basis matrix for natural cubic splines

### Usage

```
ncs_basis(x, knots)
```

### Arguments

x	Predictor variable vector.
knots	Knots location in terms of real values of x.

### Value

Basis matrix evaluated at each x value.



## Examples

```
x<-seq(0, 1, 0.001)
knots <- seq(0, 1, 0.1)

basis<-ncs_basis(x,knots)
plot(x,rep(0,length(x)),type="l",ylim=c(0,1))
for (i in 1: (length(knots))){
  lines(x,basis[,i])
}
```

---

ncs_predict	<i>Prediction using regression spline with natural cubic spline.</i>
-------------	--

---

## Description

Prediction using regression spline with natural cubic spline.

## Usage

```
ncs_predict(x_test, knots = NULL, beta = NULL, basis = NULL)
```

## Arguments

x_test	The input values at which evaluations are required.
knots	Breakpoints that define the spline. knots should be in terms of real-values of x It can be the return value of generate_knots.
beta	The coefficients of nonparametric regression.
basis	The return value of function ncs_train. Instead of specify knots and beta,One can supply basis directly.

## Value

y_pred	A vector of dimension length(x), the prediction vector evaluated at x_test values.
--------	--

---

ncs_train	<i>Train regression coefficients for natural cubic splines.</i>
-----------	---

---

## Description

Train regression coefficients for natural cubic splines.

## Usage

```
ncs_train(x_train, y_train, df = NULL, knots = NULL, q = FALSE)
```

**Arguments**

x_train	The input vector of training dataset.
y_train	The output vector of training dataset.
df	Degrees of freedom. One can supply df rather than knots; ncs() then chooses (df + 1) knots at uniform quantiles of x. The default, df = 4, sets 5 knots with 3 inner knots at uniform quantiles of x.
knots	Breakpoints that define the spline, in terms of quantiles of x or real values of x. The default is five knots at uniform quantiles c(0, .25, .5, .75, 1). Typical values are the mean or median for one knot, quantiles for more knots.
q	A boolean variable define whether knots provided are quantiles or real values. When q=TRUE, knots provided are quantiles of x. When q=FALSE, knots provided are real values of x. Default is FALSE.

**Value**

A list of following components:

nknots	Number of knots.
knots	A vector of knot locations.
N	Basis matrix evaluated at each x value.
betas	Least square fit parameters.

**Examples**

```
x_train <- seq(1, 10, 0.1)
y_train <- cos(x_train)^3 * 3 - sin(x_train)^2 * 2 + x_train + exp(1)+rnorm(length(x_train),0,1)
plot(x_train,y_train)
x_test <- seq(1, 10, 0.1)
df <- 10
train_result <- ncs_train(x_train, y_train, df)
print(train_result$betas)
print(train_result$N[1:5,1:5])
```

---

np\_reg

---

*Nonparametric Regression using spline based methods*


---

**Description**

This function provides regression using spline based methods. It finish both training procedure and predicting procedure. Only univariate input can be used.

**Usage**

```
np_reg(x_train, y_train, x_test, func = "bs", order = 4, df = NULL,
       knots = NULL, lambda = 0.001, q = FALSE)
```

**Arguments**

<code>x_train</code>	The input vector of training dataset.
<code>y_train</code>	The output vector of training dataset.
<code>x_test</code>	The input values at which evaluations are required.
<code>func</code>	The name of regression functions. It can be "pbs" for power basis spline, "ncs" for natural cubic spline, "css" for cubic smoothing spline, "bs" for B-spline. Default is "bs".
<code>order</code>	The order that defines the spline. Default is 4.
<code>df</code>	Degrees of freedom. One can supply df rather than knots.
<code>knots</code>	The innerknots and boundary knots that define the spline. The knots provided can be quantiles of x or real values. More explanation of knots, df, q can be seen in <code>generate_knots</code> .
<code>lambda</code>	The smoothing parameter for css. Default is 0.001.
<code>q</code>	A boolean variable define whether knots provided are quantiles or real values. When q=TRUE, knots provided are quantiles of x. When q=FALSE, knots provided are real values of x. Default is FALSE.

**Value**

<code>y_pred</code>	A vector of dimension <code>length(x)</code> , the prediction vector evaluated at <code>x_test</code> values.
---------------------	---

**See Also**

`generate_knots`.

**Examples**

```
x_train <- seq(1, 10, 0.1)
y_train <- cos(x_train)^3 * 3 - sin(x_train)^2 * 2 + x_train + exp(1)+rnorm(length(x_train),0,1)
plot(x_train,y_train)
title('Comparison of Different Degrees of Freedom')
x_test <- seq(1, 10, 0.1)
lines(x_test,cos(x_test)^3 * 3 - sin(x_test)^2 * 2 + x_test + exp(1),col="red")

df <- 2
y_pred <- np_reg(x_train, y_train, x_test,func="ncs", df=df)
lines(x_test,y_pred, col='blue')
df <- 4
y_pred <- np_reg(x_train, y_train, x_test,func="ncs", df=df)
lines(x_test,y_pred, col='green')
df <- 10
y_pred <- np_reg(x_train, y_train, x_test,func="ncs", df=df)
lines(x_test,y_pred, col='black')
legends <- c("Actual", "Prediction: 2 df", "Prediction: 4 df", "Prediction: 10 df")
legend('topleft', legend=legends, col=c('red', 'blue', 'green', 'black'), lty=1, cex=0.8)
```

---

pbs_basis	<i>Evaluate basis functions as each x and return the evaluated basis matrix N</i>
-----------	---

---

### Description

Evaluate basis functions as each x and return the evaluated basis matrix N

### Usage

```
pbs_basis(x, order, knots)
```

### Arguments

x	Predictor variable vector.
order	The order that defines the power basis spline.
knots	The innerknots and boundary knots that define the spline. The knots should be real values of x. The knots can be generated by generate_knots.

### Value

Basis matrix evaluated at each x value.

### See Also

generate\_knots.

### Examples

```
x<-seq(0, 1, 0.001)
knots <- seq(0, 1, 0.1)
order<-4

basis<-pbs_basis(x,order,knots)
plot(x,rep(0,length(x)),type="l",ylim=c(0,1))
for (i in 1: (length(knots)+order)){
  lines(x,basis[,i])
}
```

---

pbs_predict	<i>Prediction using regression spline with truncated power basis</i>
-------------	--

---

### Description

This function provides prediction at value of interest using regression spline with truncated power basis. The truncated power basis are generated by pbs\_basis and trained by the pbs\_train. The return value of pbs\_train can be used as an argument of pbs\_predict

**Usage**

```
pbs_predict(x_test, order = NULL, knots = NULL, beta = NULL,
            basis = NULL)
```

**Arguments**

x_test	The input values at which evaluations are required.
order	The order of basis functions. order=degree+1
knots	Breakpoints that define the spline, in terms of real values of input. It can be the return value of generate_knots.
beta	The coefficients of nonparametric regression.
basis	The return value of function pbs_train. Instead of specify knots, order and beta, One can supply basis directly.

**Value**

The evaluated output at x\_test.

**See Also**

pbs\_basis, pbs\_train, generate\_knots.

**Examples**

```
n <- 100
t <- seq(0, 2*pi, length.out = 100)
a <- 3
b <- 2
c.unif <- runif(n)
amp <- 2
set.seed(1)
y1 <- a*sin(b*t) + c.unif*amp # uniform error
knots <- c(min(t), 2*pi*c(1/4, 2/4, 3/4), max(t))
order <- 4
basis <- pbs_train(t, y1, order, knots=knots)
fit <- pbs_predict(t, basis=basis)
y.hat <- fit
plot(t, y1, t="l")
lines(t, y.hat, col=2)
```

---

pbs\_train

*Regression using Power Basis spline*

---

**Description**

This function provides regressions using Power Basis splines. The basis are defined as  $1, x, x^2, \dots, x^m, (x-k_1)^{+(m-1)}, (x-k_2)^{+(m-1)}, \dots, (x-k_n)^{+(m-1)}$  where  $m$  is the order,  $k_1, k_2$  and  $k_n$  are  $n$  knots, '+' denotes the positive part.

**Usage**

```
pbs_train(x, y, order, df = NULL, knots = NULL, q = FALSE)
```

**Arguments**

x	The input vector of training dataset.
y	The output vector of training dataset.
order	The order that defines the spline.
df	Degrees of freedom. One can supply df rather than knots.
knots	The innerknots and boundary knots that define the spline. The knots provided can be quantiles of x or real values. More explanation of knots, df, q can be seen in generate_knots.
q	A boolean variable define whether knots provided are quantiles or real values. When q=TRUE, knots provided are quantiles of x. When q=FALSE, knots provided are real values of x. Default is FALSE.
x_test	The input values at which evaluations are required.

**Details**

Only univariate input can be used.

**Value**

A list with the following components:

beta	The coefficients of nonparametric regression.
basis	The spline basis matrix of dimension $c(\text{length}(x), \text{length}(\text{knots}) + \text{order})$
knots	The knots used to construct the power basis splines
order	The order of basis functions. $\text{order} = \text{degree} + 1$

**See Also**

generate\_knots.

**Examples**

```
n <- 100
t <- seq(0, 2*pi, length.out = 100)
a <- 3
b <- 2
c.unif <- runif(n)
amp <- 2
set.seed(1)
y1 <- a*sin(b*t) + c.unif*amp # uniform error
knots <- c(min(t), 2*pi*c(1/4, 2/4, 3/4), max(t))
order <- 4
basis <- pbs_train(t, y1, order, knots=knots)
cat("trained coefficients for every spline are", basis$beta)
```

---

place_knots	<i>Find evenly spaced knots by quantile</i>
-------------	---

---

**Description**

Knots found include boundary knots at 0th and 100th quantile.

**Usage**

```
place_knots(nknots, x)
```

**Arguments**

nknots	Number of knots to be located.
x	Data vector on which knots are placed.

**Value**

A named vector with knot quantiles and values.

---

sel_smoothing_para	<i>Select smoothing parameter based on leave-one-out CV error</i>
--------------------	---

---

**Description**

Select smoothing parameter based on leave-one-out CV error  
 Select smoothing parameter based on leave-one-out CV error

**Usage**

```
sel_smoothing_para(x, y, cv_lambda)

sel_smoothing_para(x, y, cv_lambda)
```

**Arguments**

x	predictor variable
y	response variable
cv_lambda	vector of candidate lambda values
x	predictor variable.
y	response variable.
cv_lambda	vector of candidate lambda values, must be between 0 and 1.

**Value**

lambda value that minimizes leave-one-out CV error  
 lambda value that minimizes leave-one-out CV error.

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