

# Package ‘lrspline’

May 30, 2021

**Type** Package

**Title** Low-rank Approximation of Smoothing Splines for Massive Data

**Version** 0.1.0

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**Description** Functions for approximating the smoothing spline estimates: (1) Low-rank Approximation via Eigensystem Truncation, (2) Nystrom Methods.

**License** GPL-3

**Encoding** UTF-8

**LazyData** true

**Imports** assist,  
mgcv

**RoxygenNote** 7.1.1

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eigenM	<i>An example of grid points and eigendecomposition for reprodcng kernel of cubic spline.</i>
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## Description

It contains e a list of eigenvalues and eigenfunctions at grid points, xg a vector of 1000 evenly spaced points.

## Usage

```
data(eigenM)
```

**Format**

An object of class list of length 2.

**Examples**

```
data(eigenM)
## Not run:
# check the number of eigenvalues
length(eigenM$e$values)

## End(Not run)
```

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generate.eigen.cubic	<i>Grid Points and Eigendecomposition of Reproducing Kernal for Cubic Splines</i>
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**Description**

Generates eigendecomposition from grid points from an interval.

**Usage**

```
generate.eigen.cubic(N = 1000, a = 0, b = 1)
```

**Arguments**

N	The number of grid points. It should be an integer. The default value is 1000.
a	The lower limit of the interval used for grid points. The default value is 0.
b	The upper limit of the interval used for grid points. The default value is 1.

**Value**

It returns (and saves) a list of following components:

e	A list of two elements of "values" and "vectors", which refer respectively, the eigenvalues and eigenfunctions of reproducing kernel for cubic splines at the pre-selected grid points.
xg	A vector of grid points.

**Examples**

```
## Not run:
eigen_res <- generate.eigen.cubic(N=1000,a=2,b=3)

## End(Not run)
```

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hello	<i>Hello, World!</i>
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**Description**

Prints 'Hello, world!'.

**Usage**

```
hello()
```

**Examples**

```
hello()
```

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lrspline.cubic	<i>Low-rank Approximation Based on Eigenspaces for Cubic Smoothing Spline Estimates</i>
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**Description**

Computes a low-rank approximation based on eigenspaces, where estimation utilizes functions for linear mixed effect model (LME).

**Usage**

```
lrspline.cubic(x, y, xg, e, K = 30, method = "REML", pstd = FALSE)
```

**Arguments**

x	The values of independent variable. It should be a vector.
y	The values of dependent variable. It should be a vector.
xg	The grid points used for approximation of eigensystem.
e	A list of two elements of "values" and "vectors", which refer respectively, the eigenvalues and eigenfunctions of reproducing kernel for cubic splines at the pre-selected grid points (must agree with xg).
K	An integer value. The truncation parameter indicates the number of eigenvalues/eigenfunctions used in approximation. The default value is 30.
method	A character string. If "REML" the LME model is fit by maximizing the restricted log-likelihood. If "ML" the log-likelihood is maximized. Defaults to "REML".
pstd	An indicator of whether standard deviation is desired. The default value is FALSE.

**Value**

A vector(s) of following component(s):

fit	The low-rank approximation of cubic smoothing spline estimate.
pstd	The corresponding posterior standard deviation.

**Examples**

```
## Not run:
data(eigenM)
x <- runif(1000)
y <- sin(32*pi*x)-8*(x-.5)^2 + rnorm(1000)
lrspline.cubic(x,y,eigenM$xg,eigenM$e,K,method="REML",pstd=FALSE)

## End(Not run)
```

nystrom.cubic

*Nystrom Approximation for Cubic Smoothing Spline Estimates***Description**

Computes a Nystrom approximation based on randomly selected columns of  $\Sigma$  matrix, where estimation utilizes functions for linear mixed effect model (LME).

**Usage**

```
nystrom.cubic(x, y, p = 30, method = "REML", pstd = FALSE)
```

**Arguments**

x	The values of independent variable. It should be a vector.
y	The values of dependent variable. It should be a vector.
p	An integer value. The selection parameter indicates the number of columns for random selection and approximation. The default value is 30.
method	A character string. If "REML" the LME model is fit by maximizing the restricted log-likelihood. If "ML" the log-likelihood is maximized. Defaults to "REML".
pstd	An indicator of whether standard deviation is desired. The default value is FALSE.
e	A list of two elements of "values" and "vectors", which refer respectively, the eigenvalues and eigenfunctions of reproducing kernel for cubic splines at the pre-selected gridpoints (must agree with xg).

**Value**

A vector(s) of following component(s):

fit	The Nystrom approximation of cubic smoothing spline estimate.
pstd	The corresponding posterior standard deviation.

**Examples**

```
## Not run:
data(eigenM)
x <- runif(1000)
y <- sin(32*pi*x)-8*(x-.5)^2 + rnorm(1000)
nystrom.cubic(x,y,xg,e,K,method="REML",pstd=FALSE)

## End(Not run)
```

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