

# Package ‘interpolators’

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**Title** Some Interpolation Methods

**Version** 1.0.1

**Description** Some interpolation methods taken from 'Boost': barycentric rational interpolation, modified Akima interpolation, PCHIP (piecewise cubic Hermite interpolating polynomial) interpolation, and Catmull-Rom splines.

**License** GPL-3

**URL** <https://github.com/stla/interpolators>

**BugReports** <https://github.com/stla/interpolators/issues>

**Imports** Rcpp

**LinkingTo** BH, Rcpp

**Encoding** UTF-8

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**Repository** CRAN

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evalInterpolator      *Interpolator evaluation*

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**Description**

Evaluation of an interpolator at some given values.

**Usage**

```
evalInterpolator(ipr, x, derivative = 0)
```

**Arguments**

ipr	an interpolator
x	numeric vector giving the values to be interpolated; missing values are not allowed; for Catmull-Rom splines, the values must be between 0 and 1
derivative	order of differentiation, 0 or 1

**Value**

Numeric vector of interpolated values, or numeric matrix of interpolated points for the Catmull-Rom interpolator.

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iprBarycentricRational  
*Barycentric rational interpolator*

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**Description**

Barycentric rational interpolator.

**Usage**

```
iprBarycentricRational(x, y, ao = 3)
```

**Arguments**

x, y	numeric vectors giving the coordinates of the known points, without missing value
ao	approximation order, an integer greater than or equal to 3

**Details**

See [Barycentric rational interpolation](#).

**Value**

An interpolator, for usage in [evalInterpolator](#).

**Examples**

```
library(interpolators)
x <- c(1, 2, 4, 5)
y <- x^2
ipr <- iprBarycentricRational(x, y)
evalInterpolator(ipr, c(2, 3))
evalInterpolator(ipr, c(2, 3), derivative = 1)
```

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iprCatmullRom	<i>Catmull-Rom interpolator</i>
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**Description**

Catmull-Rom interpolator for 2-dimensional or 3-dimensional points.

**Usage**

```
iprCatmullRom(points, closed = FALSE, alpha = 0.5)
```

**Arguments**

points	numeric matrix of 2D or 3D points, one point per row
closed	Boolean, whether the curve is closed
alpha	parameter between 0 and 1; the default value 0.5 is recommended

**Details**

See [Catmull-Rom splines](#).

**Value**

An interpolator, for usage in [evalInterpolator](#).

**Examples**

```
library(interpolators)
points <- rbind(
  c(0, 2.5),
  c(2, 4),
  c(3, 2),
  c(4, 1.5),
  c(5, 6),
  c(6, 5),
  c(7, 3),
```

```

    c(9, 1),
    c(10, 2.5),
    c(11, 7),
    c(9, 5),
    c(8, 6),
    c(7, 5.5)
  )
  ipr <- iprCatmullRom(points)
  s <- seq(0, 1, length.out = 400)
  Curve <- evalInterpolator(ipr, s)
  head(Curve)
  plot(Curve, type = "l", lwd = 2)
  points(points, pch = 19)

# a closed example (pentagram) ####
rho <- sqrt((5 - sqrt(5))/10)
R <- sqrt((25 - 11*sqrt(5))/10)
points <- matrix(NA_real_, nrow = 10L, ncol = 2L)
points[c(1, 3, 5, 7, 9), ] <- t(vapply(0:4, function(i){
  c(rho*cospi(2*i/5), rho*sinpi(2*i/5))
}, numeric(2L)))
points[c(2, 4, 6, 8, 10), ] <- t(vapply(0:4, function(i){
  c(R*cospi(2*i/5 + 1/5), R*sinpi(2*i/5 + 1/5))
}, numeric(2L)))
ipr <- iprCatmullRom(points, closed = TRUE)
s <- seq(0, 1, length.out = 400L)
Curve <- evalInterpolator(ipr, s)
plot(Curve, type = "l", lwd = 2, asp = 1)
points(points, pch = 19)

```

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iprMakima

*Modified Akima interpolator*


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### Description

Modified Akima interpolator.

### Usage

```
iprMakima(x, y)
```

### Arguments

x, y	numeric vectors giving the coordinates of the known points, without missing value
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### Details

See [Modified Akima interpolation](#).

**Value**

An interpolator, for usage in [evalInterpolator](#).

**Examples**

```
library(interpolators)
x <- seq(0, 4*pi, length.out = 9L)
y <- x - sin(x)
ipr <- iprMakima(x, y)
curve(x - sin(x), from = 0, to = 4*pi, lwd = 2)
curve(
  evalInterpolator(ipr, x),
  add = TRUE, col = "blue", lwd = 3, lty = "dashed"
)
points(x, y, pch = 19)
```

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iprPCHIP

*PCHIP interpolator*


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**Description**

PCHIP interpolator. It is monotonic.

**Usage**

```
iprPCHIP(x, y)
```

**Arguments**

`x, y` numeric vectors giving the coordinates of the known points, without missing value

**Details**

See [PCHIP interpolation](#).

**Value**

An interpolator, for usage in [evalInterpolator](#).

**Examples**

```
library(interpolators)
x <- seq(0, 4*pi, length.out = 9L)
y <- x - sin(x)
ipr <- iprPCHIP(x, y)
curve(x - sin(x), from = 0, to = 4*pi, lwd = 2)
curve(
  evalInterpolator(ipr, x),
```

```
    add = TRUE, col = "blue", lwd = 3, lty = "dashed"  
  )  
points(x, y, pch = 19)
```

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