

# Package ‘rsegfit’

September 7, 2017

**Type** Package

**Title** An r package for segfit

**Version** 0.1

**Date** 2017-09-05

**Description** An r interface to segfit algorithm

**Depends** R (>= 2.15.0)

**License** GPL (>= 2)

**NeedsCompilation** yes

**RoxygenNote** 6.0.1

## R topics documented:

fitted.segfit . . . . .	1
plot.segfit . . . . .	2
residuals.segfit . . . . .	3
segfit . . . . .	3
summary.segfit . . . . .	4
<b>Index</b>	<b>5</b>

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fitted.segfit	<i>Extract the fitted values from a segfit object</i>
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## Description

Extract the fitted values from a segfit object

## Usage

```
## S3 method for class 'segfit'
fitted(sf, concat = TRUE)
```

**Arguments**

<code>sf</code>	The segfit object
<code>concat</code>	Whether the fitted values of different segments will be concatenated.

**Details**

Extract the fitted values from a segfit object

**Value**

If `concat==TRUE`, the fitted values of different segments will be concatenated into one vector. Otherwise, the returned value will be a list of fitted values of each segment.

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<code>plot.segfit</code>	<i>Plot a segfit object</i>
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**Description**

Plot a segfit object

**Usage**

```
## S3 method for class 'segfit'
plot(x, y = "", col.data = "black", col.seg = "red",
     legend.pos = "topleft")
```

**Arguments**

<code>x</code>	If x is a segfit object, input y is ignored.
<code>y</code>	A segfit object, If not provided, x must be a segfit object.
<code>col.data</code>	color of data
<code>col.seg</code>	color of segments
<code>legend.pos</code>	legend positions

**Details**

Plot a segfit object

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residuals.segfit	<i>Extract the residuals from a segfit object</i>
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**Description**

Extract the residuals from a segfit object

**Usage**

```
## S3 method for class 'segfit'
residuals(sf, concat = TRUE)
```

**Arguments**

sf	The segfit object
concat	Whether the residuals of different segments will be concatenated.

**Details**

Extract the residuals from a segfit object

**Value**

If concat==TRUE, the residuals of different segments will be concatenated into one vector. Otherwise, the returned value will be a list of residuals of each segment.

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segfit	<i>segfit a sequence</i>
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**Description**

Do segmentation on "data"

**Usage**

```
segfit(data, smp = 2.3, lb = -6, ub = 6, maxiter = 1000, factr = 5000,
       pgtol = 1e-04)
```

**Arguments**

data	The series to be segfitted
smp	The smaller, the more segments will be found
lb	lower bound of parameter \$b\$ of each segment
ub	upper bound of parameter \$b\$ of each segment
maxiter	maximum iteration in optimisation
factr	maximum function evaluation
pgtol	tolerance used in optimisation

**Details**

"Segment and fit" a sequence. Each sequence will be described by  $y=a*x^b+c$  where  $x=1:\text{seg.length}$  ( $\text{order}==0$ ) or  $x = \text{seg.length}:1$  ( $\text{order}==1$ ). The segmentation is to minimise the sum of mse of each segments plus  $\text{smp}*\text{numSegment}$ .

**Value**

A "segfit" object. It contains the params of all the segments. For each segment, it contains the head index (hi), tail index (ei), parameter \$a\$ (a) parameter \$b\$ (b), parameter \$c\$ (c), fitting order (order), fitted values (fit), and fitting residuals (residual). It also has a attribute "data" for the original data

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summary.segfit	<i>Summarise a segfit object</i>
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**Description**

Summarise a segfit object

**Usage**

```
## S3 method for class 'segfit'
summary(sf)
```

**Arguments**

sf                      The segfit object to summarise

**Details**

Summarise a segfit object

# Index

`fitted.segfit`, [1](#)

`plot.segfit`, [2](#)

`residuals.segfit`, [3](#)

`segfit`, [3](#)

`summary.segfit`, [4](#)