

# Package ‘pacviz’

August 25, 2020

**Title** Pac-Man Residual Function

**Version** 1.0.0.0

**Description** TBA

**License** MIT + file LICENSE

**Depends** R (>= 3.3.3)

**Imports** circlize,  
e1071,  
graphics,  
plotrix,  
stats,  
utils

**Suggests** knitr,  
rmarkdown

**VignetteBuilder** knitr

**Encoding** UTF-8

**LazyData** true

**RoxygenNote** 7.1.1

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deg2rad	<i>Pac-Man SVM</i>
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**Description**

Conversion between degrees and radians

**Usage**

deg2rad(deg)

**Arguments**

deg                      Angle in degrees

**Value**

Pac-Man SVM

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linMap	<i>Pac-Man SVM</i>
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**Description**

linear map

**Usage**

linMap(x, i, f)

**Arguments**

x                      Range of values to be mapped  
i                      Lowest value  
f                      Largest value

**Value**

Pac-Man SVM

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`lsvm`*Pac-Man SVM*

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**Description**

A visualization technique in R for regression analysis results, specifically residual values, based on a restricted radial coordinate system. It provides a broad view perspective on the performance of regression models, and supports most model inputs. See the `pacviz` documentation page for more information: <https://pharaohcola13.github.io/pacviz/book/>

**Usage**

```
lsvm(x, y, l, title, train_size = 0.7, rand_state = sample(1:2^15, 1))
```

**Arguments**

<code>x, y</code>	Numeric data
<code>l</code>	Numeric labels data
<code>title</code>	Figure title
<code>train_size</code>	Fraction of total data that the SVM will train on
<code>rand_state</code>	Value of the random state used to set the seed

**Value**

Pac-Man SVM

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`pac.lsvm`*Pac-Man SVM*

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**Usage**

```
pac.lsvm(  
  x,  
  y,  
  l,  
  title,  
  axis_label,  
  train_size = 0.7,  
  rand_state = sample(1:2^15, 1)  
)
```

**Arguments**

<code>x, y</code>	Numeric data
<code>l</code>	Numeric labels data
<code>title</code>	Figure title
<code>axis_label</code>	Label for the axis
<code>train_size</code>	Fraction of total data that the SVM will train on
<code>rand_state</code>	Value of the random state used to set the seed

**Value**

Pac-Man SVM

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<code>pac.plot</code>	<i>Pac-Man SVM</i>
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**Usage**

```
pac.plot(x, y, title, xaxislabel, yaxislablel, xunits, yunits, color1 = "gold")
```

**Arguments**

<code>x, y</code>	Numeric data
<code>title</code>	Figure title
<code>xaxislabel</code>	Angular axis label
<code>yaxislabel</code>	Radial axis label
<code>xunits</code>	String to define units on the angular axis (For temperature measurements use 'degC' or 'degF')
<code>yunits</code>	String to define units on the radial axis (For temperature measurements use 'degC' or 'degF')
<code>color1</code>	Color value as string or rgb

**Value**

Pac-Man SVM

**Examples**

```
# Generic Pac-Man residual
data("cars")
pac.plot(cars$dist, cars$speed, 'Example 1', "Distance", "Speed", 'm', 'm/s')
```

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pac.resid	<i>Pac-Man Residual Function</i>
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## Description

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## Usage

```
pac.resid(  
  x,  
  y,  
  title,  
  unit,  
  axis_label,  
  model = lm(y ~ x, data = data.frame(x, y)),  
  color1 = "gold",  
  standardize = FALSE  
)
```

## Arguments

x, y	Numeric data
title	Figure title
unit	String to define units on the angular axis (For temperature measurements use 'degC' or 'degF')
axis_label	Angular axis label
model	An object for which the extraction of model residuals is meaningful.
color1	Color value as string or rgb
standardize	Boolean to standardize the residual value

## Value

Pac-Man residual plot

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rad2deg

*Pac-Man SVM*


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**Description**

Conversion between radians and degrees

**Usage**

```
rad2deg(rad)
```

**Arguments**

rad	Angle in radians
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**Value**

Pac-Man SVM

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svm.partition

*Pac-Man SVM*


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**Description**

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**Usage**

```
## S3 method for class 'partition'
svm(x, y, l, train_size = 0.7, rand_state = sample(1:2^15, 1))
```

**Arguments**

x, y	Numeric data
l	Numeric labels data
train_size	Fraction of total data that the SVM will train on
rand_state	Value of the random state used to set the seed

**Value**

Data Partition

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