

pacviz

Spencer Riley

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About this document

This is the documentation of the **pacviz** package. Examples in the book are generated under version 1.0.0.0.

Session info:

```
sessionInfo()
```

```
## R version 3.6.1 (2019-07-05)
## Platform: x86_64-pc-linux-gnu (64-bit)
## Running under: Linux Mint 20
##
## Matrix products: default
## BLAS: /usr/local/lib/R/lib/libRblas.so
## LAPACK: /usr/local/lib/R/lib/libRlapack.so
##
## locale:
##  [1] LC_CTYPE=en_US.UTF-8      LC_NUMERIC=C
##  [3] LC_TIME=en_US.UTF-8      LC_COLLATE=en_US.UTF-8
##  [5] LC_MONETARY=en_US.UTF-8  LC_MESSAGES=en_US.UTF-8
##  [7] LC_PAPER=en_US.UTF-8     LC_NAME=C
##  [9] LC_ADDRESS=C             LC_TELEPHONE=C
## [11] LC_MEASUREMENT=en_US.UTF-8 LC_IDENTIFICATION=C
##
## attached base packages:
## [1] stats      graphics  grDevices  utils      datasets  methods    base
##
## loaded via a namespace (and not attached):
##  [1] compiler_3.6.1  magrittr_1.5    bookdown_0.20   htmltools_0.5.0
##  [5] tools_3.6.1     yaml_2.2.1      stringi_1.4.6   rmarkdown_2.3
##  [9] knitr_1.29      stringr_1.4.0   digest_0.6.25   xfun_0.16
## [13] rlang_0.4.7     evaluate_0.14
```

The Author

Chapter 1

Introduction to the package

See the pacviz documentation page for more information: <https://pharaohcola13.github.io/pacviz/book/>.

The functions that are enclosed in this package include:

- pacman

1.1 Installation Guide

For the most up-to-date version of the package, install it directly from GitHub.

```
devtools::install_github("PharaohCola13/pacviz")
```

Through CRAN (Not yet available)

```
install.packages('pacviz')
```

1.2 Package Dependencies

R ($\geq 3.3.3$)

Packages: circlize, graphics, plotrix, stats, utils

Chapter 2

Pac-Man Residual Plot

2.1 Description

The results of a regression algorithm typically takes the form of a residual plot, showing the relationship (or lack thereof) between the domain and the residual values of the data associated with the model. From the residual, a broad scope of the model's performance can be determined.

`pacviz` contributes a simple approach for looking at the broad view performance of the regression model by constructing a 'Pac-Man' residual plot.

2.1.1 Formalism

This visualization technique applies a bijective map from the domain of the data to angular values between 40 and 320 degrees,

$$X \mapsto [40, 320]. \quad (2.1)$$

This restriction is applied to allow space for radial labels. By taking the absolute value of the residual values on the radial coordinate system, we can observe the overall performance of the model with relative ease.

In addition, we have added the residual standard deviation for the model both in its numerical form and graphically as a dashed line at one σ . This circular segment was created by the `circlize` package .

There are shortcomings for a visualization that views big-picture components of a model. In the case of the 'Pac-Man' residual plot we lose the ability to determine the dependence of the relationship. Through a traditional residual plot, it would be simple to determine if there was systematic or random error based on the relationship between the standard deviation and the domain of the data.

2.2 Recommendations

The discussions in this section will revolve around preferred color schemes and helpful character codes for UTF-8 symbols that can be used as units.

2.2.1 Color Scheme

Since one of the two colors in the visualization is white, the other is a user input with the default being `gold`. The following colors are predefined in R, with the whole list available here¹, and are a good fit in terms of contrast and readability.

- `lightskyblue`
- `lightsteelblue`
- `darksalmon`
- `palegreen`
- `gray86`
- `plum`

2.2.2 Characters

- Angstrom: `\uc5`
- More will be added

2.3 Usage

The function is setup to implement an arbitrary regression model and supports residual standardization. As we have discussed above,

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

2.4 Examples

For the following examples, the domain and range that will be processed by the function will be:

```
x <- rnorm(20, mean=0, sd=10)  
y <- log(rnorm(20, mean=0, sd=10), base=exp(1))
```

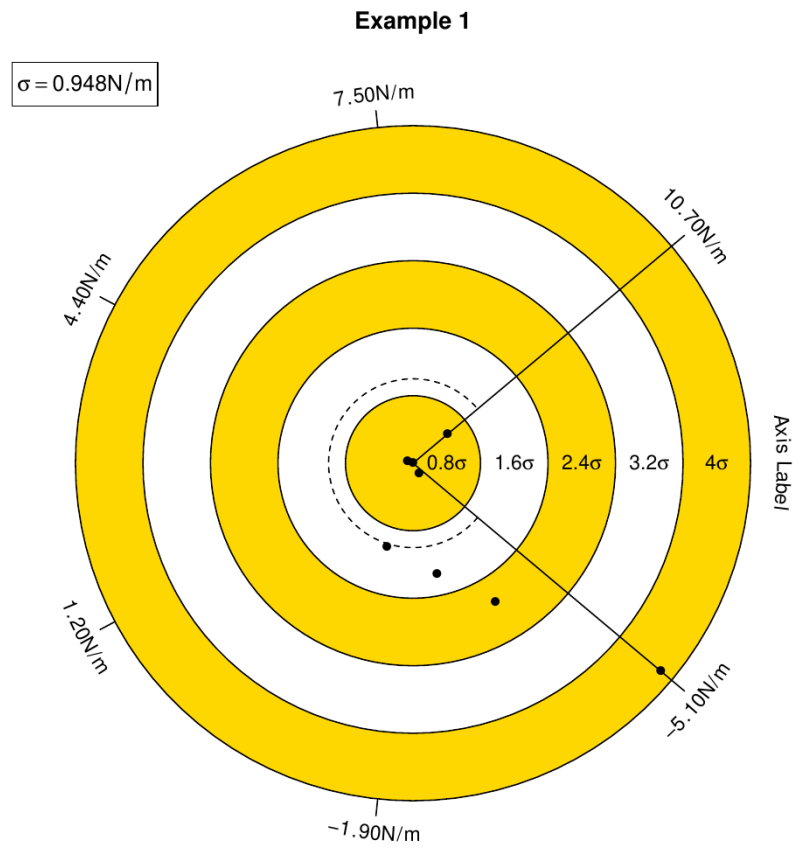
¹<http://www.stat.columbia.edu/tzheng/files/Rcolor.pdf>

It should be noted that these values do change after every run.

2.4.1 Example 1

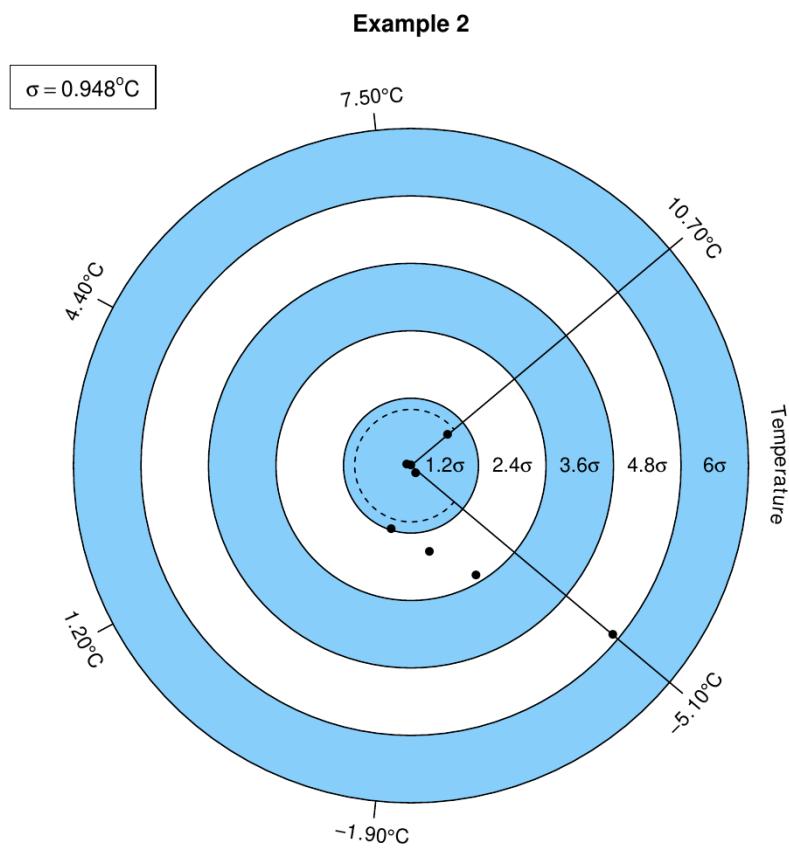
In the below snippet, we use

```
# Generic Pac-Man residual
pacman(x,y,'Example 1',
      'units',
      'Axis Label')
```



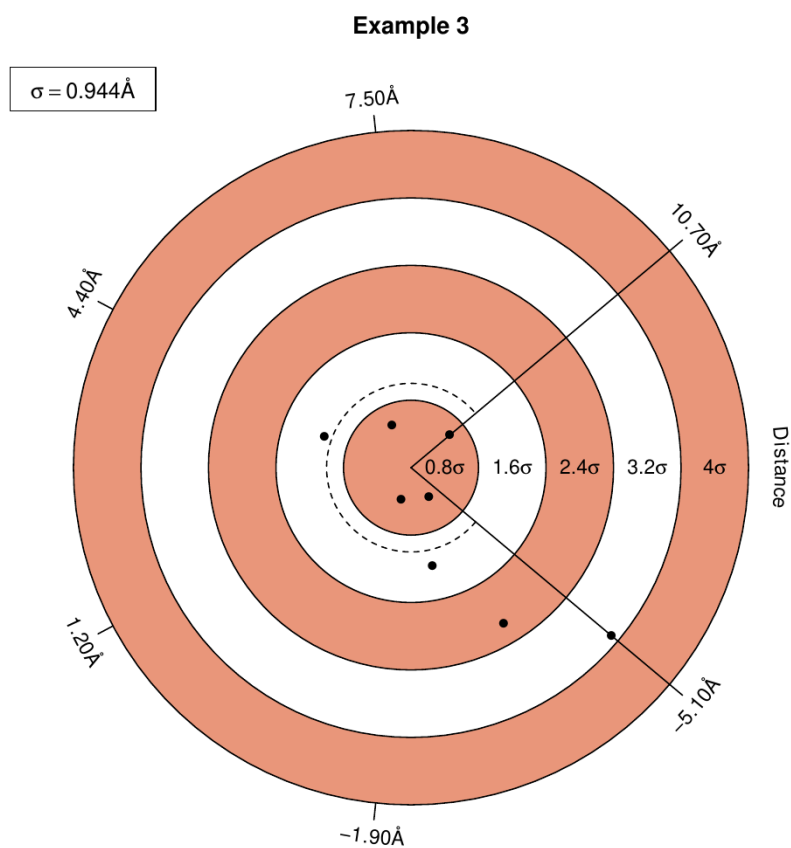
2.4.2 Example 2

```
# Pac-Man residual using alternate color,
# residual standardization, and temperature units
pacman(x,y, 'Example 2',
        'degC',
        "Temperature",
        color1="lightblue",
        standardize=TRUE)
```



2.4.3 Example 3

```
# Pac-Man residual using alternate color,  
# a quadratic model, and a UTF8 character for units  
pacman(x,y, 'Example 3',  
        "\uc5",  
        "Distance",  
        model=lm(y~poly(x,2)),  
        color1="darksalmon")
```



Chapter 3

Releases

V1.0 *Blinky*

This is the initial release of the `pacviz` R package.

Coming Soon