

# Package: andrews (via r-universe)

September 17, 2024

**Type** Package

**Title** Various Andrews Curves

**Version** 1.1.2

**Maintainer** Sigbert Klinke <sigbert@hu-berlin.de>

**Depends** R (>= 2.10)

**Description** Visualisation of multidimensional data through different Andrews curves: Andrews, D. F. (1972) Plots of High-Dimensional Data. *Biometrics*, 28(1), 125-136. <doi:10.2307/2528964>.

**License** GPL-3

**URL** <https://github.com/sigbertklinke/andrews> (development version)

**Encoding** UTF-8

**LazyData** true

**Roxygen** list(markdown = TRUE)

**Imports** grDevices, graphics, gmp

**Suggests** knitr, rmarkdown, robustbase, mclust

**VignetteBuilder** knitr

**RoxygenNote** 7.2.3

**Repository** <https://sigbertklinke.r-universe.dev>

**RemoteUrl** <https://github.com/sigbertklinke/andrews>

**RemoteRef** HEAD

**RemoteSha** 75fdbe2e26620f9f05d9330a48675818c13c5ba0

## Contents

andrews	2
andrews0	4
banknote	5
deftype	6
generate_n_primes	7

normalize . . . . .	7
numarray . . . . .	8
outlyingness . . . . .	9
selectand . . . . .	10
zzz . . . . .	11

## Index 12

andrews *Andrews curves*

### Description

Andrews curves for visualization of multidimensional data. For colouring the curves see the details. For differences between `andrews` and `andrews0` see the vignette("andrews"). With the same parameters called both functions should create the same plot. `type==5` is a modification of `type==3` and `type==6` is a modification of `type==4`.

### Usage

```
andrews(
  df,
  type = 1,
  clr = NULL,
  step = 100,
  ymax = 10,
  alpha = NULL,
  palcol = NULL,
  lwd = 1,
  lty = "solid",
  ...
)
```

### Arguments

<code>df</code>	data frame or an R object that can be converted into a data frame with <code>as.data.frame</code>
<code>type</code>	type of curve <ul style="list-style-type: none"> <li>• 1: <math>f(t) = x_1/\sqrt{2} + x_2 \sin(t) + x_3 \cos(t) + x_4 \sin(2t) + x_5 \cos(2t) + \dots</math></li> <li>• 2: <math>f(t) = x_1 \sin(t) + x_2 \cos(t) + x_3 \sin(2t) + x_4 \cos(2t) + \dots</math></li> <li>• 3: <math>f(t) = x_1 \cos(t) + x_2 \cos(\sqrt{2}t) + x_3 \cos(\sqrt{3}t) + \dots</math></li> <li>• 4: <math>f(t) = 0.5^{p/2}x_1 + 0.5^{(p-1)/2}x_2(\sin(t) + \cos(t)) + 0.5^{(p-2)/2}x_3(\sin(t) - \cos(t)) + 0.5^{(p-3)/2}x_4(\sin(2t) + \cos(2t)) + 0.5^{(p-4)/2}x_5(\sin(2t) - \cos(2t)) + \dots</math> with <math>p</math> the number of variables</li> <li>• 5: <math>f(t) = x_1 \cos(\sqrt{p_0}t) + x_2 \cos(\sqrt{p_1}t) + x_3 \cos(\sqrt{p_2}t) + \dots</math> with <math>p_0 = 1</math> and <math>p_i</math> the <math>i</math>-th prime number</li> <li>• 6: <math>f(t) = 1/\sqrt{2}(x_1 + x_2(\sin(t) + \cos(t)) + x_3(\sin(t) - \cos(t)) + x_4(\sin(2t) + \cos(2t)) + x_5(\sin(2t) - \cos(2t)) + \dots)</math></li> </ul>

<code>clr</code>	number/name of column in the data frame for color of curves
<code>step</code>	smoothness of curves
<code>ymin</code>	minimum of y coordinate
<code>ymax</code>	maximum of y coordinate
<code>alpha</code>	semi-transparent color ( $0 < \alpha < 1$ ) which are supported only on some devices
<code>palcol</code>	a function which generates a set of colors, see details
<code>lwd</code>	line width, a positive number, defaulting to 1.
<code>lty</code>	line type, can either be specified as an integer (0=blank, 1=solid (default), 2=dashed, 3=dotted, 4=dotdash, 5=longdash, 6=twodash) or as one of the character strings "blank", "solid", "dashed", "dotted", "dotdash", "longdash", or "twodash", where "blank" uses 'invisible lines' (i.e., does not draw them).
<code>...</code>	further named parameters given to <code>graphics::plot.default()</code> except x, y, and type.

### Details

If `clr` has length one then it is used as column number or column name for coloring the curves:

- If `df[, clr]` is numeric then `palcol` must be function which returns colors for values in the range  $\in [0, 1]$  using normalized variable. The default is function `function(v) { hsv(0, 1, v) }`.
- Otherwise `df[, clr]` is converted to a factor and `palcol` must be a function which returns for each level a color. The parameter for `palcol` is the number of levels and the default is `grDevices::rainbow()`. If the length of `clr` is the number of rows of `df` then `clr` is interpreted as colors.

Andrews curves transform multidimensional data into curves. This package presents four types of curves.

### Value

nothing

### Author(s)

Sigbert Klinke [sigbert@hu-berlin.de](mailto:sigbert@hu-berlin.de), Jaroslav Myslivec [jaroslav.myslivec@upce.cz](mailto:jaroslav.myslivec@upce.cz)

### References

- Andrews, D. F. (1972) Plots of High-Dimensional Data. *Biometrics*, vol. 28, no. 1, pp. 125-136.
- Khattree, R., Naik, D. N. (2002) Andrews Plots for Multivariate Data: Some New Suggestions and Applications. *Journal of Statistical Planning and Inference*, vol. 100, no. 2, pp. 411-425.

**Examples**

```
data(iris)
op <- par(mfrow=c(1,2))
andrews0(iris,clr=5,ymax=3)
andrews(iris,clr=5,ymax=3)
par(op)
andrews(iris,type=4,clr=5,ymax=NA)
```

andrews0

*Andrews curves***Description**

Andrews curves for visualization of multidimensional data. For differences between `andrews` and `andrews2` see the `vignette("andrews")`. For colouring the curves see the details.

**Usage**

```
andrews0(
  df,
  type = 1,
  clr = NULL,
  step = 100,
  ymax = 10,
  main = NULL,
  sub = NULL
)
```

**Arguments**

<code>df</code>	data frame
<code>type</code>	type of curve <ul style="list-style-type: none"> <li>1: <math>f(t) = x_1/\sqrt{2} + x_2 \sin(t) + x_3 \cos(t) + x_4 \sin(2t) + x_5 \cos(2t) + \dots</math></li> <li>2: <math>f(t) = x_1 \sin(t) + x_2 \cos(t) + x_3 \sin(2t) + x_4 \cos(2t) + \dots</math></li> <li>3: <math>f(t) = 0.5^{p/2}x_1 + 0.5^{(p-1)/2}x_2(\sin(t) + \cos(t)) + 0.5^{(p-2)/2}x_3(\sin(t) - \cos(t)) + 0.5^{(p-3)/2}x_4(\sin(2t) + \cos(2t)) + 0.5^{(p-6)/2}x_5(\sin(2t) - \cos(2t)) + \dots</math> with <math>p</math> the number of variables</li> <li>4: <math>f(t) = 1/\sqrt{2}(x_1 + x_2(\sin(t) + \cos(t)) + x_3(\sin(t) - \cos(t)) + x_4(\sin(2t) + \cos(2t)) + x_5(\sin(2t) - \cos(2t)) + \dots)</math></li> </ul>
<code>clr</code>	number/name of column in the date frame for color of curves
<code>step</code>	smoothness of curves
<code>ymax</code>	maximum of y coordinate.
<code>main</code>	main title for the plot
<code>sub</code>	sub title for the plot

## Details

Andrews curves transform multidimensional data into curves. This package presents four types of curves

If `df[,clr]` is numeric then `hsv(1,1,v)` with the normalized values (on  $[0, 1]$ ) of `df[,clr]` is used. Otherwise the number of unique values in `nuv <- unique(df[,clr])` is used in connection with `rainbow(nuv)`.

## Value

nothing

## Author(s)

Jaroslav Myslivec [jaroslav.myslivec@upce.cz](mailto:jaroslav.myslivec@upce.cz)

## References

- Andrews, D. F. (1972) Plots of High-Dimensional Data. *Biometrics*, vol. 28, no. 1, pp. 125-136.
- Khatree, R., Naik, D. N. (2002) Andrews Plots for Multivariate Data: Some New Suggestions and Applications. *Journal of Statistical Planning and Inference*, vol. 100, no. 2, pp. 411-425.

## Examples

```
data(iris)
andrews0(iris,clr=5,ymax=3)
andrews0(iris,type=4,clr=5,ymax=2)
```

---

banknote

*Swiss banknotes data*

---

## Description

The data set contains six measurements made on 100 genuine and 100 counterfeit old-Swiss 1000-franc bank notes. The data frame and the documentation is a copy of [mclust::banknote](#).

## Usage

```
banknote
```

## Format

A data frame with 200 rows and 7 columns:

**Status** the status of the banknote: genuine or counterfeit

**Length** Length of bill (mm)

**Left** Width of left edge (mm)

**Right** Width of right edge (mm)  
**Bottom** Bottom margin width (mm)  
**Top** Top margin width (mm)  
**Diagonal** Length of diagonal (mm)

### Source

Flury, B. and Riedwyl, H. (1988). Multivariate Statistics: A practical approach. London: Chapman & Hall, Tables 1.1 and 1.2, pp. 5-8.

---

deftype	<i>deftype</i>
---------	----------------

---

### Description

Defines a function which can be used as basis for Andrews curves  $f_i(t) = \sum_{j=1}^p x_{ij} f_j(t)$ .

### Usage

```
deftype(index = NULL, FUN = NULL, xlim = c(-pi, pi))
```

### Arguments

index	index/name of the function
FUN	function of the form function(n, t) {...}
xlim	default range for displaying curves (default: c(-pi, pi))

### Value

either a list of all functions or a single function

### Examples

```
# define a new andrews curve, just with sine curves
deftype("sine", function(n, t) {
  n <- as.integer(if (n<1) 1 else n)
  m <- matrix(NA, nrow=length(t), ncol=n)
  for (i in 1:n) m[,i] <- sin(i*t)
  m
})
andrews(iris, "sine")
# query
deftype()
deftype("sine")
```

---

generate_n_primes	<i>Generate a Sequence of Prime Numbers</i>
-------------------	---

---

**Description**

Generates a vector of the first  $n$  primes using `gmp::nextprime()`.

**Usage**

```
generate_n_primes(n, one = FALSE)
```

**Arguments**

n	the number of primes to generate.
one	should 1 included in the sequence (default: FALSE)

**Value**

an integer vector of prime numbers

**Examples**

```
generate_n_primes(5)
generate_n_primes(5, TRUE)
```

---

normalize	<i>Normalization</i>
-----------	----------------------

---

**Description**

Normalization of a variable:

- `type==1`: ar normalized into  $[0, 1]$ ,
- `type==2`: ar is standardized,
- otherwise no normalization is done.

**Usage**

```
normalize(ar, type = 1)
```

**Arguments**

ar	numeric variable.
type	integer: type of normalization (default: 1)

**Details**

Normalization of variable:  $ar \leftarrow (ar - \min(ar)) / (\max(ar) - \min(ar))$

**Value**

Returns normalized variable.

**Author(s)**

Jaroslav Myslivec [jaroslav.myslivec@upce.cz](mailto:jaroslav.myslivec@upce.cz), Sigbert Klinke [sigbert@hu-berlin.de](mailto:sigbert@hu-berlin.de)

**Examples**

```
normalize(iris[,1])
```

---

numarray

*Numeric array*

---

**Description**

Extracts numeric array from data frame.

**Usage**

```
numarray(df)
```

**Arguments**

df                    data frame.

**Details**

Extracts numeric array from data frame.

**Value**

Returns numeric array.

**Author(s)**

Jaroslav Myslivec [jaroslav.myslivec@upce.cz](mailto:jaroslav.myslivec@upce.cz), Sigbert Klinke [sigbert@hu-berlin.de](mailto:sigbert@hu-berlin.de)

**Examples**

```
numarray(iris)
```



---

outlyingness	<i>outlyingness</i>
--------------	---------------------

---

### Description

Computes the Stahel-Donoho outlyingness. If `type` is any of the available types by `andrews()` then the projection vectors are generated along the andrews curves. Otherwise step random directions will be used. Note that the projection vectors are always normalized to length one.

### Usage

```
outlyingness(x, type = 1, step = 100, xlim = NULL, normalize = 1)
```

### Arguments

<code>x</code>	data frame
<code>type</code>	type of curve, see <code>andrews()</code>
<code>step</code>	step smoothness of curves
<code>xlim</code>	the x limits (x1, x2)
<code>normalize</code>	type of normalization, see <code>normalize()</code>

### Value

the Stahel-Donoho outlyingness

### References

- Stahel, W. (1981), Robuste Schätzungen: infinitesimale Optimalität und Schätzungen von Kovarianzmatrizen, PhD thesis, ETH Zürich.
- Donoho, D. (1982), Breakdown properties of multivariate location estimators, Ph.D. Qualifying paper, Dept. Statistics, Harvard University, Boston.

### Examples

```
# use projection vectors from the Andrews curve
sdo <- outlyingness(iris)
col <- gray(1-sdo/max(sdo))
andrews(iris, clr=col, ymax=NA)
# use 1000 random projection vectors
sdo <- outlyingness(iris, type=0, step=1000)
col <- gray(1-sdo/max(sdo))
andrews(iris, clr=col, ymax=NA)
# use 1000 random projection vectors with adjusted outlyingness
library("robustbase")
x <- numarray(iris)
x <- scale(x, center=apply(x, 2, min), scale=apply(x, 2, max)-apply(x, 2, min))
sdo <- adjOutlyingness(x, ndir=1000, only.outlyingness=TRUE)
```

```
col <- gray(1-sdo/max(sdo))
andrews(as.data.frame(x), clr=col, ymax=NA)
```

---

selectand	<i>Selecting in Andrews curves</i>
-----------	------------------------------------

---

### Description

Selecting object utility in Andrews curves

### Usage

```
selectand(df, type = 1, step = 100, ncol = 0, from = 0, to = 1, col = 2)
```

### Arguments

df	data frame.
type	type of curve.
step	smoothness of curves.
ncol	number of column in data frame for selection.
from	from value.
to	to value.
col	color of selected objects.

### Details

Define which objects will be selected (colored) in Andrews curves.

### Value

Nothing

### Author(s)

Jaroslav Myslivec [jaroslav.myslivec@upce.cz](mailto:jaroslav.myslivec@upce.cz)

### Examples

```
data(iris)
andrews(iris,clr=5,ymax=3)
selectand(iris,ncol=1,from=5,to=5.5,col=1)
```

---

*zzz**Comparison*

---

**Description**

Creates and displays a temporary PDF file with different diagrams comparing `andrews` and `andrews0` plots.

**Usage**

```
zzz()
```

**Value**

nothing

**Examples**

```
if (interactive()) zzz()
```

# Index

## \* datasets

banknote, 5

## \* hplot

andrews, 2

andrews0, 4

selectand, 10

andrews, 2

andrews(), 9

andrews0, 4

banknote, 5

deftype, 6

generate\_n\_primes, 7

gmp::nextprime(), 7

graphics::plot.default(), 3

grDevices::rainbow(), 3

mclust::banknote, 5

normalize, 7

normalize(), 9

numarray, 8

outlyingness, 9

selectand, 10

zzz, 11