

# Package ‘ggESDA’

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**Type** Package

**Title** Exploratory Symbolic Data Analysis with 'ggplot2'

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**Description** Implements an extension of 'ggplot2' and visualizes the symbolic data with multiple plot which can be adjusted by more general and flexible input arguments. It also provides a function to transform the classical data to symbolic data by both clustering algorithm and customized method.

**Depends** ggplot2, R (>= 3.5.0), tidyverse, RSDA

**Suggests** testthat (>= 2.1.0), knitr, rmarkdown

**Imports** rlang, R6, dplyr, tidyr, gridExtra, gtools, stringr, proclim, ggforce, ggpubr, ggthemes, tibble, magrittr, vctrs

**License** GPL (>= 2)

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**LazyData** true

**RoxygenNote** 7.1.2

**VignetteBuilder** knitr

**URL** <https://github.com/kiangkiangkiang/ggESDA>

**BugReports** <https://github.com/kiangkiangkiang/ggESDA/issues>

**NeedsCompilation** no

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**R topics documented:**

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AbaloneIdt	<i>AbaloneIdt data example</i>
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**Description**

AbaloneIdt interval data example.

**Usage**

```
data(AbaloneIdt)
```

**Format**

An object of class `data.frame` (inherits from `symbolic_tbl`) with 24 rows and 7 columns.

**References**

Billard L. and Diday E. (2006). Symbolic data analysis: Conceptual statistics and data mining. Wiley, Chichester.

**Examples**

```
data(AbaloneIdt)
ggInterval_index(AbaloneIdt, aes(x = Length))
```

---

BLOOD

*BLOOD data example*

---

**Description**

BLOOD interval data example.

**Usage**

```
data(BLOOD)
```

**Format**

An object of class `tbl_df` (inherits from `tbl`, `data.frame`, `symbolic_tbl`) with 14 rows and 3 columns.

**References**

Billard L. and Diday E. (2006). Symbolic data analysis: Conceptual statistics and data mining. Wiley, Chichester.

**Examples**

```
data(BLOOD)
ggInterval_minmax(BLOOD, aes(x = Hematocrit))
```

Cardiological

*Cardiological data example*

---

**Description**

Cardiological interval data example.

**Usage**

```
data(Cardiological)
```

**Format**

An object of class `symbolic_tbl` (inherits from `tbl_df`, `tbl`, `data.frame`) with 11 rows and 3 columns.

**Source**

<https://CRAN.R-project.org/package=RSDA>

**References**

Billard L. and Diday E. (2006). Symbolic data analysis: Conceptual statistics and data mining. Wiley, Chichester.

**Examples**

```
data(Cardiological)
ggInterval_index(Cardiological, aes(x = Syst))
```

---

Cardiological2

*Cardiological data example*

---

**Description**

Cardiological interval data example.

**Usage**

```
data(Cardiological2)
```

**Format**

An object of class `symbolic_tbl` (inherits from `tbl_df`, `tbl`, `data.frame`) with 15 rows and 3 columns.

## References

Billard L. and Diday E. (2006). Symbolic data analysis: Conceptual statistics and data mining. Wiley, Chichester.

## Examples

```
data(Cardiological2)
ggInterval_index(Cardiological2, aes(x = Syst))
```

---

classic2sym	<i>Convert classical data frame into a symbolic data.</i>
-------------	---

---

## Description

A function for converting a classical data, which may present as a data frame or a matrix with one entry one value, into a symbolic data, which is shown as an interval or a set in an entry. Object after converting is ggESDA class containing interval data and raw data (if it exist) and typically statistics.

## Usage

```
classic2sym(data=NULL, groupby = "kmeans", k=5, minData=NULL, maxData=NULL,
modalData = NULL)
```

## Arguments

data	A classical data frame that you want to be converted into an interval data
groupby	A way to aggregate. It can be either a clustering method or a variable name which exist in input data (necessary factor type). Default "kmeans".
k	A number of groups, which is used by clustering. Default k = 5.
minData	if choose groupby parameter as 'customize', user need to define which data is min data or max data.
maxData	if choose groupby parameter as 'customize', user need to define which data is min data or max data.
modalData	list, each cell of list contain a set of column index of its modal multi-valued data of the input data. the value of it is a proportion presentation, and sum of each row in these columns must be equal to 1. ex 0,1,0 or 0.2,0.3,0.5. the input type of modalData for example is modalData[[1]] = c(2, 3), modalData[[2]] = c(7:10), that 2, 3, 7, 8, 9, 10 columns are modal type of the data. Note: the option is only valid when groupby == "customize".

## Value

classic2sym returns an object of class "ggESDA", which have an interval data and others as follows.

- intervalData - The Interval data after converting also known as a RSDA object.
- rawData - Classical data that user input.
- clusterResult - Cluster results. If the groupby method is a clustering method then it will exist.
- statisticsDF - A list contains data frame including some typically statistics in each group.

**Examples**

```

#classical data to symbolic data
classic2sym(iris)
classic2sym(mtcars, groupby = "kmeans", k = 10)
classic2sym(iris, groupby = "hclust", k = 7)
classic2sym(iris, groupby = "Species")

x1<-runif(10, -30, -10)
y1<-runif(10, -10, 30)
x2<-runif(10, -5, 5)
y2<-runif(10, 10, 50)
x3<-runif(10, -50, 30)
y3<-runif(10, 31, 60)

d<-data.frame(min1=x1,max1=y1,min2=x2,max2=y2,min3=x3,max3=y3)
classic2sym(d,groupby="customize",minData=d[,c(1,3,5)],maxData=d[,c(2,4,6)])
classic2sym(d,groupby="customize",minData=d$min1,maxData=d$min2)

#example for build modal data
#for the first modal data proportion
a1 <- runif(10, 0,0.4) %>% round(digits = 1)
a2 <- runif(10, 0,0.4) %>% round(digits = 1)

#for the second modal data proportion
b1 <- runif(10, 0,0.4) %>% round(digits = 1)
b2 <- runif(10, 0,0.4) %>% round(digits = 1)

#for interval-valued data
c1 <- runif(10, 10, 20) %>% round(digits = 0)
c2 <- runif(10, -50, -10) %>% round(digits = 0)

#build simulated data
d <- data.frame(a1 = a1, a2 = a2, a3 = 1-(a1+a2),
               c1 = c1, c2 = c2,
               b1 = b1, b2 = b2, b3 = 1-(b1+b2))

#transformation
classic2sym(d, groupby = "customize",
           minData = d$c2,
           maxData = d$c1,
           modalData = list(1:3, 6:8))#two modal data

#extract the data
symObj<-classic2sym(iris)
symObj$intervalData      #interval data
symObj$rawData           #raw data
symObj$clusterResult     #cluster result
symObj$statisticsDF      #statistics

```

---

`cor`*Generic function for the correlation*

---

## Description

This function compute the symbolic correlation

## Usage

```
cor(x, ...)  
  
## Default S3 method:  
cor(  
  x,  
  y = NULL,  
  use = "everything",  
  method = c("pearson", "kendall", "spearman"),  
  ...  
)  
  
## S3 method for class 'symbolic_tbl'  
cor(x, ...)  
  
## S3 method for class 'symbolic_interval'  
cor(x, y, method = c("centers", "B", "BD", "BG"), ...)
```

## Arguments

<code>x</code>	First symbolic variables.
<code>...</code>	As in R <code>cor</code> function.
<code>y</code>	Second symbolic variables.
<code>use</code>	an optional character string giving a method for computing correlation in the presence of missing values. This must be (an abbreviation of) one of the strings 'everything', 'all.obs', 'complete.obs', 'na.or.complete', or 'pairwise.complete.obs'.
<code>method</code>	The method to be use.

## Value

Return a real number.

## Author(s)

Oldemar Rodriguez Rojas

## References

- Billard L. and Diday E. (2006). Symbolic data analysis: Conceptual statistics and data mining. Wiley, Chichester.
- Rodriguez, O. (2000). Classification et Modeles Lineaires en Analyse des Donnees Symboliques. Ph.D. Thesis, Paris IX-Dauphine University.

---

cov

*Generic function for the covariance*

---

## Description

This function compute the symbolic covariance.

## Usage

```
cov(x, ...)
```

```
## Default S3 method:
cov(
  x,
  y = NULL,
  use = "everything",
  method = c("pearson", "kendall", "spearman"),
  ...
)
```

```
## S3 method for class 'symbolic_tbl'
cov(x, ...)
```

```
## S3 method for class 'symbolic_interval'
cov(x, y = NULL, method = c("centers", "B", "BD", "BG"), na.rm = FALSE, ...)
```

## Arguments

x	First symbolic variables.
...	As in R cov function.
y	Second symbolic variables.
use	an optional character string giving a method for computing covariances in the presence of missing values. This must be (an abbreviation of) one of the strings 'everything', 'all.obs', 'complete.obs', 'na.or.complete', or 'pairwise.complete.obs'.
method	The method to be use.
na.rm	As in R cov function.

## Value

Return a real number.

**Author(s)**

Oldemar Rodriguez Rojas

**References**

Billard L. and Diday E. (2006). Symbolic data analysis: Conceptual statistics and data mining. Wiley, Chichester.

Rodriguez, O. (2000). Classification et Modeles Lineaires en Analyse des Donnees Symboliques. Ph.D. Thesis, Paris IX-Dauphine University.

---

Environment

*Environment data example*

---

**Description**

Environment interval and modal data example.

**Usage**

```
data(Environment)
```

**Format**

An object of class `symbolic_tbl` (inherits from `tbl_df`, `tbl`, `data.frame`) with 14 rows and 17 columns.

**Examples**

```
data(Environment)
ggInterval_radar(Environment,
  plotPartial = 2,
  showLegend = FALSE,
  base_circle = TRUE,
  base_lty = 2,
  addText = FALSE)
```

---

 facedata

 Face Data Example
 

---

**Description**

Symbolic data matrix with all the variables of interval type.

**Usage**

```
data('facedata')
```

**Format**

```
$I;AD;AD;$I;BC;BC;.....
```

```
HUS1;$I;168.86;172.84;$I;58.55;63.39;.....
HUS2;$I;169.85;175.03;$I;60.21;64.38;.....
HUS3;$I;168.76;175.15;$I;61.4;63.51;.....
INC1;$I;155.26;160.45;$I;53.15;60.21;.....
INC2;$I;156.26;161.31;$I;51.09;60.07;.....
INC3;$I;154.47;160.31;$I;55.08;59.03;.....
ISA1;$I;164;168;$I;55.01;60.03;.....
ISA2;$I;163;170;$I;54.04;59;.....
ISA3;$I;164.01;169.01;$I;55;59.01;.....
JPL1;$I;167.11;171.19;$I;61.03;65.01;.....
JPL2;$I;169.14;173.18;$I;60.07;65.07;.....
JPL3;$I;169.03;170.11;$I;59.01;65.01;.....
KHA1;$I;149.34;155.54;$I;54.15;59.14;.....
KHA2;$I;149.34;155.32;$I;52.04;58.22;.....
KHA3;$I;150.33;157.26;$I;52.09;60.21;.....
LOT1;$I;152.64;157.62;$I;51.35;56.22;.....
LOT2;$I;154.64;157.62;$I;52.24;56.32;.....
LOT3;$I;154.83;157.81;$I;50.36;55.23;.....
PHI1;$I;163.08;167.07;$I;66.03;68.07;.....
PHI2;$I;164;168.03;$I;65.03;68.12;.....
PHI3;$I;161.01;167;$I;64.07;69.01;.....
ROM1;$I;167.15;171.24;$I;64.07;68.07;.....
ROM2;$I;168.15;172.14;$I;63.13;68.07;.....
ROM3;$I;167.11;171.19;$I;63.13;68.03;.....
```

**Source**

<https://CRAN.R-project.org/package=RSDA>

## References

Billard L. and Diday E. (2006). Symbolic data analysis: Conceptual statistics and data mining. Wiley, Chichester.

---

ggESDA

*A symbolic object by R6 class for interval analysis and ggplot*

---

## Description

This is an object that will be used to make a ggplot object. A ggESDA object contains both classic data that user have and interval data which we transform. More over, some basic statistics from row data will also be recorded in this object, and the interval data which is from RSDA transformation will still contain RSDA properties.

## Public fields

`rawData` the data from user.

`statisticsDF` contains min max mean median dataframe for each group of symbolic data

`intervalData` interval data from RSDA type

`clusterResult` clustering result

## Methods

### Public methods:

- [ggESDA\\$new\(\)](#)
- [ggESDA\\$clone\(\)](#)

**Method** `new()`: initialize all data, check whether satisfy theirs form

*Usage:*

```
ggESDA$new(  
  rawData = NULL,  
  statisticsDF = NULL,  
  intervalData = NULL,  
  clusterResult = NULL  
)
```

**Method** `clone()`: The objects of this class are cloneable with this method.

*Usage:*

```
ggESDA$clone(deep = FALSE)
```

*Arguments:*

`deep` Whether to make a deep clone.

---

ggInterval_2Dhist	<i>visualize a 2-dimension histogram by symbolic data with ggplot package.</i>
-------------------	--

---

### Description

Visualize the two continuous variable distribution by dividing both the x axis and y axis into bins, and calculating the frequency of observation interval in each bin.

### Usage

```
ggInterval_2Dhist(data = NULL, mapping = aes(NULL)
, xBins = 14, yBins = 16, removeZero = FALSE,
addFreq = TRUE)
```

### Arguments

data	A ggESDA object. It can also be either RSDA object or classical data frame, which will be automatically convert to ggESDA data.
mapping	Set of aesthetic mappings created by aes() or aes_(). If specified and inherit.aes = TRUE (the default), it is combined with the default mapping at the top level of the plot. You must supply mapping if there is no plot mapping. It is the same as the mapping of ggplot2.
xBins	x axis bins, which mean how many partials x variable will be separate into.
yBins	y axis bins. It is the same as xBins.
removeZero	whether remove data whose frequency is equal to zero
addFreq	where add frequency text in each cells.

### Value

Return a ggplot2 object.

### Examples

```
ggInterval_2Dhist(oils, aes(x = GRA, y = FRE),
xBins = 5, yBins = 5)
```

---

ggInterval\_2DhistMatrix  
*2-Dimension histogram matrix*

---

### Description

Visualize the all continuous variable distribution by dividing both the x axis and y axis into bins, and calculating the frequency of observation interval in each bin. Eventually show it by a matrix plot. Note: this function will automatically filter out the discrete variables, and plot all continuous in input data, so it can not be necessary that give the particularly variables in aes such like (aes(x = x, y = y)). It isn't also recommended to deal with too many variables because the big O in calculating full matrix will be too large.

### Usage

```
ggInterval_2DhistMatrix(data = NULL, mapping = aes(NULL)
, xBins = 8, yBins=8, removeZero = FALSE,
addFreq = TRUE)
```

### Arguments

data	A ggESDA object. It can also be either RSDA object or classical data frame, which will be automatically convert to ggESDA data.
mapping	Set of aesthetic mappings created by aes() or aes_(). If specified and inherit. aes = TRUE (the default), it is combined with the default mapping at the top level of the plot. You must supply mapping if there is no plot mapping. It is the same as the mapping of ggplot2.
xBins	x axis bins, which mean how many bins x variable will be separate into
yBins	y axis bins. It is the same as xBins
removeZero	whether remove data whose frequency is equal to zero
addFreq	where add frequency text in each cells.

### Value

Return a plot with ggplot2 object

### Examples

```
ggInterval_2DhistMatrix(oils, xBins = 5, yBins = 5)
```

---

ggInterval\_3Dscatter    *3D scatter plot for interval data*

---

### Description

Visualize the three continuous variable distribution by collecting all vertices in each interval to form a shape of cube. Also show the difference between each group.

### Usage

```
ggInterval_3Dscatter(data = NULL, mapping = aes(NULL), scale=FALSE)
```

### Arguments

data	A ggSDA object. It can also be either RSDA object or classical data frame, which will be automatically convert to ggSDA data.
mapping	Set of aesthetic mappings created by aes() or aes_(). If specified and inherit.aes = TRUE (the default), it is combined with the default mapping at the top level of the plot. You must supply mapping if there is no plot mapping. It is the same as the mapping of ggplot2.
scale	A boolean variable, TRUE, standardize data. FALSE, not standardize. If variance is too large (or small) or the difference between two variables are too large, it will be distortion or unseeable, which may happen in different units or others. So, a standardize way is necessary.

### Value

Return a ggplot2 object (It will still be 2-Dimension).

### Examples

```
ggInterval_3Dscatter(facedata[1:5, ], aes(x = BC, y = EH, z = GH))
```

---

ggInterval\_boxplot    *A interval Box plot*

---

### Description

Visualize the one continuous variable distribution by box represented by multiple rectangles.

### Usage

```
ggInterval_boxplot(data = NULL, mapping = aes(NULL), plotAll=FALSE)
```

**Arguments**

data	A ggESDA object. It can also be either RSDA object or classical data frame, which will be automatically convert to ggESDA data.
mapping	Set of aesthetic mappings created by aes() or aes_(). If specified and inherit.aes = TRUE (the default), it is combined with the default mapping at the top level of the plot. You must supply mapping if there is no plot mapping. It is the same as the mapping of ggplot2.
plotAll	booleans, if TRUE, plot all variable together

**Value**

Return a ggplot2 object.

**Examples**

```
p<-ggInterval_boxplot(iris,aes(iris$Petal.Length))
p
p+scale_fill_manual(values = c("red","yellow",
  "green","blue","black"),
  labels=c("0%","25%","50%","75%","100%"),
  name="quantile")

mydata<-ggESDA::facedata
ggInterval_boxplot(mydata,aes(AD,col="black",alpha=0.5))

myMtcars<-classic2sym(mtcars)
myMtcars<-myMtcars$intervalData
ggInterval_boxplot(myMtcars,aes(dis))
```

---

ggInterval\_centerRange

*Figure with x-axis = center y-axis = range*

---

**Description**

Visualize the relation between center and range.

**Usage**

```
ggInterval_centerRange(data = NULL,mapping = aes(NULL),plotAll=FALSE)
```

**Arguments**

data	A ggESDA object. It can also be either RSDA object or classical data frame,which will be automatically convert to ggESDA data.
------	--

mapping	Set of aesthetic mappings created by aes() or aes_(). If specified and inherit. aes = TRUE (the default), it is combined with the default mapping at the top level of the plot. You must supply mapping if there is no plot mapping. It is the same as the mapping of ggplot2.
plotAll	booleans, if TRUE, plot all variable together

**Value**

Return a ggplot2 object.

**Examples**

```
ggInterval_centerRange(iris,aes(iris$Sepal.Length))

mydata<-ggESDA::facedata
ggInterval_centerRange(mydata,aes(AD,col="blue",pch=2))
```

---

ggInterval_hist	<i>Histogram for symbolic data with equal-bin or unequal-bin.</i>
-----------------	---

---

**Description**

Visualize the continuous variable distribution by dividing the x axis into bins,and calculating the frequency of observation interval in each bin.

**Usage**

```
ggInterval_hist(data = NULL,mapping = aes(NULL),method="equal-bin",bins=10,
  plotAll = FALSE, position = "identity", alpha = 0.5)
```

**Arguments**

data	A ggESDA object.It can also be either RSDA object or classical data frame,which will be automatically convert to ggESDA data.
mapping	Set of aesthetic mappings created by aes() or aes_(). If specified and inherit. aes = TRUE (the default), it is combined with the default mapping at the top level of the plot. You must supply mapping if there is no plot mapping. It is the same as the mapping of ggplot2.
method	It can be equal-bin(default) or unequal-bin.Enqual-bin means the width in histogram is equal, which represent all intervals divided have the same range. unequal-bin means the range of intervals are not the same,and it can be more general on data. Thus, the bins of unequal-bin method depends on the data, and the argument "bins" will be unused.
bins	x axis bins,which mean how many partials the variable
plotAll	boolean, whether plot all variables, default FALSE. will be separate into.
position	"stack" or "identity"
alpha	fill alpha

**Value**

Return a ggplot2 object.

**Examples**

```
ggInterval_hist(mtcars, aes(x=wt))

ggInterval_hist(iris, aes(iris$Petal.Length, col="blue", alpha=0.2,
  fill="red"), bins=30)

d<-data.frame(x=rnorm(1000, 0, 1))
p<-ggInterval_hist(d, aes(x=x), bins=40, method="equal-bin")$plot
p

p+scale_fill_manual(values=rainbow(40))+labs(title="myNorm")
```

---

ggInterval_index	<i>Plot the range of each observations</i>
------------------	--

---

**Description**

Visualize the range of the variables of each observations by using a kind of margin bar that indicate the minimal and maximal of observations.

**Usage**

```
ggInterval_index(data = NULL, mapping = aes(NULL),
  plotAll = FALSE)
```

**Arguments**

data	A ggESDA object. It can also be either RSDA object or classical data frame, which will be automatically convert to ggESDA data.
mapping	Set of aesthetic mappings created by aes() or aes_(). If specified and inherit. aes = TRUE (the default), it is combined with the default mapping at the top level of the plot. You must supply mapping if there is no plot mapping. It is the same as the mapping of ggplot2.
plotAll	plot all variables

**Value**

Return a ggplot2 object.

**Examples**

```
#the observations show on the y-axis .values on x-axis
ggInterval_index(iris,aes(x=iris$Sepal.Length))

#change above axis
ggInterval_index(mtcars,aes(y=disp,col="red",fill="grey"))

#symbolic data
mydata <- ggESDA::facedata
ggInterval_index(mydata,aes(x=3:13,y=AD))
```

---

ggInterval\_indexImage *An index plot presented by color image for interval data.*

---

**Description**

Visualize the range of the variables of each observations by using color image.The index image replace margin bar by color,thus it will be more visible for data.

**Usage**

```
ggInterval_indexImage(data = NULL,mapping = aes(NULL),
column_condition=TRUE,full_strip=FALSE, plotAll = FALSE)
```

**Arguments**

data	A ggESDA object. It can also be either RSDA object or classical data frame,which will be automatically convert to ggESDA data.
mapping	Set of aesthetic mappings created by aes() or aes_(). If specified and inherit. aes = TRUE (the default), it is combined with the default mapping at the top level of the plot. You must supply mapping if there is no plot mapping.
column_condition	Boolean variables, which mean the color present by column condition (if TRUE) or matrix condition (if FALSE)
full_strip	Boolean variables, which mean the strip present in full figure-width (if TRUE) or only in its variable values(if FALSE).
plotAll	Boolean, which determine if the heatmap type for visualizing full variables is used. default FALSE.

**Value**

Return a ggplot2 object.

**Examples**

```
d<-data.frame(qq=rnorm(1000,0,1))
ggInterval_indexImage(d,aes(qq))

mydata<-ggESDA::facedata
p<-ggInterval_indexImage(mydata,aes(AD),full_strip=TRUE,column_condition = TRUE)
#Recommend to add coord_flip() to make the plot more visible
p+coord_flip()

myIris<-classic2sym(iris,groupby="Species")
myIris<-myIris$intervalData
p<-ggInterval_indexImage(myIris,aes(myIris$Petal.Length),full_strip=FALSE,column_condition=TRUE)
p

ggInterval_indexImage(mtcars,aes(dis)))+labs(x="anything")
```

---

ggInterval\_minmax      *A min-max plot for interval data*

---

**Description**

Visualize the range of the variables of each observations by marking minimal and maximal point.

**Usage**

```
ggInterval_minmax(data = NULL,mapping = aes(NULL),
                  scaleXY = "local",plotAll=FALSE)
```

**Arguments**

data	A ggESDA object. It can also be either RSDA object or classical data frame,which will be automatically convert to ggESDA data.
mapping	Set of aesthetic mappings created by aes() or aes_(). If specified and inherit. aes = TRUE (the default), it is combined with the default mapping at the top level of the plot. You must supply mapping if there is no plot mapping.
scaleXY	default "local", which means limits of x-axis and y-axis depend on their own variable. "global" means limits of them depend on all variables that user input.
plotAll	booleans, if TRUE, plot all variable together

**Value**

Return a ggplot2 object.

**Examples**

```
ggInterval_minmax(mtcars,aes(disp))

mydata2<-ggESDA::Cardiological
ggInterval_minmax(mydata2,aes(mydata2$Pulse,size=3))

d<-mapply(c(10,20,40,80,160),c(20,40,80,160,320),FUN=runif,n=1000)
d<-data.frame(qq=matrix(d,ncol=1))
ggInterval_minmax(d,aes(qq))

myIris<-classic2sym(iris,groupby="Species")
myIris<-myIris$intervalData
ggInterval_minmax(myIris,aes(myIris$Petal.Length))+
  theme_classic()
```

---

ggInterval_PCA	<i>Vertice-PCA for interval data</i>
----------------	--------------------------------------

---

**Description**

ggInterval\_PCA performs a principal components analysis on the given numeric interval data and returns the results like princomp, ggplot object and a interval scores.

**Usage**

```
ggInterval_PCA(data = NULL,mapping = aes(NULL),plot=TRUE,
               concepts_group=NULL, poly = FALSE, adjust = TRUE)
```

**Arguments**

data	A ggESDA object. It can also be either RSDA object or classical data frame, which will be automatically convert to ggESDA data.
mapping	Set of aesthetic mappings created by aes() or aes_(). If specified and inherit. aes = TRUE (the default), it is combined with the default mapping at the top level of the plot. You must supply mapping if there is no plot mapping. It is the same as the mapping of ggplot2.
plot	Boolean variable,Auto plot (if TRUE).It can also plot by its inner object
concepts_group	color with each group of concept
poly	if plot a poly result
adjust	adjust sign of the principal component

**Value**

A ggplot object for PC1,PC2,and a interval scores and others.

- scores\_interval - The interval scores after PCA.
- ggplotPCA - a ggplot object with x-axis and y-axis are PC1 and PC2.
- others - others are the returns values of princomp.

**Examples**

```

ggInterval_PCA(iris)

mydata2<-ggESDA::Cardiological
ggInterval_PCA(mydata2,aes(col="red",alpha=0.2))

d<-mapply(c(10,20,40,80,160),c(20,40,80,160,320),FUN=runif,n=1000)
d<-data.frame(qq=matrix(d,ncol=4))
ggInterval_PCA(d)

myIris<-classic2sym(iris,"Species")
p<-ggInterval_PCA(myIris,plot=FALSE)
p$ggplotPCA
p$scores_interval

```

---

ggInterval\_radar      *A interval Radar plot*

---

**Description**

Using ggplot2 package to make a radar plot with multiple variables.Each variables contains min values and max values as a symbolic data.

**Usage**

```

ggInterval_radar(data=NULL,layerNumber=3,
inOneFig=TRUE,showLegend=TRUE,showXYLabs=FALSE,
plotPartial=NULL,
alpha=0.5,
base_circle=TRUE,
base_lty=2,
addText=TRUE,
type="default",
quantileNum=4,
Drift=0.5,
addText_modal=TRUE,
addText_modal.p=FALSE)

```

**Arguments**

data	A ggESDA object. It can also be either RSDA object or classical data frame(not recommended),which will be automatically convert to ggESDA data.
layerNumber	number of layer of a concentric circle,usually to visuallize the reach of a obser- vation in particularly variable.
inOneFig	whether plot all observations in one figure.if not, it will generate a new windows containing distinct observations.

<code>showLegend</code>	whether show the legend.
<code>showXYLabs</code>	whether show the x,y axis labels.
<code>plotPartial</code>	a numeric vector,which is the row index from the data.if it is not null, it will extract the row user deciding to draw a radar plot from original data.Notes : the data must be an interval data if the plotPartial is not null.
<code>alpha</code>	aesthetic alpha of fill color
<code>base_circle</code>	boolean, if true, it will generate inner circle.
<code>base_lty</code>	line type in base figure
<code>addText</code>	add the value of interval-valued variables in figure
<code>type</code>	different type of radar,it can be "default","rect","quantile"
<code>quantileNum</code>	if type == "quantile", it will provide the number of percentage
<code>Drift</code>	The drift term, which determines the radar values beginning.
<code>addText_modal</code>	add the factor of modal multi-valued variables in figure..
<code>addText_modal.p</code>	add the value of modal multi-valued variables in figure..

## Examples

```
# must specify plotPartial to some certain rows you want to plot
Environment.n <- Environment[, 5:17]
ggInterval_radar(Environment.n,
                  plotPartial = 2,
                  showLegend = FALSE,
                  base_circle = TRUE,
                  base_lty = 2,
                  addText = FALSE
) +
  labs(title = "") +
  scale_fill_manual(values = c("gray50")) +
  scale_color_manual(values = c("red"))

ggInterval_radar(Environment,
                  plotPartial = 2,
                  showLegend = FALSE,
                  base_circle = FALSE,
                  base_lty = 1,
                  addText = TRUE
) +
  labs(title = "") +
  scale_fill_manual(values = c("gray50")) +
  scale_color_manual(values = c("gray50"))
```

---

ggInterval\_scaMatrix *scatter plot for all variable by interval data.*

---

### Description

Visualize the all continuous variable distribution by rectangle for both x-axis and y-axis with a matrix grid. Note: this function will automatically filter out the discrete variables, and plot all continuous in input data, so it can not be necessary that give the particularly variables in aes such like (aes(x = x, y = y)). It isn't also recommended to deal with too many variables because the big O in calculating full matrix will be too large.

### Usage

```
ggInterval_scaMatrix(data = NULL, mapping = aes(NULL), showLegend=FALSE)
```

### Arguments

data	A ggESDA object. It can also be either RSDA object or classical data frame, which will be automatically convert to ggESDA data.
mapping	Set of aesthetic mappings created by aes() or aes_(). If specified and inherit.aes = TRUE (the default), it is combined with the default mapping at the top level of the plot. You must supply mapping if there is no plot mapping.
showLegend	whether show the legend.

### Value

Return a plot with no longer a ggplot2 object, instead of a arrangeGrob object.

### Examples

```
a<-rnorm(1000,0,5)
b<-runif(1000,-20,-10)
c<-rgamma(1000,10,5)
d<-as.data.frame(cbind(norm=a,unif=b,gamma_10_5=c))
ggInterval_scaMatrix(d)
```

```
ggInterval_scaMatrix(mtcars[,c("mpg", "wt", "qsec")],
  aes(col="red", lty=2, fill="blue", alpha=0.3))
```

```
myIris <- classic2sym(iris, groupby = "Species")$intervalData
ggInterval_scaMatrix(myIris[,1:3])
```

```
mydata <- ggESDA::Cardiological
ggInterval_scaMatrix(mydata[,1:3], aes(fill="black", alpha=0.2))
```

---

ggInterval\_scatter     *scatter plot for two continuous interval data*

---

### Description

Visualize the two continuous variable distribution by rectangle and each of its width and height represents a interval of the data.

### Usage

```
ggInterval_scatter(data = NULL, mapping = aes(NULL), ...)
```

### Arguments

data	A ggESDA object. It can also be either RSDA object or classical data frame, which will be automatically convert to ggESDA data.
mapping	Set of aesthetic mappings created by aes() or aes_(). If specified and inherit.aes = TRUE (the default), it is combined with the default mapping at the top level of the plot. You must supply mapping if there is no plot mapping.
...	Others in ggplot2.

### Value

Return a ggplot2 object.

### Examples

```
a<-rnorm(1000,0,5)
b<-runif(1000,-20,-10)
d<-as.data.frame(cbind(norm=a,unif=b))
ggInterval_scatter(d,aes(a,b))
```

```
ggInterval_scatter(mtcars[,c("mpg","wt","qsec")],
  aes(x=mpg,y=wt,
    col="red",lty=2,fill="blue",alpha=0.3))
```

```
myIris <- classic2sym(iris,groupby = "Species")$intervalData
p<-ggInterval_scatter(myIris,aes(myIris$Petal.Length,myIris$Petal.Width))
p
p+scale_fill_manual(labels=rownames(myIris),
  values=c("red","blue","green"),
  name="Group")
```

```
mydata <- ggESDA::facedata
p<-ggInterval_scatter(mydata[1:10,],aes(AD,BC,alpha=0.2))
```

```
p+scale_fill_manual(labels=rownames(mydata)[1:10],
                    values=rainbow(10),
                    name="Group")
```

---

iris.i	<i>iris.i data example</i>
--------	----------------------------

---

**Description**

iris.i interval data example.

**Usage**

```
data(iris.i)
```

**Format**

An object of class `data.frame` (inherits from `symbolic_tbl`) with 3 rows and 4 columns.

**Examples**

```
data(iris.i)
ggInterval_index(iris.i, aes(x = Sepal.Length))
```

---

mtcars.i	<i>mtcars.i data example</i>
----------	------------------------------

---

**Description**

mtcars.i interval and modal data example.

**Usage**

```
data(mtcars.i)
```

**Format**

An object of class `symbolic_tbl` (inherits from `tbl_df`, `tbl`, `data.frame`, `symbolic_tbl`) with 5 rows and 11 columns.

**Examples**

```
data(mtcars.i)
ggInterval_index(mtcars.i, aes(x = mpg))
```

---

mushroom	<i>mushroom data example</i>
----------	------------------------------

---

**Description**

mushroom interval data example.

**Usage**

```
data(mushroom)
```

**Format**

An object of class `tbl_df` (inherits from `tbl`, `data.frame`, `symbolic_tbl`) with 23 rows and 3 columns.

**References**

Billard L. and Diday E. (2006). Symbolic data analysis: Conceptual statistics and data mining. Wiley, Chichester.

**Examples**

```
data(mushroom)
ggInterval_scatter(mushroom, aes(x = Cap.Widths, y = Stipe.Lengths))
```

---

oils	<i>oils data example</i>
------	--------------------------

---

**Description**

oils interval data example.

**Usage**

```
data(oils)
```

**Format**

An object of class `symbolic_tbl` (inherits from `tbl_df`, `tbl`, `data.frame`) with 8 rows and 4 columns.

**References**

Billard L. and Diday E. (2006). Symbolic data analysis: Conceptual statistics and data mining. Wiley, Chichester.

**Examples**

```
data(oils)
ggInterval_scatter(oils, aes(x = GRA, y = IOD))
```

---

RSDA2sym

*RSDA object to symbolic object for ggplot*


---

**Description**

It will be a good way to unify all symbolic data object in R that collects all useful symbolic analysis tools such like RSDA into the same class for management. In this way, user who wants to do some study in symbolic data will be more convenient for searching packages. Thus, RSDA2sym collecting RSDA object into ggESDA object will do for plot(ggplot) and RSDA's analysis.

**Usage**

```
RSDA2sym(data=NULL, rawData=NULL)
```

**Arguments**

data	an interval data, which may transform by <code>RSDA::classic.to.sym</code> . Note: data is a necessary parameter, and must have <code>symbolic_tbl</code> class.
rawData	rawData, which can be transformed to interval data, must be a data frame and match to data.

**Value**

Return an object of class "ggESDA", which have a interval data and others as follows.

- intervalData - The Interval data after converting also known as a RSDA object.
- rawData - Classical data that user input.
- clusterResult - Cluster results .If the groupby method is a clustering method then it will exist.
- statisticsDF - A list contains data frame including some typically statistics in each group.

```
#'
```

**Examples**

```
r<-ggESDA::Cardiological
mySym<-RSDA2sym(r)
mySym$intervalData
```

---

scale	<i>scale for symbolic data table</i>
-------	--------------------------------------

---

**Description**

scale for symbolic data table

**Usage**

```
scale(x, ...)
```

```
## Default S3 method:
scale(x, center = TRUE, scale = TRUE, ...)
```

```
## S3 method for class 'symbolic_tbl'
scale(x, ...)
```

```
## S3 method for class 'symbolic_interval'
scale(x, ...)
```

**Arguments**

x	A ggESDA object. It can also be either RSDA object or classical data frame, which will be automatically convert to ggESDA data.
...	Used by other R function.
center	same as base::scale, either a logical value or numeric-alike vector of length equal to the number of columns of x, where ‘numeric-alike’ means that as.numeric(.) will be applied successfully if is.numeric(.) is not true.
scale	same as base::scale, either a logical value or a numeric-alike vector of length equal to the number of columns of x.

**Value**

Return a scale ggESDA object.

**Examples**

```
#For all interval-valued
scale(facedata)
```

```
#For both interval-valued and modal multi-valued
scale(mtcars.i)
```

---

summary	<i>summary for symbolic data table</i>
---------	--

---

## Description

summary for symbolic data table

## Usage

```
summary(object, ...)  
  
## Default S3 method:  
summary(object, ...)  
  
## S3 method for class 'symbolic_tbl'  
summary(object, ...)  
  
## S3 method for class 'symbolic_interval'  
summary(object, ...)  
  
## S3 method for class 'symbolic_modal'  
summary(object, summary_fun = "mean", ...)
```

## Arguments

object	an object for which a summary is desired.
...	additional arguments affecting the summary produced.
summary_fun	only works when the symbolic_modal class input, it determine which summary function be applied for each modal.

## Value

Return a summary table.

## Examples

```
#For all interval-valued  
summary(facedata)  
  
#For both interval-valued and modal multi-valued  
summary(Environment)  
  
summary(Environment$URBANICITY, summary_fun = "quantile")
```

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