

# Package ‘SCOUTer’

June 29, 2020

**Title** Simulate Controlled Outliers

**Version** 1.0.0

**Description** Using Principal Component Analysis as a base model, 'SCOUTer' offers a new approach to simulate outliers in a simple and precise way. The user can generate new observations defining them by a pair of well-known statistics: the Squared Prediction Error and the Hotelling's  $T^2$  statistics. Just by introducing the target values of the SPE and  $T^2$ , 'SCOUTer' returns a new set of observations with the desired target properties. Authors: Alba González, Abel Folch-Fortuny, Francisco Arteaga and Alberto Ferrer (2020).

**License** GPL-3

**Encoding** UTF-8

**LazyData** true

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**BugReports** <https://github.com/albagc/SCOUTerRpack>

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ggplot2,  
ggpubr,  
stats

**Suggests** knitr,  
rmarkdown

**VignetteBuilder** knitr

## R topics documented:

barwithucl . . . . .	2
custombar . . . . .	3
distplot . . . . .	3
distplotsimple . . . . .	4
dotag . . . . .	5
dscplot . . . . .	5
ht2info . . . . .	6
obscontribpanel . . . . .	7
pcamb_classic . . . . .	7
pcame . . . . .	8

scoreplot . . . . .	9
scoreplotsimple . . . . .	9
scout . . . . .	10
scoutgrid . . . . .	11
scoutsimple . . . . .	12
scoutsteps . . . . .	13
speinfo . . . . .	14
X . . . . .	14
xshift . . . . .	15
<b>Index</b>	<b>16</b>

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barwithucl	<i>barwithucl</i>
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**Description**

Single bar plot with Upper Control Limis. Customized title and labels. Y-Axis limits are fixed according to the range of the values in x.

**Usage**

```
barwithucl(  
  x,  
  iobs,  
  ucl,  
  plotname = "",  
  ylabelname = "",  
  xlabelname = "Obs. Index"  
)
```

**Arguments**

- x                    vector with the values of the statistic.
- iobs                index of the observations whose value will be displayed.
- ucl                 Upper Control Limit of the statistic.
- plotname           string with the title of the plot. Set to "" by default.
- ylabelname         string with the y-axis label. Set to "" by default.
- xlabelname         string with the y-axis label. Set to "Obs. Index" by default.

**Value**

ggplot object with the individual value of a variable as a geom\_col with an horizontal line reference.

---

`custombar`*custombar*

---

**Description**

Bar plot with customized title and labels. Y-Axis limits are fixed according to the range of the values in X.

**Usage**

```
custombar(  
  X,  
  iobs,  
  plotname = "",  
  ylabelname = "Contribution",  
  xlabelname = "ggplot2:"  
)
```

**Arguments**

X	matrix with observations as row vectors.
iobs	index of the observations whose value will be displayed.
plotname	string with the title of the plot. Set to "" by default.
ylabelname	string with the y-axis label. Set to "Contribution" by default.
xlabelname	string with the y-axis label. Set to "Variables" by default.

**Value**

ggplot object with the values of a vector with a customized geom\_col layer.

---

`distplot`*distplot*

---

**Description**

Returns the distance plot providing a dataset and a pca model.

**Usage**

```
distplot(  
  X,  
  pcaref,  
  obstag = matrix(0, nrow(X), 1),  
  plottitle = "Distance plot\n"  
)
```

**Arguments**

<code>X</code>	data matrix with observations to be displayed in the distance plot.
<code>pcaref</code>	list with the information of the PCA model.
<code>obstag</code>	Optional column vector of integers indicating the group of each observation (0 or 1). Default value set to <code>matrix(0, nrow(X), 1)</code> .
<code>plottitle</code>	Optional string with the plot title. Set to "Distance plot" by default.

**Details**

Coordinates are expressed in terms of the Hotelling's  $T^2$  (x-axis) and the Squared Prediction Error (y-axis) obtained projecting  $X$  on the provided `pca` model. Observations can be identified by the `obstag` input argument.

**Value**

ggplot object with the distance plot.

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<code>distplotsimple</code>	<i>displotsimple</i>
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**Description**

Returns the distance plot directly providing the coordinates and Upper Control Limits.

**Usage**

```
distplotsimple(
  T2,
  SPE,
  lim.t2,
  lim.spe,
  ncomp,
  obstag = matrix(0, length(T2), 1),
  alpha = 0.05,
  plottitle = "Distance plot\n"
)
```

**Arguments**

<code>T2</code>	Vector with the Hotelling's $T^2$ values for each observation.
<code>SPE</code>	Vector with the SPE values for each observation.
<code>lim.t2</code>	Value of the Upper Control Limit for the $T^2$ statistic.
<code>lim.spe</code>	Value of the Upper Control Limit for the SPE.
<code>ncomp</code>	An integer indicating the number of PCs.
<code>obstag</code>	Optional column vector of integers indicating the group of each observation (0 or 1). Default value set to <code>matrix(0, nrow(X), 1)</code> .
<code>alpha</code>	Optional number between 0 and 1 expressing the type I risk assumed in the computation of the Upper Control Limits set to 0.05 (5 %) by default.
<code>plottitle</code>	Optional string with the plot title, "Distance plot" by default.

**Details**

Coordinates are expressed in terms of the Hotelling's  $T^2$  (x-axis) and the Squared Prediction Error (y-axis) obtained projecting X on the provided pca model. Observations can be identified by the obstag input argument.

**Value**

distplotobj ggplot object with the generated distance plot.

---

dotag	<i>dotag</i>
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**Description**

Returns the tag vector to identify two different data sets

**Usage**

```
dotag(X.zeros = NA, X.ones = NA)
```

**Arguments**

X.zeros	Matrix with the tag 0.
X.ones	Matrix with the tag 1.

**Value**

tag.all vector with 0-tags for observations in X.zeros and 1-tags for observations in X.ones.

---

dscplot	<i>dscplot</i>
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**Description**

Returns the distance plot and the score plot providing a dataset and a pca model. Observations can be identified by the obstag input argument.

**Usage**

```
dscplot(
  X,
  pcamodel,
  obstag = matrix(0, nrow(X), 1),
  pcx = 1,
  pcy = 2,
  alpha = 0.05,
  nrow = 1,
  ncol = 2,
  legpos = "bottom"
)
```

**Arguments**

<code>X</code>	Matrix with the data to be displayed.
<code>pcamodel</code>	List with the PCA model elements.
<code>obstag</code>	Optional column vector of integers indicating the group of each observation (0 or 1). Default value set to <code>matrix(0, nrow(X), 1)</code> .
<code>pcx</code>	Optional integer with the number of the PC in the horizontal axis. Set to 1 by default.
<code>pcy</code>	Optional integer with the number of the PC in the vertical axis. Set to 2 by default.
<code>alpha</code>	Optional number between 0 and 1 expressing the type I risk assumed in the computation of the confidence ellipse, set to 0.05 (5 %) by default.
<code>nrow</code>	Optional number of rows the plot layout. Set to 1 by default.
<code>ncol</code>	Optional number of columns the plot layout. Set to 2 by default.
<code>legpos</code>	Optional string with the position of the legend. Set to "bottom" by default.

**Value**

ggplot object with the generated score plot.

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ht2info	<i>ht2info</i>
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**Description**

Returns information about  $T^2$  statistic for an observation. Two subplots show the information of an observation regarding its  $T^2$  statistic, i.e.: a bar plot indicating the value of the statistic for the observation, and a bar plot with the contribution that each component had for the  $T^2$  value

**Usage**

```
ht2info(HT2, T2matrix, limht2, iobs = NA)
```

**Arguments**

<code>HT2</code>	A vector with values of the Hotelling's $T^2_A$ statistic.
<code>T2matrix</code>	A matrix with the contributions of each PC (A columns) for each observation (rows) to the Hotelling's $T^2_A$ statistic.
<code>limht2</code>	Upper Control Limit for the Hotelling's $T^2_A$ statistic, at a certain confidence level $(1-\alpha)*100$ %.
<code>iobs</code>	Integer with the index of the observation of interest. Default value set to NA.

**Value**

ggplot object with the generated bar plots.

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obscontribpanel	<i>obscontribpanel</i>
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### Description

Information about T<sup>2</sup> and SPE statistics of an observation.

### Usage

```
obscontribpanel(pcax, pcaref, obsid = NA)
```

### Arguments

pcax	A list with the elements of the PCA model that will be displayed: SPE, T <sup>2</sup> _A and their contributions (E and T2matrix).
pcaref	A list with the PCA model according to which the distance and contributions are expressed.
obsid	Integer with the index of the observation of interest. Default set to NA.

### Value

ggplot object with the generated bar plots in a 1 x 4 subplots layout.

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pcamb_classic	<i>pcamb_classic</i>
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### Description

PCA model fitting according to a matrix X using svd.

### Usage

```
pcamb_classic(X, ncomp, alpha, prepro)
```

### Arguments

X	Matrix with observations that will used to fit the PCA model.
ncomp	An integer indicating the number of PCs that the model will have.
alpha	A number between 0 and 1 indicating the type I risk assumed to calculate the Upper Control Limits for the SPE, the T <sup>2</sup> _A and the scores. The confidence level of these limits will be (1-alpha)*100.
prepro	A string indicating the preprocessing to be performed on X. Its possible values are: "none", for any preprocessing, "cent", for a mean-centering, or "autosc", for a mean-centering and unitary variance scaling (autoscaling).

**Value**

list with elements containing information about PCA model: *m* (mean vector), *s* (standard deviation vector), *P* (loading matrix with the loadings of each PC stored as columns), *Pfull* (full loading matrix obtained by the *svd*), *lambda* (vector with the variance of each PC), *limspe* (Upper Control Limit for the SPE with a confidence level  $(1-\alpha)*100\%$ ), *limt2* (Upper Control Limit for the  $T^2_A$  with a confidence level  $(1-\alpha)*100\%$ ), *limits\_t* (Upper control Limits for the scores with a confidence level  $(1-\alpha)*100\%$ ), *prepro* (string indicating the type of preprocessing performed on *X*), *ncomp* (number of PCs of the PCA model, *A*), *alpha* (value of the type I risk assumed to calculate the Upper Control Limits of the SPE,  $T^2_A$  and scores), *n* (dimension of the number of rows in *X*), *S* (covariance matrix of *X*).

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pcame	<i>pcame</i>
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**Description**

Projection of *X* onto a PCA model.

**Usage**

```
pcame(X, pcaref)
```

**Arguments**

<i>X</i>	Matrix with observations that will be projected onto the PCA model.
<i>pcaref</i>	A list with the elements of a PCA model: <i>m</i> (mean), <i>s</i> (standard deviation), <i>prepro</i> (preprocessing: "none", "cent" or "autosc"), <i>P</i> (loading matrix), <i>lambda</i> (vector with variances of each PC).

**Details**

*pcame* performs the projection of the data in *X* onto the PCA model stored as a list of parameters. It returns the projection of the observations in *X*, along with the SPE, Hotelling's  $T^2_A$ , contribution elements and the reconstruction of *X* obtained by the PCA model.

**Value**

list with elements containing information about *X* in the PCA model: *Xpreprocessed* (matrix *X* preprocessed), *Tscores* (score matrix with the projection of *X* on each one of the *A* PCs), *E* (error matrix with the part of *X* not explained by the PCA model), *SPE* (vector with the SPE for each observation of *X*), *T2* (vector with the  $T^2_A$  for each observation of *X*), *T2matrix* (matrix with the contributions of each PC to the  $T^2_A$  for each observation of *X*) and *Xrec* (matrix with the reconstructed part of *X*, i.e. the part of *X* explained by the PCA model).



scoreplot

*scoreplot***Description**

Returns the score plot providing a dataset and a pca model. Observations can be identified by the obstag input argument.

**Usage**

```
scoreplot(
  X,
  pcamodel,
  obstag = matrix(0, nrow(X), 1),
  pcx = 1,
  pcy = 2,
  alpha = 0.05,
  plottitle = "Score plot\n"
)
```

**Arguments**

X	Matrix with the data to be displayed.
pcamodel	List wiht the PCA model elements.
obstag	Optional column vector of integers indicating the group of each observation (0 or 1). Default value set to <code>matrix(0, nrow(X), 1)</code> .
pcx	Optional integer with the number of the PC in the horizontal axis. Set to 1 by default.
pcy	Optional integer with the number of the PC in the vertical axis. Set to 2 by default.
alpha	Optional number between 0 and 1 expressing the type I risk assumed in the computation of the confidence ellipse, set to 0.05 (5 %) by default.
plottitle	Optional string with the plot title. Set to "Score plot" by default.

**Value**

ggplot object with the generated score plot.

scoreplotsimple

*scoreplotsimple***Description**

Returns the score plot providing the scores matrix, **T**. Observations can be identified by the obstag input argument.

**Usage**

```
scoreplotsimple(
  Tscores,
  pcx = 1,
  pcy = 2,
  obstag = matrix(0, nrow(Tscores), 1),
  alpha = 0.05,
  varT = stats::var(Tscores),
  plottitle = "Score plot\n"
)
```

**Arguments**

Tscores	Matrix with the scores to be displayed, with the information of each PC stored by columns.
pcx	Optional integer with the number of the PC in the horizontal axis. Set to 1 by default.
pcy	Optional integer with the number of the PC in the vertical axis. Set to 2 by default.
obstag	Optional column vector of integers indicating the group of each observation (0 or 1). Default value set to <code>matrix(0, nrow(X), 1)</code> .
alpha	Optional number between 0 and 1 expressing the type I risk assumed in the computation of the confidence ellipse, set to 0.05 (5 %) by default.
varT	Optional parameter expressing the variance of each PC. Set to <code>var(Tscores)</code> by default.
plottitle	Optional string with the plot title. Set to "Score plot" by default.

**Value**

ggplot object with the generated score plot.

---

scout	<i>scout</i>
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---

**Description**

Shift of an observation following a selected pattern.

**Usage**

```
scout(
  X,
  pcaref,
  T2.y = NA,
  SPE.y = NA,
  nsteps = 1,
  nsteps.spe = 1,
  nsteps.t2 = 1,
  gspe = 1,
```

```

    gt2 = 1,
    mode = "simple"
)

```

### Arguments

<code>X</code>	Matrix with observations that will be shifted as rows.
<code>pcaref</code>	List with the elements of a PCA model: <code>m</code> (mean), <code>s</code> (standard deviation), <code>prepro</code> (preprocessing: "none", "cent" or "autosc"), <code>P</code> (loading matrix), <code>lambda</code> (vector with variances of each PC).
<code>T2.y</code>	A number indicating the target value for the $T^2_A$ after the shift. Set to NA by default.
<code>SPE.y</code>	A number indicating the target value for the SPE after the shift. Set to NA by default.
<code>nsteps</code>	A number indicating the number of steps between the reference and target values of the SPE and the $T^2$ . Set to 1 by default.
<code>nsteps.spe</code>	An integer indicating the number of steps in which the shift from the reference to the target value of the SPE will be performed. Set to 1 by default
<code>nsteps.t2</code>	An integer indicating the number of steps in which the shift from the reference to the target value of the $T^2_A$ will be performed. Set to 1 by default
<code>gspe</code>	A number indicating the term that will tune the spacing between steps for the SPE. Set to 1 by default (linear spacing).
<code>gt2</code>	A number indicating the term that will tune the spacing between steps for the $T^2$ . Set to 1 by default (linear spacing).
<code>mode</code>	A character indicating the type of shift that will be performed: "simple", "steps" or "grid".

### Value

list with elements: `X`, matrix with the new and shifted data, `SPE` and `T2` vectors with the statistic values of each one of the new generated outliers or observations, elements `step.spe` and `step.t2` make reference to the step of each observation. Finally, the element `tag`, is a vector of ones as long as the number of generated observations.

---

scoutgrid

*scoutgrid*


---

### Description

Shift of an array following a grid pattern.

### Usage

```

scoutgrid(
  X,
  pcaref,
  T2.target = NA,
  SPE.target = NA,
  nsteps.t2 = 1,

```

```

nsteps.spe = 1,
gspe = 1,
gt2 = 1
)

```

### Arguments

<code>X</code>	Matrix with observations that will be shifted as rows.
<code>pcaref</code>	List with the elements of a PCA model: <code>m</code> (mean), <code>s</code> (standard deviation), <code>prepro</code> (preprocessing: "none", "cent" or "autosc"), <code>P</code> (loading matrix), <code>lambda</code> (vector with variances of each PC).
<code>T2.target</code>	A number indicating the target value for the $T^2_A$ after the shift. Set to NA by default.
<code>SPE.target</code>	A number indicating the target value for the SPE after the shift. Set to NA by default.
<code>nsteps.t2</code>	An integer indicating the number of steps in which the shift from the reference to the target value of the $T^2_A$ will be performed. Set to 1 by default.
<code>nsteps.spe</code>	An integer indicating the number of steps in which the shift from the reference to the target value of the SPE will be performed. Set to 1 by default.
<code>gspe</code>	A number indicating the term that will tune the spacing between steps for the SPE. Set to 1 by default (linear spacing).
<code>gt2</code>	A number indicating the term that will tune the spacing between steps for the SPE. Set to 1 by default (linear spacing).

### Value

list with elements: `X`, matrix with the new and shifted data, `SPE` and `T2` vectors with the statistic values of each one of the new generated outliers or observations, elements `step.spe` and `step.t2` make reference to the step of each observation. Finally, the element `tag`, is a vector of ones as long as the number of generated observations.

---

scoutsimple

*scoutsimple*


---

### Description

Shift of an array with a single step.

### Usage

```
scoutsimple(X, pcaref, T2.target = NA, SPE.target = NA)
```

### Arguments

<code>X</code>	Matrix with observations that will be shifted as rows.
<code>pcaref</code>	List with the elements of a PCA model: <code>m</code> (mean), <code>s</code> (standard deviation), <code>prepro</code> (preprocessing: "none", "cent" or "autosc"), <code>P</code> (loading matrix), <code>lambda</code> (vector with variances of each PC).

T2.target	A number indicating the target value for the $T^2_A$ after the shift. Set to NA by default.
SPE.target	A number indicating the target value for the SPE after the shift. Set to NA by default.

### Value

list with elements: X, matrix with the new and shifted data, SPE and T2 vectors with the statistic values of each one of the new generated outliers or observations, elements step.spe and step.t2 make reference to the step of each observation. Finally, the element tag, is a vector of ones as long as the number of generated observations.

---

scoutsteps	<i>scoutsteps</i>
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---

### Description

Shift of an array following a step-wise pattern.

### Usage

```
scoutsteps(
  X,
  pcaref,
  T2.target = NA,
  SPE.target = NA,
  nsteps = 1,
  gspe = 1,
  gt2 = 1
)
```

### Arguments

X	Matrix with observations that will be shifted as rows.
pcaref	List with the elements of a PCA model: m (mean), s (standard deviation), prepro (preprocessing: "none", "cent" or "autosc"), P (loading matrix), lambda (vector with variances of each PC).
T2.target	A number indicating the target value for the $T^2_A$ after the shift. Set to NA by default.
SPE.target	A number indicating the target value for the SPE after the shift. Set to NA by default.
nsteps	An integer indicating the number of steps in which the shift from the reference to the target values of the SPE and the $T^2_A$ will be performed. Set to 1 by default.
gspe	A number indicating the term that will tune the spacing between steps for the SPE. Set to 1 by default (linear spacing).
gt2	A number indicating the term that will tune the spacing between steps for the SPE. Set to 1 by default (linear spacing).

**Value**

list with elements: X, matrix with the new and shifted data, SPE and T2 vectors with the statistic values of each one of the new generated outliers or observations, elements `step.spe` and `step.t2` make reference to the step of each observation. Finally, the element `tag`, is a vector of ones as long as the number of generated observations.

---

<code>speinfo</code>	<i>speinfo</i>
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---

**Description**

Information about the Squared Prediction Error (SPE) of an observation. Two subplots show the information of an observation regarding its SPE statistic, i.e.: a bar plot indicating the value of the statistic for the observation, and a bar plot with the contribution that each component had for the SPE value

**Usage**

```
speinfo(SPE, E, limspe, iobs = NA)
```

**Arguments**

SPE	Vector with values of the SPE statistic.
E	Matrix with the contributions of each variable (columns) for each observation (rows) to the SPE. It is the error term obtained from the unexplained part of X by the PCA model.
limspe	Upper Control Limit for the SPE, at a certain confidence level $(1-\alpha)*100\%$ .
iobs	Integer with the index of the observation of interest. Default value set to NA.

**Value**

ggplot object with the generated bar plots.

---

X	<i>Demo dataset with 50 observations and 5 normally distributed variables with two Principal Components explaining the 80% of the total variance.</i>
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---

**Description**

It is a small data set to use as a demo for the SCOUTer package.

**Usage**

```
X
```

**Format**

A data frame with 50 rows and 5 normally distributed variables.

---

xshift	<i>xshift</i>
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---

**Description**

Shift of an observation. The performed operation results as a combination of two main directions: the direction of maximum gradient for the SPE (weighted by the parameter b) and the direction of the projection of the observation on the model (weighted by the parameter a).

**Usage**

```
xshift(X, P, a, b)
```

**Arguments**

X	Matrix with observations that will be shifted
P	Loading matrix of the PCA model according to which the shift will be performed.
a	A number or vector tuning the shift in the direction of its projection.
b	A number or vector tuning the shift in the direction of its residual.

**Value**

Matrix with shifted observation as rows, keeping the order of the input matrix X.

# Index

\*Topic **datasets**

[X](#), [14](#)

[barwithucl](#), [2](#)

[custombar](#), [3](#)

[distplot](#), [3](#)

[distplotsimple](#), [4](#)

[dotag](#), [5](#)

[dscplot](#), [5](#)

[ht2info](#), [6](#)

[obscontribpanel](#), [7](#)

[pcamb\\_classic](#), [7](#)

[pcame](#), [8](#)

[scoreplot](#), [9](#)

[scoreplotsimple](#), [9](#)

[scout](#), [10](#)

[scoutgrid](#), [11](#)

[scoutsimple](#), [12](#)

[scoutsteps](#), [13](#)

[speinfo](#), [14](#)

[X](#), [14](#)

[xshift](#), [15](#)