## Package 'Umatrix'

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```
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Description By gaining the property of emergence through self-organization, the enhance-
     ment of SOMs(self organizing maps) is called Emergent SOM (ESOM). The result of the projec-
     tion by ESOM is a grid of neurons which can be visualised as a three dimensional land-
     scape in form of the Umatrix. Further details can be found in the referenced publica-
     tions (see url). This package offers tools for calculating and visualis-
     ing the ESOM as well as Umatrix, Pmatrix and UStarMatrix. All the functionality is also avail-
     able through graphical user interfaces implemented in 'shiny'. Based on the recog-
     nized data structures, the method can be used to generate new data.
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     tools, grid, abind, deldir, geometry, pdist, AdaptGauss,
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Title Visualization of Structures in High-Dimensional Data

Type Package

2 Umatrix-package

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#### **Description**

The **ESOM**(emergent self organizing map) is an improvement of the regular **SOM**(self organizing map) which allows for toroid grids of neurons and is intended to be used in combination with the **Umatrix**. The set of neurons is referred to as **weights** within this package, as they represent the values within the high dimensional space. The neuron with smallest distance to a datapoint is called a **Bestmatch** and can be considered as projection of said datapoint. As the Umatrix is usually toroid, it is drawn four consecutive times to remove border effects. An island, or Imx, is a filter mask, which cuts out a subset of the Umatrix, which shows every point only a single time while avoiding border effects cutting through potential clusters. Finally the Pmatrix shows the density structures within the grid, by a set radius. It can be combined with the Umatrix resulting in the UStarMatrix, which is therefore a combination of density based structures as well as clearly divided ones.

#### References

Ultsch, A.: Data mining and knowledge discovery with emergent self-organizing feature maps for multivariate time series, In Oja, E. & Kaski, S. (Eds.), Kohonen maps, (1 ed., pp. 33-46), Elsevier, 1999.

Ultsch, A.: Maps for the visualization of high-dimensional data spaces, Proc. Workshop on Self organizing Maps (WSOM), pp. 225-230, Kyushu, Japan, 2003.

Ultsch, A.: U\* C: Self-organized Clustering with Emergent Feature Maps, Lernen, Wissensent-deckung und Adaptivitaet (LWA), pp. 240-244, Saarbruecken, Germany, 2005.

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Lotsch, J., Ultsch, A.: Exploiting the Structures of the U-Matrix, in Villmann, T., Schleif, F.-M., Kaden, M. & Lange, M. (eds.), Proc. Advances in Self-Organizing Maps and Learning Vector Quantization, pp. 249-257, Springer International Publishing, Mittweida, Germany, 2014.

Ultsch, A., Behnisch, M., Lotsch, J.: ESOM Visualizations for Quality Assessment in Clustering, In Merenyi, E., Mendenhall, J. M. & O'Driscoll, P. (Eds.), Advances in Self-Organizing Maps and Learning Vector Quantization: Proceedings of the 11th International Workshop WSOM 2016, pp. 39-48, Houston, Texas, USA, January 6-8, 2016, (10.1007/978-3-319-28518-4\_3), Cham, Springer International Publishing, 2016.

Thrun, M. C., Lerch, F., Lotsch, J., Ultsch, A.: Visualization and 3D Printing of Multivariate Data of Biomarkers, in Skala, V. (Ed.), International Conference in Central Europe on Computer Graphics, Visualization and Computer Vision, Plzen, 2016.

BMUHepta

Best matching units (BMU) of Hepta from FCPS (Fundamental Clustering Problem Suite)

#### Description

Best matching units (BMU) of an ESOM projection of the Hepta data set from FCPS (Fundamental Clustering Problem Suite) on an 80 x 40 planar grid of artifical neurons.

#### Usage

```
data("BMUHepta")
```

#### Details

Size 212, Dimensions 3 (key, linecoordinates, columncoorditaes)

Classes 7, stored in Hepta\$Cls

#### References

Ultsch A, Lotsch J: Machine-learned cluster identification in high-dimensional data. J Biomed Inform. 2017 Feb;66:95-104. doi: 10.1016/j.jbi.2016.12.011. Epub 2016 Dec 28.

```
data("BMUHepta")
str("BMUHepta")
```

calculate\_Delauny\_radius

Calculate the Delauny graph based radius

#### **Description**

Function to calculate the radius for data generation.

#### Usage

```
calculate_Delauny_radius(Data, BestMatches,
Columns = 80, Lines = 50, Toroid = TRUE)
```

#### **Arguments**

Data Matrix of data (as submitted to Umatrix generation)

BestMatches Array with positions of Bestmatches
Columns Number of columns of the Umatrix
Lines Number of columns of the Umatrix
Toroid Whether a toroid Umatrx was used

#### Value

Returns a list of results.

neighbourDistances

Distances on the Umatrix neigborhood matrix.

RadiusByEM Radius suggested by EM algorithm.

#### References

Ultsch A, Lotsch J: Machine-learned cluster identification in high-dimensional data. J Biomed Inform. 2017 Feb;66:95-104. doi: 10.1016/j.jbi.2016.12.011. Epub 2016 Dec 28.

```
## Not run:
data("Hepta")
data("HeptaBMU")
DelaunyHepta <- calculate_Delauny_radius(Data = Hepta$Data, BestMatches = HeptaBMU, Toroid = FALSE)
## End(Not run)</pre>
```

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esomTrain	Train an ESOM (emergent self organizing map) and project data

#### **Description**

The ESOM (emergent self organizing map) algorithm as defined by [Ultsch 1999]. A set of weights(neurons) on a two-dimensional grid get trained to adapt the given datastructure. The weights will be used to project data on a two-dimensional space, by seeking the BestMatches for every datapoint.

#### **Arguments**

Data that will be used for training and projection

Lines Height of grid
Columns Width of grid

Epochs Number of Epochs the ESOM will run
Toroid If TRUE, the grid will be toroid

NeighbourhoodFunction

Type of Neighbourhood; Possible values are: "cone", "mexicanhat" and "gauss"

StartLearningRate

Initial value for LearningRate

EndLearningRate

Final value for LearningRate

Start Radius Start value for the Radius in which will be searched for neighbours EndRadius End value for the Radius in which will be searched for neighbours

NeighbourhoodCooling

Cooling method for radius; "linear" is the only available option at the moment LearningRateCooling

Cooling method for LearningRate; "linear" is the only available option at the

shinyProgress Generate progress output for shiny if Progress Object is given ShiftToHighestDensity

If True, the Umatrix will be shifted so that the point with highest density will be

at the center

InitMethod name of the method that will be used to choose initializations Valid Inputs:

"uni\_min\_max": uniform distribution with minimum and maximum from sampleData "norm\_mean\_std": normal distribuation based on mean and standard

deviation of sampleData

Key Vector of numeric keys matching the datapoints. Will be added to Bestmatches

UmatrixForEsom If TRUE, Umatrix based on resulting ESOM is calculated and returned

#### Details

On a toroid grid, opposing borders are connected.

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#### Value

List with

BestMatches BestMatches of datapoints

Weights Trained weights

Lines Height of grid

Columns Width of grid

Toroid TRUE if grid is a toroid

JumpingDataPointsHist

Nr of DataPoints that jumped to a different BestMatch in every epoch

#### References

Kohonen, T., Self-organized formation of topologically correct feature maps. Biological cybernetics, 1982. 43(1): p. 59-69.

Ultsch, A., Data mining and knowledge discovery with emergent self-organizing feature maps for multivariate time series. Kohonen maps, 1999. 46: p. 33-46.

#### **Examples**

```
data('Hepta')
res=esomTrain(Hepta$Data, Key = 1:nrow(Hepta$Data))
```

generate\_data

Generative ESOM

#### **Description**

Function to generate new data with the same structure as the input data.

#### Usage

```
generate_data(Data, density_radius, Cls = NULL, gen_per_data = 10)
```

#### **Arguments**

Data Matrix of data (as submitted to Umatrix generation)

density\_radius Numeric value of data generation radius
Cls Classification of the data as a vector

gen\_per\_data New isntances per original iunstance to be generated

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#### Value

Returns a list of results.

#### References

Ultsch A, Lotsch J: Machine-learned cluster identification in high-dimensional data. J Biomed Inform. 2017 Feb;66:95-104. doi: 10.1016/j.jbi.2016.12.011. Epub 2016 Dec 28.

#### **Examples**

```
## Not run:
data("Hepta")
data("HeptaBMU")
HeptaData <- Hepta$Data
HeptaCls <y- Hepta$Cls
HeptaGenerated <- generate_data(HeptaData, 1, HeptaCls )
## End(Not run)</pre>
```

Hepta

Hepta from FCPS (Fundamental Clustering Problem Suite)

#### Description

Dataset with 7 easily seperable classes.

#### Usage

```
data("Hepta")
```

#### **Details**

```
Size 212, Dimensions 3, stored in Hepta$Data Classes 7, stored in Hepta$Cls
```

#### References

Ultsch, A.: U\* C: Self-organized Clustering with Emergent Feature Maps, Lernen, Wissensent-deckung und Adaptivitaet (LWA), pp. 240-244, Saarbruecken, Germany, 2005.

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#### **Examples**

```
data("Hepta")
str("Hepta")
```

iClassification

GUI for manual classification

#### **Description**

This tool is a 'shiny' GUI that visualizes a given Umatrix and allows the user to select areas and mark them as clusters.

#### **Arguments**

Umatrix Matrix of Umatrix Heights

BestMatches Array with positions of Bestmatches
Classification of the Bestmatches

Imx Matrix of an island that will be cut out of the Umatrix

Toroid Are BestMatches placed on a toroid grid? TRUE by default

#### Value

A vector containing the selected class ids. The order is corresponding to the given Bestmatches

#### References

Thrun, M. C., Lerch, F., Loetsch, J., Ultsch, A.: Visualization and 3D Printing of Multivariate Data of Biomarkers, in Skala, V. (Ed.), International Conference in Central Europe on Computer Graphics, Visualization and Computer Vision, Plzen, 2016.

```
## Not run:
data("Hepta")
e = esomTrain(Hepta$Data, Key = 1:nrow(Hepta$Data))
cls = iClassification(e$Umatrix, e$BestMatches)
## End(Not run)
```

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iEsomTrain	iEsomTrain	
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#### **Description**

Trains the ESOM and shows the Umatrix.

#### **Arguments**

Data Matrix of Data that will be used to learn. One DataPoint per row

BestMatches Array with positions of Bestmatches

Cls Classification of the Bestmatches as a vector

Key Numeric vector of keys matching the Bestmatches

Toroid Are BestMatches placed on a toroid grid? TRUE by default

#### Value

List with

Umatrix matrix with height values of the umatrix

BestMatches matrix containing the bestmatches

Lines number of lines of the chosen ESOM

Columns of the chosen ESOM

Epochs number of epochs of the chosen ESOM

Weights List of weights

Toroid True if a toroid grid was used

EsomDetails Further details describing the chosen ESOM parameters

JumpingDataPointsHist

Number of Datapoints that jumped to another neuron in each epoch

#### References

Thrun, M. C., Lerch, F., Loetsch, J., Ultsch, A.: Visualization and 3D Printing of Multivariate Data of Biomarkers, in Skala, V. (Ed.), International Conference in Central Europe on Computer Graphics, Visualization and Computer Vision, Plzen, 2016.

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iUmapIsland	iUmapIsland
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#### Description

The toroid Umatrix is usually drawn 4 times, so that connected areas on borders can be seen as a whole. An island is a manual cutout of such a tiled visualization, that is selected such that all connected areas stay intact. This 'shiny' tool allows the user to do this manually.

#### **Arguments**

Umatrix Matrix of Umatrix Heights

BestMatches Array with positions of BestMatches
Classification of the BestMatches

#### Value

Boolean Matrix that represents the island within the tiled Umatrix

#### References

Thrun, M. C., Lerch, F., Loetsch, J., Ultsch, A.: Visualization and 3D Printing of Multivariate Data of Biomarkers, in Skala, V. (Ed.), International Conference in Central Europe on Computer Graphics, Visualization and Computer Vision, Plzen, 2016.

#### **Examples**

```
## Not run:
data("Hepta")
e = esomTrain(Hepta$Data, Key = 1:nrow(Hepta$Data))
Imx = iUmapIsland(e$Umatrix, e$BestMatches)
plotMatrix(e$Umatrix, e$BestMatches, Imx = Imx$Imx)
## End(Not run)
```

iUstarmatrix

iUstarmatrix

#### Description

Calculates the Ustarmatrix by combining a Umatrix with a Pmatrix.

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#### **Arguments**

Weights Weights that were trained by the ESOM algorithm

Lines Height of the used grid
Columns Width of the used grid

Data Matrix of Data that was used to train the ESOM. One datapoint per row

Imx Island mask that will be cut out from displayed Umatrix

Cls Classification of the Bestmatches

Toroid Are weights placed on a toroid grid?

#### Value

Ustarmatrix matrix with height values of the Ustarmatrix

#### References

Thrun, M. C., Lerch, F., Loetsch, J., Ultsch, A.: Visualization and 3D Printing of Multivariate Data of Biomarkers, in Skala, V. (Ed.), International Conference in Central Europe on Computer Graphics, Visualization and Computer Vision, Plzen, 2016.

#### **Description**

Draws a plot based of given Umatrix or Pmatrix.

## Arguments

Matrix	Umatrix or Pmatrix to be plotted
BestMatches	Positions of BestmMtches to be plotted onto the Umatrix
Cls	Class identifier for the BestMatches
ClsColors	Vector of colors that will be used to colorize the different classes
ColorStyle	If "Umatrix" the colors of a Umatrix (Blue -> Green -> Brown -> White) will be used; If "Pmatrix" the colors of a Pmatrix (White -> Yellow -> Red) will be used
Toroid	Should the Umatrix be drawn 4times?
BmSize	Integer between 0.1 and 5, magnification factor of the drawn BestMatch circles
DrawLegend	If TRUE, a color legend will be drawn next to the plot
FixedRatio	If TRUE, the plot will be drawn with a fixed ratio of x and y axis
CutoutPol	Only draws the area within given polygon
Nrlevels	Number of height levels that will be used within the Umatrix

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TransparentContours

Use half transparent contours. Looks better but is slow

Imx Mask to cut out an island. Every value should be either 1 (stays in) or 0 (gets

cut out)

Clean If TRUE axis, margins, ... surrounding the Umatrix image will be removed

RemoveOcean If TRUE, the surrounding blue area around an island will be reduced as much as

possible (while still maintaining a rectangular form)

Transparent0cean

If TRUE, the surrounding blue area around an island will be transparent

Title A title that will be drawn above the plot

BestMatchesLabels

Vector of strings corresponding to the order of BestMatches which will be drawn

on the plot as labels

BestMatchesShape

Numeric value of Shape that will be used. Responds to the usual shapes of

ggplot

MarkDuplicatedBestMatches

If TRUE, BestMatches that are shown more than once within an island, will be

marked

YellowCircle If TRUE, a yellow circle is drawn around Bestmatches to distinct them better

from background

#### **Details**

The heightScale (nrlevels) is set at the proportion of the 1 percent quantile against the 99 percent quantile of the matrix values.

#### Value

A 'ggplot' of a Matrix

#### References

Thrun, M. C., Lerch, F., Loetsch, J., Ultsch, A.: Visualization and 3D Printing of Multivariate Data of Biomarkers, in Skala, V. (Ed.), International Conference in Central Europe on Computer Graphics, Visualization and Computer Vision, Plzen, 2016.

Ultsch, A.: Maps for the visualization of high-dimensional data spaces, Proc. Workshop on Self organizing Maps (WSOM), pp. 225-230, Kyushu, Japan, 2003.

Siemon, H.P., Ultsch, A.: Kohonen Networks on Transputers: Implementation and Animation, in: Proceedings Intern. Neural Networks, Kluwer Academic Press, Paris, pp. 643-646, 1990.

```
data("Hepta")
e = esomTrain(Hepta$Data, Key = 1:nrow(Hepta$Data))
plotMatrix(e$Umatrix,e$BestMatches)
```

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#### Description

Generates a Pmatrix based on the weights of an ESOM.

#### **Arguments**

Data	A [n,k] matrix containing the data
Weights	Weights stored as a list in a 2D matrix
Lines	Number of lines of the SOM that is described by weights
Columns	Number of columns of the SOM that is described by weights
Radius	The radius for measuring the density within the hypersphere
PlotIt	If set the Pmatrix will also be plotted
Toroid	Are BestMatches placed on a toroid grid? TRUE by default

#### Value

UstarMatrix

#### References

Ultsch, A.: Maps for the visualization of high-dimensional data spaces, Proc. Workshop on Self organizing Maps (WSOM), pp. 225-230, Kyushu, Japan, 2003.

Ultsch, A., Loetsch, J.: Computed ABC Analysis for Rational Selection of Most Informative Variables in Multivariate Data, PloS one, Vol. 10(6), pp. e0129767. doi 10.1371/journal.pone.0129767, 2015.

Thrun, M. C., Lerch, F., Loetsch, J., Ultsch, A.: Visualization and 3D Printing of Multivariate Data of Biomarkers, in Skala, V. (Ed.), International Conference in Central Europe on Computer Graphics, Visualization and Computer Vision, Plzen, 2016.

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#### **Description**

Visualizes the matrix(Umatrix/Pmatrix) in an interactive window in 3D.

#### **Arguments**

Matrix

Toroid

BestMatches	Positions of BestMatches to be plotted onto the matrix
Cls	Class identifier for the BestMatch at the given point
Imx	a mask (island) that will be used to cut out the Umatrix

HeightScale Optional. Scaling Factor for Mountain Height

BmSize Size of drawn BestMatches

RemoveOcean Remove as much area sourrounding an island as possible

ColorStyle Either "Umatrix" or "Pmatrix" respectively for their colors

ShowAxis Draw an axis arround the drawn matrix

Matrix to be plotted

SmoothSlope Try to increase the island size, to get smooth slopes around the island

Should the Matrix be drawn 4 times (in a toroid view)

ClsColors Vector of colors that will be used for classes
FileName Name for a stl file to write the Matrix to

#### **Details**

The heightScale is set at the proportion of the 1 percent quantile against the 99 percent quantile of the Matrix values.

#### References

Thrun, M. C., Lerch, F., Loetsch, J., Ultsch, A.: Visualization and 3D Printing of Multivariate Data of Biomarkers, in Skala, V. (Ed.), International Conference in Central Europe on Computer Graphics, Visualization and Computer Vision, Plzen, 2016.

```
## Not run:
data("Hepta")
e = esomTrain(Hepta$Data, Key = 1:nrow(Hepta$Data))
showMatrix3D(e$Umatrix)
## End(Not run)
```

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umatrixForEsom	umatrixForEsom	

#### **Description**

Calculate the Umatrix for given ESOM projection

#### **Arguments**

Weights	Weights from which the Umatrix will be calculated
Lines	Number of lines of the SOM that is described by weights
Columns	Number of columns of the SOM that is described by weights
Toroid	Boolean describing if the neural grid should be borderless

#### Value

Umatrix

#### References

Ultsch, A. and H.P. Siemon, Kohonen's Self Organizing Feature Maps for Exploratory Data Analysis. 1990.

#### **Examples**

ustarmatrixCalc

ustarmatrixCalc

#### Description

The UStarMatrix is a combination of the Umatrix (average distance to neighbours) and Pmatrix (density in a point). It can be used to improve the Umatrix, if the dataset contains density based structures.

#### Arguments

 $\begin{array}{ll} \mbox{Umatrix} & \mbox{A given Umatrix} \\ \mbox{Pmatrix} & \mbox{A density matrix} \end{array}$ 

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#### Value

**UStarMatrix** 

#### References

Ultsch, A. U\* C: Self-organized Clustering with Emergent Feature Maps. in Lernen, Wissensent-deckung und Adaptivitaet (LWA). 2005. Saarbruecken, Germany.

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