Package: CluMP (via r-universe)

October 20, 2026

Title Clustering of Micro Panel Data

Version 0.8.1

Description Two-step feature-based clustering method designed for micro panel (longitudinal) data with the artificial panel data generator. See Sobisek, Stachova, Fojtik (2018) <doi:10.48550/arXiv.1807.05926>.

URL https://arxiv.org/ftp/arxiv/papers/1807/1807.05926.pdf

Depends R (>= 3.4.0)

License GPL (>= 3)

Encoding UTF-8

LazyData true

RoxygenNote 7.1.1

Imports MASS, ggplot2 (>= 3.0.0), dplyr (>= 0.7.6), NbClust (>= 3.0), amap (>= 0.8-16), tableone, stats, data.table, rlang

Suggests knitr, rmarkdown

NeedsCompilation no

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Config/pak/sysreqs make libicu-dev libx11-dev zlib1g-dev

Repository https://9afojtik.r-universe.dev

Date/Publication 2020-11-27 01:20:29 UTC

RemoteUrl https://github.com/cran/CluMP

RemoteRef HEAD

RemoteSha 3ba9a1efec9ef83c782f5d8ada76e225ac9edd91

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`CluMP` *Cluster Micro-Panel (longitudinal) Data employing the CluMP algorithm*

Description

This function clusters Micro-Panel (longitudinal) Data (or trajectories) to a pre-defined number of clusters by employing Feature-Based Clustering of Micro-Panel (longitudinal) Data algorithm called CluMP (see Reference). Currently, only univariate clustering analysis is available.

Usage

```
CluMP(formula, group, data, cl_numb = NA, base_val = FALSE, method = "ward.D")
```

Arguments

`formula`	A two-sided `formula` object with a numeric clustering variable (Y) on the left of a ~ separator and the time (numeric) variable on the right. Time is measured from the start of the follow-up period (baseline). Any time units are possible.
`group`	A grouping factor variable (vector), i.e. single identifier for each individual (trajectory).
`data`	A data frame containing the variables named in the `formula` and `group` arguments.
`cl_numb`	An integer, positive number (scalar) specifying the number of clusters. The `OptiNum` function can be used to determine the optimal number of clusters according to common evaluation criteria (indices).
`base_val`	Indicates whether include a value at zero time point as an additional clustering variable. Default is *FALSE* and the standard number (7) of clustering parameters is used.
`method`	A method which use in hierarhical clustering, same as in `hclust` function, namely "ward.D", "ward.D2", "single", "complete", "average", "mcquitty", "median", "centroid". Default is "ward.D".

Value

Cluster Micro-Panel data. The output is the list of 5 components which contain results from clustering.

Source

Sobisek, L., Stachova, M., Fojtik, J. (2018) Novel Feature-Based Clustering of Micro-Panel Data (CluMP). Working paper version online: www.arxiv.org

Examples

```
data <- GeneratePanel(n = 100, Param = ParamLinear, NbVisit = 10)
CluMP(formula = Y ~ Time, group = "ID", data = data, cl_numb = 3,
base_val = FALSE, method = "ward.D")

CluMP(formula = Y ~ Time, group = "ID", data = data, cl_numb = 3,
base_val = TRUE, method = "ward.D")
```

CluMP_profiles *Summary characteristics of identified clusters via CluMP*

Description

The function CluMP_profiles provides a description (profile) for each cluster. The description is in the form of a summary list containing descriptive statistics of a cluster variable, time variable, cluster parameters and other variables (covariates), both continuous and categorical.

Usage

```
CluMP_profiles(CluMPoutput, cont_vars = NULL, cat_vars = NULL, show_NA = FALSE)
```

Arguments

CluMPoutput	An object (output) from the CluMP function.
cont_vars	An optional single character or a character vector of continuous variables' names (from the original dataset).
cat_vars	An optional single character or a character vector of categorical variables' names (from the original dataset).
show_NA	Logical scalar. Should be calculated and shown descriptive statistics for *NA* cluster if exists? Default is *FALSE*. *NA* cluster gathers improper individuals (trajectories with < 3 not missing observations) for longitudinal clustering.

Value

Returns a list with cluster variable (Y) summary, both baseline and changes; time and a summary of the number of observations (visits); clustering parameters summary and optional continuous variables summary (baseline and changes) and categorical variables summary (baseline and end).

Examples

```
set.seed(123)
dataMale <- GeneratePanel(n = 50, Param = ParamLinear, NbVisit = 10)
dataMale$Gender <- "M"
dataFemale <- GeneratePanel(n = 50, Param = ParamLinear, NbVisit = 10)
dataFemale$ID <- dataFemale$ID + 50
dataFemale$Gender <- "F"
data <- rbind(dataMale, dataFemale)

CluMPoutput <- CluMP(formula = Y ~ Time, group = "ID", data = data, cl_numb = 3)
CluMP_profiles(CluMPoutput, cat_vars = "Gender")
```

CluMP_view *Cluster profiles' (CluMP results) visualisation*

Description

This graphical function enables to visualise cluster profiles (mean representatives of each cluster). Available are three types of plots: non-parametric (LOESS method for small/medium or GAM method for complex data of large size. Both methods are applied from ggplot2 representatives (mean within-cluster trajectories) with/without all individual (original) trajectories, and nonparametric mean trajectories with error bars.

Usage

```
CluMP_view(
  CluMPoutput,
  type = "all",
  nb_intervals = NULL,
  return_table = FALSE,
  title = NULL,
  x_title = NULL,
  y_title = NULL,
  plot_NA = FALSE
)
```

Arguments

CluMPoutput	An object (output) from the CluMP function.

type	String. Indicates which type of graph is required. Possible values for this argument are: *"all"* (plots all data with non-parametric mean trajectories), *"cont"* (only non-parametric mean trajectories) or *"breaks"* (mean trajectories with error bars).
nb_intervals	An integer, positive number (scalar) specifying the number of regular timepoints into which should be follow-up period split. This argument works only with graph type = *"breaks"*. In case of other graph types the argument is ignored. The number of error bars is equal to the number of timepoints specified by this argument.
return_table	Logical scalar indicating if the summary table of plotted values in the graph of type = *"breaks"* should be returned. Default is *FALSE*.
title	String. Optional title for a plot. If undefined, no title will used.
x_title	String. An optional title for *x* axis. If undefined, the variable name after ~ in `formula` will used.
y_title	String. An optional title for *y* axis. If undefined, the variable name before ~ in `formula` will used.
plot_NA	Plot *NA* cluster if exists. Default is *FALSE*. *NA* cluster gathers improper individuals (< 3 observations) for longitudinal clustering.

Value

Returns graph for type *"all"* and *"cont"* or (list with) graph and table of mean trajectories (if specified) for type = *"breaks"*.

Examples

```
set.seed(123)
dataMale <- GeneratePanel(n = 50, Param = ParamLinear, NbVisit = 10)
dataMale$Gender <- "M"
dataFemale <- GeneratePanel(n = 50, Param = ParamLinear, NbVisit = 10)
dataFemale$ID <- dataFemale$ID + 50
dataFemale$Gender <- "F"
data <- rbind(dataMale, dataFemale)

CluMPoutput <- CluMP(formula = Y ~ Time, group = "ID", data = data, cl_numb = 3)
title <- "Plotting clusters' representatives with error bars"
CluMP_view(CluMPoutput, type = "all" , return_table = TRUE)
CluMP_view(CluMPoutput, type = "cont")
CluMP_view(CluMPoutput, type = "breaks", nb_intervals = 5, return_table=TRUE, title = title)
```

`GeneratePanel` *Generate an artificial Micro-Panel (longitudinal) Data*

Description

This function creates artificial linear or non-linear micro-panel (longitudinal) data coming from generating process with a certain function (linear, quadratic, cubic, exponencial) set of parameters (fixed and random (intercept, slope) effects of time).

Usage

```
GeneratePanel(
  n,
  Param,
  NbVisit,
  VisitFreq = NULL,
  TimeVar = NULL,
  RegModel = NULL,
  ClusterProb = NULL,
  Rho = NULL,
  units = NULL
)
```

Arguments

n	An integer specifying the number of individuals (trajectories) being observed.
Param	Object of data.frame containing regression parameters for each cluster. The dimensions are the various number of generating clusters and the fixed number of parameters. The second dimension (the fixed number of parameters) is given by the type of regression model specified by the argument "RegModel". For more information about the parameters, see documentation of: ParamLinear for linear model, ParamQuadrat for quadratic, ParamCubic for cubic model and ParamExpon for exponential model.
NbVisit	A positive integer numeric input defining expected number of visits. Option is *Fixed* or *Random*. Number of visits given by the argument VisitFreq. If VisitFreq is *Fixed*, the NbVisits defines exact number of visits for all individuals. If VisitFreq is *Random* then each individual has different number of visits. The number of visits is then generated from the poisson distribution with the mean (lambda) equal to NbVisits.
VisitFreq	String that defines the frequency of visits for each individual. Option is *Random* or *Fixed*. If set to *Fixed* or not defined, each individual has the same number of visits given by NbVisits. If set as *Random* the number of visits is generated from poisson distribution for each individual with the mean equal to the argument NbVisits. For example if this parameter is set as 5 then the random integer from interval of -5 to 5 is drawned and added to the time variable. Make sure that TimeVar must be lower then the number of days in parameter units.
TimeVar	A positive integer representing daily, time variability of the occurrence of repeated measurement (timepoint) from the regular, fixed occurrence (visit) given by the argument units. For example, if this argument is set to 5 then the random integer from interval of -5 to 5 is drawn and added to the time variable. TimeVar must be lower than the regular frequency of repeat measurement given by the argument units.

RegModel	String specifying the mathematical function for generating trajectory for each of n individuals. Options are *linear*, *quadratic*, *cubic* or *exponential*. If set to *linear* or not defined, then each trajectory has a linear trend. If set to *quadratic*, then each trajectory has a quadratic development in time. If set to *cubic* then each trajectory has cubic development. If set to *exponential*, then each trajectory has exponential development.
ClusterProb	Numeric scalar (for 2 clusters) or a vector of numbers (for >2 clusters) defining the probability of each cluster. If not defined, then each cluster has the same occurrence probability.
Rho	A numeric scalar specifying autocorrelation parameter with the values from range 0 to 1. If set as 0 or not define then there is no autocorrelation between the within-individual repeated observations.
units	String defining the units of time series. Options are *day*, *week*, *month* or *year*.

Value

Generates artificial panel data.

Examples

```
set.seed(123)
#Simple Linear model where each individual has 10 observations.
data <- GeneratePanel(n = 100, Param = ParamLinear, NbVisit = 10)

#Exponential model where each individual has 10 observations.
data <- GeneratePanel(100, ParamExpon, NbVisit = 10, VisitFreq = "Fixed", RegModel = "exponential")
PanelPlot(data)

#Cubic model where each individual has random number of observations on daily basis.
#Average number of observation is given by parameter NbVisit.
data <- GeneratePanel(n = 100, Param = ParamCubic, NbVisit = 100, RegModel = "cubic", units = "day")
PanelPlot(data)

#Quadratic model where each individual has random number of observations.
#Each object is observede weekly with variability 2 days.
data <- GeneratePanel(5,ParamQuadrat,NbVisit=50,RegModel="quadratic",units="week",TimeVar=2)
PanelPlot(data)

#Generate panel data with linear trend with 75% objects in first cluster and 25% in the second.
data <- GeneratePanel(n = 100, Param = ParamLinear, NbVisit = 10, ClusterProb = c(0.75, 0.25))
PanelPlot(data, colour = "Cluster")
```

OptiNum *Finding an optimal number of clusters*

Description

This function finds optimal number of clusters based on evaluation criteria (indices) available from the NbClust package.

Usage

```
OptiNum(
  formula,
  group,
  data,
  index = c("silhouette", "ch", "db"),
  max_clust = 10,
  base_val = FALSE
)
```

Arguments

formula	A two-sided formula object, with a numeric, clustering variable (Y) on the left of a ~ separator and the time (numeric) variable on the right. Time is measured from the start of the follow-up period (baseline).
group	A grouping factor variable (vector), i.e. single identifier for each individual (trajectory).
data	A data frame containing the variables named in formula and group arguments.
index	String vector of indices to be computed. Default is c("silhouette", "ch", "db"). See NbClust package for available indices and their description.
max_clust	An integer, positive number (scalar) defining the maximum number of clusters to check. Default value of this argument is 10 or maximum number of individuals.
base_val	Indicates whether include a value at zero time point as an additional clustering variable. Default is *FALSE* and the standard number (7) of clustering parameters is used.

Value

Determine the optimal number of clusters, returns graphical output (red dot in plot indicates the recommended number of clusters according to that index) and table with indices.

Source

Malika Charrad, Nadia Ghazzali, Veronique Boiteau, Azam Niknafs (2014). NbClust: An R Package for Determining the Relevant Number of Clusters in a Data Set. Journal of Statistical Software, 61(6), 1-36. URL http://www.jstatsoft.org/v61/i06/.

Examples

```
set.seed(123)
data <- GeneratePanel(n = 100, Param = ParamLinear, NbVisit = 10)
OptiNum(data = data, formula = Y ~ Time, group = "ID")
```

PanelPlot *Plot Micro-Panel (longitudinal) Data*

Description

This function plots micro-panel (longitudinal) data from stored `data.frame` or randomly generated panel data from `GeneratePanel` function.

Usage

```
PanelPlot(
  data,
  formula = Y ~ Time,
  group = "ID",
  colour = NA,
  mean_traj_all = FALSE,
  mean_traj_group = FALSE,
  show_legend = TRUE,
  title = NULL,
  x_title = NULL,
  y_title = NULL
)
```

Arguments

`data`	A data frame containing the variables named in `formula` and `group` arguments.
`formula`	A two-sided `formula` object, with a numeric, clustering variable (Y) on the left of a ~ separator and the time (numeric) variable on the right. Time is measured from the start of the follow-up period (baseline).
`group`	A grouping factor variable (vector), i.e. single identifier for each (trajectory).
`colour`	Character, which is a variable's name in data. The trajectories are distinguished by colour according to this variable.
`mean_traj_all`	Logical scalar. It indicates whether to show mean overall trajectory. Default is *FALSE*.
`mean_traj_group`	Logical scalar. It indicates whether to show mean trajectory by group. Default is *FALSE*.
`show_legend`	Logical scalar. It indicates whether to show cluster legend. Default is *TRUE*.
`title`	String. Is an optional title for a plot. Otherwise no title will used.
`x_title`	String. Is an optional title for x axis. Otherwise variable name after ~ in `formula` will used.
`y_title`	String. Is an optional title for y axis. Otherwise variable name before ~ in `formula` will used.

Value

Returns plot using package ggplot2.

Examples

```
set.seed(123)
dataMale <- GeneratePanel(n = 50, Param = ParamLinear, NbVisit = 10)
dataMale$Gender <- "M"
dataFemale <- GeneratePanel(n = 50, Param = ParamLinear, NbVisit = 10)
dataFemale$ID <- dataFemale$ID + 50
dataFemale$Gender <- "F"
data <- rbind(dataMale, dataFemale)

PanelPlot(data = data, formula = Y ~ Time, group = "ID", colour = "Gender")
PanelPlot(data = data, formula = Y ~ Time, group = "ID", colour = "Gender", mean_traj_all = TRUE)
PanelPlot(data = data, formula = Y ~ Time, group = "ID", colour = "Gender", mean_traj_group = TRUE)
```

ParamCubic *Parameters of cubic model*

Description

Default parameters to generate micro-panel (longitudinal) data with quadratic trend. The parameters may differ per each cluster. The parameters of each cluster are in rows. Number of rows denotes the number of clusters. Fixed effects are taken from Allen et al. (2005), and the source for random effects is Uher et al. (2017).

Usage

```
ParamCubic
```

Format

Its adviced to keep parameters in `data.frame`. The Parameters structure is as follows:

b0 fixed parameter of intercept

b1 fixed parameter of slope

b2 fixed parameter of defining the quadraticity

b3 fixed parameter of defining the cubicity

varU0 variance of random factor U0 given to fixed parameter b0

varU1 variance of random factor U1 given to fixed parameter b1

corr correlation between random factors U0 and U1

varE the variability of the residuals

Source

Allen, JS, Bruss, J, Brown, CK, Damasio, H. Normal neuroanatomical variation due to age: the major lobes and a parcellation of the temporal region. Neurobiol Aging. 2005 Oct;26(9):1245-60; discussion 1279-82.

Uher T, Vaneckova M, Krasensky J, Sobisek L, Tyblova M, Volna J, Seidl Z, Bergsland N, Dwyer MG, Zivadinov R, De Stefano N, Sormani MP, Havrdova EK, Horakova D. Pathological cut-offs of global and regional brain volume loss in multiple sclerosis. Mult Scler. 2017 Nov 1:1352458517742739. doi: 10.1177/1352458517742739.

`ParamExpon` *Parameters of exponential model*

Description

Default parameters to generate micro-panel (longitudinal) data with exponencial trend. The parameters may differ per each cluster. The parameters of each cluster are in rows. Number of rows denotes the number of clusters. Fixed effects are taken from Jones et al. (2013).

Usage

```
ParamExpon
```

Format

It is adviced to keep parameters in `data.frame`. The Parameters structure is as follows:

b0 fixed parameter of intercept

b1 fixed parameter of slope

b2 fixed parameter of defining the decay

varU0 variance of random factor U0 given to fixed parameter b0

varU1 variance of random factor U1 given to fixed parameter b1

corr correlation between random factors U0 and U1

varE the variability of the residuals

Source

Jones BC, Nair G, Shea CD, Crainiceanu CM, Cortese IC, Reich DS. Quantification of multiple-sclerosis-related brain atrophy in two heterogeneous MRI datasets using mixed-effects modeling. Neuroimage Clin. 2013 Aug 13;3:171-9. doi: 10.1016/j.nicl.2013.08.001.

ParamLinear *Parameters of linear model*

Description

Default parameters to generate micro-panel (longitudinal) data with linear trend. The parameters may differ per each cluster. The parameters of each cluster are in rows. Number of rows denotes the number of clusters. Fixed and random effects are taken from Uher et al. (2017).

Usage

```
ParamLinear
```

Format

It is adviced to keep parameters in `data.frame`. The Parameters structure is as follows:

b0 fixed parameter of intercept

b1 fixed parameter of slope

varU0 variance of random factor U0 given to fixed parameter b0

varU1 variance of random factor U1 given to fixed parameter b1

corr correlation between random factors U0 and U1

varE the variability of the residuals

Source

Uher T, Vaneckova M, Krasensky J, Sobisek L, Tyblova M, Volna J, Seidl Z, Bergsland N, Dwyer MG, Zivadinov R, De Stefano N, Sormani MP, Havrdova EK, Horakova D. Pathological cut-offs of global and regional brain volume loss in multiple sclerosis. Mult Scler. 2017 Nov 1:1352458517742739. doi: 10.1177/1352458517742739.

ParamQuadrat *Parameters of quadratic model*

Description

Parameters to generate panel data with quadratic trend. The parameters may differ per each cluster. The parameters of each cluster are in rows. Number of rows denotes the number of clusters. Fixed effects are taken from Allen et al. (2005), and the source for random effects is Uher et al. (2017).

Usage

```
ParamQuadrat
```

Format

It is adviced to keep parameters in `data.frame`. The Parameters structure is as follows:

b0 fixed parameter of intercept

b1 fixed parameter of slope

b2 fixed parameter of defining the quadraticity

varU0 variance of random factor U0 given to fixed parameter b0

varU1 variance of random factor U1 given to fixed parameter b1

corr correlation between random factors U0 and U1

varE the variability of the residuals

Source

Allen, JS, Bruss, J, Brown, CK, Damasio, H. Normal neuroanatomical variation due to age: the major lobes and a parcellation of the temporal region. Neurobiol Aging. 2005 Oct;26(9):1245-60; discussion 1279-82.

Uher T, Vaneckova M, Krasensky J, Sobisek L, Tyblova M, Volna J, Seidl Z, Bergsland N, Dwyer MG, Zivadinov R, De Stefano N, Sormani MP, Havrdova EK, Horakova D. Pathological cut-offs of global and regional brain volume loss in multiple sclerosis. Mult Scler. 2017 Nov 1:1352458517742739. doi: 10.1177/1352458517742739.

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