

# Package: trajeR (via r-universe)

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**Type** Package

**Title** Group Based Modeling Trajectory

**Description** Estimation of group-based trajectory models, including finite mixture models for longitudinal data, supporting censored normal, zero-inflated Poisson, logit, and beta distributions, using expectation-maximization and quasi-Newton methods, with tools for model selection, diagnostics, and visualization of latent trajectory groups, [doi:10.4159/9780674041318](https://doi.org/10.4159/9780674041318), Nagin, D. (2005). Group-Based Modeling of Development. Cambridge, MA: Harvard University Press. and Noel (2022), <https://orbilu.uni.lu/>, thesis.

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**Suggests** spelling

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adequacy	<i>Adequacy of the model</i>
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---

**Description**

Calculate the summary of the five methods : assignment proportion, average posterior probability, confidence interval, odds of Correct Classification.

**Usage**

```
adequacy(sol, Y, A, nb = 10000, alpha = 0.98)
```

**Arguments**

sol	Trajectory's object. An object of type Trajectory.
Y	Matrix. A matrix containing the variables in the model.
A	Matrix. A matrix containing the time variable data.
nb	Integer. The numbers of repetitions in the bootstrap method.
alpha	Real. The degree of confidence of the interval.

**Value**

A table of reals. A table with 5 rows: the estimate probabilities, the two bounds of the confidence interval, the proportion of assignment, the Average Posterior Probability and the Odds of Correct Classification.

**Examples**

```
data <- read.csv(system.file("extdata", "CNORM2gr.csv", package = "trajeR"))
data <- as.matrix(data)
sol <- trajeR(Y = data[, 2:6], A = data[, 7:11], degre = c(2, 2), Model = "CNORM", Method = "EM")
adequacy(sol, Y = data[, 2:6], A = data[, 7:11])
```

---

AvePP	<i>Average Posterior Probability</i>
-------	--------------------------------------

---

**Description**

Calculate the Average Posterior Probability. Average Posterior Probability (AvePP) is the average posterior probability of membership for each group for those individuals that were assigned to.

**Usage**

```
AvePP(sol, Y, A, X = NULL)
```

**Arguments**

sol	Trajectory's object. An object of type Trajectory.
Y	Matrix. A matrix containing the variables in the model.
A	Matrix. A matrix containing the time variable data.
X	Matrix. An optional matrix that modifies the probability of belong to group. By default its value is a one column matrix with value 1.

**Value**

A vector of reals. The average posterior probability.

**Examples**

```
data <- read.csv(system.file("extdata", "CNORM2gr.csv", package = "trajeR"))
data <- as.matrix(data)
sol <- trajeR(Y = data[, 2:6], A = data[, 7:11], degree = c(2, 2), Model = "CNORM", Method = "EM")
AvePP(sol, Y = data[, 2:6], A = data[, 7:11])
```

---

ConfIntT

*Confidence interval*

---

**Description**

Calculate the confidence interval of the probabilities with bootstrap method. We have to specify the number of the repetitions of bootstrap and the degree of confidence.

**Usage**

```
ConfIntT(sol, Y, A, nb = 10000, alpha = 0.98)
```

**Arguments**

sol	Trajectory's object. An object of type Trajectory.
Y	Matrix. A matrix containing the variables in the model.
A	Matrix. A matrix containing the time variable data.
nb	An integer. The number of repetitions in the bootstrap method.
alpha	A number. The degree of confidence of the interval.

**Value**

A vector of reals. The two bounds of the confidence interval given a degree of confidence.

**Examples**

```
data <- read.csv(system.file("extdata", "CNORM2gr.csv", package = "trajeR"))
data <- as.matrix(data)
sol <- trajeR(Y = data[, 2:6], A = data[, 7:11], degree = c(2, 2), Model = "CNORM", Method = "EM")
ConfIntT(sol, Y = data[, 2:6], A = data[, 7:11])
```

dataNORM01

*Simulated CNORM Dataset (data01)***Description**

The data01 dataset is a simulated dataset included in the package. It contains 500 trajectories with 10 time-points each, simulating a 3-group solution. The parameters for the groups are summarized in the table below:

Parameter	Group 1	Group 2	Group 3
$\beta_{k0}$	2.797	7.000	19.545
$\beta_{k1}$	8.809	-0.297	-0.407
$\beta_{k2}$	-3.201	0.463	0.026
$\beta_{k3}$	0.000	-0.021	0.000
$\pi_k$	0.32	0.54	0.14
$\sigma_k$	4.000	4.000	4.000

**Usage**

dataNORM01

**Format**

A data frame with 500 rows and 43 columns:

**Columns 2:11** Matrix of real numbers (variable of interest, Y).

**Columns 12:21** Matrix of time points (1 to 10, A).

**Columns 22:31** Matrix of binary values (0 or 1, time-dependent covariate W).

**Columns 32:41** Matrix of real numbers (additional time-dependent covariate W).

**Columns 42:43** Matrix of binary values (0 or 1, covariate X).

**Details**

The dataset includes the following variables:

- Y (columns 2:11): Matrix of real numbers representing the variable of interest.
- A (columns 12:21): Matrix of time points (1 to 10).
- W (columns 22:31): Matrix with values 0 and 1, indicating the presence or absence of a characteristic.
- W (columns 32:41): Matrix of real numbers (additional time-dependent covariate).
- X (columns 42:43): Matrix with values 0 and 1, influencing group membership probability.

**Source**

Simulated data for illustrative purposes.

---

diffaitbeta                      *Differential*

---

### Description

Differential

### Usage

diffaitbeta(betak, i, t, A, TCOV, fct, diffct)

### Arguments

betak	Vector of integer.
i	Integer.
t	Real.
A	Matrix of real.
TCOV	Matrix of real.
fct	Function.
diffct	Function.

### Value

real. Compute the value of the differential function fct for individual i, time t and group k.

---

fait                                      *Function fait*

---

### Description

Function fait

### Usage

fait(betak, i, t, A, TCOV, fct, diffct)

### Arguments

betak	Vector of integer.
i	Integer.
t	Real.
A	Matrix of real.
TCOV	Matrix of real.
fct	Function.
diffct	Function.

**Value**

real. Compute the value of the function fct for individual i, time t and group k.

---

GroupProb	<i>Membership's probabilities</i>
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---

**Description**

GroupProb calculate the membership probability of each value of the data.

**Usage**

```
GroupProb(Obj, Y, A, TCOV = NULL, X = NULL)
```

**Arguments**

Obj	Trajectory's object. A trajectory object that is return by trajeR function.
Y	Matrix. A real matrix. The data.
A	Matrix. A real matrix. The time variable.
TCOV	Matrix. A real matrix. Optional, by default the value is NULL. It contained the time dependent covariate.
X	Matrix. A real matrix. Optional, by default the value is NULL. It contained a covariate that modify the probability membership.

**Value**

a real matrix. For each individual i in the data, this matrix contained the membership probability of each group.

**Examples**

```
data <- read.csv(system.file("extdata", "CNORM2gr.csv", package = "trajeR"))
data <- as.matrix(data)
sol <- trajeR(Y = data[, 2:6], A = data[, 7:11], degre = c(2, 2), Model = "CNORM", Method = "EM")
GroupProb(sol, Y = data[, 2:6], A = data[, 7:11])
```

---

GroupProfiles	<i>Profiles of each group</i>
---------------	-------------------------------

---

### Description

GroupProfiles calculate the profile of a group regarding covariate. It is a cross tabulation of individual level trajectory group assignments with individual level characteristic that might be associated with trajectory group membership.

### Usage

```
GroupProfiles(sol, Y, A, X)
```

### Arguments

sol	Trajectory's object. A object of type trajectory.
Y	Matrix. A matrix containing the variables in the model.
A	Matrix. A matrix containing the time variable data.
X	Matrix. An optional matrix that modify the probability of belong to group. By default its value is a matrix with one column with value 1.

### Value

A table of real.

### Examples

```
data <- read.csv(system.file("extdata", "CNORM2gr.csv", package = "trajeR"))
data <- as.matrix(data)
sol <- trajeR(
  Y = data[, 2:6], A = data[, 7:11], Risk = data[, 12],
  degree = c(2, 2), Model = "CNORM", Method = "L"
)
GroupProfiles(sol, Y = data[, 2:6], A = data[, 7:11], X = data[, 12])
```

---

OCC	<i>Odds of Correct Classification</i>
-----	---------------------------------------

---

### Description

Calculate Odds of Correct Classification. The Odds of Correct Classification for group k (OCC<sub>j</sub>) is the ratio between the odds of a correct classification into group j on the basis of the posterior probability rule and the odds of correct assignment based on random assignments with the probability of assignment to group j is the probability estimate by the model.

**Usage**

```
OCC(sol, Y, A)
```

**Arguments**

<code>sol</code>	Trajectory's object. An object of type Trajectory.
<code>Y</code>	Matrix. A matrix containing the variables in the model.
<code>A</code>	Matrix. A matrix containing the time variable data.

**Value**

A vector of reals. The Odds of Correct Classification.

**Examples**

```
data <- read.csv(system.file("extdata", "CNORM2gr.csv", package = "trajeR"))
data <- as.matrix(data)
sol <- trajeR(Y = data[, 2:6], A = data[, 7:11], degree = c(2, 2), Model = "CNORM", Method = "EM")
OCC(sol, Y = data[, 2:6], A = data[, 7:11])
```

---

plotrajeR

*plot trajectory*

---

**Description**

plot trajectory

**Usage**

```
plotrajeR(Obj, ...)
```

**Arguments**

<code>Obj</code>	an object of class "Trajectory".
<code>...</code>	optional parameters

**Value**

a graphic.

**Examples**

```
data <- read.csv(system.file("extdata", "CNORM2gr.csv", package = "trajeR"))
data <- as.matrix(data)
sol <- trajeR(Y = data[, 2:6], A = data[, 7:11], degree = c(2, 2), Model = "CNORM", Method = "EM")
plotrajeR(sol)
```

---

```
plotrajeR.Trajectory.BETA
      plot BETA trajectory
```

---

**Description**

plot BETA trajectory

**Usage**

```
## S3 method for class 'Trajectory.BETA'
plotrajeR(
  Obj,
  plotcov = NULL,
  col = "black",
  Y = NULL,
  A = NULL,
  Risk = NULL,
  TCOV = NULL,
  mean = FALSE,
  alpha = 1,
  ...
)
```

**Arguments**

Obj	an object of class "Trajectory.LOGIT".
plotcov	an optional vector or matrix with the same length as the time period. Default value is NULL.
col	an optional vector. The vector of colors. It must contain a color for each trajectory and each points of groups. Its length is the double of the number of group. Default value is a grayscale.
Y	Matrix. A matrix containing the variables in the model.
A	Matrix. A matrix containing the time variable data.
Risk	Matrix. An optional matrix that modify the probability of belong to group. By default its value is a matrix with one column with value 1.
TCOV	Matrix. An optional matrix containing the time covariate that influence the trajectory themselves. By default its value is NULL.
mean	an optional logical. Indicate if the mean of ech group and time value must be draw.
alpha	on optional real. Indicate the alpha channel of the points color.
...	optional parameters

**Value**

a graphic.

---

```
plotrajeR.Trajectory.CNORM
  plot CNORM trajectory
```

---

**Description**

plot CNORM trajectory

**Usage**

```
## S3 method for class 'Trajectory.CNORM'
plotrajeR(
  Obj,
  plotcov = NULL,
  col = "black",
  Y = NULL,
  A = NULL,
  Risk = NULL,
  mean = FALSE,
  alpha = 1,
  ...
)
```

**Arguments**

Obj	an object of class "Trajectory.CNORM".
plotcov	an optional vector or matrix with the same length as the time period. Default value is NULL.
col	an optional vector. The vector of colors. It must contain a color for each trajectory and each points of groups. Its length is the double of the number of group. Default value is a grayscale.
Y	Matrix. A matrix containing the variables in the model.
A	Matrix. A matrix containing the time variable data.
Risk	Matrix. An optional matrix that modify the probability of belong to group. By default its value is a matrix with one column with value 1.
mean	an optional logical. Indicate if the mean of ech group and time value must be draw.
alpha	on optional real. Indicate the alpha channel of the points color.
...	optional parameters

**Value**

a graphic.

---

```
plotrajeR.Trajectory.LOGIT
      plot LOGIT trajectory
```

---

**Description**

plot LOGIT trajectory

**Usage**

```
## S3 method for class 'Trajectory.LOGIT'
plotrajeR(
  Obj,
  plotcov = NULL,
  dec = 0,
  col = "black",
  Y = NULL,
  A = NULL,
  Risk = NULL,
  mean = FALSE,
  alpha = 1,
  ...
)
```

**Arguments**

Obj	an object of class "Trajectory.LOGIT".
plotcov	an optional vector or matrix with the same length as the time period. Default value is NULL.
dec	an optional real. It precise the shift to draw the data points.
col	an optional vector. The vector of colors. It must contain a color for each trajectory and each points of groups. Its length is the double of the number of group. Default value is a grayscale.
Y	Matrix. A matrix containing the variables in the model.
A	Matrix. A matrix containing the time variable data.
Risk	Matrix. An optional matrix that modify the probability of belong to group. By default its value is a matrix with one column with value 1.
mean	an optional logical. Indicate if the mean of ech group and time value must be draw.
alpha	on optional real. Indicate the alpha channel of the points color.
...	optional parameters

**Value**

a graphic.

---

```
plotrajeR.Trajectory.NL
  plot Non Linear trajectory
```

---

**Description**

plot Non Linear trajectory

**Usage**

```
## S3 method for class 'Trajectory.NL'
plotrajeR(
  Obj,
  plotcov = NULL,
  col = "black",
  Y = NULL,
  A = NULL,
  Risk = NULL,
  mean = FALSE,
  alpha = 1,
  TCOV = NULL,
  ...
)
```

**Arguments**

Obj	an object of class "Trajectory.LOGIT".
plotcov	an optional vector or matrix with the same length as the time period. Default value is NULL.
col	an optional vector. The vector of colors. It must contain a color for each trajectory and each points of groups. Its length is the double of the number of group. Default value is a grayscale.
Y	Matrix. A matrix containing the variables in the model.
A	Matrix. A matrix containing the time variable data.
Risk	Matrix. An optional matrix that modify the probability of belong to group. By default its value is a matrix with one column with value 1.
mean	an optional logical. Indicate if the mean of ech group and time value must be draw.
alpha	on optional real. Indicate the alpha channel of the points color.
TCOV	Matrix. An optional matrix containing the time covariate that influence the trajectory themselves. By default its value is NULL.
...	optional parameters

**Value**

a graphic.

---

```
plotrajeR.Trajectory.POIS
  plot POIS trajectory
```

---

## Description

plot POIS trajectory

## Usage

```
## S3 method for class 'Trajectory.POIS'
plotrajeR(
  Obj,
  plotcov = NULL,
  dec = 0,
  col = "black",
  Y = NULL,
  A = NULL,
  Risk = NULL,
  TCOV = NULL,
  mean = FALSE,
  alpha = 1,
  ...
)
```

## Arguments

Obj	an object of class "Trajectory.POIS".
plotcov	an optional vector or matrix with the same length as the time period. Default value is NULL.
dec	an optional real. It precise the shift to draw the data points.
col	an optional vector. The vector of colors. It must contain a color for each trajectory and each points of groups. Its length is the double of the number of group. Default value is a grayscale.
Y	Matrix. A matrix containing the variables in the model.
A	Matrix. A matrix containing the time variable data.
Risk	Matrix. An optional matrix that modify the probability of belong to group. By default its value is a matrix with one column with value 1.
TCOV	Matrix. An optional matrix containing the time covariate that influence the trajectory themselves. By default its value is NULL.
mean	an optional logical. Indicate if the mean of ech group and time value must be draw.
alpha	on optional real. Indicate the alpha channel of the points color.
...	optional parameters

**Value**

a graphic.

---

```
plotrajeR.Trajectory.ZIP
  plot ZIP trajectory
```

---

**Description**

plot ZIP trajectory

**Usage**

```
## S3 method for class 'Trajectory.ZIP'
plotrajeR(
  Obj,
  plotcov = NULL,
  dec = 0,
  col = "black",
  Y = NULL,
  A = NULL,
  Risk = NULL,
  TCOV = NULL,
  mean = FALSE,
  alpha = 1,
  ...
)
```

**Arguments**

Obj	an object of class "Trajectory.LOGIT".
plotcov	an optional vector or matrix with the same length as the time period. Default value is NULL.
dec	an optional real. It precise the shift to draw the data points.
col	an optional vector. The vector of colors. It must contain a color for each trajectory and each points of groups. Its length is the double of the number of group. Default value is a grayscale.
Y	Matrix. A matrix containing the variables in the model.
A	Matrix. A matrix containing the time variable data.
Risk	Matrix. An optional matrix that modify the probability of belong to group. By default its value is a matrix with one column with value 1.
TCOV	Matrix. An optional matrix containing the time covariate that influence the trajectory themselves. By default its value is NULL.

mean	an optional logical. Indicate if the mean of each group and time value must be drawn.
alpha	an optional real. Indicate the alpha channel of the points color.
...	optional parameters

**Value**

a graphic.

---

```
print.Trajectory.BETA Print BETA
```

---

**Description**

Print method for an object of class "Trajectory.BETA".

**Usage**

```
## S3 method for class 'Trajectory.BETA'
print(x, ...)
```

**Arguments**

x	Trajectory's object. An object of class "Trajectory.BETA".
...	optional parameters

**Value**

The print of Obj.

**Examples**

```
data <- read.csv(system.file("extdata", "BETA2gr.csv", package = "trajeR"))
data <- as.matrix(data)
data[, 2:6] <- data[, 2:6] * (nrow(data[, 2:6]) - 1 + 0.5) / nrow(data[, 2:6])
sol <- trajeR(
  Y = data[, 2:6], A = data[, 7:11], itermax = 50,
  degree = c(2, 2), degree.phi = c(1, 1), Model = "BETA", Method = "L"
)
sol
```

---

```
print.Trajectory.CNORM
      Print CNORM
```

---

**Description**

Print method for an object of class "Trajectory.CNORM".

**Usage**

```
## S3 method for class 'Trajectory.CNORM'
print(x, ...)
```

**Arguments**

x                   Trajectory's object. An object of class "Trajectory.CNORM".  
...                   optional parameters

**Value**

The print of Obj.

**Examples**

```
data <- read.csv(system.file("extdata", "CNORM2gr.csv", package = "trajeR"))
data <- as.matrix(data)
sol <- trajeR(Y = data[, 2:6], A = data[, 7:11], degree = c(2, 2), Model = "CNORM", Method = "EM")
sol
```

---

```
print.Trajectory.LOGIT
      Print LOGIT
```

---

**Description**

Print method for an object of class "Trajectory.LOGIT".

**Usage**

```
## S3 method for class 'Trajectory.LOGIT'
print(x, ...)
```

**Arguments**

x                   Trajectory's object. . An object of class "Trajectory.LOGIT".  
...                   optional parameters

**Value**

The print of Obj.

**Examples**

```
data <- read.csv(system.file("extdata", "LOGIT2gr.csv", package = "trajeR"))
data <- as.matrix(data)
sol <- trajeR(Y = data[, 2:6], A = data[, 7:11], degree = c(1, 2), Model = "LOGIT", Method = "L")
sol
```

---

```
print.Trajectory.NL   print NL trajectory
```

---

**Description**

Print method for an object of class "Trajectory.NL".

**Usage**

```
## S3 method for class 'Trajectory.NL'
print(x, ...)
```

**Arguments**

x                   Trajectory's object. . An object of class "Trajectory.NL".  
...                   optional parameters

**Value**

The print of Obj.

---

```
print.Trajectory.POIS Print POIS
```

---

**Description**

Print method for an object of class "Trajectory.POIS".

**Usage**

```
## S3 method for class 'Trajectory.POIS'
print(x, ...)
```

**Arguments**

x                   Trajectory's object. . An object of class "Trajectory.POIS".  
...                   optional parameters

**Value**

The print of Obj.

**Examples**

```
data <- read.csv(system.file("extdata", "POIS2gr.csv", package = "trajeR"))
data <- as.matrix(data)
sol <- trajeR(
  Y = data[, 2:6], A = data[, 7:11],
  degre = c(2, 2), Model = "POIS", Method = "L", hessian = FALSE
)
sol
```

---

`print.Trajectory.ZIP` *Print ZIP*

---

**Description**

Print method for an object of class "Trajectory.ZIP".

**Usage**

```
## S3 method for class 'Trajectory.ZIP'
print(x, ...)
```

**Arguments**

x                   Trajectory's object. An object of class "Trajectory.ZIP".  
...                   optional parameters

**Value**

The print of Obj.

**Examples**

```
data <- read.csv(system.file("extdata", "ZIP2gr.csv", package = "trajeR"))
data <- as.matrix(data)
sol <- trajeR(
  Y = data[, 2:6], A = data[, 7:11],
  degre = c(1, 2), degre.nu = c(1, 1), Model = "ZIP", Method = "L"
)
sol
```

---

propAssign	<i>Assignment proportion</i>
------------	------------------------------

---

**Description**

Calculate the proportion of individuals in a given group. That is the ratio of the number of individuals in one group and all the individuals.

**Usage**

```
propAssign(sol, Y, A)
```

**Arguments**

sol	Trajectory's object. An object of type Trajectory.
Y	Matrix. A matrix containing the variables in the model.
A	Matrix. A matrix containing the time variable data.

**Value**

A vector of real. The proportion.

**Examples**

```
data <- read.csv(system.file("extdata", "CNORM2gr.csv", package = "trajeR"))
data <- as.matrix(data)
sol <- trajeR(Y = data[, 2:6], A = data[, 7:11], degre = c(2, 2), Model = "CNORM", Method = "EM")
propAssign(sol, Y = data[, 2:6], A = data[, 7:11])
```

---

trajeR	<i>Fitting longitudinal mixture models</i>
--------	--

---

**Description**

trajeR is used to fit longitudinal mixture models. It used 3 types of mixture models : LOGIT, ZIP and censored Normal.

**Usage**

```
trajeR(
  Y,
  A,
  Risk = NULL,
  TCOV = NULL,
  degre = NULL,
```

```

degre.nu = 0,
degre.phi = 0,
Model,
Method = "L",
ssigma = FALSE,
ymax = max(Y, na.rm = TRUE) + 1,
ymin = min(Y, na.rm = TRUE) - 1,
hessian = TRUE,
itermax = 100,
paraminit = NULL,
ProbIRLS = TRUE,
refgr = 1,
fct = NULL,
diffct = NULL,
nbvar = NULL,
ng.nl = NULL,
nls.limiter = 50
)

```

### Arguments

Y	Matrix. A matrix containing the variables in the model.
A	Matrix. A matrix containing the time variable data.
Risk	Matrix. An optional matrix that modify the probability of belong to group. By default its value is a matrix with one column with value 1.
TCOV	Matrix. An optional matrix containing the time covariate that influence the trajectory themselves. By default its value is NULL.
degre	Vector of integer. The degree of every polynomial function.
degre.nu	Vector of integer. The degree of all Poisson part for a ZIP model.
degre.phi	Vector of integer. The degree of beta parameter for a BETA model.
Model	String. The model used. The value are LOGIT for a Logit Mixture model, CNORM for a Censored Normal Mixture Model or ZIP for Zero Inflated Poisson Mixture model.
Method	String. Determine the method used for find the parameters of the model. The value are L for the Maximum Likelihood Estimation, EM for Expectation Maximization method with quasi newton method inside, EMIWRLS for Expectation Maximization method with Iterative Weighted Least Square.
ssigma	Logical. By default its value is FALSE. For the CNORM model, indicate if we want the same sigma for all normal density function.
ymax	Real. For the CNORM model, indicate the maximum value of the data. It concern only the model with censored data. By default its value is the maximum value of the data plus 1.
ymin	Real. For the CNORM model, indicate the minimum value of the data. It concern only the model with censored data. By default its value is the maximum value of the data minus 1.

<code>hessian</code>	Logical. Indicate if we want calculate the hessian matrix. Default is FALSE. If the method use is Likelihood, the hessian is calculated by inverting the Information's Fisher Matrix. To avoid numerically singular matrix we find the pseudo inverse matrix by using the <code>ginv</code> function in the package MASS. If the method is EM or EMIWRLS, the hessian is calculated by using Louis method.
<code>itermax</code>	Integer. Indicate the maximal number of iteration for <code>optim</code> function or for the EM algorithm.
<code>paraminit</code>	Vector. The vector of initial parameters. By default <code>trajeR</code> calculate the initial value based of the range or the standard deviation.
<code>ProbIRLS</code>	Logical. Indicate the method to sue in the search of predictor's probability. If TRUE (by default) we use IRLS method and if FALSE we use optimization method.
<code>refgr</code>	Integer. The number of reference group. By default is 1.
<code>fct</code>	Function. The definition of the function <code>f</code> in the definition in nonlinear model.
<code>diffct</code>	Function. The differential of the function <code>f</code> in the nonlinear model.
<code>nbvar</code>	Integer. The number of variable in the nonlinear model.
<code>ng.nl</code>	Integer. The number of group for a non linear model.
<code>nls.limiter</code>	Integer. In the case of non linear model, the maximum number of iterations allowed.

## Details

Models for `trajeR` is, by default, a polynomial regression of the time value parameters for each groups. The number fo group is controlled by the integer `ng`. We can specify the degree of the polynomial shape for each groups by the vector `degre`.

## Value

return an object of class "Trajectory.LOGIT". The generic accessor functions `beta`, `delta`, `theta`, `sd`, `tab`, `Likelihood`, `ng`, `model` and `method` extract various useful features of the value returned by `trajeR`.

An object of class "Trajectory.LOGIT" is a list containing at least the following components:

`beta` a vector of the parameters `beta`.

`delta` a vector of the parameter `delta`. Only if we use time covariate.

`theta` a vector with the parameter `theta` if there exist a covariate `X` that modify the probability or the probability of group membership.

`sd` a vector of the standard deviation of the parameters.

`tab` a matrix with all the parameters and standard deviation.

`Likelihood` a real with the Likelihood obtained by the parameters.

`ng` a integer with the number of group.

`model` a string with the model used.

`method` a string with the method used.

**Examples**

```
data("dataNORM01")
soll <- trajeR(dataNORM01[, 1:5], dataNORM01[, 6:10],
  ng = 3, degre = c(2, 2, 2),
  Model = "CNORM", Method = "L", ssigma = FALSE,
  hessian = TRUE
)
```

---

trajeR.BETA

*Internal function to fit Beta regression*

---

**Description**

Internal function to fit Beta regression

**Usage**

```
trajeR.BETA(
  Y,
  A,
  X,
  TCOV,
  ng,
  nx,
  n,
  nbeta,
  nphi,
  nw,
  ntheta,
  period,
  degre,
  theta,
  beta,
  phi,
  delta,
  pi,
  Method,
  hessian,
  itermax,
  paraminit,
  EMIRLS,
  refgr
)
```

**Arguments**

Y	Matrix. A matrix containing the variables in the model.
A	Matrix. A matrix containing the time variable data.
X	Matrix. An optional matrix that modify the probability of belong to group. By default its value is a matrix with one column with value 1.
TCOV	Matrix. An optional matrix containing the time covariate that influence the trajectory themselves. By default its value is NULL.
ng	Integer. The number of groups.
nx	Integer. The number of covariates.
n	Integer. Number of individuals.
nbeta	Vector of integers. Number of beta parameters for each group.
nphi	Vector of integers. Number of phi parameters for each group.
nw	Integer. Number of time dependent covariate.
ntheta	Vector of integers. Number of theta parameters for each group.
period	Integer.
degre	Vector of integer. The degree of every polynomial function.
theta	Vector of real. The parameter for calculated the group membership probability.
beta	Vector of real. The beta parameter.
phi	Vector of real. The phi parameter.
delta	Vector of real. The delta parameter.
pi	Vector of real. The group membership probability.
Method	String. Determine the method used for find the parameters of the model. The value are L for the Maximum Likelihood Estimation, EM for Expectation Maximization method with quasi newton method inside, EMIWRLS for Expectation Maximization method with Iterative Weighted Least Square.
hessian	Logical. Indicate if we want calculate the hessian matrix. Default is FALSE. If the method use is Likelihood, the hessian is calculated by inverting the Information's Fisher Matrix. To avoid numerically singular matrix we find the pseudo inverse matrix by using the <code>ginv</code> function in the package MASS. If the method is EM or EMIWRLS, the hessian is calculated by using Louis method.
itermax	Integer. Indicate the maximal number of iteration for <code>optim</code> function or for the EM algorithm.
paraminit	Vector. The vector of initial parameters. By default trajeR calculate the initial value based of the range or the standard deviation.
EMIRLS	Boolean. True if we use EMIRLS method.
refgr	Integer. The number of reference group. By default is 1.

**Value**

return a object of class Trajectory.NL

- beta - vector of the parameter beta.
- sigma - vector of the parameters sigma.
- delta - vector of the parameter delta. Only if we use time covariate.
- theta - vector with the parameter theta if there exist a covariate X that modify the probability or the probability of group membership.
- sd - vector of the standard deviation of the parameters.
- tab - a matrix with all the parameters and standard deviation.
- Model - a string with the model used.
- groups - a integer with the number of group.
- Names - strings with the name of the parameters.
- Method - a string with the method used.
- Size - a integer with the number of individuals.
- Likelihood - a real with the Likelihood obtained by the parameters.
- Time - a vector with the first row of time values.
- degre - a vector with the degree of the polynomial shape.

---

trajeR.CNORM

*Internal function to fit CNORM Model*

---

**Description**

Internal function to fit CNORM Model

**Usage**

```
trajeR.CNORM(  
  Y,  
  A,  
  X,  
  TCOV,  
  ng,  
  nx,  
  n,  
  nbeta,  
  nw,  
  ntheta,  
  period,  
  degre,  
  theta,  
  beta,
```

```

    sigma,
    delta,
    pi,
    Method,
    sigma,
    ymax,
    ymin,
    hessian,
    itermax,
    paraminit,
    EMIRLS,
    refgr
)

```

### Arguments

Y	Matrix. A matrix containing the variables in the model.
A	Matrix. A matrix containing the time variable data.
X	Matrix. An optional matrix that modify the probability of belong to group. By default its value is a matrix with one column with value 1.
TCOV	Matrix. An optional matrix containing the time covariate that influence the trajectory themselves. By default its value is NULL.
ng	Integer. The number of groups.
nx	Integer. The number of covariates.
n	Integer. Number of individuals.
nbeta	Vector of integers. Number of beta parameters for each group.
nw	Integer. Number of time dependent covariate.
ntheta	Vector of integers. Number of theta parameters for each group.
period	Integer.
degre	Vector of integer. The degree of every polynomial function.
theta	Vector of real. The parameter for calculated the group membership probability.
beta	Vector of real. The beta parameter.
sigma	Vector of real. The sigma parameter.
delta	Vector of real. The delta parameter.
pi	Vector of real. The group membership probability.
Method	String. Determine the method used for find the parameters of the model. The value are L for the Maximum Likelihood Estimation, EM for Expectation Maximization method with quasi newton method inside, EMIWRLS for Expectation Maximization method with Iterative Weighted Least Square.
sigma	Logical. By default its value is FALSE. For the CNORM model, indicate if we want the same sigma for all normal density function.
ymax	Real. For the CNORM model, indicate the maximum value of the data. It concern only the model with censored data. By default its value is the maximum value of the data plus 1.

ymin	Real. For the CNORM model, indicate the minimum value of the data. It concern only the model with censored data. By default its value is the maximum value of the data minus 1.
hessian	Logical. Indicate if we want calculate the hessian matrix. Default is FALSE. If the method use is Likelihood, the hessian is calculated by inverting the Information's Fisher Matrix. To avoid numerically singular matrix we find the pseudo inverse matrix by using the <code>ginv</code> function in the package MASS. If the method is EM or EMIWRLS, the hessian is calculated by using Louis method.
itermax	Integer. Indicate the maximal number of iteration for <code>optim</code> function or for the EM algorithm.
paraminit	Vector. The vector of initial parameters. By default trajeR calculate the initial value based of the range or the standard deviation.
EMIRLS	Boolean. True if we use EMIRLS method.
refgr	Integer. The number of reference group. By default is 1.

### Value

return a object of class Trajectory.CNORM

- beta - vector of the parameter beta.
- sigma - vector of the parameters sigma.
- delta - vector of the parameter delta. Only if we use time covariate.
- theta - vector with the parameter theta if there exist a covariate X that modify the probability or the probability of group membership.
- sd - vector of the standard deviation of the parameters.
- tab - a matrix with all the parameters and standard deviation.
- Model - a string with the model used.
- groups - a integer with the number of group.
- Names - strings with the name of the parameters.
- Method - a string with the method used.
- Size - a integer with the number of individuals.
- Likelihood - a real with the Likelihood obtained by the parameters.
- Time - a vector with the first row of time values.
- degre - a vector with the degree of the polynomial shape.
- min - a real with the minimum value for censored data.
- max - a real with the maximum value for censored data.

trajeR.LOGIT

*Internal function to fit LOGIT Model***Description**

Internal function to fit LOGIT Model

**Usage**

```
trajeR.LOGIT(
  Y,
  A,
  X,
  TCOV,
  ng,
  nx,
  n,
  nbeta,
  nw,
  ntheta,
  period,
  degre,
  theta,
  beta,
  delta,
  pi,
  Method,
  hessian,
  itermax,
  paraminit,
  EMIRLS,
  refgr
)
```

**Arguments**

Y	Matrix. A matrix containing the variables in the model.
A	Matrix. A matrix containing the time variable data.
X	Matrix. An optional matrix that modify the probability of belong to group. By default its value is a matrix with one column with value 1.
TCOV	Matrix. An optional matrix containing the time covariate that influence the trajectory themselves. By default its value is NULL.
ng	Integer. The number of groups.
nx	Integer. The number of covariates.
n	Integer. Number of individuals.

nbeta	Vector of integers. Number of beta parameters for each group.
nw	Integer. Number of time dependent covariate.
ntheta	Vector of integers. Number of theta parameters for each group.
period	Integer.
degre	Vector of integer. The degree of every polynomial function.
theta	Vector of real. The parameter for calculated the group membership probability.
beta	Vector of real. The beta parameter.
delta	Vector of real. The delta parameter.
pi	Vector of real. The group membership probability.
Method	String. Determine the method used for find the parameters of the model. The value are L for the Maximum Likelihood Estimation, EM for Expectation Maximization method with quasi newton method inside, EMIWRLS for Expectation Maximization method with Iterative Weighted Least Square.
hessian	Logical. Indicate if we want calculate the hessian matrix. Default is FALSE. If the method use is Likelihood, the hessian is calculated by inverting the Information's Fisher Matrix. To avoid numerically singular matrix we find the pseudo inverse matrix by using the <code>ginv</code> function in the package MASS. If the method is EM or EMIWRLS, the hessian is calculated by using Louis method.
itermax	Integer. Indicate the maximal number of iteration for <code>optim</code> function or for the EM algorithm.
paraminit	Vector. The vector of initial parameters. By default trajeR calculate the initial value based of the range or the standard deviation.
EMIRLS	Boolean. True if we use EMIRLS method.
refgr	Integer. The number of reference group. By default is 1.

### Value

return a object of class Trajectory.LOGIT

- beta - vector of the parameter beta.
- delta - vector of the parameter delta. Only if we use time covariate.
- theta - vector with the parameter theta if there exist a covariate X that modify the probability or the probability of group membership.
- sd - vector of the standard deviation of the parameters.
- tab - a matrix with all the parameters and standard deviation.
- Model - a string with the model used.
- groups - a integer with the number of group.
- Names - strings with the name of the parameters.
- Method - a string with the method used.
- Size - a integer with the number of individuals.
- Likelihood - a real with the Likelihood obtained by the parameters.
- Time - a vector with the first row of time values.
- degre - a vector with the degree of the polynomial shape.

---

`trajeR.NL`*Internal function to fit Non Linear Model*

---

**Description**

Internal function to fit Non Linear Model

**Usage**

```
trajeR.NL(  
  Y,  
  A,  
  X,  
  TCOV,  
  ng,  
  nx,  
  n,  
  nbeta,  
  nw,  
  ntheta,  
  period,  
  degre,  
  theta,  
  beta,  
  sigma,  
  pi,  
  Method,  
  ssigma,  
  hessian,  
  itermax,  
  paraminit,  
  EMIRLS,  
  refgr,  
  fct,  
  diffct,  
  nls.limiter  
)
```

**Arguments**

Y	Matrix. A matrix containing the variables in the model.
A	Matrix. A matrix containing the time variable data.
X	Matrix. An optional matrix that modify the probability of belong to group. By default its value is a matrix with one column with value 1.
TCOV	Matrix. An optional matrix containing the time covariate that influence the trajectory themselves. By default its value is NULL.

ng	Integer. The number of groups.
nx	Integer. The number of covariates.
n	Integer. Number of individuals.
nbeta	Vector of integers. Number of beta parameters for each group.
nw	Integer. Number of time dependent covariate.
ntheta	Vector of integers. Number of theta parameters for each group.
period	Integer.
degre	Vector of integer. The degree of every polynomial function.
theta	Vector of real. The parameter for calculated the group membership probability.
beta	Vector of real. The beta parameter.
sigma	Vector of real. The sigma parameter.
pi	Vector of real. The group membership probability.
Method	String. Determine the method used for find the parameters of the model. The value are L for the Maximum Likelihood Estimation, EM for Expectation Maximization method with quasi newton method inside, EMIWRLS for Expectation Maximization method with Iterative Weighted Least Square.
ssigma	Logical. By default its value is FALSE. For the CNORM model, indicate if we want the same sigma for all normal density function.
hessian	Logical. Indicate if we want calculate the hessian matrix. Default is FALSE. If the method use is Likelihood, the hessian is calculated by inverting the Information's Fisher Matrix. To avoid numerically singular matrix we find the pseudo inverse matrix by using the <code>ginv</code> function in the package MASS. If the method is EM or EMIWRLS, the hessian is calculated by using Louis method.
itermax	Integer. Indicate the maximal number of iteration for <code>optim</code> function or for the EM algorithm.
paraminit	Vector. The vector of initial parameters. By default trajeR calculate the initial value based of the range or the standard deviation.
EMIRLS	Boolean. True if we use EMIRLS method.
refgr	Integer. The number of reference group. By default is 1.
fct	Function. The definition of the function f in the definition in nonlinear model.
diffct	Function. The differential of the function f in the nonlinear model.
nls.limiter	Integer. In the case of non linear model, the maximum number of iterations allowed.

### Value

return a object of class Trajectory.NL

- beta - vector of the parameter beta.
- sigma - vector of the parameters sigma.
- delta - vector of the parameter delta. Only if we use time covariate.

- theta - vector with the parameter theta if there exist a covariate X that modify the probability or the probability of group membership.
- sd - vector of the standard deviation of the parameters.
- tab - a matrix with all the parameters and standard deviation.
- Model - a string with the model used.
- groups - a integer with the number of group.
- Names - strings with the name of the parameters.
- Method - a string with the method used.
- Size - a integer with the number of individuals.
- Likelihood - a real with the Likelihood obtained by the parameters.
- Time - a vector with the first row of time values.
- degre - a vector with the degree of the polynomial shape.
- fct - the definition of the function used int this model.

---

trajeR.POIS

*Internal function to fit poisson Model*

---

## Description

Internal function to fit poisson Model

## Usage

```
trajeR.POIS(  
  Y,  
  A,  
  X,  
  TCOV,  
  ng,  
  nx,  
  n,  
  nbeta,  
  nw,  
  ntheta,  
  period,  
  degre,  
  theta,  
  beta,  
  delta,  
  pi,  
  Method,  
  hessian,  
  itermax,
```

```

    paraminit,
    EMIRLS,
    refgr
)

```

### Arguments

Y	Matrix. A matrix containing the variables in the model.
A	Matrix. A matrix containing the time variable data.
X	Matrix. An optional matrix that modify the probability of belong to group. By default its value is a matrix with one column with value 1.
TCOV	Matrix. An optional matrix containing the time covariate that influence the trajectory themselves. By default its value is NULL.
ng	Integer. The number of groups.
nx	Integer. The number of covariates.
n	Integer. Number of individuals.
nbeta	Vector of integers. Number of beta parameters for each group.
nw	Integer. Number of time dependent covariate.
ntheta	Vector of integers. Number of theta parameters for each group.
period	Integer.
degre	Vector of integer. The degree of every polynomial function.
theta	Vector of real. The parameter for calculated the group membership probability.
beta	Vector of real. The beta parameter.
delta	Vector of real. The delta parameter.
pi	Vector of real. The group membership probability.
Method	String. Determine the method used for find the parameters of the model. The value are L for the Maximum Likelihood Estimation, EM for Expectation Maximization method with quasi newton method inside, EMIWRLS for Expectation Maximization method with Iterative Weighted Least Square.
hessian	Logical. Indicate if we want calculate the hessian matrix. Default is FALSE. If the method use is Likelihood, the hessian is calculated by inverting the Information's Fisher Matrix. To avoid numerically singular matrix we find the pseudo inverse matrix by using the <code>ginv</code> function in the package MASS. If the method is EM or EMIWRLS, the hessian is calculated by using Louis method.
itermax	Integer. Indicate the maximal number of iteration for <code>optim</code> function or for the EM algorithm.
paraminit	Vector. The vector of initial parameters. By default trajeR calculate the initial value based of the range or the standard deviation.
EMIRLS	Boolean. True if we use EMIRLS method.
refgr	Integer. The number of reference group. By default is 1.

**Value**

return a object of class Trajectory.Pois

- beta - vector of the parameter beta.
- delta - vector of the parameter delta. Only if we use time covariate.
- theta - vector with the parameter theta if there exist a covariate X that modify the probability or the probability of group membership.
- sd - vector of the standard deviation of the parameters.
- tab - a matrix with all the parameters and standard deviation.
- Model - a string with the model used.
- groups - a integer with the number of group.
- Names - strings with the name of the parameters.
- Method - a string with the method used.
- Size - a integer with the number of individuals.
- Likelihood - a real with the Likelihood obtained by the parameters.
- Time - a vector with the first row of time values.
- degre - a vector with the degree of the polynomial shape for the Poisson part.

---

trajeR.ZIP

*Internal function to fit ZIP Model*

---

**Description**

Internal function to fit ZIP Model

**Usage**

```
trajeR.ZIP(  
  Y,  
  A,  
  X,  
  TCOV,  
  ng,  
  nx,  
  n,  
  nbeta,  
  nw,  
  ntheta,  
  period,  
  degre,  
  degre.nu,  
  theta,  
  beta,
```

```

    nu,
    delta,
    pi,
    Method,
    hessian,
    itermx,
    paraminit,
    EMIRLS,
    refgr
)

```

### Arguments

Y	Matrix. A matrix containing the variables in the model.
A	Matrix. A matrix containing the time variable data.
X	Matrix. An optional matrix that modify the probability of belong to group. By default its value is a matrix with one column with value 1.
TCOV	Matrix. An optional matrix containing the time covariate that influence the trajectory themselves. By default its value is NULL.
ng	Integer. The number of groups.
nx	Integer. The number of covariates.
n	Integer. Number of individuals.
nbeta	Vector of integers. Number of beta parameters for each group.
nw	Integer. Number of time dependent covariate.
ntheta	Vector of integers. Number of theta parameters for each group.
period	Integer.
degre	Vector of integer. The degree of every polynomial function.
degre.nu	Vector of integer. The degree of all Poisson part for a ZIP model.
theta	Vector of real. The parameter for calculated the group membership probability.
beta	Vector of real. The beta parameter.
nu	Vector of real. The nu parameter.
delta	Vector of real. The delta parameter.
pi	Vector of real. The group membership probability.
Method	String. Determine the method used for find the parameters of the model. The value are L for the Maximum Likelihood Estimation, EM for Expectation Maximization method with quasi newton method inside, EMIWRLS for Expectation Maximization method with Iterative Weighted Least Square.
hessian	Logical. Indicate if we want calculate the hessian matrix. Default is FALSE. If the method use is Likelihood, the hessian is calculated by inverting the Information's Fisher Matrix. To avoid numerically singular matrix we find the pseudo inverse matrix by using the <code>ginv</code> function in the package MASS. If the method is EM or EMIWRLS, the hessian is calculated by using Louis method.

itermax	Integer. Indicate the maximal number of iteration for optim function or for the EM algorithm.
paraminit	Vector. The vector of initial parameters. By default trajeR calculate the initial value based of the range or the standard deviation.
EMIRLS	Boolean. True if we use EMIRLS method.
refgr	Integer. The number of reference group. By default is 1.

**Value**

return a object of class Trajectory.ZIP

- beta - vector of the parameter beta.
- delta - vector of the parameter delta. Only if we use time covariate.
- theta - vector with the parameter theta if there exist a covariate X that modify the probability or the probability of group membership.
- nu - vector of the parameters nu.
- sd - vector of the standard deviation of the parameters.
- tab - a matrix with all the parameters and standard deviation.
- Model - a string with the model used.
- groups - a integer with the number of group.
- Names - strings with the name of the parameters.
- Method - a string with the method used.
- Size - a integer with the number of individuals.
- Likelihood - a real with the Likelihood obtained by the parameters.
- Time - a vector with the first row of time values.
- degre - a vector with the degree of the polynomial shape for the Poisson part.
- degre.nu - a vector with the degree of the polynomial shape for the exceeded zero state.

---

trajeRAIC

*AIC function to an trajectory object*


---

**Description**

Calculate the AIC value to an trajectory object.

**Usage**

```
trajeRAIC(sol)
```

**Arguments**

sol                   Trajectory's object. An object of type trajectory.

**Value**

A real.

**Examples**

```
data <- read.csv(system.file("extdata", "CNORM2gr.csv", package = "trajeR"))
data <- as.matrix(data)
sol <- trajeR(Y = data[, 2:6], A = data[, 7:11], degre = c(2, 2), Model = "CNORM", Method = "EM")
trajeRAIC(sol)
```

---

trajeRBIC

*BIC function to an trajectory object*

---

**Description**

Calculate the BIC value to an trajectory object.

**Usage**

```
trajeRBIC(sol)
```

**Arguments**

sol                   Trajectory's object. An object of type trajectory.

**Value**

A real.

**Examples**

```
data <- read.csv(system.file("extdata", "CNORM2gr.csv", package = "trajeR"))
data <- as.matrix(data)
sol <- trajeR(Y = data[, 2:6], A = data[, 7:11], degre = c(2, 2), Model = "CNORM", Method = "EM")
trajeRBIC(sol)
```

---

`trajeRSH`*SH function to an trajectory object*

---

**Description**

Calculate the Slope Heuristic value to a list of trajectory objects.

**Usage**

```
trajeRSH(l)
```

**Arguments**

l List. A list of objects of type trajectory.

**Value**

A vector of real.

**Examples**

```
data <- read.csv(system.file("extdata", "CNORM2gr.csv", package = "trajeR"))
data <- as.matrix(data)
degre <- list(c(2, 2), c(1, 1), c(1, 2), c(2, 1), c(0, 0),
c(0, 1), c(1, 0), c(0, 0), c(0, 2), c(2, 0))
sol <- list()
for (i in 1:10) {
  sol[[i]] <- trajeR(
    Y = data[, 2:6], A = data[, 7:11],
    degre = degre[[i]], Model = "CNORM", Method = "EM"
  )
}
trajeRSH(sol)
```

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