

Package ‘spm’

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Type Package

Title Stochastic Process Model (SPM)

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Description Stochastic Process Modeling

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Imports Rcpp (>= 0.11.1), RcppArmadillo (>= 0.4.200.0)

LinkingTo Rcpp, RcppArmadillo

Depends deSolve,mice,sas7bdat,RcppArmadillo

Suggests knitr

VignetteBuilder knitr

RoxygenNote 5.0.1

R topics documented:

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| | |
|--------------|--|
| prepare_data | <i>Data pre-processing for analysis with stochastic process model methodology.</i> |
|--------------|--|

Description

Data pre-processing for analysis with stochastic process model methodology.

Usage

```
prepare_data(longdat, vitstat, interval = 1, col.status = "IsDead",
  col.id = "ID", col.age = "Age", col.age.event = "LSmort",
  covariates = c("DBP", "BMI", "DBP1", "DBP2", "Weight", "Height"),
  verbose = TRUE)
```

Arguments

| | |
|---------------|--|
| longdat | A table with longitude records. |
| vitstat | A table with vital statistics (mortality). |
| interval | A number of breaks between observations for discrete simulation. Default = 1 (no breaks). |
| col.status | A name of column containing status variable (0/1 which indicate alive/dead). |
| col.id | A name of column containing patient ID. This ID should be the same in both longdat and vitstat tables. |
| col.age | A name of age column. |
| col.age.event | A name of event column. |
| covariates | A list of covariates. |
| verbose | A verbosing output indicator. Default=TRUE. |

Value

A list of two elements: first element contains a data table for continuous case, with arbitrary intervals between observations and second element contains a data table for a discrete case (fixed intervals between observations).

Examples

```
library(spm)
#Reading longitude data:
ldat <- read.csv(system.file("data", "longdat.csv", package="spm"))
# Prepare data for optimization:
vdat <- read.csv(system.file("data", "vitstat.csv", package="spm"))
data <- prepare_data(longdat=ldat, vitstat=vdat, interval=1, col.status="IsDead", col.id="ID", col.age="Age",
  # Parameters estimation:
  pars <- spm(data, k = 1)
pars
```

| | |
|-----------------|--|
| simdata_cont_1D | <i>One-dimensional simulation function for continuous time (arbitrary intervals between observations).</i> |
|-----------------|--|

Description

One-dimensional simulation function for continuous time (arbitrary intervals between observations).

Usage

```
simdata_cont_1D(N = 10, aH = -0.05, f1H = 80, QH = 2e-07, fH = 80,
  bH = 5, mu0H = 2e-05, thetaH = 0.08, step = 0.05, tstart = 30,
  tend = 105, ystart = 80, sd0 = 4)
```

Arguments

| | |
|--------|---|
| N | Number of individuals. |
| tstart | A number that defines starting time (30 by default). |
| tend | A number, defines final time (105 by default). |
| ystart | A vector with length equal to number of dimensions used, defines starting values of covariates. |
| a | A k by k matrix, which characterize the rate of the adaptive response. |
| f1 | A particular state, which if a deviation from the normal (or optimal). This is a vector with length of k. |
| Q | A matrix k by k, which is a non-negative-definite symmetric matrix. |
| f | A vector-function (with length k) of the normal (or optimal) state. |
| b | A diffusion coefficient, k by k matrix. |
| mu0 | mortality at start period of time. |
| theta | A displacement coefficient of the Gompertz function. |

Value

A table with simulated data.

Examples

```
library(spm)
dat <- simdata_cont_1D(N=2500)
dat
```

simdata_cont_MD

*Multi-dimensional simulation function for continuous trait.***Description**

Multi-dimensional simulation function for continuous trait.

Usage

```
simdata_cont_MD(N = 100, a = -0.05, f1 = 80, Q = 2e-07, f = 80,
  b = 5, mu0 = 2e-05, theta = 0.08, step = 0.05, tstart = 30,
  tend = 105, ystart = 80, sd0 = 4, k = 1)
```

Arguments

| | |
|--------|---|
| N | Number of individuals. |
| a | A k by k matrix, which characterize the rate of the adaptive response. |
| f1 | A particular state, which if a deviation from the normal (or optimal). This is a vector with length of k. |
| Q | A matrix k by k, which is a non-negative-definite symmetric matrix. |
| f | A vector-function (with length k) of the normal (or optimal) state. |
| b | A diffusion coefficient, k by k matrix. |
| mu0 | mortality at start period of time. |
| theta | A displacement coefficient of the Gompertz function. |
| tstart | A number that defines starting time (30 by default). |
| tend | A number, defines final time (105 by default). |
| ystart | A vector with length equal to number of dimensions used, defines starting values of covariates. |
| k | number of dimensions (k = 1 by default). |

Value

A table with simulated data.

Examples

```
library(spm)
dat <- simdata_cont_MD(N=2500)
dat
```

simdata_discr_MD

*Multi-dimension simulation function***Description**

Multi-dimension simulation function

Usage

```
simdata_discr_MD(N = 100, a = -0.05, f1 = 80, Q = 2e-08, f = 80,
  b = 5, mu0 = 1e-05, theta = 0.08, ystart = 80, tstart = 30,
  tend = 105, dt = 1, k = 1)
```

Arguments

| | |
|--------|---|
| N | Number of individuals |
| a | A k by k matrix, which characterize the rate of the adaptive response. |
| f1 | A particular state, which is a deviation from the normal (or optimal). This is a vector with length of k. |
| Q | A matrix k by k, which is a non-negative-definite symmetric matrix. |
| f | A vector-function (with length k) of the normal (or optimal) state. |
| b | A diffusion coefficient, k by k matrix. |
| mu0 | mortality at start period of time. |
| theta | A displacement coefficient of the Gompertz function. |
| ystart | A vector with length equal to number of dimensions used, defines starting values of covariates. |
| tstart | A number that defines starting time (30 by default). |
| tend | A number, defines final time (105 by default). |
| dt | A time step (1 by default). |
| k | number of dimensions (k = 1 by default). |

Value

A table with simulated data.

Examples

```
library(spm)
data <- simdata_discr_MD(N=1000, ystart=80, k=1)
head(data)
```

| | |
|------------------|---|
| simdata_time_dep | <i>Simulation function for continuous trait with time-dependant coefficients.</i> |
|------------------|---|

Description

Simulation function for continuous trait with time-dependant coefficients.

Usage

```
simdata_time_dep(N = 10, formulas = list(at = "-0.05", flt = "80", Qt =
  "2e-7*exp(0.08*t)", ft = "80", bt = "5", mu0t = "2e-5*exp(0.08*t)"),
  step = 0.05, tstart = 30, tend = 105, ystart = 80, sd0 = 4, k = 1)
```

Arguments

| | |
|----------|---|
| N | Number of individuals. |
| formulas | : a list of formulas that define age (time) - dependency. Default: list(at="a", flt="f1", Qt="Q*exp(theta*t)", ft="f", bt="b", mu0t="mu0*exp(theta*t)") |
| tstart | A number that defines starting time (30 by default). |
| tend | A number, defines final time (105 by default). |
| ystart | A starting value of covariates. |

Value

A table with simulated data.

Examples

```
library(spm)
dat <- simdata_time_dep(N=2500)
dat
```

| | |
|-----|--|
| spm | <i>Stochastic Process Modelling (SPM) A main function that estimates parameters a, fl, Q, f, b, mu0, theta from given dataset.</i> |
|-----|--|

Description

Stochastic Process Modelling (SPM) A main function that estimates parameters a, fl, Q, f, b, mu0, theta from given dataset.

Usage

```
spm(dat, k = 2, verbose = F, tol = NULL)
```

Arguments

| | |
|---------|---|
| dat | A dataset. |
| k | Number of dimensions. |
| verbose | A verbosing output indicator. |
| tol | A tolerance threshold for matrix inversion. |

Value

A list of (1) Estimated starting point (from quick discrete optimization) and (2) Estimated coefficients.

Examples

```
library(spm)
#Prepare data for optimization
longdat <- read.csv(system.file("data", "longdat.csv", package="spm"))
vitstat <- read.csv(system.file("data", "vitstat.csv", package="spm"))
data=prepare_data(longdat=longdat, vitstat=vitstat, interval=1, col.status="IsDead", col.id="ID", col.age="Age")
#Parameters estimation:
pars=spm(data,k = 1)
pars
```

| | |
|-------------------|--|
| spm_continuous_1D | <i>Continuous one-dimensional optimization</i> |
|-------------------|--|

Description

Continuous one-dimensional optimization

Usage

```
spm_continuous_1D(dat, a = -0.05, f1 = 80, Q = 2e-08, f = 80, b = 5,
  mu0 = 2e-05, theta = 0.08)
```

Arguments

| | |
|---------|---|
| dat | A data table. |
| a | A starting value of the rate of adaptive response to any deviation of Y from f1(t). |
| f1 | A starting value of the average age trajectories of the variables which process is forced to follow. |
| Q | Starting values of the quadratic hazard term. |
| f | A starting value of the "optimal" value of variable which corresponds to the minimum of hazard rate at a respective time. |
| b | A starting value of a diffusion coefficient representing a strength of the random disturbance from Wiener Process. |
| mu0 | A starting value of the baseline hazard. |
| theta | A starting value of the parameter theta (axe displacement of Gompertz function). |
| k | A number of dimensions. |
| verbose | An indicator of verbosing output. |
| tol | A tolerance threshold for matrix inversion. |

Details

spm_integral_1D runs much slower than discrete but more precise and can handle time intervals with different lengths.

Value

A set of estimated parameters a , f_1 , Q , f , b , μ_0 , θ .

Examples

```
library(spm)
# Reading the data:
longdat <- read.csv(system.file("data", "longdat.csv", package="spm"))
vitstat <- read.csv(system.file("data", "vitstat.csv", package="spm"))
dd <- prepare_data(longdat=longdat, vitstat=vitstat, interval=1, col.status="IsDead", col.id="ID", col.age="Age")
data <- dd[[1]][,2:6]
#Parameters estimation:
pars <- spm_continuous_1D(dat=data, a=-0.05, f1=80, Q=2e-8, f=80, b=5, mu0=2e-5, theta=0.08)
pars
```

spm_continuous_MD

Continuous multi-dimensional optimization

Description

Continuous multi-dimensional optimization

Usage

```
spm_continuous_MD(dat, a = 0.05, f1 = 80, Q = 2e-08, f = 81, b = 5,
  mu0 = 2e-05, theta = 0.08, k = 1, verbose = F)
```

Arguments

| | |
|----------------------|---|
| <code>dat</code> | A data table. |
| <code>a</code> | A starting value of the rate of adaptive response to any deviation of Y from $f_1(t)$. |
| <code>f1</code> | A starting value of the average age trajectories of the variables which process is forced to follow. |
| <code>Q</code> | Starting values of the quadratic hazard term. |
| <code>f</code> | A starting value of the "optimal" value of variable which corresponds to the minimum of hazard rate at a respective time. |
| <code>b</code> | A starting value of a diffusion coefficient representing a strength of the random disturbance from Wiener Process. |
| <code>mu0</code> | A starting value of the baseline hazard. |
| <code>theta</code> | A starting value of the parameter θ (axe displacement of Gompertz function). |
| <code>k</code> | A number of dimensions. |
| <code>verbose</code> | An indicator of verbosing output. |
| <code>tol</code> | A tolerance threshold for matrix inversion. |

Details

spm_integral_MD runs much slower than discrete but more precise and can handle time intervals with different lengths.

Value

A set of estimated parameters a , f_1 , Q , f , b , μ_0 , θ .

Examples

```
library(spm)
# Reading the data:
longdat <- read.csv(system.file("data", "longdat.csv", package="spm"))
vitstat <- read.csv(system.file("data", "vitstat.csv", package="spm"))
dd <- prepare_data(longdat=longdat, vitstat=vitstat, interval=1, col.status="IsDead", col.id="ID", col.age="Age")
data <- dd[[1]][,2:6]
#Parameters estimation:
pars <- spm_continuous_MD(dat=data, a=-0.05, f1=80, Q=2e-8, f=80, b=5, mu0=2e-5, theta=0.08, k = 1)
pars
```

| | |
|-----------------|--|
| spm_discrete_MD | <i>Discrete multi-dimensional optimization</i> |
|-----------------|--|

Description

Discrete multi-dimensional optimization

Usage

```
spm_discrete_MD(dat, k = 1, theta_range = seq(0.001, 0.09, by = 0.001),
  tol = NULL)
```

Arguments

| | |
|-------------|--|
| dat | A data table. |
| k | A number of dimensions. |
| theta_range | A range of θ parameter (axe displacement of Gompertz function), default: from 0.001 to 0.09 with step of 0.001. |
| tol | A tolerance threshold for matrix inversion (NULL by default). |

Details

This function is way much faster than continuous `spm_continuous_MD(...)` (but less precise) and used mainly in estimation a starting point for the `spm_continuous_MD(...)`.

Value

A list of two elements: (1) estimated parameters u , R , b , ϵ , Q , μ_0 , θ and (2) estimated parameters a , f_1 , Q , f , b , μ_0 , θ . Note: b and μ_0 from first list are different from b and μ_0 from the second list.

Examples

```
library(spm)
# Reading longitudinal data
longdat <- read.csv(system.file("data", "longdat.csv", package="spm"))
# Prepare data for optimization
vitstat <- read.csv(system.file("data", "vitstat.csv", package="spm"))
data <- prepare_data(longdat=longdat, vitstat=vitstat, interval=1, col.status="IsDead", col.id="ID", col.age=
# Parameters estimation
pars <- spm_discrete_MD(data[[2]], k=1, theta_range=seq(0.001, 0.09, by=0.001), tol=NULL)
pars
```

| | |
|--------------|---|
| spm_time_dep | <i>spm_time_dep : a function that can handle time-dependant coefficients:</i> |
|--------------|---|

Description

spm_time_dep : a function that can handle time-dependant coefficients:

Usage

```
spm_time_dep(data, start = list(a1 = -0.5, a2 = 0.2, f1 = 80, Q = 2e-08, f = 80, b = 5, mu0 = 1e-05, theta = 0.08), formulas = list(at = "a1*t+a2", f1t = "f1", Qt = "Q*exp(theta*t)", ft = "f", bt = "b", mu0t = "mu0*exp(theta*t)"), verbose = TRUE, lower_bound = NULL, upper_bound = NULL, factr = 1e-16, lmult = 0.5, umult = 2)
```

Arguments

| | |
|----------|---|
| start | : a list of starting parameters, default: llist(a=-0.5, f1=80, Q=2e-8, f=80, b=5, mu0=1e-5, theta=0.08), |
| formulas | : a list of formulas that define age (time) - dependency. Default: list(at="a1*t+a2", f1t="f1", Qt="Q*exp(theta*t)", ft="f", bt="b", mu0t="mu0*exp(theta*t)") |

Value

optimal coefficients

Examples

```
library(spm)
#Data preparation:
N <- 1000
data <- simdata_time_dep(N=2500)
opt.par <- spm_time_dep(data[,2:6], formulas=list(at="a", f1t="f1", Qt="Q*exp(theta*t)", ft="f", bt="b", mu0t="mu0*exp(theta*t)"))
opt.par
```

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