# spm: an R-package that implements a Stochastic Process Model (SPM)

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# Overview

The R-package spm (https://github.com/izhbannikov/spm) is developed for modeling aging-related changes and it allows (1) data simulation and (2) estimating the process parameters using maximum likelihood estimation by optimizing parameters used in the model. Specifically, developed R-package spm allows (i) one-dimensional SPM; (ii) multiple dimensional SPM; (iii) data simulation for one- and multiple dimensions.

#### Introduction

General description of age-related changes.

Model description

What we propose

# Data description

Data represents a typical clinical trait data and must be presented in form of two datasets: longitudinal dataset (follow-up studies), in which one record represents a single observation, and vital statistics, where one record represents all patient information. Longitudinal dataset must contain person ID (Identification number), status (dead(1)/alive(0)), time and measurements across the covariates. The spm can handle an infinite number of covariates but in practice, 5-7 covariates is enough.

Below there is an example of clinical data that can be used in spm and we will discuss the field later. Longitudinal studies:

##		X	ID	${\tt IndicatorDeath}$	Age	AgeNext	DBP	BMI
##	1	1	1	0	30	32	80.00000	25.00000
##	2	2	1	0	32	34	80.51659	26.61245
##	3	3	1	0	34	36	77.78412	29.16790
##	4	4	1	0	36	38	77.86665	32.40359
##	5	5	1	0	38	40	96.55673	31.92014
##	6	6	1	0	40	42	94.48616	32.89139

Vital statistics:

```
X ID IsDead
                   LSmort
## 1 1
       1
               1 85.34578
       2
               1 80.55053
## 3 3
       3
               1 98.07315
               1 81.29779
## 5 5
      5
               1 89.89829
## 6 6 6
               1 72.47687
```

# Data fields description

### Longitude studies

- ID subject unique identificatin number.
- IndicatorDeath 0/1, indicates death of a subject.
- Age current age of subjects.
- AgeNext next age of subject he will attend to the survey/exam.
- DBP, BMI covariates, here "DBP" represents a diastolic blood pressure, "BMI" a body-mass index.

#### Vital statistics

- ID subject's unique ID.
- IsDead death indicator, 0 alive, 1 dead.
- LSmort age at death of stopping observations.

# Discrete and Continuous cases

There are two main SPM types in the package: discrete model and continuous model. Discrete model assumes equal intervals between follow-up observations. The example of discrete dataset is given below.

```
library(spm)
data <- simdata_discr_MD(N=10, ystart=c(80), k=1)
head(data)</pre>
```

```
##
        id xi t1 t2
                      par1_1
                               par1 2
            0 30 31 80.00000 66.66658
  [2,]
         1
            0 31 32 66.66658 63.38874
## [3,]
         1
            0 32 33 63.38874 64.77483
            0 33 34 64.77483 67.15203
## [5,]
         1
            0 34 35 67.15203 74.06955
           0 35 36 74.06955 75.32344
## [6,]
         1
```

In this case there are equal intervals between t1 and t2 (Age and Age.next).

The opposite is continuous case, in which intervals between observations are not equal. The example of continuous case dataset is shown below:

```
library(spm)
data <- simdata_cont_MD(N=5,ystart = c(50))
head(data)</pre>
```

#### Discrete case

In discrete case, we use the following assumptions:

$$\bar{y}(t+1) = \bar{u} + \bar{R} \times \bar{y}(t) + \bar{\epsilon}$$
$$\mu(t) = \mu_0(t) + \bar{b}(t) \times \bar{y}(t) + \bar{Q} \times \bar{y}(t)^2$$

Where:

$$\mu_0(t) = \mu_0 e^{\theta t}$$
$$\bar{b}(t) = \bar{b}e^{\theta t}$$
$$\bar{Q}(t) = \bar{Q}e^{\theta t}$$

# Continuous case

$$\mu(u) = \mu_0(u) + (\bar{m}(u) - \bar{f}(u)^* \times \bar{Q}(u) \times (\bar{m}(u) - \bar{f}(u)) + Tr(\bar{Q}(u) \times \bar{\gamma}(u))$$

$$dm(t)/dt = \bar{a}(t) \times (\bar{m}(t) - \bar{f}_1(t)) - 2\bar{\gamma}(t) \times \bar{Q}(t) \times (\bar{m}(t) - \bar{f}(t))$$

$$d\bar{\gamma}(t)/dt = \bar{a}(t) \times \bar{\gamma}(t) + \bar{\gamma}(t) \times \bar{a}(t)^* + \bar{b}(t) \times \bar{b}(t)^* - 2\bar{\gamma}t \times \bar{Q}(t) \times \bar{\gamma}(t)$$

# Coefficient conversion between continuous and discrete cases

$$\begin{split} Q &= Q \\ \bar{a} &= \bar{R} - diag(k) \\ \bar{b} &= \bar{\epsilon} \\ \bar{f}1 &= -1 \times \bar{u} \times a^{-1} \\ \bar{f} &= -0.5 \times \bar{b} \times Q^{-1} \\ mu_0 &= mu_0 - \bar{f} \times \bar{Q} \times t(\bar{f}) \\ \theta &= \theta \end{split}$$

# Case with time-dependent coefficients

In two previous cases, we assumed that coefficients is sort of time-dependant: we multiplied them on to

$$e^{\theta t}$$

. In general, this may not be the case. We extend this to a general case, i.e. (we consider one-dimensional case):

$$a(t) = par_1t + par_2$$

- linear function.

The corresponding equations will be equivalent to one-dimensional continuous case described above.

# Simulation

We added one- and multi- dimensional simulation to be able to generate test data for hyphotesis testing. Data, which can be simulated can be discrete (equal intervals between observations) and continuous (with arbitrary intervals).

# Discrete

For discrete case:

```
library(spm)
data <- simdata_discr_MD(N=100, ystart=c(75, 94), k=2)
head(data)</pre>
```

```
##
        id xi t1 t2
                     par1_1
                              par1_2
                                         par2_1
                                                   par2_2
## [1,]
        1 0 30 31 75.00000 69.29038
                                      94.00000
                                                 96.27888
## [2,]
        1 0 31 32 69.29038 63.42717
                                                 94.22123
                                      96.27888
## [3,]
        1
           0 32 33 63.42717 54.07290
                                      94.22123
                                                 95.23642
## [4,]
        1 0 33 34 54.07290 49.58226
                                      95.23642 100.91389
        1 0 34 35 49.58226 47.11312 100.91389
        1 0 35 36 47.11312 40.63211 98.87588
## [6,]
                                                86.45995
```

# Continuous

For continuous case:

```
library(spm)
data <- simdata_cont_MD(N=100)
head(data)</pre>
```

```
## id xi t1 t2 y1 y1.next
## 1 1 0 30.97811 31.42534 85.74200 87.40471
## 2 1 0 31.42534 31.95067 87.40471 87.31503
## 3 1 0 31.95067 32.52584 87.31503 87.43195
## 4 1 0 32.52584 33.68741 87.43195 90.09190
## 5 1 0 33.68741 33.97101 90.09190 87.47674
## 6 1 0 33.97101 35.74413 87.47674 84.42130
```

# More Examples

[TODO]