Multiple decrements tables with lifecontingencies package

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Abstract

This paper introduces the mdt class within lifecontingencies R package, that handles multiple decrements models for actuarial applications. Applied examples will show how to perform demographic and actuarial calculation with the package.

Keywords: actuarial mathematics, multiple decrement models, lifecontingencies.

1. Introduction

1.1. Multiple Decrement Models within R

As of 2014 no R package provides easy tools to manage multiple decrement tables for actuarial applications. Multiple decrement tables are the basis of many applications, as demography, medicine and actuarial science.

Until now no R package provides a good tool to manage multiple decrement tables, even if Deshmukh (2012) provides an R based focus on multiple decrement tables with applications in R. The topic is deeply related to multistate analysis of life histories on which Willekens (2014) provide a very good introduction.

This paper introduces the mdt class that has been specifically engineered to manage multiple decrements models with R. Applied examples will follows.

1.2. A review of Discrete Multiple Decrement Mathematics

Following notation in Finan (2014), we provide definitions of the key quantities that allow to understand the main concepts regarding Multiple Decrement (MD) theory. Be $l_x^{(\tau)} = \sum_{j=1...m} l_x^{(j)}$ survivors to age x that will, at future ages, be fully depleted by m causes of decrement. $d_x^{(j)} = l_x^{(j)} - l_{x+1}^{(j)}$ represents the expected number of lives exiting from the population between ages x and x+1 due to decrement j. Therefore it follows that $nd_x^{(j)} = \sum_{t=0...n-1} d_{x+t}^{(j)}$. The probability that a life x will leave the group within one year as a result of decrement j is $nq_x^{(j)} = \frac{nd_x^{(j)}}{l_x^{(\tau)}}$. It follows that $q_x^{(\tau)} = \sum_{i=1}^m q_x^{(j)}$ and that $tq_x^{(\tau)} = 1 - tp_x^{(\tau)} = \sum_{j} tq_x^{(\tau)}$.

2. Multiple Decrement Models within lifecontingencies Package

2.1. The mdt class

Examples in this paper are worked on slides provided in Valdez (2011). First of all, we load the R package.

R> library(lifecontingencies)

Then we create a mdt class object. We can use the first example found on (?, p. 4).

Added fictional decrement below last x and completed x and lx until zero.... Completed the table at top, all decrements on first cause

The mdt class is an S4 class object (Chambers 2008) comprised by a character slot name and a data.frame slot table that is composed by following columns:

- 1. **x**: the age, from 0 to ω .
- 2. lx: the subject living (at risk) at the beginning of age.
- 3. one or more colums for different causes of decrements.

Values within table item represents absolute number of subjects at risk at the beginning of age x and dying for cause j during period x - x + 1.

Within the various methods defined within the mdt class, setValidity performs consistency checks to properly create the mdt object. In particular, it verifies whether:

- 1. \mathbf{x} and $\mathbf{l}\mathbf{x}$ exist and that they are consistent. \mathbf{x} should start from 0 and flows by increments of one. The first $\mathbf{l}\mathbf{x}$ value should be equal to the sum of all decrements and that $l_x = l_{x-1} (d_{x-1,1} + d_{x-1,2} + \ldots + d_{x-1,k})$ for any x.
- 2. If the decrements (or x and lx) have been provided only for partial ages, the table is completed below (from 0 to l_{x-1}) assuming a decrement rate of 0.01 for the first cause of death.
- 3. if the decrements at last provided age, ω , do not sum to l_{ω} , the table is incremented by one row such as $lx_{\omega+1} = lx_{\omega} (d_{\omega,1} + d_{\omega,2} + \ldots + d_{\omega,j})$.

As shown, when the table is sanitized the operations performed are reported on logs.

An internal function, .tableSanitizer tries to fix the limitations on the input table in order it to meet the class definition requirements.

Table can be viewed thanks to a print and show method (output omitted for simplicity). Similarly, it is possible to export a mdt to a data.frame or to a markovchainList object (from markovchain package).

```
R> print(valdezMdt)

R> valdezDf<-as(valdezMdt, "data.frame")
R> require(markovchain)
R> valdezMarkovChainList<-as(valdezMdt, "markovchainList")</pre>
```

Two specific methods have been defined for mdt class objects: getOmega, that returns the maximum attainable age (similar to the one of lifetable class), and getDecrements, that returns the decrements (by means of the names within table slot different from x and lx).

```
[1] 55
R> getDecrements(valdezMdt)
[1] "hearth" "accidents" "other"
A summary method is available as well.
R> summary(valdezMdt)
This is Multiple Decrements Table: ValdezExample Omega age is: 55
Stored decrements are: hearth accidents other
```

R> getOmega(valdezMdt)

2.2. Decrement probabilities calculation

The lifecontingencies package makes easy to compute $d_x^{(j)}$, $_nd_x^{(j)}$ as well as $_nd_x^{(\tau)}$ quantities thanks to dxt function.

```
R> dxt(valdezMdt,x=51,decrement="other")
[1] 5162
R> dxt(valdezMdt,x=51,t=2, decrement="other")
[1] 11122
```

```
R> dxt(valdezMdt,x=51)
```

[1] 11731

Probabilities could be computed as well.

```
R> dxt(valdezMdt,x=51,t=2, decrement="other")
```

[1] 11122

R> pxt(valdezMdt,x=50,t=3)

[1] 0.9926809

R> qxt(valdezMdt, x=53, t=2, decrement=1)

[1] 0.002544409

It is possible to generate random traiectories of a life subject to multiple cause of decrements as the following code shows.

```
R > rmdt(n = 2, object = valdezMdt, x = 50, t = 2)
```

```
1 2
50 "alive" "alive"
51 "alive" "alive"
```

52 "alive" "alive"

3. Actuarial Applications

TO BE WRITTEN

Acknowledgments

TBD.

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