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Discretizing the Distribution of a Population following a Discrete-time Markov Chain on a Continuous State Space

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Abstract

Package discretizeCtsDTMC creates a discrete approximation to a Markov process defined on a continuous state space in discrete time. Once the state space is discretized, discretizeCtsDTMC provides tools to estimate the transition matrices and analyze the Markov process. It is used to model a population of individuals, each following a continuous-state Markov process in discrete time.

We'll choose a better name later.

Keywords: Markov chain, Markov process, discretization.

1. Markov Chains in R

This article illustrates how to create a discrete approximation to the distribution function of a population of individuals following a Markov process defined on a continuous state space in discrete time. Once the state space is discretized, **discretizeCtsDTMC** provides tools to estimate the transition matrices and analyze the Markov process. It is used to model a population of individuals, each following a continuous-state Markov process in discrete time.

There are several R packages (R Core Team 2017) available for working with Markov processes. The packages **markovchain** (Spedicato, Kang, Yalamanchi, Yadav, and Cordon 2020) and **DTMCPack** (Nicholson 2013) provide tools for basic computations with Markov chains. The **mcmc** is designed for working with Monte Carlo Markov Chains. Packages **HMM** (Geyer and Johnson 2020) and **depmixS4** (Visser and Speekenbrink 2010) are designed for fitting Hidden Markov Models.

There are several packages aimed at specific applications of Markov chains. Some of these include packages designed for application in health care. There is **TPmsm** (Araújo, Meira-

Machado, and Roca-Pardiñas 2014) for estimating transition probabilities for 3-state progressive disease models and **heemod** (Filipović-Pierucci, Zarca, and Durand-Zaleski 2017) for applying Markov models to health care economic applications. Aimed at specific applications, some of these packages assume a considerable knowledge of the relevant subject matter and theory behind those applications.

The packages msm (Jackson 2011) and SemiMarkov (Król and Saint-Pierre 2015) are used for fitting multistate models to panel data, along with mstate (de Wreede, Fiocco, and Putter 2011) for survival analysis applications. These packages are designed to model discrete-state Markov processes in continuous time, with transitions taking place at a stochastic arrival times.



Figure 1: Caption goes here

2. Model

Harry, put all that good stuff here.

3. Example

A demonstration of analysis is shown in discCtsDTMC_demo.R and it serves as an example of what a typical session of model specification, estimation and testing can include. This procedure includes the following steps:

- 1. Organizing data
- 2. Choosing estimation options
- 3. Lag selection
- 4. Model estimation
- 5. Hypothesis testing
- 3.1. Organizing data
- 3.2. Choosing options
- 3.3. Lag-order selection
- 3.4. Model estimation
- 3.5. Hypothesis testing

4. Summary and discussion

This is a good package because...

Computational details

The results in this paper were obtained using R 3.5.1. with the **discreteCtsDTMC** package Version 0.0.0.9000. R itself and all packages used are available from the Comprehensive R Archive Network (CRAN) at https://CRAN.R-project.org/.

The development version of this package is available by using the **devtools** package, with which the latest version can be installed by

devtools::install_github(LeeMorinUCF/discreteCtsDTMC).

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