# tmon-p1

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## 1 Montecarlo Techniques - Practice 1

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## 1.1 Old car simulation process

### 1.1.1 Implement the simulation model of the old car problem in R or python.

```
[1]: import numpy as np
     import random
     ### Implement the simulation model of the old car problem in R or python.
     def dtmc(N = 10**4, transitionMatrix = [[1, 0, 0, 0],
                                                 [0.05, 0.50, 0.20, 0.25],
                                                 [0.10, 0.75, 0, 0.15],
                                                 [0, 0, 0, 1]],
               initialState = 1, finalState = 0):
         Simulate a Discrete-Time Markov Chain (DTMC).
         Parameters:
          - N (int): The number of steps to simulate the Markov Chain. Default is \sqcup
      →1000.
          - transitionMatrix (list of lists): The transition matrix representing the ⊔
      \hookrightarrowDTMC. Default is a 4x4 matrix.
          - initialState (int): The initial state from which the simulation starts.
       \hookrightarrow Default is state 1.
          - finalState (int): The absorption state, if any, at which the simulation ⊔
       ⇒should stop. Default is state 0.
         Returns:
          - t: A list which contains the number of steps from the initialState to the _{\sqcup}
       \hookrightarrow finalState in each iteration.
```

```
- p: A list which indicates whether the iteration has reach to the _{\! \sqcup}
\hookrightarrow finalState (1) or not (0).
   - states: A list that saves all the states it has gone through since the 
⇔initial state for each iteration.
   # list where we will save our states
  states = [initialState]
  np.random.seed(123)
                                 # final state reached.
  p = np.zeros(N)
  t = np.full(N,np.nan)
                                       # simulation time.
  states = [[] for _ in range(N)]
                                       # states.
  # check if we have an absorption state
  absorption_state = []
  for tm in range(0,len(transitionMatrix)):
      for m in transitionMatrix[tm]:
           if (m == 1)&(transitionMatrix[tm].index(m) == tm):
               absorption_state.append(tm)
  # simulation
  for n in range(N):
      i = initialState
      states[n].append(i)
      while (i not in absorption_state):
           j = -1; # columns of our transition matrix (this is the 'next_\subseteq
⇔state' column)
           pj = 0; # transition probability from state 'i' to state 'j'.
           threshold = np.random.uniform(0, 1)
           while (pj < threshold):</pre>
               j += 1
               pj = pj + transitionMatrix[i][j]
           i = j
           states[n].append(i)
           if i == finalState:
               t[n] = len(states[n])-1
               p[n] += 1
  return(t,p,states)
```

### [2]: help(dtmc)

```
Help on function dtmc in module __main__:

dtmc(N=10000, transitionMatrix=[[1, 0, 0, 0], [0.05, 0.5, 0.2, 0.25], [0.1, 0.75, 0, 0.15], [0, 0, 0, 1]], initialState=1, finalState=0)
    Simulate a Discrete-Time Markov Chain (DTMC).
```

#### Parameters:

- N (int): The number of steps to simulate the Markov Chain. Default is 1000.
- transitionMatrix (list of lists): The transition matrix representing the DTMC. Default is a 4x4 matrix.
- initial $\operatorname{State}$  (int): The initial state from which the simulation starts. Default is state 1.
- finalState (int): The absorption state, if any, at which the simulation should stop. Default is state 0.

#### Returns:

- $\mbox{-}$  t: A list which contains the number of steps from the initialState to the finalState in each iteration.
- p: A list which indicates whether the iteration has reach to the final $\operatorname{State}$  (1) or not (0).
- states: A list that saves all the states it has gone through since the initial state for each iteration.

#### 1.1.2 Example of use and some metrics

The number of steps consumed to reach to the final state in each iteration is [nan 24. nan ... nan nan nan]

The times that our function has reached to the final state is 19887.0 in 100000 simulations

```
The history of states in 100000 simulations is [[1, 2, 1, 1, 2, 1, 1, 3], [1, 2, 1, 1, 1, 2, 1, 1, 1, 2, 1, 1, 1, 2, 1, 2, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 3], [1, 3], [1, 1, 2, 1, 1, 1, 3]]
```

- 1.1.3 Approximate the absorption time from Operativo to Siniestrado through simulation (years).
- [4]: np.nanmean(dtmc(N=N, initialState = 1, finalState = 0)[0])
- [4]: 3.571227435007794
  - 1.1.4 Approximate through simulation the probability of absorption from Averiado to Vendido (%).
- [5]: round(sum(dtmc(N=N,initialState = 2, finalState = 3)[1])\*100/N,4)
- [5]: 75.168