# Microsimulation Sick-Sicker model with time dependency with PSA

Includes individual characteristics: age, age dependent mortality probabilities, individual treatment effect modifyer, time dependency for the sick (S1) state, increasing change of death in the first 6 year of sickness (tunnel)

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- Jalal H, Pechlivanoglou P, Krijkamp E, Alarid-Escudero F, Enns E, Hunink MG. An Overview of R in Health Decision Sciences. Med Decis Making. 2017; 37(3): 735-746. https://journals.sagepub.com/doi/abs/10.1177/0272989X16686559
- Krijkamp EM, Alarid-Escudero F, Enns EA, Jalal HJ, Hunink MGM, Pechlivanoglou P. Microsimulation modeling for health decision sciences using R: A tutorial. Med Decis Making. 2018;38(3):400–22. https://journals.sagepub.com/doi/abs/10.1177/0272989X18754513
- Krijkamp EM, Alarid-Escudero F, Enns E, Pechlivanoglou P, Hunink MM, Jalal H. A Multidimensional Array Representation of State-Transition Model Dynamics. Med Decis Making. 2020 Online first. https://doi.org/10.1177/0272989X19893973

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```
rm(list = ls())  # clear memory (removes all the variables from the workspace)
```

## 01 Load packages

```
if (!require('pacman')) install.packages('pacman'); library(pacman) # use this package to conveniently
# load (install if required) packages from CRAN
p_load("here", "dplyr", "devtools", "scales", "ellipse", "ggplot2", "lazyeval", "igraph", "r
# load (install if required) packages from GitHub
# install_github("DARTH-git/dampack", force = TRUE) Uncomment if there is a newer version
p_load_gh("DARTH-git/dampack")
```

#### 02 Load functions

```
source(here("functions", "Functions.R"))
```

# 03 Input model parameters

```
set.seed(1) # set the seed
# Model structure
n_t <- 30
                                  # time horizon, 30 cycles
n_i <- 100000
                                  # number of simulated individuals
v_n <- c("H", "S1", "S2", "D") # the model states names
n_s <- length(v_n)
                                  # the number of health states
                                  # discount rate of 3% per cycle
d r
     <- 0.03
v_dwe \leftarrow v_dwc \leftarrow 1 / ((1 + d_r) ^ (0:n_t))
                                             # discount weight
v_names_str <- c("no treatment", "treatment") # strategy names</pre>
n_str <- length(v_names_str)</pre>
                                 # number of strategies
### Event probabilities (per cycle)
# Annual transition probabilities
      <- 0.15
                                  # probability of becoming sick when healthy
p_HS1
p S1H <- 0.5
                                  # probability of recovering to healthy when sick
p S1S2 <- 0.105
                                  # probability of becoming sicker when sick
# Annual probabilities of death
# load age dependent probability
p_mort <- read.csv(here("data", "mortProb_age.csv"))</pre>
# load age distribution
dist_Age <- read.csv(here("data", "MyPopulation-AgeDistribution.csv"))</pre>
# probability to die in S1 by cycle
        <- c(0.0149, 0.018, 0.021, 0.026, 0.031, rep(0.037, n_t - 5))
p_S1D
p_S2D
         <- 0.048
                            # probability to die in S2
# Cost inputs
сН
       <- 2000
                            # cost of one cycle in the healthy state
c_S1
       <- 4000
                            # cost of one cycle in the sick state
c_S2 <- 15000
                            # cost of one cycle in the sicker state
```

```
c_D <- 0
                           # cost of one cycle in the dead state
       <- 12000
                           # cost of treatment (per cycle)
c_Trt
# Utility inputs
      <- 1
                           # utility when healthy
u_H
       <- 0.75
u_S1
                           # utility when sick
u_S2
      <- 0.5
                           # utility when sicker
       <- 0
u_D
                           # utility when dead
u_Trt <- 0.95
                           # utility when sick(er) and being treated
```

# 04 Sample individual level characteristics

#### 04.1 Static characteristics

```
# your turn
```

### 04.2 Dynamic characteristics

```
# your turn
```

#### 05 Define Simulation Functions

#### 05.1 Probability function

The function that updates the transition probabilities of every cycle is shown below.

```
# your turn
```

#### 05.2 Cost function

The Costs function estimates the costs at every cycle.

```
# your turn
```

#### 05.3 Health outcome function

The Effs function to update the utilities at every cycle.

```
# your turn
```

#### 06 Run Microsimulation

```
# your turn
```

### 07 Visualize results

```
# your turn
```

08	Cost	Effectiveness	Anal	lysis
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# your turn

# 09 Probabilistic Sensitivity Analysis (PSA)

# your turn

#### 09.1 Load function of microsimulation model

# your turn

#### 09.2 Run microsimulation model on each parameter set of PSA input dataset

# your turn

#### 09.3 Cost Effectiveness Analysis

Vector with willingness-to-pay (WTP) thresholds

# your turn

#### 09.3.1 ICER

# your turn

### 09.3.2 Cost-Effectiveness Acceptability Curves (CEAC) and Frontier (CEAF)

# your turn

#### 09.3.3 Cost-Effectiveness Scatter plot

# your turn

### 09.3.4 Expected Value of Perfect Information (EVPI)

# your turn