# Microsimulation Sick-Sicker model with time dependency

Includes individual characteristics: age, age dependent mortality, individual treatment effect modifer, state-residency for the sick (S1) state, increasing change of death in the first 6 year of sickness

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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 EA, et al. A Multidimensional Array Representation of State-Transition Model Dynamics. Med Decis Mak. https://doi.org/10.1177/0272989X19893973

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Change eval to TRUE if you want to knit this document.

## 01 Load packages

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

### 02 Load functions

```
# No functions needed
```

## 03 Model input

```
## General setup
set.seed(1)
                          # set the seed
cycle_length <- 1
                          # cycle length equal to one year (use 1/12 for monthly)
                          # time horizon, number of cycles
n_cycles
             <- 30
               <- 100000 # number of individuals
n_i
# the 4 health states of the model:
v_names_states <- c("H", # Healthy (H)</pre>
                    "S1", # Sick (S1)
                    "S2", # Sicker (S2)
                    "D") # Dead (D)
               <- length(v_names_states) # number of health states</pre>
n states
### Discounting factors
d_c <- 0.03 # annual discount rate for costs</pre>
d_e <- 0.03 # annual discount rate for QALYs</pre>
### Strategies
v_names_str <- c("Standard of care", # store the strategy names
                   "Strategy AB")
             <- length(v_names_str) # number of strategies</pre>
n_str
# (all non-probabilities are conditional on survival)
              <- 0.15 # probability of becoming sick when healthy
p_HS1
p_S1H
              <- 0.5
                          # probability of recovering to healthy when sick
p_S1S2_SoC
             <- 0.105 # probability of becoming sicker when sick under standard of care
p_S1S2_trtAB <- 0.05
                          # probability of becoming sicker when sick under treatment AB
# Annual probabilities of death
```

```
# load age dependent probability
p_mort <- read.csv("mortProb_age.csv")</pre>
# load age distribution
dist Age <- read.csv("MyPopulation-AgeDistribution.csv")</pre>
# probability to die in S1 by cycle (is increasing)
p_S1D \leftarrow c(0.0149, 0.018, 0.021, 0.026, 0.031, rep(0.037, n_cycles - 5))
p_S2D <- 0.048 # probability to die in S2
### State rewards
#### Costs
c_H <- 2000 # annual cost of being Healthy
c_S1 <- 4000 # annual cost of being Sick
       <- 15000 # annual cost of being Sicker
c_D <- 0 # annual cost of being dead</pre>
c_trtAB <- 25000 # annual cost of receiving treatment AB when in Sick
#### Utilities
uН
      <- 1 # annual utility of being Healthy
u_S1 <- 0.75 # annual utility of being Sick
u_S2 <- 0.5 # annual utility of being Sicker
u D <- 0 # annual utility of being dead
u trtAB <- 0.95 # annual utility when receiving treatment AB when in Sick
### Discount weight for costs and effects
v_dwc <- 1 / ((1 + (d_e * cycle_length)) ^ (0:n_cycles))</pre>
v_{dwe} \leftarrow 1 / ((1 + (d_c * cycle_length)) ^ (0:n_cycles))
```

## 04 Sample individual level characteristics

#### 04.1 Static characteristics

```
v_x <- runif(n_i, min = 0.95, max = 1.05) # treatment effect modifier at baseline
# sample from age distribution an initial age for every individual
v_age0 <- sample(x = dist_Age$age, prob = dist_Age$prop, size = n_i, replace = TRUE)</pre>
```

#### 04.2 Dynamic characteristics

```
# Your turn
```

#### 04.3 Create a dataframe with the individual characteristics

```
# Your turn
```

### 05 Define Simulation Functions

### 05.1 Probability function

The Probs function 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

#### 05.4 The Microsimulation function

# Your turn

## 06 Run Microsimulation

# Your turn

### 07 Visualize results

```
# Standard of care
# Your turn
```

# Strategy AB
# Your turn

# 08 Cost-effectiveness analysis (CEA)

# Your turn

## CEA table in proper format
# Your turn

## CEA frontier
# Your turn