

Cohort State-Transition Models in R

The DARTH workgroup

Developed by the Decision Analysis in R for Technologies in Health (DARTH) workgroup:

Fernando Alarid-Escudero, PhD (1)

Eva A. Enns, MS, PhD (2)

M.G. Myriam Hunink, MD, PhD (3,4)

Hawre J. Jalal, MD, PhD (5)

Eline M. Krijkamp, MSc (3)

Petros Pechlivanoglou, PhD (6,7)

Alan Yang, MSc (7)

In collaboration of:

1. Division of Public Administration, Center for Research and Teaching in Economics (CIDE), Aguascalientes, Mexico
2. University of Minnesota School of Public Health, Minneapolis, MN, USA
3. Erasmus MC, Rotterdam, The Netherlands
4. Harvard T.H. Chan School of Public Health, Boston, USA
5. University of Pittsburgh Graduate School of Public Health, Pittsburgh, PA, USA
6. University of Toronto, Toronto ON, Canada
7. The Hospital for Sick Children, Toronto ON, Canada

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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*. Feb;40(2):242-248. <https://doi.org/10.1177/0272989X19893973>
- Alarid-Escudero, F., Krijkamp, E. M., Enns, E. A., Hunink, M. G. M., Pechlivanoglou, P., & Jalal, H. (2020). Cohort state-transition models in R: From conceptualization to implementation. *ArXiv:2001.07824v1*, 1-31. <http://arxiv.org/abs/2001.07824>

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Code of Appendix

Implements a time-independent Sick-Sicker cohort state-transition model (cSTM).

- *Standard of Care (SoC)*: current available care for the patients with the disease. This strategy reflects the natural history of the disease progression.
- *Strategy A*: treatment A is given to all sick patients, patients in the Sick and Sicker, but only improves the utility of those being sick.
- *Strategy B*: treatment B reduces disease progression from the Sick to Sicker states. However, it is not possible to distinguish those sick from sicker, and therefore all individuals in one of the two sick states get the treatment.
- *Strategy AB*: combines treatment A and treatment B. The disease progression is reduced, and Sick individuals have an improved utility.

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

```
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 install other packages
}
# load (install if required) packages from CRAN
pacman::p_load("dplyr", "data.table", "devtools", "scales", "ellipse", "ggplot2",
               "lazyeval", "igraph", "truncnorm", "ggraph", "reshape2", "knitr",
               "stringr", "gridExtra", "diagram", "dampack", "boot", "doParallel")
# load (install if required) packages from GitHub
# install_github("DARTH-git/darthtools", force = TRUE) Uncomment if there is a newer version
pacman::p_load_gh("DARTH-git/darthtools")
```

02 Load functions

```
source("Functions.R")
```

03 Input model parameters

```
# Strategy names
v_names_str <- c("Standard of care", # store the strategy names
                "Strategy A",
                "Strategy B",
                "Strategy AB")

# Markov model parameters
age      <- 25                # age at baseline
max_age  <- 100              # maximum age of follow up
n_cycles <- max_age - age    # time horizon, number of cycles
# the 4 states of the model:
v_names_states <- c("H", # Healthy (H)
                   "S1", # Sick (S1),
                   "S2", # Sicker (S2)
                   "D") # Dead (D)
# initial cohort distribution (everyone allocated to the "healthy" state)
v_m_init <- c("H" = 1,
              "S1" = 0,
              "S2" = 0,
              "D" = 0)

## Transition probabilities (per cycle), hazard ratios
r_HD      <- 0.002 # constant rate of dying when Healthy (all-cause mortality)
p_HS1     <- 0.15  # probability to become Sick when Healthy conditional on surviving
p_S1H     <- 0.5   # probability to become Healthy when Sick conditional on surviving
p_S1S2    <- 0.105 # probability to become Sicker when Sick conditional on surviving
hr_S1     <- 3     # hazard ratio of death in Sick vs Healthy
```

```

hr_S2      <- 10      # hazard ratio of death in Sicker vs Healthy

# Effectiveness of treatment B
hr_S1S2_trtB <- 0.6   # hazard ratio of becoming Sicker when Sick under treatment B

## State rewards
# Costs
c_H      <- 2000      # cost of remaining one cycle in Healthy
c_S1     <- 4000      # cost of remaining one cycle in Sick
c_S2     <- 15000     # cost of remaining one cycle in Sicker
c_D      <- 0         # cost of being dead (per cycle)
c_trtA   <- 12000     # cost of treatment A
c_trtB   <- 13000     # cost of treatment B
# Utilities
u_H      <- 1         # utility when Healthy
u_S1     <- 0.75      # utility when Sick
u_S2     <- 0.5       # utility when Sicker
u_D      <- 0         # utility when Dead
u_trtA   <- 0.95      # utility when being treated with A

n_str     <- length(v_names_str)      # number of strategies
n_states  <- length(v_names_states)   # number of states

# Discounting factors
d_c       <- 0.03                  # discount rate for costs
d_e       <- 0.03                  # discount rate for QA

# Discount weights for costs and effects
v_dw      <- 1 / (1 + d_c) ^ (0:n_cycles)
v_dwe     <- 1 / (1 + d_e) ^ (0:n_cycles)

# Within-cycle correction (WCC) using Simpson's 1/3 rule
v_wcc     <- darthtools::gen_wcc(n_cycles = n_cycles,
                                method = "Simpson1/3") # vector of wcc

### Process model inputs
## Transition probabilities to the Dead state
# compute mortality rates
r_S1D     <- r_HD * hr_S1          # Mortality in the Sick state
r_S2D     <- r_HD * hr_S2          # Mortality in the Sick state
# transform rates to probabilities
p_HD      <- rate_to_prob(r_HD)    # Mortality risk in the Healthy state
p_S1D     <- rate_to_prob(r_S1D)   # Mortality risk in the Sick state
p_S2D     <- rate_to_prob(r_S2D)   # Mortality risk in the Sicker state

## Transition probability of becoming Sicker when Sick for treatment B
# transform probability to rate
r_S1S2     <- prob_to_rate(p = p_S1S2)
# apply hazard ratio to rate to obtain transition rate of becoming Sicker when
# Sick for treatment B
r_S1S2_trtB <- r_S1S2 * hr_S1S2_trtB
# transform rate to probability
p_S1S2_trtB <- rate_to_prob(r = r_S1S2_trtB) # probability to become Sicker when Sick

```

```
# under treatment B conditional on surviving
```

Create a state-transition diagram of the cohort model

```
m_P_diag <- matrix(0,
                    nrow = n_states, ncol = n_states,
                    dimnames = list(v_names_states, v_names_states))
m_P_diag["H" , "S1"] = ""
m_P_diag["H" , "D" ] = ""
m_P_diag["H" , "H" ] = ""
m_P_diag["S1", "H" ] = ""
m_P_diag["S1", "S2"] = ""
m_P_diag["S1", "D" ] = ""
m_P_diag["S1", "S1"] = ""
m_P_diag["S2", "D" ] = ""
m_P_diag["S2", "S2"] = ""
m_P_diag["D" , "D" ] = ""
layout.fig <- c(3, 1)

plotmat(t(m_P_diag), t(layout.fig), self.cex = 0.5, curve = 0, arr.pos = 0.7,
        latex = T, arr.type = "curved", relsize = 0.9, box.prop = 0.8,
        cex = 0.8, box.cex = 0.9, lwd = 1)
```

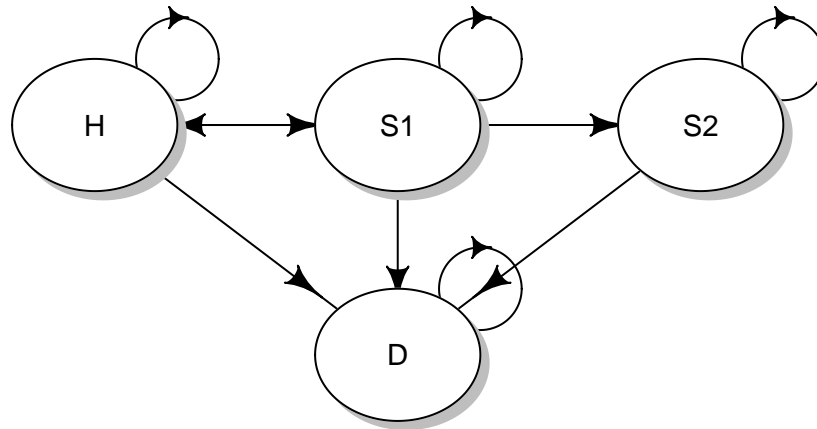


Figure 1: State-transition diagram of the time-independent Sick-Sicker cohort state-transition model.

04 Define and initialize matrices and vectors

04.1 Cohort trace

```
## Initialize cohort trace for SoC
m_M <- matrix(NA,
              nrow = (n_cycles + 1), ncol = n_states,
              dimnames = list(0:n_cycles, v_names_states))
# Store the initial state vector in the first row of the cohort trace
m_M[1, ] <- v_m_init
## Initialize cohort trace for strategies A, B, and AB
# Structure and initial states are the same as for SoC
```

```

m_M_strA <- m_M # Strategy A
m_M_strB <- m_M # Strategy B
m_M_strAB <- m_M # Strategy AB

```

04.2 Transition probability matrices

```

## Initialize transition probability matrix for strategy SoC
# all transitions to a non-death state are assumed to be conditional on survival
m_P <- matrix(0,
              nrow = n_states, ncol = n_states,
              dimnames = list(v_names_states,
                              v_names_states)) # define row and column names

```

```
m_P
```

```

##      H S1 S2 D
## H   0  0  0  0
## S1  0  0  0  0
## S2  0  0  0  0
## D   0  0  0  0

```

Fill in the transition probability matrix:

```

# From H
m_P["H", "H"] <- (1 - p_HD) * (1 - p_HS1)
m_P["H", "S1"] <- (1 - p_HD) * p_HS1
m_P["H", "D"] <- p_HD
# From S1
m_P["S1", "H"] <- (1 - p_S1D) * p_S1H
m_P["S1", "S1"] <- (1 - p_S1D) * (1 - (p_S1H + p_S1S2))
m_P["S1", "S2"] <- (1 - p_S1D) * p_S1S2
m_P["S1", "D"] <- p_S1D
# From S2
m_P["S2", "S2"] <- 1 - p_S2D
m_P["S2", "D"] <- p_S2D
# From D
m_P["D", "D"] <- 1

## Initialize transition probability matrix for strategy A as a copy of SoC's
m_P_strA <- m_P

## Initialize transition probability matrix for strategy B
m_P_strB <- m_P
# Update only transition probabilities from S1 involving p_S1S2
m_P_strB["S1", "S1"] <- (1 - p_S1D) * (1 - (p_S1H + p_S1S2_trtB))
m_P_strB["S1", "S2"] <- (1 - p_S1D) * p_S1S2_trtB

## Initialize transition probability matrix for strategy AB as a copy of B's
m_P_strAB <- m_P_strB

### Check if transition probability matrices are valid
## Check that transition probabilities are [0, 1]
check_transition_probability(m_P, verbose = TRUE)
check_transition_probability(m_P_strA, verbose = TRUE)
check_transition_probability(m_P_strB, verbose = TRUE)

```

```

check_transition_probability(m_P_strAB, verbose = TRUE)
## Check that all rows sum to 1
check_sum_of_transition_array(m_P,          n_states = n_states, verbose = TRUE)
check_sum_of_transition_array(m_P_strA,     n_states = n_states, verbose = TRUE)
check_sum_of_transition_array(m_P_strB,     n_states = n_states, verbose = TRUE)
check_sum_of_transition_array(m_P_strAB,    n_states = n_states, verbose = TRUE)

```

05 Run Markov model

```

# Iterative solution of time-independent cSTM
for(t in 1:n_cycles){
  # For SoC
  m_M[t + 1, ] <- m_M[t, ] %*% m_P
  # For strategy A
  m_M_strA[t + 1, ] <- m_M_strA[t, ] %*% m_P_strA
  # For strategy B
  m_M_strB[t + 1, ] <- m_M_strB[t, ] %*% m_P_strB
  # For strategy AB
  m_M_strAB[t + 1, ] <- m_M_strAB[t, ] %*% m_P_strAB
}

## Store the cohort traces in a list
l_m_M <- list(m_M,
              m_M_strA,
              m_M_strB,
              m_M_strAB)
names(l_m_M) <- v_names_str

```

06 Compute and Plot Epidemiological Outcomes

06.1 Cohort trace

```

# create a plot of the data
plot_trace(l_m_M$`Standard of care`)

```

06.2 Overall Survival (OS)

```

# calculate the overall survival (OS) probability for no treatment
v_os_SoC <- 1 - l_m_M$`Standard of care`[, "D"]
# alternative way of calculating the OS probability
v_os_SoC <- rowSums(l_m_M$`Standard of care`[, 1:3])
# create a simple plot showing the OS
plot(0:n_cycles, v_os_SoC, type = 'l',
     ylim = c(0, 1),
     ylab = "Survival probability",
     xlab = "Cycle",
     main = "Overall Survival")
# add grid
grid(nx = n_cycles, ny = 10, col = "lightgray", lty = "dotted", lwd = par("lwd"),
     equilogs = TRUE)

```

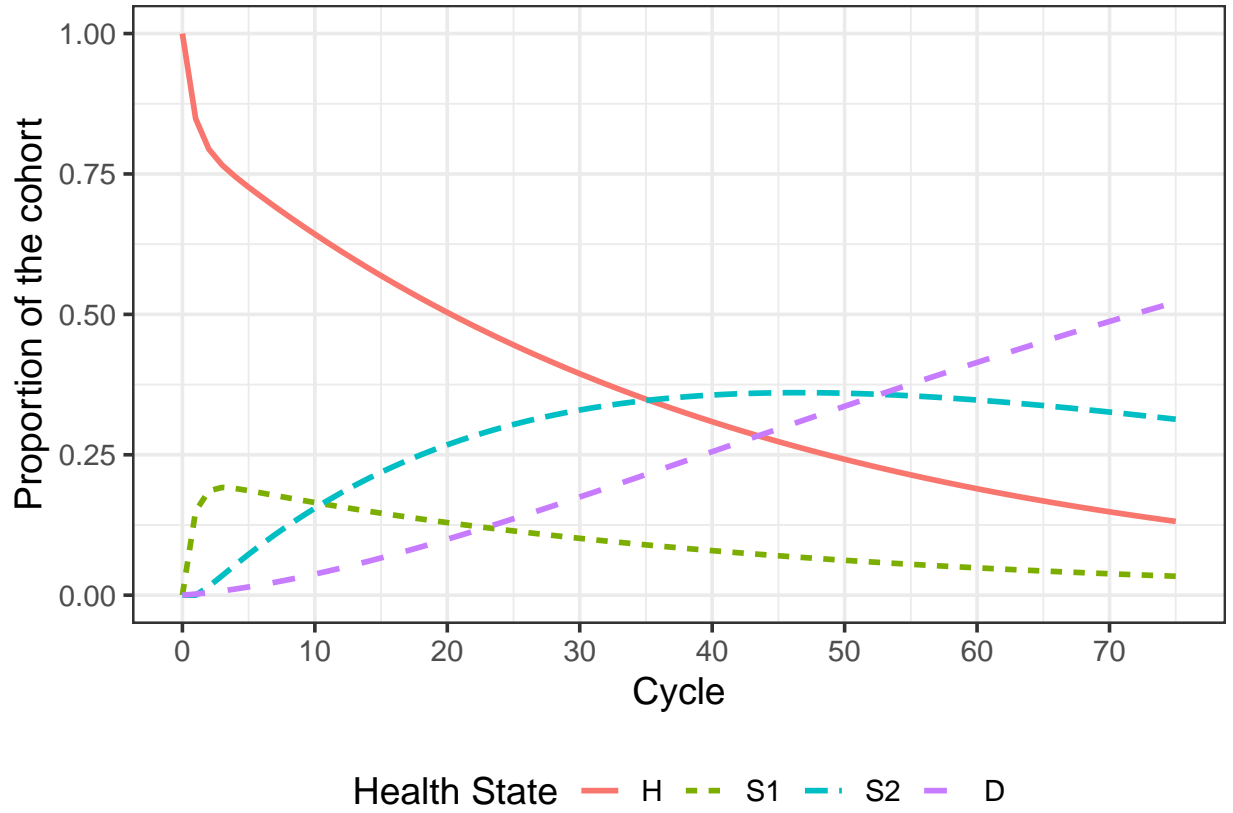


Figure 2: Cohort trace of the time-independent cSTM under standard of care

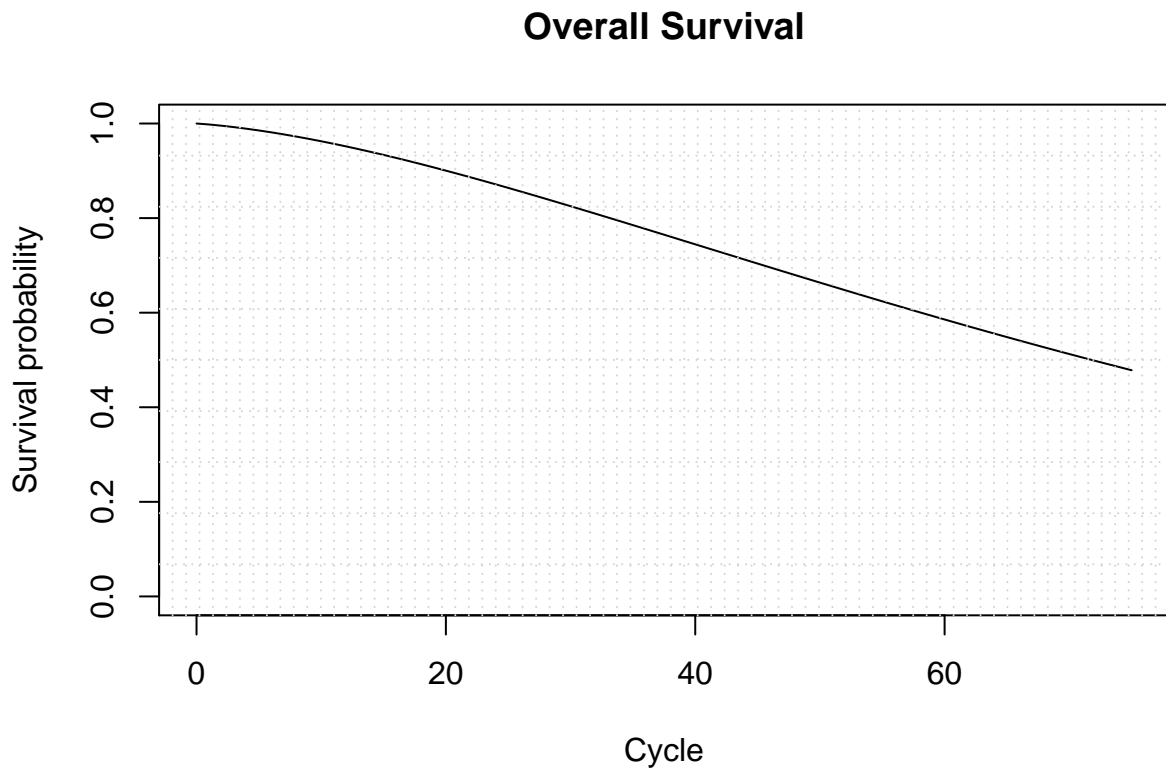


Figure 3: Overall survival of the time-independent cSTM under standard of care.

06.2.1 Life Expectancy (LE)

```
v_le <- sum(v_os_SoC) # summing probability of OS over time (i.e. life expectancy)
```

06.3 Disease prevalence

```
v_prev <- rowSums(l_m_M$`Standard of care`[, c("S1", "S2")]) / v_os_SoC
plot(v_prev,
     ylim = c(0, 1),
     ylab = "Prevalence",
     xlab = "Cycle",
     main = "Disease prevalence")
```

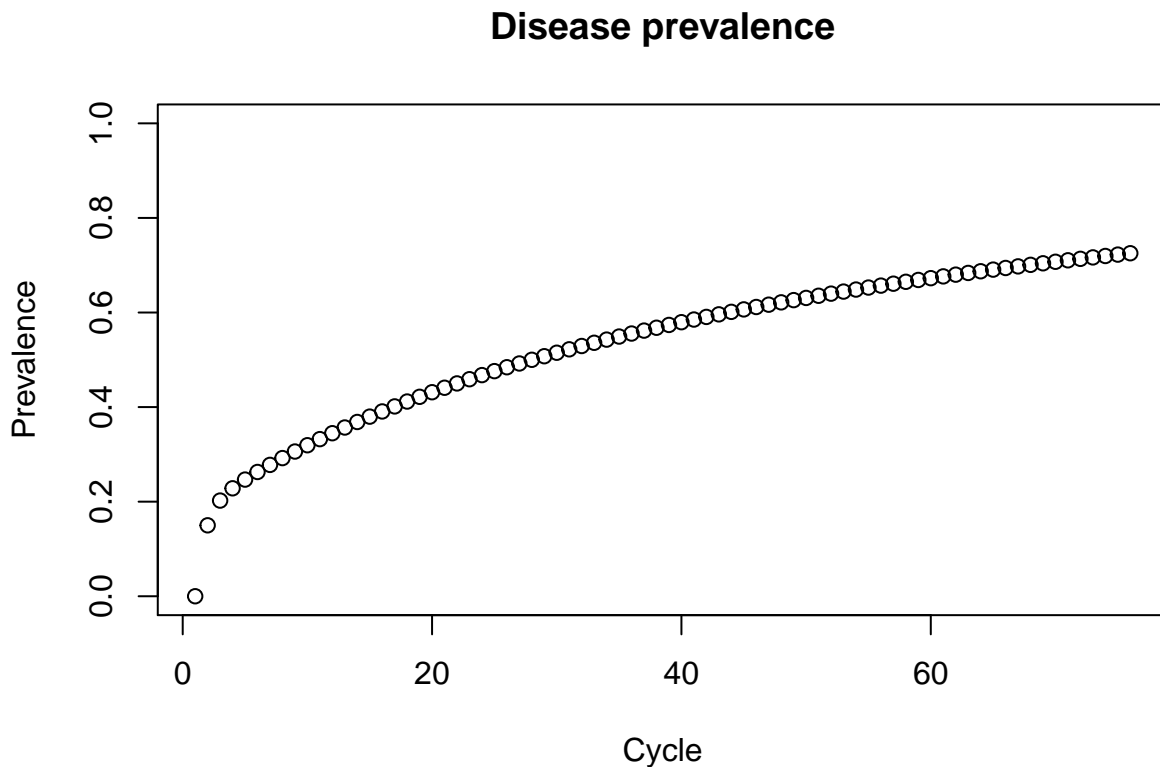


Figure 4: Disease prevalence of the time-independent cSTM under standard of care.

06.4 Proportion of sick in S1 state

```
v_prop_S1 <- l_m_M$`Standard of care`[, "S1"] / v_prev
plot(0:n_cycles, v_prop_S1,
     xlab = "Cycle",
     ylab = "Proportion",
     main = "Proportion of sick in S1 state",
     col = "black", type = "l")
```

07 Compute Cost-Effectiveness Outcomes

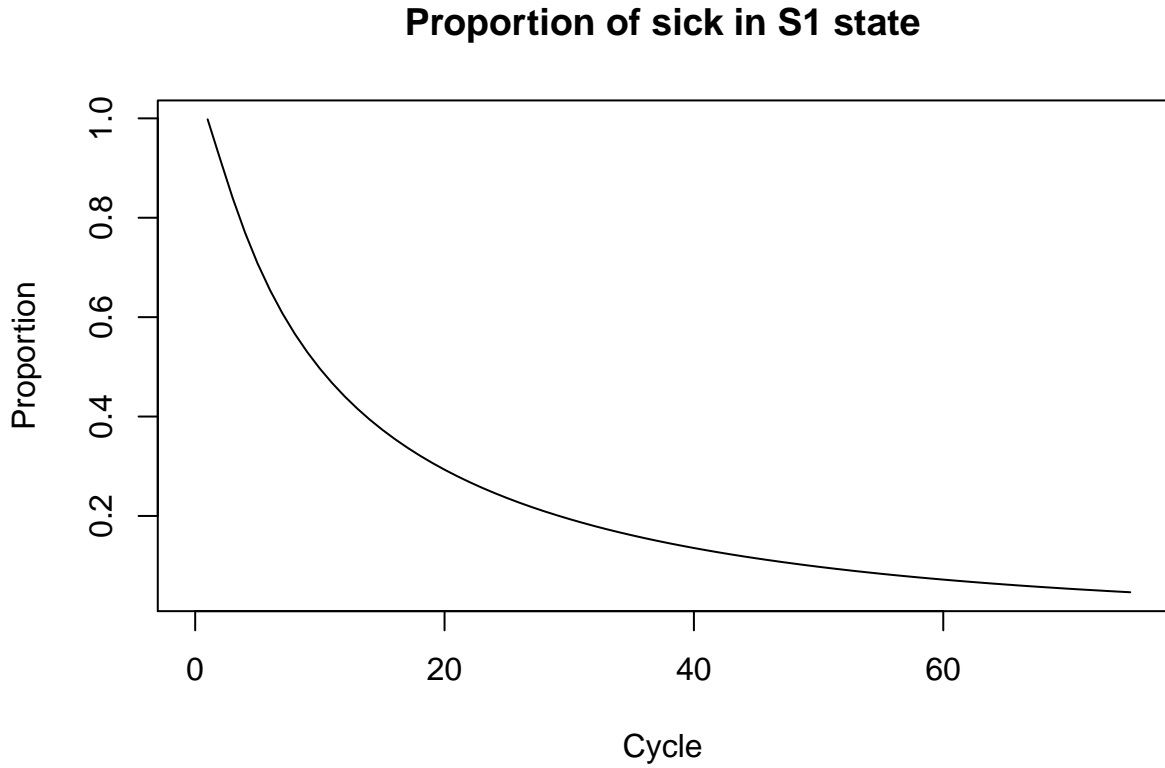


Figure 5: Proportion of sick in S1 state of the time-independent cSTM under standard of care.

07.1 State rewards for each strategy

```
## Vector of state utilities under strategy SoC
v_u_SoC    <- c(H = u_H,
               S1 = u_S1,
               S2 = u_S2,
               D = u_D)

## Vector of state costs under strategy SoC
v_c_SoC    <- c(H = c_H,
               S1 = c_S1,
               S2 = c_S2,
               D = c_D)

## Vector of state utilities under strategy A
v_u_strA    <- c(H = u_H,
               S1 = u_trtA,
               S2 = u_S2,
               D = u_D)

## Vector of state costs under strategy A
v_c_strA    <- c(H = c_H,
               S1 = c_S1 + c_trtA,
               S2 = c_S2 + c_trtA,
               D = c_D)

## Vector of state utilities under strategy B
v_u_strB    <- c(H = u_H,
               S1 = u_S1,
               S2 = u_S2,
```

```

        D = u_D)
## Vector of state costs under strategy B
v_c_strB <- c(H = c_H,
             S1 = c_S1 + c_trtB,
             S2 = c_S2 + c_trtB,
             D = c_D)
## Vector of state utilities under strategy AB
v_u_strAB <- c(H = u_H,
              S1 = u_trtA,
              S2 = u_S2,
              D = u_D)
## Vector of state costs under strategy AB
v_c_strAB <- c(H = c_H,
              S1 = c_S1 + (c_trtA + c_trtB),
              S2 = c_S2 + (c_trtA + c_trtB),
              D = c_D)

## Store the vectors of state utilities for each strategy in a list
l_u <- list(SQ = v_u_SoC,
           A = v_u_strA,
           B = v_u_strB,
           AB = v_u_strAB)
## Store the vectors of state cost for each strategy in a list
l_c <- list(SQ = v_c_SoC,
           A = v_c_strA,
           B = v_c_strB,
           AB = v_c_strAB)

# assign strategy names to matching items in the lists
names(l_u) <- names(l_c) <- v_names_str

```

07.2 Mean Costs and QALYs for each strategy

```

# create empty vectors to store total utilities and costs
v_tot_qaly <- v_tot_cost <- vector(mode = "numeric", length = n_str)
names(v_tot_qaly) <- names(v_tot_cost) <- v_names_str

#### Loop through each strategy and calculate total utilities and costs ####
for (i in 1:n_str) {
  v_u_str <- l_u[[i]] # select the vector of state utilities for the i-th strategy
  v_c_str <- l_c[[i]] # select the vector of state costs for the i-th strategy

  #### Expected QALYs and costs per cycle ####
  ## Vector of QALYs and Costs
  ## Apply state rewards ###
  v_qaly_str <- l_m_M[[i]] %*% v_u_str # sum the utilities of all states for each cycle
  v_cost_str <- l_m_M[[i]] %*% v_c_str # sum the costs of all states for each cycle

  #### Discounted total expected QALYs and Costs per strategy and apply half-cycle correction ####
  ## QALYs
  v_tot_qaly[i] <- t(v_qaly_str) %*% (v_dwe * v_wcc)
  ## Costs
  v_tot_cost[i] <- t(v_cost_str) %*% (v_dwc * v_wcc)
}

```

```
}
```

07.3 Compute ICERs of the Markov model

```
# Calculate incremental cost-effectiveness ratios (ICERs)
df_cea <- calculate_icers(cost      = v_tot_cost,
                        effect     = v_tot_qaly,
                        strategies = v_names_str)

df_cea
```

	Strategy	Cost	Effect	Inc_Cost	Inc_Effect
## Standard of care	Standard of care	148657.5	20.99026	NA	NA
## Strategy B	Strategy B	248570.6	22.48240	99913.15	1.4921371
## Strategy AB	Strategy AB	361341.5	23.35420	112770.85	0.8718013
## Strategy A	Strategy A	275936.5	21.71749	NA	NA

```
##
## ICER Status
## Standard of care      NA      ND
## Strategy B           66959.76    ND
## Strategy AB          129353.84    ND
## Strategy A           NA         D
```

07.4 CEA results

```
# Create CEA table in proper format
table_cea <- format_table_cea(df_cea)
table_cea
```

	Strategy	Costs (\$)	QALYs	Incremental Costs (\$)
## Standard of care	Standard of care	148,657	20.99	<NA>
## Strategy B	Strategy B	248,571	22.48	99,913
## Strategy AB	Strategy AB	361,341	23.35	112,771
## Strategy A	Strategy A	275,937	21.72	<NA>

```
##
## Incremental QALYs ICER ($/QALY) Status
## Standard of care      NA      <NA>      ND
## Strategy B           1.49      66,960      ND
## Strategy AB           0.87     129,354      ND
## Strategy A           NA      <NA>         D
```

07.5 Plot frontier of the Markov model

```
plot(df_cea, label = "all") +
  expand_limits(x = max(table_cea$QALYs) + 0.5)
```

08 Deterministic Sensitivity Analysis

08.1 List of input parameters

Create list `l_params_all` with all input probabilities, cost and utilities.

```
l_params_all <- list(
  # Transition probabilities (per cycle), hazard ratios
  r_HD   = 0.002, # constant rate of dying when Healthy (all-cause mortality)
  p_HS1  = 0.15,  # probability to become Sick when Healthy conditional on surviving
```

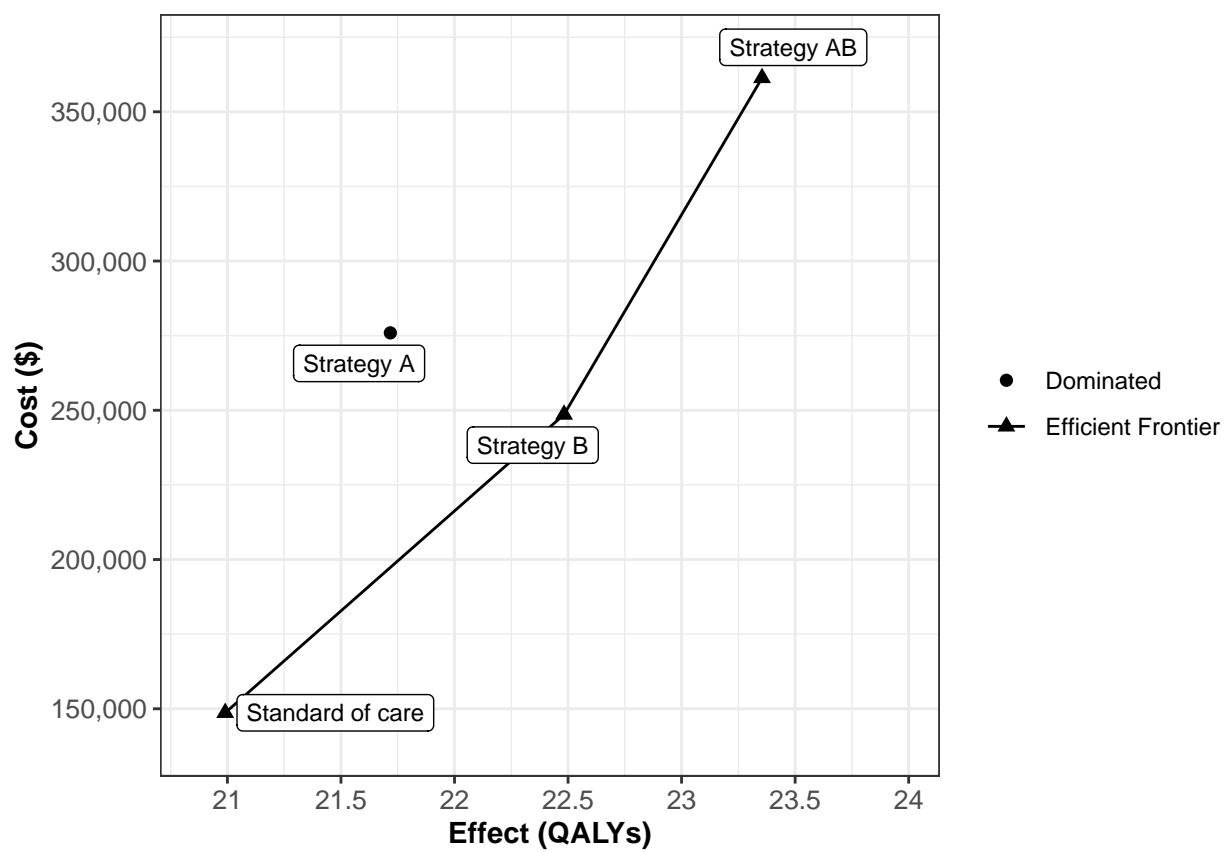


Figure 6: Cost-effectiveness efficient frontier for the time-independent Sick-Sicker model.

```

p_S1H = 0.5, # probability to become Healthy when Sick conditional on surviving
p_S1S2 = 0.105, # probability to become Sicker when Sick conditional on surviving
hr_S1 = 3, # hazard ratio of death in Sick vs Healthy
hr_S2 = 10, # hazard ratio of death in Sicker vs Healthy
# Effectiveness of treatment B
hr_S1S2_trtB = 0.6, # hazard ratio of becoming Sicker when Sick under B under treatment B
## State rewards
# Costs
c_H = 2000, # cost of remaining one cycle in Healthy
c_S1 = 4000, # cost of remaining one cycle in Sick
c_S2 = 15000, # cost of remaining one cycle in Sicker
c_D = 0, # cost of being dead (per cycle)
c_trtA = 12000, # cost of treatment A
c_trtB = 13000, # cost of treatment B
# Utilities
u_H = 1, # utility when Healthy
u_S1 = 0.75, # utility when Sick
u_S2 = 0.5, # utility when Sicker
u_D = 0, # utility when Dead
u_trtA = 0.95 # utility when being treated with A
)

# store the parameter names into a vector
v_names_params <- names(l_params_all)

```

08.2 Load Sick-Sicker Markov model function

```

source("Functions_markov_sick-sicker_intro_tutorial.R")
# Test function to compute CE outcomes
calculate_ce_out(l_params_all)

##               Strategy      Cost   Effect   NMB
## Standard of care Standard of care 148657.5 20.99026 1950369
## Strategy A      Strategy A 275936.5 21.71749 1895812
## Strategy B      Strategy B 248570.6 22.48240 1999669
## Strategy AB     Strategy AB 361341.5 23.35420 1974079

```

08.3 One-way sensitivity analysis (OWSA)

```

options(scipen = 999) # disabling scientific notation in R
# data.frame containing all parameters, their base-case values, and the min and
# max values of the parameters of interest
df_params_owsa <- data.frame(pars = c("hr_S1S2_trtB", "c_trtA", "u_S1", "u_trtA"),
                             min = c(0.10, 6000, 0.65, 0.80), # min parameter values
                             max = c(1.00, 18000, 0.85, 0.98) # max parameter values
                             )

owsa_nmb <- run_owsa_det(params_range = df_params_owsa, # data.frame with parameters for OWSA
                        params_basecase = l_params_all, # list with all parameters
                        nsamp = 100, # number of parameter values
                        FUN = calculate_ce_out, # function to compute outputs
                        outcomes = c("NMB"), # output to do the OWSA on
                        strategies = v_names_str, # names of the strategies

```

```

n_wtp = 120000) # extra argument to pass to FUN

## |
plot(owsa_nmb, txtsize = 10, n_x_ticks = 4,
     facet_scales = "free") +
  theme(legend.position = "bottom")

```

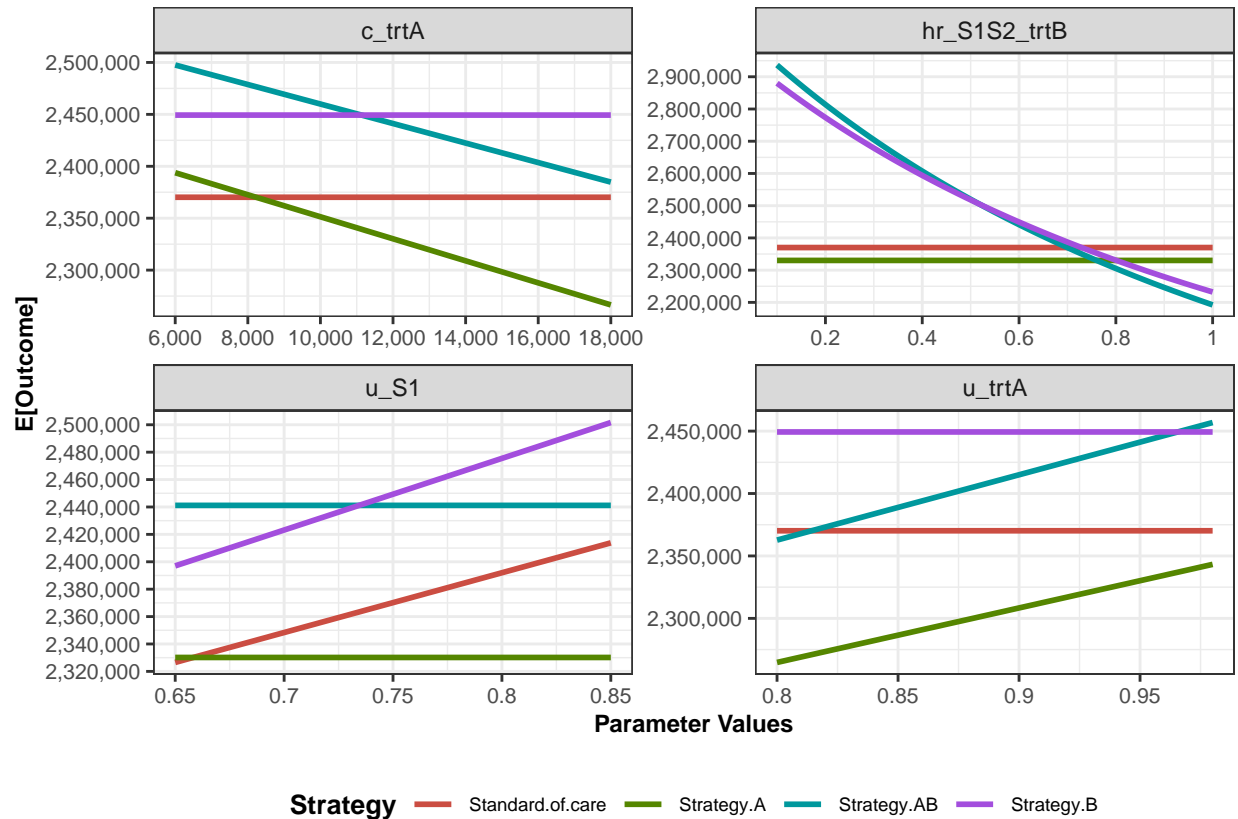


Figure 7: One-way sensitivity analysis for the time-independent Sick-Sicker model.

08.3.2 Optimal strategy with OWSA

```
owsa_opt_strat(owsa = owsa_nmb, txtsize = 10)
```

08.3.3 Tornado plot

```
owsa_tornado(owsa = owsa_nmb, txtsize = 10)
```

08.4 Two-way sensitivity analysis (TWSA)

```

# dataframe containing all parameters, their basecase values, and the min and
# max values of the parameters of interest
df_params_twsa <- data.frame(pars = c("hr_S1S2_trtB", "u_trtA"),
                             min = c(0.10, 0.80), # min parameter values
                             max = c(1.00, 0.98)  # max parameter values
                             )

```

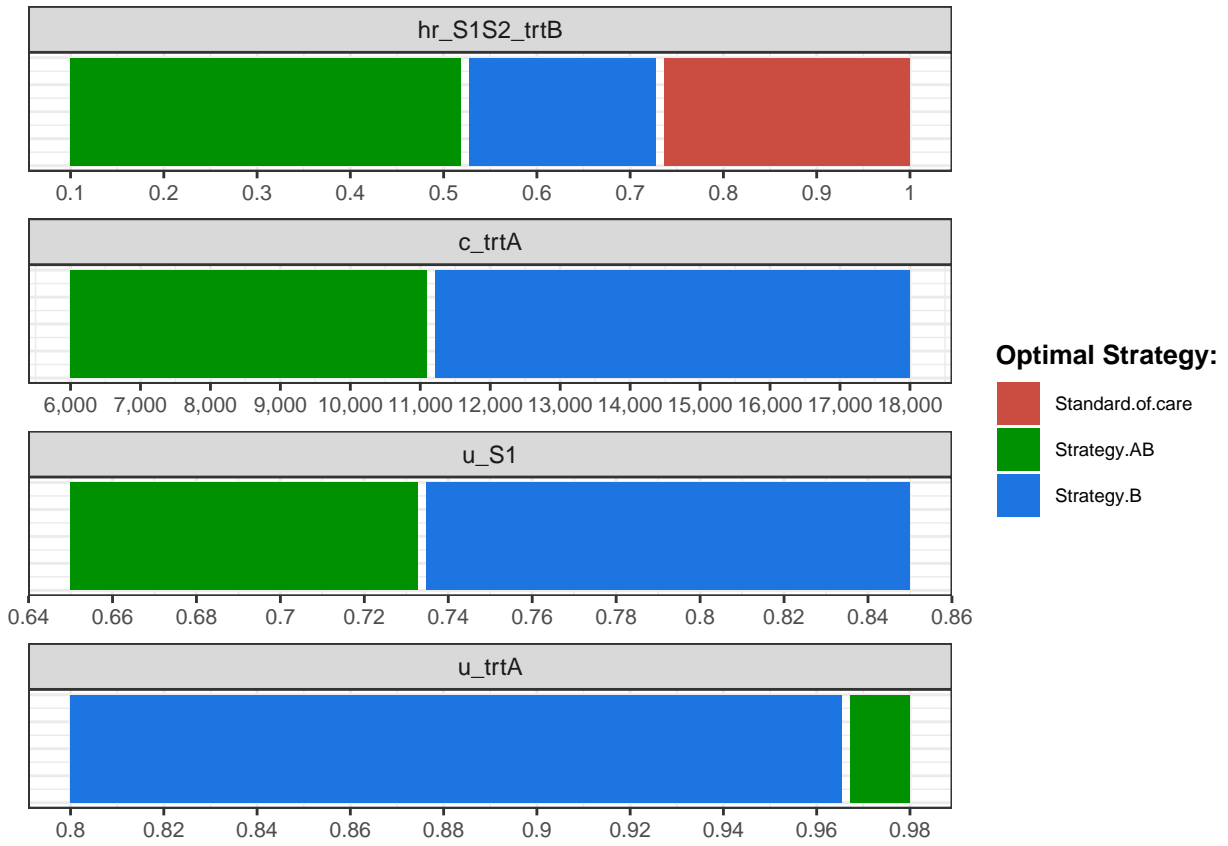


Figure 8: Optimal strategy with one-way sensitivity analysis for the time-independent Sick-Sicker model.

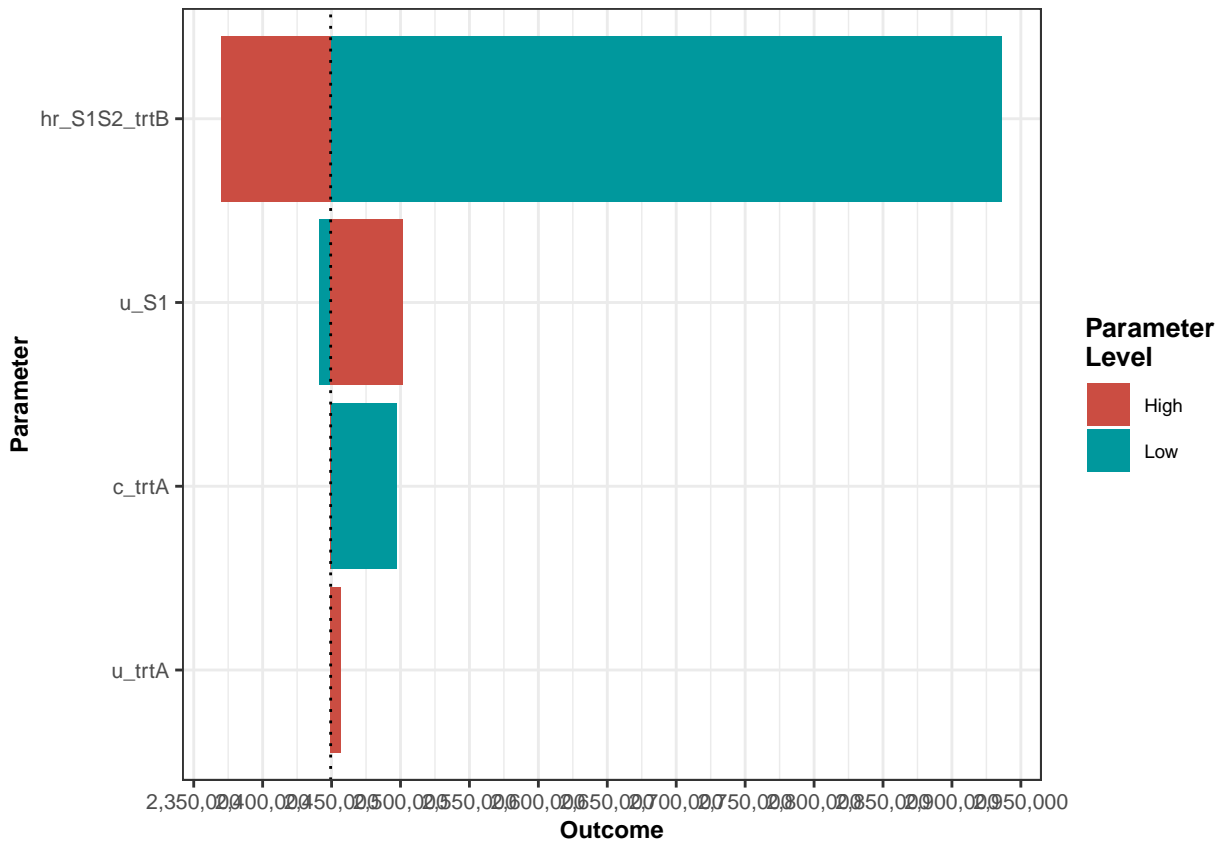


Figure 9: Tornado plot for the time-independent Sick-Sicker model.

```

twsa_nmb <- run_twsa_det(params_range = df_params_twsa,      # data.frame with parameters for TWSA
                        params_basecase = l_params_all,    # list with all parameters
                        nsamp           = 40,              # number of parameter values
                        FUN              = calculate_ce_out, # function to compute outputs
                        outcomes         = c("NMB"),        # output to do the TWSA on
                        strategies      = v_names_str,     # names of the strategies
                        n_wtp           = 120000)          # extra argument to pass to FUN

## |

```

```
plot(twsa_nmb)
```

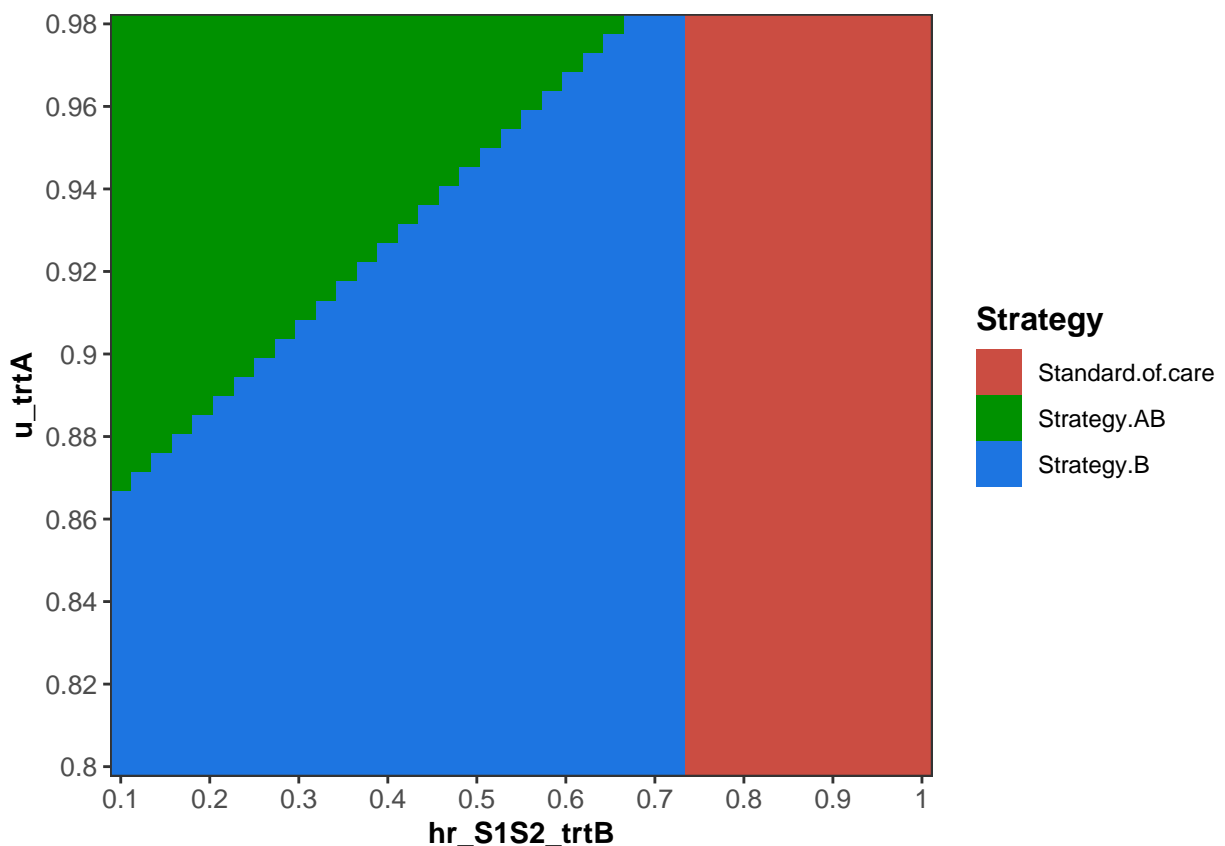


Figure 10: Two-way sensitivity analysis for the time-independent Sick-Sicker model.

09 Probabilistic Sensitivity Analysis (PSA)

```

# Function to generate PSA input dataset
generate_psa_params <- function(n_sim = 1000, seed = 071818){
  set.seed(seed) # set a seed to be able to reproduce the same results
  df_psa <- data.frame(
    # Transition probabilities (per cycle), hazard ratios
    r_HD      = rlnorm(n_sim, meanlog = log(0.002), sdlog = 0.01), # constant rate of dying when Healthy

```

```

p_HS1 = rbeta(n_sim, shape1 = 30, shape2 = 170), # probability to become Sick when Healthy
p_S1H = rbeta(n_sim, shape1 = 60, shape2 = 60), # probability to become Healthy when Sick
p_S1S2 = rbeta(n_sim, shape1 = 84, shape2 = 716), # probability to become Sicker when Sick
hr_S1 = rlnorm(n_sim, meanlog = log(3), sdlog = 0.01), # hazard ratio of death in Sick vs Healthy
hr_S2 = rlnorm(n_sim, meanlog = log(10), sdlog = 0.02), # hazard ratio of death in Sicker vs Healthy

# Effectiveness of treatment B
hr_S1S2_trtB = rlnorm(n_sim, meanlog = log(0.6), sdlog = 0.02), # hazard ratio of becoming Sicker w

# State rewards
# Costs
c_H = rgamma(n_sim, shape = 100, scale = 20), # cost of remaining one cycle in Healthy
c_S1 = rgamma(n_sim, shape = 177.8, scale = 22.5), # cost of remaining one cycle in Sick
c_S2 = rgamma(n_sim, shape = 225, scale = 66.7), # cost of remaining one cycle in Sicker
c_D = 0, # cost of being dead (per cycle)
c_trtA = rgamma(n_sim, shape = 73.5, scale = 163.3), # cost of treatment A
c_trtB = rgamma(n_sim, shape = 86.2, scale = 150.8), # cost of treatment B

# Utilities
u_H = rbeta(n_sim, shape1 = 200, shape2 = 3), # utility when Healthy
u_S1 = rbeta(n_sim, shape1 = 130, shape2 = 45), # utility when Sick
u_S2 = rbeta(n_sim, shape1 = 230, shape2 = 230), # utility when Sicker
u_D = 0, # utility when Dead
u_trtA = rbeta(n_sim, shape1 = 300, shape2 = 15) # utility when being treated with A
)
return(df_psa)
}

```

```

# Try it
generate_psa_params(10)

```

```

##          r_HD      p_HS1      p_S1H      p_S1S2      hr_S1      hr_S2 hr_S1S2_trtB
## 1  0.002028152  0.09824029  0.5127637  0.09753778  3.055410  10.148768    0.5937824
## 2  0.002000921  0.14028588  0.5472914  0.10372958  3.005252   9.928044    0.5950295
## 3  0.001982103  0.11912372  0.5563895  0.10145687  2.982392  10.021998    0.6131617
## 4  0.002004616  0.16814566  0.4513392  0.13328700  2.981918  10.167486    0.6066259
## 5  0.001973180  0.12386074  0.5451874  0.11709463  3.024975  10.006505    0.5839462
## 6  0.001994982  0.20236810  0.4959245  0.09820372  2.971345  10.012715    0.5861758
## 7  0.002029022  0.12808209  0.4467345  0.10922783  2.989181  10.001860    0.5838734
## 8  0.001995376  0.17956555  0.5889240  0.11869605  2.984080   9.919332    0.6081428
## 9  0.001986669  0.14451930  0.4441287  0.09900264  2.988974  10.142331    0.5784071
## 10 0.002004075  0.19389383  0.5076377  0.07977953  2.968692   9.749869    0.6141500
##          c_H      c_S1      c_S2 c_D      c_trtA      c_trtB      u_H      u_S1
## 1  1738.307  3448.146  15401.50   0  11760.17  12545.42  0.9936986  0.7921002
## 2  1934.791  4656.666  15198.97   0  10216.98  16021.51  0.9890348  0.8069535
## 3  2113.222  3879.536  14070.22   0  13948.46  11876.24  0.9943986  0.7796555
## 4  2123.062  3911.558  16006.75   0  12209.42  13905.64  0.9903866  0.7634364
## 5  2387.026  4497.693  15402.85   0  13348.28  13951.11  0.9922080  0.6960808
## 6  1600.231  4198.111  16322.33   0  10982.80  14379.26  0.9923096  0.7201886
## 7  1916.467  4087.810  15848.52   0  11359.07  13869.89  0.9951053  0.7368862
## 8  1977.534  4088.065  16378.74   0  15146.19  14538.77  0.9838664  0.7871166
## 9  1785.850  4110.476  15114.98   0  12174.76  14097.89  0.9789327  0.7801267
## 10 1852.581  4181.534  15081.51   0  10827.73  11962.58  0.9888362  0.6948902
##          u_S2 u_D      u_trtA

```

```
## 1 0.5120295 0 0.9618487
## 2 0.5596476 0 0.9588147
## 3 0.5227040 0 0.9500272
## 4 0.4972854 0 0.9503212
## 5 0.4890811 0 0.9527867
## 6 0.5479399 0 0.9474899
## 7 0.4880988 0 0.9272372
## 8 0.5286497 0 0.9504067
## 9 0.5192718 0 0.9562053
## 10 0.4905318 0 0.9443718
```

```
# Number of simulations
n_sim <- 1000

# Generate PSA input dataset
df_psa_input <- generate_psa_params(n_sim = n_sim)
# First six observations
head(df_psa_input)
```

```
##          r_HD      p_HS1      p_S1H      p_S1S2      hr_S1      hr_S2 hr_S1S2_trtB
## 1 0.002028152 0.1602643 0.4968849 0.09399301 3.021339 10.423829 0.6174156
## 2 0.002000921 0.1185122 0.5553071 0.09355730 2.954005 10.122297 0.5953304
## 3 0.001982103 0.1239255 0.5275639 0.11118352 3.040920 10.466481 0.6037374
## 4 0.002004616 0.1823176 0.5340465 0.10680270 3.008736 9.836515 0.6013755
## 5 0.001973180 0.1250214 0.5540891 0.08518676 3.009890 10.315554 0.5977386
## 6 0.001994982 0.1437542 0.4900105 0.08858359 2.983996 10.204962 0.5847355
##          c_H      c_S1      c_S2 c_D      c_trtA      c_trtB      u_H      u_S1
## 1 2024.100 4111.210 15516.92 0 14536.17 12753.80 0.9884166 0.7043466
## 2 2269.089 3825.014 15036.51 0 11084.58 15073.93 0.9782958 0.8001897
## 3 2020.638 4091.684 14959.30 0 12668.41 14888.98 0.9752874 0.7430207
## 4 1945.772 3989.084 13147.33 0 13583.26 11832.31 0.9856703 0.7422257
## 5 2089.839 3663.629 15669.36 0 13687.70 10075.88 0.9811165 0.7274420
## 6 2107.511 3553.373 15294.76 0 12141.25 12343.99 0.9894955 0.7223727
##          u_S2 u_D      u_trtA
## 1 0.5079267 0 0.9745802
## 2 0.5128133 0 0.9452539
## 3 0.4741263 0 0.9521755
## 4 0.4981919 0 0.9490015
## 5 0.4955725 0 0.9571041
## 6 0.5264675 0 0.9504759
```

```
# Histogram of parameters
ggplot(melt(df_psa_input, variable.name = "Parameter"), aes(x = value)) +
  facet_wrap(~Parameter, scales = "free") +
  geom_histogram(aes(y = ..density..)) +
  scale_x_continuous(breaks = scales::pretty_breaks(n = 3)) +
  theme_bw(base_size = 16) +
  theme(axis.text = element_text(size=6))
```

09.1 Conduct probabilistic sensitivity analysis

```
# Initialize data.frames with PSA output
# data.frame of costs
df_c <- as.data.frame(matrix(0,
                             nrow = n_sim,
```

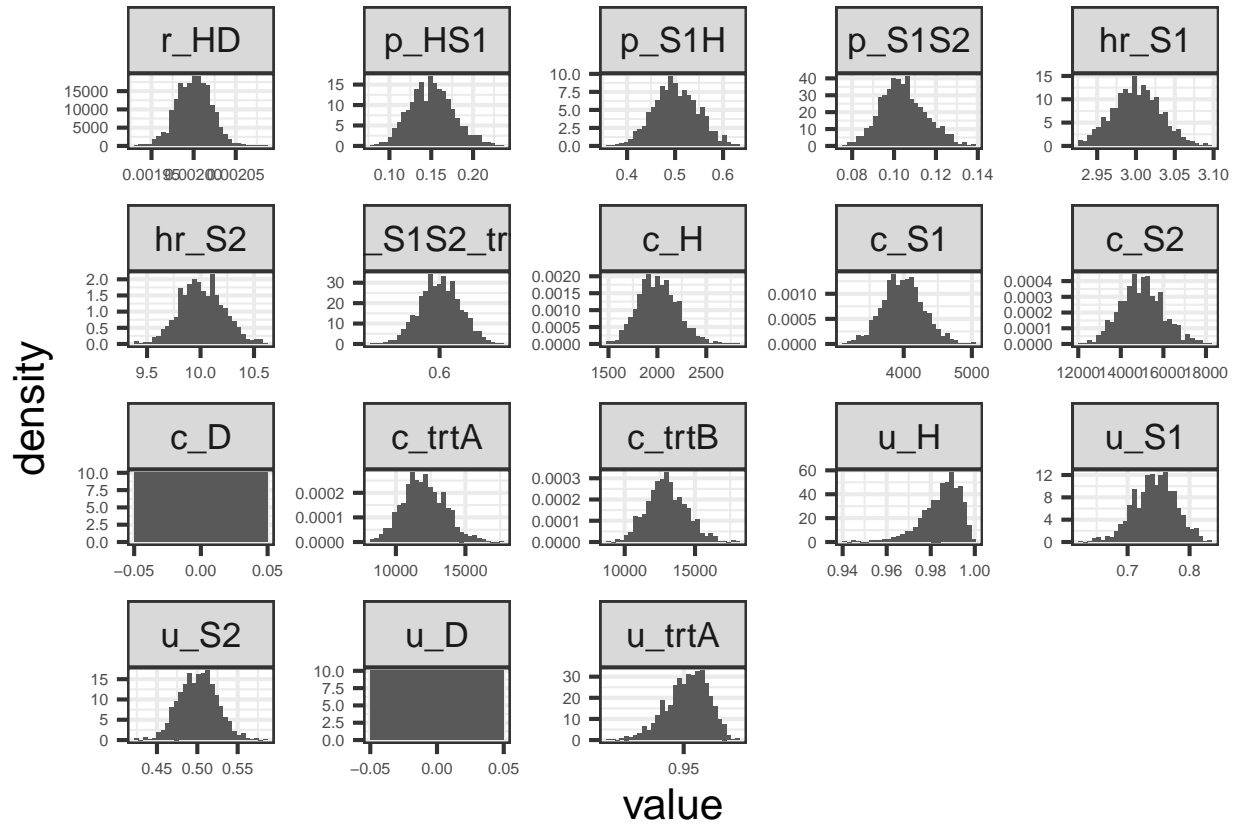


Figure 11: Histogram of parameters distributions for the time-independent Sick-Sicker model.

```

                                ncol = n_str))
colnames(df_c) <- v_names_str
# data.frame of effectiveness
df_e <- as.data.frame(matrix(0,
                                nrow = n_sim,
                                ncol = n_str))

colnames(df_e) <- v_names_str
# Run Markov model on each parameter set of PSA input dataset
n_time_init_psa_series <- Sys.time()
for(i in 1:n_sim){
  l_out_temp <- calculate_ce_out(df_psa_input[i, ])
  df_c[i, ] <- l_out_temp$Cost
  df_e[i, ] <- l_out_temp$Effect
  # Display simulation progress
  if(i/(n_sim/10) == round(i/(n_sim/10), 0)) { # display progress every 10%
    cat('\r', paste(i/n_sim * 100, "% done", sep = " "))
  }
}

```

```
## 10 % done 20 % done 30 % done 40 % done 50 % done 60 % done 70 % done 80 % done 90 % done 100 % done
```

```

n_time_end_psa_series <- Sys.time()
n_time_total_psa_series <- n_time_end_psa_series - n_time_init_psa_series
print(paste0("PSA with ", comma(n_sim), " simulations run in series in ",
             round(n_time_total_psa_series, 2), " ",
             units(n_time_total_psa_series)))

```

```
## [1] "PSA with 1,000 simulations run in series in 1.68 secs"
```

09.2 Create PSA object for dampack

```

l_psa <- make_psa_obj(cost      = df_c,
                     effectiveness = df_e,
                     parameters  = df_psa_input,
                     strategies  = v_names_str)

l_psa$strategies <- v_names_str
colnames(l_psa$effectiveness) <- v_names_str
colnames(l_psa$cost) <- v_names_str

```

09.2.1 Save PSA objects

```

save(df_psa_input, df_c, df_e, v_names_str, n_str, l_psa,
     file = "markov_sick-sicker_intro_tutorial_PSA_dataset.RData")

```

09.3 Create probabilistic analysis graphs

```
load(file = "markov_sick-sicker_intro_tutorial_PSA_dataset.RData")
```

Vector with willingness-to-pay (WTP) thresholds.

```
v_wtp <- seq(0, 250000, by = 5000)
```

09.3.1 Cost-Effectiveness scatter plot

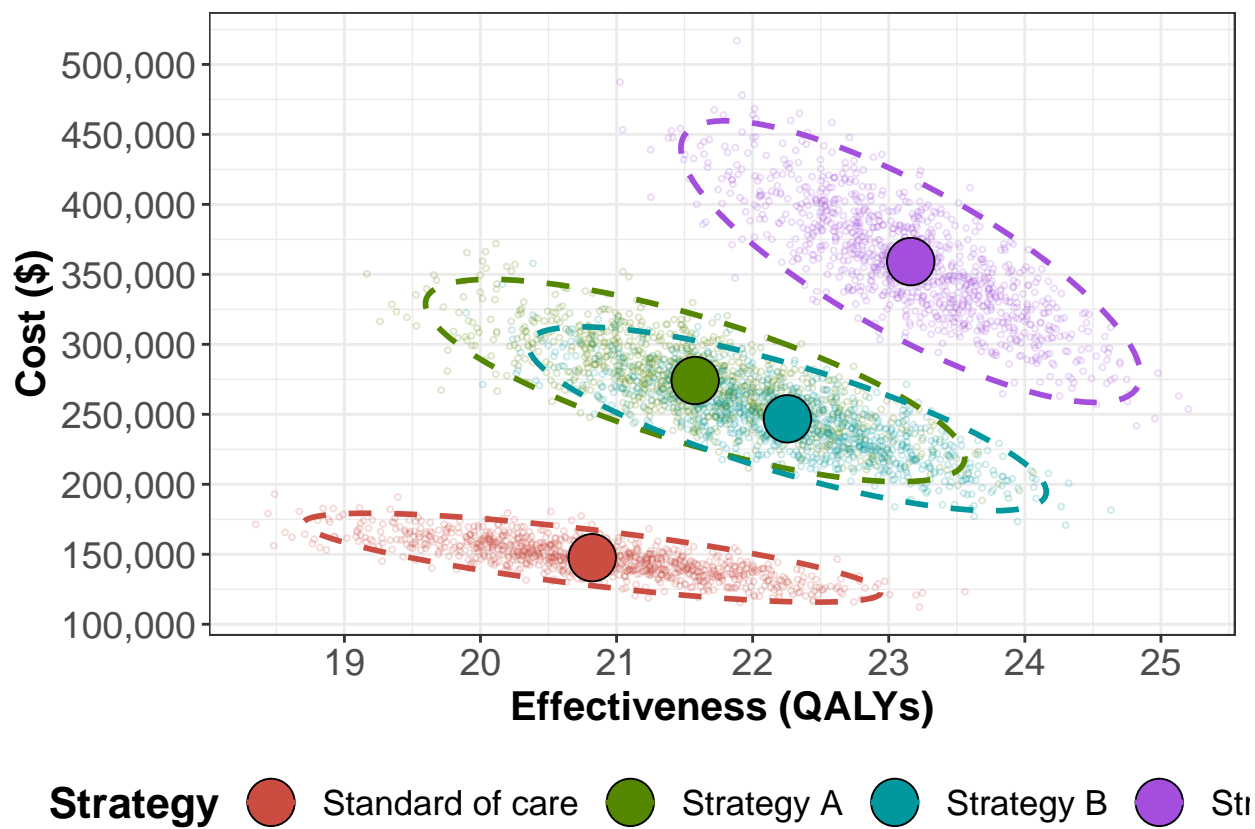


Figure 12: Cost-effectiveness scatter plot.

09.4 Conduct CEA with probabilistic output

```
# Compute expected costs and effects for each strategy from the PSA
df_out_ce_psa <- summary(l_psa)

# Calculate incremental cost-effectiveness ratios (ICERs)
df_cea_psa <- calculate_icers(cost      = df_out_ce_psa$meanCost,
                             effect    = df_out_ce_psa$meanEffect,
                             strategies = df_out_ce_psa$Strategy)

df_cea_psa

##           Strategy      Cost      Effect  Inc_Cost  Inc_Effect      ICER Status
## 1 Standard of care 147556.9 20.82141         NA         NA         NA      ND
## 2 Strategy B      246796.5 22.25558  99239.59  1.4341754  69196.27      ND
## 3 Strategy AB     359123.0 23.16154 112326.53  0.9059587 123986.38      ND
## 4 Strategy A      274149.6 21.57760         NA         NA         NA      D

# Save CEA table with ICERs
# As .RData
save(df_cea_psa,
     file = "markov_sick-sicker_intro_tutorial_probabilistic_CEA_results.RData")
# As .csv
write.csv(df_cea_psa,
          file = "markov_sick-sicker_intro_tutorial_probabilistic_CEA_results.csv")
```

09.4.1 Plot cost-effectiveness frontier

```
plot(df_cea_psa)
```

09.4.2 Cost-effectiveness acceptability curves (CEACs) and frontier (CEAF)

```
##   range_min range_max  cost_eff_strat
## 1         0    70000 Standard of care
## 2         0   125000 Strategy B
## 3    70000   250000 Strategy B
## 4   125000   250000 Strategy AB
## 5   125000        NA Strategy AB
```

09.4.3 Expected Loss Curves (ELCs)

The expected loss is the the quantification of the foregone benefits when choosing a suboptimal strategy given current evidence.

```
##      WTP      Strategy  Expected_Loss  On_Frontier
## 1      0 Standard of care    0.0000000      TRUE
## 2      0 Strategy A    126592.7446448     FALSE
## 3      0 Strategy B    99239.5859762     FALSE
## 4      0 Strategy AB   211566.1182551     FALSE
## 5   5000 Standard of care    0.0000000      TRUE
## 6   5000 Strategy A   122811.8022610     FALSE
## 7   5000 Strategy B    92068.7090661     FALSE
## 8   5000 Strategy AB   199865.4479683     FALSE
## 9  10000 Standard of care    0.0000000      TRUE
## 10 10000 Strategy A   119030.8598772     FALSE
## 11 10000 Strategy B    84897.8321559     FALSE
```

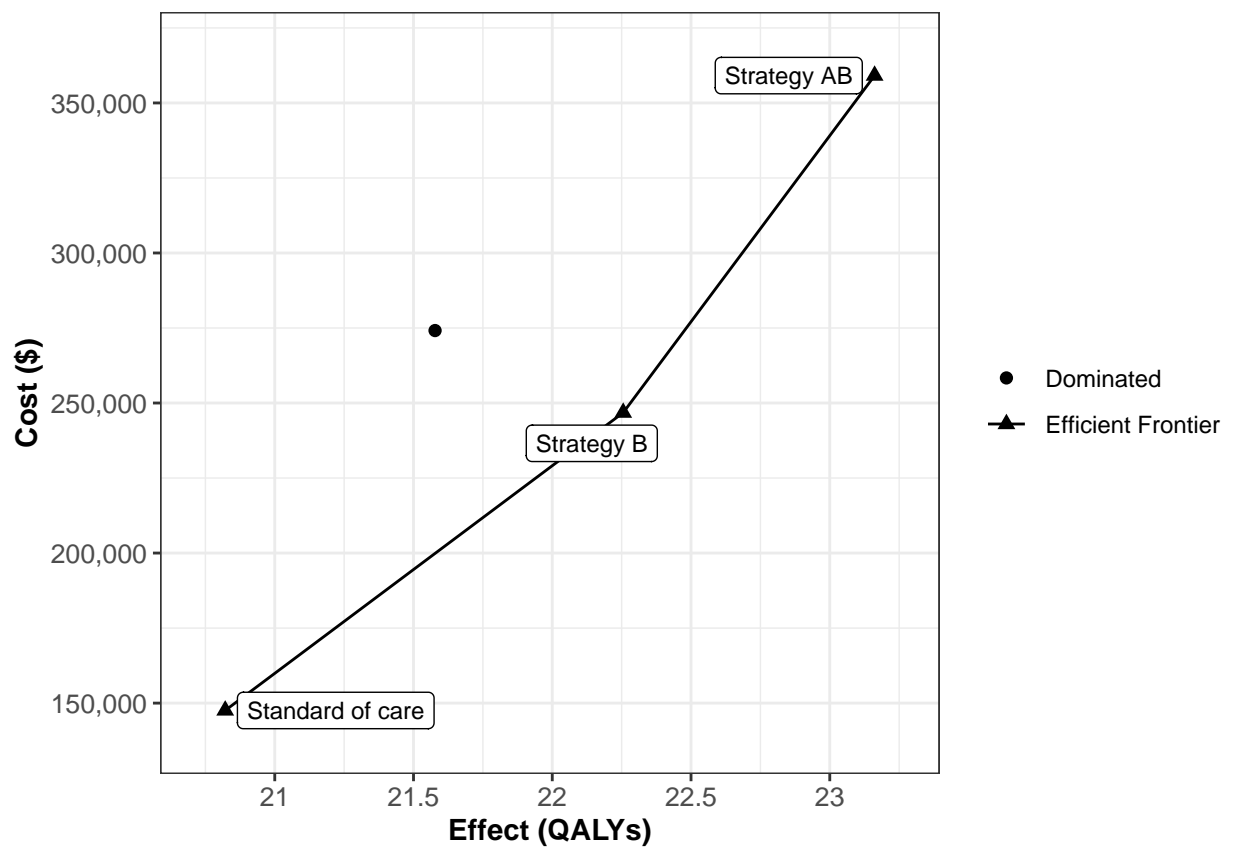



Figure 13: Cost-effectiveness efficient frontier from probabilistic outputs for the time-independent Sick-Sicker model.

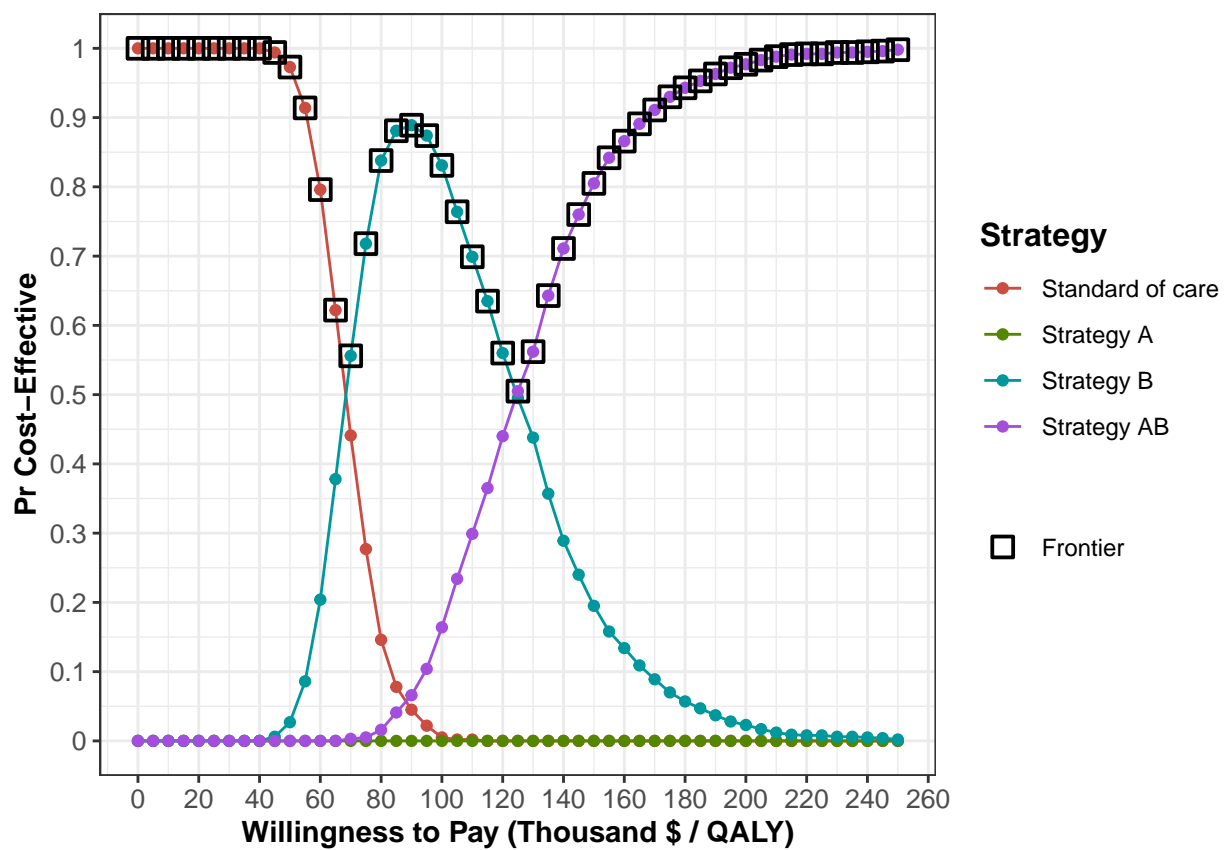


Figure 14: Cost-effectiveness acceptability curves (CEACs) and frontier (CEAF).

## 12	10000	Strategy AB	188164.7776815	FALSE
## 13	15000	Standard of care	0.0000000	TRUE
## 14	15000	Strategy A	115249.9174934	FALSE
## 15	15000	Strategy B	77726.9552457	FALSE
## 16	15000	Strategy AB	176464.1073947	FALSE
## 17	20000	Standard of care	0.0000000	TRUE
## 18	20000	Strategy A	111468.9751096	FALSE
## 19	20000	Strategy B	70556.0783355	FALSE
## 20	20000	Strategy AB	164763.4371079	FALSE
## 21	25000	Standard of care	0.0000000	TRUE
## 22	25000	Strategy A	107688.0327258	FALSE
## 23	25000	Strategy B	63385.2014254	FALSE
## 24	25000	Strategy AB	153062.7668211	FALSE
## 25	30000	Standard of care	0.0000000	TRUE
## 26	30000	Strategy A	103907.0903420	FALSE
## 27	30000	Strategy B	56214.3245152	FALSE
## 28	30000	Strategy AB	141362.0965343	FALSE
## 29	35000	Standard of care	0.0000000	TRUE
## 30	35000	Strategy A	100126.1479582	FALSE
## 31	35000	Strategy B	49043.4476050	FALSE
## 32	35000	Strategy AB	129661.4262475	FALSE
## 33	40000	Standard of care	0.0000000	TRUE
## 34	40000	Strategy A	96345.2055744	FALSE
## 35	40000	Strategy B	41872.5706949	FALSE
## 36	40000	Strategy AB	117960.7559607	FALSE
## 37	45000	Standard of care	20.4105025	TRUE
## 38	45000	Strategy A	92584.6736931	FALSE
## 39	45000	Strategy B	34722.1042872	FALSE
## 40	45000	Strategy AB	106280.4961764	FALSE
## 41	50000	Standard of care	127.3151594	TRUE
## 42	50000	Strategy A	88910.6359661	FALSE
## 43	50000	Strategy B	27658.1320339	FALSE
## 44	50000	Strategy AB	94686.7305464	FALSE
## 45	55000	Standard of care	484.6710630	TRUE
## 46	55000	Strategy A	85487.0494860	FALSE
## 47	55000	Strategy B	20844.6110274	FALSE
## 48	55000	Strategy AB	83343.4161633	FALSE
## 49	60000	Standard of care	1443.8027334	TRUE
## 50	60000	Strategy A	82665.2387725	FALSE
## 51	60000	Strategy B	14632.8657876	FALSE
## 52	60000	Strategy AB	72601.8775469	FALSE
## 53	65000	Standard of care	3565.3369993	TRUE
## 54	65000	Strategy A	81005.8306546	FALSE
## 55	65000	Strategy B	9583.5231433	FALSE
## 56	65000	Strategy AB	63022.7415260	FALSE
## 57	70000	Standard of care	6910.0632690	FALSE
## 58	70000	Strategy A	80569.6145406	FALSE
## 59	70000	Strategy B	5757.3725029	TRUE
## 60	70000	Strategy AB	54666.7975089	FALSE
## 61	75000	Standard of care	11505.0686181	FALSE
## 62	75000	Strategy A	81383.6775058	FALSE
## 63	75000	Strategy B	3181.5009418	TRUE
## 64	75000	Strategy AB	47561.1325712	FALSE
## 65	80000	Standard of care	17238.3093138	FALSE

## 66	80000	Strategy A	83335.9758178	FALSE
## 67	80000	Strategy B	1743.8647274	TRUE
## 68	80000	Strategy AB	41593.7029801	FALSE
## 69	85000	Standard of care	23755.3452939	FALSE
## 70	85000	Strategy A	86072.0694140	FALSE
## 71	85000	Strategy B	1090.0237973	TRUE
## 72	85000	Strategy AB	36410.0686734	FALSE
## 73	90000	Standard of care	30806.4109083	FALSE
## 74	90000	Strategy A	89342.1926446	FALSE
## 75	90000	Strategy B	970.2125015	TRUE
## 76	90000	Strategy AB	31760.4640010	FALSE
## 77	95000	Standard of care	38230.5409662	FALSE
## 78	95000	Strategy A	92985.3803187	FALSE
## 79	95000	Strategy B	1223.4656493	TRUE
## 80	95000	Strategy AB	27483.9237721	FALSE
## 81	100000	Standard of care	46063.3396948	FALSE
## 82	100000	Strategy A	97037.2366635	FALSE
## 83	100000	Strategy B	1885.3874676	TRUE
## 84	100000	Strategy AB	23616.0522139	FALSE
## 85	105000	Standard of care	54281.8445962	FALSE
## 86	105000	Strategy A	101474.7991811	FALSE
## 87	105000	Strategy B	2933.0154589	TRUE
## 88	105000	Strategy AB	20133.8868285	FALSE
## 89	110000	Standard of care	62817.0836292	FALSE
## 90	110000	Strategy A	106229.0958303	FALSE
## 91	110000	Strategy B	4297.3775817	TRUE
## 92	110000	Strategy AB	16968.4555747	FALSE
## 93	115000	Standard of care	71721.1628151	FALSE
## 94	115000	Strategy A	111352.2326324	FALSE
## 95	115000	Strategy B	6030.5798575	TRUE
## 96	115000	Strategy AB	14171.8644738	FALSE
## 97	120000	Standard of care	80956.0704381	FALSE
## 98	120000	Strategy A	116806.1978715	FALSE
## 99	120000	Strategy B	8094.6105703	TRUE
## 100	120000	Strategy AB	11706.1018100	FALSE
## 101	125000	Standard of care	90527.9620256	FALSE
## 102	125000	Strategy A	122597.1470752	FALSE
## 103	125000	Strategy B	10495.6252476	FALSE
## 104	125000	Strategy AB	9577.3231107	TRUE
## 105	130000	Standard of care	100377.6364126	FALSE
## 106	130000	Strategy A	128665.8790785	FALSE
## 107	130000	Strategy B	13174.4227245	FALSE
## 108	130000	Strategy AB	7726.3272109	TRUE
## 109	135000	Standard of care	110517.5028876	FALSE
## 110	135000	Strategy A	135024.8031696	FALSE
## 111	135000	Strategy B	16143.4122893	FALSE
## 112	135000	Strategy AB	6165.5233991	TRUE
## 113	140000	Standard of care	120962.5035865	FALSE
## 114	140000	Strategy A	141688.8614847	FALSE
## 115	140000	Strategy B	19417.5360780	FALSE
## 116	140000	Strategy AB	4909.8538112	TRUE
## 117	145000	Standard of care	131681.1198459	FALSE
## 118	145000	Strategy A	148626.5353603	FALSE
## 119	145000	Strategy B	22965.2754272	FALSE

## 120	145000	Strategy AB	3927.7997838	TRUE
## 121	150000	Standard of care	142590.1349892	FALSE
## 122	150000	Strategy A	155754.6081198	FALSE
## 123	150000	Strategy B	26703.4136603	FALSE
## 124	150000	Strategy AB	3136.1446403	TRUE
## 125	155000	Standard of care	153664.6205903	FALSE
## 126	155000	Strategy A	163048.1513371	FALSE
## 127	155000	Strategy B	30607.0223513	FALSE
## 128	155000	Strategy AB	2509.9599546	TRUE
## 129	160000	Standard of care	164850.4494507	FALSE
## 130	160000	Strategy A	170453.0378137	FALSE
## 131	160000	Strategy B	34621.9743015	FALSE
## 132	160000	Strategy AB	1995.1185282	TRUE
## 133	165000	Standard of care	176129.4072113	FALSE
## 134	165000	Strategy A	177951.0531905	FALSE
## 135	165000	Strategy B	38730.0551519	FALSE
## 136	165000	Strategy AB	1573.4060020	TRUE
## 137	170000	Standard of care	187487.7778505	FALSE
## 138	170000	Strategy A	185528.4814459	FALSE
## 139	170000	Strategy B	42917.5488810	FALSE
## 140	170000	Strategy AB	1231.1063544	TRUE
## 141	175000	Standard of care	198912.2934561	FALSE
## 142	175000	Strategy A	193172.0546677	FALSE
## 143	175000	Strategy B	47171.1875764	FALSE
## 144	175000	Strategy AB	954.9516732	TRUE
## 145	180000	Standard of care	210399.8906218	FALSE
## 146	180000	Strategy A	200878.7094495	FALSE
## 147	180000	Strategy B	51487.9078319	FALSE
## 148	180000	Strategy AB	741.8785521	TRUE
## 149	185000	Standard of care	221925.4154186	FALSE
## 150	185000	Strategy A	208623.2918626	FALSE
## 151	185000	Strategy B	55842.5557186	FALSE
## 152	185000	Strategy AB	566.7330621	TRUE
## 153	190000	Standard of care	233493.9174456	FALSE
## 154	190000	Strategy A	216410.8515057	FALSE
## 155	190000	Strategy B	60240.1808354	FALSE
## 156	190000	Strategy AB	434.5648023	TRUE
## 157	195000	Standard of care	245091.7033528	FALSE
## 158	195000	Strategy A	224227.6950291	FALSE
## 159	195000	Strategy B	64667.0898324	FALSE
## 160	195000	Strategy AB	331.6804227	TRUE
## 161	200000	Standard of care	256711.3313377	FALSE
## 162	200000	Strategy A	232066.3806303	FALSE
## 163	200000	Strategy B	69115.8409072	FALSE
## 164	200000	Strategy AB	250.6381209	TRUE
## 165	205000	Standard of care	268349.6561853	FALSE
## 166	205000	Strategy A	239923.7630940	FALSE
## 167	205000	Strategy B	73583.2888446	FALSE
## 168	205000	Strategy AB	188.2926816	TRUE
## 169	210000	Standard of care	280011.0751160	FALSE
## 170	210000	Strategy A	247804.2396409	FALSE
## 171	210000	Strategy B	78073.8308652	FALSE
## 172	210000	Strategy AB	149.0413256	TRUE
## 173	215000	Standard of care	291683.9400780	FALSE

## 174	215000	Strategy A	255696.1622191	FALSE
## 175	215000	Strategy B	82575.8189170	FALSE
## 176	215000	Strategy AB	121.2360008	TRUE
## 177	220000	Standard of care	303360.5755906	FALSE
## 178	220000	Strategy A	263591.8553479	FALSE
## 179	220000	Strategy B	87081.5775194	FALSE
## 180	220000	Strategy AB	97.2012265	TRUE
## 181	225000	Standard of care	315038.0048799	FALSE
## 182	225000	Strategy A	271488.3422534	FALSE
## 183	225000	Strategy B	91588.1298985	FALSE
## 184	225000	Strategy AB	73.9602290	TRUE
## 185	230000	Standard of care	326717.9270273	FALSE
## 186	230000	Strategy A	279387.3220169	FALSE
## 187	230000	Strategy B	96097.1751357	FALSE
## 188	230000	Strategy AB	53.2120896	TRUE
## 189	235000	Standard of care	338400.8501989	FALSE
## 190	235000	Strategy A	287289.3028048	FALSE
## 191	235000	Strategy B	100609.2213972	FALSE
## 192	235000	Strategy AB	35.4649745	TRUE
## 193	240000	Standard of care	350084.9926938	FALSE
## 194	240000	Strategy A	295192.5029159	FALSE
## 195	240000	Strategy B	105122.4869819	FALSE
## 196	240000	Strategy AB	18.9371825	TRUE
## 197	245000	Standard of care	361773.7469258	FALSE
## 198	245000	Strategy A	303100.3147641	FALSE
## 199	245000	Strategy B	109640.3643038	FALSE
## 200	245000	Strategy AB	7.0211278	TRUE
## 201	250000	Standard of care	373467.7937629	FALSE
## 202	250000	Strategy A	311013.4192174	FALSE
## 203	250000	Strategy B	114163.5342307	FALSE
## 204	250000	Strategy AB	0.3976781	TRUE

09.4.4 Expected value of perfect information (EVPI)

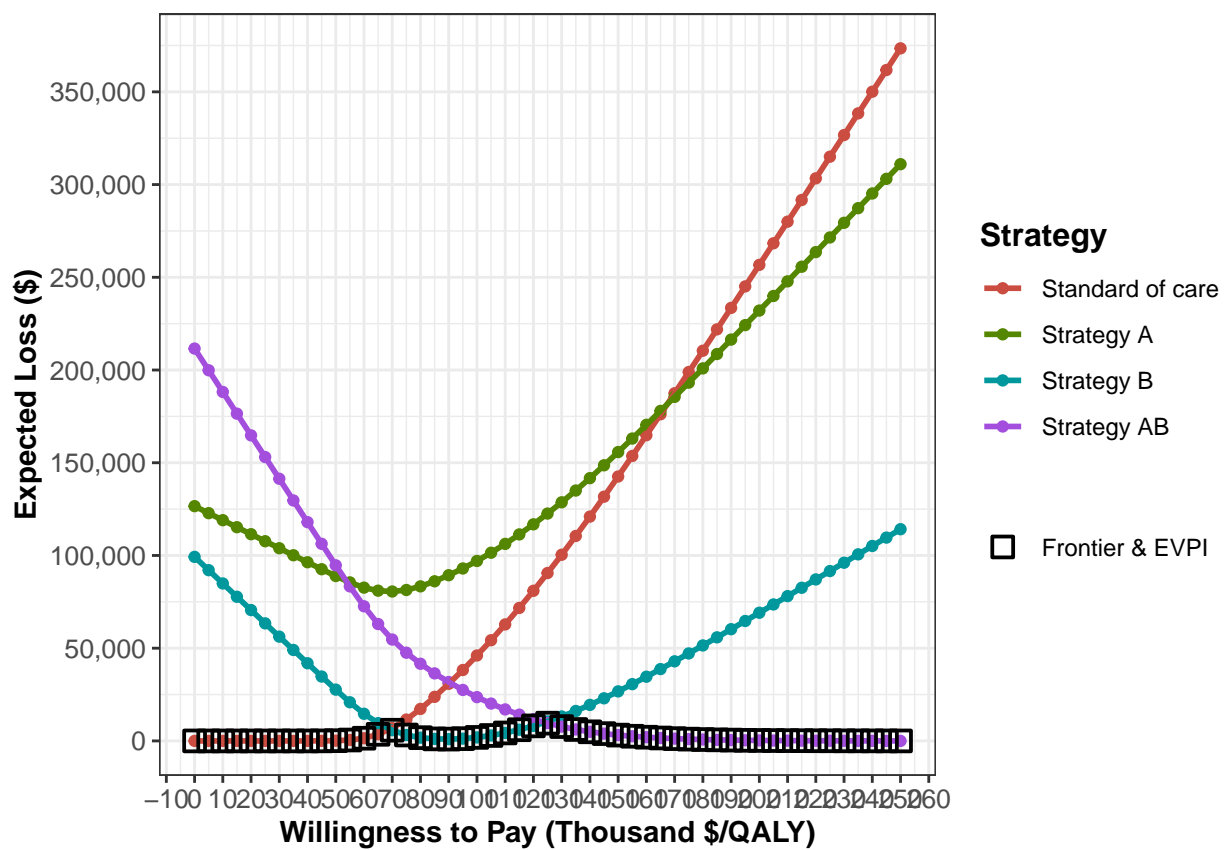


Figure 15: Expected loss curves (ELCs).

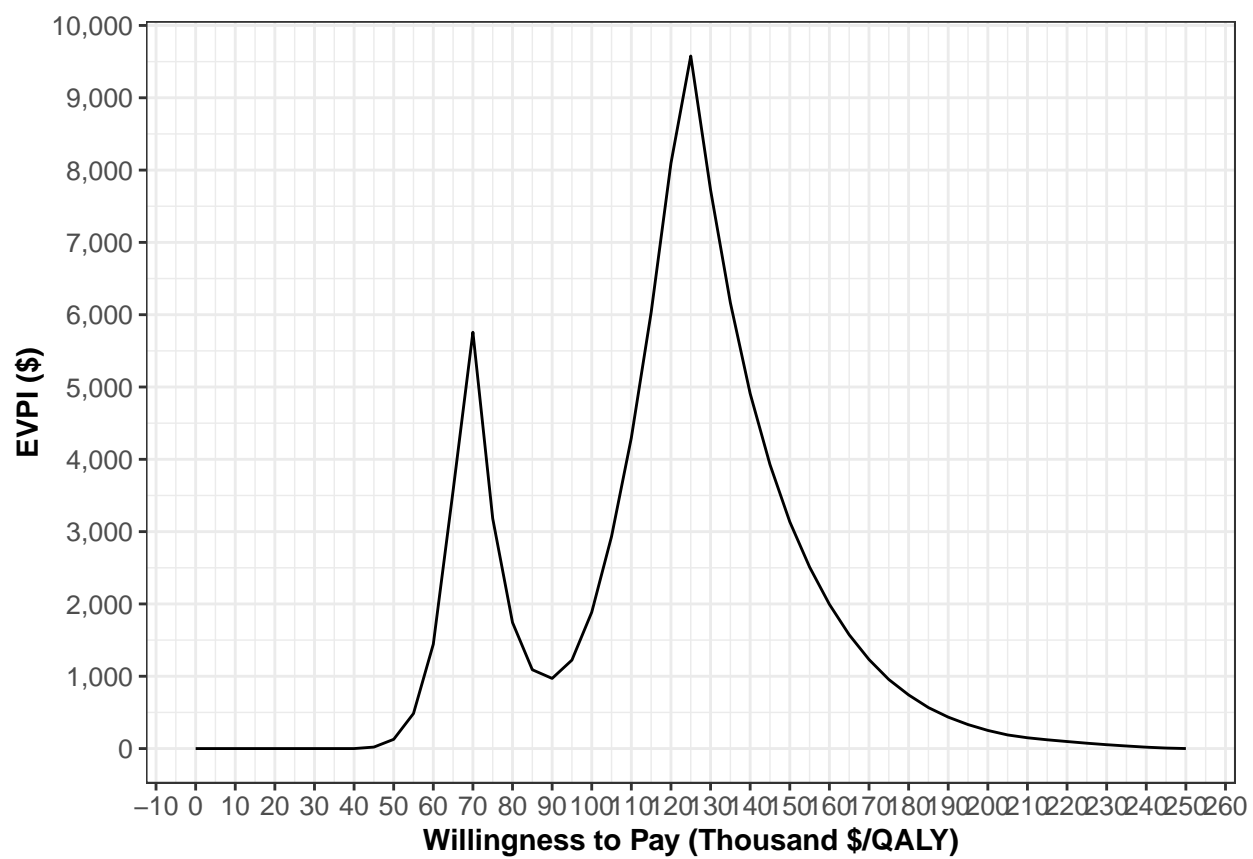


Figure 16: Expected value of perfect information (EVPI).