Decision Tree Example in R - No Spray, Spray, Test

The DARTH workgroup

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Please cite our publications when using this code:

- Alarid-Escudero F, Krijkamp EM, Enns EA, Yang A, Hunink MGM Pechlivanoglou P, Jalal H. An Introductory Tutorial on Cohort State-Transition Models in R Using a Cost-Effectiveness Analysis Example. Medical Decision Making, 2022 (Epub). https://doi.org/10.1177/0272989X221103163
- Alarid-Escudero F, Krijkamp EM, Enns EA, Yang A, Hunink MGM Pechlivanoglou P, Jalal H. A
 Tutorial on Time-Dependent Cohort State-Transition Models in R using a Cost-Effectiveness Analysis
 Example. Medical Decision Making, 2022 (Epub). https://doi.org/10.1177/0272989X221121747
- 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 Mak. 2020;40(2):242-248. https://doi.org/10.1177/0272989X19893973

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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

```
# no packages needed
```

02 Load functions

```
# no functions needed
```

03 Model input

```
### Strategies
                  <- c("Do not spray",
v_names_strat
                                          # store the strategy names
                      "Spray")
# your turn #
# add a strategy name "Test"
                 n_strat
### Branch probabilities
p_outbreak <- 0.20 # probability that there is an outbreak</pre>
{\tt p\_die\_tox} \, \mathrel{<\!\!\!\!-} \, 0.001 \quad \textit{\# probability of dying due to exposure to spray}
p_die_inf <- 0.33  # probability of dying if infected</pre>
# probability of infection under each strategy
p_inf_outbreak_nospray <- 0.02 # probability of becoming infected if did not spray
p_inf_outbreak_spray <- 0.003 # probability of becoming infected if did spray (right away)</pre>
# your turn #
# add parameter here: # probability of becoming infected if did spray after test
### Terminal node values
# Number of deaths
n_death_die <- 1  # terminal value if pathway results in death
n_death_survive <- 0 # terminal value if pathway does not result in death
# Costs
c_inf_survive <- 10000</pre>
c_inf_die <- 20000</pre>
```

```
c_tox_die <- 5000
c_spray <- 1500

# your turn #
# add parameter here: Cost of test #</pre>
```

04 Construct and evaluate decision tree model equations

```
# Vector of expected values for each strategy
# One vector for deaths, the other for costs
v_EV_death <- v_EV_cost <- rep(NA,n_strat)</pre>
names(v_EV_death) <- v_names_strat # attach strategy name</pre>
names(v_EV_cost) <- v_names_strat # attach strategy name</pre>
# Do not spray
v_EV_death["Do not spray"] <- p_outbreak*p_inf_outbreak_nospray*p_die_inf*n_death_die
v_EV_cost["Do not spray"] <- p_outbreak*p_inf_outbreak_nospray*p_die_inf*c_inf_die +
                              p_outbreak*p_inf_outbreak_nospray*(1-p_die_inf)*c_inf_survive
# Spray
v_EV_death["Spray"] <- p_die_tox*n_death_die +</pre>
                       (1-p_die_tox)*p_outbreak*p_inf_outbreak_spray*p_die_inf*n_death_die
v_EV_cost["Spray"] <- c_spray +</pre>
                      p_die_tox*c_tox_die +
                       (1-p_die_tox)*p_outbreak*p_inf_outbreak_spray*p_die_inf*c_inf_die +
                       (1-p_die_tox)*p_outbreak*p_inf_outbreak_spray*(1-p_die_inf)*c_inf_survive
# your turn #
# Test
# Fill in equation to calculate expected deaths and costs under test strategy
```

05 Summarize Output