

An Introductory Tutorial on Cohort State-Transition Models in R Using a Cost-Effectiveness Analysis Example

Appendix

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Cohort tutorial model components

Table I

This table contains an overview of the key model components used in the code for the Sick-Sicker example from the DARTH manuscript: “An Introductory Tutorial to Cohort State-Transition Models in R”. The first column gives the mathematical notation for some of the model components that are used in the equations in the manuscript. The second column gives a description of the model component with the R name in the third column. The forth gives the data structure, e.g. scalar, list, vector, matrix etc, with the according dimensions of this data structure in the fifth column. The final column indicated the type of data that is stored in the data structure, e.g. numeric (5,2,6,3,7,4), category (A,B,C), integer (5,6,7), logical (TRUE, FALSE).

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Element	Description	R name	Data structure	Dimensions	Data type
n_t	Time horizon	<code>n_cycles</code>	scalar		numeric
	Cycle length	<code>cycle_length</code>	scalar		numeric
v_s	Names of the health states	<code>v_names_states</code>	vector	$\mathbf{n_states} \times 1$	character
n_s	Number of health states	<code>n_states</code>	scalar		numeric
v_{str}	Names of the strategies	<code>v_names_str</code>	scalar		character
n_{str}	Number of strategies	<code>n_str</code>	scalar		character
$\mathbf{d_c}$	Discount rate for costs	<code>d_c</code>	scalar		numeric
$\mathbf{d_e}$	Discount rate for effects	<code>d_e</code>	scalar		numeric
	Discount weights for costs	<code>v_dwc</code>	vector	$(\mathbf{n_t} \times 1) + 1$	numeric
	Discount weights for effects	<code>v_dwe</code>	vector	$(\mathbf{n_t} \times 1) + 1$	numeric
	Sequence of cycle numbers	<code>v_cycles</code>	vector	$(\mathbf{n_t} \times 1) + 1$	numeric
\mathbf{wcc}	Within-cycle correction weights using Simpson's 1/3 rule	<code>v_wcc</code>	vector	$(\mathbf{n_t} \times 1) + 1$	numeric
age_0	Age at baseline	<code>n_age_init</code>	scalar		numeric
age	Maximum age of follow up	<code>n_age_max</code>	scalar		numeric
M	Cohort trace	<code>m_M</code>	matrix	$(\mathbf{n_t} + 1) \times \mathbf{n_states}$	numeric
m_0	Initial state vector	<code>v_m_init</code>	vector	$1 \times \mathbf{n_states}$	numeric
m_t	State vector in cycle t	<code>v_mt</code>	vector	$1 \times \mathbf{n_states}$	numeric
Transition probabilities and rates					
$p_{[H,S1]}$	From Healthy to Sick conditional on surviving	<code>p_HS1</code>	scalar		numeric
$p_{[S1,H]}$	From Sick to Healthy conditional on surviving	<code>p_S1H</code>	scalar		numeric
$p_{[S1,S2]}$	From Sick to Sicker conditional on surviving	<code>p_S1S2</code>	scalar		numeric
$p_{[S1,S2]_{trtB}}$	From Sicker to Sick under treatment B conditional on surviving	<code>p_S1S2_trtB</code>	scalar		numeric
$r_{[H,D]}$	Constant rate of dying when Healthy (all-cause mortality rate)	<code>r_HD</code>	scalar		numeric
$r_{[S1,S2]}$	Constant rate of becoming Sicker when Sick	<code>r_S1S2</code>	scalar		numeric
$r_{[S1,S2]_{trtB}}$	Constant rate of becoming Sicker when Sick for treatment B	<code>r_S1S2_trtB</code>	scalar		numeric
$hr_{[S1,H]}$	Hazard ratio of death in Sick vs Healthy	<code>hr_S1</code>	scalar		numeric
$hr_{[S2,H]}$	Hazard ratio of death in Sicker vs Healthy	<code>hr_S2</code>	scalar		numeric
$hr_{[S1,S2]_{trtB}}$	Hazard ratio of becoming Sicker when Sick under treatment B	<code>hr_S1S2_trtB</code>	scalar		numeric
P	Time-independent transition probability matrix* * <code>_trtX</code> is used to specify for which strategy the transition probability matrix is	<code>m_P</code>	matrix	$\mathbf{n_states} \times \mathbf{n_states}$	numeric

Element	Description	R name	Data structure	Dimensions	Data type
Annual costs					
	Healthy individuals	c_H	scalar		numeric
	Sick individuals in Sick	c_S1	scalar		numeric
	Sick individuals in Sicker	c_S2	scalar		numeric
	Dead individuals	c_D	scalar		numeric
	Additional costs treatment A	c_trtA	scalar		numeric
	Additional costs treatment B	c_trtB	scalar		numeric
	Vector of state costs for a strategy	v_c_str	vector	1 x n_states	numeric
	list that stores the vectors of state costs for each strategy	l_c	list		numeric
Utility weights					
	Healthy individuals	u_H	scalar		numeric
	Sick individuals in Sick	u_S1	scalar		numeric
	Sick individuals in Sicker	u_S2	scalar		numeric
	Dead individuals	u_D	scalar		numeric
	Treated with treatment A	u_trtA	scalar		numeric
	Vector of state utilities for a strategy	v_u_str	vector	1 x n_states	numeric
	List that stores the vectors of state utilities for each strategy	l_u	list		numeric
Outcome structures					
	Expected QALYs per cycle under a strategy	v_qaly_str	vector	1 x (n_t + 1)	numeric
	Expected costs per cycle under a strategy	v_cost_str	vector	1 x (n_t + 1)	numeric
	Vector of expected discounted QALYs for each strategy	v_tot_qaly	vector	1 x n_states	numeric
	Vector of expected discounted costs for each strategy	v_tot_cost	vector	1 x n_states	numeric
	Summary matrix with costs and QALYS per strategy	m_outcomes	table	n_states x 2	
	Summary of the model outcomes	df_cea	data frame		
	Summary of the model outcomes	table_cea	table		
Probabilistic analysis structures					
	Number of PSA iterations	n_sim	scalar		numeric

Element	Description	R name	Data structure	Dimensions	Data type
	List that stores all the values of the input parameters	<code>l_params_all</code>	list		numeric
	Vector with the names of all the input parameters	<code>v_names_params</code>	vector		character
	List with the model outcomes of the PSA for all strategies	<code>l_psa</code>	list		numeric
	Vector with a sequence of relevant willingness-to-pay values	<code>v_wtp</code>	vector		numeric
	Data frame to store expected costs and effects for each strategy from the PSA	<code>df_out_ce_psa</code>	table		numeric
	Data frame to store incremental cost-effectiveness ratios (ICERs) from the PSA	<code>df_cea_psa</code>	table		numeric
	For more details about the PSA structures read the vignettes of <code>dampack</code>				