## Cohort State-Transition Models in R: A Tutorial 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: "Implementation of 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).

Element	Description	R name	Data structure	Dimensions	Data type
$\overline{n_t}$	Time horizon	n_t	scalar		numeric
$v_s$	Names of the health states	v_n	vector	${ t n\_states} \ { t x} \ 1$	categoric
$n_s$	Number of health states	n_states	$\operatorname{scalar}$		numeric
$v_{str}$	Names of the strategies	v_names_str	$\operatorname{scalar}$		categoric

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Element	Description	R name	Data structure	Dimensions	Data type
$n_{str}$	Number of strategies	n_str	scalar		categoric
$ ext{d}_{\mathbf{c}}$	Discount rate for costs	d_c	scalar		numeric
$\mathbf{d_e}$	Discount rate for effects	d_e	scalar		numeric
	Discount weights for costs	v_dwc	vector	$(\mathtt{n\_t} \ \mathtt{x} \ 1 \ ) + 1$	numeric
	Discount weights for effects	v_dwe	vector	$(n_t x 1) + 1$	numeric
$age_{_0}$	Age at baseline	n_age_init	scalar		numeric
age	Maximum age of follow up	n_age_max	scalar		numeric
M	Cohort trace	m_M	matrix	$(\mathtt{n\_t}+1) \ge \mathtt{n\_states}$	numeric
$m_0$	Initial state vector	v_s_init	vector	$1 \times n$ _states	numeric
$m_t$	State vector in cycle t	v_mt	vector	$1 \times n_s$ tates	numeric
	Transition probabilities				
$p_{[H,S1]}$	From Healthy to Sick conditional on surviving	p_HS1	scalar		numeric
$p_{[S1,H]}$	From Sick to Healthy conditional on surviving	p_S1H	scalar		numeric
$p_{[S1,S2]}$	From Sick to Sicker conditional on surviving	p_S1S2	scalar		numeric
$r_{[H,D]}$	Constant rate of dying when Healthy (all-cause mortality rate)	r_HD	scalar		numeric
$hr_{[S1,H]}$	Hazard ratio of death in Sick vs Healthy	hr_S1	scalar		numeric
$hr_{[S2,H]}$	Hazard ratio of death in Sicker vs Healthy	hr_S2	scalar		numeric
$hr_{[S1,S2]_{trtB}}$		hr_S1S2_trtB	scalar		numeric
P	Time-independent transition probability matrix	m_P	matrix	n_states x n_states	numeric
	Annual costs				
	Healthy individuals	c_H	scalar		numeric
	Sick individuals in S1	c_S1	scalar		numeric
	Sick individuals in S2	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
	Utility weights				
	Healthy individuals	u_H	scalar		numeric
	Sick individuals in S1	u_S1	scalar		numeric
	Sick individuals in S2	_ u_S2	scalar		numeric
	Dead individuals	u_D	scalar		numeric
	Treated with treatment A	u_trtA	scalar		$_{ m numeric}$

Element	Description	R name	Data structure	Dimensions	Data type
A	Outcome structures Transition dynamics array	a_A	array	${ t n\_states \ x \ n\_states \ x} \ ({ t n\_t+1})$	numeric
	Expected QALYs per cycle under a strategy Expected costs per cycle under a strategy Total expected discounted QALYs for a strategy Total expected discounted costs for a strategy Summary of the model outcomes Summary of the model outcomes	<pre>v_qaly v_cost n_tot_qaly n_tot_cost df_cea table_cea</pre>	vector vector scalar scalar data frame table	$(n_{t} + 1)$ $1 \times (n_{t} + 1)$ $1 \times (n_{t} + 1)$	numeric numeric numeric numeric