

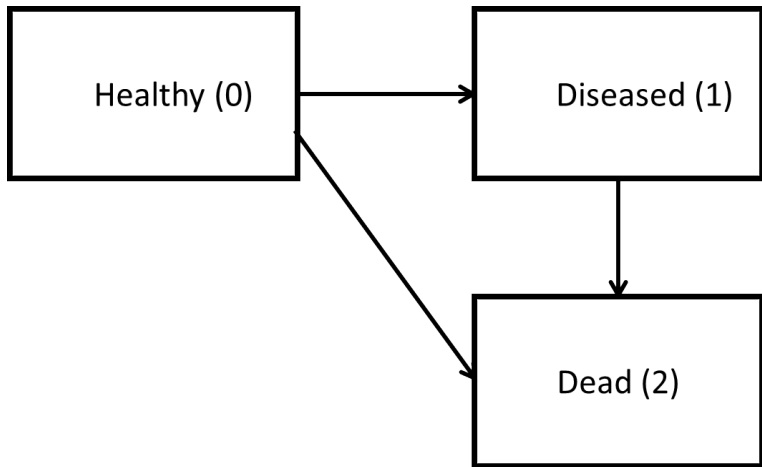
# SURVIVAL ANALYSIS

Event-history models  
PART VI, HY 2019

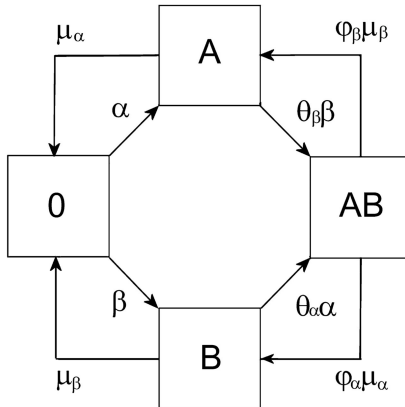
# Outline

- ▶ Examples on multi-state models
- ▶ Model specification
- ▶ Likelihood function
- ▶ Health-illness model
- ▶ Reference: Andersen et al. Multi-state models for event history analysis, Stat Methods Med Res, 2012

# Health-illness model



# Infection model for two competing strains



Auranen K et al. Am. J. Epidemiol. 2009;171:169-176

# Event-history models

- ▶ Event-history models or multi-state models are generalisations of simple survival or competing risks models
- ▶ There are a number of different states between which the individual can make transitions
- ▶ The model can contain an absorbing state(s) and/or it can accommodate repeated transitions between the model states
- ▶ The interest is in the estimation of transition-specific hazard functions and quantities that can be derived from those
- ▶ Numerous uses in epidemiology, medicine, biosciences, social sciences, engineering, ...

# Model specification

- ▶  $X_i(t) \in \{1, \dots, k\}$  is the state of individual  $i$  at time  $t$ ,  $t > 0$
- ▶ Time  $\tau_i$  is the termination of observation for individual  $i$ 
  - ▶ This can be the transition time to an absorbing state or censoring time
- ▶ Observations on  $X_i(t)$  are equivalent to observations from counting processes for each possible transition in the model

$$N_{hj}^i = \#(\text{direct transitions from } h \text{ to } j \text{ in } [0, t] \text{ for } i), \text{ with data} \\ 0 < T_{hj}^{i1} < \dots < T_{hj}^{i, N_{hj}^i(\tau_i)} \leq \tau_i$$

- ▶ The hazard for individual  $i$  to make transition  $h \rightarrow j$  at time  $t$ , conditionally on being in state  $h$  at  $t-$ , is denoted by  $\alpha_{hj}^i(t)$ , for all  $h, j \in \{1, \dots, k\}$ 
  - ▶ Some of the hazards can be 0 (i.e. for those transitions that do not exist in the model)

# Likelihood function

- ▶ The likelihood is constructed from individual and transition specific contributions
- ▶ For each individual  $i$  and transition  $h \rightarrow j$ , the likelihood expression consists of contributions from all transition events  $h \rightarrow j$  of the individual, evaluated at the times of those transitions, as well as from the 'at-risk' time spent in state  $h$  before those transitions:

$$\prod_{h \neq j} \prod_{k=1}^{N_{hj}^i(\tau_i)} \alpha_{hj}^i(T_{hj}^{ik}) \exp \left( - \int_0^{\tau_i} \alpha_{hj}^i(t) Y_h^i(t) dt \right)$$

- ▶ For simplicity of notation, it is assumed above that the terms  $\alpha_{hj}^i(T_{hj}^{ik})$  appear only if such transitions occur
- ▶ Note how the the risk indicator takes care of the appropriate at-risk time in each of the states

# Remarks

- ▶ In the above presentation, only one time variable is used; however, the hazard on making a specific transition may well depend on e.g. the *duration of stay* in that state, or on more than one time variable
- ▶ The likelihood function guides in writing the data matrix for statistical analysis
  - ▶ Each individual's each transition type ( $h \rightarrow j$ ) produces as many data matrix rows as there are arrows from  $h$  to any other states of the model
  - ▶ In fact, this is exactly the same situation than we have already seen in competing risks models: the states *to which* there is an arrow *from* state  $h$  are competing over which of these events actually occur



## Example: health-illness model

- ▶ In the model of page 3, there are three possible transitions, with hazard functions  $\alpha_{01}(t)$ ,  $\alpha_{02}(t)$  and  $\alpha_{12}(t)$
- ▶ The likelihood contribution from individual  $i$  is now

$$\begin{aligned} L_i(\text{parameters}) = & (\alpha_{01}^i(T_{01}))^{d_{01}^i} \exp\left(-\int_0^{\tau_i} Y_{i0}(t)\alpha_{01}(t)dt\right) \times \\ & (\alpha_{02}^i(T_{02}))^{d_{02}^i} \exp\left(-\int_0^{\tau_i} Y_{i0}(t)\alpha_{02}(t)dt\right) \times \\ & (\alpha_{12}^i(T_{12}))^{d_{12}^i} \exp\left(-\int_0^{\tau_i} Y_{i1}(t)\alpha_{12}(t)dt\right) \end{aligned}$$

- ▶ The likelihood function based on data from  $n$  independent individuals is now

$$L(\text{parameters}) = \prod_{i=1}^n L_i(\text{parameters})$$

## Example continues

- ▶ A toy example modified from Andersen et al.
- ▶ Health-illness model parameterised as follows:

$$\alpha_{01}^i(t) = \alpha_{01}(t)e^{\beta_1 Z_{1i} + \beta_2 Z_{2i}}$$

$$\alpha_{02}^i(t) = \alpha_{02}(t)e^{\beta_3 Z_{1i} + \beta_4 Z_{2i}}$$

$$\alpha_{12}^i(t) = \alpha_{02}(t)e^{\gamma + \beta_1 Z_{1i}}$$

- ▶ The death rates are proportional, with the hazard ratio  $\exp(\gamma)$
- ▶ The covariates  $Z_{1i}$  and  $Z_{2i}$  have different proportional effects on the baseline rates  $\alpha_{01}(t)$  and  $\alpha_{02}(t)$
- ▶ Covariate  $Z_{1i}$  has the same effect on  $\alpha_{01}(t)$  and  $\alpha_{12}(t)$

## Example continues

- ▶ For any one individual  $i$ , the data matrix contains three rows, where three is now the number of possible transitions in the model
- ▶ For fitting the Cox proportional hazards model, the three data matrix rows from individual  $i$  are

Time	Status	Stratum	$\log(\gamma)$	$\log(\beta_1)$	$\log(\beta_2)$	$\log(\beta_3)$	$\log(\beta_4)$
$T_{01}^i$	$d_{01}^i$	0	0	$Z_{1i}$	$Z_{2i}$	0	0
$T_{02}^i$	$d_{02}^i$	1	0	0	0	$Z_{1i}$	$Z_{2i}$
$T_{12}^i$	$d_{12}^i$	1	1	$Z_{1i}$	0	0	0

```
coxph(Surv(Time,Status)~  
logg + logb1 + logb2 + logb3 + logb4 + strata(Stratum), data = A)
```