Olli Saarela

Likelihood construction under independent and noninformative censoring

constant hazard mode

Survival Analysis I (CHL5209H)

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Piecewise constant hazard mode

- ▶ Previously we noted that the hazard function specifies the event time distribution in the absence of censoring.
- What about in the presence of censoring?
- We need assumptions about the censoring.
- ▶ To express these, suppose that the observed time is given by $T_i = \min\{\tilde{T}_i, C_i\}$, where \tilde{T}_i and C_i are latent event and censoring times.
- ► Further, we can define the event indicator as $E_i = \mathbf{1}_{\{T_i = \tilde{T}_i\}}$.

Parametrized hazard function

- Often we are interested in modeling the relationship between survival and some covariates X_i .
- ▶ This dependency is parametrized through a vector of parameters θ .
- ► The resulting individual-level hazard function may be defined as

$$\lambda_i(t;\theta) \equiv \frac{P(t \leq \tilde{T}_i < t + dt \mid \tilde{T}_i \geq t, x_i; \theta)}{dt}$$

$$= \frac{f_i(t;\theta)}{S_i(t;\theta)},$$

where f_i and S_i are the density and survival functions characterizing the latent event time distribution.

▶ How to estimate θ in the presence of censoring?

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Bring in the censoring times

▶ We can write this as

$$\lambda_{i}(t;\theta) = \frac{f_{i}(t;\theta)}{S_{i}(t;\theta)}$$

$$= \frac{P(t \leq \tilde{T}_{i} < t + dt \mid x_{i};\theta)/dt}{P(\tilde{T}_{i} \geq t \mid x_{i};\theta)}$$

$$= \frac{P(C_{i} > t \mid x_{i};\theta)P(t \leq \tilde{T}_{i} < t + dt \mid x_{i};\theta)/dt}{P(C_{i} \geq t \mid x_{i};\theta)P(\tilde{T}_{i} \geq t \mid x_{i};\theta)}$$

Independent censoring

Assume that given the covariates X_i , the latent event and censoring times are independent:

$$\lambda_{i}(t;\theta) = \frac{P(C_{i} > t \mid x_{i};\theta)P(t \leq \tilde{T}_{i} < t + dt \mid x_{i};\theta)/dt}{P(C_{i} \geq t \mid x_{i};\theta)P(\tilde{T}_{i} \geq t \mid x_{i};\theta)}$$

$$= \frac{P(t \leq \tilde{T}_{i} < t + dt, C_{i} > t \mid x_{i};\theta)/dt}{P(\tilde{T}_{i} \geq t, C_{i} \geq t \mid x_{i};\theta)}$$

$$= \frac{P(t \leq T_{i} < t + dt, E_{i} = 1 \mid x_{i};\theta)/dt}{P(T_{i} \geq t \mid x_{i};\theta)}$$

$$= \frac{P(t \leq T_{i} < t + dt, E_{i} = 1 \mid T_{i} \geq t, x_{i};\theta)}{dt}.$$

Thus, under independent censoring, the hazard functions defined in terms of the latent and observed times are equivalent.

Likelihood contributions

- ▶ The observed data consist of realizations of random vectors (T_i, E_i, X_i) , i = 1, ..., n.
- ► Conditional on X_i , the likelihood contribution for a non-censored individual i with $E_i = 1$ is

$$P(t_i \leq T_i < t_i + dt, E_i = 1 \mid x_i; \theta)$$

$$= P(C_i > t_i \mid x_i; \theta) P(t_i \leq \tilde{T}_i < t_i + dt \mid x_i; \theta)$$

$$= P(C_i > t_i \mid x_i; \theta) \lambda_i(t_i; \theta) dt S_i(t_i; \theta).$$

Similarly, the likelihood contribution for a censored individual i with $E_i = 0$ is

$$P(t_i \leq T_i < t_i + dt, E_i = 0 \mid x_i; \theta)$$

$$= P(t_i \leq C_i < t_i + dt \mid x_i; \theta) P(\tilde{T}_i > t_i \mid x_i; \theta)$$

$$= P(t_i \leq C_i < t_i + dt \mid x_i; \theta) S_i(t_i; \theta).$$

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Non-informative censoring

- Assume further that the distribution of the latent censoring time does not involve θ .
- \blacktriangleright We can now write the full likelihood for θ as

$$\prod_{i=1}^{n} P(t_{i} \leq T_{i} < t_{i} + dt, E_{i} = e_{i} \mid x_{i}; \theta)$$

$$\stackrel{\theta}{\propto} \prod_{i=1}^{n} [\lambda_{i}(t_{i}; \theta)^{e_{i}} S_{i}(t_{i}; \theta)]$$

$$= \prod_{i=1}^{n} \left[\lambda_{i}(t_{i}; \theta)^{e_{i}} \exp \left\{ - \int_{0}^{t_{i}} \lambda_{i}(u; \theta) du \right\} \right].$$

- ▶ Note that with these assumptions, the likelihood is fully specified by the hazard function for the event of interest.
- Maximum likelihood criterion can now be applied as usual to estimate θ .

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Interpretation?

- ▶ What did we have to assume to obtain the previous likelihood expression?
- Independent and non-informative censoring are rather abstract properties.
- Note that the independence of the censoring mechanism was conditional on the covariates X_i .
- Such independence is more believable if we can condition on all common determinants of the censoring events and the events of interest.

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Parametric survival models

- We can parametrize the hazard function through a regression equation.
- For example,

$$\lambda_i(u;\theta) = \exp\{\alpha + \beta' X_i\},\,$$

where $\theta = (\alpha, \beta)$, would specify a Poisson regression model, with the baseline rate given by $\exp{\{\alpha\}}$ and the regression coefficients having interpretation as log-rate ratios (this is a special case of a proportional hazards model).

- Usually, we would not want to assume the hazard to be constant over time.
- ▶ A generalization of this model is obtained if we assume that hazard to be constant over pre-specified intervals.
- ► This also allows us to easily incorporate more than one time scale.

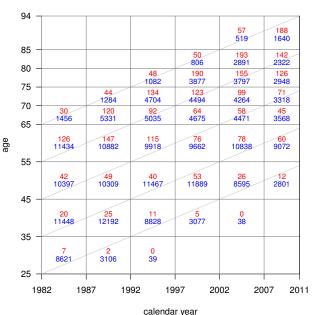
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Piecewise constant hazard model

Lexis diagram



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Piecewise constant hazard model

Notation for grouped follow-up data

- ▶ The Lexis diagram depicted the follow-up for total mortality of 9029 individuals recruited as a cross-sectional cohort in 1982 (then of age 25-65) until the end of year 2010.
- Assume that the mortality rate is constant within the agegroups $k=1,\ldots,9$ in the Lexis diagram, and within one-year calendar time intervals $l=1,\ldots,29$.
- Let $d_{ikl} \in \{0,1\}$ denote whether individual i died at age k in year l.
- Let y_{ikl} denote the person-years individual i contributed in age group k and year l.
- If we have no other individual level information, the hazard rate of any individual i in age group k and year l is assumed to be λ_{kl} .
- ▶ This is why the model is called piecewise constant.

Piecewise constant hazard model ▶ The likelihood contribution of individual *i* is given by

$$\begin{split} &\prod_{k=1}^{9} \prod_{l=1}^{29} \lambda_{kl}^{d_{ikl}} \exp \left\{ -\sum_{k=1}^{9} \sum_{l=1}^{29} \int_{0}^{y_{ikl}} \lambda_{kl} \, \mathrm{d}t \right\} \\ &= \prod_{k=1}^{9} \prod_{l=1}^{29} \left[\lambda_{kl}^{d_{ikl}} \exp \left\{ -\lambda_{kl} y_{ikl} \right\} \right]. \end{split}$$

▶ The likelihood expression from *n* individuals is then

$$\prod_{i=1}^{n} \prod_{k=1}^{9} \prod_{l=1}^{29} \left[\lambda_{kl}^{d_{ikl}} \exp \left\{ -\lambda_{kl} y_{ikl} \right\} \right].$$

Connection to Poisson model

▶ Since λ_{kl} does not depend on i, we get

$$\begin{split} & \prod_{i=1}^{n} \prod_{k=1}^{9} \prod_{l=1}^{29} \left[\lambda_{kl}^{d_{ikl}} \exp \left\{ -\lambda_{kl} y_{ikl} \right\} \right] \\ & = \prod_{k=1}^{9} \prod_{l=1}^{29} \left[\lambda_{kl}^{\sum_{i=1}^{n} d_{ikl}} \exp \left\{ -\lambda_{kl} \sum_{i=1}^{n} y_{ikl} \right\} \right]. \end{split}$$

We would get the same likelihood expression if we assume the total number of deaths in each age group/year to be independently Poisson distributed as

$$\sum_{i=1}^n d_{ikl} \sim \text{Poisson}\left(\lambda_{kl} \sum_{i=1}^n y_{ikl}\right).$$

► Thus, the model can be fitted using any available Poisson regression software such as the glm function in R.

More general models

We can allow the piecewise constant mortality rates to further depend on individual-level covariates X_i , in which case the likelihood expression is of the form

$$\prod_{i=1}^{n}\prod_{k=1}^{9}\prod_{l=1}^{29}\left[\lambda_{ikl}^{d_{ikl}}\exp\left\{-\lambda_{ikl}y_{ikl}\right\}\right].$$

While there are no Poisson distributed counts here, the model can still be fitted as a Poisson regression. (Why?)

Model parametrization

- Even without further individual-level characteristics, the example model involved $9 \times 29 = 261$ mortality rate parameters λ_{kl} .
- Not all of these can be estimated, since some age group/year combinations have no observed deaths. (Why?)
- Estimating this many parameters would also be inefficient, and interpretation of the results would be difficult.
- ▶ Suppose that we are mainly interested in the calendar time trend in mortality, while removing the age effect.
- Further, we allow the mortality rate to depend covariates sex $(x_{i1} \in \{0,1\}, \text{ men/women})$ and region $(x_{i2} \in \{0,1\}, \text{ eastern/western Finland})$.
- ▶ A more parsimonious parametrization can now be specified through a regression equation.

Regression equation

▶ For example, we can specify the model as

$$\log(\lambda_{ikl}) = \alpha_k + \beta_l + \gamma_1 x_{i1} + \gamma_2 x_{i2}.$$

- ▶ This model involves only 9 + 29 + 2 = 40 parameters.
- Now we are mainly interested in the calendar time effect parameters β_I , $I=1,\ldots,29$.
- Adjustment for age through the parameters α_k , $k=1,\ldots,9$ is needed to exctract the calendar time effect. (What would happen to the calendar time effect if we did not adjust for age?)
- Note that for this model to be identifiable, one parameter restriction, for example

$$\alpha_1 = 0$$

is needed.

Parametrization with an intercept term

▶ An alternative way to parametrize the same model would be

$$\log(\lambda_{ikl}) = \mu + \alpha_k + \beta_l + \gamma_1 x_{i1} + \gamma_2 x_{i2},$$

which includes a separate intercept term μ .

Now two parameter restrictions are required, for example

$$\alpha_1 = 0 \text{ and } \beta_1 = 0.$$

▶ The interpretation of the calendar time effects β_I , $l=2,\ldots,29$ is now different, they represent log-rate ratios w.r.t. the first year.