

# Part IIA: Introduction to parametric survival models and the survival likelihood

- ▶ The exponential model
  - ▶ Properties of the constant-hazard model
  - ▶ Maximum likelihood estimation of the hazard
  - ▶ Intro to exponential regression models and Poisson regression
- ▶ Survival likelihood under the general hazard model
  - ▶ Survival (Poisson) likelihood
  - ▶ Cumulative hazards
- ▶ Examples of survival models
  - ▶ Weibull
  - ▶ Log-logistic

# The exponential model

- ▶ The exponential model is the simplest survival model
- ▶ It is characterised by one parameter, a constant hazard  $\lambda$
- ▶ An analogy to radioactive decay
  - ▶ Each nucleus has, irrespective of how long it has remained intact, a constant rate  $\lambda$  to decay
  - ▶ Starting with  $N$  nuclei at time  $t = 0$ , the number of nuclei remaining at time  $t$  is  $N \exp(-\lambda t)$
- ▶ In other words, the survival function of the exponential distribution is  $S(t) = P(T > t) = \exp(-\lambda t)$ ,  $t \geq 0$
- ▶ Although the exponential model may appear too simplistic in most applications, it is sometimes useful on its own right and quite often serves as a building block for more realistic models (e.g. models with pieewise constant hazards; see part IV)

# Characteristics of the exponential distribution

- ▶ The survival function is  $S(t) = \exp(-\lambda t)$ ,  $t \geq 0$
- ▶ The density function is  $f(t) = \lambda \exp(-\lambda t)$ ,  $t \geq 0$
- ▶ The expected time until failure is  $1/\lambda$ 
  - ▶ This is called *scale* ( $= 1/\text{hazard}$ ) and sometimes used as the parameter of the exponential model
  - ▶ The dimension of hazard is “1/time” and that of scale “time”
- ▶ The hazard does not depend on the history of the individual, not even on how long the individual has survived so far
  - ▶ This is why the distribution is said to be memoryless
  - ▶ The expected remaining lifetime at any(!) time  $t$  is always the same, i.e.  $1/\lambda$
- ▶ N.B. Terms *hazard*, *hazard rate* or even *rate* are often used interchangeably

# Survival likelihood under the exponential model

- ▶ Next we construct the likelihood function for a constant hazard  $\lambda$ , based on a follow-up of a study cohort with the following observations
  - ▶ occurrence times (failures and censorings)  $t_i$ ,  $i = 1, \dots, N$
  - ▶ failure (event) indicators  $d_i$ ,  $i = 1, \dots, N$
- ▶ Assuming independent observations across different individuals, the likelihood function based on the above data is a product over individual contributions:

$$\begin{aligned} L(\lambda) &= \prod_{i=1}^N L_i(\lambda) = \prod_{i=1}^N \lambda^{d_i} S(t_i) = \lambda^{\sum_{i=1}^N d_i} \times \prod_{i=1}^N \exp(-\lambda t_i) \\ &= \lambda^D \exp(-\lambda \sum_{i=1}^N t_i) = \lambda^D \exp(-\lambda Y) \end{aligned}$$

- ▶  $L(\lambda)$  is often called the Poisson likelihood (see page 14)

# Maximum likelihood estimation

- ▶ Above,  $Y = \sum_{i=1}^N t_i$  is the total person-time in the study cohort
  - ▶ It is the time spent 'at risk' for failure in the study cohort under observation.
- ▶  $D$  is the total number of failures observed in the study cohort.
- ▶ A sufficient summary (statistics) of the data to estimate a constant hazard  $\lambda$  thus is
  - ▶ person-time  $Y = \sum_{i=1}^N t_i$
  - ▶ the total number of failures  $D = \sum_{i=1}^N d_i$
- ▶ The maximum likelihood estimate is easily found from the log likelihood  $\log L(\lambda) = D \log(\lambda) - \lambda Y$ :

$$\frac{d \log(L(\lambda))}{d\lambda} = \frac{D}{\lambda} - Y = 0 \Rightarrow \hat{\lambda} = D/Y$$

# Standard error of the hazard

- ▶ Next, the standard error is derived for the log hazard  $\beta = \log(\lambda)$

$$\beta = \log(\lambda) \Leftrightarrow \exp(\beta) = \lambda$$

- ▶ The log-likelihood of parameter  $\beta$  is

$$\log(L(\beta)) = D\beta - Y \exp(\beta)$$

- ▶ The derivatives are

$$\begin{aligned}\frac{d}{d\beta}(\log(L)) &= D - Y \exp(\beta) \\ \frac{d^2}{d\beta^2}(\log(L)) &= -Y \exp(\beta)\end{aligned}$$

- ▶ The standard error is found to be

$$S_{\beta} = \sqrt{1/(Y \exp(\hat{\beta}))} = 1/\sqrt{Y \exp(\log(D/Y))} = 1/\sqrt{D}$$

## Example: leukaemia remission

- ▶ Time of remission in the treatment group of leukaemia patients (\* = censored):

6\*, 6, 6, 6, 7, 9\*, 10\*, 10, 11\*, 13, 16, 17\*, 19\*, 20\*, 22, 23, 25\*, 32\*, 32\*, 34\*, 35\* (weeks)

- ▶ The sufficient data summary is ( $D = 9, Y = 359$ ) from which the point estimate of the rate is  $\hat{\lambda} = D/Y = 9/359 = 0.025$  (per week)
- ▶ 90% CI based on the normal approximation of the log likelihood for parameter  $\beta (= \log \lambda)$  and its standard error  $S_{\beta} = 1/\sqrt{D}$ :

$$\log(\hat{\lambda}) \pm 1.645/\sqrt{D}$$

- ▶ This leads to the following 90% confidence interval for  $\lambda$ :

$$\exp\left(\log(\lambda) \pm 1.645/\sqrt{D}\right) = 0.025 \div \exp(1.645/\sqrt{9}) = [0.015, 0.043]$$

# About parameter transformations

- ▶ There is often need to transform parameters estimates (including confidence intervals) between different one-to-one parameterisations of the model
- ▶ Above we calculated the confidence interval of the constant rate for logarithm, before transforming it back to the “absolute” scale
- ▶ N.B. Most statistical programmes tend to output logarithms of parameters
- ▶ Another example, should we be interested in the scale parameter (scale =  $1/\text{rate}$ ):
  - ▶ The hazard rate is 0.025 (per week), with a 90% confidence interval  $[0.015, 0.043]$
  - ▶ The point estimate of the scale parameter is  $1/0.025 = 40$  (weeks) and the 90% confidence interval is  $[1/0.043, 1/0.015] = [23.3, 66.7]$



# Exponential regression

- ▶ The hazard can be allowed to depend on explanatory variables (covariates)  $Z_1, \dots, Z_k$
- ▶ The hazard is then specified as follows:

$$\lambda(t; Z_1, \dots, Z_k) = \lambda(Z_1, \dots, Z_k) = \lambda_0 \exp(Z_1\beta_1 + \dots + Z_k\beta_k)$$

- ▶ The baseline hazard  $\lambda_0$  is constant over time, pertaining to those with baseline values of the explanatory variables
- ▶ For any set of explanatory variables, the hazard remains a constant but a different from the baseline
- ▶ We return to regression models in general in Part 2A of the lectures

## Poisson likelihood

- ▶ It was already mentioned that the survival likelihood is often called Poisson likelihood
- ▶ This is based on the notion that considering the observed number of failures,  $D$ , as the observation when a fixed total of  $Y$  time units of person-time (unit-time) has accrued in the study cohort, the distribution of  $D$  is proportional to the Poisson distribution with expectation  $\lambda Y$ :

$$P(D = d; \lambda) = \lambda^D \exp(-\lambda Y) \propto \frac{(\lambda Y)^D}{D!} \exp(-\lambda Y)$$

- ▶ It follows that exponential regression models can be fitted as Poisson regression because the expected counts of events in individuals with explanatory variables  $Z_1, \dots, Z_k$  is  $\lambda(Z_1, \dots, Z_k) Y(Z_1, \dots, Y_k)$ , where  $Y(Z_1, \dots, Y_k)$  is the person-time (unit-time) accrued in those individuals (see part IV)

# Hazard revisited

- ▶ Recall that, in general, the hazard function is defined as the rate of change of the conditional failure probability:

$$\lambda(t) = \lim_{h \rightarrow 0} \frac{P(T \in [t, t+h[ | T \geq t)}{h}$$

- ▶ Assuming that  $h$  is short, the conditional failure probability over the time interval  $[t, t+h[$ , given survival until  $t$ , is

$$\pi_t = P(T \in [t, t+h[ | T \geq t) \simeq \lambda(t)h$$

- ▶ In the exponential model, the hazard (rate) is constant:  
 $\lambda(t) = \lambda$  for all times  $t$

# Survival function and density function

- ▶ A survival model (i.e. distribution) can be defined terms of three alternative and equivalent ways: the hazard function  $\lambda(t)$ , the survival function  $S(t)$  or the density function  $f(t)$
- ▶ It was shown in Part I lectures that the following identities hold between the three functions:

$$S(t) = \exp\left(-\int_0^t \lambda(u) du\right)$$

$$f(t) = -\frac{dS(t)}{dt} = \lambda(t)S(t)$$

- ▶ In particular, knowing any one function, the two other functions can be calculated

# Survival models

- ▶ There are a large number of possible models for survival times
- ▶ In addition the exponential (constant and piece-wise constant hazard) models, some of the more commonly used ones include
  - ▶ Weibull distribution
  - ▶ Log-logistic distribution
  - ▶ Log-normal distribution
  - ▶ Gamma distribution
  - ▶ And more distributions: see e.g. Kalbfleisch and Prentice
- ▶ These distributions and others aim to add more flexibility in how the hazard may vary over time

# Weibull model

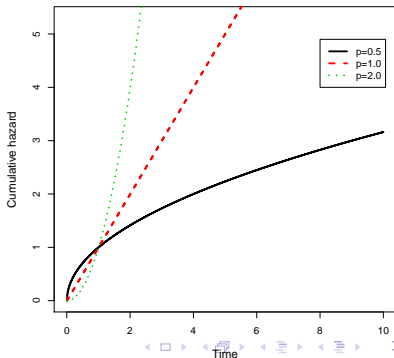
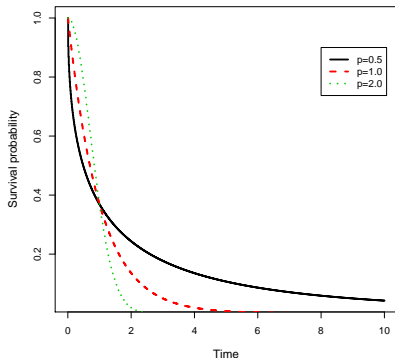
- ▶ The Weibull model is a more flexible family of survival distributions than the exponential model
- ▶ With one more parameter, it can account for hazard rates that either increase or decrease over time
  - ▶ Such models are required if, for example, the failure rate increases as the individual/unit grows older
- ▶ The model has two parameters. We use the following parameterisation for the hazard  $\lambda(t)$  in terms of parameters  $\alpha$  (rate) and  $p$  (shape):

$$\lambda(t) = p\alpha^p t^{p-1}, \quad t > 0$$

- ▶ If  $p > 1$  ( $< 1$ ), the hazard increases (decreases) over time. If  $p = 1$ , we have the standard exponential model
- ▶ Warning: there are many alternative ways to parameterise the Weibull distribution. Always check which parameterisation is being used.

# The survival function under the Weibull model

- ▶ The survival function is now  $S(t) = \exp(-(\alpha t)^p)$  and the cumulative hazard  $\Lambda(t) = (\alpha t)^p$
- ▶ The mean and the median of the Weibull distribution are  $\Gamma(1 + 1/p)/\lambda$  and  $(\ln 2)^{1/p}/\lambda$
- ▶ In the figures below,  $\alpha = 1$  and the shape parameter is given three alternative values:  $p = 0.5, 1.0, 2.0$



# Checking the Weibull assumption

- ▶ Taking the log-log transformation of the survival function under the Weibull model we obtain

$$\log(-\log(S(t))) = p \log(\alpha) + p \log(t)$$

- ▶ In other words, the log-log transformed survival function is linear over logarithmic time under the Weibull model
  - ▶ Remember also that  $-\log S(t)$  is the cumulative hazard
- ▶ This can be used to data exploration and/or model checking, based on the Kaplan-Meier estimate of the survival function



# Log-logistic model

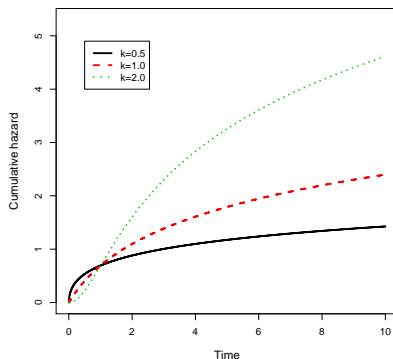
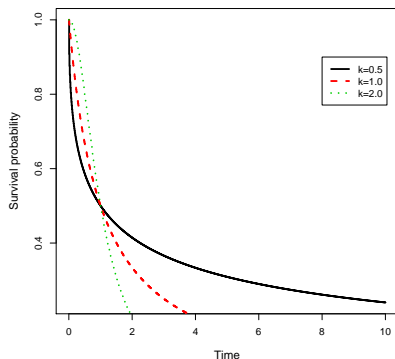
- ▶ The model has two parameters, rate  $\alpha$  and shape  $k$
- ▶ The hazard function with shape  $k$  and rate  $\alpha$  is defined as

$$f(t) = \frac{k\alpha(\alpha t)^{(k-1)}}{[1 + (\alpha t)^k]^2}$$

- ▶ The survival function is simple (next page)
- ▶ The mean is  $(k/\alpha)\pi/\sin(\pi/k)$  (if  $k > 1$ )
- ▶ The median is  $1/\alpha$

# The survival function in the log-logistic model

- ▶ The survival function is now  $S(t) = 1/[1 + (\alpha t)^k]$  and the cumulative hazard  $\Lambda(t) = \log[1 + (\alpha t)^k]$



# Survival likelihood: the general case

- ▶ In what follows we take that  $\lambda(t)$  is the parameter of the survival distribution
- ▶ Later, the explicit dependence of  $\lambda(t)$  on explanatory variables will be made explicit (see, however, also p. 13 for the exponential model)
- ▶ Survival data are presented as  $(t_i, d_i)$ ,  $i = 1, \dots, N$
- ▶ The likelihood function for  $\lambda(t)$  is again a product over individual contributions:

$$\begin{aligned} L(\lambda(t)) &= \prod_{i=1}^N L_i(\lambda(t)) = \prod_{i=1}^N \lambda(t_i)^{d_i} S(t_i) \\ &= \prod_{i=1}^N \left[ \lambda(t_i)^{d_i} \exp\left(-\int_0^{t_i} \lambda(u) du\right) \right] \\ &= \prod_{i=1}^N \left[ \lambda(t_i)^{d_i} \right] \times \exp\left(-\int_0^\infty \lambda(u) Y(u) du\right) \end{aligned}$$

where  $Y(u) = \sum_{i=1}^N \mathbf{1}(t_i \geq u)$  is the size of the risk set at time  $u$

## More on censoring

- ▶ Censoring: individuals leave the study cohort still facing the risk of the event of interest
- ▶ Above we assumed that right censoring was *uninformative*:
  - ▶ the censored individuals are statistically alike with those remaining in the study
  - ▶ in particular, the reason of leaving the study is independent of the event of interest
- ▶ If censoring is *informative*, the analysis may lead to biased estimates
  - ▶ For example, if those that remain in the study are healthier, estimates of survival may be biased upwards
  - ▶ It is sometimes very difficult to know whether censoring could be informative if the censoring mechanism is not under the control of the investigator
- ▶ N.B. When the study is predefined to end at a certain time, censoring can safely be taken to be uninformative