STAT 6390: Analysis of Survival Data

Textbook coverage: Chapter 8

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Parametric models

- Methods described previously are non-parametric; no distributional assumptions were made.
- Non-parametric (and semi-parametric) methods have the flexibility to accommodate a wide range of applications.
- If the assumption of a particular probability distribution for the data is valid, inferences based on such an assumption will be more precise.
- The validity of the parametric methods depends heavily on the appropriateness of the distributional assumption.
- Parametric models are often much easier to work with.

- Suppose actual survival times observed for n individuals are $\{t_1, \ldots, t_n\}$.
- If the probability density function of the random variable associated with theos survival times is f(t), the likelihood of the n observations is

$$\prod_{i=1}^n f(t_i).$$

• If a distributional assumption is made (e.g., $f(t) = \lambda e^{-t\lambda}$.), the unkonwon parameters (λ) can be estimated by maximzing the likelihood.

- Now suppose a the survival data includes (right) censored data.
- In this case, n pairs of observations are observed (\tilde{t}_i, Δ_i) , $i = 1, \dots, n$, where \tilde{T}_i is the observed survival time.
- When $\Delta_i = 0$, t_i is right-censored.
- The likelihood then takes the form

$$L = \prod_{i=1}^{n} [f_{\mathcal{T}}(t_i)]^{\Delta_i} \cdot [S_{\mathcal{T}}(t_i)]^{1-\Delta_i} = \prod_{i=1}^{n} [h_{\mathcal{T}}(t_i)]^{\Delta_i} \cdot S_{\mathcal{T}}(t_i). \tag{1}$$

- The last equation follows from the property h(t) = f(t)/S(t).
- Note that the derivation of of L does not require a distributional assumption.

- A more careful derivation of the likelihood function in (1) is to assume the censoring times to be random.
- Let C_i be the random variable associated with the censoring time.
- Let \tilde{T}_i be the observed survival time, $\tilde{T}_i = \min(C_i, T_i)$.
- We will consider censored and uncensored cases separately.
- For the censored observation:

$$P(\tilde{T}_i = t, \Delta_i = 0) = P(C_i = t, T_i > t).$$

For the uncensored observation:

$$P(\tilde{T}_i = t, \Delta_i = 1) = P(T_i = t, C_i > t).$$

The likelihood is then

$$L^* = \prod_{i=1}^n [P(T_i = t, C_i > t)]^{\Delta_i} \cdot [P(C_i = t, T_i > t)]^{1-\Delta_i}.$$

• Under the assumption that C_i and T_i are independent, L^* becomes

$$L^* = \prod_{i=1}^n \left[f_T(t_i)S_C(t_i)\right]^{\Delta_i} \cdot \left[f_C(t_i)S_T(t_i)\right]^{1-\Delta_i}.$$

- If the interest is in the parameter estimation in $f_T(\cdot)$, e.g., the λ in the exponential assumption, $f_C(\cdot)$ and $S_C(\cdot)$ can be considered as constant in the maximum likelihood estimation and L^* reduces to L.
- This construction shows the relevance of the assumption of independent censoring.

Exponential model

• If a random variable, T, follows an exponential distribution with rate λ , then

$$f(t) = \lambda e^{-\lambda t}$$
, $S(t) = e^{-\lambda t}$, and $h(t) = \lambda$.

• If we are willing to assumption $\{t_1, \dots, t_n\}$ are iid samples from an exponential distribution with rate λ , then the likelihood $L(\lambda)$ is

$$L(\lambda) = \prod_{i=1}^{n} \left[\lambda e^{-\lambda t_i} \right]^{\Delta_i} \cdot \left[e^{-\lambda t_i} \right]^{1-\Delta_i} = \prod_{i=1}^{n} \lambda^{\Delta_i} \cdot e^{-\lambda t_i}.$$

The log-likelihood is

$$\log L(\lambda) = \ell(\lambda) = \log(\lambda) \left(\sum_{i=1}^{n} \Delta_{i} \right) - \lambda \sum_{i=1}^{n} t_{i}.$$

Exponential model

Solving for

$$\frac{\mathrm{d} \log L(\lambda)}{\mathrm{d} \lambda} = \ell'(\lambda) = 0 \text{ gives } \hat{\lambda} = \frac{\sum_{i=1}^{n} \Delta_i}{\sum_{i=1}^{n} t_i}.$$

- The maximum likelihood estimator (MLE), $\hat{\lambda}$, is the *number of deaths* divides by the total survival time (*number of person-years*).
- The MLE for the average survival time is $1/\hat{\lambda}$, which is the total survival time divides by the number of deaths.
- With $\hat{\lambda}$, other quantities like the MLE for median survival times, can be derived.

Exponential model

- The second derivative of $\ell(\lambda)$ gives the *information*.
- In the exponential model, we have

$$\ell''(\lambda) = -\frac{1}{\lambda^2} \sum_{i=1}^n \Delta_i.$$

The standard MLE theory implies

$$\operatorname{Var}(\hat{\lambda}) \approx \frac{\hat{\lambda}^2}{\sum_{i=1}^n \Delta_i}.$$

- The 100(1 α)% confidence interval can be constructed accordingly.
- The Delta method can be applied to obtain standard errors for $g(\lambda)$, e.g., average survival time, median survival time, etc.

- The simplicity of the exponential distribution makes it attractive for some specialized applications.
- A more flexibility alternative is modeling with the Weibull distribution.
- If T follows a Weibull distribution with scale parameters λ and shape parameter γ , then

$$f(t) = \lambda \gamma t^{\gamma - 1} e^{-\lambda t^{\gamma}}, S(t) = e^{-\lambda t^{\gamma}}, \text{ and } h(t) = \lambda \gamma t^{\gamma - 1}.$$

• It is easy to see that when $\gamma =$ 1, Weibull reduces to an exponential distribution with rate λ .

• Following the similar procedure as before, the likelihood $L(\lambda, \gamma)$ is

$$\prod_{i=1}^{n} \left\{ \lambda \gamma t_i^{\gamma - 1} \right\}^{\Delta_i} e^{-\lambda t_i^{\gamma}}.$$

• Let $\ell(\lambda, \gamma) = \log L(\lambda, \gamma)$, the MLE for λ turns out to be

$$\hat{\lambda} = \frac{\sum_{i=1}^{n} \Delta_i}{\sum_{i=1}^{n} t_i^{\hat{\gamma}}},$$

but there is no close-form solution for $\hat{\gamma}$.

- The MLE $\hat{\theta} \equiv (\hat{\lambda}, \hat{\gamma})$ can be obtained directly implementing the likelihood and optimized with optim.
- Numerical method like the Newton-Raphson procedure can also be used.
- The basic idea of the Newton-Raphson procedure iterates

$$\hat{\theta}_{n+1} = \hat{\theta}_n + \left(-\frac{\mathrm{d}^2\ell(\hat{\theta}_n)}{\mathrm{d}\theta^2}\right)^{-1} \cdot \frac{\mathrm{d}\ell(\hat{\theta}_n)}{\mathrm{d}\theta}.$$

The variance-covariance matrix comes as a by-product.

- Since parametric models are sensitive to the distributional assumption, it is important to have a diagnostic tool.
- A diagnostic tool for Weibull model is derived from its survival curve.
- The log-log transformation of the Weibull survival function gives

$$\log[-\log S(t)] = \log(\lambda) - \gamma \log(t).$$

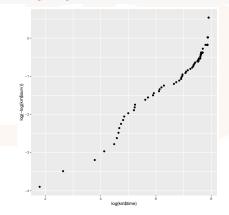
- This suggest that if $\log[-\log S(t)]$ is plotted against $\log(t)$, we would expect to see a straight line if the Weibull assumption is valid.
- S(t) can be replaced with $\widehat{S}_{KM}(t)$ or $\widehat{S}_{NA}(t)$.

• Recall that $\hat{S}_{KM}(t)$ for the whas 100 can be obtained with the survfit:

```
> km <- survfit(Surv(lenfol, fstat) ~ 1, whas100)
```

• We plot $\log[-\log S(t)]$ against $\log(t)$ via the qplot function of gqplot2:

```
> qplot(log(km$time), log(-log(km$surv)))
```



The Weibull assumption might be questionable.

- An alternative way is to select λ and γ to match the survival data at two specified time points.
- This approach is motivated by the linear relationship between log [-log S(t)] and log(t).
- Suppose we have (t_1, s_1) , and (t_2, s_2) that are two time points on a estimated survival curve (e.g., set $s_i = \widehat{S}_{KM}(t_i)$ for i = 1, 2).
- Then $\hat{\lambda}$ and $\hat{\gamma}$ can be obtained by solving the system of equation

$$\begin{cases} \log\left(-\log s_1\right) = \log(\lambda) - \gamma \log(t_1) \\ \log\left(-\log s_2\right) = \log(\lambda) - \gamma \log(t_2) \end{cases},$$

for λ and γ .



- The Weibull2 function in the **Hmisc** package can be used to produce a Weibull function that matches the two points (t_1, s_1) , and (t_2, s_2) .
- Suppose we want to find a Weibull distribution that matches the KM estimator at the 1st and the 6th year (t = 365 and t = 2190).

```
> summary(km, time = c(365, 2190))
Call: survfit(formula = Surv(lenfol, fstat) ~ 1, data = whas100)
time n.risk n.event survival std.err lower 95% CI upper 95% CI
365 80 20 0.800 0.0400 0.725 0.882
2190 15 27 0.505 0.0537 0.410 0.622
```

• The two points we what the Weibull curve to pass through are (365, 0.8) and (2190, 0.505).

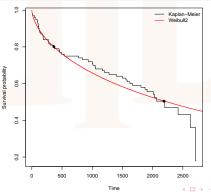
Matching the two points with

```
> weiSurv <- Weibull2(c(365, 2190), c(.8, .505))
> str(weiSurv)
function (times = NULL, alpha = 0.00560289516761242, gamma = 0.624507785991088)
> class(weiSurv)
[1] "function"
```

- Weibull2 returns a function.
- The parameters are $\lambda = 0.006$ and $\gamma = 0.625$.
- The "alpha" used in Weibull2 is equivalent to our λ .

weiSurv returns survival probability depending on inputs.

```
> weiSurv(365)
[1] 0.8
> weiSurv(2190)
[1] 0.505
> weiSurv(0:13)
[1] 1.0000000 0.9944128 0.9913993 0.9889347 0.9867714 0.9848084 0.9829920
[8] 0.9812894 0.9796788 0.9781448 0.9766759 0.9752633 0.9739001 0.9725807
```



- Recall that if T follows an exponential distribution with rate λ , T has the hazard function $h(t) = \lambda$.
- If one wants to construct a regression model under the exponential assumption, it is natural to model the exponential parameter λ .
- Suppose a covariates vector $X = (X_1, \dots, X_p)'$ is available for an individual.
- The hazard at time t for an individual can be written as

$$\lambda(t; \mathbf{x}) = \lambda \cdot r(\mathbf{X}'\beta),$$

where $\beta = (\beta_1, \dots, \beta_p)'$ is the regression coefficient, λ is a constant, and $r(\cdot)$ is a specified functional form.

- A few choices of $r(\cdot)$ have been proposed:

 - 1 r(u) = u2 $r(u) = u^{-1}$
 - $r(u) = e^{u}$
- The first two forms suffer from the disadvantage that 4ta must be restricted to guarantee $r(X'\beta) > 0$ for all possible X.
- The third form is commonly considered and will be used here.

• A few choices of $r(\cdot)$ have been proposed:

```
1 r(u) = u
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```

- The first two forms suffer from the disadvantage that 4ta must be restricted to guarantee $r(X'\beta) > 0$ for all possible X.
- The third form is commonly considered and will be used here, but we should keep in mind that there may be more appropriate forms in specific settings.

Working with model with hazard function

$$\lambda(t;x) = \lambda e^{X'\beta}. (2)$$

- This model specifies that log failure rate is a linear function of X.
- Setting $Y = \log(T)$, the model (2) implies

$$Y = \alpha - X'\beta + \epsilon, \tag{3}$$

where $\alpha = -\log(\lambda)$ and ϵ follows an extreme value distribution.

The model (3) is a log-linear model.

• The model (2) also implies

$$S(t;x) = \lambda t e^{X'\beta}$$

and the conditional density function of T given X is then

$$f(t;x) = \lambda e^{X'\beta} \cdot e^{\{-\lambda t e^{X'\beta}\}}.$$

Parameters can be solved by maximizing the likelihood.

The similar idea can be applied to Weibull assumption.

$$\lambda(t; \mathbf{x}) = \lambda \gamma t^{\gamma - 1} e^{X'\beta}. \tag{4}$$

• Setting $Y = \log(T)$, model 4 implies

$$Y = \alpha - X'\beta^* + \sigma\epsilon, \tag{5}$$

where $\alpha = -\log(\lambda)/\gamma$, $\beta^* = \beta/\gamma$, $\sigma = 1/\gamma$, and ϵ is an extreme value distribution.

- The (Weibull) relationship suggests the effect of the covariates
 - 1 act multiplicatively on the hazard function.
 - 2 act additively on Y; the general model has a log-linear models.
- The conditional density function and the survival function can be derived for likelihood estimation

 The survreg function in survival package covers a large family of parametric models.

```
> library(survival)
> args(survreg)
function (formula, data, weights, subset, na.action, dist = "weibull",
    init = NULL, scale = 0, control, parms = NULL, model = FALSE,
    x = FALSE, y = TRUE, robust = FALSE, score = FALSE, ...)
NULL
```

- Suppose we want to fit a parametric model using covariates:
 - gender
 - age
 - gender-age interaction
 - body mass index (BMI)
- We can create a Surv formula as

```
> fm <- Surv(lenfol, fstat) ~ (age + gender)^2 + bmi
```

Exponential regression model:

```
> fit.exp <- survreg(fm, data = whas100, dist = "exp")
> summary(fit.exp)
Call:
survreg(formula = fm, data = whas100, dist = "exp")
            Value Std. Error z
(Intercept) 9.2897 1.6200 5.73 9.8e-09
   -0.0532 0.0157 -3.39 0.0007
age
gender -3.9324 1.8098 -2.17 0.0298
bmi 0.0935 0.0376 2.49 0.0128
age:gender 0.0498 0.0241 2.06 0.0394
Scale fixed at 1
Exponential distribution
Loglik(model) = -444.4 Loglik(intercept only) = -458.5
Chisa= 28.25 on 4 degrees of freedom, p= 1.1e-05
Number of Newton-Raphson Iterations: 5
n = 100
```

- This is equivalent to the Weibull regression model when λ (scale) = 1.
- The same result is presented in Table 8.2.

- Since Weibull relationship suggests two kinds of covariates effects (see page 23), the regression coefficient can be interpret in two ways.
- For one unit increase in bmi ($\hat{\beta}_{bmi} = 0.0935$):
 - the risk of death is expected to increase by $e^{0.0935} = 1.098$ times.
 - the log of survival time is expected to decrease by 0.0935 (days).
- For one unit increase in age among females (gender = 1):
 - the risk of death is expected to increase by $e^{-0.0532+0.0498} = 0.997$ times.
 - the log of survival time is expected to decrease by -0.0532 + 0.0498 = -0.0034.
 - The Wald p-value of $\hat{\beta}_{age} + \hat{\beta}_{gender=1}$ can be computed as following

- Since the exponential distribution is a special case of the Weibull distribution, fitting a exponential regression model is equivalent to fitting a Weibull regression model with $\gamma=1$.
- The intercept defines the exponential parameter, λ.
- The likelihood, Loglik (model), is available because we are fitting a
 parametric model.
- Chisq is likelihood ratio statistics:

```
> 2 * log(exp(-444.4) / exp(-458.5))
[1] 28.2
```

The likelihood ratio test gives a p-value of

```
> 1 - pchisq(28.25, 4)
[1] 1.109905e-05
```

Weibull regression model:

```
> summary(survreg(fm, data = whas100))
Call:
survreg(formula = fm, data = whas100)
            Value Std. Error z
(Intercept) 9.8727 2.0470 4.82 1.4e-06
          -0.0639 0.0206 -3.10 0.0019
age
gender -4.6895 2.2848 -2.05 0.0401
hmi
     0.1055 0.0465 2.27 0.0232
age:gender 0.0592 0.0304 1.94 0.0518
Log(scale) 0.2254 0.1242 1.81 0.0695
Scale = 1.25
Weibull distribution
Loglik(model) = -442.6 Loglik(intercept only) = -455.3
Chisq= 25.36 on 4 degrees of freedom, p= 4.3e-05
Number of Newton-Raphson Iterations: 5
n = 100
```

• The same result is presented in Table 8.5.

- The interpretation of the regression parameters is similar to that in exponential regression model.
- In addition to the intercept and parameter estimates, surverg also gives Log(scale), which corresponds to $log(\gamma)$.
- The *p*-value for Log(scale) is at the borderline of $\alpha = 0.05$, suggesting that adding an extra parameter (γ) to the model does not improve the overall fit significantly.

Log-normal regression model

Suppose we have a common form

$$Y = X'\beta + \epsilon$$
,

different parametric model can be specified through the distribution of ϵ .

• Another common parametric model is when ϵ follows a standard normal, this also implies the survival times follow a log-normal distribution.

Log-normal regression model

 In the log-normal case, the parameters are not in the form of a proportional hazards model.

```
> summary(survreg(fm, data = whas100, dist = "lognormal"))
Call:
survreg(formula = fm, data = whas100, dist = "lognormal")
              Value Std. Error
(Intercept) 10.3193 2.2278 4.63 3.6e-06
age
           -0.0737 0.0233 -3.16 0.0016
gender -4.9028 2.5880 -1.89 0.0582
bmi 0.0969 0.0500 1.94 0.0525
age:gender 0.0626 0.0354 1.77 0.0774
Log(scale) 0.6871 0.1066 6.44 1.2e-10
Scale= 1.99
Log Normal distribution
Loglik (model) = -446.5 Loglik (intercept only) = -457.1
Chisq= 21.22 on 4 degrees of freedom, p= 0.00029
Number of Newton-Raphson Iterations: 4
n = 100
```

Other parametric models

 Here is a list of distributions ε can be specified via survreg.distributions.

```
> names(survreg.distributions)
[1] "extreme" "logistic" "gaussian" "weibull" "exponential"
[6] "rayleigh" "loggaussian" "lognormal" "loglogistic" "t"
```