

Lecture fifteen: Parametric PH Models: The Weibull Model(II)

1. The Weibull proportional hazards model

(a) The proportional hazards property

i. One dummy variable (two groups)

$$h_i(t) = e^{\beta x_i} h_0(t),$$

where x_i is dummy variable, and the baseline hazard is from Weibull distribution (known instead of arbitrary in Cox model).

$$h_0(t) = \lambda \gamma t^{\gamma-1}. \quad (1)$$

Let $\psi = \exp(\beta)$, then $\psi h_0(t)$ is hazard function for a Weibull distribution with scale parameter $\psi \lambda$ and the same shape parameter γ (*the proportional hazards property*).

ii. More than one covariates

$$h_i(t) = \exp(\beta_1 x_{i1} + \beta_2 x_{i2} + \dots + \beta_p x_{ip}) h_0(t),$$

where $h_0(t)$ is from (1). The corresponding survival function and density function are still from Weibull distribution with scale parameter $\lambda \exp(\beta' x_i)$ and shape parameter γ . Namely

$$S_i(t) = \exp\{-\lambda \exp(\beta' x_i) t^\gamma\}. \quad (2)$$

$$f_i(t) = \lambda \gamma \exp(\beta' x_i) t^{\gamma-1} \exp\{-\lambda \exp(\beta' x_i) t^\gamma\}.$$

iii. Maximum Likelihood Estimation (**Not MPLE as in Cox model**)

iv. Formulas for median and p th percentile of survival time.

See the ones in lecture one of this chapter.

(b) An alternative form of the Weibull PH model

Consider following transformation of r.v. T_i of survival time:

$$\log T_i = \mu + \alpha_1 x_{i1} + \alpha_2 x_{i2} + \dots + \alpha_p x_{ip} + \sigma \epsilon_i.$$

If ϵ has the *Gumbel* distribution, also known as *extreme value* distribution: see Kalbfleisch and Prentice's book (at page 33) for details, with pdf:

$$f(\epsilon) = \exp(\epsilon - e^\epsilon),$$

then, $\xi = e^\epsilon$ has exponential distribution $\text{Exp}(1)$.

Follow steps at page 237- 238, then we have

$$S_i(t) = \exp[-\exp\{\frac{\log t - \mu - \alpha'x_i}{\sigma}\}]. \quad (3)$$

In model (3), μ is *intercept* and σ is *scale parameter*.

Let (2) and (3) equal, then we have

$$\lambda = \exp(-\mu/\sigma), \quad \gamma = \sigma^{-1}, \quad \beta_j = -\alpha_j/\sigma,$$

for $j = 1, 2, \dots, p$ (page 203, and pages 237-238).

In SAS, the output from **PROC LIFEREG** is expressed in terms of $\mu, \sigma, \alpha_1, \alpha_2, \dots, \alpha_p$ in equation (3) (still use maximum likelihood estimation, but different parameterization).

2. Example 5.3: IUD data (one sample, intercept only)

For exponential distribution, the score equation has closed form solution, but for Weibull distribution, the score equations are non-linear in γ , thus, iterative procedures, such as the Newton-Raphson algorithm, are required.

```
/* SAS program */
options ls=80;
libname fu '..../sdata';
data w;
    set fu.iud;
proc lifereg;
    model survt*censor(0) = / covb dist=weibull;
run;
```

```
/* SAS output */
```

Analysis of Parameter Estimates

Parameter	DF	Estimate	Standard Error	95% Confidence Limits		Chi-Square	Pr>ChiSq
Intercept	1	4.5915	0.2044	4.1910	4.9921	504.74	<.0001
Scale	1	0.5965	0.1638	0.3482	1.0218		
Weibull Scale	1	98.6440	20.1601	66.0859	147.2425		
Weibull Shape	1	1.6764	0.4604	0.9786	2.8717		

3. The log-cumulative plot: Example 5.4: Breast cancer study

Is the Weibull PH model appropriate for the data?

(a) Log-cumulative hazard plot

(b) SAS program

```
options ls=80;
libname fu '.././sdata';
data w;
    set fu.hpa;
filename gsf 'ex54.gsf';
goptions reset=all gunit=pct border ftext=swissb htitle=6 htext=2.5
gaccess=gsf ROTATE=LANDSCAPE gsfmode=replace device=ps;
title1 'Log-cumulative hazard plot for HPA data';
footnote1 c=black f=special h=4 'L L'
        f=swiss h=4 ' Negative'
        c=blue f=special h=3 ' m m '
        f=swissb h=4 ' Positive';
symbol1 c=black i=join v=triangle height=3;
symbol2 c=blue i=join v=star height=3;

proc lifetest notable plot = (lls) method=pl;
    time survt*censor(0);
    strata group;
run;
```

4. Example 5.5: Breast cancer study

Fitting exponential model; calculate the risk ratio and its variance. We use output from sas instead of formulas at page 183.

```
/* SAS program */
options ls=80;
libname fu '.././sdata';
data w;
    set fu.hpa;
proc lifereg;
    model survt*censor(0) = group / covb dist = exponential;
run;
/* SAS output */
    Analysis of Parameter Estimates
```

Parameter	DF	Estimate	Standard Error	95% Confidence Limits		Chi-Square	Pr>ChiSq
Intercept	1	5.8003	0.4472	4.9238	6.6768	168.22	<.0001
GROUP	1	-0.9516	0.4976	-1.9269	0.0237	3.66	0.0558
Scale	0	1.0000	0.0000	1.0000	1.0000		
Weibull Shape	0	1.0000	0.0000	1.0000	1.0000		

5. Example 5.6: Breast cancer study

Fitting Weibull model; calculate the risk ratio and its variance using output from sas program.

```
/* SAS program */
options ls=80;
libname fu '.././sdata';
data w;
    set fu.hpa;
proc lifereg;
    model survt*censor(0) = group / covb dist = weibull;
run;
```

```
/* SAS output */
```

Analysis of Parameter Estimates

Parameter	DF	Estimate	Standard Error	95% Confidence Limits		Chi-Square	Pr>ChiSq
Intercept	1	5.8544	0.4989	4.8766	6.8321	137.71	<.0001
GROUP	1	-0.9967	0.5441	-2.0631	0.0697	3.36	0.0670
Scale	1	1.0668	0.1786	0.7684	1.4810		
Weibull Shape	1	0.9374	0.1569	0.6752	1.3014		

Estimated Covariance Matrix

	Intercept	GROUP	Scale
Intercept	0.248878	-0.245010	0.026044
GROUP	-0.245010	0.296036	-0.021310
Scale	0.026044	-0.021310	0.031880

What's the variance of $\hat{t}(50)$? Recall

$$\hat{t}(p) = \left[\frac{1}{\hat{\lambda}} \log\left(\frac{100}{100-p}\right) \right]^{1/\hat{\gamma}},$$

6. Example 5.7: Breast cancer study: CI for hazard ratio

- (a) Delta method: A function of two estimates (two random variables)

The general formula for $g(\hat{\theta}_1, \hat{\theta}_2)$ (see the eq. (5.24) at p. 188)

- (b) Illustration: CI for $\hat{\beta}$, i.e.

$$g(\hat{\alpha}, \hat{\sigma}) = -\frac{\hat{\alpha}}{\hat{\sigma}}.$$

Log – cumulative hazard plot for HPA data

