# Lecture Three: Standard Error of KM Estimate and Estimating Hazard Function

# **1. Standard Error and Confidence interval for** $\hat{S}(t)$

We also need to know about how good it's the (KM) estimate. A common way is to estimate the sample variation or standard error of the estimate  $\hat{S}(t)$ .

### Use the derivation at page 22-23: Steps:

- Take log transformation of KM estimate
- # of survivals,  $n_j d_j$ , through the interval beginning at  $t_{(j)}$  has Binomial $(n_j, \hat{p}_j)$ , where  $\hat{p}_j = 1 d_j/n_j$
- Obtain the variance of  $\log \hat{p}_i$  by the delta-method:

$$\operatorname{var}\{g(X)\} \approx \left\{\frac{dg(X)}{dX}\right\}^2 \operatorname{var}(X),$$

which is known as the *Taylor series approximation* to the variance of a function of a random variable.

• Standard error (S.E.): square-root of variance estimate.

With the estimated standard error, a  $(1 - \alpha)100\%$  confidence interval for S(t) at each time point t can be easily constructed, based on a typical normal approximation (meaning?). When we link the upper and lower confidence limits together along the time axis, we form a so-called confidence band. This can be done on different scales as implemented in Splus and SAS (PROC LIFETEST: conftype, confband options in SURVIVAL statement).

#### • Original scale: S (t).

- Confidence interval for  $\hat{S}(t_j)$  at  $t_j$   $CI = \hat{S}(t_j) \pm z_{\alpha/2} * S.E. (\hat{S}(t_j))$
- Although S (t) should be in [0, 1], the lower and upper limit can be out of the range. A practical solution to this problem is to replace any limit that greater than 1 by 1, and any limit that is less than zero by 0.0.

# • Log-scale: log S (t).

• Confidence interval for  $\log \hat{S}(t_j)$   $\operatorname{CI}_{\log} = \log \hat{S}(t_j) \pm z_{\alpha/2} * \operatorname{S.E.} (\log \hat{S}(t_j))$ 

• Converting  $CI_{log}$  back to the original scale to have CI for  $\hat{S}(t_j)$   $CI = \exp(CI_{log}) = ?$ 

- Where the lower bound is always nonnegative, the upper bound may exceed 1
- Log-log scale:  $\log (-\log \hat{S}(t_i))$ .
  - Obtain the standard error for  $\log(-\log \hat{S}(t_i))$  by the delta-method
  - Confidence interval  $CI_{log-log}$  for  $log(-log \hat{S}(t_j))$  by the normal approximation
  - Convert  $CI_{log-log}$  to have CI for  $\hat{S}(t_i)$

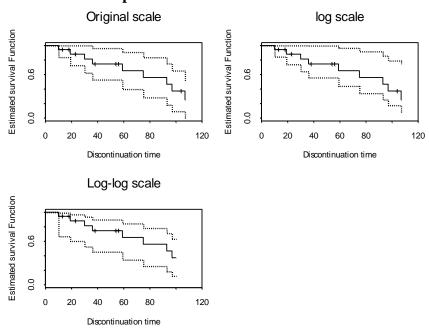
$$CI = exp (-exp (CI_{log-log}))$$

- Lower limit >= 0 and upper limit <= 1
- Appropriate with moderate to large sample size because of repeated use of the delta-method.

The Greenwood variance estimate is appropriate only when the expected risk set size  $n_j$  is fairly large at each time point  $t_{(j)}$  because the use of the delta-method requires large sample size. As  $n_j$  gets smaller with increasing time, the Greenwood estimate becomes unstable at the tail. (Cut the tail out requested by investigators, reasonable?)

- In Splus, use option "conf.type" in "survfit()" to choose different methods
- In SAS, use *conftype* option in the PROC LIFETEST statement.

### • Example: IUD



#### **Splus code:**

```
iud.s<-function (){</pre>
tmpdf <- importData("../sdata/iud.sas7bdat")</pre>
par(mfrow=c(2,2))
iud.km1 <- survfit(Surv(survt, censor), conf.type="plain",
        type=''kaplan-meier'', data=tmpdf)
plot(iud.km1,xlab="Discontinuation time",
      ylab="Estimated survival Function", xlim=c(0, 120),
         ylim=c(0,1), mark.time=T, conf.int=T,
           main="Original scale")
iud.km2 <- survfit(Surv(survt, censor), conf.type="log",
          type="kaplan-meier", data=tmpdf)
plot(iud.km2,xlab="Discontinuation time",
          ylab="Estimated survival Function", xlim=c(0, 120),
       vlim=c(0,1),mark.time=T, conf.int=T, main="log scale")
iud.km3 <- survfit(Surv(survt, censor), conf.type="log-log",
      type="kaplan-meier", data=tmpdf)
plot(iud.km3,xlab="Discontinuation time",
    ylab="Estimated survival Function", xlim=c(0, 120),
    vlim=c(0,1), mark.time=T, xmax=100, conf.int=T,
     main="log-log scale")
```

### 2. Estimating the hazard function

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#### Life-table estimate of the hazard function

- O Dividing the period of observation into a series of time intervals:  $t_j$  to  $t_{j+1}$ , j = 1, 2, ..., m, with length  $\tau_j$
- o  $d_j$  deaths,  $c_j$  censored in  $(t_j, t_{j+1}]$  and  $n_j$  at risk at the start of the j'th interval
- O Assume censored times occur uniformly (i.e. U(0,  $c_j$ )) through the j'th interval, then average number of individual at risk is  $n'_j = n_j c_j/2$
- Assuming the death rate is constant during the j'th interval
- o The average hazard of death per unit time can be estimated by

$$h^*(t) = \frac{d_j}{(n_j - d_j/2)\tau_j},$$

for  $t_{j} \le t < t_{j+1}, j = 1, 2, ..., m$ , where  $(n_{j} - d_{j}/2)\tau_{j}$  is the average time survived in  $(t_{j}, t_{j+1}]$ .

# • Kaplan-Meier Type Estimate

Let the observed survival times:  $t_1, t_2, \ldots, t_n$  and r ordered death times:  $t_{(1)} < t_{(2)} < \ldots < t_{(r)}$ ;  $n_j$  at risk just before  $t_{(j)}$ ,  $d_j$  deaths at the j'th death time

- o Assuming constant hazard between successive death times
- o The hazard can be estimated by

$$\hat{h}(t) = \frac{d_j}{n_j \tau_j},$$
 for  $t_{(j)} \le t < t_{(j+1)}$ , where  $\tau_j = t_{(j+1)} - t_{(j)}$ 

- No estimate for  $t > t_{(r)}$
- 0 Proof: The conditional death probability for  $t_j \leq T < t_{(j+1)}$  is  $\hat{h}(t)\tau_j$ , which is  ${\rm d_j/n_j}$

#### Kernel-smoothed estimate

- Above estimates are rather irregular
- Using smoothing techniques (ref: Smoothing Methods in Statistics, 1996, Simonoff JS).
- O A weighted average of values of the estimated hazard  $\hat{h}(t)$  at death times in the neighborhood of t.

## Estimating the cumulative hazard function

- O Use relation  $H(t) = -\log S(t)$  and KM estimate of survivor function  $\hat{S}(t) = \prod_{j=1}^{k} \frac{n_j d_j}{n_j}$ , for
- $\circ$  Use Taylor series expansion of log(1 x), and ignore higher-order terms when x is small
- $\circ \quad \hat{H}(t) \approx \sum_{j=1}^{k} \frac{d_j}{n_j}, \text{ which is called Nelson-Aalen estimate.}$

#### 3. Estimating the median, mean and percentiles of survival times

- **Median survival time**: defined as smallest observed survival time for which the value of the estimated survival function is less than 0.5
- In math term

$$\hat{t}(50) = \min\{t_i \mid \hat{S}(t_i) \le 0.5\}$$

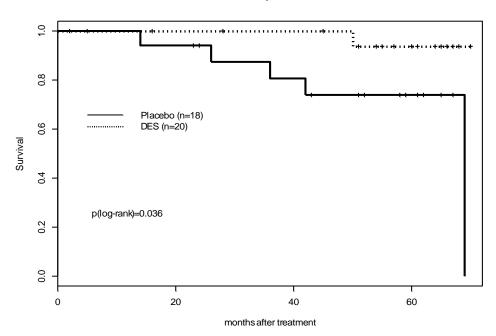
where  $t_i$  is the observed survival time for the *i*'th individual, I = 1, ..., n

- What if  $\hat{S}(t) > 0.5$  for any t > 0?
- Mean:  $E(T) = \int_{0}^{\infty} tf(t)dt = -\int_{0}^{\infty} tdS(t) = -tS(t)|_{0}^{\infty} + \int_{0}^{\infty} S(t)dt = \int_{0}^{\infty} S(t)dt$
- **p'th percentile:** Defined to be the value t(p), such that  $F\{t(p)\} = p/100$ . In terms of survival, t(p) is such that  $S\{t(p)\} = 1 (p/100)$
- The p'th percentile of the estimated survival:

$$\hat{t}(p) = \min\{t_i \mid \hat{S}(t_i) \le 1 - (p/100)\}$$

• Example: Medians of two treatment groups of prostatic cancer patients (Table 1.4, p10). Use the plot from lecture one

#### Survival by treatment



• Confidence intervals for the median and percentiles by the deltamethod.