

Survival Analysis: Kaplan-Meier Method

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What we will learn ...

- **Cox's proportional hazards model**
- **Computing the hazard ratio**
- **Adjusted survival curves using the Cox PH model**
- **The meaning of the PH assumption**

Leukemia remission data

Leukemia Remission Data

| Group 1($n = 21$) | | Group 2($n = 21$) | |
|---------------------|---------|---------------------|---------|
| t (weeks) | log WBC | t (weeks) | log WBC |
| 6 | 2.31 | 1 | 2.80 |
| 6 | 4.06 | 1 | 5.00 |
| 6 | 3.28 | 2 | 4.91 |
| 7 | 4.43 | 2 | 4.48 |
| 10 | 2.96 | 3 | 4.01 |
| 13 | 2.88 | 4 | 4.36 |
| 16 | 3.60 | 4 | 2.42 |
| 22 | 2.32 | 5 | 3.49 |
| 23 | 2.57 | 5 | 3.97 |
| 6+ | 3.20 | 8 | 3.52 |
| 9+ | 2.80 | 8 | 3.05 |
| 10+ | 2.70 | 8 | 2.32 |
| 11+ | 2.60 | 8 | 3.26 |
| 17+ | 2.16 | 11 | 3.49 |
| 19+ | 2.05 | 11 | 2.12 |
| 20+ | 2.01 | 12 | 1.50 |
| 25+ | 1.78 | 12 | 3.06 |
| 32+ | 2.20 | 15 | 2.30 |
| 32+ | 2.53 | 17 | 2.95 |
| 34+ | 1.47 | 22 | 2.73 |
| 35+ | 1.45 | 23 | 1.97 |

+ denotes censored observation

```
group = c(1,1,1,1,1, 1,1,1,1,1, 1,1,1,1,1,
1,1,1,1,1,1, 0,0,0,0,0, 0,0,0,0,0,
0,0,0,0,0, 0,0,0,0,0,0)
```

```
time =
c(6,6,6,7,10,13,16,22,23,6,9,10,11,17,19,2
0,25,32,32,34,35,1,1,2,2,3,4,4,5,5,8,8,8,8
,11,11,12,12,15,17,22,23)
```

```
status =
c(0,0,0,0,0,0,0,0,0,1,1,1,1,1,1,1,1,1,1,1,
1, 0,0,0,0,0, 0,0,0,0,0, 0,0,0,0,0,
0,0,0,0,0,0)
```

```
wbc =
c(2.31,4.06,3.28,4.43,2.96,2.88,3.60,2.32,
2.57,3.20,2.80,2.70,2.60,2.16,2.05,2.01,1.
78,2.20,2.53,1.47,1.45,
2.80,5.00,4.91,4.48,4.01,4.36,2.42,3.49,3.
97,3.52,3.05,2.32,3.26,3.49,2.12,1.50,3.06
,2.30,2.95,2.73,1.97)
```

```
dat=data.frame(group,time,status,wbc)
```

Kaplan-Meier curves

```
km = survfit(Surv(time, status==0) ~ group); km
plot(km, lty=c(1,4), lwd=2, xlab="Weeks", ylab="S(t)")
legend("topright", c("Placebo", "Treatment"),
lty=c(1,4), lwd=2)
```

Leukemia remission data

- **Outcome: time (+ status)**
- **Covariates / predictors: group, wbc**

Log-rank test and Cox's model

- **Log-rank test:**
 - No covariates

- **Cox's proportional hazards model**
 - Adjusted for covariates

Cox's proportional hazards model

$$h(t, \mathbf{X}) = h_0(t) e^{\sum_{i=1}^p \beta_i X_i}$$

- $\mathbf{X} = (X_1, X_2, \dots, X_p)$, explanatory/predictor variables
- $h_0(t)$: **baseline hazard**, involves t but not X's
- $\exp(\beta_i X_i)$: exponential, involves X, but not t (Xs can be time-dependent)

Cox's proportional hazards model

- **Hazard(t) = (baseline risk) x (effects of covariates)**
- **Semi-parametric model**

Proportional hazards model

- Let $X = (x_1, \dots, x_p)$ - the set of predictors
- $h(t|x)$: hazard of someone with predictors x
- $h(t|x) = h_0(t) \exp(\beta_1 x_1 + \dots + \beta_p x_p)$
- $h(t|x)/h_0(t) = \exp(\beta_1 x_1 + \dots + \beta_p x_p)$
- $\log(h(t|x)) = \log(h_0(t)) + \beta_1 x_1 + \dots + \beta_p x_p$ because $\log(a/b) = \log(a) - \log(b)$
- Much like logistic regression but change *odds* to *hazards*

Cox's model

- The “baseline” hazard $h_0(t)$ is unspecified
plays the role of intercept
- Predictor effects in terms of hazard ratios
relative rates of failure
- Don't need to know $h_0(t)$
to understand these predictor effects
- Effect of one unit increase in predictor x_p is to multiply hazard by $\exp(\beta_p)$
holding all other predictors constant

+1 unit change in x_p

$$h(t|x) = h_0(t) \exp(\beta_1 x_1 + \dots + \beta_p V) \quad x_p = V$$

versus

$$h(t|x) = h_0(t) \exp(\beta_1 x_1 + \dots + \beta_p (V+1)) \quad x_p = V+1$$

$$\text{Ratio} = \frac{h_0(t) \exp(\beta_1 x_1 + \dots + \beta_p (V+1))}{h_0(t) \exp(\beta_1 x_1 + \dots + \beta_p V)}$$

+1 unit change in x_p

$$\begin{aligned} \text{Ratio} &= \frac{\cancel{h_0(t)} \exp(\beta_1 x_1 + \dots + \beta_p (V+1))}{\cancel{h_0(t)} \exp(\beta_1 x_1 + \dots + \beta_p V)} \\ &= \frac{\exp(\beta_1 x_1 + \dots + \beta_p (V+1))}{\exp(\beta_1 x_1 + \dots + \beta_p V)} \end{aligned}$$

Ratio does not depend on t !

+1 unit change in x_p

$$\begin{aligned} \text{Ratio} &= \frac{\exp(\beta_1 x_1 + \dots + \beta_p (V+1))}{\exp(\beta_1 x_1 + \dots + \beta_p V)} \\ &= \exp(\beta_1 x_1 + \dots + \beta_p (V+1) - (\beta_1 x_1 + \dots + \beta_p V)) \\ &\quad \text{because } \exp(a) / \exp(b) = \exp(a-b) \\ &= \exp(\beta_p (V+1) - \beta_p V) \quad \text{b/c same other predictors} \\ &= \exp(\beta_p) \quad \text{b/c } \beta_p V \text{ terms cancel} \end{aligned}$$

Hazard ratio

- β is the regression coefficient
If no effect of a predictor variable then $\beta=0$
- HR for a unit increase in a predictor is $\exp(\beta)$
If no effect of variable then $\exp(\beta)=1$
- A useful way to discuss predictor effects
(increases or decreases Hazard by a factor)

Why Cox model?

- Can be fitted without an explicit model for the hazard
- Can model the effect of a continuous predictor
- Can model multiple predictors: *continuous, binary, categorical*
- Can adjust for confounders: *adjust by adding confounders to the model*
- Can incorporate interaction, mediation: *create and add product terms*
- Can detect and estimate predictors for patient-level prognosis

Comparison with other forms of regression

- *Same issues as in linear and logistic regression:*
predictor selection
- *Differences:*
interpretation, assumptions, model checking

Baseline data

```
group = c(1,1,1,1,1, 1,1,1,1,1, 1,1,1,1,1, 1,1,1,1,1,1,  
0,0,0,0,0, 0,0,0,0,0, 0,0,0,0,0, 0,0,0,0,0,0)
```

```
time =  
c(6,6,6,7,10,13,16,22,23,6,9,10,11,17,19,20,25,32,32,34,35,1,1,  
,2,2,3,4,4,5,5,8,8,8,8,11,11,12,12,15,17,22,23)
```

```
status = c(0,0,0,0,0,0,0,0,0,1,1,1,1,1,1,1,1,1,1,1,1,  
0,0,0,0,0, 0,0,0,0,0, 0,0,0,0,0, 0,0,0,0,0,0)
```

```
wbc =  
c(2.31,4.06,3.28,4.43,2.96,2.88,3.60,2.32,2.57,3.20,2.80,2.70,  
2.60,2.16,2.05,2.01,1.78,2.20,2.53,1.47,1.45,  
2.80,5.00,4.91,4.48,4.01,4.36,2.42,3.49,3.97,3.52,3.05,2.32,3.  
26,3.49,2.12,1.50,3.06,2.30,2.95,2.73,1.97)
```

```
dat=data.frame(group,time,status,wbc)
```

```
baseline = Surv(time, status==0)
```

```
km = survfit(baseline ~ 1)
```

```
summary(km)
```

Baseline data

| time | n.risk | n.event | survival | std.err | lower 95% CI | upper 95% CI |
|------|--------|---------|----------|---------|--------------|--------------|
| 1 | 42 | 2 | 0.952 | 0.0329 | 0.8901 | 1.000 |
| 2 | 40 | 2 | 0.905 | 0.0453 | 0.8202 | 0.998 |
| 3 | 38 | 1 | 0.881 | 0.0500 | 0.7883 | 0.985 |
| 4 | 37 | 2 | 0.833 | 0.0575 | 0.7279 | 0.954 |
| 5 | 35 | 2 | 0.786 | 0.0633 | 0.6709 | 0.920 |
| 6 | 33 | 3 | 0.714 | 0.0697 | 0.5899 | 0.865 |
| 7 | 29 | 1 | 0.690 | 0.0715 | 0.5628 | 0.845 |
| 8 | 28 | 4 | 0.591 | 0.0764 | 0.4588 | 0.762 |
| 10 | 23 | 1 | 0.565 | 0.0773 | 0.4325 | 0.739 |
| 11 | 21 | 2 | 0.512 | 0.0788 | 0.3783 | 0.692 |
| 12 | 18 | 2 | 0.455 | 0.0796 | 0.3227 | 0.641 |
| 13 | 16 | 1 | 0.426 | 0.0795 | 0.2958 | 0.615 |
| 15 | 15 | 1 | 0.398 | 0.0791 | 0.2694 | 0.588 |
| 16 | 14 | 1 | 0.369 | 0.0784 | 0.2437 | 0.560 |
| 17 | 13 | 1 | 0.341 | 0.0774 | 0.2186 | 0.532 |
| 22 | 9 | 2 | 0.265 | 0.0765 | 0.1507 | 0.467 |
| 23 | 7 | 2 | 0.189 | 0.0710 | 0.0909 | 0.395 |

Cox's model using R

```
dat=data.frame(group,time,status,wbc)
```

```
baseline = Surv(time, status==0)
```

```
km = survfit(baseline)
```

```
cox = coxph(Surv(time, status==0) ~ group + wbc)
```

```
summary(cox)
```

Cox's model using R

n= 42, number of events= 30

| | coef | exp(coef) | se(coef) | z | Pr(> z) | |
|-------|---------|-----------|----------|--------|----------|-----|
| group | -1.3861 | 0.2501 | 0.4248 | -3.263 | 0.0011 | ** |
| wbc | 1.6909 | 5.4243 | 0.3359 | 5.034 | 4.8e-07 | *** |

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

| | exp(coef) | exp(-coef) | lower .95 | upper .95 |
|-------|-----------|------------|-----------|-----------|
| group | 0.2501 | 3.9991 | 0.1088 | 0.5749 |
| wbc | 5.4243 | 0.1844 | 2.8082 | 10.4776 |

Concordance= 0.852 (se = 0.062)

Rsquare= 0.671 (max possible= 0.988)

Likelihood ratio test= 46.71 on 2 df, p=7.187e-11

Wald test = 33.6 on 2 df, p=5.061e-08

Score (logrank) test = 46.07 on 2 df, p=9.921e-11

Interpretation of output

n= 42, number of events= 30

| | coef | exp(coef) | se(coef) | z | Pr(> z) | |
|-------|---------|-----------|----------|--------|----------|-----|
| group | -1.3861 | 0.2501 | 0.4248 | -3.263 | 0.0011 | ** |
| wbc | 1.6909 | 5.4243 | 0.3359 | 5.034 | 4.8e-07 | *** |

- **Model**

$$h(t) = L_t \times e^{b1 \times \text{group} + b2 \times \text{wbc}}$$

- **Risk(t) = $L_t \times e^{-1.3861 \times \text{group} + 1.6909 \times \text{wbc}}$**

group: 0 = control, 1 = treatment

Interpretation of output

| | exp(coef) | exp(-coef) | lower .95 | upper .95 |
|-------|-----------|------------|-----------|-----------|
| group | 0.2501 | 3.9991 | 0.1088 | 0.5749 |
| wbc | 5.4243 | 0.1844 | 2.8082 | 10.4776 |

- Risk of remission in the treatment group is 75% lower than that in the controls (hazard ratio: 0.25; 95% CI 0.11 to 0.57)
- Risk of remission increased by 5.42 folds (95% CI 2.81 to 10.48) for each unit increase in wbc

Interpretation of output

```
Concordance= 0.852 (se = 0.062 )  
Rsquare= 0.671 (max possible= 0.988 )  
Likelihood ratio test= 46.71 on 2 df, p=7.187e-11  
Wald test = 33.6 on 2 df, p=5.061e-08  
Score (logrank) test = 46.07 on 2 df, p=9.921e-11
```

- Predicted risk and observed risk agree 85.2% of the times
- Group and wbc “explained” 67.1% variance in the risk of remission

What about “baseline risk” ?

- $\text{Risk}(t) = L_t \times e^{-1.3861 \times \text{group} + 1.6909 \times \text{wbc}}$

group: 0 = control, 1 = treatment

- What is L_t ?

- L_t = baseline risk

- Can be estimated from

```
baseline = Surv(time, status==0)
```

```
km = survfit(baseline ~ 1)
```

```
summary(km)
```


Baseline probability of remission

| time | n.risk | n.event | survival | std.err | lower 95% CI | upper 95% CI |
|------|--------|---------|----------|---------|--------------|--------------|
| 1 | 42 | 2 | 0.952 | 0.0329 | 0.8901 | 1.000 |
| 2 | 40 | 2 | 0.905 | 0.0453 | 0.8202 | 0.998 |
| 3 | 38 | 1 | 0.881 | 0.0500 | 0.7883 | 0.985 |
| 4 | 37 | 2 | 0.833 | 0.0575 | 0.7279 | 0.954 |
| 5 | 35 | 2 | 0.786 | 0.0633 | 0.6709 | 0.920 |
| 6 | 33 | 3 | 0.714 | 0.0697 | 0.5899 | 0.865 |
| 7 | 29 | 1 | 0.690 | 0.0715 | 0.5628 | 0.845 |
| 8 | 28 | 4 | 0.591 | 0.0764 | 0.4588 | 0.762 |
| 10 | 23 | 1 | 0.565 | 0.0773 | 0.4325 | 0.739 |
| 11 | 21 | 2 | 0.512 | 0.0788 | 0.3783 | 0.692 |
| 12 | 18 | 2 | 0.455 | 0.0796 | 0.3227 | 0.641 |
| 13 | 16 | 1 | 0.426 | 0.0795 | 0.2958 | 0.615 |
| 15 | 15 | 1 | 0.398 | 0.0791 | 0.2694 | 0.588 |
| 16 | 14 | 1 | 0.369 | 0.0784 | 0.2437 | 0.560 |
| 17 | 13 | 1 | 0.341 | 0.0774 | 0.2186 | 0.532 |
| 22 | 9 | 2 | 0.265 | 0.0765 | 0.1507 | 0.467 |
| 23 | 7 | 2 | 0.189 | 0.0710 | 0.0909 | 0.395 |

Probability of remission with covariates

- Can estimate the probability of remission for
 - Any time point
 - A given group
 - AND a given wbc level

$$\text{Risk}(t) = L_t \times e^{-1.3861 \times \text{group} + 1.6909 \times \text{wbc}}$$

Estimation of risk probability

- **Step 1: determine baseline risk during a period**
- **Step 2: calculate the average “linear term” (M)**
- **Step 3: calculate the individual “linear term” (L)**
- **Step 4: calculate $D = L - M$**
- **Step 5: calculate the risk of event**

Estimation of risk probability: example

Example: An individual on treatment =1 with wbc = 3.0

What is the individual's probability of remission at 10 weeks?

Step 1 – determine baseline risk

- 10-week baseline “survival” is 0.565

$$S_0(10) = 0.565$$

| time | n.risk | n.event | survival | std.err | lower 95% CI | upper 95% CI |
|------|--------|---------|----------|---------|--------------|--------------|
| 1 | 42 | 2 | 0.952 | 0.0329 | 0.8901 | 1.000 |
| 2 | 40 | 2 | 0.905 | 0.0453 | 0.8202 | 0.998 |
| 3 | 38 | 1 | 0.881 | 0.0500 | 0.7883 | 0.985 |
| 4 | 37 | 2 | 0.833 | 0.0575 | 0.7279 | 0.954 |
| 5 | 35 | 2 | 0.786 | 0.0633 | 0.6709 | 0.920 |
| 6 | 33 | 3 | 0.714 | 0.0697 | 0.5899 | 0.865 |
| 7 | 29 | 1 | 0.690 | 0.0715 | 0.5628 | 0.845 |
| 8 | 28 | 4 | 0.591 | 0.0764 | 0.4588 | 0.762 |
| 10 | 23 | 1 | 0.565 | 0.0773 | 0.4325 | 0.739 |
| ... | | | | | | |

Step 2 – calculate the average “linear term” (M)

- Using the mean of risk factors to calculate M
- Mean group = 0.5
- Mean wbc = 2.93
- $M = (-1.3861 \times 0.5) + (1.6909 \times 2.93)$
= 4.261

Step 3 - calculate the individual “linear term” (L)

- Using the individual's group and wbc to calculate L
- group = 1 (on treatment)
- wbc = 3.0
- $L = (-1.3861 \times 1) + (1.6909 \times 3)$
= 3.6866

Step 4 - calculate $D = L - M$

- Difference between the individual's linear term and average linear term

$$D = L - M$$

$$= 3.6866 - 4.261$$

$$= -0.5744$$

Step 5 - calculate the risk of event

We want to estimate the 10-week risk for the individual:

$$\begin{aligned}\text{Risk}(10) &= 1 - [S_0]^{\exp(d)} \\ &= 1 - 0.565^{\exp(-0.5744)} \\ &= 0.682\end{aligned}$$

The risk of remission at week 10 is 68.2%.

What is the risk of remission at week 10 for a control patient?

Exponential and Weibull models

Comparison

```
exponential = survreg(Surv(time, status==0) ~ group + wbc,  
dist="exponential")
```

```
summary(exponential)
```

```
weibull = survreg(Surv(time, status==0) ~ group + wbc)
```

```
summary(weibull)
```

Summary

- **Time-to-event data**
- **Kaplan-Meier analysis (actuarial analysis)**
- **Cox' s regression allows an assessment of risk factors**
- **Cox' s regression provides a very useful prognostic model in clinical medicine**