

BIOS 6312: Modern Regression Analysis

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Set 5 supplementary slides for R enthusiasts

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EXAMPLES FOR SET 5

Examples for R enthusiasts:

- ▶ Kaplan-Meier by gender (Slide 517)
- ▶ Log-rank test for gender (Slide 530)
- ▶ Cox regression by gender (Slide 541)
- ▶ Time-dependent Cox for heart transplant (Slide 565)
- ▶ Cumulative incidence of cardiovascular death (Slide 577)

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- ▶ **Kaplan-Meier by gender (Slide 517)**
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KAPLAN-MEIER BY GENDER

Reading in the MRI data:

- ▶ Read in data:

```
mri.data <- read.csv("mri.csv",  
                    header = TRUE,  
                    stringsAsFactors = FALSE)
```

KAPLAN-MEIER BY GENDER

Survival library:

- ▶ Many methods to model time-to-event data rely on the `survival` package.

```
library("survival")
```

KAPLAN-MEIER BY GENDER

Kaplan-Meier estimation:

- ▶ The `survfit` function allows us to perform Kaplan-Meier estimation.

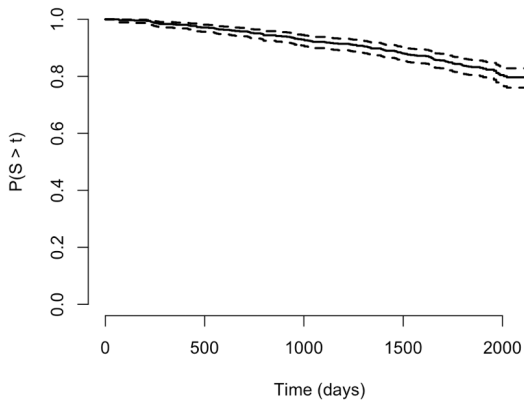
```
s.overall <- survfit(Surv(obstime, death) ~ 1,  
                    conf.type = "log-log",  
                    data = mri.data)
```

```
plot(s.overall,  
     frame.plot = FALSE,  
     col = c("black"),  
     lwd = 2,  
     xlab = "Time (days)",  
     ylab = "P(S > t)")
```

- ▶ Confidence intervals from a log-log transformation will allow us to come close to Stata's results.

KAPLAN-MEIER BY GENDER

Kaplan-Meier estimation: Overall



- Confidence intervals included if only presenting one group.

KAPLAN-MEIER BY GENDER

Kaplan-Meier estimation:

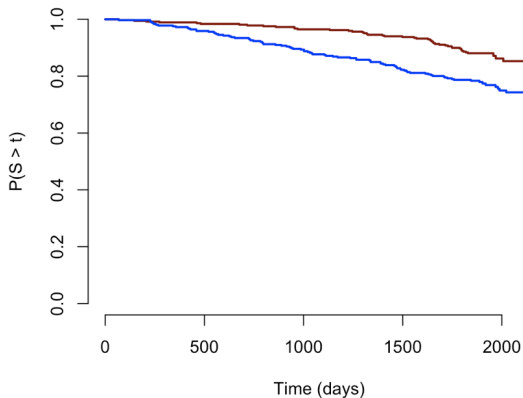
- ▶ Curves stratified by gender.

```
s.gender <- survfit(Surv(obstime, death) ~ male,  
                    conf.type = "log-log",  
                    data = mri.data)
```

```
plot(s.gender,  
     frame.plot = FALSE,  
     conf.int = FALSE,  
     col = c("darkred", "blue"),  
     lwd = c(2,2),  
     xlab = "Time (days)",  
     ylab = "P(S > t)")
```


KAPLAN-MEIER BY GENDER

Kaplan-Meier estimation: By gender



- Confidence intervals not included if presenting 2+ groups.

KAPLAN-MEIER BY GENDER

Kaplan-Meier estimation:

- ▶ Extracting restricted mean (overall).

```
summary(s.overall)$table[5:6]
```

```
## Output
```

```
  *rmean *se(rmean)
```

```
1974.46913  16.56625
```

- ▶ Agrees with Stata output (Slide 520).

KAPLAN-MEIER BY GENDER

Kaplan-Meier estimation:

- ▶ Extracting restricted mean (by gender).

```
summary(s.gender)$table[,5:6]
```

```
## Output
```

```
      *rmean *se(rmean)
male=0 2049.954   17.55878
male=1 1899.064   27.59734
```

- ▶ Approximately agrees with Stata output (Slide 521).

KAPLAN-MEIER BY GENDER

Kaplan-Meier estimation:

- ▶ Extracting quantiles (overall).

```
quantile(s.overall, 0.10)
```

```
## Output
```

```
$quantile
```

```
10
```

```
1338
```

```
$lower
```

```
10
```

```
1045
```

```
$upper
```

```
10
```

```
1519
```

- ▶ Agrees with Stata output (Slide 522).

KAPLAN-MEIER BY GENDER

Kaplan-Meier estimation:

- ▶ Extracting quantiles (by gender).

```
quantile(s.gender, 0.20)

## Output
$quantile
      20
male=0  NA
male=1 1707

$lower
      20
male=0  NA
male=1 1457

$upper
      20
male=0  NA
male=1 1988
```

- ▶ Approximately agrees with Stata output (Slide 523).

EXAMPLES FOR SET 5

Examples for R enthusiasts:

- ▶ *Kaplan-Meier by gender (Slide 517)*
- ▶ **Log-rank test for gender (Slide 530)**
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- ▶ Cumulative incidence of cardiovascular death (Slide 577)

LOG-RANK TEST FOR GENDER

Log-rank test:

- ▶ Function for log-rank test: `survdiff`.

```
logrank.gender

## Output
Call:
survdiff(formula = Surv(obstime, death) ~ male, data = mri.data)

      N Observed Expected (O-E)^2/E (O-E)^2/V
male=0 369      47    68.8      6.89    14.3
male=1 366      86    64.2      7.38    14.3

Chisq= 14.3 on 1 degrees of freedom, p= 2e-04
```

- ▶ Agrees with Stata output (Slide 530).

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COX REGRESSION BY GENDER

Proportional hazards regression:

- ▶ Function for Cox model: `coxph`.

```
model.gen <- coxph(Surv(obstime, death) ~ male,  
                  ties = "breslow",  
                  data = mri.data)  
N <- length(model.gen$residuals)  
robust.var <- sandwich(model.gen) * N/(N - 1)
```

COX REGRESSION BY GENDER

Proportional hazards regression:

► Results:

```
exp(c(HR = model.gen$coef,  
      CI.Low = model.gen$coef - qnorm(0.975) * sqrt(robust.var),  
      CI.Hi = model.gen$coef + qnorm(0.975) * sqrt(robust.var)))  
  
## Output  
HR.male  CI.Low  CI.Hi  
1.961765 1.378102 2.792624
```

► Agrees with Stata output (Slide 543).

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TIME-DEPENDENT COX FOR HEART TRANSPLANT

Reading in the transplant data:

- ▶ Read in data:

```
heart.data <- read.csv("transplant.csv",  
                      header = TRUE,  
                      stringsAsFactors = FALSE)
```

TIME-DEPENDENT COX FOR HEART TRANSPLANT

Structuring data:

- ▶ Create variable for initial time windows.

```
N <- dim(heart.data)[1]
heart.data$ptime <- c(0, heart.data$time[1:(N - 1)])
heart.data$ptime[heart.data$time == 1] <- 0
```

TIME-DEPENDENT COX FOR HEART TRANSPLANT

Time-dependent covariates:

- ▶ Account for clustering.

```
model.heart <- coxph(Surv(ptime, time, death) ~ transplant + cluster(id),  
                    method = "breslow",  
                    data = heart.data)  
robust.var <- summary(model.heart)$coef[4]^2 * 21/20
```

TIME-DEPENDENT COX FOR HEART TRANSPLANT

Time-dependent covariates:

► Results:

```
exp(c(HR = model.heart$coef,  
      CI.Low = model.heart$coef - qnorm(0.975) * sqrt(robust.var),  
      CI.High = model.heart$coef + qnorm(0.975) * sqrt(robust.var)))  
  
## Output  
      HR.transplant  CI.Low.transplant  CI.High.transplant  
      0.261495      0.086369      0.791713
```

► Agrees with Stata output (Slide 566).

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- ▶ **Cumulative incidence of cardiovascular death (Slide 577)**

CUMULATIVE INCIDENCE OF CARDIOVASCULAR DEATH

Subdistribution hazard in R:

- ▶ Competing risks regression can be performed using the `crr` function in the library `cmprsk`.
- ▶ We again use the MRI data.

```
library("cmprsk")
```

```
mri.data <- read.csv("mri.csv",  
                    header = TRUE,  
                    stringsAsFactors = FALSE)
```

CUMULATIVE INCIDENCE OF CARDIOVASCULAR DEATH

Structuring data:

- ▶ Create variable indicating censoring, event of interest, and competing event(s).

```
mri.data$event <- 0
mri.data$event[mri.data$death == 1 & mri.data$cvd == 1] <- 1
mri.data$event[mri.data$death == 1 & mri.data$cvd == 0] <- 2
```

CUMULATIVE INCIDENCE OF CARDIOVASCULAR DEATH

Subdistribution hazard regression:

- ▶ Account for competing risks.

```
model.diab <- crr(mri.data$obstime,  
                 mri.data$event,  
                 mri.data$diabetes)  
N <- model.diab$n  
robust.var <- summary(model.diab)$coef[3]^2 * N / (N - 1)
```

CUMULATIVE INCIDENCE OF CARDIOVASCULAR DEATH

Subdistribution hazard regression:

- ▶ Results:

```
exp(c(summary(model.diab)$coef[1],
      summary(model.diab)$coef[1] - qnorm(0.975) * sqrt(robust.var),
      summary(model.diab)$coef[1] + qnorm(0.975) * sqrt(robust.var)))

## Output
[1] 2.44409 1.34119 4.45391
```

- ▶ Agrees with Stata output (Slide 582).

CUMULATIVE INCIDENCE OF CARDIOVASCULAR DEATH

Plot cumulative incidence:

► Create figures:

```
plot(predict(model.diab, cov1 = 0),
      frame.plot = FALSE,
      ylim = c(0, 0.2),
      col = "darkblue",
      xlab = "Analysis time",
      ylab = "Cumulative incidence")

lines(predict(model.diab, cov1 = 1),
       col = "darkred")

legend(0, 0.20, col = c("darkblue", "darkred"),
       lwd = c(1,1), lty = c(1,1),
       c("diabetes = 0", "diabetes = 1"))
```

CUMULATIVE INCIDENCE OF CARDIOVASCULAR DEATH

Cumulative incidence: By diabetes status

