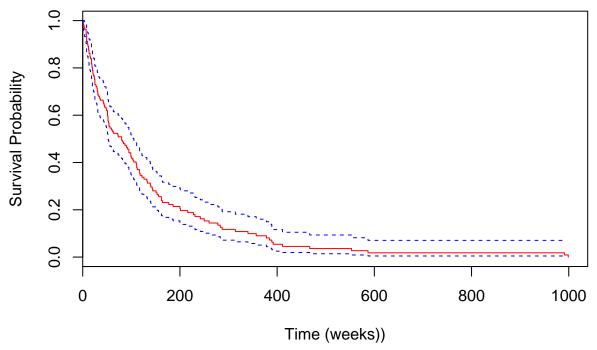
PSTAT175 Lab B

Mujie Wang 2019/10/21

```
#1
#a)
vet <- read.table("/Users/wangmujie/Desktop/vets.txt")</pre>
library(survival)
vet.time <- vet$V1</pre>
vet.event <- vet$V2</pre>
vet.surv <- Surv(vet.time,vet.event)</pre>
vet.fit <- survfit(vet.surv ~ 1)</pre>
vet.fit
## Call: survfit(formula = vet.surv ~ 1)
##
            events median 0.95LCL 0.95UCL
         n
       137
                128
                                  52
plot(vet.fit, main= "Kaplan-Meier Curves for subject in VA medical",
     xlab = "Time (weeks))",
     ylab = "Survival Probability",
     col = c("red","blue","blue")
```

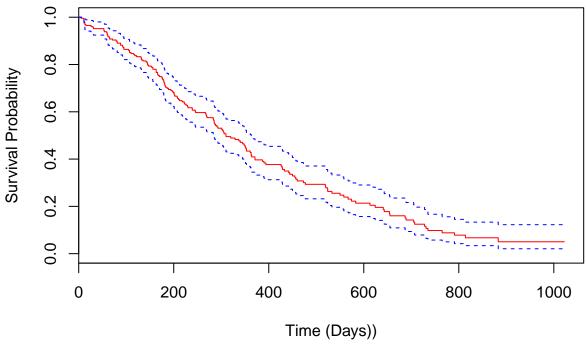
Kaplan-Meier Curves for subject in VA medical



```
#b)
q1tm <- min(vet.fit$time[vet.fit$surv < 0.75])
q2tm <- min(vet.fit$time[vet.fit$surv < 0.5])</pre>
```

```
q3tm <- min(vet.fit$time[vet.fit$surv < 0.25])
c(q1tm, q2tm, q3tm)
## [1] 25 80 162
quantile(vet.fit,c(0.25,0.5,0.75))$q
##
        50 75
    25
##
    25
        80 162
#2
#a)
library(survival)
data(lung)
lung.fit <- survfit(Surv(lung$time,lung$status) ~ 1)</pre>
plot(lung.fit, main= "Kaplan-Meier Curves for lung",
     xlab = "Time (Days))",
     ylab = "Survival Probability",
     conf.int = TRUE,col=c("red","blue","blue")
```

Kaplan-Meier Curves for lung



```
#b)
summary(lung.fit,times=150)

## Call: survfit(formula = Surv(lung$time, lung$status) ~ 1)
##
## time n.risk n.event survival std.err lower 95% CI upper 95% CI
## 150 179 47 0.793 0.0269 0.742 0.848

#According to the data, (0.741,0.848)is the 95% Confidence Interval
#for the survivial function at 150 days
```

```
#c)
med.tm <- min(lung.fit$time[lung.fit$surv < 0.5])</pre>
med.low.tm <- min(lung.fit$time[lung.fit$lower < 0.5])</pre>
med.upper.tm <- min(lung.fit$time[lung.fit$upper < 0.5])</pre>
c(med.low.tm, med.tm, med.upper.tm)
## [1] 285 310 363
#According to the data, (285,363) is the 95% Confidence Interval for the median survivial time
#d)
sex <- as.factor(lung$sex)</pre>
lung.fit.sex <- survfit(Surv(lung$time,lung$status) ~ sex)</pre>
plot(lung.fit.sex, col=c(2,4),lwd=2,conf.int= FALSE, xlab = "Days", ylab = "Survival Function")
legend(0,0.3,c("Male","Female"), fill= c(2,4))
      0.8
Survival Function
      9.0
      0.4
      0.2
                  Male
                  Female
      0.0
                         200
                                        400
                                                                                    1000
           0
                                                       600
                                                                      800
                                                 Days
```

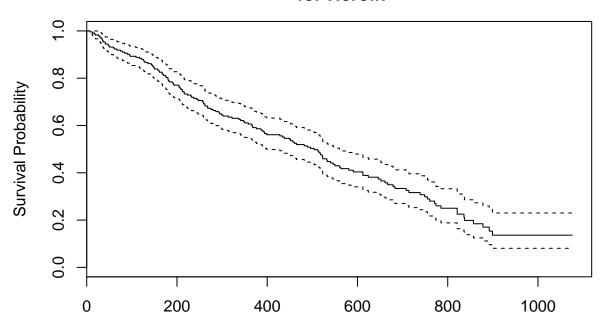
#Generally speaking, women have better suivival rates. By looking through the graph, it is clear that #the women's curve are above men'a curve. Since a higher survival function means a longer time until #failure or death, we can say women have better suivival rates.

```
#e)
quantile(lung.fit.sex,0.5)
```

```
## $quantile
## 50
## sex=1 270
## sex=2 426
##
## $lower
## 50
## sex=1 212
## sex=2 348
```

```
## $upper
##
          50
## sex=1 310
## sex=2 550
#The interval for women is (348,550) while the interval for men is (212,310);
#the median survival time for women is 426 days while for men is 270 days;
#This result agrees with our assesment that women have better suivival rates than men.
#But it may not tell the whole story since the curves come back together at the end which
#indicates the benifit for women may diminish overtime.
#3
#a)
library(survival)
heroin <- read.table("/Users/wangmujie/desktop/Heroin.txt")
heroin.fit <- survfit(Surv(heroin$Time,heroin$Status) ~ 1)</pre>
plot(heroin.fit, main= "Kaplan-Meier Curves \n for Heroin",
     xlab = "Days each subject spent in the clinics)",
     ylab = "Survival Probability",
     conf.int = TRUE
```

Kaplan-Meier Curves for Heroin

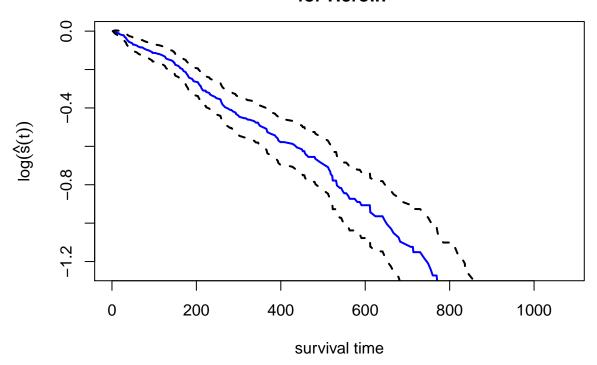


Days each subject spent in the clinics)

```
#b)
mj = heroin.fit$n.event
nj = heroin.fit$n.risk
Vj=mj/nj/(nj-mj)
cVj=cumsum(Vj)
lowerCI = log(heroin.fit$surv)-1.96*sqrt(cVj)
upperCI = log(heroin.fit$surv)+1.96*sqrt(cVj)
par(mar=c(5,5,4,2))
```

```
plot(heroin.fit$time,log(heroin.fit$surv),lwd =2, type = "l",
    ylim = c(-1.25,0),
    main= "Kaplan-Meier Curves \n for Heroin",
    xlab = " survival time ",
    ylab = expression(log(hat(s)(t))),col="blue")
lines(heroin.fit$time,lowerCI,lty=2,col=1,lwd=2)
lines(heroin.fit$time,upperCI,lty=2,col=1,lwd=2)
```

Kaplan-Meier Curves for Heroin



```
0; - (1)S - (20) - (20) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30) - (30)
```

#d)

```
summary(heroin.fit,times = 365)
## Call: survfit(formula = Surv(heroin$Time, heroin$Status) ~ 1)
##
##
   time n.risk n.event survival std.err lower 95% CI upper 95% CI
                     87
                           0.606 0.0331
                                                 0.545
#NULL Hypothesis(H0)is 50% or less of the patients are discharged by the one year mark;
#Estimate is that nearly less than 40% have been discharged at 365 days.
#Therefore, ther is no evidencr for Ha that the survival function is less than 50%
#two-sided test:
heroin.tk <- max(heroin.fit$time[heroin.fit$time <365])
heroin.fit$surv[heroin.fit$time == heroin.tk ]
## [1] 0.6060647
zstats <- (log(heroin.fit$surv[heroin.fit$time == heroin.tk]) - log(0.5))/heroin.fit$std.err[heroin.fit
zstats
## [1] 3.524093
# zstats > 1.96(critical value while standard error=0.05)
# we can conclude that a statistically significant fewer than 50% are being discharged by the end of th
pnorm(-zstats)
## [1] 0.0002124677
\#p\text{-value} = 0.0002124677
#95% CI for the 70th percentile
heroin.LI <- min(heroin.fit$time[heroin.fit$lower < 0.3])
```

heroin.UI <- max(heroin.fit\$time[heroin.fit\$upper > 0.3])

```
summary(heroin.fit,times = c(heroin.LI,heroin.UI))
## Call: survfit(formula = Surv(heroin$Time, heroin$Status) ~ 1)
##
## time n.risk n.event survival std.err lower 95% CI upper 95% CI
##
     661
             46
                    130
                           0.357 0.0359
                                                0.294
                                                              0.435
##
     826
             18
                     14
                           0.225 0.0368
                                                 0.163
                                                              0.310
c(heroin.LI,heroin.UI)
## [1] 661 826
#95% CI for the 70th percentile is (661,826)
#95% CI for the 80th percentile
heroin.LI <- min(heroin.fit$time[heroin.fit$lower < 0.2])</pre>
heroin.UI <- max(heroin.fit$time[heroin.fit$upper > 0.2])
summary(heroin.fit,times = c(heroin.LI,1050,heroin.UI))
## Call: survfit(formula = Surv(heroin$Time, heroin$Status) ~ 1)
##
## time n.risk n.event survival std.err lower 95% CI upper 95% CI
##
    774
             27
                    141
                           0.260 0.0364
                                               0.1978
                                                              0.342
## 1050
              2
                      9
                           0.136 0.0364
                                                0.0807
                                                              0.230
                      0
## 1076
              1
                           0.136 0.0364
                                               0.0807
                                                              0.230
#95% CI for the 80th percentile is (774,1076)
\# tk=1076 is the last time in the whole data set, it is a censored observation.
```