## HW2

#### Bulun Te

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```
library(dplyr)
library(survival)
library(kableExtra)
library(knitr)
library(ggplot2)
# Reading the data
data = read.table("Breast_cancer_Table_1_2_Collet.txt")
# Changing the column names
data = data %>% rename(
 id = V1,
 x = V2,
 delta = V3,
 z = V4
)
data %>% head()
##
    id x delta z
## 1 1 224
               0 0
## 2 2 212
               0 0
## 3 3 208
               0 0
## 4 4 198
               0 0
## 5 5 181
               1 0
## 6 6 148
               1 0
data %>% psych::describe()
                         sd median trimmed
##
        vars n mean
                                             mad min max range skew kurtosis
## id
          1 45 23.00 13.13
                                23
                                     23.00 16.31
                                                  1 45
                                                           44 0.00
                                                                       -1.28
           2 45 96.22 69.35
                                71
                                     92.32 66.72
                                                  5 225
                                                          220 0.51
                                                                       -1.09
## delta
           3 45 0.58 0.50
                               1
                                   0.59 0.00
                                                  0 1
                                                           1 -0.30
                                                                       -1.95
                               1
                                     0.76 0.00
           4 45 0.71 0.46
                                                            1 -0.90
                                                                       -1.21
##
           se
## id
         1.96
## x
        10.34
## delta 0.07
         0.07
## z
```

# Answer without using survival package

### Question 1(a)

```
\hat{\Lambda}_{NA}(t) = \sum_{t_j \leq t} \frac{D_j}{Y_j}
```

where  $D_j$  is the number of events at time  $t_j$  and  $Y_j$  is the number of individuals at risk at time  $t_j$ .

```
# sort data with x column with aescending order

data_sorted = data %>% arrange(x)

data_sorted
```

```
##
          x delta z
     id
## 1
     14
          5
## 2
     15
          8
                1 1
## 3
     16 10
## 4 17
         13
                1 1
## 5
     18 18
## 6
     13 23
                1 0
## 7
     19 24
## 8
     20 25
## 9
     21 26
## 10 22
         31
                1 1
## 11 23 35
                1 1
## 12 24 40
                1 1
## 13 25 41
                1 1
## 14 12 47
                1 0
## 15 26 48
                1 1
## 16 27
         50
## 17 28
         59
                1 1
## 18 29
         61
## 19 30
        68
                1 1
## 20 11
         69
## 21 10
         70
                0 0
## 22 9 71
## 23 31 71
                1 1
## 24 32 76
## 25 8 100
                0 0
## 26 7 101
                0 0
## 27 33 105
                0 1
## 28 34 107
## 29 35 109
## 30 36 113
## 31 37 116
## 32 38 118
## 33 39 143
## 34 6 148
                1 0
## 35 40 154
## 36 41 162
                0 1
## 37 5 181
## 38 42 188
                0 1
```

```
## 39 4 198
                 0 0
## 40 3 208
                 0 0
## 41 2 212
## 42 43 212
                 0 1
## 43 44 217
                 0 1
## 44 1 224
                 0 0
## 45 45 225
                 0 1
data_sorted$y = sapply(data_sorted$x,function(u){sum(data_sorted$x >= u)})
data_sorted$lambda = data_sorted$delta / data_sorted$y
data_sorted$cumulative_hazard = cumsum(data_sorted$lambda)
cumulative_hazard = data_sorted %>%
  select(x,delta,y,lambda,cumulative_hazard) %>%
  filter(delta ==1) %>%
  select(cumulative hazard)
ppl_at_risk = data_sorted %>%
  select(x,delta,y,lambda,cumulative_hazard) %>%
  filter(delta >=1) %>% select(x,y)
data2 = data.frame(time = ppl_at_risk$x,
                    Y = ppl_at_risk$y,
                   Nelson_Alan_Cumaltive= cumulative_hazard)
data2
```

```
##
      time Y cumulative_hazard
## 1
         5 45
                     0.0222222
## 2
         8 44
                     0.04494949
## 3
        10 43
                     0.06820531
## 4
        13 42
                     0.09201483
## 5
        18 41
                     0.11640508
## 6
        23 40
                     0.14140508
## 7
        24 39
                     0.16704610
## 8
        25 38
                     0.19336189
## 9
        26 37
                     0.22038892
        31 36
## 10
                     0.24816670
## 11
        35 35
                     0.27673813
## 12
        40 34
                     0.30614989
## 13
        41 33
                     0.33645292
## 14
        47 32
                     0.36770292
## 15
        48 31
                     0.39996098
## 16
        50 30
                     0.43329432
## 17
        59 29
                      0.46777708
## 18
        61 28
                     0.50349136
## 19
        68 27
                      0.54052840
        69 26
## 20
                     0.57898994
## 21
        71 24
                     0.62065660
## 22 113 16
                     0.68315660
## 23 118 14
                     0.75458518
## 24 143 13
                     0.83150825
```

```
## 25 148 12 0.91484159
## 26 181 9 1.02595270
```

### Question 1(b)

Assuming the  $\hat{\Lambda}(t)$  follows normal distribution, and the variance of  $\hat{\Lambda}(t)$  is estimated as  $\hat{V}(t) = \sum_{t_j \leq t} \frac{D_j}{Y_j^2}$ , then the confidence interval is estimated as  $\hat{\Lambda}(t) \pm z_{0.975} \sqrt{\hat{V}(t)}$ .

```
# Computing 95% confidence interval for the cumulative hazard assuming lambda(t) follows normal distrib

V_estimate = ((data_sorted$delta)/(data_sorted$y^2)) %>% cumsum()

CI_upper = data_sorted$cumulative_hazard + qnorm(0.975)*sqrt(V_estimate)

CI_lower = data_sorted$cumulative_hazard - qnorm(0.975)*sqrt(V_estimate)

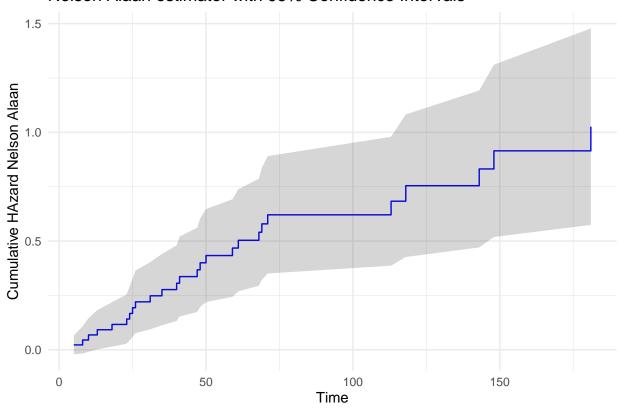
result_1_b = data_sorted %>% mutate(CI_upper = CI_upper, CI_lower = CI_lower)%>%
    filter(delta>=1) %>%
    select(x,cumulative_hazard,CI_upper,CI_lower) %>%
    round(.,6)

result_1_b
```

```
##
        x cumulative_hazard CI_upper CI_lower
## 1
                   0.022222 0.065777 -0.021333
## 2
       8
                   0.044949 0.107249 -0.017350
## 3
       10
                   0.068205 0.145399 -0.008988
## 4
       13
                   0.092015 0.182218 0.001812
## 5
       18
                   0.116405 0.218492
                                      0.014318
## 6
       23
                   0.141405 0.254642
                                      0.028168
## 7
       24
                   0.167046 0.290934
                                      0.043158
## 8
       25
                   0.193362 0.327558
                                     0.059166
## 9
       26
                   0.220389 0.364662
                                      0.076116
## 10
       31
                   0.248167 0.402370 0.093963
## 11
       35
                   0.276738 0.440795 0.112682
## 12
       40
                   0.306150 0.480040 0.132260
                   0.336453 0.520206 0.152700
## 13
       41
## 14
       47
                   0.367703 0.561395 0.174011
## 15
       48
                   0.399961 0.603711 0.196211
## 16
       50
                   0.433294 0.647262 0.219327
## 17
       59
                   0.467777 0.692165 0.243389
                   0.503491 0.738544 0.268439
## 18
       61
## 19
       68
                   0.540528 0.786535 0.294522
## 20
       69
                   0.578990 0.836287
                                      0.321693
## 21
       71
                   0.620657 0.890603
                                      0.350710
## 22 113
                   0.683157 0.979597
                                      0.386716
## 23 118
                   0.754585 1.082421
                                      0.426750
## 24 143
                   0.831508 1.192350
                                      0.470667
## 25 148
                   0.914842 1.310927
                                      0.518757
## 26 181
                   1.025953 1.477958 0.573947
```

```
ggplot(result_1_b, aes(x = x)) +
  geom_step(aes(y = cumulative_hazard), direction = "hv", col = "blue") +
  geom_ribbon(aes(ymin = CI_lower, ymax = CI_upper), alpha = 0.2) +
  labs(x = "Time", y = "Cumulative HAzard Nelson Alaan") +
  ggtitle("Nelson Alaan estimator with 95% Confidence Intervals") +
  theme_minimal()
```

#### Nelson Alaan estimator with 95% Confidence Intervals



#### Question 1(c)

Assuming the  $\hat{\Lambda}(t)$  follows normal distribution. The variance of  $\hat{\Lambda(t)}$ , using Delta method is estimated as  $\hat{V}(\log(\hat{\Lambda}(t)) = \sum_{t_j \leq t} \frac{D_j}{Y_i^2 \hat{\Lambda}(t)^2}$ , then the confidence interval is estimated as  $\exp(\log(\hat{\Lambda}(t) \pm z_{0.975} \sqrt{\hat{V}(\log(\hat{\Lambda}(t)))})$ .

```
# Computing 95% confidence interval for the cumulative hazard assuming
# log(lambda(t)) follows normal distribution

# Computing using delta mathod

V_estimate = ((data_sorted$delta)/(data_sorted$y^2)) %>% cumsum()

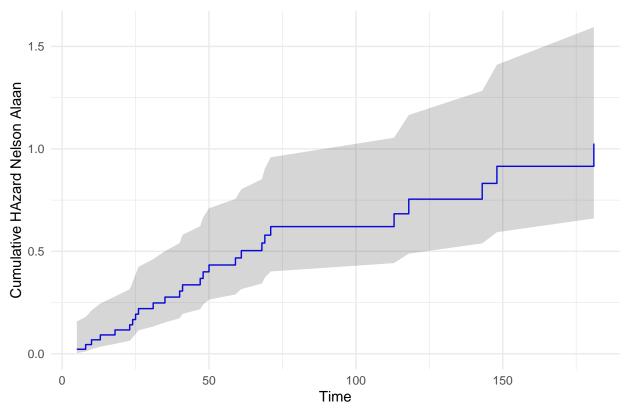
V_estimate = V_estimate / (data_sorted$cumulative_hazard^2)

CI_upper = data_sorted$cumulative_hazard * exp(qnorm(0.975)*sqrt(V_estimate))
```

```
CI_lower = data_sorted$cumulative_hazard * exp(-qnorm(0.975)*sqrt(V_estimate))
result_1_c = data_sorted %>%
  mutate(CI_upper = CI_upper, CI_lower = CI_lower)%>%
  filter(delta>=1) %>%
  select(x,cumulative_hazard,CI_upper,CI_lower) %>%
 round(.,6)
result_1_c
##
        x cumulative_hazard CI_upper CI_lower
## 1
                   0.022222 0.157757 0.003130
                   0.044949 0.179743 0.011241
## 2
       8
                   0.068205 0.211517 0.021993
## 3
       10
## 4
       13
                   0.092015 0.245245 0.034524
## 5
       18
                   0.116405 0.279800 0.048428
## 6
       23
                   0.141405 0.314955 0.063487
## 7
                   0.167046 0.350693 0.079569
       24
## 8
       25
                   0.193362 0.387059 0.096597
## 9
       26
                   0.220389 0.424122 0.114522
## 10 31
                   0.248167 0.461957 0.133317
## 11
       35
                   0.276738 0.500643 0.152971
## 12
       40
                   0.306150 0.540267 0.173484
## 13
      41
                   0.336453 0.580916 0.194866
## 14
      47
                   0.367703 0.622683 0.217134
## 15
       48
                   0.399961 0.665668 0.240313
## 16
     50
                   0.433294 0.709978 0.264436
## 17
      59
                   0.467777 0.755728 0.289542
                   0.503491 0.803046 0.315677
## 18
     61
## 19
       68
                   0.540528 0.852071 0.342895
## 20 69
                   0.578990 0.902957 0.371257
## 21 71
                   0.620657 0.958831 0.401754
## 22 113
                   0.683157 1.054320 0.442658
## 23 118
                   0.754585 1.165174 0.488681
## 24 143
                   0.831508 1.283314 0.538766
## 25 148
                   0.914842 1.410509 0.593357
## 26 181
                   1.025953 1.593914 0.660374
```

```
ggplot(result_1_c, aes(x = x)) +
  geom_step(aes(y = cumulative_hazard), direction = "hv", col = "blue") +
  geom_ribbon(aes(ymin = CI_lower, ymax = CI_upper), alpha = 0.2) +
  labs(x = "Time", y = "Cumulative HAzard Nelson Alaan") +
  ggtitle("Nelson Alaan estimator with 95% Confidence Intervals") +
  theme_minimal()
```





#### Question 1(d)

Assuming the  $log(\Lambda(t))$  follows normal distribution. The variance of  $log(\Lambda(t))$  is estimated as above. As  $S(t) = exp(-\Lambda(t))$ , thus  $P\{\hat{\Lambda}(t) \in (a,b)\} = P\{\hat{S}(t) \in (e^{-b},e^{-a})\} = 0.95$  And the CI are computed as follow:

```
V_estimate_temp = ((data_sorted$delta)/(data_sorted$y^2)) %>% cumsum()

V_estimate = V_estimate_temp / (data_sorted$cumulative_hazard^2)

S_estimate = exp(-data_sorted$cumulative_hazard)

CI_lower = exp(-data_sorted$cumulative_hazard * exp(qnorm(0.975)*sqrt(V_estimate)))

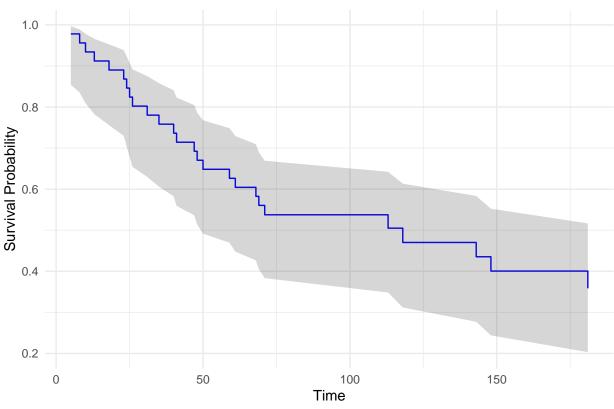
CI_upper = exp(-data_sorted$cumulative_hazard * exp(-qnorm(0.975)*sqrt(V_estimate)))

result_1_d = data_sorted %>%
    mutate(CI_upper = CI_upper, CI_lower = CI_lower,S_estimate=S_estimate)%>%
    filter(delta>=1) %>%
    select(x,CI_upper,S_estimate,CI_lower) %>%
    round(.,6)
```

## x CI\_upper S\_estimate CI\_lower

```
## 1
       5 0.996875
                   0.978023 0.854057
## 2
      8 0.988822
                   0.956046 0.835484
## 3
     10 0.978247
                   0.934069 0.809356
## 4
     13 0.966066
                   0.912092 0.782513
## 5
      18 0.952726
                   0.890115 0.755935
## 6
      23 0.938487
                   0.868138 0.729822
## 7
      24 0.923514 0.846161 0.704200
## 8
     25 0.907922
                   0.824184 0.679051
## 9
      26 0.891792
                   0.802207 0.654344
## 10 31 0.875188 0.780230 0.630050
## 11 35 0.858155
                   0.758253 0.606140
## 12 40 0.840731
                   0.736276 0.582593
## 13 41 0.822945
                   0.714300 0.559386
## 14 47 0.804822
                   0.692323 0.536503
## 15 48 0.786382
                   0.670346 0.513930
## 16 50 0.767638
                   0.648370 0.491655
## 17 59 0.748606
                   0.626393 0.469669
## 18 61 0.729295
                   0.604417 0.447962
## 19 68 0.709713
                   0.582440 0.426531
## 20 69 0.689866
                   0.560464 0.405369
## 21 71 0.669145
                   0.537591 0.383341
## 22 113 0.642327
                   0.505020 0.348429
## 23 118 0.613435
                   0.470206 0.311868
## 24 143 0.583468
                   0.435392 0.277117
## 25 148 0.552470 0.400580 0.244019
# plotting the survival function and its confidence interval from result_1_d as step function
ggplot(result_1_d, aes(x = x)) +
 geom_step(aes(y = S_estimate), direction = "hv", col = "blue") +
  geom_ribbon(aes(ymin = CI_lower, ymax = CI_upper), alpha = 0.2) +
 labs(x = "Time", y = "Survival Probability") +
 ggtitle("Survival Function with 95% Confidence Intervals") +
 theme minimal()
```





## Question 1(e)

Quartile of 0.75 is unable to compute due to the estimation of survival function ends in 0.358. However, the quartiles of 0.25 and 0.5 are computed as follow:

```
print("quartiles of 0.25,0.5 of survival estimations are")
## [1] "quartiles of 0.25,0.5 of survival estimations are"
```

```
c(min(result_1_d$x[which(result_1_d$S_estimate <= (1-0.25))]),
min(result_1_d$x[which(result_1_d$S_estimate <= 0.5)]))</pre>
```

**##** [1] 40 118