

Cox regression

A manually worked out, simple example: two groups

Load libraries

```
library(tidyverse)
```

```
## -- Attaching packages ----- tidyverse_

## v ggplot2 3.1.0      v purrr  0.3.0
## v tibble  2.0.1      v dplyr  0.7.8
## v tidyr   0.8.2      v stringr 1.3.1
## v readr   1.3.1      v forcats 0.3.0

## -- Conflicts ----- tidyverse_
## x dplyr::filter() masks stats::filter()
## x dplyr::lag()     masks stats::lag()
```

```
library(maxLik)
```

```
## Loading required package: miscTools

##
## Please cite the 'maxLik' package as:
## Henningsen, Arne and Toomet, Ott (2011). maxLik: A package for maximum likelihood estimation in R. C
##
## If you have questions, suggestions, or comments regarding the 'maxLik' package, please use a forum o
## https://r-forge.r-project.org/projects/maxlik/
```

```
library(survival)
```

```
##
## Attaching package: 'survival'

## The following object is masked from 'package:rpart':
##
## solder
```

Data definition

Lets enter the data in R:

```
dat <- data.frame(ratID = paste0("rat", 1:5),
                  time = c(55, 50, 70, 120, 110),
                  failure = c(0, 1, 1, 0, 1),
                  group = c(0, 1, 0, 1, 1))
```

Total number of failures D:

```
sum(dat$failure)
```

```
## [1] 3
```

For convenience, rename 'group' to 'x':

```
dat <- rename(dat, x = group)
dat
```

```
##   ratID time failure x
## 1  rat1   55        0 0
## 2  rat2   50        1 1
## 3  rat3   70        1 0
## 4  rat4  120        0 1
## 5  rat5  110        1 1
```

We also define an auxiliary data.frame containing events only:

```
dat.events <- subset(dat, failure == 1)
dat.events
```

```
##   ratID time failure x
## 2  rat2   50        1 1
## 3  rat3   70        1 0
## 5  rat5  110        1 1
```

Partial log-likelihood function

Lets define the partial (log-)likelihood function

```
pLogLik <- function(beta) {
  numerator <- with(dat.events, x * beta)
  denominator <- rep(NA_real_, length(numerator))
  for(j in seq_along(denominator))
  {
    t_j <- dat.events[j, "time"]
    risk_set <- subset(dat, time >= t_j)
    theta_j <- with(risk_set, exp(x * beta))
    denominator[j] <- log(sum(theta_j))
  }
  return(sum(numerator - denominator))
}
```

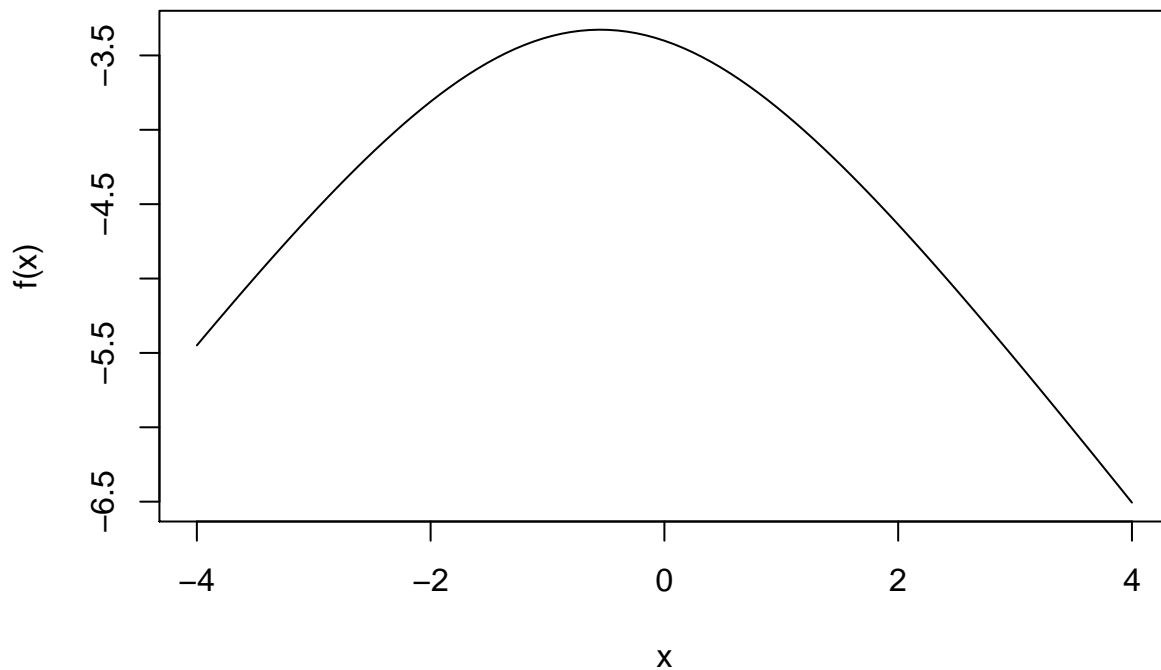
```
debugonce(pLogLik)
```

We can plot it:

```
f <- Vectorize(pLogLik)
curve(f, from = -4, to = 4)
```

```
## debugging in: (function (beta)
## {
##     numerator <- with(dat.events, x * beta)
##     denominator <- rep(NA_real_, length(numerator))
##     for (j in seq_along(denominator)) {
##         t_j <- dat.events[j, "time"]
##         risk_set <- subset(dat, time >= t_j)
##         theta_j <- with(risk_set, exp(x * beta))
##         denominator[j] <- log(sum(theta_j))
##     }
##     return(sum(numerator - denominator))
## }(beta = dots[[1L]][[1L]])
## debug at <text>#1: {
##     numerator <- with(dat.events, x * beta)
##     denominator <- rep(NA_real_, length(numerator))
##     for (j in seq_along(denominator)) {
##         t_j <- dat.events[j, "time"]
##         risk_set <- subset(dat, time >= t_j)
##         theta_j <- with(risk_set, exp(x * beta))
##         denominator[j] <- log(sum(theta_j))
##     }
##     return(sum(numerator - denominator))
## }
## debug at <text>#2: numerator <- with(dat.events, x * beta)
## debug at <text>#3: denominator <- rep(NA_real_, length(numerator))
## debug at <text>#4: for (j in seq_along(denominator)) {
##     t_j <- dat.events[j, "time"]
##     risk_set <- subset(dat, time >= t_j)
##     theta_j <- with(risk_set, exp(x * beta))
##     denominator[j] <- log(sum(theta_j))
## }
## debug at <text>#6: t_j <- dat.events[j, "time"]
## debug at <text>#7: risk_set <- subset(dat, time >= t_j)
## debug at <text>#8: theta_j <- with(risk_set, exp(x * beta))
## debug at <text>#9: denominator[j] <- log(sum(theta_j))
## debug at <text>#6: t_j <- dat.events[j, "time"]
## debug at <text>#7: risk_set <- subset(dat, time >= t_j)
## debug at <text>#8: theta_j <- with(risk_set, exp(x * beta))
## debug at <text>#9: denominator[j] <- log(sum(theta_j))
## debug at <text>#6: t_j <- dat.events[j, "time"]
## debug at <text>#7: risk_set <- subset(dat, time >= t_j)
## debug at <text>#8: theta_j <- with(risk_set, exp(x * beta))
## debug at <text>#9: denominator[j] <- log(sum(theta_j))
## debug at <text>#11: return(sum(numerator - denominator))
## exiting from: (function (beta)
## {
##     numerator <- with(dat.events, x * beta)
##     denominator <- rep(NA_real_, length(numerator))
##     for (j in seq_along(denominator)) {
##         t_j <- dat.events[j, "time"]
```

```
##      risk_set <- subset(dat, time >= t_j)
##      theta_j <- with(risk_set, exp(x * beta))
##      denominator[j] <- log(sum(theta_j))
##    }
##    return(sum(numerator - denominator))
## }(beta = dots[[1L]][[1L]])
```



Maximum partial-Likelihood estimation

```
fit.ML <- maxLik(pLogLik, start = c(beta = 0))
summary(fit.ML)
```

```
## -----
## Maximum Likelihood estimation
## Newton-Raphson maximisation, 2 iterations
## Return code 1: gradient close to zero
## Log-Likelihood: -3.327063
## 1 free parameters
## Estimates:
##      Estimate Std. error t value Pr(> t)
## beta  -0.5493    1.4179  -0.387  0.698
## -----
```

Beta is not significant different from 0 and 1

With the `coxph` function:

```
fit.cph <- coxph(Surv(time, failure) ~ x, data = dat)
summary(fit.cph)
```

```
## Call:
## coxph(formula = Surv(time, failure) ~ x, data = dat)
##
##      n= 5, number of events= 3
##
##      coef exp(coef) se(coef)      z Pr(>|z|)
## x -0.5493   0.5774   1.4179 -0.387   0.698
##
##      exp(coef) exp(-coef) lower .95 upper .95
## x   0.5774      1.732   0.03585   9.297
##
## Concordance= 0.5 (se = 0.202 )
## Rsquare= 0.029 (max possible= 0.743 )
## Likelihood ratio test= 0.15 on 1 df,  p=0.7
## Wald test               = 0.15 on 1 df,  p=0.7
## Score (logrank) test = 0.15 on 1 df,  p=0.7
```

Lower risk means longer time $\text{Beta} = -0.549 \pm 1.418 * 1.96 \exp(\text{Beta}) = 0.577$

```
confint(fit.cph)
```

```
##      2.5 %   97.5 %
## x -3.328286 2.229673
```

We can reproduce the Likelihood-ratio test:

```
LRT <- 2 * (fit.ML$maximum - pLogLik(0))
data.frame(LRT = LRT,
            pvalue = pchisq(LRT, df = 1, lower.tail = FALSE))
```

```
##      LRT    pvalue
## 1 0.1482688 0.7001953
```

The Wald test is already in the `maxLik` summary output.

A manually worked out, simple example: one continuous covariate

```
dat <- data.frame(time = c(6, 7, 10, 15, 19, 25),
                  event = c(1, 0, 1, 1, 0, 1),
                  age = c(67, 62, 34, 41, 46, 28))
```

```
fit <- coxph(Surv(time, event) ~ age, data = dat)
summary(fit)
```

```
## Call:
## coxph(formula = Surv(time, event) ~ age, data = dat)
##
##      n= 6, number of events= 4
##
##              coef exp(coef) se(coef)      z Pr(>|z|)
## age 0.07606    1.07903  0.07316 1.04    0.298
##
##      exp(coef) exp(-coef) lower .95 upper .95
## age      1.079      0.9268    0.9349    1.245
##
## Concordance= 0.7 (se = 0.237 )
## Rsquare= 0.209 (max possible= 0.76 )
## Likelihood ratio test= 1.41 on 1 df,  p=0.2
## Wald test              = 1.08 on 1 df,  p=0.3
## Score (logrank) test = 1.33 on 1 df,  p=0.2
```

We might express age in decades:

```
dat <- mutate(dat, age_dec = age / 10)
summary(coxph(Surv(time, event) ~ age_dec, data = dat))
```

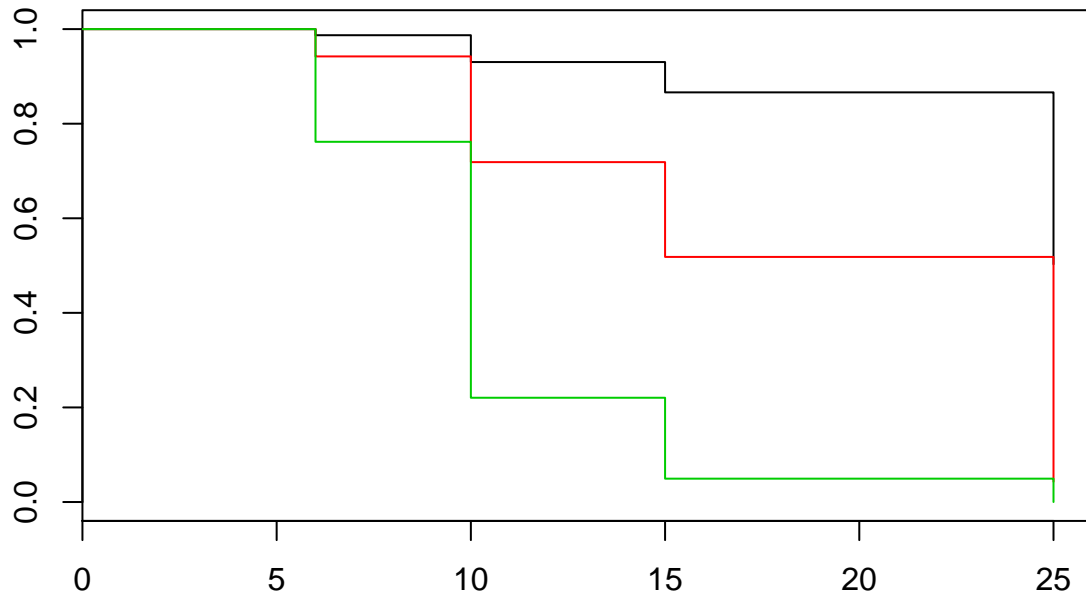
```
## Call:
## coxph(formula = Surv(time, event) ~ age_dec, data = dat)
##
##      n= 6, number of events= 4
##
##              coef exp(coef) se(coef)      z Pr(>|z|)
## age_dec 0.7606    2.1397  0.7316 1.04    0.298
##
##      exp(coef) exp(-coef) lower .95 upper .95
## age_dec      2.14      0.4674    0.51    8.976
##
## Concordance= 0.7 (se = 0.237 )
## Rsquare= 0.209 (max possible= 0.76 )
## Likelihood ratio test= 1.41 on 1 df,  p=0.2
## Wald test              = 1.08 on 1 df,  p=0.3
## Score (logrank) test = 1.33 on 1 df,  p=0.2
```

Try to do some predictions

```
pred <- survfit(fit, newdata = data.frame(age = c(20, 40, 60)))
pred
```

```
## Call: survfit(formula = fit, newdata = data.frame(age = c(20, 40, 60)))
##
##      n events median 0.95LCL 0.95UCL
## 1 6      4      NA      25      NA
## 2 6      4      25      10      NA
## 3 6      4      10       6      NA
```

```
plot(pred, col = 1:3)
```



Case study: the pharmacoSmoking dataset, DAP (Data Analysis Plan)

1. Model for TTR given: TRT, Age, EMPL, Gender, Race
 - Table of estimates
 - With commentary
2. Risk stratification/ Data segmentation:
 - Based on the model identify subject. At high risk Top 10
3. Predictions:
 - Median TTR
 - Survival ("6 months"/ covariates)
4. Compare TRT efficacy in FT vs NON-FT employment

Load the data

```
library(asauro)
d_raw <- pharmacoSmoking
head(d_raw)
```

```
##      id ttr relapse      grp age gender      race employment yearsSmoking
## 1  21 182      0  patchOnly 36  Male    white      ft          26
## 2 113  14      1  patchOnly 41  Male    white    other          27
## 3  39   5      1 combination 25 Female  white    other          12
## 4  80  16      1 combination 54  Male    white      ft          39
## 5  87   0      1 combination 45  Male    white    other          30
## 6  29 182      0 combination 43  Male  hispanic      ft          30
##      levelSmoking ageGroup2 ageGroup4 priorAttempts longestNoSmoke
## 1      heavy      21-49      35-49          0          0
## 2      heavy      21-49      35-49          3          90
## 3      heavy      21-49      21-34          3          21
## 4      heavy      50+      50-64          0          0
## 5      heavy      21-49      35-49          0          0
## 6      heavy      21-49      35-49          2         1825
```

```
table(d_raw$employment, useNA = "always")
```

```
##
##      ft other      pt <NA>
##      72   39      14     0
```

```
dat <- mutate(d_raw,
              employment = ifelse(employment == "ft", "ft", "other"),
              grp = relevel(grp, ref = "patchOnly"),
              race = ifelse(race == "white", "white",
                           ifelse(race == "black", "black", "other")))
table(dat$employment)
```

```
##
##      ft other
##      72   53
```

```
table(dat$race)
```

```
##
## black other white
##      38   10   77
```

Fit the Cox model

```
fit <- coxph(Surv(ttr, relapse) ~ grp + age + gender + employment + race, data = dat)
summary(fit)
```



```
## Call:
## coxph(formula = Surv(ttr, relapse) ~ grp + age + gender + employment +
##       race, data = dat)
##
##      n= 125, number of events= 89
##
##              coef exp(coef) se(coef)      z Pr(>|z|)
## grpcombination -0.60461   0.54629  0.21830 -2.770 0.005613 **
## age            -0.03575   0.96488  0.01069 -3.345 0.000821 ***
## genderMale     -0.04312   0.95780  0.23711 -0.182 0.855691
## employmentother 0.73991   2.09574  0.24132  3.066 0.002168 **
## raceother      -0.68519   0.50400  0.45634 -1.501 0.133232
## racewhite      -0.25824   0.77241  0.23268 -1.110 0.267065
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
##              exp(coef) exp(-coef) lower .95 upper .95
## grpcombination    0.5463    1.8305    0.3561    0.8380
## age                0.9649    1.0364    0.9449    0.9853
## genderMale         0.9578    1.0441    0.6018    1.5244
## employmentother    2.0957    0.4772    1.3060    3.3631
## raceother          0.5040    1.9841    0.2061    1.2327
## racewhite          0.7724    1.2947    0.4895    1.2187
##
## Concordance= 0.649  (se = 0.03 )
## Rsquare= 0.18  (max possible= 0.998 )
## Likelihood ratio test= 24.8  on 6 df,  p=4e-04
## Wald test              = 24.37  on 6 df,  p=4e-04
## Score (logrank) test = 25.07  on 6 df,  p=3e-04
```

We can change the contrasts as we see fit:

```
dat <- mutate(dat, grp = relevel(grp, ref = "patchOnly"))
fit <- update(fit)
summary(fit)
```

```
## Call:
## coxph(formula = Surv(ttr, relapse) ~ grp + age + gender + employment +
##       race, data = dat)
##
##      n= 125, number of events= 89
##
##              coef exp(coef) se(coef)      z Pr(>|z|)
## grpcombination -0.60461   0.54629  0.21830 -2.770 0.005613 **
## age            -0.03575   0.96488  0.01069 -3.345 0.000821 ***
## genderMale     -0.04312   0.95780  0.23711 -0.182 0.855691
## employmentother 0.73991   2.09574  0.24132  3.066 0.002168 **
## raceother      -0.68519   0.50400  0.45634 -1.501 0.133232
## racewhite      -0.25824   0.77241  0.23268 -1.110 0.267065
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
##              exp(coef) exp(-coef) lower .95 upper .95
```

```
## grpcombination      0.5463      1.8305      0.3561      0.8380
## age                 0.9649      1.0364      0.9449      0.9853
## genderMale          0.9578      1.0441      0.6018      1.5244
## employmentother     2.0957      0.4772      1.3060      3.3631
## raceother           0.5040      1.9841      0.2061      1.2327
## racewhite           0.7724      1.2947      0.4895      1.2187
##
## Concordance= 0.649 (se = 0.03 )
## Rsquare= 0.18 (max possible= 0.998 )
## Likelihood ratio test= 24.8 on 6 df, p=4e-04
## Wald test = 24.37 on 6 df, p=4e-04
## Score (logrank) test = 25.07 on 6 df, p=3e-04
```

Export the result table

```
broom::tidy(fit) %>%
  write_csv("coefficients_table.csv")
```

Data segmentation

Remove outcomes

```
d_new <- select(dat, -ttr, -relapse)
```

```
d_segmented <- d_new %>% mutate(risk_score = predict(fit, newdata = d_new, type = "lp"))
head(d_segmented)
```

```
##      id      grp age gender  race employment yearsSmoking levelSmoking
## 1  21  patchOnly 36  Male white      ft          26          heavy
## 2 113  patchOnly 41  Male white    other          27          heavy
## 3  39 combination 25 Female white    other          12          heavy
## 4  80 combination 54  Male white      ft          39          heavy
## 5  87 combination 45  Male white    other          30          heavy
## 6  29 combination 43  Male other      ft          30          heavy
##      ageGroup2 ageGroup4 priorAttempts longestNoSmoke risk_score
## 1      21-49      35-49           0           0 0.3681131
## 2      21-49      35-49           3           90 0.9292519
## 3      21-49      21-34           3           21 0.9398153
## 4       50+      50-64           0           0 -0.8800627
## 5      21-49      35-49           0           0 0.1816247
## 6      21-49      35-49           2          1825 -0.9137190
```

```
d_segmented %>% arrange(desc(risk_score)) %>% head(10)
```

```
##      id      grp age gender  race employment yearsSmoking levelSmoking
## 1   91  patchOnly 22 Female white    other          10          heavy
## 2   95  patchOnly 22 Female white    other           9          heavy
## 3   94  patchOnly 43 Female black    other          24          light
## 4   37  patchOnly 44 Female black    other          31          heavy
## 5   25  patchOnly 37 Female white    other          23          light
## 6   88  patchOnly 41 Female white    other          26          heavy
```

```
## 7 63 patchOnly 40 Male white other 22 heavy
## 8 62 patchOnly 49 Female black other 35 light
## 9 39 combination 25 Female white other 12 heavy
## 10 113 patchOnly 41 Male white other 27 heavy
## ageGroup2 ageGroup4 priorAttempts longestNoSmoke risk_score
## 1 21-49 21-34 3 2 1.6516891
## 2 21-49 21-34 2 3 1.6516891
## 3 21-49 35-49 2 3 1.1591080
## 4 21-49 35-49 2 2 1.1233545
## 5 21-49 35-49 5 1095 1.1153871
## 6 21-49 35-49 1 7 0.9723732
## 7 21-49 35-49 2 2 0.9650054
## 8 21-49 35-49 4 540 0.9445872
## 9 21-49 21-34 3 21 0.9398153
## 10 21-49 35-49 3 90 0.9292519
```

Predicting Median TTR and Survival (6 months | covariates)

```
predict_survival <- survfit(fit, new_data = d_new[1, ])
summary(predict_survival)
```

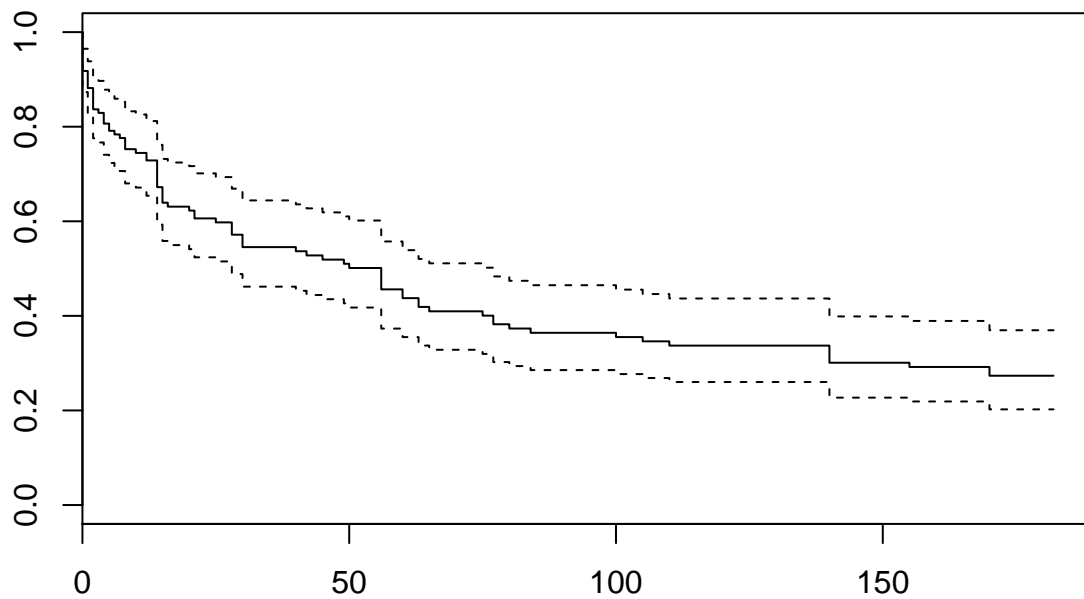
```
## Call: survfit(formula = fit, new_data = d_new[1, ])
##
## time n.risk n.event survival std.err lower 95% CI upper 95% CI
## 0 125 12 0.918 0.0234 0.873 0.965
## 1 113 5 0.882 0.0279 0.829 0.938
## 2 108 6 0.837 0.0325 0.775 0.903
## 3 102 1 0.829 0.0332 0.767 0.897
## 4 101 3 0.807 0.0351 0.741 0.878
## 5 98 2 0.791 0.0362 0.723 0.866
## 6 96 1 0.784 0.0368 0.715 0.859
## 7 95 1 0.776 0.0373 0.706 0.853
## 8 94 3 0.752 0.0388 0.680 0.833
## 10 91 1 0.745 0.0393 0.671 0.826
## 12 90 2 0.729 0.0402 0.654 0.812
## 14 88 7 0.672 0.0429 0.593 0.762
## 15 81 4 0.639 0.0441 0.558 0.732
## 16 77 1 0.631 0.0444 0.550 0.724
## 20 76 1 0.623 0.0447 0.541 0.717
## 21 75 2 0.606 0.0451 0.524 0.701
## 25 73 1 0.598 0.0453 0.515 0.693
## 28 72 3 0.572 0.0459 0.488 0.669
## 30 69 3 0.545 0.0463 0.462 0.644
## 40 66 1 0.537 0.0464 0.453 0.636
## 42 65 1 0.528 0.0465 0.444 0.627
## 45 64 1 0.519 0.0466 0.435 0.619
## 49 63 1 0.510 0.0466 0.426 0.610
## 50 62 1 0.501 0.0467 0.418 0.602
## 56 61 5 0.456 0.0467 0.373 0.557
## 60 56 2 0.437 0.0466 0.355 0.539
## 63 54 2 0.419 0.0464 0.337 0.520
## 65 52 1 0.410 0.0463 0.328 0.511
```

##	75	51	1	0.400	0.0461	0.320	0.502
##	77	50	2	0.382	0.0458	0.302	0.483
##	80	48	1	0.373	0.0456	0.294	0.474
##	84	47	1	0.364	0.0453	0.285	0.465
##	100	46	1	0.355	0.0451	0.277	0.455
##	105	45	1	0.346	0.0448	0.269	0.446
##	110	44	1	0.337	0.0446	0.260	0.437
##	140	43	4	0.301	0.0432	0.227	0.399
##	155	39	1	0.292	0.0429	0.219	0.389
##	170	38	2	0.273	0.0420	0.202	0.370

```
summary(predict_survival, times = 180)
```

```
## Call: survfit(formula = fit, new_data = d_new[1, ])
##
##   time n.risk n.event survival std.err lower 95% CI upper 95% CI
##   180     36     89   0.273   0.042   0.202      0.37
```

```
plot(predict_survival)
```



```
predict_survival
```

```
## Call: survfit(formula = fit, new_data = d_new[1, ])
##
```

```
##      n  events  median 0.95LCL 0.95UCL
##    125     89     56     28     77
```

```
individual_prediction <- function(grp, age, employment, race) {
  tibble(median_ttr = ..., surv_6_months = ...)
}
```

TRT.Efficacy in FT vs other

I Part

	Beta	SE	CI	P
FT	Beta FT			
OTH	Beta OTH			

```
library(tidyverse)
d_ft <- filter(dat, employment == "ft")
d_nft <- filter(dat, employment != "ft")
head(d_ft)
```

```
##   id ttr relapse      grp age gender  race employment yearsSmoking
## 1 21 182      0  patchOnly 36  Male white      ft           26
## 2 80 16      1  combination 54  Male white      ft           39
## 3 29 182      0  combination 43  Male other      ft           30
## 4 54 2       1  patchOnly 40  Female black      ft           25
## 5 70 0       1  patchOnly 38  Male black      ft           23
## 6 85 182      0  combination 51  Male black      ft           35
##   levelSmoking ageGroup2 ageGroup4 priorAttempts longestNoSmoke
## 1      heavy    21-49    35-49          0           0
## 2      heavy      50+    50-64          0           0
## 3      heavy    21-49    35-49          2          1825
## 4      heavy    21-49    35-49          4           7
## 5     light    21-49    35-49         10           90
## 6      heavy      50+    50-64          1           7
```

```
head(d_nft)
```

```
##   id ttr relapse      grp age gender  race employment yearsSmoking
## 1 113 14      1  patchOnly 41  Male white    other           27
## 2 39 5       1  combination 25  Female white    other           12
## 3 87 0       1  combination 45  Male white    other           30
## 4 16 14      1  patchOnly 66  Male black    other           54
## 5 35 77      1  patchOnly 78  Female black    other           56
## 6 84 12      1  patchOnly 64  Female black    other           30
##   levelSmoking ageGroup2 ageGroup4 priorAttempts longestNoSmoke
## 1      heavy    21-49    35-49          3           90
## 2      heavy    21-49    21-34          3           21
## 3      heavy    21-49    35-49          0           0
## 4      heavy      50+    65+          0           0
```

## 5	light	50+	65+	10	15
## 6	heavy	50+	50-64	12	365

```
coxph(Surv(ttr, relapse) ~ grp, data = d_ft) %>% summary()
```

```
## Call:
## coxph(formula = Surv(ttr, relapse) ~ grp, data = d_ft)
##
## n= 72, number of events= 49
##
##               coef exp(coef) se(coef)      z Pr(>|z|)
## grpcombination -0.3580   0.6991  0.2891 -1.238   0.216
##
##               exp(coef) exp(-coef) lower .95 upper .95
## grpcombination   0.6991      1.43   0.3967   1.232
##
## Concordance= 0.551 (se = 0.038 )
## Rsquare= 0.021 (max possible= 0.995 )
## Likelihood ratio test= 1.55 on 1 df,  p=0.2
## Wald test               = 1.53 on 1 df,  p=0.2
## Score (logrank) test = 1.55 on 1 df,  p=0.2
```

```
d_ft$grp <- relevel(d_ft$grp, ref = "combination")
coxph(Surv(ttr, relapse) ~ grp, data = d_ft) %>% summary()
```

```
## Call:
## coxph(formula = Surv(ttr, relapse) ~ grp, data = d_ft)
##
## n= 72, number of events= 49
##
##               coef exp(coef) se(coef)      z Pr(>|z|)
## grppatchOnly 0.3580   1.4304  0.2891 1.238   0.216
##
##               exp(coef) exp(-coef) lower .95 upper .95
## grppatchOnly   1.43   0.6991   0.8117   2.521
##
## Concordance= 0.551 (se = 0.038 )
## Rsquare= 0.021 (max possible= 0.995 )
## Likelihood ratio test= 1.55 on 1 df,  p=0.2
## Wald test               = 1.53 on 1 df,  p=0.2
## Score (logrank) test = 1.55 on 1 df,  p=0.2
```

```
coxph(Surv(ttr, relapse) ~ grp, data = d_nft) %>% summary()
```

```
## Call:
## coxph(formula = Surv(ttr, relapse) ~ grp, data = d_nft)
##
## n= 53, number of events= 40
##
##               coef exp(coef) se(coef)      z Pr(>|z|)
## grpcombination -0.9847   0.3736  0.3288 -2.995 0.00274 **
## ---
```

```
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
##               exp(coef) exp(-coef) lower .95 upper .95
## grpcombination    0.3736      2.677    0.1961    0.7115
##
## Concordance= 0.624  (se = 0.039 )
## Rsquare= 0.16   (max possible= 0.994 )
## Likelihood ratio test= 9.22  on 1 df,   p=0.002
## Wald test          = 8.97  on 1 df,   p=0.003
## Score (logrank) test = 9.64  on 1 df,   p=0.002
```

II Part

H0: Beta FT = Beta OTH? <-> H0: interact beta = 0 p-value = ?

Interaction Test: beta in interaction term between trt and emp

```
fit_int <- coxph(Surv(ttr, relapse) ~ grp + employment + grp:employment, data = dat)
summary(fit_int)
```

```
## Call:
## coxph(formula = Surv(ttr, relapse) ~ grp + employment + grp:employment,
##       data = dat)
##
##    n= 125, number of events= 89
##
##               coef exp(coef) se(coef)      z Pr(>|z|)
## grpcombination   -0.3554    0.7009  0.2889 -1.230  0.2186
## employmentother    0.6361    1.8892  0.2802  2.270  0.0232
## grpcombination:employmentother -0.6587    0.5175  0.4343 -1.517  0.1294
##
## grpcombination
## employmentother      *
## grpcombination:employmentother
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
##               exp(coef) exp(-coef) lower .95 upper .95
## grpcombination    0.7009    1.4268    0.3978    1.235
## employmentother    1.8892    0.5293    1.0908    3.272
## grpcombination:employmentother 0.5175    1.9323    0.2209    1.212
##
## Concordance= 0.601  (se = 0.03 )
## Rsquare= 0.099   (max possible= 0.998 )
## Likelihood ratio test= 12.99  on 3 df,   p=0.005
## Wald test          = 14.22  on 3 df,   p=0.003
## Score (logrank) test = 15.13  on 3 df,   p=0.002
```

p-value = 0.13, so it not significant different