

Cox regression

A manually worked out, simple example: two groups

Load libraries

```
library(tidyverse)
```

```
## -- Attaching packages ----- tidyverse 1.2.1 --

## v ggplot2 3.2.0      v purrr   0.3.2
## v tibble  2.1.3      v dplyr  0.8.3
## v tidyr   0.8.3      v stringr 1.4.0
## v readr   1.3.1      v forcats 0.4.0

## Warning: package 'dplyr' was built under R version 3.6.1

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

```
library(maxLik)
```

```
## Warning: package 'maxLik' was built under R version 3.6.1

## Loading required package: miscTools

## Warning: package 'miscTools' was built under R version 3.6.1

##
## 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)
```

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)
```

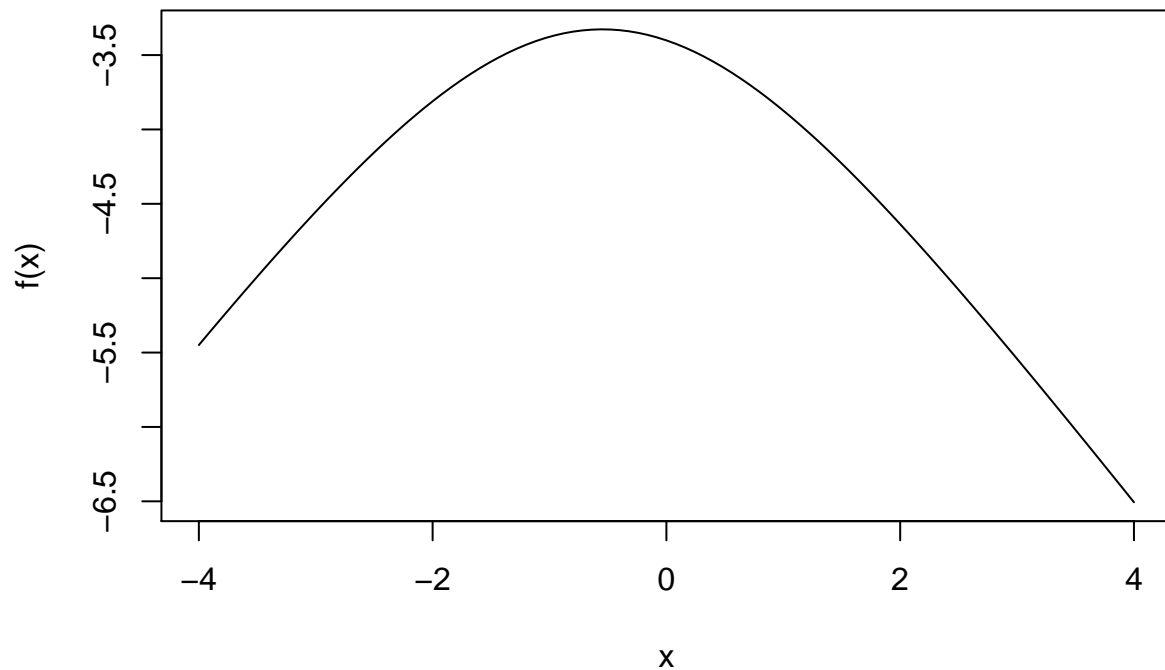
Partial log-likelihood function

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)) {
    risk_set <- subset(dat, time >= dat.events[j, "time"])
    theta_j <- with(risk_set, exp(x * beta)) # within the risk set, we compute the function for each rat
    denominator[j] <- log(sum(theta_j))
  }
  #with log, we only need to do sum, not product, to easier computation
  return(sum(numerator - denominator))
}
```

We can plot it:

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



Maximum partial-Likelihood estimation

interpretation:

x_i

- 0: normal sleep pattern
- 1: sleep deprived

$$h_i(t) = h_0(t) \exp(x_i B) \quad \hat{B} = -0.55 (SE = 1.4)$$

Hazard ratio: (between 2 group)

$$\frac{h_{SD}(t) = h_0 \exp(1 \cdot -0.55)}{h_{NSD}(t) = h_0 \exp(0 \cdot -0.55)} = \exp(-0.55)$$

```
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
## -----
```

```
#hazard ratio
exp(-0.55)
```

```
## [1] 0.5769498
```

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 )
## 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
```

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 )
## 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 )
## 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
```

Case study: the pharmacoSmoking dataset

Load the data

```
library(asaur)
dat <- pharmacoSmoking
head(dat)
```

```
##    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
```

grp is not 0,1. R would transform it to be 0,1 (alph ordering combination:0, patchOnly:1), so that we see the risk $H_1 > H_0$, the risk in pathOnly is higher so it means the time is shorter. P-value is small, so there's a significant difference.

```
summary(coxph(Surv(ttr,relapse)~grp,data=dat))
```

```
## Call:
## coxph(formula = Surv(ttr, relapse) ~ grp, data = dat)
##
## n= 125, number of events= 89
##
##              coef exp(coef) se(coef)  z Pr(>|z|)
## grppatchOnly 0.6050    1.8313  0.2161 2.8 0.00511 **
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
##              exp(coef) exp(-coef) lower .95 upper .95
## grppatchOnly    1.831    0.5461    1.199    2.797
##
## Concordance= 0.581 (se = 0.027 )
## Likelihood ratio test= 7.99 on 1 df,  p=0.005
## Wald test              = 7.84 on 1 df,  p=0.005
## Score (logrank) test = 8.07 on 1 df,  p=0.004
```

Fit the Cox model

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

```
## Call:
## coxph(formula = Surv(ttr, relapse) ~ grp + age + gender + priorAttempts,
## data = dat)
##
## n= 125, number of events= 89
##
##              coef exp(coef) se(coef)  z Pr(>|z|)
## grppatchOnly 0.5656340 1.7605636 0.2181634 2.593 0.00952 **
```

```
## age          -0.0220948  0.9781475  0.0097572 -2.264  0.02355 *
## genderMale   -0.1215514  0.8855455  0.2334349 -0.521  0.60257
## priorAttempts 0.0002078  1.0002079  0.0010898  0.191  0.84876
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
##              exp(coef) exp(-coef) lower .95 upper .95
## grppatchOnly    1.7606    0.5680    1.1480    2.700
## age             0.9781    1.0223    0.9596    0.997
## genderMale      0.8855    1.1292    0.5604    1.399
## priorAttempts   1.0002    0.9998    0.9981    1.002
##
## Concordance= 0.623 (se = 0.031 )
## Likelihood ratio test= 14.14 on 4 df,  p=0.007
## Wald test              = 13.87 on 4 df,  p=0.008
## Score (logrank) test = 14.12 on 4 df,  p=0.007
```

We can change the contrasts as we see fit:

```
dat <- mutate(dat, grp = relevel(grp, ref = "patchOnly")) #change patchOnly to 0
fit <- update(fit)
summary(fit)
```

```
## Call:
## coxph(formula = Surv(ttr, relapse) ~ grp + age + gender + priorAttempts,
##       data = dat)
##
##      n= 125, number of events= 89
##
##              coef  exp(coef)    se(coef)      z Pr(>|z|)
## grpcombination -0.5656340  0.5679999  0.2181634 -2.593  0.00952 **
## age            -0.0220948  0.9781475  0.0097572 -2.264  0.02355 *
## genderMale     -0.1215514  0.8855455  0.2334349 -0.521  0.60257
## priorAttempts  0.0002078  1.0002079  0.0010898  0.191  0.84876
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
##              exp(coef) exp(-coef) lower .95 upper .95
## grpcombination    0.5680    1.7606    0.3704    0.8711
## age               0.9781    1.0223    0.9596    0.9970
## genderMale        0.8855    1.1292    0.5604    1.3993
## priorAttempts     1.0002    0.9998    0.9981    1.0023
##
## Concordance= 0.623 (se = 0.031 )
## Likelihood ratio test= 14.14 on 4 df,  p=0.007
## Wald test              = 13.87 on 4 df,  p=0.008
## Score (logrank) test = 14.12 on 4 df,  p=0.007
```

encoding for categorical varianve more then 2 category (we can also change the reverance)

```
summary(coxph(Surv(ttr,relapse)~employment,data=dat)) #default fulltime is reference (decide alphabetic
```

```
## Call:
```

```
## coxph(formula = Surv(ttr, relapse) ~ employment, data = dat)
##
##   n= 125, number of events= 89
##
##               coef exp(coef) se(coef)      z Pr(>|z|)
## employmentother 0.1982    1.2192  0.2371 0.836   0.403
## employmentpt    0.4500    1.5683  0.3229 1.394   0.163
##
##               exp(coef) exp(-coef) lower .95 upper .95
## employmentother    1.219    0.8202   0.7661   1.940
## employmentpt      1.568    0.6376   0.8328   2.953
##
## Concordance= 0.541 (se = 0.028 )
## Likelihood ratio test= 2.06 on 2 df,  p=0.4
## Wald test               = 2.17 on 2 df,  p=0.3
## Score (logrank) test = 2.2 on 2 df,  p=0.3
```

```
#if we want to change the reference
dat1<-mutate(dat,employment =relevel(employment, ref="pt"))
summary(coxph(Surv(ttr,relapse)~employment,data=dat1))
```

```
## Call:
## coxph(formula = Surv(ttr, relapse) ~ employment, data = dat1)
##
##   n= 125, number of events= 89
##
##               coef exp(coef) se(coef)      z Pr(>|z|)
## employmentft   -0.4500    0.6376  0.3229 -1.394   0.163
## employmentother -0.2518    0.7774  0.3455 -0.729   0.466
##
##               exp(coef) exp(-coef) lower .95 upper .95
## employmentft    0.6376    1.568   0.3386   1.201
## employmentother 0.7774    1.286   0.3949   1.530
##
## Concordance= 0.541 (se = 0.028 )
## Likelihood ratio test= 2.06 on 2 df,  p=0.4
## Wald test               = 2.17 on 2 df,  p=0.3
## Score (logrank) test = 2.2 on 2 df,  p=0.3
```

Case study: the lung cancer dataset

Load the data

```
library(survival)

dat <- lung
dat$delta<-dat$status-1
dat$S <-with(dat,Surv(time/365.25,delta))
#equivalent as Surv(dat$time,dat$delta)
#for S it store 2 value, but when we print it out when it's censoring it put a plus in the end
```



```
#rescale time to year
head(dat)
```

```
##   inst time status age sex ph.ecog ph.karno pat.karno meal.cal wt.loss
## 1    3  306      2  74  1      1      90      100     1175      NA
## 2    3  455      2  68  1      0      90      90     1225      15
## 3    3 1010      1  56  1      0      90      90        NA      15
## 4    5  210      2  57  1      1      90      60     1150      11
## 5    1  883      2  60  1      0     100      90        NA       0
## 6   12 1022      1  74  1      1      50      80      513       0
##   delta      S
## 1      1 0.8377823
## 2      1 1.2457221
## 3      0 2.7652293+
## 4      1 0.5749487
## 5      1 2.4175222
## 6      0 2.7980835+
```

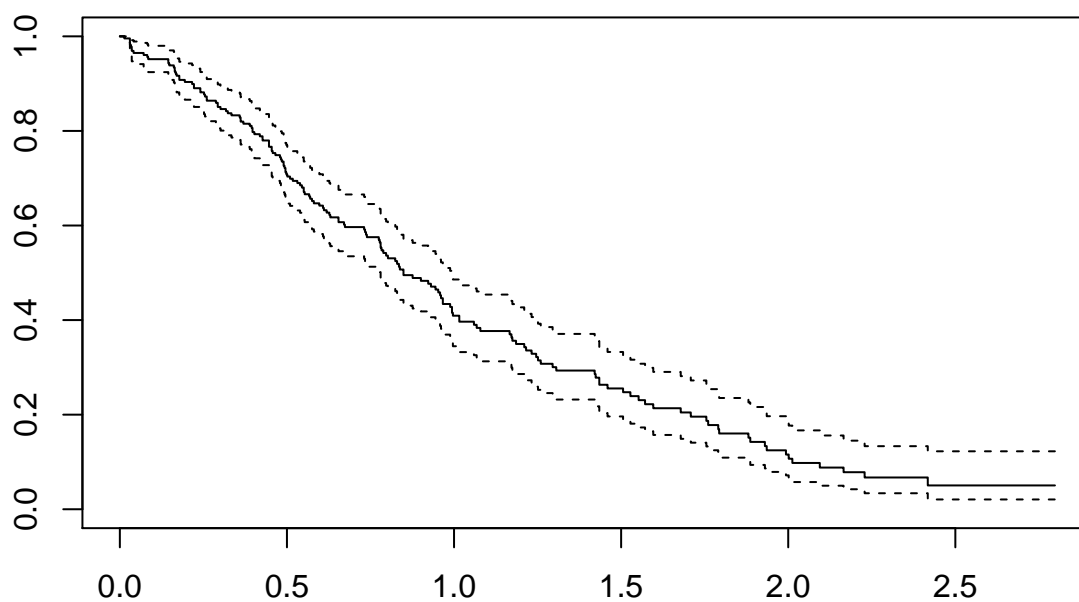
exercise:

Q1: median survival and confidence interval Q2: survival of men vs woman

- median survival with each group
- test & the diff
- HR?

Q3: is self-evaluate karno score equivalent to physician's score?

```
#Q1 medium survival
fit.KM <-survfit(S~1, data=dat) #survival curve go to 0, after 2.5 years almost every patients dead
plot(fit.KM)
```



```
fit.KM # we can get the median and confidence interval
```

```
## Call: survfit(formula = S ~ 1, data = dat)
##
##      n  events  median 0.95LCL 0.95UCL
## 228.000 165.000   0.849   0.780   0.994
```

```
#Q2
table(dat$sex, useNA = "always")# report also missing
```

```
##
##      1      2 <NA>
## 138    90      0
```

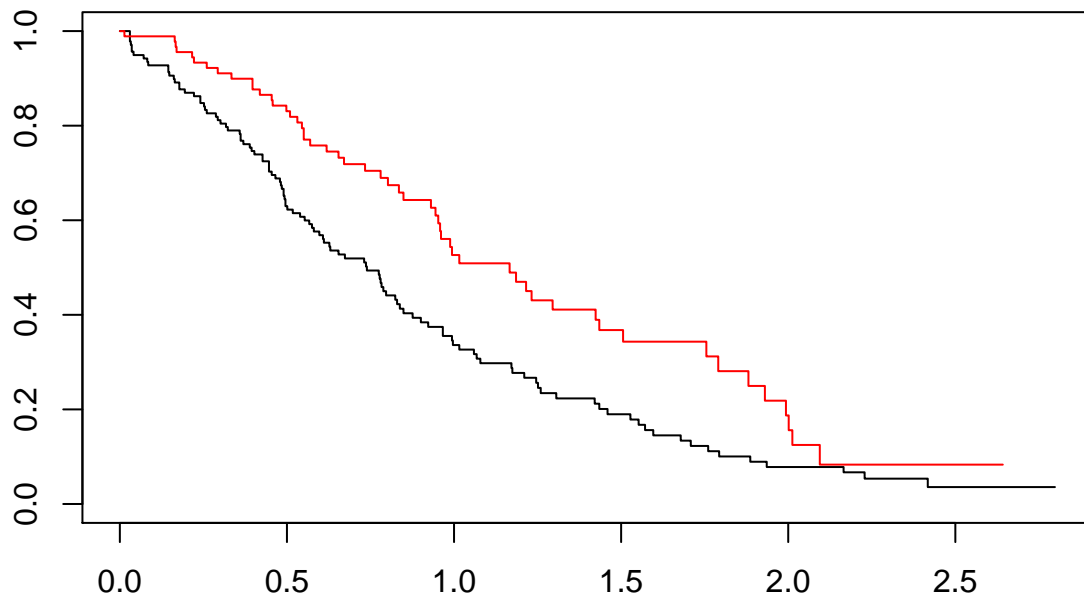
```
dat$sex <- factor(dat$sex, levels=1:2, labels=c("m", "f"))
table(dat$sex, useNA = "always")
```

```
##
##      m      f <NA>
## 138    90      0
```

```
fit.KM <- survfit(S~sex, data=dat)
fit.KM
```

```
## Call: survfit(formula = S ~ sex, data = dat)
##
##           n events median 0.95LCL 0.95UCL
## sex=m 138     112  0.739   0.580   0.849
## sex=f  90      53  1.166   0.953   1.506
```

```
plot(fit.KM,col=1:2) #black:m red:female
```



```
#survival for man is worse then women
survdif(S~sex, data=dat)
```

```
## Call:
## survdiff(formula = S ~ sex, data = dat)
##
##           N Observed Expected (O-E)^2/E (O-E)^2/V
## sex=m 138      112      91.6      4.55      10.3
## sex=f  90       53      73.4      5.68      10.3
##
##   Chisq= 10.3  on 1 degrees of freedom, p= 0.001
```

```
#there's the significance difference
```

```
#HR?
summary(coxph(S~sex,data = dat))
```

```
## Call:
## coxph(formula = S ~ sex, data = dat)
##
## n= 228, number of events= 165
##
##      coef exp(coef) se(coef)      z Pr(>|z|)
## sexf -0.5310    0.5880   0.1672 -3.176  0.00149 **
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
##      exp(coef) exp(-coef) lower .95 upper .95
## sexf      0.588      1.701    0.4237    0.816
##
## Concordance= 0.579 (se = 0.021 )
## Likelihood ratio test= 10.63 on 1 df,  p=0.001
## Wald test               = 10.09 on 1 df,  p=0.001
## Score (logrank) test = 10.33 on 1 df,  p=0.001
```

*#risk for man is higher, it's significantly different
#confirm the result of the survival test*

Q3:

`summary(dat)` *# supposedly they should be similar, self one has larger range*

```
##      inst      time      status      age      sex
## Min.   : 1.00   Min.   : 5.0   Min.   :1.000   Min.   :39.00   m:138
## 1st Qu.: 3.00   1st Qu.: 166.8   1st Qu.:1.000   1st Qu.:56.00   f: 90
## Median :11.00   Median : 255.5   Median :2.000   Median :63.00
## Mean   :11.09   Mean   : 305.2   Mean   :1.724   Mean   :62.45
## 3rd Qu.:16.00   3rd Qu.: 396.5   3rd Qu.:2.000   3rd Qu.:69.00
## Max.   :33.00   Max.   :1022.0   Max.   :2.000   Max.   :82.00
## NA's    :1
##      ph.ecog      ph.karno      pat.karno      meal.cal
## Min.   :0.0000   Min.   : 50.00   Min.   : 30.00   Min.   : 96.0
## 1st Qu.:0.0000   1st Qu.: 75.00   1st Qu.: 70.00   1st Qu.: 635.0
## Median :1.0000   Median : 80.00   Median : 80.00   Median : 975.0
## Mean   :0.9515   Mean   : 81.94   Mean   : 79.96   Mean   : 928.8
## 3rd Qu.:1.0000   3rd Qu.: 90.00   3rd Qu.: 90.00   3rd Qu.:1150.0
## Max.   :3.0000   Max.   :100.00   Max.   :100.00   Max.   :2600.0
## NA's    :1      NA's    :1      NA's    :3      NA's    :47
##      wt.loss      delta
## Min.   : -24.000   Min.   : 0.0000
## 1st Qu.:  0.000   1st Qu.: 0.0000
## Median :  7.000   Median : 1.0000
## Mean   :  9.832   Mean   : 0.7237
## 3rd Qu.: 15.750   3rd Qu.: 1.0000
## Max.   : 68.000   Max.   : 1.0000
## NA's    :14
##      S.time      S.status
## Min.   :0.0136893   Min.   :0.0000000
## 1st Qu.:0.4565366   1st Qu.:0.0000000
## Median :0.6995209   Median :1.0000000
```

```
## Mean :0.8356809 Mean :0.7236842
## 3rd Qu.:1.0855578 3rd Qu.:1.0000000
## Max. :2.7980835 Max. :1.0000000
##
```

Doctor HR: 0.9837 (0.9725,0.995) Patient HR: 0.980 (0.970,0.991) so we can kind of assume that they are redundant

- it's better to reverse the ratio. because saying it 2 times more is better then saying it's 0.5 times: $1/0.983=1.017$
- it's not significant 1.017 to explain so we can also do some transformation
- we can now explain now the observation is 1.17 per 10 units **decrease**(we use the $\exp(-\text{coef})$ as reference) of the score

```
#summary(coxph(S~ph.karno,data=dat))
#summary(coxph(S~pat.karno,data=dat))
summary(coxph(S~I(pat.karno/10),data=dat))
```

```
## Call:
## coxph(formula = S ~ I(pat.karno/10), data = dat)
##
## n= 225, number of events= 162
## (3 observations deleted due to missingness)
##
##               coef exp(coef) se(coef)      z Pr(>|z|)
## I(pat.karno/10) -0.19850    0.81996  0.05467 -3.631 0.000282 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
##               exp(coef) exp(-coef) lower .95 upper .95
## I(pat.karno/10)      0.82      1.22    0.7366    0.9127
##
## Concordance= 0.607 (se = 0.025 )
## Likelihood ratio test= 12.47 on 1 df,  p=4e-04
## Wald test               = 13.18 on 1 df,  p=3e-04
## Score (logrank) test = 13.23 on 1 df,  p=3e-04
```

what if we try another model with both?

```
summary(coxph(S~I(pat.karno/10)+I(ph.karno/10),data=dat))
```

```
## Call:
## coxph(formula = S ~ I(pat.karno/10) + I(ph.karno/10), data = dat)
##
## n= 224, number of events= 161
## (4 observations deleted due to missingness)
##
##               coef exp(coef) se(coef)      z Pr(>|z|)
## I(pat.karno/10) -0.16275    0.84980  0.06372 -2.554  0.0107 *
## I(ph.karno/10)  -0.07404    0.92863  0.06959 -1.064  0.2873
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

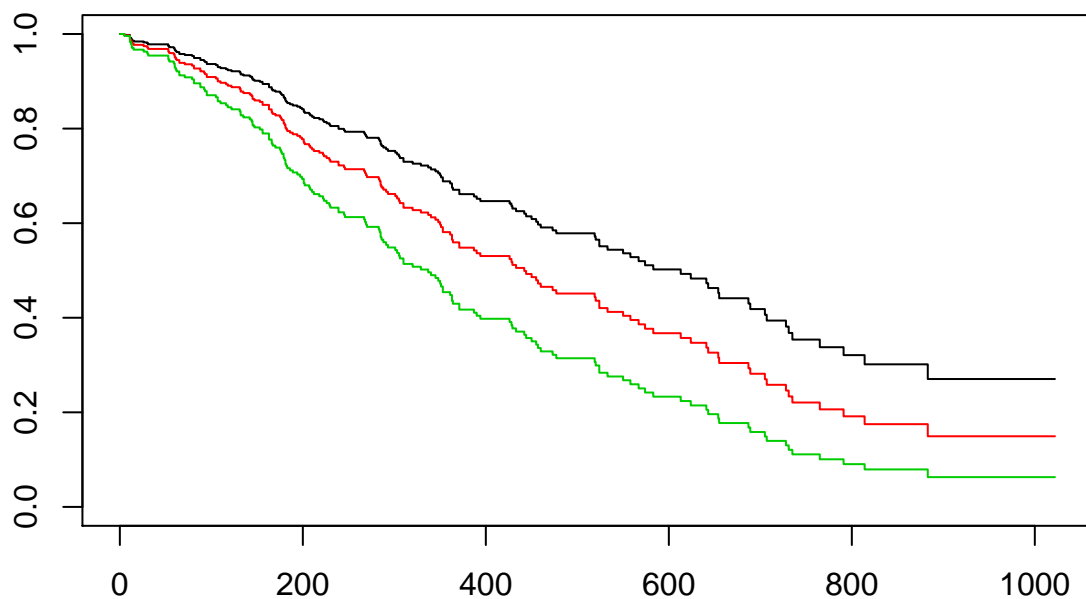
```
##
##               exp(coef) exp(-coef) lower .95 upper .95
## I(pat.karno/10)  0.8498      1.177   0.7500   0.9629
## I(ph.karno/10)   0.9286      1.077   0.8102   1.0643
##
## Concordance= 0.616 (se = 0.025 )
## Likelihood ratio test= 13.3 on 2 df,  p=0.001
## Wald test            = 14.01 on 2 df,  p=9e-04
## Score (logrank) test = 14.13 on 2 df,  p=9e-04
```

*#taking each by each, the effect is very close
 #but if we put them together, the doctor one drop
 #the model is telling us it's redundant to one and another
 #the pvalue if we fix one, the another would not have much impact on the model*

Cox regression: predictions

generally we don't use it in practice

```
fit.cph <- coxph(Surv(time, status) ~ age, data = dat)
pred.cph <- survfit(fit.cph, newdata = data.frame(age = c(20,40,60)))
plot(pred.cph, col = 1:3)
```



```
#one cure for each indivisual datadrame
```

```
print(pred.cph)# base on cox model fix, we have a median of the point estimator
```

```
## Call: survfit(formula = fit.cph, newdata = data.frame(age = c(20, 40,  
##      60)))
```

```
##
```

```
##      n events median 0.95LCL 0.95UCL
```

```
## 1 228      165      613      363      NA
```

```
## 2 228      165      442      329      705
```

```
## 3 228      165      337      288      371
```