Variable Selection

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```
rm(list=ls())
library(dplyr)
##
## Attaching package: 'dplyr'
## The following objects are masked from 'package:stats':
##
##
       filter, lag
## The following objects are masked from 'package:base':
##
##
       intersect, setdiff, setequal, union
library(survival)
library(survMisc)
data <- read.csv("../data/survDataCleaned.csv")</pre>
Which race variable should we use?
data.raceCleaned <- data %>% select(-race_collapsed)
data.raceCollapsed <- data %>% select(-race_cleaned)
```

Variable Selection.

Fit Cox-PH models for both the data.raceCleaned and data.raceCollapsed.

```
data.raceCleaned$race_cleaned <- relevel(factor(data.raceCleaned$race_cleaned), ref = "white")</pre>
data.raceCleaned$FIGO <- relevel(factor(data.raceCleaned$FIGO), ref = "Stage I")</pre>
fullCoxMod <- coxph(Surv(time, delta) ~ size.intermediate + factor(race_cleaned) + factor(FIGO)+age_at_
fullCoxMod
## Call:
## coxph(formula = Surv(time, delta) ~ size.intermediate + factor(race_cleaned) +
##
       factor(FIGO) + age_at_diagnosis, data = data.raceCleaned)
##
##
                                             coef exp(coef) se(coef)
## size.intermediate
                                         0.151974 1.164130 0.147632 1.029
                                        -0.914762 0.400612 0.413712 -2.211
## factor(race_cleaned)asian
## factor(race_cleaned)black
                                         0.280853 1.324259 0.221897 1.266
                                         0.093532 1.098046 0.384043 0.244
## factor(race_cleaned)hispanic
```

factor(race_cleaned)unreported/other -0.896446 0.408017 0.383582 -2.337

```
## factor(FIGO)Stage II
                                         0.578492 1.783347 0.660023 0.876
## factor(FIGO)Stage III
                                        1.546288 4.694015 0.581007 2.661
## factor(FIGO)Stage IV
                                        2.019321 7.533212 0.591642 3.413
## age_at_diagnosis
                                        0.021200 1.021426 0.004712 4.499
                                               p
## size.intermediate
                                        0.303286
## factor(race cleaned)asian
                                        0.027028
## factor(race_cleaned)black
                                        0.205624
## factor(race_cleaned)hispanic
                                        0.807583
## factor(race_cleaned)unreported/other 0.019437
## factor(FIGO)Stage II
                                        0.380773
## factor(FIGO)Stage III
                                        0.007782
## factor(FIGO)Stage IV
                                        0.000642
## age_at_diagnosis
                                        6.83e-06
##
## Likelihood ratio test=78.46 on 9 df, p=3.264e-13
## n= 598, number of events= 359
data.raceCollapsed$race_cleaned <- relevel(factor(data.raceCollapsed$race_collapsed), ref = "white")</pre>
data.raceCollapsed$FIGO <- relevel(factor(data.raceCollapsed$FIGO), ref = "Stage I")</pre>
fullCoxMod.r <- coxph(Surv(time, delta) ~ size.intermediate + factor(race_cleaned) + factor(FIGO)+age_a
fullCoxMod.r
## Call:
## coxph(formula = Surv(time, delta) ~ size.intermediate + factor(race_cleaned) +
       factor(FIGO) + age_at_diagnosis, data = data.raceCollapsed)
##
##
                                             coef exp(coef) se(coef)
## size.intermediate
                                         0.153985 1.166473 0.147402 1.045
## factor(race_cleaned)asian
                                        -0.913906 0.400955 0.413715 -2.209
## factor(race_cleaned)black/hisp.
                                         0.232315 1.261517
                                                             0.195233 1.190
## factor(race_cleaned)other/unreported -0.896253  0.408096
                                                             0.383577 -2.337
## factor(FIGO)Stage II
                                         0.583044 1.791483
                                                             0.659915 0.884
## factor(FIGO)Stage III
                                         1.546116 4.693204
                                                             0.581015 2.661
## factor(FIGO)Stage IV
                                         2.017950 7.522891
                                                             0.591655 3.411
                                         0.021367 1.021597 0.004697 4.549
## age_at_diagnosis
##
                                        0.296180
## size.intermediate
## factor(race cleaned)asian
                                        0.027173
## factor(race_cleaned)black/hisp.
                                        0.234071
## factor(race_cleaned)other/unreported 0.019462
## factor(FIGO)Stage II
                                        0.376959
## factor(FIGO)Stage III
                                        0.007790
## factor(FIGO)Stage IV
                                        0.000648
## age_at_diagnosis
                                        5.39e-06
##
## Likelihood ratio test=78.27 on 8 df, p=1.089e-13
## n= 598, number of events= 359
```

Above is proof that collapsing the race category doesn't improve or make inference any better. I will use the data.raceCleaned variable going forward.

Stepwise AIC Selection.

```
# import package
library(MASS)
##
## Attaching package: 'MASS'
## The following object is masked from 'package:dplyr':
##
##
      select
best.model.aic <- stepAIC(fullCoxMod, direction="both",</pre>
       k=2, trace=1)
## Start: AIC=4247.61
## Surv(time, delta) ~ size.intermediate + factor(race_cleaned) +
##
      factor(FIGO) + age_at_diagnosis
##
##
                         Df
                               AIC
## - size.intermediate
                          1 4246.6
## <none>
                            4247.6
## - factor(race_cleaned) 4 4255.3
## - age_at_diagnosis
                          1 4265.7
## - factor(FIGO)
                          3 4279.7
## Step: AIC=4246.65
## Surv(time, delta) ~ factor(race_cleaned) + factor(FIGO) + age_at_diagnosis
##
##
                         Df
                               AIC
## <none>
                            4246.6
## + size.intermediate
                          1 4247.6
## - factor(race_cleaned) 4 4254.9
## - age_at_diagnosis
                          1 4265.7
## - factor(FIGO)
                          3 4279.5
# Display the best model obtained by AIC
best.model.aic
## Call:
## coxph(formula = Surv(time, delta) ~ factor(race_cleaned) + factor(FIGO) +
      age_at_diagnosis, data = data.raceCleaned)
##
##
                                            coef exp(coef)
                                                            se(coef)
## factor(race_cleaned)asian
                                       -0.928315 0.395219 0.413392 -2.246
## factor(race_cleaned)black
                                        0.287141 1.332613 0.221873 1.294
## factor(race_cleaned)hispanic
                                        0.085670 1.089447 0.383921 0.223
## factor(race_cleaned)unreported/other -0.907432 0.403559 0.383402 -2.367
## factor(FIGO)Stage II
                                        0.579955 1.785958 0.660003 0.879
## factor(FIGO)Stage III
                                       1.544113 4.683814 0.580996 2.658
## factor(FIGO)Stage IV
                                      2.035096 7.652983 0.591420 3.441
## age_at_diagnosis
                                       0.021626 1.021861 0.004692 4.609
##
                                              р
## factor(race_cleaned)asian
                                       0.024729
## factor(race_cleaned)black
                                       0.195606
## factor(race_cleaned)hispanic
                                       0.823422
```

```
## factor(race_cleaned)unreported/other 0.017943
## factor(FIGO)Stage II
                                        0.379556
## factor(FIGO)Stage III
                                        0.007868
## factor(FIGO)Stage IV
                                        0.000579
## age_at_diagnosis
                                         4.04e-06
##
## Likelihood ratio test=77.43 on 8 df, p=1.608e-13
## n= 598, number of events= 359
Ok, AIC stepwise selection only selected race_cleaned, FIGO and age_at_diagnosis.
Stepwise BIC Selection.
n <- nrow(data) # number of rows</pre>
best.model.bic <- stepAIC(fullCoxMod, direction="both",</pre>
                          k=log(n), trace=1)
## Start: AIC=4287.15
## Surv(time, delta) ~ size.intermediate + factor(race_cleaned) +
##
       factor(FIGO) + age_at_diagnosis
##
##
                          Df
                                AIC
## - factor(race cleaned) 4 4277.3
## - size.intermediate
                           1 4281.8
## <none>
                             4287.2
## - age_at_diagnosis
                           1 4300.8
## - factor(FIGO)
                           3 4306.0
##
## Step: AIC=4277.28
## Surv(time, delta) ~ size.intermediate + factor(FIGO) + age_at_diagnosis
##
                                AIC
                          Df
## - size.intermediate
                           1 4272.5
                             4277.3
## <none>
## + factor(race_cleaned) 4 4287.2
## - age_at_diagnosis
                           1 4293.9
## - factor(FIGO)
                           3 4298.4
##
## Step: AIC=4272.49
## Surv(time, delta) ~ factor(FIGO) + age_at_diagnosis
##
##
                          Df
                                AIC
## <none>
                             4272.5
## + size.intermediate
                           1 4277.3
## + factor(race_cleaned) 4 4281.8
## - age_at_diagnosis
                           1 4290.5
## - factor(FIGO)
                           3 4294.6
best.model.bic
## Call:
## coxph(formula = Surv(time, delta) ~ factor(FIGO) + age_at_diagnosis,
```

p

coef exp(coef) se(coef)

data = data.raceCleaned)

##

BIC selected FIGO and age_at_diagonosis to obtain the best model.

AFT Models.

Exponential AFT.

```
full.expAFT <- survreg(Surv(time, delta) ~ size.intermediate + factor(race_cleaned) +</pre>
    factor(FIGO) + age_at_diagnosis, data = data.raceCleaned, dist = "exponential")
expAFT.aic <- stepAIC(full.expAFT, direction = "both", k=2, trace=1)</pre>
## Start: AIC=6673.96
## Surv(time, delta) ~ size.intermediate + factor(race_cleaned) +
##
       factor(FIGO) + age_at_diagnosis
##
##
                           Df
                                 AIC
## <none>
                              6674.0
## - size.intermediate
                            1 6674.9
## - factor(race_cleaned) 4 6687.1
## - age_at_diagnosis
                            1 6692.7
## - factor(FIGO)
                            3 6717.2
expAFT.aic
## survreg(formula = Surv(time, delta) ~ size.intermediate + factor(race_cleaned) +
       factor(FIGO) + age_at_diagnosis, data = data.raceCleaned,
##
##
       dist = "exponential")
##
## Coefficients:
##
                             (Intercept)
                                                             size.intermediate
##
                              11.5786835
                                                                     -0.2600141
              {\tt factor(race\_cleaned)asian}
##
                                                     factor(race_cleaned)black
##
                               0.9821891
                                                                     -0.3971982
           factor(race_cleaned)hispanic factor(race_cleaned)unreported/other
##
##
                              -0.1746070
                                                                      1.0148055
##
                   factor(FIGO)Stage II
                                                         factor(FIGO)Stage III
##
                              -0.6588743
                                                                     -1.7113719
##
                   factor(FIGO)Stage IV
                                                              age_at_diagnosis
##
                              -2.2627941
                                                                     -0.0213537
##
## Scale fixed at 1
##
## Loglik(model) = -3327 Loglik(intercept only) = -3377.4
## Chisq= 100.91 on 9 degrees of freedom, p= <2e-16
## n= 598
```

```
expAFT.bic <- stepAIC(full.expAFT, direction = "both", k=log(n), trace=1)
## Start: AIC=6717.9
## Surv(time, delta) ~ size.intermediate + factor(race_cleaned) +
##
       factor(FIGO) + age_at_diagnosis
##
##
                          Df
                                AIC
## - factor(race_cleaned) 4 6713.5
## - size.intermediate
                           1 6714.5
## <none>
                             6717.9
## - age_at_diagnosis
                           1 6732.2
## - factor(FIGO)
                           3 6748.0
##
## Step: AIC=6713.49
## Surv(time, delta) ~ size.intermediate + factor(FIGO) + age_at_diagnosis
##
##
                         Df
                                AIC
## - size.intermediate
                           1 6711.3
## <none>
                             6713.5
## + factor(race_cleaned) 4 6717.9
## - age_at_diagnosis
                          1 6731.0
## - factor(FIGO)
                           3 6746.6
##
## Step: AIC=6711.3
## Surv(time, delta) ~ factor(FIGO) + age_at_diagnosis
##
##
                          Df
                                AIC
## <none>
                             6711.3
## + size.intermediate
                           1 6713.5
## + factor(race_cleaned) 4 6714.5
## - age_at_diagnosis 1 6731.1
## - factor(FIGO)
                           3 6746.3
expAFT.bic
## Call:
## survreg(formula = Surv(time, delta) ~ factor(FIGO) + age_at_diagnosis,
       data = data.raceCleaned, dist = "exponential")
##
## Coefficients:
##
             (Intercept) factor(FIGO)Stage II factor(FIGO)Stage III
             11.52387703
                                   -0.64591491
                                                        -1.71559843
##
                              age_at_diagnosis
## factor(FIGO)Stage IV
            -2.32689756
                                  -0.02338033
##
##
## Scale fixed at 1
## Loglik(model) = -3339.7 Loglik(intercept only) = -3377.4
## Chisq= 75.54 on 4 degrees of freedom, p= 1.53e-15
## n= 598
```

Weibull AFT

```
full.weibullAFT <- survreg(Surv(time, delta) ~ size.intermediate + factor(race_cleaned) +
    factor(FIGO) + age_at_diagnosis, data = data.raceCleaned, dist = "weibull")
weibullAFT.aic <- stepAIC(full.weibullAFT, direction = "both", k=2, trace=1)</pre>
## Start: AIC=6640.12
## Surv(time, delta) ~ size.intermediate + factor(race_cleaned) +
##
       factor(FIGO) + age_at_diagnosis
##
##
                          Df
                                AIC
## <none>
                             6640.1
## - size.intermediate
                           1 6640.4
## - factor(race_cleaned) 4 6650.7
## - age_at_diagnosis
                           1 6656.5
## - factor(FIGO)
                           3 6676.7
weibullAFT.aic
## Call:
## survreg(formula = Surv(time, delta) ~ size.intermediate + factor(race_cleaned) +
       factor(FIGO) + age_at_diagnosis, data = data.raceCleaned,
       dist = "weibull")
##
## Coefficients:
##
                            (Intercept)
                                                            size.intermediate
                            12.36301594
                                                                   -0.28985830
##
##
              factor(race cleaned)asian
                                                    factor(race_cleaned)black
##
                             1.22733827
                                                                   -0.44097742
##
           factor(race_cleaned)hispanic factor(race_cleaned)unreported/other
##
                            -0.20784887
                                                                    1.24874844
##
                   factor(FIGO)Stage II
                                                        factor(FIGO)Stage III
##
                            -0.82430720
                                                                  -2.11463119
##
                   factor(FIGO)Stage IV
                                                             age_at_diagnosis
##
                            -2.76756691
                                                                  -0.02613169
##
## Scale= 1.294644
## Loglik(model) = -3309.1 Loglik(intercept only) = -3352.5
## Chisq= 86.94 on 9 degrees of freedom, p= 6.68e-15
## n= 598
weibullAFT.bic <- stepAIC(full.weibullAFT, direction = "both", k=log(n), trace=1)</pre>
## Start: AIC=6688.45
## Surv(time, delta) ~ size.intermediate + factor(race_cleaned) +
##
       factor(FIGO) + age_at_diagnosis
##
                                AIC
## - factor(race_cleaned) 4 6681.4
## - size.intermediate
                           1 6684.3
                             6688.4
## <none>
## - age_at_diagnosis
                           1 6700.4
## - factor(FIGO)
                           3 6711.8
## Step: AIC=6681.41
```

```
## Surv(time, delta) ~ size.intermediate + factor(FIGO) + age_at_diagnosis
##
##
                          Df
                                AIC
## - size.intermediate
                           1 6678.2
## <none>
                             6681.4
## + factor(race_cleaned) 4 6688.4
## - age at diagnosis
                           1 6696.2
## - factor(FIGO)
                           3 6707.3
##
## Step: AIC=6678.2
## Surv(time, delta) ~ factor(FIGO) + age_at_diagnosis
##
                          Df
##
                                AIC
## <none>
                             6678.2
## + size.intermediate
                           1 6681.4
## + factor(race_cleaned) 4 6684.3
## - age_at_diagnosis
                           1 6694.8
## - factor(FIGO)
                           3 6705.6
weibullAFT.bic
## Call:
## survreg(formula = Surv(time, delta) ~ factor(FIGO) + age_at_diagnosis,
       data = data.raceCleaned, dist = "weibull")
##
##
## Coefficients:
##
             (Intercept) factor(FIGO)Stage II factor(FIGO)Stage III
##
             12.35580783
                                   -0.80343728
                                                         -2.13537950
   factor(FIGO)Stage IV
##
                              age_at_diagnosis
                                   -0.02884478
##
             -2.86714404
##
## Scale= 1.311245
##
## Loglik(model) = -3319.9 Loglik(intercept only) = -3352.5
## Chisq= 65.22 on 4 degrees of freedom, p= 2.31e-13
## n= 598
Log-Logistic AFT.
full.loglogisticAFT <- survreg(Surv(time, delta) ~ size.intermediate + factor(race_cleaned) +</pre>
    factor(FIGO) + age_at_diagnosis, data = data.raceCleaned, dist = "loglogistic")
loglogisticAFT.aic <- stepAIC(full.loglogisticAFT, direction = "both", k=2, trace=1)</pre>
## Start: AIC=6595.7
## Surv(time, delta) ~ size.intermediate + factor(race_cleaned) +
##
       factor(FIGO) + age_at_diagnosis
##
##
                          Df
                                AIC
                           1 6594.6
## - size.intermediate
## <none>
                             6595.7
## - factor(race_cleaned) 4 6605.0
## - age_at_diagnosis
                           1 6618.7
## - factor(FIGO)
                           3 6630.9
##
```

```
## Step: AIC=6594.55
## Surv(time, delta) ~ factor(race_cleaned) + factor(FIGO) + age_at_diagnosis
##
##
                                AIC
                          Df
## <none>
                              6594.6
## + size.intermediate
                           1 6595.7
## - factor(race cleaned) 4 6604.2
## - age_at_diagnosis
                           1 6618.3
## - factor(FIGO)
                           3 6630.5
loglogisticAFT.aic
## Call:
## survreg(formula = Surv(time, delta) ~ factor(race_cleaned) +
       factor(FIGO) + age_at_diagnosis, data = data.raceCleaned,
##
       dist = "loglogistic")
##
##
## Coefficients:
##
                             (Intercept)
                                                    factor(race_cleaned)asian
##
                            11.75168203
                                                                    1.24102582
              factor(race_cleaned)black
##
                                                 factor(race_cleaned)hispanic
##
                            -0.48585138
                                                                   -0.17884472
## factor(race cleaned)unreported/other
                                                         factor(FIGO)Stage II
##
                              1.09502164
                                                                   -0.72098231
##
                  factor(FIGO)Stage III
                                                         factor(FIGO)Stage IV
##
                                                                   -2.69683380
                            -1.96859618
##
                       age_at_diagnosis
##
                            -0.03253153
##
## Scale= 0.9524551
## Loglik(model) = -3287.3 Loglik(intercept only) = -3330.7
## Chisq= 86.94 on 8 degrees of freedom, p= 1.94e-15
## n= 598
loglogisticAFT.bic <- stepAIC(full.loglogisticAFT, direction = "both", k=log(n), trace=1)</pre>
## Start: AIC=6644.03
## Surv(time, delta) ~ size.intermediate + factor(race_cleaned) +
##
       factor(FIGO) + age_at_diagnosis
##
##
                          Df
                                ATC
## - factor(race cleaned) 4 6635.7
## - size.intermediate
                           1 6638.5
## <none>
                              6644.0
## - age_at_diagnosis
                           1 6662.6
## - factor(FIGO)
                           3 6666.1
##
## Step: AIC=6635.75
## Surv(time, delta) ~ size.intermediate + factor(FIGO) + age_at_diagnosis
##
                          Df
                                AIC
## - size.intermediate
                           1 6630.6
## <none>
                              6635.7
## + factor(race_cleaned) 4 6644.0
```

```
## - age_at_diagnosis
                           1 6657.3
## - factor(FIGO)
                           3 6659.9
##
## Step: AIC=6630.58
## Surv(time, delta) ~ factor(FIGO) + age_at_diagnosis
##
##
                          Df
                                AIC
## <none>
                             6630.6
## + size.intermediate
                           1 6635.7
## + factor(race_cleaned) 4 6638.5
## - age_at_diagnosis
                           1 6653.1
## - factor(FIGO)
                           3 6655.7
loglogisticAFT.bic
## Call:
## survreg(formula = Surv(time, delta) ~ factor(FIGO) + age_at_diagnosis,
       data = data.raceCleaned, dist = "loglogistic")
##
##
## Coefficients:
             (Intercept) factor(FIGO)Stage II factor(FIGO)Stage III
##
##
             11.93419596
                                   -0.69099201
                                                          -1.98625823
##
  factor(FIGO)Stage IV
                              age_at_diagnosis
##
             -2.75224858
                                   -0.03446027
##
## Scale= 0.9666854
                            Loglik(intercept only) = -3330.7
## Loglik(model) = -3296.1
## Chisq= 69.27 on 4 degrees of freedom, p= 3.24e-14
## n= 598
Log-Normal AFT
full.lognormalAFT <- survreg(Surv(time, delta) ~ size.intermediate + factor(race_cleaned) +</pre>
    factor(FIGO) + age_at_diagnosis, data = data.raceCleaned, dist = "lognormal")
lognormalAFT.aic <- stepAIC(full.lognormalAFT, direction = "both", k=2, trace=1)</pre>
## Start: AIC=6604.08
## Surv(time, delta) ~ size.intermediate + factor(race_cleaned) +
##
       factor(FIGO) + age_at_diagnosis
##
##
                                AIC
## - size.intermediate
                           1 6602.9
                             6604.1
## <none>
## - factor(race_cleaned) 4 6611.1
## - age_at_diagnosis
                           1 6629.1
## - factor(FIGO)
                           3 6638.9
## Step: AIC=6602.87
## Surv(time, delta) ~ factor(race_cleaned) + factor(FIGO) + age_at_diagnosis
##
##
                          Df
                                AIC
## <none>
                             6602.9
## + size.intermediate
                           1 6604.1
```

```
## - factor(race_cleaned) 4 6610.4
## - age_at_diagnosis
                           1 6628.8
## - factor(FIGO)
                           3 6638.3
lognormalAFT.aic
## Call:
## survreg(formula = Surv(time, delta) ~ factor(race_cleaned) +
##
       factor(FIGO) + age_at_diagnosis, data = data.raceCleaned,
##
       dist = "lognormal")
##
## Coefficients:
                                                    factor(race_cleaned)asian
##
                             (Intercept)
##
                             11.60283189
                                                                    1.32670368
##
              factor(race_cleaned)black
                                                 factor(race_cleaned)hispanic
##
                             -0.60372256
                                                                   -0.21390418
## factor(race_cleaned)unreported/other
                                                         factor(FIGO)Stage II
##
                             0.71779553
                                                                   -0.27403185
##
                  factor(FIGO)Stage III
                                                         factor(FIGO)Stage IV
##
                            -1.53718723
                                                                   -2.39908988
##
                       age_at_diagnosis
##
                            -0.03567528
##
## Scale= 1.698155
## Loglik(model) = -3291.4 Loglik(intercept only) = -3333.2
## Chisq= 83.54 on 8 degrees of freedom, p= 9.43e-15
## n= 598
lognormalAFT.bic <- stepAIC(full.lognormalAFT, direction = "both", k=log(n), trace=1)</pre>
## Start: AIC=6652.41
## Surv(time, delta) ~ size.intermediate + factor(race_cleaned) +
##
       factor(FIGO) + age_at_diagnosis
##
##
                          Df
                                AIC
## - factor(race_cleaned) 4 6641.9
## - size.intermediate
                           1 6646.8
## <none>
                              6652.4
## - age_at_diagnosis
                           1 6673.0
## - factor(FIGO)
                           3 6674.0
##
## Step: AIC=6641.89
## Surv(time, delta) ~ size.intermediate + factor(FIGO) + age_at_diagnosis
##
##
                                 AIC
                          Df
                           1 6636.8
## - size.intermediate
## <none>
                             6641.9
## + factor(race_cleaned) 4 6652.4
## - factor(FIGO)
                           3 6663.8
## - age_at_diagnosis
                           1 6664.4
## Step: AIC=6636.78
## Surv(time, delta) ~ factor(FIGO) + age_at_diagnosis
##
```

```
##
                          Df
                                AIC
## <none>
                             6636.8
## + size.intermediate
                           1 6641.9
## + factor(race_cleaned) 4 6646.8
## - factor(FIGO)
                           3 6659.4
## - age_at_diagnosis
                           1 6660.5
lognormalAFT.bic
## Call:
## survreg(formula = Surv(time, delta) ~ factor(FIGO) + age_at_diagnosis,
       data = data.raceCleaned, dist = "lognormal")
##
## Coefficients:
##
             (Intercept) factor(FIGO)Stage II factor(FIGO)Stage III
##
              11.6248024
                                    -0.1457258
                                                           -1.4340866
##
   factor(FIGO)Stage IV
                              age_at_diagnosis
                                     -0.0371437
##
              -2.3211559
##
## Scale= 1.716158
##
## Loglik(model) = -3299.2
                           Loglik(intercept only) = -3333.2
## Chisq= 68 on 4 degrees of freedom, p= 6.01e-14
## n= 598
```

Conclusions.

For both the AIC and BIC Stepwise selection models, FIGO and age_at_diagnosis were selected. In both models, the Biopsy size variable size.intermediate was not considered significant. Going forward, it would be prudent to use the BIC chosen model since there is a larger penalty which adjusts for the large sample size (n=598).

Model Diagnostics.

Cox-Snell Residual Plot.

```
plotCoxSnellCPH<- function(survFit, delta, fitType){

# get Cox-Snell Residual based on Martingale Residuals
mg.residual <- resid(survFit, type = "martingale")

cs.residual <- delta - mg.residual

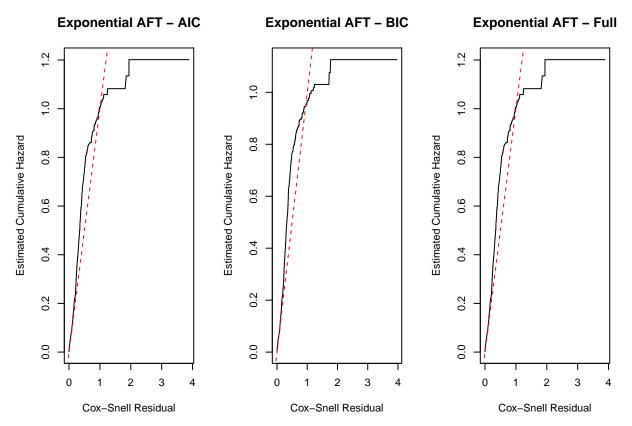
# Graphical Plot
fit.cs <- survfit(Surv(cs.residual, delta) ~ 1) # get KM estimates
H.cs <- cumsum(fit.cs$n.event/fit.cs$n.risk)

plot(fit.cs$time, H.cs, type='s', col='blue',
    main = paste0('Cox-PH - ', fitType),
    xlab = 'Residual', ylab = 'Nelson-Aalen Cum. Hazard')
#Note here that 'time' is the value of the Cox-Snell residual
abline(0, 1, col='red', lty = 2)
}</pre>
```

```
plotCSExpAFT <- function(survFit, data, fitType){</pre>
  sigma <- survFit$scale</pre>
  eta <- -survFit$linear.predictors/sigma
 r.exp <- data$time * exp(eta)</pre>
 fit <- survfit(Surv(r.exp, data$delta) ~ 1)</pre>
 H.exp <- cumsum(fit$n.event / fit$n.risk)</pre>
 plot(H.exp ~ fit$time, type = 'l', main = paste0('Exponential AFT - ', fitType),
       ylab = 'Estimated Cumulative Hazard', xlab = 'Cox-Snell Residual')
  abline(0, 1, col='red', lty=2)
plotCSWeibullAFT <- function(survFit, data, fitType){</pre>
  sigma <- survFit$scale</pre>
  alpha <- 1 / sigma
      <- -survFit$linear.predictors / sigma</pre>
 r.wb <- data$time^alpha * exp(eta)</pre>
 fit <- survfit(Surv(r.wb, data$delta) ~ 1)</pre>
 H.wb <- cumsum(fit$n.event/fit$n.risk)
 plot(H.wb ~ fit$time, type = 'l', main = paste0('Weibull AFT - ', fitType),
       ylab = 'Estimated Cumulative Hazard', xlab = 'Cox-Snell Residual')
  abline(0, 1, col='red', lty=2)
plotCSLogLogisticAFT <- function(survFit, data, fitType){</pre>
  sigma <- data$scale
  alpha <- 1 / sigma
      <- -data$linear.predictors / sigma</pre>
  eta
 r.ll <- -log(1/(1 + data$time^alpha*exp(eta)))</pre>
 fit <- survfit(Surv(r.ll, data$delta) ~ 1)
 H.ll <- cumsum(fit$n.event / fit$n.risk)</pre>
 plot(H.11 ~ fit$time, type = '1', main = paste0('Log-Logistic AFT- ', fitType),
       ylab = 'Estimated Cumulative Hazard', xlab = 'Cox-Snell Residual')
  abline(0, 1, col='red', lty=2)
}
plotCSLogNormalAFT <- function(survFit, data, fitType){</pre>
        <- -survFit$linear.predictors / survFit$scale</pre>
  r.ln <- -log(1 - pnorm((log(data$time) - survFit$linear.predictors) / survFit$scale))
  fit <- survfit(Surv(r.ln, data$delta) ~ 1)</pre>
 H.ln <- cumsum(fit$n.event/fit$n.risk)</pre>
  plot(H.ln ~ fit$time, type = 'l', main = paste0('Log-Normal AFT - ', fitType),
       ylab = 'Estimated Cumulative Hazard', xlab = 'Cox-Snell Residual')
  abline(0, 1, col='red', lty = 2)
```

```
bic.modfit <- coxph(Surv(time, delta)~factor(FIGO)+age_at_diagnosis, data = data.raceCleaned, method =
aic.modfit <- coxph(Surv(time,delta)~factor(race_cleaned)+factor(FIGO)+age_at_diagnosis, data = data.ra
par(mfrow = c(1,3))
plotCoxSnellCPH(aic.modfit, data.raceCleaned$delta, "AIC")
plotCoxSnellCPH(bic.modfit, data.raceCleaned$delta, "BIC")
plotCoxSnellCPH(fullCoxMod, data.raceCleaned$delta, "Full Variable")
                                               Cox-PH - BIC
           Cox-PH - AIC
                                                                             Cox-PH - Full Variable
                                       2.0
    2.0
                                                                          2.0
                                       1.5
    1.5
Nelson-Aalen Cum. Hazard
                                   Nelson-Aalen Cum. Hazard
                                                                      Nelson-Aalen Cum. Hazard
                                       0.1
    1.0
                                                                          1.0
                                       0.5
    0.5
    0.0
                                       0.0
                                                                          0.0
       0.0
                1.0
                        2.0
                                           0.0
                                                   1.0
                                                           2.0
                                                                              0.0 0.5 1.0 1.5 2.0 2.5
                                                   Residual
               Residual
                                                                                      Residual
par(mfrow = c(1,3))
plotCSExpAFT(expAFT.aic, data.raceCleaned, "AIC")
plotCSExpAFT(expAFT.bic, data.raceCleaned, "BIC")
```

plotCSExpAFT(full.expAFT, data.raceCleaned, "Full")



Ok both models fit similarly, which isn't helpful.