

Math 150 Survival Analysis Project Test Code

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```
knitr::opts_chunk$set(message=FALSE, warning=FALSE, fig.height=4, fig.width=5,
                        fig.align = "center")
library(tidyverse)
library(broom)
library(survival)
library(survminer)
library(coxed)
```

Importing data:

```
AD = read.csv("AIDSdata.csv")
```

Trying to do power analysis

```
null = sim.survdata(851, 100, x = 1, beta = 0, censor = (851 - 69)/851)
alt = sim.survdata(851, 100, x = 1, beta = -1.454, censor = (851 - 69)/851)
#alt$data$y
cx = coxph(Surv(y, failed) ~ X, data = alt$data)
#coxph(Surv(y, failed) ~ X, data = null$data)
scx = summary(cx)
pval = scx$waldtest[3]
```

X might be the number of covariates?

```
loopVals = c(1:50)
output = rep(NA, length(loopVals))
#output
for(i in loopVals){
  loopSim = sim.survdata(851, 100, x = 1, beta = -1.454, censor = (851 - 69)/851)
  pval = summary(coxph(Surv(y, failed) ~ X, data = loopSim$data))$waldtest[3]
  output[i] = pval < 0.05
}
sum(output) / length(output)
```

```
betaVals = c(0, -0.2, -0.5, -1, -1.454, -1.5, -2, -3)
powersForBeta = rep(NA, length(betaVals))
outerLoop = c(1:length(betaVals)) #loop through betas
innerLoop = c(1:1000) #replicates
for(i in outerLoop){
  pBools = rep(NA, length(innerLoop))
  for(j in innerLoop){
    loopSim = sim.survdata(851, 100, x = 1, beta = betaVals[i], censor = (851 - 69)/851)
```

```

    pval = summary(coxph(Surv(y, failed) ~ X, data = loopSim$data))$waldtest[3]
    pBools[j] = pval < 0.05
  }
  powersForBeta[i] = sum(pBools) / length(pBools)
}

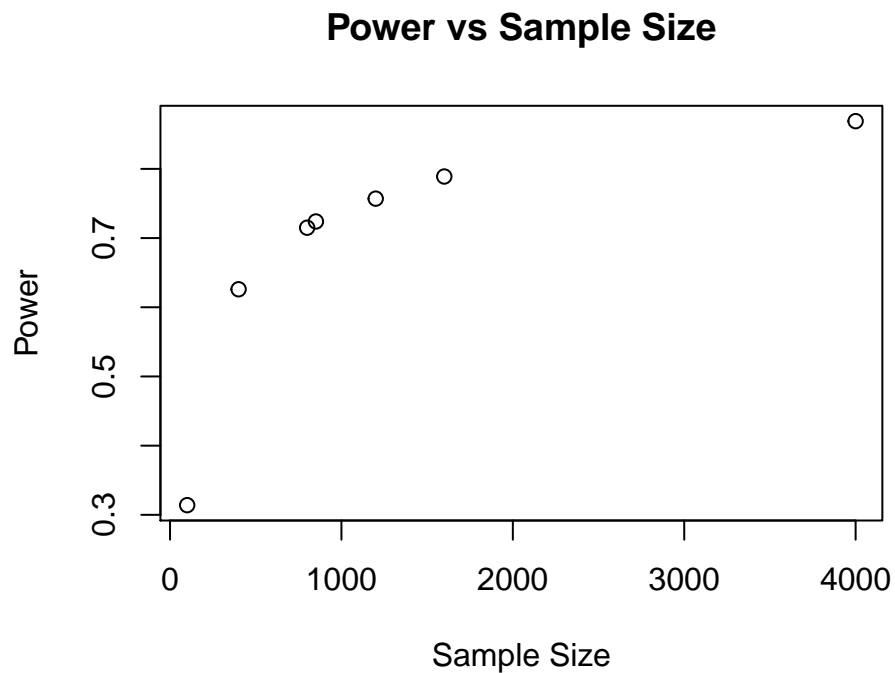
powersForBeta

popVals = c(100, 400, 800, 851, 1200, 1600, 4000)
powersForPop = rep(NA, length(popVals))
outerLoop = c(1:length(popVals)) #loop through betas
innerLoop = c(1:1000) #replicates
for(i in outerLoop){
  pBools = rep(NA, length(innerLoop))
  for(j in innerLoop){
    loopSim = sim.survdata(popVals[i], 100, x= 1, beta = -1.454, censor = (851 - 69)/851)
    pval = summary(coxph(Surv(y, failed) ~ X, data = loopSim$data))$waldtest[3]
    pBools[j] = pval < 0.05
  }
  powersForPop[i] = sum(pBools) / length(pBools)
}

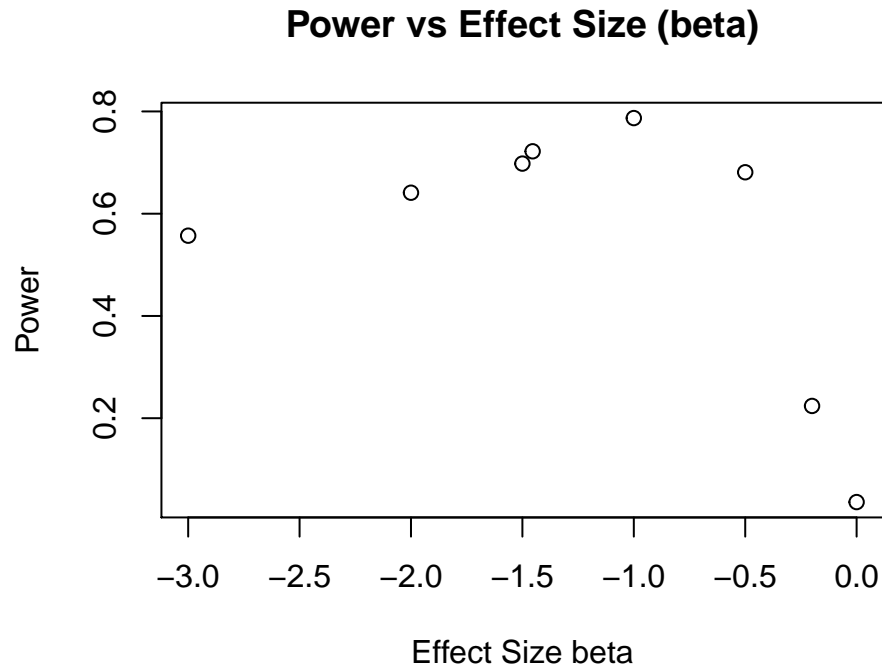
powersForPop

popx = c(100, 400, 800, 851, 1200, 1600, 4000)
popy = c(0.314, 0.626, 0.715, 0.724, 0.757, 0.789, 0.869)
plot(popx, popy, main = "Power vs Sample Size", xlab = "Sample Size", ylab = "Power")

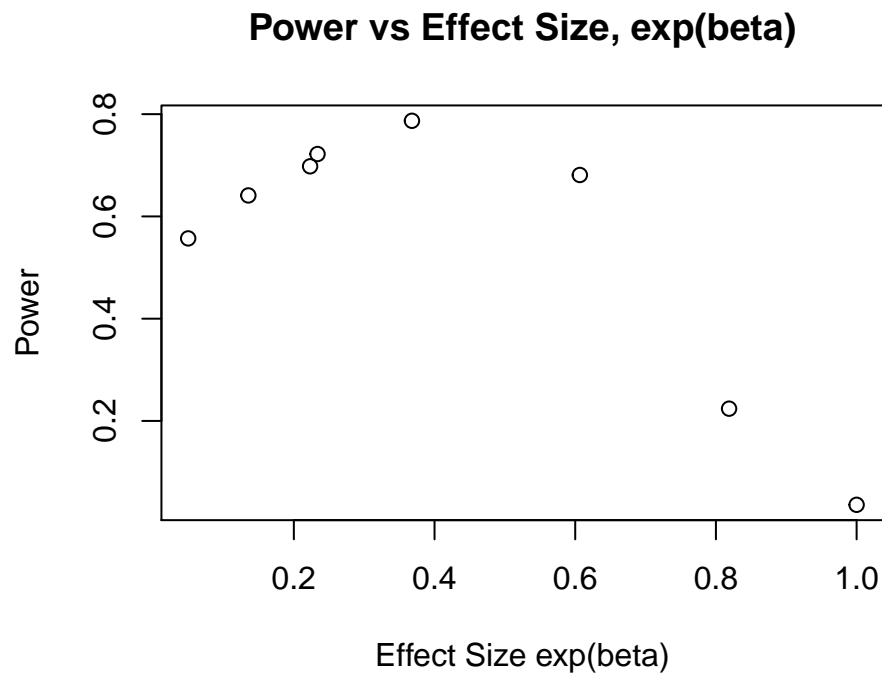
```



```
betax = c(0, -0.2, -0.5, -1, -1.454, -1.5, -2, -3)
betay = c(0.036, 0.224, 0.681, 0.787, 0.722, 0.698, 0.641, 0.557)
plot(betax, betay, main = "Power vs Effect Size (beta)", xlab = "Effect Size beta", ylab = "Power")
```

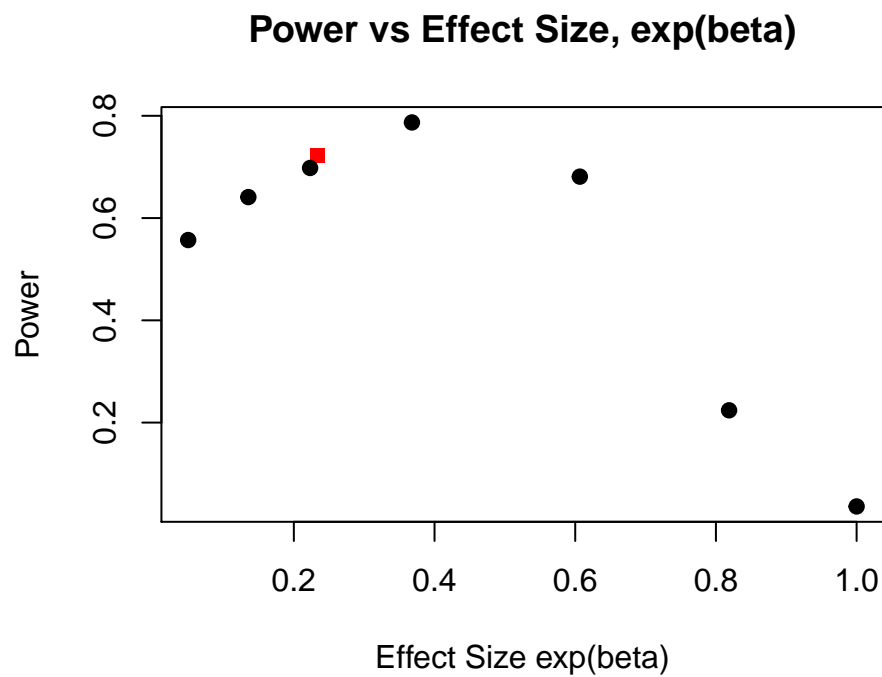


```
expbetax = c(1.00000000, 0.81873075, 0.60653066, 0.36787944, 0.23363388, 0.22313016, 0.13533528, 0.04978707)
plot(expbetax, betay, main = "Power vs Effect Size, exp(beta)", xlab = "Effect Size exp(beta)", ylab = "Power")
```



Updated power plot

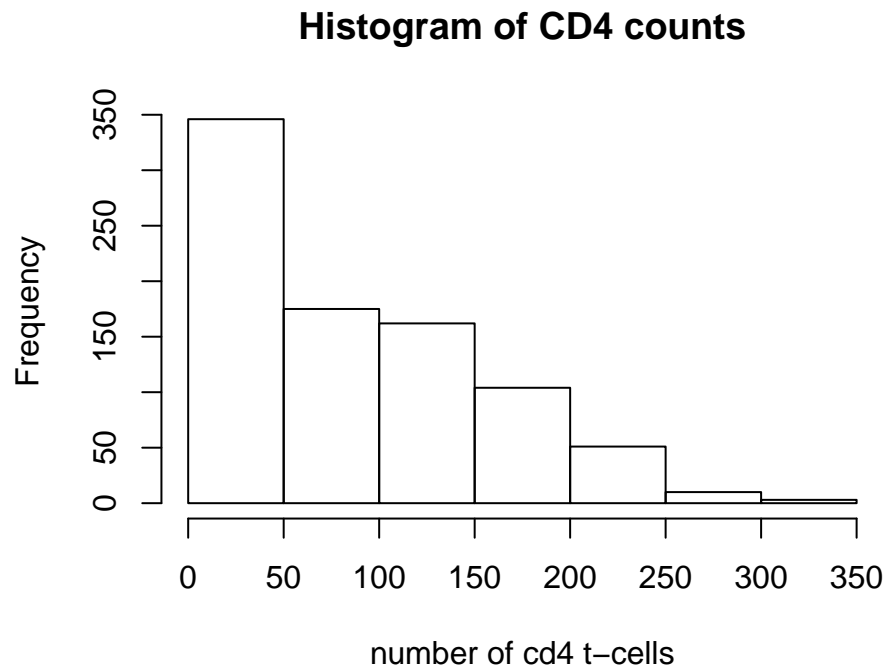
```
plot(expbetax, betay, main = "Power vs Effect Size,  $\exp(\beta)$ ", xlab = "Effect Size  $\exp(\beta)$ ", ylab =
```



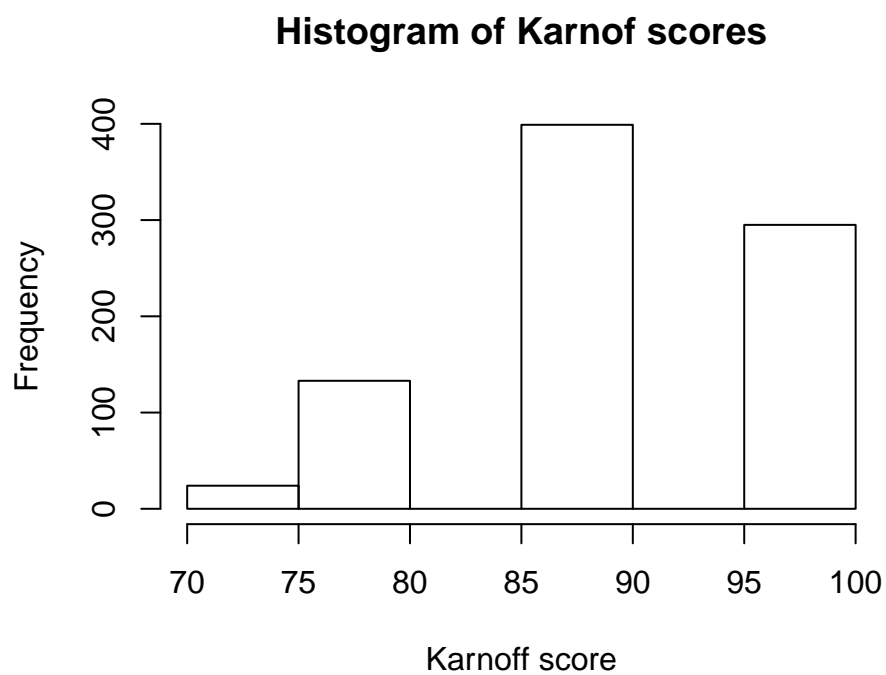
Beta	Exp(Beta)	Power
0	1	0.036
-0.2	0.8187	0.224
-0.5	0.6065	0.681
-1	0.3679	0.787
-1.454	0.2336	0.722
-1.5	0.2231	0.698
-2	0.1353	0.641
-3	0.0498	0.557

Some exploratory plots:

```
hist(AD$cd4, main = "Histogram of CD4 counts", xlab = "number of cd4 t-cells")
```



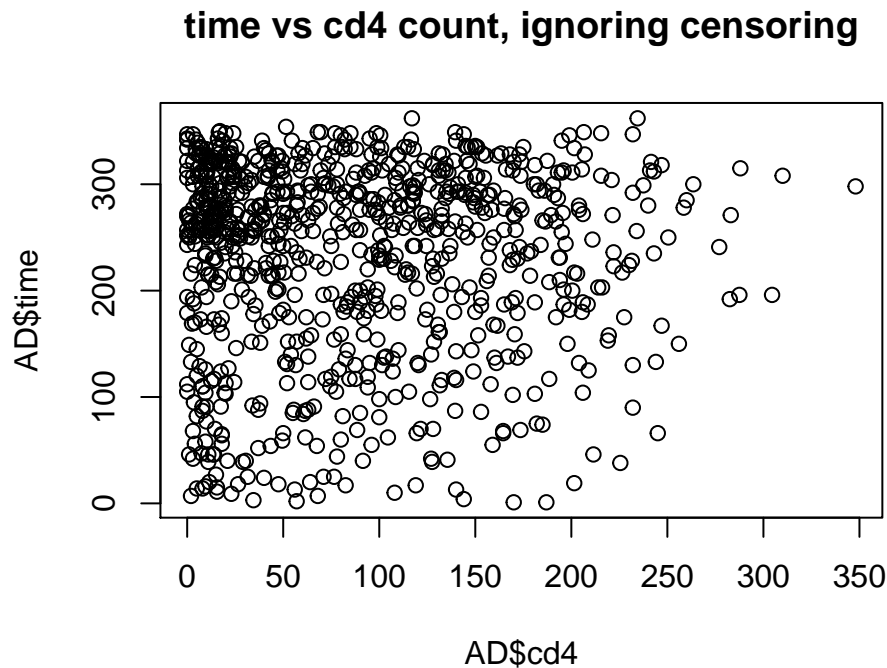
```
hist(AD$karnof, main = "Histogram of Karnof scores", xlab = "Karnoff score", breaks = 5)
```



#I'm having a hard time getting the histogram breaks right - karnof scores are multiples of 10 and it
`plot(ADage, ADkarnof, main = "plot of karnof score vs age")`



```
plot(AD$cd4, AD$time, main = "time vs cd4 count, ignoring censoring")
```



Working on COX model

Variables: Response: time, censor Not using: time.d, censor.d (because we're using "aids defining event or death", not just death) Explanatory: tx, txgrp, strat2, sex, raceth, ivdrug, hemophil, karnof, cd4, priorzdv, age

Let's check if we need interaction. Likelihood ratio test is $2\ln(\text{Likelihood of full}) - 2\ln(\text{Likelihood of reduced})$

```
full = coxph(Surv(time, censor) ~ (tx+strat2+sex+raceth+ivdrug+hemophil+karnof+cd4+priorzdv+age)^2, data = AD)
full$loglik[2]
```

```
## [1] -380.5977
```

```
red = coxph(Surv(time, censor) ~ tx+strat2+sex+raceth+ivdrug+hemophil+karnof+cd4+priorzdv+age, data = AD)
red$loglik[2]
```

```
## [1] -411.5144
```

```
length(red$coefficients)
```

```
## [1] 10
```

```
1 - pchisq(2 * full$loglik[2] - 2 * red$loglik[2], df = length(full$coefficients) - length(red$coefficients))
```

```
## [1] 0.04845761
```

```
full
```

```
## Call:
```

```
## coxph(formula = Surv(time, censor) ~ (tx + strat2 + sex + raceth +
```

```
##      ivdrug + hemophil + karnof + cd4 + priorzdv + age)^2, data = AD)
##
##              coef exp(coef)   se(coef)      z      p
## tx            -5.501e+00  4.084e-03  4.369e+00 -1.259 0.20807
## strat2        -9.622e-01  3.820e-01  7.284e+00 -0.132 0.89490
## sex            1.137e+01  8.710e+04  5.589e+00  2.035 0.04182
## raceth         7.870e+00  2.617e+03  2.418e+00  3.255 0.00113
## ivdrug        -7.581e+00  5.101e-04  3.252e+00 -2.331 0.01976
## hemophil       1.996e+01  4.646e+08  2.107e+04  0.001 0.99924
## karnof         1.860e-01  1.204e+00  1.246e-01  1.493 0.13539
## cd4           -1.073e-01  8.983e-01  8.413e-02 -1.275 0.20221
## priorzdv       6.378e-02  1.066e+00  8.320e-02  0.767 0.44336
## age           4.436e-01  1.558e+00  2.030e-01  2.185 0.02890
## tx:strat2      1.039e-02  1.010e+00  9.648e-01  0.011 0.99141
## tx:sex         1.969e+00  7.163e+00  7.915e-01  2.487 0.01287
## tx:raceth      -6.084e-02  9.410e-01  3.554e-01 -0.171 0.86408
## tx:ivdrug      9.461e-02  1.099e+00  5.560e-01  0.170 0.86488
## tx:hemophil    -1.611e+01  1.008e-07  4.131e+03 -0.004 0.99689
## tx:karnof      2.166e-02  1.022e+00  3.952e-02  0.548 0.58369
## tx:cd4         5.997e-03  1.006e+00  1.114e-02  0.538 0.59049
## tx:priorzdv    -3.839e-02  9.623e-01  1.685e-02 -2.278 0.02273
## tx:age         3.409e-02  1.035e+00  3.266e-02  1.044 0.29656
## strat2:sex     -2.387e-01  7.876e-01  1.326e+00 -0.180 0.85716
## strat2:raceth  -4.696e-01  6.253e-01  6.182e-01 -0.760 0.44747
## strat2:ivdrug  5.624e-01  1.755e+00  7.919e-01  0.710 0.47756
## strat2:hemophil 1.885e+00  6.585e+00  3.504e+00  0.538 0.59061
## strat2:karnof  -1.333e-02  9.868e-01  5.889e-02 -0.226 0.82091
## strat2:cd4     -2.740e-03  9.973e-01  1.214e-02 -0.226 0.82142
## strat2:priorzdv 4.170e-02  1.043e+00  2.261e-02  1.845 0.06510
## strat2:age     4.497e-02  1.046e+00  4.961e-02  0.906 0.36468
## sex:raceth     -1.886e+00  1.516e-01  6.562e-01 -2.875 0.00404
## sex:ivdrug     4.405e-01  1.553e+00  6.784e-01  0.649 0.51617
## sex:hemophil   -2.067e+01  1.060e-09  2.085e+04 -0.001 0.99921
## sex:karnof     -7.926e-02  9.238e-01  5.430e-02 -1.460 0.14439
## sex:cd4        -8.461e-03  9.916e-01  1.769e-02 -0.478 0.63250
## sex:priorzdv   -3.032e-04  9.997e-01  2.286e-02 -0.013 0.98941
## sex:age        -5.357e-02  9.478e-01  4.356e-02 -1.230 0.21880
## raceth:ivdrug  2.133e-01  1.238e+00  2.426e-01  0.879 0.37933
## raceth:hemophil -1.350e+01  1.373e-06  3.044e+03 -0.004 0.99646
## raceth:karnof  -5.435e-02  9.471e-01  2.404e-02 -2.261 0.02374
## raceth:cd4     7.225e-03  1.007e+00  6.483e-03  1.114 0.26509
## raceth:priorzdv -3.059e-03  9.969e-01  9.559e-03 -0.320 0.74898
## raceth:age     -3.873e-02  9.620e-01  2.056e-02 -1.884 0.05957
## ivdrug:hemophil 2.633e+01  2.712e+11  6.088e+03  0.004 0.99655
## ivdrug:karnof  5.604e-02  1.058e+00  2.796e-02  2.004 0.04506
## ivdrug:cd4     -1.876e-02  9.814e-01  1.333e-02 -1.408 0.15921
## ivdrug:priorzdv 2.655e-02  1.027e+00  1.055e-02  2.517 0.01182
## ivdrug:age     2.986e-02  1.030e+00  3.208e-02  0.931 0.35186
## hemophil:karnof -8.703e-02  9.167e-01  1.461e-01 -0.596 0.55134
## hemophil:cd4   -9.746e-03  9.903e-01  4.981e-02 -0.196 0.84486
## hemophil:priorzdv -4.504e-02  9.560e-01  3.127e-02 -1.440 0.14975
## hemophil:age   -7.388e-02  9.288e-01  1.262e-01 -0.586 0.55820
## karnof:cd4     1.096e-03  1.001e+00  6.856e-04  1.599 0.10983
## karnof:priorzdv -4.181e-04  9.996e-01  7.074e-04 -0.591 0.55454
```



```
## karnof:age      -4.056e-03  9.960e-01  2.148e-03 -1.888 0.05905
## cd4:priorzdv   -2.991e-04  9.997e-01  2.646e-04 -1.130 0.25830
## cd4:age        3.926e-04  1.000e+00  5.166e-04  0.760 0.44732
## priorzdv:age   -1.207e-03  9.988e-01  8.244e-04 -1.464 0.14323
##
## Likelihood ratio test=144.1 on 55 df, p=6.536e-10
## n= 851, number of events= 69
```

This is close enough to 0.05 that I'm not comfortable eliminating interaction.

lets eliminate hemophil entirely

```
full = coxph(Surv(time, censor) ~ (tx+strat2+sex+raceth+ivdrug+hemophil+karnof+cd4+priorzdv+age)^2, data = AD)
full$loglik[2]
```

```
## [1] -380.5977
```

```
red = coxph(Surv(time, censor) ~ (tx+strat2+sex+raceth+ivdrug+karnof+cd4+priorzdv+age)^2, data = AD)
red$loglik[2]
```

```
## [1] -384.3829
```

```
1 - pchisq(2 * full$loglik[2] - 2 * red$loglik[2] , df = length(full$coefficients) - length(red$coefficients))
```

```
## [1] 0.6707304
```

```
#red
```

We can eliminate hemophil entirely. Let's try that with strat 2

```
full = coxph(Surv(time, censor) ~ (tx+strat2+sex+raceth+ivdrug+karnof+cd4+priorzdv+age)^2, data = AD)
full$loglik[2]
```

```
## [1] -384.3829
```

```
red = coxph(Surv(time, censor) ~ (tx+sex+raceth+ivdrug+karnof+cd4+priorzdv+age)^2, data = AD)
red$loglik[2]
```

```
## [1] -387.8855
```

```
1 - pchisq(2 * full$loglik[2] - 2 * red$loglik[2] , df = length(full$coefficients) - length(red$coefficients))
```

```
## [1] 0.6365669
```

```
red
```

```
## Call:
```

```
## coxph(formula = Surv(time, censor) ~ (tx + sex + raceth + ivdrug +  
##       karnof + cd4 + priorzdv + age)^2, data = AD)
```

```
##
```

	coef	exp(coef)	se(coef)	z	p
## tx	-6.310e+00	1.818e-03	4.296e+00	-1.469	0.14191
## sex	1.144e+01	9.324e+04	5.571e+00	2.054	0.03999
## raceth	7.255e+00	1.415e+03	2.301e+00	3.153	0.00162
## ivdrug	-6.516e+00	1.480e-03	2.981e+00	-2.186	0.02881
## karnof	1.679e-01	1.183e+00	1.184e-01	1.418	0.15611
## cd4	-7.608e-02	9.267e-01	5.281e-02	-1.441	0.14969
## priorzdv	-3.132e-03	9.969e-01	6.658e-02	-0.047	0.96248
## age	4.069e-01	1.502e+00	1.893e-01	2.150	0.03156
## tx:sex	2.125e+00	8.372e+00	7.668e-01	2.771	0.00558
## tx:raceth	1.504e-01	1.162e+00	3.420e-01	0.440	0.66003

```
## tx:ivdrug      4.191e-02  1.043e+00  5.375e-01  0.078 0.93784
## tx:karnof      2.505e-02  1.025e+00  3.819e-02  0.656 0.51179
## tx:cd4         4.788e-03  1.005e+00  7.088e-03  0.676 0.49930
## tx:priorzdvdv -3.927e-02  9.615e-01  1.679e-02 -2.339 0.01933
## tx:age        3.413e-02  1.035e+00  3.090e-02  1.104 0.26946
## sex:raceth     -1.675e+00  1.874e-01  6.383e-01 -2.624 0.00870
## sex:ivdrug     3.202e-01  1.377e+00  6.562e-01  0.488 0.62555
## sex:karnof     -7.869e-02  9.243e-01  5.442e-02 -1.446 0.14816
## sex:cd4        -1.099e-02  9.891e-01  1.298e-02 -0.847 0.39703
## sex:priorzdvdv -4.393e-03  9.956e-01  2.031e-02 -0.216 0.82876
## sex:age        -6.038e-02  9.414e-01  4.138e-02 -1.459 0.14451
## raceth:ivdrug  1.385e-03  1.001e+00  2.362e-01  0.006 0.99532
## raceth:karnof  -5.064e-02  9.506e-01  2.266e-02 -2.235 0.02543
## raceth:cd4     2.794e-03  1.003e+00  4.207e-03  0.664 0.50658
## raceth:priorzdvdv -1.678e-03  9.983e-01  8.282e-03 -0.203 0.83939
## raceth:age     -3.409e-02  9.665e-01  1.899e-02 -1.795 0.07264
## ivdrug:karnof  5.035e-02  1.052e+00  2.597e-02  1.939 0.05256
## ivdrug:cd4     -9.323e-03  9.907e-01  7.417e-03 -1.257 0.20874
## ivdrug:priorzdvdv 2.362e-02  1.024e+00  9.567e-03  2.469 0.01356
## ivdrug:age     2.753e-02  1.028e+00  3.120e-02  0.883 0.37747
## karnof:cd4     6.475e-04  1.001e+00  4.480e-04  1.445 0.14839
## karnof:priorzdvdv -2.237e-05  1.000e+00  6.636e-04 -0.034 0.97311
## karnof:age     -3.687e-03  9.963e-01  1.942e-03 -1.898 0.05769
## cd4:priorzdvdv -4.921e-05  1.000e+00  1.366e-04 -0.360 0.71869
## cd4:age        4.985e-04  1.000e+00  3.177e-04  1.569 0.11670
## priorzdvdv:age -2.041e-04  9.998e-01  5.395e-04 -0.378 0.70521
##
## Likelihood ratio test=129.5 on 36 df, p=1.777e-12
## n= 851, number of events= 69
```

Okay let's eliminate anything with a p-value of 0.1 or larger.

```
full = coxph(Surv(time, censor) ~ (tx+sex+raceth+ivdrug+karnof+cd4+priorzdvdv+age)^2, data = AD)
full$loglik[2]
```

```
## [1] -387.8855
```

```
red = coxph(Surv(time, censor) ~ tx*(sex + priorzdvdv) + raceth*(sex + karnof + age) + ivdrug*(karnof + priorzdvdv) +
red$loglik[2]
```

```
## [1] -395.5924
```

```
1 - pchisq(2 * full$loglik[2] - 2 * red$loglik[2] , df = length(full$coefficients) - length(red$coefficients))
```

```
## [1] 0.7522683
```

```
red
```

```
## Call:
```

```
## coxph(formula = Surv(time, censor) ~ tx * (sex + priorzdvdv) +
##       raceth * (sex + karnof + age) + ivdrug * (karnof + priorzdvdv) +
##       karnof * age + cd4, data = AD)
```

```
##
```

	coef	exp(coef)	se(coef)	z	p
## tx	-1.773877	0.169674	0.886051	-2.002	0.04528
## sex	2.217867	9.187711	0.876329	2.531	0.01138
## priorzdvdv	-0.013380	0.986709	0.010620	-1.260	0.20770
## raceth	5.731699	308.492931	1.817565	3.154	0.00161

```
## karnof      0.027056  1.027426  0.081709  0.331  0.74055
## age        0.195550  1.215980  0.163624  1.195  0.23204
## ivdrug     -2.935066  0.053127  1.896507 -1.548  0.12171
## cd4        -0.016165  0.983965  0.003300 -4.899 9.63e-07
## tx:sex      1.644482  5.178329  0.681884  2.412  0.01588
## tx:priorzdv -0.035672  0.964957  0.014254 -2.503  0.01233
## sex:raceth  -1.480031  0.227631  0.588021 -2.517  0.01184
## raceth:karnof -0.036726  0.963940  0.019241 -1.909  0.05629
## raceth:age  -0.028096  0.972295  0.017905 -1.569  0.11660
## karnof:ivdrug 0.025423  1.025749  0.021618  1.176  0.23958
## priorzdv:ivdrug 0.016103  1.016233  0.006869  2.344  0.01906
## karnof:age   -0.001477  0.998524  0.001803 -0.819  0.41276
##
## Likelihood ratio test=114.1 on 16 df, p=< 2.2e-16
## n= 851, number of events= 69
```

We can do that, so let's eliminate karnof*age and karnof*ivdrug

```
full = coxph(Surv(time, censor) ~ tx*(sex + priorzdv) + raceth*(sex + karnof + age) + ivdrug*(karnof + age) + cd4)
full$loglik[2]
```

```
## [1] -395.5924
```

```
red = coxph(Surv(time, censor) ~ tx*(sex + priorzdv) + raceth*(sex + karnof + age) + ivdrug*priorzdv + cd4)
red$loglik[2]
```

```
## [1] -396.5796
```

```
1 - pchisq(2 * full$loglik[2] - 2 * red$loglik[2], df = length(full$coefficients) - length(red$coefficients))
```

```
## [1] 0.3726069
```

```
red
```

```
## Call:
```

```
## coxph(formula = Surv(time, censor) ~ tx * (sex + priorzdv) +
##       raceth * (sex + karnof + age) + ivdrug * priorzdv + cd4,
##       data = AD)
```

```
##
```

	coef	exp(coef)	se(coef)	z	p
tx	-1.651633	0.191736	0.877322	-1.883	0.05976
sex	2.340187	10.383173	0.875181	2.674	0.00750
priorzdv	-0.013415	0.986674	0.010817	-1.240	0.21490
raceth	5.275991	195.584220	1.762487	2.993	0.00276
karnof	-0.006901	0.993123	0.030802	-0.224	0.82274
age	0.064565	1.066695	0.031193	2.070	0.03847
ivdrug	-0.804226	0.447434	0.354498	-2.269	0.02329
cd4	-0.015752	0.984371	0.003258	-4.834	1.34e-06
tx:sex	1.551639	4.719197	0.674424	2.301	0.02141
tx:priorzdv	-0.035633	0.964994	0.014349	-2.483	0.01302
sex:raceth	-1.504677	0.222089	0.586714	-2.565	0.01033
raceth:karnof	-0.031977	0.968528	0.019002	-1.683	0.09240
raceth:age	-0.025912	0.974421	0.017915	-1.446	0.14806
priorzdv:ivdrug	0.015861	1.015988	0.007212	2.199	0.02786

```
##
```

```
## Likelihood ratio test=112.1 on 14 df, p=< 2.2e-16
```

```
## n= 851, number of events= 69
```

Let's try raceth*age

```
full = coxph(Surv(time, censor) ~ tx*(sex + priorzdv) + raceth*sex + raceth*karnof + raceth*age + ivdrug*age, data = AD)
full$loglik[2]
```

```
## [1] -396.5796
```

```
red = coxph(Surv(time, censor) ~ tx*(sex + priorzdv) + raceth*sex + raceth*karnof + ivdrug*priorzdv + cd4, data = AD)
red$loglik[2]
```

```
## [1] -398.9897
```

```
1 - pchisq(2 * full$loglik[2] - 2 * red$loglik[2], df = length(full$coefficients) - length(red$coefficients))
```

```
## [1] 0.08981025
```

```
red
```

```
## Call:
```

```
## coxph(formula = Surv(time, censor) ~ tx * (sex + priorzdv) +  
##       raceth * sex + raceth * karnof + ivdrug * priorzdv + cd4,  
##       data = AD)
```

```
##
```

	coef	exp(coef)	se(coef)	z	p
## tx	-1.555087	0.211171	0.864569	-1.799	0.07207
## sex	2.283299	9.808991	0.877912	2.601	0.00930
## priorzdv	-0.013985	0.986112	0.010731	-1.303	0.19251
## raceth	4.287019	72.749312	1.625738	2.637	0.00837
## karnof	-0.009523	0.990523	0.030620	-0.311	0.75581
## ivdrug	-0.837568	0.432762	0.351801	-2.381	0.01728
## cd4	-0.015009	0.985103	0.003152	-4.762	1.91e-06
## tx:sex	1.473724	4.365463	0.668784	2.204	0.02755
## tx:priorzdv	-0.034477	0.966111	0.014294	-2.412	0.01587
## sex:raceth	-1.480506	0.227523	0.587672	-2.519	0.01176
## raceth:karnof	-0.032507	0.968016	0.018972	-1.713	0.08663
## priorzdv:ivdrug	0.016117	1.016247	0.007143	2.256	0.02405

```
##
```

```
## Likelihood ratio test=107.3 on 12 df, p=< 2.2e-16
```

```
## n= 851, number of events= 69
```

Let's try raceth*karnof

```
full = coxph(Surv(time, censor) ~ tx*(sex + priorzdv) + raceth*sex + raceth*karnof + ivdrug*priorzdv + cd4, data = AD)
full$loglik[2]
```

```
## [1] -398.9897
```

```
red = coxph(Surv(time, censor) ~ tx*(sex + priorzdv) + raceth*sex + karnof + ivdrug*priorzdv + cd4, data = AD)
red$loglik[2]
```

```
## [1] -400.5166
```

```
1 - pchisq(2 * full$loglik[2] - 2 * red$loglik[2], df = length(full$coefficients) - length(red$coefficients))
```

```
## [1] 0.08054651
```

```
red
```

```
## Call:
```

```
## coxph(formula = Surv(time, censor) ~ tx * (sex + priorzdv) +  
##       raceth * sex + karnof + ivdrug * priorzdv + cd4, data = AD)
```

```
##
##               coef exp(coef) se(coef)      z      p
## tx            -1.453900  0.233657  0.862695 -1.685 0.091931
## sex            2.469187 11.812844  0.866826  2.849 0.004392
## priorzdv      -0.015350  0.984767  0.010419 -1.473 0.140691
## raceth         1.670770  5.316262  0.625436  2.671 0.007554
## karnof         -0.056287  0.945268  0.014591 -3.857 0.000115
## ivdrug         -0.855328  0.425144  0.337158 -2.537 0.011185
## cd4            -0.015267  0.984849  0.003145 -4.854 1.21e-06
## tx:sex          1.392238  4.023845  0.666413  2.089 0.036694
## tx:priorzdv     -0.034642  0.965951  0.014195 -2.441 0.014666
## sex:raceth      -1.599834  0.201930  0.578075 -2.768 0.005648
## priorzdv:ivdrug  0.017395  1.017547  0.006697  2.597 0.009395
##
## Likelihood ratio test=104.2 on 11 df, p=< 2.2e-16
## n= 851, number of events= 69
```

Okay all the interaction terms are now below 0.05 and the only p-values above 0.05 are variables that are also in interaction terms. So, I have a model. Let's check against the full model with all interaction just to be sure.

```
full = coxph(Surv(time, censor) ~ (tx+strat2+sex+raceth+ivdrug+hemophil+karnof+cd4+priorzdv+age)^2, data=AD)
red = coxph(Surv(time, censor) ~ tx*(sex + priorzdv) + raceth*sex + ivdrug*priorzdv + karnof + cd4, data=AD)
1 - pchisq(2 * full$loglik[2] - 2 * red$loglik[2], df = length(full$coefficients) - length(red$coefficients))
```

```
## [1] 0.6505501
```

Alright that works then.

The model is:

```
model = coxph(Surv(time, censor) ~ tx*(sex + priorzdv) + raceth*sex + ivdrug*priorzdv + karnof + cd4, data=AD)
model
```

```
## Call:
## coxph(formula = Surv(time, censor) ~ tx * (sex + priorzdv) +
##       raceth * sex + ivdrug * priorzdv + karnof + cd4, data = AD)
##
##               coef exp(coef) se(coef)      z      p
## tx            -1.453900  0.233657  0.862695 -1.685 0.091931
## sex            2.469187 11.812844  0.866826  2.849 0.004392
## priorzdv      -0.015350  0.984767  0.010419 -1.473 0.140691
## raceth         1.670770  5.316262  0.625436  2.671 0.007554
## ivdrug         -0.855328  0.425144  0.337158 -2.537 0.011185
## karnof         -0.056287  0.945268  0.014591 -3.857 0.000115
## cd4            -0.015267  0.984849  0.003145 -4.854 1.21e-06
## tx:sex          1.392238  4.023845  0.666413  2.089 0.036694
## tx:priorzdv     -0.034642  0.965951  0.014195 -2.441 0.014666
## sex:raceth      -1.599834  0.201930  0.578075 -2.768 0.005648
## priorzdv:ivdrug  0.017395  1.017547  0.006697  2.597 0.009395
##
## Likelihood ratio test=104.2 on 11 df, p=< 2.2e-16
## n= 851, number of events= 69
```

Let's test removing tx from the model to see what that does to the p-value so I can talk about it in my results.

```

model = coxph(Surv(time, censor) ~ tx*(sex + priorzdv) + raceth*sex + ivdrug*priorzdv + karnof + cd4, data = AD)
modelNoTx = coxph(Surv(time, censor) ~ raceth*sex + ivdrug*priorzdv + karnof + cd4, data = AD)
1 - pchisq(2 * model$loglik[2] - 2 * modelNoTx$loglik[2] , df = length(model$coefficients) - length(modelNoTx$coefficients))

```

```
## [1] 0.000307816
```

P value of 0.000307816 that model is the same without tx (so model is not the same)