

Multi-Task Neural Learning for Survival Analysis: from traditional to machine learning models

Diego Vallarino

Junio 2022

Contents

1	Descriptive	1
2	Train & Test data	6
3	Primeros analisis	6
4	Relevancia de las variables a utilizar usando XGBoost	7
5	Modelo de Tobit - parametric survival model	9
6	Kaplan-Meier Model - non parametric survival model	10
7	Cox models - semi parametric survival model	11
8	MTLR Model - machine learning model	14
9	Cantidad de observaciones censuradas a la derecha	15
10	Conclusion	16

1 Descriptive

```
head(veteran)
```

```
##   trt celltype time status karno diagtime age prior
## 1   1 squamous  72      1   60        7  69     0
## 2   1 squamous 411      1   70        5  64    10
## 3   1 squamous 228      1   60        3  38     0
## 4   1 squamous 126      1   60        9  63    10
## 5   1 squamous 118      1   70       11  65    10
## 6   1 squamous  10      1   20        5  49     0
```

```
str(veteran)
```

```
## 'data.frame': 137 obs. of 8 variables:
## $ trt : num 1 1 1 1 1 1 1 1 1 1 ...
## $ celltype: Factor w/ 4 levels "squamous","smallcell",...: 1 1 1 1 1 1 1 1 1 1 ...
## $ time : num 72 411 228 126 118 10 82 110 314 100 ...
## $ status : num 1 1 1 1 1 1 1 1 1 0 ...
## $ karno : num 60 70 60 60 70 20 40 80 50 70 ...
## $ diagtime: num 7 5 3 9 11 5 10 29 18 6 ...
## $ age : num 69 64 38 63 65 49 69 68 43 70 ...
## $ prior : num 0 10 0 10 10 0 10 0 0 0 ...
```

```
describeBy(veteran, veteran$celltype)
```

```
##
## Descriptive statistics by group
## group: squamous
##      vars  n   mean    sd median trimmed   mad min max range  skew
## trt      1 35   1.57   0.50      2    1.59   0.00   1  2     1 -0.28
## celltype* 2 35   1.00   0.00      1    1.00   0.00   1  1     0  NaN
## time      3 35 200.20 248.23    111 152.41 142.33   1 999   998  1.91
## status     4 35   0.89   0.32      1    0.97   0.00   0  1     1 -2.32
## karno      5 35  60.86  20.49     60  62.07  14.83   20  90    70 -0.49
## diagtime   6 35  11.03  11.53      7    8.90   5.93   1  58    57  2.32
## age        7 35  58.46  10.37     62  59.03  10.38  35  81    46 -0.40
## prior      8 35   4.00   4.97      0    3.79   0.00   0  10    10  0.39
##      kurtosis  se
## trt          -1.98 0.08
## celltype*      NaN 0.00
## time           3.35 41.96
## status          3.49 0.05
## karno          -0.69 3.46
## diagtime        5.92 1.95
## age            -0.41 1.75
## prior          -1.90 0.84
## -----
## group: smallcell
##      vars  n   mean    sd median trimmed   mad min max range  skew kurtosis
## trt      1 48   1.38   0.49     1.0    1.35   0.00   1  2     1  0.50    -1.79
## celltype* 2 48   2.00   0.00     2.0    2.00   0.00   2  2     0  NaN      NaN
## time      3 48  71.67  85.77    51.0   55.10  50.41   2 392   390  2.35     5.68
## status     4 48   0.94   0.24     1.0    1.00   0.00   0  1     1 -3.50    10.49
## karno      5 48  53.54  19.10    60.0   53.88  29.65  20  85    65 -0.15    -1.25
## diagtime   6 48   9.25  13.91     4.0    6.58   2.97   1  87    86  3.85    17.80
## age        7 48  59.88   9.92    62.5   60.83   8.90  35  72    37 -0.88    -0.25
## prior      8 48   2.29   4.25     0.0    1.75   0.00   0  10    10  1.25    -0.45
##      se
## trt      0.07
## celltype* 0.00
## time     12.38
## status    0.04
## karno     2.76
## diagtime  2.01
```

```
## age      1.43
## prior    0.61
## -----
## group: adeno
##      vars  n  mean    sd median trimmed   mad min max range  skew kurtosis
## trt      1 27  1.67  0.48     2    1.70  0.00    1  2     1 -0.67   -1.61
## celltype* 2 27  3.00  0.00     3    3.00  0.00    3  3     0  NaN     NaN
## time      3 27 64.11 50.59    51   59.70 48.93    3 186   183  0.71   -0.50
## status     4 27  0.96  0.19     1    1.00  0.00    0  1     1 -4.63   20.22
## karno      5 27 58.11 22.12    60   58.70 29.65   10  99    89 -0.20   -0.83
## diagtime   6 27  5.63  4.76     4    4.65  1.48    2  22    20  2.29    4.57
## age        7 27 57.41 11.32    61   57.70  5.93   34  81    47 -0.43   -0.54
## prior      8 27  1.85  3.96     0    1.30  0.00    0  10    10  1.53    0.36
##      se
## trt      0.09
## celltype* 0.00
## time     9.74
## status    0.04
## karno     4.26
## diagtime  0.92
## age       2.18
## prior     0.76
## -----
## group: large
##      vars  n  mean    sd median trimmed   mad min max range  skew
## trt      1 27  1.44  0.51     1    1.43  0.00    1  2     1  0.21
## celltype* 2 27  4.00  0.00     4    4.00  0.00    4  4     0  NaN
## time      3 27 166.11 124.22   156  153.35 111.19   12 553   541  1.13
## status     4 27  0.96  0.19     1    1.00  0.00    0  1     1 -4.63
## karno      5 27  65.00 17.49    70   65.87 14.83   30  90    60 -0.66
## diagtime   6 27  8.15  4.99     8    7.96  5.93    1  18    17  0.27
## age        7 27  56.22 11.16    62   56.87  7.41   37  68    31 -0.57
## prior      8 27  3.70  4.92     0    3.48  0.00    0  10    10  0.51
##      kurtosis    se
## trt      -2.03  0.10
## celltype*   NaN  0.00
## time       1.36 23.91
## status     20.22 0.04
## karno      -0.57 3.37
## diagtime   -1.32 0.96
## age        -1.38 2.15
## prior      -1.81 0.95
```

```
describeBy(veteran, veteran$trt)
```

```
##
## Descriptive statistics by group
## group: 1
##      vars  n  mean    sd median trimmed   mad min max range  skew
## trt      1 69  1.00  0.00     1    1.00  0.00    1  1     0  NaN
## celltype* 2 69  2.35  1.05     2    2.32  1.48    1  4     3  0.40
## time      3 69 115.14 112.74   97   97.61 97.85    3 553   550  1.55
## status     4 69  0.93  0.26     1    1.00  0.00    0  1     1 -3.23
## karno      5 69  59.20 18.74    60   60.26 29.65   20  90    70 -0.40
```

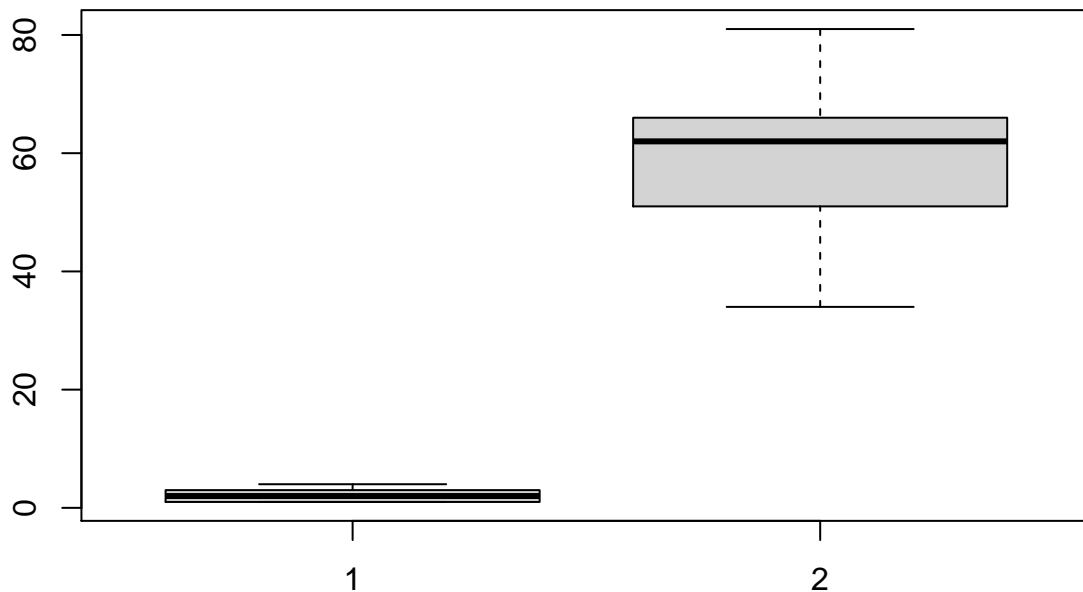
```
## diagtime      6 69    8.65    8.76      5    7.11  4.45    1 58    57 3.02
## age           7 69   57.51   10.81     62   58.21  8.90   34 81    47 -0.54
## prior         8 69    3.04    4.64      0    2.63  0.00    0 10    10 0.83
##               kurtosis    se
## trt              NaN    0.00
## celltype*       -1.09  0.13
## time            2.57 13.57
## status          8.54  0.03
## karno          -0.92  2.26
## diagtime       12.79  1.05
## age            -0.74  1.30
## prior         -1.33  0.56
## -----
## group: 2
##      vars  n   mean      sd median trimmed   mad min max range skew
## trt      1 68    2.00    0.00     2.0    2.00  0.00    2  2     0   NaN
## celltype* 2 68    2.32    1.09     2.0    2.29  1.48    1  4     3  0.17
## time      3 68 128.21 193.83    52.5   87.16 55.60    1 999   998  2.95
## status     4 68    0.94    0.24     1.0    1.00  0.00    0  1     1 -3.67
## karno      5 68   57.93   21.40    60.0   58.39 29.65   10 99    89 -0.27
## diagtime   6 68    8.90   12.27     4.5    6.52  3.71    1 87    86  4.16
## age        7 68   59.12   10.28    62.0   59.96  8.15   35 81    46 -0.70
## prior      8 68    2.79    4.52     0.0    2.32  0.00    0 10    10  0.96
##               kurtosis    se
## trt              NaN    0.00
## celltype*       -1.30  0.13
## time            9.56 23.50
## status         11.62  0.03
## karno          -0.89  2.59
## diagtime       21.99  1.49
## age            -0.28  1.25
## prior         -1.09  0.55
```

```
describeBy(veteran, veteran$prior)
```

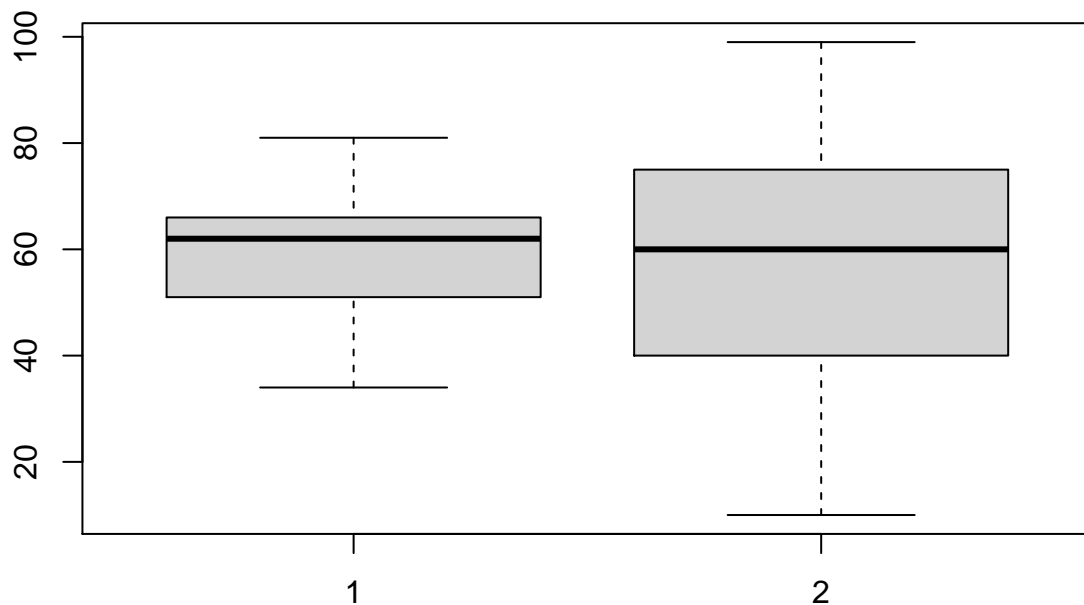
```
##
## Descriptive statistics by group
## group: 0
##      vars  n   mean      sd median trimmed   mad min max range skew
## trt      1 97    1.51    0.50     2    1.51  0.00    1  2     1 -0.02
## celltype* 2 97    2.36    1.01     2    2.33  1.48    1  4     3  0.26
## time      3 97 112.15 121.94    80   88.65 78.58    1 587   586  1.88
## status     4 97    0.94    0.24     1    1.00  0.00    0  1     1 -3.58
## karno      5 97   59.32   20.50    60   60.13 29.65   10 99    89 -0.38
## diagtime   6 97    5.94    5.87     4    4.81  2.97    1 36    35  2.77
## age        7 97   58.98   10.71    62   59.82  7.41   34 81    47 -0.70
## prior      8 97    0.00    0.00     0    0.00  0.00    0  0     0   NaN
##               kurtosis    se
## trt          -2.02  0.05
## celltype*    -1.05  0.10
## time          3.42 12.38
## status       10.94  0.02
## karno        -0.77  2.08
## diagtime      9.23  0.60
```

```
## age          -0.32  1.09
## prior         NaN  0.00
## -----
## group: 10
##      vars  n   mean    sd median trimmed   mad min max range skew
## trt      1 40   1.48   0.51     1    1.47  0.00   1  2     1  0.10
## celltype* 2 40   2.28   1.20     2    2.22  1.48   1  4     3  0.35
## time      3 40 144.60 222.45    69   94.50 84.51   1 999   998  2.78
## status     4 40   0.92   0.27     1    1.00  0.00   0  1     1 -3.11
## karno      5 40  56.75 19.00    60   57.50 29.65  20  90    70 -0.24
## diagtime   6 40  15.65 15.48    12   12.59  6.67   2  87    85  2.94
## age        7 40  56.67 10.07    60   57.47 10.38  36  70    34 -0.51
## prior      8 40  10.00  0.00    10   10.00  0.00  10  10     0   NaN
##      kurtosis    se
## trt      -2.04  0.08
## celltype* -1.46  0.19
## time       7.80 35.17
## status     7.85  0.04
## karno     -1.13  3.00
## diagtime   9.86  2.45
## age       -1.01  1.59
## prior      NaN  0.00
```

```
boxplot(veteran$celltype, veteran$age)
```



```
boxplot(veteran$age, veteran$karno)
```



```
any(is.na(veteran))
```

```
## [1] FALSE
```

2 Train & Test data

```
set.seed(123)
data.train <- sample_frac(veteran, 0.7)
train_index <- as.numeric(rownames(data.train))
data.test <- veteran [-train_index, ]
```

3 Primeros analisis

```
survdif(Surv(time, status) ~ trt, data = data.train)
```

```
## Call:
```

```
## survdiff(formula = Surv(time, status) ~ trt, data = data.train)
##
##           N Observed Expected (O-E)^2/E (O-E)^2/V
## trt=1 44      39      33.6      0.851      1.53
## trt=2 52      49      54.4      0.527      1.53
##
## Chisq= 1.5  on 1 degrees of freedom, p= 0.2
```

```
survdiff(Surv(time, status) ~ celltype, data = data.train)
```

```
## Call:
## survdiff(formula = Surv(time, status) ~ celltype, data = data.train)
##
##           N Observed Expected (O-E)^2/E (O-E)^2/V
## celltype=squamous 26      23      38.2      6.046      13.02
## celltype=smallcell 36      33      21.0      6.923      10.14
## celltype=adeno    18      17      10.8      3.518       4.22
## celltype=large    16      15      18.0      0.506       0.66
##
## Chisq= 21.1  on 3 degrees of freedom, p= 1e-04
```

```
survdiff(Surv(time, status) ~ prior + status, data = data.train)
```

```
## Call:
## survdiff(formula = Surv(time, status) ~ prior + status, data = data.train)
##
##           N Observed Expected (O-E)^2/E (O-E)^2/V
## prior=0, status=0  6       0      5.51  5.50599      6.0703
## prior=0, status=1 63      63     55.94  0.89111      2.5941
## prior=10, status=0  2       0      1.13  1.13051      1.1684
## prior=10, status=1 25      25     25.42  0.00707      0.0109
##
## Chisq= 7.8  on 3 degrees of freedom, p= 0.05
```

4 Relevancia de las variables a utilizar usando XGBoost

```
require(xgboost)
require(Matrix)
require(data.table)

df <- data.table(data.train, keep.rownames = FALSE)
sparse_matrix <- sparse.model.matrix(status~., data = data.train)
head(sparse_matrix)
```

```
## 6 x 10 sparse Matrix of class "dgCMatrix"
##
## 14  1 1 . . . 25 80  9 52 10
## 50  1 . . 1 . 132 80  5 50 .
## 118 2 . . 1 . 48 10  4 81 .
```

```
## 43  1 . 1 . . 63 50 11 48 .
## 137 2 . . . 1 49 30 3 37 .
## 135 2 . . . 1 231 70 18 67 10
```

```
output_vector = df[,status]
```

```
# desarrollo el modelo de relevancia
```

```
bst <- xgboost(data = sparse_matrix, label = output_vector, max.depth = 4, eta = 1, nthread = 2, nround
```

```
## [10:59:59] WARNING: amalgamation/./src/learner.cc:1115: Starting in XGBoost 1.3.0, the default eval
```

```
## [1] train-logloss:0.310176
## [2] train-logloss:0.202168
## [3] train-logloss:0.148494
## [4] train-logloss:0.123007
## [5] train-logloss:0.104148
## [6] train-logloss:0.089287
## [7] train-logloss:0.083504
## [8] train-logloss:0.074024
## [9] train-logloss:0.067894
## [10] train-logloss:0.064258
```

```
# Medimos los la importancia de la variables.
```

```
importance <- xgb.importance(feature_names = sparse_matrix@Dimnames[[2]], model = bst)
head(importance)
```

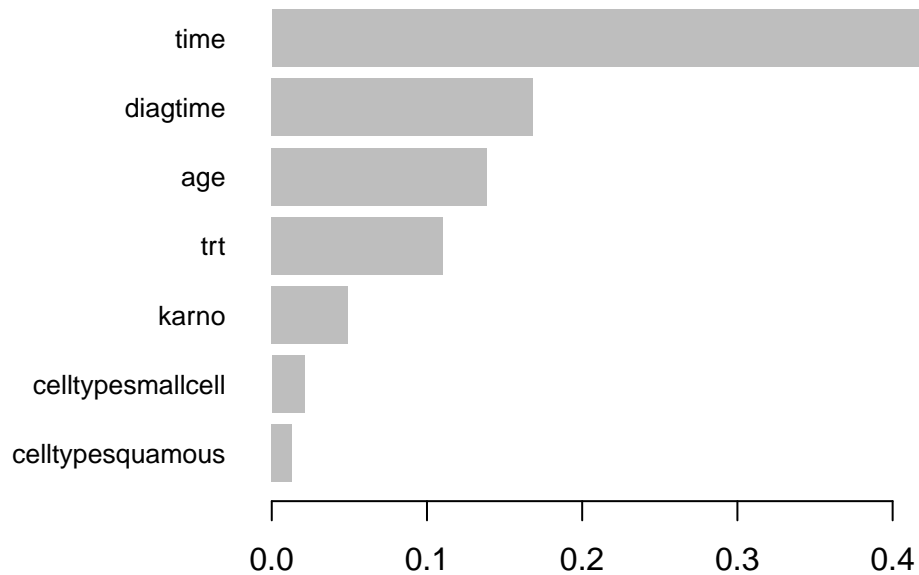
```
##           Feature      Gain      Cover  Frequency
## 1:           time 0.49967836 0.34390974 0.32258065
## 2:        diagtime 0.16815368 0.12704402 0.16129032
## 3:            age 0.13858913 0.14225801 0.19354839
## 4:            trt 0.11037545 0.10794435 0.12903226
## 5:           karno 0.04906092 0.21078392 0.09677419
## 6: celltypesmallcell 0.02099943 0.02759307 0.06451613
```

```
# Mejora en la interpretabilidad de la tabla de datos de importancia de características
```

```
importanceRaw <- xgb.importance(feature_names = sparse_matrix@Dimnames[[2]], model = bst, data = sparse
importanceClean <- importanceRaw[,`:=`(Cover=NULL, Frequency=NULL)]
head(importanceClean)
```

```
##           Feature      Gain
## 1:           time 0.49967836
## 2:        diagtime 0.16815368
## 3:            age 0.13858913
## 4:            trt 0.11037545
## 5:           karno 0.04906092
## 6: celltypesmallcell 0.02099943
```

```
xgb.plot.importance(importance_matrix = importanceRaw)
```

5 Modelo de Tobit - parametric survival model

```
surv_obj = Surv(data.test$time, data.test$status)
fit2 <- survreg(Surv(time, status) ~ karno + age + trt, data=data.train)

predictfit2<-predict(fit2, data.test)
metrics_fit2<-Cindex(surv_obj, predicted = predictfit2)

summary(fit2)
```

```
##
## Call:
## survreg(formula = Surv(time, status) ~ karno + age + trt, data = data.train)
##              Value Std. Error      z      p
## (Intercept)  2.76122    0.75410  3.66 0.00025
## karno        0.03556    0.00519  6.85 7.3e-12
## age         -0.00525    0.01073 -0.49 0.62432
## trt          0.12805    0.20984  0.61 0.54170
## Log(scale)  -0.05965    0.07847 -0.76 0.44718
##
## Scale= 0.942
##
## Weibull distribution
```

```
## Loglik(model)= -496.5   Loglik(intercept only)= -517.3
##  Chisq= 41.4 on 3 degrees of freedom, p= 5.4e-09
## Number of Newton-Raphson Iterations: 5
## n= 96
```

6 Kaplan-Meier Model - non parametric survival model

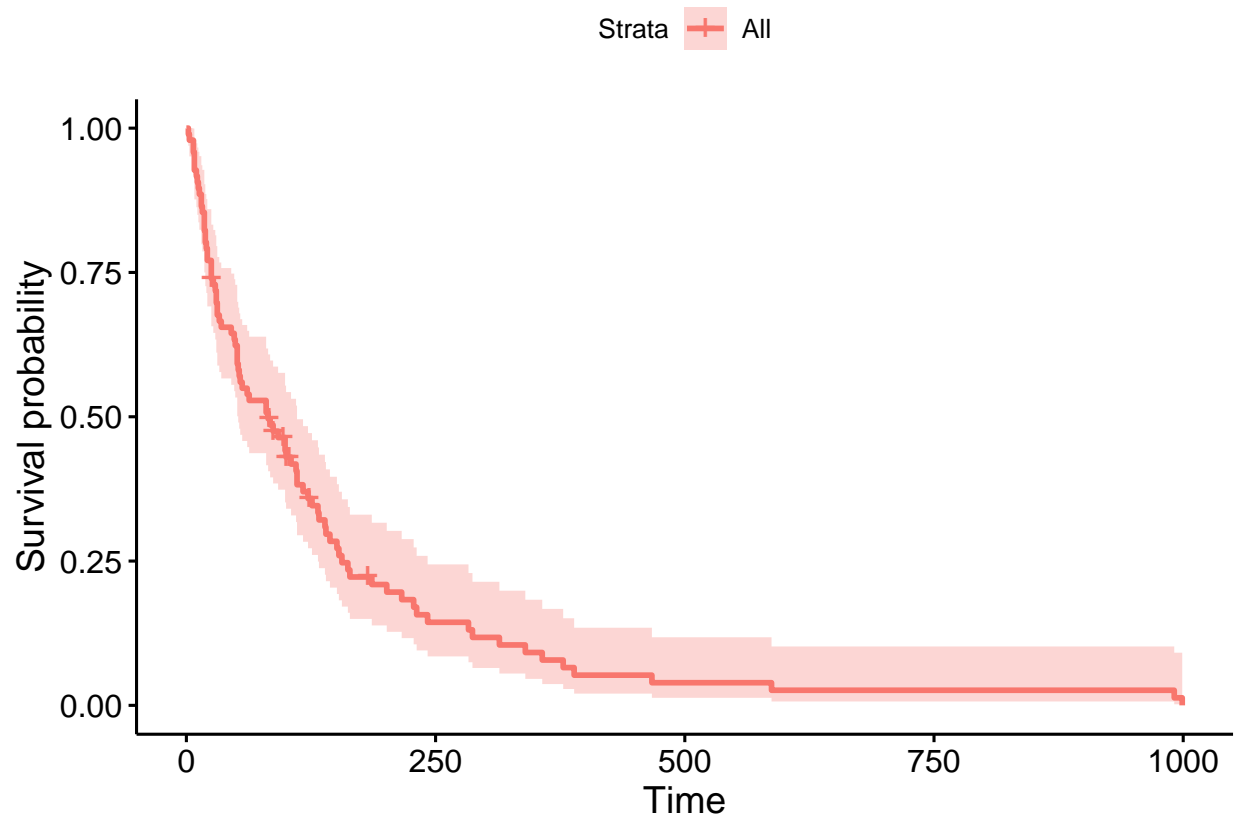
```
fit3<-survfit(Surv(time, status) ~ 1, data = data.train)
dis_timefit3 = fit3$time

med_indexfit3 = median(1:length(dis_timefit3))

predictfit3<-predictSurvProb(fit3, data.test, dis_timefit3)

metrics_fit3 = Cindex(surv_obj, predicted = predictfit3[, med_indexfit3])

ggsurvplot(fit3, data = veteran, pval = TRUE)
```



```
print(fit3, print.rmean=TRUE)
```

```
## Call: survfit(formula = Surv(time, status) ~ 1, data = data.train)
##
```

```
##           n      events      *rmean *se(rmean)      median      0.95LCL      0.95UCL
##          96         88         137         20         82         52         111
##      * restricted mean with upper limit = 999
```

```
summary(fit3, times=c(20, 50, 100, 350))
```

```
## Call: survfit(formula = Surv(time, status) ~ 1, data = data.train)
##
##   time n.risk n.event survival std.err lower 95% CI upper 95% CI
##    20     77     20   0.7917  0.0414   0.7145   0.877
##    50     59     16   0.6234  0.0496   0.5333   0.729
##   100     39     18   0.4300  0.0511   0.3406   0.543
##   350      7     27   0.0916  0.0324   0.0458   0.183
```

7 Cox models - semi parametric survival model

```
fit4 <- coxph(Surv(time, status) ~ ., data=data.train, x = TRUE)
shapiro.test(fit4$residuals)
```

```
##
##  Shapiro-Wilk normality test
##
## data:  fit4$residuals
## W = 0.83037, p-value = 4.037e-09
```

```
anova(fit4)
```

```
## Analysis of Deviance Table
## Cox model: response is Surv(time, status)
## Terms added sequentially (first to last)
##
##           loglik    Chisq Df Pr(>|Chi|)
## NULL          -316.20
## trt           -315.47  1.4742  1  0.2246861
## celltype     -305.38 20.1670  3  0.0001567 ***
## karno        -286.84 37.0863  1  1.13e-09 ***
## diagtime     -286.77  0.1366  1  0.7116438
## age          -285.91  1.7250  1  0.1890525
## prior        -285.91  0.0008  1  0.9779353
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

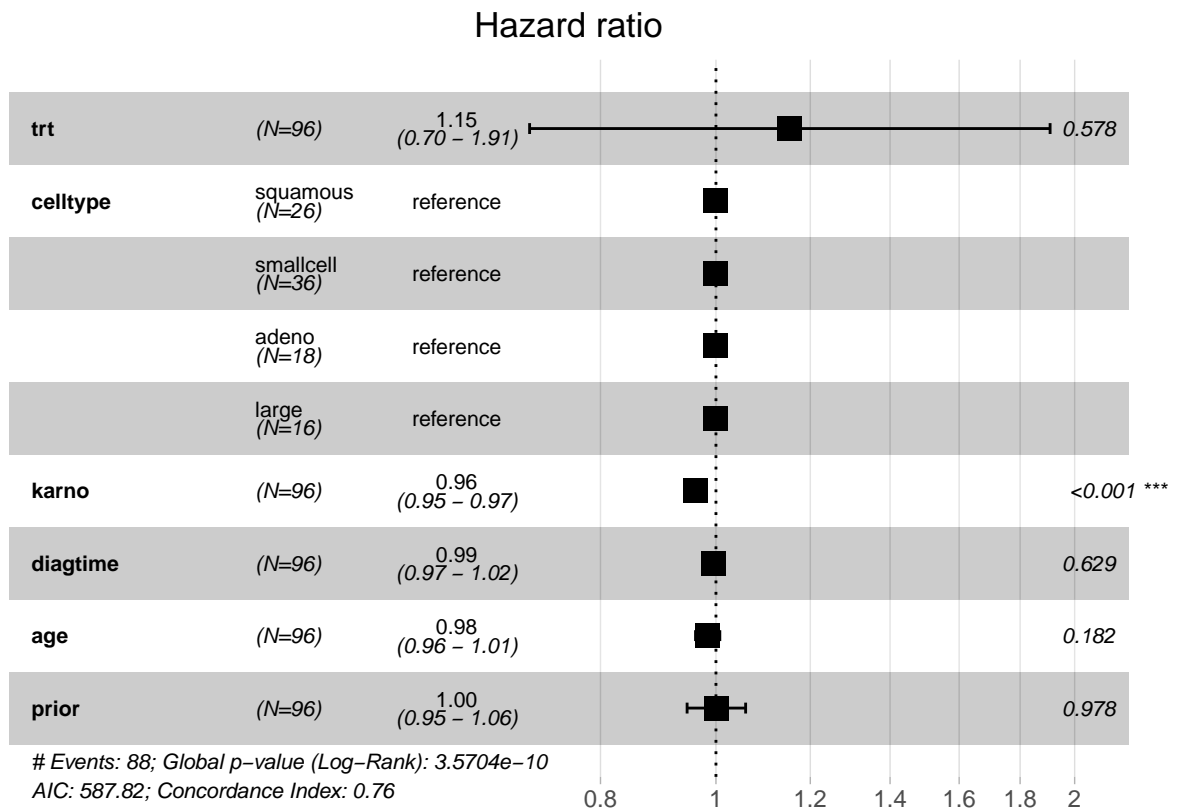
```
dis_timefit4 = fit3$time
```

```
med_indexfit4 = median(1:length(dis_timefit4))
predictfit4<-predictSurvProb(fit4, data.test, dis_timefit4)
metrics_fit4 = Cindex(surv_obj, predicted = predictfit4[, med_indexfit4])

summary(fit4)
```

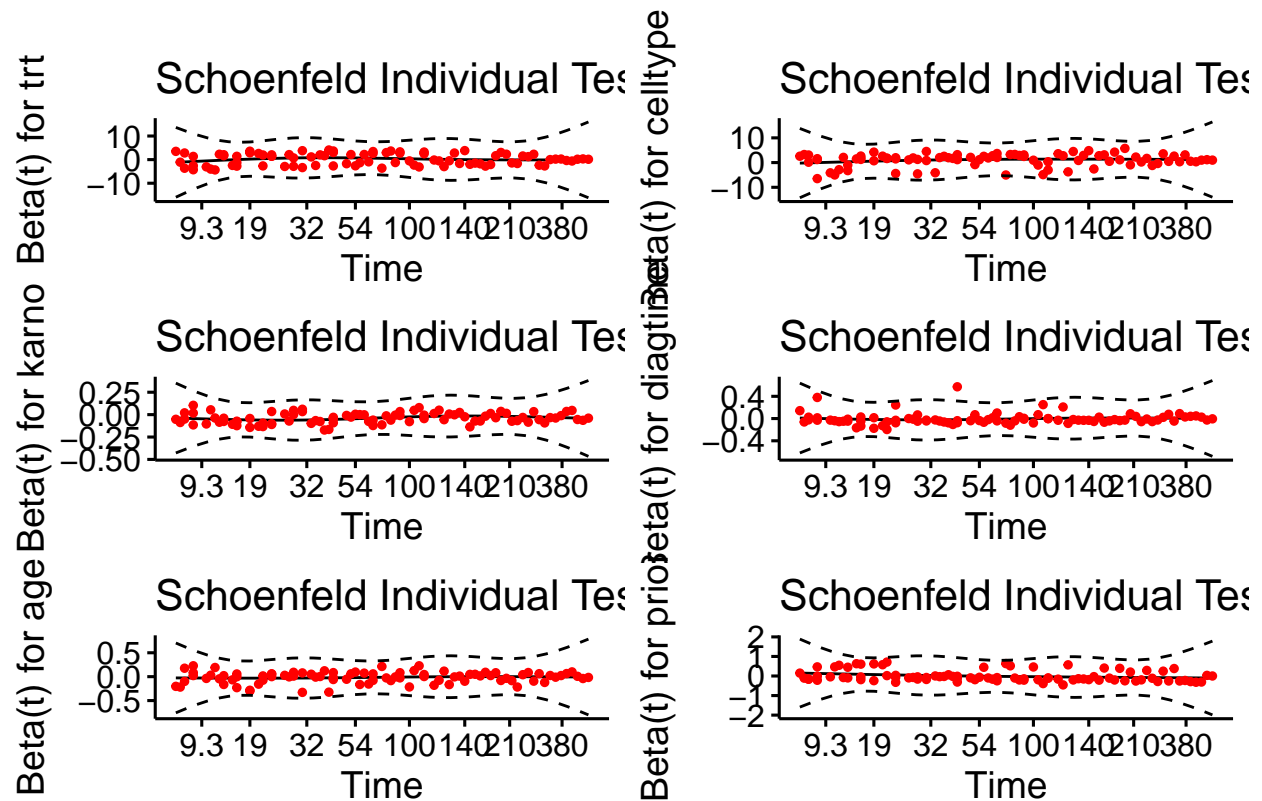
```
## Call:
## coxph(formula = Surv(time, status) ~ ., data = data.train, x = TRUE)
##
##    n= 96, number of events= 88
##
##              coef exp(coef)    se(coef)      z Pr(>|z|)
## trt           0.1428218  1.1535242  0.2565892  0.557  0.57779
## celltype1    -0.9161660  0.4000499  0.2341873 -3.912 9.15e-05 ***
## celltype2     0.4729743  1.6047601  0.1992676  2.374  0.01762 *
## celltype3     0.6220558  1.8627536  0.2321140  2.680  0.00736 **
## karno        -0.0394631  0.9613054  0.0064027 -6.164 7.11e-10 ***
## diagtime    -0.0050613  0.9949515  0.0104799 -0.483  0.62913
## age         -0.0159867  0.9841404  0.0119828 -1.334  0.18216
## prior         0.0007983  1.0007986  0.0288504  0.028  0.97793
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
##              exp(coef) exp(-coef) lower .95 upper .95
## trt              1.1535    0.8669    0.6976    1.9074
## celltype1         0.4000    2.4997    0.2528    0.6331
## celltype2         1.6048    0.6231    1.0859    2.3715
## celltype3         1.8628    0.5368    1.1819    2.9358
## karno             0.9613    1.0403    0.9493    0.9734
## diagtime          0.9950    1.0051    0.9747    1.0156
## age               0.9841    1.0161    0.9613    1.0075
## prior             1.0008    0.9992    0.9458    1.0590
##
## Concordance= 0.759 (se = 0.029 )
## Likelihood ratio test= 60.59 on 8 df,  p=4e-10
## Wald test              = 55.67 on 8 df,  p=3e-09
## Score (logrank) test = 60.99 on 8 df,  p=3e-10
```

```
ggforest(fit4)
```



```
test_cox<-cox.zph(fit4)
ggcoxzph(test_cox)
```

Global Schoenfeld Test p: 0.02501



8 MTLR Model - machine learning model

```
fit7 <- mtlr(Surv(time, status)~., data = data.train, nintervals = 9)
dis_timefit7 = fit7$time_points
med_indexfit7 = median(1:length(dis_timefit7))

predictfit7<-predict(fit7, data.test, type = "mean_time")
metrics_fit7 = Cindex(surv_obj, predicted = predictfit7)

fit7
```

```
##
## Call: mtlr(formula = Surv(time, status) ~ ., data = data.train, nintervals = 9)
##
## Time points:
## [1] 11.6 19.0 26.8 46.6 56.9 87.0 107.3 133.5 177.1 296.8
##
##
## Weights:
##      Bias      trt celltype1 celltype2 celltype3  karno diagtime    age
## 11.64  0.1171 -0.03339  0.03081  0.000274  0.03571 -0.0197  0.03465 -0.01826
## 19     -0.0507 -0.03835  0.00333 -0.007206  0.00663 -0.0780  0.01243 -0.02928
## 26.82 -0.1685  0.01514 -0.00729  0.014363 -0.00533 -0.1320  0.00240 -0.03835
```

```
## 46.64 -0.0351 0.01607 0.01185 0.031709 0.03268 -0.1219 -0.02626 -0.02049
## 56.91 0.4227 0.01776 -0.03017 0.038213 0.02317 -0.1353 0.01017 -0.01955
## 87 -0.1226 0.03394 -0.02180 0.059672 0.05490 -0.1428 0.00457 0.00763
## 107.27 -0.2863 0.02987 -0.05675 0.059685 0.03929 -0.1099 -0.00350 0.00468
## 133.55 -0.0104 0.00147 -0.04636 0.049062 0.02854 -0.0682 0.00628 -0.00473
## 177.09 -0.0166 -0.04860 -0.06836 0.057887 0.03886 -0.0690 -0.00596 0.03760
## 296.82 -0.0524 -0.04662 -0.05469 0.035507 0.02726 -0.0490 0.01090 0.01674
## prior
## 11.64 0.01958
## 19 0.05089
## 26.82 0.06572
## 46.64 0.01226
## 56.91 0.00542
## 87 0.01663
## 107.27 0.00601
## 133.55 -0.01814
## 177.09 -0.02078
## 296.82 -0.00646
```

9 Cantidad de observaciones censuradas a la derecha

```
table(fit3$n.censor)
```

```
##
## 0 1
## 68 8
```

```
fit2$y
```

```
## 14 50 118 43 137 135 90 91 130 57 92 121 9 93 99 72
## 25+ 132 48 63 49 231 25 103+ 15 216 21 186 314 13 99 87+
## 26 7 42 125 83 36 78 81 134 103 76 15 32 106 120 132
## 16 82 7 80 467 287 587 25 111 25 111 11 139 51 140 340
## 41 74 23 27 60 53 107 100 102 96 38 89 34 69 122 111
## 54 242 153 151 12 3 29 8 61 20 51 15 31 100 84 31
## 63 13 82 25 95 21 79 105 47 101 16 6 129 39 31 136
## 156 144 357 117 2 123+ 389 80 92 99 30 10 53 122 18 378
## 124 4 88 127 86 52 22 109 70 112 35 40 48 30 12 75
## 45 126 283 164 30 162 97+ 18 999 51 52 27 35 21 8 991
## 128 46 80 94 133 29 66 123 3 64 110 84 37 8 10 119
## 19 8 33 87 133 56 105 19 228 182+ 83+ 201 18 110 100+ 7
```

```
fit4$y
```

```
## 14 50 118 43 137 135 90 91 130 57 92 121 9 93 99 72
## 25+ 132 48 63 49 231 25 103+ 15 216 21 186 314 13 99 87+
## 26 7 42 125 83 36 78 81 134 103 76 15 32 106 120 132
## 16 82 7 80 467 287 587 25 111 25 111 11 139 51 140 340
## 41 74 23 27 60 53 107 100 102 96 38 89 34 69 122 111
```

```
## 54 242 153 151 12 3 29 8 61 20 51 15 31 100 84 31
## 63 13 82 25 95 21 79 105 47 101 16 6 129 39 31 136
## 156 144 357 117 2 123+ 389 80 92 99 30 10 53 122 18 378
## 124 4 88 127 86 52 22 109 70 112 35 40 48 30 12 75
## 45 126 283 164 30 162 97+ 18 999 51 52 27 35 21 8 991
## 128 46 80 94 133 29 66 123 3 64 110 84 37 8 10 119
## 19 8 33 87 133 56 105 19 228 182+ 83+ 201 18 110 100+ 7
```

```
fit7$response
```

```
## 14 50 118 43 137 135 90 91 130 57 92 121 9 93 99 72
## 25+ 132 48 63 49 231 25 103+ 15 216 21 186 314 13 99 87+
## 26 7 42 125 83 36 78 81 134 103 76 15 32 106 120 132
## 16 82 7 80 467 287 587 25 111 25 111 11 139 51 140 340
## 41 74 23 27 60 53 107 100 102 96 38 89 34 69 122 111
## 54 242 153 151 12 3 29 8 61 20 51 15 31 100 84 31
## 63 13 82 25 95 21 79 105 47 101 16 6 129 39 31 136
## 156 144 357 117 2 123+ 389 80 92 99 30 10 53 122 18 378
## 124 4 88 127 86 52 22 109 70 112 35 40 48 30 12 75
## 45 126 283 164 30 162 97+ 18 999 51 52 27 35 21 8 991
## 128 46 80 94 133 29 66 123 3 64 110 84 37 8 10 119
## 19 8 33 87 133 56 105 19 228 182+ 83+ 201 18 110 100+ 7
```

10 Conclusion

```
metrics_fit2
```

```
## C index
## 0.6857319
```

```
metrics_fit3
```

```
## C index
## 0.50246
```

```
metrics_fit4
```

```
## C index
## 0.6863469
```

```
metrics_fit7
```

```
## C index
## 0.7207872
```

```
data_CI = data.frame(Cindex = c(metrics_fit2, metrics_fit3, metrics_fit4, metrics_fit7),
                     Model = c(rep('Tobit', 1), rep('KM', 1), rep('Cox', 1), rep('MTLR', 1)))

ggplot(data_CI, aes(x = Model, y = Cindex)) + geom_boxplot()
```