

Practical_survival_1_R.R

classale

Tue May 10 15:14:55 2016

```
#####  
## load packages ##  
#####  
#install.packages("survival")  
#install.packages("ggplot2")  
library(survival)  
  
## Loading required package: splines  
  
library(ggplot2)  
#####  
## 1. Set working directory and get data ##  
#####  
setwd("C:/Users/classale/STATS/")  
load("elsa_cf.rdata")  
attach(elsa_cf)  
  
#####  
## 2. Getting a sense of the data ##  
#####  
# Get list of variables in the dataset 'elsa_cf'  
names(elsa_cf)  
  
## [1] "cf1" "cancer1" "cigst1" "sex" "age1"  
## [6] "dodmnth" "dodyr" "educ1" "chd1" "totwq5_bu1"  
## [11] "iintdtm1" "iintdty1" "death" "alcohol1" "physinact1"  
## [16] "time" "id"  
  
# Get summary statistic for 'age1' and 'time'  
summary(elsa_cf$age1)  
  
## Min. 1st Qu. Median Mean 3rd Qu. Max.  
## 50.00 57.00 63.00 64.31 71.00 90.00  
  
summary(elsa_cf$time)  
  
## Min. 1st Qu. Median Mean 3rd Qu. Max.  
## 0.04107 10.20000 10.62000 9.60500 10.79000 11.04000  
  
# Histogram separately for people with an event and survivors #  
  
#Create a subset#  
elsadead <-subset(elsa_cf, elsa_cf$death==1)  
elsalive <-subset(elsa_cf, elsa_cf$death==0)  
  
summary(elsadead$time)
```

```
##      Min. 1st Qu.  Median    Mean 3rd Qu.    Max.
## 0.04107 3.20500 5.62400 5.38000 7.70700 9.96300
```

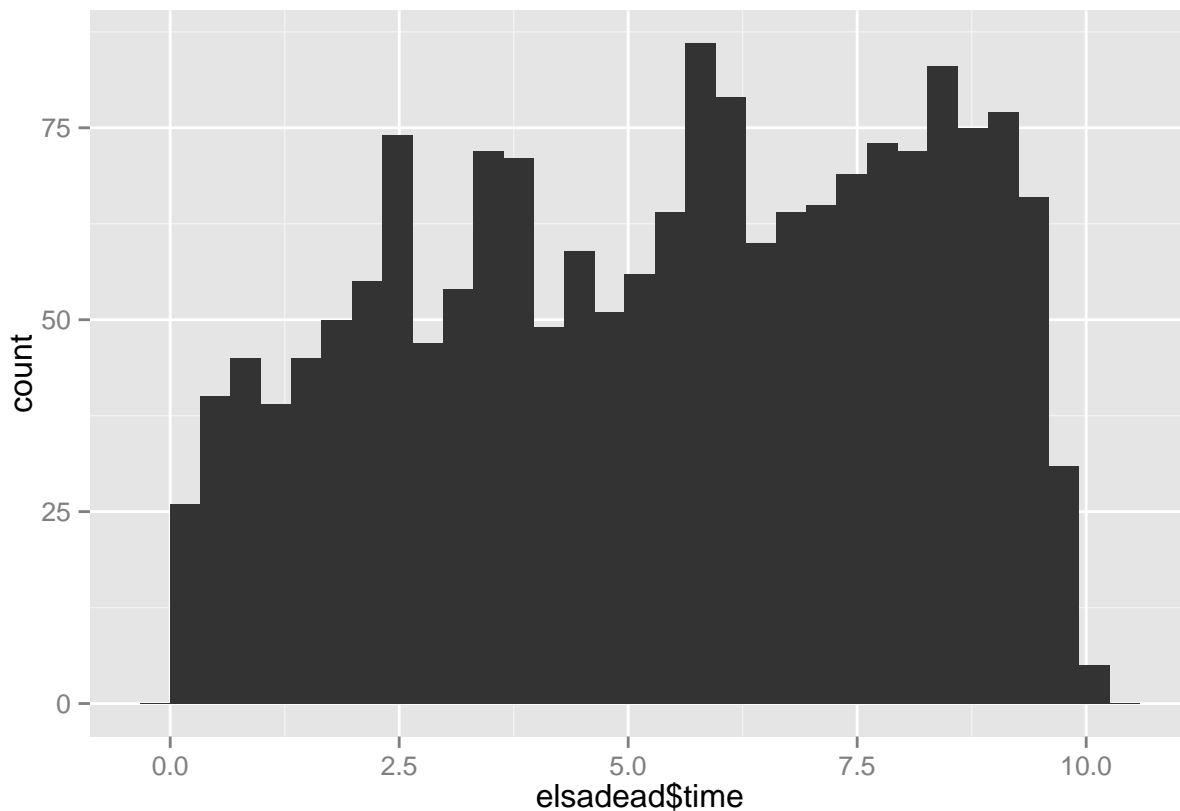
```
summary(elsalive$time)
```

```
##      Min. 1st Qu.  Median    Mean 3rd Qu.    Max.
## 10.04 10.46 10.71 10.62 10.79 11.04
```

```
# Histogram in each subset#
```

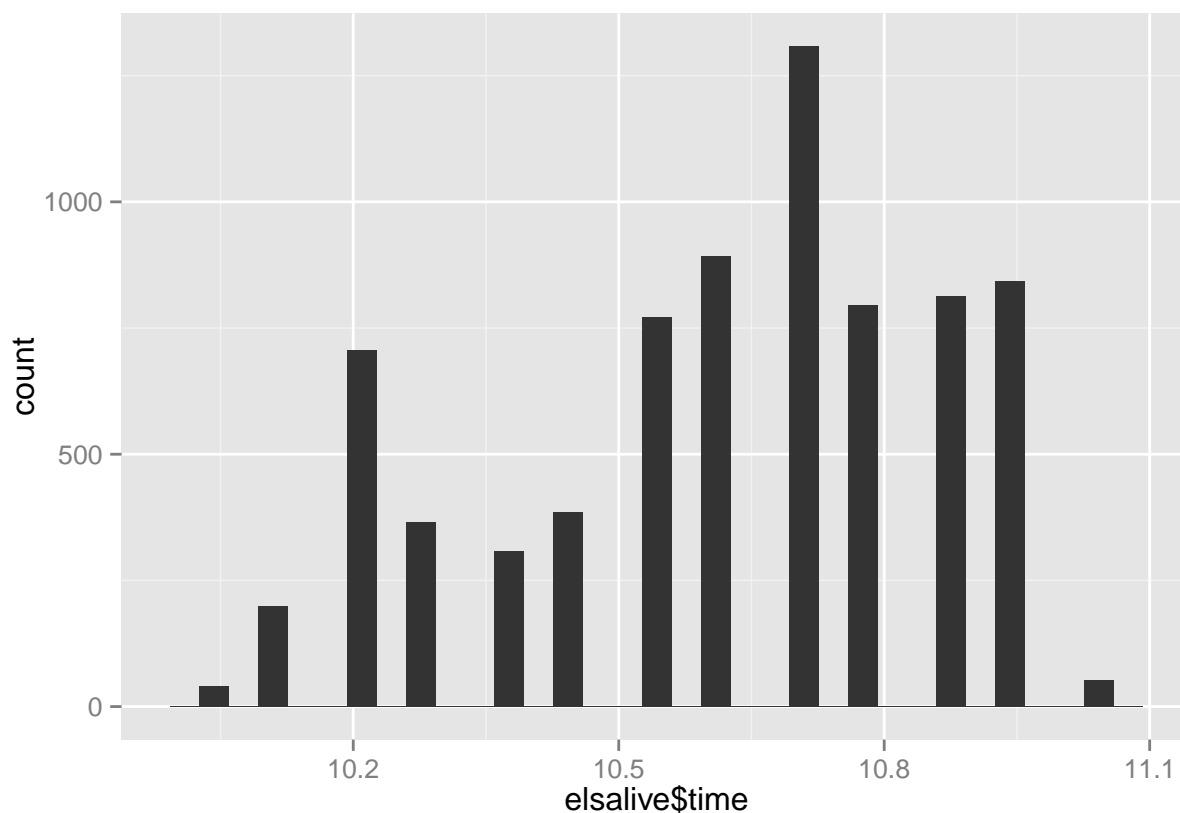
```
qplot(elsadead$time, geom = 'histogram', bins = 40)
```

```
## stat_bin: binwidth defaulted to range/30. Use 'binwidth = x' to adjust this.
```



```
qplot(elsalive$time, geom = 'histogram', bins = 40)
```

```
## stat_bin: binwidth defaulted to range/30. Use 'binwidth = x' to adjust this.
```



```
# Tabulate the outcome 'death'
table(elsa_cf$death)
```

```
##
##      0      1
## 7480 1802
```

```
prop.table(table(elsa_cf$death))
```

```
##
##           0           1
## 0.8058608 0.1941392
```

```
#####
## 3. Survival and KM curve                                     ##
#####
# Calculate person-years and incidence rate #
pyear <- sum(elsa_cf$time)
incidencerate <- 1802/pyear
print(pyear)
```

```
## [1] 89155.01
```

```
print(incidencerate)
```

```
## [1] 0.02021199
```

```
# This gives a summary of survival time again  
summary(Surv(time, death))
```

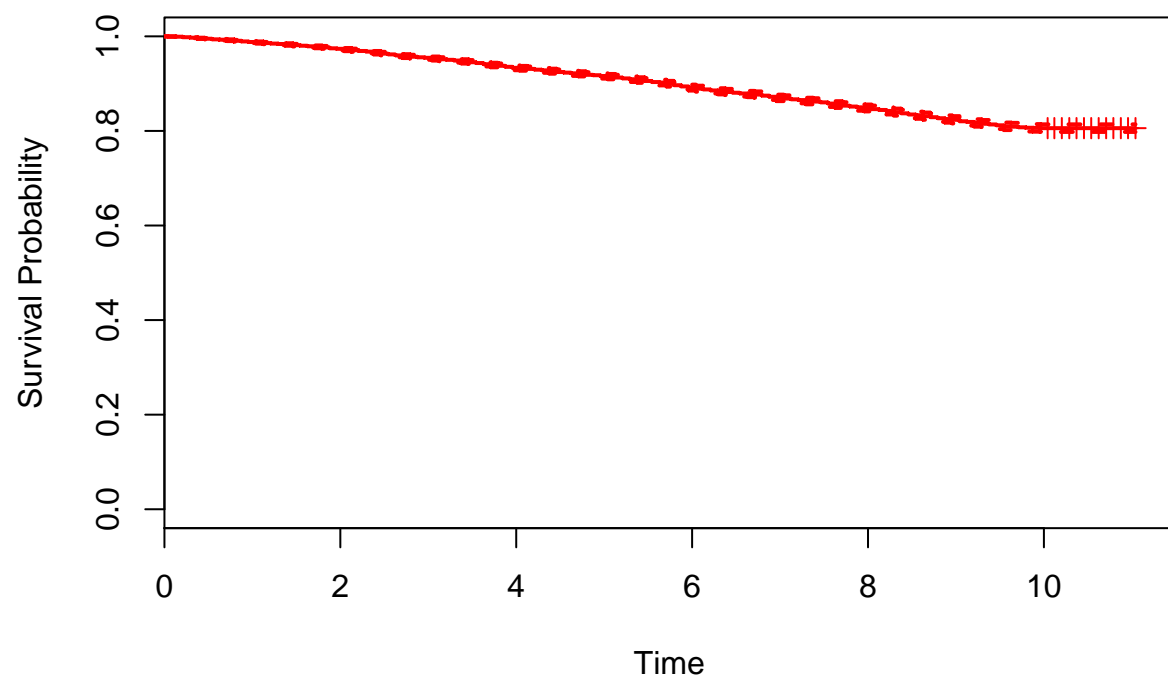
```
##      time      status  
## Min.   : 0.04107   Min.   :0.0000  
## 1st Qu.:10.20397   1st Qu.:0.0000  
## Median :10.62286   Median :0.0000  
## Mean   : 9.60515   Mean    :0.1941  
## 3rd Qu.:10.78987   3rd Qu.:0.0000  
## Max.   :11.04175   Max.    :1.0000
```

```
# Create a survival object and return summary #  
km <- survfit(Surv(time, death)~ 1)  
km
```

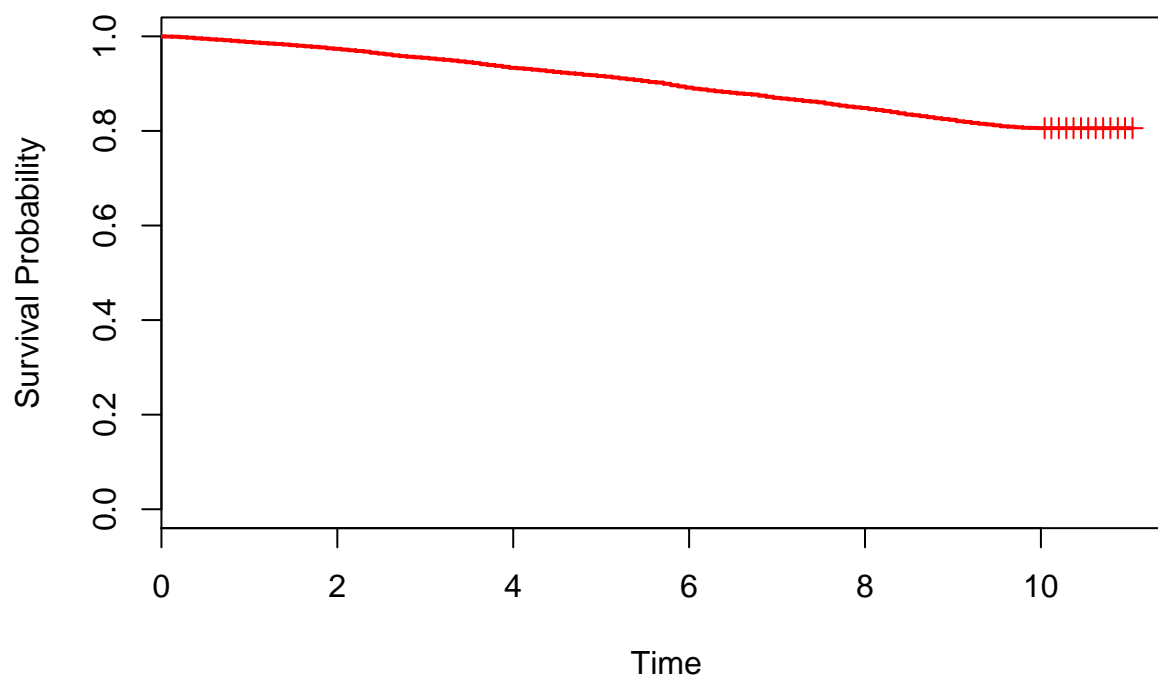
```
## Call: survfit(formula = Surv(time, death) ~ 1)  
##  
## records    n.max n.start  events  median 0.95LCL 0.95UCL  
##    9282     9282    9282    1802     NA      NA      NA
```

```
# It doesn't give any value for the median survival
```

```
# Let's assess this graphically  
# Draw the KM plot with CI  
plot(km, lty=1, lwd=2, xlab="Time", ylab="Survival Probability",  
      col=rainbow(1))
```



```
# If we don't want Confidence limits  
# We have to specify it in survfit  
kmnoci <- survfit(Surv(time, death)~ 1, conf.type="none")  
plot (kmnoci,lty=1, lwd=2, xlab="Time", ylab="Survival Probability",  
      col=rainbow(1))
```



```
# Quantiles of survival #
quantile(kmnoci)
```

```
## 25 50 75
## NA NA NA
```

```
# No value for the 25% either
```

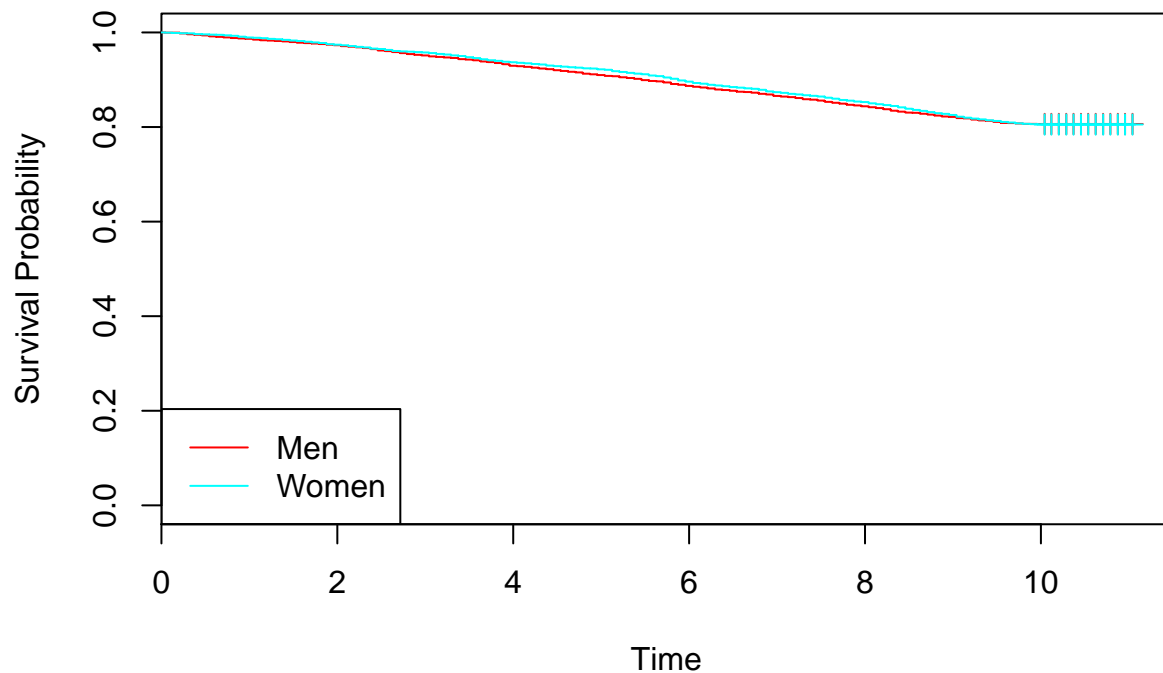
```
# We can return the survival table at every time
# or specify at a specific time point, here 11y
# which is the end of follow-up
summary(kmnoci, times=11)
```

```
## Call: survfit(formula = Surv(time, death) ~ 1, conf.type = "none")
##
##   time n.risk n.event survival std.err
##    11     52   1802    0.806 0.00411
```

```
# We see here that at the end of follow-up
# the survival was ~80% hence less than 25%
# developed the event, which is why R can't estimate
# a 25% survival
```

```
#####
## 4. Group comparisons                                ##
```

```
#####
# Draw the KM plot by sex
kmsex <- survfit( Surv(time, death)~ strata(sex), data=elsa_cf,
                  conf.type="none")
plot(kmsex, lty=1, lwd=1, xlab="Time", ylab="Survival Probability",
      col=rainbow(2))
legend("bottomleft", c("Men", "Women") , lty=1, lwd=1,
      col=rainbow(2) )
```



```
#This function implements the G-rho family of Harrington and Fleming (1982),
# with weights on each death of  $S(t)^\rho$ , where  $S$  is the Kaplan-Meier
#estimate of survival.
# With  $\rho = 0$  this is the log-rank or Mantel-Haenszel test,
# and with  $\rho = 1$  it is equivalent to the Peto & Peto modification
# of the Gehan-Wilcoxon test.
survdif(Surv(time, death) ~ sex, rho=0)
```

```
## Call:
## survdif(formula = Surv(time, death) ~ sex, rho = 0)
##
##           N Observed Expected (O-E)^2/E (O-E)^2/V
## sex=0 4484      868      867  0.000575  0.00111
## sex=1 4798      934      935  0.000534  0.00111
##
## Chisq= 0 on 1 degrees of freedom, p= 0.973
```

```

# It is not significant: no difference between men and women
#####
## 5. Exercise                                     ##
#####
# Divide cognitive function score (tertiles)
quantile(elsa_cf$cf1, prob=c(0.33, 0.66))

```

```

## 33% 66%
## 42 52

```

```

elsa_cf$t_cf1 <-cut(elsa_cf$cf1, breaks=c(0, 42, 52, 194))
# Divide cognitive function score (quintiles)
quantile(elsa_cf$cf1, prob=c(0.20, 0.40, 0.60, 0.80))

```

```

## 20% 40% 60% 80%
## 38 45 50 57

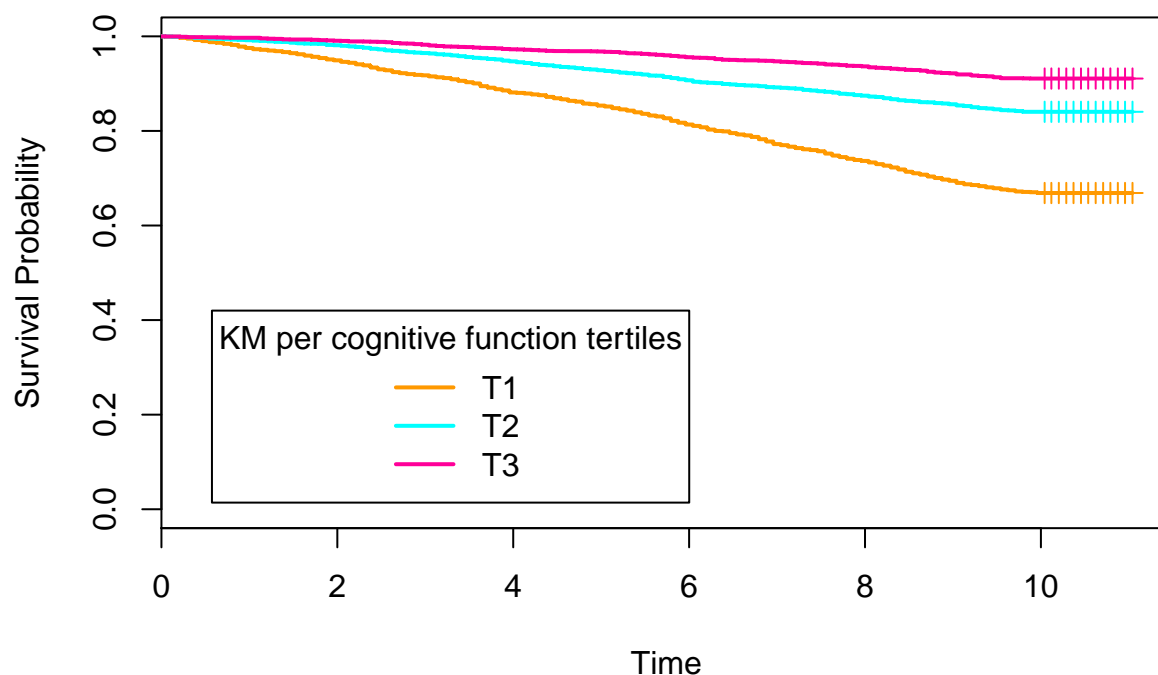
```

```

elsa_cf$q_cf1 <-cut(elsa_cf$cf1, breaks=c(0, 38, 45, 50, 57, 194))

# Draw the KM plot by tertiles t_cf1
kmcft <- survfit( Surv(time, death)~ strata(t_cf1), data=elsa_cf,
                 conf.type="none")
plot(kmcft, lty=1, lwd=2, xlab="Time", ylab="Survival Probability",
      col=rainbow(3, start=0.1, end=0.9) )
legend("bottomleft", inset=.05, title="KM per cognitive function tertiles",
      c("T1", "T2", "T3"), lty=1, lwd=2, col = rainbow(3, start=0.1, end=0.9) )

```

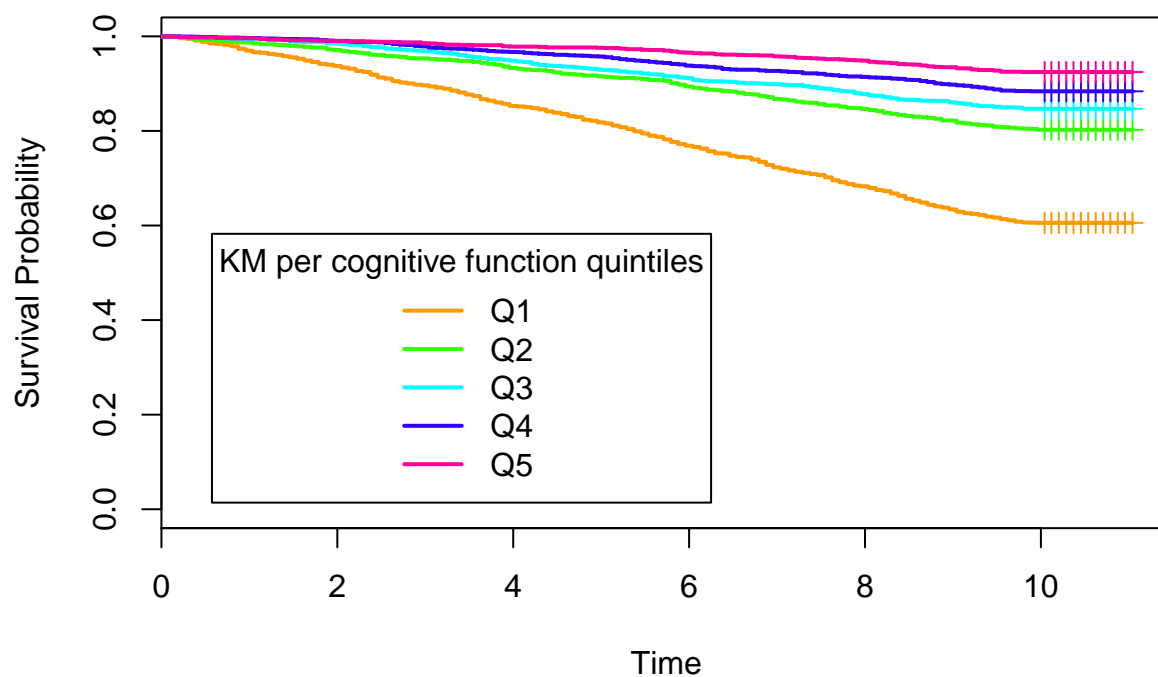



```
#logrank
survdif(Surv(elsa_cf$time, elsa_cf$death) ~ elsa_cf$t_cf1, rho=0)
```

```
## Call:
## survdiff(formula = Surv(elsa_cf$time, elsa_cf$death) ~ elsa_cf$t_cf1,
##          rho = 0)
##
##               N Observed Expected (O-E)^2/E (O-E)^2/V
## elsa_cf$t_cf1=(0,42] 3094      1025      551    407.5    588.4
## elsa_cf$t_cf1=(42,52] 3196       510      632     23.7     36.6
## elsa_cf$t_cf1=(52,194] 2992       267      618    199.7    304.6
##
## Chisq= 632 on 2 degrees of freedom, p= 0
```

```
# Significant difference
```

```
# Draw the KM plot by quintiles q_cf1
kmcfq <- survfit( Surv(time, death)~ strata(q_cf1), data=elsa_cf,
                 conf.type="none")
plot(kmcfq, lty=1,lwd=2, xlab="Time", ylab="Survival Probability",
      col=rainbow(5,start=0.1, end=0.9) )
legend("bottomleft", inset=.05, title="KM per cognitive function quintiles",
      c("Q1", "Q2", "Q3", "Q4", "Q5"), lty=1, lwd=2,
      col = rainbow(5,start=0.1, end=0.9) )
```



```
#logrank
survdif(Surv(elsa_cf$time, elsa_cf$death) ~ elsa_cf$q_cf1, rho=0)
```

```
## Call:
## survdiff(formula = Surv(elsa_cf$time, elsa_cf$death) ~ elsa_cf$q_cf1,
##          rho = 0)
##
##               N Observed Expected (O-E)^2/E (O-E)^2/V
## elsa_cf$q_cf1=(0,38] 2046      807      348    607.11   754.127
## elsa_cf$q_cf1=(38,45] 1978      391      384      0.14     0.178
## elsa_cf$q_cf1=(45,50] 1632      250      324    16.97    20.712
## elsa_cf$q_cf1=(50,57] 1970      229      402    74.13    95.496
## elsa_cf$q_cf1=(57,194] 1656      125      345   140.31   173.776
##
## Chisq= 841  on 4 degrees of freedom, p= 0
```

```
# Significant difference
```