Survival Analysis of OV with Taxol

This is for survival analysis of OV data from TCGA.

Date: 05/05/2015

380 patients have been treated with Taxol.

Time measured by month.

1. Load Data:

```
load("/Users/zhenyu/Downloads/working/Steph/ov/taxol.data.rda")
head(taxol.data)
##
                contact.days death.days days death months taxol taxotere
## TCGA-25-2042
                         396
                                     396
                                          396
                                                    13.01
                                                            TRUE
                                                                     FALSE
                                               TRUE
                                                            TRUE
## TCGA-25-2399
                         608
                                     608
                                          608
                                               TRUE
                                                     19.98
                                                                     FALSE
                        1900
                                                     62.42
                                                            TRUE
## TCGA-29-2427
                                     NA 1900 FALSE
                                                                     FALSE
## TCGA-59-2355
                                               TRUE
                                                            TRUE
                          NA
                                           65
                                                      2.14
                                                                     FALSE
                         165
## TCGA-59-2363
                                      NA
                                          165 FALSE
                                                      5.42 TRUE
                                                                     FALSE
## TCGA-25-2391
                        1492
                                    1492 1492
                                               TRUE 49.02 TRUE
                                                                      TRUE
##
                carboplatin cisplatin other taxol.prediction
## TCGA-25-2042
                       TRUE
                                FALSE TRUE
                                                    -2.455937 -2.455937
## TCGA-25-2399
                       TRUE
                                FALSE TRUE
                                                    -3.603826 -3.603826
## TCGA-29-2427
                                                    -2.493162 -2.493162
                       TRUE
                                 TRUE TRUE
## TCGA-59-2355
                       TRUE
                                FALSE FALSE
                                                    -3.449695 -3.449695
## TCGA-59-2363
                       TRUE
                                FALSE FALSE
                                                    -3.116783 -3.116783
## TCGA-25-2391
                       TRUE
                                FALSE FALSE
                                                    -2.985817 -2.985817
##
                quantile
## TCGA-25-2042
                      Q1
## TCGA-25-2399
                      Q4
## TCGA-29-2427
                      Q1
## TCGA-59-2355
                      Q4
## TCGA-59-2363
                      Q3
## TCGA-25-2391
cat = table(with(taxol.data, paste(taxol, taxotere, carboplatin, cisplatin, other, sep="|")))
cat2 = cbind(rownames(cat), cat)
rownames(cat2)=NULL
colnames(cat2) = c("Taxol|Taxotere|Carboplatin|Cisplatin|Other", "count")
print(cat2)
```

```
## Taxol|Taxotere|Carboplatin|Cisplatin|Other count
## [1,] "TRUE|FALSE|FALSE|TRUE|FALSE" "17"

## [2,] "TRUE|FALSE|FALSE|TRUE|TRUE" "9"

## [3,] "TRUE|FALSE|TRUE|FALSE|FALSE" "97"

## [4,] "TRUE|FALSE|TRUE|FALSE|TRUE" "121"

## [5,] "TRUE|FALSE|TRUE|TRUE|FALSE" "29"
```

```
## [6,] "TRUE|FALSE|TRUE|TRUE" "40"

## [7,] "TRUE|TRUE|FALSE|TRUE|FALSE" "3"

## [8,] "TRUE|TRUE|FALSE|TRUE|TRUE" "1"

## [9,] "TRUE|TRUE|TRUE|FALSE|FALSE" "11"

## [10,] "TRUE|TRUE|TRUE|FALSE|TRUE" "24"

## [11,] "TRUE|TRUE|TRUE|TRUE|FALSE" "5"

## [12,] "TRUE|TRUE|TRUE|TRUE|TRUE" "23"
```

2. Predicted sensitivity by death, and one-sided t-test:

```
qplot(death, pred, data=taxol.data, geom=c("boxplot", "jitter"),
  fill=death, main="Predicted Taxol Sensitivity by Death",
  xlab="", ylab="Predicted Taxol Sensitivity")
```

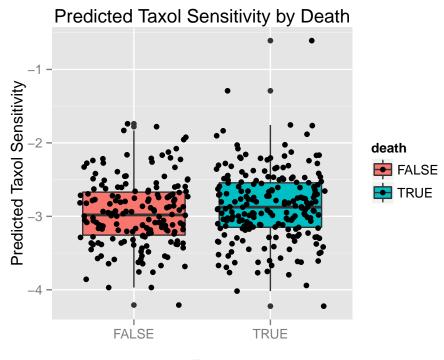


Figure 1:

```
with(taxol.data, t.test(taxol.prediction[death], taxol.prediction[!death], "greater"))
```

```
##
## Welch Two Sample t-test
##
## data: taxol.prediction[death] and taxol.prediction[!death]
## t = 1.9609, df = 370.98, p-value = 0.02532
## alternative hypothesis: true difference in means is greater than 0
## 95 percent confidence interval:
## 0.01521272 Inf
```

```
## sample estimates:
## mean of x mean of y
## -2.856808 -2.952432
```

3. KM Curve by Sensitivity Quantile:

```
ggsurv(survfit(with(taxol.data, Surv(months,death) ~ quantile)))
```

Loading required package: scales

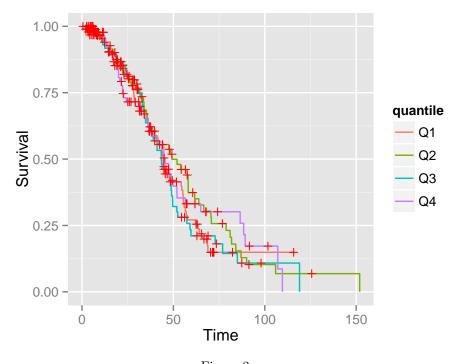


Figure 2:

```
#ggsurv(survfit(surv.data))
#survdiff(surv.data)
```

4. KM Curve and Survial test by Q1 and Q4:

```
surv.data = with(taxol.data[which(taxol.data$quantile %in% c("Q1", "Q4")), ], Surv(months,death) ~ quantile
ggsurv(survfit(surv.data))
```

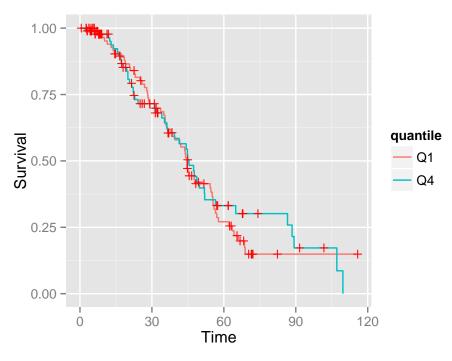


Figure 3:

survdiff(surv.data)