Survival

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code for fitting Kaplan-Meier and log-rank test and for displaying survival curves for each type of media

```
data <- read.table(here('data','Worm_9days.csv'), header = TRUE, sep = ";", dec = ",")
head(data)</pre>
```

```
##
     ï..Time Status Group Replicate
## 1
           7
                   1 FG13
           7
## 2
                   1 FG13
                                    1
## 3
           7
                      0P50
                                    1
## 4
           7
                      0P50
                                    1
                   1
## 5
           9
                      FG13
                                    1
## 6
           7
                     FG13
                                    1
```

here() should show that your position is in the folder 7.semester else select the active project to be 7.semester in the upper right corner.

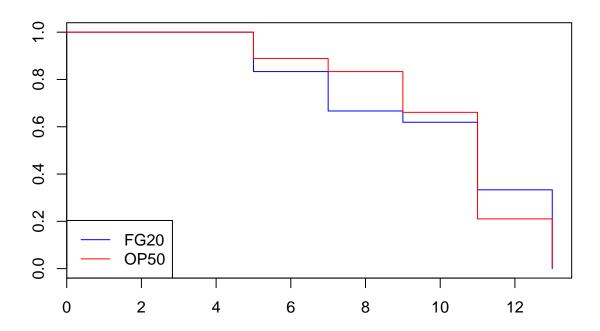
Or make a new .Rproj file in the folder 7.semester. There is a problem with the numbers of the worms so we change that with col.names()

```
colnames(data)<-c("Time", "status", "media", 'Replicate')
head(data)</pre>
```

```
##
     Time status media Replicate
## 1
        7
                   FG13
## 2
        7
                   FG13
                1
        7
## 3
                   0P50
## 4
        7
                   OP50
                                 1
                1
## 5
        9
                1
                   FG13
                                 1
## 6
        7
                0 FG13
                                 1
```

Then we create a survival object with the survival package and make a kaplan-meier curve

```
data$Survobj <- with(data,Surv(data$Time, event = data$status))
km <- survfit(Survobj ~ media, data = data,conf.type = "log-log")
plot(km, col = c("blue","red") )
legend("bottomleft",legend = c("FG20","OP50"), col = c("blue","red"), lwd=1)</pre>
```

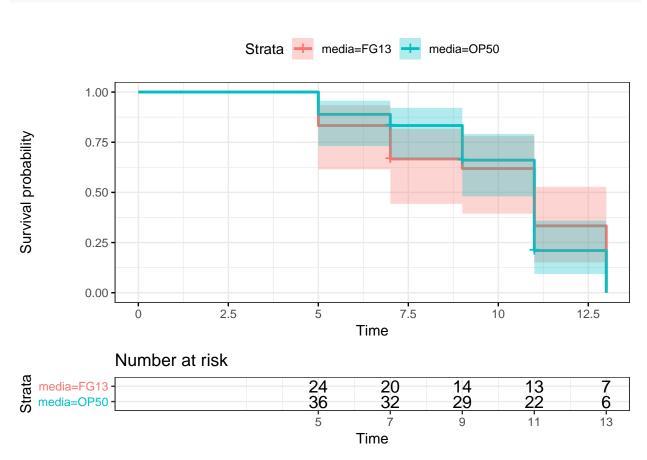


Now we test for difference between the curves with both log-rank and gehan-wilcoxon

```
survdiff(Survobj ~media, data = data, rho = 0)
## Call:
  survdiff(formula = Survobj ~ media, data = data, rho = 0)
##
               N Observed Expected (O-E)^2/E (O-E)^2/V
##
  media=FG13 24
                        22
                               22.6
                                       0.0152
                                                  0.0546
##
  media=OP50 36
                        33
                               32.4
                                       0.0106
                                                  0.0546
##
    Chisq= 0.1 on 1 degrees of freedom, p= 0.8
survdiff(Survobj ~media, data = data, rho = 1)
## Call:
## survdiff(formula = Survobj ~ media, data = data, rho = 1)
##
##
               N Observed Expected (O-E)^2/E
                                              (0-E)^2/V
## media=FG13 24
                      13.9
                               13.8
                                      0.00124
                                                 0.00401
## media=0P50 36
                      21.5
                                      0.00079
                                                 0.00401
                               21.6
##
    Chisq= 0 on 1 degrees of freedom, p= 0.9
```

Which both don't find a significant difference between the groups But we can make a nicer survival curve with a different package survminer and add readable confidence intervals





We then try to split up in replicates to see if there are any differences between scorers

```
d1 <- data[data$Replicate == 1,]
d2 <- data[data$Replicate == 2,]
d1$Survobj <- with(d1,Surv(d1$Time, event = d1$status))
d2$Survobj <- with(d2,Surv(d2$Time, event = d2$status))</pre>
```

Then we make kaplan-meier curves

```
par(mfrow = c(1,2))
km1 <- survfit(Survobj ~ media, data = d1,conf.type = "log-log")
plot(km1, col = c("blue","red") )
legend("bottomleft",legend = c("FG20","OP50"), col = c("blue","red"), lwd=1)
km2 <- survfit(Survobj ~ media, data = d2,conf.type = "log-log")
plot(km2, col = c("blue","red"))
legend("bottomleft",legend = c("FG20","OP50"), col = c("blue","red"), lwd=1)</pre>
```

