Survival of C. elegans with three different media

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

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

```
##
     ï..Time Status Replicate Group Mutant
## 1
                   1
                            NA FG13
## 2
           6
                            NA FG13
                   1
                                         PMK
## 3
           8
                   1
                            NA FG13
                                         PMK
## 4
           8
                            NA FG13
                                         PMK
                   1
## 5
          10
                   1
                            NA FG13
                                         PMK
                            NA FG13
                                         PMK
## 6
          10
                   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()

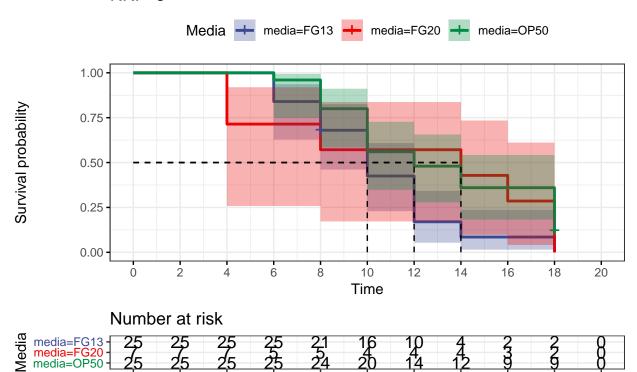
```
Time status Replicate media mutant
##
## 1
               1
                         NA FG13
## 2
        6
               1
                         NA FG13
                                     PMK
## 3
        8
               1
                         NA FG13
                                     PMK
## 4
                         NA FG13
                                     PMK
        8
               1
## 5
       10
               1
                         NA FG13
                                     PMK
                         NA FG13
                                     PMK
## 6
       10
               1
```

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

First we see for each mutant

```
d_RRF <- subset(data, mutant == 'RRF')
d_PMK <- subset(data, mutant == 'PMK')
d_DAF <- subset(data, mutant == 'DAF')</pre>
```

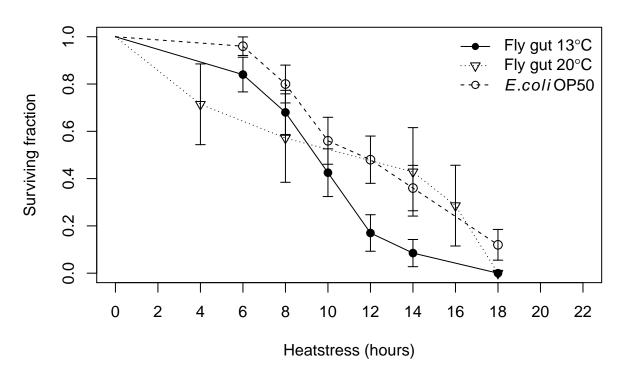
RRF-3



```
df_fly_13_r \leftarrow data.frame(c(0,s_km_RRF\$time[1:6]),c(1,s_km_RRF\$surv[1:6]),
                                                                                                       c(0,s
colnames(df_fly_13_r) <- c('Time', 'Surv', 'Std.error')</pre>
df_fly_20_r <- data.frame(c(0,s_km_RRF$time[7:11])</pre>
                         ,c(1,s_km_RRF$surv[7:11]),
                                                        c(0,s_km_RRF$std.err[7:11]))
colnames(df_fly_20_r) <- c('Time','Surv','Std.error')</pre>
df_OP50_r \leftarrow data.frame(c(0,s_km_RRF\$time[12:17])
                       ,c(1,s_km_RRF\$surv[12:17]),
                         c(0,s_km_RRF$std.err[12:17]))
colnames(df_OP50_r) <- c('Time', 'Surv', 'Std.error')</pre>
plot(df_fly_13_r$Time[2:7],df_fly_13_r$Surv[2:7], pch = 16, cex = 1.2, xlab = 'Heatstress (hours)',
     ylab = 'Surviving fraction',
     main = expression('Survival heat stress for'~italic(C.elegans)),
     xlim = c(0,22), ylim = c(0,1), xaxp = c(0,22,11))
lines(df_fly_13_r$Time,df_fly_13_r$Surv)
arrows(df_fly_13_r$Time, df_fly_13_r$Surv-df_fly_13_r$Std.error, df_fly_13_r$Time,
       df_fly_13_r$Surv+df_fly_13_r$Std.error, length=0.05, angle=90, code=3, col = 'black')
## Warning in arrows(df_fly_13_r$Time, df_fly_13_r$Surv -
## df_fly_13_r$Std.error, : zero-length arrow is of indeterminate angle and so
## skipped
points(df_OP50_r$Time[2:9],df_OP50_r$Surv[2:9], pch = 1, cex = 1.2)
lines(df_OP50_r$Time,df_OP50_r$Surv, lty = 2)
arrows(df OP50 r$Time, df OP50 r$Surv-df OP50 r$Std.error,
```

10 Time

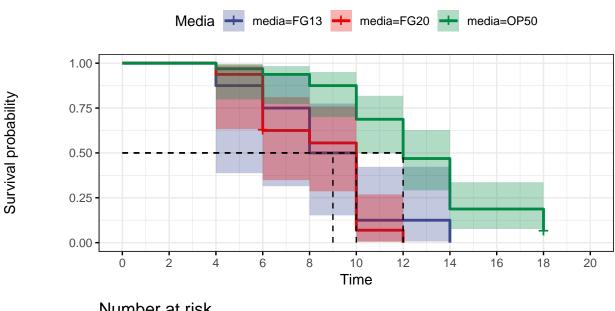
Survival heat stress for *C.elegans*



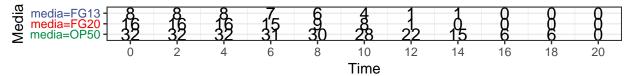
```
d_PMK$Survobj <- with(d_PMK,Surv(d_PMK$Time, event = d_PMK$status))
km_PMK <- survfit(Survobj ~ media, data = d_PMK,conf.type = "log-log", error = "greenwood")
s_km_PMK <- summary(km_PMK)
ggsurvplot(km_PMK,data = d_PMK, conf.int = TRUE,</pre>
```

```
ggtheme = theme_bw(),risk.table = 0.25,
palette ='aaas', surv.median.line = 'hv',
legend.title = 'Media', break.x.by = 2,
title = 'PMK-1')
```

PMK-1



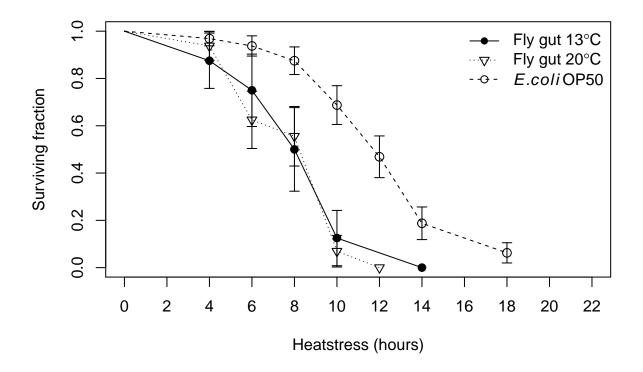
Number at risk



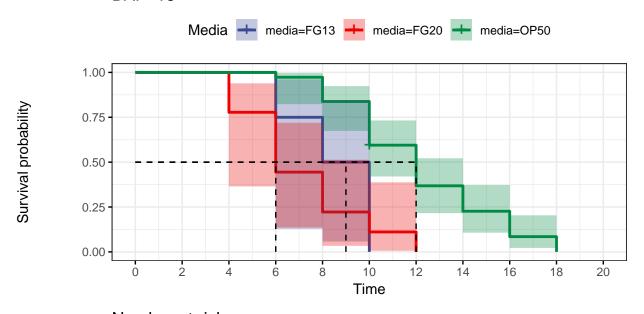
```
df fly 13 P <- data.frame(c(0,s km PMK$time[1:5]),</pre>
                         c(1,s_km_PMK$surv[1:5]),
                         c(0,s km PMK$std.err[1:5]))
colnames(df_fly_13_P) <- c('Time', 'Surv', 'Std.error')</pre>
df_fly_20_P \leftarrow data.frame(c(0,s_km_PMK_time[6:10]),
                         c(1,s_km_PMK$surv[6:10]),
                         c(0,s km PMK$std.err[6:10]))
colnames(df_fly_20_P) <- c('Time', 'Surv', 'Std.error')</pre>
df_0P50_P \leftarrow data.frame(c(0,s_km_PMK\$time[11:17]),
                       c(1,s_km_PMK$surv[11:17]),
                         c(0,s_km_PMK$std.err[11:17]))
colnames(df_OP50_P) <- c('Time', 'Surv', 'Std.error')</pre>
plot(df_fly_13_P$Time[2:9],df_fly_13_P$Surv[2:9], pch = 16, cex = 1.2, xlab = 'Heatstress (hours)',
     ylab = 'Surviving fraction',
     main = expression('Survival heat stress for'~italic(C.elegans)),
     xlim = c(0,22), ylim = c(0,1), xaxp = c(0,22,11))
lines(df_fly_13_P$Time,df_fly_13_P$Surv)
arrows(df_fly_13_P$Time, df_fly_13_P$Surv-df_fly_13_P$Std.error, df_fly_13_P$Time,df_fly_13_P$Surv+df_f
```

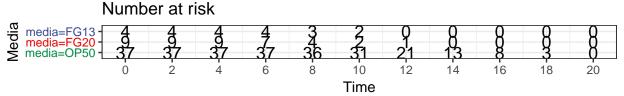
```
## Warning in arrows(df_fly_13_P$Time, df_fly_13_P$Surv -
## df_fly_13_P$Std.error, : zero-length arrow is of indeterminate angle and so
## skipped
points(df_0P50_P$Time[2:9],df_0P50_P$Surv[2:9], pch = 1, cex = 1.2)
lines(df_OP50_P$Time,df_OP50_P$Surv, lty = 2)
arrows(df OP50 P$Time, df OP50 P$Surv-df OP50 P$Std.error,
      df_OP50_P$Time, df_OP50_P$Surv+df_OP50_P$Std.error,
      length=0.05, angle=90, code=3, col = 'black')
## Warning in arrows(df_OP50_P$Time, df_OP50_P$Surv - df_OP50_P$Std.error, :
## zero-length arrow is of indeterminate angle and so skipped
points(df_fly_20_P$Time[2:9],df_fly_20_P$Surv[2:9], pch = 6)
lines(df_fly_20_P$Time,df_fly_20_P$Surv,lty = 3)
arrows(df_fly_20_P$Time, df_fly_20_P$Surv-df_fly_20_P$Std.error, df_fly_20_P$Time,df_fly_20_P$Surv+df_f
## Warning in arrows(df_fly_20_P$Time, df_fly_20_P$Surv -
## df_fly_20_P$Std.error, : zero-length arrow is of indeterminate angle and so
## skipped
legend('topright', pch = c(16,6,1), lty = c(1,3,2),
       legend = c(expression('Fly gut 13'*degree*C),
                  expression('Fly gut 20'*degree*C),
                  expression(italic(E.coli) ~ OP50)), bty = 'n')
```

Survival heat stress for C.elegans



DAF-16

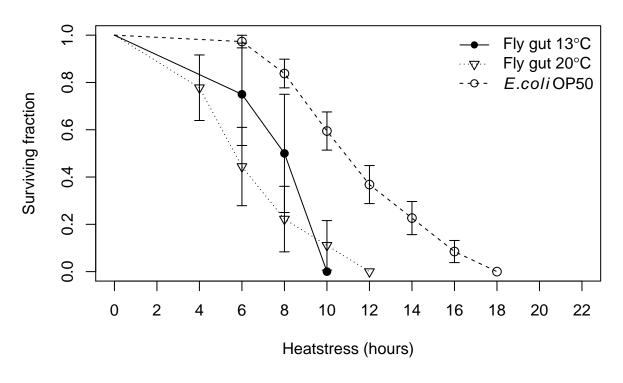




```
xlim = c(0,22), ylim = c(0,1), xaxp = c(0,22,11))
lines(df_fly_13_d$Time,df_fly_13_d$Surv)
arrows(df_fly_13_d$Time, df_fly_13_d$Surv-df_fly_13_d$Std.error, df_fly_13_d$Time,df_fly_13_d$Surv+df_f
## Warning in arrows(df_fly_13_d$Time, df_fly_13_d$Surv -
## df_fly_13_d$Std.error, : zero-length arrow is of indeterminate angle and so
## skipped
points(df OP50 d$Time[2:9],df OP50 d$Surv[2:9], pch = 1, cex = 1.2)
lines(df_OP50_d$Time,df_OP50_d$Surv, lty = 2)
arrows(df_OP50_d$Time, df_OP50_d$Surv-df_OP50_d$Std.error,
      df_OP50_d$Time, df_OP50_d$Surv+df_OP50_d$Std.error,
      length=0.05, angle=90, code=3, col = 'black')
## Warning in arrows(df_0P50_d$Time, df_0P50_d$Surv - df_0P50_d$Std.error, :
## zero-length arrow is of indeterminate angle and so skipped
points(df_fly_20_d$Time[2:9],df_fly_20_d$Surv[2:9], pch = 6)
lines(df_fly_20_d$Time,df_fly_20_d$Surv,lty = 3)
arrows(df_fly_20_d$Time, df_fly_20_d$Surv-df_fly_20_d$Std.error, df_fly_20_d$Time,df_fly_20_d$Surv+df_f
## Warning in arrows(df_fly_20_d$Time, df_fly_20_d$Surv -
## df fly 20 d$Std.error, : zero-length arrow is of indeterminate angle and so
## skipped
legend('topright', pch = c(16,6,1), lty = c(1,3,2),
       legend = c(expression('Fly gut 13'*degree*C),
                  expression('Fly gut 20'*degree*C),
```

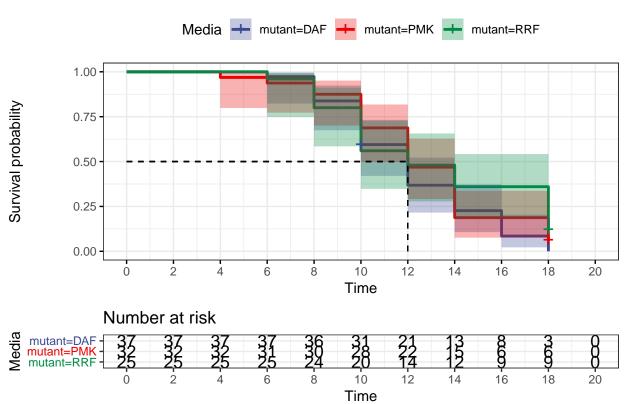
expression(italic(E.coli) ~ OP50)), bty = 'n')

Survival heat stress for *C.elegans*

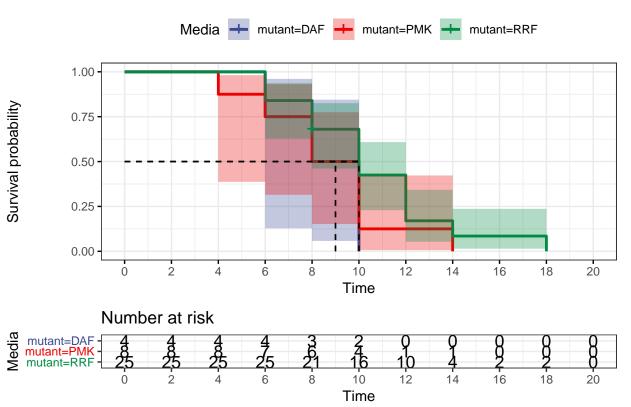


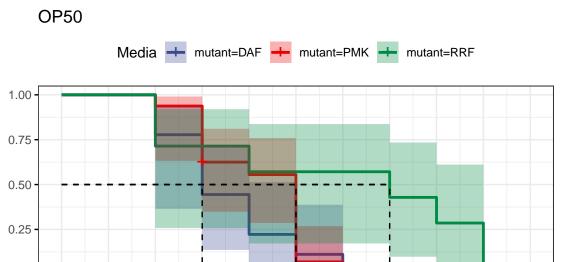
Then we compare by medium

OP50



FG13





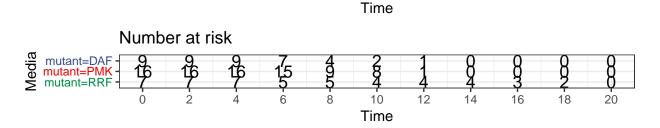
10

12

14

16

20



8

Shows errorbars and lineplot based on one of Anders papers https://onlinelibrary.wiley.com/doi/full/10. 1111/acel.12165 This is done by extracting the surviving proportion, standard errors and time from survfit(). and adding a startpoint where the survival is 100% at 0 hours.

Now we test for difference between the curves with both log-rank and gehan-wilcoxon and with an cox proportional hazard model

```
data$Survobj <- with(data,Surv(data$Time, event = data$status))</pre>
survdiff(Survobj ~media + mutant, data = data, rho = 0)
```

```
## Call:
## survdiff(formula = Survobj ~ media + mutant, data = data, rho = 0)
##
                            N Observed Expected (O-E)^2/E (O-E)^2/V
##
  media=FG13, mutant=DAF
##
                            4
                                      4
                                            1.78
                                                      2.762
                                                                 3.545
                                      8
                                            4.22
## media=FG13, mutant=PMK
                            8
                                                      3.384
                                                                 4.615
## media=FG13, mutant=RRF
                           25
                                     24
                                           19.78
                                                      0.901
                                                                 1.483
## media=FG20, mutant=DAF
                                      9
                                            2.82
                                                     13.539
                                                                17.336
## media=FG20, mutant=PMK 16
                                     15
                                            6.88
                                                      9.595
                                                                13.238
  media=FG20, mutant=RRF
                                      7
                                            8.29
                                                      0.200
                                                                 0.341
## media=OP50, mutant=DAF 37
                                     36
                                           38.84
                                                      0.208
                                                                 0.404
  media=OP50, mutant=PMK 32
                                     30
                                           39.06
                                                      2.100
                                                                 4.263
                                     22
##
  media=OP50, mutant=RRF 25
                                           33.34
                                                      3.855
                                                                 7.900
##
##
```

Chisq= 51.7 on 8 degrees of freedom, p= 2e-08

Survival probability

0.00

0

2

```
survdiff(Survobj ~media + mutant, data = data, rho = 1)
## Call:
## survdiff(formula = Survobj ~ media + mutant, data = data, rho = 1)
##
                          N Observed Expected (O-E)^2/E (O-E)^2/V
## media=FG13, mutant=DAF
                               3.24
                                        1.50
                                                2.0432
                         4
                                                          2.9683
## media=FG13, mutant=PMK 8
                               6.12
                                        3.18
                                                2.7304
                                                          4.3317
## media=FG13, mutant=RRF 25
                                       12.93
                                                0.4151
                              15.24
                                                          0.8063
## media=FG20, mutant=DAF 9
                                        2.32
                               7.76
                                               12.7917
                                                         18.3933
## media=FG20, mutant=PMK 16
                              12.12
                                        5.63
                                                7.4975
                                                        11.8513
## media=FG20, mutant=RRF 7
                               3.62
                                        3.99
                                              0.0348
                                                        0.0719
## media=OP50, mutant=DAF 37
                             18.26
                                       23.74
                                               1.2635
                                                          2.9998
## media=OP50, mutant=PMK 32
                               14.58
                                       22.00
                                                2.5059
                                                          6.0333
## media=OP50, mutant=RRF 25
                               11.37
                                       17.04
                                                1.8896
                                                          4.3076
##
## Chisq= 48 on 8 degrees of freedom, p= 1e-07
fit_both<- coxph(Survobj ~media + mutant, data = data)</pre>
summary(fit_both)
## Call:
## coxph(formula = Survobj ~ media + mutant, data = data)
##
    n= 163, number of events= 155
##
##
                 coef exp(coef) se(coef)
## mediaFG20 0.005922 1.005940 0.250871 0.024 0.981166
## mediaOP50 -1.147199 0.317525 0.231637 -4.953 7.32e-07 ***
## mutantPMK -0.253909 0.775762 0.202013 -1.257 0.208791
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
##
##
            exp(coef) exp(-coef) lower .95 upper .95
## mediaFG20
               1.0059
                          0.9941
                                   0.6152
                                             1.6448
## mediaOP50
               0.3175
                          3.1494
                                             0.5000
                                    0.2017
## mutantPMK
               0.7758
                          1.2891
                                    0.5221
                                             1.1526
## mutantRRF
               0.4187
                          2.3885
                                   0.2669
                                             0.6568
##
## Concordance= 0.67 (se = 0.028)
## Rsquare= 0.211
                   (max possible= 1 )
## Likelihood ratio test= 38.69 on 4 df,
                                          p=8e-08
## Wald test
                       = 38.5 \text{ on } 4 \text{ df},
                                         p = 9e - 08
## Score (logrank) test = 39.91 on 4 df,
                                         p=5e-08
AIC(fit_both)
```

[1] 1273.218

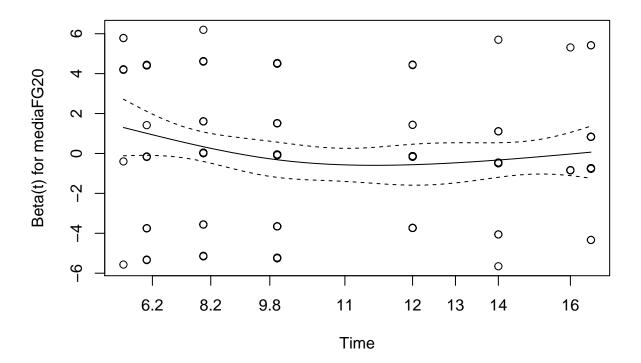
```
fit_int <- coxph(Survobj ~media + mutant + mutant:media, data = data)</pre>
AIC(fit int)
## [1] 1274.41
fit_med <- coxph(Survobj ~media, data = data)</pre>
summary(fit_med)
## Call:
## coxph(formula = Survobj ~ media, data = data)
##
##
    n= 163, number of events= 155
##
##
                coef exp(coef) se(coef)
                                             z Pr(>|z|)
                       1.2178
                                 0.2456 0.802 0.422369
## mediaFG20 0.1971
## mediaOP50 -0.7195
                        0.4870
                                 0.2019 -3.564 0.000365 ***
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
##
             exp(coef) exp(-coef) lower .95 upper .95
##
## mediaFG20
                1.218
                           0.8211
                                     0.7525
                                               1.9709
## mediaOP50
                 0.487
                           2.0535
                                     0.3278
                                               0.7234
##
## Concordance= 0.649 (se = 0.025)
## Rsquare= 0.131
                   (max possible= 1)
                                          p=1e-05
## Likelihood ratio test= 22.98 on 2 df,
                                           p=5e-06
## Wald test
                       = 24.52 on 2 df,
## Score (logrank) test = 25.84 on 2 df,
                                          p=2e-06
AIC(fit med)
## [1] 1284.932
fit_mut <- coxph(Survobj ~mutant, data = data)</pre>
summary(fit_mut)
## Call:
## coxph(formula = Survobj ~ mutant, data = data)
##
    n= 163, number of events= 155
##
##
##
                 coef exp(coef) se(coef)
                                              z Pr(>|z|)
                      0.91979 0.19910 -0.420
                                                   0.675
## mutantPMK -0.08361
## mutantRRF -0.33095
                      0.71824 0.20139 -1.643
                                                   0.100
##
             exp(coef) exp(-coef) lower .95 upper .95
                            1.087
## mutantPMK
                0.9198
                                     0.6226
                                                1.359
## mutantRRF
                0.7182
                            1.392
                                     0.4840
                                                1.066
##
## Concordance= 0.526 (se = 0.029)
## Rsquare= 0.018 (max possible= 1 )
```

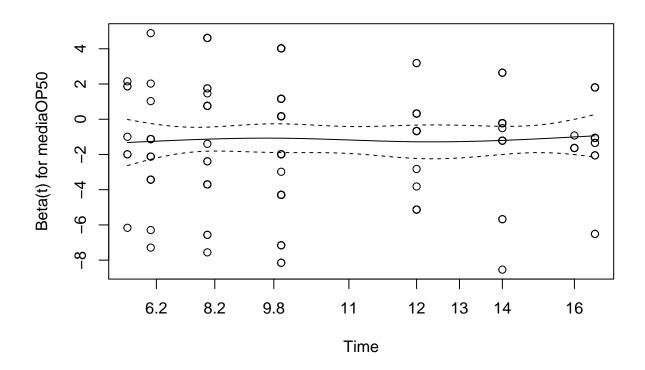
```
## Likelihood ratio test= 3.03 on 2 df, p=0.2
## Wald test = 2.97 on 2 df, p=0.2
## Score (logrank) test = 2.99 on 2 df, p=0.2
AIC(fit_mut)
```

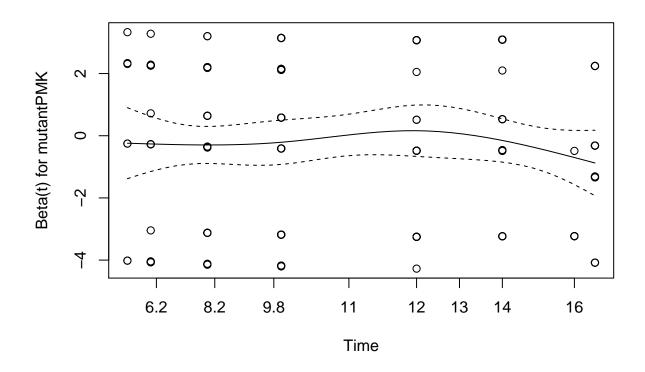
[1] 1304.882

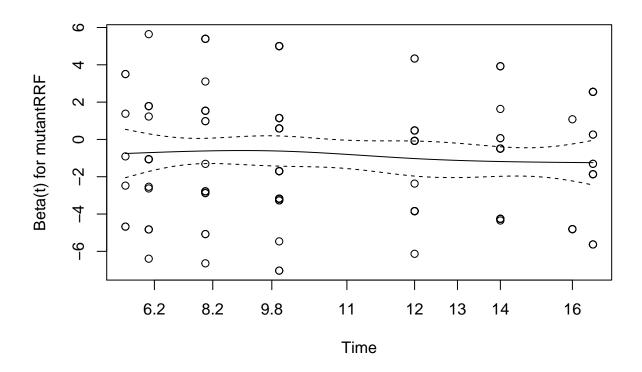
Best model must include both media and mutant strain but not necessarily an interaction between them. This could be driven by few observations for FG20 in RRF-3 Then we do a check on the proportional hazard assumption

```
prop_both <- cox.zph(fit_both)
prop_int <- cox.zph(fit_int)
plot(prop_both)</pre>
```

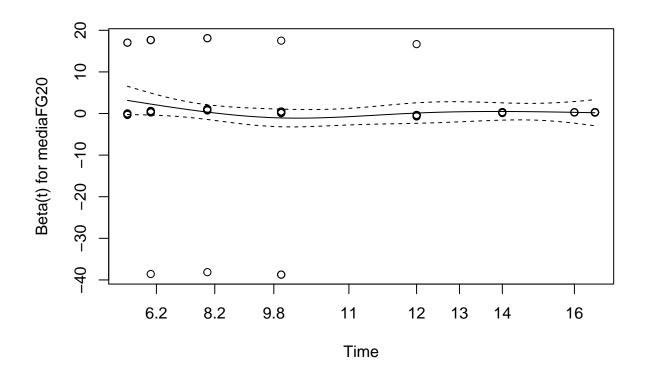


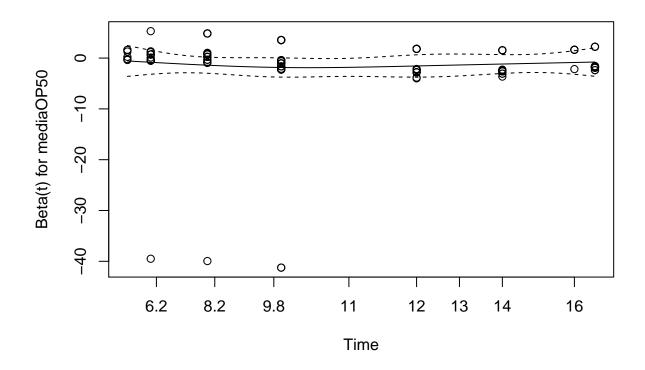


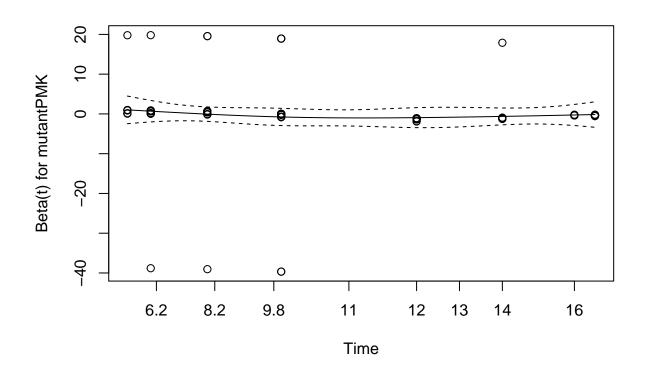


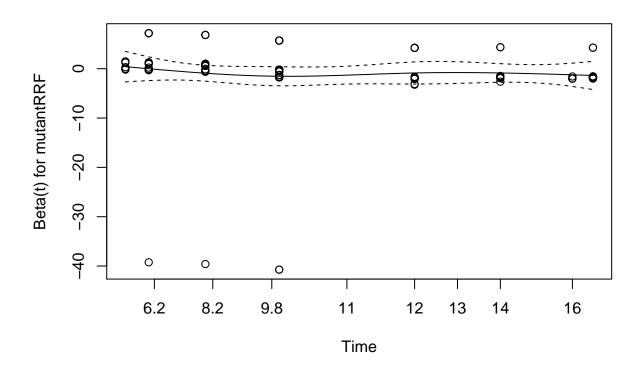


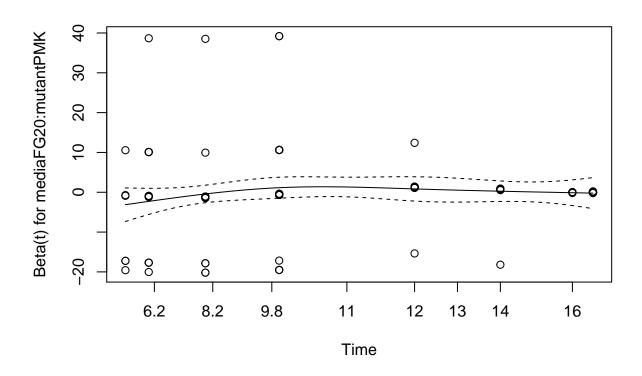
plot(prop_int)

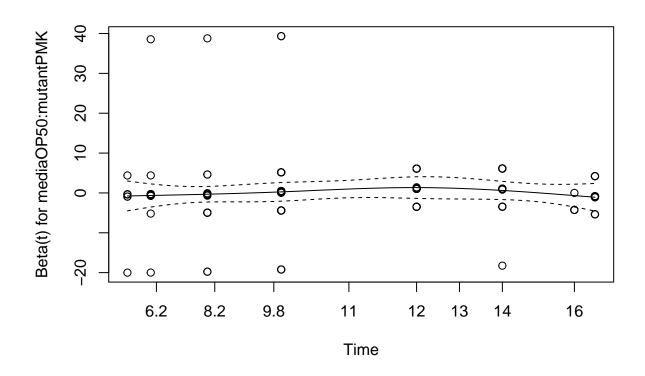


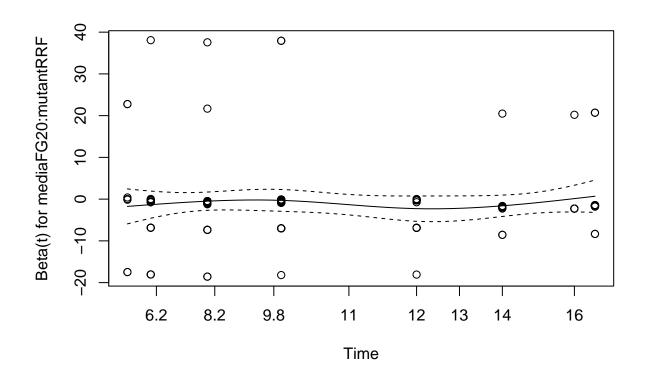


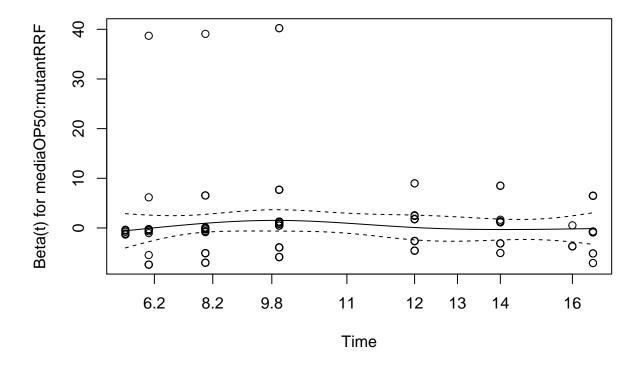








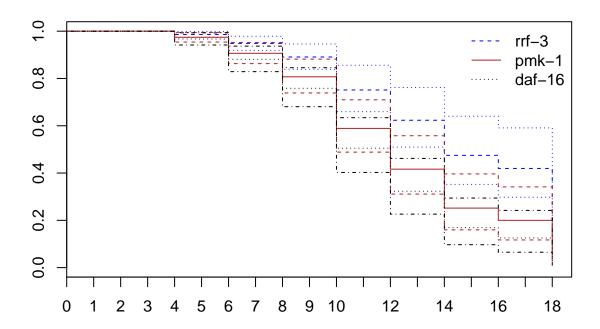




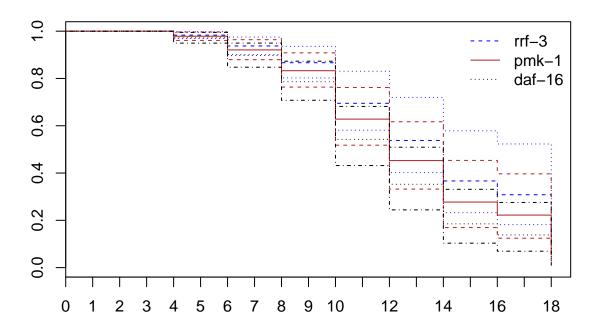
Both schoenfield residuals and the test is not significant so we proceed as if the assumptions is true for our dataset.

Then we plot predictions for both models

Predictions for model without interactions OP50

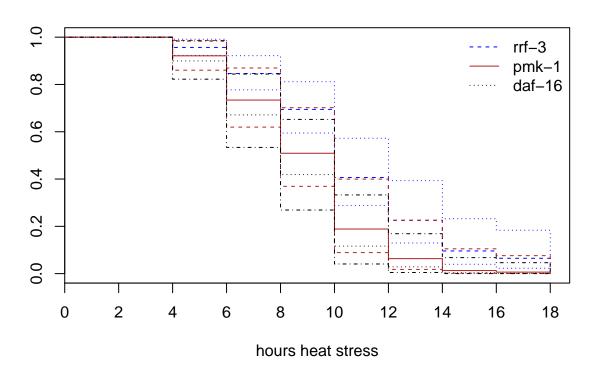


Predictions for model with interactions OP50

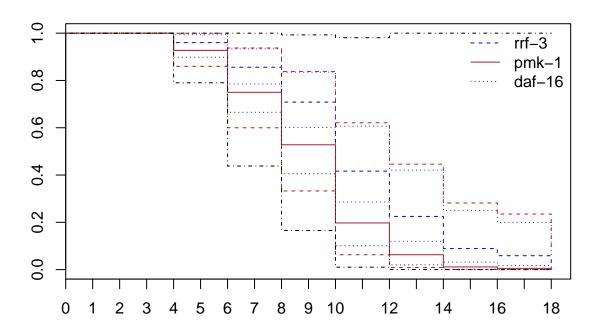


```
plot(survfit(fit_both, newdata = data.frame(mutant = 'RRF', media = 'FG13')), lty = 2, xaxp = c(0,18,9)
    main = 'Predictions for model without interactions FG13',
    xlab = 'hours heat stress')
lines(survfit(fit_both, newdata = data.frame(mutant = 'PMK', media = 'FG13')), col = 'firebrick')
lines(survfit(fit_both, newdata = data.frame(mutant = 'DAF', media = 'FG13')), lty =3)
legend('topright', legend = c('rrf-3','pmk-1','daf-16'),
    lty = c(2,1,3), col = c('blue','firebrick','black'),
    bty = 'n')
```

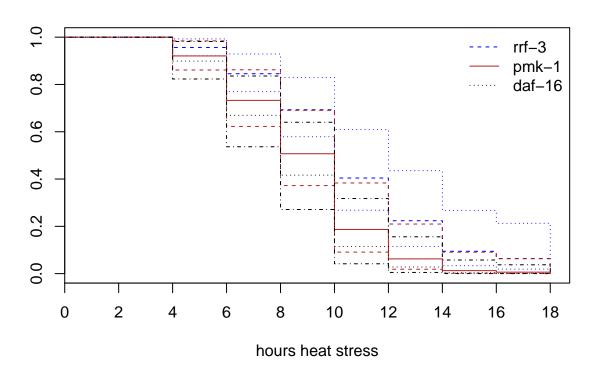
Predictions for model without interactions FG13



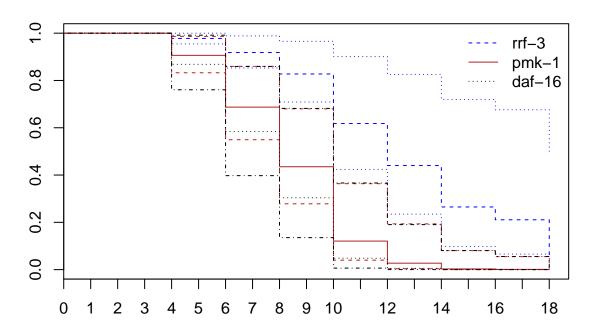
Predictions for model with interactions FG13



Predictions for model without interactions FG20

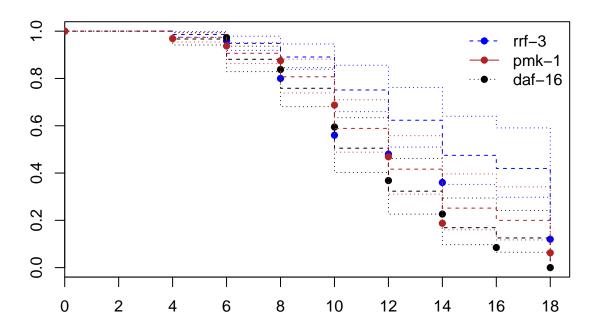


Predictions for model with interactions FG20



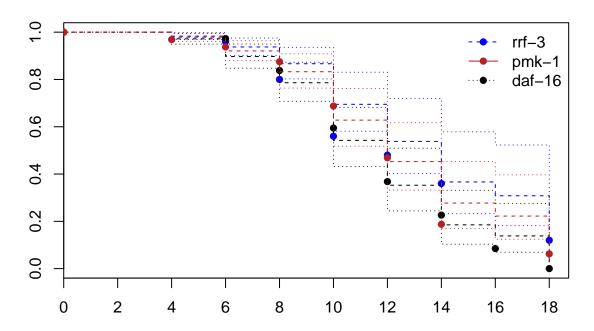
We check how well predictions line up with reality

Predictions for model without interactions OP50



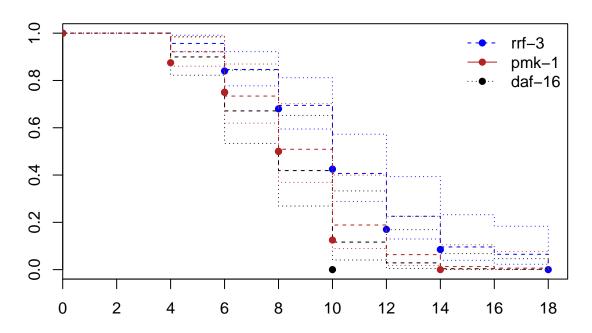
Fit is very bad as is clear both concordance and R^2 from the model. But there is far too little uncertainty in the confidence intervals. Which overestimates survival heavily for this group. Lets try the other model

Predictions for model with interactions OP50



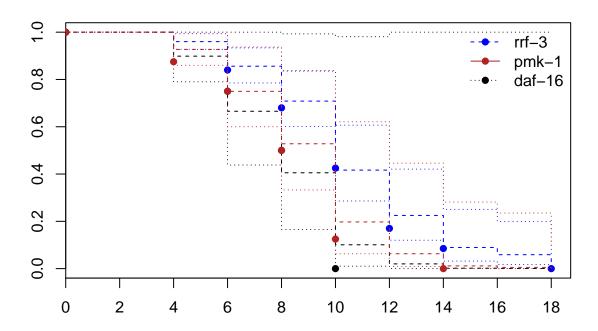
Still on the low end but a lot better. Seems that without interactions it overestimates survival on OP50.

Predictions for model with interactions FG13



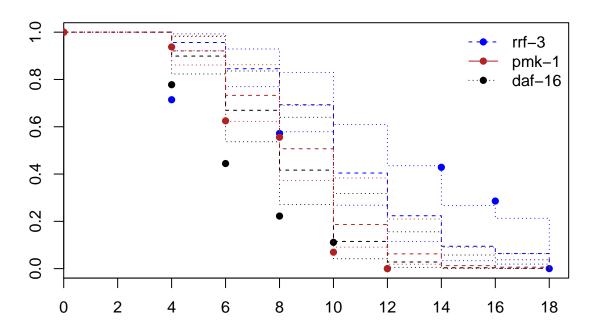
Fits notably better

Predictions for model with interactions FG13



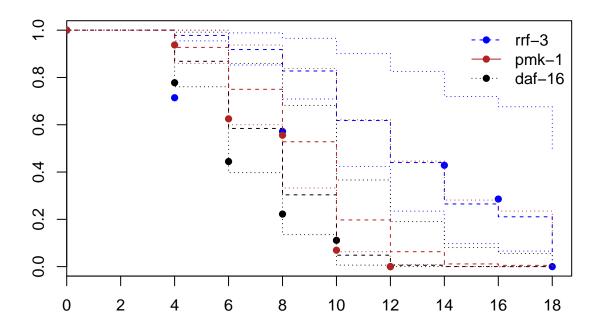
Change opacity on some points to fix overlap

Predictions for model without interactions FG20



Fitsvery bad with far too much uncertainty

Predictions for model with interactions FG20



The model has a problem with catching that all worms die around the same time and may therefore overestimate survival as it fits okay until the worms start dying. Some other model structure is maybe necessary or not all important covariates of survival are measured.