# Mortality of zoea 1 larvae of *Taliepus dentatus* reared at different temperatures

Reproducible report

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This document reproduces and supports all data work and statistical analysis of the larval mortality in the paper: swimming performance and morphometrics of the kelp-crab *Taliepus dentatus* zoea 1 exposed to different rearing temperatures.

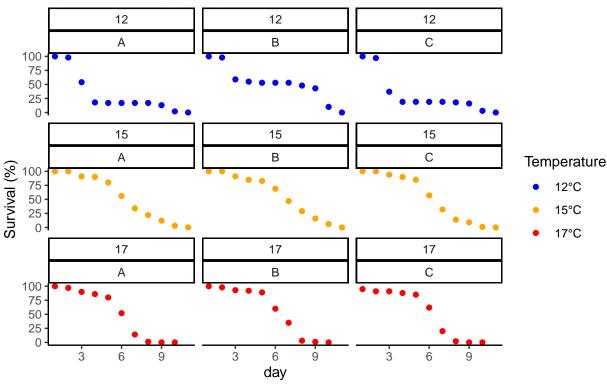
#### 1. Libraries

```
library(cowplot)
library(scales)
library(dplyr)
library(gridExtra)
library(tidyverse)
library(ggsci)
library(Rmisc)
library(survminer)
library(survival)
```

#### 2. Import of data base and definition of factors

```
c("blue", "orange", "red"),
labels=c("12°C","15°C","17°C"))
```

## survival replicates (A–B–C) by temperature



#### 2.1 table with statistics for LT50 and Kaplan-meier Analysis

```
##
    Temperatura Dia N
                         bino
                                    sd
                                              se
                                                       ci
                ## 1
            12
                2 3 0.976666667 0.005773503 0.003333333 0.01434218
## 2
            12
                3 3 0.500000000 0.115325626 0.066583281 0.28648474
## 3
            12
            12
                4 3 0.306666667 0.210792157 0.121700908 0.52363675
## 4
## 5
            12
                5 3 0.296666667 0.202319879 0.116809436 0.50259044
                6 3 0.296666667 0.202319879 0.116809436 0.50259044
## 6
            12
## 7
            12
                7 3 0.296666667 0.202319879 0.116809436 0.50259044
                8 3 0.276666667 0.176162803 0.101707642 0.43761266
## 8
            12
## 9
            12
                9 3 0.240000000 0.165227116 0.095393920 0.41044691
               10 3 0.050000000 0.043588989 0.025166115 0.10828105
## 10
            12
## 11
               ## 12
            15
```

```
## 14
             15
                 3 3 0.920000000 0.017320508 0.010000000 0.04302653
## 15
                 4 3 0.883333333 0.028867513 0.016666667 0.07171088
                 5 3 0.826666667 0.025166115 0.014529663 0.06251609
## 16
            15
## 17
             15
                 6 3 0.606666667 0.072341781 0.041766547 0.17970695
                 7 3 0.376666667 0.081445278 0.047022453 0.20232129
## 18
            15
## 19
                 8 3 0.216666667 0.075055535 0.043333333 0.18644828
## 20
            15
                 9 3 0.123333333 0.035118846 0.020275875 0.08724005
## 21
            15 10 3 0.033333333 0.025166115 0.014529663 0.06251609
## 22
            ## 23
            17
                 1 3 0.983333333 0.028867513 0.016666667 0.07171088
## 24
            17
                 2 3 0.953333333 0.037859389 0.021858128 0.09404794
## 25
            17
                 3 3 0.913333333 0.015275252 0.008819171 0.03794583
## 26
            17
                 4 3 0.886666667 0.030550505 0.017638342 0.07589166
## 27
            17
                 5 3 0.846666667 0.045092498 0.026034166 0.11201597
## 28
             17
                 6 3 0.580000000 0.052915026 0.030550505 0.13144821
            17
## 29
                 7 3 0.230000000 0.108166538 0.062449980 0.26870058
## 30
            17
                 8 3 0.020000000 0.010000000 0.005773503 0.02484138
                 9 3 0.003333333 0.005773503 0.003333333 0.01434218
## 31
            17
## 32
```

#### 3. GLM analysis and LT50% plot.

```
#separation of the table by temp
mort_12<-LT50_summary[1:11, ]
mort_15<-LT50_summary[12:22, ]
mort_17<-LT50_summary[23:32,]

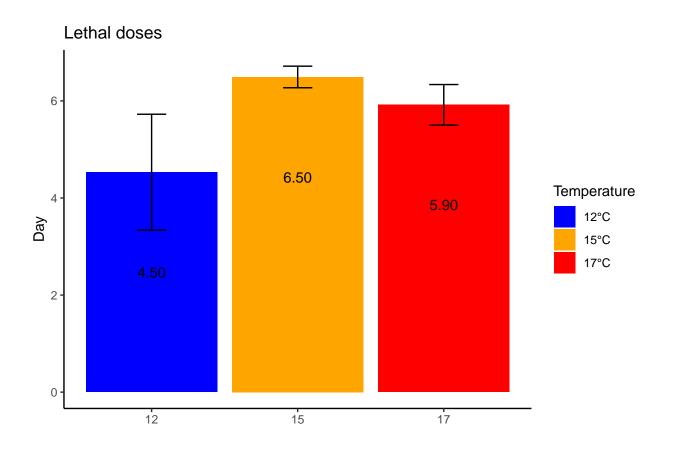
#GLM analysis for each temp

model1<-glm(data=mort_12, bino ~ Dia, family=quasibinomial())
summary(model1)</pre>
```

```
## Call:
  glm(formula = bino ~ Dia, family = quasibinomial(), data = mort_12)
## Deviance Residuals:
                1Q
                     Median
                                   3Q
                                          Max
      Min
## -0.5179 -0.2999 -0.0738
                              0.3400
                                        0.5964
##
## Coefficients:
              Estimate Std. Error t value Pr(>|t|)
                           0.6380
                                    3.403 0.00784 **
## (Intercept)
                2.1710
                -0.4791
                           0.1117 -4.289 0.00202 **
## Dia
##
## Signif. codes: 0 '*** 0.001 '** 0.01 '* 0.05 '.' 0.1 ' 1
##
## (Dispersion parameter for quasibinomial family taken to be 0.1436587)
##
      Null deviance: 5.4993 on 10 degrees of freedom
## Residual deviance: 1.5399 on 9 degrees of freedom
```

```
## AIC: NA
##
## Number of Fisher Scoring iterations: 5
model2<-glm(data=mort_15, bino ~ Dia, family=quasibinomial())</pre>
summary(model2)
##
## Call:
## glm(formula = bino ~ Dia, family = quasibinomial(), data = mort_15)
## Deviance Residuals:
        Min
                     1Q
                            Median
                                           3Q
                                                     Max
## -0.185582 -0.053923 -0.005199
                                     0.088830
                                                0.186640
##
## Coefficients:
              Estimate Std. Error t value Pr(>|t|)
##
## (Intercept) 5.84038
                          0.37027
                                     15.77 7.28e-08 ***
                           0.05508 -16.33 5.39e-08 ***
## Dia
              -0.89940
## Signif. codes: 0 '*** 0.001 '** 0.01 '* 0.05 '.' 0.1 ' 1
## (Dispersion parameter for quasibinomial family taken to be 0.01151942)
##
##
      Null deviance: 8.21286 on 10 degrees of freedom
## Residual deviance: 0.13672 on 9 degrees of freedom
## AIC: NA
##
## Number of Fisher Scoring iterations: 6
model3<-glm(data=mort_17, bino ~ Dia, family=quasibinomial())</pre>
summary(model3)
##
## Call:
## glm(formula = bino ~ Dia, family = quasibinomial(), data = mort_17)
## Deviance Residuals:
        Min
                    1Q
                            Median
                                           30
                                                     Max
## -0.287545 -0.221565 -0.146333 -0.006668
                                                0.263467
##
## Coefficients:
##
              Estimate Std. Error t value Pr(>|t|)
                                   6.974 0.000116 ***
## (Intercept) 6.6331
                            0.9512
               -1.1205
                            0.1559 -7.186 9.37e-05 ***
## Dia
## Signif. codes: 0 '*** 0.001 '** 0.01 '* 0.05 '.' 0.1 ' 1
## (Dispersion parameter for quasibinomial family taken to be 0.05103717)
##
      Null deviance: 8.41356 on 9 degrees of freedom
## Residual deviance: 0.38809 on 8 degrees of freedom
## AIC: NA
```

```
##
## Number of Fisher Scoring iterations: 6
#Dose.p function to find moment at which 50% of the population has
#died based in the qlm models made above
print(dose.p(model1, p = 0.5))
                Dose
                             SE
## p = 0.5: 4.531844 0.6121778
print(dose.p(model2, p = 0.5))
                Dose
## p = 0.5: 6.493628 0.11405
print(dose.p(model3, p = 0.5))
                Dose
## p = 0.5: 5.919705 0.2142654
# LT50% plot with values given by dose.p
lt50<-c(4.531844, 6.493628, 5.919705)
CI<-c(0.6121778*1.95, 0.11405*1.95, 0.2142654*1.95)
temp < -c(12, 15, 17)
data final<-data.frame(lt50, CI, temp)</pre>
data_final$temp<-as.factor(data_final$temp)</pre>
lt50_plot<-ggplot(data_final, aes(x=temp, y=lt50, fill=temp))+</pre>
  geom_bar(stat="identity") + theme_classic() +
  labs(title = "Lethal doses",
       x="", y="Day", fill= "Temperature") +
  scale_fill_manual(values =c("blue", "orange", "red"),
                    labels=c("12°C","15°C","17°C"))
lt50_plot + geom_errorbar(aes(ymin=lt50-CI, ymax=lt50+CI),
                          width=.2, position=position_dodge(0.05)) +
  geom_text(aes(label=c("4.50","6.50","5.90")),position =
              position_dodge(0.9), vjust = 10)
```



#### 3.1 Kaplan-Meier Analysis

```
#Vector for creating correct data table
repV<- Mort$Muertas</pre>
  repV[repV=="0"]<-1
#Datatable KAP for analysis
KAP<-Mort %>%
  mutate(count = repV) %>%
  uncount(count)
# Adding Status column (binary in which 1=no death; 2=death on that day)
KAP$STATUS <- ifelse (KAP$Muertas==0, 1, 2)</pre>
survObj <- Surv(time=KAP$Dia, event=KAP$STATUS==2)</pre>
#Kaplan-Meier analysis
sfit <- survfit(Surv(Dia, STATUS)~Temperatura, data=KAP)</pre>
#Ploting Kaplan-meier results
ggsurvplot(sfit, conf.int=TRUE, pval=TRUE, risk.table=FALSE,
           legend.labs=c("12°C", "15°C", "17°C"),
           legend.title="Temperature",
           palette=(c("blue", "orange", "red")),
```

### Kaplan-Meier Curve for Zoea 1 Survival to Temperature

