

2017_08_26 Lifehist glm

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R Markdown

Description of columns:

Independent variables

- Biome [Amazon, Cerrado, Mata Atlantica]- Categorical/nominal
 - State [Amazonas, Rondonia, Tocantins, Mata Atlantica]-Ordinal
 - Locality [ARS, APR, RPV, RMO, TLC, TPN, SJU]-Ordinal
 - Latitude [-2.864, -3.028, -8.742, -9.223, -10.7, -10.796, -22.611]- Ordinal
- Family [for lab reared, which female within the biome/locality they came from]- Categorical
- Well grouping [1,2,3]- Ordinal (5 larvae raised in each of 3 wells per female per temperature)
- Temperature [20 (A), 24 (B), 28C (C)]- Ordinal
- Sex [M/F]- Categorical

Dependent variables

- Death status (0,1,2) - Ordinal 0=died as larvae, 1= died as pupae, 2=died as adult
- LL- Larvae life length (days)- Discrete, all
- sLL- Larvae life length (days)- Discrete, only ones that went to adult
- PL- Pupal life length (days)- Discrete (Emergence.date-pupation.date)
- AL- Adult life length (days)- Discrete (Deathtime-Emergence.date)
- EmTime- time to emergence (Emergence.date-Hatch.day)
- Wing length (mm)- Continuous
- Rate of development
- Sex ratio

```
print("Gamma distribution results")
```

```
## [1] "Gamma distribution results"
```

```
print(modelAIC)
```

```
##      df      AIC
## mod1 45 -19713.44
## mod2  4 -18832.49
## mod3  9 -18834.34
## mod4 22 -19719.68
## mod5 10 -19343.92
## mod6 21 -19352.36
## mod7 27 -19655.53
## mod8 13 -19651.75
```

```
print("mod1<-glm(rate~Temp*Locality*Sex,data=lifedat,family=Gamma);summary(mod1)")
```

```
## [1] "mod1<-glm(rate~Temp*Locality*Sex,data=lifedat,family=Gamma);summary(mod1)"
```

```
print(summary(mod1))
```

```
##
## Call:
## glm(formula = rate ~ Temp * Locality * Sex, family = Gamma, data = lifedat)
##
## Deviance Residuals:
##      Min       1Q   Median       3Q      Max
## -0.34794 -0.07935 -0.00933  0.07091  0.44500
##
## Coefficients: (19 not defined because of singularities)
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)    16.975641    2.345447   7.238 5.97e-13 ***
## TempB          -3.337201    0.358109  -9.319 < 2e-16 ***
## TempC          -4.041481    2.922904  -1.383 0.166877
## LocalityAPR     1.146346    0.366982   3.124 0.001805 **
## LocalityRPV     2.558976    0.393079   6.510 8.98e-11 ***
## LocalityRMO     3.837704    0.406206   9.448 < 2e-16 ***
## LocalityTLC     3.024359    0.485761   6.226 5.56e-10 ***
## LocalityTPN     4.101536    0.801693   5.116 3.35e-07 ***
## LocalitySJU     5.911875    0.472918  12.501 < 2e-16 ***
## SexF            3.776063    2.363964   1.597 0.110310
## SexM            3.403779    2.328496   1.462 0.143918
## TempB:LocalityAPR -0.612066    0.474985  -1.289 0.197651
## TempC:LocalityAPR  0.168012    0.451899   0.372 0.710079
## TempB:LocalityRPV -1.077469    0.506006  -2.129 0.033318 *
## TempC:LocalityRPV -0.437372    0.486425  -0.899 0.368653
## TempB:LocalityRMO -2.026473    0.510236  -3.972 7.33e-05 ***
## TempC:LocalityRMO -1.771864    0.495576  -3.575 0.000356 ***
## TempB:LocalityTLC -0.003076    0.638906  -0.005 0.996159
## TempC:LocalityTLC  0.419779    0.604596   0.694 0.487548
## TempB:LocalityTPN -0.031690    0.992508  -0.032 0.974531
## TempC:LocalityTPN  0.795714    1.012193   0.786 0.431864
## TempB:LocalitySJU -2.571974    0.596253  -4.314 1.67e-05 ***
## TempC:LocalitySJU -2.238695    0.713524  -3.138 0.001723 **
## TempB:SexF       -0.593561    0.529894  -1.120 0.262753
## TempC:SexF       -1.724324    2.947086  -0.585 0.558535
## TempB:SexM              NA              NA      NA      NA
## TempC:SexM       -2.040178    2.903228  -0.703 0.482288
## LocalityAPR:SexF -0.585215    0.538355  -1.087 0.277118
## LocalityRPV:SexF  0.003271    0.567543   0.006 0.995402
## LocalityRMO:SexF -0.313481    0.572939  -0.547 0.584326
## LocalityTLC:SexF -0.219749    0.696766  -0.315 0.752495
## LocalityTPN:SexF -1.076552    1.069265  -1.007 0.314117
## LocalitySJU:SexF  0.095863    0.673549   0.142 0.886834
## LocalityAPR:SexM              NA              NA      NA      NA
## LocalityRPV:SexM              NA              NA      NA      NA
## LocalityRMO:SexM              NA              NA      NA      NA
## LocalityTLC:SexM              NA              NA      NA      NA
## LocalityTPN:SexM              NA              NA      NA      NA
## LocalitySJU:SexM              NA              NA      NA      NA
## TempB:LocalityAPR:SexF 1.303061    0.699049   1.864 0.062427 .
## TempC:LocalityAPR:SexF -0.084661    0.672263  -0.126 0.899794
## TempB:LocalityRPV:SexF 0.248134    0.731583   0.339 0.734506
```

```
## TempC:LocalityRPV:SexF -0.265909 0.718615 -0.370 0.711390
## TempB:LocalityRMO:SexF 0.649245 0.733667 0.885 0.376275
## TempC:LocalityRMO:SexF -0.091857 0.712794 -0.129 0.897471
## TempB:LocalityTLC:SexF 1.963332 0.934158 2.102 0.035674 *
## TempC:LocalityTLC:SexF 0.158383 0.913394 0.173 0.862351
## TempB:LocalityTPN:SexF 1.798372 1.400330 1.284 0.199169
## TempC:LocalityTPN:SexF 1.525218 1.457403 1.047 0.295413
## TempB:LocalitySJU:SexF 1.249390 0.870449 1.435 0.151310
## TempC:LocalitySJU:SexF -1.922464 1.013394 -1.897 0.057931 .
## TempB:LocalityAPR:SexM NA NA NA NA
## TempC:LocalityAPR:SexM NA NA NA NA
## TempB:LocalityRPV:SexM NA NA NA NA
## TempC:LocalityRPV:SexM NA NA NA NA
## TempB:LocalityRMO:SexM NA NA NA NA
## TempC:LocalityRMO:SexM NA NA NA NA
## TempB:LocalityTLC:SexM NA NA NA NA
## TempC:LocalityTLC:SexM NA NA NA NA
## TempB:LocalityTPN:SexM NA NA NA NA
## TempC:LocalityTPN:SexM NA NA NA NA
## TempB:LocalitySJU:SexM NA NA NA NA
## TempC:LocalitySJU:SexM NA NA NA NA
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for Gamma family taken to be 0.01316289)
##
## Null deviance: 102.052 on 2652 degrees of freedom
## Residual deviance: 33.967 on 2609 degrees of freedom
## (778 observations deleted due to missingness)
## AIC: -19713
##
## Number of Fisher Scoring iterations: 4
```

```
print("Binomial distribution results")
```

```
## [1] "Binomial distribution results"
```

```
print(modelAICb)
```

```
##      df      AIC
## mod1b 44 372.2764
## mod2b  3 290.2323
## mod3b  8 300.2330
## mod4b 21 326.2748
## mod5b  9 302.2572
## mod6b 20 324.2586
## mod7b 26 336.2721
## mod8b 12 308.2708
```

```
print("mod2b<-glm(rate~Temp,data=lifedat,family=binomial);summary(mod2b)")
```

```
## [1] "mod2b<-glm(rate~Temp,data=lifedat,family=binomial);summary(mod2b)"
```

```
print(summary(mod2b))
```

```
##
## Call:
## glm(formula = rate ~ Temp, family = binomial, data = lifedat)
##
## Deviance Residuals:
##      Min       1Q   Median       3Q      Max
## -0.103906  -0.024918   0.000161   0.020172   0.123856
##
## Coefficients:
##              Estimate Std. Error z value Pr(>|z|)
## (Intercept)  -3.0918     0.1585 -19.509  <2e-16 ***
## TempB         0.2155     0.2141   1.006   0.3142
## TempC         0.3723     0.2201   1.691   0.0908 .
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for binomial family taken to be 1)
##
##      Null deviance: 5.6422  on 2652  degrees of freedom
## Residual deviance: 2.7366  on 2650  degrees of freedom
## (778 observations deleted due to missingness)
## AIC: 290.23
##
## Number of Fisher Scoring iterations: 6
```

Including Plots

You can also embed plots, for example:

Note that the `echo = FALSE` parameter was added to the code chunk to prevent printing of the R code that generated the plot.