

Paper Figures

Table 1

Act	Travel	<i>add desc</i>
	Play	<i>add desc</i>
	Rest	<i>add desc</i>
	Eat	<i>add desc</i>
	Cling	<i>add desc</i>
Body Position	Hang	<i>add desc</i>
	Bipedal	<i>add desc</i>
	Lie	<i>add desc</i>
	Quadrupedal	<i>add desc</i>
	Sit	<i>add desc</i>
	Brachiate	<i>add desc</i>
	Slow Clamber	<i>add desc</i>
Tree Position	Ibu	<i>add desc</i>
	Branch	<i>add desc</i>
	Crotch	<i>add desc</i>
	Ground	<i>add desc</i>
	Liana	<i>add desc</i>
	Nest	<i>add desc</i>
	Trunk	<i>add desc</i>

CATEGORIES OF BEHAVIOR

Nonpoled	RestSitTrunk	RestHangTrunk	RestLieDownBranch
TravelHangBranch	EatBipedalTrunk	RestLieDownNest	RestLieDownCrotch
EatHangGround	TravelBrachiateLiana	EatBipedalLiana	EatHangLiana
RestHangGround	RestSitGround	EatBipedalCrotch	EatSitLiana
RestBipedalIbu	RestSitIbu	TravelSlowClamberLiana	PlayHangBranch
TravelSlowClamberNest	EatSitGround	EatHangTrunk	EatBipedalBranch
EatSitIbu	TravelSlowClamberTrunk	TravelSlowClamberCrotch	RestHangBranch
TravelTreeSwayTrunk	PlayHangIbu	PlaySitCrotch	RestHangIbu
EatQuadBranch	RestQuadIbu	PlayHangCrotch	EatHangBranch
RestHangNest	EatSitTrunk	RestHangCrotch	TravelSlowClamberBranch
TravelBrachiateGround	RestQuadBranch	PlayHangLiana	RestSitBranch
EatLieDownBranch	RestBipedalLiana	TravelTreeSwayBranch	
RestBipedalGround	EatHangIbu	PlaySitBranch	
RestBipedalTrunk	TravelQuadGround	RestSitLiana	
TravelSlowClamberGround	PlayBipedalBranch	RestSitLiana	
TravelQuadBranch	EatSitNest	RestBipedalBranch	
RestBipedalNest	RestBipedalCrotch	ClingHangIbu	

POSITIONAL BEHAVIOR WITH AGE

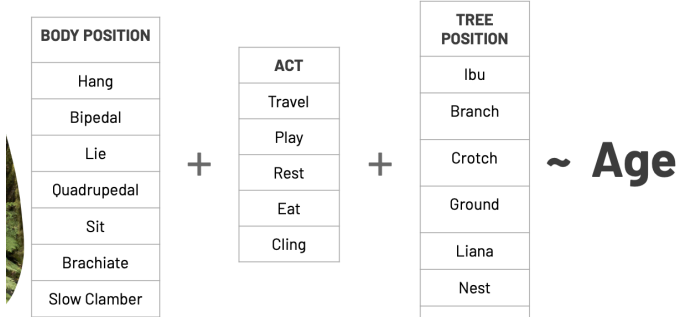
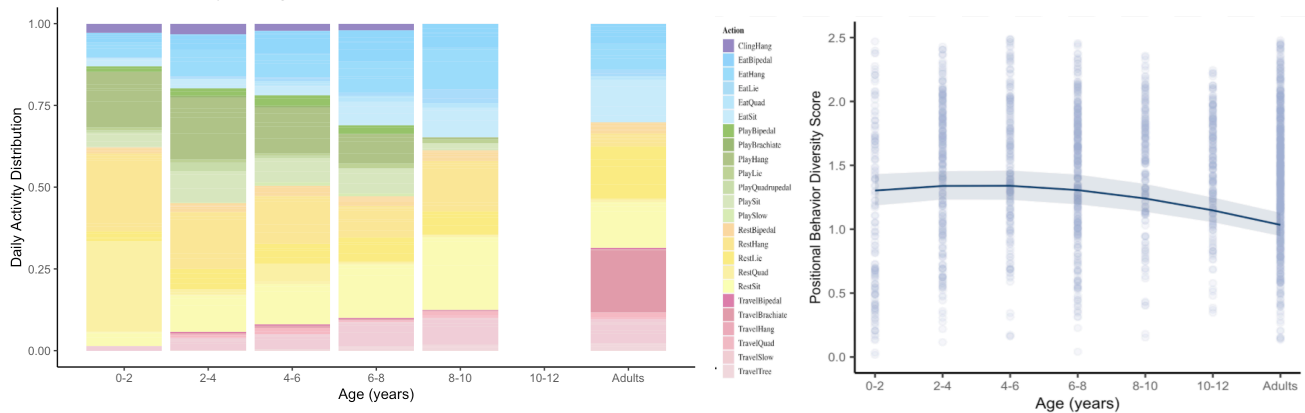


Figure 1



Statistic details:

Model Information (right graph)

```
> summary(model)
Family: gaussian ( log )
Formula: DiversityScore ~ Age + I(Age^2) + (1 | Name) + (1 | FollowNumber)
Data: dive_d

      AIC      BIC    logLik deviance df.resid
 2858.6   2893.2  -1423.3   2846.6     2346

Random effects:
Conditional model:
Groups      Name      Variance Std.Dev.
Name      (Intercept) 9.888e-02 3.144e-01
FollowNumber (Intercept) 6.059e-10 2.462e-05
Residual              1.826e-01 4.273e-01
Number of obs: 2352, groups: Name, 83; FollowNumber, 2339

Dispersion estimate for gaussian family (sigma^2): 0.183

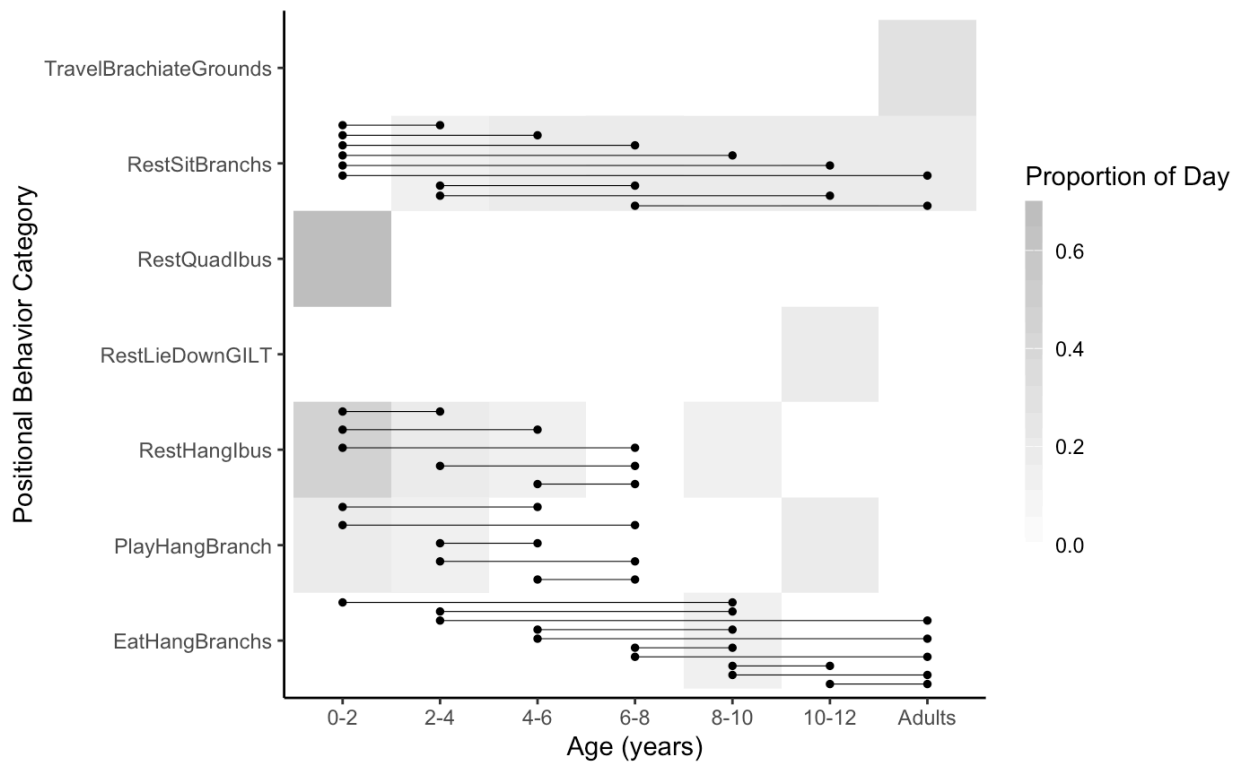
Conditional model:
              Estimate Std. Error z value Pr(>|z|)
(Intercept)  0.210762   0.066167   3.185 0.001446 **
Age          0.066736   0.032439   2.057 0.039663 *
I(Age^2)     -0.013153   0.003802  -3.459 0.000542 ***
---
Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

Age makes the model better (anova):

```
> # Test that adding age actually made the model better
> anova(model.test1,model2a)
Data: dive_d
Models:
model.test1: DiversityScore ~ (1 | Name) + (1 | FollowNumber), zi=0, disp=-1
model2a: DiversityScore ~ Age + (1 | Name) + (1 | FollowNumber), zi=0, disp=-1
      Df    AIC      BIC    logLik deviance Chisq Chi Df Pr(>Chisq)
model.test1  4 2953.2 2976.2 -1472.6  2945.2
model2a      5 2868.8 2897.6 -1429.4  2858.8 86.379      1 < 2.2e-16 ***
---
Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
> anova(model2a,model2)
Data: dive_d
Models:
model2a: DiversityScore ~ Age + (1 | Name) + (1 | FollowNumber), zi=0, disp=-1
model2: DiversityScore ~ Age + I(Age^2) + (1 | Name) + (1 | FollowNumber), zi=0, disp=-1
      Df    AIC      BIC    logLik deviance Chisq Chi Df Pr(>Chisq)
model2a      5 2868.8 2897.6 -1429.4  2858.8
model2       6 2858.6 2893.2 -1423.3  2846.6 12.139      1 0.0004938 ***
---
Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

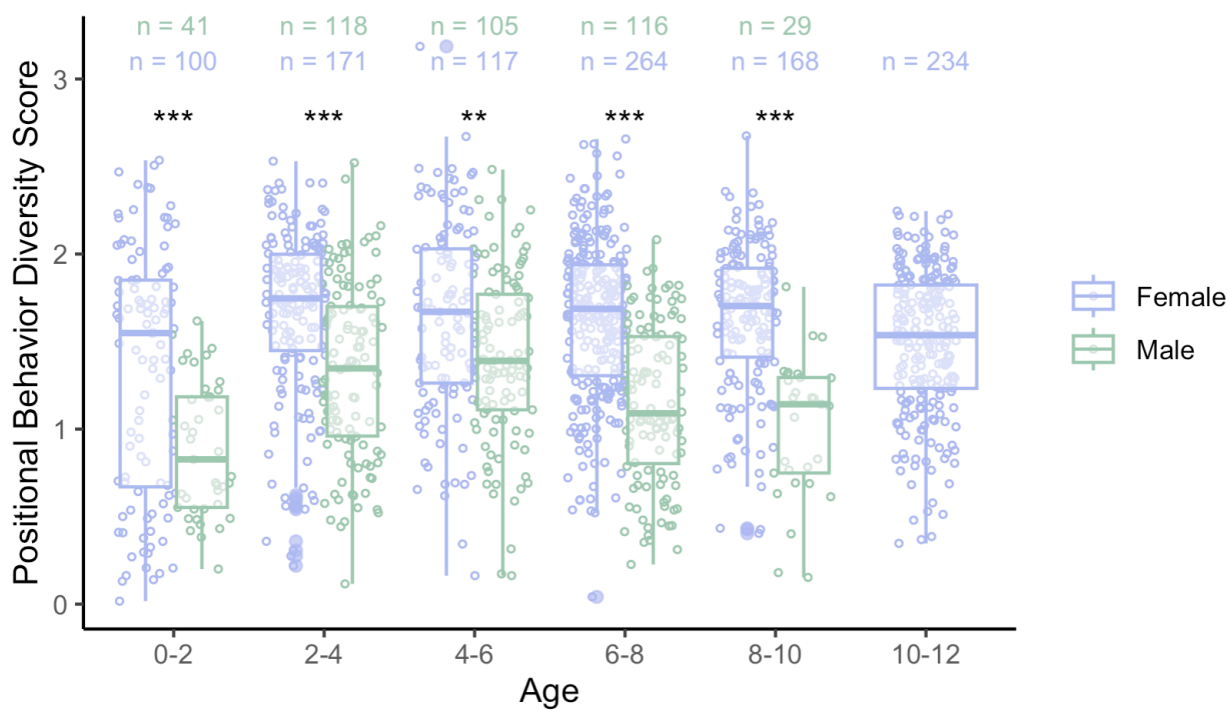
Figure 2

Only showing > 0.1 proportion of day spent, and p tests are all below 0.002 (bonferoni)



All data in Figure2_SummaryMatrix.xlsx. True if less than 0.002.

Figure 3

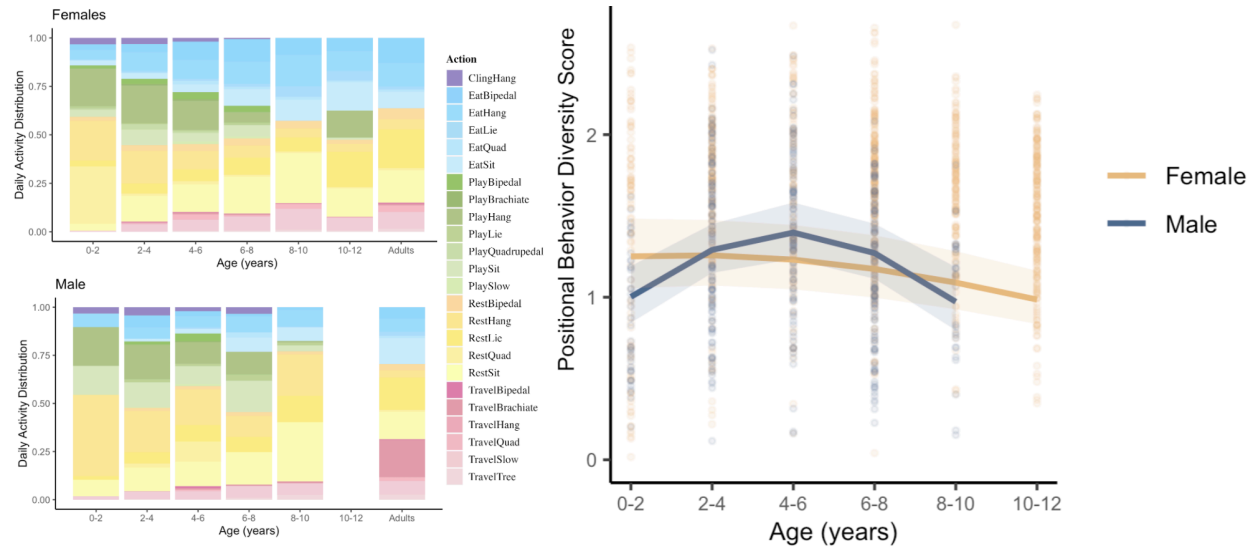


Statistic results:

T-test pvalues (1 = Age 0-2)

	x
1	3.342075e-06
2	2.645750e-08
3	1.092318e-03
4	7.151492e-19
5	2.902526e-09

Figure 4



Statistic results:

Females

Model Information (right graph)

```
> summary(model_F)
Family: gaussian ( log )
Formula: DiversityScore ~ Age + I(Age^2) + (1 | Name) + (1 | FollowNumber)
Data: dive_d

            AIC      BIC    logLik deviance df.resid
1122.9    1152.6   -555.4    1110.9      1048

Random effects:
Groups Name Variance Std.Dev.
Name (Intercept) 1.436e-01 3.789e-01
FollowNumber (Intercept) 3.435e-10 1.853e-05
Residual 1.566e-01 3.957e-01
Number of obs: 1054, groups: Name, 28; FollowNumber, 1054

Dispersion estimate for gaussian family (sigma^2): 0.157

Conditional model:
            Estimate Std. Error z value Pr(>|z|)
(Intercept)  0.192654    0.104806   1.838  0.06603 .
Age          0.044793    0.040423   1.108  0.26782
I(Age^2)     -0.013219    0.004911  -2.692  0.00711 **
---
Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

Age makes the model better (anova):

```
Data: dive_d
Models:
model.1: DiversityScore ~ (1 | Name) + (1 | FollowNumber), zi=0, disp=-1
model.2: DiversityScore ~ Age + (1 | Name) + (1 | FollowNumber), zi=0, disp=-1
            Df      AIC      BIC    logLik deviance Chisq Chi Df Pr(>Chisq)
model.1  4 1189.0 1208.8 -590.49  1181.0
model.2  5 1128.3 1153.1 -559.14  1118.3 62.704      1 2.402e-15 ***
---
Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

Data: dive_d
Models:
model.2: DiversityScore ~ Age + (1 | Name) + (1 | FollowNumber), zi=0, disp=-1
model.3: DiversityScore ~ Age + I(Age^2) + (1 | Name) + (1 | FollowNumber), zi=0, disp=-1
            Df      AIC      BIC    logLik deviance Chisq Chi Df Pr(>Chisq)
model.2  5 1128.3 1153.1 -559.14  1118.3
model.3  6 1122.9 1152.6 -555.44  1110.9 7.3992      1 0.006525 **
---
Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

Males

Model Information (right graph)

```
> summary(model_M)
Family: gaussian ( log )
Formula: DiversityScore ~ Age + I(Age^2) + (1 | Name) + (1 | FollowNumber)
Data: dive_d

            AIC      BIC    logLik deviance df.resid
507.1    531.2   -247.5    495.1      403

Random effects:
Groups Name Variance Std.Dev.
Name (Intercept) 3.430e-02 1.852e-01
FollowNumber (Intercept) 3.143e-09 5.607e-05
Residual 1.848e-01 4.299e-01
Number of obs: 409, groups: Name, 14; FollowNumber, 409

Dispersion estimate for gaussian family (sigma^2): 0.185

Conditional model:
            Estimate Std. Error z value Pr(>|z|)
(Intercept) -0.42457    0.17578  -2.415  0.0157 *
Age          0.51400    0.12120   4.241 2.23e-05 ***
I(Age^2)     -0.08695    0.01993  -4.363 1.28e-05 ***
---
Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

Age makes the model better (anova):

```
Data: dive_d
Models:
model.1: DiversityScore ~ (1 | Name) + (1 | FollowNumber), zi=0, disp=-1
model.2: DiversityScore ~ Age + (1 | Name) + (1 | FollowNumber), zi=0, disp=-1
            Df      AIC      BIC    logLik deviance Chisq Chi Df Pr(>Chisq)
model.1  4 524.10 540.15 -258.05  516.10
model.2  5 525.98 546.05 -257.99  515.98 0.1171      1 0.7322
Data: dive_d
Models:
model.2: DiversityScore ~ Age + (1 | Name) + (1 | FollowNumber), zi=0, disp=-1
model.3: DiversityScore ~ Age + I(Age^2) + (1 | Name) + (1 | FollowNumber), zi=0, disp=-1
            Df      AIC      BIC    logLik deviance Chisq Chi Df Pr(>Chisq)
model.2  5 525.98 546.05 -257.99  515.98
model.3  6 507.07 531.15 -247.54  495.07 20.908      1 4.818e-06 ***
---
Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

Figure 5

Only showing rows where atleast one comparison was significant
Bonferoni corrections done. All code in Figure5.rmd

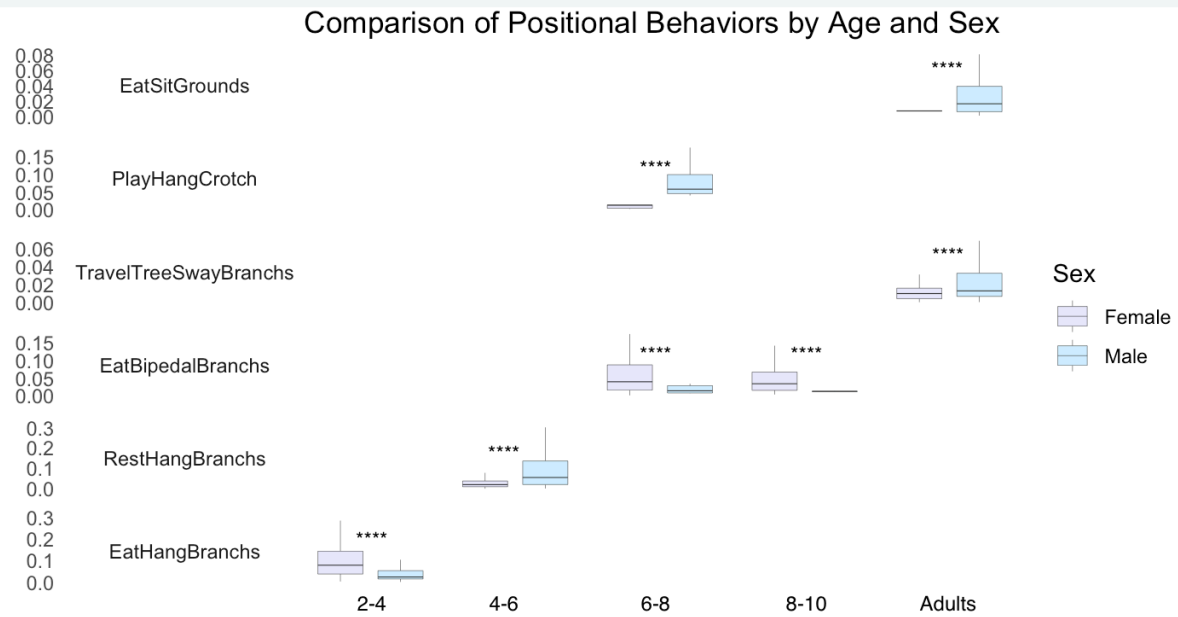
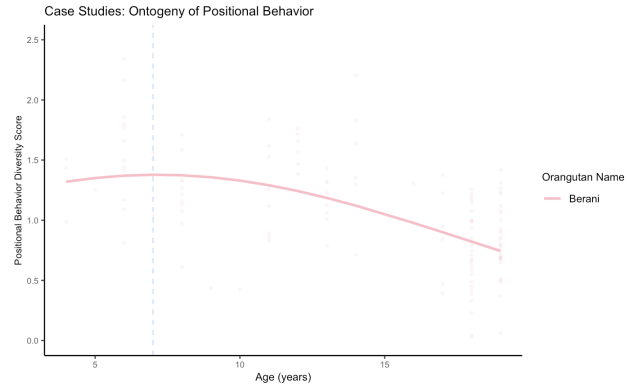
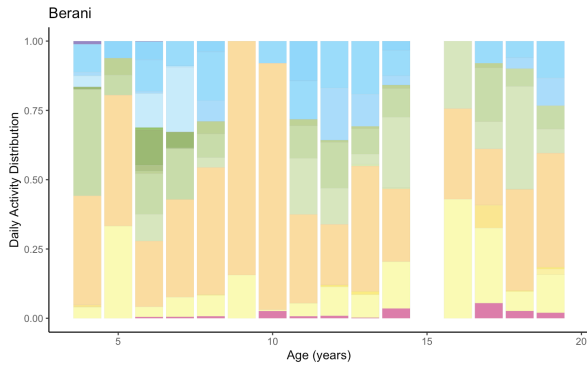


Figure 6



Statistic results:

Berani

Model Information (right graph)

|

Age makes the model better (anova):

```
> summary(model_berani)
Family: gaussian ( log )
Formula: DiversityScore ~ Age + I(Age^2) + (1 | Name) + (1 | FollowNumber)
Data: dive_d

      AIC      BIC logLik deviance df.resid
118.1    135.3   -53.0    106.1      125

Random effects:
Conditional model:
Groups      Name      Variance Std.Dev.
Name      (Intercept) 5.364e-12 2.316e-06
FollowNumber (Intercept) 3.569e-02 1.889e-01
Residual      9.216e-02 3.036e-01
Number of obs: 131, groups: Name, 1; FollowNumber, 131

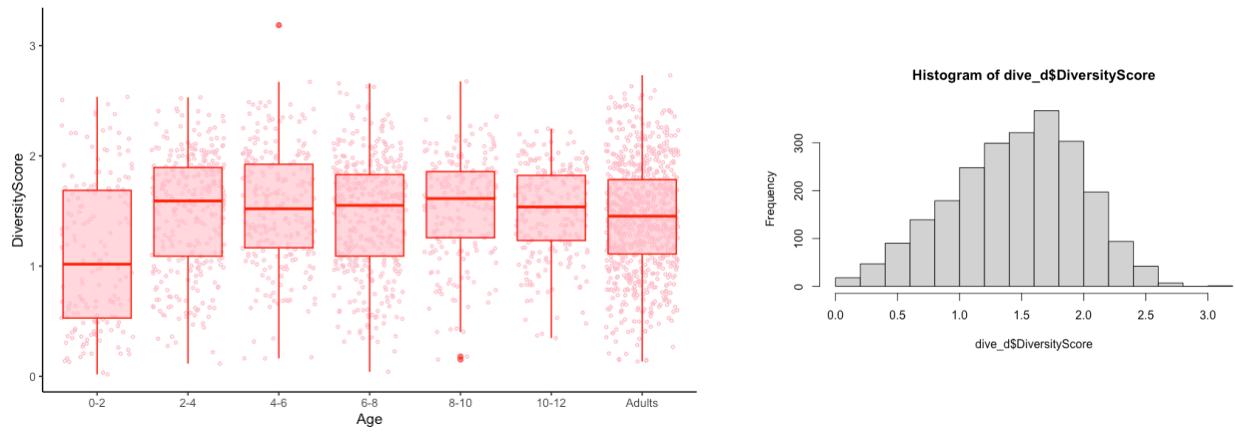
Dispersion estimate for gaussian family (sigma^2): 0.0922

Conditional model:
      Estimate Std. Error z value Pr(>|z|)
(Intercept)  0.098249   0.227714   0.432  0.66614
Age          0.062407   0.041346   1.509  0.13120
I(Age^2)     -0.004374   0.001650  -2.651  0.00803 **
---
Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

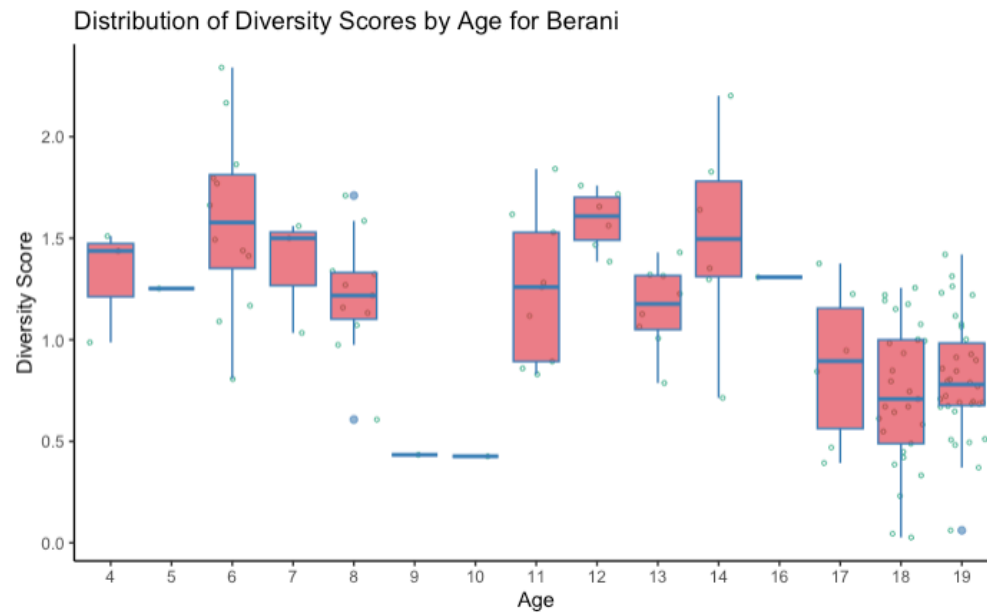
```
> model_berani <- best_model(subset(dive_d, Name == "berani"))
Data: dive_d
Models:
model.1: DiversityScore ~ (1 | Name) + (1 | FollowNumber), zi=~0, disp=~1
model.2: DiversityScore ~ Age + (1 | Name) + (1 | FollowNumber), zi=~0, disp=~1
      Df    AIC    BIC logLik deviance Chisq Chi Df Pr(>Chisq)
model.1  4 169.47 180.97 -80.733   161.47
model.2  5 122.93 137.30 -56.464   112.93 48.537      1 3.241e-12 ***
---
Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
Data: dive_d
Models:
model.2: DiversityScore ~ Age + (1 | Name) + (1 | FollowNumber), zi=~0, disp=~1
model.3: DiversityScore ~ Age + I(Age^2) + (1 | Name) + (1 | FollowNumber), zi=~0, disp=~1
      Df    AIC    BIC logLik deviance Chisq Chi Df Pr(>Chisq)
model.2  5 122.93 137.30 -56.464   112.93
model.3  6 118.08 135.33 -53.037   106.08 6.8535      1 0.008847 **
---
Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

Supplemental Information

Distribution of diversity scores for combined dataset (both sex)



Distribution of diversity scores for Berani



Chi-squared testing of all ages

Code in Figure1.Rmd

	0-2	2-4	4-6	6-8	8-10	10-12	Adults
0-2	ns	***	***	***	***	***	***
2-4	***	ns	***	***	***	***	***
4-6	***	***	ns	***	***	***	***
6-8	***	***	***	ns	***	***	***
8-10	***	***	***	***	ns	***	***
10-12	***	***	***	***	***	ns	***
Adults	***	***	***	***	***	***	ns

Expected distribution = top row

Chi-squared testing of all ages of Female vs Male

In Figure 3&4.Rmd

	Class	stat	pval
1	Age 0-2	Inf	0
2	Age 2-4	Inf	0
3	Age 4-6	Inf	0
4	Age 6-8	Inf	0
5	Age 8-10	Inf	0

List of positional behavior categories

Nonpooled

TravelHangBranch
EatHangGround
RestHangGround
RestBipedallbu
TravelSlowClamberNest
EatSitlbu
TravelTreeSwayTrunk
EatQuadBranch
RestHangNest
TravelBrachiateGround
EatLieDownBranch
RestBipedalGround
RestBipedalTrunk
TravelSlowClamberGround
TravelQuadBranch
RestBipedalNest

RestSitTrunk

EatBipedalTrunk
TravelBrachiateLiana
RestSitGround
RestSitlbu
EatSitGround
TravelSlowClamberTrunk
PlayHanglbu
RestQuadlbu
EatSitTrunk
RestQuadBranch
RestBipedalLiana
EatHanglbu
TravelQuadGround
PlayBipedalBranch
EatSitNest
RestBipedalCrotch

RestHangTrunk

RestLieDownNest
EatBipedalLiana
EatBipedalCrotch
TravelSlowClamberLiana
EatHangTrunk
TravelSlowClamberCrotch
PlaySitCrotch
PlayHangCrotch
RestHangCrotch
PlayHangLiana
TravelTreeSwayBranch
PlaySitBranch
RestHangLiana
RestSitLiana
RestBipedalBranch
ClingHanglbu

RestLieDownBranch

RestLieDownCrotch
EatHangLiana
EatSitLiana
PlayHangBranch
EatBipedalBranch
RestHangBranch
RestHanglbu
EatHangBranch
TravelSlowClamberBranch
RestSitBranch

Pooled

PlayBrachiate BL
PlayQuadrupedal BCLINT
PlayHang GNT
TravelBrachiate NT
PlaySlowClamber BCLT
TravelTreeSway CGLN
PlayLieDown BCLINT
EatBipedal GIN
TravelQuad CLNT
TravelBipedal BCGLNT
EatQuad CGILT
PlayBipedal CILNT
EatLieDown CGILNT
PlaySit ILNT
RestLieDown GILT
RestQuad CGLNT