• The original analysis on the untransformed data:

```
Fixed effects:
                  Estimate Std. Error
                                         df t value Pr(>ItI)
                               76.24
                                      50.07 20.577
(Intercept)
                   1568.75
                                                      <2e-16 ***
Context1
                   -36.20
                               86.01
                                      29.77 -0.421
                                                      0.6768
                   -69.01
                               39.87
                                      25.93 -1.731
                                                      0.0954 .
Sentence1
Context1:Sentence1 -168.73
                               80.36
                                      25.51 -2.100
                                                      0.0458 *
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

The new analysis on the log transformed data:

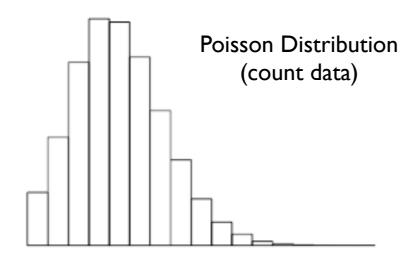
```
Fixed effects:
                  Estimate Std. Error
                                            df t value
(Intercept)
                   7.23975
                              0.04967 49.13000 145.761
Sentence1
                              0.05278 29.03000
                                                 0.264
                   0.01392
                                                 1.911
Context1
                   0.04316
                              0.02258 28.62000
Sentence1:Context1 -0.09333
                              0.04618 25.55000
                                                -2.021
```

t-value of the interaction smaller than in analysis over untransformed data. With similar dfs, p will be bigger.

Other distributions under the GLMM via the function glmer are available...

Usage

```
family(object, ...)
binomial(link = "logit")
gaussian(link = "identity")
Gamma(link = "inverse")
inverse.gaussian(link = "1/mu^2")
poisson(link = "log")
quasi(link = "identity", variance = "constant")
quasibinomial(link = "logit")
quasipoisson(link = "log")
```



Normal (Gaussian) Distribution



