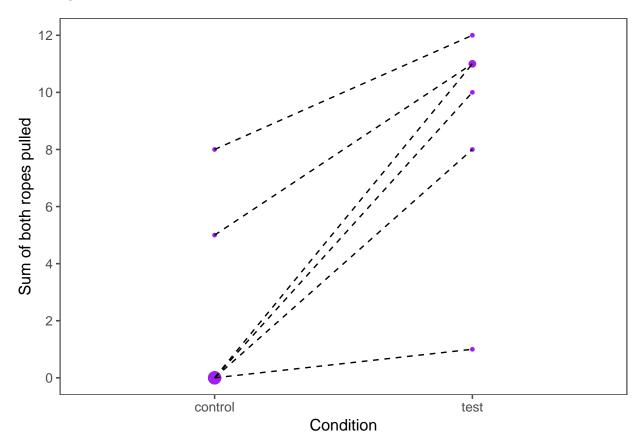
$multiple_possibilities_GLMM$

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Summary

Both the paired-samples t-test and a mixed model show a significant effect for condition. There was no significant interaction between session and condition.

Plotting data



Paired-sample t-test

```
contr_data <- plot_individual %>%
  filter(condition == "control")
exp_data <- plot_individual %>%
  filter(condition == "test")
t.test(exp_data$correct_resp, contr_data$correct_resp, paired = TRUE, alternative = "two.sided")
##
## Paired t-test
##
## data: exp_data$correct_resp and contr_data$correct_resp
## t = 4.3234, df = 5, p-value = 0.007546
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
   2.702818 10.630515
## sample estimates:
## mean of the differences
                  6.666667
```

GLMM

GLMM 01 with session * condition interaction

Confidence intervals

Confidence intervals for the binomial models were derived using the function bootMer of the R package lme4, using 1,000 parametric bootstraps and bootstrapping over the random effects.

```
boot.res=boot.glmm(model.res=mm.1, excl.warnings=T,nboots=1000, para=F)
saveRDS(boot.res, "multpos_model.rds")
```

Null model

singular fit

Full-null model comparison

```
anova(mm.1, mm.1.null, test="Chisq")
## Data: model.data
## Models:
## mm.1.null: both_ropes ~ 1 + (1 + z.session | chimp)
## mm.1: both_ropes ~ condition * z.session + z.trial + (1 + z.session |
            chimp)
## mm.1:
##
            Df
                  AIC
                         BIC logLik deviance Chisq Chi Df Pr(>Chisq)
## mm.1.null 4 183.37 195.25 -87.686
                                     175.373
             8 111.67 135.43 -47.833
                                       95.667 79.706
                                                            < 2.2e-16 ***
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
```

Model output

<none>

z.trial

• Coefficients

```
round(summary(mm.1)$coefficients, 3)
```

```
Estimate Std. Error z value Pr(>|z|)
##
## (Intercept)
                              -3.901
                                          1.827 -2.135
                                                            0.033
                               6.366
                                                            0.001
## conditiontest
                                          1.945
                                                 3.272
## z.session
                               1.608
                                          1.291
                                                  1.246
                                                            0.213
## z.trial
                               0.714
                                          0.312
                                                  2.288
                                                            0.022
## conditiontest:z.session
                             -2.368
                                          1.853 -1.278
                                                            0.201
```

111.67 1 115.65 5.981

• Individual predictors: likelihood ratio tests

Drop1: P values for the individual effects were based on likelihood ratio tests comparing the full with respective reduced models (Barr et al., 2013; R function drop1 with argument 'test' set to "Chisq").

```
xdrop1=drop1(mm.1, test="Chisq")

## singular fit
## singular fit

round(xdrop1,3)

## Single term deletions
##
## Model:
## both_ropes ~ condition * z.session + z.trial + (1 + z.session |
## chimp)
## Df AIC LRT Pr(Chi)
```

0.014 *

```
## condition:z.session 1 113.25 3.580 0.058 .
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
```

GLMM02 without the interaction

Confidence intervals

Confidence intervals for the binomial models were derived using the function bootMer of the R package lme4, using 1,000 parametric bootstraps and bootstrapping over the random effects.

```
boot.res2=boot.glmm(model.res=mm.2, excl.warnings=T,nboots=1000, para=F)
saveRDS(boot.res2, "multpos_model2.rds")

## orig X2.5. X97.5.
## (Intercept) -2.688 -6.605 -0.204
## conditiontest 4.389 2.969 7.883
## z.session 0.256 -0.335 0.950
## z.trial 0.622 0.063 1.438
```

Null model

Full-null model comparison

```
anova(mm.2, mm.2.null, test="Chisq")

## Data: model.data
## Models:
## mm.2.null: both_ropes ~ 1 + (1 | chimp)
## mm.2: both_ropes ~ condition + z.session + z.trial + (1 | chimp)
## Df AIC BIC logLik deviance Chisq Chi Df Pr(>Chisq)
## mm.2.null 2 179.38 185.32 -87.691 175.382
## mm.2 5 109.25 124.10 -49.626 99.253 76.129 3 < 2.2e-16 ***
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1</pre>
```

Model output

Coefficients

```
round(summary(mm.2)$coefficients, 3)
```

```
Estimate Std. Error z value Pr(>|z|)
##
## (Intercept)
                    -3.069
                                 1.248 - 2.460
                                                  0.014
                     5.095
                                0.980
                                         5.198
                                                  0.000
## conditiontest
## z.session
                     0.156
                                 0.281
                                         0.555
                                                  0.579
## z.trial
                     0.681
                                0.300
                                         2.268
                                                  0.023
```

• Individual predictors: likelihood ratio tests

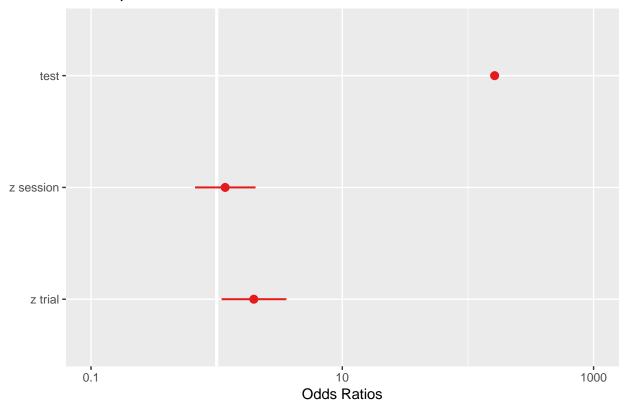
Drop1: P values for the individual effects were based on likelihood ratio tests comparing the full with

```
respective reduced models (Barr et al., 2013; R function drop1 with argument 'test' set to "Chisq").
xdrop1=drop1(mm.2, test="Chisq")
round(xdrop1,3)
## Single term deletions
##
## Model:
## both_ropes ~ condition + z.session + z.trial + (1 | chimp)
##
             Df
                    AIC
                           LRT Pr(Chi)
                 109.25
## <none>
## condition 1 180.44 73.184
                               <2e-16 ***
## z.session 1 107.56 0.310
                                 0.578
## z.trial
              1 113.00 5.743
                                 0.017 *
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
check for colinearity in the previous model.
## condition
                z.trial z.session
## 1.042645 1.040646 1.002028
-> no collinearity
```

Model output

```
plot_model(mm.2, type='est')
```

both ropes



plot_model(mm.2, type='re')

Random effects

