Binary Choice Analysis

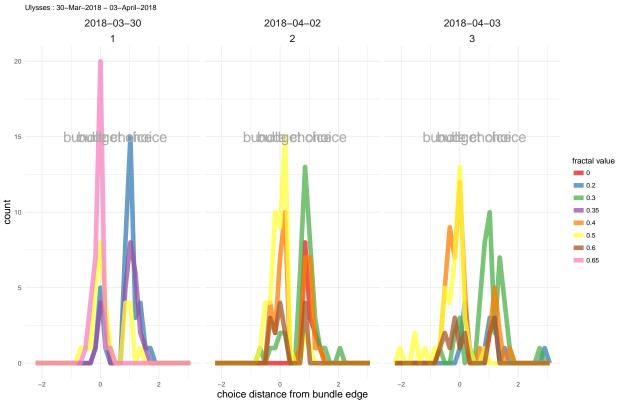
Robert Hickman 22 February 2018

```
monkey <- "Ulysses"
today <- "03-April-2018"
look_back <- "30-Mar-2018"

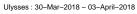
start_trial <- 0
stop_trial <- "all"

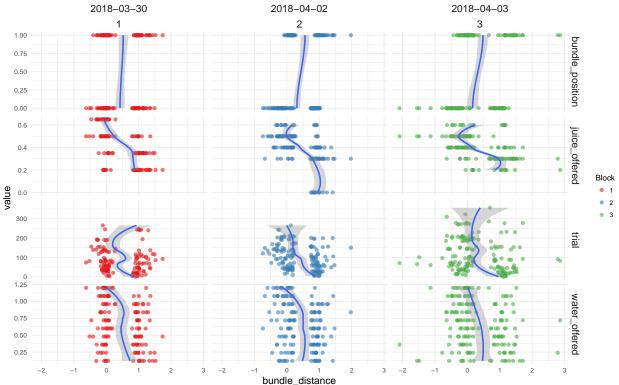
merge_days <- TRUE</pre>
p1
```

Monkey Choice Distance From Bundle on Binary Choice Task



Monkey Choice Distance From Bundle on Binary Choice Task





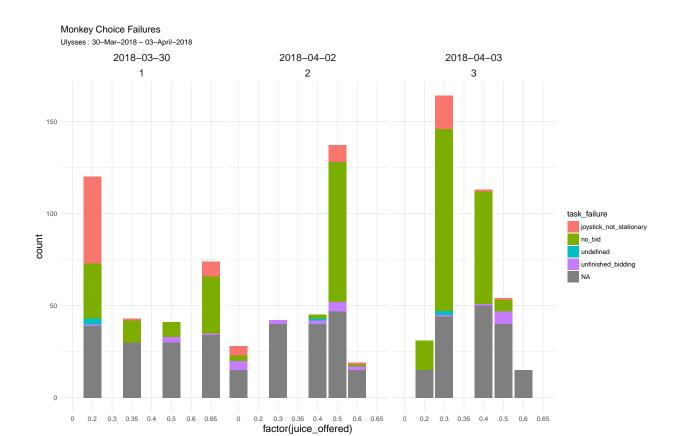
```
##
## Call:
   glm(formula = fractal_choice ~ bundle_position + water_offered +
       juice_offered + trial + date, family = "binomial", data = task_data)
##
##
## Deviance Residuals:
##
       Min
                 10
                      Median
                                   3Q
                                           Max
## -2.8842 -0.4709
                      0.1323
                               0.5323
                                         2.4840
##
## Coefficients:
                     Estimate Std. Error z value Pr(>|z|)
##
## (Intercept)
                   -1.704e+03 1.538e+03
                                          -1.107 0.26808
## bundle position 8.268e-01
                              2.878e-01
                                           2.872 0.00407 **
## water_offered
                    3.607e+00 4.879e-01
                                           7.392 1.44e-13 ***
## juice_offered
                    1.650e+01
                               1.624e+00
                                          10.159 < 2e-16 ***
## trial
                    1.373e-02 2.226e-03
                                           6.171 6.80e-10 ***
## date
                    9.610e-02 8.730e-02
                                           1.101 0.27097
## ---
```

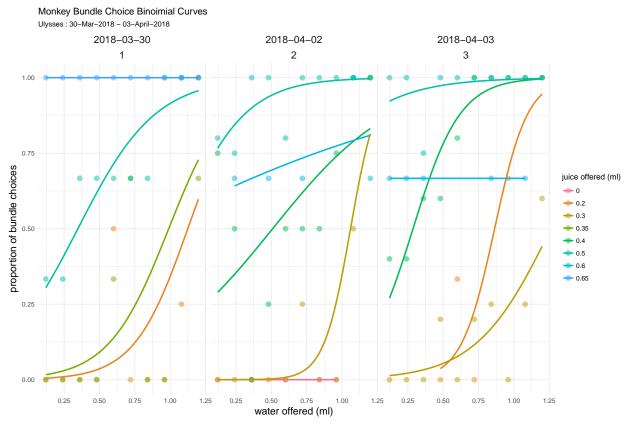
```
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for binomial family taken to be 1)
##
      Null deviance: 621.95 on 453 degrees of freedom
## Residual deviance: 326.20 on 448 degrees of freedom
     (472 observations deleted due to missingness)
## AIC: 338.2
## Number of Fisher Scoring iterations: 6
#test for side bias with an exact binomial test
binom.test(c(nrow(task_data %>%
                    .[c(bundle_position != fractal_choice)]),
             nrow(task_data %>%
                    .[c(bundle_position == fractal_choice)])))
##
## Exact binomial test
## data: c(nrow(task_data %% .[c(bundle_position != fractal_choice)]),
                                                                              nrow(task_data %>% .[c(bunder)]
## number of successes = 221, number of trials = 454, p-value =
## alternative hypothesis: true probability of success is not equal to 0.5
## 95 percent confidence interval:
## 0.4399220 0.5338195
## sample estimates:
## probability of success
                0.4867841
#generate a model of likelihood to choice for the fractal dependent on it's position,
#value and associated water
model <- glm(data = dplyr::filter(task_data, block_no == max(block_no)),</pre>
             fractal_choice ~ bundle_position + water_offered + as.factor(juice_offered) + trial + date
             family = "binomial")
#summarise the parameters
summary(model)
##
## Call:
## glm(formula = fractal_choice ~ bundle_position + water_offered +
       as.factor(juice_offered) + trial + date, family = "binomial",
##
       data = dplyr::filter(task_data, block_no == max(block_no)))
##
## Deviance Residuals:
##
        Min
                         Median
                                       3Q
                                                Max
## -2.59408 -0.29321
                        0.08482
                                  0.36811
                                            1.98060
## Coefficients: (1 not defined because of singularities)
                                Estimate Std. Error z value Pr(>|z|)
                                           1.431300 -4.400 1.08e-05 ***
## (Intercept)
                               -6.297877
## bundle_position
                                0.770928
                                           0.576296
                                                      1.338 0.180985
## water_offered
                                           1.057643
                                                      4.390 1.14e-05 ***
                                4.642533
```

0.863501 -1.178 0.238663

as.factor(juice_offered)0.3 -1.017494

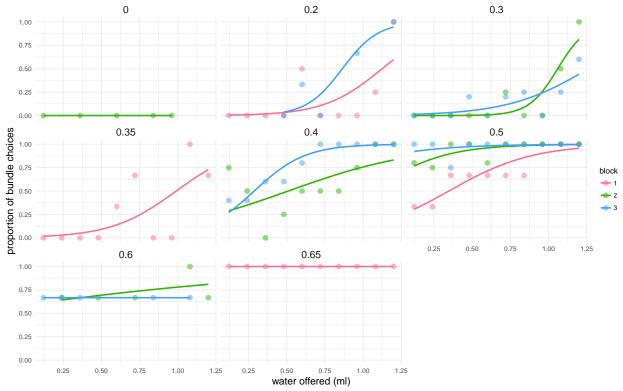
```
## as.factor(juice_offered)0.4 3.235756
                                           0.948393
                                                      3.412 0.000645 ***
## as.factor(juice_offered)0.5 6.501167
                                           1.440674
                                                      4.513 6.40e-06 ***
                                           1.094002
## as.factor(juice_offered)0.6 3.279973
                                                      2.998 0.002716 **
                                           0.003394
                                                      3.793 0.000149 ***
## trial
                                0.012875
## date
                                      NA
                                                 NA
                                                         NA
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## (Dispersion parameter for binomial family taken to be 1)
##
##
       Null deviance: 218.47 on 163 degrees of freedom
## Residual deviance: 94.32 on 156 degrees of freedom
     (213 observations deleted due to missingness)
## AIC: 110.32
##
## Number of Fisher Scoring iterations: 6
#test for side bias with an exact binomial test
binom.test(c(nrow(task_data %>%
                    .[c(bundle_position != fractal_choice & block_no == max(block_no))]),
             nrow(task_data %>%
                    .[c(bundle_position == fractal_choice & block_no == max(block_no))])))
##
##
   Exact binomial test
##
## data: c(nrow(task\_data \%)\% .[c(bundle\_position != fractal\_choice & 
                                                                           block_no == max(block_no))])
## number of successes = 74, number of trials = 164, p-value = 0.2414
## alternative hypothesis: true probability of success is not equal to 0.5
## 95 percent confidence interval:
## 0.3735194 0.5307146
## sample estimates:
## probability of success
##
                0.4512195
рЗ
```





Monkey Bundle Choice Binoimial Curves Ulysses: 30-Mar-2018 - 03-April-2018

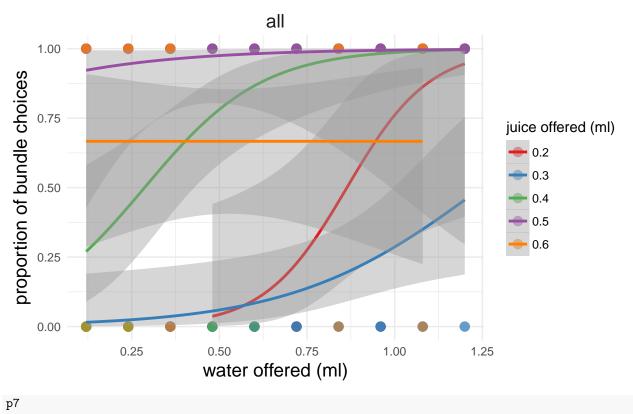




р6

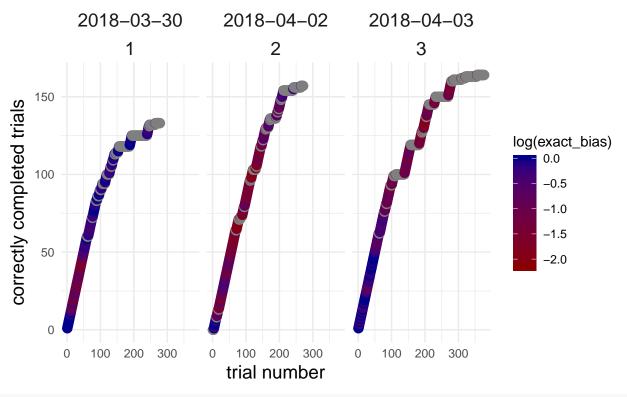
Today's Monkey Bundle Choice Binoimial Curves

Ulysses: 03-April-2018



Monkey Trial Progression and Bias

Ulysses: 30-Mar-2018 - 03-April-2018



Monkey Trial Progression and Bias

Ulysses: 30-Mar-2018 - 03-April-2018

