

Binary Choice Analysis

Robert Hickman

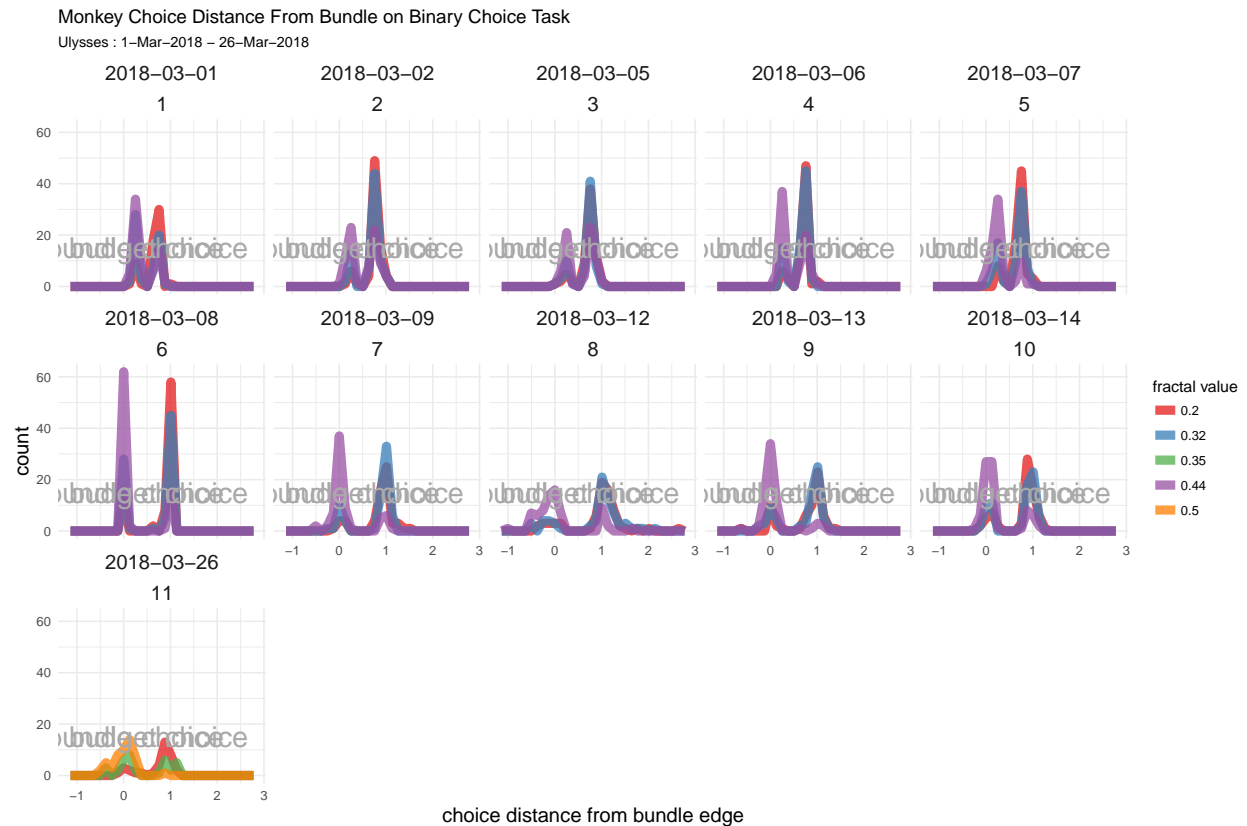
22 February 2018

```
monkey <- "Ulysses"  
today <- "26-Mar-2018"  
look_back <- "1-Mar-2018"
```

```
start_trial <- 0  
stop_trial <- "all"
```

```
merge_days <- TRUE
```

p1

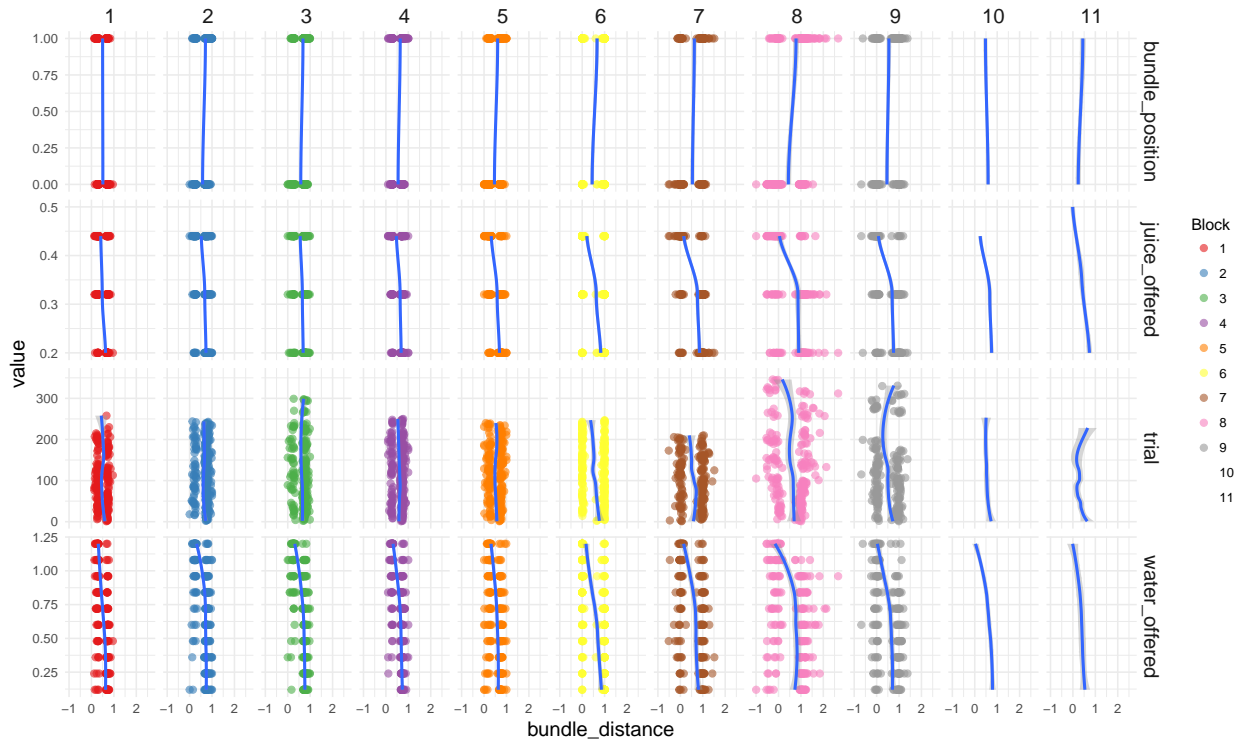


p2

Monkey Choice Distance From Bundle on Binary Choice Task

Ulysses : 1-Mar-2018 – 26-Mar-2018

018-03-0 018-03-0 018-03-0 018-03-0 018-03-0 018-03-0 018-03-0 018-03-1 018-03-1 018-03-1 018-03-2



*#generate a model of likelihood to choice for the fractal dependent on it's position,
#value and associated water*

```
model <- glm(data = task_data,
             fractal_choice ~ bundle_position + water_offered + juice_offered + trial + date,
             family = "binomial")
```

#summarise the parameters

```
summary(model)
```

```
##
## Call:
## glm(formula = fractal_choice ~ bundle_position + water_offered +
##     juice_offered + trial + date, family = "binomial", data = task_data)
##
## Deviance Residuals:
##      Min       1Q   Median       3Q      Max
## -2.7723  -0.5273  -0.1401   0.4882   3.2252
##
## Coefficients:
##              Estimate Std. Error z value Pr(>|z|)
## (Intercept)  -1.518e+03  2.021e+02  -7.514 5.73e-14 ***
## bundle_position -9.326e-01  1.277e-01  -7.303 2.82e-13 ***
## water_offered  5.450e+00  2.548e-01  21.391 < 2e-16 ***
## juice_offered  1.874e+01  8.705e-01  21.532 < 2e-16 ***
## trial         4.780e-03  8.391e-04   5.697 1.22e-08 ***
## date          8.567e-02  1.148e-02   7.465 8.32e-14 ***
## ---
```

```

## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for binomial family taken to be 1)
##
##    Null deviance: 3106.0  on 2305  degrees of freedom
## Residual deviance: 1645.4  on 2300  degrees of freedom
##    (850 observations deleted due to missingness)
## AIC: 1657.4
##
## 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(bun
## number of successes = 1272, number of trials = 2306, p-value =
## 7.831e-07
## alternative hypothesis: true probability of success is not equal to 0.5
## 95 percent confidence interval:
##  0.5310344 0.5720433
## sample estimates:
## probability of success
##          0.5516045
#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)),
             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       1Q   Median       3Q      Max
## -3.13406  -0.25233   0.05239   0.31819   2.28487
##
## Coefficients: (1 not defined because of singularities)
##              Estimate Std. Error z value Pr(>|z|)
## (Intercept)    -6.698980    1.650324  -4.059 4.92e-05 ***
## bundle_position     0.395591    0.687430   0.575 0.564977
## water_offered      6.185261    1.481074   4.176 2.96e-05 ***
## as.factor(juice_offered)0.35  3.193386    0.921399   3.466 0.000529 ***

```

```

## as.factor(juice_offered)0.5    7.686337    1.519365    5.059 4.22e-07 ***
## trial                        0.003018    0.005533    0.546 0.585395
## date                         NA          NA          NA      NA
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for binomial family taken to be 1)
##
##    Null deviance: 157.472  on 120  degrees of freedom
## Residual deviance:  60.274  on 115  degrees of freedom
##    (128 observations deleted due to missingness)
## AIC: 72.274
##
## Number of Fisher Scoring iterations: 7
#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 = 62, number of trials = 121, p-value = 0.8558
## alternative hypothesis: true probability of success is not equal to 0.5
## 95 percent confidence interval:
##  0.4198745 0.6042979
## sample estimates:
## probability of success
##      0.5123967
p3

```



p4

Monkey Bundle Choice Binoimial Curves

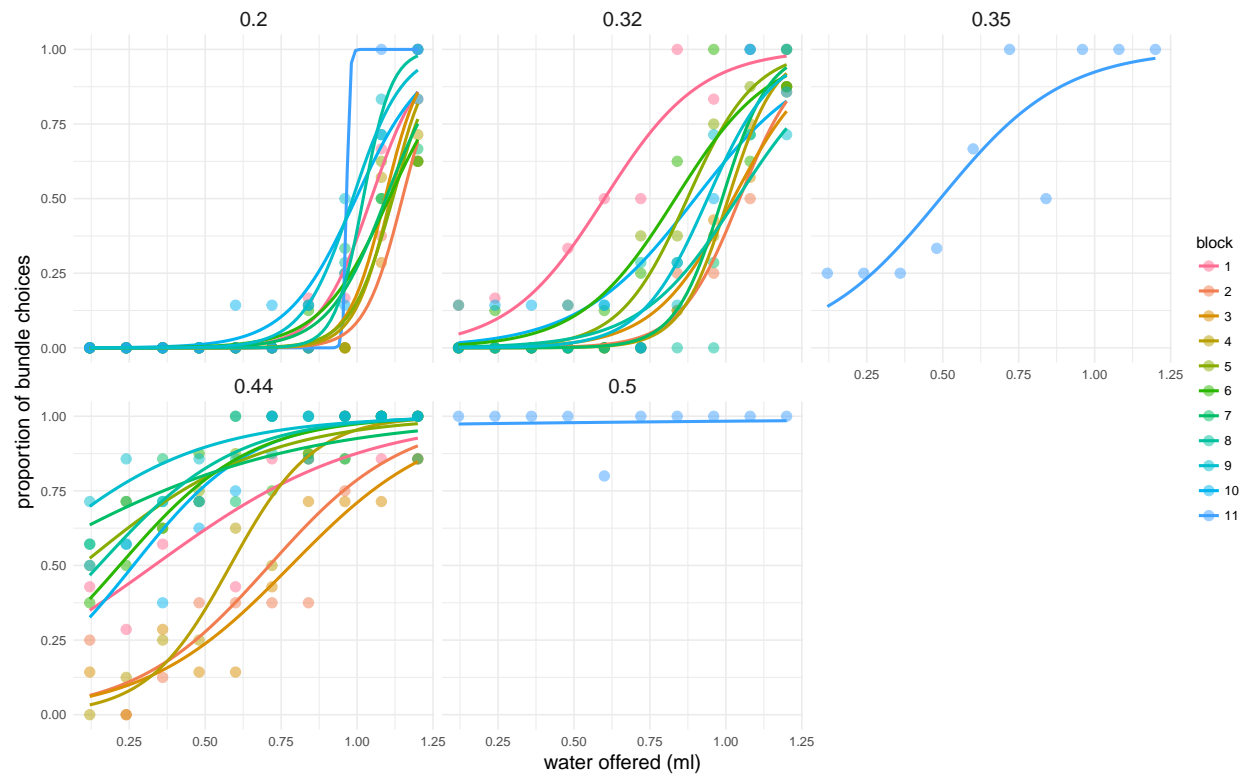
Ulysses : 1-Mar-2018 – 26-Mar-2018



p5

Monkey Bundle Choice Binoimial Curves

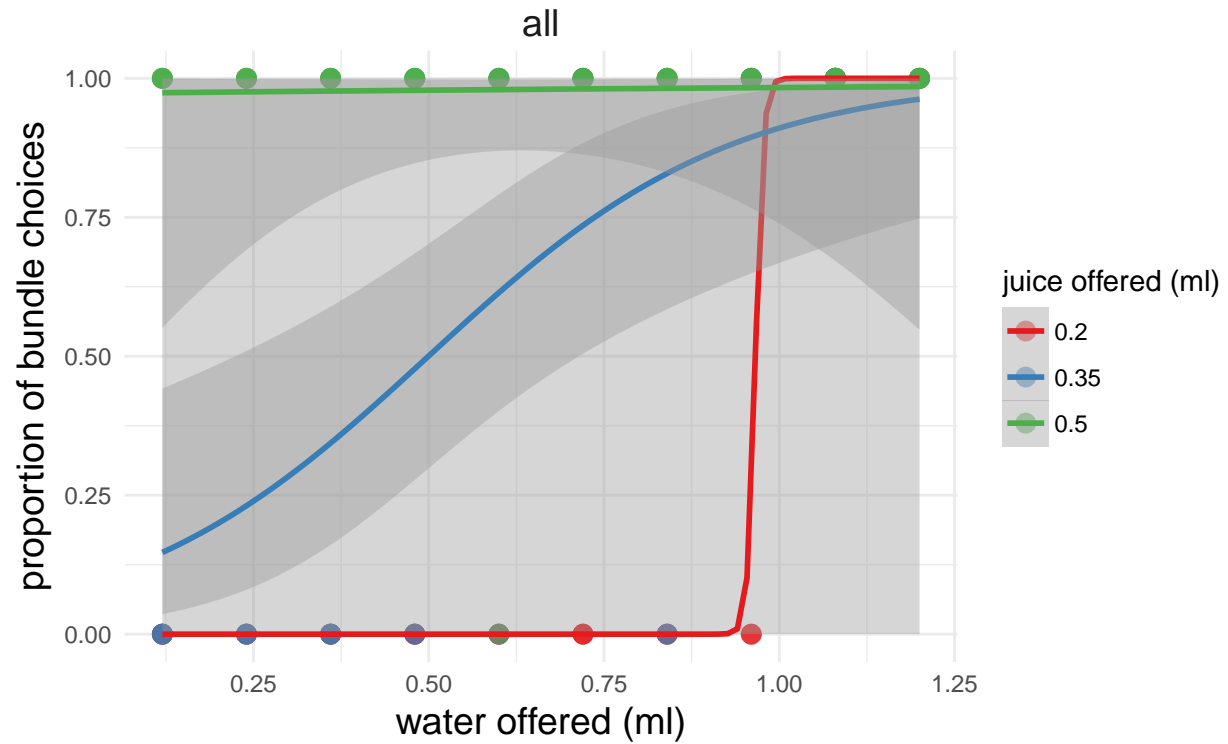
Ulysses : 1-Mar-2018 – 26-Mar-2018



p6

Today's Monkey Bundle Choice Binoimial Curves

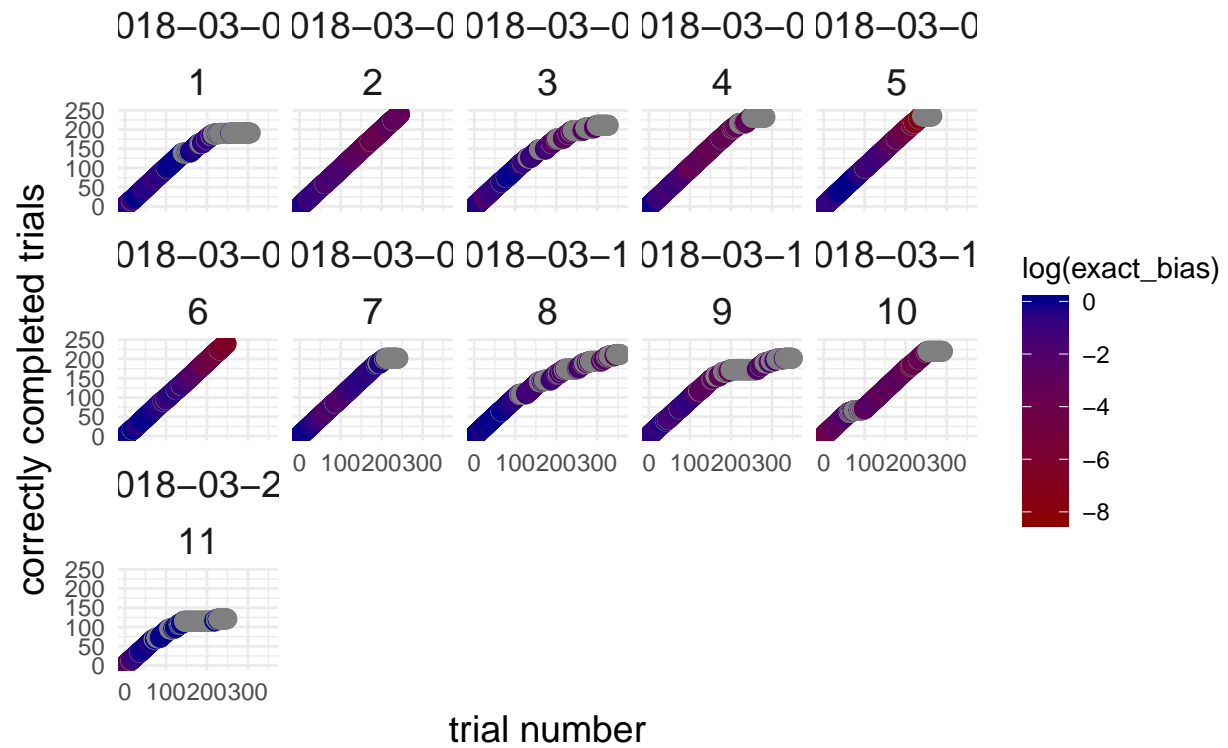
Ulysses : 26-Mar-2018



p7

Monkey Trial Progression and Bias

Ulysses : 1-Mar-2018 – 26-Mar-2018



p8

Monkey Trial Progression and Bias

Ulysses : 1–Mar–2018 – 26–Mar–2018

