

BCb Analysis- Early March

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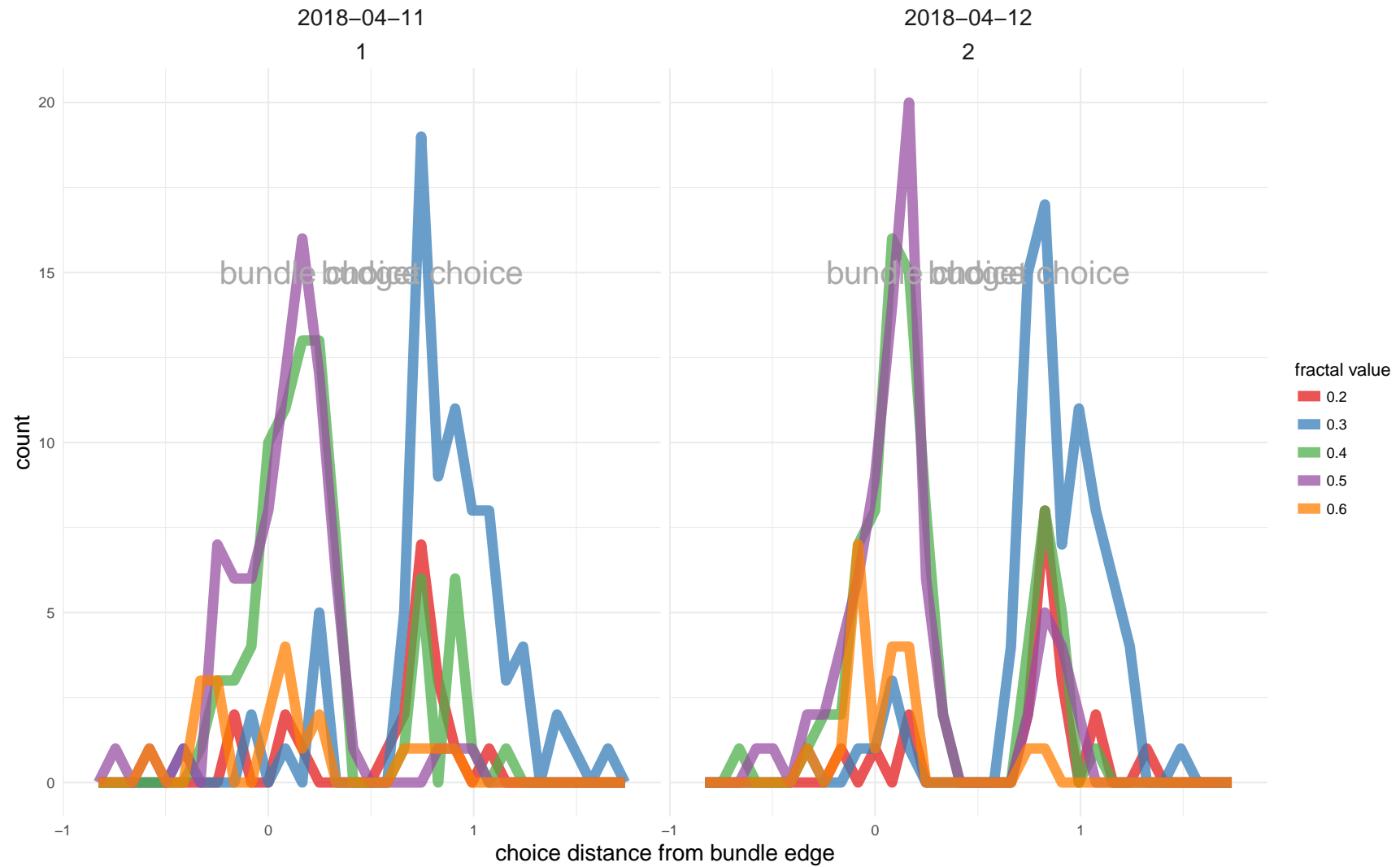
05 April 2018

```
monkey <- "Ulysses"  
today <- "12-Apr-2018"  
look_back <- "11-Apr-2018"  
  
start_trial <- 0  
stop_trial <- "all"  
  
merge_days <- TRUE
```

p1

Monkey Choice Distance From Bundle on Binary Choice Task

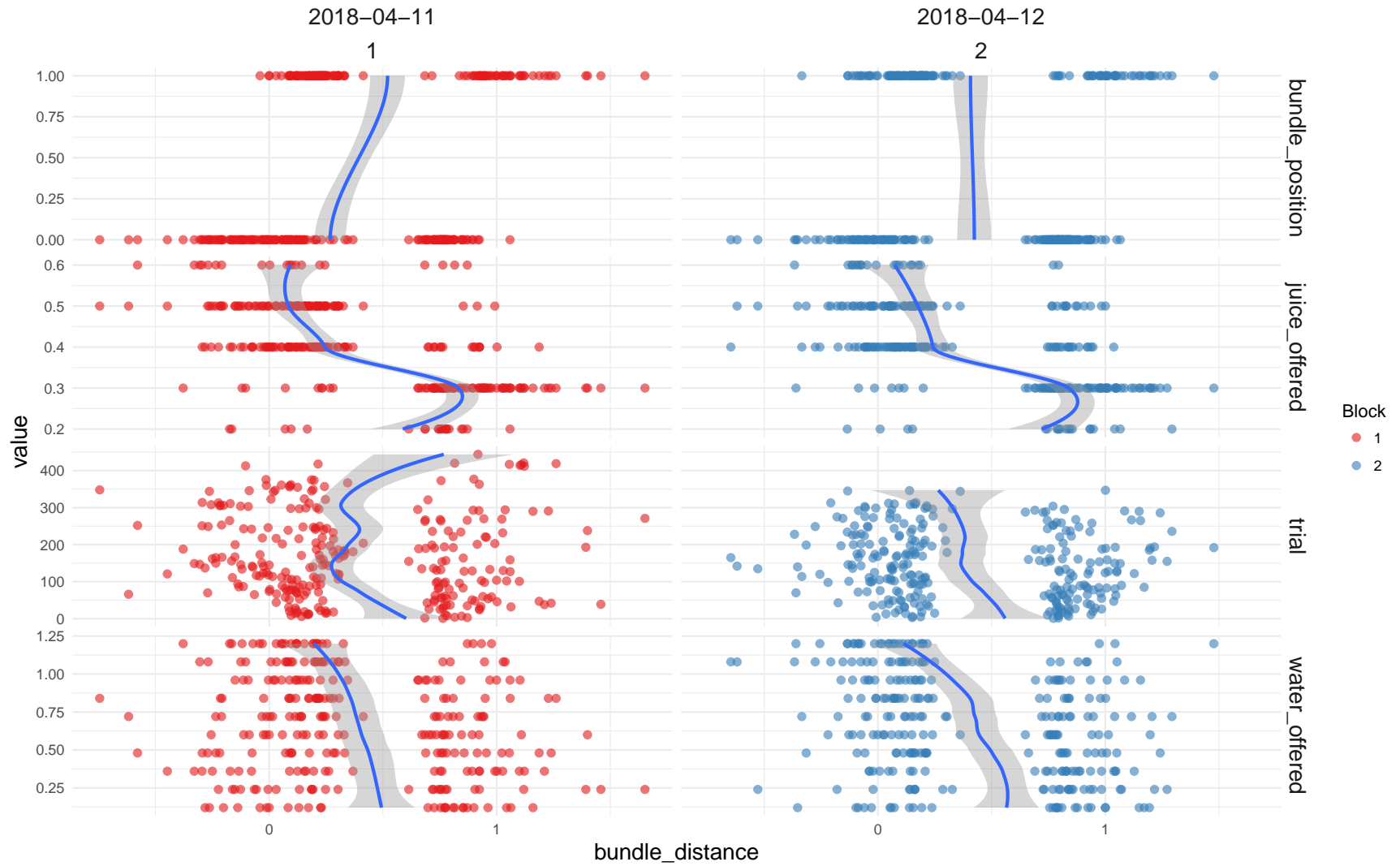
Ulysses : 11-Apr-2018 – 12-Apr-2018



p2

Monkey Choice Distance From Bundle on Binary Choice Task

Ulysses : 11-Apr-2018 – 12-Apr-2018



```

#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
## -3.3328  -0.5078   0.1694   0.5637   2.5535
##
## Coefficients:
##              Estimate Std. Error z value Pr(>|z|)
## (Intercept)    5.524e+03  4.362e+03   1.266 0.205336
## bundle_position  4.719e-01  2.475e-01   1.907 0.056532 .
## water_offered    3.293e+00  4.185e-01   7.870 3.55e-15 ***
## juice_offered    2.105e+01  1.808e+00  11.643 < 2e-16 ***
## trial           4.304e-03  1.210e-03   3.558 0.000374 ***
## date            -3.139e-01  2.474e-01  -1.269 0.204456
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for binomial family taken to be 1)
##
##      Null deviance: 758.37  on 559  degrees of freedom
## Residual deviance: 417.74  on 554  degrees of freedom
## (231 observations deleted due to missingness)
## AIC: 429.74
##
## 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(bundle_position == fractal_choice)]))
## number of successes = 250, number of trials = 560, p-value =
## 0.01259
## alternative hypothesis: true probability of success is not equal to 0.5
## 95 percent confidence interval:
##  0.4047392 0.4886859
## sample estimates:
## probability of success
##      0.4464286

```

```

#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
## -2.45713  -0.18833   0.06506   0.27717   2.31811
##
## Coefficients: (1 not defined because of singularities)
##              Estimate Std. Error z value Pr(>|z|)
## (Intercept)    -8.826124    1.429421  -6.175 6.63e-10 ***
## bundle_position    2.264338    0.534549   4.236 2.28e-05 ***
## water_offered     6.540830    1.092013   5.990 2.10e-09 ***
## as.factor(juice_offered)0.3 -2.208347    0.875427  -2.523 0.01165 *
## as.factor(juice_offered)0.4  4.671066    0.960376   4.864 1.15e-06 ***
## as.factor(juice_offered)0.5  5.426584    1.007813   5.385 7.26e-08 ***
## as.factor(juice_offered)0.6  5.386787    1.321249   4.077 4.56e-05 ***
## trial           0.008131    0.002557   3.180 0.00147 **
## 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: 383.52  on 279  degrees of freedom
## Residual deviance: 135.86  on 272  degrees of freedom
## (67 observations deleted due to missingness)
## AIC: 151.86

```

```
##
```

```
## 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))]), nrow(task_data %>% .[c(bundle_pos
```

```
## number of successes = 107, number of trials = 280, p-value =
```

```
## 9.575e-05
```

```
## alternative hypothesis: true probability of success is not equal to 0.5
```

```
## 95 percent confidence interval:
```

```
## 0.3249597 0.4418490
```

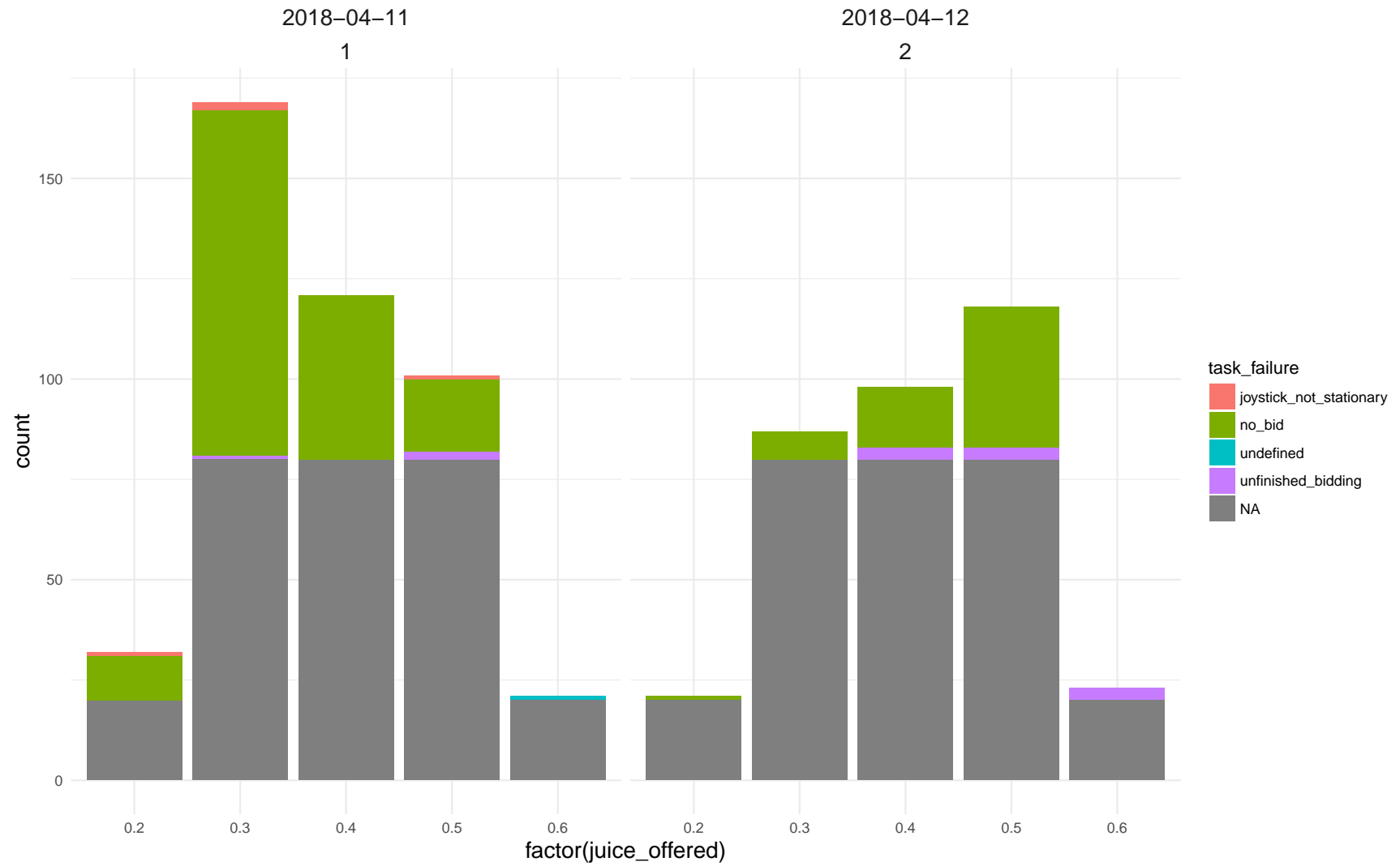
```
## sample estimates:
```

```
## probability of success
```

```
## 0.3821429
```

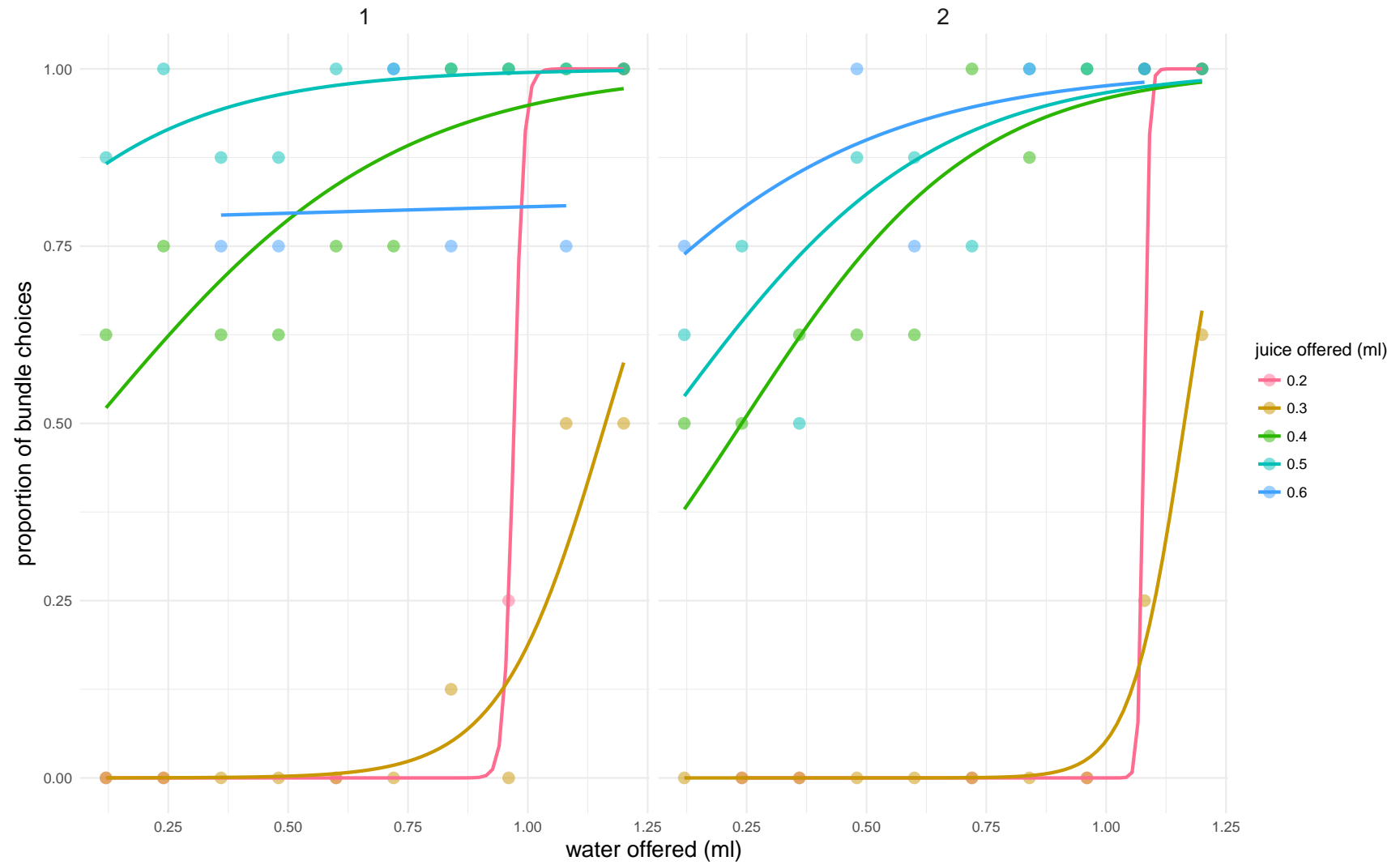
Monkey Choice Failures

Ulysses : 11-Apr-2018 – 12-Apr-2018



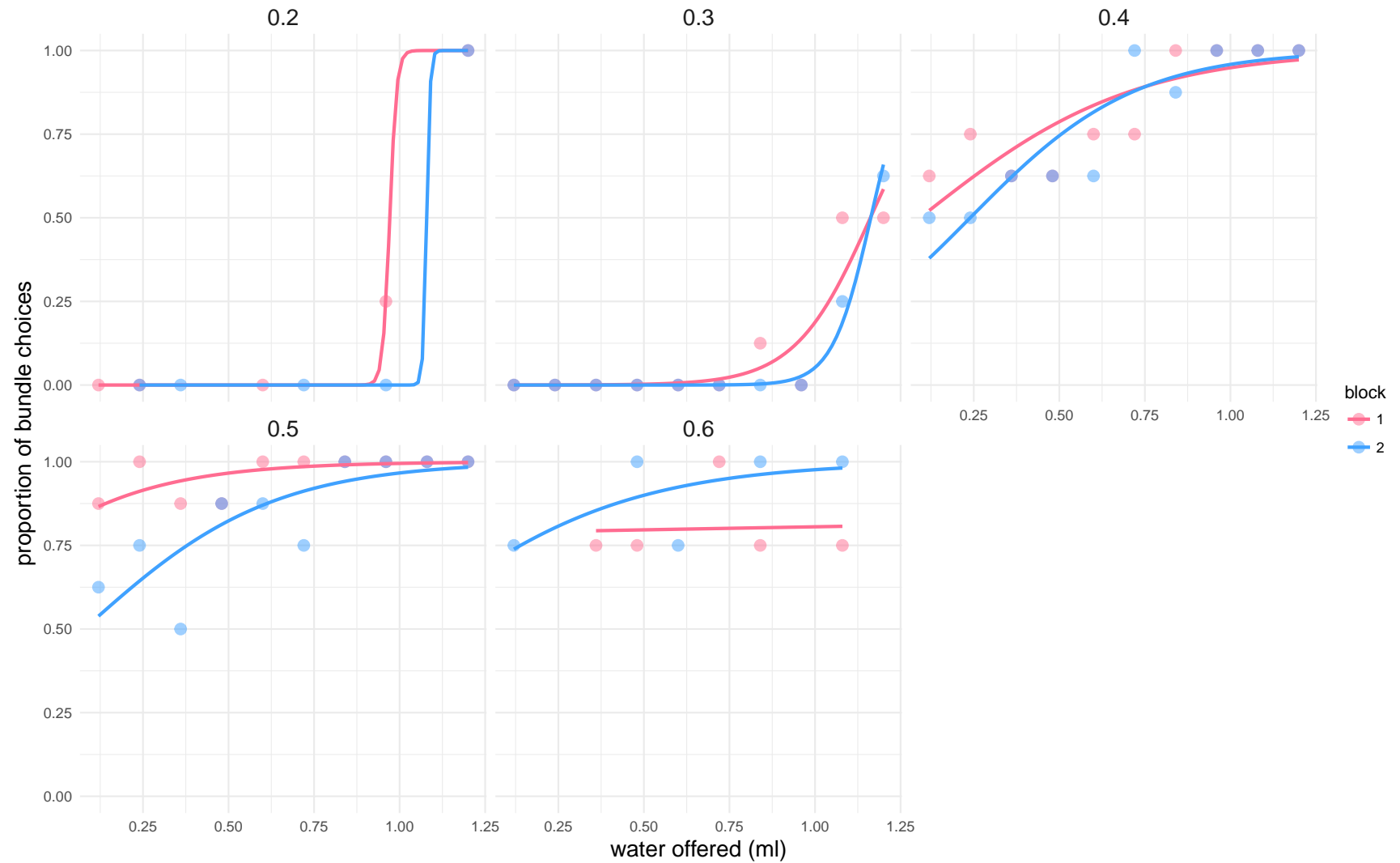
Monkey Bundle Choice Binoimial Curves

Ulysses : 11-Apr-2018 – 12-Apr-2018



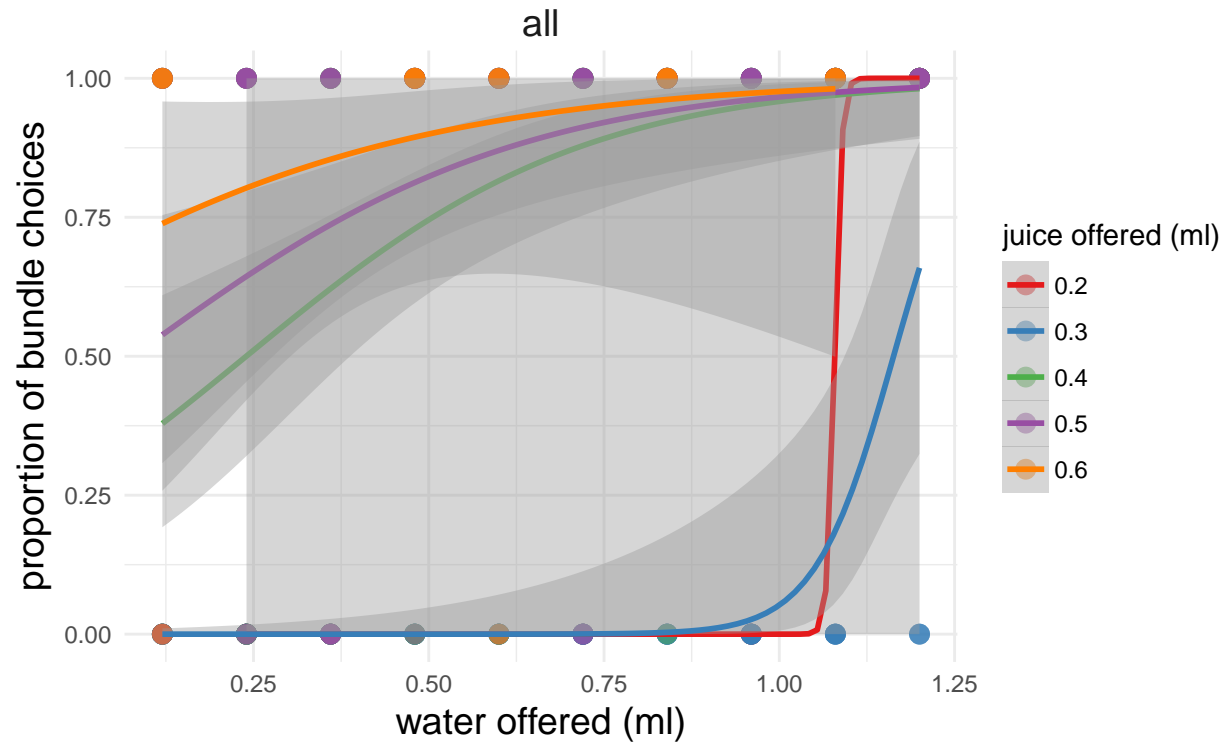
Monkey Bundle Choice Binoimial Curves

Ulysses : 11-Apr-2018 – 12-Apr-2018



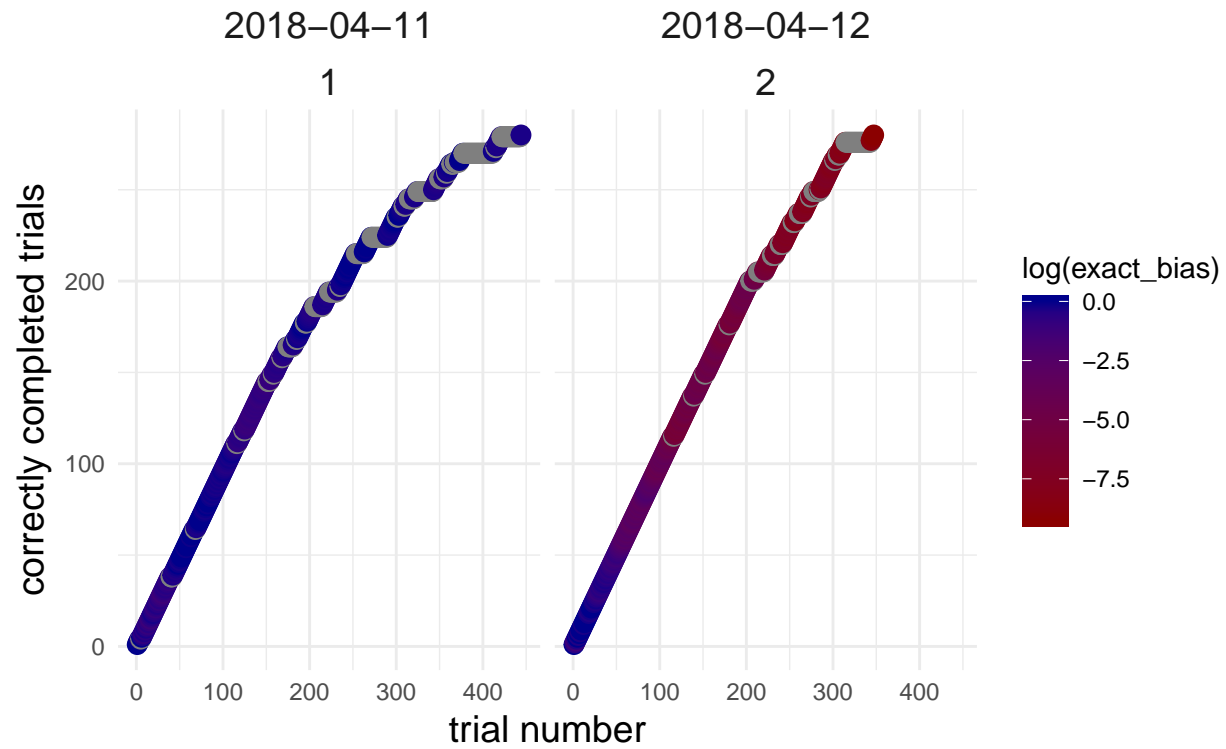
Today's Monkey Bundle Choice Binoimial Curves

Ulysses : 12-Apr-2018



Monkey Trial Progression and Bias

Ulysses : 11-Apr-2018 – 12-Apr-2018



Monkey Trial Progression and Bias

Ulysses : 11-Apr-2018 – 12-Apr-2018

