

BCb Analysis- Early March

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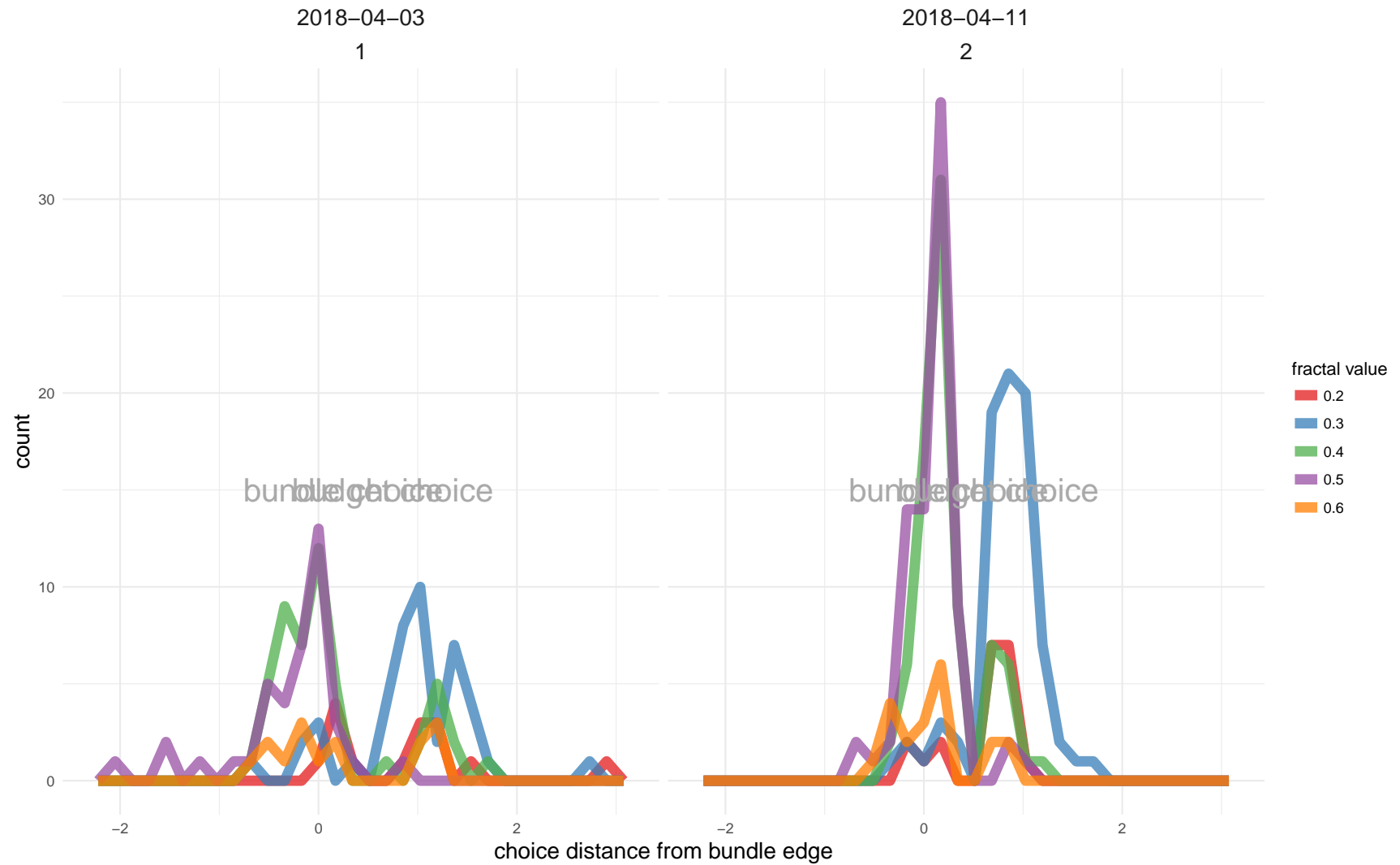
05 April 2018

```
monkey <- "Ulysses"  
today <- "11-Apr-2018"  
look_back <- "3-Apr-2018"  
  
start_trial <- 0  
stop_trial <- "all"  
  
merge_days <- TRUE
```

p1

Monkey Choice Distance From Bundle on Binary Choice Task

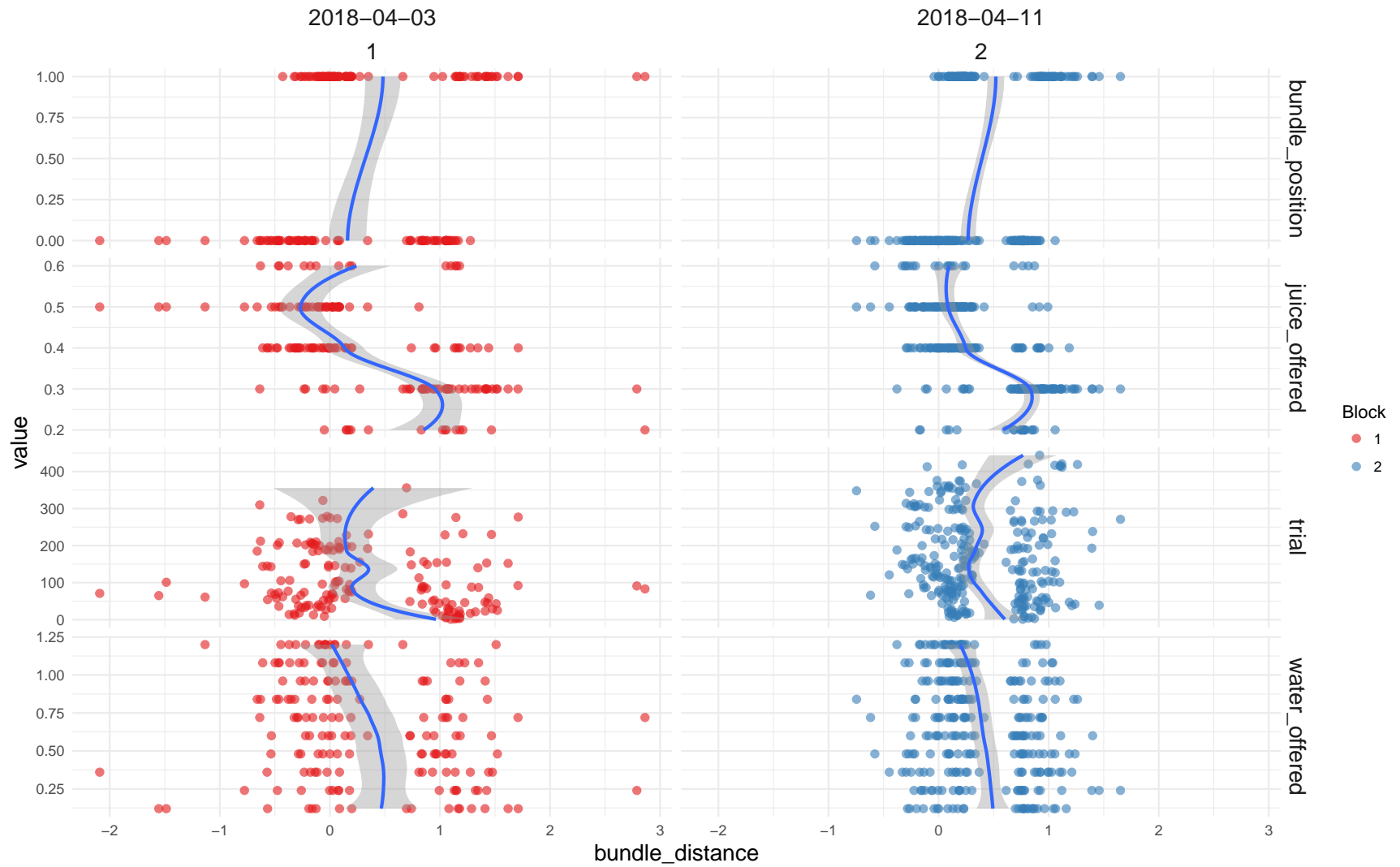
Ulysses : 3-Apr-2018 – 11-Apr-2018



p2

Monkey Choice Distance From Bundle on Binary Choice Task

Ulysses : 3-Apr-2018 – 11-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
## -3.2286  -0.5768   0.2469   0.6175   2.4911
##
## Coefficients:
##              Estimate Std. Error z value Pr(>|z|)
## (Intercept)    8.064e+02  6.242e+02   1.292  0.19638
## bundle_position -7.411e-04  2.644e-01  -0.003  0.99776
## water_offered   2.911e+00  4.365e-01   6.668 2.59e-11 ***
## juice_offered   1.816e+01  1.770e+00  10.263 < 2e-16 ***
## trial           5.005e-03  1.294e-03   3.867  0.00011 ***
## date           -4.626e-02  3.541e-02  -1.306  0.19148
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for binomial family taken to be 1)
##
##      Null deviance: 591.87  on 443  degrees of freedom
## Residual deviance: 358.91  on 438  degrees of freedom
## (377 observations deleted due to missingness)
## AIC: 370.91
##
## Number of Fisher Scoring iterations: 5

```

```

#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 = 217, number of trials = 444, p-value =
## 0.6693
## alternative hypothesis: true probability of success is not equal to 0.5
## 95 percent confidence interval:
##  0.4413258 0.5363026
## sample estimates:
## probability of success
##      0.4887387

```

```

#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
## -2.6594  -0.2399   0.1137   0.3700   2.0269
##
## Coefficients: (1 not defined because of singularities)
##              Estimate Std. Error z value Pr(>|z|)
## (Intercept)    -5.426890    1.094773  -4.957 7.16e-07 ***
## bundle_position -0.333946    0.460184  -0.726  0.46804
## water_offered    4.562174    0.866711   5.264 1.41e-07 ***
## as.factor(juice_offered)0.3 -1.359720    0.815218  -1.668  0.09533 .
## as.factor(juice_offered)0.4  3.863619    0.898663   4.299 1.71e-05 ***
## as.factor(juice_offered)0.5  6.019091    1.071595   5.617 1.94e-08 ***
## as.factor(juice_offered)0.6  3.474090    0.956214   3.633  0.00028 ***
## trial           0.005266    0.001955   2.693  0.00708 **
## 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: 373.40  on 279  degrees of freedom
## Residual deviance: 150.08  on 272  degrees of freedom
##      (164 observations deleted due to missingness)
## AIC: 166.08

```

```
##
```

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

```
## number of successes = 143, number of trials = 280, p-value =
```

```
## 0.7651
```

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

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

```
## 0.4505330 0.5706664
```

```
## sample estimates:
```

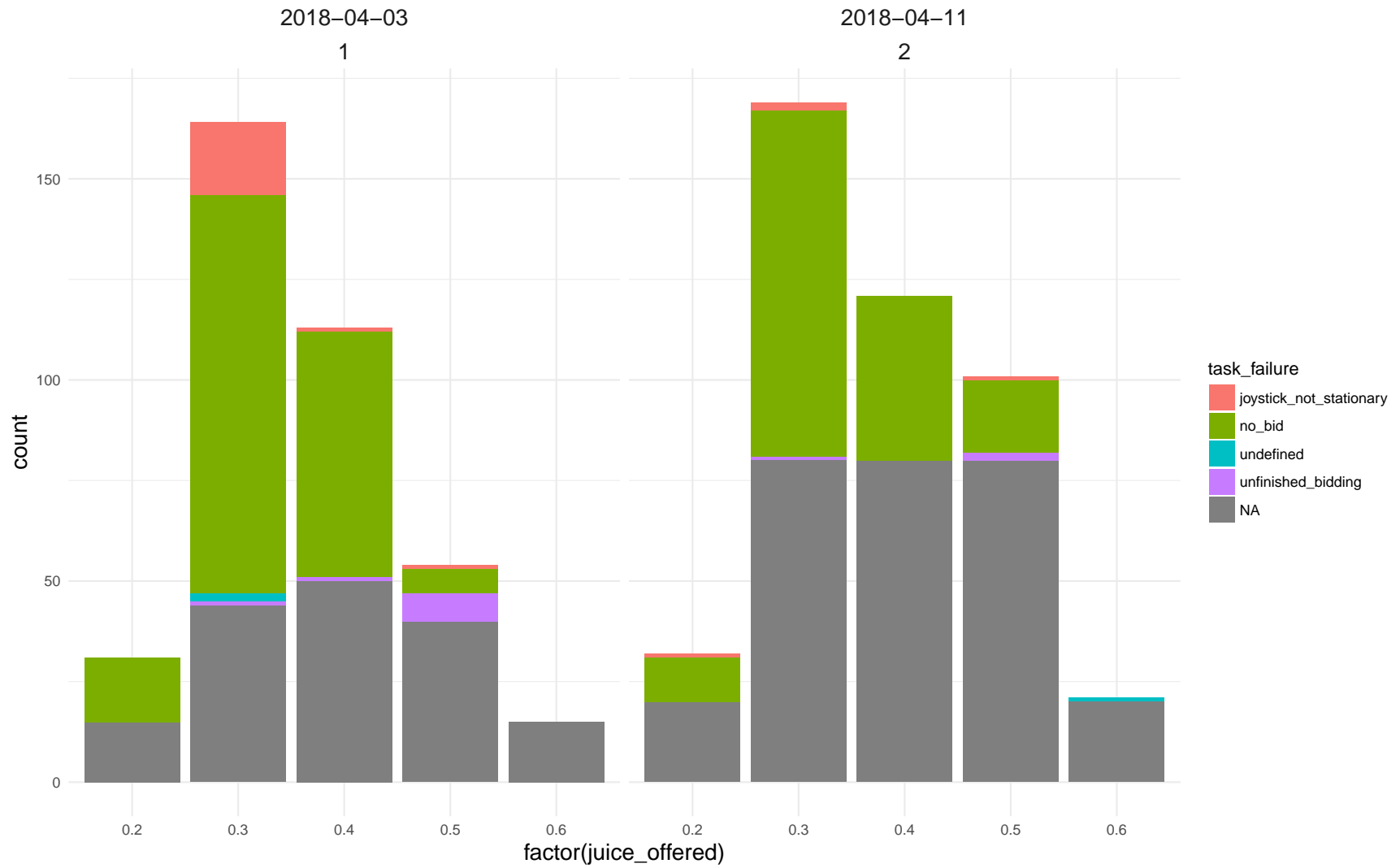
```
## probability of success
```

```
## 0.5107143
```

p3

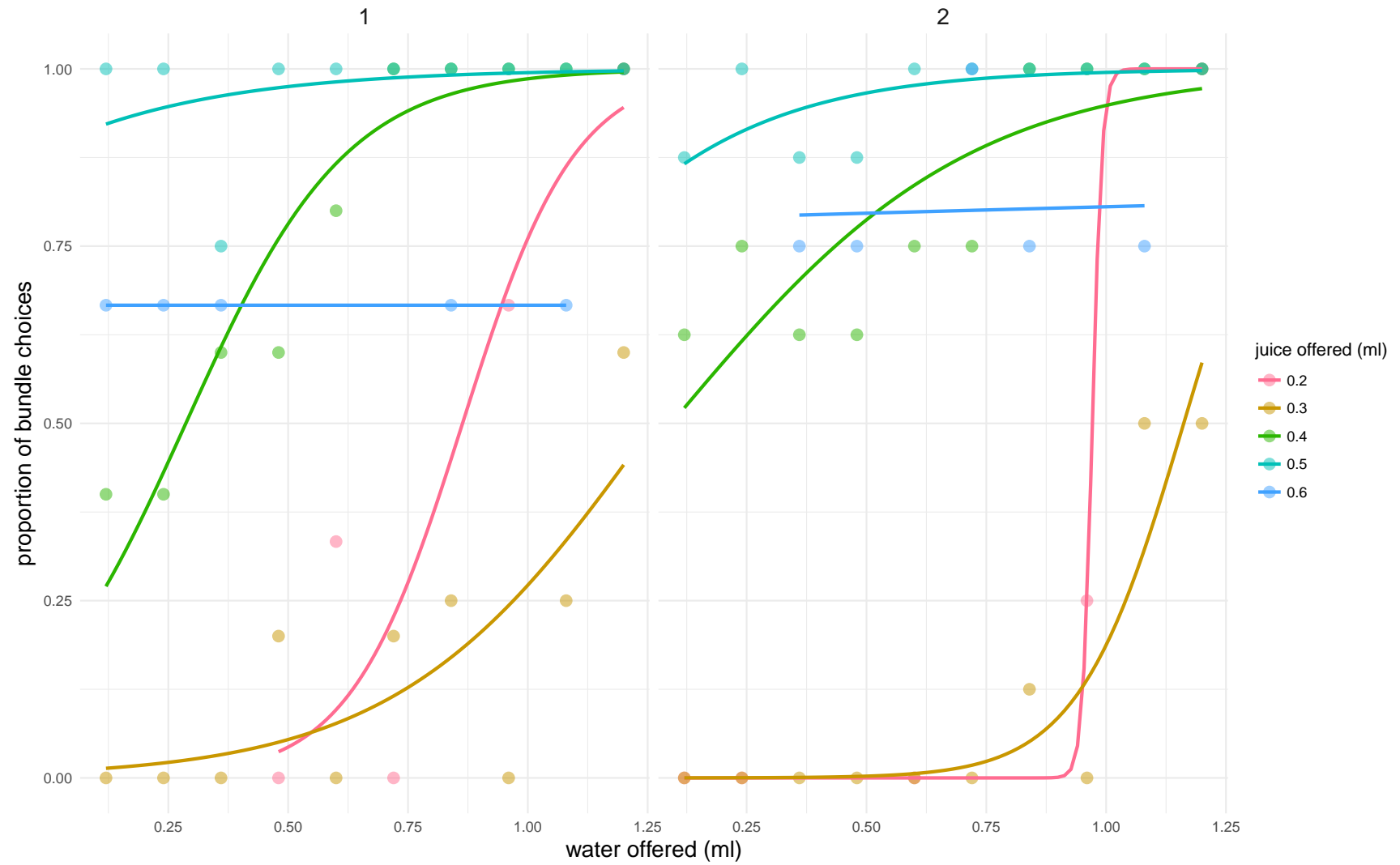
Monkey Choice Failures

Ulysses : 3-Apr-2018 – 11-Apr-2018



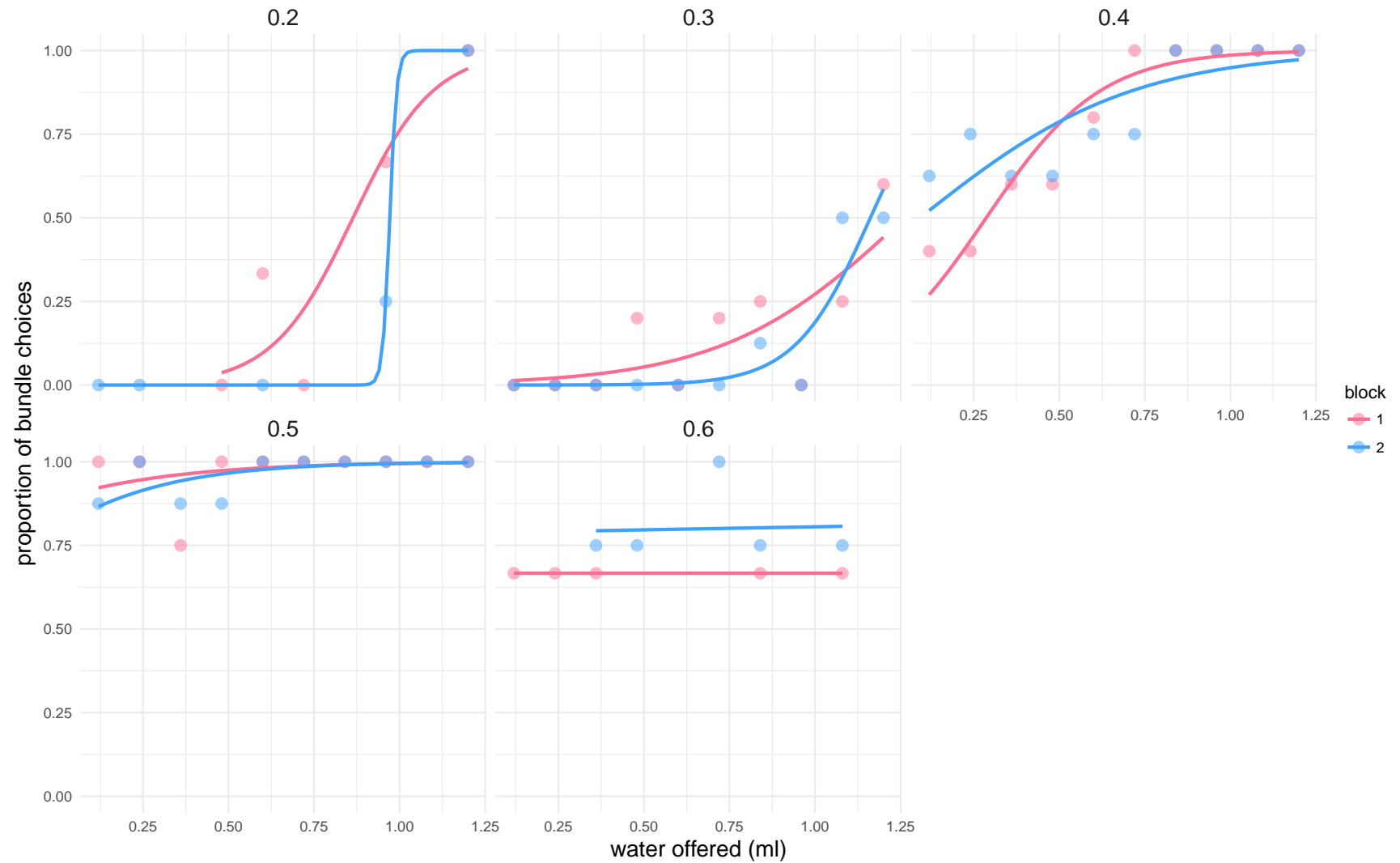
Monkey Bundle Choice Binoimial Curves

Ulysses : 3-Apr-2018 – 11-Apr-2018



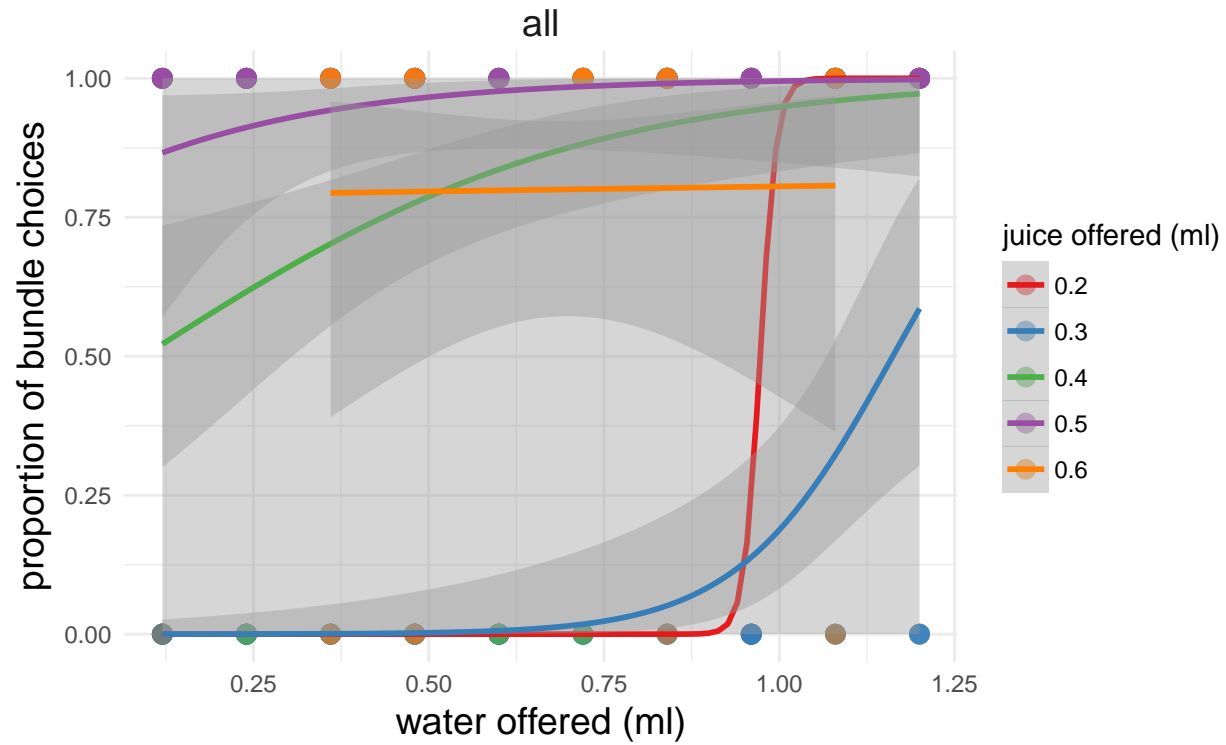
Monkey Bundle Choice Binoimial Curves

Ulysses : 3-Apr-2018 – 11-Apr-2018



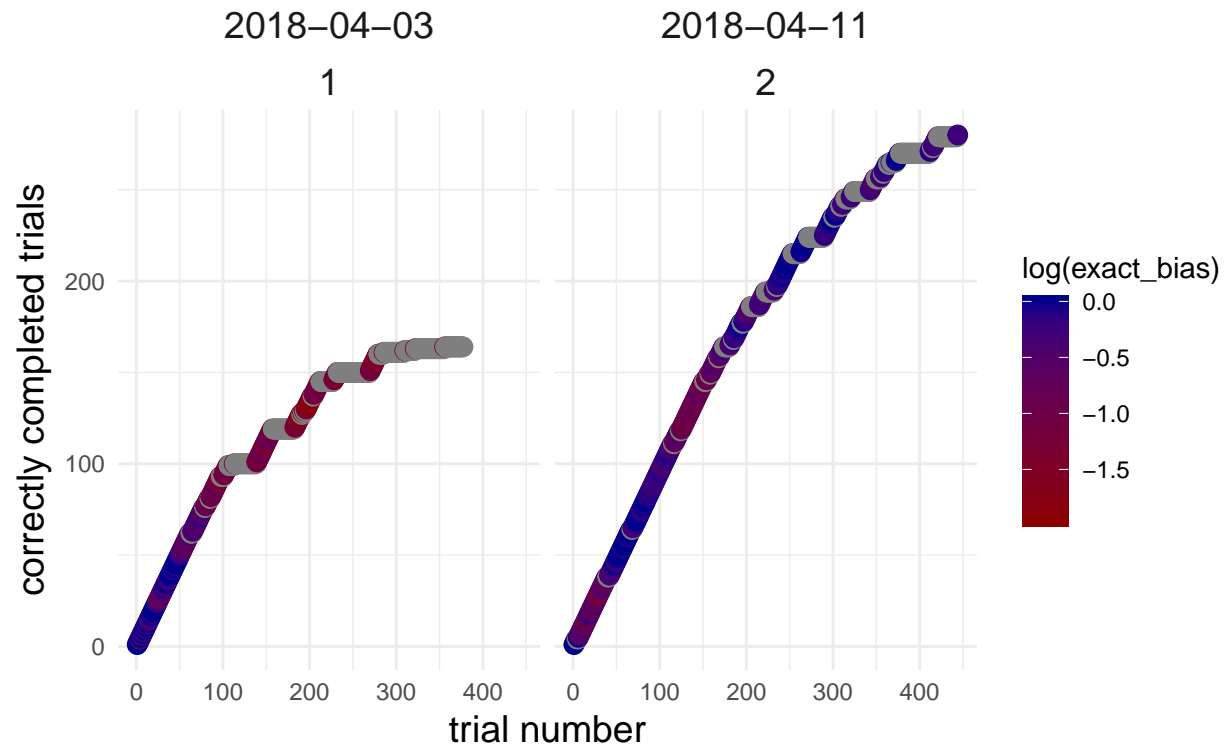
Today's Monkey Bundle Choice Binoimial Curves

Ulysses : 11-Apr-2018



Monkey Trial Progression and Bias

Ulysses : 3-Apr-2018 – 11-Apr-2018



Monkey Trial Progression and Bias

Ulysses : 3-Apr-2018 – 11-Apr-2018

