

BCb Analysis- Early March

Robert Hickman

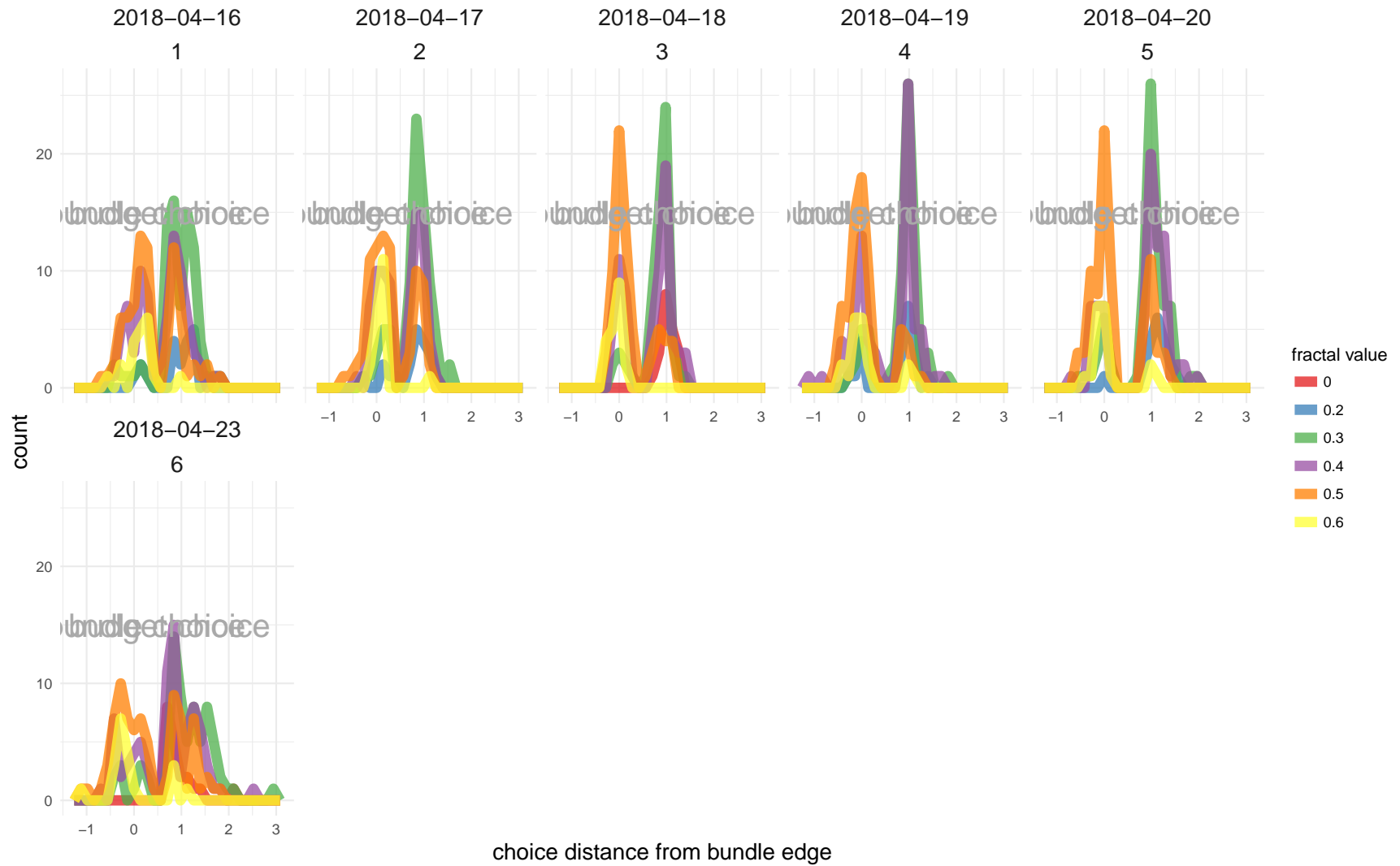
05 April 2018

```
monkey <- "Ulysses"  
today <- "23-Apr-2018"  
look_back <- "16-Apr-2018"  
  
start_trial <- 0  
stop_trial <- 220  
  
merge_days <- TRUE
```

p1

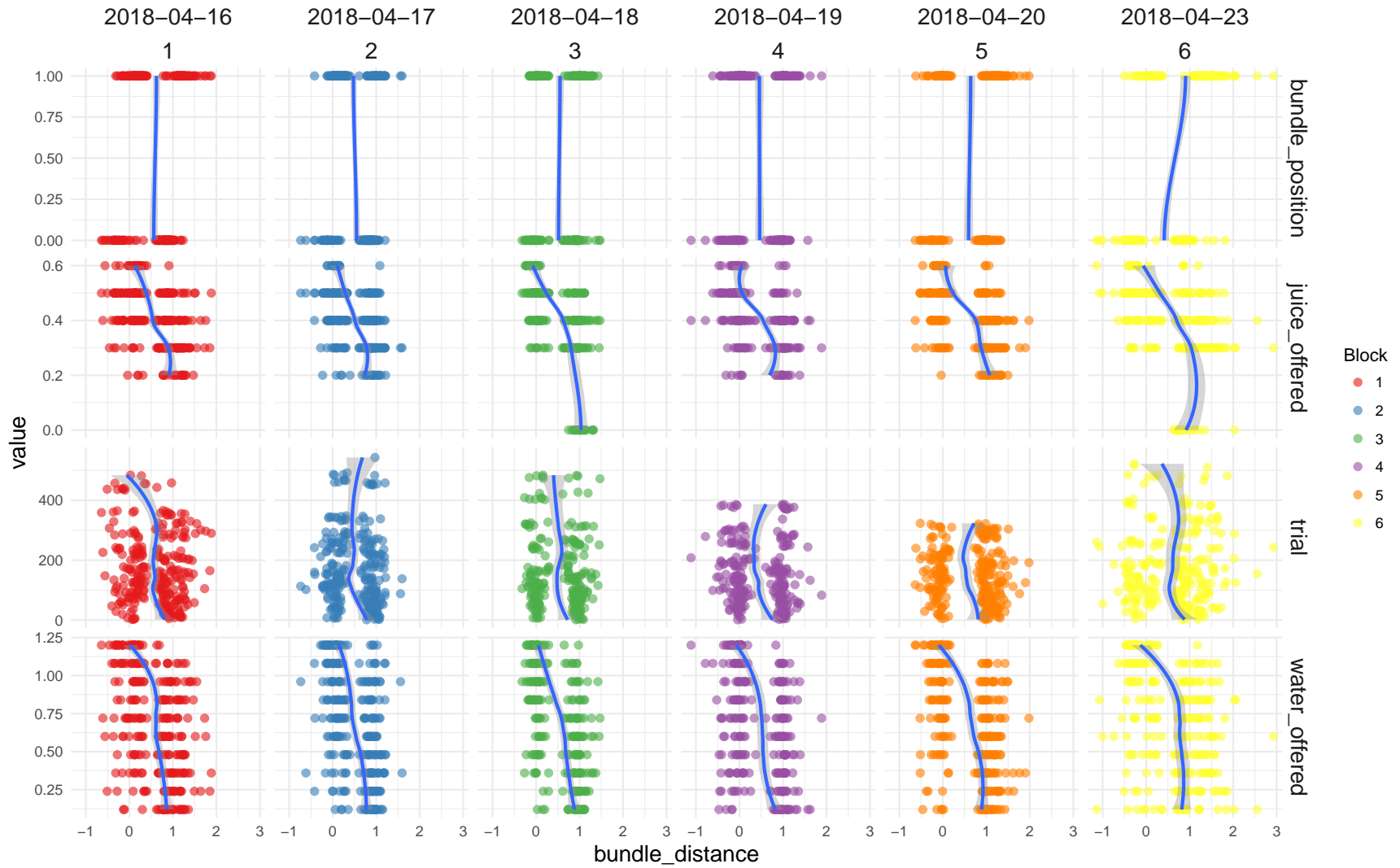
Monkey Choice Distance From Bundle on Binary Choice Task

Ulysses : 16-Apr-2018 – 23-Apr-2018



Monkey Choice Distance From Bundle on Binary Choice Task

Ulysses : 16-Apr-2018 – 23-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
## -2.6968  -0.5679  -0.1393   0.5533   3.2731
##
## Coefficients:
##              Estimate Std. Error z value Pr(>|z|)
## (Intercept)   1.295e+03  5.521e+02   2.346   0.019 *
## bundle_position 1.252e+00  1.482e-01   8.448 < 2e-16 ***
## water_offered  4.525e+00  2.651e-01  17.071 < 2e-16 ***
## juice_offered  1.706e+01  9.613e-01  17.747 < 2e-16 ***
## trial          3.241e-03  6.170e-04   5.252 1.5e-07 ***
## date          -7.406e-02  3.130e-02  -2.366   0.018 *
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for binomial family taken to be 1)
##
##      Null deviance: 2210.2  on 1614  degrees of freedom
## Residual deviance: 1232.4  on 1609  degrees of freedom
## (1213 observations deleted due to missingness)
## AIC: 1244.4
##
## 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(bundle_position == fractal_choice)]))
## number of successes = 669, number of trials = 1615, p-value =
## 5.802e-12
## alternative hypothesis: true probability of success is not equal to 0.5
## 95 percent confidence interval:
##  0.3900822 0.4387131
## sample estimates:
## probability of success
##          0.4142415

```

```

#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
## -2.5107  -0.6802  -0.2495   0.6776   2.4688
##
## Coefficients: (1 not defined because of singularities)
##              Estimate Std. Error z value Pr(>|z|)
## (Intercept)    -19.959422  847.236776  -0.024   0.981
## bundle_position     0.207185   0.324801   0.638   0.524
## water_offered     3.445791   0.545740   6.314 2.72e-10 ***
## as.factor(juice_offered)0.3 15.157811  847.236737   0.018   0.986
## as.factor(juice_offered)0.4 16.406327  847.236700   0.019   0.985
## as.factor(juice_offered)0.5 17.973381  847.236699   0.021   0.983
## as.factor(juice_offered)0.6 19.340251  847.236927   0.023   0.982
## trial              0.001171   0.001225   0.956   0.339
## 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: 360.83  on 272  degrees of freedom
## Residual deviance: 234.73  on 265  degrees of freedom
## (277 observations deleted due to missingness)
## AIC: 250.73

```

```
##
```

```
## Number of Fisher Scoring iterations: 16
```

```
#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 = 131, number of trials = 273, p-value =
```

```
## 0.5451
```

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

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

```
## 0.419261 0.540888
```

```
## sample estimates:
```

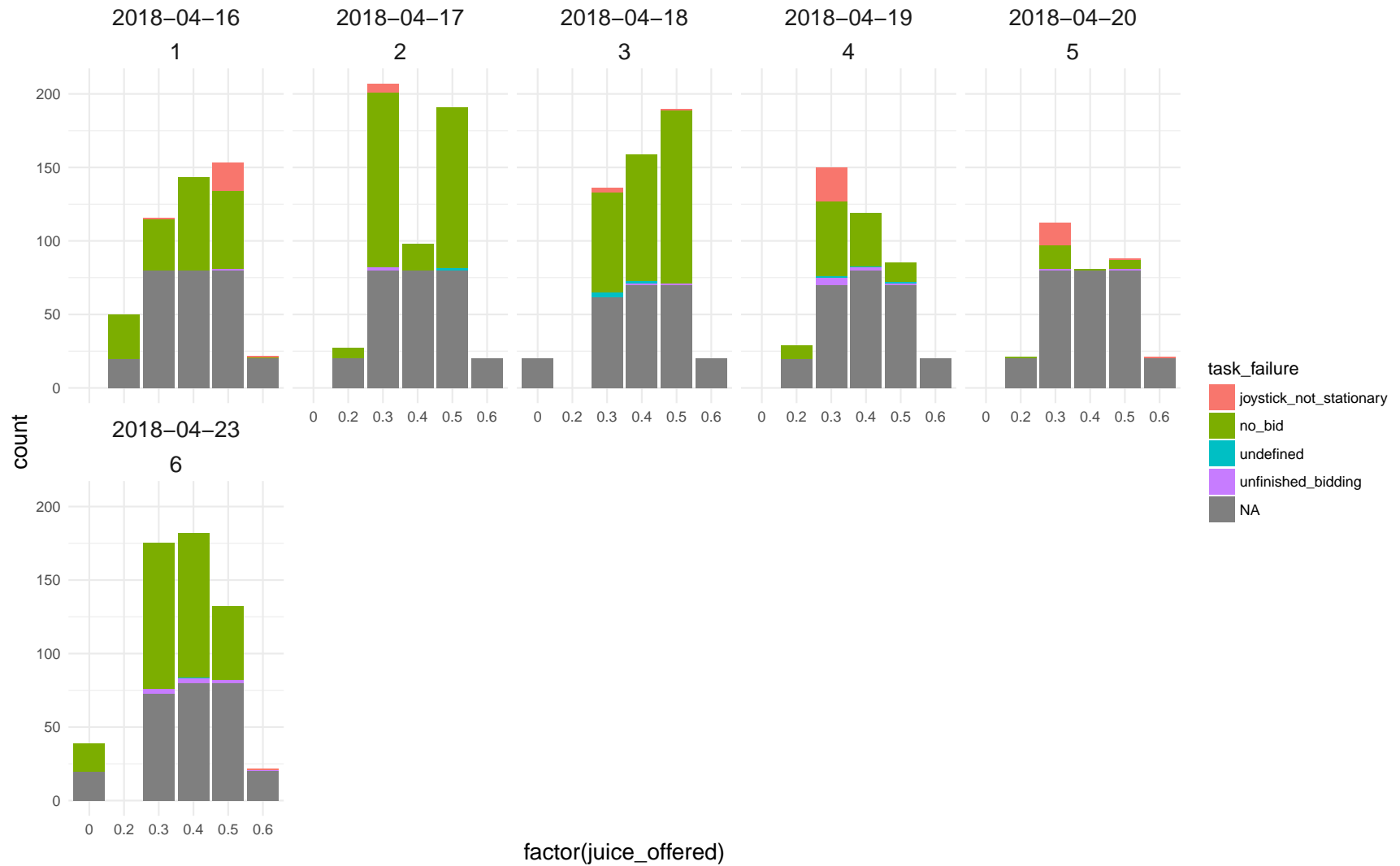
```
## probability of success
```

```
## 0.4798535
```

p3

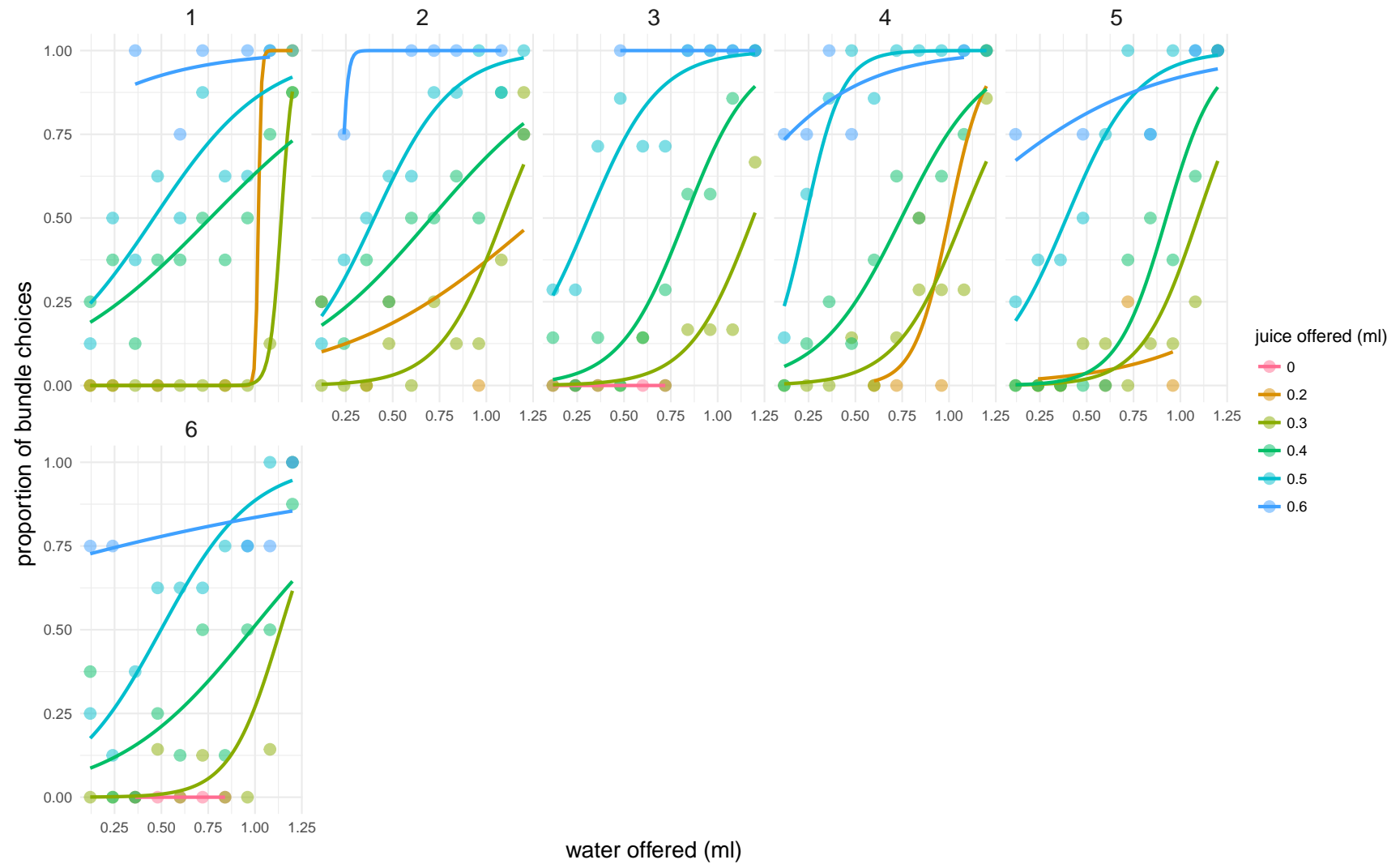
Monkey Choice Failures

Ulysses : 16-Apr-2018 – 23-Apr-2018



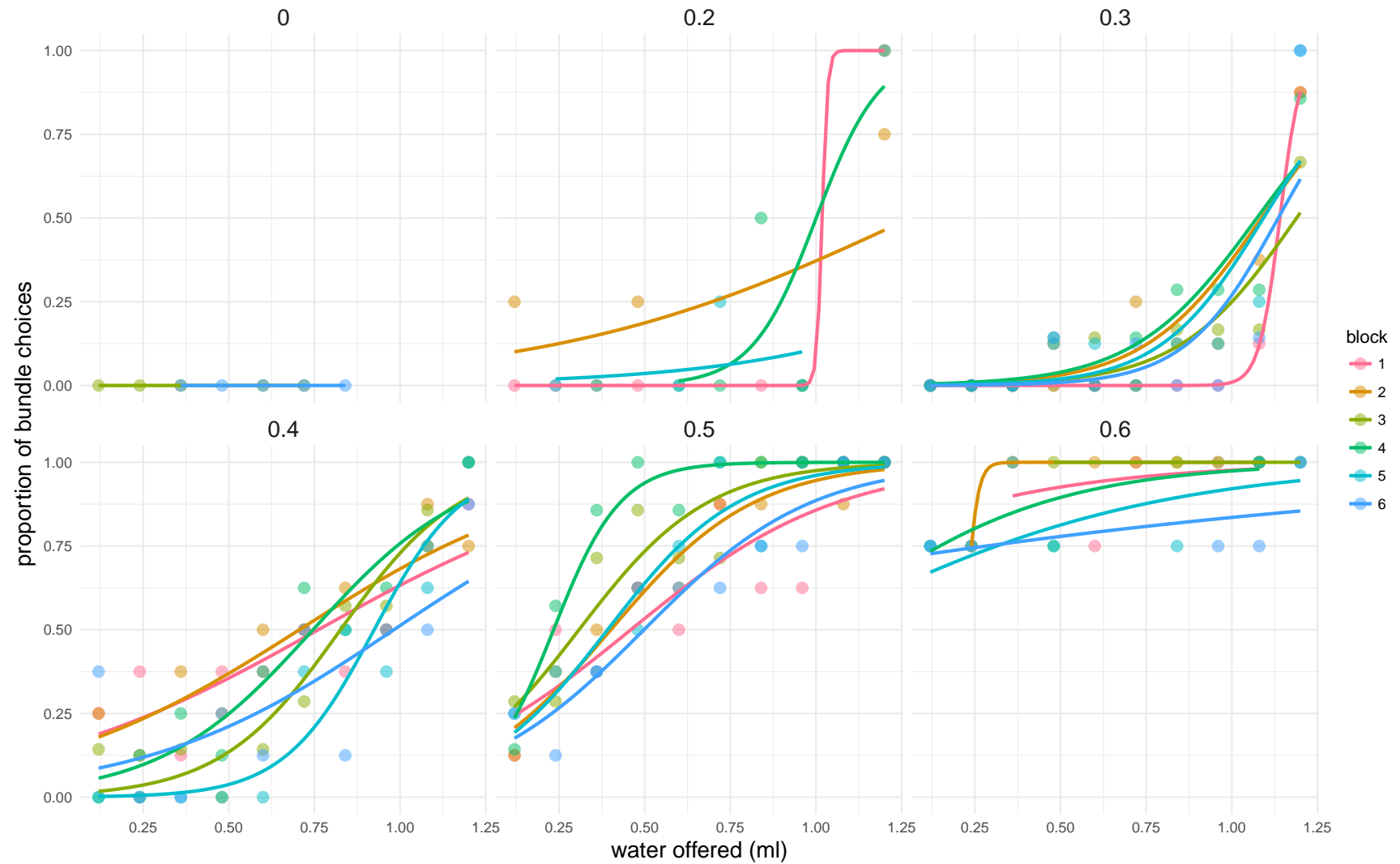
Monkey Bundle Choice Binoimial Curves

Ulysses : 16-Apr-2018 – 23-Apr-2018



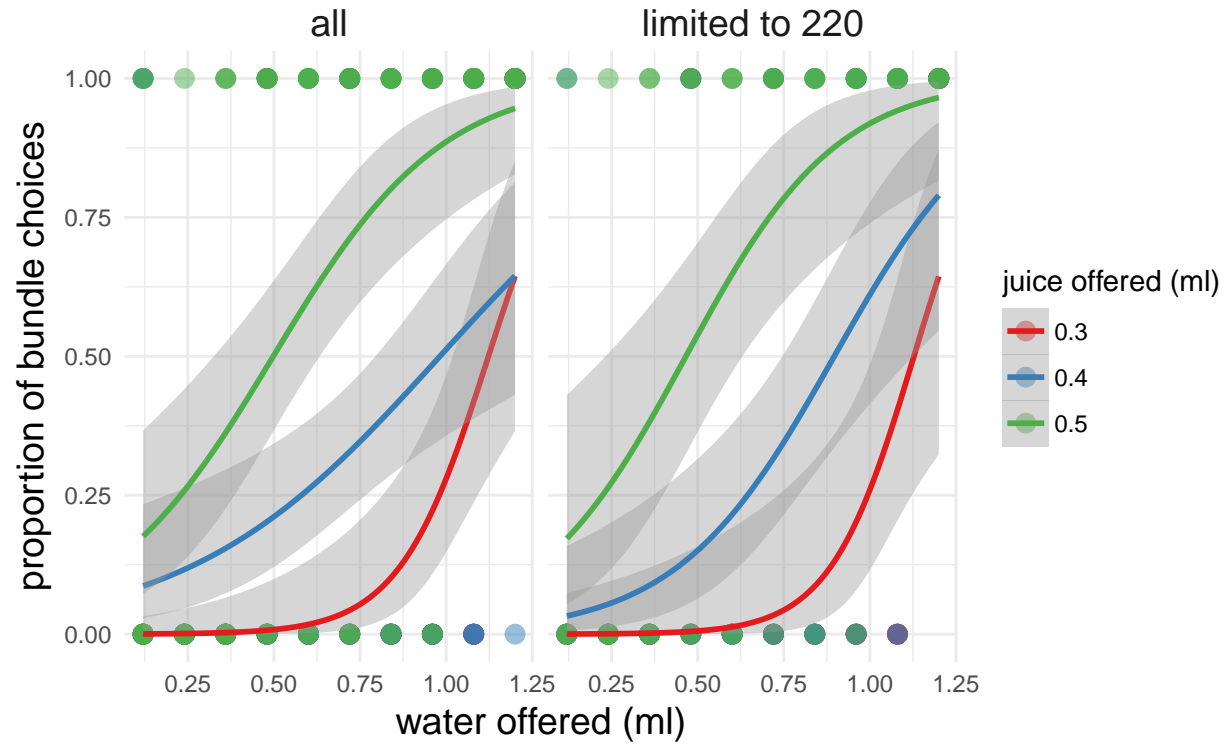
Monkey Bundle Choice Binoimial Curves

Ulysses : 16-Apr-2018 – 23-Apr-2018



Today's Monkey Bundle Choice Binoimial Curves

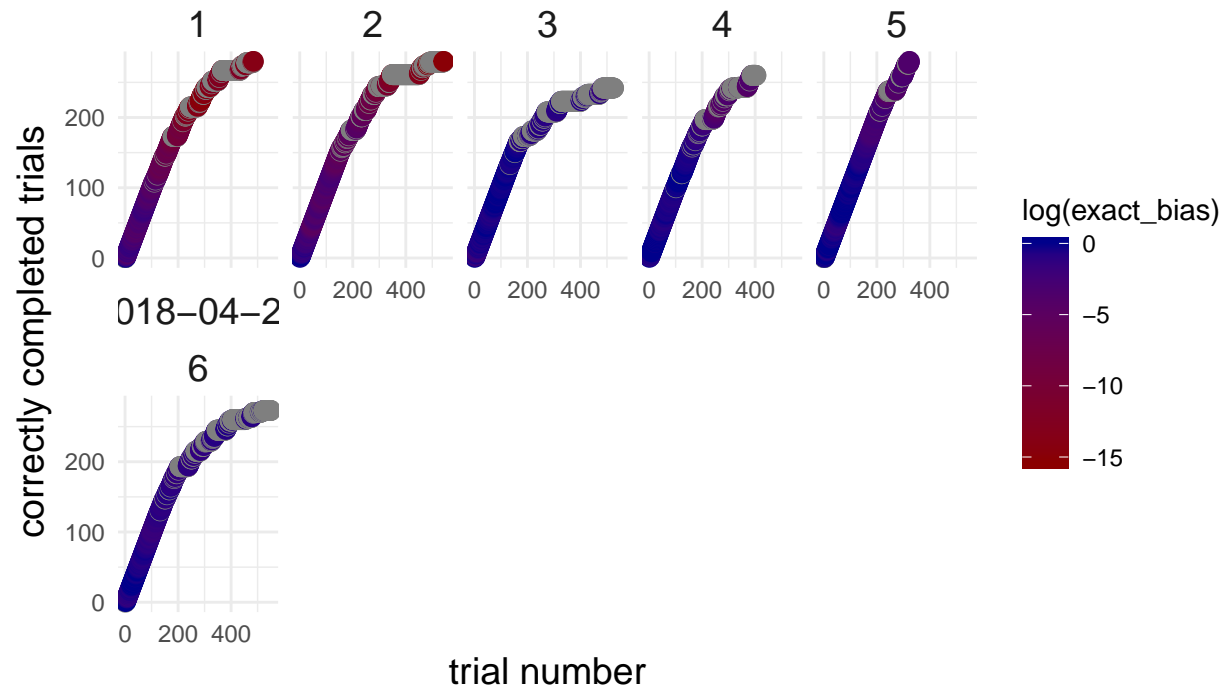
Ulysses : 23-Apr-2018



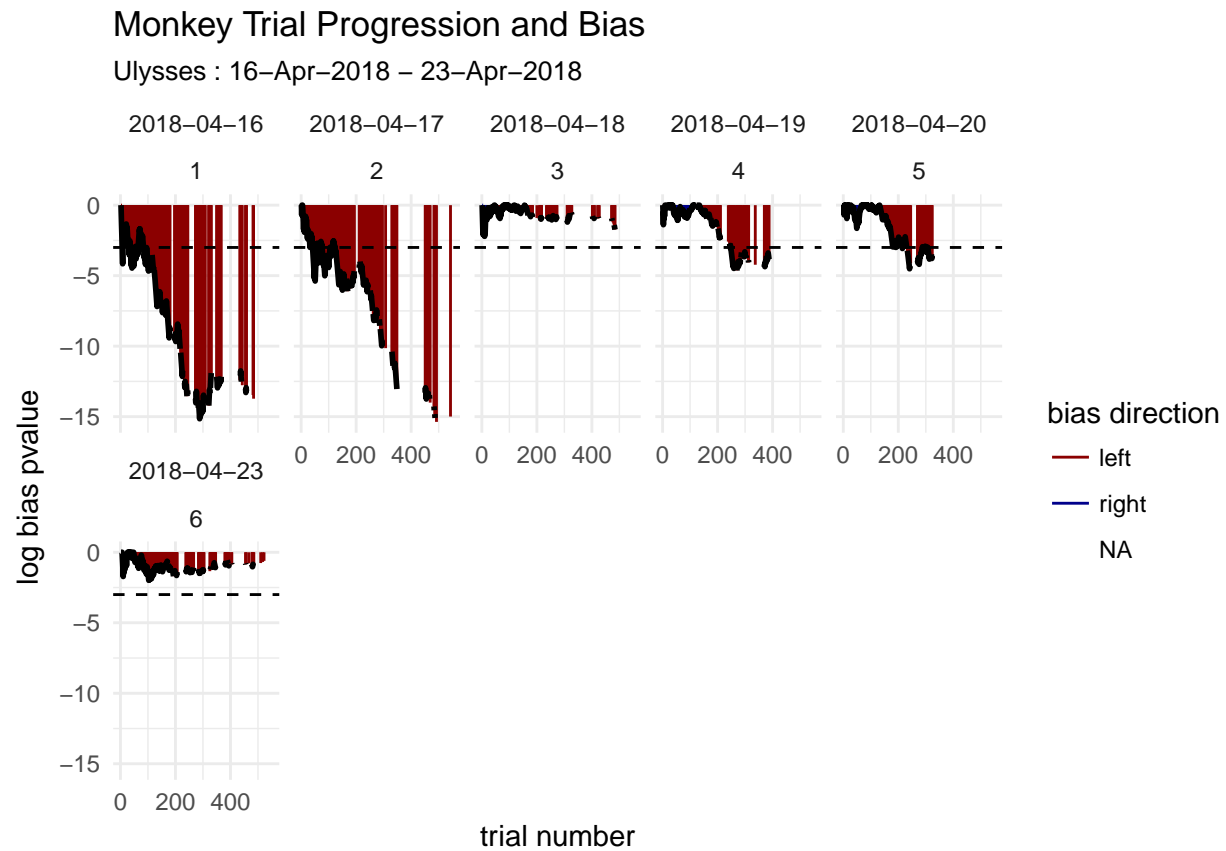
Monkey Trial Progression and Bias

Ulysses : 16-Apr-2018 – 23-Apr-2018

018-04-1 018-04-1 018-04-1 018-04-1 018-04-2



p8



p9

Pooled Monkey Bundle Choice Binoimial Curves

Ulysses : 23-Apr-2018

