

BCb Analysis- Early March

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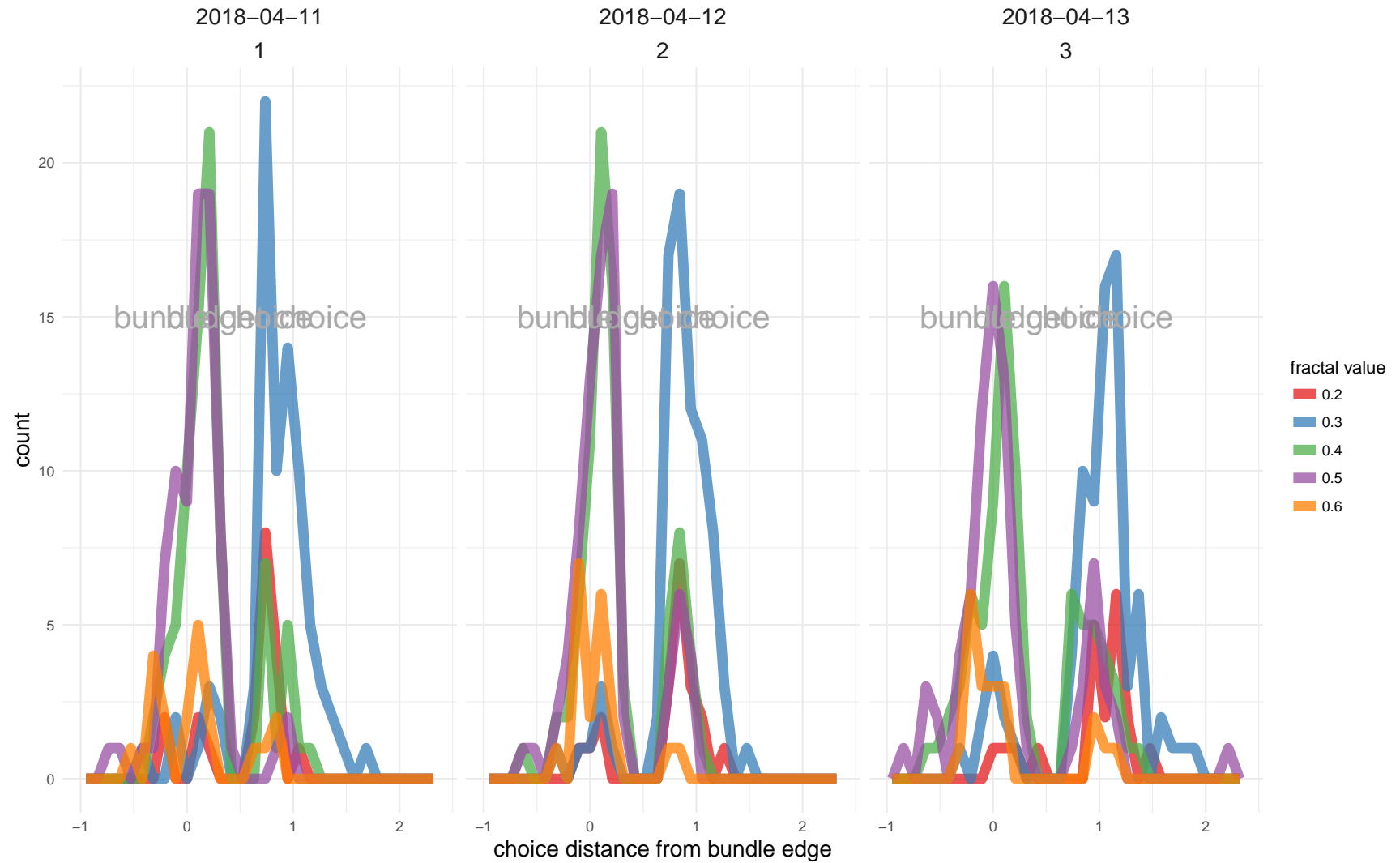
05 April 2018

```
monkey <- "Ulysses"  
today <- "13-Apr-2018"  
look_back <- "11-Apr-2018"  
  
start_trial <- 0  
stop_trial <- "all"  
  
merge_days <- TRUE
```

p1

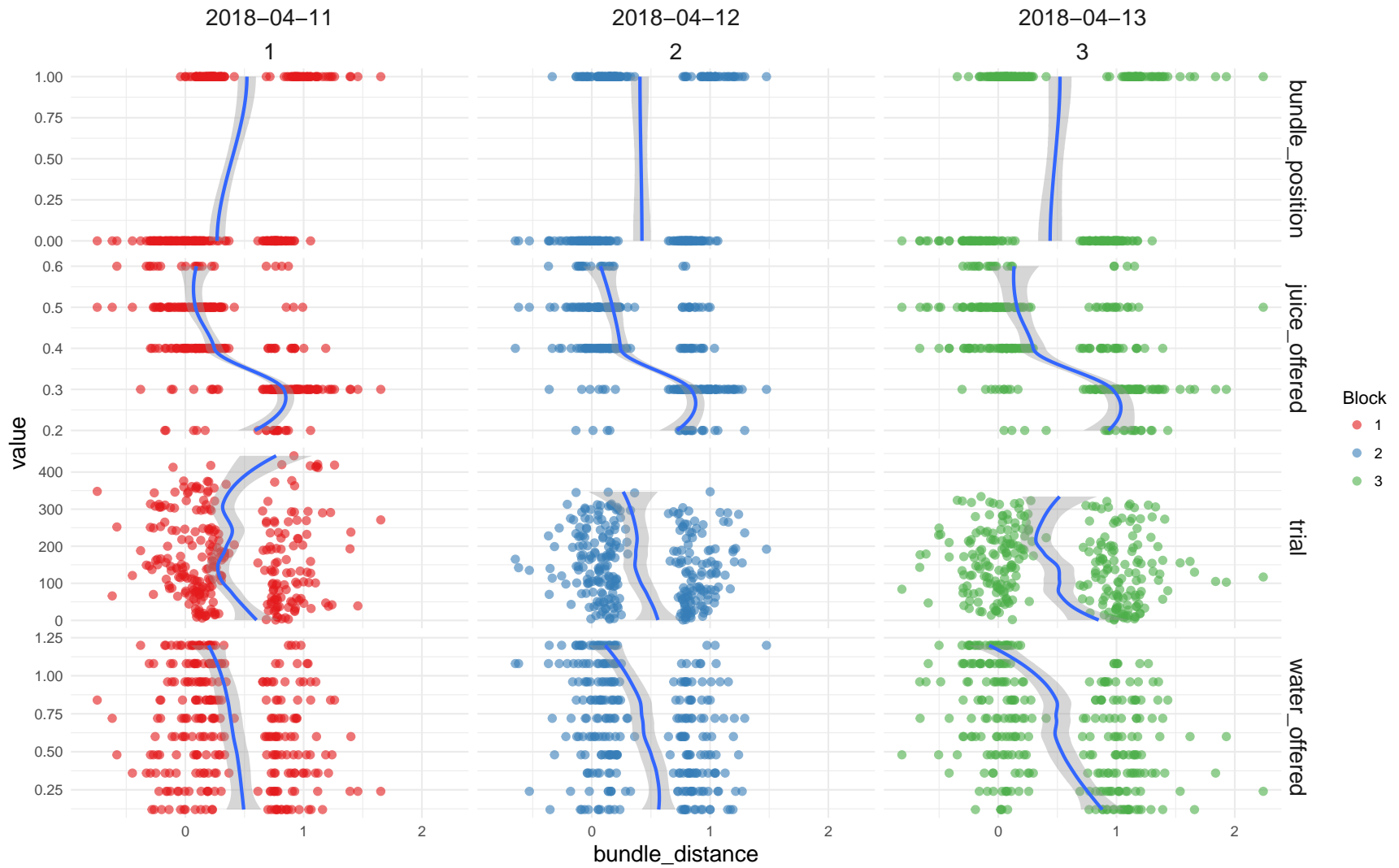
Monkey Choice Distance From Bundle on Binary Choice Task

Ulysses : 11-Apr-2018 – 13-Apr-2018



Monkey Choice Distance From Bundle on Binary Choice Task

Ulysses : 11-Apr-2018 – 13-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
## -3.2006  -0.5305   0.1684   0.5750   2.6413
##
## Coefficients:
##              Estimate Std. Error z value Pr(>|z|)
## (Intercept)    5.375e+03  2.168e+03   2.480   0.0131 *
## bundle_position  9.381e-01  2.036e-01   4.607 4.09e-06 ***
## water_offered    3.568e+00  3.439e-01  10.373 < 2e-16 ***
## juice_offered    1.982e+01  1.420e+00  13.957 < 2e-16 ***
## trial           5.232e-03  1.036e-03   5.050 4.43e-07 ***
## date            -3.054e-01  1.229e-01  -2.485   0.0130 *
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for binomial family taken to be 1)
##
##      Null deviance: 1148.42  on 839  degrees of freedom
## Residual deviance:  634.41  on 834  degrees of freedom
##      (285 observations deleted due to missingness)
## AIC: 646.41
##
## 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 = 363, number of trials = 840, p-value =
## 9.446e-05
## alternative hypothesis: true probability of success is not equal to 0.5
## 95 percent confidence interval:
##  0.3983365 0.4664269
## sample estimates:
## probability of success
##      0.4321429

```

```

#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
## -2.51100  -0.30530   0.04333   0.31742   2.27344
##
## Coefficients: (1 not defined because of singularities)
##              Estimate Std. Error z value Pr(>|z|)
## (Intercept)    -9.561610   1.326896  -7.206 5.76e-13 ***
## bundle_position    2.718804   0.508331   5.348 8.87e-08 ***
## water_offered     6.485957   0.952612   6.809 9.86e-12 ***
## as.factor(juice_offered)0.3 -1.819730   0.841420  -2.163  0.0306 *
## as.factor(juice_offered)0.4  3.915466   0.846126   4.628 3.70e-06 ***
## as.factor(juice_offered)0.5  5.012243   0.917542   5.463 4.69e-08 ***
## as.factor(juice_offered)0.6  4.721712   1.018095   4.638 3.52e-06 ***
## trial            0.010986   0.002642   4.158 3.21e-05 ***
## 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: 387.25  on 279  degrees of freedom
## Residual deviance: 148.28  on 272  degrees of freedom
## (54 observations deleted due to missingness)
## AIC: 164.28

```

```
##
```

```
## 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 = 113, number of trials = 280, p-value =
```

```
## 0.001493
```

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

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

```
## 0.3456128 0.4635936
```

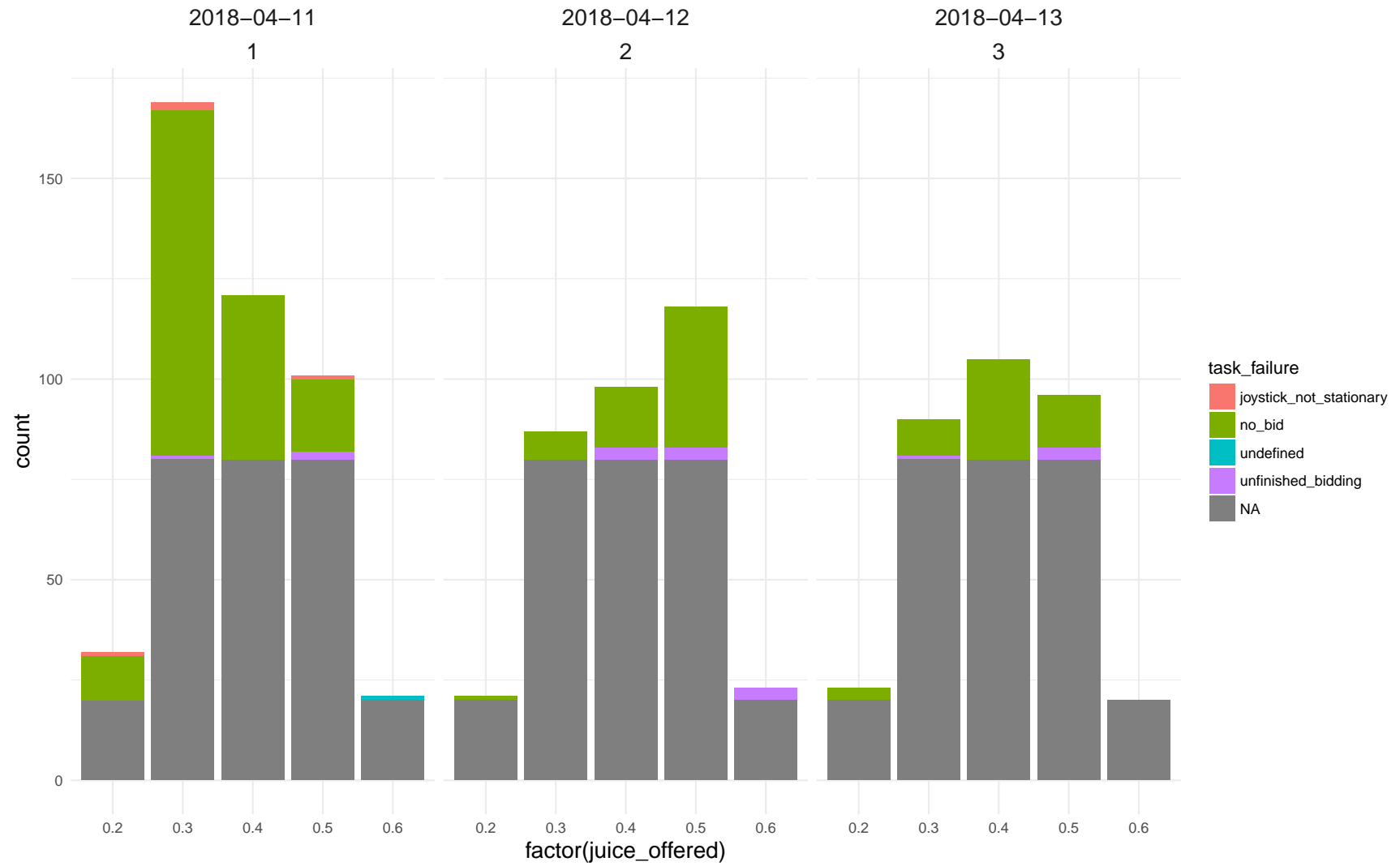
```
## sample estimates:
```

```
## probability of success
```

```
## 0.4035714
```

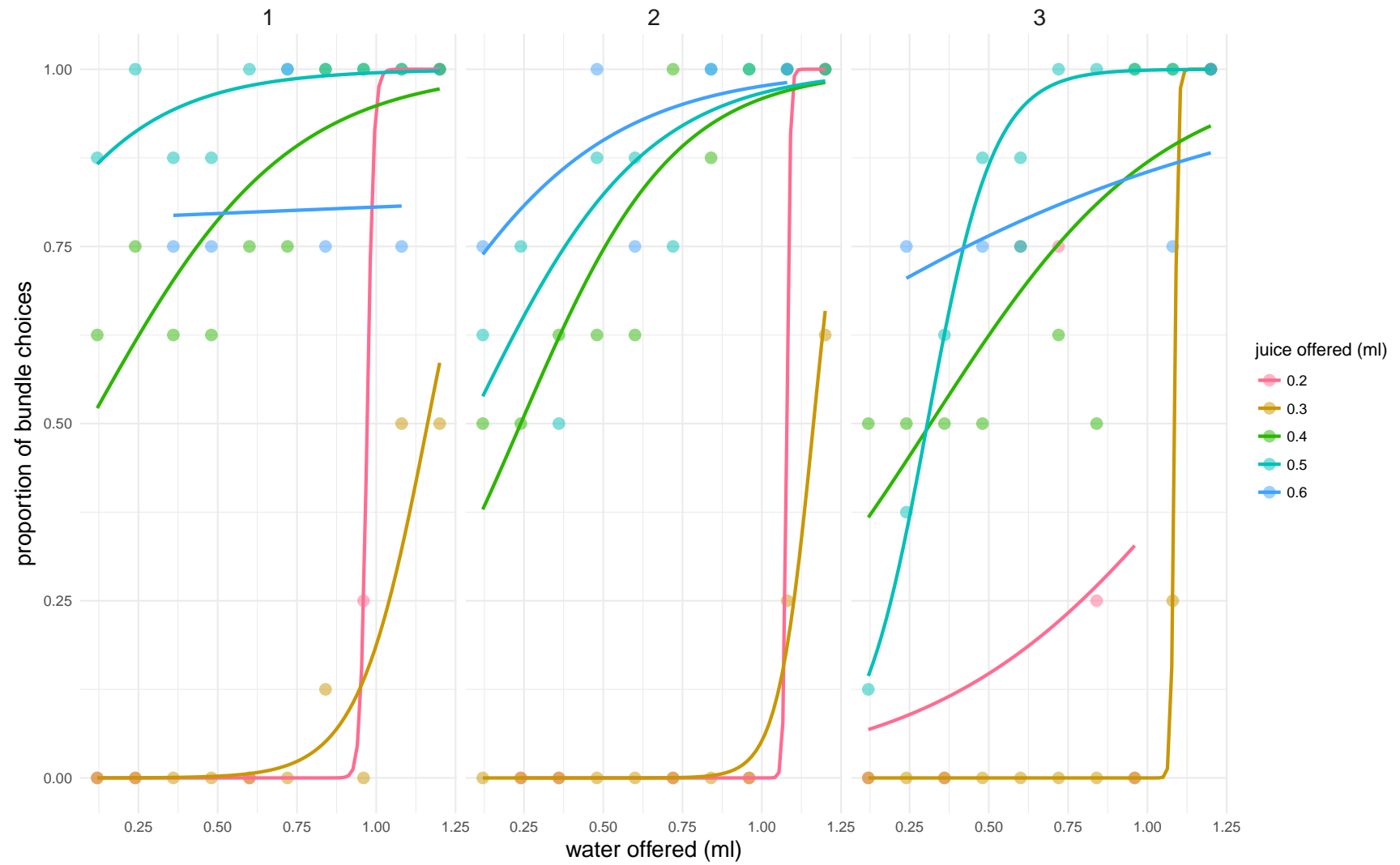
Monkey Choice Failures

Ulysses : 11-Apr-2018 – 13-Apr-2018



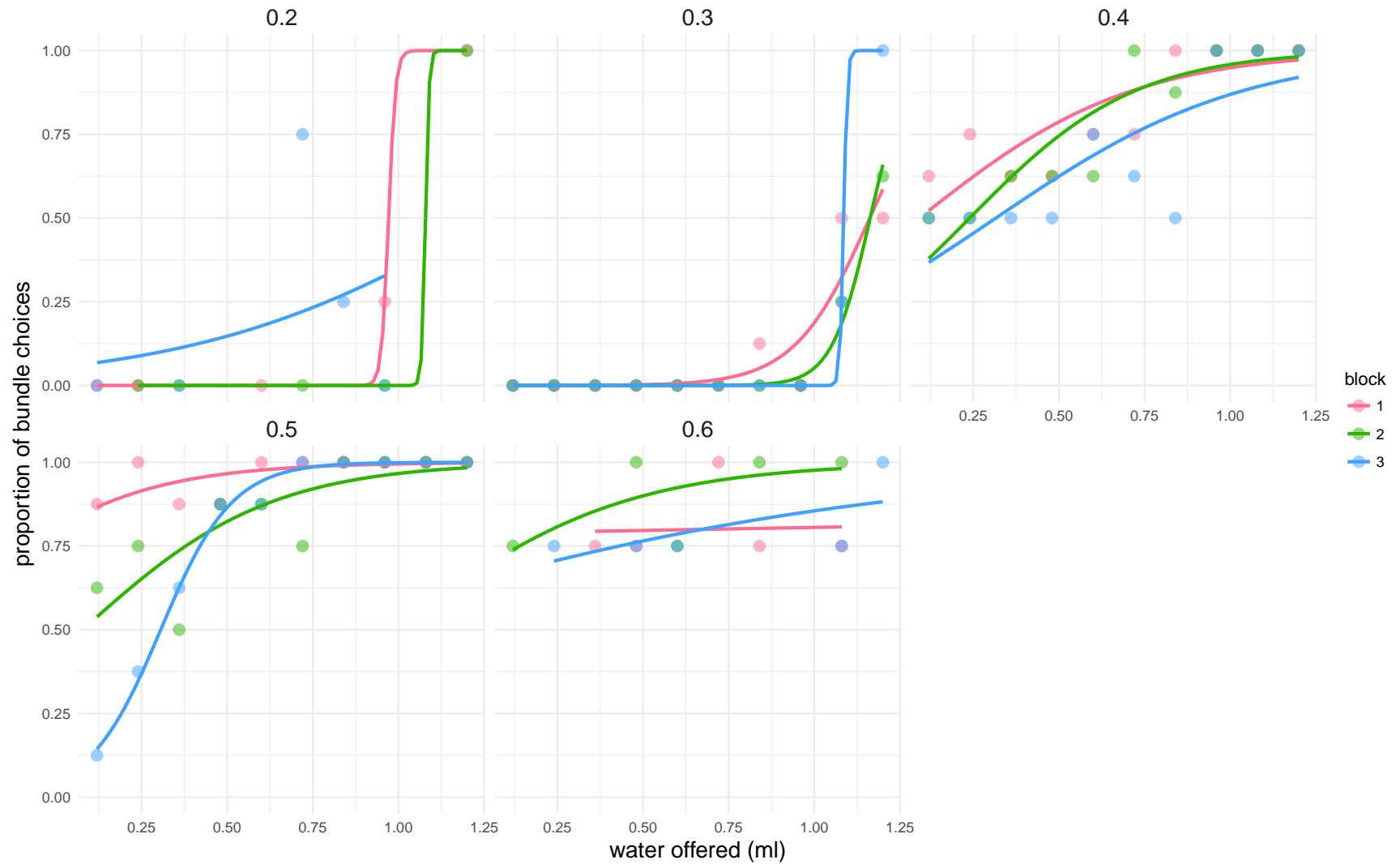
Monkey Bundle Choice Binoimial Curves

Ulysses : 11-Apr-2018 – 13-Apr-2018



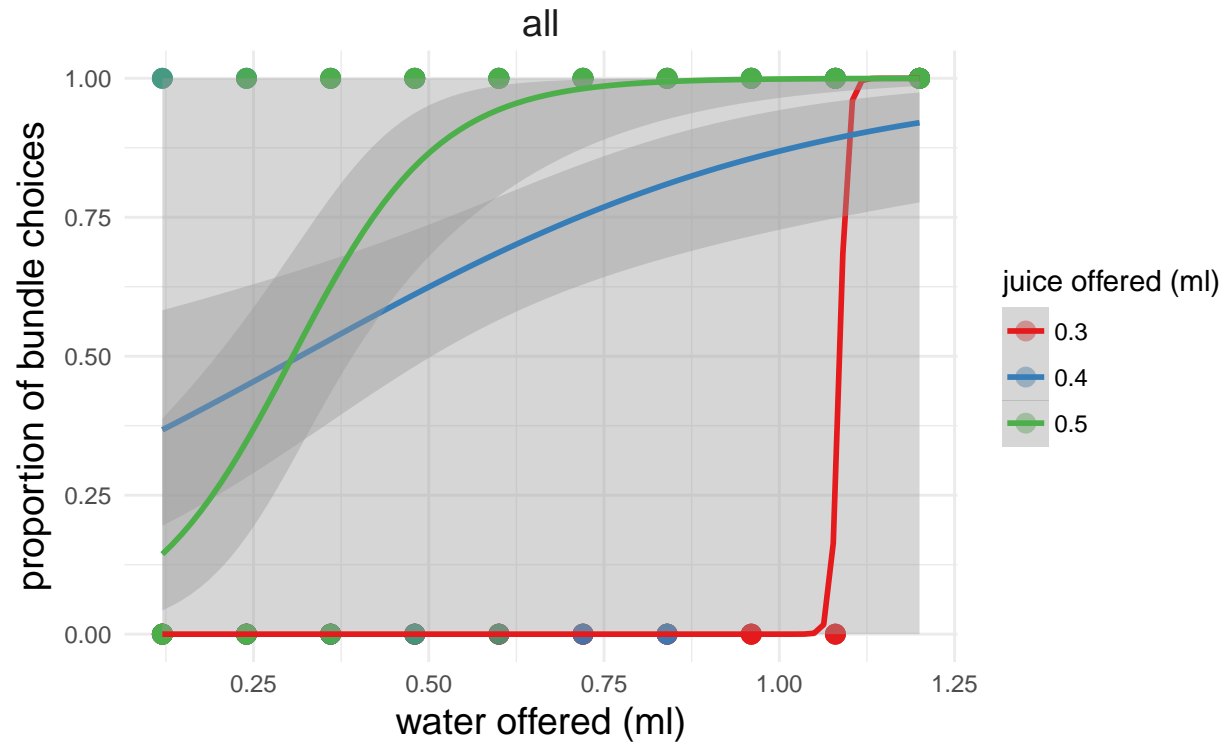
Monkey Bundle Choice Binoimial Curves

Ulysses : 11-Apr-2018 – 13-Apr-2018



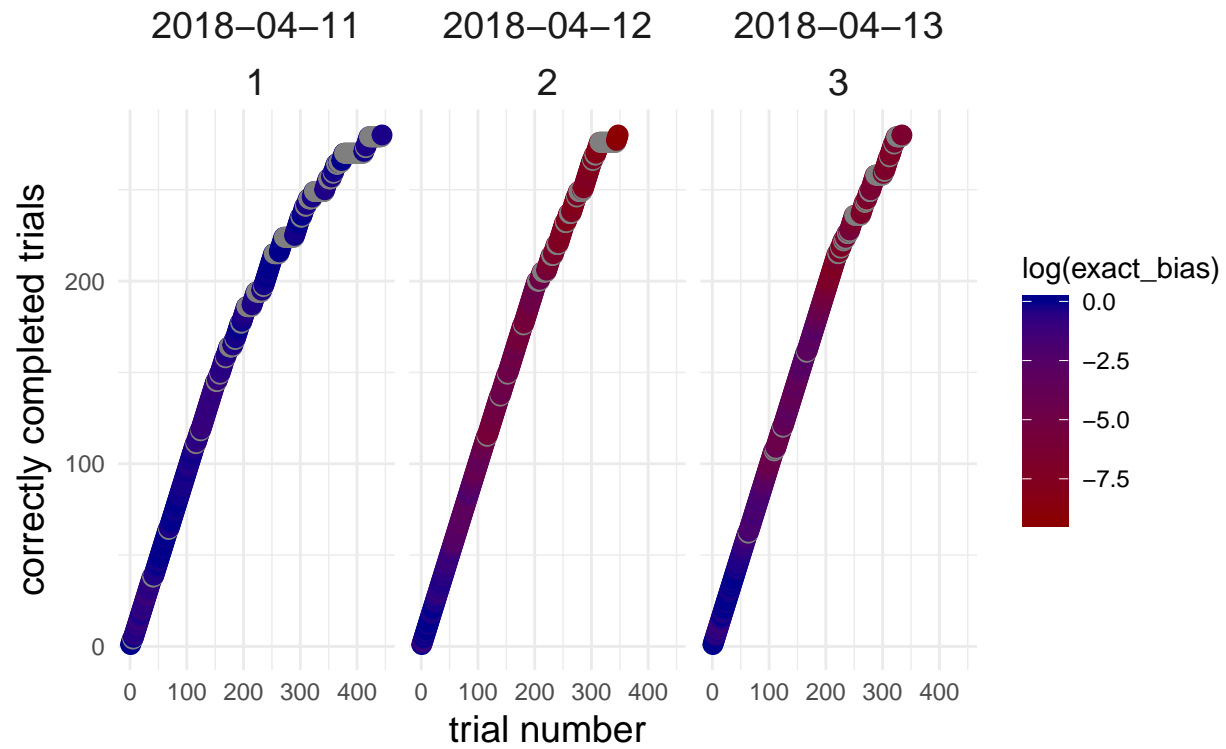
Today's Monkey Bundle Choice Binoimial Curves

Ulysses : 13-Apr-2018



Monkey Trial Progression and Bias

Ulysses : 11-Apr-2018 – 13-Apr-2018



Monkey Trial Progression and Bias

Ulysses : 11-Apr-2018 – 13-Apr-2018

