

# Binary Choice Analysis

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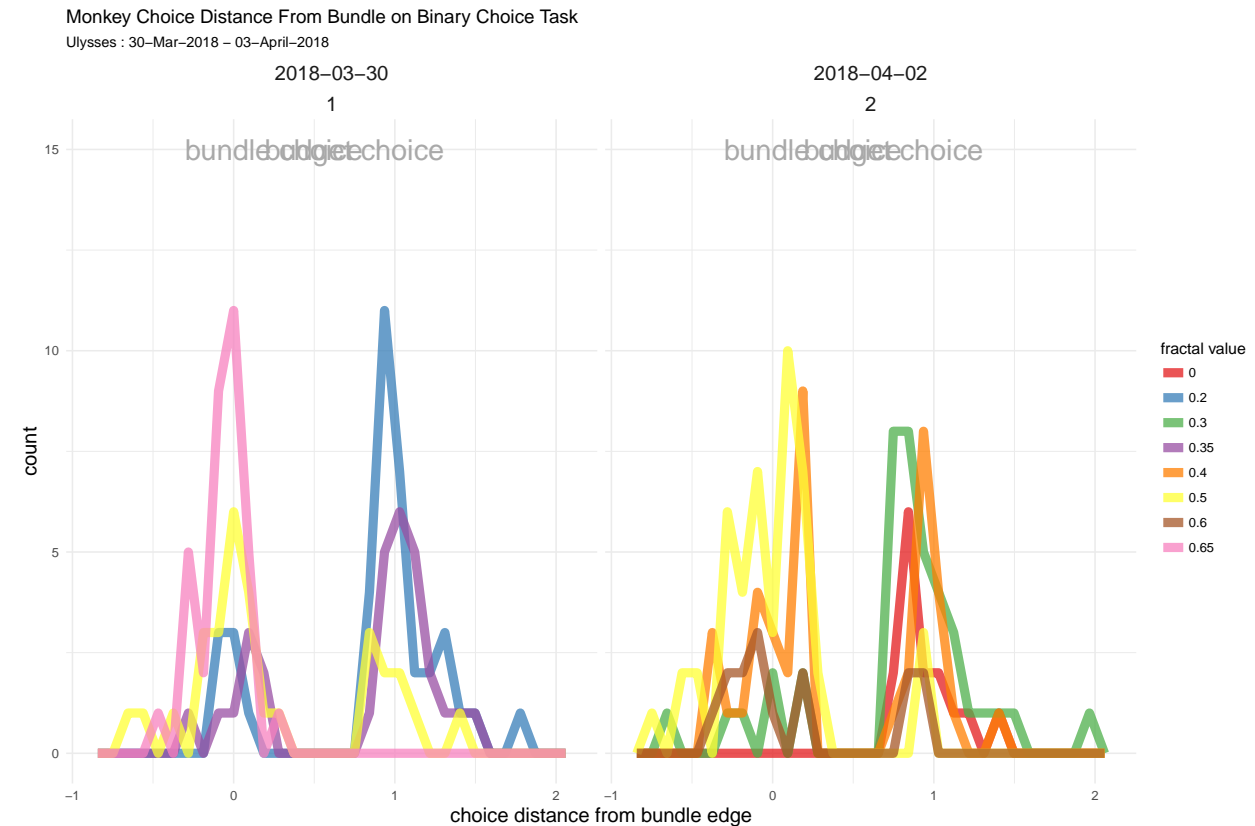
*22 February 2018*

```
monkey <- "Ulysses"  
today <- "03-April-2018"  
look_back <- "30-Mar-2018"
```

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

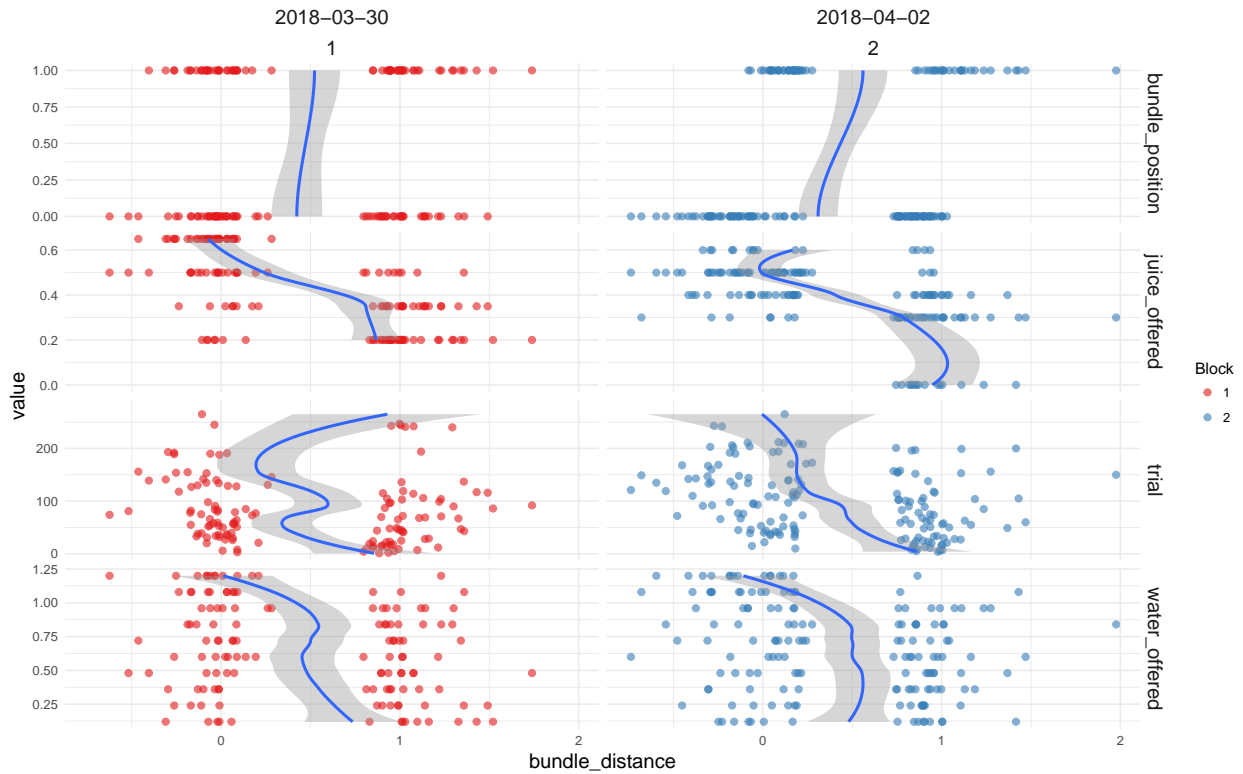
```
merge_days <- TRUE
```

p1



p2

Monkey Choice Distance From Bundle on Binary Choice Task  
Ulysses : 30-Mar-2018 – 03-April-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
## -2.86408  -0.41222   0.06723   0.42985   2.51665
##
## Coefficients:
##              Estimate Std. Error z value Pr(>|z|)
## (Intercept)  -552.10820  2175.40028  -0.254   0.7997
## bundle_position    1.04577    0.39179   2.669   0.0076 **
## water_offered     3.88733    0.65896   5.899 3.65e-09 ***
## juice_offered     17.68980    2.20919   8.007 1.17e-15 ***
## trial            0.01640    0.00342   4.795 1.63e-06 ***
## date             0.03069    0.12345   0.249   0.8037
## ---
```

```

## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for binomial family taken to be 1)
##
##    Null deviance: 400.64  on 289  degrees of freedom
## Residual deviance: 190.34  on 284  degrees of freedom
##    (259 observations deleted due to missingness)
## AIC: 202.34
##
## 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 = 147, number of trials = 290, p-value =
## 0.8602
## alternative hypothesis: true probability of success is not equal to 0.5
## 95 percent confidence interval:
##  0.4478168 0.5658340
## sample estimates:
## probability of success
##          0.5068966
#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
## -2.46219  -0.26865   0.03381   0.35023   2.07437
##
## Coefficients: (1 not defined because of singularities)
##              Estimate Std. Error z value Pr(>|z|)
## (Intercept)    -2.657e+01  1.295e+03  -0.021   0.9836
## bundle_position    1.542e+00  6.362e-01   2.424   0.0154 *
## water_offered     4.399e+00  1.057e+00   4.161 3.17e-05 ***
## as.factor(juice_offered)0.3  1.807e+01  1.295e+03   0.014   0.9889

```

```
## as.factor(juice_offered)0.4 2.126e+01 1.295e+03 0.016 0.9869
## as.factor(juice_offered)0.5 2.425e+01 1.295e+03 0.019 0.9851
## as.factor(juice_offered)0.6 2.267e+01 1.295e+03 0.018 0.9860
## trial 2.632e-02 6.632e-03 3.968 7.23e-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: 216.571 on 156 degrees of freedom
```

```
## Residual deviance: 83.649 on 149 degrees of freedom
```

```
## (114 observations deleted due to missingness)
```

```
## AIC: 99.649
```

```
##
```

```
## Number of Fisher Scoring iterations: 17
```

```
#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 = 77, number of trials = 157, p-value = 0.8732
```

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

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

```
## 0.4099059 0.5713532
```

```
## sample estimates:
```

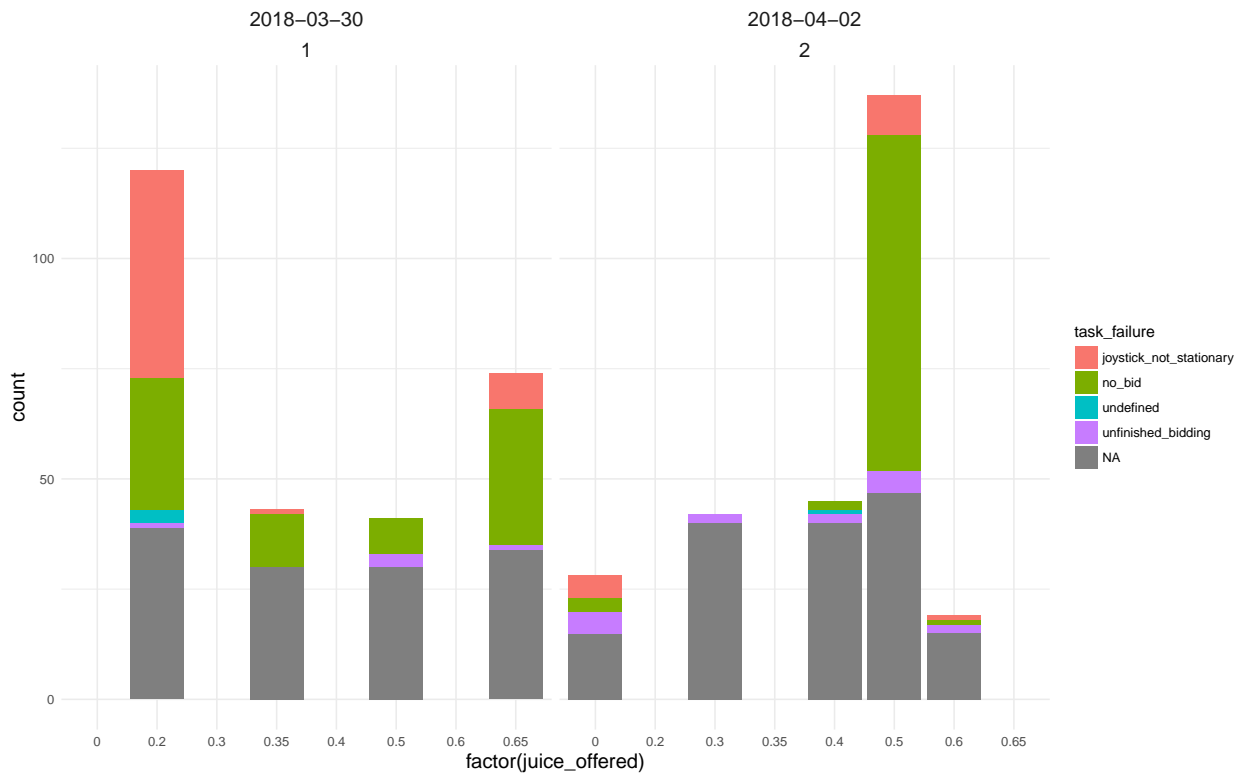
```
## probability of success
```

```
## 0.4904459
```

```
p3
```

# Monkey Choice Failures

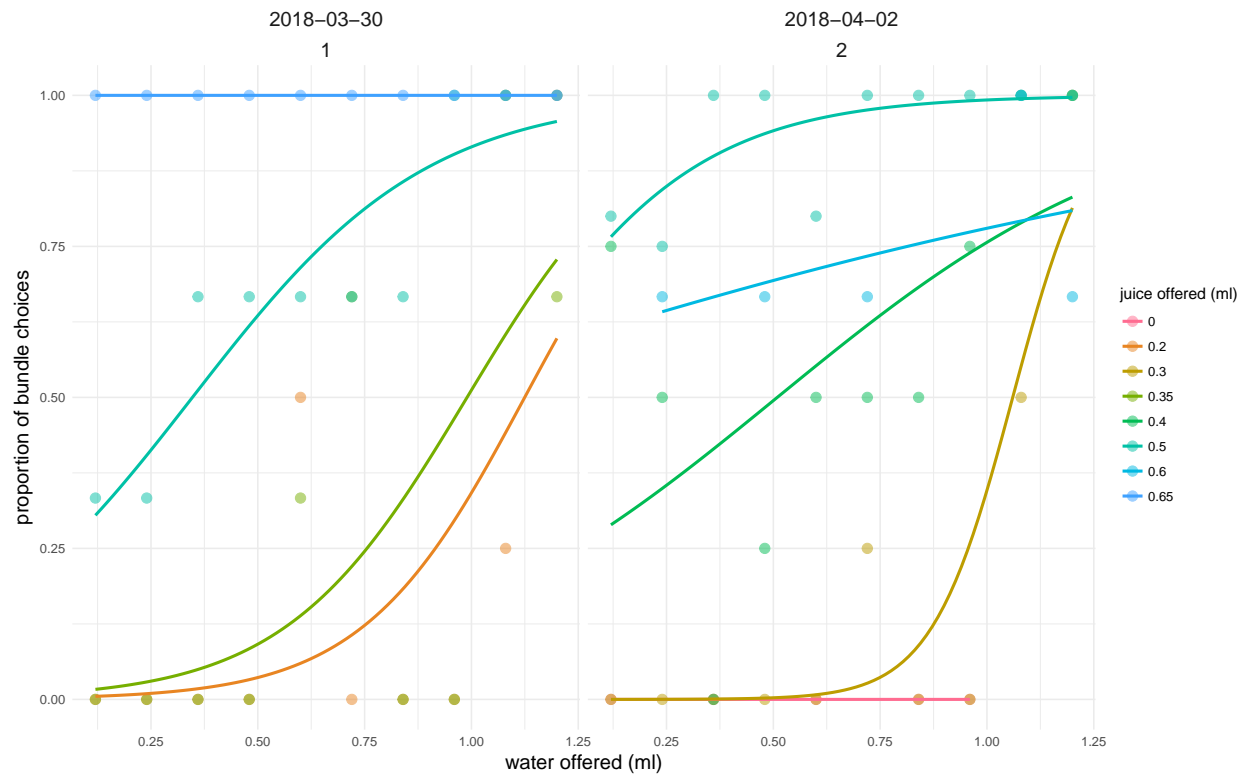
Ulysses : 30-Mar-2018 – 03-April-2018



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# Monkey Bundle Choice Binoimial Curves

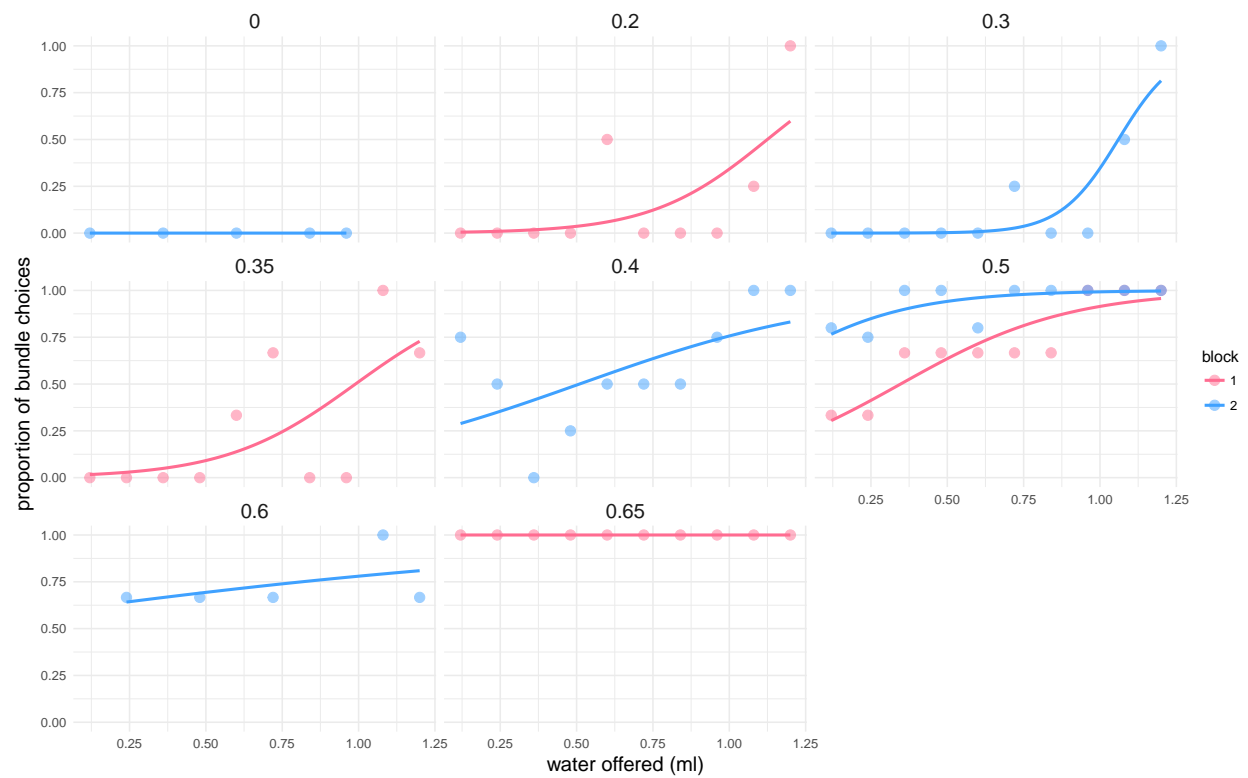
Ulysses : 30-Mar-2018 – 03-April-2018



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Ulysses : 30-Mar-2018 – 03-April-2018

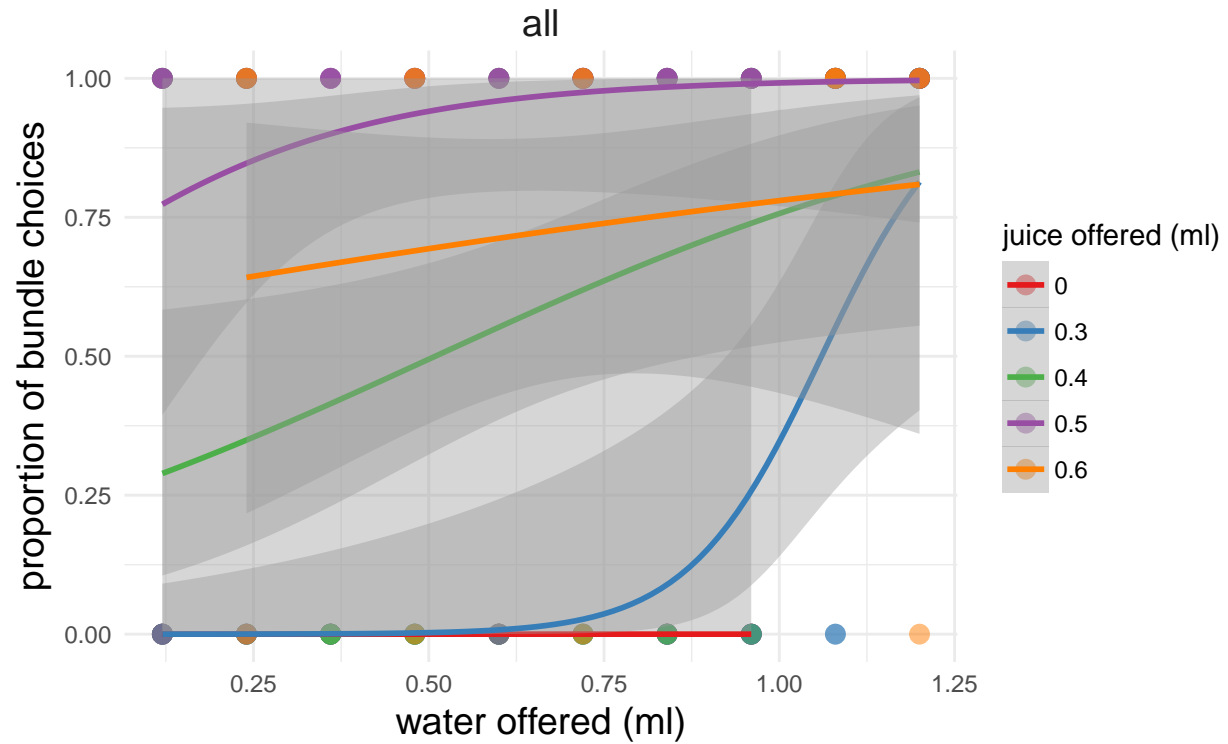
Ulysses : 30-Mar-2018 – 03-April-2018



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## Today's Monkey Bundle Choice Binoimial Curves

Ulysses : 03-April-2018

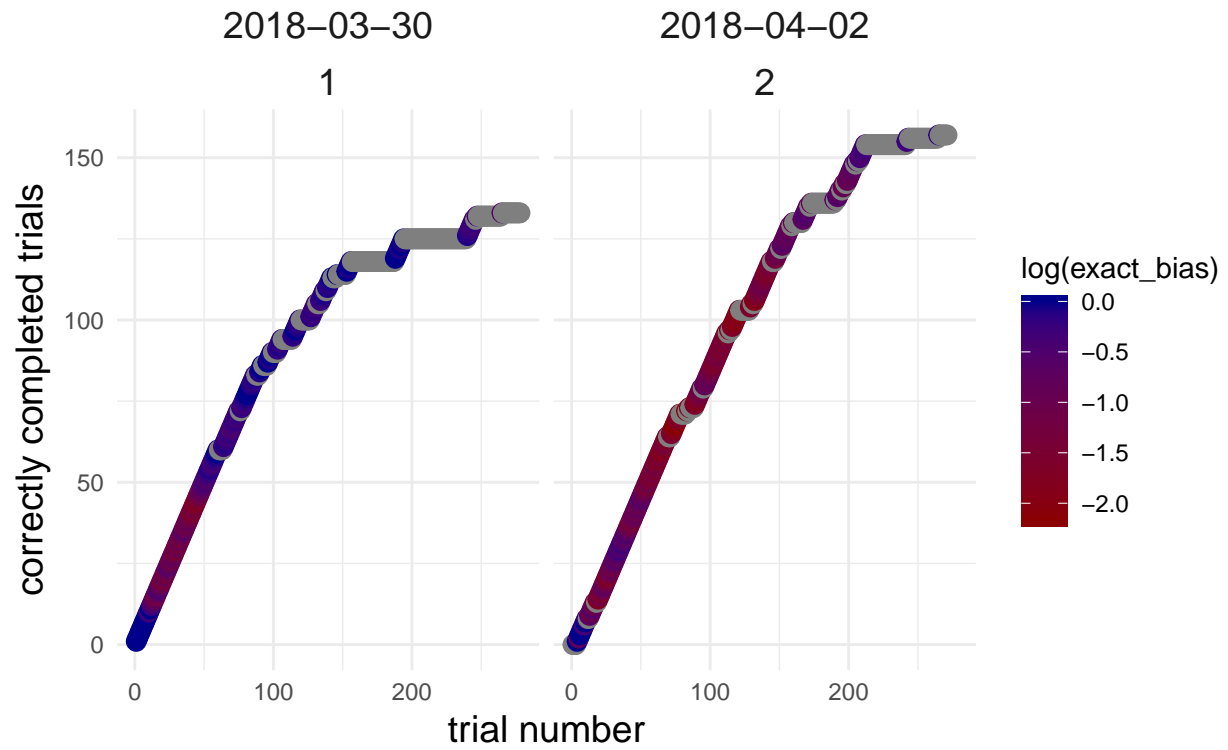


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# Monkey Trial Progression and Bias

Ulysses : 30-Mar-2018 – 03-April-2018



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# Monkey Trial Progression and Bias

Ulysses : 30-Mar-2018 – 03-April-2018

