

Binary Choice Analysis

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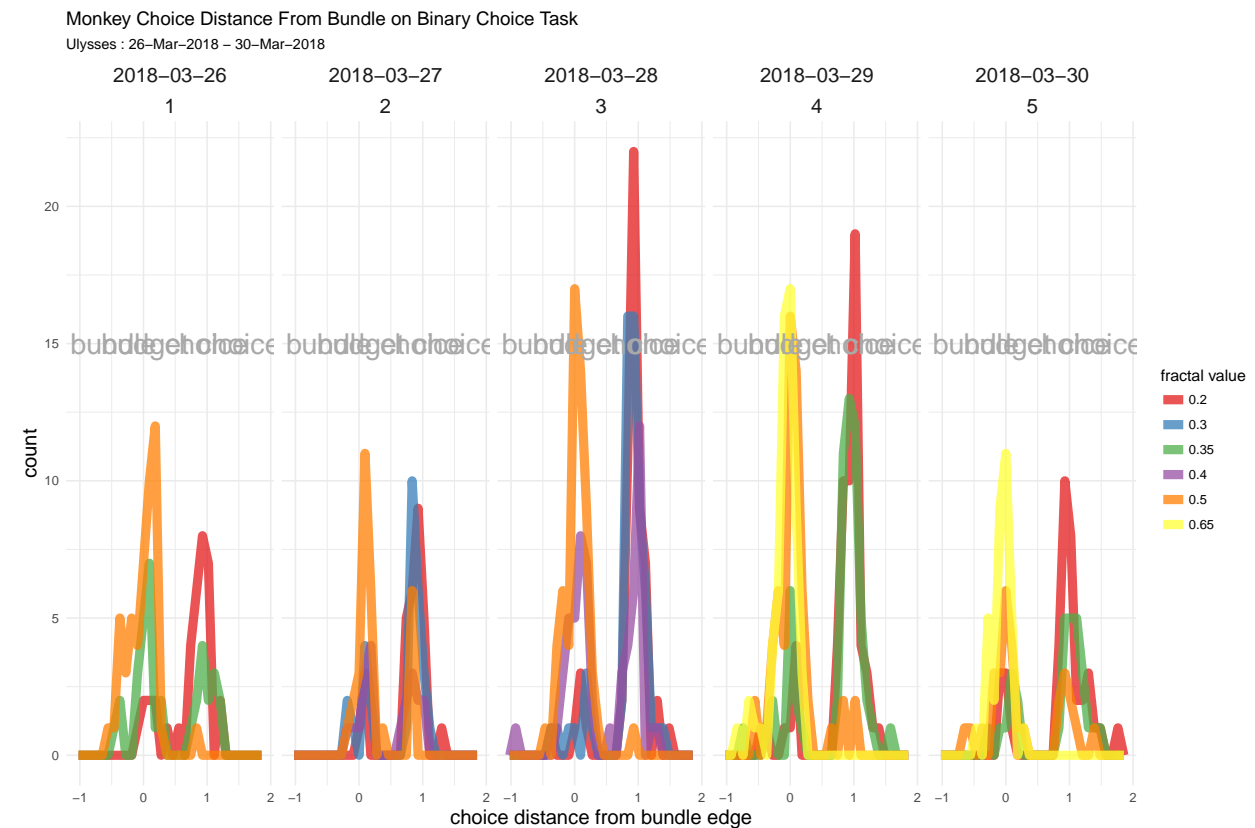
22 February 2018

```
monkey <- "Ulysses"  
today <- "30-Mar-2018"  
look_back <- "26-Mar-2018"
```

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

```
merge_days <- TRUE
```

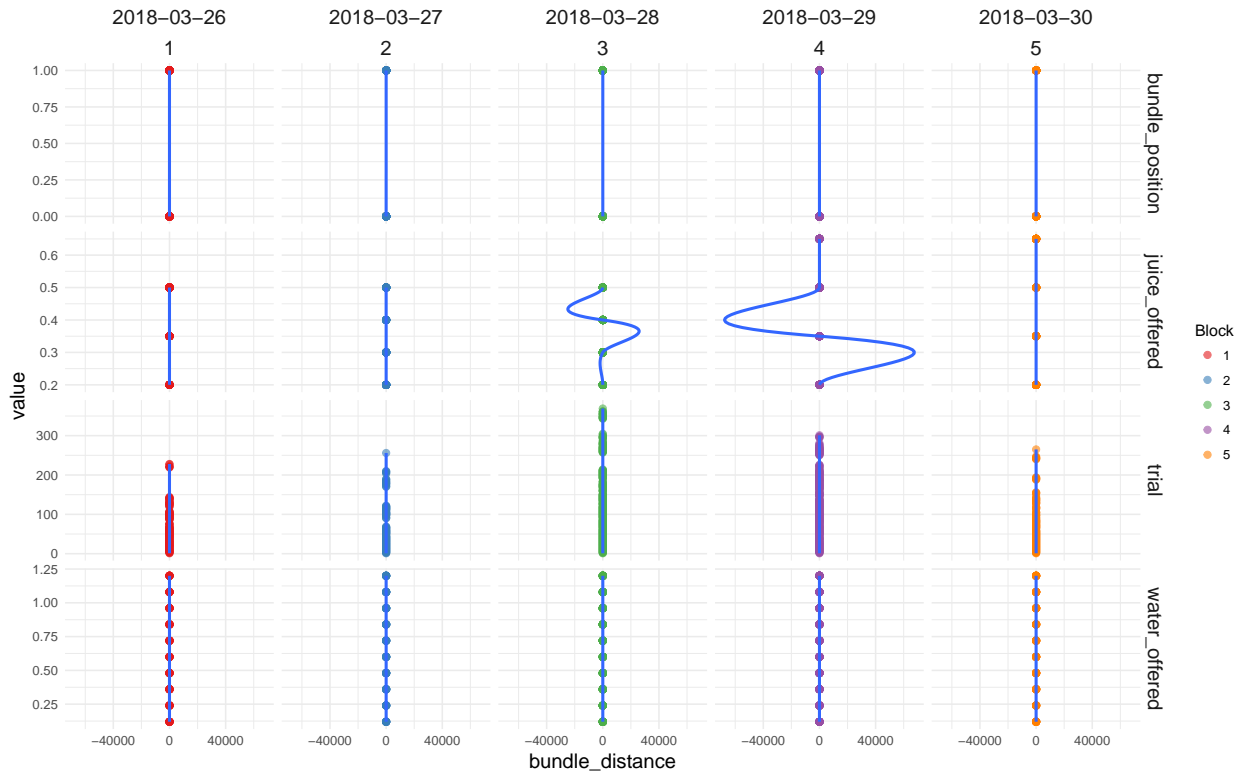
p1



p2

Monkey Choice Distance From Bundle on Binary Choice Task

Ulysses : 26-Mar-2018 - 30-Mar-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
## -2.62666  -0.40026   0.03784   0.37645   3.11618
##
## Coefficients:
##              Estimate Std. Error z value Pr(>|z|)
## (Intercept)  7.322e+03  1.698e+03  4.313 1.61e-05 ***
## bundle_position 1.956e-01  2.240e-01  0.873  0.383
## water_offered  4.851e+00  4.459e-01 10.877 < 2e-16 ***
## juice_offered  2.049e+01  1.413e+00 14.498 < 2e-16 ***
## trial        -1.103e-03  1.269e-03 -0.869  0.385
## date         -4.162e-01  9.637e-02 -4.319 1.57e-05 ***
## ---
```

```
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for binomial family taken to be 1)
##
##    Null deviance: 1169.10  on 843  degrees of freedom
## Residual deviance:  513.55  on 838  degrees of freedom
##    (613 observations deleted due to missingness)
## AIC: 525.55
##
## 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 = 422, number of trials = 844, p-value = 1
## alternative hypothesis: true probability of success is not equal to 0.5
## 95 percent confidence interval:
##  0.4657256 0.5342744
## sample estimates:
## probability of success
##                0.5
#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)

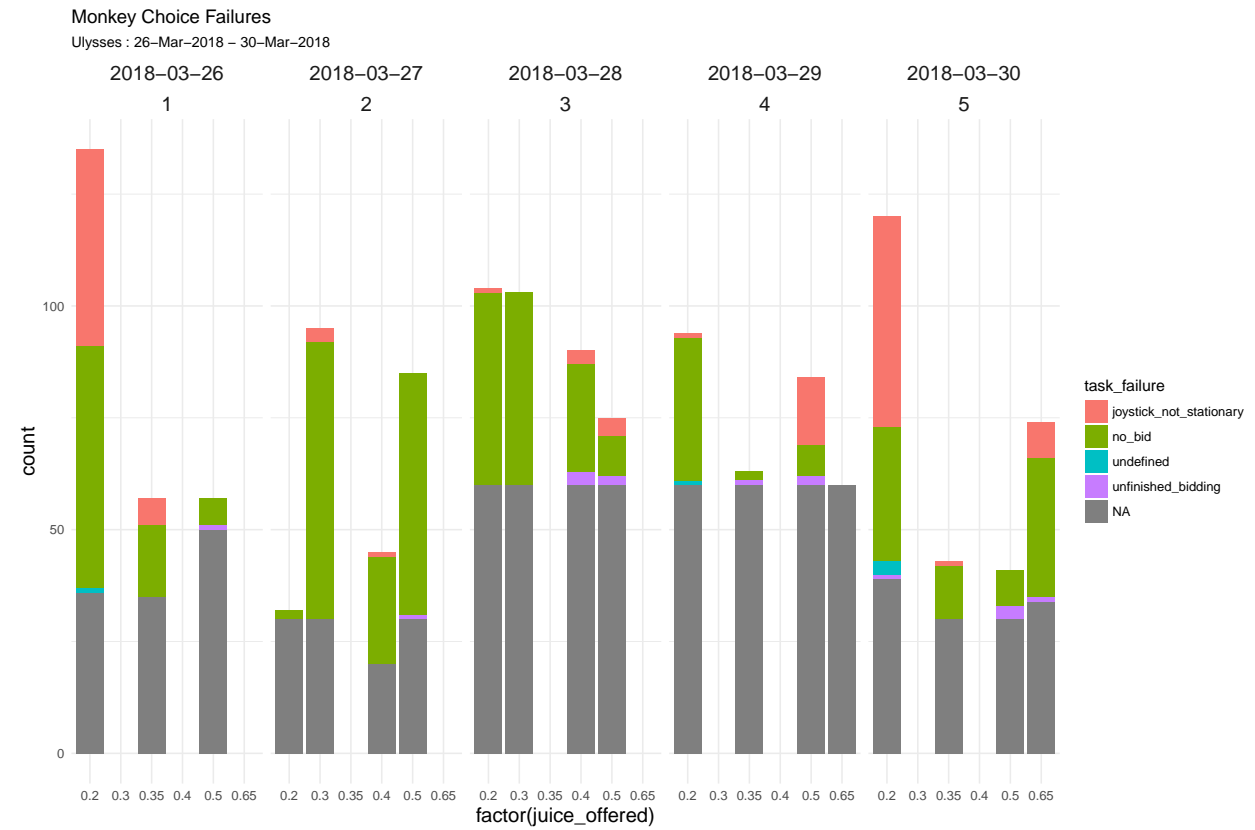
##
## Call:
## glm(formula = fractal_choice ~ bundle_position + water_offered +
##      as.factor(juice_offered) + trial + date, family = "binomial",
##      data = dplyr::filter(task_data, block_no == max(block_no)))
##
## Deviance Residuals:
##      Min       1Q   Median       3Q      Max
## -1.77312  -0.48334   0.00001   0.23731   2.33908
##
## Coefficients: (1 not defined because of singularities)
##              Estimate Std. Error z value Pr(>|z|)
## (Intercept)    -7.076e+00  1.573e+00  -4.498 6.87e-06 ***
## bundle_position    8.699e-01  6.229e-01   1.397  0.1625
## water_offered     5.127e+00  1.226e+00   4.183 2.88e-05 ***
## as.factor(juice_offered)0.35  1.120e+00  7.411e-01   1.511  0.1308
## as.factor(juice_offered)0.5   4.101e+00  9.375e-01   4.374 1.22e-05 ***
```

```

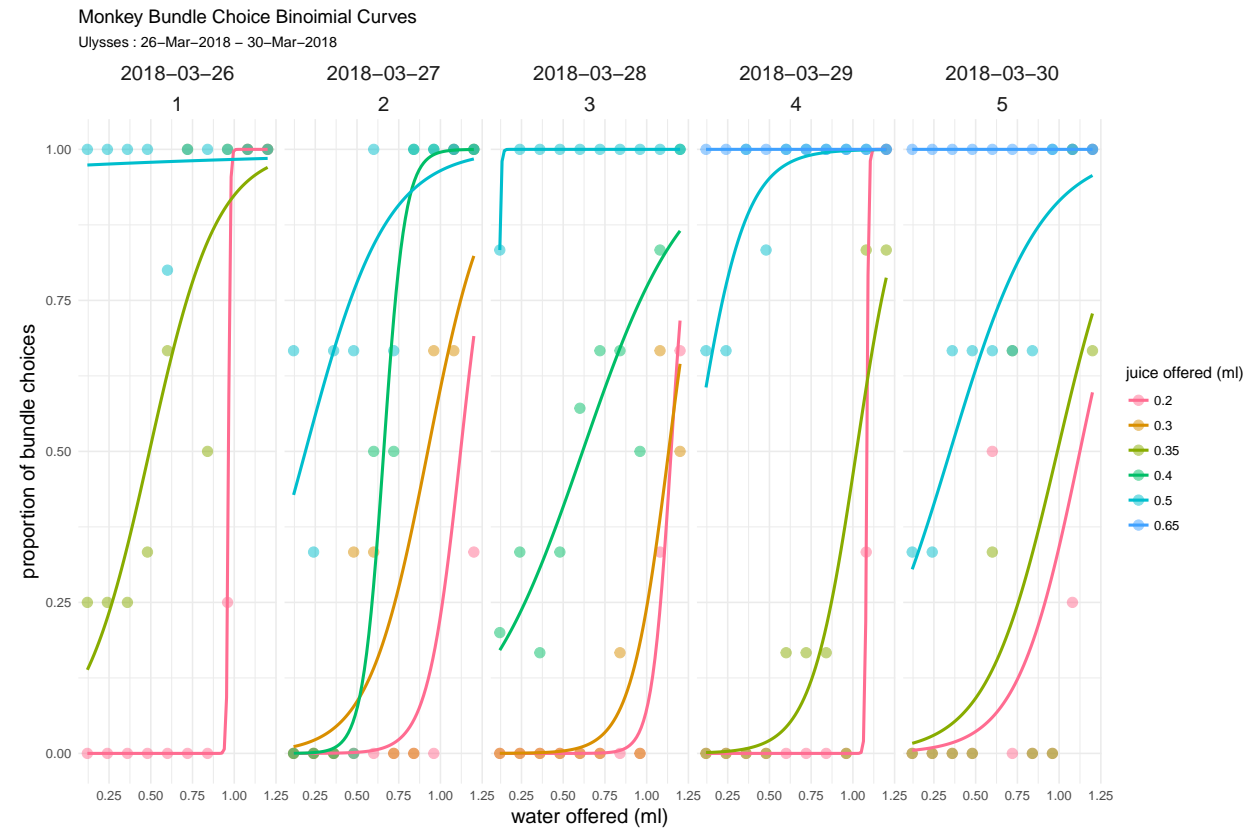
## as.factor(juice_offered)0.65  2.321e+01  1.570e+03  0.015  0.9882
## trial                        7.826e-03  4.526e-03  1.729  0.0838 .
## date                        NA          NA      NA      NA
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for binomial family taken to be 1)
##
##    Null deviance: 184.009  on 132  degrees of freedom
## Residual deviance:  76.516  on 126  degrees of freedom
## (145 observations deleted due to missingness)
## AIC: 90.516
##
## Number of Fisher Scoring iterations: 18
#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 = 70, number of trials = 133, p-value = 0.603
## alternative hypothesis: true probability of success is not equal to 0.5
## 95 percent confidence interval:
##  0.4379476 0.6134864
## sample estimates:
## probability of success
##          0.5263158
p3

```



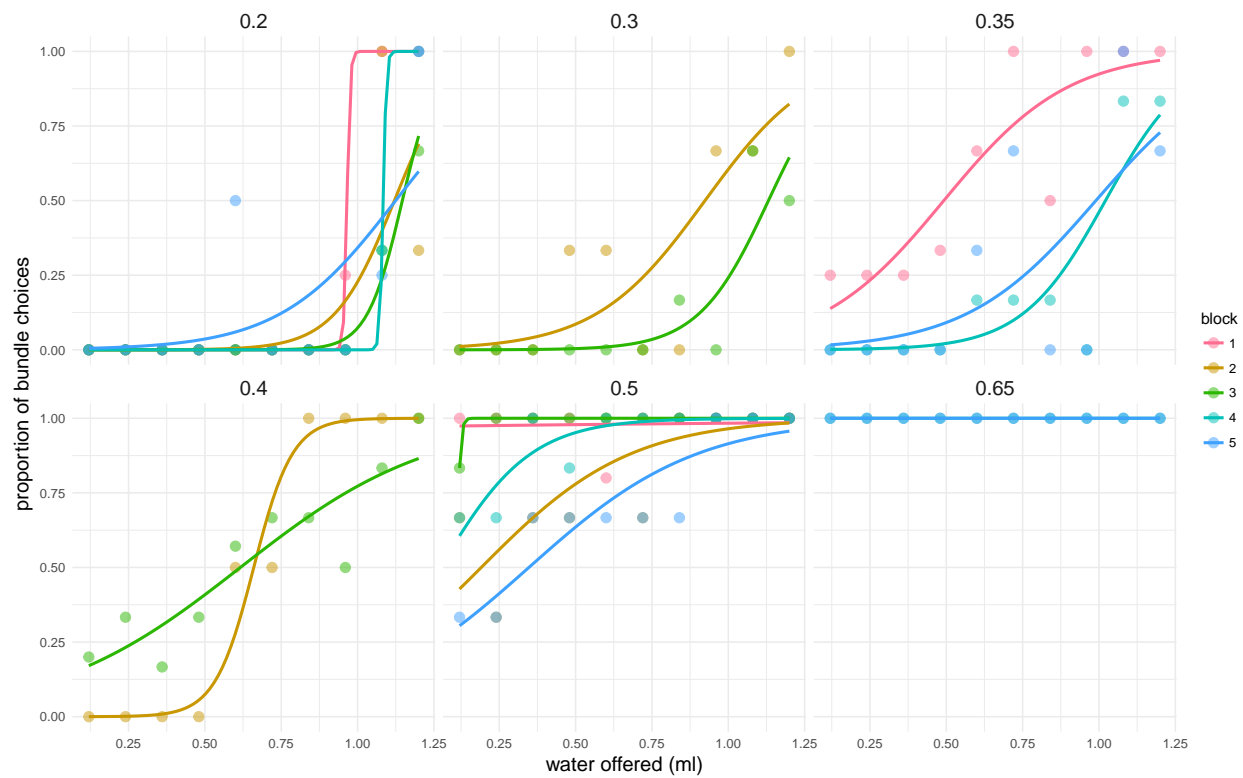
p4



p5

Monkey Bundle Choice Binoimial Curves

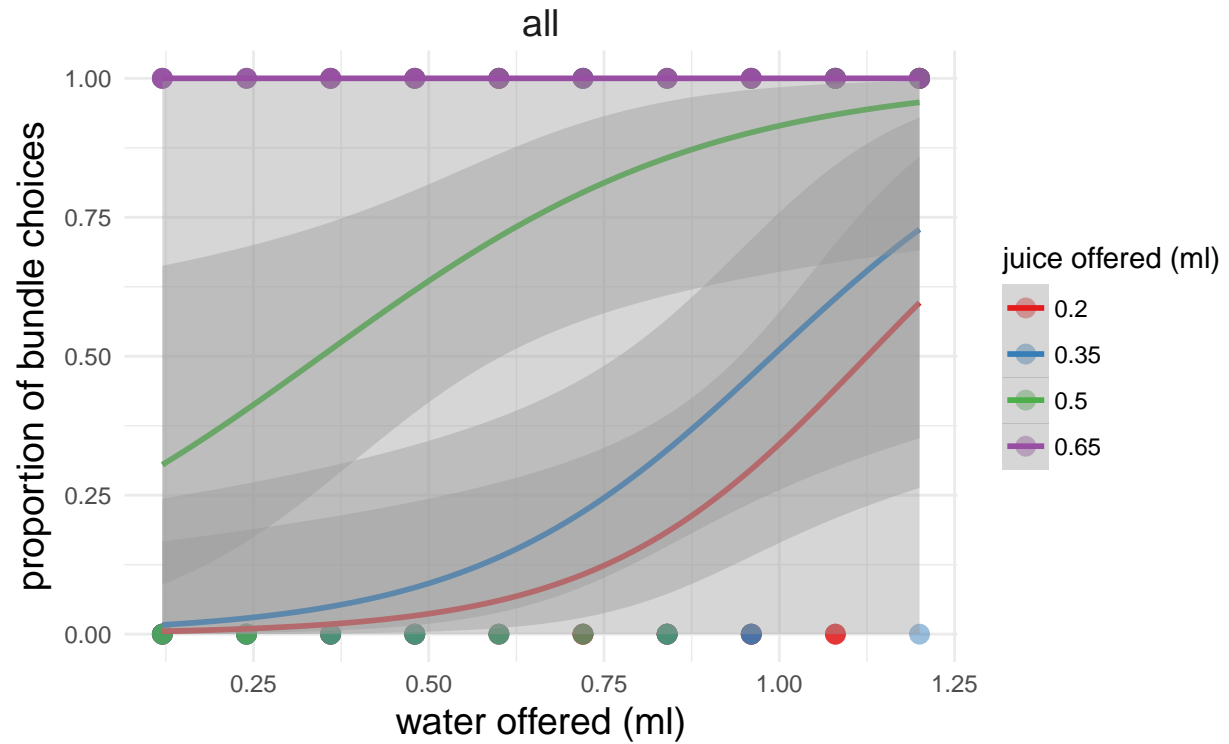
Ulysses : 26-Mar-2018 – 30-Mar-2018



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Today's Monkey Bundle Choice Binoimial Curves

Ulysses : 30-Mar-2018

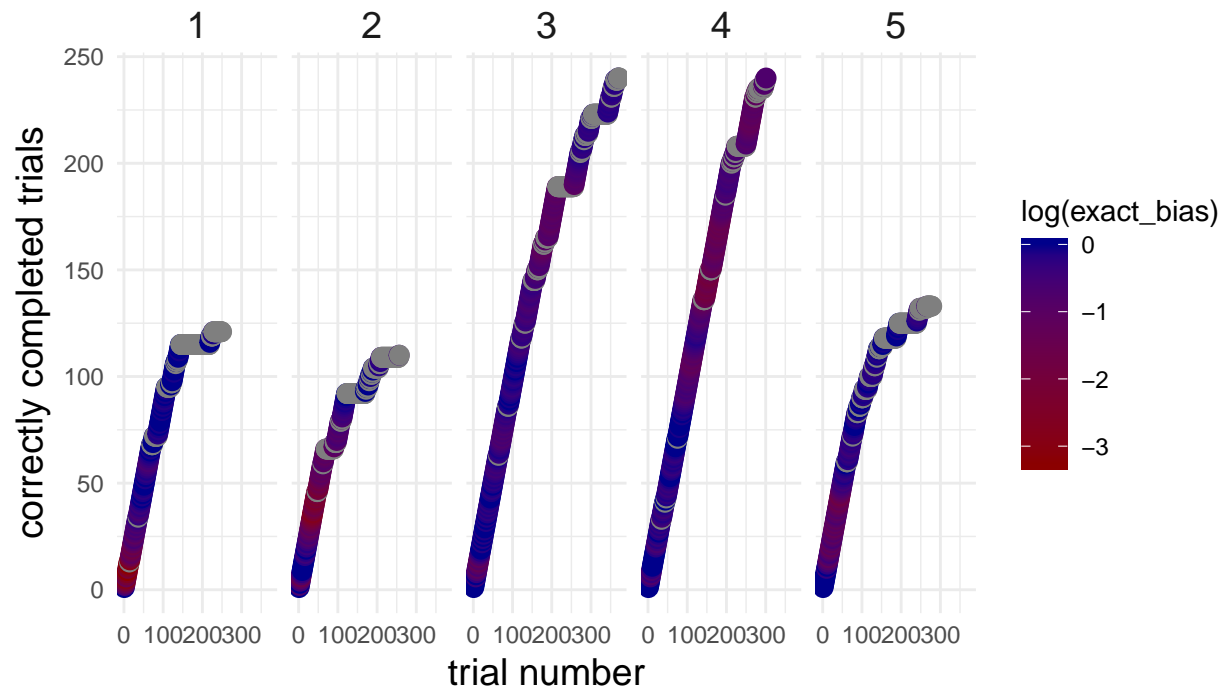


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Monkey Trial Progression and Bias

Ulysses : 26-Mar-2018 – 30-Mar-2018

018-03-2 018-03-2 018-03-2 018-03-2 018-03-3



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Monkey Trial Progression and Bias

Ulysses : 26-Mar-2018 – 30-Mar-2018

