

Binary Choice Analysis

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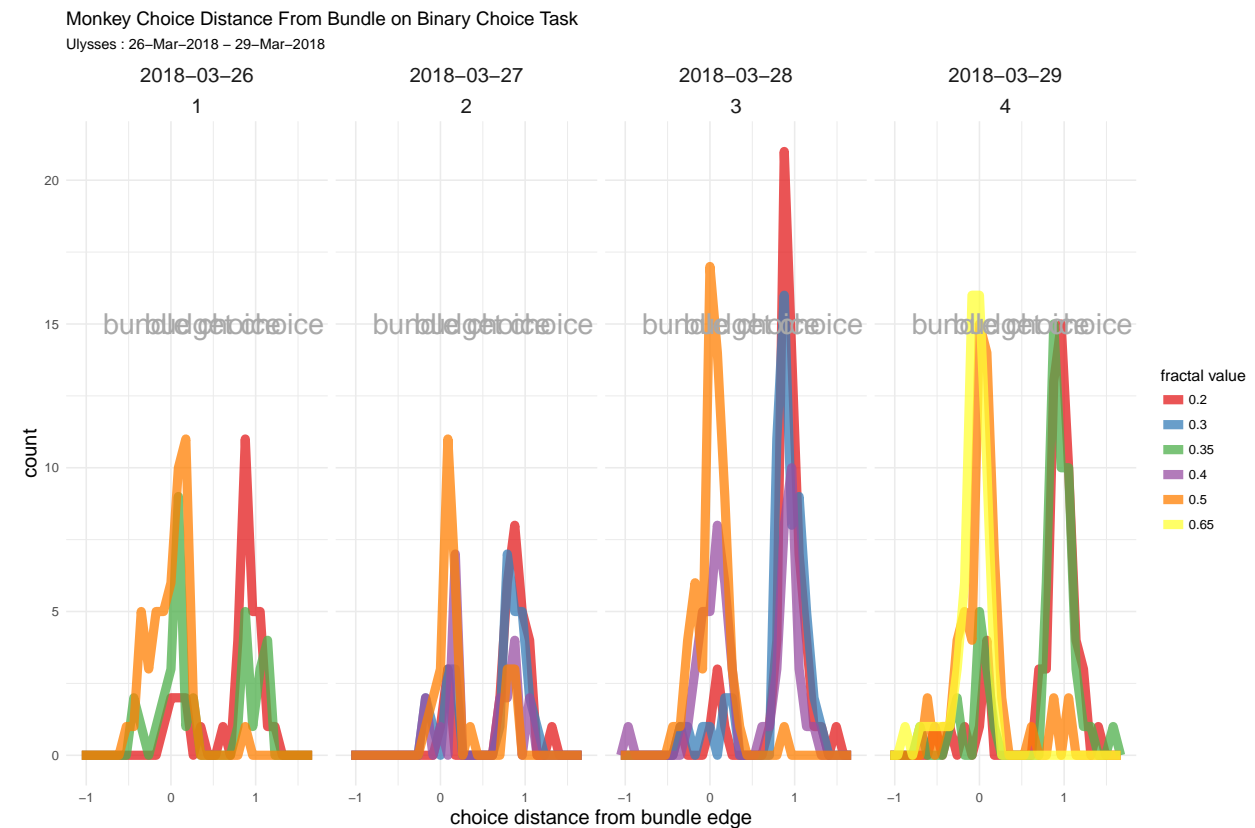
22 February 2018

```
monkey <- "Ulysses"  
today <- "29-Mar-2018"  
look_back <- "26-Mar-2018"
```

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

```
merge_days <- TRUE
```

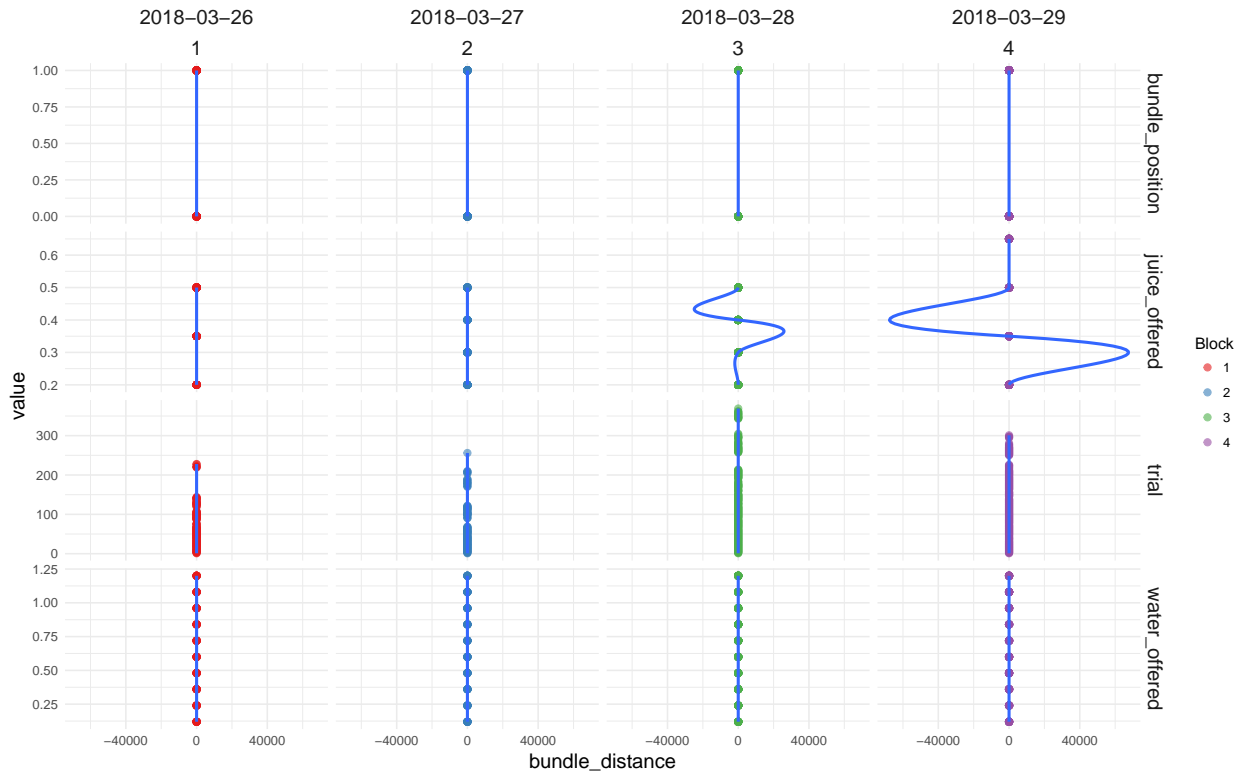
p1



p2

Monkey Choice Distance From Bundle on Binary Choice Task

Ulysses : 26-Mar-2018 - 29-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
## -2.80340  -0.37118   0.02806   0.35127   2.21960
##
## Coefficients:
##              Estimate Std. Error z value Pr(>|z|)
## (Intercept)  8.990e+03  2.325e+03  3.866 0.000111 ***
## bundle_position  9.108e-02  2.503e-01  0.364 0.715925
## water_offered   5.142e+00  5.131e-01  10.022 < 2e-16 ***
## juice_offered   2.266e+01  1.726e+00  13.129 < 2e-16 ***
## trial          -1.908e-03  1.413e-03  -1.350 0.177077
## date           -5.110e-01  1.320e-01  -3.871 0.000109 ***
## ---
```

```
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for binomial family taken to be 1)
##
##    Null deviance: 985.03  on 710  degrees of freedom
## Residual deviance: 412.71  on 705  degrees of freedom
##    (468 observations deleted due to missingness)
## AIC: 424.71
##
## 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 = 352, number of trials = 711, p-value = 0.822
## alternative hypothesis: true probability of success is not equal to 0.5
## 95 percent confidence interval:
##  0.4577096 0.5324862
## sample estimates:
## probability of success
##      0.4950774
#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)

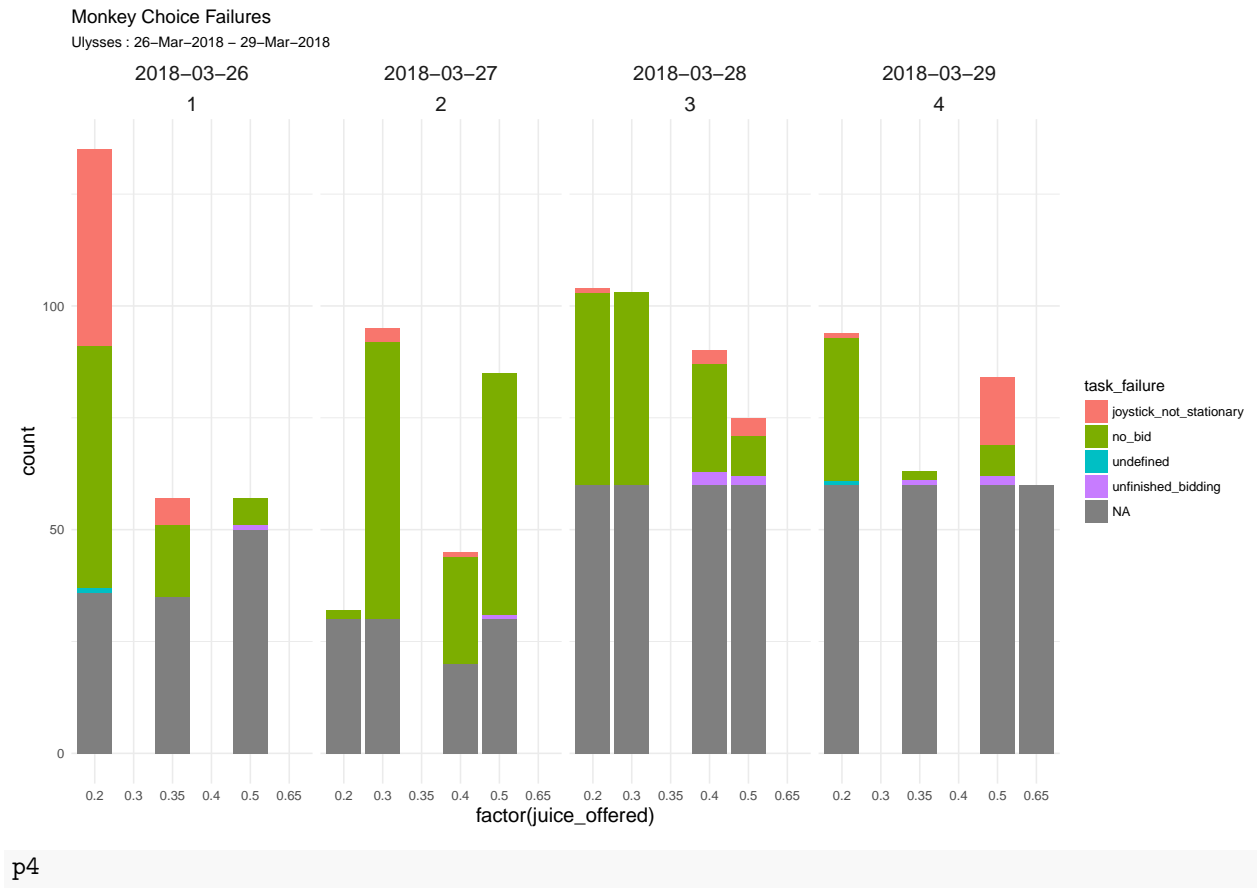
##
## Call:
## glm(formula = fractal_choice ~ bundle_position + water_offered +
##      as.factor(juice_offered) + trial + date, family = "binomial",
##      data = dplyr::filter(task_data, block_no == max(block_no)))
##
## Deviance Residuals:
##      Min       1Q   Median       3Q      Max
## -2.38418  -0.08472   0.00000   0.02614   2.65470
##
## Coefficients: (1 not defined because of singularities)
##              Estimate Std. Error z value Pr(>|z|)
## (Intercept)    -1.129e+01  2.315e+00  -4.879 1.06e-06 ***
## bundle_position    1.018e+00  6.025e-01   1.690   0.091 .
## water_offered     9.830e+00  2.056e+00   4.780 1.75e-06 ***
## as.factor(juice_offered)0.35  1.132e+00  7.010e-01   1.615   0.106
## as.factor(juice_offered)0.5   1.000e+01  1.888e+00   5.300 1.16e-07 ***
```

```

## as.factor(juice_offered)0.65  2.816e+01  1.799e+03   0.016   0.988
## trial                        -2.147e-03  3.745e-03  -0.573   0.566
## 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: 328.431  on 239  degrees of freedom
## Residual deviance:  74.759  on 233  degrees of freedom
##    (61 observations deleted due to missingness)
## AIC: 88.759
##
## Number of Fisher Scoring iterations: 19
#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))])))

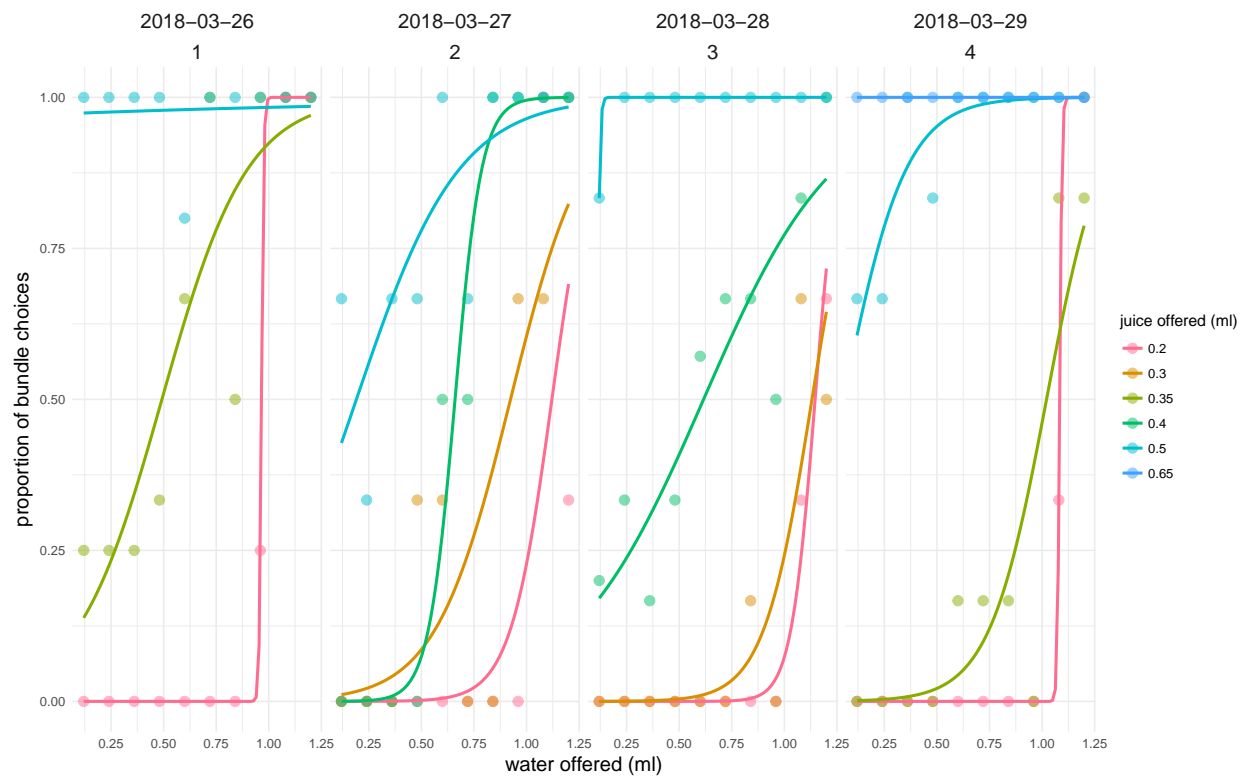
##
## Exact binomial test
##
## data:  c(nrow(task_data %>% .[c(bundle_position != fractal_choice &      block_no == max(block_no))])
## number of successes = 114, number of trials = 240, p-value =
## 0.4778
## alternative hypothesis: true probability of success is not equal to 0.5
## 95 percent confidence interval:
##  0.4103929 0.5402322
## sample estimates:
## probability of success
##          0.475
p3

```



Monkey Bundle Choice Binoimial Curves

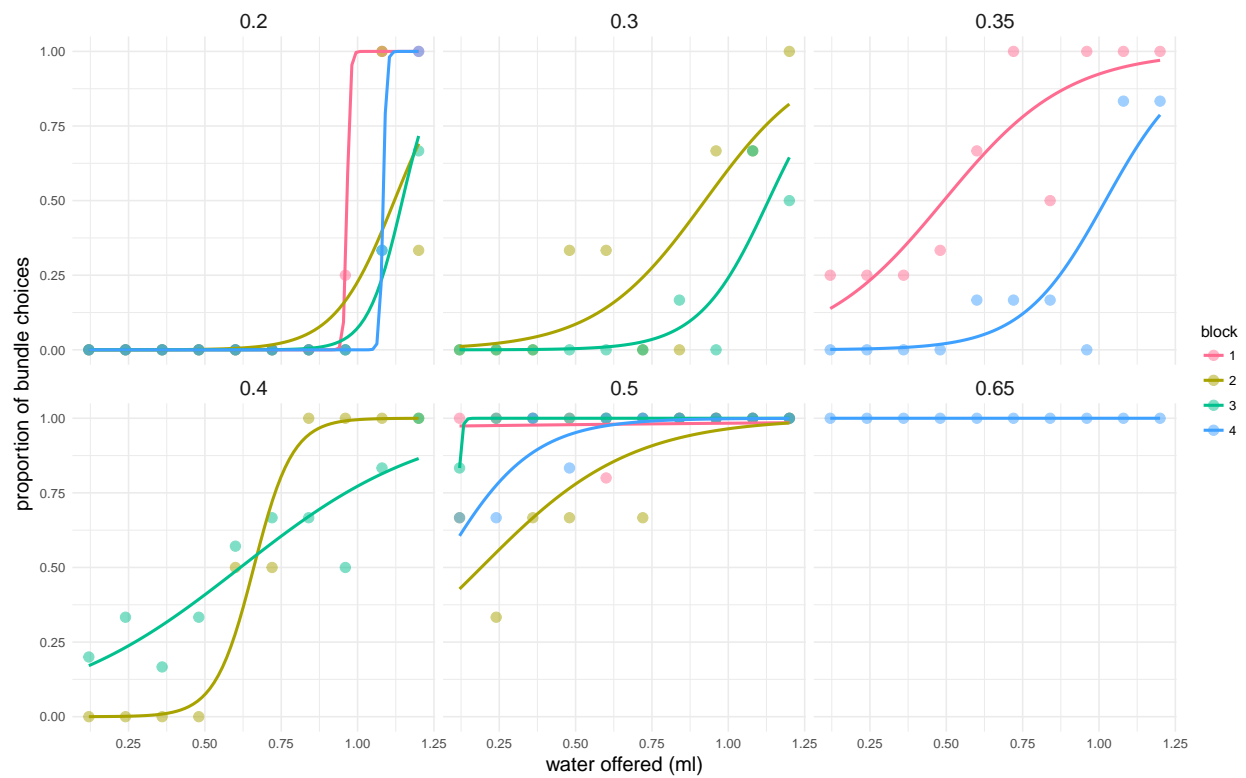
Ulysses : 26-Mar-2018 - 29-Mar-2018



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Monkey Bundle Choice Binoimial Curves

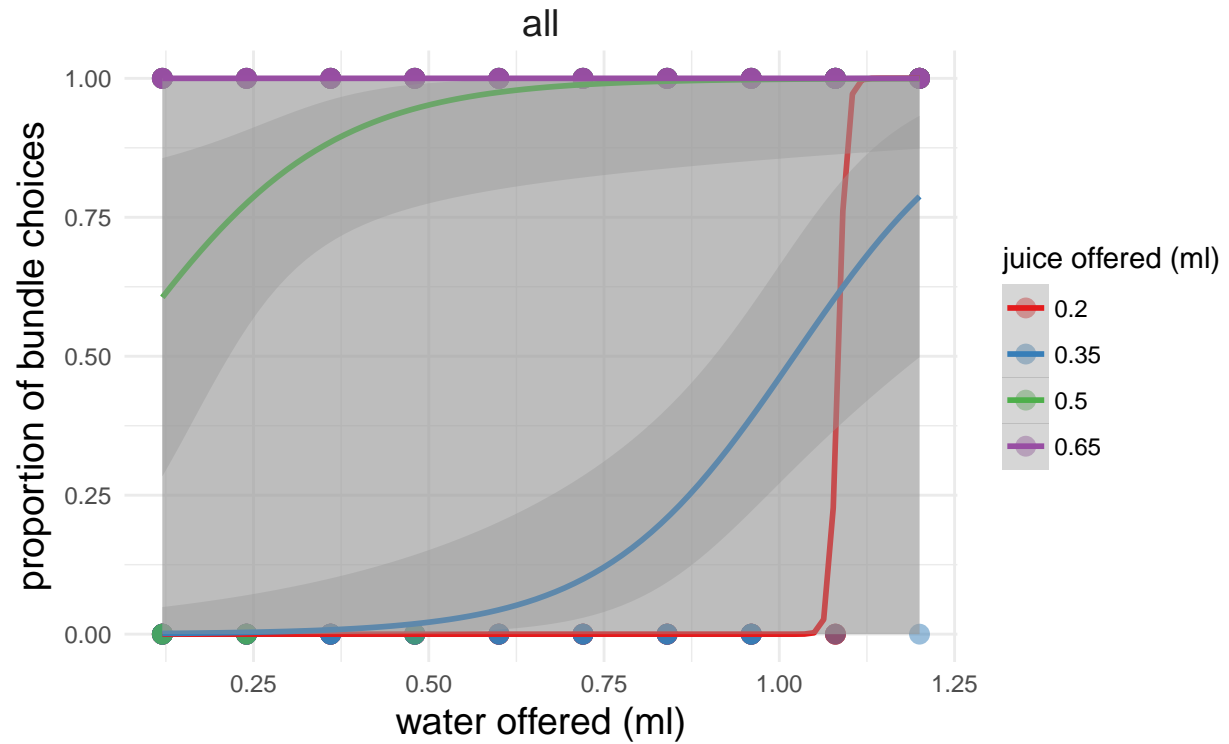
Ulysses : 26-Mar-2018 – 29-Mar-2018



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Today's Monkey Bundle Choice Binoimial Curves

Ulysses : 29-Mar-2018

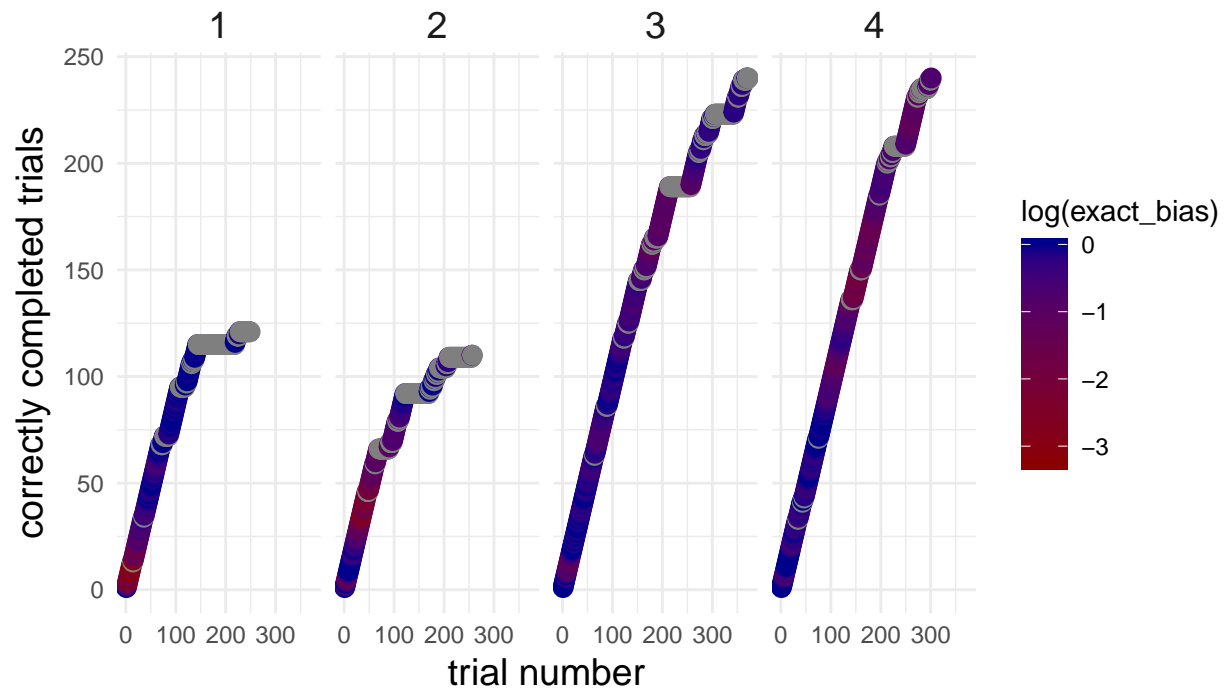


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Monkey Trial Progression and Bias

Ulysses : 26-Mar-2018 – 29-Mar-2018

2018-03-26 2018-03-27 2018-03-28 2018-03-29



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Monkey Trial Progression and Bias

Ulysses : 26-Mar-2018 – 29-Mar-2018

