

BCb Analysis- Early March

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05 April 2018

```
monkey <- "Vicer"  
today <- "13-September-2018"  
look_back <- "05-September-2018"
```

```
start_trial <- 0  
stop_trial <- 400
```

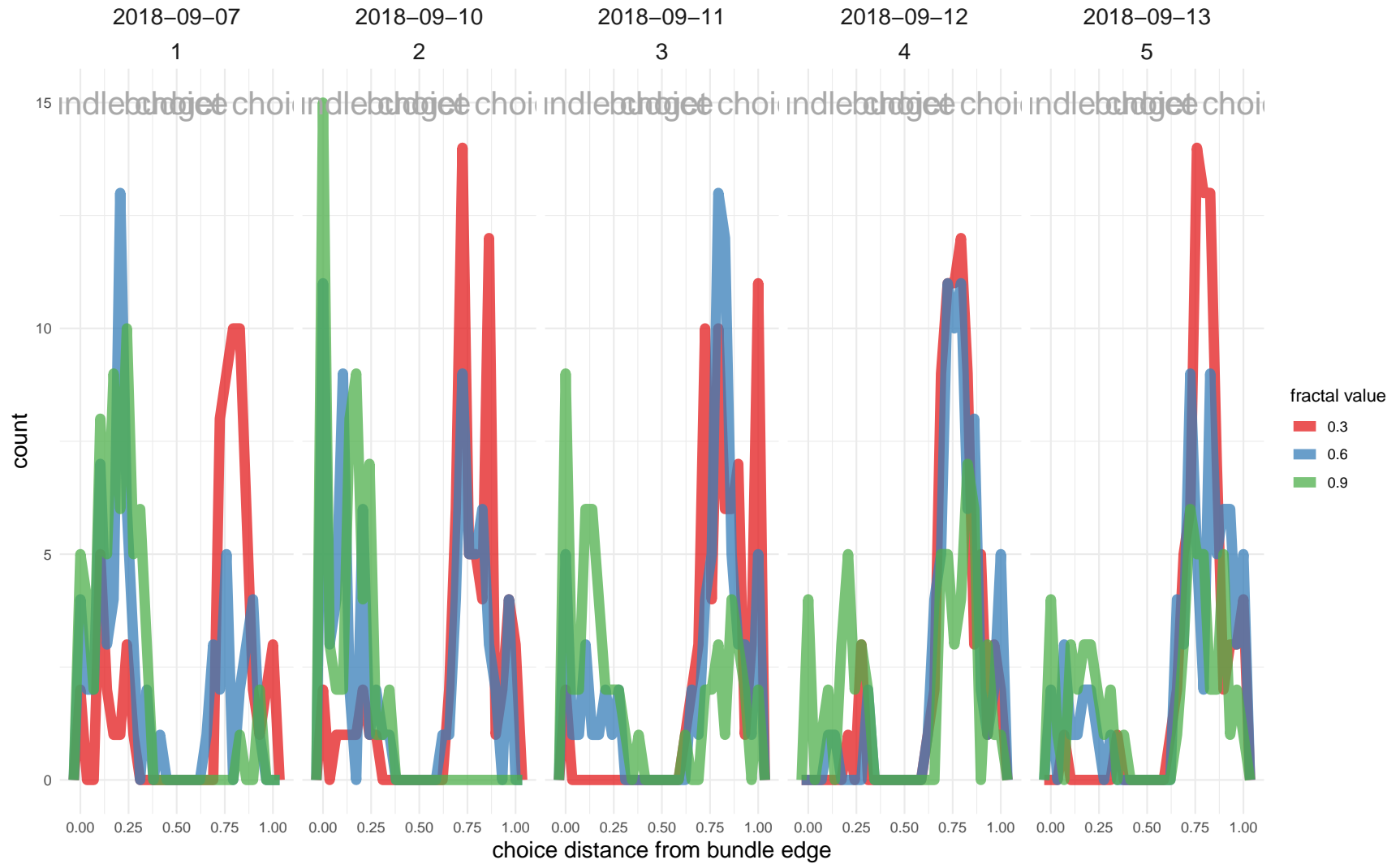
```
merge_days <- TRUE
```

```
#task_data %<>% .[is.na(task_failure), completed_trials := 1:.N, by = "block_no"] %>%  
# .[completed_trials < 181] %>%  
# .[completed_trials > 59]
```

p1

Monkey Choice Distance From Bundle on Binary Choice Task

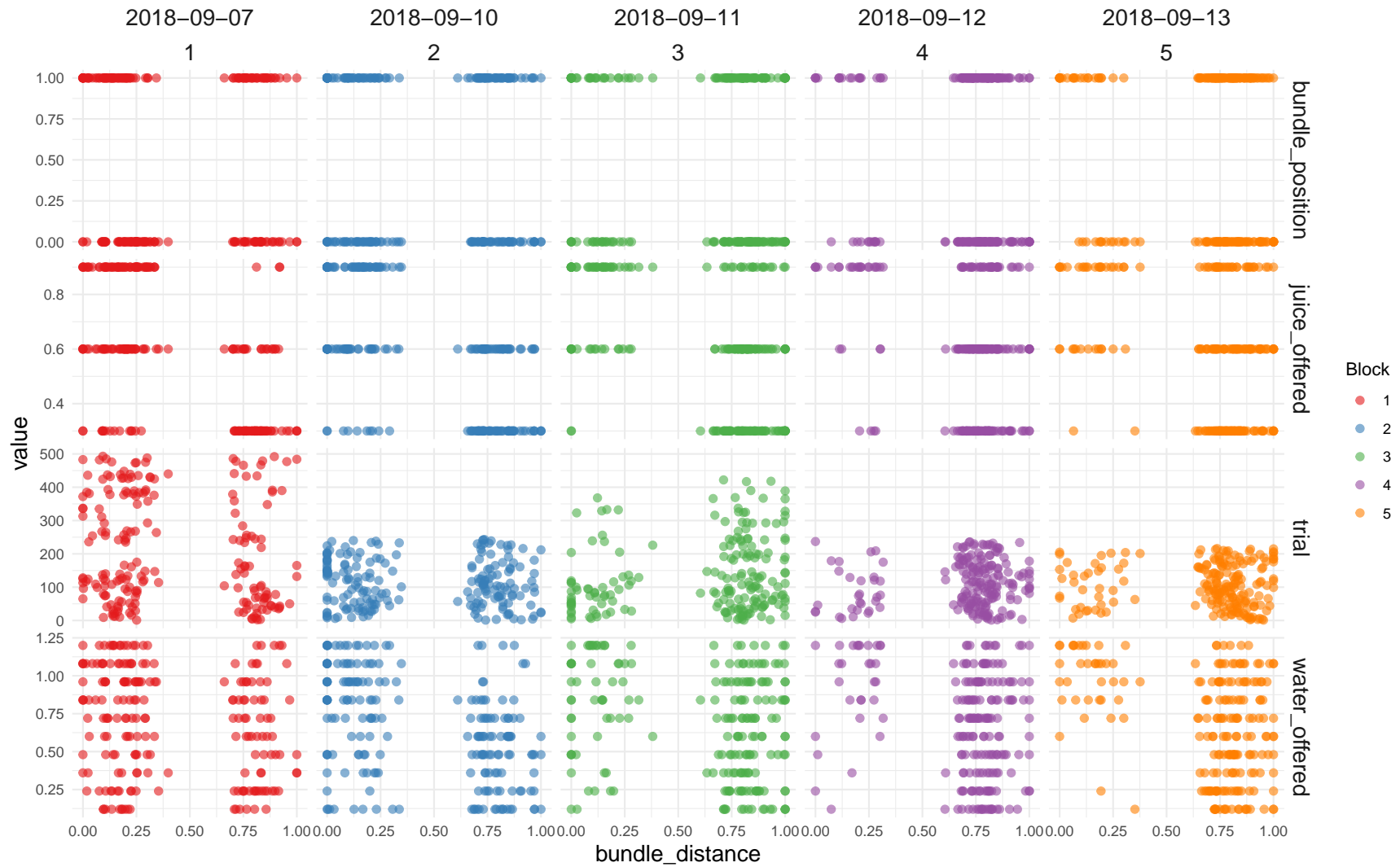
Vicer : 05–September–2018 – 13–September–2018



p2

Monkey Choice Distance From Bundle on Binary Choice Task

Vicer : 05–September–2018 – 13–September–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
## -3.5236  -0.5198  -0.1756   0.4117   3.8882
##
## Coefficients:
##              Estimate Std. Error z value Pr(>|z|)
## (Intercept)   1.228e+04  1.048e+03  11.717  <2e-16 ***
## bundle_position  3.091e-01  1.943e-01   1.591   0.1116
## water_offered   4.080e+00  3.633e-01  11.231  <2e-16 ***
## juice_offered   8.587e+00  6.032e-01  14.236  <2e-16 ***
## trial          -2.350e-03  9.384e-04  -2.505   0.0123 *
## date           -6.912e-01  5.896e-02 -11.722  <2e-16 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for binomial family taken to be 1)
##
##      Null deviance: 1293.87  on 995  degrees of freedom
## Residual deviance:  682.97  on 990  degrees of freedom
##      (640 observations deleted due to missingness)
## AIC: 694.97
##
## 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 = 488, number of trials = 996, p-value =
## 0.5472
## alternative hypothesis: true probability of success is not equal to 0.5
## 95 percent confidence interval:
##  0.4584811 0.5214981
## sample estimates:
## probability of success
##      0.4899598

```

```

#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
## -2.2666  -0.3425  -0.1435  -0.0221   4.6452
##
## Coefficients: (1 not defined because of singularities)
##              Estimate Std. Error z value Pr(>|z|)
## (Intercept)    -12.090458    1.970015  -6.137 8.40e-10 ***
## bundle_position     0.781906    0.561728   1.392 0.163933
## water_offered      7.601928    1.435435   5.296 1.18e-07 ***
## as.factor(juice_offered)0.6  3.091471    0.860159   3.594 0.000326 ***
## as.factor(juice_offered)0.9  5.407780    0.995542   5.432 5.57e-08 ***
## trial            0.006179    0.004335   1.425 0.154102
## 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: 191.556  on 199  degrees of freedom
## Residual deviance:  93.067  on 194  degrees of freedom
##      (16 observations deleted due to missingness)
## AIC: 105.07
##
## Number of Fisher Scoring iterations: 7

```

```

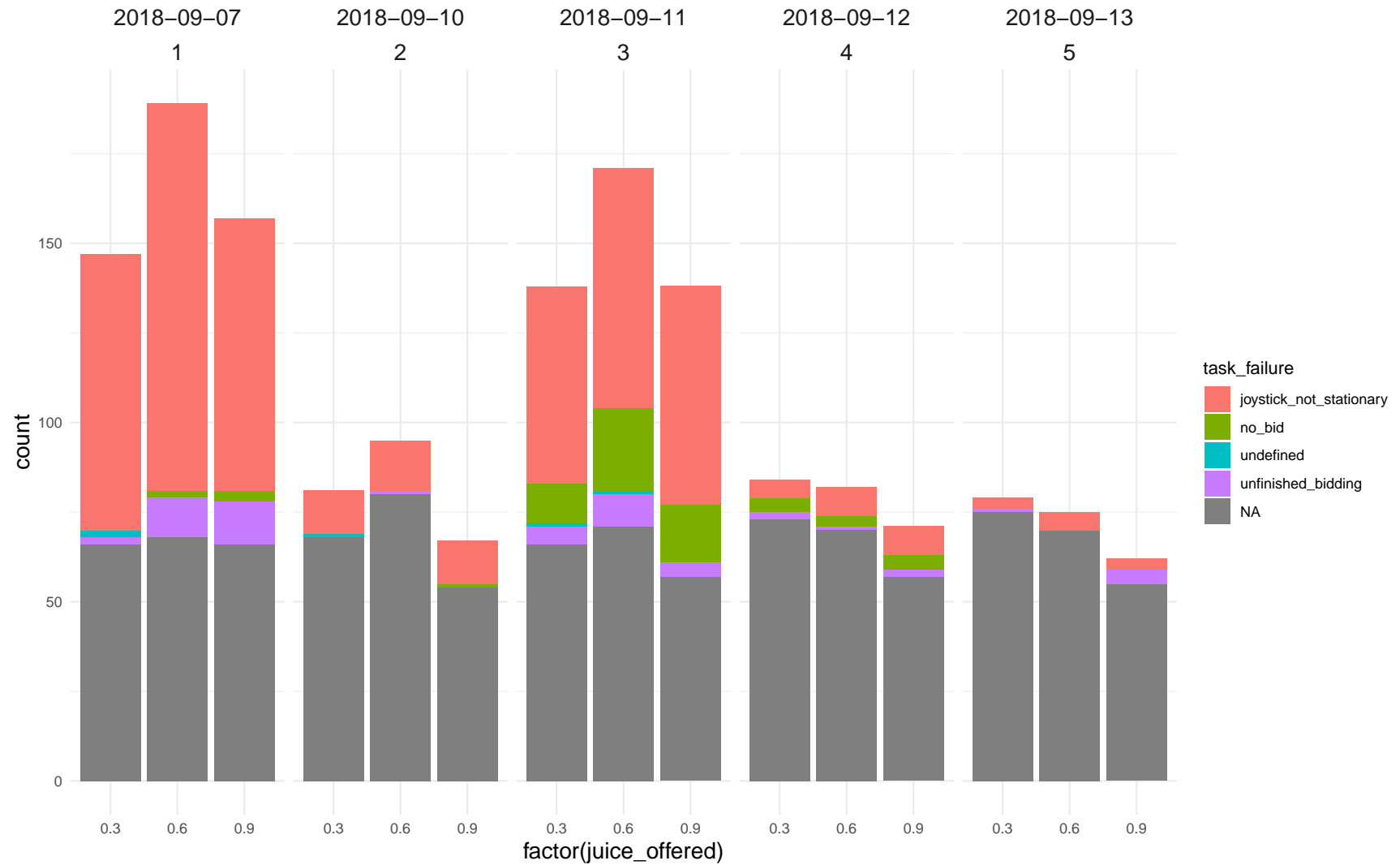
#test for side bias with an exact binomial test
binom.test(c(nrow(task_data %>%
  .[c(bundle_position != fractal_choice & block_no == max(block_no))]),
  nrow(task_data %>%
  .[c(bundle_position == fractal_choice & block_no == max(block_no))])))

##
## 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 = 101, number of trials = 200, p-value =
## 0.9437
## alternative hypothesis: true probability of success is not equal to 0.5
## 95 percent confidence interval:
##  0.4335867 0.5762629
## sample estimates:
## probability of success
##           0.505

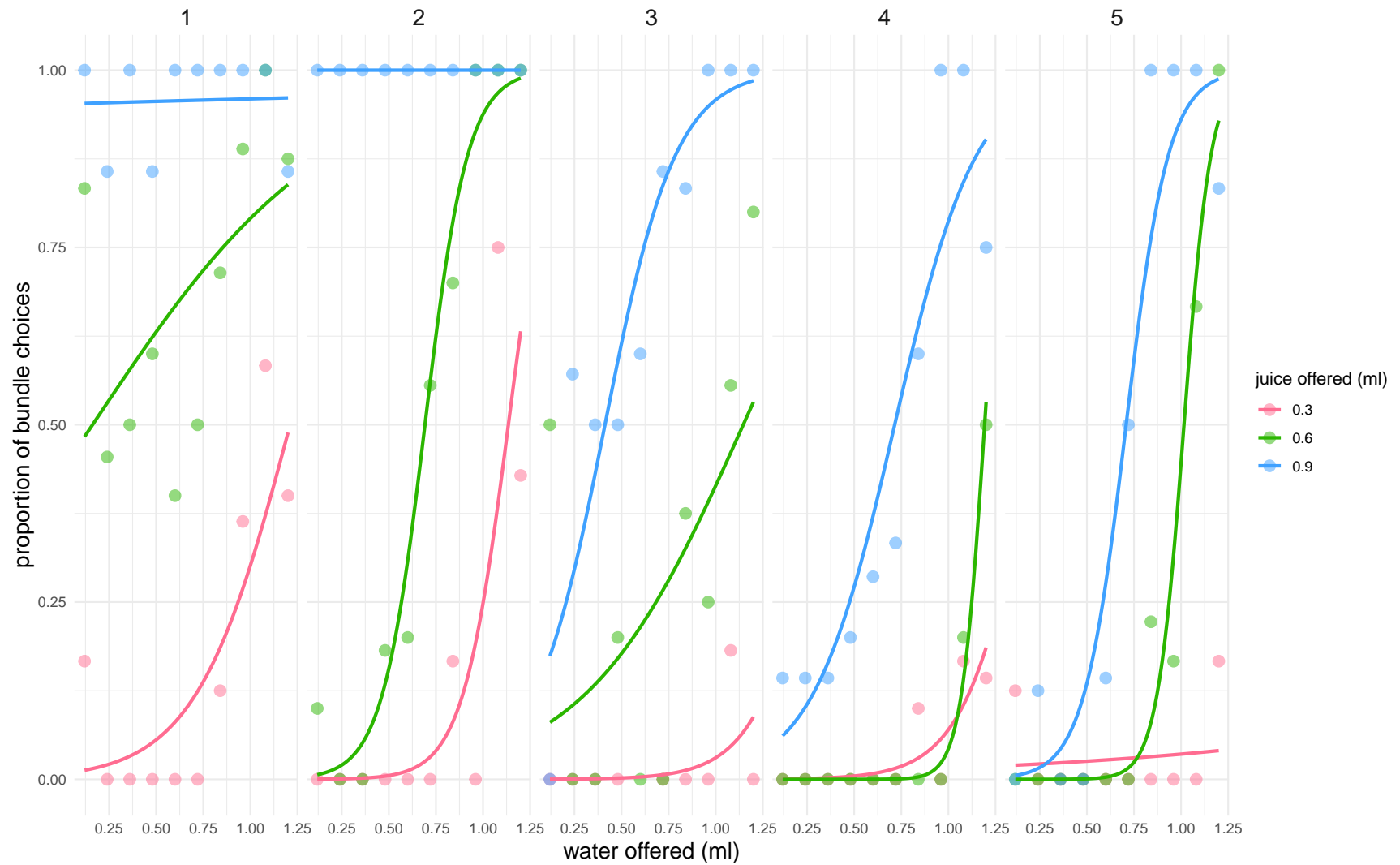
```

Monkey Choice Failures

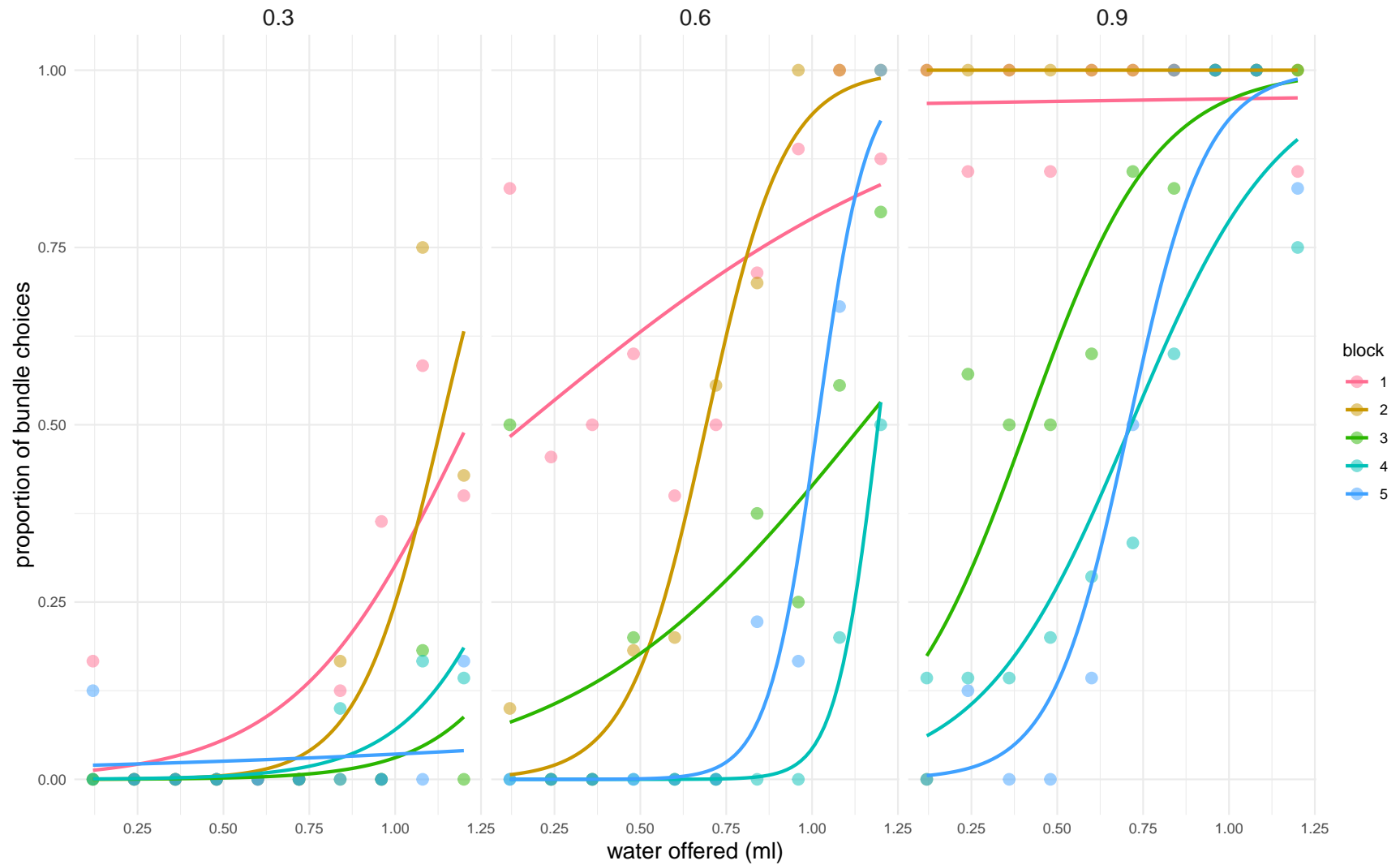
Vicer : 05–September–2018 – 13–September–2018



Monkey Bundle Choice Binoimial Curves
Vicer : 05–September–2018 – 13–September–2018

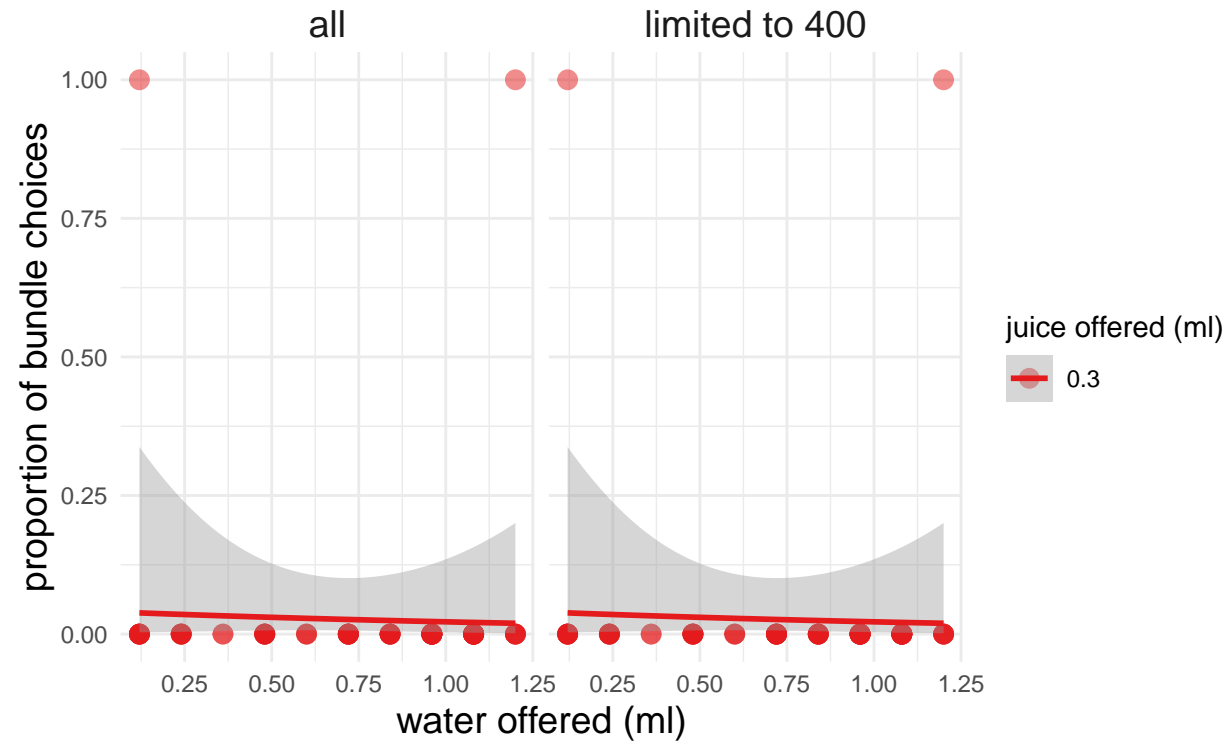


Monkey Bundle Choice Binoimial Curves
Vicer : 05–September–2018 – 13–September–2018



Today's Monkey Bundle Choice Binoimial Curves

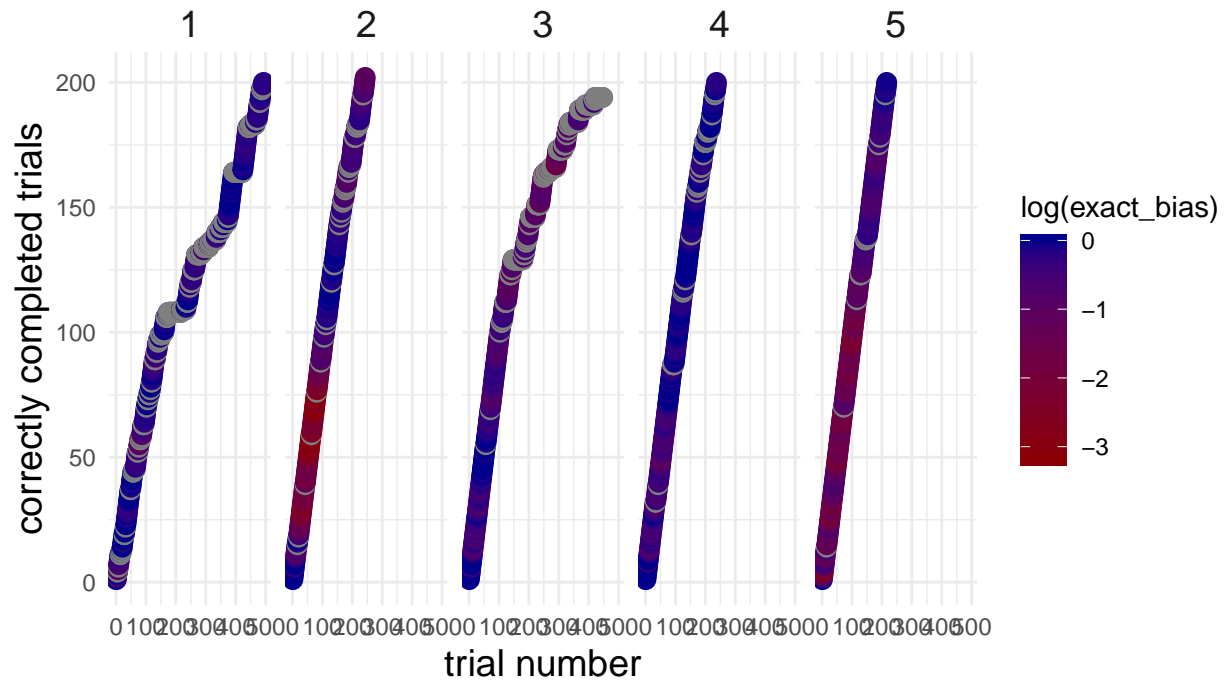
Vicer : 13–September–2018



Monkey Trial Progression and Bias

Vicer : 05–September–2018 – 13–September–2018

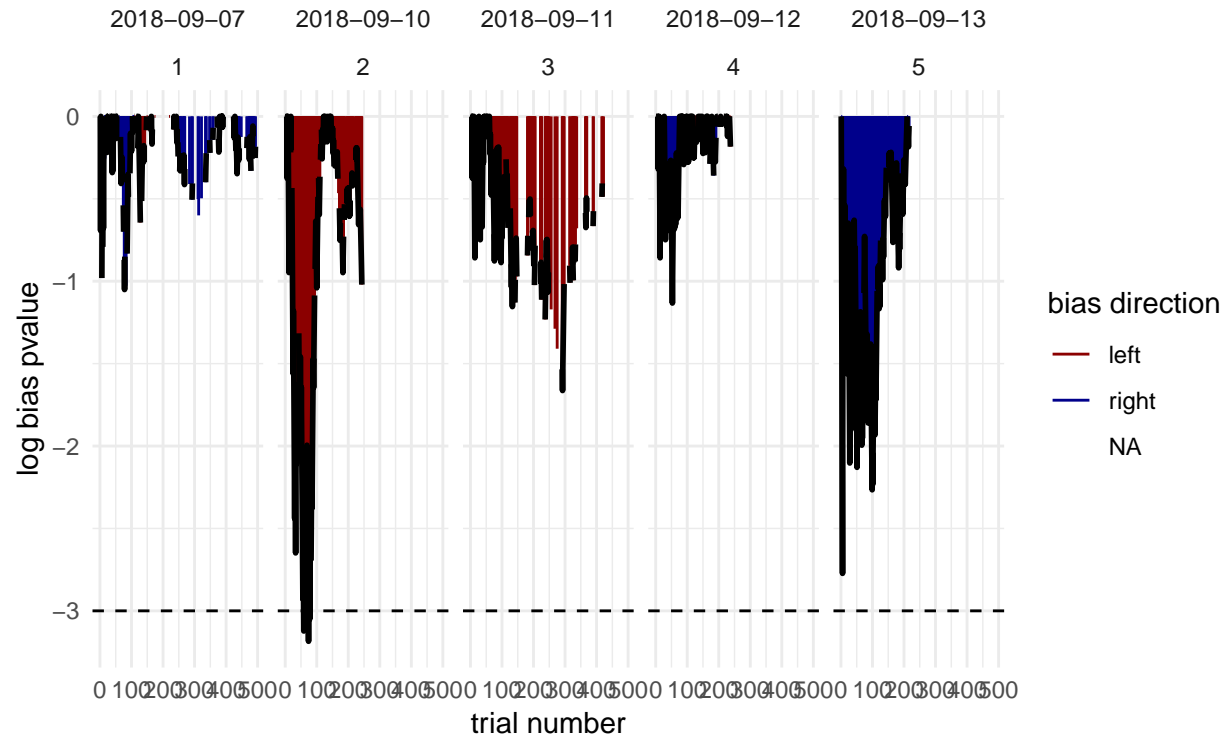
018–09–0 018–09–1 018–09–1 018–09–1 018–09–1



p8

Monkey Trial Progression and Bias

Vicer : 05–September–2018 – 13–September–2018



p9

Pooled Monkey Bundle Choice Binoimial Curves

Vicer : 13–September–2018

