

Exp 2 - Control-Response Analysis

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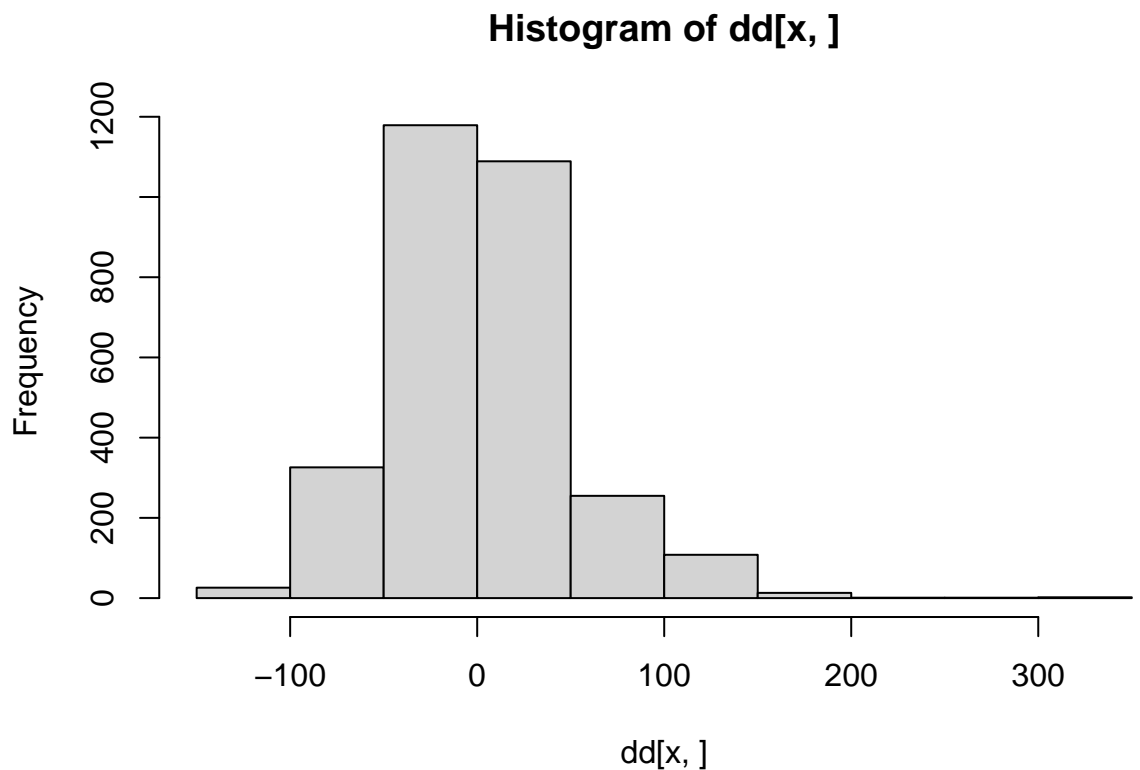
```
## Data: CC2Target
## Models:
## e2.ControlvTarget1: RespRate ~ Phase + ResponseType + (1 | ID)
## e2.ControlvTarget2: RespRate ~ Phase * ResponseType + (1 | ID)
##           npar    AIC    BIC logLik deviance Chisq Df Pr(>Chisq)
## e2.ControlvTarget1     6 47027 47066 -23508    47015
## e2.ControlvTarget2     8 46492 46543 -23238    46476 539.43  2  < 2.2e-16 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

## Linear mixed model fit by maximum likelihood ['lmerMod']
## Formula: RespRate ~ Phase * ResponseType + (1 | ID)
## Data: CC2Target
##
##           AIC          BIC    logLik deviance df.resid
## 46491.8 46542.9 -23237.9 46475.8      4392
##
## Scaled residuals:
##      Min       1Q   Median       3Q      Max
## -3.0463 -0.6917  0.0075  0.4262  6.8982
##
## Random effects:
## Groups Name Variance Std.Dev.
## ID      (Intercept) 798.3 28.25
## Residual 2121.1 46.05
## Number of obs: 4400, groups: ID, 100
##
## Fixed effects:
##
##              Estimate Std. Error t value
## (Intercept)    19.9933    3.0654  6.522
## Phase2         2.3667    2.3783  0.995
## Phase3         0.4817    3.4669  0.139
## ResponseTypeTarget 81.3000    1.6817 48.344
## Phase2:ResponseTypeTarget -74.1500    3.3634 -22.046
## Phase3:ResponseTypeTarget -64.0250    4.9029 -13.059
##
## Correlation of Fixed Effects:
##              (Intr) Phase2 Phase3 RspnTT P2:RTT
## Phase2      -0.194
## Phase3      -0.133 0.171
## RspnsTypTrg -0.274 0.354 0.243
## Phs2:RspnTT 0.137 -0.707 -0.121 -0.500
## Phs3:RspnTT 0.094 -0.121 -0.707 -0.343 0.171
```

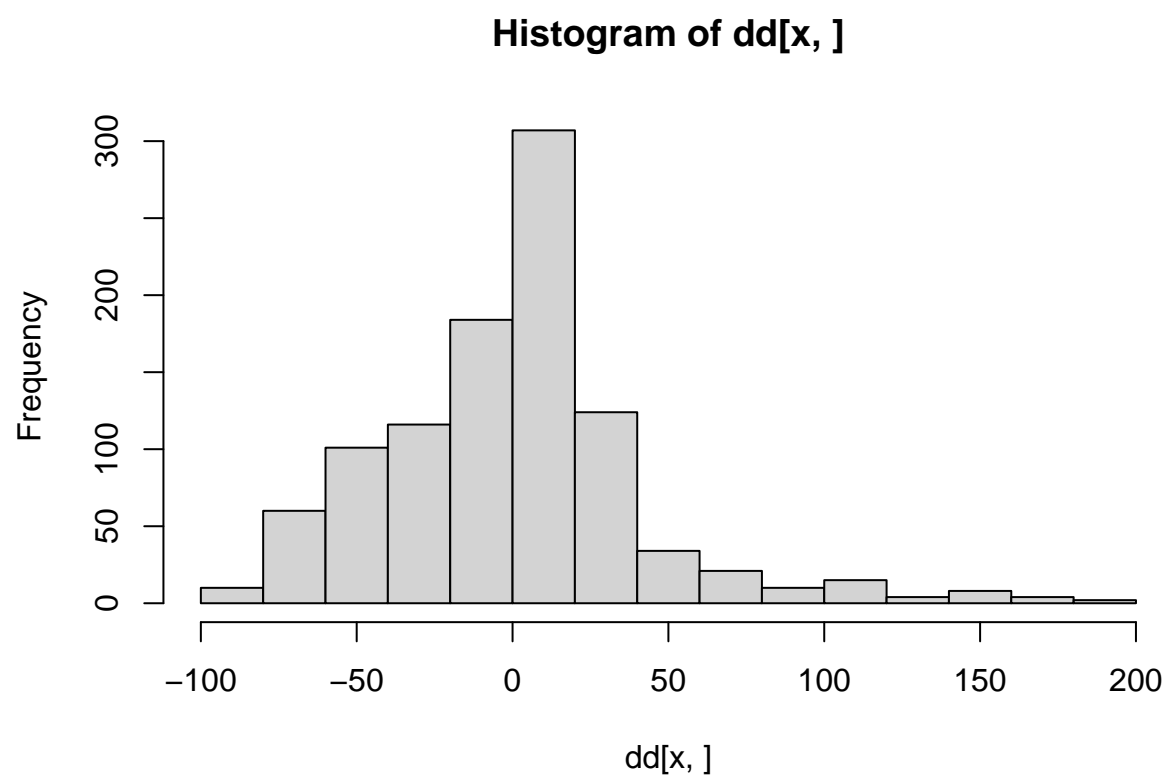
Fixed effects

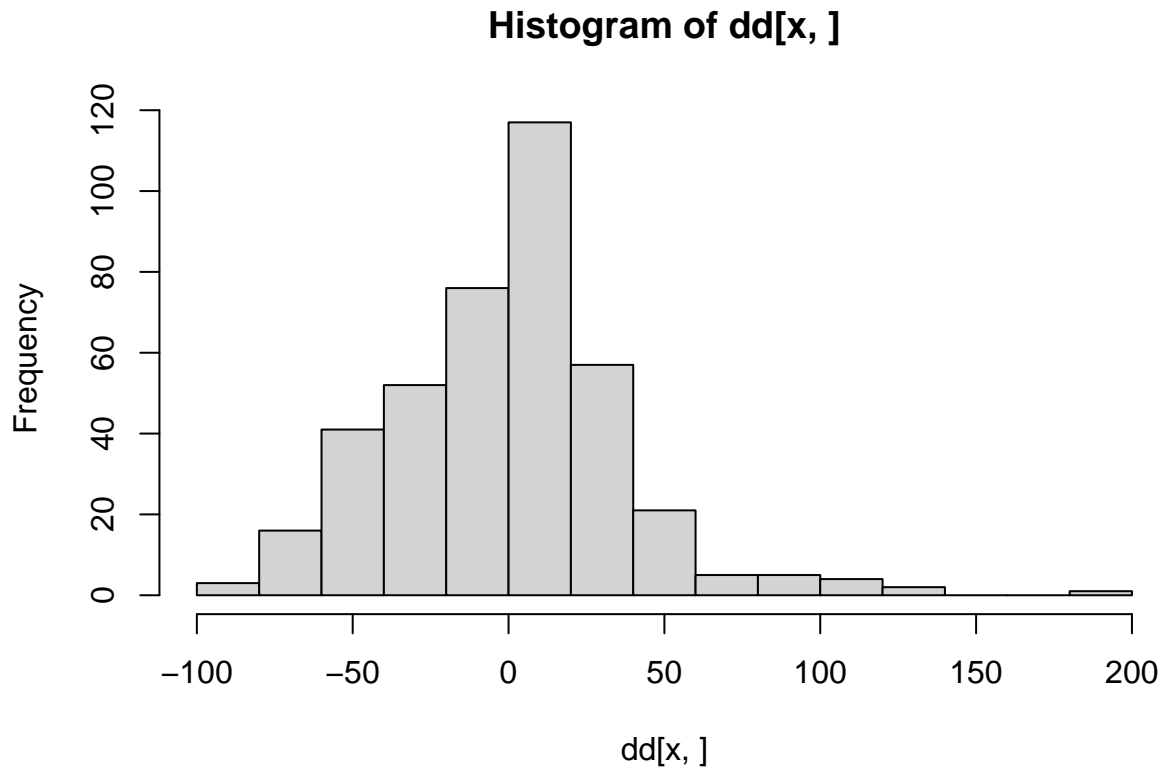
```
## Registered S3 methods overwritten by 'car':
##   method                from
##   influence.merMod       lme4
##   cooks.distance.influence.merMod lme4
##   dfbeta.influence.merMod lme4
##   dfbetas.influence.merMod lme4

## Analysis of Deviance Table (Type II Wald chisquare tests)
##
## Response: RespRate
##               Chisq Df Pr(>Chisq)
## Phase          515.51  2 < 2.2e-16 ***
## ResponseType   1782.54  1 < 2.2e-16 ***
## Phase:ResponseType 574.72  2 < 2.2e-16 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```



Checking residuals



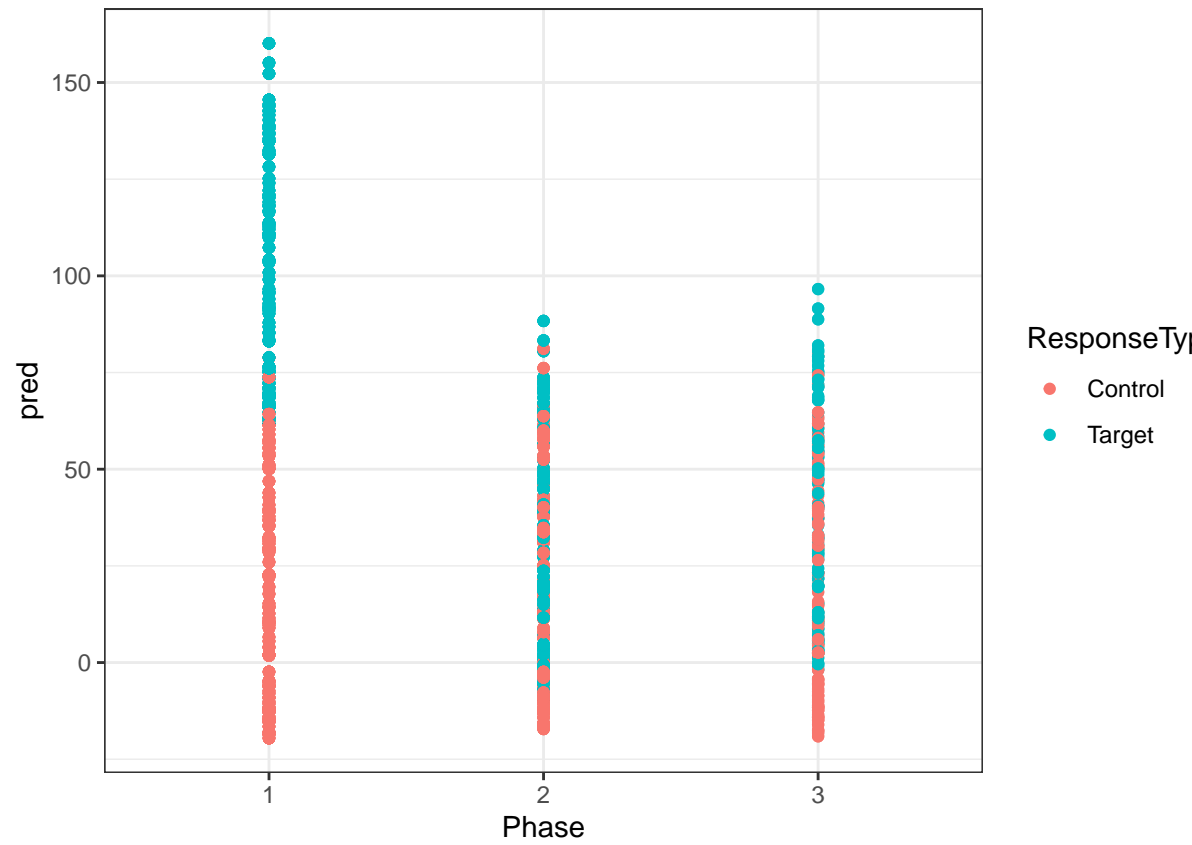


```
## CC2Target$Phase: 1
## $breaks
## [1] -150 -100 -50 0 50 100 150 200 250 300 350
##
## $counts
## [1] 26 326 1179 1089 255 108 13 1 1 2
##
## $density
## [1] 1.733333e-04 2.173333e-03 7.860000e-03 7.260000e-03 1.700000e-03
## [6] 7.200000e-04 8.666667e-05 6.666667e-06 6.666667e-06 1.333333e-05
##
## $mids
## [1] -125 -75 -25 25 75 125 175 225 275 325
##
## $xname
## [1] "dd[x, ]"
##
## $equidist
## [1] TRUE
##
## attr(,"class")
## [1] "histogram"
## -----
## CC2Target$Phase: 2
## $breaks
## [1] -100 -80 -60 -40 -20 0 20 40 60 80 100 120 140 160 180
```

```

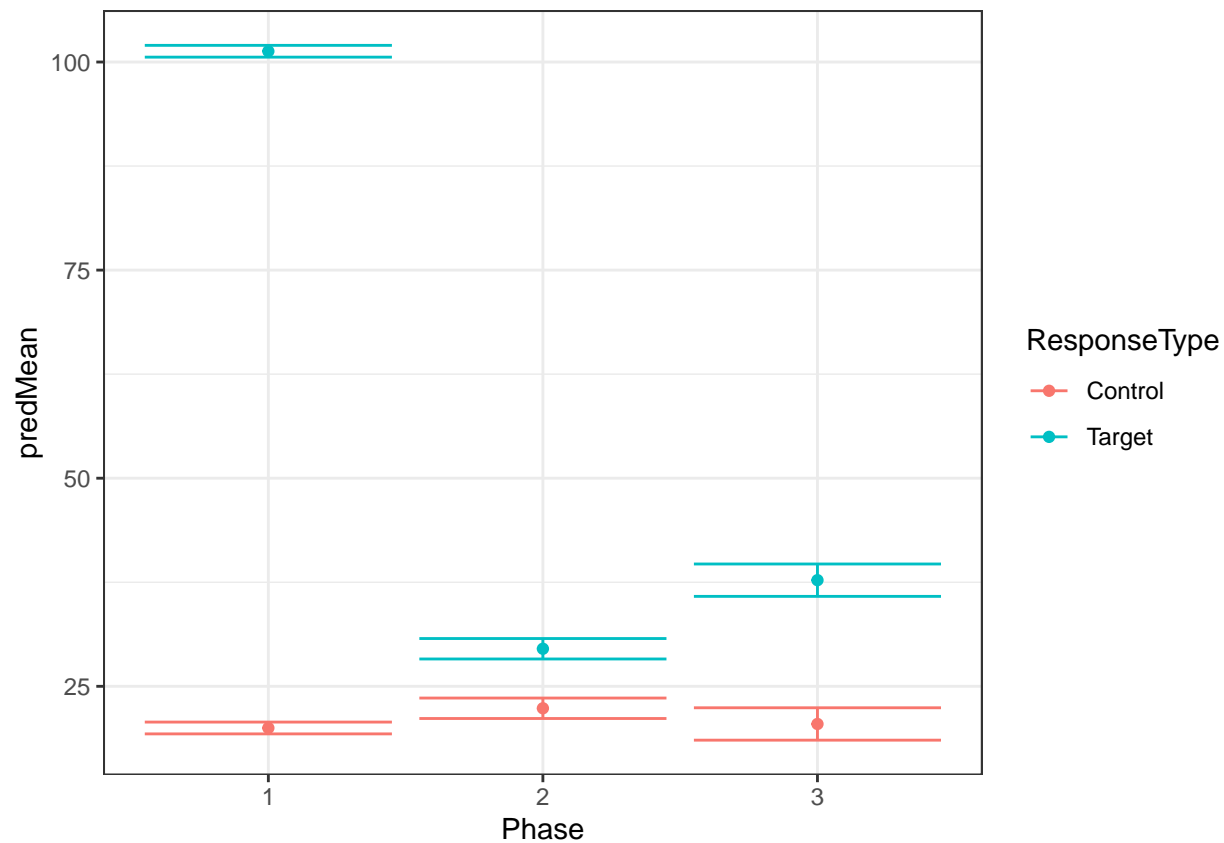
## [16] 200
##
## $counts
## [1] 10 60 101 116 184 307 124 34 21 10 15 4 8 4 2
##
## $density
## [1] 0.00050 0.00300 0.00505 0.00580 0.00920 0.01535 0.00620 0.00170 0.00105
## [10] 0.00050 0.00075 0.00020 0.00040 0.00020 0.00010
##
## $mids
## [1] -90 -70 -50 -30 -10 10 30 50 70 90 110 130 150 170 190
##
## $xname
## [1] "dd[x,]"
##
## $equidist
## [1] TRUE
##
## attr("class")
## [1] "histogram"
## -----
## CC2Target$Phase: 3
## $breaks
## [1] -100 -80 -60 -40 -20 0 20 40 60 80 100 120 140 160 180
## [16] 200
##
## $counts
## [1] 3 16 41 52 76 117 57 21 5 5 4 2 0 0 1
##
## $density
## [1] 0.000375 0.002000 0.005125 0.006500 0.009500 0.014625 0.007125 0.002625
## [9] 0.000625 0.000625 0.000500 0.000250 0.000000 0.000000 0.000125
##
## $mids
## [1] -90 -70 -50 -30 -10 10 30 50 70 90 110 130 150 170 190
##
## $xname
## [1] "dd[x,]"
##
## $equidist
## [1] TRUE
##
## attr("class")
## [1] "histogram"

```



looking @ predictions

'summarise()' has grouped output by 'Phase'. You can override using the '.groups' argument.



specific comparisons

```
## $emmeans
## ResponseType = Control:
## Phase emmean SE df lower.CL upper.CL
## 1 20.0 3.08 124 13.9 26.1
## 2 22.4 3.51 210 15.4 29.3
## 3 20.5 4.32 471 12.0 29.0
##
## ResponseType = Target:
## Phase emmean SE df lower.CL upper.CL
## 1 101.3 3.08 124 95.2 107.4
## 2 29.5 3.51 210 22.6 36.4
## 3 37.8 4.32 471 29.3 46.2
##
## Degrees-of-freedom method: kenward-roger
## Confidence level used: 0.95
##
## $contrasts
## ResponseType = Control:
## contrast estimate SE df t.ratio p.value
## 1 - 2 -2.367 2.38 4305 -0.995 0.3200
## 1 - 3 -0.482 3.47 4305 -0.139 0.8896
## 2 - 3 1.885 3.86 4305 0.489 0.6249
##
## ResponseType = Target:
## contrast estimate SE df t.ratio p.value
```

```

## 1 - 2      71.783 2.38 4305 30.165 <.0001
## 1 - 3      63.543 3.47 4305 18.318 <.0001
## 2 - 3      -8.240 3.86 4305 -2.137 0.0326
##
## Degrees-of-freedom method: kenward-roger

## $emmeans
## Phase = 1:
## ResponseType emmean SE df lower.CL upper.CL
## Control      20.0 3.08 124 13.9 26.1
## Target       101.3 3.08 124 95.2 107.4
##
## Phase = 2:
## ResponseType emmean SE df lower.CL upper.CL
## Control      22.4 3.51 210 15.4 29.3
## Target       29.5 3.51 210 22.6 36.4
##
## Phase = 3:
## ResponseType emmean SE df lower.CL upper.CL
## Control      20.5 4.32 471 12.0 29.0
## Target       37.8 4.32 471 29.3 46.2
##
## Degrees-of-freedom method: kenward-roger
## Confidence level used: 0.95
##
## $contrasts
## Phase = 1:
## contrast      estimate SE df t.ratio p.value
## Control - Target -81.30 1.68 4305 -48.316 <.0001
##
## Phase = 2:
## contrast      estimate SE df t.ratio p.value
## Control - Target -7.15 2.91 4305 -2.453 0.0142
##
## Phase = 3:
## contrast      estimate SE df t.ratio p.value
## Control - Target -17.27 4.61 4305 -3.749 0.0002
##
## Degrees-of-freedom method: kenward-roger

```

Conclusion: No statistically significant increases in control responses from Phase 2 to Phase 3. Control responses remained low throughout the experiment.