

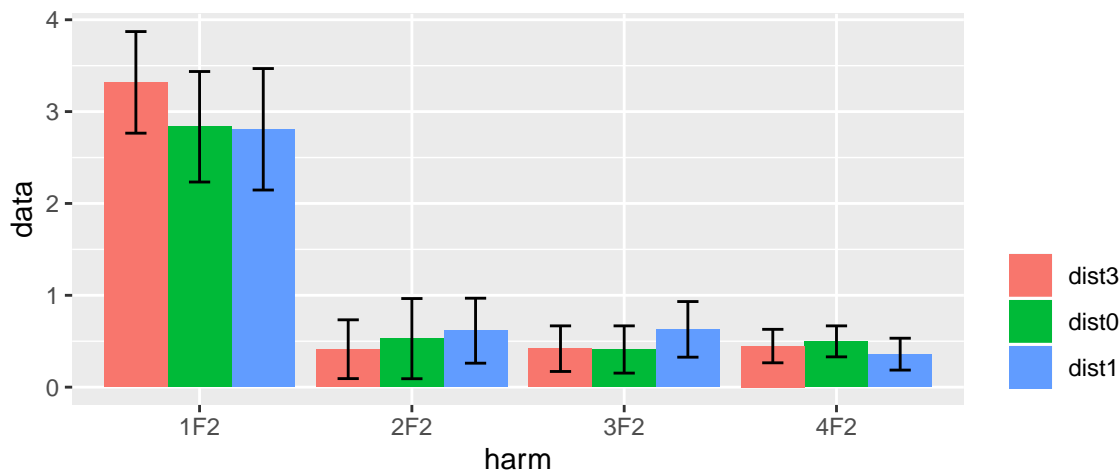
NUMEROSITY ANALYSIS RESULTS

```
top_folder = '/Users/kohler/Google Drive/WRITING/Articles/2019_KohlerNumerositySSVEP/figures/results/experiment2
counter = 0
for (q in c(1,2,3,4)) {
  for (c in c(6,8)) {
    cur_file = switch(q, "RLS_carrier_rc1_carr", "RLS_oddball_rc1_carr", "RLS_carrier_rc2_carr", "RLS_oddball_rc2_carr")
    cur_csv <- sprintf('%s/%s%d_full_projected_all_trials.csv', top_folder, cur_file, c)
    cur_data <- data.frame( read.csv(file = cur_csv) )
    cur_data$cond <- factor(cur_data$condition, levels(cur_data$condition)[c(3,1,2)])
    cur_data$harm <- cur_data$harmonic
    cur_data$harm_alt <- factor(cur_data$harmonic, levels(cur_data$harmonic)[c(2,1,3,4)])
    g <- ggplot(cur_data, aes(harm, data, fill = cond)) +
      stat_summary(geom = "bar", fun.y = mean, position=position_dodge()) +
      stat_summary(geom = "errorbar", fun.data = mean_se, width=.3, position=position_dodge(.9))
    g <- g + theme(legend.title=element_blank(),
      legend.justification=c(1,0),
      legend.background = element_blank()) +
      ggtitle(toupper(sprintf('%s%d\n', cur_file, c)))
    if (q == 1 && c == 6) {
      cat("RESULTS BELOW\n ")
    }
    print(g)
    m1 <- lmer(data ~ cond * harm + (1|subject), cur_data)
    emm = emmeans(m1, ~ cond * harm, lmer.df = "satterthwaite")
    m2 <- lmer(data ~ cond + harm + (1|subject), cur_data)
    if (isSingular(m1)) {
      if (isSingular(m2)) {
        cat("WARNING: BOTH MODELS ARE SINGULAR! ")
      } else {
        cat("WARNING: MODEL1 IS SINGULAR, BUT MODEL2 IS NOT! ")
      }
    } else {
      cat("LOVELY: NONE OF THE MODELS ARE SINGULAR! ")
    }
    if (!performance::check_convergence(m1)) {
      if (!performance::check_convergence(m2)) {
        cat("WARNING: BOTH MODELS DID NOT CONVERGE!\n\n")
      } else {
        cat("WARNING: MODEL1 DID NOT CONVERGE, BUT MODEL2 DID!\n\n")
      }
    } else {
      cat("LOVELY: BOTH MODELS CONVERGED!\n\n")
    }
    cat("ANOVA TEST FOR MAIN EFFECTS AND INTERACTIONS\n")
    print(anova(m1), type='pdf')
    write.csv(as.matrix(anova(m1)),
      file = sprintf('%s/%s%d_full_projected_all_results.csv', top_folder, cur_file, c), na = "")
    cat("\nESTIMATED MARGINAL MEANS, SIMPLE MAIN EFFECTS OF CONDITION WITHOUT CORRECTION\n")
    print(prettify(summary(pairs(emm, simple = "cond", adjust = "none"))))
    cat("\nTEST OF WHETHER OR NOT THE INTERACTION PROVIDES A BETTER FIT\n")
    print(anova(m1, m2))
  }
}
```

RESULTS BELOW

##

RLS_CARRIER_RC1_CARR6



LOVELY: NONE OF THE MODELS ARE SINGULAR! LOVELY: BOTH MODELS CONVERGED!

##

ANOVA TEST FOR MAIN EFFECTS AND INTERACTIONS

Type III Analysis of Variance Table with Satterthwaite's method

	Sum Sq	Mean Sq	NumDF	DenDF	F value	Pr(>F)
cond	0.200	0.100	2	154	0.0836	0.9199
harm	212.272	70.757	3	154	59.0031	<2e-16 ***
cond:harm	3.195	0.532	6	154	0.4440	0.8484

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

##

ESTIMATED MARGINAL MEANS, SIMPLE MAIN EFFECTS OF CONDITION WITHOUT CORRECTION

	contrast	harm	estimate	SE	df	t.ratio	p.value
## 1	1 dist3 - dist0	1F2	0.483408479	0.3998686	154	1.2089183	0.2285471
## 2	2 dist3 - dist1	1F2	0.511023124	0.3998686	154	1.2779776	0.2031803
## 3	3 dist0 - dist1	1F2	0.027614645	0.3998686	154	0.0690593	0.9450320
## 4	4 dist3 - dist0	2F2	-0.115142962	0.3998686	154	-0.2879520	0.7737706
## 5	5 dist3 - dist1	2F2	-0.201511228	0.3998686	154	-0.5039436	0.6150212
## 6	6 dist0 - dist1	2F2	-0.086368266	0.3998686	154	-0.2159916	0.8292801
## 7	7 dist3 - dist0	3F2	0.008830539	0.3998686	154	0.0220836	0.9824099
## 8	8 dist3 - dist1	3F2	-0.210272088	0.3998686	154	-0.5258530	0.5997468
## 9	9 dist0 - dist1	3F2	-0.219102627	0.3998686	154	-0.5479366	0.5845288
## 10	10 dist3 - dist0	4F2	-0.051334788	0.3998686	154	-0.1283791	0.8980164
## 11	11 dist3 - dist1	4F2	0.087975623	0.3998686	154	0.2200113	0.8261537
## 12	12 dist0 - dist1	4F2	0.139310411	0.3998686	154	0.3483905	0.7280228

##

TEST OF WHETHER OR NOT THE INTERACTION PROVIDES A BETTER FIT

Data: cur_data

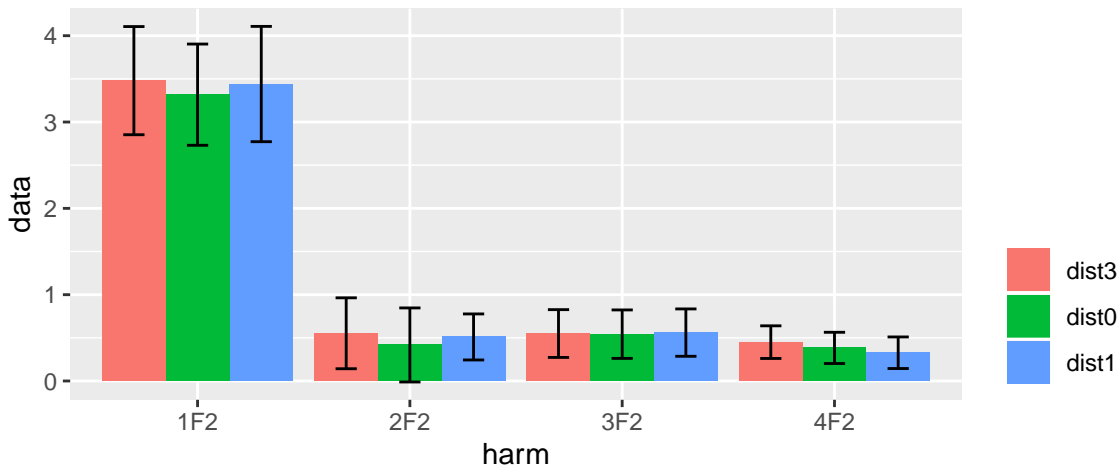
Models:

m2: data ~ cond + harm + (1 | subject)

m1: data ~ cond * harm + (1 | subject)

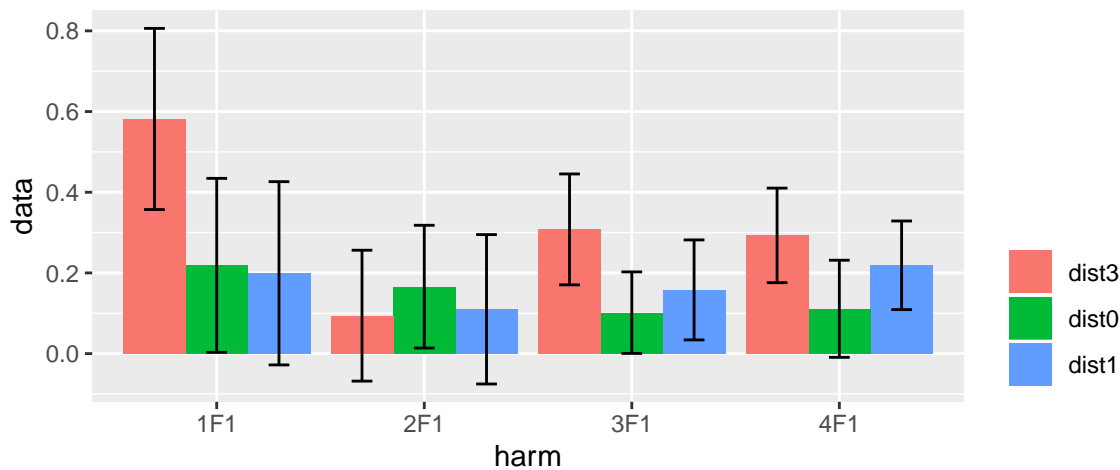
	Df	AIC	BIC	logLik	deviance	Chisq	Chi	Df	Pr(>Chisq)
## m2	8	587.13	612.67	-285.56	571.13				
## m1	14	596.30	641.00	-284.15	568.30	2.8297		6	0.8299

RLS_CARRIER_RC1_CARR8



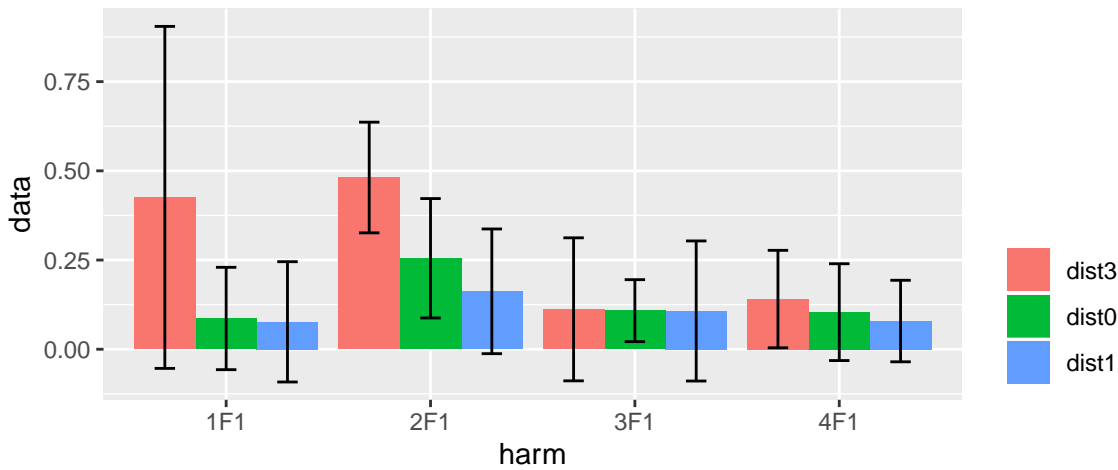
```
## LOVELY: NONE OF THE MODELS ARE SINGULAR! LOVELY: BOTH MODELS CONVERGED!
##
## ANOVA TEST FOR MAIN EFFECTS AND INTERACTIONS
## Type III Analysis of Variance Table with Satterthwaite's method
##          Sum Sq Mean Sq NumDF DenDF F value Pr(>F)
## cond          0.259   0.129     2   154   0.1019 0.9032
## harm        291.384  97.128     3   154  76.4352 <2e-16 ***
## cond:harm    0.216   0.036     6   154   0.0283 0.9999
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## ESTIMATED MARGINAL MEANS, SIMPLE MAIN EFFECTS OF CONDITION WITHOUT CORRECTION
##          contrast harm estimate SE df t.ratio p.value
## 1 1 dist3 - dist0 1F2 0.16243456 0.4116188 154 0.39462378 0.6936665
## 2 2 dist3 - dist1 1F2 0.03923738 0.4116188 154 0.09532456 0.9241810
## 3 3 dist0 - dist1 1F2 -0.12319718 0.4116188 154 -0.29929922 0.7651153
## 4 4 dist3 - dist0 2F2 0.13536052 0.4116188 154 0.32884923 0.7427164
## 5 5 dist3 - dist1 2F2 0.04253188 0.4116188 154 0.10332832 0.9178368
## 6 6 dist0 - dist1 2F2 -0.09282864 0.4116188 154 -0.22552091 0.8218731
## 7 7 dist3 - dist0 3F2 0.00724635 0.4116188 154 0.01760452 0.9859771
## 8 8 dist3 - dist1 3F2 -0.01072360 0.4116188 154 -0.02605225 0.9792494
## 9 9 dist0 - dist1 3F2 -0.01796995 0.4116188 154 -0.04365677 0.9652346
## 10 10 dist3 - dist0 4F2 0.06643266 0.4116188 154 0.16139367 0.8719949
## 11 11 dist3 - dist1 4F2 0.12264568 0.4116188 154 0.29795939 0.7661358
## 12 12 dist0 - dist1 4F2 0.05621301 0.4116188 154 0.13656572 0.8915524
##
## TEST OF WHETHER OR NOT THE INTERACTION PROVIDES A BETTER FIT
## Data: cur_data
## Models:
## m2: data ~ cond + harm + (1 | subject)
## m1: data ~ cond * harm + (1 | subject)
##      Df      AIC      BIC logLik deviance Chisq Chi Df Pr(>Chisq)
## m2   8 594.84 620.38 -289.42  578.84
## m1  14 606.66 651.36 -289.33  578.66 0.1819     6    0.9999
```

RLS_ODDBALL_RC1_CARR6



```
## WARNING: BOTH MODELS ARE SINGULAR! LOVELY: BOTH MODELS CONVERGED!
##
## ANOVA TEST FOR MAIN EFFECTS AND INTERACTIONS
## Type III Analysis of Variance Table with Satterthwaite's method
##      Sum Sq Mean Sq NumDF DenDF F value Pr(>F)
## cond      1.0229  0.51147      2    168   1.2910  0.2777
## harm      1.0389  0.34631      3    168   0.8741  0.4558
## cond:harm  1.0040  0.16734      6    168   0.4224  0.8634
##
## ESTIMATED MARGINAL MEANS, SIMPLE MAIN EFFECTS OF CONDITION WITHOUT CORRECTION
##      contrast harm estimate      SE df    t.ratio    p.value
## 1  1 dist3 - dist0 1F1  0.36281403 0.2298395 168  1.57855387 0.11631947
## 2  2 dist3 - dist1 1F1  0.38249444 0.2298395 168  1.66418061 0.09794042
## 3  3 dist0 - dist1 1F1  0.01968041 0.2298395 168  0.08562675 0.93186510
## 4  4 dist3 - dist0 2F1 -0.07189634 0.2298395 168 -0.31281108 0.75481217
## 5  5 dist3 - dist1 2F1 -0.01579473 0.2298395 168 -0.06872072 0.94529365
## 6  6 dist0 - dist1 2F1  0.05610161 0.2298395 168  0.24409036 0.80745867
## 7  7 dist3 - dist0 3F1  0.20638085 0.2298395 168  0.89793466 0.37050544
## 8  8 dist3 - dist1 3F1  0.15005721 0.2298395 168  0.65287824 0.51472720
## 9  9 dist0 - dist1 3F1 -0.05632364 0.2298395 168 -0.24505642 0.80671183
## 10 10 dist3 - dist0 4F1  0.18194455 0.2298395 168  0.79161568 0.42970106
## 11 11 dist3 - dist1 4F1  0.07415749 0.2298395 168  0.32264904 0.74736224
## 12 12 dist0 - dist1 4F1 -0.10778706 0.2298395 168 -0.46896664 0.63970148
##
## TEST OF WHETHER OR NOT THE INTERACTION PROVIDES A BETTER FIT
## Data: cur_data
## Models:
## m2: data ~ cond + harm + (1 | subject)
## m1: data ~ cond * harm + (1 | subject)
##      Df      AIC      BIC logLik deviance Chisq Chi Df Pr(>Chisq)
## m2   8 350.44 375.99 -167.22  334.44
## m1  14 359.75 404.45 -165.87  331.75 2.6949      6    0.8461
```

RLS_ODDBALL_RC1_CARR8



```
## LOVELY: NONE OF THE MODELS ARE SINGULAR! LOVELY: BOTH MODELS CONVERGED!
##
## ANOVA TEST FOR MAIN EFFECTS AND INTERACTIONS
## Type III Analysis of Variance Table with Satterthwaite's method
##
```

	Sum Sq	Mean Sq	NumDF	DenDF	F value	Pr(>F)
cond	1.15285	0.57642	2	154	0.9350	0.3948
harm	1.11449	0.37150	3	154	0.6026	0.6143
cond:harm	0.86813	0.14469	6	154	0.2347	0.9646

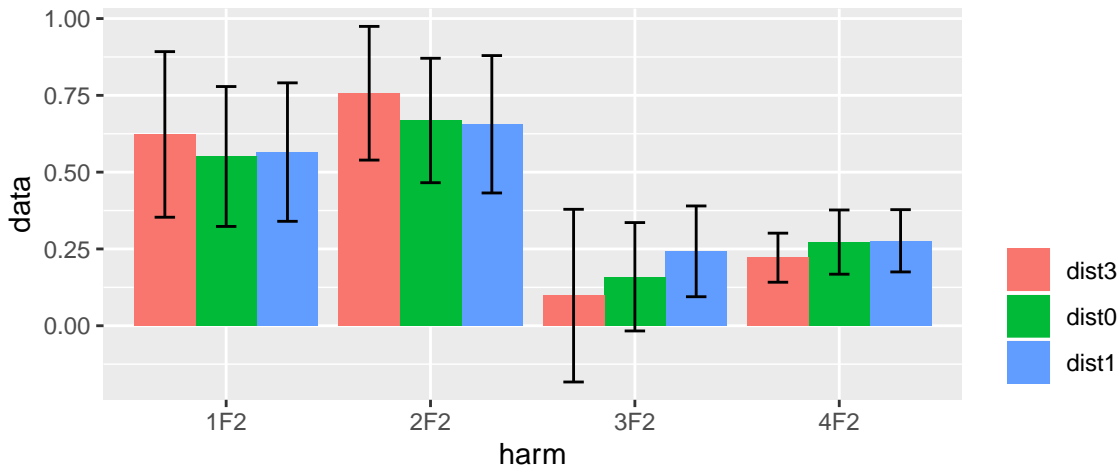
```
##
## ESTIMATED MARGINAL MEANS, SIMPLE MAIN EFFECTS OF CONDITION WITHOUT CORRECTION
##
```

	contrast	harm	estimate	SE	df	t.ratio	p.value
## 1	1 dist3 - dist0	1F1	0.339370833	0.2867019	154	1.183706287	0.2383533
## 2	2 dist3 - dist1	1F1	0.348783386	0.2867019	154	1.216536739	0.2256418
## 3	3 dist0 - dist1	1F1	0.009412553	0.2867019	154	0.032830452	0.9738523
## 4	4 dist3 - dist0	2F1	0.226285282	0.2867019	154	0.789270275	0.4311675
## 5	5 dist3 - dist1	2F1	0.318910055	0.2867019	154	1.112340250	0.2677253
## 6	6 dist0 - dist1	2F1	0.092624773	0.2867019	154	0.323069974	0.7470805
## 7	7 dist3 - dist0	3F1	0.003657791	0.2867019	154	0.012758167	0.9898372
## 8	8 dist3 - dist1	3F1	0.004662915	0.2867019	154	0.016263984	0.9870448
## 9	9 dist0 - dist1	3F1	0.001005124	0.2867019	154	0.003505817	0.9972073
## 10	10 dist3 - dist0	4F1	0.036675729	0.2867019	154	0.127922871	0.8983769
## 11	11 dist3 - dist1	4F1	0.061591786	0.2867019	154	0.214828670	0.8301851
## 12	12 dist0 - dist1	4F1	0.024916057	0.2867019	154	0.086905799	0.9308593

```
##
## TEST OF WHETHER OR NOT THE INTERACTION PROVIDES A BETTER FIT
## Data: cur_data
## Models:
## m2: data ~ cond + harm + (1 | subject)
## m1: data ~ cond * harm + (1 | subject)
##
```

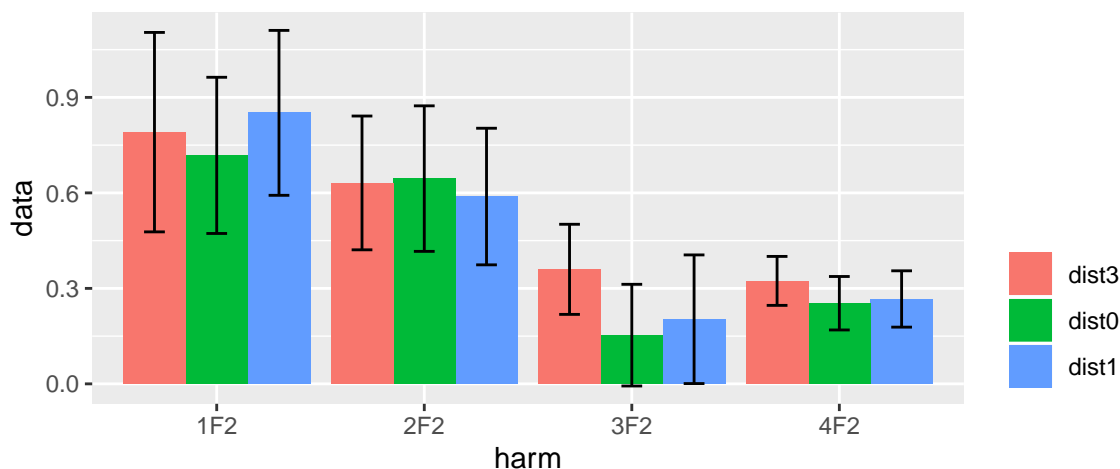
	Df	AIC	BIC	logLik	deviance	Chisq	Chi Df	Pr(>Chisq)
## m2	8	430.35	455.90	-207.18	414.35			
## m1	14	440.85	485.55	-206.43	412.85	1.5019	6	0.9594

RLS_CARRIER_RC2_CARR6



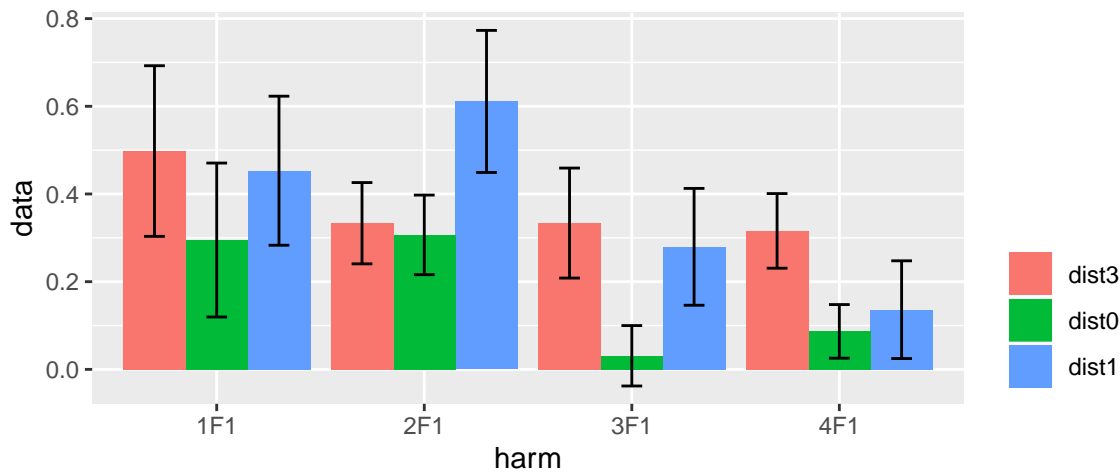
```
## LOVELY: NONE OF THE MODELS ARE SINGULAR! LOVELY: BOTH MODELS CONVERGED!
##
## ANOVA TEST FOR MAIN EFFECTS AND INTERACTIONS
## Type III Analysis of Variance Table with Satterthwaite's method
##      Sum Sq Mean Sq NumDF DenDF F value    Pr(>F)
## cond      0.0149  0.00744      2    154   0.0167  0.9834820
## harm      8.6039  2.86798      3    154   6.4250  0.0003964 ***
## cond:harm  0.3049  0.05081      6    154   0.1138  0.9946983
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## ESTIMATED MARGINAL MEANS, SIMPLE MAIN EFFECTS OF CONDITION WITHOUT CORRECTION
##      contrast harm      estimate      SE df    t.ratio    p.value
## 1  1 dist3 - dist0  1F2  0.071828598  0.2439611 154   0.29442649  0.7688285
## 2  2 dist3 - dist1  1F2  0.057472989  0.2439611 154   0.23558264  0.8140695
## 3  3 dist0 - dist1  1F2 -0.014355609  0.2439611 154  -0.05884385  0.9531528
## 4  4 dist3 - dist0  2F2  0.088687900  0.2439611 154   0.36353302  0.7167053
## 5  5 dist3 - dist1  2F2  0.101003648  0.2439611 154   0.41401545  0.6794388
## 6  6 dist0 - dist1  2F2  0.012315747  0.2439611 154   0.05048243  0.9598034
## 7  7 dist3 - dist0  3F2 -0.061486917  0.2439611 154  -0.25203579  0.8013495
## 8  8 dist3 - dist1  3F2 -0.144248607  0.2439611 154  -0.59127718  0.5552016
## 9  9 dist0 - dist1  3F2 -0.082761689  0.2439611 154  -0.33924139  0.7348901
## 10 10 dist3 - dist0  4F2 -0.050824245  0.2439611 154  -0.20832934  0.8352471
## 11 11 dist3 - dist1  4F2 -0.054976600  0.2439611 154  -0.22534990  0.8220058
## 12 12 dist0 - dist1  4F2 -0.004152355  0.2439611 154  -0.01702057  0.9864422
##
## TEST OF WHETHER OR NOT THE INTERACTION PROVIDES A BETTER FIT
## Data: cur_data
## Models:
## m2: data ~ cond + harm + (1 | subject)
## m1: data ~ cond * harm + (1 | subject)
##      Df      AIC      BIC logLik deviance Chisq Chi Df Pr(>Chisq)
## m2   8 393.79 419.34 -188.90  377.79      0    6 0.9938
## m1  14 405.06 449.76 -188.53  377.06  0.7302    6 0.9938
```

RLS_CARRIER_RC2_CARR8



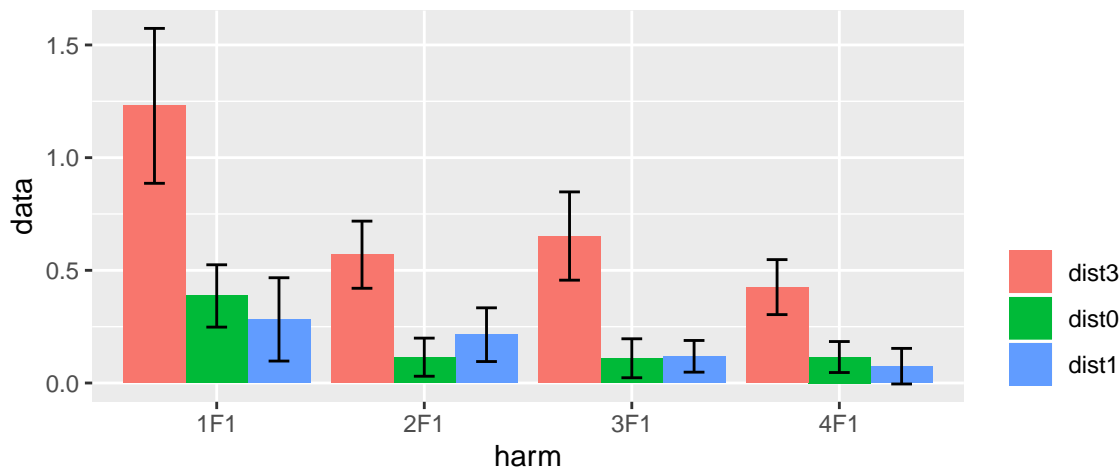
```
## LOVELY: NONE OF THE MODELS ARE SINGULAR! LOVELY: BOTH MODELS CONVERGED!
##
## ANOVA TEST FOR MAIN EFFECTS AND INTERACTIONS
## Type III Analysis of Variance Table with Satterthwaite's method
##      Sum Sq Mean Sq NumDF DenDF F value    Pr(>F)
## cond      0.2140  0.1070      2    154  0.2579    0.7730
## harm      9.5334  3.1778      3    154  7.6616 8.321e-05 ***
## cond:harm  0.3370  0.0562      6    154  0.1354    0.9915
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## ESTIMATED MARGINAL MEANS, SIMPLE MAIN EFFECTS OF CONDITION WITHOUT CORRECTION
##      contrast harm      estimate      SE df    t.ratio  p.value
## 1  1 dist3 - dist0  1F2  0.07297505 0.2351658 154  0.31031317 0.7567424
## 2  2 dist3 - dist1  1F2 -0.06065915 0.2351658 154 -0.25794201 0.7967960
## 3  3 dist0 - dist1  1F2 -0.13363420 0.2351658 154 -0.56825519 0.5706894
## 4  4 dist3 - dist0  2F2 -0.01358039 0.2351658 154 -0.05774815 0.9540241
## 5  5 dist3 - dist1  2F2  0.04277160 0.2351658 154  0.18187848 0.8559173
## 6  6 dist0 - dist1  2F2  0.05635199 0.2351658 154  0.23962663 0.8109384
## 7  7 dist3 - dist0  3F2  0.20674243 0.2351658 154  0.87913465 0.3806980
## 8  8 dist3 - dist1  3F2  0.15674794 0.2351658 154  0.66654216 0.5060620
## 9  9 dist0 - dist1  3F2 -0.04999449 0.2351658 154 -0.21259249 0.8319260
## 10 10 dist3 - dist0 4F2  0.07016915 0.2351658 154  0.29838157 0.7658142
## 11 11 dist3 - dist1 4F2  0.05682165 0.2351658 154  0.24162376 0.8093931
## 12 12 dist0 - dist1 4F2 -0.01334750 0.2351658 154 -0.05675781 0.9548117
##
## TEST OF WHETHER OR NOT THE INTERACTION PROVIDES A BETTER FIT
## Data: cur_data
## Models:
## m2: data ~ cond + harm + (1 | subject)
## m1: data ~ cond * harm + (1 | subject)
##      Df      AIC      BIC logLik deviance Chisq Chi Df Pr(>Chisq)
## m2   8 384.19 409.73 -184.09  368.19      0      6 0.9901
## m1  14 395.32 440.02 -183.66  367.32 0.8683      6 0.9901
```

RLS_ODDBALL_RC2_CARR6



```
## LOVELY: NONE OF THE MODELS ARE SINGULAR! LOVELY: BOTH MODELS CONVERGED!
##
## ANOVA TEST FOR MAIN EFFECTS AND INTERACTIONS
## Type III Analysis of Variance Table with Satterthwaite's method
##      Sum Sq Mean Sq NumDF DenDF F value  Pr(>F)
## cond      1.44769  0.72384      2    154   3.7696 0.02522 *
## harm      2.18758  0.72919      3    154   3.7975 0.01159 *
## cond:harm  0.96378  0.16063      6    154   0.8365 0.54346
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## ESTIMATED MARGINAL MEANS, SIMPLE MAIN EFFECTS OF CONDITION WITHOUT CORRECTION
##      contrast harm estimate      SE df  t.ratio  p.value
## 1  1 dist3 - dist0 1F1  0.20294695 0.1600089 154  1.2683479 0.20658774
## 2  2 dist3 - dist1 1F1  0.04487824 0.1600089 154  0.2804734 0.77949074
## 3  3 dist0 - dist1 1F1 -0.15806871 0.1600089 154 -0.9878745 0.32476368
## 4  4 dist3 - dist0 2F1  0.02661440 0.1600089 154  0.1663307 0.86811486
## 5  5 dist3 - dist1 2F1 -0.27776301 0.1600089 154 -1.7359223 0.08457766
## 6  6 dist0 - dist1 2F1 -0.30437741 0.1600089 154 -1.9022531 0.05900488
## 7  7 dist3 - dist0 3F1  0.30270420 0.1600089 154  1.8917961 0.06039538
## 8  8 dist3 - dist1 3F1  0.05418727 0.1600089 154  0.3386516 0.73533348
## 9  9 dist0 - dist1 3F1 -0.24851693 0.1600089 154 -1.5531445 0.12244130
## 10 10 dist3 - dist0 4F1  0.22919965 0.1600089 154  1.4324182 0.15405088
## 11 11 dist3 - dist1 4F1  0.17968990 0.1600089 154  1.1229995 0.26318541
## 12 12 dist0 - dist1 4F1 -0.04950975 0.1600089 154 -0.3094187 0.75742128
##
## TEST OF WHETHER OR NOT THE INTERACTION PROVIDES A BETTER FIT
## Data: cur_data
## Models:
## m2: data ~ cond + harm + (1 | subject)
## m1: data ~ cond * harm + (1 | subject)
##      Df      AIC      BIC logLik deviance Chisq Chi Df Pr(>Chisq)
## m2   8 246.20 271.74 -115.10  230.20
## m1  14 252.91 297.61 -112.45  224.91 5.2918     6     0.507
```


RLS_ODDBALL_RC2_CARR8



```
## LOVELY: NONE OF THE MODELS ARE SINGULAR! LOVELY: BOTH MODELS CONVERGED!
##
## ANOVA TEST FOR MAIN EFFECTS AND INTERACTIONS
## Type III Analysis of Variance Table with Satterthwaite's method
##      Sum Sq Mean Sq NumDF DenDF F value    Pr(>F)
## cond      11.7624   5.8812     2    154 18.8575 4.727e-08 ***
## harm       4.7903   1.5968     3    154  5.1199 0.002103 **
## cond:harm   2.0517   0.3419     6    154  1.0964 0.367033
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## ESTIMATED MARGINAL MEANS, SIMPLE MAIN EFFECTS OF CONDITION WITHOUT CORRECTION
##      contrast harm estimate      SE df    t.ratio    p.value
## 1 1 dist3 - dist0 1F1 0.84346855 0.2039207 154  4.13625725 5.786860e-05
## 2 2 dist3 - dist1 1F1 0.94756777 0.2039207 154  4.64674592 7.190578e-06
## 3 3 dist0 - dist1 1F1 0.10409922 0.2039207 154  0.51048868 6.104401e-01
## 4 4 dist3 - dist0 2F1 0.45468754 0.2039207 154  2.22972706 2.721218e-02
## 5 5 dist3 - dist1 2F1 0.35488872 0.2039207 154  1.74032694 8.379916e-02
## 6 6 dist0 - dist1 2F1 -0.09979882 0.2039207 154 -0.48940011 6.252548e-01
## 7 7 dist3 - dist0 3F1 0.54233203 0.2039207 154  2.65952390 8.652841e-03
## 8 8 dist3 - dist1 3F1 0.53364313 0.2039207 154  2.61691470 9.757156e-03
## 9 9 dist0 - dist1 3F1 -0.00868890 0.2039207 154 -0.04260921 9.660683e-01
## 10 10 dist3 - dist0 4F1 0.31024409 0.2039207 154  1.52139564 1.302112e-01
## 11 11 dist3 - dist1 4F1 0.35090445 0.2039207 154  1.72078862 8.729762e-02
## 12 12 dist0 - dist1 4F1 0.04066036 0.2039207 154  0.19939298 8.422184e-01
##
## TEST OF WHETHER OR NOT THE INTERACTION PROVIDES A BETTER FIT
## Data: cur_data
## Models:
## m2: data ~ cond + harm + (1 | subject)
## m1: data ~ cond * harm + (1 | subject)
##      Df      AIC      BIC logLik deviance Chisq Chi Df Pr(>Chisq)
## m2  8 328.19 353.73 -156.09 312.19
## m1 14 333.29 377.99 -152.64 305.29 6.9019 6 0.33
```