Numbers-Experment-MML

Behavioural analysis

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Table of contents

```
Analysis based on Persons experiment: Middleton and Moitra 2023
library("tidyverse")
library("plotrix")
library("lme4")
library("lmerTest")
library("report")
library("Hmisc")
library("multcomp")
library("emmeans")
source('ggplot_theme_Publication-2.R')
# Load data
df_slet_1 <- read_csv("data/N72/slet.csv")</pre>
df_98zo_1 <- read_csv("data/N72/98zo.csv")</pre>
df_goty_1 <- read_csv("data/N72/goty.csv")</pre>
df_hebw_1 <- read_csv("data/N72/hebw.csv")</pre>
df_ajpl_1 <- read_csv("data/N72/ajpl.csv")</pre>
df_ld3j_1 <- read_csv("data/N72/ld3j.csv")</pre>
df_slet_2 <- read_csv("data/Rest/slet.csv")</pre>
df 98zo 2 <- read csv("data/Rest/98zo.csv")</pre>
df_goty_2 <- read_csv("data/Rest/goty.csv")</pre>
df hebw 2 <- read csv("data/Rest/hebw.csv")</pre>
df ajpl 2 <- read csv("data/Rest/ajpl.csv")</pre>
df_ld3j_2 <- read_csv("data/Rest/ld3j.csv")</pre>
```

```
df_slet_3 <- read_csv("data/Rest_new/slet.csv")</pre>
df_98zo_3 <- read_csv("data/Rest_new/98zo.csv")</pre>
df_goty_3 <- read_csv("data/Rest_new/goty.csv")</pre>
df_hebw_3 <- read_csv("data/Rest_new/hebw.csv")</pre>
df ajpl 3 <- read csv("data/Rest new/ajpl.csv")</pre>
df_ld3j_3 <- read_csv("data/Rest_new/ld3j.csv")</pre>
# Custom Function
data_cleaning <- function(arg1){</pre>
arg1 <- arg1
arg1 <- arg1 %>%
         dplyr::select(`Participant Private ID`, `Trial Number`, 'Tree Node Key', 'Reaction Time
         rename(Subject = `Participant Private ID`,
                     Item = `Trial Number`,
                     Condition = 'Task Name',
                     RT = 'Reaction Time',
                     Accuracy = Correct) %>%
         filter(Screen == "Testing",
                 `Response Type` == "response") %>%
         mutate(LogRT = log(RT),
         Condition = gsub(" B Testing", "", Condition))
return(arg1)
}
## Use the custom function to clean the data
df_slet_1 <- data_cleaning(df_slet_1)</pre>
df_98zo_1 <- data_cleaning(df_98zo_1)
df_goty_1 <- data_cleaning(df_goty_1)</pre>
df_hebw_1 <- data_cleaning(df_hebw_1)</pre>
df_ajpl_1 <- data_cleaning(df_ajpl_1)</pre>
df_ld3j_1 <- data_cleaning(df_ld3j_1)</pre>
df_slet_2 <- data_cleaning(df_slet_2)</pre>
df 98zo 2 <- data cleaning(df 98zo 2)
df_goty_2 <- data_cleaning(df_goty_2)</pre>
df_hebw_2 <- data_cleaning(df_hebw_2)</pre>
df_ajpl_2 <- data_cleaning(df_ajpl_2)</pre>
df_ld3j_2 <- data_cleaning(df_ld3j_2)</pre>
df_slet_3 <- data_cleaning(df_slet_3)</pre>
df_98zo_3 <- data_cleaning(df_98zo_3)</pre>
```

```
Subject
                       Item
                                  Tree Node Key
                                                           RT
Min. : 9898448
                         : 1.00
                                  Length:19392
                                                                 16.63
                  Min.
                                                     Min.
                                                          :
1st Qu.: 9992482
                  1st Qu.:12.75
                                                     1st Qu.: 1793.88
                                  Class :character
Median :10004852
                  Median :24.50
                                  Mode :character
                                                     Median :
                                                               3318.40
                        :24.50
Mean :10383072
                  Mean
                                                     Mean : 5322.75
3rd Qu.:10847984
                  3rd Qu.:36.25
                                                     3rd Qu.:
                                                               6334.27
Max.
      :10990139
                  Max.
                         :48.00
                                                     Max.
                                                            :205295.10
   Accuracy
                Spreadsheet: display
                                        Screen
                                                         Condition
      :0.0000
                Length: 19392
                                     Length: 19392
                                                        Length: 19392
Min.
1st Qu.:0.0000
                Class : character
                                     Class :character
                                                        Class : character
Median :1.0000
                Mode :character
                                     Mode :character
                                                        Mode :character
Mean
      :0.6358
3rd Qu.:1.0000
Max.
      :1.0000
Response Type
                      LogRT
Length: 19392
                        : 2.811
                  Min.
                  1st Qu.: 7.492
Class : character
Mode :character
                  Median: 8.107
                  Mean : 8.027
                  3rd Qu.: 8.754
                  Max. :12.232
```

#Stats

```
HP_data <- data %>%
  mutate(
    Subject = as.factor(Subject),
   Condition = as.factor(Condition),
    Item = as.factor(Item)
    #Cond_Type = as.factor(Cond_Type),
    #Response = as.factor(Response)
  )
str(HP_data)
tibble [19,392 x 10] (S3: tbl_df/tbl/data.frame)
                      : Factor w/ 404 levels "9898448", "9898449", ...: 13 13 13 13 13 13 13 13
 $ Subject
                       : Factor w/ 48 levels "1", "2", "3", "4", ...: 1 2 3 4 5 6 7 8 9 10 ...
 $ Item
                       : chr [1:19392] "task-slet" "task-slet" "task-slet" "task-slet" ...
 $ Tree Node Key
                       : num [1:19392] 10352 4506 6214 5762 7446 ...
 $ RT
 $ Accuracy
                      : num [1:19392] 1 1 1 1 1 1 1 1 1 1 ...
 $ Spreadsheet: display: chr [1:19392] "Testing02" "Testing02" "Testing02" "Testing02" ...
                      : chr [1:19392] "Testing" "Testing" "Testing" "Testing" ...
 $ Screen
                      : Factor w/ 6 levels "1e-1i", "1e-2", ...: 1 1 1 1 1 1 1 1 1 1 ...
 $ Condition
                      : chr [1:19392] "response" "response" "response" "response" ...
 $ Response Type
 $ LogRT
                       : num [1:19392] 9.24 8.41 8.73 8.66 8.92 ...
HP_data$Condition <-relevel(HP_data$Condition, ref="1e-1i")</pre>
RT_model_1 <- lmer(LogRT ~ Condition + (1|Subject) + (1|Item), data = HP_data, REML = F)</pre>
summary(RT_model_1)
Linear mixed model fit by maximum likelihood . t-tests use Satterthwaite's
  method [lmerModLmerTest]
Formula: LogRT ~ Condition + (1 | Subject) + (1 | Item)
   Data: HP_data
     AIC
              BIC logLik deviance df.resid
 37339.6 37410.5 -18660.8 37321.6
                                       19383
Scaled residuals:
    Min 1Q Median 3Q
                                    Max
-9.8912 -0.5815 -0.0555 0.5167 7.8253
Random effects:
 Groups
                 Variance Std.Dev.
        Name
```

```
Subject (Intercept) 0.7329
                              0.8561
          (Intercept) 0.1251
                              0.3537
 Item
 Residual
                     0.3602
                              0.6002
Number of obs: 19392, groups:
                             Subject, 404; Item, 48
Fixed effects:
               Estimate Std. Error
                                          df t value Pr(>|t|)
(Intercept)
               8.080737 0.121428 426.374017 66.547
                                                       <2e-16 ***
Condition1e-2 0.050243 0.154565 403.653064 0.325
                                                       0.745
Condition1e-3 -0.220706 0.147919 403.653066 -1.492
                                                        0.136
Condition1i-2 0.009701 0.154565 403.653064
                                              0.063
                                                        0.950
Condition1i-3 -0.034886 0.151223 403.653066 -0.231
                                                        0.818
Condition2-3 -0.085942 0.149738 403.653067 -0.574
                                                        0.566
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
Correlation of Fixed Effects:
             (Intr) Condition1e-2 Condition1e-3 Condition1i-2 Condition1i-3
Condition1e-2 -0.647
Condition1e-3 -0.676 0.531
Condition1i-2 -0.647 0.508
                                   0.531
Condition1i-3 -0.661 0.519
                                   0.543
                                                0.519
Conditin2-3 -0.668 0.524
                                  0.548
                                                0.524
                                                              0.536
ACC_model_1 <- glmer(Accuracy ~ Condition + (1|Subject) + (1|Item), data = HP_data, family =
summary(ACC_model_1)
Generalized linear mixed model fit by maximum likelihood (Laplace
  Approximation) [glmerMod]
 Family: binomial (logit)
Formula: Accuracy ~ Condition + (1 | Subject) + (1 | Item)
   Data: HP_data
             BIC logLik deviance df.resid
    AIC
 24420.9 24483.9 -12202.5 24404.9
                                     19384
Scaled residuals:
            1Q Median
                            3Q
                                  Max
-3.2172 -1.0560 0.5304 0.7839 1.5866
```

Variance Std.Dev.

Random effects: Groups Name

```
Subject (Intercept) 0.459757 0.67805
        (Intercept) 0.003554 0.05961
 Item
Number of obs: 19392, groups: Subject, 404; Item, 48
Fixed effects:
             Estimate Std. Error z value Pr(>|z|)
(Intercept)
           Condition1e-2 -0.079669 0.135745 -0.587 0.55727
Condition1i-2 0.005684 0.135773 0.042 0.96661
Condition1i-3 -0.328282 0.132309 -2.481 0.01310 *
Condition2-3 -0.302998 0.131022 -2.313 0.02075 *
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
Correlation of Fixed Effects:
            (Intr) Condition1e-2 Condition1e-3 Condition1i-2 Condition1i-3
Condition1e-2 -0.712
Condition1e-3 -0.748 0.536
Condition1i-2 -0.712 0.511
                                0.537
Condition1i-3 -0.731 0.524
                                0.551
                                             0.524
Conditin2-3 -0.738 0.530
                                0.556
                                             0.530
                                                          0.544
ACC_model_2 <- glmer(Accuracy ~ Condition + (1|Subject) , data = HP_data, family = "binomial"
summary(ACC_model_2)
Generalized linear mixed model fit by maximum likelihood (Laplace
  Approximation) [glmerMod]
Family: binomial (logit)
Formula: Accuracy ~ Condition + (1 | Subject)
  Data: HP_data
                 logLik deviance df.resid
    AIC
            BIC
24420.8 24475.9 -12203.4 24406.8
Scaled residuals:
           1Q Median
                          3Q
-3.2007 -1.0587 0.5328 0.7836 1.5544
Random effects:
Groups Name
                   Variance Std.Dev.
```

Subject (Intercept) 0.4589 0.6775

```
Number of obs: 19392, groups: Subject, 404
Fixed effects:
            Estimate Std. Error z value Pr(>|z|)
          0.82088 0.09711 8.453 < 2e-16 ***
(Intercept)
Condition1e-2 -0.07961
                     0.13564 -0.587 0.55728
Condition1i-2 0.00568 0.13568 0.042 0.96661
0.13093 -2.313 0.02074 *
Condition2-3 -0.30281
___
Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
Correlation of Fixed Effects:
            (Intr) Condition1e-2 Condition1e-3 Condition1i-2 Condition1i-3
Condition1e-2 -0.714
Condition1e-3 -0.750 0.536
Condition1i-2 -0.715 0.511
                               0.537
Condition1i-3 -0.733 0.524
                               0.551
                                           0.525
Conditin2-3 -0.741 0.530
                               0.556
                                           0.530
                                                       0.544
anova(ACC_model_1,ACC_model_2)
Data: HP_data
Models:
ACC_model_2: Accuracy ~ Condition + (1 | Subject)
ACC_model_1: Accuracy ~ Condition + (1 | Subject) + (1 | Item)
          npar AIC BIC logLik deviance Chisq Df Pr(>Chisq)
ACC model 2
            7 24421 24476 -12203
                                  24407
ACC_model_1 8 24421 24484 -12202
                                  24405 1.8305 1
                                                   0.1761
0.1 Plot
Data$Condition <- as.factor(Data$Condition)</pre>
```

```
Data$Condition <- as.factor(Data$Condition)

# RT <- ggplot(Data, aes(x=Condition, y=LogRT)) +

# geom_violin(aes(fill = Condition), trim = FALSE, show.legend = FALSE) +

# ylab("Log RT") +

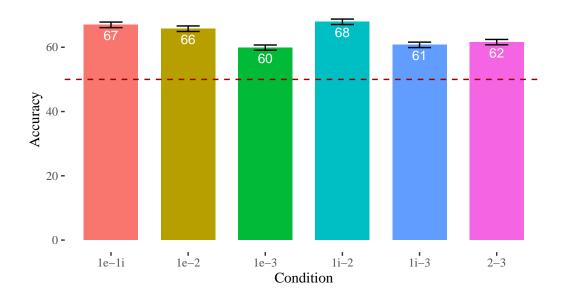
# xlab("Conditions") +</pre>
```

```
#
#
   # geom_signif(
   # comparisons = list(c("Grammatical", "Pseudowords")),
#
   # margin_top = 0.20,
   # step_increase = 0.05,
#
   # tip_length = 0.01,
#
   # map_signif_level = TRUE
   # )+
#
#
   theme_Publication()+
#
#
#
#
   # Add geom_boxplot() to include box plot
    geom_boxplot(width = 0.2, fill = "white", color = "blue")
#
# RT + stat_summary(fun.data=mean_sdl, mult=1,
                   geom="pointrange", color="red")
```

```
ACC_plot<- ggplot(data_group,
               aes(x=Condition,y=ACC,fill=Condition)) +
  # geom_bar function is used to plot bars of barplot
  geom_bar(stat = "identity", width = 0.7, position = position_dodge(0.7), show.legend = FAL
  #scale_x_discrete(limits = Conditions) + facet_wrap( ~Prefix) +
  ylab("% Accept") +
  xlab("Conditions") +
  geom_text(aes(label=round(ACC)),vjust=1.6, color="white", size=3.5) +
  theme_Publication()+
  #theme_minimal() +
  #theme_clean() +
  #theme_wsj() +
  theme tufte() +
  geom_hline(aes(yintercept=50), colour="#990000", linetype="dashed") +
  coord_cartesian(ylim = c(0, 75)) +
  # scale_y_continuous(expand = expansion(mult = c(0, 0.05)))+
  # geom_signif(
     comparisons = list(c("Grammatical", "Pseudowords")),
  # margin_top = 0.12,
  # step_increase = 0.09,
  # tip_length = 0.05,
    annotation = c("***")
  # )+
```

```
ACC_plot + labs(title="Accuracy | Condition",
    x = "Condition", y = "Accuracy")
```

Accuracy | Condition



report(ACC_model_1)

We fitted a logistic mixed model (estimated using ML and Nelder-Mead optimizer) to predict Accuracy with Condition (formula: Accuracy ~ Condition). The model included Subject as random effects (formula: list(~1 | Subject, ~1 | Item)). The model's total explanatory power is weak (conditional R2 = 0.13) and the part related to the fixed effects alone (marginal R2) is of 6.77e-03. The model's intercept, corresponding to Condition = 1e-1i, is at 0.82 (95% CI [0.63, 1.01], p < .001). Within this model:

- The effect of Condition [1e-2] is statistically non-significant and negative (beta = -0.08, 95% CI [-0.35, 0.19], p = 0.557; Std. beta = -0.08, 95% CI [-0.35, 0.19])

```
- The effect of Condition [1e-3] is statistically significant and negative (beta = -0.38, 95% CI [-0.63, -0.13], p = 0.003; Std. beta = -0.38, 95% CI [-0.63, -0.13])
```

- The effect of Condition [1i-2] is statistically non-significant and positive (beta = 5.68e-03, 95% CI [-0.26, 0.27], p = 0.967; Std. beta = 5.68e-03, 95% CI [-0.26, 0.27])
- The effect of Condition [1i-3] is statistically significant and negative (beta = -0.33, 95% CI [-0.59, -0.07], p = 0.013; Std. beta = -0.33, 95% CI [-0.59, -0.07])
- The effect of Condition [2-3] is statistically significant and negative (beta = -0.30, 95% CI [-0.56, -0.05], p = 0.021; Std. beta = -0.30, 95% CI [-0.56, -0.05])

Standardized parameters were obtained by fitting the model on a standardized version of the dataset. 95% Confidence Intervals (CIs) and p-values were computed using a Wald z-distribution approximation.

```
BAN_ACC_Model1.emm <- emmeans(ACC_model_1, ~ Condition, lmer.df = "asymp")
pairs(BAN ACC Model1.emm, simple = "each")</pre>
```

```
contrast
               estimate
                           SE df z.ratio p.value
(1e-1i) - (1e-2) 0.07967 0.136 Inf
                                   0.587 0.9919
(1e-1i) - (1e-3) 0.37922 0.129 Inf
                                    2.930 0.0397
(1e-1i) - (1i-2) -0.00568 0.136 Inf -0.042 1.0000
(1e-1i) - (1i-3) 0.32828 0.132 Inf
                                    2.481 0.1296
(1e-1i) - (2-3) 0.30300 0.131 Inf
                                    2.313 0.1888
(1e-2) - (1e-3) 0.29955 0.128 Inf
                                   2.344 0.1764
(1e-2) - (1i-2) -0.08535 0.134 Inf -0.636 0.9883
(1e-2) - (1i-3) 0.24861 0.131 Inf 1.901 0.4011
(1e-2) - (2-3)
               0.22333 0.129 Inf
                                   1.726 0.5147
(1e-3) - (1i-2) -0.38490 0.128 Inf -3.012 0.0311
(1e-3) - (1i-3) -0.05094 0.124 Inf -0.410 0.9985
(1e-3) - (2-3)
               -0.07622 0.123 Inf -0.621 0.9895
(1i-2) - (1i-3) 0.33397 0.131 Inf
                                   2.554 0.1089
(1i-2) - (2-3)
               0.30868 0.129 Inf
                                    2.385 0.1614
(1i-3) - (2-3)
               -0.02528 0.126 Inf -0.201 1.0000
```

Results are given on the log odds ratio (not the response) scale. P value adjustment: tukey method for comparing a family of 6 estimates