

M21 RT Orthographic Sensitivity

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Setup

Load libraries

1. Set `ggplot2` parameters

Load Files and Format Files

Load Files

```
#DIR <- "csv_files"
df_a <- read_csv("rt_data_hc_A.csv")
df_b <- read_csv("rt_data_hc_B_fixed.csv")
frq_w <- read_csv("frq_cw.csv")
frq_nw <- read_csv("frq_nw.csv")
dmg <- read_csv("demo_lang_vsl_pca_hc.csv")
```

Format Files

```
# Concatenate datasets
rt <- bind_rows(AB = df_a,
               BA = df_b,
               .id = "List")
rt_dmg <- right_join(dmg, rt, join_by(SubjID == subject_nr)) |> # Join Participant Demographic and Lang Data
  mutate(target = tolower(target)) |>
  filter(correct == 1)

# Divide into Experimental and Filler Items
rt_fill <- rt_dmg |> filter(str_detect(targ_type, "^FILL"))
rt_exp <- rt_dmg |> filter(!str_detect(targ_type, "^FILL"))

# Define Factors and Conditions
rt_exp_format <- rt_exp |>
  separate(targ_type, into = c("trial_type", "family_size", "complexity"), sep = "_",
           remove = TRUE, extra = "drop", fill = "right")

# Divide into Words and Nonwords
rt_words <- rt_exp_format |> filter(trial_type == "CW") |> select(- complexity)
rt_nwords <- rt_exp_format |> filter(trial_type == "NW")

# Join Stimulus Frequency Data
rt_words_frq <- left_join(rt_words, frq_w, join_by(target)) |>
  select(-cond_trig.y, -word_trig.y) |>
  rename(cond_trig = cond_trig.x, word_trig = word_trig.x) # remove duplicate columns
rt_nwords_frq <- left_join(rt_nwords, frq_nw, join_by(target==word)) |>
  select(-cond_trig.y, -word_trig.y) |>
  rename(cond_trig = cond_trig.x, word_trig = word_trig.x)

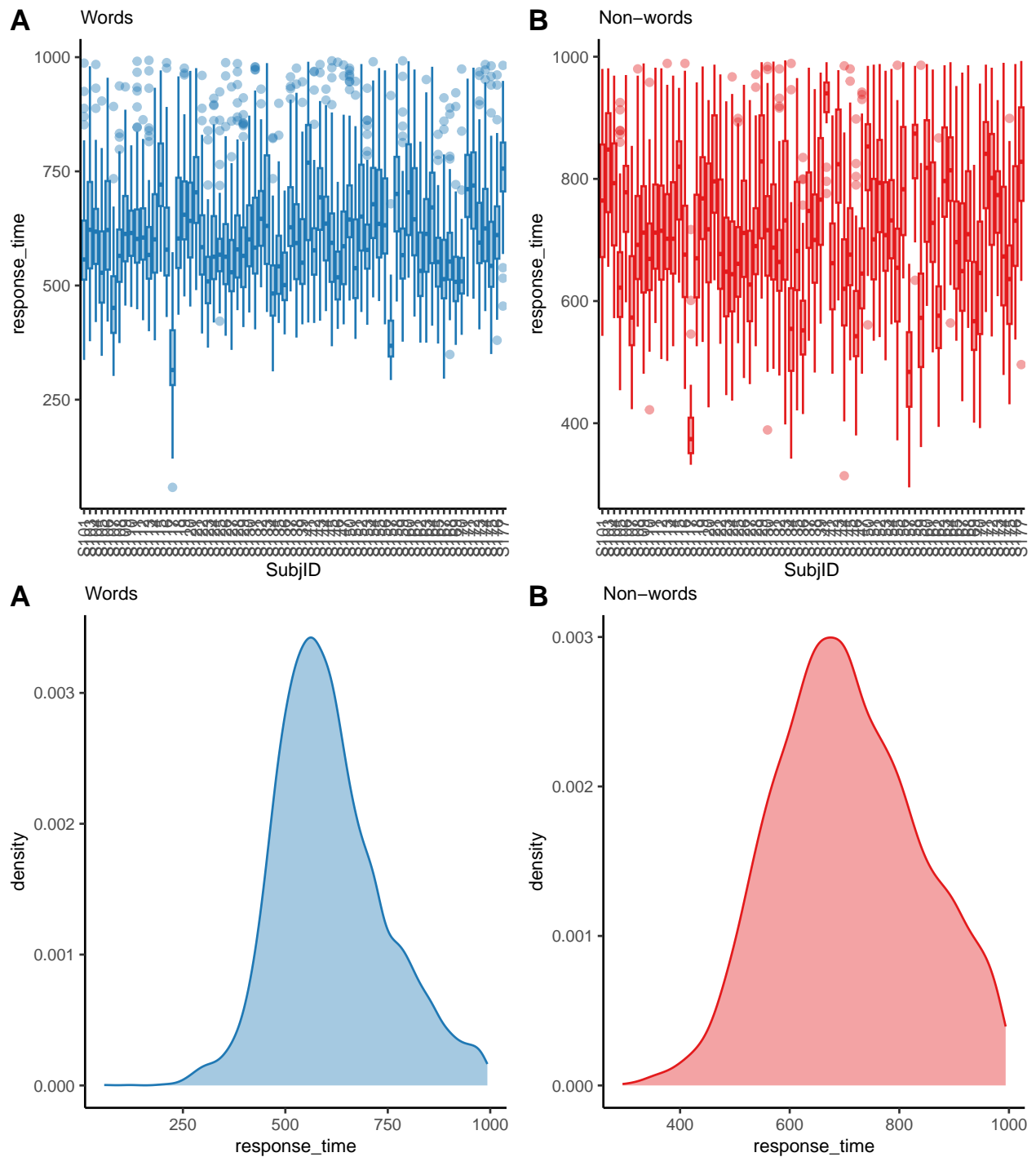
# Rename BF_Split and FS_Split columns
rt_words_frq <- rt_words_frq |> rename(Base_Frequency = BF_Split, Family_Size = FS_Split) # Rename BF_Split and FS_Split columns
rt_nwords_frq <- rt_nwords_frq |> rename(Base_Frequency = BF_Split, Family_Size = FS_Split)

# Recode factor levels
# rt_words_frq <- rt_words_frq |>
#   mutate(Base_Frequency = case_match(Base_Frequency, "Low" ~ "Low BF", "High" ~ "High BF"),
#          Family_Size = case_match(Family_Size, "Small" ~ "Small Family", "Large" ~ "Large Family"))
# rt_nwords_frq <- rt_nwords_frq |> mutate(Base_Frequency = case_match(Base_Frequency, "Low" ~ "Low BF", "High" ~ "High BF"),
#          Family_Size = case_match(Family_Size, "Small" ~ "Small Family", "Large" ~ "Large Family"))
#
```

```
# rt_words_freq$Orthographic_Sensitivity[rt_words_freq$Orthographic_Sensitivity == "Low"] <- "Low Sensitivity"
# rt_words_freq$Orthographic_Sensitivity[rt_words_freq$Orthographic_Sensitivity == "High"] <- "High Sensitivity"
```

Analyse Data

Plot RT distributions

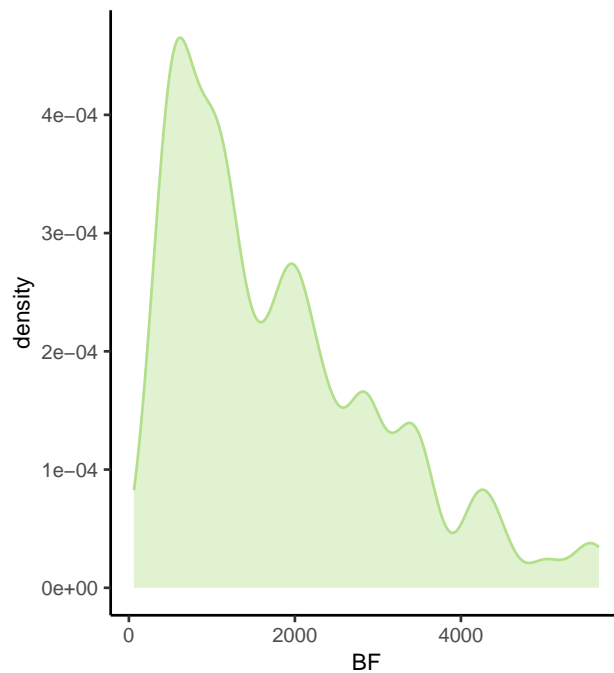


Test for Skewness

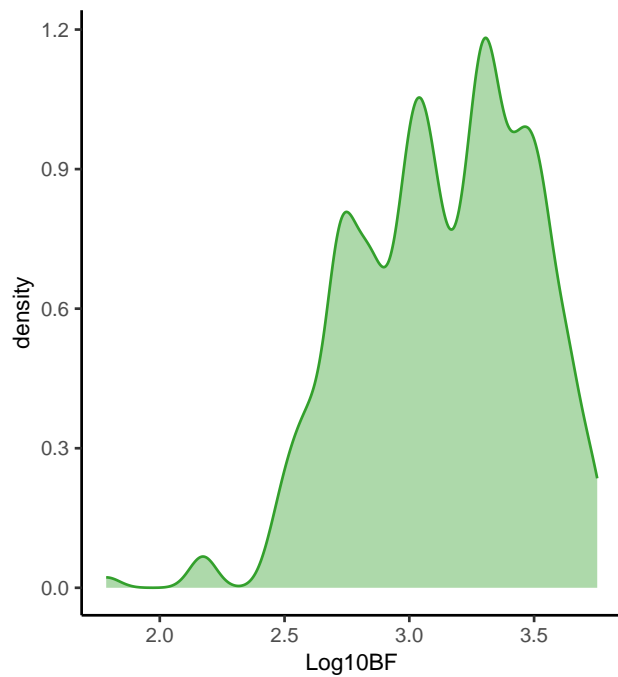
Base Frequency

...

A Distribution of Raw Base Frequency



B Distribution of Log10 Base Frequency



```
|| [1] 0.9852864
```

```
|| [1] -0.4180109
```

Family Size

...

Skewness values

```
rt_words_frq <- rt_words_frq |> mutate(Log10FS = log10(FS))  
skewness(rt_words_frq$FS, na.rm = TRUE)
```

```
|| [1] 1.104411
```

```
skewness(rt_words_frq$Log10FS, na.rm = TRUE)
```

```
|| [1] 0.05939575
```

Raw FS

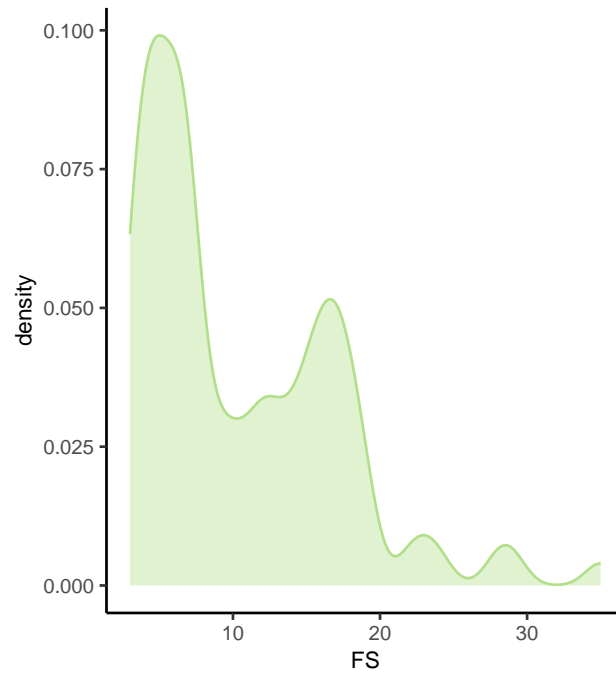
```
p1 <- ggplot(rt_words_frq, aes(x = FS)) +  
  geom_density(colour = "#B2DF8A", fill = "#B2DF8A", alpha = .4) +  
  labs(title = "Distribution of Family Size")
```

Log10 FS

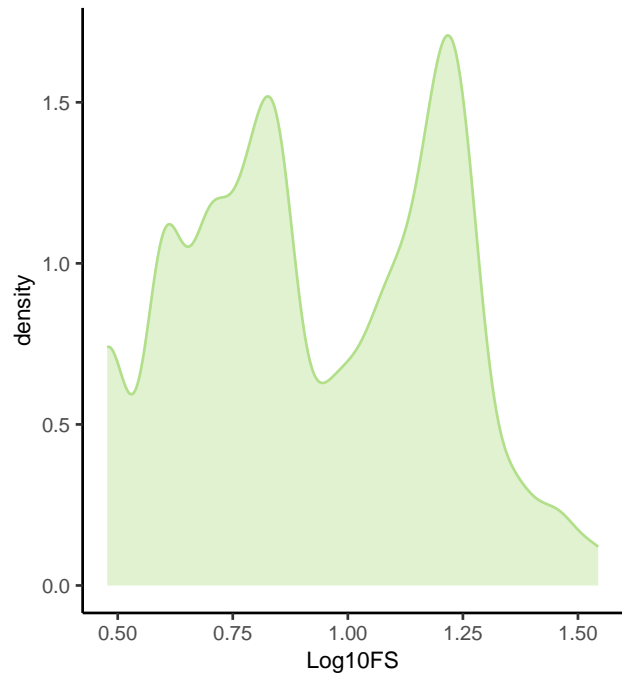
```
p2 <- ggplot(rt_words_frq, aes(x = Log10FS)) +  
  geom_density(colour = "#B2DF8A", fill = "#B2DF8A", alpha = .4) +  
  labs(title = "Distribution of Log10 Family Size")
```

```
plot_grid(p1, p2, ncol = 2, labels = "AUTO")
```

A Distribution of Family Size



B Distribution of Log10 Family Size



Word Data

Use `complete.cases()` to find which rows have missing data in the model-relevant variables:

```
rt_words_cmpl %>%
  summarise(
    n_subjects = n_distinct(SubjID),
    n_items = n_distinct(STRING))
```

```
|| # A tibble: 1 x 2
||   n_subjects n_items
||   <int>     <int>
|| 1         67      100
```

Count trials per subject

```
rt_words_cmpl %>%
  count(SubjID, name = "n_trials") %>%
  summarise(
    min_trials = min(n_trials),
    max_trials = max(n_trials),
    mean_trials = mean(n_trials))
```

```
|| # A tibble: 1 x 3
||   min_trials max_trials mean_trials
||   <int>      <int>      <dbl>
|| 1         67         97         87.3
```

```
(trial_count_by_subj <- rt_words_cmpl %>%
  count(SubjID, name = "n_trials") %>%
  arrange(desc(n_trials)))
```

```
|| # A tibble: 67 x 2
||   SubjID n_trials
||   <chr>   <int>
|| 1 S113      97
|| 2 S130      96
|| 3 S142      96
|| 4 S168      96
|| 5 S110      95
|| 6 S129      95
|| 7 S104      94
|| 8 S114      94
|| 9 S123      94
|| 10 S125     94
|| # i 57 more rows
```

```
rt_words_cmpl %>%
  count(Family_Size, Base_Frequency, Orthographic_Sensitivity)
```

```
|| # A tibble: 8 x 4
||   Family_Size Base_Frequency Orthographic_Sensitivity     n
||   <chr>      <chr>          <chr>                <int>
|| 1 Large      High           High                 773
|| 2 Large      High           Low                  684
|| 3 Large      Low            High                 774
|| 4 Large      Low            Low                  684
|| 5 Small      High           High                 782
|| 6 Small      High           Low                  655
|| 7 Small      Low            High                 813
|| 8 Small      Low            Low                  682
```

Anova

```
anova_model_words <- mixed(
  response_time ~ Base_Frequency * Family_Size * Orthographic_Sensitivity +
    (1 | SubjID) +
    (1 | STRING),
  data = rt_words_cmpl,
  method = "S")
anova_model_words
```

|| Mixed Model Anova Table (Type 3 tests, S-method)

```
||
|| Model: response_time ~ Base_Frequency * Family_Size * Orthographic_Sensitivity +
|| Model:      (1 | SubjID) + (1 | STRING)
|| Data: rt_words_cmpl
||
||           Effect      df      F p.value
|| 1      Base_Frequency    1, 92.45 10.29 **   .002
|| 2      Family_Size      1, 92.44  9.41 **   .003
|| 3      Orthographic_Sensitivity 1, 64.87  3.83 +   .055
|| 4      Base_Frequency:Family_Size 1, 92.45  1.08   .300
|| 5      Base_Frequency:Orthographic_Sensitivity 1, 5682.28  0.05   .817
```

```

|| 6          Family_Size:Orthographic_Sensitivity 1, 5682.26      0.06      .809
|| 7 Base_Frequency:Family_Size:Orthographic_Sensitivity 1, 5682.09      0.16      .691
|| ---
|| Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
m1 <- anova_model_words$full_model      # Extract the lmer model
ranova(m1) # formally test whether adding each random effect improves fit

|| ANOVA-like table for random-effects: Single term deletions
||
|| Model:
|| response_time ~ Base_Frequency + Family_Size + Orthographic_Sensitivity + (1 | SubjID) + (1 | STRING) + Base_Frequency:Family_Size + Base_Frequency:Orthographic_Sensitivity
||               npar logLik   AIC    LRT Df Pr(>Chisq)
|| <none>          11 -35809 71639
|| (1 | SubjID)     10 -36718 73455 1817.77  1 < 2.2e-16 ***
|| (1 | STRING)     10 -35899 71817  179.65  1 < 2.2e-16 ***
|| ---
|| Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
# Extract effect sizes from your ANOVA model
eta_squared(anova_model_words, partial = TRUE)

|| # Effect Size for ANOVA (Type III)
||
|| Parameter | Eta2 (partial) | 95% CI
|| -----|-----|-----
|| Base_Frequency | 0.10 | [0.02, 1.00]
|| Family_Size | 0.09 | [0.02, 1.00]
|| Orthographic_Sensitivity | 0.06 | [0.00, 1.00]
|| Base_Frequency:Family_Size | 0.01 | [0.00, 1.00]
|| Base_Frequency:Orthographic_Sensitivity | 9.41e-06 | [0.00, 1.00]
|| Family_Size:Orthographic_Sensitivity | 1.03e-05 | [0.00, 1.00]
|| Base_Frequency:Family_Size:Orthographic_Sensitivity | 2.78e-05 | [0.00, 1.00]
||
|| - One-sided CIs: upper bound fixed at [1.00].
# Compute Marginal(fixed effects only) and Conditional(fixed + random effects) R^2
r2(anova_model_words)

|| # R2 for Mixed Models
||
|| Conditional R2: 0.360
|| Marginal R2: 0.028

```

Concise

Models including random slopes for Base Frequency and Family Size by subject failed to converge or produced singular fits, indicating that the data did not support estimation of these additional variance components. Consequently, we report results from a simpler model with random intercepts for subjects and items (STRING), which converged cleanly and provided stable estimates.

Fuller explanation

We initially attempted to fit a maximal random-effects structure following Barr et al. (2013), including random slopes for Base Frequency and Family Size by subject. However, these models yielded singular fits (zero variance estimates and perfect correlations among random effects). Because such structures can produce unreliable standard errors and inflated Type I error rates, we adopted the maximal non-singular model, containing random intercepts for both subjects and items (STRING). All reported statistics are based on this model.

Brief

(A more complex model including by-subject random slopes failed to converge; results from the non-singular intercept-only model are reported.)

Main Findings

Effect	df	F	p.value
Base_Frequency	1, 92.45	10.29 **	.002
Family_Size	1, 92.44	9.41 **	.003

Plots

```

|| Base_Frequency  emmean      SE df asymp.LCL asymp.UCL
|| High           602.4149  9.881996 Inf  583.0465  621.7832
|| Low            621.9280  9.826032 Inf  602.6694  641.1867
||
|| Results are averaged over the levels of: Family_Size, Orthographic_Sensitivity
|| Degrees-of-freedom method: asymptotic
|| Confidence level used: 0.95

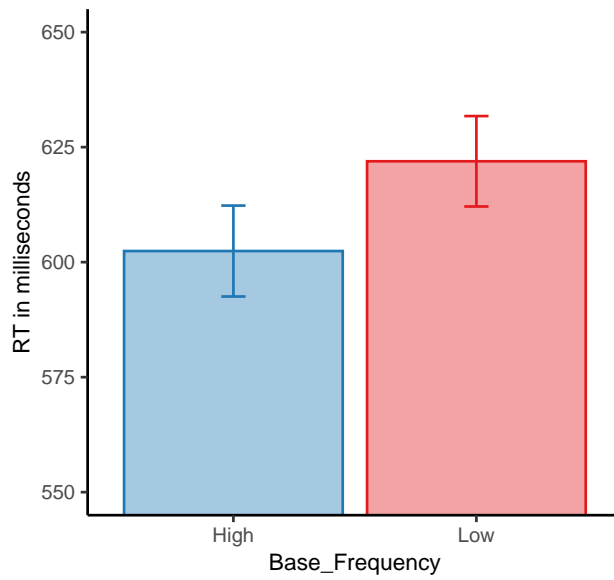
```

```

|| Family_Size  emmean      SE  df asymp.LCL asymp.UCL
|| Large       602.8423  9.856568 Inf  583.5238  622.1608
|| Small       621.5006  9.851504 Inf  602.1920  640.8092
||
|| Results are averaged over the levels of: Base_Frequency, Orthographic_Sensitivity
|| Degrees-of-freedom method: asymptotic
|| Confidence level used: 0.95

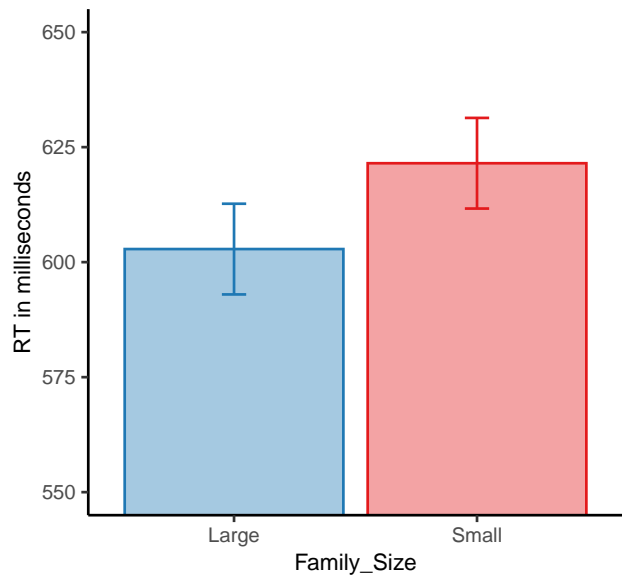
```

A Base Frequency Effect (Words)



Base_Frequency ■ High ■ Low

B Family Size Effect (Words)



Family_Size ■ Large ■ Small

Non-word Data

Use `complete.cases()` to find which rows had missing data in the model-relevant variables:

Anova Family Size

```
rt_nwords_cmpl %>%
  count(Complexity, Base_Frequency, Orthographic_Sensitivity)

|| # A tibble: 8 x 4
||   Complexity Base_Frequency Orthographic_Sensitivity     n
||   <chr>      <chr>          <chr>                <int>
|| 1 Complex    High             High                 522
|| 2 Complex    High             Low                  384
|| 3 Complex    Low              High                 617
|| 4 Complex    Low              Low                  458
|| 5 Simple     High             High                 754
|| 6 Simple     High             Low                  513
|| 7 Simple     Low              High                 752
|| 8 Simple     Low              Low                  607

temp <- rt_nwords_cmpl |> filter(is.na(Complexity) & is.na(Base_Frequency))
# write_csv(temp, "temp.csv")

anova_model_nwords_fs <- mixed(
  response_time ~ Complexity * Family_Size * Orthographic_Sensitivity +
    (1 | SubjID) +
    (1 | ItemID),
  data = rt_nwords_cmpl,
  method = "S")
anova_model_nwords_fs

|| Mixed Model Anova Table (Type 3 tests, S-method)
||
|| Model: response_time ~ Complexity * Family_Size * Orthographic_Sensitivity +
|| Model:      (1 | SubjID) + (1 | ItemID)
|| Data: rt_nwords_cmpl
||
||              Effect              df              F p.value
|| 1              Complexity 1, 4529.58 124.76 *** <.001
|| 2              Family_Size  1, 95.23   1.10   .297
|| 3      Orthographic_Sensitivity  1, 63.59   5.37 *   .024
|| 4      Complexity:Family_Size  1, 4525.57   0.92   .338
|| 5      Complexity:Orthographic_Sensitivity  1, 4512.75   0.96   .327
|| 6      Family_Size:Orthographic_Sensitivity  1, 4448.54   0.09   .770
|| 7      Complexity:Family_Size:Orthographic_Sensitivity  1, 4508.97   0.06   .809
|| ---
|| Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

m2 <- anova_model_nwords_fs$full_model # Extract the lmer model
ranova(m2) # Run random effects comparison

|| ANOVA-like table for random-effects: Single term deletions
||
|| Model:
|| response_time ~ Complexity + Family_Size + Orthographic_Sensitivity + (1 | SubjID) + (1 | ItemID) + Complexity:Family_Size + Complexity:Orthographic_Sensitivity
||      npar logLik  AIC    LRT Df Pr(>Chisq)
|| <none>      11 -28033 56089
|| (1 | SubjID)  10 -28862 57743 1656.71  1 < 2.2e-16 ***
|| (1 | ItemID)  10 -28105 56230  142.95  1 < 2.2e-16 ***
|| ---
|| Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

Main Findings

Effect	df	F	p.value
Complexity	1, 4529.58	124.76 ***	<.001
Orthographic_Sensitivity	1, 63.59	5.37 *	.024

Non-word complexity had a robust effect; complex non-words (e.g., pseudoderived forms) elicited longer response times than simple ones. Participants with higher **orthographic sensitivity** responded significantly faster overall, suggesting more efficient processing of letter patterns even in non-words. **Morphological family size** did not modulate non-word RTs, nor did it interact with complexity or orthographic sensitivity. Interpretation: In the absence of lexical representations, apparent “family size” (based on real-word analogues) does not measurably influence non-word recognition.

Plots

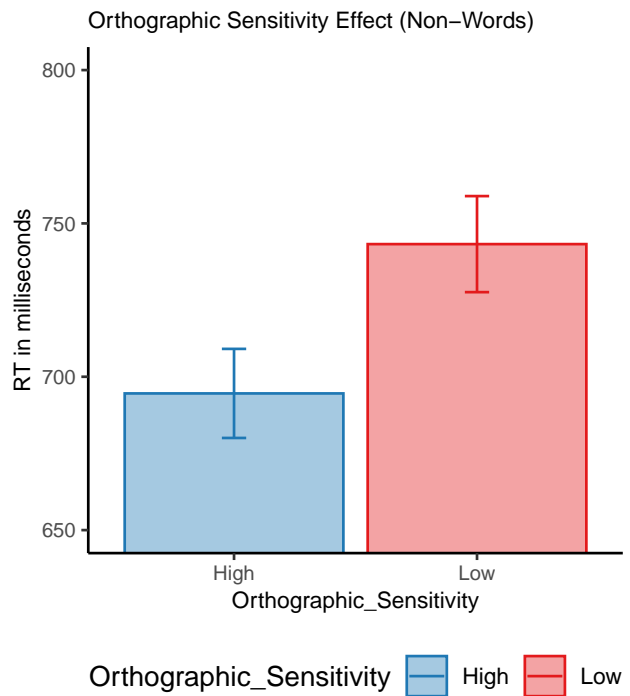
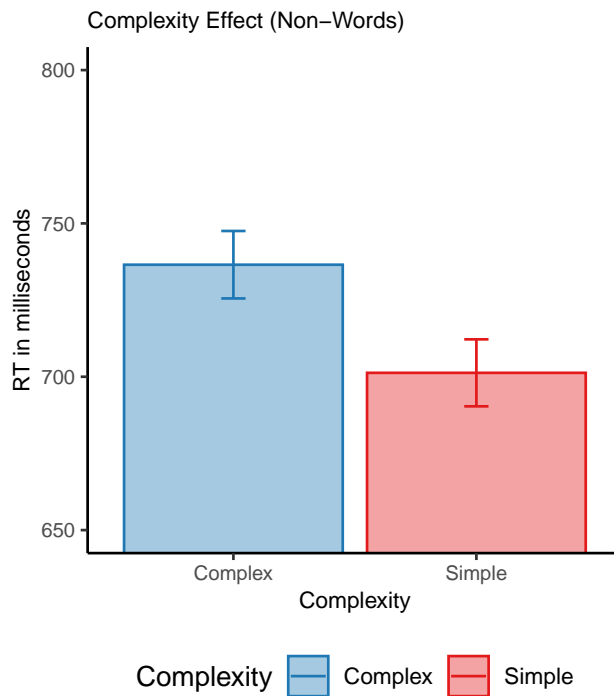
```
|| Complexity  emmean      SE df asymp.LCL asymp.UCL
```



```

|| Complex    736.5355 11.00010 Inf  714.9757  758.0953
|| Simple     701.2803 10.92368 Inf  679.8703  722.6904
||
|| Results are averaged over the levels of: Family_Size, Orthographic_Sensitivity
|| Degrees-of-freedom method: asymptotic
|| Confidence level used: 0.95
||
|| Orthographic_Sensitivity  emmean      SE  df  asymp.LCL  asymp.UCL
|| High                     694.5669 14.52299 Inf   666.1023   723.0314
|| Low                      743.2490 15.65686 Inf   712.5621   773.9359
||
|| Results are averaged over the levels of: Complexity, Family_Size
|| Degrees-of-freedom method: asymptotic
|| Confidence level used: 0.95

```



Anova Base Frequency

```
anova_model_nwords_bf <- mixed(
  response_time ~ Complexity * Base_Frequency * Orthographic_Sensitivity +
    (1 | SubjID) +
    (1 | ItemID),
  data = rt_nwords_cmpl,
  method = "S")
anova_model_nwords_bf

|| Mixed Model Anova Table (Type 3 tests, S-method)
||
|| Model: response_time ~ Complexity * Base_Frequency * Orthographic_Sensitivity +
|| Model: (1 | SubjID) + (1 | ItemID)
|| Data: rt_nwords_cmpl
||
||          Effect          df          F p.value
|| 1          Complexity 1, 4534.35 127.54 *** <.001
|| 2          Base_Frequency 1, 95.95 12.99 *** <.001
|| 3      Orthographic_Sensitivity 1, 63.60 5.32 * .024
|| 4      Complexity:Base_Frequency 1, 4535.47 4.26 * .039
|| 5      Complexity:Orthographic_Sensitivity 1, 4517.84 0.86 .353
|| 6      Base_Frequency:Orthographic_Sensitivity 1, 4452.86 0.04 .838
|| 7      Complexity:Base_Frequency:Orthographic_Sensitivity 1, 4515.97 0.35 .555
|| ---
|| Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

m3 <- anova_model_nwords_bf$full_model # Extract the lmer model
ranova(m3) # Run random effects comparison

|| ANOVA-like table for random-effects: Single term deletions
||
|| Model:
|| response_time ~ Complexity + Base_Frequency + Orthographic_Sensitivity + (1 | SubjID) + (1 | ItemID) + Complexity:Base_Frequency + Complexity:Orthographic_Sensitivity
||          npar logLik  AIC    LRT Df Pr(>Chisq)
|| <none>         11 -28026 56074
|| (1 | SubjID)    10 -28857 57734 1662.17 1 < 2.2e-16 ***
|| (1 | ItemID)    10 -28087 56195 122.89 1 < 2.2e-16 ***
|| ---
|| Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

Main Findings

Effect	df	F	p.value
Complexity	1, 4534.35	127.54 ***	<.001
Base_Frequency	1, 177.77	12.99 **	<.001
Orthographic_Sensitivity	1, 63.60	5.32 *	.024
Complexity:Base_Frequency	1, 4535.47	4.26 *	.039

- Complexity ($F = 127.5$, $p < .001$): complex > simple non-words → slower responses.
- Base Frequency ($F = 13.0$, $p < .001$): non-words derived from high-frequency bases were processed faster than those from low-frequency bases — an echo of lexical familiarity effects even though the items are illegal.
- Orthographic Sensitivity ($F = 5.32$, $p = .024$): same direction as before.
- Complexity × Base Frequency ($F = 4.26$, $p = .039$): The effect of complexity was larger for high-frequency bases than for low-frequency ones.

Interaction Effects: Complexity x Base_Frequency

```
# Estimated marginal means for the family_size x base frequency interaction
(emml <- emmeans(anova_model_nwords_bf, ~ Complexity * Base_Frequency))
```

Simple Contrasts

```
|| Complexity Base_Frequency emmean SE df asymp.LCL asymp.UCL
|| Complex High 751 11.6 Inf 728 773
|| Simple High 709 11.4 Inf 686 731
|| Complex Low 723 11.5 Inf 700 745
|| Simple Low 694 11.4 Inf 671 716
||
|| Results are averaged over the levels of: Orthographic_Sensitivity
|| Degrees-of-freedom method: asymptotic
|| Confidence level used: 0.95

# Get all pairwise contrasts
emml_contrasts <- contrast(emml, method = "pairwise", by = NULL, adjust = "none")
emml_contrasts
```

```

|| contrast          estimate    SE  df z.ratio p.value
|| Complex High - Simple High    42.3 4.68 Inf   9.033 <.0001
|| Complex High - Complex Low    28.0 6.95 Inf   4.021 0.0001
|| Complex High - Simple Low     57.1 6.78 Inf   8.429 <.0001
|| Simple High - Complex Low    -14.3 6.69 Inf  -2.136 0.0326
|| Simple High - Simple Low     14.9 6.51 Inf   2.285 0.0223
|| Complex Low - Simple Low     29.2 4.27 Inf   6.834 <.0001
||
|| Results are averaged over the levels of: Orthographic_Sensitivity
|| Degrees-of-freedom method: asymptotic
# Keep only the contrasts you want
# Simple effects of Complexity at each level of Base_Frequency
# Simple effects of Base_Frequency at each level of Complexity
keep <- c("Complex High - Simple High",
          "Complex Low - Simple Low",
          "Complex High - Complex Low",
          "Simple High - Simple Low")
(emmi1_contrasts_filtered <- subset(emmi1_contrasts, contrast %in% keep))

```

```

|| contrast          estimate    SE  df z.ratio p.value
|| Complex High - Simple High    42.3 4.68 Inf   9.033 <.0001
|| Complex High - Complex Low    28.0 6.95 Inf   4.021 0.0001
|| Simple High - Simple Low     14.9 6.51 Inf   2.285 0.0223
|| Complex Low - Simple Low     29.2 4.27 Inf   6.834 <.0001
||
|| Results are averaged over the levels of: Orthographic_Sensitivity
|| Degrees-of-freedom method: asymptotic

```

```

# Get Confidence Intervals
(emmi1_contrasts_filtered_ci <- confint(emmi1_contrasts_filtered))

```

```

|| contrast          estimate    SE  df asymp.LCL asymp.UCL
|| Complex High - Simple High    42.3 4.68 Inf   33.09    51.4
|| Complex High - Complex Low    28.0 6.95 Inf   14.34    41.6
|| Simple High - Simple Low     14.9 6.51 Inf    2.12    27.6
|| Complex Low - Simple Low     29.2 4.27 Inf   20.81    37.5
||
|| Results are averaged over the levels of: Orthographic_Sensitivity
|| Degrees-of-freedom method: asymptotic
|| Confidence level used: 0.95

```

```

# Get effect sizes
# Get all pairwise effect sizes
effs1 <- eff_size(emmi1, sigma = sigma(m3), edf = df.residual(m3))
effs1

```

```

|| contrast          effect.size    SE  df asymp.LCL asymp.UCL
|| Complex High - Simple High    0.414 0.0460 Inf   0.3235    0.5038
|| Complex High - Complex Low    0.274 0.0681 Inf   0.1402    0.4073
|| Complex High - Simple Low     0.559 0.0666 Inf   0.4288    0.6899
|| Simple High - Complex Low    -0.140 0.0655 Inf  -0.2682   -0.0115
|| Simple High - Simple Low     0.146 0.0638 Inf   0.0207    0.2707
|| Complex Low - Simple Low     0.286 0.0419 Inf   0.2035    0.3677
||
|| Results are averaged over the levels of: Orthographic_Sensitivity
|| sigma used for effect sizes: 102.2
|| Degrees-of-freedom method: inherited from asymptotic when re-gridding
|| Confidence level used: 0.95

```

```

# Remove the two redundant rows (rows 3 and 4)
(effs1_filtered <- subset(effs1, !contrast %in% c("Complex High - Simple Low",
          "Simple High - Complex Low")))

```

```

|| contrast          effect.size    SE  df asymp.LCL asymp.UCL
|| Complex High - Simple High    0.414 0.0460 Inf   0.3235    0.504
|| Complex High - Complex Low    0.274 0.0681 Inf   0.1402    0.407
|| Simple High - Simple Low     0.146 0.0638 Inf   0.0207    0.271
|| Complex Low - Simple Low     0.286 0.0419 Inf   0.2035    0.368
||
|| Results are averaged over the levels of: Orthographic_Sensitivity
|| sigma used for effect sizes: 102.2
|| Degrees-of-freedom method: inherited from asymptotic when re-gridding
|| Confidence level used: 0.95

```

```

# Interaction contrasts (difference-of-differences)
# Compare base frequency effect in large vs small family
contrast(emmi1, interaction = "pairwise", by = NULL, adjust = "holm")

```

Interaction Contrasts

```

|| Complexity_pairwise Base_Frequency_pairwise estimate    SE  df z.ratio p.value

```

```

|| Complex - Simple      High - Low                13.1 6.34 Inf    2.063 0.0391
||
|| Results are averaged over the levels of: Orthographic_Sensitivity
|| Degrees-of-freedom method: asymptotic
# Get confidence intervals, for each base frequency effect for each family size and then for interaction effect
confint(contrast(emmeans(m3, ~ Complexity | Base_Frequency), "pairwise"))

|| Base_Frequency = High:
|| contrast      estimate    SE df asymp.LCL asymp.UCL
|| Complex - Simple      42.3 4.68 Inf      33.1      51.4
||
|| Base_Frequency = Low:
|| contrast      estimate    SE df asymp.LCL asymp.UCL
|| Complex - Simple      29.2 4.27 Inf      20.8      37.5
||
|| Results are averaged over the levels of: Orthographic_Sensitivity
|| Degrees-of-freedom method: asymptotic
|| Confidence level used: 0.95
confint(contrast(emm1, interaction = c("pairwise", "pairwise")))

|| Complexity_pairwise Base_Frequency_pairwise estimate    SE df asymp.LCL asymp.UCL
|| Complex - Simple      High - Low                13.1 6.34 Inf    0.655 25.5
||
|| Results are averaged over the levels of: Orthographic_Sensitivity
|| Degrees-of-freedom method: asymptotic
|| Confidence level used: 0.95

```

Table 4. Estimated Marginal Means and Pairwise Contrasts for the Complexity × Base Frequency Interaction

Contrast	Delta RT (ms)	p	Effect size (d)	95% CI for Delta RT (ms)	Interpretation
Complex High (751) – Simple High (709)	+42.3	< .001	0.41	[33.1, 51.4]	Strong complexity cost at high base frequency
Complex Low (723) – Simple Low (694)	+29.2	< .001	0.29	[20.8, 37.5]	Moderate complexity cost at low base frequency
Complex High (751) – Complex Low (723)	+28.0	.0001	0.27	[14.3, 41.6]	Complex items slower when based on high-frequency stems
Simple High (709) – Simple Low (694)	+14.9	.022	0.15	[2.1, 27.6]	Small frequency benefit among simple items

Table 5. Summary of Main Effects and Interactions

Effect	Direction / Pattern	Interpretation
Complexity	Complex > Simple across all conditions	Morphological structure slows non-word rejection (decomposition cost).
Base Frequency	High > Low (faster RTs)	Familiar letter sequences facilitate processing even for non-words.
Complexity × Base Frequency	Larger complexity cost for high-frequency bases (Delta ~ 13 ms)	Morphological activation stronger for familiar bases.
Family Size	n.s.	No measurable influence; non-words lack real morphological families.
Orthographic Sensitivity	High > Low (faster RTs overall)	Readers with higher orthographic sensitivity are generally more efficient.

Main Effects Plots

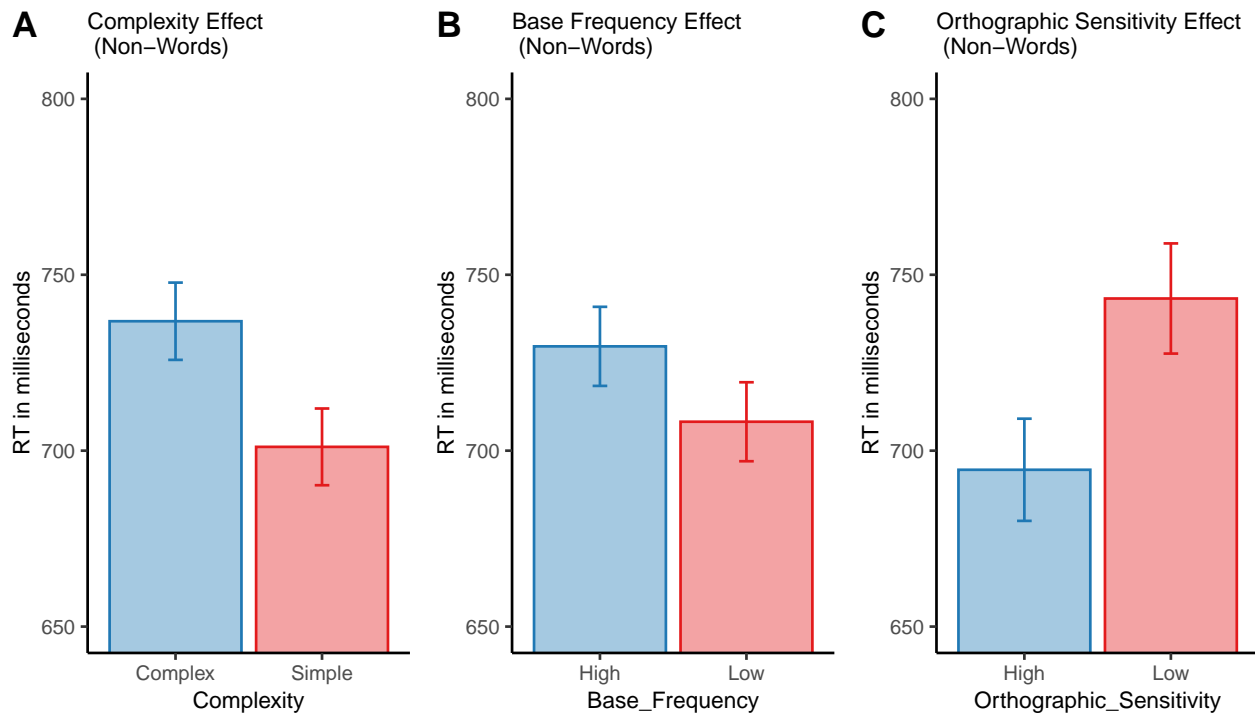
```

|| Complexity      emmean      SE df asymp.LCL asymp.UCL
|| Complex      736.7913 10.98256 Inf 715.2659 758.3168
|| Simple      701.0784 10.90537 Inf 679.7042 722.4525
||
|| Results are averaged over the levels of: Base_Frequency, Orthographic_Sensitivity
|| Degrees-of-freedom method: asymptotic
|| Confidence level used: 0.95

|| Base_Frequency      emmean      SE df asymp.LCL asymp.UCL
|| High      729.6473 11.23700 Inf 707.6232 751.6715
|| Low      708.2224 11.22244 Inf 686.2268 730.2180
||
|| Results are averaged over the levels of: Complexity, Orthographic_Sensitivity
|| Degrees-of-freedom method: asymptotic
|| Confidence level used: 0.95

|| Orthographic_Sensitivity      emmean      SE df asymp.LCL asymp.UCL
|| High      694.6539 14.52304 Inf 666.1892 723.1185
|| Low      743.2159 15.66167 Inf 712.5196 773.9122
||
|| Results are averaged over the levels of: Complexity, Base_Frequency
|| Degrees-of-freedom method: asymptotic
|| Confidence level used: 0.95

```



Interaction Plots

```
# Compute means for each level of Complexity and Base Frequency
(emm_nw_bf_cmpXbf_df <- as.data.frame(emmeans(anova_model_nwords_bf, ~ Complexity * Base_Frequency)))
```

```
|| Complexity Base_Frequency emmean SE df asymp.LCL asymp.UCL
|| Complex High 750.7745 11.56040 Inf 728.1165 773.4324
|| Simple High 708.5202 11.39462 Inf 686.1871 730.8532
|| Complex Low 722.8082 11.47929 Inf 700.3092 745.3072
|| Simple Low 693.6366 11.36764 Inf 671.3564 715.9167
||
|| Results are averaged over the levels of: Orthographic_Sensitivity
|| Degrees-of-freedom method: asymptotic
|| Confidence level used: 0.95
```

```
p8<-emm_nw_bf_cmpXbf_df |> ggplot(aes(x = Base_Frequency, y = emmean,
color = Complexity, group = Complexity)) +
  geom_line(position = position_dodge(0.2)) +
  geom_point(position = position_dodge(0.2)) +
  geom_errorbar(aes(ymin = emmean - SE, ymax = emmean + SE),
width = 0.1, position = position_dodge(0.2)) +
  labs(x = "Base Frequency", y = "RT in milliseconds",
color = "Complexity",
title = "Complexity x Base Frequency") +
  scale_color_custom() +
  scale_fill_custom()
# p8

p9 <- emm_nw_bf_cmpXbf_df |> ggplot(aes(x = Complexity, y = emmean,
color = Base_Frequency, group = Base_Frequency)) +
  geom_line(position = position_dodge(0.2)) +
  geom_point(position = position_dodge(0.2)) +
  geom_errorbar(aes(ymin = emmean - SE, ymax = emmean + SE),
width = 0.1, position = position_dodge(0.2)) +
  labs(x = "Complexity", y = "RT in milliseconds",
color = "Base Frequency",
title = "Base Frequency x Complexity") +
  scale_color_custom() +
  scale_fill_custom()
# p9

plot_grid(p8, p9, ncol = 2, labels = "AUTO")
```

