

Data analysis Morphological masked priming experiment on L1-ITA, L2-ENG bilingual speakers/readers

Paper titled *Masked morphological priming tracks the development of a fully mature lexical system in L2* - Submitted to JML in June 2019

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Contents

clean WS, set WD and load data	3
Participants	4
Stimuli	4
Target features, ITA	5
Prime features, ITA	5
Outliers trimming, ITA	6
Outliers trimming, ENG	9
Raw means	12
Modelling, ITA	13
Summary of the results for ITA dataset	16
Modelling, ENG	17
Summary of the results for ENG dataset	20
Cross language interaction	20
Proficiency modelling	24
Phonemic fluency	24
Phonemic comprehension	25
Morphologic awareness	25

Spelling	25
Reading comprehension	26
Vocabulary	26
Oral comprehension	27
Does proficiency specifically interact with priming?	27
Phonemic fluency	27
Phonemic comprehension	29
Spelling	32
Reading comprehension	34
Vocabulary	36
Oral comprehension	37
AoA	39
correlation	39
scores, modelling	40
OSC	51
modelling	52
Plots	52
Figure 1 - plot of estimated RTs	52
Figure 2 - Participants' score distributions for each proficiency subtest	54
Figure 3 - Interaction by phonemic comprehension by relatedness and morphotype	55
Figure 4 - Interaction by morphological awareness/comprehension by relatedness and morphotype	56
Figure 5 - Interaction by vocabulary by relatedness and morphotype	57
Figure 6 - Interaction by phonemic Fluency by relatedness and morphotype	58
Figure 7 - Scores distributions in the AoA questionnaire	59
Figure 8 - Interaction by AoA1 by relatedness and morphotype	60
Figure 9 - Interaction by AoA2 by relatedness and morphotype	61
Figure 10 - Interaction by AoA5 by relatedness and morphotype	62
Figure 11 - OSC by morphotype	63
Figure 12 - OSC by phonemic fluency by relatedness	64
Figure 13 - OSC by vocabulary by relatedness	65
Figure 14 - morphComprehension by relatedness	66

This script takes preprocessed data and produces all the analyses that are reported in the paper.

clean WS, set WD and load data

Set your local working directory. This should be (and is assumed to be in the rest of the code) the highest point in your local folder:

```
localGitDir <- 'C:/Users/eva_v/Documents/GitHub/M2-maskedprimingBilinguals'
setwd(localGitDir);
```

This script works on the outcome of preProcessing.R, which you can upload here:

```
read.table(paste(localGitDir, '/preprocessedData.txt', sep = ''), header = T, sep='\t', dec='.',
           colClasses=c("prime"="factor",
                        "target"="factor",
                        "lexicality"="factor",
                        "morphType"="factor",
                        "relatedness"="factor",
                        "aoa3.context"="factor",
                        "aoa4.contextMultling"="factor",
                        "aoa6.otherLang"="factor",
                        "language"="factor",
                        "gender"="factor",
                        "handedness"="factor"
                       )) -> masterFile;

head(masterFile);
summary(masterFile);
```

Load packages and create functions

```
library(ggplot2);
library(rms);
library(effects);
library(plyr);
library(corrplot);
library(lme4);
library(ggpubr);

inv <- function(x) {-1000/x};
```

GLMMs takes a lot of time to run, depending on the machine and the dataset. We have saved the models that successfully converged in a folder.

```
df <- list.files(paste(localGitDir, "/LMMs and GLMMs/", sep = ""));
length(df);
```

```
## [1] 45
```

```
for (i in 1:length(df)){
  gsub(".rds$", "", df[i]) -> id
  assign(id, data.frame())
  readRDS(paste(localGitDir, "/LMMs and GLMMs/", df[i], sep = "")) -> temp
```

```

    assign(paste0(id), temp)
  };
  rm(temp)

```

Participants

```
temp <- unique(masterFile[,c('subject','age','gender','handedness')]);
```

Number of participants: 81

age, education and handedness:

```
summary(temp)
```

```
##      subject      age      gender handedness
##  Min.   : 1.00  Min.   :18.00  Female:56   Left  :11
##  1st Qu.:21.00  1st Qu.:22.00  Male  :28   Right:73
##  Median :41.50  Median :24.00
##  Mean   :41.65  Mean   :24.32
##  3rd Qu.:62.25  3rd Qu.:26.00
##  Max.   :84.00  Max.   :34.00
```

Stimuli

```
temp <- unique(masterFile[,c('target','prime','lexicality','morphType','relatedness','freqTarget','freqPrime')]);
summary(temp);
```

```
##      target      prime      lexicality morphType relatedness
##  abito   : 2  abbazia   : 1  nonword:300  op   :200  ctrl:450
##  acid    : 2  ability   : 1  word   :600  or   :200  rel :450
##  acre    : 2  abitudine: 1                      tr   :200
##  adopt   : 2  absence   : 1                      NA's:300
##  again   : 2  abstain   : 1
##  agree   : 2  abstract  : 1
##  (Other):888  (Other)  :894
##  freqTarget  freqPrime  lengthTarget  lengthPrime
##  Min.   :1.367  Min.   :1.170  Min.   :3.00  Min.   : 5.000
##  1st Qu.:3.382  1st Qu.:2.630  1st Qu.:4.00  1st Qu.: 6.000
##  Median :3.913  Median :3.155  Median :5.00  Median : 7.000
##  Mean   :3.875  Mean   :3.233  Mean   :4.92  Mean   : 7.423
##  3rd Qu.:4.446  3rd Qu.:3.860  3rd Qu.:5.00  3rd Qu.: 8.000
##  Max.   :5.840  Max.   :6.290  Max.   :8.00  Max.   :13.000
##  NA's    :300   NA's    :2
##  nTarget     nPrime     language
##  Min.   : 0.00  Min.   : 0.000  eng:450
##  1st Qu.: 5.00  1st Qu.: 1.000  ita:450
##  Median :10.00  Median : 2.000
```

```
## Mean :13.43 Mean : 3.131
## 3rd Qu.:19.00 3rd Qu.: 4.000
## Max. :54.00 Max. :36.000
##
```

Target features, ITA

Frequency of the targets, mean and sd

```
## morphType freqTarget
## 1 op 3.63032
## 2 or 3.94728
## 3 tr 3.96898
```

```
## morphType freqTarget
## 1 op 0.8779550
## 2 or 0.8523670
## 3 tr 0.6718303
```

Length of the targets, mean and sd

```
## morphType lengthTarget
## 1 op 5.08
## 2 or 4.94
## 3 tr 5.16
```

```
## morphType lengthTarget
## 1 op 0.8490042
## 2 or 0.8855746
## 3 tr 1.0704478
```

Coltheart's N, mean and sd

```
## morphType nTarget
## 1 op 20.12
## 2 or 21.56
## 3 tr 18.14
```

```
## morphType nTarget
## 1 op 11.91492
## 2 or 13.45274
## 3 tr 11.32265
```

Prime features, ITA

Frequency of the primes, mean and sd

```
## relatedness morphType freqPrime
## 1 ctrl op 3.095800
## 2 rel op 3.157653
```

## 3	ctrl	or	3.198040
## 4	rel	or	3.228920
## 5	ctrl	tr	2.913200
## 6	rel	tr	2.923460

##	relatedness	morphType	freqPrime
## 1	ctrl	op	0.8518056
## 2	rel	op	0.7863899
## 3	ctrl	or	0.6784778
## 4	rel	or	0.6984253
## 5	ctrl	tr	0.6843910
## 6	rel	tr	0.8404621

Length of the primes, mean and sd

##	relatedness	morphType	lengthPrime
## 1	ctrl	op	7.96
## 2	rel	op	7.96
## 3	ctrl	or	7.52
## 4	rel	or	7.52
## 5	ctrl	tr	7.70
## 6	rel	tr	7.70

##	relatedness	morphType	lengthPrime
## 1	ctrl	op	1.211509
## 2	rel	op	1.211509
## 3	ctrl	or	1.182181
## 4	rel	or	1.182181
## 5	ctrl	tr	1.249490
## 6	rel	tr	1.249490

Coltheart's N, mean and sd

##	relatedness	morphType	nPrime
## 1	ctrl	op	3.84
## 2	rel	op	3.52
## 3	ctrl	or	3.54
## 4	rel	or	4.24
## 5	ctrl	tr	3.82
## 6	rel	tr	3.62

##	relatedness	morphType	nPrime
## 1	ctrl	op	2.937165
## 2	rel	op	2.628261
## 3	ctrl	or	2.525058
## 4	rel	or	6.169576
## 5	ctrl	tr	2.973969
## 6	rel	tr	2.905940

Outliers trimming, ITA

```
subset(masterFile, language=="ita") -> masterFileIta;
```

The following code generates target and sbj means and SDs, and the outlier graphs in the file 'ita.jpg'

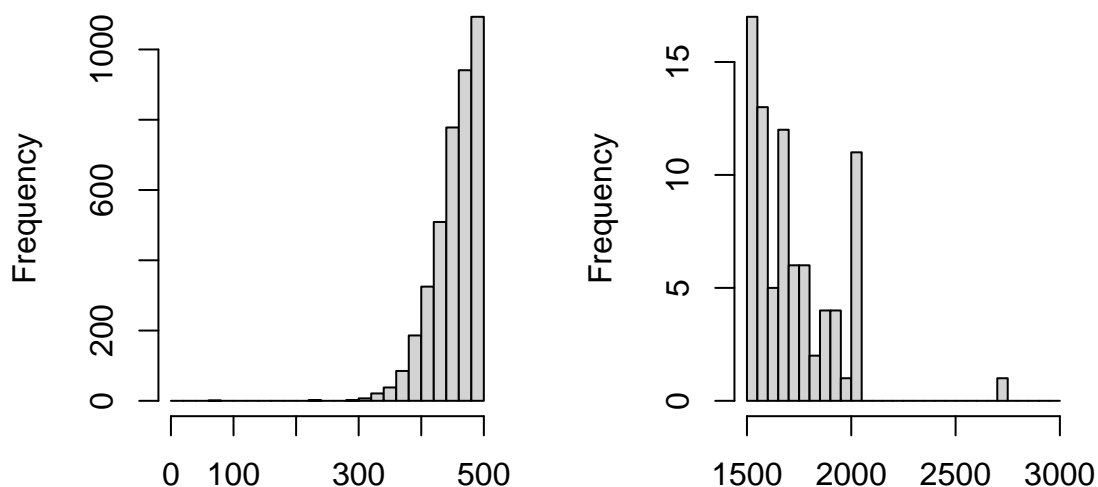
```
sbj.id <- masterFileIta$subject;
acc <- masterFileIta$accuracy;
lexicality <- tolower(masterFileIta$lexicality);
target <- masterFileIta$target;
rt <- masterFileIta$rt;

source(paste(localGitDir, "/tools/diagnostics.R", sep=''));
outlierGraphStore <- 'Desktop';
diagnostics.f(rt = rt, acc = acc, sbj.id = sbj.id, target = target, lexicality = lexicality, paste(outl.
rm(outlierGraphStore, rt, target, lexicality, acc, sbj.id);
```

Based on the graphs in 'ita.jpg': we exclude **sbj 2** and **31** for an abnormal error rate on nonwords (<80%); and target words **GUANO**, **UGGIA** and **VELLO** with abnormally low accuracy (<60%). **sbj 15** is excluded because s/he reported having seen the primes.

Individual RTs distribution seems fine, but let's check the tails more carefully:

```
par(mfrow=c(1,2))
hist(masterFileIta$rt[masterFileIta$rt<500], breaks=seq(0,500,20), main = " ", xlab = " ");
hist(masterFileIta$rt[masterFileIta$rt>1500], breaks=seq(1500,3000,50), main = " ", xlab = " ");
```



```
par(mfrow=c(1,1))
```

Based on these graph we cut distributions at **2500ms** and **280ms**

```
dataItaAcc <- subset(masterFileIta, lexicality=="word");
dataItaTemp <- subset(dataItaAcc, accuracy==1);
dataIta <- subset(dataItaTemp, rt>280 & rt<2000 & subject!=15 & subject!=2 & subject!=31 & target!= "gu")
```

- Number of datapoints trimmed: 526
- Percentage of datapoints trimmed: 4.56%
- Number of datapoints left: 11009

Summary of the dataset trimmed:

```
summary(dataIta)
```

```
##      subject      age      gender handedness  trialCount
## Min.   : 1.00  Min.   :18.00  Female:7362  Left :1284  Min.   : 1.0
## 1st Qu.:22.00  1st Qu.:22.00  Male  :3647  Right:9725  1st Qu.: 77.0
## Median :43.00  Median :24.00                                Median :150.0
## Mean   :42.19  Mean   :24.44                                Mean   :150.5
## 3rd Qu.:62.00  3rd Qu.:27.00                                3rd Qu.:225.0
## Max.   :84.00  Max.   :34.00                                Max.   :300.0
##
##      rt      resp  trialType      lexicality  morphType
## Min.   : 284.0  Min.   :2  Length:11009  nonword: 0  op:3717
## 1st Qu.: 499.0  1st Qu.:2  Class :character  word  :11009  or:3621
## Median : 564.0  Median :2  Mode  :character              tr:3671
## Mean   : 594.5  Mean   :2
## 3rd Qu.: 652.0  3rd Qu.:2
## Max.   :1960.0  Max.   :2
##
##      target      prime      relatedness  freqTarget  freqPrime
## abito   : 78  abitudine: 39  ctrl:5498  Min.   :1.367  Min.   :1.191
## ballo   : 78  albergo  : 39  rel :5511  1st Qu.:3.435  1st Qu.:2.562
## banca   : 78  alleanza : 39              Median :3.989  Median :3.039
## cambio  : 78  areola   : 39              Mean   :3.915  Mean   :3.093
## corda   : 78  ballatoio: 39              3rd Qu.:4.450  3rd Qu.:3.606
## corte   : 78  bancario : 39              Max.   :5.671  Max.   :4.909
## (Other):10541 (Other) :10775              NA's    :37
## lengthTarget  lengthPrime      nTarget      nPrime
## Min.   :3.000  Min.   : 5.000  Min.   : 0.00  Min.   : 0.000
## 1st Qu.:4.000  1st Qu.: 7.000  1st Qu.:11.00  1st Qu.: 2.000
## Median :5.000  Median : 8.000  Median :18.00  Median : 3.000
## Mean   :5.064  Mean   : 7.756  Mean   :20.13  Mean   : 3.723
## 3rd Qu.:6.000  3rd Qu.: 9.000  3rd Qu.:30.00  3rd Qu.: 5.000
## Max.   :8.000  Max.   :10.000  Max.   :47.00  Max.   :36.000
##
##      rotation      language      oscTarget      accuracy phonemicFluency
## Length:11009      eng: 0  Min.   : NA  Min.   :1  Min.   : 0.00
## Class :character  ita:11009  1st Qu.: NA  1st Qu.:1  1st Qu.:17.00
## Mode  :character              Median : NA  Median :1  Median :23.00
```



```
##                               Mean   :NaN      Mean   :1      Mean   :22.52
##                               3rd Qu.: NA       3rd Qu.:1      3rd Qu.:27.00
##                               Max.    : NA       Max.    :1      Max.    :45.00
##                               NA's    :11009
## phonemicComprehension morphComprehension spelling readingComprehension
## Min.    : 0.000      Min.    : 4.000      Min.    : 0.000      Min.    :1.000
## 1st Qu.: 8.000      1st Qu.: 8.000      1st Qu.: 4.000      1st Qu.:3.000
## Median : 9.000      Median : 9.000      Median : 7.000      Median :5.000
## Mean    : 8.828      Mean    : 8.929      Mean    : 8.155      Mean    :4.564
## 3rd Qu.:10.000      3rd Qu.:10.000      3rd Qu.:11.000      3rd Qu.:6.000
## Max.    :13.000      Max.    :10.000      Max.    :18.000      Max.    :7.000
##
## vocabulary oralComprehension aoa1.Aoa aoa2.usage
## Min.    : 8.00      Min.    :1.000      Min.    : 0.000      Min.    :1.00
## 1st Qu.:14.00      1st Qu.:3.000      1st Qu.: 6.000      1st Qu.:2.00
## Median :15.00      Median :5.000      Median : 6.000      Median :3.00
## Mean    :15.28      Mean    :4.593      Mean    : 6.544      Mean    :3.12
## 3rd Qu.:17.00      3rd Qu.:6.000      3rd Qu.: 8.000      3rd Qu.:4.00
## Max.    :19.00      Max.    :6.000      Max.    :15.000      Max.    :5.00
##
## aoa3.context aoa4.contextMultling aoa5.selfRatedProf aoa6.otherLang
## home :1433 no :8446 Min.    :1.00 no :2673
## school:9576 yes:2563 1st Qu.:3.00 yes:8336
## Median :4.00
## Mean    :3.54
## 3rd Qu.:4.00
## Max.    :5.00
##
```

Outliers trimming, ENG

```
subset(masterFile, language=="eng") -> masterFileEng;
```

The following code generates target and sbj means and SDs, and the outlier graphs in the file 'eng.jpg'

```
outlierGraphStore <- 'Desktop';
sbj.id <- masterFileEng$subject;
acc <- masterFileEng$accuracy;
lexicality <- masterFileEng$lexicality;
lexicality <- tolower(masterFileEng$lexicality);
target <- masterFileEng$target;
rt <- masterFileEng$rt;

diagnostics.f(rt = rt, acc = acc, sbj.id = sbj.id, target = target, lexicality = lexicality, paste(outl.
rm(outlierGraphStore, rt, target, lexicality, acc, sbj.id);
```

sbj 26 likely confused YES/NO buttons. Let's check the frequency effect, just to confirm:

```
cor(masterFileEng[masterFileEng$subject==26 & masterFileEng$lexicality=='word', c('rt','freqTarget')], v
```

```
##          rt freqTarget
## rt      1.0000000 -0.2116401
## freqTarget -0.2116401  1.0000000
```

```
masterFileEng$accuracy[masterFileEng$subject==26] <- car::recode(masterFileEng$accuracy[masterFileEng$subject==26], "1.0000000")
```

ok, we can now rerun diagnostics:

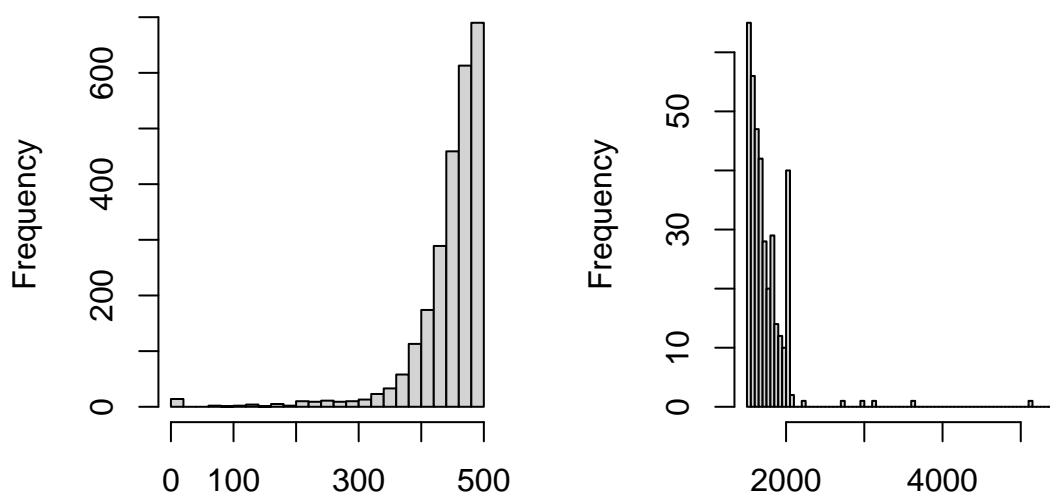
```
outlierGraphStore <- 'Desktop';
sbj.id <- masterFileEng$subject;
acc <- masterFileEng$accuracy;
lexicity <- masterFileEng$lexicity;
lexicity <- tolower(masterFileEng$lexicity);
target <- masterFileEng$target;
rt <- masterFileEng$rt;

diagnostics.f(rt = rt, acc = acc, sbj.id = sbj.id, target = target, lexicity = lexicity, paste(outlierGraphStore, sbj.id, ".jpg"),
rm(outlierGraphStore, rt, target, lexicity, acc, sbj.id);
```

Based on the graphs in 'eng.jpg': we exclude **sbj 22** for a very atypical performance (average **RT<200ms**). We would exclude no target, even though some of them do elicit bad performance—the distribution is very continuous, no sign of glaring outliers. Plus, this is L2, so low performance is to be expected. **sbj 15** and **43** reported having seen the primes.

Individual RTs distribution seems fine, but let's check the tails more carefully:

```
par(mfrow=c(1,2))
hist(masterFileEng$rt[masterFileEng$rt<500], breaks=seq(0,500,20), main = "", xlab = "");
hist(masterFileEng$rt[masterFileEng$rt>1500], breaks=seq(1500,5500,50), main = "", xlab = "");
```



- First graph: very continuous towards zero; perhaps some technical problem with the response box? Weird though, only in English. Anyway, deflection in the curve around 300ms, so let's cut there.
- Second graph: clear outliers over 2000ms

```
dataEngAcc <- subset(masterFileEng, lexicality=="word");
dataEngTemp <- subset(dataEngAcc, accuracy==1);
dataEng <- subset(dataEngTemp, rt>300 & rt<2000 & subject!=15 & subject!=22 & subject!=43);
```

- Number of datapoints trimmed: 281
- Percentage of datapoints trimmed: 3.05%
- Number of datapoints left: 8938

Summary of the trimmed dataset:

```
summary(dataEng);
```

```
##      subject      age      gender handedness  trialCount
## Min.   : 1.00   Min.   :18.00 Female:6170 Left : 907   Min.   : 1.0
## 1st Qu.:20.00  1st Qu.:22.00 Male :2768 Right:8031 1st Qu.: 74.0
## Median :42.00  Median :24.00          :          :          Median :150.0
## Mean   :41.76  Mean   :24.19          :          :          Mean   :149.8
## 3rd Qu.:63.00  3rd Qu.:26.00          :          :          3rd Qu.:224.0
## Max.   :84.00  Max.   :34.00          :          :          Max.   :300.0
##
##      rt      resp      trialType      lexicality morphType
## Min.   : 312.0   Min.   :0.000 Length:8938 nonword: 0 op:2946
## 1st Qu.: 544.0   1st Qu.:2.000 Class :character word :8938 or:2762
## Median : 622.0   Median :2.000 Mode :character          tr:3230
## Mean   : 672.1   Mean   :1.986
## 3rd Qu.: 742.0   3rd Qu.:2.000
## Max.   :1998.0   Max.   :2.000
##
##      target      prime relatedness freqTarget freqPrime
## again : 78   against : 40   ctrl:4388 Min.   :1.650 Min.   :1.170
## angel : 78   angelic : 40   rel :4550 1st Qu.:3.555 1st Qu.:2.820
## legend : 78   cloudless: 40          Median :4.050 Median :3.410
## unit : 78   dreamer : 40          Mean :4.058 Mean :3.472
## blood : 77   drunkard : 40          3rd Qu.:4.680 3rd Qu.:4.080
## fruit : 77   extract : 40          Max.   :5.840 Max.   :6.290
## (Other):8472 (Other) :8698
## lengthTarget lengthPrime nTarget nPrime
## Min.   :3.000 Min.   : 5.000 Min.   : 0.000 Min.   : 0.000
## 1st Qu.:4.000 1st Qu.: 6.000 1st Qu.: 4.000 1st Qu.: 0.000
## Median :5.000 Median : 7.000 Median : 7.000 Median : 1.000
## Mean   :4.828 Mean   : 7.215 Mean   : 8.785 Mean   : 2.485
## 3rd Qu.:5.000 3rd Qu.: 8.000 3rd Qu.:12.000 3rd Qu.: 3.000
## Max.   :7.000 Max.   :13.000 Max.   :44.000 Max.   :24.000
##
##      rotation      language oscTarget accuracy phonemicFluency
## Length:8938 eng:8938 Min.   : -0.0940 Min.   :1 Min.   : 0.00
## Class :character ita: 0 1st Qu.: 0.2630 1st Qu.:1 1st Qu.:17.00
## Mode :character Median : 0.6230 Median :1 Median :23.00
```

```
##              Mean   : 0.5767   Mean   :1   Mean   :22.61
##              3rd Qu.: 0.8990   3rd Qu.:1   3rd Qu.:27.00
##              Max.    : 1.0000   Max.    :1   Max.    :45.00
##              NA's    :155
## phonemicComprehension morphComprehension   spelling   readingComprehension
## Min.    : 0.000      Min.    : 4.000      Min.    : 0.000   Min.    :1.000
## 1st Qu.: 8.000      1st Qu.: 8.000      1st Qu.: 5.000   1st Qu.:3.000
## Median : 9.000      Median : 9.000      Median : 8.000   Median :5.000
## Mean    : 8.866      Mean    : 8.975      Mean    : 8.342   Mean    :4.478
## 3rd Qu.:11.000      3rd Qu.:10.000     3rd Qu.:12.000   3rd Qu.:6.000
## Max.    :13.000      Max.    :10.000     Max.    :18.000   Max.    :7.000
##
##      vocabulary      oralComprehension      aoa1.Aoa      aoa2.usage
## Min.    : 8.00      Min.    :1.00      Min.    : 0.000   Min.    :1.000
## 1st Qu.:14.00      1st Qu.:3.00      1st Qu.: 6.000   1st Qu.:2.000
## Median :16.00      Median :5.00      Median : 6.000   Median :3.000
## Mean    :15.34      Mean    :4.59      Mean    : 6.433   Mean    :3.174
## 3rd Qu.:17.00      3rd Qu.:6.00      3rd Qu.: 8.000   3rd Qu.:4.000
## Max.    :19.00      Max.    :6.00      Max.    :15.000   Max.    :5.000
##
##      aoa3.context  aoa4.contextMultling  aoa5.selfRatedProf  aoa6.otherLang
## home   :1231      no :6811              Min.    :1.000      no :2062
## school:7707      yes:2127              1st Qu.:3.000      yes:6876
##                                     Median :3.000
##                                     Mean    :3.526
##                                     3rd Qu.:4.000
##                                     Max.    :5.000
##
```

Raw means

- Mean accuracy ITA: 95%
- Mean RT ITA: 594.51 ms
- Mean accuracy ENG: 76%
- Mean RT ENG: 672.1 ms

Mean RT and sd by relatedness and morphType - ITA dataset:

```
aggregate(rt ~ relatedness + morphType, FUN=mean, data=dataIta);
```

```
##      relatedness morphType      rt
## 1          ctrl         op 614.4428
## 2           rel         op 598.8104
## 3          ctrl         or 607.0192
## 4           rel         or 606.5746
## 5          ctrl         tr 589.1723
## 6           rel         tr 551.3402
```

```
aggregate(rt ~ relatedness + morphType, FUN=sd, data=dataIta);
```

```
##      relatedness morphType      rt
```

```
## 1      ctrl      op 150.3067
## 2      rel      op 157.5980
## 3      ctrl      or 145.9696
## 4      rel      or 155.8299
## 5      ctrl      tr 131.0569
## 6      rel      tr 126.2728
```

Mean RT and sd by relatedness and morphotype - ENG dataset:

```
aggregate(rt ~ relatedness + morphType, FUN=mean, data=dataEng);
```

```
##   relatedness morphType      rt
## 1         ctrl      op 685.2774
## 2         rel      op 666.6475
## 3         ctrl      or 703.1108
## 4         rel      or 688.2460
## 5         ctrl      tr 667.2606
## 6         rel      tr 630.8267
```

```
aggregate(rt ~ relatedness + morphType, FUN=sd, data=dataEng);
```

```
##   relatedness morphType      rt
## 1         ctrl      op 196.6292
## 2         rel      op 214.0158
## 3         ctrl      or 214.4669
## 4         rel      or 207.4222
## 5         ctrl      tr 177.6199
## 6         rel      tr 183.6509
```

Modelling, ITA

```
dataIta$morphType <- relevel(dataIta$morphType, "or");
contrasts(dataIta$relatedness);
```

```
##      rel
## ctrl   0
## rel    1
```

```
contrasts(dataIta$morphType);
```

```
##      op tr
## or   0  0
## op   1  0
## tr   0  1
```

```
itaglmer0<- glmer(rt ~ 1 + (1|subject) + (1|target), data= dataIta, family=Gamma(link="identity"));
```

```
itaglmer1<- glmer(rt ~ trialCount + rotation + (1|subject) + (1|target), data= dataIta, family=Gamma(link="log"))
```

```
anova(itaglmer0, itaglmer1);
```

```
## Data: dataIta
## Models:
## itaglmer0: rt ~ 1 + (1 | subject) + (1 | target)
## itaglmer1: rt ~ trialCount + rotation + (1 | subject) + (1 | target)
##          npar      AIC      BIC logLik deviance Chisq Df Pr(>Chisq)
## itaglmer0      4 134591 134620 -67291   134583
## itaglmer1      6 134593 134637 -67291   134581 1.4915  2    0.4744
```

no effect of rotation here

```
itaglmer1a<- glmer(rt ~ freqTarget + lengthTarget + nTarget + (1|subject) + (1|target), data= dataIta, family=Gamma(link="log"))
```

```
anova(itaglmer0, itaglmer1a);
```

```
## Data: dataIta
## Models:
## itaglmer0: rt ~ 1 + (1 | subject) + (1 | target)
## itaglmer1a: rt ~ freqTarget + lengthTarget + nTarget + (1 | subject) + (1 | target)
## itaglmer1a:          target)
##          npar      AIC      BIC logLik deviance Chisq Df Pr(>Chisq)
## itaglmer0      4 134591 134620 -67291   134583
## itaglmer1a      7 134540 134592 -67263   134526 56.328  3 3.576e-12 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

strong improvement in GoF

```
car::Anova(itaglmer1a)
```

```
## Analysis of Deviance Table (Type II Wald chisquare tests)
##
## Response: rt
##          Chisq Df Pr(>Chisq)
## freqTarget 234.6265 1    <2e-16 ***
## lengthTarget  1.6941 1    0.1931
## nTarget      0.2406 1    0.6238
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

To which only freq seems to contribute. We introduce the variables of interest now:

```
itaglmer2<- glmer(rt ~ relatedness * morphType + freqTarget + (1|subject) + (1|target), data= dataIta, family=Gamma(link="log"))
```

```
summary(itaglmer2)-> modelsum2
car::Anova(itaglmer2)-> itaglmer2.anova
itaglmer2.anova
```

```
## Analysis of Deviance Table (Type II Wald chisquare tests)
##
## Response: rt
##               Chisq Df Pr(>Chisq)
## relatedness    63.282  1  1.791e-15 ***
## morphType      70.279  2  5.484e-16 ***
## freqTarget    420.382  1  < 2.2e-16 ***
## relatedness:morphType 228.836  2  < 2.2e-16 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

With “orthographic” and “unrelated/ctrl” as base contrast:

```
knitr::kable(round(modelsum2$coefficients, 4))
```

	Estimate	Std. Error	t value	Pr(> z)
(Intercept)	800.1365	3.9876	200.6554	0.0000
relatednessrel	-1.3974	1.8425	-0.7584	0.4482
morphTypeop	-11.6087	2.6859	-4.3221	0.0000
morphTypetr	-17.7294	2.7439	-6.4614	0.0000
freqTarget	-44.2176	2.1566	-20.5032	0.0000
relatednessrel:morphTypeop	-14.2715	2.2082	-6.4628	0.0000
relatednessrel:morphTypetr	-36.8151	2.5344	-14.5262	0.0000

Let's relevel for transparent versus opaque condition:

```
dataIta$morphType <- relevel(dataIta$morphType, "op");
```

```
itaglmer2c<- glmer(rt ~ relatedness * morphType + freqTarget + (1|subject) + (1|target), data= dataIta,
```

```
summary(itaglmer2c)->modelsum2c
car::Anova(itaglmer2c)-> itaglmer2c.anova
itaglmer2c.anova
```

```
## Analysis of Deviance Table (Type II Wald chisquare tests)
##
## Response: rt
##               Chisq Df Pr(>Chisq)
## relatedness    112.873  1  < 2.2e-16 ***
## morphType      13.006  2  0.001499 **
## freqTarget     488.101  1  < 2.2e-16 ***
## relatedness:morphType 100.900  2  < 2.2e-16 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

```
knitr::kable(round(modelsum2c$coefficients, 4))
```

	Estimate	Std. Error	t value	Pr(> z)
(Intercept)	788.5277	4.0865	192.9579	0.0000
relatednessrel	-15.6690	2.1883	-7.1604	0.0000
morphTypeeor	11.6097	4.3057	2.6964	0.0070
morphTypetr	-6.1195	3.3664	-1.8178	0.0691
freqTarget	-44.2179	2.0014	-22.0930	0.0000
relatednessrel:morphTypeeor	14.2720	2.7368	5.2148	0.0000
relatednessrel:morphTypetr	-22.5436	3.2858	-6.8608	0.0000

Is proficiency in L2 (ENG) able to modulate priming in L1 (ITA)?

Phonemic fluency:

```
itaglmer_phonFlu<- glmer(rt ~ relatedness * morphType * phonemicFluency + freqTarget + (1|subject) + (1|target))
```

Phonemic comprehension:

```
itaglmer_phonCom<- glmer(rt ~ relatedness * morphType * phonemicComprehension + freqTarget + (1|subject) + (1|target))
```

Unfortunately the other proficiency tests the model don't converge. Morphological awareness:

```
itaglmer_morphAw<- glmer(rt ~ relatedness * morphType * morphComprehension + freqTarget + (1|subject) + (1|target))
```

Spelling:

```
itaglmer_spell<- glmer(rt ~ relatedness * morphType * spelling + freqTarget + (1|subject) + (1|target),  
car::Anova(itaglmer_spell))
```

Vocabulary:

```
itaglmer_voc<- glmer(rt ~ relatedness * morphType * vocabulary + freqTarget + (1|subject) + (1|target),  
car::Anova(itaglmer_voc))
```

Reading comprehension:

```
itaglmer_read<- glmer(rt ~ relatedness * morphType * readingComprehension + freqTarget + (1|subject) + (1|target),  
car::Anova(itaglmer_read))
```

Oral comprehension:

```
itaglmer_oral<- glmer(rt ~ relatedness * morphType * oralComprehension + freqTarget + (1|subject) + (1|target),  
car::Anova(itaglmer_oral))
```

Summary of the results for ITA dataset

The model itaglmer2 found a main effect of relatedness [$\chi^2 = 63.28$, $p < 0.0001$]. A main effect of morphType [$\chi^2 = 70.28$, $p < 0.0001$], and a main effect of frequency [$\chi^2 = 420.38$, $p < 0.0001$]. More importantly, itaglmer2 shows a significant interaction between relatedness and morphotype [$\chi^2 = 228.84$, $p < 0.0001$], such

that among related trials there were faster reaction times to orthographic trials compared to opaque trials [$\beta = -14.27$, $p < 0.0001$], and to transparent trials [$\beta = -36.82$, $p < 0.0001$]. By setting the base contrast to opaque trials, itaglm2c shows that opaque trials are faster than transparent trials [$\beta = -22.54$, $p < 0.0001$].

We have also checked whether proficiency tests obtained on the second language, English, would influence the pattern of results in Italian. Only two tests showed convergence: phonological comprehension and phonological fluency, however both showed non significant results [both $p > 1$].

Modelling, ENG

Set contrasts:

```
dataEng$morphType <- relevel(dataEng$morphType, "or");
contrasts(dataEng$relatedness);
```

```
##      rel
## ctrl   0
## rel    1
```

```
contrasts(dataEng$morphType);
```

```
##      op tr
## or   0  0
## op   1  0
## tr   0  1
```

```
engglmer0 <- glmer(rt ~ 1 + (1|subject) + (1|target), data = dataEng,
  family=Gamma(link="identity"));
```

```
engglmer1 <- glmer(rt ~ trialCount + rotation + (1|subject) + (1|target), data = dataEng, family=Gamma(1.
```

```
anova(engglmer0, engglmer1);
```

```
## Data: dataEng
## Models:
## engglmer0: rt ~ 1 + (1 | subject) + (1 | target)
## engglmer1: rt ~ trialCount + rotation + (1 | subject) + (1 | target)
##      npar      AIC      BIC logLik deviance  Chisq Df Pr(>Chisq)
## engglmer0    4 113778 113806 -56885   113770
## engglmer1    6 113778 113820 -56883   113766 3.8608  2    0.1451
```

Again like in the ITA dataset, no effect of rotation here.

```
engglmer1c <- glmer(rt ~ freqTarget + lengthTarget + nTarget + (1|subject) + (1|target), data = dataEng
```

```
anova(engglmer0, engglmer1c);
```

```
## Data: dataEng
## Models:
## engglm0: rt ~ 1 + (1 | subject) + (1 | target)
## engglm1c: rt ~ freqTarget + lengthTarget + nTarget + (1 | subject) + (1 |
## engglm1c: target)
##          npar      AIC      BIC logLik deviance  Chisq Df Pr(>Chisq)
## engglm0      4 113778 113806 -56885    113770
## engglm1c     7 113719 113769 -56853    113705 64.523  3 6.344e-14 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

strong improvement in GoF

```
car::Anova(engglm1c)
```

```
## Analysis of Deviance Table (Type II Wald chisquare tests)
##
## Response: rt
##          Chisq Df Pr(>Chisq)
## freqTarget  301.5128 1    <2e-16 ***
## lengthTarget  92.6719 1    <2e-16 ***
## nTarget       0.0036 1    0.9523
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

Frequency and length contribute.

We introduce our variable of interest now:

```
engglm2 <- glmer(rt ~ relatedness * morphType + freqTarget + lengthTarget + (1|subject) + (1|target),
```

```
engglm2.anova <- car::Anova(engglm2)
engglm2.modelsum <- summary(engglm2)
engglm2.anova
```

```
## Analysis of Deviance Table (Type II Wald chisquare tests)
##
## Response: rt
##          Chisq Df Pr(>Chisq)
## relatedness  115.6393 1 < 2.2e-16 ***
## morphType     3.3445 2    0.1878
## freqTarget   663.8526 1 < 2.2e-16 ***
## lengthTarget  97.4373 1 < 2.2e-16 ***
## relatedness:morphType 32.9506 2 6.996e-08 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

Summary of the engglm2 model:

```
knitr::kable(round(engglm2.modelsum$coefficients, 4))
```

	Estimate	Std. Error	t value	Pr(> z)
(Intercept)	1148.3132	3.0478	376.7719	0.0000
relatednessrel	-19.0099	2.5847	-7.3548	0.0000
morphTypeop	0.0623	3.5746	0.0174	0.9861
morphTypetr	-0.7171	4.8117	-0.1490	0.8815
freqTarget	-72.1990	2.8022	-25.7653	0.0000
lengthTarget	-26.4356	2.6781	-9.8710	0.0000
relatednessrel:morphTypeop	-5.6052	2.9651	-1.8904	0.0587
relatednessrel:morphTypetr	-17.1599	3.1044	-5.5275	0.0000

We set the base contrast on the opaque condition and re-run the model:

```
dataEng$morphType <- relevel(dataEng$morphType, "op");

engglmer2c <- glmer(rt ~ relatedness * morphType + freqTarget + lengthTarget + (1|subject) + (1|target))

engglmer2c.anova <- car::Anova(engglmer2c)
engglmer2c.modelsum <- summary(engglmer2c)
engglmer2c.anova

## Analysis of Deviance Table (Type II Wald chisquare tests)
##
## Response: rt
##               Chisq Df Pr(>Chisq)
## relatedness    107.6610  1 < 2.2e-16 ***
## morphType       1.4125  2  0.4934930
## freqTarget     398.3679  1 < 2.2e-16 ***
## lengthTarget    54.1887  1  1.821e-13 ***
## relatedness:morphType 16.7656  2  0.0002288 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

knitr::kable(round(engglmer2c.modelsum$coefficients, 4))
```

	Estimate	Std. Error	t value	Pr(> z)
(Intercept)	1148.3958	4.1030	279.8895	0.0000
relatednessrel	-24.6153	2.7411	-8.9800	0.0000
morphTypeop	-0.0644	4.3638	-0.0148	0.9882
morphTypetr	-0.7773	8.3555	-0.0930	0.9259
freqTarget	-72.2010	3.6174	-19.9592	0.0000
lengthTarget	-26.4380	3.5915	-7.3613	0.0000
relatednessrel:morphTypeop	5.6057	4.1280	1.3580	0.1745
relatednessrel:morphTypetr	-11.5547	3.0514	-3.7866	0.0002

Summary of the results for ENG dataset

The model `engglmer2` found a main effect of relatedness [$\chi^2 = 115.64$, $p < 0.0001$]. A main effect of frequency [$\chi^2 = 663.85$, $p < 0.0001$], and also a main effect of length [$\chi^2 = 97.44$, $p < 0.0001$]. The effect of MorphType was not significant [$\chi^2 = 3.34$, $p = 0.1878$]. However, `itaglmer2` shows a significant interaction between relatedness and morphtype [$\chi^2 = 32.95$, $p < 0.0001$], such that among related trials there were faster reaction times to orthographic trials compared to opaque trials, although only marginally significant [$\beta = -5.61$, $p = 0.0587$], and to transparent trials [$\beta = -17.16$, $p < 0.0001$]. By setting the base contrast to opaque trials, `engglmer2c` shows that opaque trials are faster than transparent trials [$\beta = -11.55$, $p < 0.0002$].

Cross language interaction

```
rbind(dataEng, dataIta) -> crossExp
summary(crossExp)
```

```
##      subject      age      gender      handedness      trialCount
## Min.      : 1    Min.      :18.00    Female:13532    Left : 2191    Min.      : 1.0
## 1st Qu.:21    1st Qu.:22.00    Male  : 6415    Right:17756    1st Qu.: 75.0
## Median :42    Median :24.00                                Median :150.0
## Mean      :42    Mean      :24.33                                Mean      :150.2
## 3rd Qu.:62    3rd Qu.:26.00                                3rd Qu.:225.0
## Max.      :84    Max.      :34.00                                Max.      :300.0
##
##      rt      resp      trialType      lexicality      morphType
## Min.      : 284.0    Min.      :0.000    Length:19947    nonword: 0    op:6663
## 1st Qu.: 517.0    1st Qu.:2.000    Class :character    word :19947    or:6383
## Median : 588.0    Median :2.000    Mode  :character                                tr:6901
## Mean      : 629.3    Mean      :1.994
## 3rd Qu.: 693.0    3rd Qu.:2.000
## Max.      :1998.0    Max.      :2.000
##
##      target      prime      relatedness      freqTarget      freqPrime
## abito      : 78    against      : 40    ctrl: 9886    Min.      :1.367    Min.      :1.170
## again      : 78    angelic      : 40    rel :10061    1st Qu.:3.470    1st Qu.:2.670
## angel      : 78    cloudless: 40                                Median :4.011    Median :3.180
## ballo      : 78    dreamer      : 40                                Mean      :3.979    Mean      :3.263
## banca      : 78    drunkard      : 40                                3rd Qu.:4.520    3rd Qu.:3.867
## cambio      : 78    extract      : 40                                Max.      :5.840    Max.      :6.290
## (Other):19479    (Other) :19707                                NA's      :37
## lengthTarget    lengthPrime      nTarget      nPrime
## Min.      :3.000    Min.      : 5.000    Min.      : 0.00    Min.      : 0.000
## 1st Qu.:4.000    1st Qu.: 7.000    1st Qu.: 6.00    1st Qu.: 1.000
## Median :5.000    Median : 7.000    Median :12.00    Median : 2.000
## Mean      :4.958    Mean      : 7.514    Mean      :15.04    Mean      : 3.169
## 3rd Qu.:5.000    3rd Qu.: 8.000    3rd Qu.:22.00    3rd Qu.: 4.000
## Max.      :8.000    Max.      :13.000    Max.      :47.00    Max.      :36.000
##
##      rotation      language      oscTarget      accuracy phonemicFluency
## Length:19947      eng: 8938    Min.      :-0.094    Min.      :1    Min.      : 0.00
## Class :character    ita:11009    1st Qu.: 0.263    1st Qu.:1    1st Qu.:17.00
```

```
## Mode :character           Median : 0.623   Median :1   Median :23.00
##                               Mean  : 0.577   Mean  :1   Mean  :22.56
##                               3rd Qu.: 0.899   3rd Qu.:1   3rd Qu.:27.00
##                               Max.   : 1.000   Max.   :1   Max.   :45.00
##                               NA's    :11164
## phonemicComprehension morphComprehension spelling readingComprehension
## Min. : 0.000      Min. : 4.00      Min. : 0.000   Min. :1.000
## 1st Qu.: 8.000      1st Qu.: 8.00      1st Qu.: 5.000   1st Qu.:3.000
## Median : 9.000      Median : 9.00      Median : 8.000   Median :5.000
## Mean : 8.845      Mean : 8.95      Mean : 8.239   Mean :4.525
## 3rd Qu.:10.000      3rd Qu.:10.00      3rd Qu.:12.000   3rd Qu.:6.000
## Max. :13.000      Max. :10.00      Max. :18.000   Max. :7.000
##
## vocabulary oralComprehension aoa1.Aoa aoa2.usage
## Min. : 8.00      Min. :1.000      Min. : 0.000   Min. :1.000
## 1st Qu.:14.00      1st Qu.:3.000      1st Qu.: 6.000   1st Qu.:2.000
## Median :15.00      Median :5.000      Median : 6.000   Median :3.000
## Mean :15.31      Mean :4.591      Mean : 6.494   Mean :3.145
## 3rd Qu.:17.00      3rd Qu.:6.000      3rd Qu.: 8.000   3rd Qu.:4.000
## Max. :19.00      Max. :6.000      Max. :15.000   Max. :5.000
##
## aoa3.context aoa4.contextMultling aoa5.selfRatedProf aoa6.otherLang
## home : 2664   no :15257      Min. :1.000      no : 4735
## school:17283   yes: 4690      1st Qu.:3.000      yes:15212
##                               Median :4.000
##                               Mean :3.534
##                               3rd Qu.:4.000
##                               Max. :5.000
##
```

Setting out base contrast to orthographic, default of relatedness is “unrelated/ctrl”:

```
crossExp$morphType <- relevel(crossExp$morphType, "or");
```

```
crossglmer <- glmer(rt ~ relatedness * morphType * language + freqTarget + lengthTarget + (1|subject) +
```

```
crossglmer.modelsum <- summary(crossglmer)
```

```
crossglmer.anova <- car::Anova(crossglmer)
```

```
crossglmer.anova
```

```
## Analysis of Deviance Table (Type II Wald chisquare tests)
```

```
##
```

```
## Response: rt
```

```
##                               Chisq Df Pr(>Chisq)
## relatedness                 200.067  1 < 2.2e-16 ***
## morphType                   39.044  2 3.325e-09 ***
## language                   1765.815  1 < 2.2e-16 ***
## freqTarget                  842.282  1 < 2.2e-16 ***
## lengthTarget                38.695  1 4.956e-10 ***
## relatedness:morphType       136.567  2 < 2.2e-16 ***
## relatedness:language        60.080  1 9.108e-15 ***
## morphType:language          32.571  2 8.459e-08 ***
## relatedness:morphType:language 61.727  2 3.946e-14 ***
```

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

Summary:

```
knitr::kable(round(crossglmer.modelsum$coefficients, 4))
```

	Estimate	Std. Error	t value	Pr(> z)
(Intercept)	1015.4226	2.0309	499.9902	0.0000
relatednessrel	-18.3962	1.8571	-9.9060	0.0000
morphTypeop	-4.7311	2.3889	-1.9804	0.0477
morphTypetr	-11.5795	2.1260	-5.4465	0.0000
languageita	-89.1579	2.2066	-40.4049	0.0000
freqTarget	-59.0016	2.0330	-29.0221	0.0000
lengthTarget	-10.1169	1.6264	-6.2205	0.0000
relatednessrel:morphTypeop	-5.9379	2.1402	-2.7744	0.0055
relatednessrel:morphTypetr	-18.5042	2.1512	-8.6019	0.0000
relatednessrel:languageita	17.0474	1.7390	9.8030	0.0000
morphTypeop:languageita	-10.7867	2.8390	-3.7995	0.0001
morphTypetr:languageita	-3.0255	2.1983	-1.3763	0.1687
relatednessrel:morphTypeop:languageita	-8.8894	2.3634	-3.7613	0.0002
relatednessrel:morphTypetr:languageita	-18.1998	2.5149	-7.2368	0.0000

```
crossExp$morphType <- relevel(crossExp$morphType, "op");
```

```
crossglmerc <- glmer(rt ~ relatedness * morphType * language + freqTarget + lengthTarget + (1|subject) .
```

```
crossglmerc.modelsum <- summary(crossglmerc)
crossglmerc.anova <- car::Anova(crossglmerc)
crossglmerc.anova
```

```
## Analysis of Deviance Table (Type II Wald chisquare tests)
##
## Response: rt
##
##           Chisq Df Pr(>Chisq)
## relatedness    347.480  1 < 2.2e-16 ***
## morphType      71.744  2  2.636e-16 ***
## language     2418.082  1 < 2.2e-16 ***
## freqTarget     943.784  1 < 2.2e-16 ***
## lengthTarget   29.549  1  5.451e-08 ***
## relatedness:morphType    59.972  2  9.490e-14 ***
## relatedness:language    21.906  1  2.864e-06 ***
## morphType:language     42.098  2  7.221e-10 ***
## relatedness:morphType:language  44.627  2  2.039e-10 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

```
knitr::kable(round(crossglmerc.modelsum$coefficients, 4))
```

	Estimate	Std. Error	t value	Pr(> z)
(Intercept)	1010.7039	1.8982	532.4559	0.0000
relatednessrel	-24.3339	1.4798	-16.4438	0.0000
morphTypeor	4.7242	2.0056	2.3555	0.0185
morphTypetr	-6.8469	3.7620	-1.8200	0.0688
languageita	-99.9490	2.1581	-46.3134	0.0000
freqTarget	-59.0028	1.9206	-30.7211	0.0000
lengthTarget	-10.1180	1.8613	-5.4359	0.0000
relatednessrel:morphTypeor	5.9377	1.7657	3.3628	0.0008
relatednessrel:morphTypetr	-12.5666	2.6633	-4.7183	0.0000
relatednessrel:languageita	8.1579	2.0027	4.0735	0.0000
morphTypeor:languageita	10.8020	2.3890	4.5216	0.0000
morphTypetr:languageita	7.7589	2.7490	2.8225	0.0048
relatednessrel:morphTypeor:languageita	8.8895	1.8392	4.8333	0.0000
relatednessrel:morphTypetr:languageita	-9.3100	2.3367	-3.9842	0.0001
# Proficiency scores, correlation and distribution				
Create a dataframe with one row per participant				

```
pptFeatures <- unique(dataEng[,c('subject', 'age', 'gender', 'handedness', 'rotation', 'phonemicFluency', 'p
summary(pptFeatures)
```

```
##      subject      age      gender handedness  rotation
## Min.   : 1.00   Min.   :18.00   Female:54   Left : 8   Length:78
## 1st Qu.:21.50   1st Qu.:22.00   Male :24   Right:70   Class :character
## Median :41.50   Median :24.00                                Mode  :character
## Mean   :41.74   Mean    :24.28
## 3rd Qu.:61.75   3rd Qu.:26.00
## Max.    :84.00   Max.     :34.00
## phonemicFluency phonemicComprehension morphComprehension  spelling
## Min.    : 0.00   Min.    : 0.000   Min.    : 4.000   Min.    : 0.000
## 1st Qu.:16.25   1st Qu.: 8.000   1st Qu.: 8.000   1st Qu.: 4.000
## Median :22.00   Median : 9.000   Median : 9.000   Median : 7.000
## Mean    :22.17   Mean    : 8.769   Mean    : 8.897   Mean    : 8.026
## 3rd Qu.:27.00   3rd Qu.:10.000   3rd Qu.:10.000   3rd Qu.:11.000
## Max.    :45.00   Max.    :13.000   Max.    :10.000   Max.    :18.000
## readingComprehension vocabulary oralComprehension  aoa1.Aoa
## Min.    :1.000   Min.    : 8.00   Min.    :1.0   Min.    : 0.000
## 1st Qu.:3.000   1st Qu.:14.00   1st Qu.:3.0   1st Qu.: 6.000
## Median :5.000   Median :15.00   Median :5.0   Median : 6.000
## Mean    :4.423   Mean    :15.24   Mean    :4.5   Mean    : 6.487
## 3rd Qu.:6.000   3rd Qu.:17.00   3rd Qu.:6.0   3rd Qu.: 8.000
## Max.    :7.000   Max.    :19.00   Max.    :6.0   Max.    :15.000
## aoa2.usage aoa3.context aoa4.contextMultling aoa5.selfRatedProf
## Min.    :1.000   home :10   no :60   Min.    :1.000
## 1st Qu.:2.000   school:68   yes:18   1st Qu.:3.000
## Median :3.000                                Median :3.000
## Mean    :3.115                                Mean    :3.487
## 3rd Qu.:4.000                                3rd Qu.:4.000
## Max.    :5.000                                Max.    :5.000
## aoa6.otherLang
## no :18
## yes:60
```

```
##
##
##
##
```

```
knitr::kable(round(cor(pptFeatures[,c(6:12)], use='pairwise.complete.obs'), digits=2))
```

	phonemicFluency	phonemicComprehension	morphComprehension	spelling	readingComprehension
phonemicFluency	1.00	0.25	0.54	0.61	
phonemicComprehension	0.25	1.00	0.43	0.46	
morphComprehension	0.54	0.43	1.00	0.64	
spelling	0.61	0.46	0.64	1.00	
readingComprehension	0.35	0.44	0.40	0.49	
vocabulary	0.45	0.44	0.54	0.65	
oralComprehension	0.43	0.45	0.68	0.62	

Proficiency modelling

Set base contrasts to orthographic

```
dataEng$morphType <- relevel(dataEng$morphType, "or");
```

This model establishes the baseline model, with no proficiency score:

```
proficiencyglmer0 <- glmer(rt ~ relatedness * morphType + freqTarget + lengthTarget + (1|subject) + (1|target), data = dataEng, family=Gamma(link="identity"), control=glmerControl(optimizer="bobyqa"))
```

Now we test whether each individual proficiency score guarantees a better overall fit:

Phonemic fluency

```
proficiencyglmer1 <- glmer(rt ~ relatedness * morphType * phonemicFluency + lengthTarget + freqTarget + (1|subject) + (1|target), data = dataEng, family=Gamma(link="identity"), control=glmerControl(optimizer="bobyqa"))
```

```
anova(proficiencyglmer0, proficiencyglmer1);
```

```
## Data: dataEng
## Models:
## proficiencyglmer0: rt ~ relatedness * morphType + freqTarget + lengthTarget + (1 | subject) + (1 | target)
## proficiencyglmer1: rt ~ relatedness * morphType * phonemicFluency + lengthTarget + freqTarget + (1 | subject) + (1 | target)
##               npar      AIC      BIC logLik deviance  Chisq Df Pr(>Chisq)
## proficiencyglmer0   11 113629 113707 -56803   113607
## proficiencyglmer1   17 113621 113742 -56793   113587 19.855  6   0.002939 **
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

Increase in GoF

Phonemic comprehension

```
proficiencyglmer2<- glmer(rt ~ relatedness * morphType * phonemicComprehension + lengthTarget + freqTa  
+ (1|target), data = dataEng, family=Gamma(link="identity"), control=glmerControl(optimizer="b
```

```
anova(proficiencyglmer0, proficiencyglmer2);
```

```
## Data: dataEng  
## Models:  
## proficiencyglmer0: rt ~ relatedness * morphType + freqTarget + lengthTarget + (1 |  
## proficiencyglmer0:      subject) + (1 | target)  
## proficiencyglmer2: rt ~ relatedness * morphType * phonemicComprehension + lengthTarget +  
## proficiencyglmer2:      freqTarget + (1 | subject) + (1 | target)  
##               npar      AIC      BIC logLik deviance  Chisq Df Pr(>Chisq)  
## proficiencyglmer0   11 113629 113707 -56803    113607  
## proficiencyglmer2   17 113619 113740 -56793    113585 21.519  6    0.00148 **  
## ---  
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

Increase in GoF

Morphologic awareness

```
proficiencyglmer3<- glmer(rt ~ relatedness * morphType * morphComprehension + lengthTarget + freqTarget
```

```
anova(proficiencyglmer0, proficiencyglmer3);
```

```
## Data: dataEng  
## Models:  
## proficiencyglmer0: rt ~ relatedness * morphType + freqTarget + lengthTarget + (1 |  
## proficiencyglmer0:      subject) + (1 | target)  
## proficiencyglmer3: rt ~ relatedness * morphType * morphComprehension + lengthTarget +  
## proficiencyglmer3:      freqTarget + (1 | subject) + (1 | target)  
##               npar      AIC      BIC logLik deviance  Chisq Df Pr(>Chisq)  
## proficiencyglmer0   11 113629 113707 -56803    113607  
## proficiencyglmer3   17 113614 113734 -56790    113580 27.227  6 0.0001313 ***  
## ---  
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

Increase in GoF

Spelling

```
proficiencyglmer4<- glmer(rt ~ relatedness * morphType * spelling + lengthTarget + freqTarget + (1|sub
```

```
anova(proficiencyglmer0, proficiencyglmer4);
```

```
## Data: dataEng
## Models:
## proficiencyglmer0: rt ~ relatedness * morphType + freqTarget + lengthTarget + (1 |
## proficiencyglmer0:      subject) + (1 | target)
## proficiencyglmer4: rt ~ relatedness * morphType * spelling + lengthTarget + freqTarget +
## proficiencyglmer4:      (1 | subject) + (1 | target)
##
##          npar      AIC      BIC logLik deviance  Chisq Df Pr(>Chisq)
## proficiencyglmer0   11 113629 113707 -56803    113607
## proficiencyglmer4   17 113621 113741 -56793    113587 20.196  6   0.002556 **
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

Increase in GoF

Reading comprehension

```
proficiencyglmer5<- glmer(rt ~ relatedness * morphType * readingComprehension + lengthTarget + freqTar
```

```
anova(proficiencyglmer0, proficiencyglmer5);
```

```
## Data: dataEng
## Models:
## proficiencyglmer0: rt ~ relatedness * morphType + freqTarget + lengthTarget + (1 |
## proficiencyglmer0:      subject) + (1 | target)
## proficiencyglmer5: rt ~ relatedness * morphType * readingComprehension + lengthTarget +
## proficiencyglmer5:      freqTarget + (1 | subject) + (1 | target)
##
##          npar      AIC      BIC logLik deviance  Chisq Df Pr(>Chisq)
## proficiencyglmer0   11 113629 113707 -56803    113607
## proficiencyglmer5   17 113622 113743 -56794    113588 18.739  6   0.004628 **
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

Increase in GoF

Vocabulary

```
proficiencyglmer6<- glmer(rt ~ relatedness * morphType * vocabulary + lengthTarget + freqTarget + (1|s
```

```
anova(proficiencyglmer0, proficiencyglmer6)
```

```
## Data: dataEng
## Models:
## proficiencyglmer0: rt ~ relatedness * morphType + freqTarget + lengthTarget + (1 |
## proficiencyglmer0:      subject) + (1 | target)
## proficiencyglmer6: rt ~ relatedness * morphType * vocabulary + lengthTarget + freqTarget +
```

```
## proficiencyglmer6:      (1 | subject) + (1 | target)
##               npar      AIC      BIC logLik deviance  Chisq Df Pr(>Chisq)
## proficiencyglmer0     11 113629 113707 -56803    113607
## proficiencyglmer6     17 113624 113744 -56795    113590 16.974  6   0.009381 **
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

Increase in GoF

Oral comprehension

```
proficiencyglmer7<- glmer(rt ~ relatedness * morphType * oralComprehension + lengthTarget + freqTarget
anova(proficiencyglmer0, proficiencyglmer7)
```

```
## Data: dataEng
## Models:
## proficiencyglmer0: rt ~ relatedness * morphType + freqTarget + lengthTarget + (1 |
## proficiencyglmer0:      subject) + (1 | target)
## proficiencyglmer7: rt ~ relatedness * morphType * oralComprehension + lengthTarget +
## proficiencyglmer7:      freqTarget + (1 | subject) + (1 | target)
##               npar      AIC      BIC logLik deviance  Chisq Df Pr(>Chisq)
## proficiencyglmer0     11 113629 113707 -56803    113607
## proficiencyglmer7     17 113628 113749 -56797    113594 12.826  6   0.04587 *
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

Increase in GoF

All tests seem to increase the goodness of fit of the base model proficiencyglmer0.

Does proficiency specifically interact with priming?

We're going to inspect the anova and summary of each model first with the base contrast set to orthographic, and then with base contrast set to opaque.

Phonemic fluency

```
proficiencyglmer1.anova <- car::Anova(proficiencyglmer1)
proficiencyglmer1.modelsum <- summary(proficiencyglmer1)
proficiencyglmer1.anova
```

```
## Analysis of Deviance Table (Type II Wald chisquare tests)
##
## Response: rt
##               Chisq Df Pr(>Chisq)
## relatedness    149.9141  1 < 2.2e-16 ***
## morphType       55.5984  2  8.452e-13 ***
```

```
## phonemicFluency          0.1490  1    0.69946
## lengthTarget            59.0381  1  1.546e-14 ***
## freqTarget             247.7524  1 < 2.2e-16 ***
## relatedness:morphType    12.7430  2    0.00171 **
## relatedness:phonemicFluency  6.4054  1    0.01138 *
## morphType:phonemicFluency  20.1663  2  4.178e-05 ***
## relatedness:morphType:phonemicFluency  4.6787  2    0.09639 .
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

```
knitr::kable(round(proficiencyglmer1.modelsum$coefficients, 4))
```

	Estimate	Std. Error	t value	Pr(> z)
(Intercept)	1134.8547	4.0888	277.5538	0.0000
relatednessrel	-31.6121	3.6873	-8.5732	0.0000
morphTypeop	4.7598	4.9646	0.9588	0.3377
morphTypetr	24.7012	3.6178	6.8277	0.0000
phonemicFluency	0.7018	1.0012	0.7009	0.4834
lengthTarget	-26.8585	3.4955	-7.6836	0.0000
freqTarget	-72.2249	4.5886	-15.7402	0.0000
relatednessrel:morphTypeop	-14.2252	5.1627	-2.7554	0.0059
relatednessrel:morphTypetr	-6.9186	4.5080	-1.5347	0.1248
relatednessrel:phonemicFluency	0.5560	0.2662	2.0889	0.0367
morphTypeop:phonemicFluency	-0.2189	0.3966	-0.5519	0.5810
morphTypetr:phonemicFluency	-1.1214	0.3811	-2.9425	0.0033
relatednessrel:morphTypeop:phonemicFluency	0.4064	0.3690	1.1012	0.2708
relatednessrel:morphTypetr:phonemicFluency	-0.4492	0.3411	-1.3169	0.1879

```
dataEng$morphType <- relevel(dataEng$morphType, "op");
```

```
proficiencyglmer1b <- glmer(rt ~ relatedness * morphType * phonemicFluency + lengthTarget + freqTarget
+ (1|target), data = dataEng, family=Gamma(link="identity"), control=glmerControl(optimizer="bobyq"))
```

```
proficiencyglmer1b.anova <- car::Anova(proficiencyglmer1b)
proficiencyglmer1b.modelsum <- summary(proficiencyglmer1b)
proficiencyglmer1b.anova
```

```
## Analysis of Deviance Table (Type II Wald chisquare tests)
##
## Response: rt
##
##           Chisq Df Pr(>Chisq)
## relatedness    288.0903  1 < 2.2e-16 ***
## morphType       4.6475  2  0.0979071 .
## phonemicFluency  0.0169  1  0.8964806
## lengthTarget    66.6933  1  3.172e-16 ***
## freqTarget     353.1834  1 < 2.2e-16 ***
## relatedness:morphType    81.2126  2 < 2.2e-16 ***
## relatedness:phonemicFluency  11.8989  1  0.0005617 ***
## morphType:phonemicFluency  16.7373  2  0.0002320 ***
## relatedness:morphType:phonemicFluency  50.4393  2  1.115e-11 ***
```

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

```
knitr::kable(round(proficiencyglmer1b.modelsum$coefficients, 4))
```

	Estimate	Std. Error	t value	Pr(> z)
(Intercept)	1117.5070	4.5333	246.5109	0.0000
relatednessrel	-56.4754	3.5793	-15.7782	0.0000
morphTypeor	-3.4588	3.1128	-1.1111	0.2665
morphTypetr	3.9184	3.9380	0.9950	0.3197
phonemicFluency	0.0543	0.9325	0.0582	0.9536
lengthTarget	-27.7346	3.3961	-8.1666	0.0000
freqTarget	-67.8810	3.6120	-18.7932	0.0000
relatednessrel:morphTypeor	24.8902	3.1630	7.8691	0.0000
relatednessrel:morphTypetr	32.1827	3.9124	8.2257	0.0000
relatednessrel:phonemicFluency	1.4174	0.2242	6.3229	0.0000
morphTypeor:phonemicFluency	0.2584	0.3291	0.7852	0.4324
morphTypetr:phonemicFluency	-0.0995	0.3190	-0.3119	0.7551
relatednessrel:morphTypeor:phonemicFluency	-0.7568	0.2845	-2.6605	0.0078
relatednessrel:morphTypetr:phonemicFluency	-2.0029	0.2823	-7.0955	0.0000

Phonemic comprehension

```
proficiencyglmer2.anova <- car::Anova(proficiencyglmer2)
proficiencyglmer2.modelsum <- summary(proficiencyglmer2)
proficiencyglmer2.anova
```

```
## Analysis of Deviance Table (Type II Wald chisquare tests)
##
## Response: rt
##
##           Chisq Df Pr(>Chisq)
## relatedness    190.6468  1 < 2.2e-16 ***
## morphType      38.4328  2 4.513e-09 ***
## phonemicComprehension  4.2647  1  0.03891 *
## lengthTarget   38.6991  1 4.944e-10 ***
## freqTarget     247.5258  1 < 2.2e-16 ***
## relatedness:morphType    94.7005  2 < 2.2e-16 ***
## relatedness:phonemicComprehension  4.5962  1  0.03204 *
## morphType:phonemicComprehension  23.3758  2 8.395e-06 ***
## relatedness:morphType:phonemicComprehension  42.5724  2 5.695e-10 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

```
knitr::kable(round(proficiencyglmer2.modelsum$coefficients, 4))
```

	Estimate	Std. Error	t value	Pr(> z)
(Intercept)	1081.7928	10.7410	100.7165	0.0000
relatednessrel	-52.2509	4.2269	-12.3616	0.0000

	Estimate	Std. Error	t value	Pr(> z)
morphTypeop	0.1424	5.7203	0.0249	0.9801
morphTypetr	35.9996	4.4364	8.1146	0.0000
phonemicComprehension	7.6788	2.6089	2.9433	0.0032
lengthTarget	-26.5325	4.2651	-6.2209	0.0000
freqTarget	-72.3439	4.5982	-15.7330	0.0000
relatednessrel:morphTypeop	43.9669	3.9545	11.1183	0.0000
relatednessrel:morphTypetr	8.5439	4.1758	2.0461	0.0408
relatednessrel:phonemicComprehension	3.7867	0.7156	5.2916	0.0000
morphTypeop:phonemicComprehension	-0.0161	1.1564	-0.0139	0.9889
morphTypetr:phonemicComprehension	-4.1631	1.1514	-3.6157	0.0003
relatednessrel:morphTypeop:phonemicComprehension	-5.6581	0.8770	-6.4517	0.0000
relatednessrel:morphTypetr:phonemicComprehension	-2.9423	0.8527	-3.4507	0.0006

```
dataEng$morphType <- relevel(dataEng$morphType, "op");
```

```
proficiencyglmer2b <- glmer(rt ~ relatedness * morphType * phonemicComprehension + lengthTarget + freqTarget + (1|target), data = dataEng, family=Gamma(link="identity"), control=glmerControl(optimizer="bobyqa"))
```

```
proficiencyglmer2b.anova <- car::Anova(proficiencyglmer2b)
proficiencyglmer2b.modelsum <- summary(proficiencyglmer2b)
proficiencyglmer2b.anova
```

```
## Analysis of Deviance Table (Type II Wald chisquare tests)
##
## Response: rt
##
##              Chisq Df Pr(>Chisq)
## relatedness    174.9357  1 < 2.2e-16 ***
## morphType       53.2608  2 2.720e-12 ***
## phonemicComprehension 11.0448  1 0.0008894 ***
## lengthTarget    64.0166  1 1.234e-15 ***
## freqTarget     552.0537  1 < 2.2e-16 ***
## relatedness:morphType 34.1079  2 3.922e-08 ***
## relatedness:phonemicComprehension 1.3885  1 0.2386511
## morphType:phonemicComprehension 22.5979  2 1.239e-05 ***
## relatedness:morphType:phonemicComprehension 9.0798  2 0.0106747 *
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

```
knitr::kable(round(proficiencyglmer2b.modelsum$coefficients, 4))
```

	Estimate	Std. Error	t value	Pr(> z)
(Intercept)	1057.2324	3.4268	308.5158	0.0000
relatednessrel	-24.0579	3.1487	-7.6405	0.0000
morphTypeop	-5.4762	3.9364	-1.3911	0.1642
morphTypetr	26.6542	3.2615	8.1724	0.0000
phonemicComprehension	6.9144	2.2068	3.1333	0.0017
lengthTarget	-27.4608	3.4322	-8.0010	0.0000
freqTarget	-68.1731	2.9015	-23.4958	0.0000
relatednessrel:morphTypeop	-10.1777	3.9825	-2.5556	0.0106

	Estimate	Std. Error	t value	Pr(> z)
relatednessrel:morphTypetr	-16.3573	3.7107	-4.4082	0.0000
relatednessrel:phonemicComprehension	-0.2154	0.5638	-0.3820	0.7025
morphTypeop:phonemicComprehension	0.8294	0.9615	0.8626	0.3883
morphTypetr:phonemicComprehension	-2.8532	0.9344	-3.0536	0.0023
relatednessrel:morphTypeop:phonemicComprehension	2.3120	0.7692	3.0059	0.0026
relatednessrel:morphTypetr:phonemicComprehension	0.5461	0.7292	0.7489	0.4539
### Morphological awareness				

```
proficiencyglmer3.anova <- car::Anova(proficiencyglmer3)
proficiencyglmer3.modelsum <- summary(proficiencyglmer3)
proficiencyglmer3.anova
```

```
## Analysis of Deviance Table (Type II Wald chisquare tests)
##
## Response: rt
##
##           Chisq Df Pr(>Chisq)
## relatedness      152.563  1 < 2.2e-16 ***
## morphType        524.480  2 < 2.2e-16 ***
## morphComprehension  31.407  1 2.092e-08 ***
## lengthTarget     51.218  1 8.265e-13 ***
## freqTarget       215.887  1 < 2.2e-16 ***
## relatedness:morphType 108.661  2 < 2.2e-16 ***
## relatedness:morphComprehension 23.800  1 1.069e-06 ***
## morphType:morphComprehension  54.872  2 1.215e-12 ***
## relatedness:morphType:morphComprehension 16.297  2 0.0002891 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

```
knitr::kable(round(proficiencyglmer3.modelsum$coefficients, 4))
```

	Estimate	Std. Error	t value	Pr(> z)
(Intercept)	1038.9463	6.1748	168.2550	0.0000
relatednessrel	-32.4576	4.5194	-7.1818	0.0000
morphTypeop	33.8259	5.4833	6.1689	0.0000
morphTypetr	97.3078	4.9178	19.7870	0.0000
morphComprehension	12.4166	2.4773	5.0121	0.0000
lengthTarget	-26.4624	3.6976	-7.1567	0.0000
freqTarget	-72.5024	4.9345	-14.6931	0.0000
relatednessrel:morphTypeop	-43.0410	6.4120	-6.7126	0.0000
relatednessrel:morphTypetr	-25.1422	3.9168	-6.4191	0.0000
relatednessrel:morphComprehension	1.5220	0.7987	1.9055	0.0567
morphTypeop:morphComprehension	-3.8026	1.4157	-2.6860	0.0072
morphTypetr:morphComprehension	-10.9238	1.4274	-7.6530	0.0000
relatednessrel:morphTypeop:morphComprehension	4.2340	1.0488	4.0370	0.0001
relatednessrel:morphTypetr:morphComprehension	0.8449	0.8661	0.9755	0.3293

```
dataEng$morphType <- relevel(dataEng$morphType, "op");
```

```
proficiencyglmer3b<- glmer(rt ~ relatedness * morphType * morphComprehension + lengthTarget + freqTarget)
```

```
proficiencyglmer3b.anova <- car::Anova(proficiencyglmer3b)
proficiencyglmer3b.modelsum <- summary(proficiencyglmer3b)
proficiencyglmer3b.anova
```

```
## Analysis of Deviance Table (Type II Wald chisquare tests)
##
## Response: rt
##
##           Chisq Df Pr(>Chisq)
## relatedness      308.0157  1 < 2.2e-16 ***
## morphType        78.0208  2 < 2.2e-16 ***
## morphComprehension  4.8901  1  0.02701 *
## lengthTarget     68.2622  1 < 2.2e-16 ***
## freqTarget      244.9858  1 < 2.2e-16 ***
## relatedness:morphType  48.4653  2 2.991e-11 ***
## relatedness:morphComprehension  33.4050  1 7.483e-09 ***
## morphType:morphComprehension  60.6706  2 6.692e-14 ***
## relatedness:morphType:morphComprehension  45.9456  2 1.055e-10 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

```
knitr::kable(round(proficiencyglmer3b.modelsum$coefficients, 4))
```

	Estimate	Std. Error	t value	Pr(> z)
(Intercept)	1054.6027	4.5686	230.8358	0.0000
relatednessrel	-71.0514	4.3546	-16.3166	0.0000
morphTypeeor	-30.1665	6.1510	-4.9044	0.0000
morphTypetr	49.5455	4.6677	10.6145	0.0000
morphComprehension	6.9006	2.5011	2.7590	0.0058
lengthTarget	-27.0451	3.2734	-8.2621	0.0000
freqTarget	-68.1457	4.3538	-15.6520	0.0000
relatednessrel:morphTypeeor	44.3050	7.0870	6.2516	0.0000
relatednessrel:morphTypetr	37.1531	4.9787	7.4624	0.0000
relatednessrel:morphComprehension	5.1647	0.6525	7.9158	0.0000
morphTypeeor:morphComprehension	3.5995	1.2520	2.8749	0.0040
morphTypetr:morphComprehension	-5.3315	1.2085	-4.4116	0.0000
relatednessrel:morphTypeeor:morphComprehension	-3.9468	1.0220	-3.8618	0.0001
relatednessrel:morphTypetr:morphComprehension	-5.5851	0.8487	-6.5805	0.0000

Spelling

```
proficiencyglmer4.anova <- car::Anova(proficiencyglmer4)
proficiencyglmer4.modelsum <- summary(proficiencyglmer4)
proficiencyglmer4.anova
```

```
## Analysis of Deviance Table (Type II Wald chisquare tests)
##
```



```
## Response: rt
##
##           Chisq Df Pr(>Chisq)
## relatedness    134.1645  1 < 2.2e-16 ***
## morphType      27.4114  2  1.116e-06 ***
## spelling        1.7117  1   0.19076
## lengthTarget   43.1821  1  4.987e-11 ***
## freqTarget     484.5839  1 < 2.2e-16 ***
## relatedness:morphType  30.6629  2  2.196e-07 ***
## relatedness:spelling   4.7912  1   0.02861 *
## morphType:spelling     19.0341  2  7.359e-05 ***
## relatedness:morphType:spelling  1.4888  2   0.47502
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

```
knitr::kable(round(proficiencyglmer4.modelsum$coefficients, 4))
```

	Estimate	Std. Error	t value	Pr(> z)
(Intercept)	1171.0568	3.5566	329.2639	0.0000
relatednessrel	-35.2546	3.7784	-9.3305	0.0000
morphTypeor	-9.2844	3.9463	-2.3527	0.0186
morphTypetr	17.2039	3.2317	5.3234	0.0000
spelling	-3.0051	2.2525	-1.3341	0.1822
lengthTarget	-26.3075	4.0034	-6.5713	0.0000
freqTarget	-72.2192	3.2807	-22.0133	0.0000
relatednessrel:morphTypeor	13.1761	3.5253	3.7376	0.0002
relatednessrel:morphTypetr	-11.4206	4.1772	-2.7341	0.0063
relatednessrel:spelling	1.2902	0.6058	2.1296	0.0332
morphTypeor:spelling	1.0854	0.7777	1.3956	0.1628
morphTypetr:spelling	-2.0594	0.7404	-2.7815	0.0054
relatednessrel:morphTypeor:spelling	-0.9101	0.7961	-1.1432	0.2530
relatednessrel:morphTypetr:spelling	-0.0719	0.7685	-0.0936	0.9254

```
dataEng$morphType <- relevel(dataEng$morphType, "op");
```

```
proficiencyglmer4b<- glmer(rt ~ relatedness * morphType * spelling + lengthTarget + freqTarget + (1|subject))
```

```
proficiencyglmer4b.anova <- car::Anova(proficiencyglmer4b)
proficiencyglmer4b.modelsum <- summary(proficiencyglmer4b)
proficiencyglmer4b.anova
```

```
## Analysis of Deviance Table (Type II Wald chisquare tests)
##
## Response: rt
##
##           Chisq Df Pr(>Chisq)
## relatedness    190.7043  1 < 2.2e-16 ***
## morphType       6.2936  2  0.042990 *
## spelling        1.8417  1  0.174746
## lengthTarget    66.9171  1  2.832e-16 ***
## freqTarget     305.4612  1 < 2.2e-16 ***
## relatedness:morphType  33.7450  2  4.703e-08 ***
```

```
## relatedness:spelling          1.0397  1  0.307902
## morphType:spelling           25.7411  2  2.573e-06 ***
## relatedness:morphType:spelling 10.2678  2  0.005894 **
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

```
knitr::kable(round(proficiencyglmer4b.modelsum$coefficients, 4))
```

	Estimate	Std. Error	t value	Pr(> z)
(Intercept)	1137.4869	5.3156	213.9913	0.0000
relatednessrel	-37.9314	3.5927	-10.5578	0.0000
morphTypeor	-3.6070	3.6578	-0.9861	0.3241
morphTypetr	13.6607	3.9968	3.4180	0.0006
spelling	-2.8889	1.8754	-1.5404	0.1235
lengthTarget	-27.1608	3.3203	-8.1803	0.0000
freqTarget	-67.8268	3.8808	-17.4774	0.0000
relatednessrel:morphTypeor	18.3245	3.7504	4.8860	0.0000
relatednessrel:morphTypetr	5.3533	3.8062	1.4065	0.1596
relatednessrel:spelling	1.4781	0.5415	2.7298	0.0063
morphTypeor:spelling	0.6788	0.6788	1.0001	0.3173
morphTypetr:spelling	-1.3550	0.6364	-2.1291	0.0332
relatednessrel:morphTypeor:spelling	-1.1394	0.7143	-1.5950	0.1107
relatednessrel:morphTypetr:spelling	-2.1023	0.6678	-3.1482	0.0016

Reading comprehension

```
proficiencyglmer5.anova <- car::Anova(proficiencyglmer5)
proficiencyglmer5.modelsum <- summary(proficiencyglmer5)
proficiencyglmer5.anova
```

```
## Analysis of Deviance Table (Type II Wald chisquare tests)
##
## Response: rt
##
##           Chisq Df Pr(>Chisq)
## relatedness    107.3151  1 < 2.2e-16 ***
## morphType      30.6589  2  2.200e-07 ***
## readingComprehension  2.8110  1  0.09362 .
## lengthTarget   47.2183  1  6.350e-12 ***
## freqTarget    238.0712  1 < 2.2e-16 ***
## relatedness:morphType  2.9111  2  0.23327
## relatedness:readingComprehension  0.4667  1  0.49450
## morphType:readingComprehension  24.5211  2  4.735e-06 ***
## relatedness:morphType:readingComprehension  4.7872  2  0.09130 .
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

```
knitr::kable(round(proficiencyglmer5.modelsum$coefficients, 4))
```

	Estimate	Std. Error	t value	Pr(> z)
(Intercept)	1120.0242	4.9253	227.4028	0.0000
relatednessrel	-30.2895	6.6045	-4.5862	0.0000
morphTypeop	7.7289	8.2889	0.9324	0.3511
morphTypetr	25.3555	5.1036	4.9681	0.0000
readingComprehension	6.5530	3.7596	1.7430	0.0813
lengthTarget	-26.4444	3.8484	-6.8716	0.0000
freqTarget	-72.3485	4.6890	-15.4296	0.0000
relatednessrel:morphTypeop	6.5187	10.0629	0.6478	0.5171
relatednessrel:morphTypetr	2.8760	7.9395	0.3622	0.7172
relatednessrel:readingComprehension	2.5117	1.6692	1.5047	0.1324
morphTypeop:readingComprehension	-1.7530	1.9071	-0.9192	0.3580
morphTypetr:readingComprehension	-5.8572	1.8995	-3.0835	0.0020
relatednessrel:morphTypeop:readingComprehension	-2.6925	2.4439	-1.1017	0.2706
relatednessrel:morphTypetr:readingComprehension	-4.5258	2.1070	-2.1480	0.0317

```
dataEng$morphType <- relevel(dataEng$morphType, "op");
```

```
proficiencyglmer5b<- glmer(rt ~ relatedness * morphType * readingComprehension + lengthTarget + freqTa
```

```
proficiencyglmer5b.anova <- car::Anova(proficiencyglmer5b)
proficiencyglmer5b.modelsum <- summary(proficiencyglmer5b)
proficiencyglmer5b.anova
```

```
## Analysis of Deviance Table (Type II Wald chisquare tests)
##
## Response: rt
##
##           Chisq Df Pr(>Chisq)
## relatedness    111.7743  1 < 2.2e-16 ***
## morphType       5.4697  2  0.064902 .
## readingComprehension  3.9589  1  0.046625 *
## lengthTarget    30.1253  1  4.050e-08 ***
## freqTarget     276.6214  1 < 2.2e-16 ***
## relatedness:morphType  11.1129  2  0.003862 **
## relatedness:readingComprehension  0.1612  1  0.688045
## morphType:readingComprehension  20.9944  2  2.761e-05 ***
## relatedness:morphType:readingComprehension  1.3950  2  0.497816
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

```
knitr::kable(round(proficiencyglmer5b.modelsum$coefficients, 4))
```

	Estimate	Std. Error	t value	Pr(> z)
(Intercept)	1012.7534	31.7721	31.8756	0.0000
relatednessrel	-21.3113	12.1704	-1.7511	0.0799
morphTypeop	2.4386	14.3158	0.1703	0.8647
morphTypetr	21.0404	13.8040	1.5242	0.1275
readingComprehension	7.5404	3.1820	2.3697	0.0178
lengthTarget	-23.3681	4.2575	-5.4887	0.0000
freqTarget	-64.4392	3.8744	-16.6319	0.0000

	Estimate	Std. Error	t value	Pr(> z)
relatednessrel:morphTypeop	-3.1886	17.7521	-0.1796	0.8575
relatednessrel:morphTypetr	-5.2726	16.7712	-0.3144	0.7532
relatednessrel:readingComprehension	-0.9684	2.5839	-0.3748	0.7078
morphTypeop:readingComprehension	0.3061	2.7209	0.1125	0.9104
morphTypetr:readingComprehension	-5.2326	2.5825	-2.0262	0.0427
relatednessrel:morphTypeop:readingComprehension	2.9554	3.7715	0.7836	0.4333
relatednessrel:morphTypetr:readingComprehension	-1.3077	3.5386	-0.3695	0.7117

Vocabulary

```
proficiencyglmer6.anova <- car::Anova(proficiencyglmer6)
proficiencyglmer6.modelsum <- summary(proficiencyglmer6)
proficiencyglmer6.anova
```

```
## Analysis of Deviance Table (Type II Wald chisquare tests)
##
## Response: rt
##
##           Chisq Df Pr(>Chisq)
## relatedness    363.795  1 < 2.2e-16 ***
## morphType     100.233  2 < 2.2e-16 ***
## vocabulary       1.540  1    0.2146
## lengthTarget    26.524  1 2.603e-07 ***
## freqTarget     403.518  1 < 2.2e-16 ***
## relatedness:morphType    51.289  2 7.290e-12 ***
## relatedness:vocabulary   64.775  1 8.397e-16 ***
## morphType:vocabulary    25.618  2 2.737e-06 ***
## relatedness:morphType:vocabulary  39.369  2 2.826e-09 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

```
knitr::kable(round(proficiencyglmer6.modelsum$coefficients, 4))
```

	Estimate	Std. Error	t value	Pr(> z)
(Intercept)	1188.2727	4.3258	274.6936	0.0000
relatednessrel	-83.6997	4.0119	-20.8628	0.0000
morphTypeop	-2.8159	3.4086	-0.8261	0.4087
morphTypetr	35.8139	3.8939	9.1975	0.0000
vocabulary	-2.5755	1.8260	-1.4105	0.1584
lengthTarget	-26.6260	5.1700	-5.1501	0.0000
freqTarget	-72.2009	3.5943	-20.0878	0.0000
relatednessrel:morphTypeop	21.4149	3.8695	5.5343	0.0000
relatednessrel:morphTypetr	33.0798	4.5225	7.3145	0.0000
relatednessrel:vocabulary	4.2189	0.4231	9.9711	0.0000
morphTypeop:vocabulary	0.1896	0.8300	0.2285	0.8193
morphTypetr:vocabulary	-2.3500	0.8294	-2.8334	0.0046
relatednessrel:morphTypeop:vocabulary	-1.7518	0.5288	-3.3129	0.0009
relatednessrel:morphTypetr:vocabulary	-3.2847	0.5285	-6.2147	0.0000

```
dataEng$morphType <- relevel(dataEng$morphType, "op");
```

```
proficiencyglmer6b<- glmer(rt ~ relatedness * morphType * vocabulary + lengthTarget + freqTarget + (1|
```

```
proficiencyglmer6b.anova <- car::Anova(proficiencyglmer6b)
proficiencyglmer6b.modelsum <- summary(proficiencyglmer6b)
proficiencyglmer6b.anova
```

```
## Analysis of Deviance Table (Type II Wald chisquare tests)
##
## Response: rt
##
##               Chisq Df Pr(>Chisq)
## relatedness    306.0907  1 < 2.2e-16 ***
## morphType      96.1525  2 < 2.2e-16 ***
## vocabulary      0.0783  1    0.7796
## lengthTarget   48.7342  1 2.931e-12 ***
## freqTarget    370.9151  1 < 2.2e-16 ***
## relatedness:morphType  2.9168  2    0.2326
## relatedness:vocabulary 36.7855  1 1.319e-09 ***
## morphType:vocabulary 29.5597  2 3.812e-07 ***
## relatedness:morphType:vocabulary 34.7464  2 2.850e-08 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

```
knitr::kable(round(proficiencyglmer6b.modelsum$coefficients, 4))
```

	Estimate	Std. Error	t value	Pr(> z)
(Intercept)	1137.8423	5.6940	199.8331	0.0000
relatednessrel	-54.5748	4.0569	-13.4522	0.0000
morphTypeeor	6.7985	4.8077	1.4141	0.1573
morphTypetr	47.1190	4.8970	9.6219	0.0000
vocabulary	-1.4249	1.7199	-0.8285	0.4074
lengthTarget	-27.7565	3.9760	-6.9810	0.0000
freqTarget	-67.3156	3.4953	-19.2592	0.0000
relatednessrel:morphTypeeor	-9.4822	4.4362	-2.1374	0.0326
relatednessrel:morphTypetr	23.8131	8.1969	2.9051	0.0037
relatednessrel:vocabulary	1.8978	0.3746	5.0662	0.0000
morphTypeeor:vocabulary	-0.3569	0.7363	-0.4847	0.6279
morphTypetr:vocabulary	-2.9586	0.7072	-4.1839	0.0000
relatednessrel:morphTypeeor:vocabulary	1.2299	0.4838	2.5420	0.0110
relatednessrel:morphTypetr:vocabulary	-2.3265	0.6233	-3.7328	0.0002

Oral comprehension

```
proficiencyglmer7.anova <- car::Anova(proficiencyglmer7)
proficiencyglmer7.modelsum <- summary(proficiencyglmer7)
proficiencyglmer7.anova
```

```
## Analysis of Deviance Table (Type II Wald chisquare tests)
##
## Response: rt
##
##           Chisq Df Pr(>Chisq)
## relatedness      148.4503  1 < 2.2e-16 ***
## morphType        138.1402  2 < 2.2e-16 ***
## oralComprehension    0.4328  1  0.51063
## lengthTarget       91.6203  1 < 2.2e-16 ***
## freqTarget        352.8560  1 < 2.2e-16 ***
## relatedness:morphType    20.9182  2  2.869e-05 ***
## relatedness:oralComprehension    3.0700  1  0.07975 .
## morphType:oralComprehension    21.3570  2  2.303e-05 ***
## relatedness:morphType:oralComprehension    0.7190  2  0.69802
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

```
knitr::kable(round(proficiencyglmer7.modelsum$coefficients, 4))
```

	Estimate	Std. Error	t value	Pr(> z)
(Intercept)	1151.4971	5.3597	214.8429	0.0000
relatednessrel	-23.9274	3.6114	-6.6255	0.0000
morphTypeop	19.2799	4.7451	4.0631	0.0000
morphTypetr	35.4399	3.6540	9.6989	0.0000
oralComprehension	-0.9326	3.9715	-0.2348	0.8144
lengthTarget	-26.2976	2.7474	-9.5718	0.0000
freqTarget	-72.3171	3.8498	-18.7845	0.0000
relatednessrel:morphTypeop	-12.8680	7.3714	-1.7457	0.0809
relatednessrel:morphTypetr	-19.5969	4.6647	-4.2011	0.0000
relatednessrel:oralComprehension	1.1364	1.1931	0.9524	0.3409
morphTypeop:oralComprehension	-4.1494	1.7829	-2.3274	0.0199
morphTypetr:oralComprehension	-7.7514	1.7610	-4.4018	0.0000
relatednessrel:morphTypeop:oralComprehension	1.5364	1.8156	0.8462	0.3974
relatednessrel:morphTypetr:oralComprehension	0.4198	1.5258	0.2751	0.7832

```
dataEng$morphType <- relevel(dataEng$morphType, "op");
```

```
proficiencyglmer7b<- glmer(rt ~ relatedness * morphType * oralComprehension + lengthTarget + freqTarget
```

```
proficiencyglmer7b.anova <- car::Anova(proficiencyglmer7b)
proficiencyglmer7b.modelsum <- summary(proficiencyglmer7b)
proficiencyglmer7b.anova
```

```
## Analysis of Deviance Table (Type II Wald chisquare tests)
##
## Response: rt
##
##           Chisq Df Pr(>Chisq)
## relatedness      242.9122  1 < 2.2e-16 ***
## morphType        12.6312  2  0.001808 **
## oralComprehension    1.2854  1  0.256893
## lengthTarget       67.0173  1  2.691e-16 ***
```

```
## freqTarget                330.2657  1 < 2.2e-16 ***
## relatedness:morphType     35.5801  2 1.879e-08 ***
## relatedness:oralComprehension 2.8583  1 0.090904 .
## morphType:oralComprehension 19.5765  2 5.611e-05 ***
## relatedness:morphType:oralComprehension 7.1928  2 0.027422 *
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

```
knitr::kable(round(proficiencyglmer7b.modelsum$coefficients, 4))
```

	Estimate	Std. Error	t value	Pr(> z)
(Intercept)	1131.9377	4.2024	269.3556	0.0000
relatednessrel	-39.2651	3.3166	-11.8391	0.0000
morphTypeeor	-7.6802	3.9429	-1.9478	0.0514
morphTypetr	21.1512	4.8673	4.3456	0.0000
oralComprehension	-3.8344	2.9304	-1.3085	0.1907
lengthTarget	-27.2475	3.3284	-8.1864	0.0000
freqTarget	-67.6627	3.7232	-18.1732	0.0000
relatednessrel:morphTypeeor	22.5775	4.2902	5.2626	0.0000
relatednessrel:morphTypetr	1.7790	3.5574	0.5001	0.6170
relatednessrel:oralComprehension	2.8856	1.0144	2.8445	0.0044
morphTypeeor:oralComprehension	2.1804	1.5563	1.4010	0.1612
morphTypetr:oralComprehension	-4.0872	1.5430	-2.6489	0.0081
relatednessrel:morphTypeeor:oralComprehension	-2.7044	1.4128	-1.9142	0.0556
relatednessrel:morphTypetr:oralComprehension	-2.9646	1.2412	-2.3886	0.0169

AoA

correlation

Correlation between the individual scores, and between aoa and proficiency:

```
round(cor(pptFeatures[,c(13,14,17)], use='pairwise.complete.obs', method='spearman'), digits=2);
```

```
##          aoa1.Aoa  aoa2.usage  aoa5.selfRatedProf
## aoa1.Aoa          1.00      -0.15              0.04
## aoa2.usage       -0.15          1.00              0.49
## aoa5.selfRatedProf 0.04          0.49              1.00
```

```
round(cor(pptFeatures[,c(6:12, 13)], use='pairwise.complete.obs', method='spearman'), digits=2)[8,];
```

```
##      phonemicFluency  phonemicComprehension  morphComprehension
##              0.04              -0.21              -0.13
##      spelling      readingComprehension      vocabulary
##      -0.23              -0.15              -0.13
##      oralComprehension      aoa1.Aoa
##      -0.19              1.00
```

aoa2 and aoa5 are quite correlated (unsurprisingly)

scores, modelling

Set base contrast to “orthographic”

```
dataEng$morphType <- relevel(dataEng$morphType, "or");
```

```
aoaglm1<- glmer(rt ~ relatedness * morphType*aoa1.Aoa + freqTarget + (1|subject) + (1|target), data=
```

```
anova(proficiencyglmer0, aoaglm1)
```

```
## Data: dataEng
## Models:
## proficiencyglmer0: rt ~ relatedness * morphType + freqTarget + lengthTarget + (1 |
## proficiencyglmer0:      subject) + (1 | target)
## aoaglm1: rt ~ relatedness * morphType * aoa1.Aoa + freqTarget + (1 | subject) +
## aoaglm1:      (1 | target)
##               npar    AIC    BIC logLik deviance  Chisq Df Pr(>Chisq)
## proficiencyglmer0   11 113629 113707 -56803    113607
## aoaglm1             16 113636 113750 -56802    113604 2.3202  5      0.8033
```

No increase in GoF

```
aoaglm2<- glmer(rt ~ relatedness * morphType*aoa2.usage + freqTarget + (1|subject) + (1|target), data=
```

```
anova(proficiencyglmer0, aoaglm2)
```

```
## Data: dataEng
## Models:
## proficiencyglmer0: rt ~ relatedness * morphType + freqTarget + lengthTarget + (1 |
## proficiencyglmer0:      subject) + (1 | target)
## aoaglm2: rt ~ relatedness * morphType * aoa2.usage + freqTarget + (1 |
## aoaglm2:      subject) + (1 | target)
##               npar    AIC    BIC logLik deviance  Chisq Df Pr(>Chisq)
## proficiencyglmer0   11 113629 113707 -56803    113607
## aoaglm2             16 113631 113745 -56800    113599 7.3367  5      0.1968
```

No increase in GoF

```
aoaglm3<- glmer(rt ~ relatedness * morphType*aoa3.context + freqTarget + (1|subject) + (1|target), da
```

```
anova(proficiencyglmer0, aoaglm3)
```

```
## Data: dataEng
## Models:
## proficiencyglmer0: rt ~ relatedness * morphType + freqTarget + lengthTarget + (1 |
## proficiencyglmer0:      subject) + (1 | target)
## aoaglm3: rt ~ relatedness * morphType * aoa3.context + freqTarget + (1 |
## aoaglm3:      subject) + (1 | target)
##               npar    AIC    BIC logLik deviance  Chisq Df Pr(>Chisq)
## proficiencyglmer0   11 113629 113707 -56803    113607
## aoaglm3             16 113641 113755 -56805    113609      0  5      1
```


No increase in GoF

```
aoaglm4<- glmer(rt ~ relatedness * morphType*aoa4.contextMultling + freqTarget + (1|subject) + (1|target))
```

```
anova(proficiencyglmer0, aoaglm4)
```

```
## Data: dataEng
## Models:
## proficiencyglmer0: rt ~ relatedness * morphType + freqTarget + lengthTarget + (1 |
## proficiencyglmer0:      subject) + (1 | target)
## aoaglm4: rt ~ relatedness * morphType * aoa4.contextMultling + freqTarget +
## aoaglm4:      (1 | subject) + (1 | target)
##
##           npar      AIC      BIC logLik deviance Chisq Df Pr(>Chisq)
## proficiencyglmer0  11 113629 113707 -56803    113607
## aoaglm4           16 113641 113755 -56805    113609      0  5          1
```

No Increase in GoF

```
aoaglm5<- glmer(rt ~ relatedness * morphType*aoa5.selfRatedProf + freqTarget + lengthTarget + (1|subject))
```

```
anova(proficiencyglmer0, aoaglm5)
```

```
## Data: dataEng
## Models:
## proficiencyglmer0: rt ~ relatedness * morphType + freqTarget + lengthTarget + (1 |
## proficiencyglmer0:      subject) + (1 | target)
## aoaglm5: rt ~ relatedness * morphType * aoa5.selfRatedProf + freqTarget +
## aoaglm5:      lengthTarget + (1 | subject) + (1 | target)
##
##           npar      AIC      BIC logLik deviance Chisq Df Pr(>Chisq)
## proficiencyglmer0  11 113629 113707 -56803    113607
## aoaglm5           17 113623 113744 -56794    113589 17.787  6  0.006789 **
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

Self rated proficiency appears to increase GoF

```
aoaglm6<- glmer(rt ~ relatedness * morphType*aoa6.otherLang + freqTarget + lengthTarget + (1|subject))
```

```
anova(proficiencyglmer0, aoaglm6)
```

```
## Data: dataEng
## Models:
## proficiencyglmer0: rt ~ relatedness * morphType + freqTarget + lengthTarget + (1 |
## proficiencyglmer0:      subject) + (1 | target)
## aoaglm6: rt ~ relatedness * morphType * aoa6.otherLang + freqTarget +
## aoaglm6:      (1 | subject) + (1 | target)
##
##           npar      AIC      BIC logLik deviance Chisq Df Pr(>Chisq)
## proficiencyglmer0  11 113629 113707 -56803    113607
## aoaglm6           16 113644 113758 -56806    113612      0  5          1
```

No Increase in GoF

Let's change the contrast to "opaque" now:

```
dataEng$morphType <- relevel(dataEng$morphType, "op");

aoaglm1c<- glmer(rt ~ relatedness * morphType*aoa1.Aoa + freqTarget + (1|subject) + (1|target), data=
aoaglm2c<- glmer(rt ~ relatedness * morphType*aoa2.usage + freqTarget + (1|subject) + (1|target), data=
aoaglm3c<- glmer(rt ~ relatedness * morphType*aoa2.context + freqTarget + (1|subject) + (1|target), data=
aoaglm4c<- glmer(rt ~ relatedness * morphType*aoa4.contextMultling + freqTarget + (1|subject) + (1|target), data=
aoaglm5c<- glmer(rt ~ relatedness * morphType*aoa5.selfRatedProf + freqTarget + (1|subject) + (1|target), data=
aoaglm6c<- glmer(rt ~ relatedness * morphType*aoa6.otherLang + freqTarget + (1|subject) + (1|target), data=

aoaglm1.anova <- car::Anova(aoaglm1)
aoaglm1c.anova <- car::Anova(aoaglm1c)
aoaglm1.modelsum <- summary(aoaglm1)
aoaglm1c.modelsum <- summary(aoaglm1c)
```

Anova of aoaglm1 - or contrast

```
aoaglm1.anova
```

```
## Analysis of Deviance Table (Type II Wald chisquare tests)
##
## Response: rt
##
##           Chisq Df Pr(>Chisq)
## relatedness    147.1853  1 < 2.2e-16 ***
## morphType      136.6736  2 < 2.2e-16 ***
## aoa1.Aoa         7.2479  1  0.007098 **
## freqTarget      390.3539  1 < 2.2e-16 ***
## relatedness:morphType    30.6241  2 2.239e-07 ***
## relatedness:aoa1.Aoa     1.0688  1  0.301216
## morphType:aoa1.Aoa       7.1315  2  0.028276 *
## relatedness:morphType:aoa1.Aoa 36.7215  2 1.062e-08 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

```
knitr::kable(round(aoaglm1.modelsum$coefficients, 4))
```

	Estimate	Std. Error	t value	Pr(> z)
(Intercept)	1065.3439	4.0794	261.1534	0.0000
relatednessrel	-37.6734	4.5342	-8.3087	0.0000
morphTypeeop	-6.9483	3.3388	-2.0810	0.0374
morphTypetr	-45.8923	4.0326	-11.3803	0.0000

	Estimate	Std. Error	t value	Pr(> z)
aoa1.Aoa	-7.2188	2.4008	-3.0068	0.0026
freqTarget	-69.4579	3.5155	-19.7574	0.0000
relatednessrel:morphTypeeop	-0.3865	6.1742	-0.0626	0.9501
relatednessrel:morphTypetr	22.3288	3.0399	7.3452	0.0000
relatednessrel:aoa1.Aoa	2.8539	0.9182	3.1081	0.0019
morphTypeeop:aoa1.Aoa	0.3297	1.3148	0.2508	0.8020
morphTypetr:aoa1.Aoa	5.5318	1.2664	4.3682	0.0000
relatednessrel:morphTypeeop:aoa1.Aoa	-0.8039	1.3168	-0.6104	0.5416
relatednessrel:morphTypetr:aoa1.Aoa	-6.0672	1.0510	-5.7726	0.0000

Anova of aoaglmr1c - op contrast

```
aoaglmr1c.anova
```

```
## Analysis of Deviance Table (Type II Wald chisquare tests)
##
## Response: rt
##
##           Chisq Df Pr(>Chisq)
## relatedness    149.2574  1 < 2.2e-16 ***
## morphType      13.6248  2  0.001100 **
## aoa1.Aoa        6.2726  1  0.012262 *
## freqTarget     332.2366  1 < 2.2e-16 ***
## relatedness:morphType    11.7886  2  0.002755 **
## relatedness:aoa1.Aoa     1.4647  1  0.226176
## morphType:aoa1.Aoa      4.9247  2  0.085232 .
## relatedness:morphType:aoa1.Aoa 34.2866  2 3.587e-08 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

```
knitr::kable(round(aoaglmr1c.modelsum$coefficients, 4))
```

	Estimate	Std. Error	t value	Pr(> z)
(Intercept)	1058.3942	4.0485	261.4303	0.0000
relatednessrel	-38.0602	4.4646	-8.5249	0.0000
morphTypeeop	6.9564	3.9917	1.7427	0.0814
morphTypetr	-38.9403	12.0542	-3.2304	0.0012
aoa1.Aoa	-6.8888	2.4528	-2.8085	0.0050
freqTarget	-69.4589	3.8107	-18.2274	0.0000
relatednessrel:morphTypeeop	0.3829	3.8127	0.1004	0.9200
relatednessrel:morphTypetr	22.7143	3.9820	5.7042	0.0000
relatednessrel:aoa1.Aoa	2.0500	0.8996	2.2790	0.0227
morphTypeeop:aoa1.Aoa	-0.3303	1.2881	-0.2564	0.7976
morphTypetr:aoa1.Aoa	5.2019	1.3206	3.9392	0.0001
relatednessrel:morphTypeeop:aoa1.Aoa	0.8044	1.0975	0.7329	0.4636
relatednessrel:morphTypetr:aoa1.Aoa	-5.2632	1.0522	-5.0020	0.0000

```
aoaglmr2.anova <- car::Anova(aoaglmr2)
aoaglmr2c.anova <- car::Anova(aoaglmr2c)
```

```
aoaglm2.modelsum <- summary(aoaglm2)
aoaglm2c.modelsum <- summary(aoaglm2c)
```

Anova of aoaglm2 - or contrast

```
aoaglm2.anova
```

```
## Analysis of Deviance Table (Type II Wald chisquare tests)
##
## Response: rt
##
##               Chisq Df Pr(>Chisq)
## relatedness    221.0385  1 < 2.2e-16 ***
## morphType       0.8731  2  0.646273
## aoa2.usage       0.2344  1  0.628264
## freqTarget     594.1524  1 < 2.2e-16 ***
## relatedness:morphType  9.5127  2  0.008597 **
## relatedness:aoa2.usage 19.1904  1  1.183e-05 ***
## morphType:aoa2.usage 12.4446  2  0.001985 **
## relatedness:morphType:aoa2.usage 19.8405  2  4.917e-05 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

```
knitr::kable(round(aoaglm2.modelsum$coefficients, 4))
```

	Estimate	Std. Error	t value	Pr(> z)
(Intercept)	1008.8140	4.3403	232.4319	0.0000
relatednessrel	-51.5715	3.7709	-13.6762	0.0000
morphTypeop	1.8958	3.3523	0.5655	0.5717
morphTypetr	5.9059	4.6465	1.2710	0.2037
aoa2.usage	2.8479	3.3890	0.8404	0.4007
freqTarget	-69.2448	2.8408	-24.3752	0.0000
relatednessrel:morphTypeop	10.2784	3.4085	3.0155	0.0026
relatednessrel:morphTypetr	6.5114	3.9510	1.6480	0.0993
relatednessrel:aoa2.usage	10.3616	1.7518	5.9149	0.0000
morphTypeop:aoa2.usage	-2.1007	2.5761	-0.8155	0.4148
morphTypetr:aoa2.usage	-4.9514	2.1994	-2.2512	0.0244
relatednessrel:morphTypeop:aoa2.usage	-5.0028	2.0351	-2.4582	0.0140
relatednessrel:morphTypetr:aoa2.usage	-7.5540	1.8106	-4.1721	0.0000

Anova of aoaglm2c - op contrast

```
aoaglm2c.anova
```

```
## Analysis of Deviance Table (Type II Wald chisquare tests)
##
## Response: rt
##
##               Chisq Df Pr(>Chisq)
## relatedness    204.6584  1 < 2.2e-16 ***
## morphType       0.2466  2  0.88401
```

```
## aoa2.usage          0.0949  1    0.75803
## freqTarget         531.4092  1 < 2.2e-16 ***
## relatedness:morphType 6.2196  2    0.04461 *
## relatedness:aoa2.usage 20.5887  1 5.693e-06 ***
## morphType:aoa2.usage  11.1510  2    0.00379 **
## relatedness:morphType:aoa2.usage 10.6047  2    0.00498 **
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

```
knitr::kable(round(aoaglmr2c.modelsum$coefficients, 4))
```

	Estimate	Std. Error	t value	Pr(> z)
(Intercept)	1010.6973	3.2757	308.5464	0.0000
relatednessrel	-41.2928	3.4545	-11.9532	0.0000
morphTypeeor	-1.8942	3.0906	-0.6129	0.5400
morphTypetr	4.0100	4.1432	0.9678	0.3331
aoa2.usage	0.7503	2.9865	0.2512	0.8016
freqTarget	-69.2444	3.0038	-23.0523	0.0000
relatednessrel:morphTypeeor	-10.2732	4.0885	-2.5127	0.0120
relatednessrel:morphTypetr	-3.7669	3.3324	-1.1304	0.2583
relatednessrel:aoa2.usage	5.3587	1.4626	3.6637	0.0002
morphTypeeor:aoa2.usage	2.1010	2.3084	0.9102	0.3627
morphTypetr:aoa2.usage	-2.8509	2.1377	-1.3336	0.1823
relatednessrel:morphTypeeor:aoa2.usage	5.0014	2.0999	2.3818	0.0172
relatednessrel:morphTypetr:aoa2.usage	-2.5512	1.8387	-1.3875	0.1653

```
aoaglmr3.anova <- car::Anova(aoaglmr3)
aoaglmr3c.anova <- car::Anova(aoaglmr3c)
aoaglmr3.modelsum <- summary(aoaglmr3)
aoaglmr3c.modelsum <- summary(aoaglmr3c)
```

Anova of aoaglmr3 - or contrast

```
aoaglmr3.anova
```

```
## Analysis of Deviance Table (Type II Wald chisquare tests)
##
## Response: rt
##              Chisq Df Pr(>Chisq)
## relatedness    112.2561  1 < 2.2e-16 ***
## morphType       58.1050  2 2.414e-13 ***
## aoa3.context     5.6946  1 0.0170176 *
## freqTarget     554.3539  1 < 2.2e-16 ***
## relatedness:morphType 17.3889  2 0.0001675 ***
## relatedness:aoa3.context 1.6211  1 0.2029450
## morphType:aoa3.context 53.9894  2 1.889e-12 ***
## relatedness:morphType:aoa3.context 12.7531  2 0.0017010 **
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

```
knitr::kable(round(aoaglm3c.modelsum$coefficients, 4))
```

	Estimate	Std. Error	t value	Pr(> z)
(Intercept)	1010.3281	4.0408	250.0303	0.0000
relatednessrel	-15.7640	2.5784	-6.1139	0.0000
morphTypeop	-20.2492	4.6900	-4.3175	0.0000
morphTypetr	-29.6723	5.4359	-5.4586	0.0000
aoa3.contextschool	8.6139	4.0124	2.1468	0.0318
freqTarget	-69.3765	2.9466	-23.5447	0.0000
relatednessrel:morphTypeop	-7.8865	3.5802	-2.2028	0.0276
relatednessrel:morphTypetr	-9.4440	2.9994	-3.1487	0.0016
relatednessrel:aoa3.contextschool	-3.7625	2.9816	-1.2619	0.2070
morphTypeop:aoa3.contextschool	18.1619	3.4790	5.2204	0.0000
morphTypetr:aoa3.contextschool	23.2368	3.8477	6.0391	0.0000
relatednessrel:morphTypeop:aoa3.contextschool	2.6407	2.7957	0.9446	0.3449
relatednessrel:morphTypetr:aoa3.contextschool	-8.9994	2.6306	-3.4211	0.0006

Anova of aoaglm3c - op contrast

```
aoaglm3c.anova
```

```
## Analysis of Deviance Table (Type II Wald chisquare tests)
##
## Response: rt
##
##           Chisq Df Pr(>Chisq)
## relatedness    124.8011  1 < 2.2e-16 ***
## morphType      45.5467  2  1.287e-10 ***
## aoa3.context    48.2869  1  3.682e-12 ***
## freqTarget     481.4402  1 < 2.2e-16 ***
## relatedness:morphType    6.4028  2  0.0407056 *
## relatedness:aoa3.context    0.8070  1  0.3690075
## morphType:aoa3.context    16.9403  2  0.0002096 ***
## relatedness:morphType:aoa3.context    8.0212  2  0.0181227 *
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

```
knitr::kable(round(aoaglm3c.modelsum$coefficients, 4))
```

	Estimate	Std. Error	t value	Pr(> z)
(Intercept)	990.0474	11.1235	89.0047	0.0000
relatednessrel	-23.6486	3.1099	-7.6043	0.0000
morphTypeop	20.2524	3.7346	5.4229	0.0000
morphTypetr	-9.4203	4.7280	-1.9925	0.0463
aoa3.contextschool	26.8035	3.6248	7.3945	0.0000
freqTarget	-69.3751	3.1618	-21.9417	0.0000
relatednessrel:morphTypeop	7.8848	4.1938	1.8801	0.0601
relatednessrel:morphTypetr	-1.5600	3.5857	-0.4351	0.6635
relatednessrel:aoa3.contextschool	-1.1238	3.6388	-0.3088	0.7575
morphTypeop:aoa3.contextschool	-18.1640	4.5644	-3.9795	0.0001

	Estimate	Std. Error	t value	Pr(> z)
morphTypetr:aoa3.contextschool	5.0720	3.9001	1.3005	0.1934
relatednessrel:morphTypeop:aoa3.contextschool	-2.6390	5.9862	-0.4408	0.6593
relatednessrel:morphTypetr:aoa3.contextschool	-11.6374	4.1993	-2.7713	0.0056

```
aoaglm4.anova <- car::Anova(aoaglm4)
aoaglm4c.anova <- car::Anova(aoaglm4c)
aoaglm4.modelsum <- summary(aoaglm4)
aoaglm4c.modelsum <- summary(aoaglm4c)
```

Anova of aoaglm4 - or contrast

```
aoaglm4.anova
```

```
## Analysis of Deviance Table (Type II Wald chisquare tests)
##
## Response: rt
##
##               Chisq Df Pr(>Chisq)
## relatedness    138.033  1 < 2.2e-16 ***
## morphType       11.109  2  0.0038704 **
## aoa4.contextMultling 13.357  1  0.0002575 ***
## freqTarget     487.766  1 < 2.2e-16 ***
## relatedness:morphType 34.899  2  2.641e-08 ***
## relatedness:aoa4.contextMultling 16.465  1  4.955e-05 ***
## morphType:aoa4.contextMultling 37.648  2  6.680e-09 ***
## relatedness:morphType:aoa4.contextMultling 89.579  2 < 2.2e-16 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

```
knitr::kable(round(aoaglm4.modelsum$coefficients, 4))
```

	Estimate	Std. Error	t value	Pr(> z)
(Intercept)	1014.8046	5.7263	177.2179	0.0000
relatednessrel	-16.0639	2.6856	-5.9814	0.0000
morphTypeop	-0.1632	3.9786	-0.0410	0.9673
morphTypetr	-7.7779	4.5384	-1.7138	0.0866
aoa4.contextMultlingyes	14.7674	3.3393	4.4224	0.0000
freqTarget	-69.3824	3.1415	-22.0854	0.0000
relatednessrel:morphTypeop	-14.2395	3.0575	-4.6573	0.0000
relatednessrel:morphTypetr	-19.4036	3.5524	-5.4621	0.0000
relatednessrel:aoa4.contextMultlingyes	-12.5803	3.0768	-4.0887	0.0000
morphTypeop:aoa4.contextMultlingyes	-19.0786	3.5764	-5.3346	0.0000
morphTypetr:aoa4.contextMultlingyes	-8.9828	3.4055	-2.6378	0.0083
relatednessrel:morphTypeop:aoa4.contextMultlingyes	37.3186	4.0395	9.2383	0.0000
relatednessrel:morphTypetr:aoa4.contextMultlingyes	9.3316	4.3502	2.1451	0.0319

Anova of aoaglm4c - op contrast

```
aoaglm4c.anova
```

```
## Analysis of Deviance Table (Type II Wald chisquare tests)
##
## Response: rt
##
##               Chisq Df Pr(>Chisq)
## relatedness    118.4592  1 < 2.2e-16 ***
## morphType       8.0640  2  0.0177388 *
## aoa4.contextMultling 0.0178  1  0.8938698
## freqTarget    381.5908  1 < 2.2e-16 ***
## relatedness:morphType 21.4154  2  2.237e-05 ***
## relatedness:aoa4.contextMultling 12.8175  1  0.0003434 ***
## morphType:aoa4.contextMultling 67.6195  2  2.073e-15 ***
## relatedness:morphType:aoa4.contextMultling 112.2455  2 < 2.2e-16 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

```
knitr::kable(round(aoaglm4c.modelsum$coefficients, 4))
```

	Estimate	Std. Error	t value	Pr(> z)
(Intercept)	1014.6335	3.7310	271.9502	0.0000
relatednessrel	-30.3036	2.6323	-11.5120	0.0000
morphTypeeor	0.1644	3.5029	0.0469	0.9626
morphTypetr	-7.6135	3.6554	-2.0828	0.0373
aoa4.contextMultlingyes	-4.3090	4.1699	-1.0333	0.3014
freqTarget	-69.3810	3.5517	-19.5343	0.0000
relatednessrel:morphTypeeor	14.2400	2.9286	4.8624	0.0000
relatednessrel:morphTypetr	-5.1641	4.0898	-1.2627	0.2067
relatednessrel:aoa4.contextMultlingyes	24.7394	3.6402	6.7962	0.0000
morphTypeeor:aoa4.contextMultlingyes	19.0799	3.5817	5.3270	0.0000
morphTypetr:aoa4.contextMultlingyes	10.0963	3.3269	3.0347	0.0024
relatednessrel:morphTypeeor:aoa4.contextMultlingyes	-37.3209	4.4065	-8.4695	0.0000
relatednessrel:morphTypetr:aoa4.contextMultlingyes	-27.9879	4.0259	-6.9520	0.0000

```
aoaglm5.anova <- car::Anova(aoaglm5)
aoaglm5c.anova <- car::Anova(aoaglm5c)
aoaglm5.modelsum <- summary(aoaglm5)
aoaglm5c.modelsum <- summary(aoaglm5c)
```

Anova of aoaglm5 - or contrast

```
aoaglm5.anova
```

```
## Analysis of Deviance Table (Type II Wald chisquare tests)
##
## Response: rt
##
##               Chisq Df Pr(>Chisq)
## relatedness    208.3417  1 < 2.2e-16 ***
## morphType       7.5328  2  0.023135 *
## aoa5.selfRatedProf 7.0812  1  0.007790 **
```



```
## freqTarget                371.8232  1 < 2.2e-16 ***
## lengthTarget              78.5755  1 < 2.2e-16 ***
## relatedness:morphType    126.4083  2 < 2.2e-16 ***
## relatedness:aoa5.selfRatedProf  30.3694  1 3.571e-08 ***
## morphType:aoa5.selfRatedProf   10.1555  2  0.006234 **
## relatedness:morphType:aoa5.selfRatedProf  68.0810  2 1.646e-15 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

```
knitr::kable(round(aoaglmr5.modelsum$coefficients, 4))
```

	Estimate	Std. Error	t value	Pr(> z)
(Intercept)	1190.5452	6.3788	186.6397	0.0000
relatednessrel	-70.9797	5.0746	-13.9872	0.0000
morphTypeop	15.0837	3.9889	3.7814	0.0002
morphTypetr	8.5210	3.9509	2.1568	0.0310
aoa5.selfRatedProf	-12.1673	4.3049	-2.8264	0.0047
freqTarget	-72.1470	3.7415	-19.2827	0.0000
lengthTarget	-26.5652	2.9969	-8.8643	0.0000
relatednessrel:morphTypeop	41.7713	3.9409	10.5995	0.0000
relatednessrel:morphTypetr	31.2858	4.0719	7.6834	0.0000
relatednessrel:aoa5.selfRatedProf	14.7631	1.7101	8.6330	0.0000
morphTypeop:aoa5.selfRatedProf	-4.1976	2.7160	-1.5455	0.1222
morphTypetr:aoa5.selfRatedProf	-2.5407	2.3469	-1.0826	0.2790
relatednessrel:morphTypeop:aoa5.selfRatedProf	-13.4543	2.0818	-6.4629	0.0000
relatednessrel:morphTypetr:aoa5.selfRatedProf	-13.7726	1.9811	-6.9521	0.0000

Anova of aoaglmr5c - op contrast

```
aoaglmr5c.anova
```

```
## Analysis of Deviance Table (Type II Wald chisquare tests)
##
## Response: rt
##
##           Chisq Df Pr(>Chisq)
## relatedness    178.378  1 < 2.2e-16 ***
## morphType      15.949  2 0.0003442 ***
## aoa5.selfRatedProf  18.979  1 1.321e-05 ***
## freqTarget     551.733  1 < 2.2e-16 ***
## relatedness:morphType  107.112  2 < 2.2e-16 ***
## relatedness:aoa5.selfRatedProf  16.315  1 5.363e-05 ***
## morphType:aoa5.selfRatedProf   12.829  2 0.0016376 **
## relatedness:morphType:aoa5.selfRatedProf  57.396  2 3.441e-13 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

```
knitr::kable(round(aoaglmr5c.modelsum$coefficients, 4))
```

	Estimate	Std. Error	t value	Pr(> z)
(Intercept)	1069.7679	4.3337	246.8492	0.0000
relatednessrel	-28.9715	2.9596	-9.7891	0.0000
morphTypeeor	-10.2669	4.1874	-2.4519	0.0142
morphTypetr	-10.9334	4.1105	-2.6599	0.0078
aoa5.selfRatedProf	-16.3513	3.6982	-4.4214	0.0000
freqTarget	-69.2447	2.9480	-23.4890	0.0000
relatednessrel:morphTypeeor	-41.7143	3.5639	-11.7047	0.0000
relatednessrel:morphTypetr	-10.5767	3.8383	-2.7556	0.0059
relatednessrel:aoa5.selfRatedProf	1.2412	1.3897	0.8932	0.3718
morphTypeeor:aoa5.selfRatedProf	4.2182	2.6033	1.6203	0.1052
morphTypetr:aoa5.selfRatedProf	1.6447	2.3242	0.7076	0.4792
relatednessrel:morphTypeeor:aoa5.selfRatedProf	13.4287	1.8466	7.2723	0.0000
relatednessrel:morphTypetr:aoa5.selfRatedProf	-0.3072	1.8570	-0.1654	0.8686

```
aoaglm6.anova <- car::Anova(aoaglm6)
aoaglm6c.anova <- car::Anova(aoaglm6c)
aoaglm6.modelsum <- summary(aoaglm6)
aoaglm6c.modelsum <- summary(aoaglm6c)
```

Anova of aoaglm6 - or contrast

```
aoaglm6.anova
```

```
## Analysis of Deviance Table (Type II Wald chisquare tests)
##
## Response: rt
##
##           Chisq Df Pr(>Chisq)
## relatedness    94.3669  1 < 2.2e-16 ***
## morphType      6.1065  2  0.04721 *
## aoa6.otherLang  24.5541  1  7.225e-07 ***
## freqTarget    492.3427  1 < 2.2e-16 ***
## relatedness:morphType  54.5036  2  1.461e-12 ***
## relatedness:aoa6.otherLang  3.1366  1  0.07656 .
## morphType:aoa6.otherLang  1.0621  2  0.58797
## relatedness:morphType:aoa6.otherLang  25.4509  2  2.974e-06 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

```
knitr::kable(round(aoaglm6.modelsum$coefficients, 4))
```

	Estimate	Std. Error	t value	Pr(> z)
(Intercept)	1005.2142	4.7982	209.4997	0.0000
relatednessrel	-8.9288	2.9942	-2.9820	0.0029
morphTypeop	1.5979	3.8290	0.4173	0.6765
morphTypetr	-12.2422	3.8726	-3.1612	0.0016
aoa6.otherLangyes	16.7862	6.1024	2.7507	0.0059
freqTarget	-69.3289	3.1245	-22.1888	0.0000
relatednessrel:morphTypeop	-19.3675	4.0133	-4.8258	0.0000
relatednessrel:morphTypetr	-25.3511	3.3966	-7.4638	0.0000

	Estimate	Std. Error	t value	Pr(> z)
relatednessrel:aoa6.otherLangyes	-13.3232	3.5199	-3.7851	0.0002
morphTypeop:aoa6.otherLangyes	-8.4471	4.2259	-1.9989	0.0456
morphTypetr:aoa6.otherLangyes	2.8512	3.5963	0.7928	0.4279
relatednessrel:morphTypeop:aoa6.otherLangyes	18.1754	4.4657	4.0700	0.0000
relatednessrel:morphTypetr:aoa6.otherLangyes	10.8099	3.4435	3.1392	0.0017

Anova of aoaglmr5c - op contrast

aoaglmr6c.anova

```
## Analysis of Deviance Table (Type II Wald chisquare tests)
##
## Response: rt
##               Chisq Df Pr(>Chisq)
## relatedness    157.6142  1 < 2.2e-16 ***
## morphType       24.3774  2  5.088e-06 ***
## aoa6.otherLang    2.3460  1  0.1256046
## freqTarget     470.5704  1 < 2.2e-16 ***
## relatedness:morphType    17.0403  2  0.0001994 ***
## relatedness:aoa6.otherLang    1.0043  1  0.3162694
## morphType:aoa6.otherLang    16.6182  2  0.0002463 ***
## relatedness:morphType:aoa6.otherLang    28.3287  2  7.055e-07 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

```
knitr::kable(round(aoaglmr6c.modelsum$coefficients, 4))
```

	Estimate	Std. Error	t value	Pr(> z)
(Intercept)	1006.8022	3.7388	269.2883	0.0000
relatednessrel	-28.2969	3.0307	-9.3367	0.0000
morphTypeop	-1.6004	3.6196	-0.4421	0.6584
morphTypetr	-13.8407	3.2983	-4.1963	0.0000
aoa6.otherLangyes	8.3424	3.4133	2.4440	0.0145
freqTarget	-69.3273	3.1959	-21.6926	0.0000
relatednessrel:morphTypeop	19.3686	3.7056	5.2268	0.0000
relatednessrel:morphTypetr	-5.9826	3.5372	-1.6913	0.0908
relatednessrel:aoa6.otherLangyes	4.8529	3.3157	1.4636	0.1433
morphTypeop:aoa6.otherLangyes	8.4488	3.1001	2.7253	0.0064
morphTypetr:aoa6.otherLangyes	11.2997	3.2771	3.4481	0.0006
relatednessrel:morphTypeop:aoa6.otherLangyes	-18.1763	3.6070	-5.0392	0.0000
relatednessrel:morphTypetr:aoa6.otherLangyes	-7.3672	3.0128	-2.4453	0.0145

OSC

First, let's try to pit OSC against priming condition – these two are typically confounded:

```
temp <- unique(masterFile[masterFile$lexicality=='word' & masterFile$language=='eng',c('target','prime')
aggregate(oscTarget ~ morphType, FUN=fivenum, data=temp)
```

```
##   morphType oscTarget.1 oscTarget.2 oscTarget.3 oscTarget.4 oscTarget.5
## 1      op      -0.0940      0.2300      0.4580      0.8130      0.9750
## 2      or      -0.0020      0.1220      0.2625      0.6950      0.9990
## 3      tr       0.3640      0.6040      0.8260      0.9180      1.0000
```

Indeed they are. This tests it via NHST

```
summary(aov(oscTarget~morphType, data=subset(temp, relatedness=='rel')));
```

```
##              Df Sum Sq Mean Sq F value    Pr(>F)
## morphType      2   3.433   1.7163    21.02 9.75e-09 ***
## Residuals    144  11.755   0.0816
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
## 3 observations deleted due to missingness
```

modelling

```
osc1 <- glmer(rt ~ relatedness * oscTarget * phonemicFluency + freqTarget + lengthTarget + (1|subject)
```

```
osc1.anova <- car::Anova(osc1)
```

```
osc2 <- glmer(rt ~ relatedness * oscTarget * vocabulary + freqTarget + lengthTarget + (1|subject) + (1|subject:oscTarget)
```

```
osc2.anova<-car::Anova(osc2)
```

```
osc3 <- glmer(rt ~ relatedness * oscTarget * phonemicComprehension + freqTarget + lengthTarget + (1|subject)
```

```
osc3.anova<-car::Anova(osc3)
```

```
osc4 <- glmer(rt ~ relatedness * oscTarget * morphComprehension + freqTarget + lengthTarget + (1|subject)
```

```
osc4.anova<-car::Anova(osc4)
```

Plots

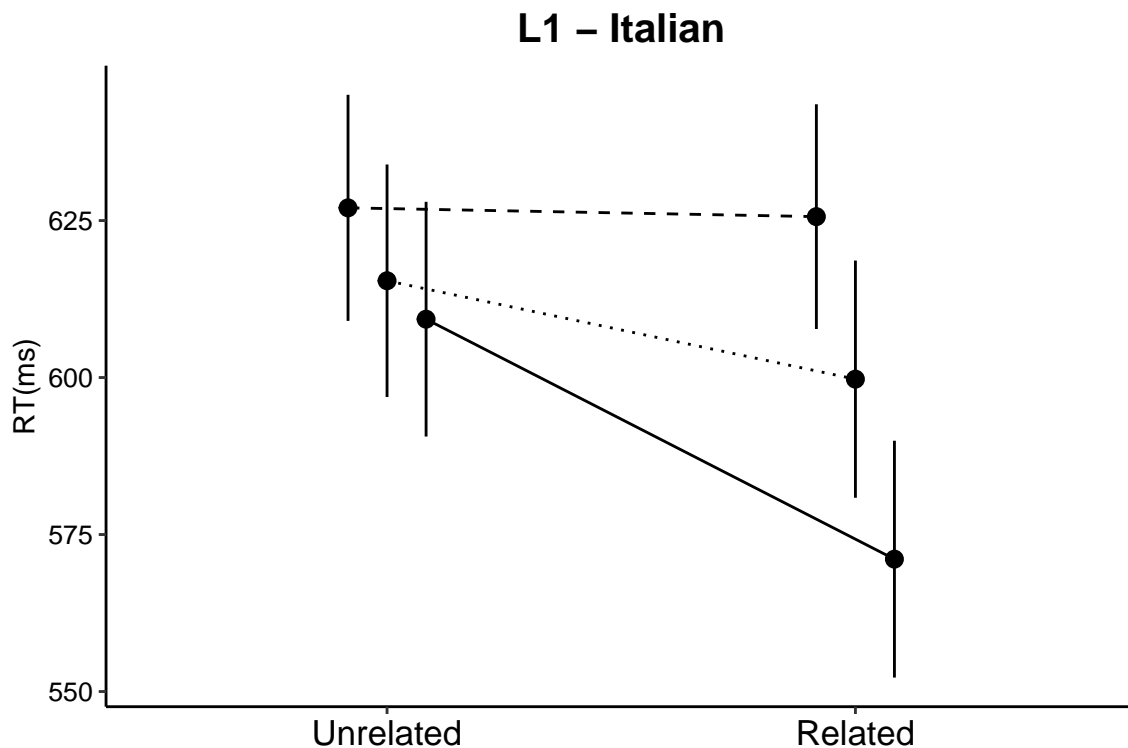
Figure 1 - plot of estimated RTs

```
df <- effect("relatedness:morphType", itaglm2);
df <- as.data.frame(df);
revalue(df$relatedness, c("ctrl"="Unrelated"))-> df$relatedness;
revalue(df$relatedness, c("rel"="Related"))-> df$relatedness;
```

```

dodge1 <- position_dodge(width = 0.25);
bb <- ggplot(data = df, aes(x = relatedness, y = fit, group = morphType)) +
  geom_point(size = 2, position = dodge1) +
  geom_line(aes(linetype=morphType), position = dodge1) +
  scale_linetype_manual(values=c("dashed", "dotted", "solid")) +
  theme_classic();
bb <- bb + geom_pointrange(aes(ymin = df$lower, ymax = df$upper), position = dodge1) ;
bb <- bb + scale_y_continuous("RT(ms)") ;
bb <- bb + theme(axis.title.y = element_text(size = rel(1), angle = 90));
bb <- bb + theme(axis.text.y = element_text(angle = 00, hjust = 1, size=10, colour = 'black'));
bb <- bb + theme(axis.title.x = element_blank()) + theme(axis.text.x = element_text(size=13, colour = 'black'));
bb <- bb + labs(title='L1 - Italian');
bb <- bb + theme(plot.title= element_text(angle = 00, hjust=0.5, size=15, face = 'bold', colour = 'black'));
bb <- bb + theme(legend.position="none")
bb

```



```

df <- effect("relatedness:morphType", enggler2);
df <- as.data.frame(df);
revalue(df$relatedness, c("ctrl"="Unrelated"))-> df$relatedness;
revalue(df$relatedness, c("rel"="Related"))-> df$relatedness;

dodge <- position_dodge(width = 0.25);
gg <- ggplot(data = df, aes(x = relatedness, y = fit, group = morphType)) +
  geom_point(size = 2, position = dodge) +
  geom_line(aes(linetype=morphType), position = dodge) +
  scale_linetype_manual(values=c("dotted", "dashed", "solid")) +
  theme_classic();

```

```

gg <- gg + geom_pointrange(aes(ymin = df$lower, ymax = df$upper), position = dodge);
gg <- gg + scale_y_continuous("RT (ms)" );
gg <- gg + theme(axis.text.y = element_text(angle = 00, hjust = 1, size=10, colour = 'black'));
gg <- gg + theme(axis.title.x = element_blank()) + theme(axis.text.x = element_text(size=13, colour = 'black'));
gg <- gg + labs(title='L2 - English');
gg <- gg + theme(plot.title= element_text(angle = 00, hjust=0.5, size=15, face = 'bold', colour = 'black'));
gg<- gg + theme(legend.position="none")
gg

```

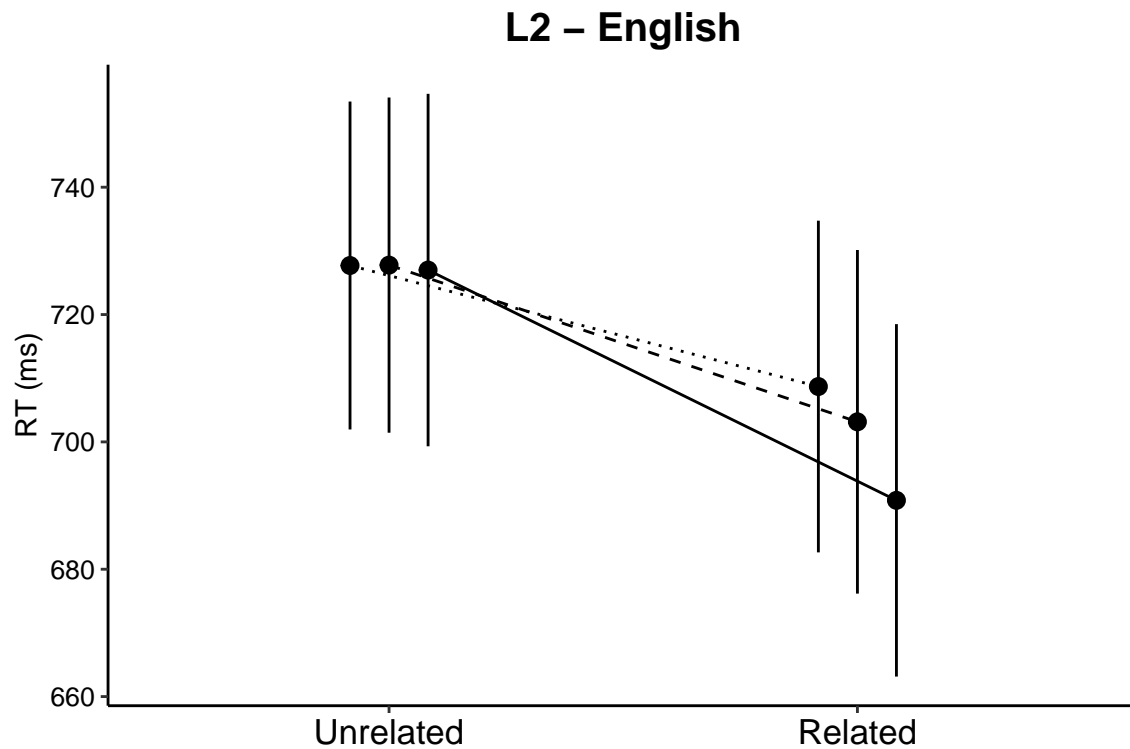


Figure 2 - Participants' score distributions for each proficiency subtest

```

par(mfrow=c(2,4));
par(mar=c(5,5,4,.5)+.1);
par(lwd=2);

attach(pptFeatures);

hist(phonemicFluency, breaks = seq(0,50,5), main = '(a) Phon Fluency', cex.main=2, xlab = 'Scores', ylab = 'Frequency');
axis(1, cex.axis=2);
axis(2, at=c(0,50), cex.axis=2, las=1);

hist(phonemicComprehension, breaks = seq(0,13,1), main = '(b) Phon Comprehension', cex.main=2, xlab = 'Scores', ylab = 'Frequency');
axis(1, cex.axis=2);
axis(2, at=c(0,50), cex.axis=2, las=1);

hist(morphComprehension, breaks = seq(0,10,1), main = '(c) Morph Awareness', cex.main=2, xlab = 'Scores', ylab = 'Frequency');
axis(1, cex.axis=2);
axis(2, at=c(0,50), cex.axis=2, las=1);

```

```

axis(1, cex.axis=2);
axis(2, at=c(0,50), cex.axis=2, las=1);

hist(spelling, breaks = seq(0,20,2), main = '(d) Spelling', cex.main=2, xlab = 'Scores', ylab = 'N of p
axis(1, cex.axis=2);
axis(2, at=c(0,50), cex.axis=2, las=1);

hist(readingComprehension, breaks = seq(0,7,1), main = '(e) Read Comprehension', cex.main=2, xlab = 'Sc
axis(1, cex.axis=2);
axis(2, at=c(0,50), cex.axis=2, las=1);

hist(vocabulary, breaks = seq(0,20,2), main = '(f) Vocabulary', cex.main=2, xlab = 'Scores', ylab = 'N
axis(1, cex.axis=2);
axis(2, at=c(0,50), cex.axis=2, las=1);

hist(oralComprehension, breaks = seq(0,6,1), main = '(g) Oral comprehension', cex.main=2, xlab = 'Scores
axis(1, cex.axis=2);
axis(2, at=c(0,50), cex.axis=2, las=1);
#dev.off()
detach(pptFeatures);

```

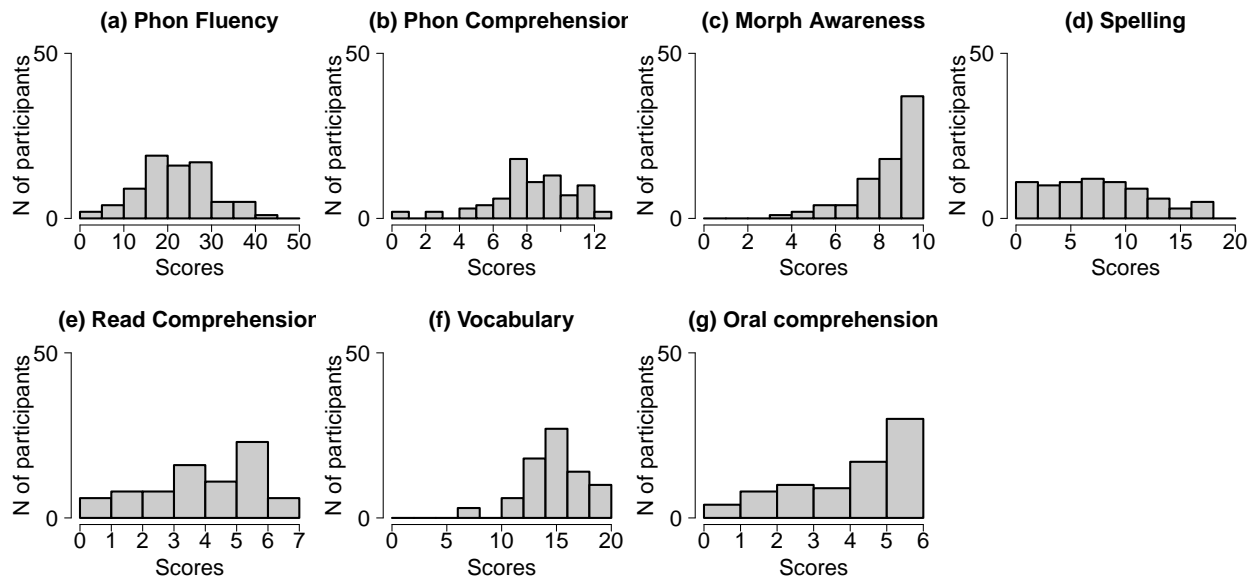


Figure 3 - Interaction by phonemic comprehension by relatedness and morphtype

```

temp <- data.frame(effect('relatedness:morphType:phonemicComprehension', proficiencyglmer2, se=list(lev
revalue(temp$relatedness, c("rel"="Related"))-> temp$relatedness;
revalue(temp$relatedness, c("ctrl"="Unrelated"))-> temp$relatedness;

phonComprehension_names <- c(
  "5" = "Low phonComprehension",
  "9" = "Medium phonComprehension",

```

```

"12" = "High phonComprehension");

ggplot(data = temp, aes(x=relatedness, y=fit, group=morphType)) +
  geom_point(size = 2, position = position_dodge(width = 0.25)) +
  geom_line(aes(linetype=morphType), position = position_dodge(width = 0.25)) +
  scale_linetype_manual(values=c("dashed", "dotted", "solid")) +
  theme_bw() +
  theme(panel.grid.major = element_blank()) +
  ylab('RTs (ms)') + xlab('') +
  theme(axis.text.y = element_text(angle = 00, hjust = 1, size=8, colour = 'black'))+
  theme(axis.text.x = element_text(size=13, colour = 'black'))+
  geom_pointrange(aes(ymin = lower, ymax = upper), position = position_dodge(width = 0.25)) +
  facet_grid(~ phonemicComprehension,
             labeller = labeller(phonemicComprehension = as_labeller(phonComprehension_names))) +
  theme(strip.text = element_text(size=12)) +
  theme(legend.position="none");

```

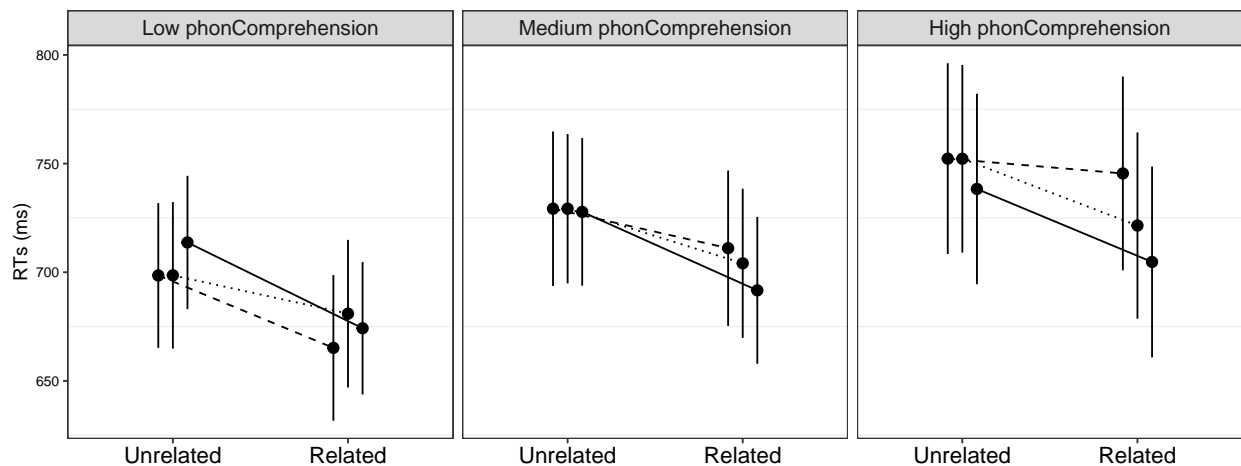


Figure 4 - Interaction by morphological awareness/comprehension by relatedness and morphtype

```

temp <- data.frame(effect('relatedness:morphType:morphComprehension', proficiencyglmer3, se=list(level=
revalue(temp$relatedness, c("rel"="Related"))-> temp$relatedness;
revalue(temp$relatedness, c("ctrl"="Unrelated"))-> temp$relatedness;

morphComprehension_names <- c(
  "6" = "Low morphComprehension",
  "9" = "Medium morphComprehension",
  "10" = "High morphComprehension");

ggplot(data = temp, aes(x=relatedness, y=fit, group=morphType)) +
  geom_point(size = 2, position = position_dodge(width = 0.25)) +
  geom_line(aes(linetype=morphType), position = position_dodge(width = 0.25)) +
  scale_linetype_manual(values=c("dashed", "dotted", "solid")) +

```



```

theme_bw() +
theme(panel.grid.major = element_blank()) +
ylab('RTs (ms)') + xlab('') +
theme(axis.text.y = element_text(angle = 00, hjust = 1, size=8, colour = 'black'))+
theme(axis.text.x = element_text(size=13, colour = 'black'))+
geom_pointrange(aes(ymin = lower, ymax = upper), position = position_dodge(width = 0.25)) +
facet_grid(~ morphComprehension,
            labeller = labeller(morphComprehension = as_labeller(morphComprehension_names))) +
theme(strip.text = element_text(size=12)) +
theme(legend.position="none");

```

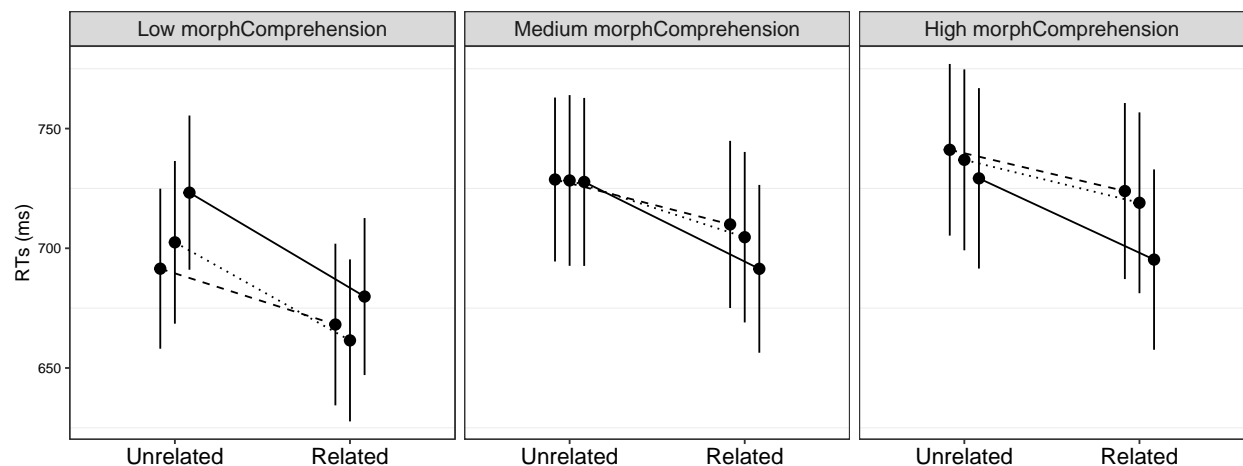


Figure 5 - Interaction by vocabulary by relatedness and morphotype

```

temp <- data.frame(effect('relatedness:morphType:vocabulary', proficiencyglmer6, se=list(level=.95), xlab=''))
revalue(temp$relatedness, c("rel"="Related"))-> temp$relatedness;
revalue(temp$relatedness, c("ctrl"="Unrelated"))-> temp$relatedness;

vocabulary_names <- c(
  "11" = "Low vocabulary",
  "16" = "Medium vocabulary",
  "19" = "High vocabulary");

ggplot(data = temp, aes(x=relatedness, y=fit, group=morphType)) +
  geom_point(size = 2, position = position_dodge(width = 0.25)) +
  geom_line(aes(linetype=morphType), position = position_dodge(width = 0.25)) +
  scale_linetype_manual(values=c("dashed", "dotted", "solid")) +
  theme_bw() +
  theme(panel.grid.major = element_blank()) +
  ylab('RTs (ms)') + xlab('') +
  theme(axis.text.y = element_text(angle = 00, hjust = 1, size=8, colour = 'black'))+
  theme(axis.text.x = element_text(size=13, colour = 'black'))+
  geom_pointrange(aes(ymin = lower, ymax = upper), position = position_dodge(width = 0.25)) +
  facet_grid(~ vocabulary,

```

```
labeller = labeller(vocabulary = as_labeller(vocabulary_names))) +
theme(strip.text = element_text(size=12)) +
theme(legend.position="none");
```

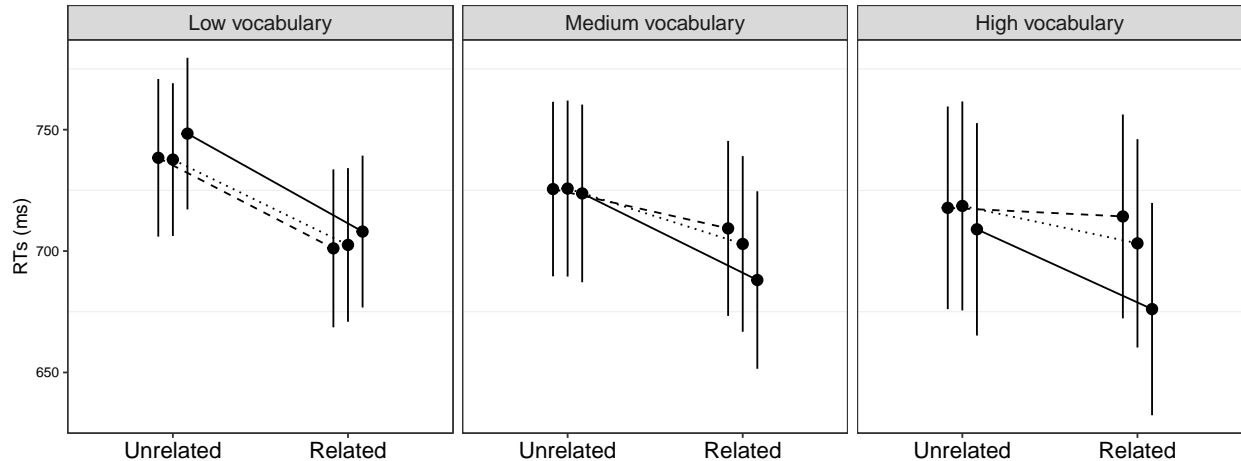


Figure 6 - Interaction by phonemic Fluency by relatedness and morphotype

```
temp <- data.frame(effect('relatedness:morphType:phonemicFluency', proficiencyglmer1, se=list(level=.95),
revalue(temp$relatedness, c("rel"="Related"))-> temp$relatedness;
revalue(temp$relatedness, c("ctrl"="Unrelated"))-> temp$relatedness;

phonemicFluency_names <- c(
  "10" = "Low phonFluency",
  "23" = "Medium phonFluency",
  "39" = "High phonFluency");

ggplot(data = temp, aes(x=relatedness, y=fit, group=morphType)) +
  geom_point(size = 2, position = position_dodge(width = 0.25)) +
  geom_line(aes(linetype=morphType), position = position_dodge(width = 0.25)) +
  scale_linetype_manual(values=c("dashed", "dotted", "solid")) +
  theme_bw() +
  theme(panel.grid.major = element_blank()) +
  ylab('RTs (ms)') + xlab('') +
  theme(axis.text.y = element_text(angle = 00, hjust = 1, size=8, colour = 'black'))+
  theme(axis.text.x = element_text(size=13, colour = 'black'))+
  geom_pointrange(aes(ymin = lower, ymax = upper), position = position_dodge(width = 0.25)) +
  facet_grid(~ phonemicFluency,
    labeller = labeller(phonemicFluency = as_labeller(phonemicFluency_names))) +
  theme(strip.text = element_text(size=12)) +
  theme(legend.position="none");
```

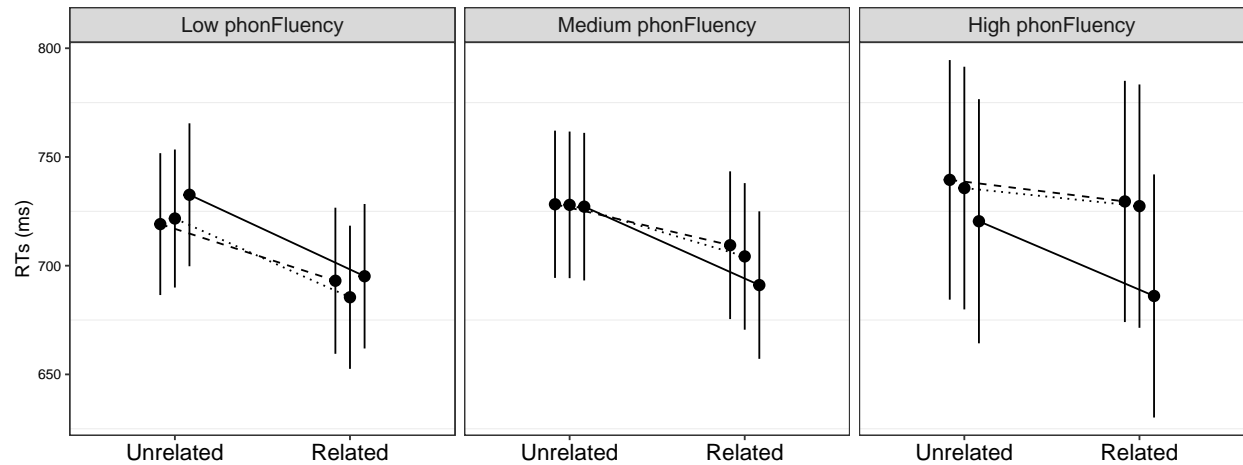


Figure 7 - Scores distributions in the AoA questionnaire

```
#jpeg(filename = paste(localGitDir, '/aoaScores.jpg', sep = ''), res=300, height=2200, width=4400);
par(mfrow=c(2,3));
par(mar=c(5,5,4,.5)+.1);
par(lwd=2);

attach(pptFeatures);

hist(aoa1.Aoa, breaks = seq(-.5,15.5,1), main = '(a) Age first exposed', cex.main=2, xlab = 'Scores', ylab = 'N of participants');
axis(1, cex.axis=2);
axis(2, at=c(0,30), cex.axis=2, las=1);

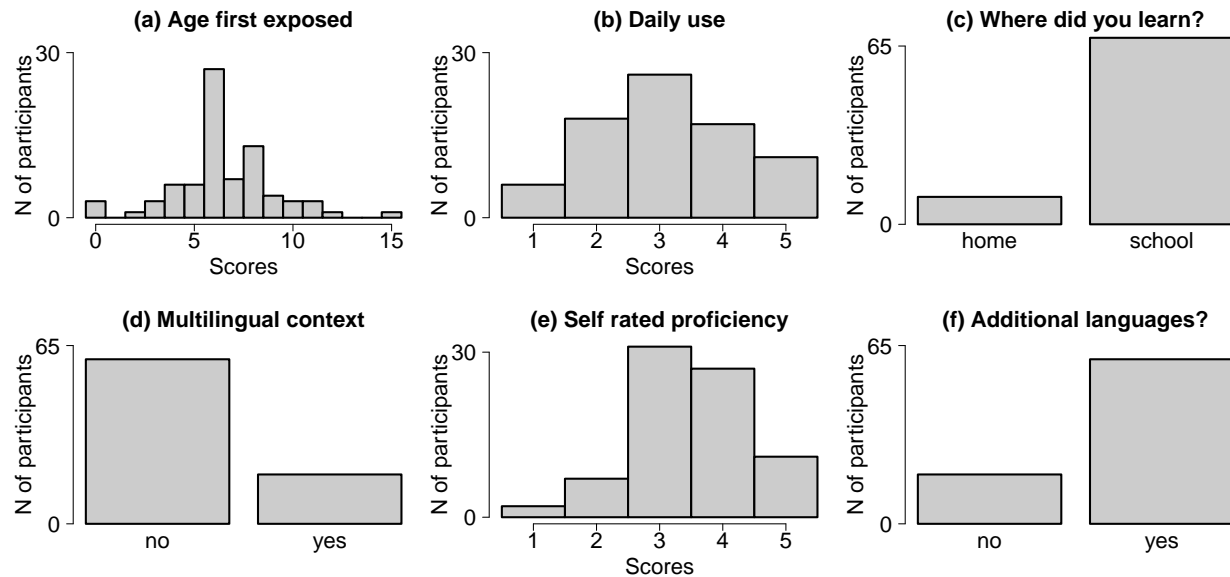
hist(aoa2.usage, breaks = seq(.5,5.5,1), main = '(b) Daily use', cex.main=2, xlab = 'Scores', ylab = 'N of participants');
axis(1, cex.axis=2);
axis(2, at=c(0,30), cex.axis=2, las=1);

barplot(table(aoa3.context), main = '(c) Where did you learn?', cex.main=2, ylab = 'N of participants');
axis(2, at=c(0,65), cex.axis=2, las=1);

barplot(table(aoa4.contextMultling), main = '(d) Multilingual context', cex.main=2, ylab = 'N of participants');
axis(2, at=c(0,65), cex.axis=2, las=1);

hist(aoa5.selfRatedProf, breaks = seq(.5,5.5,1), main = '(e) Self rated proficiency', cex.main=2, xlab = 'Scores', ylab = 'N of participants');
axis(1, cex.axis=2);
axis(2, at=c(0,30), cex.axis=2, las=1);

barplot(table(aoa6.otherLang), main = '(f) Additional languages?', cex.main=2, ylab = 'N of participants');
axis(2, at=c(0,65), cex.axis=2, las=1);
```



```
detach(pptFeatures);
```

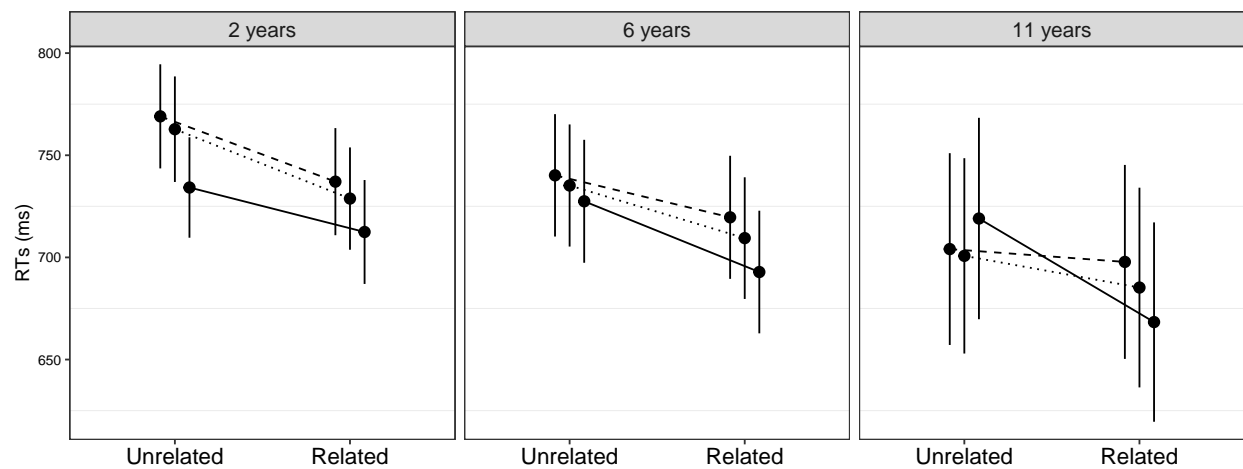
```
par(mfrow=c(1,1));  
#dev.off();
```

Figure 8 - Interaction by AoA1 by relatedness and morphtype

```
temp <- data.frame(effect('relatedness:morphType:aoa1.Aoa', aoaglm1, se=list(level=.95), xlevels=list(
revalue(temp$relatedness, c("rel"="Related"))-> temp$relatedness;
revalue(temp$relatedness, c("ctrl"="Unrelated"))-> temp$relatedness;

aoa1.Aoa_names <- c(
  "2" = "2 years",
  "6" = "6 years",
  "11" = "11 years");

ggplot(data = temp, aes(x=relatedness, y=fit, group=morphType)) +
  geom_point(size = 2, position = position_dodge(width = 0.25)) +
  geom_line(aes(linetype=morphType), position = position_dodge(width = 0.25)) +
  scale_linetype_manual(values=c("dashed", "dotted", "solid")) +
  theme_bw() +
  theme(panel.grid.major = element_blank() +
  ylab('RTs (ms)') + xlab('') +
  theme(axis.text.y = element_text(angle = 00, hjust = 1, size=8, colour = 'black'))+
  theme(axis.text.x = element_text(size=13, colour = 'black'))+
  geom_pointrange(aes(ymin = lower, ymax = upper), position = position_dodge(width = 0.25)) +
  facet_grid(~ aoa1.Aoa,
    labeller = labeller(aoa1.Aoa = as_labeller(aoa1.Aoa_names))) +
  theme(strip.text = element_text(size=12)) +
  theme(legend.position="none");
```



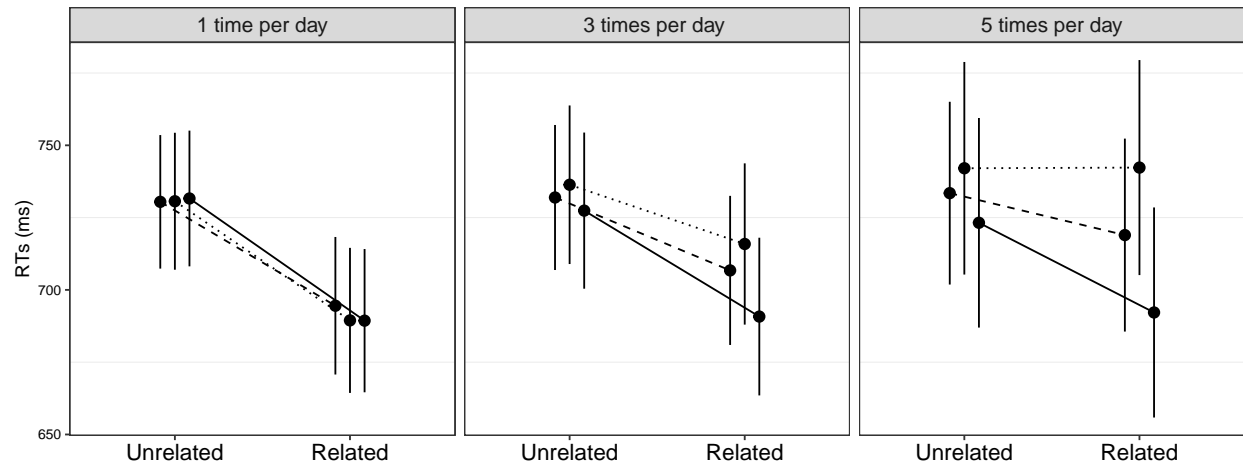
```
#ggsave("aoa1.AoA.jpg", width = 7, height = 3, dpi = 300);
```

Figure 9 - Interaction by AoA2 by relatedness and morphotype

```
temp <- data.frame(effect('relatedness:morphType:aoa2.usage', aoa1mer2c, se=list(level=.95), xlevels=1,
revalue(temp$relatedness, c("rel"="Related"))-> temp$relatedness;
revalue(temp$relatedness, c("ctrl"="Unrelated"))-> temp$relatedness;

aoa2.usage_names <- c(
  "1" = "1 time per day",
  "3" = "3 times per day ",
  "5" = "5 times per day");

ggplot(data = temp, aes(x=relatedness, y=fit, group=morphType)) +
  geom_point(size = 2, position = position_dodge(width = 0.25)) +
  geom_line(aes(linetype=morphType), position = position_dodge(width = 0.25)) +
  scale_linetype_manual(values=c("dashed", "dotted", "solid")) +
  theme_bw() +
  theme(panel.grid.major = element_blank()) +
  ylab('RTs (ms)') + xlab('') +
  theme(axis.text.y = element_text(angle = 00, hjust = 1, size=8, colour = 'black'))+
  theme(axis.text.x = element_text(size=13, colour = 'black'))+
  geom_pointrange(aes(ymin = lower, ymax = upper), position = position_dodge(width = 0.25)) +
  facet_grid(~ aoa2.usage,
             labeller = labeller(aoa2.usage = as_labeller(aoa2.usage_names))) +
  theme(strip.text = element_text(size=12)) +
  theme(legend.position="none");
```



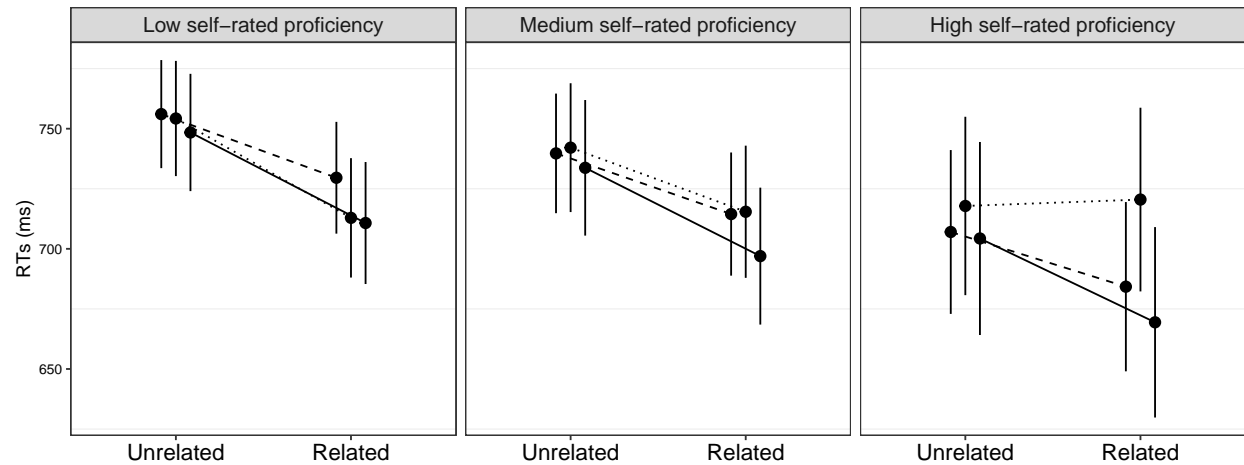
```
#ggsave("aoa2.usage.jpg", width = 7, height = 3, dpi = 300);
```

Figure 10 - Interaction by AoA5 by relatedness and morphtype

```
temp <- data.frame(effect('relatedness:morphType:aoa5.selfRatedProf', aoa1mer5c, se=list(level=.95), x=
revalue(temp$relatedness, c("rel"="Related"))-> temp$relatedness;
revalue(temp$relatedness, c("ctrl"="Unrelated"))-> temp$relatedness;

aoa5.selfRatedProf_names <- c(
  "2" = "Low self-rated proficiency",
  "3" = "Medium self-rated proficiency",
  "5" = "High self-rated proficiency");

ggplot(data = temp, aes(x=relatedness, y=fit, group=morphType)) +
  geom_point(size = 2, position = position_dodge(width = 0.25)) +
  geom_line(aes(linetype=morphType), position = position_dodge(width = 0.25)) +
  scale_linetype_manual(values=c("dashed", "dotted", "solid")) +
  theme_bw() +
  theme(panel.grid.major = element_blank()) +
  ylab('RTs (ms)') + xlab('') +
  theme(axis.text.y = element_text(angle = 00, hjust = 1, size=8, colour = 'black'))+
  theme(axis.text.x = element_text(size=13, colour = 'black'))+
  geom_pointrange(aes(ymin = lower, ymax = upper), position = position_dodge(width = 0.25)) +
  facet_grid(~ aoa5.selfRatedProf,
    labeller = labeller(aoa5.selfRatedProf = as_labeller(aoa5.selfRatedProf_names))) +
  theme(strip.text = element_text(size=12)) +
  theme(legend.position="none");
```



```
#ggsave("aoa5.selfratedProf.jpg", width = 7.5, height = 3, dpi = 300);
```

Figure 11 - OSC by morphotype

```
temp <- unique(masterFile[masterFile$lexicality=='word' & masterFile$language=='eng',c('target','prime'
revalue(temp$morphType, c("or"="Orthographic", 'op'='Opaque', 'tr'='Transparent'))-> temp$morphType;
library(ggpubr);
ggboxplot(subset(temp, oscTarget>0), "morphType", "oscTarget",
          color = "black", fill = grey(.80),
          width = 0.5, ylab = 'OSC', xlab = '');
```

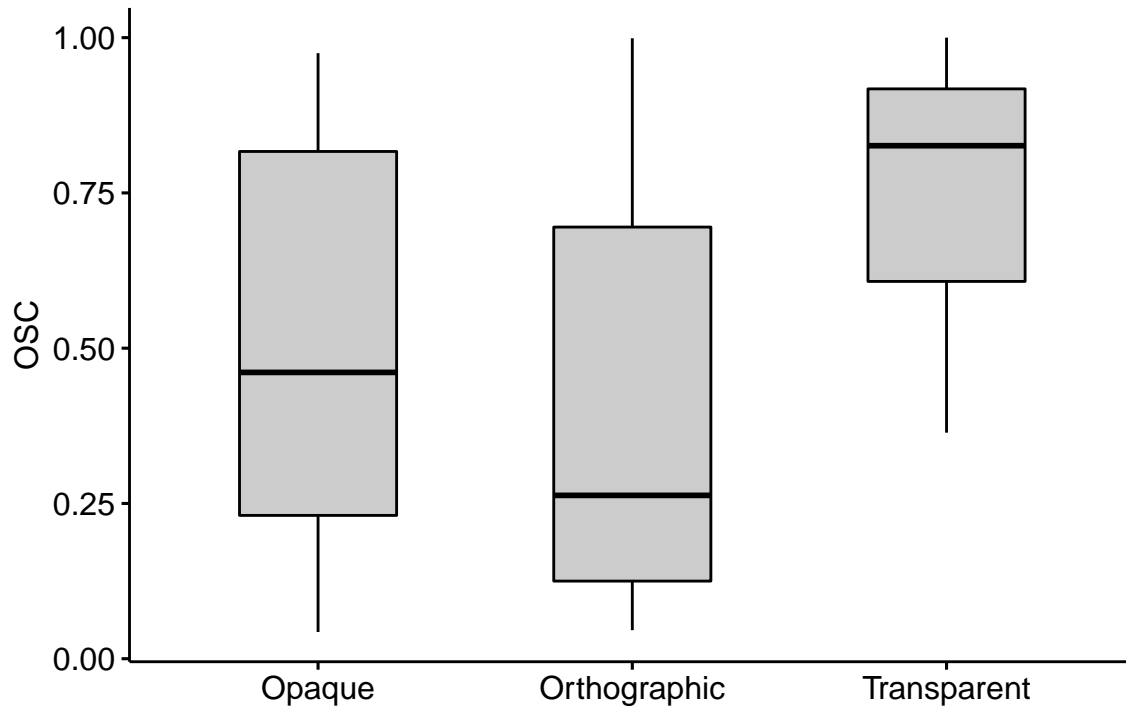
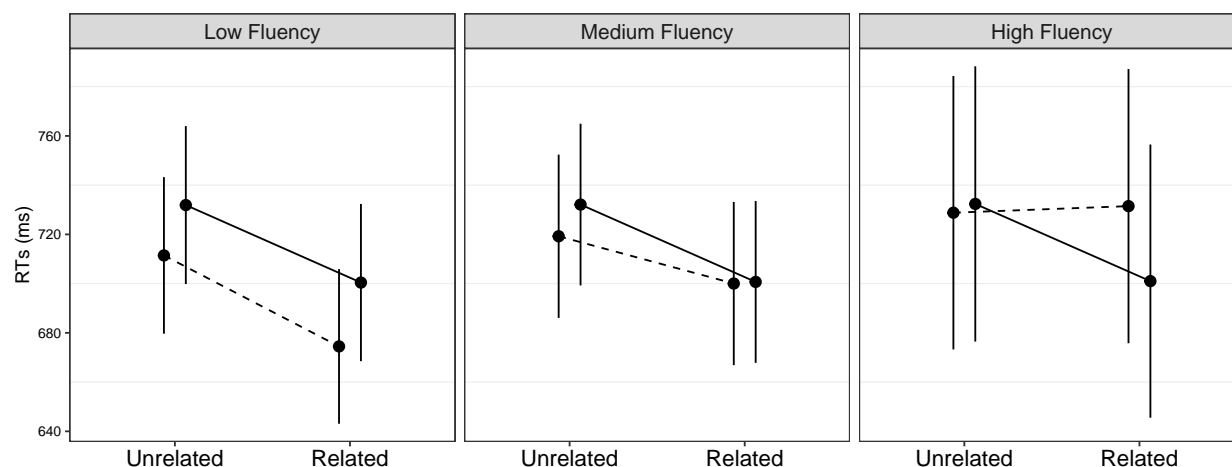


Figure 12 - OSC by phonemic fluency by relatedness

```
temp <- data.frame(effect('relatedness:oscTarget:phonemicFluency', osc1, se=list(level=.95), xlevels=li
revalue(temp$relatedness, c("rel"="Related"))-> temp$relatedness;
revalue(temp$relatedness, c("ctrl"="Unrelated"))-> temp$relatedness;

phonemicFluency_names <- c(
  "10" = "Low Fluency",
  "23" = "Medium Fluency",
  "39" = "High Fluency"
);

temp$oscTarget <- as.factor(temp$oscTarget);
ggplot(data = temp, aes(x=relatedness, y=fit, group=oscTarget)) +
  geom_point(position = position_dodge(width = 0.25)) +
  geom_line(aes(linetype = oscTarget), position = position_dodge(width = 0.25)) +
  scale_linetype_manual(values=c("dashed", "solid")) +
  theme_bw() +
  theme(panel.grid.major = element_blank()) +
  ylab('RTs (ms)') + xlab('') +
  theme(axis.text.y = element_text(angle = 00, hjust = 1, size=8, colour = 'black'))+
  theme(axis.text.x = element_text(size=13, colour = 'black'))+
  geom_pointrange(aes(ymin = lower, ymax = upper), position = position_dodge(width = 0.25)) +
  facet_grid(~ phonemicFluency,
             labeller = labeller(phonemicFluency = as_labeller(phonemicFluency_names))) +
  theme(strip.text = element_text(size=12))+
  theme(legend.position="none");
```

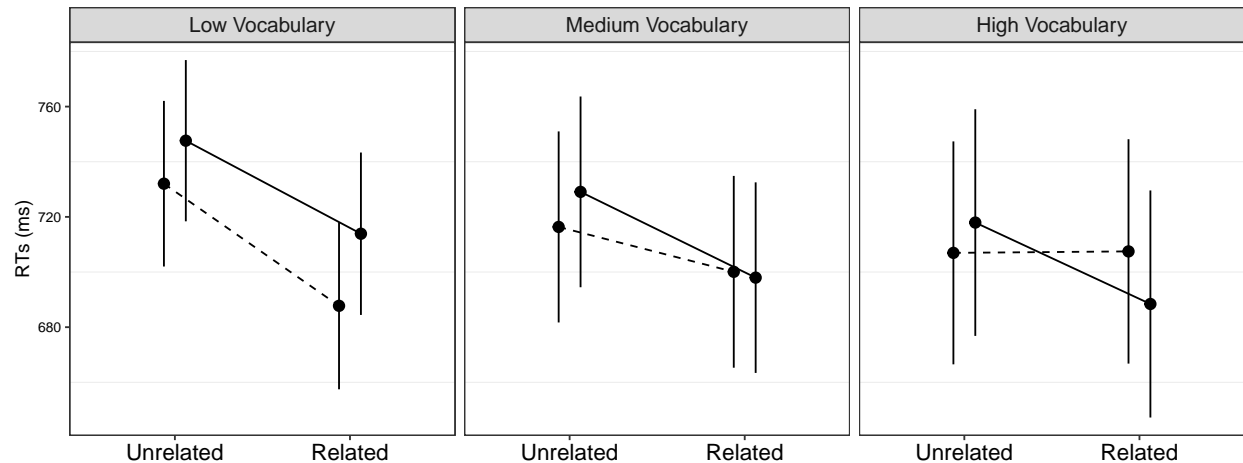
```
ggsave("oscModel.jpg", width = 7, height = 3, dpi = 300);
```

Figure 13 - OSC by vocabulary by relatedness

```
temp <- data.frame(effect('relatedness:oscTarget:vocabulary', osc2, se=list(level=.95), xlevels=list(osc2,
revalue(temp$relatedness, c("rel"="Related"))-> temp$relatedness;
revalue(temp$relatedness, c("ctrl"="Unrelated"))-> temp$relatedness;

vocabulary_names <- c(
  "11" = "Low Vocabulary",
  "16" = "Medium Vocabulary",
  "19" = "High Vocabulary"
);

temp$oscTarget <- as.factor(temp$oscTarget);
ggplot(data = temp, aes(x=relatedness, y=fit, group=oscTarget)) +
  geom_point(position = position_dodge(width = 0.25)) +
  geom_line(aes(linetype = oscTarget), position = position_dodge(width = 0.25)) +
  scale_linetype_manual(values=c("dashed", "solid")) +
  theme_bw() +
  theme(panel.grid.major = element_blank()) +
  ylab('RTs (ms)') + xlab('') +
  theme(axis.text.y = element_text(angle = 00, hjust = 1, size=8, colour = 'black'))+
  theme(axis.text.x = element_text(size=13, colour = 'black'))+
  geom_pointrange(aes(ymin = lower, ymax = upper), position = position_dodge(width = 0.25)) +
  facet_grid(~ vocabulary,
    labeller = labeller(vocabulary = as_labeller(vocabulary_names))) +
  theme(strip.text = element_text(size=12))+
  theme(legend.position="none");
```



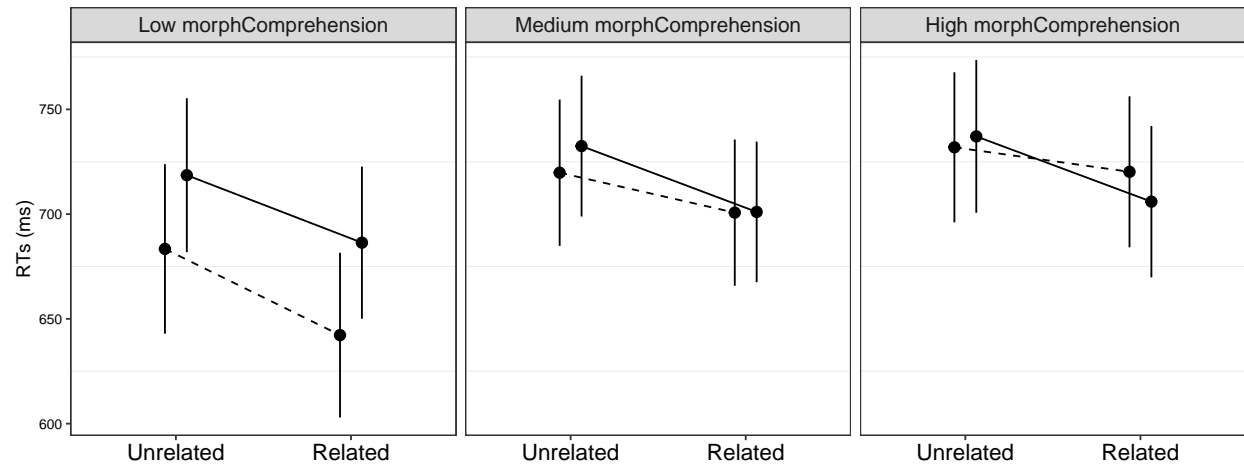
```
ggsave("vocabulary.jpg", width = 7.5, height = 3, dpi = 300);
```

Figure 14 - morphComprehension by relatedness

```
temp <- data.frame(effect('relatedness:oscTarget:morphComprehension', osc4, se=list(level=.95), xlevels=
revalue(temp$relatedness, c("rel"="Related"))-> temp$relatedness;
revalue(temp$relatedness, c("ctrl"="Unrelated"))-> temp$relatedness;

morphComprehension_names <- c(
  "6" = "Low morphComprehension",
  "9" = "Medium morphComprehension",
  "10" = "High morphComprehension"
);

temp$oscTarget <- as.factor(temp$oscTarget);
ggplot(data = temp, aes(x=relatedness, y=fit, group=oscTarget)) +
  geom_point(position = position_dodge(width = 0.25)) +
  geom_line(aes(linetype = oscTarget), position = position_dodge(width = 0.25)) +
  scale_linetype_manual(values=c("dashed", "solid")) +
  theme_bw() +
  theme(panel.grid.major = element_blank()) +
  ylab('RTs (ms)') + xlab('') +
  theme(axis.text.y = element_text(angle = 00, hjust = 1, size=8, colour = 'black'))+
  theme(axis.text.x = element_text(size=13, colour = 'black'))+
  geom_pointrange(aes(ymin = lower, ymax = upper), position = position_dodge(width = 0.25)) +
  facet_grid(~ morphComprehension,
    labeller = labeller(morphComprehension = as_labeller(morphComprehension_names))) +
  theme(strip.text = element_text(size=12))+
  theme(legend.position="none");
```



```
ggsave("figure14.jpg", width = 7.7, height = 3, dpi = 300);
```